GLUFOSINATE-AMMONIUM (175)

EXPLANATION

Glufosinate-ammonium is a herbicide and desiccant. It was first evaluated for residues and toxicology by the 1991 JMPR and subsequently for residues in 1994.

Glufosinate-tolerant crops have now been developed with new use patterns necessitating revised MRLs. The manufacturer has provided many new reports for evaluation dealing with animal and plant metabolism, environmental fate in soil and water, methods of residue analysis, stability in stored analytical samples, supervised residue trials, animal transfer studies, fate of residues in processing and national residue limits.

The delegation of The Netherlands was requested by the 1996 CCPR to send their comments on the definition of the residue to the JMPR (ALINORM 97/24 para 73).

Information was provided to the Meeting by Germany, The Netherlands, Poland and the basic manufacturer.

METABOLISM AND ENVIRONMENTAL FATE

Information was made available on the metabolism and environmental fate of glufosinate in animals, crops, soils and water-sediment systems.

A variety of code numbers and abbreviations have been used for the metabolites and degradation products of glufosinate-ammonium. There are nine different code numbers for glufosinate itself, covering the salts and free acid, and the racemate and separate stereoisomers. Earlier publications used HOE numbers but they have recently been replaced by AEF, AEC or AE numbers. The names, structures and code numbers of glufosinate and its metabolites and degradation products are listed below.

In this evaluation "glufosinate" and the abbreviations NAG, MPP, etc. refer to the compounds without specifying their stereoisomerism or whether they are presents as salts or free acid. Where such extra information is important, e.g. because of molecular weight considerations, extra information will be given, e.g. "glufosinate, expressed as the free acid".

Structures, names and code numbers of glufosinate, metabolites and degradation products

Glufosinate

	racemate	L-isomer	R-isomer	MW	O U
free acid	AE F035956	AE F057740	AE F090532	181.1	 D ===================================
Ammonium	AE F039866	AE F058192	AE F093854	198.2	CH ₃ COOH
salt					он /
HCl salt	AE F035125	AE F057742	AE F057741	217.6	
					glufosinate INn2

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3-methylphosphinicopropionic acid (MPP)

		MW
Free acid	AE F061517	152.1
Disodium salt	AE C527855	196.1

2-methylphosphinicoacetic acid (MPA)

		MW
Free acid	AE F064619	138.1
Disodium salt	AE F159481	182.1



2-methyl-phosphinicoacetic acid

4-methylphosphinico-2-hydroxybutanoic acid (MHB)

		MW
Free acid	AE F053705	182.1
Disodium salt	AE F042231	226.1

MW

166.1

4-methylphosphinicobutanoic acid (MPB)

AE F039046

Free acid



4-methyl-phosphinico-2hydroxy-butanoic acid



4-methyl-phosphinico-butanoic acid

free acid

Disodiu

m salt







AE 0015081 3-methyl-phosphinico acrylic acid



methylphosphinico-formic acid

Methylphosphinico-formic acid



AE 0015081

Animal metabolism

Free acid

Information on the metabolism of glufosinate-ammonium and NAG (N-acetyl-L-glufosinate) in laboratory rats, lactating goats and laying hens was reported. In summary, most of the administered dose of both compounds is rapidly excreted. NAG may be partially metabolized back to glufosinate.

Bremmer and Leist (1997) examined the possible conversion of NAG to glufosinate in rats. Up to 10% deacetylation occurred at a low dose of 3 mg/kg bw as shown by the occurrence of glufosinate in the faeces. The authors concluded however that most of the conversion was caused by bacteria in the colon and rectum although toxicity findings indicate partial bioavailability (Bremmer and Leist, 1998).

Kellner et al. (1993) showed that almost all the radiolabel was excreted in the faeces within 4 days when rats were dosed orally with single doses of ¹⁴C]NAG disodium salt at 3 mg/kg body weight. The ¹⁴C label was in positions 3 and 4.

Unchanged NAG was the main source of the radiolabel (85-89%) in faecal extracts from rats dosed orally with [14C]NAG disodium salt at 1000 mg/kg body weight (Lauck-Birkel, 1995a). Small amounts (approx. 1%) of glufosinate were produced.



When $[3,4-^{14}C]$ glufosinate-ammonium was administered in a single oral dose of 500 mg/kg bw to rats, 75% in males and 89% in females of the radiolabel was excreted in the faeces within 48 hours (Lauck-Birkel, 1995b) and 8-11% in the urine. Glufosinate was the principal labelled component in faecal extracts – 72% and 84% of the dose in males and females respectively.



Labelled glufosinate-ammonium

Lauck-Birkel (1996) identified the labelled compounds in urine and faeces from rats dosed orally with $[3,4^{-14}C]$ glufosinate-ammonium at 2 mg/kg bw. The main compound in faecal extracts was glufosinate (77% of the dose); other identified components were NAG (7.5%), MHB (4.3%) and MPP (1.3%). The main components in the urine were glufosinate (4.3% of the dose) and MPP (0.8%).

Tissue, milk and excreta residues were measured in a lactating goat weighing 60 kg dosed orally twice daily for 4 consecutive days by capsule with $[3,4-^{14}C]$ glufosinate at a rate equivalent to 101 ppm glufosinate-ammonium in the dry-weight diet and 3.0 mg/kg bw/day (Huang and Smith, 1995a). The feed intake was 1.8 kg/day. The animal was milked twice daily and slaughtered 15 hours after the final dose.

Most of the administered ¹⁴C (69%) was excreted in the faeces with 2.9% in the urine and 11% in the GI tract with contents. Less than 0.1% and 0.02% of the dose was found in the tissues and milk respectively. The levels in the kidneys were higher than in other tissues (Table 1). Levels of ¹⁴C reached a plateau in milk by day 2.

The parent compound was the main residue detected in the kidneys, liver and milk, with MPP forming a substantial part of the residue in the kidneys and liver (Table 2).

Table 1. Distribution of ¹⁴ C in the	tissues and	milk of a	goat dosed	twice da	aily for 4	days with	[3,4-
¹⁴ C]glufosinate (Huang and Smith,	1995a).						

Sample	¹⁴ C as glufosinate-ammonium, mg/kg
Kidney	0.61
Liver	0.40
Muscle	0.007
Fat	0.004
Milk day 1, am and pm	0.003 0.009
Milk day 2, am and pm	0.016 0.020
Milk day 3, am and pm	0.022 0.020
Milk day 4, am and pm	0.020 0.014

Compound	Kidney		Liver		Milk	
	% of total ¹⁴ C	mg/kg	% of total ¹⁴ C	mg/kg	% of total ¹⁴ C	mg/kg
Glufosinate	49	0.30	53	0.21	49	0.010
NAG, L-isomer	4.2	0.026	not detecte	d	2.2	< 0.001
MPP	29	0.18	37	0.15	6.3	0.001
MPA	1.2	0.008	0.4	0.001	5.3	0.001

Table 2. Compounds identified in the tissues and milk of a goat dosed twice daily for 4 days with $[3,4-{}^{14}C]$ glufosinate (Huang and Smith, 1995a). Residue levels are expressed as glufosinate-ammonium equivalents.

Tissue, milk and excreta residues were measured in a lactating goat weighing 36 kg dosed orally twice daily for 3 consecutive days by capsule with $[3,4-^{14}C]N$ -acetyl-L-glufosinate disodium salt at a rate equivalent to 84 ppm in the dry-weight diet or 3.0 mg/kg bw/day (Huang and Smith, 1995b). The feed intake was 1.4 kg/day. The animal was milked twice daily and slaughtered 16 hours after the final dose.

Most of the administered ¹⁴C was excreted in the faeces (68%) with 7.3% in the urine and 19% in the GI tract with contents. NAG and glufosinate accounted for 52% and 34% of the ¹⁴C in the faeces respectively.

Only 0.2% of the administered dose was found in the tissues and blood, with <0.1% in the milk. Levels in the kidneys were higher than in other tissues (Table 3). Levels of ¹⁴C reached a plateau in milk by day 2. Glufosinate was the main residue detected in the kidneys, liver and milk, with NAG (the administered material) and MPP forming a substantial part of the residue in the kidneys and liver (Table 4).

Table 3. Distribution of ¹⁴C in the tissues and milk of a goat dosed twice daily for 3 consecutive days with $[3,4-^{14}C]N$ -acetyl-L-glufosinate disodium salt (Huang and Smith, 1995b).

Tissue or milk	¹⁴ C as N-acetyl-L-glufosinate disodium salt, mg/kg
Kidneys	0.93
Liver	0.29
Muscle	0.007
Fat	<0.010
Milk day 1, am and pm	0.005 0.012
Milk day 2, am and pm	0.018 0.020
Milk day 3, am and pm	0.023 0.022

Table 4. Compounds identified in the tissues and milk of a goat dosed twice daily for 3 consecutive days with $[3,4-^{14}C]N$ -acetyl-L-glufosinate disodium salt (Huang and Smith, 1995b). Residue levels are expressed as *N*-acetyl-L-glufosinate disodium salt equivalents.

Compound	Kidneys		Liver		Milk	
	% of total ¹⁴ C	mg/kg	% of total ¹⁴ C	mg/kg	% of total ¹⁴ C	mg/kg
Glufosinate	40	0.37	33	0.095	40	0.009
NAG, L-isomer	32	0.30	19	0.054	9.2	0.002
MPP	20	0.19	21	0.060	14	0.003
MPA	1.6	0.015	2.0	0.006	4.8	0.001

Tissue, eggs and excreta residues were measured in 6 laying hens weighing 1.27-1.67 kg dosed orally twice daily for 14 consecutive days by capsule with $[3,4-{}^{14}C]$ glufosinate-ammonium at a

rate equivalent to 25 ppm glufosinate-ammonium in the diet (it was not clear whether the feeding level was expressed on a fresh-weight or dry-weight basis) or 2.0 mg/kg bw/day (Huang and Smith, 1995c). The feed intake was 120 g/bird/day. Eggs were collected twice daily and the birds were slaughtered 16 hours after the final dose .

92% of the administered dose was excreted with 1.3% remaining in the GI tract. Glufosinate-ammonium accounted for 81% of the 14 C in the faeces.

Less than 0.02% of the administered dose was present in the edible tissues. Levels of ¹⁴C as glufosinate-ammonium in the liver, muscle and fat were 0.11, <0.004 and 0.003 mg/kg respectively. Those in the eggs are shown in Table 5. ¹⁴C in egg whites reached a plateau by day 6, but in the last 3 days there was again a small increase. The levels in egg yolks increased very slowly throughout the 14 days.

MPP was the main residue identified in the liver with glufosinate-ammonium also a substantial component (Table 6). Glufosinate constituted most of the residue in eggs.

Table 5. ¹⁴C in eggs from hens dosed orally twice daily for 14 consecutive days by capsule with [3,4-¹⁴C]glufosinate- (Huang and Smith, 1995c).

Collection day	Mean ¹⁴ C, mg/kg as glufosinate-ammonium				
	Egg white	Egg yolk			
1	<0.003	< 0.003			
2	0.004	< 0.003			
3	0.034	0.005			
4	0.053	0.009			
5	0.049	0.012			
6	0.057	0.015			
7	0.056	0.016			
8	0.056	0.017			
9	0.058	0.019			
10	0.059	0.021			
11	0.058	0.021			
12	0.067	0.021			
13	0.065	0.022			
14	0.067	0.024			

Table 6. Compounds identified in the tissues and eggs from hens dosed orally twice daily for 14 consecutive days with [3,4-¹⁴C]glufosinate-ammonium (Huang and Smith, 1995c). Residues are expressed as glufosinate-ammonium equivalents.

Compound	Liver		Egg white (day 14)			Egg yolk (day 13)	
	% ¹⁴ C in liver	mg/kg	% ¹⁴ C ii	n egg white	mg/kg	% ¹⁴ C in egg yolk	mg/kg
glufosinate	31	0.036	78		0.052	53	0.012
NAG, L-isomer	4.9	0.006	not detected		2.4	0.001	
MPP	44	0.050	1.3		0.001	4.1	0.001
MPA	3.5	0.004	not detected		3.1	0.001	

Tissue, eggs and excreta residues were measured in 6 laying hens weighing 1.27-1.60 kg dosed orally twice daily for 14 consecutive days by capsule with $[3,4-^{14}C]N$ -acetyl-L-glufosinate disodium salt equivalent to 27 ppm *N*-acetyl-L-glufosinate disodium salt in the diet (it was not clear whether the feeding level was expressed on a fresh- or dry-weight basis) or 2.2 mg/kg bw/day (Huang and Smith, 1995d). The mean feed intake was 116 g/bird/day. Eggs were collected twice daily and the birds were slaughtered 15 hours after the final dose.

86% of the administered dose was excreted with 1.0% remaining in the GI tract. *N*-acetyl-L-glufosinate disodium salt accounted for 73% of the ¹⁴C in the faeces with glufosinate and MPP accounting for 13% and 8.6% respectively.

Less than 0.1% of the administered dose was present in the edible tissues and blood. Levels of ¹⁴C (as *N*-acetyl-L-glufosinate disodium salt) in the liver, muscle and fat were 0.076, 0.013 and 0.011 mg/kg respectively. Those in eggs are shown in Table 7. Levels in egg whites were slightly above the LOD throughout the study; those in egg yolks increased slowly but steadily throughout the 14 days.

NAG (the administered material) was the main residue identified in liver and egg yolk (Table 8). Glufosinate and MPP were also substantial components of the liver residue. Glufosinate was the main identified residue in egg whites, but the levels in eggs were quite low, making further identification difficult.

Table 7. ¹⁴C in eggs from hens dosed orally twice daily for 14 consecutive days with [3,4-¹⁴C]*N*-acetyl-L-glufosinate disodium salt (Huang and Smith, 1995d).

Collection day	Mean ¹⁴ C level, mg/kg as N-acetyl-L-glufosinate disodium salt				
	Egg white	Egg yolk			
1	<0.009	<0.002			
2	<0.009	<0.002			
3	<0.009	0.012			
4	0.010	0.020			
5	0.010	0.027			
6	0.011	0.035			
7	<0.009	0.037			
8	0.012	0.040			
9	0.014	0.042			
10	0.012	0.045			
11	0.012	0.046			
12	0.013	0.050			
13	0.015	0.049			
day 14 sac	0.014	0.052			
necropsy	<0.009	0.056			

Table 8. Compounds identified in the tissues and eggs from hens dosed orally twice daily for 14 consecutive days with $[3,4-^{14}C]N$ -acetyl-L-glufosinate disodium salt (Huang and Smith, 1995d). Residues are expressed as *N*-acetyl-L-glufosinate disodium salt equivalents.

Compound	Liver		Egg white (day 13)		Egg yolk (necropsy)	
	% ¹⁴ C	mg/kg	% ¹⁴ C	mg/kg	% ¹⁴ C	mg/kg
Glufosinate	15	0.011	14	0.002	2.8	0.002
NAG, L-isomer	27	0.020	5.1	0.001	13	0.007
MPP	17	0.013	2.0	< 0.001	2.2	0.001
MPA	not detected		1.1	< 0.001	0.6	< 0.001



Figure 1. Proposed metabolic pathways of glufosinate and *N*-acetyl glufosinate in ruminants and poultry.

Plant metabolism

Information was reported on the metabolism of glufosinate-ammonium in genetically modified rape (canola), sugar beet, maize, soya and tomatoes. The studies examined the disposition of the residue throughout the plant and its composition. In some cases the metabolism of glufosinate-ammonium in genetically modified and unmodified crops was compared.

Stumpf *et al.* (1995b) placed cut rape plants, genetically modified and unmodified, in a nutrient solution containing 4.7 mg/l of $[3,4-^{14}C]$ glufosinate-ammonium for 6 days. In the genetically modified plants the ¹⁴C represented NAG (57%) and glufosinate-ammonium (36%). In the unmodified plants the ¹⁴C was mainly due to unchanged glufosinate-ammonium (80%) with 16% MPP. The experiment demonstrated the rapid acetylation of glufosinate in genetically modified rape.



Labelled glufosinate-ammonium

Tshabalala (1993) treated canola plants (*var* 19-2XACS-N3) at the 3-5 leaf stage once with [3,4-¹⁴C]glufosinate-ammonium at a rate equivalent to 0.75 kg ai/ha and collected samples for radioanalysis on days 1, 21 and 120. The results are shown in Table 9.

The levels in the top-growth and roots were much the same after 21 days, but at maturity were much higher in the roots than in other plant parts. The residues in the canola seed were investigated by two HPLC systems but the low residue levels made further identification difficult. Glufosinate and MPP were the main constituents with considerably lower levels of NAG.

Sample	Time since treatment	¹⁴ C as glufosinate-ammonium, mg/kg
whole plant	1 hour	145
topgrowth	21 days	5.3 3.2
roots	21 days	3.8 5.2
topgrowth	120 days	0.058 0.064 0.024 0.021
roots	120 days	0.22 0.13 0.15 0.19
hulls	120 days	0.12 0.26 0.11 0.076
seed	120 days	0.054 0.11 0.056 0.045

Table 9. Levels of ¹⁴C in canola plants treated once with [3,4-¹⁴C]glufosinate-ammonium at 0.75 kg ai/ha (Tshabalala 1993).

Thalacker (1994) applied [¹⁴C]glufosinate-ammonium to glufosinate-tolerant canola and samples were taken after 1 hour and 21 days. After 1-hour 73% of the ¹⁴C was in glufosinate and 18% in NAG, and after 21-days 60, 21 and 7% of the ¹⁴C corresponded to NAG, glufosinate-ammonium and MPP respectively. Again, genetically modified rape seed (canola) produced NAG very rapidly from glufosinate-ammonium.

In another trial, genetically modified sugar beets were treated twice (22 days interval) with $[3,4-{}^{14}C]$ glufosinate-ammonium at rates equivalent to 0.6 kg ai/ha (Allan, 1996). The leaves and beets (when formed) were harvested 0, 8 and 15 days after the first treatment and 0, 21 and 146 days after the second. The leaves were rinsed with water to separate surface residues from absorbed residues

The residues after the first treatment are shown in Table 10. The glufosinate isomer composition was unchanged in the surface residue, but in the absorbed residue L-glufosinate was metabolized to NAG (*N*-acetyl-L-glufosinate).

The composition of the residue after the second treatment is shown in Table 11. Identified residues accounted for 93-98% of the total ¹⁴C in leaves + rinses and roots. After the first treatment NAG was the main identified residue except in the leaves on the day of treatment, but even after 146 days glufosinate accounted for 19% of the ¹⁴C in the roots and 26% in the leaves. The reason is presumably that L-glufosinate is rapidly converted to NAG, but D-glufosinate remains unchanged.

Table 10. Identified residues in genetically modified sugar beets after a single treatment with [3,4-¹⁴C]glufosinate-ammonium.

Time	after	Sample	Re	Residue as % of total ¹⁴ C in the sample					
treatment			Glufosinate	D-glufosinate	L-glufosinate	NAG			
3 hours		rinse	41	20	21				
8 days		rinse	18	9	9				
15 days		rinse	14	7	7				
3 hours		leaves	45.1	24.6	20.5	9.0			
8 days		leaves	35.6	28.4	5.5	39			
15 days		leaves	29.3	25.2	3.3	49			

Time after	Sample	Glufos	sinate	MI	PP	NA	G
2^{nd}		% of ¹⁴ C in	mg/kg as	% of ¹⁴ C in	mg/kg as	% of ¹⁴ C in	mg/kg as
treatment,		(rinse + leaves)	glufosinate-	(rinse + leaves)	glufosinate-	(rinse + leaves)	glufosinate-
days		or roots	ammonium	or roots	ammonium	or roots	ammonium
0	rinse	59	12	-	-	-	-
0	leaves	25	5.1	0.4	0.07	13	2.7
0	roots	31	0.62	2.2	0.04	64	1.3
21	rinse	14	1.7	-	-	-	-
21	leaves	28	3.4	1.1	0.13	55	6.8
21	roots	31	2.1	2.0	0.14	63	4.3
146	rinse	2.3	0.05	0.3	0.006	0.2	0.005
146	leaves	24	0.49	2.7	0.055	67	1.4
146	roots	19	0.18	6.0	0.055	68	0.63

Table 11. Identified residues in genetically modified sugar beets after 2 treatments with [3,4-¹⁴C]glufosinate-ammonium (Allan, 1996).

In another trial transgenic maize plants were treated twice (40 cm and 60 cm growth stages) with $[3,4^{-14}C]$ glufosinate-ammonium at rates equivalent to 0.50 kg ai/ha (Burnett, 1994). Plants were sampled on the days of treatment and at intervals until 102 days after the second treatment. The levels of ¹⁴C as glufosinate-ammonium were 23 and 5.8 mg/kg 1 hour and 5 days after the first treatment respectively and 14.5 and 9.9 mg/kg 1 hour and 5 days after the second.

Glufosinate-ammonium was generally a minor component of the residue whereas the main component in the forage, silage and fodder was NAG and the main component in the grain, cobs and husks was MPP. 73-83% of the residue was identified except in the grain where the very low residue level made further identification difficult.

A sub-sample of the maize forage was analysed by a GLC enforcement analytical method (Czarnecki and Bertrand, 1994), in which samples are extracted with distilled water, and after cleanup which includes ion-exchange chromatography, the residues are derivatized with trimethyl orthoacetate for GLC analysis. The enforcement method and the radiolabel method were in reasonable agreement (Table 13).

Table 12. Identified residues in grain and animal feeding commodities from transgenic maize treated twice with [3,4-¹⁴C]glufosinate-ammonium (Burnett, 1994). Forage and silage were sampled 28 and 55 days respectively after the second treatment and other commodities after 102 days.

Compound		Residue components as % of ¹⁴ C in the sample and as mg/kg glufosinate-ammonium											
	Forage		Silage		Foc	Fodder		Grain ¹		Cobs		Husks	
	¹⁴ C %	mg/kg	¹⁴ C %	mg/kg	¹⁴ C %	mg/kg	¹⁴ C %	mg/kg	¹⁴ C %	mg/kg	¹⁴ C %	mg/kg	
Glufosinate	13	0.35	11	0.20	9.9	0.20	1.5	0.002	2.6	0.006	2.1	0.018	
MPP	12	0.32	12	0.21	11	0.22	33	0.043	44	0.10	41	0.36	
NAG	52	1.4	55	0.98	54	1.1	9.1	0.012	20	0.046	19	0.17	
MPA	4.6	0.12	3.9	0.070	2.9	0.058	4.4	0.006	12	0.028	11	0.097	
Identified	82	2.2	82	1.5	76	1.6	58	0.076	79	0.18	73	0.64	
Total		2.6		1.8		2.0		0.13		0.25		0.87	

¹ MPB accounted for 9.8% of the ¹⁴C in the grain, equivalent to 0.013 mg/kg of glufosinate-ammonium.

Table 13. Comparison between analysis of maize forage from [3,4-¹⁴C]glufosinate-ammonium-treated maize by an enforcement GLC method (Czarnecki and Bertrand, 1994) and the ¹⁴C HPLC method (Burnett, 1994).

Analyte	Residue, mg/k	g as glufosinate-ammonium	
	HPLC ¹⁴ C GLC enforcement, not		Spike recovery, %, GLC
		adjusted for recovery	enforcement
Glufosinate	0.366	0.256	113
MPP	0.304	0.185	69.5
NAG	1.43	1.42	105

In another trial transgenic soya bean plants were treated twice at third trifoliate leaf and full bloom growth stages with $[3,4-^{14}C]$ glufosinate-ammonium at rates equivalent to 0.50 kg ai/ha (Rupprecht and Smith, 1994). Forage was sampled just before the second application and the mature crop was harvested 85 days after the second treatment.

NAG was the main residue in all the samples. The levels of MPP were greater than those of glufosinate in the pods and beans. 90-94% of the residue was identified in all the samples. The results are shown in Table 14. They were also reported by Rupprecht *et al.* (1996b).

The samples were extracted with distilled water or acetonitrile + water and analysed by the method of Czarnecki and Bertrand (1994). The GLC and HPLC radiolabel methods produced similar results (Table 15) at the higher levels but at low levels those from the enforcement method were lower.

Table 14. Identified residues in beans, pods, forage and straw from transgenic soya plants treated twice with [3,4-¹⁴C]glufosinate-ammonium (Rupprecht and Smith, 1994).

Compound		Identified residue components as % of ¹⁴ C in the sample and as glufosinate-ammonium, mg/kg								
	Forage		Straw		Pods		Beans			
	¹⁴ C %	mg/kg	¹⁴ C %	mg/kg	¹⁴ C %	mg/kg	¹⁴ C %	mg/kg		
Glufosinate	23	0.45	19	0.58	5.8	0.29	6.2	0.091		
MPP	6.5	0.13	14	0.42	22	1.1	16	0.23		
NAG	60	1.2	53	1.7	63	3.1	61	0.89		
MPA	0.7	0.014	5.7	0.18	2.9	0.142	7.1	0.10		
Identified	91	1.8	91	2.8	94	4.6	90	1.3		
Total		1.9		3.2		4.7		1.4		

Sample			Residue, mg/k	g as glufosin	ate-ammonium
	Extract	Method	Glufosinate	NAG	MPP
Forage	water	¹⁴ C HPLC	0.364	0.962	0.101
Forage	water	GLC	0.321	1.04	0.053
Straw	water	¹⁴ C HPLC	0.542	1.58	0.404
Straw	water	GLC	0.348	1.89	0.202
Pods	water	¹⁴ C HPLC	0.250	2.88	1.02
Pods	water	GLC	0.238	3.23	0.781
Beans	water	¹⁴ C HPLC	0.073	0.687	0.187
Beans	water	GLC	0.034	0.605	0.129
Forage	acetonitrile + water	¹⁴ C HPLC	0.084	0.195	0.025
Forage	acetonitrile + water	GLC	0.046	0.156	0.022

Köcher and Becker (1991) applied [¹⁴C]glufosinate-ammonium at the 7-leaf stage to the leaves of glufosinate-resistant and unmodified tomato plants. After intervals of 1 and four days foliar absorption of the ¹⁴C was the same in both types of plant, but translocation of the ¹⁴C from the treated leaves to shoots, other leaves and roots was approximately four times as high in the resistant plants.

Stumpf *et al.* (1995a) treated genetically modified tomato plants at the 7-8 leaf stage with $[3,4-{}^{14}C]$ glufosinate-ammonium at a rate equivalent to 0.8 kg ai/ha. Samples of plants were taken on the day of treatment and subsequently at intervals up to maturity, 74 days after treatment. The composition of the surface residues (in the rinse) and absorbed residues is shown in Table 16.

The major part (83-97%) of the surface residue in the rinse was glufosinate itself, even after long intervals. Glufosinate was very rapidly converted to NAG once absorbed into the tomato leaves and eventually accounted for about half of the residue, which is in accord with the rapid conversion of L-glufosinate and the stability of both D-glufosinate and NAG. The composition of the residue in the stems followed a similar pattern except that NAG accumulated to almost 70% of the residue and glufosinate fell to about 30%.

There were no detectable residues on the tomato fruit surface, probably because the plants were sprayed before any fruit were formed. NAG accounted for about 90% of the residue in the fruit, with glufosinate itself at essentially negligible levels. MPP was a minor residue in the fruit. NAG is evidently sufficiently mobile to translocate to the fruit, but D-glufosinate is not.

The composition of the glufosinate surface residue on the tomato leaves and stems on day 74 was shown to consist essentially of equal amounts of D- and L-glufosinate. The composition was quite different in the absorbed residue in the leaves (90% D- and 10% L-glufosinate) and in the stems (86% D- and 14% L-glufosinate), showing that within the tissue the L-isomer was more rapidly metabolized

In a separate experiment Stumpf *et al.* (1995a) placed genetically modified and normal tomato plants in solutions of labelled glufosinate for 2 days and examined the uptake and composition of the residue in the plants. The genetically modified plants took up much more glufosinate and converted about half of it to NAG. In the normal tomato plants about 4% was converted to MPP and the D- and L-glufosinate remained approximately equal.

S	ample	Days after		% of ¹⁴ C, mg/kg as glufosinate-ammonium			nonium	
	-	treatment	glufosinate-	ammonium	M	PP	NA	AG
			¹⁴ C %	mg/kg	¹⁴ C %	mg/kg	¹⁴ C %	mg/kg
leaves	Rinse	0 (2 h)	94		1.8		0.0	
leaves	Rinse	0 (4 h)	94		1.5		0.0	
leaves	Rinse	1	95		1.5		0.0	
leaves	Rinse	4	94		1.6		0.3	
leaves	Rinse	6	93		1.8		0.6	
leaves	Rinse	32	93		1.4		1.2	
leaves	Rinse	74	86		4.1		3.3	
leaves	Rinse	74	83		7.2		3.9	
leaves	After rinsing	0 (2 h)	69		0.0		27	
leaves	After rinsing	0 (4 h)	61		0.0		36	
leaves	After rinsing	1	64		0.0		33	
leaves	After rinsing	4	52		1.0		42	
leaves	After rinsing	6	53		2.4		41	
leaves	After rinsing	32	57		3.5		37	
leaves	After rinsing	74	44		5.6		51	
leaves	After rinsing	74	42	2.2	2.5	0.13	50	2.6
leaves	Not rinsed	12	30		1.6		65	
leaves	not rinsed	14	17		0.0		83	
leaves	not rinsed	27	0.0		0.0		100	
leaves	not rinsed	32	75		2.7		20	
leaves	not rinsed	74	51	4.4	2.5	0.22	42	3.6
stems	rinse	0 (2 h)	95		1.0		0.0	
stems	rinse	0 (4 h)	97		0.5		0.0	
stems	rinse	1	93		1.9		1.2	
stems	rinse	4	94		1.1		2.0	
stems	rinse	6	94		1.1		2.3	
stems	rinse	32	88		2.1		5.7	
stems	rinse	74	87		2.2		7.4	
stems	rinse	74	92		0.0		7.7	
stems	after rinsing	0 (2 h)	75		0.0		16	
stems	after rinsing	0 (4 h)	61		0.0		39	
stems	after rinsing	1	58		0.0		39	
stems	after rinsing	4	42		0.0		58	
stems	after rinsing	6	45		0.0		53	
stems	after rinsing	32	32		6.4		61	
stems	after rinsing	74	31		-		69	
stems	after rinsing	74	26	0.51	3.8	0.07	67	1.3
stems	not rinsed	12	8.6		0.0		91	
stems	not rinsed	14	7.2		0.0		93	
stems	not rinsed	27	0.0		0.0		100	
stems	not rinsed	32	47		3.6		49	
stems	not rinsed	74	37	0.95	2.6	0.07	57	1.5
fruit	rinse	74	0		0		0	
fruit	after rinsing	74	0.0	-	6.2	0.009	94	0.14
fruit	not rinsed	74	0.0	-	9.5	0.02	85	0.17
fruit, green	not rinsed	32	1.7		4.6		88	
fruit, red	not rinsed	60	0.0		0.0		100	
fruit, red	not rinsed	74	0.0		0.0		100	

Table 16. Residues in genetically modified tomato plants at the 7-8 leaf stage after treatment with $[3,4^{-14}C]$ glufosinate-ammonium (Stumpf *et al.*, 1995a).

Figure 2. Proposed metabolic pathways of glufosinate-ammonium in plants.



Environmental fate in soil

Aerobic degradation

Allan (1995) studied the aerobic degradation of [3,4-¹⁴C]glufosinate-ammonium in a sandy loam soil (pH 6.0, organic carbon 1.8%) both when applied directly to the soil at 2 mg/kg and when incorporated as a residue in bushbean leaves which, having been desiccated with



Labelled glufosinate-ammonium

labelled glufosinate-ammonium, contained ¹⁴C equivalent to 76 mg/kg of glufosinate-ammonium. The bean leaves (1.66 g) were added to 50 g of soil giving a theoretical level of 2.5 mg/kg. The soil was adjusted to 40% of maximum water holding capacity and incubated at 20°C in the dark. The results are shown in Table 17.

Glufosinate disappeared very quickly with a half-life of 3-6 days (Table 18). The main product was MPP, reaching its peak after about 14 days. MPA was also an important product and in the absence of plant material became the main residue after long intervals. NAG was a very minor soil residue.

Days		Residues and	evolved ¹⁴ CO ₂	2 expressed as % of	applied ¹⁴ C.		
incubated		Soil treatment		Plant incorporation			
	extractable	Unextractable	$^{14}\text{CO}_2$	extractable	Unextractable	$^{14}CO_2$	
0	106	1.8	-	97	9.7	0	
1	80	8.3	0.6	90	14	0.1	
3	82	11	1.5	87	17	0.4	
7	77	15	5.6	80	20	1.8	
14	70	21	11	75	21	4.1	
21	61	21	17	69	24	7.6	
28	60	20	23	68	28	8.6	
41	46	19	26	58	20	9.5	
59	32	24	43	32	26	22	
90	8.7	25	57	23	29	26	
120	5.6	21	62	18	23	31	

Table 17. Distribution of ¹⁴Cfrom [3,4-¹⁴C]glufosinate-ammonium incubated aerobically in a soil in the dark or incorporate-d into the soil as a plant-metabolized residue (Allan 1995).

Table 18. Composition of residues resulting from aerobic incubation of [3,4-¹⁴C]glufosinate-ammonium in a soil in the dark or incorporated into the soil as a plant-metabolized residue (Allan 1995).

Days				% of ap	plied ¹⁴ C			
incubated		Soil trea	atment		Plant incorporation			
	glufosinate	MPP	NAG	MPA	glufosinate	MPP	NAG	MPA
0	104	-	-	-	75	14	3.2	-
1	77	11	2.3	-	70	18	-	-
3	56	19	2.4	2.1	47	30	4.8	2.2
7	32	31	2.2	6.4	15	52	-	7.3
14	18	37	0.7	11	2.9	62	-	6.3
21	7.8	33	0.4	15	0.6	56	-	9.2
28	3.3	34	-	19		61	-	6.9
41	0.9	22	0.3	20	-	55	-	0.3
59	0.5	13	-	17		30	-	-
90	0.1	1.4	-	6.8		23	-	-
120	0.5	1.6	-	1.6	-	17	-	-

Field dissipation

In a 1-year field dissipation study in California Belcher (1996a) applied glufosinate-ammonium three times to bare ground at rates of 1.7 kg ai/ha between rows in a level vineyard. The soil was a coarse sandy loam. Irrigation supplemented rainfall and followed the growers' standard practices. Soil samples were analysed for glufosinate, MPP and MPA, the main residues. The results are shown in Table 19.

Glufosinate dissipated rapidly, possibly owing to increasing soil moisture and temperature, with calculated half-lives of 15, 7.2 and 2.7 days after the first, second and third applications respectively.

The maximum MPP residues were 0.11, 0.16 and 0.14 mg/kg on days 30, 10 and 5 after the first, second and third applications respectively and the estimated half-lives were 38, 14 and 16 days from these times. The estimated half-lives for MPA were 25, 19 and 7 days for the successive

applications. MPA residues reached their highest levels of 0.06, 0.06 and 0.04 mg/kg on days 60, 10 and 5 after the first, second and third applications respectively.

No residues were detected below the 45-60 cm depth segment. Glufosinate was detected once at the analytical LOD (0.01 mg/kg) after the first application and at 0.02 mg/kg 5 and 10 days after the final application in the 45-60 cm samples. Glufosinate was detectable at low levels 5 and 10 days after application in the 30-45 cm samples, but not at other times. MPP occurred in the 30-45 cm segment on several occasions at 0.01-0.02 mg/kg. MPP was detected only once in the 45-60 cm depth segment, 10 days after the final application at 0.01 mg/kg. MPA was not detected in any sample from 30-45 cm or deeper.

Glufosinate and its soil degradation products have some mobility but their further degradation ensures that travel down the soil profile is limited.

Table 19. Residues of glufosinate and its degradation products at intervals after three applications of glufosinate-ammonium at 1.7 kg ai/ha to bare ground in a level California vineyard (Belcher, 1996a).

Interval after		Resid	dues, mg/kg as glut	fosinate free acid, i	n soil	
application	glufo	sinate	М	PP	M	PA
days	soil 0-15 cm	soil 15-30 cm	soil 0-15 cm	soil 15-30 cm	soil 0-15 cm	soil 15-30 cm
Application 1						
0	0.34	< 0.01	0.01	< 0.01	< 0.01	< 0.01
5	0.26	< 0.01	0.04	< 0.01	< 0.01	< 0.01
10	0.13	< 0.01	0.06	< 0.01	0.01 < 0.01	< 0.01
15	0.16	0.05 < 0.01	0.07	< 0.01	0.02	< 0.01
28	0.1	< 0.01	0.10	< 0.01	0.03	< 0.01
47	0.04	< 0.01	0.10	< 0.01	0.04	< 0.01
65	0.02	< 0.01	0.05	0.02	0.05	< 0.01
93	< 0.01	< 0.01	0.02	0.02	0.04	< 0.01
124	< 0.01	< 0.01	< 0.01	< 0.01	0.01	< 0.01
Application 2						
0	0.47	< 0.01	0.03	< 0.01	0.02	< 0.01
5	0.10	0.07	0.09	0.03	0.03	< 0.01
10	0.02	0.08	0.08	0.04	0.05	0.01
15	0.02	0.03	0.06	0.03	0.05	0.01
30	< 0.01	0.02 < 0.01	0.03	0.02	0.02	0.01
45	< 0.01	0.01	0.01	0.03	0.01 < 0.01	0.01 < 0.01
59	< 0.01	< 0.01	< 0.01	< 0.01	< 0.01	< 0.01
90	< 0.01	< 0.01	< 0.01	< 0.01	< 0.01	< 0.01
119	< 0.01	< 0.01	< 0.01	< 0.01	< 0.01	< 0.01
Application 3						
0	0.41	< 0.01	0.02	< 0.01	< 0.01	< 0.01
5	0.03	0.04	0.06	0.04	0.03	0.01
10	0.01	0.01	0.04	0.02	0.02	0.02
15	0.01	< 0.01	0.03	0.01	0.01	0.01
30	< 0.01	< 0.01	0.02	< 0.01	< 0.01	< 0.01
45	< 0.01	< 0.01	0.02	< 0.01	< 0.01	< 0.01
60	< 0.01	< 0.01	< 0.01	< 0.01	< 0.01	< 0.01
93	< 0.01	< 0.01	< 0.01	< 0.01	< 0.01	< 0.01

Stumpf (1993a) incubated 3.6 mg/kg $[3,4^{-14}C]N$ -acetyl-L-glufosinate disodium salt in a sandy loam soil (pH 5.8, organic carbon 1.1%) in the dark at 20°C under aerobic conditions for 62 days. The results are shown in Table 20.

N-acetyl-L-glufosinate was very rapidly degraded to L-glufosinate, which was then further degraded. The degree of mineralization (40%) and the level of bound residues (21%) in 62 days parallel the 43% mineralization of glufosinate-ammonium and 24% unextractable residues in 59 days reported by Allan (1995), and suggest that the degradation pathway for NAG is through glufosinate.

Table 20. Aerobic soil degradation of [3,4-¹⁴C]*N*-acetyl-L-glufosinate disodium salt in soil (Stumpf, 1993c).

Day		Resi	dues and	evolved ¹⁴	CO ₂ expre	essed as %	of applie	$d^{14}C$ and	as NAG n	ng/kg	
	$^{14}CO_2$	N	AG	L-glufosinate		MPP		MPA		bound residues	
	%	%	mg/kg	%	mg/kg	%	mg/kg	%	mg/kg	%	mg/kg
0.1	0	47	1.7	47	1.7	2.1	0.08	1.0	0.04	0.7	0.03
1	0.5	8.7	0.31	80	2.9	6.7	0.24	1.3	0.05	1.5	0.05
3	1.0	3.1	0.11	69	2.5	20	0.73	3.4	0.12	1.0	0.04
23	15	1.4	0.05	16	0.57	40	1.4	9.5	0.34	11	0.38
41	24	1.6	0.06	4.7	0.17	39	1.4	9.9	0.36	13	0.44
62	40	0	0	1.5	0.05	27	0.98	9.9	0.36	21	0.75

Stumpf *et al.* (1995c) incubated 0.5 mg/kg $[2^{-14}C]^2$ -methylphosphinicoacetic acid (MPA) in a sandy loam (pH 5.8, organic carbon 1.1%) and a loamy sand (pH 4.9, organic carbon 2.4%) in the dark at 20°C under aerobic conditions for 122 days and measured the rate of degradation and mineralization (Table 21). No important degradation products other than CO₂ were identified.

In the sandy loam bound residues accounted for 22% and 25% of the applied ¹⁴C on days 28 and 122 respectively. Estimated decline and mineralization half-lives were 24 and 74 days respectively, but the curves were not simple first-order and the rates were slower at longer intervals. The rates in the loamy sand were much slower, with only 20% mineralization after 122 days and a decline half-life of 120-160 days. Bound residues again accounted for about 25% of the applied ¹⁴C throughout the study.

$1000 21$. Refoble son degradation of $[2^{-10} \text{ C}]^2$ methylphosphineodectic dela (Stampi et al. 1995)	Table 21. Aerobic soil degradation of [2-	¹⁴ C]2-methylphosphinicoacetic	acid (Stumpf et al.	1995c).
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day		loamy sand		sandy loam				
	$^{14}CO_2$	M	PA	$^{14}CO_2$	М	PA		
	% of applied ¹⁴ C	% of applied ¹⁴ C	mg/kg	% of applied ¹⁴ C	% of applied ¹⁴ C	mg/kg		
0		87	0.47		99	0.54		
3	0.7	89	0.48	4.0	90	0.49		
7	1.8	86	0.47	6.6	89	0.49		
14	4.2	76	0.42	14.5	77	0.42		
28	7.9	69	0.38	31	47	0.26		
60	17	62	0.34	63	45	0.25		
94	23	53	0.29	75	3.8	0.002		
122	20	56	0.30	75	3.5	0.002		

Stumpf *et al.* (1995d) incubated [3,4-¹⁴C]glufosinate-ammonium at 1.9 mg/kg and [3-¹⁴C]MPP at 0.9 mg/kg in the same sandy loam soil in the dark at 10°C under aerobic conditions for 120 days. The results are shown in Table 22 and 23.

The estimated half-life for glufosinate-ammonium disappearance was 24 days; the rate of mineralization was slow with an estimated half-life of 300 days or more. MPP and subsequently MPA were major components of the residue after the longer periods of incubation. NAG and

AE F065594 (4-methylphosphinico-2-oxobutanoic acid) and possibly AE F086486 (3-methylphosphinico-3-oxopropionic acid) were identified as minor products.

The estimated half-life for MPP disappearance and mineralization were 86 days and 220 days respectively. MPA was the only product of significance accounting for 40% of the extractable residue by day 120.

Stumpf and Zumdick (1998) further investigated the nature of the minor soil product previously identified as AE F086486 and, using LC-MS-MS, identified the compound as 3-methylphosphinicoacrylic acid (AE 0015081). The structure was confirmed by synthesis.



AE 0015081 3-methyl-phosphinico acrylic acid

Table 22. Residues resulting from incubation of 1.9 mg/kg $[3,4-{}^{14}C]$ glufosinate-ammonium in a sandy loam soil in the dark at 10°C under aerobic conditions for 120 days (Stumpf *et al.*, 1995d).

Incubation,			Residues, % of	applied ¹⁴ C		
days	Extractable	Glufosinate	MPA	MPP	NAG/	AE F086486 ¹
	residue				AE F065594	
0	94	90	<0.5	1.6	1.8	<0.5
1	93	85	<0.5	5.3	2.5	<0.5
4	91	72	1.9	12	1.8	2.3
7	86	57	3.3	20	2.2	4.0
14	87	42	5.5	32	2.6	4.9
21	87	32	7.9	38	2.5	7.1
30	86	25	9.5	43	1.8	6.8
56	79	9.9	14	45	2.5	6.7
91	69	2.0	19	41	3.1	4.1
120	69	5.4	22	36	2.7	3.9

¹Subsequently identified as AE 0015081 (Stumpf and Zumdick, 1998).

Table 23. Aerobic degradation of $[3^{-14}C]3$ -methylphosphinicopropionic acid (MPP) incubated at 0.9 mg/kg in a sandy loam soil in the dark at 10°C for 120 days (Stumpf *et al.*, 1995d).

Incubation, days		Residues, %	of applied ¹⁴ C	
	Extractable	CO ₂	MPP`	MPA
	residue			
0	97	0	97	<1.0
1	95	0.1	95	<1.0
4	96	0.6	94	1.8
7	90	1.1	87	3.1
14	91	3.0	79	11.4
21	89	5.4	80	9.0
30	90	6.1	76	13.5
56	74	18	49	25
91	68	21	46	22
120	62	32	37	25

Zumdick (1995a) incubated 2.1 mg/kg [3,4-¹⁴C]L-*N*-acetyl-glufosinate in a sandy loam and a loamy sand under aerobic conditions for 120 days at 20°C in the dark (Table 24). In a second experiment glufosinate-tolerant tomato leaves containing residues equivalent to 1 mg glufosinate-

ammonium per kg soil were incorporated into the sandy loam and incubated under the same conditions (Table 25). Bound residues accounted for about 14-23% of the applied residue in each of the three incubations from days 28 to 120.

N-acetyl-glufosinate disappeared very quickly, with half-lives of only hours, producing initially L-glufosinate, which was then degraded further to MPP, MPA and CO_2 . MPP became the major component of the residue after 1-3 days.

The tolerant-tomato residue comprised mainly glufosinate and *N*-acetyl-glufosinate. After 3 days incubation in the sandy loam soil most of the residue had been converted to MPP, which itself was degraded more slowly to MPA and ultimately to CO_2 .

Table 24. Residues resulting from incubation of 2.1 mg/kg $[3,4-^{14}C]$ L-*N*-acetyl-glufosinate in a sandy loam and a loamy sand under aerobic conditions for 120 days at 20°C in the dark (Zumdick 1995a).

Days]	Residues, %	of applied 1	⁴ C			
			loamy sand					sandy loam		
	NAG	CO ₂	L-glufosinate	MPP	MPA	NAG	CO_2	L-glufosinate	MPP	MPA
0	70	-	20	8.4	0.4	61	-	30	4.7	0.0
0.25	14	0.3	58	13	1.2	29	0.2	58	8.1	0.7
1	15	1.1	23	35	4.7	9.2	0.9	49	22	2.2
2	6.2	3.6	23	38	5.9	4.3	2.3	41	29	3.4
3	5.1	2.8	16	42	7.4	3.1	4.9	24	37	5.9
7	4.1	11	6.3	36	9.6	1.8	1.1	38	34	8.1
14	2.8	21	5.2	27	7.6	1.0	10.5	2.6	39	19
28	1.3	39	2.2	5.2	0.8	0.4	23	1.0	28	18
62	0.5	49	1.3	1.9	0.0	0.2	56	1.0	2.1	1.2
90	0.5	52	1.1	0.7	0.0	0.4	67	1.0	0.7	0.3
120	0.3	72	0.4	0.1	0.0	0.2	66	0.4	0.5	0.1

Table 25. Residues resulting from incubation of [3,4-¹⁴C]glufosinate metabolized in tolerant tomato leaves, incorporated at 1 mg glufosinate-ammonium equivalent per kg soil, in a sandy loam under aerobic conditions for 120 days at 20°C in the dark (Zumdick 1995a).

Days		Res	sidues, % of applied	¹⁴ C	
	glufosinate	CO ₂	NAG	MPP	MPA
0	46	-	33	4.1	0.3
0.25	47	0.0	27	4.3	0.4
1	44	0.0	30	8.9	0.6
2	27	0.1	20	27	4.1
3	5.4	0.2	2.9	52	16
7	1.4	5.3	0.4	47	25
14	0.7	5.8	0.0	40	18
28	0.6	16	0.0	31	0.6
62	0.7	48	0.0	6.7	1.2
90	0.2	49	0.0	0.8	0.0
120	0.3	43	0.0	2.4	0.0

Zumdick (1995b) incubated 1 mg/kg $[3,4-^{14}C]L$ -*N*-acetyl-glufosinate in a sandy loam from Germany and a silt loam from Nebraska under aerobic conditions for 120 days at 20°C in the dark (Table 26). The results were consistent with other similar experiments, with the NAG rapidly forming glufosinate, which was in turn converted to MPP and MPA. An additional product (AE F130947, methylphosphinicoformic acid) was separated from the others by the HPLC systems used. Bound residues from days 14 to 90 constituted 14-22% of the applied ¹⁴C.



n												
Days					I	Residues,	% of ap	plied ¹⁴ C				
			sand	y loam			silt loam					
	NAG	CO_2	L-glufos	MPP	MPA	130947	NAG	CO ₂	L-glufos	MPP	MPA	130947
0.1	91	-	9.7	0.8	0.1	< 0.1	85	-	13	1.7	0.1	< 0.1
0.25	16	0.2	66	7.3	0.6	0.4	26	0.3	36	11	0.3	0.3
1	7.5	0.8	48	21	1.8	2.0	11	1.7	46	23	3.2	2.8
3	4.7	7.0	26	35	5.2	4.2	3.5	6.3	26	25	11	5.9
7	3.2	16	10	32	9.1	6.3	2.4	15	11	18	21	10
14	1.9	29	4.3	20	12	7.3	0.7	36	2.6	2.8	18	8.7
21	1.2	41	6.1	12	8.8	5.3	0.3	44	2.8	1.9	15	4.5
30	0.7	48	4.2	5.8	5.3	3.9	0.3	59	1.5	0.4	4.4	1.3
59	1.3	64	1.5	1.5	0.2	0.3	< 0.1	63	0.8	< 0.1	< 0.1	< 0.1
90	0.3	68	0.6	0.3	< 0.1	< 0.1		65				
120		77						67				

Table 26. Residues resulting from incubation of 1 mg/kg $[3,4-^{14}C]$ L-*N*-acetyl-glufosinate in a sandy loam and a silt loam under aerobic conditions for 120 days at 20°C in the dark (Zumdick 1995b).

Stumpf *et al.* (1989) subjected $[3,4-^{14}C]$ glufosinate-ammonium in a thin layer of a microbiologically active sandy loam soil to UV irradiation (xenon arc with 290 nm cut-off filters) for 35 days in a 12 hours light-dark cycle at 25°C and identified the products. The calculated degradation half-life was 35 days, with 7.6% production of $^{14}CO_2$ in that time. Three degradation products were identified: MPP (6-21% of the applied ^{14}C), MPA (0-5%) and NAG (0-10%). Three other products were in too small amounts (0-3% and 4-9%) to be identified. Degradation was caused by microbiological processes and photolysis in the presence of humic acids.

In a rotational crop study Campbell and Bennett (1997) treated bare ground plots in North carolina, Missouri and California twice at 10-day intervals with glufosinate-ammonium at 0.39 and 0.54 kg ai/ha and planted winter wheat 75 and 90 days after the final treatment. No residues of glufosinate or MPP were detected (LOD 0.05 mg/kg) in forage, hay, grain or straw samples. No uptake of glufosinate or MPP should occur in winter wheat following a previous use of glufosinate-ammonium.

In a confined rotational crop study, Meyer *et al.* (1995) applied [3,4-¹⁴C]glufosinateammonium to a bare sandy loam soil in a greenhouse in stainless steel tanks at 1.0 kg ai/ha and planted radishes, lettuce and wheat 28 and 119 days later. The intervals represented resowing after crop failure and immediate recropping. Residue levels are shown in Table 27. Product A was not identified, but was characterized as a relatively simple polar unconjugated compound that may be a degradation product taken up by roots. The products were similar in the root, leafy and small cereal grain crops. Levels were much lower after the longer sowing interval. Residues should be undetectable or very low in rotational crops.

Commodity			Res	idues, mg/kg, as	glufosinate fr	ee acid						
	1	28 days inter	val to sow	ving	119 days interval to sowing							
	total ¹⁴ C	total ¹⁴ C MPP MPA Comp			total ¹⁴ C	MPP	MPA	Compoun				
				А				d A				
Radish top	0.110	0.053	0.006	0.021	0.013 ¹							
Radish root	0.090	0.047	0.008	0.024	0.009^{1}							
Lettuce	0.079	0.020	0.006	0.013	0.013 ¹							
Wheat forage	0.30	0.17	0.019	0.027	0.047^{1}							
Wheat straw	0.78	0.35	0.073	0.19	0.14	0.029	0.008	0.035				
Wheat grain	0.33	0.11	0.035	0.035	0.12	0.015	< 0.001	0.016				

Table 27. ¹⁴C residues in rotational crops after application of [3,4-¹⁴C]glufosinate-ammonium at 1.0 kg ai/ha to a bare sandy loam soil (Meyer *et al.*, 1995).

¹Residue too low for characterization.

Figure 3. Proposed aerobic soil degradation pathways of glufosinate-ammonium.



Environmental fate in water/sediment systems

Stumpf and Schink (1992) subjected [3,4-¹⁴C]glufosinate-ammonium dissolved in surface water from a gravel pit to UV irradiation for 118 hours (equivalent to 33 days sunlight) at 25°C. Glufosinate suffered little degradation under these conditions, with 3-5% conversion to MPP and 0.2% mineralization. Photolytic degradation of glufosinate in surface waters is a minor degradation route.

Stumpf (1993b) incubated $[3,4-^{14}C]$ glufosinate-ammonium at 0.1 mg/kg in aerobic systems consisting of 180 ml water and 20 g sediment at 20°C and 8°C for 361 days. The sediment was sandy with 0.4% organic carbon.

Glufosinate disappeared from the water/sediment system with half-lives of 3 and 20 days at 20°C and 8°C respectively. Mineralization occurred to the extent of only 25 and 20% at the two temperatures in the 361 days of the study. Within a few days at 20°C and after 29 days at 8°C MPP became the major component of the residue. Seven other products were observed, but five were very minor. Two others amounting to 5% and 20% of the applied dose at various times were not positively identified. At all times most of the total residue was in the aqueous phase.

In sterile (autoclaved) samples incubated for 29 and 120 days up to 50% of the residue was converted to NAG in some samples, but heat resistant bacteria may have been present.

Table 28. Residues resulting from incubation of [3,4-¹⁴C]glufosinate-ammonium in a water/sediment system under aerobic conditions at 8°C and 20°C (Stumpf 1993e).

Days				Resi	dues, % of	applied ¹⁴ C				
			20°C			8°C				
	Glufosinate	CO ₂	MPP	NAG	MPA	Glufosinate	CO ₂	MPP	NAG	MPA
0	95	-	4.4	2.6	0.0	95	-	4.4	2.6	0.0
1	60	0.3	21	14	1.0	84	0.1	7.8	7.0	0.3
4	39	0.8	40	9.8	1.4	61	0.3	15	10.3	0.4
7	24	1.5	49	5.7	3.4	56	0.3	21	14	0.7
14	7.1	4.4	59	3.9	3.2	48	0.7	28	12	1.5
21	0.3	8.2	56	2.4	2.9	45	1.5	32	6.5	2.8
29	0.1	12	53	1.9	3.8	33	1.7	39	3.7	3.5
60	0.0	18	52	0.2	2.9	15	5.4	44	3.3	3.3
90	0.0	21	52	0.1	3.0	2.7	6.7	49	4.2	3.7
120	0.0	17	47	0.0	4.0	0.0	9.0	51	3.0	3.3
238	0.0	23	46	0.0	5.6	0.0	17	45	0.0	2.3
361	0.0	25	48	0.0	5.6	0.0	20	55	0.0	1.4

Stumpf (1994a) incubated [3,4-¹⁴C]glufosinate-ammonium at 1 and 0.1 mg/kg in two aerobic systems consisting of 180 ml water and 20 g sediment at 20°C in the dark for 130 days (Tables 29 and 30).

The mineralization in130 days in the gravel pit sediment was 7% at the lower dose and 12% at the higher dose. The half-lives for the degradation of glufosinate itself were 11, 91 and 1.4 days for the loamy river water sediment at 1 mg/kg and the gravel-pit sediment at 1 mg/kg and 0.1 mg/kg respectively. MPP was the main product identified and it was quite persistent in both systems. In all cases most of the residue was in the water phase.

As in a previous experiment (Stumpf, 1993b) a number of minor products were detected but not identified. Two products, sometimes constituting 10-20% of the residue, were further investigated and identified as AE F086486 and possibly AE F130947 (methylphosphinicoformic acid).

In a subsequent study, Stumpf (1994b) used glufosinate labelled in different positions and a higher dose rate to produce sufficient material for positive identification as AE F130947, also identified as a soil degradation study (Zumdick, 1995b).

Stumpf and Zumdick (1998) corrected the identification of AE F086486 (3-methylphosphinico-3-oxopropionic acid) to AE 0015081 and confirmed AE F130947.



AE 0015081 3-methyl-phosphinico acrylic acid

Days			R	esidues, %	of applied ¹	¹⁴ C in whole s	ystem			
		gravel-pit	t sediment, i	1 mg/kg		gravel-pit sediment, 0.1 mg/kg				
	Glufosinate	CO_2	MPP	NAG	MPA	Glufosinate	CO ₂	MPP	NAG	MPA
0	91	-	2.0	0.7	0.0	89	-	2.9	1.7	0.0
1	86	0.0	6.8	2.0	0.4	59	0.0	23	9.9	0.9
3	74	0.0	15	4.0	0.7	21	0.2	55	7.5	3.5
7	69	0.2	19	4.1	2.4	3.8	1.0	79	1.8	4.7
14	60	0.5	25	3.7	4.1	0.1	2.8	80	0.3	5.7
21	57	0.8	27	2.5	6.2	0.0	4.4	78	0.0	4.6
30	46	1.9	28	3.3	10.0	0.0	6.8	76	0.0	4.1
50	39	2.9	31	4.3	12	0.0	8.6	76	0.0	4.2
77	36	4.3	30	2.9	14	0.0	10.6	74	0.0	4.4
91	37	5.0	30	1.4	14	0.0	12	70	0.0	5.1
130	33	7.1	30	2.1	16	0.0	12	67	0.0	6.8

Table 29. Residues resulting from incubation of [3,4-¹⁴C]glufosinate-ammonium in a gravel pit water/sediment system at 20°C in the dark for 130 days (Stumpf, 1994a).

Table 30. Residues resulting from incubation of [3,4-¹⁴C]glufosinate-ammonium in a loamy river water sediment system at 20°C in the dark for 130 days (Stumpf, 1994a).

Days		Resid	lues, % of applie	ed ¹⁴ C	
	Glufosinate	CO ₂	MPP	NAG	MPA
0	91	?	1.3	1.0	0.0
1	87	?	5.4	2.5	0.3
3	72	?	13	6.2	3.4
7	60		19	8.2	5.9
14	46		26	7.6	10.2
21	35		29	4.5	12
30	19		37	1.5	16
50	3.0		41	0.0	20
77	0.0		48	0.0	18
91	0.0		48	0.0	17
130	0.0		45	0.0	16



Figure 4. Proposed aerobic degradation pathways of glufosinate-ammonium in water/sediment systems

METHODS OF RESIDUE ANALYSIS

Analytical methods

The main components of the residue in genetically modified tolerant crops are glufosinate, NAG (*N*-acetyl-L-glufosinate) and MPP (3-methyl-phosphinico-propionic acid). Analytical methods have been designed to measure the three components separately or, because glufosinate and NAG produce the same derivative in the analytical procedure, to measure glufosinate and NAG combined and MPP separately.

Holzwarth (1995) described the methods used for residue analysis of tolerant crops. Residues are extracted from the finely ground sample with water. The clear extract, after separation from solid material, is passed through an anion exchange resin and the residues are eluted with formic acid. After evaporation of the formic acid the residue is taken up in a 1:1 ethanol/water mixture which is then applied to a cation exchange column. NAG and MPP are eluted with ethanol/water and glufosinate with aqueous ammonia.

After evaporation to dryness the residues from both fractions are taken up in glacial acetic acid and methylated and acetylated with trimethyl orthoacetate in refluxing acetic acid. After solvent exchange and a final silica gel cartridge clean-up the residues are determined by GLC with flame photometric detection.

Variations on the extraction and initial clean-up are needed for samples such as maize oil. The lower limit of determination for crop samples is typically 0.05 mg/kg for each of the three analytes. For fats and milk a mixture of n-propanol and water is used for extraction instead of water.

Figure 5. Derivatization of glufosinate, NAG and MPP.



Czarnecki *et al.* (1989) used a method (HRAV-5A) similar in principle to that described above for the determination of glufosinate and MPP in apples, grapes, soya beans, maize and tree nuts. Czarnecki and Bertrand (1993) described a similar method (HRAV-24) for the determination of glufosinate, MPP and NAG in maize and its processed fractions. Sochor *et al.* (1991) measured residues of glufosinate, MPP and NAG in tomatoes, tobacco and potatoes by the same method and achieved practical LODs of 0.05-0.2 mg/kg.

Czarnecki (1995a) analysed animal commodities for glufosinate and MPP by a similar method but without the cation exchange column clean-up, achieving LODs of 0.02 mg/kg for milk, 0.05 mg/kg for eggs, meat and fat, and 0.1 mg/kg for kidneys and liver. Czarnecki (1995b) included NAG in the same method with the same LODs. NAG and glufosinate appear in the same GLC peak so the LOD is for either compound or the combined residue.

Czarnecki and Bertrand (1994) described a comprehensive procedure for the determination of glufosinate, MPP and NAG in many commodities. The method (AE-24) is essentially as described above with many variations on the extraction depending on the nature of the sample. Recoveries should lie within the range 70-120% at the practical LOD of 0.05 mg/kg for each compound, expressed as glufosinate free acid. The method was determined and used on maize grain, silage, forage, fodder, starch, grits, flour, meal and hulls, and soya bean seed, hay, hulls and meal. The same method was described by Czarnecki (1995c). Bertrand (1994) validated the method (Table 31). Most tests were at a fortification level of 0.05 mg/kg, but recoveries did not seem to depend on concentration. Median recoveries were 95%, 95% and 102% for glufosinate, MPP and NAG respectively.

Bertrand (1994) pointed out that part of the glufosinate added to glufosinate-tolerant plant material could be converted to NAG during recovery testing. When racemic glufosinate was added to transgenic soya bean seed at 0.5 mg/kg, 44-54% was recovered as glufosinate and 33-43% as NAG. When D-glufosinate was used, recoveries of glufosinate were 85-100%. In some transgenic materials it is necessary to use D-glufosinate, which is not subject to enzymic *N*-acetylation, to determine the analytical recovery of glufosinate. The method is not selective for either isomer.

Table 31. Recoveries of glufosinate, MPP and NAG from maize grain, forage, fodder, silage, flour, meal, hulls, starch, and crude oil, and soya bean seeds, hulls, meal, hay/fodder and crude oil (Bertrand, 1994).

Recovery range, %	Nui	mber of tests	
	Glufosinate	MPP	NAG
61-70		4	1
71-80	5	13	0
81-90	16	28	4
91-100	17	30	19
101-110	17	23	18
111-120	7	15	8
121-130		4	6
131-140		1	

Castro and Dacus (1994) validated analytical method AE-24 (suitable for enforcement) on commodities from transgenic maize and soya beans. D-glufosinate was used to determine recoveries from soya bean seed. An LOD of 0.05 mg/kg for each compound on each commodity was achieved (Table 32). The time needed to analyse a batch of six samples was about 18 working hours over 3 working days.

Table 32. Recoveries by analytical method AE-24 from transgenic maize and soya bean commodities (Castro and Dacus, 1994).

Commodity	Glufo	osinate	М	PP	NAG		
	spiking levels,	Recovery range,	spiking levels, Recovery range,		spiking levels,	Recovery range,	
	mg/kg %		mg/kg	%	mg/kg	%	
Maize grain	0.05 0.25 73-98% (n=4)		0.05 0.25 73-78% (n=4)		0.10 0.50	89-107% (n=4)	
Maize forage	0.50 2.5 83-89% (n=4)		0.10 0.50 76-87% (n=4)		3.0 15	101-110% (n=4)	
Corn oil,	0.05 0.25	57-96% (n=4)	0.05 0.25	70-82% (n=4)	0.05 0.25	113-149% (n=4)	
refined							
Soya bean	0.05 0.25 65-89% (n=4)		0.10 0.50 80-91% (n=4)		0.20 1.0	85-109% (n=4)	
seed	(D-glufosinate)						

Holzwarth (1996a) determined the suitability of analytical method AE-24 for residues in glufosinate-tolerant maize shoots. Recoveries of spiked analyte over the concentration range 0.05-5 mg/kg were glufosinate 60-99% (n=6), NAG 65-85% (n=6) and MPP 53-99% (n=12). Holzwarth (1996b) also validated method AE-24 for residues in maize grain from plants susceptible to glufosinate-ammonium. Recoveries over the concentration range 0.05-5 mg/kg were: glufosinate 81-95% (n=6), NAG 68-82% (n=6) and MPP 54-94% (n=11).

Snowdon and Taylor (1995b) validated method AE-24 for glufosinate-tolerant maize, achieving LODs of 0.05 mg/kg for each analyte. Recoveries are shown in Table 33.

Commodity	Glufo	osinate	М	PP	NAG		
	spiking levels, Recovery range,		spiking levels, Recovery range,		spiking levels,	Recovery range,	
	mg/kg %		mg/kg	%	mg/kg	%	
Maize shoots	0.05 2.0 50	61-105 (n=7)	0.05 2.0 50	69-85 (n=14)	0.05 2.0 50	76-132 (n=7)	
Cobs	0.05 0.50 64-104 (n=5)		0.05 0.50	66-89 (n=10)	0.05 0.50	80-113 (n=5)	
Cob/spadix	0.05 0.50 95-131 (n=4)		0.05 0.50 74-105 (n=10)		0.05 0.50	97-122 (n=5)	

Table 33. Recoveries by analytical method AE-24 from transgenic maize commodities (Snowdon and Taylor, 1995b).

Czarnecki (1995c) validated analytical method AE-24A for maize and its processed products. The method differs from AE-24 in omitting the cation-exchange column step before derivatization. Consequently, the compounds appear as a single peak and are reported as a combined residue. Recoveries were acceptable for maize grain, forage and fodder.

Niedzwiadek and Bertrand (1995b) validated analytical method AE-24A for residues in glufosinate-tolerant rape seed commodities (Table 34). The LOD was 0.05 mg/kg for each analyte in each sample.

Table 34. Recoveries by method AE-24A from transgenic rape seed commodities (Niedzwiadek and Bertrand 1995b).

Commodity	Glufo	osinate	М	PP	NAG		
	spiking levels, Recovery range,		spiking levels, Recovery range,		spiking levels,	Recovery range,	
	mg/kg	%	mg/kg	%	mg/kg	%	
Rape shoot	0.05 0.50 10	92-114 (n=15)	0.05 0.50 10	76-115 (n=30)	0.05 0.50 10	91-119 (n=15)	
Rape pod	0.05 0.50 5.0	87-108 (n=15)	0.05 0.50 5.0	71-109 (n=30)	0.05 0.50 5.0	88-120 (n=15	
Rape straw	0.05 0.50	95-116 (n=10)	0.05 0.50	71-108 (n=20)	0.05 0.50	101-134 (n=10)	
Rape seed	0.05 0.50	76-109 (n=10)	0.05 0.50	82-117 (n=20)	0.05 0.50	99-118 (n=10)	

Idstein *et al.* (1987b) extracted glufosinate and MPP from soya beans and derivatized the compounds with trimethyl orthoacetate. After clean-up on a silica gel cartridge, the residues were measured by GLC with a flame photometric detector. Idstein *et al.* (1987a extracted glufosinate and MPP from milk by dialysis and completed the analysis in the same way. Schuld (1988) extracted the residues from hens' eggs with water, washed the aqueous extract with dichloromethane and hexane to remove lipids and then followed a similar procedure.

Sochor *et al.* (1987a) extracted rape seed with water and removed oil with dichloromethane before proceeding with the Idstein method. Sochor *et al.* (1987b) extracted the residues from fats by an initial water/dichloromethane partition before following the remainder of the procedure. Sochor *et al.* (1988) extracted animal organs with water, precipitated high MW co-extractives with acetone and continued with the Idstein method, achieving practical LODs of 0.05-0.1 mg/kg.

The method used for glufosinate in The Netherlands relies on extraction of samples with water and clean-up by dialysis and ion-exchange. The residue is measured by GLC with an FPD after derivatization with trimethyl orthoacetate. Recoveries are 70-80% and the LOD 0.05 mg/kg.

Stability of pesticide residues in stored analytical samples

Information was made available on the frozen storage stability of glufosinate and its metabolites in genetically modified maize and processed commodities, soya bean and processed commodities, dairy

cow tissues and milk, eggs and chicken tissues, susceptible maize grain, and transgenic rape seed and sugar beet root.

Belcher (1995a) determined the stability of glufosinate and its metabolites in genetically modified maize commodities (hulls, grits, flour and refined maize oil) during frozen storage for 12 months (Table 35). The report did not state the storage conditions. Residues were generally stable for 12 months, but glufosinate decreased by 30% and NAG by 19% in refined maize oil.

Table 35. Recoveries of glufosinate, NAG and MPP from maize commodities spiked with 0.5 mg/kg and stored for periods up to 12 months under frozen conditions (Belcher 1995a). Results are means of duplicate samples, uncorrected for analytical recovery.

Storage,	glufos	glufosinate ammonium, racemate				NAG, racemate, free acid				MPP, free acid			
months	hulls	grits	flour	oil,	hulls	grits	flour	oil,	hulls	Grits	flour	oil,	
				refined				refined				refined	
0	88%	92%	94%	83%	80%	101%	87%	102%	82%	93%	73%	89%	
3	81%	84%	71%	67%	91%	89%	82%	94%	92%	81%	86%	84%	
6	76%	74%	83%	68%	90%	93%	93%	85%	87%	86%	85%	79%	
12	87%	88%	103%	58%	81%	93%	108%	83%	77%	81%	75%	76%	

Belcher (1996b) determined the stability of glufosinate and its metabolites in genetically modified maize grain, fodder and forage during frozen storage for 24 months (Table 36). Residues were generally stable. The report did not include the storage conditions.

Table 36. Recoveries of glufosinate, NAG and MPP from maize commodities spiked with 0.5 mg/kg and stored for periods up to 24 months under frozen conditions (Belcher 1996b). Results are means of duplicate samples, uncorrected for analytical recovery.

Storage, months	glufos	inate amm	onium,	NAG, r	acemate, f	free acid	MPP, free acid			
		racemate								
	grain fodder forage			grain	fodder	forage	grain	fodder	forage	
0	95%	87%	94%	99%	88%	96%	88%	91%	98%	
3	89%	97%	91%	93%	89%	90%	86%	99%	97%	
6	91%	91%	83%	109%	86%	100%	90%	91%	93%	
12	83%	85%	76%	93%	95%	88%	94%	103%	105%	
24	88%	79%	76%	101%	88%	78%	90%	89%	87%	

Belcher (1995b) determined the stability of glufosinate and its metabolites in genetically modified soya bean meal, hulls and refined oil during frozen storage for 12 months (Table 37). In some samples of meal and hulls enzyme activity converted some of the added glufosinate to NAG. In these cases the sum of glufosinate and NAG was treated as the remaining residue and used for the calculation in Table 37. Residues were generally stable for the 12 months. The report did not specify the storage conditions.

Table 37. Recoveries of glufosinate, NAG and MPP from soya commodities spiked with 0.5 mg/kg and stored for periods up to 12 months under frozen conditions (Belcher 1995b). Results are means of duplicate samples, uncorrected for analytical recovery.

Storage,	glufosina	te ammoniu	ım, racemate	NAG,	racemate,	free acid	MPP, free acid			
months	meal	hulls	oil, refined	meal	hulls	oil, refined	meal	hulls	oil, refined	
0	68%	83%	79%	84%	91%	104%	113%	90%	87%	
3	86%	86%	77%	92%	82%	86%	101%	96%	96%	
6	93%	96%	78%	100%	104%	97%	90%	93%	82%	
12	83%	94%	93%	78%	96%	103%	93%	95%	87%	

Homogenized eggs, chicken tissue, milk and cow tissue samples were spiked with glufosinateammonium, NAG and MPP at 0.25 mg/kg and stored at -12°C to -27°C for 15 months (Crotts and McKinney, 1995, 1996; McKinney and Crotts, 1997). The results are shown in Table 38 and 39.All three compounds were stable for 15 months.

Table 38. Recoveries of glufosinate, NAG and MPP from homogenized eggs and chicken tissues spiked with 0.25 mg/kg and stored for periods up to 15 months under frozen conditions (Crotts and McKinney, 1995; McKinney and Crotts, 1997). Results are means of duplicate samples, uncorrected for analytical recovery.

Storage,	Glufo	Glufosinate ammonium, racemate			L-isomer, o	disodium	MPP, free acid		
monuis	muscle Liver eggs						musala liver aggs		
	muscie	LIVEI	eggs	muscie	IIVEI	eggs	muscie	IIVEI	eggs
0	102%	106%	105%	103%	111%	102%	96%	88%	85%
1	104%	102%	109%	103%	102%	101%	89%	92%	82%
3	79%	81%	83%	76%	91%	81%	86%	87%	79%
15	102%	105%	100%	108%	108%	86%	90%	84%	87%

Table 39. Recoveries of glufosinate, NAG and MPP from milk and cow tissues spiked with 0.25 mg/kg and stored for periods up to 15 months under frozen conditions (Crotts and McKinney, 1996). Results are means of duplicate samples, uncorrected for analytical recovery.

Storage,	glufosinate ammonium, racemate				NAG, L-isomer, disodium				MPP, free acid			
months	kidneys	muscle	liver	whole	kidneys	muscle	liver	whole	kidneys muscle liver			whole
	-			milk	-			milk	-			milk
14	107			86	112			83	100			82
15		76	97			93	107			81	92	

Werner (1997a-d) spiked homogenized transgenic maize shoots, rape seed and sugar beet roots and susceptible maize grain with glufosinate-ammonium, NAG and MPP at 0.5 mg/kg (expressed as glufosinate free acid) and measured the stability of the residues stored deep-frozen (temperature not stated) for 24 months. Results are shown in Table. All three compounds were stable for 24 months.

Table 40. Recoveries of glufosinate, NAG and MPP from homogenized transgenic maize shoots, susceptible maize grain, transgenic rape seed and sugar beet root stored for periods up to 24 months under frozen conditions (Werner 1997a-d). Results are means of duplicate samples, uncorrected for analytical recovery.

Storage,	glufosi	nate amm	onium, rac	cemate, %	NAG,	L-isomer	, disodi	ium, %	MPP, fre	e acid, % 1	remaining	
months	remaini	ing			remaining	5						
	maize shoot	maize grain	rape- seed	sugar- beet	maize shoot	maize grain	rape- seed	sugar- beet	maize shoot	maize grain	rape- seed	sugar- beet
0	76	81	96	74	75	59	100	64	65	68	87	67
1		92		74		66		72		74		71
2	67		84		75		91		73		91	
3		80		67		83		67		61		63
4	75		85		60		97		51		90	
6				86	106 ¹	68^{1}	85 ¹	79	108	78	71	80
12			103	94	100^{1}	63 ¹	88	96	96	73	81	77
15	87	100			68	58			57	84		
18	92	106	92	98	77	89	91	103	77	68	86	86
23			101				97				90	
23							104 ¹				106	
24					91 ¹	91 ¹			86	88		
24	83	113		102	88	97		81	72	86		82

¹Analyses by method AE-24A. % remaining is calculated as sum of glufosinate + NAG.

Definition of the residue

The current definition of the residue for glufosinate-ammonium is *Sum of glufosinate-ammonium* and 3-[hydroxy(methyl)phosphinoyl]propionic acid, expressed as glufosinate (free acid).

When glufosinate is used on glufosinate-tolerant crops a major part of the residue is *N*-acetyl-glufosinate (NAG). It should be included in the definition of the residue for enforcement because it is generally the main component of the residue and the same GLC derivative is produced in the analytical method from glufosinate and NAG, so unless the compounds are separated before derivatization they both appear in the GLC peak for their common derivative.

A suitable revised definition would be *Sum of glufosinate-ammonium, 3-*[hydroxy(methyl)phosphinoyl]propionic acid and N-acetyl-glufosinate, expressed as glufosinate (free acid), but this definition could not be adopted until N-acetyl-glufosinate had been toxicologically evaluated.

USE PATTERN

Glufosinate-ammonium is a non-selective herbicide registered for crop uses, including desiccation, and non-crop uses. It is used to control annual and perennial grasses and broad-leaved weeds in horticultural and agricultural crops. Glufosinate acts as an inhibitor of glutamine synthetase, which leads to poisoning of the plant by ammonia. For weed control in susceptible crops it must be used either before crop emergence or as a directed spray away from foliage.

Glufosinate resistance has been imparted to several agronomic crops by insertion of a gene that enables the plant to detoxify L-glufosinate (the active isomer) by acetylation to *N*-acetyl-L-glufosinate which is not herbicidal.

The use pattern is necessarily different for conventional crops and those with glufosinate tolerance.

Table 41. Registered uses of glufosinate-ammonium.	Concentrations	and rates	are expressed	in terms
of the active ingredient glufosinate-ammonium.				

Almond USA 120 g1 SL directed Rate, kg aihn Spray conc, kg aihn stage 2 Almond USA 120 g1 SL directed 1.7 min spray vol 187 1/ha max total 5.1 kg/ha/yr 14 Apple Poland 3 150 g1 SL 0.60-12 0.004-035 2 - Apricau Germany 3 200 g1 SL spray 0.55 0.14-0.18 1 - Avocado Australia 200 g1 SL directed 0.2-1.0 spray vol 300-1000 1/ha - - Banana Australia 200 g1 SL spray HU 0.45-0.60 0.04-0.30 1 pre-emerg - Banana Malaysia 150 g1 SL foirerted 0.30-100 1ha - - - Banana Malaysia 150 g1 SL foirerted 0.30-50 min spray vol 100 1/ha - - - - - - - - - - - - - - - - - - </th <th>Crop¹</th> <th>Country</th> <th>Form</th> <th></th> <th>Applicatio</th> <th>n</th> <th>Number</th> <th>Growth</th> <th>PHI, days</th>	Crop ¹	Country	Form		Applicatio	n	Number	Growth	PHI, days
				Method	Rate, kg ai/ha	Spray conc. kg ai/hl		stage ²	
$ \begin{array}{ c c c c c c c c c c c c c c c c c c c$	Almond	USA	120 g/l SL	directed	1.7	min spray vol 187 l/ha	max total 5.1 kg/ha/yr		14
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	Apple	Poland ⁴	150 g/l SL		0.45-0.90	0.04-0.35	2		
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	Apricot	Poland ⁴	200 g/l SL		0.60-1.2	0.06-0.60	2		
Avocado Australia 200 g/l SL directed $0.2-1.0$ spray vol 300-1000 l/na spray vol 300-1000 l/na Banana Makysia 150 g/l SL directed $0.2-1.0$ spray vol 300-1000 l/na Banana Makysia 150 g/l SL directed $0.3-5$ 0.11 4 Backberries Netherlands 200 g/l directed $0.75-1.0$ <td>Asparagus</td> <td>Germany⁴</td> <td>200 g/l SL</td> <td>spray</td> <td>0.55</td> <td>0.14-0.18</td> <td>1</td> <td></td> <td></td>	Asparagus	Germany ⁴	200 g/l SL	spray	0.55	0.14-0.18	1		
Banana Australia 200 g/l SL directed $0.2-1.0$ spray vol 300-1000 l/ha spray vol 300-1000 l/ha - Banana Malaysia 150 g/l SL directed $0.3-0.5$ 0.11 4 14 Banana Poland ² SL spray HV $0.45-0.60$ $0.04-0.30$ 1 pre-emerg - Canola Canola 150 g/l SL folar $0.75-1.0$ 2 early bolting 2 Canola Canada 150 g/l SL folar 0.60 min spray vol 110 l/ha 2 early bolting 2 Caronta Canada 150 g/l SL folar 0.60 min spray vol 110 l/ha 200 g/l SL spray 0.55 $0.114-0.18$ 1 pre-emerg Carrot Germany ⁴ 200 g/l SL spray WIV $0.45-0.60$ $0.04-0.35$ 2 $-$ Carrot Germany ⁴ 200 g/l SL spray wol screening $ -$ Carrot Germany ⁴ <td< td=""><td>Avocado</td><td>Australia</td><td>200 g/l SL</td><td>directed</td><td>0.2-1.0</td><td>spray vol 300-1000 l/ha</td><td></td><td></td><td>-</td></td<>	Avocado	Australia	200 g/l SL	directed	0.2-1.0	spray vol 300-1000 l/ha			-
$\begin{array}{c c c c c c c c c c c c c c c c c c c $	Banana	Australia	200 g/l SL	directed	0.2-1.0	spray vol 300-1000 l/ha			-
	Banana	Malaysia	150 g/l SL	directed	0.3-0.5	0.11	4		14
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	Bean	Poland ⁴	SL	spray HV	0.45-0.60	0.04-0.30	1	pre-emerg	
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	Blackberries	Netherlands	200 g/l 150 g/l	directed spray	0.75-1.0		2		-
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	Canola (tolerant)	Canada	150 g/l SL	foliar	0.30-0.50	min spray vol 110 l/ha	2	early bolting	<u>3</u>
	Canola (tolerant)	Canada	150 g/l SL	foliar	0.60	min spray vol 110 l/ha	1	early bolting	<u>3</u>
$ \begin{array}{c ccccccccccccccccccccccccccccccccccc$	Carambola (Starfruit)	Malaysia	150 g/l SL	directed	0.3-0.5	0.11	4		14
$ \begin{array}{c ccccccccccccccccccccccccccccccccccc$	Carrot	Germany ⁴	200 g/l SL	spray	0.55	0.14-0.18	1	pre-emerg	
$ \begin{array}{c ccccccccccccccccccccccccccccccccccc$	Carrot	Poland ⁴	SL	spray HV	0.45-0.60	0.04-0.30	1	pre-emerg	
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	Cashew nut	Malaysia	150 g/l SL	directed	0.3-0.5	0.11	4	· · · ·	14
	Cherry	Poland ⁴	150 g/l SL		0.45-0.90	0.04-0.35	2		
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	Currants	Germany ⁴	200 g/l SL	spray with screening	0.92	0.15-0.31	1		14
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	Currants	Netherlands	200 g/l 150 g/l	directed spray	0.75-1.0		2		-
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	Durian	Malaysia	150 g/l SL	directed	0.3-0.5	0.11	4		14
$ \begin{array}{c ccccccccccccccccccccccccccccccccccc$	Dwarf French beans	Germany ⁴	200 g/l SL	spray between rows with screening	0.92	0.23-0.31	1		14
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	Dwarf French beans	Germany ⁴	200 g/l SL	spraying	0.46	0.08-0.15	1	desiccation	14
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	Feijoa	Australia	200 g/l SL	directed	0.2-1.0	spray vol 300-1000 l/ha			-
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	Field peas	Germany ⁴	200 g/l SL	spraying	0.46	0.08-0.15	1	desiccation	14
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	Fruit trees	Netherlands	200 g/l 150 g/l	directed under trees	0.75-1.0		2		-
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	Gooseberrie s	Germany ⁴	200 g/l SL	spray with screening	0.92	0.15-0.31	1		14
GuavaAustralia 200 g/l SL directed $0.2-1.0$ spray vol $300-1000 \text{ l/ha}$ $-$ GuavaMalaysia 150 g/l SL directed $0.3-0.5$ 0.11 414HazelnutItaly 120 g/l SL directed $0.5-1.6$ $0.08-0.78$ max total 2.5 kg/ha/yr -Horse beansGermany $\frac{4}{2}$ 200 g/l SL spraying 0.46 $0.08-0.15$ 1desiccation14	Gooseberrie s	Poland ⁴	200 g/l SL		0.60-1.2	0.06-0.60	2		
GuavaMalaysia150 g/l SLdirected $0.3-0.5$ 0.11 414HazelnutItaly120 g/l SLdirected $0.5-1.6$ $0.08-0.78$ max total 2.5 kg/ha/yr -Horse beansGermany $\frac{4}{2}$ 200 g/l SLspraying 0.46 $0.08-0.15$ 1desiccation14	Guava	Australia	200 g/l SL	directed	0.2-1.0	spray vol 300-1000 l/ha			-
HazelnutItaly120 g/l SLdirected $0.5-1.6$ $0.08-0.78$ max total 2.5 kg/ha/yr -Horse beansGermany 4 200 g/l SLspraying 0.46 $0.08-0.15$ 1desiccation14	Guava	Malaysia	150 g/l SL	directed	0.3-0.5	0.11	4		14
Horse beansGermany 4200 g/l SLspraying0.460.08-0.151desiccation14	Hazelnut	Italy	120 g/l SL	directed	0.5-1.6	0.08-0.78	max total 2.5 kg/ha/yr		-
	Horse beans	Germany ⁴	200 g/l SL	spraying	0.46	0.08-0.15	1	desiccation	14

Crop ¹	Country	Form		Application	n	Number	Growth	PHI, days
_			Method	Rate, kg ai/ha	Spray conc. kg ai/hl		stage ²	
Jack fruit	Malaysia	150 g/l SL	directed	0.3-0.5	0.11	4		14
Kiwifruit	Australia	200 g/l SL	directed	0.2-1.0	spray vol 300-1000 l/ha			-
Lamb's lettuce	Germany ⁴	200 g/l SL	spray	0.55	0.14-0.18	1	pre-emerg	
Leek	Germany ⁴	200 g/l SL	spray	0.55	0.14-0.18	1	pre-emerg	
Litchi	Australia	200 g/l SL	directed	0.2-1.0	spray vol 300-1000 l/ha			-
Macadamia	USA	120 g/l SL	directed	1.7	min spray vol 187 l/ha	max total 5.1 kg/ha/yr		14
Maize	Germany ⁴	200 g/l SL	spray	0.92	0.23-0.31	1	pre-sowing	
Maize	Germany ⁴	200 g/l SL	spray, under leaf	0.92	0.23-0.31	1	stage 16-20	
Maize (tolerant)	Germany ⁵	200 g/l SL	foliar	0.90	0.23-0.45	1	3-8 leaf	
Maize (tolerant)	Germany ⁵	200 g/l SL	foliar	0.45	0.11-0.23	2 ¹	8 leaf	
Maize (tolerant)	Canada	200 g/l SL	foliar	0.30-0.50	min spray vol 110 l/ha	2^{2}	8-leaf	86 grain 20 (graze)
Maize (tolerant)	Portugal	200 g/l SL	foliar	0.40-0.80	rec spray vol 200-400 l/ha	2	2-4 leaf ³	
Maize (tolerant)	USA	18.2%		0.20-0.36		2	60 cm	60 (corn forage) 70 (corn grain, fodder)
Maize (tolerant)	USA	200 g/l SL	foliar	0.23-0.41		2	60 cm	70 grain 70 maize fodder 60 maize forage
Mango	Australia	200 g/l SL	directed	0.2-1.0	spray vol 300-1000 l/ha			-
Mango	Malaysia	150 g/l SL	directed	0.3-0.5	0.11	4		14
Onion	Germany ⁴	200 g/l SL	spray	0.55	0.14-0.18	1	pre-emerg	
Onion	Poland ⁴	SL	spray HV	0.45-0.60	0.04-0.30	1	pre-emerg	
Рарауа	Australia	200 g/l SL	directed	0.2-1.0	spray vol 300-1000 l/ha			-
Parsley	Poland ⁴	SL	spray HV	0.45-0.60	0.04-0.30	1	pre-emerg	
Passion fruit	Australia	200 g/l SL	directed	0.2-1.0	spray vol 300-1000 l/ha			-
Peach	Poland ⁴	200 g/l SL		0.60-1.2	0.06-0.60	2		
Pear	Poland [±]	150 g/l SL	11 . 1	0.45-0.90	0.04-0.35	2		1.4
Pecan	USA	120 g/l SL	directed	1.7	min spray vol 187 l/ha	max total 5.1 kg/ha/yr		14
Pineapple	Australia	200 g/l SL	directed	0.2-1.0	spray vol 300-1000 l/ha			-
Plum	Poland ⁴	200 g/l SL		0.60-1.2	0.06-0.60	2		
Potato	Germany $\frac{4}{4}$	200 g/l SL	spray	0.55	0.14-0.18	1	stage 8-9	1.4
Potato	Germany ⁻	200 g/I SL	spraying	0.46	0.08-0.15	1	desiccation	14
Potato	Netherlands	200 g/l 150 g/l	aerial parts, haulm kill	0.45-0.60				-
Potato	Netherlands	200 g/l 150 g/l	directed spray	0.45-1.0		1		-
Potato	Netherlands	200 g/l 150 g/l	pre- emergence	0.45-1.0		1		-

¹ Germany, split application on maize, 6 weeks interval.

² Maximum rate in the 2nd application is 0.40 kg ai/ha.

³ For the second application, the maize should have no more than 8-10 leaves.

Crop ¹	Country	Form	Application			Number	Growth	PHI, days
			Method	Rate, kg	Spray conc. kg		stage ²	
				ai/ha	ai/hl			
Rambutan	Australia	200 g/l SL	directed	0.2-1.0	spray vol			-
					300-1000 l/ha			
Rape seed	Poland ⁴	SL	spray HV	0.38-0.50	0.04	0.25	desiccation	
Raspberries	Germany ⁴	200 g/l SL	spray with	0.92	0.15-0.31	1		14
			screening					
Raspberries	Netherlands	200 g/l	directed	0.75-1.0		2		-
		150 g/l	spray					
Raspberries	Poland ⁴	200 g/l SL		0.60-1.2	0.06-0.60	2		
Spinach	Netherlands	200 g/l	aerial parts,	0.6-0.8		1		-
		150 g/l	haulm kill					
Soya beans	USA	200 g/l SL	foliar	0.23-0.41		2	bloom	70 seed
(tolerant)							growth	forage ¹
Strawberries	Germany ⁴	200 g/l SL	spray	0.73	0.18-0.24	1	stage 59	42
			between rows					
			with					
	4		screening					
Strawberries	Germany ⁴	200 g/l SL	spray with	0.73	0.12-0.24	1	after harvest	
	4		screening					
Sugar beets	Germany ⁴	200 g/l SL	spray	0.92	0.23-0.31	1	pre-sowing	
Sunflowers	Germany ⁴	200 g/l SL	spraying	0.46	0.08-0.15	1	desiccation	14
Tree nuts	Australia	200 g/l SL	directed	0.2-1.0	spray vol			-
					300-1000 l/ha			
Walnut	Italy	120 g/l SL	directed	0.5-1.6	0.08-0.78	max total		-
						2.5 kg/ha/yr		
Walnut	USA	120 g/l SL	directed	1.7	min spray vol	max total		14
					187 l/ha	5.1 kg/ha/yr		
Winter rape	Germany $\frac{4}{2}$	200 g/l SL	spraying	0.46	0.08-0.15	1	desiccation	14

¹ tolerant: tolerant to glufosinate-ammonium
 ² growth stage at final application.
 ³ Do not graze the treated crop or cut for hay; sufficient data are not available to support such use.
 ⁴ Label not provided.
 ⁵ Label not available. Registration document 4574-00 1 Sept 1998 provided.

RESIDUES RESULTING FROM SUPERVISED TRIALS

Residue data from glufosinate-ammonium supervised residue trials on fruit, tree nuts and field crops are summarized in Tables 42-61.

Table 42	Tropical fruits -avocado, mango, guava, carambola, papaya. Australia,
	Malaysia.
Table 43	Tree nuts – pecan, walnut, almond, hazel nuts macadamia. USA, Italy,
	Australia.
Table 44	Maize. France, Germany, Italy, Spain.
Table 45	Maize. USA, Argentina.
Table 46	Maize. Canada.
Table 47	Soya beans. USA.
Table 48	Rape seed. France, Germany, UK.
Table 49	Canola. Canada.
Table 50	Canola. Australia.
Table 51	Sugar beet root. France, Germany, UK.
Table 52	Sugar beet root. USA.

¹ Do not feed treated green immature growing soya bean plants to livestock.

Table 53	Maize forage and fodder. France, Germany, Italy, Netherlands, Spain.
Table 54	Maize forage and fodder. USA, Argentina.
Table 55	Maize forage and fodder. Canada.
Table 56	Rape seed forage and fodder. France, Germany, UK.
Table 57	Canola forage and fodder. Australia.
Table 58	Soya bean forage and fodder. USA.
Table 59	Sugar beet tops and leaves. France, Germany, UK.
Table 60	Sugar beet tops. USA.
Table 61	Almond hulls. USA.

Where residues were not detected they are recorded as below the limit of determination (LOD), e.g. <0.05 mg/kg. Residue levels of glufosinate and its metabolites are generally expressed as glufosinate free acid. Residues, application rates and spray concentrations have generally been rounded to 2 significant figures or, for residues near the LOD, to 1 significant figure. Although all trials included control plots residues in control samples are recorded only when they exceeded the LOD. Residues are not corrected for recoveries. Recoveries were mainly in the range 70-120%.

Reports did not generally state whether residues in forage and fodder commodities were expressed on a fresh-weight or dry-weight basis.

Trials were fully reported as well as on summary sheets except the US trials on pecans and walnuts and the Italian trials on hazel-nuts. These trials were all reported on the detailed summary sheets used in Germany.

Glufosinate-ammonium is used as a directed spray for weed control in orchards, so that residues usually only occur in the crop through root uptake from the soil. Glufosinate-ammonium itself is not taken up, but the metabolite 3-methylphosphinicopropionic acid (MPP) can be taken up and translocated by the crop. This metabolite is not herbicidally active, so its residues may occur in the crop when glufosinate-ammonium is used as a directed spray.

In tropical fruit trials in Australia and Malaysia glufosinate-ammonium was applied to weeds around the trees as a directed spray. Plot sizes ranged from 1 tree to 1050 m^2 . Intervals of 109 to 468 days of frozen sample storage elapsed before analysis. Residues are expressed on a pulp + peel basis for the avocados and mangoes in the Australian trials, but this does not influence the results which were mainly below the LOD.

Glufosinate-ammonium was used in tree nut trials in the USA with plot sizes ranging from 1 to 9 trees. The method of spraying was not clearly expressed, but was presumably a directed spray. Samples were held in frozen storage for intervals of 400 to 700 days before analysis. The method of application was also not stated for the hazel nut trials in Italy. Samples were held 240 days in frozen storage before analysis. Directed sprays were used in the macadamia trials in Australia. Samples from the year 1992 were analysed 28 days after harvest while those from 1995 were stored for almost a year.

In Canadian trials on resistant maize in 1993-95 glufosinate-ammonium was applied by ground rigs. Two different SL formulations were used, with applications at two rates and at different growth stages.

Glufosinate-ammonium was applied to resistant maize in an extensive set of trials in France from 1993 to 1996. Plot sprayers were used on plots of 11 to 79 m². Harvested samples were held in frozen storage for 520-670 days (1993), 310-340 days (1994), 160-230 days (1995) and 50-90 days (1996).

In a similar series of resistant maize trials in Germany from 1994 to 1996, plot sprayers were used on plots of $48-64 \text{ m}^2$. Frozen storage periods were 300-400 days (1994), 150 days (1995) and 30-70 days (1996).

Further European trials on resistant maize were conducted in 1994-96 in Italy, The Netherlands and Spain. Hand-carried plot sprayers were used in Italy and Spain on plots of 40-60 m². Information on the sprayers and plot sizes was not available for The Netherlands. Analytical samples were stored frozen for 330-340 days (1994) and 50-150 days (1995-96).

Glufosinate-ammonium was extensively tested on resistant maize in US trials in 1993-1995. In 1993 it was applied by portable and tractor-mounted plot sprayers to plots of 28-186 m². In 1995 ground rigs and plot sprayers were used on plots of 110-740 m². Harvested samples were held in frozen storage for 280-330 days (1993) and 190-280 days (1995) before analysis. Some of the higher rate treatments in trial ER-93-USA-01 (Virginia and South Dakota) suffered phytotoxicity so grain samples were limited or unavailable for analysis.

Resistant soya beans were determined with glufosinate-ammonium in trials in many States of the USA in 1994-1996. Plots of 70-95 m² were sprayed with CO_2 pressurised backpack or plot sprayers in 1994. In 1995 tractor-mounted ground rigs and backpack or bicycle sprayers were used on plots of 40-130 m² and in 1996 backpack or plot sprayers were used on 90-280 m² plots. Harvested samples were held in frozen storage before analysis for 200-280 days (1994), 100-300 days (1995) and 200-230 days (1996).

Glufosinate-ammonium was determined on resistant rape seed in a supervised residue trial programme in France in 1993, 1994 and 1996. Plot sprayers were used on plots of 18-72 m². Periods of frozen storage of samples before analysis were 570-640 days (1993), 380-550 days (1994) and 112 days (1996). In trials on resistant rape seed in Germany (1994 and 1996) and the UK (1993 and 1996) glufosinate-ammonium was applied by plot sprayers to plots of 48-100 m² in Germany and 20-80 m² in the UK. Frozen storage periods for samples were 1 year and 2 months (1994 and 1996 respectively) in Germany and 600 days and 2 months (1993 and 1995 respectively) in the UK.

Glufosinate-ammonium was applied with boom sprayers to plots of $30-42 \text{ m}^2$ of resistant canola in Australian trials in 1996. Harvested samples were stored for 170-240 days before analysis.

Resistant canola was treated with glufosinate-ammonium in a series of supervised residue trials in Canada in 1993 and 1994. CO_2 powered backpack, plot sprayers and tractor-mounted rigs were used on plots of 24-80 m². Samples were held for 170-280 days (1993) and 80 days (1994) in frozen storage before analysis. No field reports were available for trials in report A53770 (Bertrand, 1993). One canola trial (A57514, MacDonald, 1996d) was not evaluated because the data sheets were from a barley trial. It was not clear in some of the canola trials whether the residues were expressed as glufosinate free acid or as the metabolites,.

Glufosinate-ammonium was applied with plot sprayers to plots of $38-76 \text{ m}^2$ of resistant sugar beet in French trials in 1995 and 1996. Harvested samples were stored for 120-300 days under frozen conditions before analysis. In similar trials in the UK in 1996-97 glufosinate-ammonium was applied by plot sprayer or knapsack to $48-64 \text{ m}^2$ plots. Frozen samples were stored for 150-280 days before analysis.

In a US trial programme on resistant sugar beet in 1995 and 1996 glufosinate-ammonium was applied by ground rig, backpack or bicycle sprayer to plots of 17-300 m². Storage periods for frozen samples were 240-320 days (1995) and 130-170 days (1996).

Table 42. Glufosinate residues in tropical fruits resulting from supervised trials in Australia and Malaysia. In the Australian trials on avocados and mangoes residues are expressed on a pulp + peel basis. Residues in samples from replicate plots are shown separately. Double-underlined residues are from treatments according to GAP and were used to estimate maximum residue levels.

FRUIT,	Application				PHI,	Residues, mg/kg	Ref		
country, year	ļ				days	free a			
(variety)	Form	kg ai/ha	kg ai/hl	no.		glufosinate	MPP		
AVOCADO									
Australia	SL	1.0	0.4	1	7	< <u>0.1</u> (2)	< 0.1 (2)	A59027	
(Qld) 1991					12	< 0.1 (2)	< 0.1 (2)	QD 27/90	
(Sharwil)					21	< 0.1 (2)	< 0.1 (2)		
Australia	SL	2.0	0.8	1	7	< 0.1 (2)	< 0.1 (2)	A59027	
(Qld) 1991					12	< 0.1 (2)	< 0.1 (2)	QD 27/90	
(Sharwil)					21	< 0.1 (2)	< 0.1 (2)		
Australia	SL	0.6	0.33	2	0	< 0.05 (2)	< 0.05 (2)	A59026	
(Qld) 1996					14	< 0.05 (2)	< 0.05 (2)	AU QD 26	
(Fuerte)					24	< 0.05 (2)	< 0.05 (2)		
					28	< 0.05 (2)	< 0.05 (2)		
Australia	SL	1.2	0.67	2	0	< <u>0.05</u> (2)	< 0.05 (2)	A59026	
(Qld) 1996					14	< 0.05 (2)	< 0.05 (2)	AU QD 26	
(Fuerte)					24	< 0.05 (2)	< 0.05 (2)		
					28	< 0.05 (2)	< 0.05 (2)		
Australia	SL	0.6	0.35	2	0	< 0.05 (2)	< 0.05 (2)	A59025	
(Qld) 1995			0.30		14	< 0.05 (2)	< 0.05 (2)	AU QD 24	
(Haas)					24	< 0.05 (2)	< 0.05 (2)		
					31	< 0.05 (2)	< 0.05 (2)		
Australia	SL	1.2	0.69	2	0	<u>0.06</u> <0.05	< 0.05 (2)	A59025	
(Qld) 1995			0.60		14	< 0.05 (2)	< 0.05 (2)	AU QD 24	
(Haas)					24	< 0.05 (2)	< 0.05 (2)		
					31	< 0.05 (2)	< 0.05 (2)		
MANGO									
Australia	SL	1.0	0.4	1	7	< <u>0.1</u>	< 0.1	A59030	
(Qld) 1991					14	< 0.1	< 0.1	QD 21/90	
(Kensington					20	< 0.1	< 0.1		
Pride)									
Australia	SL	2.0	0.8	1	7	< 0.1	< 0.1	A59030	
(Qld) 1991					14	< 0.1	< 0.1	QD 21/90	
(Kensington					20	< 0.1	< 0.1		
Pride)									
Australia	SL	0.6	0.21	2	0	< 0.05	< 0.05	A59029	
(Qld) 1994-5					14	< 0.05	< 0.05	AU QD 12	
(Fascell)					21	< 0.05	< 0.05		
					28	< 0.05	< 0.05		
Australia	SL	1.2	0.41	2	0	< <u>0.05</u>	< 0.05	A59029	
(Qld) 1994-5					14	< 0.05	< 0.05	AU QD 12	
(Fascell)					21	< 0.05	< 0.05		
					28	< 0.05	< 0.05		
Australia	SL	0.6	0.2	2	0	< 0.05	< 0.05	A59028	
(Qld) 1995					14	< 0.05	< 0.05	AU QD 13-94	
(Kensington					21	< 0.05	< 0.05		
Pride)					28	< 0.05	< 0.05		
Australia	SL	1.2	0.4	2	0	< <u>0.05</u>	< 0.05	A59028	
(Qld) 1995					14	< 0.05	< 0.05	AU QD 13-94	
(Kensington					21	< 0.05	< 0.05		
Pride)		ļ	ļ		28	< 0.05	< 0.05		
GUAVA			ļ						
Malaysia	SL	0.50	0.11	4	0	< 0.05	< 0.05	A57294	
(Perat) 1995					3	< 0.05	< 0.05	ER95MYS8800201	
("with					7	< 0.05	< 0.05	02	
seeds")					14	< <u>0.05</u>	< 0.05		
					21	< 0.05	< 0.05		
$\begin{array}{ c $	FRUIT,		App	lication		PHI,	Residues, mg/kg	, as glufosinate	Ref
---	---------------	------	----------	----------	-----	------	-----------------	------------------	--------------------------
$ \begin{array}{ c c c c c c c c c c c c c c c c c c c$	country, year				1	days	free a	icid	
Malaysia (Perat) 1995 SL 1.0 0.22 4 0 <0.05 <0.05 <0.57244 ("with seeds") - - - - - 0.05 - 0.05 0.05 0.05 0.05 0.05 Malaysia (Perat) 1995 SL 0.50 0.11 4 0 <0.05	(variety)	Form	kg ai/ha	kg ai/hl	no.		glufosinate	MPP	
	Malaysia	SL	1.0	0.22	4	0	< 0.05	< 0.05	A57294
("with seeds") - - 7 <0.05	(Perat) 1995					3	< 0.05	< 0.05	ER95MYS8800201
seeds") - </td <td>("with</td> <td></td> <td></td> <td></td> <td></td> <td>7</td> <td>< 0.05</td> <td>< 0.05</td> <td>03</td>	("with					7	< 0.05	< 0.05	03
$ \begin{array}{ c c c c c c c c c c c c c c c c c c c$	seeds")					14	< 0.05	< 0.05	
$ \begin{array}{ c c c c c c c c c c c c c c c c c c c$						21	< 0.05	< 0.05	
$ \begin{array}{ c c c c c c c c c c c c c c c c c c c$	Malaysia	SL	0.50	0.11	4	0	< 0.05	< 0.05	A57294
	(Perat) 1995					3	< 0.05	< 0.05	ER95MYS8800202
$ \begin{array}{ c c c c c c c c c c c c c c c c c c c$	("seedless")					7	< 0.05	< 0.05	02
$ \begin{array}{ c c c c c c c c c c c c c c c c c c c$	` ´					14	< 0.05	< 0.05	
$ \begin{array}{ c c c c c c c c c c c c c c c c c c c$						21	< 0.05	< 0.05	
(Perat) 1995 ("seedless") Image: height of the second sec	Malaysia	SL	1.0	0.22	4	0	< 0.05	< 0.05	A57294
("seedless") Image: Second	(Perat) 1995		110	0	•	3	<0.05	<0.05	ER95MYS8800202
Controls of products of	("seedless")					7	<0.05	<0.05	03
CARAMBOLA Image: Construct of the second secon	(securess)					14	<0.05	<0.05	05
CARAMBOLA Image: Constraint of the second seco						21	<0.05	<0.05	
Malaysia (Selangor) SL (Selangor) 0.50 0.11 3 (Selangor) 0 4 0.05 A57295 Malaysia (Selangor) SL (Selangor) 0.11 3 (Selangor) 3 (Selangor) -0.05 <0.05	CARAMBOI	Δ				21	(0.05	(0.02	
	Malausia	-A	0.50	0.11	2	0	<0.05	<0.05	157205
(Setangor) 1995 (Bio) 10 1	(Salargar)	SL	0.30	0.11	5	2	< 0.05	< 0.03	AJ7293 ED05MVC0000101
$ \begin{array}{ c c c c c c c c c c c c c c c c c c c$	(Selangor)					5	< 0.05	< 0.03	EK95W1150000101
$ \begin{array}{ c c c c c c c c c c c c c c c c c c c$	1995 (B10)					14	< 0.05	< 0.05	02
$ \begin{array}{ c c c c c c c c c c c c c c c c c c c$						14	< <u>0.05</u>	< 0.05	
$ \begin{array}{ c c c c c c c c c c c c c c c c c c c$	N 1 1	CT.	1.0	0.00	2	21	<0.05	< 0.05	157005
$ \begin{array}{ c c c c c c c c c c c c c c c c c c c$	Malaysia	SL	1.0	0.22	3	0	<0.05	< 0.05	A57295
$\begin{array}{ c c c c c c c c c c c c c c c c c c c$	(Selangor)					3	<0.05	< 0.05	ER95MY S8800101
$\begin{array}{ c c c c c c c c c c c c c c c c c c c$	1995 (B10)					7	<0.05	< 0.05	03
$ \begin{array}{ c c c c c c c c c c c c c c c c c c c$						14	<0.05	< 0.05	
$ \begin{array}{ c c c c c c c c c c c c c c c c c c c$						21	<0.05	< 0.05	
$ \begin{array}{ c c c c c c c c c c c c c c c c c c c$	Malaysia	SL	0.50	0.11	3	0	< 0.05	< 0.05	A57295
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	(Perat) 1995					3	< 0.05	< 0.05	ER95MYS8800102
$ \begin{array}{ c c c c c c c c c c c c c c c c c c c$	(Bio)					7	< 0.05	< 0.05	02
$ \begin{array}{ c c c c c c c c c c c c c c c c c c c$						14	< <u>0.05</u>	< 0.05	
$ \begin{array}{c ccccccccccccccccccccccccccccccccccc$						21	< 0.05	< 0.05	
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	Malaysia	SL	1.0	0.22	3	0	< 0.05	< 0.05	A57295
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	(Perat) 1995					3	< 0.05	< 0.05	ER95MYS8800102
$ \begin{array}{ c c c c c c c c c c c c c c c c c c c$	(Bio)					7	< 0.05	< 0.05	03
$\begin{tabular}{ c c c c c c c c c c c c c c c c c c c$						14	< 0.05	< 0.05	
PAPAYA Image: Constraint of the system SL 0.6 0.26 2 0 <0.05 <0.05 A59285 (Qld) 1996 0.6 0.26 2 0 <0.05						21	< 0.05	< 0.05	
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	PAPAYA								
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	Australia	SL	0.6	0.26	2	0	< 0.05	< 0.05	A59285
(Rictor Gold) 21 <0.05 <0.05 Australia SL 1.2 0.52 2 0 <0.05	(Qld) 1996					14	< 0.05	< 0.05	AU QD 25
Australia SL 1.2 0.52 2 0 <0.05 <0.05 (Qld) 1996 0.52 2 0 <0.05	(Rictor Gold)					21	< 0.05	< 0.05	
Australia (Qld) 1996 (Rictor Gold) SL 1.2 0.52 2 0 < <u>0.05</u> <0.05 A59285 (Qld) 1996 (Rictor Gold) 1.2 0.52 2 0 < <u>0.05</u> <0.05						30	< 0.05	< 0.05	
(Qld) 1996 (Rictor Gold) 14 <0.05	Australia	SL	1.2	0.52	2	0	< 0.05	< 0.05	A59285
$(\text{Rictor Gold}) \begin{array}{ c c c c c c c c c c c c c c c c c c c$	(Old) 1996				_	14	< 0.05	< 0.05	AU OD 25
	(Rictor Gold)					21	< 0.05	< 0.05	
	(30	< 0.05	< 0.05	

NUT,		Applic	ation		PHI, days	Residues, mg/kg	Ref	
(variety)	Form	kg ai/ha	kg ai/hl	No.	uays	glufosinate	MPP	
PECAN	1 01111	ing un inu	ing turin	1101		gratosinate		
USA (GA)	SL	1.7		3	14	< <u>0.05</u> (3)	< 0.05 (3)	A48446
1985 (Witchitz)								16-GA-85-011
(Witchita)	CI	2.4		2	1.4	.0.05 (2)	-0.05 (2)	A 40446
USA (GA) 1085	SL	3.4		3	14	< <u>0.05</u> (3)	<0.05 (3)	A48440
(Witchita)								10-0A-03-011
	SI	17		3	21	<0.05 (3)	<0.05 (3)	A48446
1985 (Cape	SE	1.7		5	21	< <u>0.05</u> (5)	(0.05 (5)	16-LA-85-002
Fear)								
USA (LA)	SL	3.4		3	21	< <u>0.05</u> (3)	< 0.05 (3)	A48446
1985 (Cape								16-LA-85-002
Fear)								
USA (NM)	SL	1.7		3	14	< <u>0.05</u> (3)	<0.05 (3)	A48446
1985								07-NM-85-032
(Western								
USA (NM)	S1	2.4		2	14	<0.05 (3)	<0.05 (2)	1 19116
1985	SL	5.4		5	14	< <u>0.05</u> (3)	<0.05 (5)	07-NM-85-032
(Western								07 100 05 052
Schley)								
WALNUT								
USA (CA)	SL	1.6		2	14	< 0.05 (3)	< 0.05 (3)	A34230
1984 (English)		+1.2						
USA (CA)	SL	3.2		2	14	<0.05 (3)	<0.05 (3)	A34231
1984 (English)		+2.4						
USA (CA)	SL	1.6		3	14	<0.05 (3)	<0.05 (3)	A34232
1984 (English)	GT	+2×1.2		2	1.4	0.05 (2)	0.05 (2)	4.0.4000
USA (CA) 1084 (English)	SL	3.2		3	14	<0.05 (3)	<0.05 (3)	A34233
1964 (English)	C1	+2×2.4		2	14	<0.05 (2)	<0.05 (2)	125706
1985 (Serr)	SL	1./		5	14	< <u>0.03</u> (3)	<0.03 (3)	A55700
USA (CA)	SL	4.5		3	14	< 0.05 (3)	< 0.05 (3)	A35707
1985 (Serr)	52	1.5		5	11	(0.05 (5)	(0.05 (5)	100707
USA (CA)	SL	1.7		3	14	< <u>0.05</u> (3)	< 0.05 (3)	A35708
1985								
USA (CA)	SL	3.4		3	14	<0.05 (3)	< 0.05 (3)	A35709
1985								
USA (CA)	SL	1.7		3	11	< <u>0.05</u> (3)	< 0.05 (3)	A35710
1985 (Serr)	GT	2.4		2		0.05 (0)	0.05 (2)	4.05711
USA (CA) 1085 (Sarr)	SL	3.4		3	11	<0.05 (3)	<0.05 (3)	A35/11
	SI	17		3	14	<0.05 (3)	<0.05 (3)	A 18116
1985 (Serr)	SL	1./		5	14	< <u>0.05</u> (3)	<0.05 (5)	07-CA-85-019
USA (CA)	SL	3.4		3	14	< 0.05 (3)	< 0.05 (3)	A48446
1985 (Serr)						- (-)	- \- /	07-CA-85-019
USA (CA)	SL	1.7		3	14	< <u>0.05</u> (3)	< 0.05 (3)	A48446
1985 (Serr)								07-CA-85-024
USA (CA)	SL	3.4		3	14	<0.05 (3)	< 0.05 (3)	A48446
1985 (Serr)	~-							07-CA-85-024
USA (CA)	SL	1.7		3	11	< <u>0.05</u> (3)	< 0.05 (3)	A48446
1985 (Serr)								0/-CA-85-038

Table 43. Glufosinate residues in tree nuts resulting from supervised trials in Australia, Italy and the USA. Analyses on samples from replicate plots are shown separately. Double-underlined residues are from treatments according to GAP and were used to estimate maximum residue levels.

NUT,		Applic	ation		PHI,	Residues, mg/kg	, as glufosinate	Ref
country, year	-	1 . 4	1 . 4 1	N	days	free a	acid	
(variety)	Form	kg ai/ha	kg ai/hl	No.		glufosinate	MPP	
USA (CA)	SL	3.4		3	11	<0.05 (3)	<0.05 (3)	A48446
1985 (Serr)								07-CA-85-038
ALMOND	CI	17		2	15	.0.05 (2)	.0.05 (2)	A 4044C
USA(CA) 1085 (Cormol)	SL	1./		3	15	< <u>0.05</u> (3)	<0.05 (3)	A48440
1985 (Carmer)	CI	2.4		2	15	(0.05.(2))	(0.05.(2))	0/-CA-85-018
USA (CA) 1985 (Carmel)	SL	5.4		3	15	<0.05 (5)	<0.03 (3)	A40440 07 CA 85 018
USA (CA)	SI	17		3	15	<0.05 (3)	<0.05 (3)	07-CA-05-010
1985 (Special)	SL	1.7		5	15	< <u>0.05</u> (3)	<0.03(3)	07-CA-85-018
	SI	3.4		3	15	<0.05 (3)	<0.05 (3)	A48446
1985 (Special)	SL	5.4		5	15	<0.05 (5)	<0.05 (5)	07-CA-85-018
USA (CA)	SL	17		3	14	<0.05 (3)	0.07.0.07.0.07	A48446
1985 (Non-	51	1.,		5	11	(<u>0.05</u> (3)	<u>0.07</u> 0.07 0.07	07-CA-85-028
Peril)								
USA (CA)	SL	3.4		3	14	< 0.05 (3)	0.14 0.22 0.17	A48446
1985 (Non-								07-CA-85-028
Peril)								
USA (CA)	SL	1.7		3	14	< <u>0.05</u> (3)	<0.05 (3)	A48446
1985 (Non-								07-CA-85-037
Peril)								
USA (CA)	SL	3.4		3	14	<0.05 (3)	<0.05 (3)	A48446
1985 (Non-								07-CA-85-037
Peril)								
HAZEL-NUT	~~~			-				
Italy 1985	SL	1.5	0.38	2	19	< <u>0.05</u>	<0.05	A35935
(Gentile Del								
Piemonte)	CI	1.5	0.20	2	10	-0.05	-0.05	1 25026
(Necehione)	SL	1.5	0.38	2	19	< <u>0.05</u>	<0.05	A35936
(Noccilione)	C1	15	0.28	2	10	<0.05	<0.05	A 25027
(Gentile Del	SL	1.5	0.58	Z	19	< <u>0.05</u>	<0.03	A55957
Piemonte)								
Italy 1985	SL	15	0.38	2	19	<0.05	<0.05	A 35938
(Gentile Del	SE	1.5	0.50	2	17	< <u>0.05</u>	<0.05	1133730
Piemonte)								
Italy 1985	SL	1.5	0.38	2	19	< 0.05	< 0.05	A35938
(Gentile Del					-			
Piemonte)								
MACADAMIA I	NUT							
Australia 1992	SL	1.0	0.8	4	1	< <u>0.1</u> (2)	< 0.1 (2)	A59031
(var 434)					7	<0.1 (2)	< 0.1 (2)	QDI/92
					14	<0.1 (2)	< 0.1 (2)	92/1222
					21	< 0.1 (2)	< 0.1 (2)	
Australia 1992	SL	2.0	1.6	4	1	<0.1 (2)	<0.1 (2)	A59031
(var 434)					7	<0.1 (2)	<0.1 (2)	QDI/92
					14	<0.1 (2)	<0.1 (2)	92/1222
Australia 1005	CI	1.2	0.62	2	21	<0.1 (2)	<0.1 (2)	4 50022
Australia 1995 $(\text{var} 344)$	SL	1.2	0.03	2	0 16	< <u>0.05</u> (2)	< 0.05(2)	A39032 OD20-94
(vai 344)					10 27	< 0.03(2)	< 0.03(2)	05/3715
Australia 1005	SI	24	13	2	0	< 0.05(2)	< 0.05(2)	Δ59032
(var 344)	5L	2.7	1.5	2	16	< 0.05(2)	< 0.05(2)	OD20-94
(27	<0.05 (2)	<0.05 (2)	95/3715

Table 44. Glufosinate residues in maize resulting from supervised trials in France, Germany, Italy and Spain. Double-underlined residues are from treatments according to GAP and were used to estimate maximum residue levels.

Variety Forme 1 bg ai/h1 loc loc performante MPP NAG ² France 1993 SL 0.45 0.23 2 145 <0.05 <0.05 AS4225 France 1993 SL 0.45 0.23 2 145 <0.05 <0.05 FR33000402 France 1993 SL 0.45 0.23 2 145 <0.05 <0.05 FR331000102 France 1993 (MLC SL 0.45 0.23 2 147 <0.05 <0.05 FR331000102 France 1993 (MLC SL 0.45 0.23 2 147 <0.05 <0.05 FR34000020 France 1993 (MLC SL 0.45 0.23 2 145 <0.05 <0.05 FR34000020 France 1993 (MLC SL 0.45 0.23 2 123 <0.05 <0.05 FR34200 C1H2824 T25) France 1994 SL 0.45 0.23 2 120 <0.05 FR3420 FR34200 <	Country, Year,	Applica	ation			PHI days	Residues as mg/kg	glufosinate	free acid,	Ref
France 1993 SL 0.45 0.23 2 145 <0.05 0.05 A54225 (R93TCNS50 (R93TCNS50) France 1993 SL 0.45 0.23 2 145 <0.05	Variety	form	kg ai/ha	kg ai/hl	no ¹	2	glufosinate	MPP	NAG ²	
	France 1993	SL	0.45	0.23	2	145	< 0.05	< 0.05		A54225
stP2cT14) RA00002 France 1993 SL 0.45 0.23 2 145 <0.05	(LH163×LH164-				GS25					ER93ECN550
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	×F2×T14)									FRA000402
$\begin{array}{c c c c c c c c c c c c c c c c c c c $	France 1993	SL	0.45	0.23	2	145	< 0.05	< 0.05		A54225
$xP2 \ge T14$) x <t< td=""><td>(LH206×LH74-</td><td></td><td></td><td></td><td>GS24</td><td></td><td></td><td></td><td></td><td>ER93ECN550</td></t<>	(LH206×LH74-				GS24					ER93ECN550
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	\times F2 \times T14)									FRA000502
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	France 1993 (MLC	SL	0.45	0.23	2	128	< 0.05	< 0.05		A54225
$ \begin{array}{ c c c c c c c c c c c c c c c c c c c$	2101/T14)	22	0.10	0.20	GS24	120	10100	10102		ER93ECN550
Frame 1993 (MLC 2101/T14) SL 0.45 0.23 2 (S24 47 $c0.05$ <0.05 A54225 (FR0300202) France 1993 (MLC 2101/T14) SL 0.45 0.23 2 (S24 45° <0.05 <0.05 $A54225$ (FR03ECN550) France 1993 (MLC 2101/T14) SL 0.45 0.23 2 (S25 <0.05 <0.05 $A54229$ (FR000002) France 1994 (F6xM1) SL 0.45 0.23 2 (S25 <0.05 <0.05 $A54226$ (FR000002) France 1994 (F6xM1) SL 0.45 0.23 2 (G252 <0.05 <0.05 $A54226$ (FR000102) France 1994 (F6xM1) SL 0.45 0.23 2 (G252 <0.05 <0.05 $A54230$ (FR000102) France 1994 (F6xM1) SL 0.45 0.23 2 (G252 <0.05 <0.05 $A54230$ (FR0400102) France 1994 (F6xM1) SL 0.45 0.23 2 (G25 <0.05 <0.05 $A54425$ (FR04ECN550) France 1994 (F6xM1) SL										FRA000102
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	France 1993 (MLC	SL	0.45	0.23	2	147	< 0.05	< 0.05		A54225
Image 1993 (MLC) Prance 1993 (MLC) (ILT4)SL 0.45 0.23 CS22 2 GS23 145 CS22 0.05 0.05 0.05 CS23 $A54225$ ER93ECN550 FRA000302France 1994 (LH82(4)T25)s/ (- Prance 1994 (F6xM1))SL 0.45 0.23 CS23 2 CS25 123 CS25 0.05 0.05 $A54229$ ER94ECN550 FRA000102France 1994 (LH824/T25)AF4x- B73)SL 0.45 0.23 CS2 2 CS25 120 CS25 0.05 0.05 $A54226$ ER93ECN550 FRA000102France 1994 (C)LH824/T25)AF4x- B73)SL 0.45 0.15 CS25 2 CS25 120 CS25 0.05 0.05 $A54230$ ER94ECN550 FRA000102France 1994 (C)LH224/T25)sSL 0.45 0.23 CS25 2 CS18 0.05 0.05 $A54230$ FRA000102France 1994 (C)LH82-T25)SL 0.45 0.23 CS18 2 CS18 0.05 0.05 $A5425$ FRA000102France 1995 (C)LSL 0.45 0.23 CS18 2 CS18 0.05 0.05 $A56445$ ER95ECN550 FRA000103France 1995 (C2)LH82-T25)SL 0.45 0.23 CS18 2 CS18 0.05 0.05 $A56445$ ER95ECN550 FRA000103France 1995 (transgenic hybrid)SL 0.45 0.23 CS18 2 CS18 0.05 0.05 $A56445$ ER95ECN550 FRA000103France 1995 (transgenic hybrid)SL 0.65 0.27 CS18 2 CS18 0.05 <	2101/T14)				GS24	-				ER93ECN550
France 1993 (MLC) SL 0.45 0.23 2 CS24 145 <0.05 <0.05 $A54225$ ER93ECN550 FRA000302 France 1994 SL 0.45 0.23 2 CS25 2 CS25 <0.05 <0.05 $A54225$ ER93ECN550 FRA000302 France 1994 GS41 0.45 0.23 2 CS25 <0.05 <0.05 $A54225$ ER93ECS50 FRA000102 France 1994 SL 0.45 0.23 2 CS25 <0.05 <0.05 $A54226$ ER93ECS50 FRA000102 France 1994 SL 0.45 0.23 2 CS25 <0.05 <0.05 $A54220$ ER93ECN550 FRA000102 France 1994 SL 0.45 0.23 2 CS2 <0.05 <0.05 $A54229$ ER93ECN550 FRA000102 France 1995 SL 0.45 0.23 2 CS18 <0.05 <0.05 $A56445$ ER95ECN550 FRA000102 France 1995 SL 0.45 0.23 2 CS18 <0.05 <0.05 $A56445$ ER95ECN550 FRA000103 France 1995 SL 0.45 <	,									FRA000202
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	France 1993 (MLC	SL	0.45	0.23	2	145	< 0.05	< 0.05		A54225
Image 1994 (LH82(4)T25)sf (- 2)sK1298SL 0.45 0.23 2 GS2 123 <0.05 <0.05 <0.05 $A54229$ FRA000202France 1994 (F6xM1)SL 0.45 0.23 2 GS25 $2S2$ 119 <0.05 <0.05 $A54230$ FRA000102France 1994 (LH824)T25)AF4x- B730SL 0.45 0.15 2 GS25 $2S2$ 119 <0.05 <0.05 $A54230$ FRA000102France 1994 (HR24xT25)AF4x- B730SL 0.45 0.23 2 GS25 119 <0.05 <0.05 $A54230$ FRA000102France 1994 (SL12)SL 0.45 0.23 2 GS18 119 <0.05 <0.05 $A55429$ FRA000102France 1995 (F2xLH82-T25)SL 0.45 0.23 2 GS18 100 <0.05 <0.05 $A56445$ FRA000202France 1995 (F2xLH82-T25)SL 0.45 0.23 2 GS18 100 <0.05 <0.05 $A56445$ FRA000202France 1995 (F2xLH82-T25)SL 0.45 0.23 2 GS18 106 <0.05 <0.05 $A56445$ FRA000103France 1995 (transgenic hybrid)SL 0.45 0.23 2 GS18 106 <0.05 <0.05 $A56445$ FRA000103France 1995 (transgenic hybrid)SL 0.45 0.23 2 GS18 106 <0.05 <0.05 $A56445$ FRA000103France 1995 (transgenic hybrid)SL 0.45 0.23 2 GS1	2101/T14)				GS24					ER93ECN550
France 1994 (ILH82(4)T25)sf (- L)SH298 SL 0.45 0.23 2 GS25 I23 <0.05 <0.05 A54229 (SH294) France 1994 (F6×M1) SL 0.45 0.23 2 GS25 I19 <0.05	,									FRA000302
$\begin{array}{c c c c c c c c c c c c c c c c c c c $	France 1994	SL	0.45	0.23	2	123	< 0.05	< 0.05		A54229
2)xSH298 Image: State 1994 (F6×M1) SL 0.45 0.23 2 GS25 19 <0.05 <0.05 A54226 (FRA000102) France 1994 (LH824XT25)AF4x- B73) SL 0.45 0.15 2 GS25 10 <0.05	((LH82(4)T25)sf (-				GS25					ER94ECN550
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	2)xSH298									FRA000202
	France 1994 (F6×M1)	SL	0.45	0.23	2	119	< 0.05	< 0.05		A54226
$ \begin{array}{ c c c c c c c c c c c c c c c c c c c$	· · · ·				GS25					ER93ECS550
$ \begin{array}{c ccccccccccccccccccccccccccccccccccc$										FRA000102
$\begin{array}{c c c c c c c c c c c c c c c c c c c $	France 1994	SL	0.45	0.15	2	120	< 0.05	< 0.05		A54230
$\begin{array}{ c c c c c c c c c c c c c c c c c c c$	(LH824xT25)AF4x-				GS25					ER94ECS550
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	B73)									FRA000102
	France 1994	SL	0.45	0.23	2	119	< 0.05	< 0.05		A54229
f(2)) Image: Constraint of the section of the sec	(SH298(LH82(4)T25)s				GS25					ER94ECN550
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	f(2))									FRA000102
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	France 1995	SL	0.45	0.23	2	100	< <u>0.05</u>	< 0.05		A56445
France 1995 (F2xLH82-T25)SL (P2xLH82-T25)0.45 (P2xLH82-T25)0.23 (P2xLH82-T25)2 (P2xLH82-T25)0.45 (P2	(F2xLH82-T25)				GS18					ER95ECN550
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $										FRA000202
	France 1995	SL	0.45	0.23	2	100	< <u>0.05</u>	< 0.05		A56445
France 1995 (1087866)SL (1087866) 0.45 (1087866) 0.23 (1087866) 2 (1087866) 106 (1087866) < 0.05 (1087866) < 0.05 (1087866) $< A56445$ (1087876760 (10878676767676767676767676767676767676767	(F2xLH82-T25)				GS19					ER95ECN550
$ \begin{array}{c ccccccccccccccccccccccccccccccccccc$										FRA000203
039866) Image: Simple state in the s	France 1995 (tolerant	SL	0.45	0.23	2	106	< <u>0.05</u>	< 0.05		A56445
France 1995 (tolerant 039866)SL 0.45 0.23 2 GS18 106 < 0.05 < 0.05 $< A56445$ ER95ECN550 FRA000102France 1995 (transgenic hybrid)SL 0.53 0.27 2 GS18 116 < 0.05 < 0.05 < 0.05 $< A54760$ ER95ECS550 FRA000103France 1995 (transgenic hybrid)SL 0.53 0.27 2 GS18 116 < 0.05 < 0.05 < 0.05 $< A54760$ ER95ECS550 FRA000103France 1995 (transgenic hybrid)SL 0.45 0.23 2 GS19 116 < 0.05 < 0.05 < 0.05 $< A54760$ ER95ECS550 FRA000102France 1995 (transgenic hybrid)SL 0.45 0.23 2 GS19 119 < 0.05 < 0.05 < 0.05 $< A54760$ ER95ECS550 FRA000203France 1995 (transgenic hybrid)SL 0.60 0.24 2 GS18 119 < 0.05 < 0.05 < 0.05 $< A54760$ ER95ECS550 FRA000203France 1996 (transgenic hybrid)SL 0.60 0.24 2 GS18 104 < 0.05 < 0.05 $< A58191$ ER96ECN550 FRA000102France 1996 (transgenic hybrid)SL 0.80 0.32 2 GS18 104 < 0.05 < 0.05 $< A58191$ ER96ECN550 FRA000102France 1996 (transgenic hybrid)SL 0.80 0.32 2 GS18 104 < 0.05 < 0.05 $< A58191$ ER96ECN550 FRA000103	039866)				GS18					ER95ECN550
$ \begin{array}{c ccccccccccccccccccccccccccccccccccc$		~-			-					FRA000103
$ \begin{array}{c ccccccccccccccccccccccccccccccccccc$	France 1995 (tolerant	SL	0.45	0.23	2	106	< <u>0.05</u>	<0.05		A56445
France 1995 (transgenic hybrid)SL 0.53 0.27 2 GS18 116 GS18 $\frac{0.05}{2}$ <0.05 <0.05 <0.05 FRA000103 $A54760$ ER95ECS550 FRA000103France 1995 (transgenic hybrid)SL 0.53 0.27 2 GS18 116 GS18 <0.05 <0.05 <0.05 <0.05 $A54760$ ER95ECS550 FRA000102France 1995 (transgenic hybrid)SL 0.45 0.23 2 GS19 119 GS19 <0.05 <0.05 <0.05 <0.05 $A54760$ ER95ECS550 FRA000102France 1995 (transgenic hybrid)SL 0.45 0.23 2 GS19 119 GS19 <0.05 <0.05 <0.05 <0.05 $A54760$ ER95ECS550 FRA000203France 1995 (transgenic hybrid)SL 0.60 0.24 2 GS18 104 GS18 <0.05 <0.05 <0.05 $A54760$ ER95ECS550 FRA000202France 1996 (transgenic hybrid)SL 0.80 0.32 2 GS18 104 <0.05 <0.05 <0.05 <0.05 France 1996 (transgenic hybrid)SL 0.80 0.32 2 GS18 104 <0.05 <0.05 <0.05 <0.05 France 1996 (transgenic hybrid)SL 0.80 0.32 2 GS18 104 <0.05 <0.05 <0.05 <0.05 France 1996 (transgenic hybrid)SL 0.80 0.32 2 GS18 104 <0.05 <0.05 <0.05 <0.05 France 1996 <br< td=""><td>039866)</td><td></td><td></td><td></td><td>GS18</td><td></td><td></td><td></td><td></td><td>ER95ECN550</td></br<>	039866)				GS18					ER95ECN550
France 1995 (transgenic hybrid)SL 0.53 0.27 2 GS18 116 GS18 < 0.05 < 0.05 < 0.05 < 0.05 FRA000103 $< A54760$ FRA000103France 1995 (transgenic hybrid)SL 0.53 0.27 2 GS18 116 GS18 < 0.05 < 0.05 < 0.05 $< A54760$ FRA000102France 1995 (transgenic hybrid)SL 0.45 0.23 2 GS19 119 GS19 < 0.05 < 0.05 < 0.05 $< A54760$ FRA000102France 1995 (transgenic hybrid)SL 0.45 0.23 2 GS19 119 GS19 < 0.05 < 0.05 < 0.05 $< A54760$ FRA000203France 1995 (transgenic hybrid)SL 0.45 0.23 2 GS19 119 GS19 < 0.05 < 0.05 < 0.05 $< A54760$ FRA00202France 1996 (transgenic hybrid)SL 0.60 0.24 2 GS18 104 GS18 < 0.05 < 0.05 $< A58191$ ER96ECN550 FRA000102France 1996 (transgenic hybrid)SL 0.80 0.32 2 GS18 104 < 0.05 < 0.05 $< A58191$ ER96ECN550 FRA000102France 1996 (transgenic hybrid)SL 0.80 0.32 2 GS18 104 < 0.05 < 0.05 $< A58191$ ER96ECN550 FRA000102France 1996 (transgenic hybrid)SL 0.80 0.32 2 GS18 104 < 0.05 < 0.05 $< A58191$ ER96ECN550 FRA000103	F 1005	GT.	0.50	0.07		116	0.05	0.05	0.07	FRA000102
(transgenic hybrid)SL 0.53 0.27 2 GS18 116 < 0.05 < 0.05 < 0.05 < 0.05 < 54760 ER95ECS550 FRA000103France 1995 (transgenic hybrid)SL 0.45 0.23 2 GS19 119 < 0.05 < 0.05 < 0.05 < 54760 ER95ECS550 FRA000102France 1995 (transgenic hybrid)SL 0.45 0.23 2 GS19 119 < 0.05 < 0.05 < 0.05 < 54760 ER95ECS550 FRA000203France 1995 (transgenic hybrid)SL 0.45 0.23 2 GS19 119 < 0.05 < 0.05 < 0.05 < 54760 ER95ECS550 FRA000203France 1996 (transgenic hybrid)SL 0.60 0.24 2 GS18 104 < 0.05 < 0.05 < 58191 ER96ECN550 FRA000102France 1996 (transgenic hybrid)SL 0.80 0.32 2 GS18 104 < 0.05 < 0.05 < 58191 ER96ECN550 FRA000102France 1996 (transgenic hybrid)SL 0.80 0.32 2 GS18 104 < 0.05 < 0.05 < 58191 ER96ECN550 FRA000102	France 1995	SL	0.53	0.27	2	116	< <u>0.05</u>	<0.05	<0.05	A54760
France 1995 (transgenic hybrid)SL 0.53 0.27 2 GS18 116 < 0.05 < 0.05 < 0.05 < 0.05 $< A54760$ ER95ECS550 FRA000102France 1995 (transgenic hybrid)SL 0.45 0.23 2 GS19 119 < 0.05 < 0.05 < 0.05 $< A54760$ ER95ECS550 FRA000102France 1995 (transgenic hybrid)SL 0.45 0.23 2 GS19 119 < 0.05 < 0.05 < 0.05 $< A54760$ ER95ECS550 FRA000203France 1995 (transgenic hybrid)SL 0.45 0.23 2 GS18 119 < 0.05 < 0.05 < 0.05 $< A54760$ ER95ECS550 FRA000202France 1996 (transgenic hybrid)SL 0.60 0.24 2 GS18 104 < 0.05 < 0.05 $< A58191$ ER96ECN550 FRA000102France 1996 (transgenic hybrid)SL 0.80 0.32 2 GS18 104 < 0.05 < 0.05 $< A58191$ ER96ECN550 FRA000102	(transgenic hybrid)				GS18					ER95EC5550
France 1995 SL 0.53 0.27 2 116 < 0.05 < 0.05 < 0.05 $< R95ECS550$ $ER95ECS550$ France 1995 SL 0.45 0.23 2 119 < 0.05 < 0.05 < 0.05 < 4.05 $A54760$ France 1995 SL 0.45 0.23 2 $B19$ < 0.05 < 0.05 < 0.05 $A54760$ France 1995 SL 0.45 0.23 2 $B19$ < 0.05 < 0.05 < 0.05 $A54760$ France 1995 SL 0.45 0.23 2 $B19$ < 0.05 < 0.05 < 0.05 $A54760$ France 1996 SL 0.60 0.24 2 $GS18$ 0.4 < 0.05 < 0.05 < 0.05 $< A58191$ France 1996 SL 0.80 0.32 2 104 < 0.05 < 0.05 $< A58191$ France 1996 SL 0.80 0.32 2 $GS18$ 104 < 0.05 < 0.05 $< A58191$ Fra	E 1005	CT	0.52	0.07	2	116	-0.05	-0.05	-0.05	FKA000103
GS18GS18GS18GS18FR35ECS350 FRA000102France 1995 (transgenic hybrid)SL 0.45 0.23 2 GS19 119 <0.05 <0.05 <0.05 $A54760$ ER95ECS550 FRA000203France 1995 (transgenic hybrid)SL 0.45 0.23 2 GS19 119 <0.05 <0.05 <0.05 $A54760$ ER95ECS550 FRA000203France 1996 (transgenic hybrid)SL 0.60 0.24 2 GS18 104 <0.05 <0.05 <0.05 $A58191$ ER96ECN550 FRA000102France 1996 (transgenic hybrid)SL 0.80 0.32 2 GS18 104 <0.05 <0.05 <0.55 <0.55 FRA000102France 1996 (transgenic hybrid)SL 0.80 0.32 2 GS18 104 <0.05 <0.05 <0.55 <0.550 FRA000102France 1996 (transgenic hybrid)SL 0.80 0.32 2 GS18 104 <0.05 <0.05 <0.55 <0.550 FRA000102	(trance 1995	SL	0.55	0.27	2	110	< <u>0.05</u>	<0.05	<0.05	A34/00
France 1995 (transgenic hybrid)SL 0.45 0.23 2 GS19 119 < 0.05 < 0.05 < 0.05 < 0.05 $= 895ECS550$ FRA000203France 1995 (transgenic hybrid)SL 0.45 0.23 2 GS19 119 < 0.05 < 0.05 < 0.05 < 50.05 $= 895ECS550$ FRA000203France 1995 (transgenic hybrid)SL 0.45 0.23 2 GS18 119 < 0.05 < 0.05 < 0.05 < 50.05 FRA000202France 1996 (transgenic hybrid)SL 0.60 0.24 2 GS18 104 < 0.05 < 0.05 < 458191 ER96ECN550 FRA000102France 1996 (transgenic hybrid)SL 0.80 0.32 2 GS18 104 < 0.05 < 0.05 < 458191 ER96ECN550 FRA000102France 1996 (transgenic hybrid)SL 0.80 0.32 2 GS18 104 < 0.05 < 0.05 < 458191 ER96ECN550 FRA000102	(transgenic hybrid)				0510					EK93EC5330
Infance 1995 SL 0.43 0.23 $\frac{2}{GS19}$ 119 $\frac{0.05}{C}$ (0.05) <th< td=""><td>Eranaa 1005</td><td>CI.</td><td>0.45</td><td>0.22</td><td>2</td><td>110</td><td><0.05</td><td><0.05</td><td><0.05</td><td>A 54760</td></th<>	Eranaa 1005	CI.	0.45	0.22	2	110	<0.05	<0.05	<0.05	A 54760
Image: Construction of the system of the	(transgenic hybrid)	SL	0.45	0.25	2 GS10	119	< <u>0.05</u>	<0.05	<0.05	A34700 ED05EC\$550
France 1995 (transgenic hybrid) SL 0.45 0.23 2 GS19 119 $< \underline{0.05}$ < 0.05 < 0.05 < 4.05 $A54760$ France 1996 (transgenic hybrid) SL 0.60 0.24 2 GS18 104 $< \underline{0.05}$ < 0.05 < 0.05 $A58191$ ER96ECN550 FRA000102 $France 1996$ SL 0.80 0.32 2 GS18 104 < 0.05 < 0.05 $A58191$ France 1996 (transgenic hybrid) SL 0.80 0.32 2 GS18 104 < 0.05 < 0.05 $A58191$ ER96ECN550 FRA000102 France 1996 (transgenic hybrid) SL 0.80 0.32 2 GS18 104 < 0.05 < 0.05 $A58191$ ER96ECN550 FRA000103 $FRA000103$ < 0.05 < 0.05 < 0.05 < 0.05 < 0.05	(transgenie nybrid)				0319					ER95EC5550 FR 4000203
Indice 1995 SL 0.45 0.25 2 119 0.05 0.05 $R.94700$ (transgenic hybrid) SL 0.60 0.24 2 119 0.05 0.05 $R.94700$ ER95ECS550 France 1996 SL 0.60 0.24 2 104 0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05	France 1995	SI	0.45	0.23	2	110	<0.05	<0.05	<0.05	A 54760
Gamma (mansgenic hybrid)SL 0.60 0.24 2 GS18 104 < 0.05 < 0.05 $< A58191$ ER96ECN550 FRA000102France 1996 (transgenic hybrid)SL 0.80 0.32 2 GS18 104 < 0.05 < 0.05 $< A58191$ ER96ECN550 FRA000102France 1996 (transgenic hybrid)SL 0.80 0.32 2 GS18 104 < 0.05 < 0.05 $< A58191$ ER96ECN550 FRA000102	(transgenic hybrid)	SL	0.45	0.25	$\frac{2}{GS19}$	11)	< <u>0.05</u>	<0.05	<0.05	FR95EC\$550
France 1996 (transgenic hybrid) SL 0.60 0.24 2 GS18 104 < 0.05 < 0.05 A58191 ER96ECN550 FRA000102 France 1996 (transgenic hybrid) SL 0.80 0.32 2 GS18 104 < 0.05 < 0.05 $< A58191$ ER96ECN550 FRA000102 France 1996 (transgenic hybrid) SL 0.80 0.32 2 GS18 104 < 0.05 < 0.05 $< A58191$ ER96ECN550 FRA000103	(dansgeme nyond)				0517					FRA000202
Induct 1996 (transgenic hybrid)SL 0.80 0.24 2 104 0.05 0.05 $R38191$ France 1996 (transgenic hybrid)SL 0.80 0.32 2 104 <0.05 <0.05 <0.05 $A58191$ ER96ECN550 FRA000102GS18 0.32 2 104 <0.05 <0.05 <0.05 <0.05	France 1996	SL	0.60	0.24	2	104	<0.05	<0.05		A 58191
France 1996 (transgenic hybrid) SL 0.80 0.32 2 104 <0.05 <0.05 A58191 ER96ECN550 FRA000103	(transgenic hybrid)		0.00	0.24		104	<u>\</u>	.0.05		ER96ECN550
France 1996 (transgenic hybrid) SL 0.80 0.32 2 104 <0.05 <0.05 A58191 ER96ECN550 FRA000103	(aunogenne ny orig)				0.010					FRA000102
(transgenic hybrid) GS18 GS18 FRA000103	France 1996	SL	0.80	0.32	2	104	<0.05	< 0.05		A58191
FRA000103	(transgenic hybrid)	~	0.00		GS18					ER96ECN550
1111000100	(FRA000103

Country, Year,	Applic	ation			PHI days	Residues as mg/kg	glufosinate	free acid,	Ref
Variety	form	kg ai/ha	kg ai/hl	no ¹		glufosinate	MPP	NAG ²	
France 1996 (transgenic hybrid)	SL	0.45	0.18	2 GS18	104	< <u>0.05</u>	<0.05		A58191 ER96ECN550 FRA000104
France 1996 (transgenic hybrid)	SL	0.60	0.24	2 GS18	105	< <u>0.05</u>	<0.05		A58191 ER96ECN550 FRA000302
France 1996 (transgenic hybrid)	SL	0.80	0.32	2 GS18	105	<0.05	<0.05		A58191 ER96ECN550 FRA000303
France 1996 (transgenic hybrid)	SL	0.45	0.18	2 GS18	105	< <u>0.05</u>	<0.05		A58191 ER96ECN550 FRA000304
France 1996 (transgenic hybrid)	SL	0.60	0.24	2 GS18	110	< <u>0.05</u>	<0.05		A58191 ER96ECN550 FRA000402
France 1996 (transgenic hybrid)	SL	0.80	0.32	2 GS18	110	<0.05	<0.05		A58191 ER96ECN550 FRA000403
France 1996 (transgenic hybrid)	SL	0.45	0.18	2 GS18	110	< <u>0.05</u>	<0.05		A58191 ER96ECN550 FRA000404
France 1996 (transgenic hybrid)	SL	0.45	0.18	2 GS18	129	< <u>0.05</u>	<0.05		A58190 ER96ECS550 FRA000302
France 1996 (transgenic hybrid)	SL	0.60	0.24	2 GS18	129	< <u>0.05</u>	<0.05		A58190 ER96ECS550 FRA000303
France 1996 (transgenic hybrid)	SL	0.80	0.32	2 GS18	129	<0.05	<0.05		A58190 ER96ECS550 FRA000304
France 1996 (transgenic hybrid)	SL	0.60	0.24	2 GS19	103	< <u>0.05</u>	<0.05		A58191 ER96ECN550 FRA000202
France 1996 (transgenic hybrid)	SL	0.80	0.32	2 GS19	103	<0.05	<0.05		A58191 ER96ECN550 FRA000203
France 1996 (transgenic hybrid)	SL	0.45	0.18	2 GS19	103	< <u>0.05</u>	<0.05		A58191 ER96ECN550 FRA000204
France 1996 (transgenic hybrid)	SL	0.45	0.18	2 GS19	130	< <u>0.05</u>	<0.05		A58190 ER96ECS550 FRA000102
France 1996 (transgenic hybrid)	SL	0.60	0.24	2 GS19	130	< <u>0.05</u>	<0.05		A58190 ER96ECS550 FRA000103
France 1996 (transgenic hybrid)	SL	0.80	0.32	2 GS19	130	<0.05	<0.05		A58190 ER96ECS550 FRA000104
France 1996 (transgenic hybrid)	SL	0.45	0.18	2 GS19	140	< <u>0.05</u>	<0.05		A58190 ER96ECS550 FRA000202
France 1996 (transgenic hybrid)	SL	0.60	0.24	2 GS19	140	< <u>0.05</u>	<0.05		A58190 ER96ECS550 FRA000203
France 1996 (transgenic hybrid)	SL	0.80	0.32	2 GS19	140	<0.05	<0.05		A58190 ER96ECS550 FRA000204
Germany 1995 (LH82^4T25sf^4F2)	SL	0.45	0.15	2 GS18	99	< <u>0.05</u> (2)	<0.05 (2)		A56445 ER95ECN550 DEU010202

Variety Imm kg ai/ha gg ai/ha	Country, Year,	Applica	ation			PHI days	Residues as mg/kg	glufosinate	free acid,	Ref
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	Variety	form	kg ai/ha	kg ai/hl	no ¹		glufosinate	MPP	NAG ²	
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	Germany 1995	SL	0.45	0.15	2	90	< <u>0.05</u>	< 0.05		A56445
2989 DEDCODIO2 Germany 1995 SL 0.45 0.15 2 113 < 0.05 2 < 0.05 2 < 0.05 2 < 0.05 2 < 0.05 < 2.05 < 2.05 < 2.05 < 2.05 < 2.05 < 2.05 < 2.05 < 2.05 < 2.05 < 2.05 < 2.05 < 2.05 < 2.05 < 2.05 < 2.05 < 2.05 < 2.05 < 2.05 < 2.05 < 2.05 < 2.05 < 2.05 < 2.05 < 2.05 < 2.05 < 2.05 < 2.05 < 2.05 < 2.05 < 2.05 < 2.05 < 2.05 < 2.05 < 2.05 < 2.05 < 2.05 < 2.05 < 2.05 < 2.05 < 2.05 < 2.05 < 2.05 < 2.05 < 2.05 < 2.05 < 2.05 < 2.05 < 2.05 < 2.05 < 2.05 < 2.05 < 2.05 < 2.05 < 2.05 < 2.05 < 2.05 < 2.05 < 2.05 < 2.05	(LH82^4T25sf^4SH-				GS18					ER95ECN550
$ \begin{array}{c} \mbox{Germany 1995} \\ (LH82 + V25sf^{+}4xF2) \\ (LH82 + V25sf^{+}4xF2) \\ (LH82 + V25sf^{+}4xF2) \\ \mbox{Germany 1996} (F2 \\ hybride - LH+82) \\ \mbox{Germany 1996} (F2 \\ hybride - LH+82) \\ \mbox{Germany 1996} (F2 \\ hybride - LH+82) \\ \mbox{Germany 1996} (F2 \\ H+82) \\ \mbox{Germany 1996} (H482 \\ T25 \\ \mbox{Fr} F2) \\ \mbox{Germany 1996} (LH82 \\ T25 \\ \mbox{Fr} F2) \\ \mbox{Germany 1996} (LH82 \\ T25 \\ \mbox{Fr} F2) \\ \mbox{Germany 1996} (LH82 \\ T25 \\ \mbox{Fr} F2) \\ \mbox{Germany 1996} (LH82 \\ T25 \\ \mbox{Fr} F2) \\ \mbox{Germany 1996} (LH82 \\ T25 \\ \mbox{Fr} F2) \\ \mbox{Germany 1996} (LH82 \\ T25 \\ \mbox{Fr} F2) \\ \mbox{Germany 1996} (LH82 \\ T25 \\ \mbox{Fr} F2) \\ \mbox{Germany 1996} (LH82 \\ T25 \\ \mbox{Fr} F2) \\ \mbox{Germany 1996} (LH82 \\ T25 \\ \mbox{Fr} F2) \\ \mbox{Germany 1996} (LH82 \\ T25 \\ \mbox{Fr} F2) \\ \mbox{Germany 1996} (LH82 \\ \mbox{Germany 1996} (LH82 \\ T25 \\ \mbox{Fr} F2) \\ \mbox{Germany 1996} (LH82 \\ \mbox{Fr} F2) \\ German$	298)									DEU050102
	Germany 1995	SL	0.45	0.15	2	113	< 0.05 (2)	< 0.05 (2)		A56445
Contrary 1996 (P) Control (P) Dependence Germany 1996 (P) SL 0.60 0.20 2 (3518 2 (0.05 <0.05	(LH82^4T25sf^4xF2)				GS18					ER95ECN550
Germany 1996 (F2 hybride - LH-82) SL 0.60 0.20 2 GS18 126 < 0.05 AS8191 ER96fCNS50 DEU050102 Germany 1996 (F2 hybride - LH-82) SL 0.80 0.27 2 GS18 126 < 0.05 $AS8191$ ER96fCNS50 DEU050102 Germany 1996 (F2 hybride - LH-82) SL 0.45 0.15 2 GS18 126 < 0.05 $AS8191$ ER96fCNS50 DEU050104 Germany 1996 (Facet - SL 0.60 0.20 2 GS18 126 < 0.05 $AS8191$ ER96fCNS50 DEU050102 Germany 1996 (Facet - SL 0.60 0.20 2 GS18 125 < 0.05 $AS8191$ ER96fCNS50 DEU060102 Germany 1996 (Facet - SL 0.45 0.15 2 GS18 125 < 0.05 $AS8191$ ER96fCNS50 DEU060103 Germany 1996 (Facet - SL 0.60 0.20 2 GS18 122 < 0.05 $AS8191$ ER96fCNS50 DEU06002 Germany 1996 (Facet - SL 0.60 0.27 2 GS18 22 < 0.05 $AS8191$ ER96fCNS50 DEU06002 Germany 1996 (Facet - SL 0.45 0.15 2	(- · · · /									DEU010102
hybride - LH-82) International Constraints GS18 International Constraints EPOGECNS50 DEU050102 Germany 1996 (F2 hybride - LH-82) SL 0.80 0.27 2 GS18 126 <0.05	Germany 1996 (F2	SL	0.60	0.20	2	126	< 0.05	< 0.05		A58191
Construction Image in the ima	hybride - LH-82)	~-			GS18		·			ER96ECN550
Germany 1996 (F2 hybride - LH-82) SL 0.80 0.27 2 GS18 126 <0.05 <0.05 AS\$191 Germany 1996 (F2 hybride - LH-82) SL 0.45 0.15 2 GS18 126 <0.05	J									DEU050102
hybride - LH-82) International Constraints GS18 International Constraints FROGECNS50 DEU050103 Germany 1996 (F2c SL 0.45 0.15 2 GS18 126 < 0.05 $< AS8191$ ERG6ECNS50 DEU050104 Germany 1996 (Facet - transgen) SL 0.60 0.20 2 GS18 125 < 0.05 $< AS8191$ ERG6ECNS50 DEU050104 Germany 1996 (Facet - transgen) SL 0.60 0.27 2 GS18 125 < 0.05 $< AS8191$ ERG6ECNS50 DEU060102 Germany 1996 (Facet - transgen) SL 0.45 0.15 2 GS18 122 < 0.05 $< AS8191$ ERG6ECNS50 DEU060103 Germany 1996 (Facet - transgen) SL 0.45 0.15 2 GS18 122 < 0.05 $< AS8191$ ERG6ECNS50 DEU060202 Germany 1996 (Facet - SL 0.45 0.15 2 GS18 122 < 0.05 $< AS8191$ ERG6ECNS50 DEU060203 Germany 1996 (LH82 SL 0.80 0.27 2 GS18 122 < 0.05 $< AS8191$ ERG6ECNS50 DEU010202 Germany 1996 (LH82 SL 0.45 0.15 2 GS19	Germany 1996 (F2	SL	0.80	0.27	2	126	< 0.05	< 0.05		A58191
Germany 1996 (F2 hybride - LH-82) SL 0.45 0.15 2 GS18 126 < 0.05 < 0.05 $< AS8191$ ER965CNS50 DEU050104 Germany 1996 (Facet - SL transgen) 0.60 0.20 2 GS18 125 < 0.05 $< AS8191$ ER966CNS50 DEU060102 Germany 1996 (Facet - SL transgen) 0.80 0.27 2 GS18 125 < 0.05 $< AS8191$ ER966CNS50 DEU060102 Germany 1996 (Facet - SL transgen) 0.45 0.15 2 GS18 125 < 0.05 $< AS8191$ ER966CNS50 DEU060103 Germany 1996 (Facet - SL transgen) 0.60 0.20 2 GS18 < 122 < 0.05 $< AS8191$ ER966CNS50 DEU060104 Germany 1996 (Facet - SL transgen) 0.60 0.27 2 GS18 < 2 < 0.05 $< AS8191$ ER966CNS50 DEU060103 Germany 1996 (Facet - SL transgen) 0.45 0.15 2 GS18 < 2 < 0.05 $< AS8191$ ER966CNS50 DEU06003 Germany 1996 (LH82 SL transgen) 0.45 0.15 2 GS19 < 0.05 $< AS8191$ ER966CNS50 DEU010020 Germany 1996 (LH82 SL transgen) 0.45 0.15	hvbride - LH-82)	~ -			GS18					ER96ECN550
	J									DEU050103
	Germany 1996 (F2	SL	0.45	0.15	2	126	< 0.05	< 0.05		A58191
Second	hvbride - LH-82)	~ -			GS18		· <u></u>			ER96ECN550
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	/									DEU050104
$ \begin{array}{c ccccccccccccccccccccccccccccccccccc$	Germany 1996 (Facet -	SL	0.60	0.20	2	125	< 0.05	<0.05		A58191
Internation Delument	transgen)	~-			GS18		·			ER96ECN550
Germany 1996 (Facet - SL 0.80 0.27 2 125 <0.05 <0.05 A58191 germany 1996 (Facet - SL 0.45 0.15 2 (S18) 2 <0.05	8)									DEU060102
$ \begin{array}{c ccccccccccccccccccccccccccccccccccc$	Germany 1996 (Facet -	SL	0.80	0.27	2	125	< 0.05	<0.05		A58191
Internation Internation Description DEU060103 Germany 1996 (Facet - SL 0.45 0.15 2 GS18 125 <0.05 <0.05 A58191 transgen) 0.60 0.20 2 GS18 122 <0.05 <0.05 A58191 transgen) 0.60 0.20 2 GS18 122 <0.05 <0.05 A58191 transgen) 0.60 0.20 2 GS18 122 <0.05 <0.05 A58191 transgen) 0.80 0.27 2 GS18 122 <0.05 <0.05 A58191 transgen) 0.45 0.15 2 GS18 122 <0.05 <0.05 A58191 transgen) 0.45 0.45 0.15 2 GS18 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0	transgen)	22	0.00		- GS18	120	10100	10100		ER96ECN550
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	8)									DEU060103
$ \begin{array}{c} \mbox{transgen} & \mbox{form} & \mbo$	Germany 1996 (Facet -	SL	0.45	0.15	2	125	< 0.05	<0.05		A58191
$\begin{array}{ c c c c c c c c c c c c c c c c c c c$	transgen)	52	0.15	0.12	GS18	120	~ <u>0.05</u>	10.02		ER96ECN550
Germany 1996 (Facet - SL 0.60 0.20 2 122 <0.05 $A38191$ germany 1996 (Facet - SL 0.80 0.27 2 $6S18$ $c0.05$ $c0.05$ $A58191$ germany 1996 (Facet - SL 0.80 0.27 2 $CS18$ $c0.05$ $c0.05$ $A58191$ germany 1996 (Facet - SL 0.45 0.15 2 $C2$ $c0.05$ $c0.05$ $A58191$ germany 1996 (LH82 SL 0.60 0.20 2 137 $c0.05$ $c0.05$ $A58191$ germany 1996 (LH82 SL 0.60 0.20 2 137 $c0.05$ $c0.05$ $A58191$ germany 1996 (LH82 SL 0.60 0.27 2 137 $c0.05$ $c0.05$ $A58191$ germany 1996 (LH82 SL 0.45 0.15 2 $C0.05$ $c0.05$ $A58191$ germany 1996 (LH82 SL 0.45 0.15 2 $C0.05$ $c0.05$ $A58191$ germany 1996 (LH82 SL 0.60 0.27 2 </td <td>8)</td> <td></td> <td></td> <td></td> <td></td> <td></td> <td></td> <td></td> <td></td> <td>DEU060104</td>	8)									DEU060104
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	Germany 1996 (Facet -	SL	0.60	0.20	2	122	< 0.05	<0.05		A58191
Integen Image of the second seco	transgen)	22	0.00	0.20	- GS18		\ <u>0.000</u>	10100		ER96ECN550
Germany 1996 (Facet - SL transgen) 0.80 0.27 2 GS18 122 <0.05 (0.05) <td>a anogen)</td> <td></td> <td></td> <td></td> <td>0010</td> <td></td> <td></td> <td></td> <td></td> <td>DEU060202</td>	a anogen)				0010					DEU060202
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	Germany 1996 (Facet -	SL	0.80	0.27	2	122	<0.05	<0.05		A58191
Imageny	transgen)	22	0.00		- GS18		10100	10100		ER96ECN550
Germany 1996 (Facet - SL transgen) 0.45 0.15 2 122 < 0.05 < 0.05 $< A58191$ Germany 1996 (LH82 SL 0.60 0.20 2 137 < 0.05 < 0.05 $< A58191$ T25 SF x F2) SL 0.60 0.20 2 137 < 0.05 < 0.05 $< A58191$ T25 SF x F2) SL 0.80 0.27 2 137 < 0.05 < 0.05 $< A58191$ T25 SF x F2) SL 0.80 0.27 2 137 < 0.05 < 0.05 $< A58191$ T25 SF x F2) SL 0.45 0.15 2 (37) < 0.05 < 0.05 $< A58191$ T25 SF x F2) SL 0.45 0.15 2 (37) < 0.05 < 0.05 $< A58191$ T25 SF x Fe) SL 0.60 0.20 2 (145) < 0.05 < 0.05 $< A58191$ T25 SF x Fe) SL 0.60 0.27 2 (145) < 0.05 < 0.05 $< A58191$	8)									DEU060203
$ \begin{array}{c ccccccccccccccccccccccccccccccccccc$	Germany 1996 (Facet -	SL	0.45	0.15	2	122	< 0.05	< 0.05		A58191
Images	transgen)	~ -			- GS18		· <u></u>			ER96ECN550DEU
Germany 1996 (LH82 T25 SF x F2) SL 0.60 0.20 2 GS19 137 $< \underline{0.05}$ < 0.05 $A58191$ ER96ECN550 DEU010202 Germany 1996 (LH82 T25 SF x F2) SL 0.80 0.27 2 GS19 137 < 0.05 < 0.05 $A58191$ ER96ECN550 DEU010203 Germany 1996 (LH82 T25 SF x F2) SL 0.45 0.15 2 GS19 137 $< \underline{0.05}$ < 0.05 $A58191$ ER96ECN550 DEU010203 Germany 1996 (LH82 T25 SF x F2) SL 0.45 0.15 2 GS19 145 $< \underline{0.05}$ < 0.05 $A58191$ ER96ECN550 DEU010204 Germany 1996 (LH82 T25 SF x Fe) SL 0.60 0.20 2 GS19 145 $< \underline{0.05}$ < 0.05 $A58191$ ER96ECN550 DEU010102 Germany 1996 (LH82 T25 SF x Fe) SL 0.80 0.27 2 GS19 145 < 0.05 < 0.05 $A58191$ ER96ECN550 DEU010103 Germany 1996 (LH82 T25 SF x Fe) SL 0.45 0.15 2 GS19 145 < 0.05 < 0.05 $A58191$ ER96ECN550 DEU010103 Germany 1996 (LH82 T25 SF x Fe) SL 0.45 0.15 2 GS19 145 </td <td>a anogen)</td> <td></td> <td></td> <td></td> <td>0.510</td> <td></td> <td></td> <td></td> <td></td> <td>060204</td>	a anogen)				0.510					060204
T25 SF x F2) GS19 GS19 GS19 ER96ECN550 ER96ECN550 DEU010202 Germany 1996 (LH82 SL 0.80 0.27 2 137 <0.05	Germany 1996 (LH82	SL	0.60	0.20	2	137	< 0.05	< 0.05		A58191
Germany 1996 (LH82 T25 SF x F2)SL 0.80 0.27 2 GS19 137 <0.05 <0.05 <0.05 $A58191$ ER96ECN550 DEU010203Germany 1996 (LH82 T25 SF x F2)SL 0.45 0.15 2 GS19 137 <0.05 <0.05 <0.57 <0.57 Germany 1996 (LH82 T25 SF x F2)SL 0.60 0.20 2 GS19 145 <0.05 <0.05 <0.57 <0.58191 ER96ECN550 DEU010204Germany 1996 (LH82 T25 SF x Fe)SL 0.60 0.20 2 GS19 145 <0.05 <0.05 <0.57 <0.58191 ER96ECN550 DEU010204Germany 1996 (LH82 T25 SF x Fe)SL 0.80 0.27 2 GS19 145 <0.05 <0.05 <0.57 <0.58191 ER96ECN550 DEU010102Germany 1996 (LH82 SF x Fe)SL 0.45 0.15 2 GS19 145 <0.05 <0.05 <0.57 <0.57 ER96ECN550 DEU010103Germany 1996 (LH82 RS19SL 0.45 0.15 2 GS19 145 <0.05 <0.05 <0.57 <0.57 ER96ECN550 DEU010103Germany 1996 (LH82 RS2SL 0.45 0.15 2 GS19 105 <0.05 <0.05 <0.57 ER96ECN550 DEU010104Italy 1994 (IB73(4)xT14]sfx - LM82)SL 0.45 0.15 2 GS18 106 <0.05 <0.05 <0.05 <0.05 <0.05 ER95ECS50 ITA000102Italy 1995 (transgenic hybrid)SL 0.45 <td>T25 SF x F2)</td> <td></td> <td></td> <td></td> <td>GS19</td> <td></td> <td></td> <td></td> <td></td> <td>ER96ECN550</td>	T25 SF x F2)				GS19					ER96ECN550
Germany 1996 (LH82 T25 SF x F2) SL 0.80 0.27 2 GS19 137 <0.05 <0.05 $A58191$ ER96ECN550 DEU010203 Germany 1996 (LH82 T25 SF x F2) SL 0.45 0.15 2 GS19 137 <0.05 <0.05 $A58191$ ER96ECN550 DEU010203 Germany 1996 (LH82 T25 SF x Fe) SL 0.60 0.20 2 GS19 145 <0.05 <0.05 $A58191$ ER96ECN550 DEU010204 Germany 1996 (LH82 T25 SF x Fe) SL 0.60 0.27 2 GS19 145 <0.05 <0.05 $A58191$ ER96ECN550 DEU01002 Germany 1996 (LH82 T25 SF x Fe) SL 0.80 0.27 2 GS19 145 <0.05 <0.05 $A58191$ ER96ECN550 DEU010102 Germany 1996 (LH82 T25 SF x Fe) SL 0.45 0.15 2 GS19 145 <0.05 <0.05 $A58191$ ER96ECN550 DEU010103 Germany 1996 (LH82 T25 SF x Fe) SL 0.45 0.15 2 GS19 145 <0.05 <0.05 $A58230$ ER96ECN550 DEU010104 Italy 1994 (B73(4)xT14]sfx - LM82) <	,									DEU010202
$\begin{array}{c c c c c c c c c c c c c c c c c c c $	Germany 1996 (LH82	SL	0.80	0.27	2	137	< 0.05	< 0.05		A58191
Germany 1996 (LH82 T25 SF x F2)SL 0.45 0.15 2 GS19 137 <0.05 <0.05 <0.05 $A58191$ ER96ECN550 DEU010204Germany 1996 (LH82 T25 SF x Fe)SL 0.60 0.20 2 GS19 145 <0.05 <0.05 <0.05 <0.05 Germany 1996 (LH82 T25 SF x Fe)SL 0.60 0.27 2 GS19 145 <0.05 <0.05 <0.05 <0.58191 Germany 1996 (LH82 T25 SF x Fe)SL 0.80 0.27 2 GS19 145 <0.05 <0.05 <0.05 <0.58191 Germany 1996 (LH82 T25 SF x Fe)SL 0.45 0.15 2 GS19 145 <0.05 <0.05 <0.05 <0.58191 Germany 1996 (LH82 T25 SF x Fe)SL 0.45 0.15 2 GS19 145 <0.05 <0.05 <0.05 <0.570 Germany 1996 (LH82 T25 SF x Fe)SL 0.45 0.15 2 GS19 145 <0.05 <0.05 <0.05 <0.05 Germany 1996 (LH82 T25 SF x Fe)SL 0.45 0.15 2 GS19 <0.05 <0.05 <0.05 <0.05 <0.05 Italy 1994 ([B73(4)xT14]sfx - LM82)SL 0.45 0.15 2 GS18 <0.05 <0.05 <0.05 <0.05 <0.05 Italy 1995 (transgenic hybrid)SL 0.45 0.15 2 GS18 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 Italy 1905 (transgenic hyb	T25 SF x F2)				GS19					ER96ECN550
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$\begin{array}{c c c c c c c c c c c c c c c c c c c $	Germany 1996 (LH82	SL	0.45	0.15	2	137	< 0.05	< 0.05		A58191
Image: constraint of the second se	T25 SF x F2)				GS19					ER96ECN550
Germany 1996 (LH82 T25 SF x Fe)SL 0.60 0.20 2 GS19 145 < 0.05 < 0.05 $< A58191$ ER96ECN550 DEU010102Germany 1996 (LH82 T25 SF x Fe)SL 0.80 0.27 2 GS19 145 < 0.05 < 0.05 $< A58191$ ER96ECN550 DEU010103Germany 1996 (LH82 T25 SF x Fe)SL 0.45 0.15 2 GS19 145 < 0.05 < 0.05 $< A58191$ ER96ECN550 DEU010103Germany 1996 (LH82 T25 SF x Fe)SL 0.45 0.15 2 GS19 145 < 0.05 < 0.05 $< A58191$ ER96ECN550 DEU010103Italy 1994 (IB73(4)xT14]sfx - LM82)SL 0.45 0.15 2 GS18 105 < 0.05 < 0.05 $< A54230$ ER94ECS550 ITA000102Italy 1995 (transgenic hybrid)SL 0.45 0.15 2 GS18 108 < 0.05 < 0.05 < 0.05 $< A54760$ ER95ECS550 ITA000104Italy 1995 (transgenic hybrid)SL 0.45 0.15 2 GS18 108 < 0.05 < 0.05 < 0.05 $< A54760$ ER95ECS550 ITA000104	,									DEU010204
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Germany 1996 (LH82 T25 SF x Fe) SL 0.80 0.27 2 GS19 145 <0.05 <0.05 A58191 ER96ECN550 DEU010103 Germany 1996 (LH82 T25 SF x Fe) SL 0.45 0.15 2 GS19 145 <0.05 <0.05 <0.05 <0.05 Germany 1996 (LH82 T25 SF x Fe) SL 0.45 0.15 2 GS19 145 <0.05 <0.05 <0.05 <0.05 Italy 1994 (IB73(4)xT14]sfx - LM82) SL 0.45 0.15 2 GS23 105 <0.05 <0.05 <0.05 <0.05 Italy 1995 (transgenic hybrid) SL 0.45 0.15 2 GS18 108 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <	T25 SF x Fe)				GS19					ER96ECN550
Germany 1996 (LH82 T25 SF x Fe)SL 0.80 0.27 2 GS19 145 <0.05 <0.05 <0.05 $A58191$ ER96ECN550 DEU010103Germany 1996 (LH82 T25 SF x Fe)SL 0.45 0.15 2 GS19 145 <0.05 <0.05 $A58191$ ER96ECN550 DEU010103Italy 1994 ([B73(4)xT14]sfx - LM82)SL 0.45 0.15 2 GS23 105 <0.05 <0.05 $A54230$ ER94ECS550 ITA000102Italy 1995 (transgenic hybrid)SL 0.45 0.15 2 GS18 108 <0.05 <0.05 <0.05 $A54760$ ER95ECS550 ITA000104										DEU010102
T25 SF x Fe) GS19 ER96ECN550 Germany 1996 (LH82 SL 0.45 0.15 2 145 < 0.05 < 0.05 < 458191 T25 SF x Fe) SL 0.45 0.15 2 145 < 0.05 < 0.05 < 458191 Italy 1994 SL 0.45 0.15 2 105 < 0.05 < 0.05 < 454230 [B73(4)xT14]sfx - LM82) SL 0.45 0.15 2 105 < 0.05 < 0.05 < 454230 Italy 1995 (transgenic hybrid) SL 0.45 0.15 2 < 108 < 0.05 < 0.05 < 4.05 $A54760$ Italy 1995 (transgenic hybrid) SL 0.45 0.15 2 < 104 < 0.05 < 0.05 < 0.05 < 10.5 Italy 1995 (transgenic hybrid) SL 0.45 0.15 2 < 104 < 0.05 < 0.05 < 0.05 < 10.5 Italy 1995 (transgenic hybrid) SL 0.45 0.15 2 < 104 < 0.05 < 0.05 < 0.05 <	Germany 1996 (LH82	SL	0.80	0.27	2	145	< 0.05	< 0.05		A58191
Germany 1996 (LH82 SL 0.45 0.15 2 GS19 145 < 0.05 < 0.05 A58191 ER96ECN550 DEU010104 Italy 1994 ([B73(4)xT14]sfx - LM82) SL 0.45 0.15 2 GS23 105 < 0.05 < 0.05 A54230 Italy 1994 ([B73(4)xT14]sfx - LM82) SL 0.45 0.15 2 GS23 105 < 0.05 < 0.05 A54230 Italy 1995 (transgenic hybrid) SL 0.45 0.15 2 GS18 108 < 0.05 < 0.05 < 0.05 A54760 Italy 1995 (transgenic hybrid) SL 0.45 0.15 2 GS18 104 < 0.05 < 0.05 < 0.05 < 10.45 < 0.05	T25 SF x Fe)				GS19					ER96ECN550
Germany 1996 (LH82 T25 SF x Fe) SL 0.45 0.15 2 GS19 145 $< \underline{0.05}$ < 0.05 A58191 ER96ECN550 DEU010104 Italy 1994 ([B73(4)xT14]sfx - LM82) SL 0.45 0.15 2 GS23 105 < 0.05 < 0.05 A54230 ER94ECS550 ITA000102 Italy 1995 (transgenic hybrid) SL 0.45 0.15 2 GS18 108 < 0.05 < 0.05 < 0.05 A54760 ER95ECS550 ITA000104 Italy 1995 (transgenic hybrid) SL 0.45 0.15 2 GS18 104 < 0.05 < 0.05 < 0.05 < 0.05	,									DEU010103
T25 SF x Fe) GS19 ER96ECN550 DEU010104 Italy 1994 ([B73(4)xT14]sfx - LM82) SL 0.45 0.15 2 GS23 105 <0.05 <0.05 A54230 ER94ECS550 ITA000102 Italy 1995 (transgenic hybrid) SL 0.45 0.15 2 GS18 108 <0.05 <0.05 <0.05 A54760 ER95ECS550 ITA000104	Germany 1996 (LH82	SL	0.45	0.15	2	145	< 0.05	< 0.05		A58191
$ \begin{array}{ c c c c c c c c c c c c c c c c c c c$	T25 SF x Fe)				GS19					ER96ECN550
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $										DEU010104
$ \begin{array}{ c c c c c c c c c c c c c c c c c c c$	Italy 1994	SL	0.45	0.15	2	105	< 0.05	< 0.05		A54230
LM82) ITA000102 Italy 1995 (transgenic hybrid) SL 0.45 0.15 2 108 <0.05 <0.05 <0.05 $A54760$ ER95ECS550 ITA000104 ITA000104 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05 <0.05	([B73(4)xT14]sfx -				GS23					ER94ECS550
Italy 1995 (transgenic hybrid) SL 0.45 0.15 2 GS18 108 <0.05 <0.05 A54760 ER95ECS550 ITA000104 tab: 1095 (transgenic SL 0.45 0.15 2 104 0.05 <0.05	LM82)									ITA000102
hybrid) GS18 ER95ECS550 ITA000104 ITA000104	Italy 1995 (transgenic	SL	0.45	0.15	2	108	< 0.05	< 0.05	< 0.05	A54760
ITA000104	hybrid)				GS18					ER95ECS550
										ITA000104
1.13 (uransgenic JSL 10.45 10.15 12 1104 1<0.05 1<0.05 1<0.05 1A54760	Italy 1995 (transgenic	SL	0.45	0.15	2	104	< 0.05	< 0.05	< 0.05	A54760
hybrid) GS18 ER95ECS550	hybrid)				GS18					ER95ECS550
ITA000204										ITA000204

Country, Year,	Application				PHI days	Ref			
Variety	form	kg ai/ha	kg ai/hl	no ¹		glufosinate	MPP	NAG ²	
Italy 1995 (transgenic hybrid)	SL	0.45	0.15	2 GS18	108	<0.05	<0.05	<0.05	A54760 ER95ECS550 ITA000103
Italy 1995 (transgenic hybrid)	SL	0.45	0.15	2 GS18	104	<0.05	<0.05	<0.05	A54760 ER95ECS550 ITA000203
Italy 1995 (transgenic hybrid)	SL	0.45	0.15	2 GS18	108	<0.05	<0.05	<0.05	A54760 ER95ECS550 ITA000102
Italy 1995 (transgenic hybrid)	SL	0.45	0.15	2 GS18	104	<0.05	<0.05	<0.05	A54760 ER95ECS550 ITA000202
Italy 1996 (transgenic hybrid)	SL	0.80	0.27	2 GS18	100	< <u>0.05</u>	<0.05		A58190 ER96ECS550 ITA000105
Italy 1996 (transgenic hybrid)	SL	0.80	0.27	2 GS18	124	< <u>0.05</u>	<0.05		A58190 ER96ECS550 ITA000205
Italy 1996 (transgenic hybrid)	SL	0.60	0.20	2 GS18	100	< <u>0.05</u>	<0.05		A58190 ER96ECS550 ITA000103
Italy 1996 (transgenic hybrid)	SL	0.80	0.27	2 GS18	100	< <u>0.05</u>	<0.05		A58190 ER96ECS550 ITA000104
Italy 1996 (transgenic hybrid)	SL	0.60	0.20	2 GS18	124	<0.05	<0.05		A58190 ER96ECS550 ITA000203
Italy 1996 (transgenic hybrid)	SL	0.80	0.27	2 GS18	124	< <u>0.05</u>	<0.05		A58190 ER96ECS550 ITA000204
Italy 1996 (transgenic hybrid)	SL	0.80	0.27	2 GS19	104	< <u>0.05</u>	<0.05		A58190 ER96ECS550 ITA000305
Italy 1996 (transgenic hybrid)	SL	0.60	0.20	2 GS19	104	<0.05	<0.05		A58190 ER96ECS550 ITA000303
Italy 1996 (transgenic hybrid)	SL	0.80	0.27	2 GS19	104	< <u>0.05</u>	<0.05		A58190 ER96ECS550 ITA000304
Spain 1994 (LH119- 6xT14/LH82)	SL	0.45	0.15	2 GS25	102	<0.05	<0.05		A54230 ER94ECS550 ESP000102
Spain 1995 (transgenic hybrid)	SL	0.45	0.15	2 GS18	95	<0.05	<0.05	<0.05	A54760 ER95ECS550 ESP000103
Spain 1995 (transgenic hybrid)	SL	0.45	0.15	2 GS18	95	<0.05	<0.05	<0.05	A54760 ER95ECS550 ESP000102
Spain 1996 (transgenic hybrid)	SL	0.45	0.15	2 GS18	95	<0.05	<0.05		A58190 ER96ECS550 ESP000102
Spain 1996 (transgenic hybrid)	SL	0.60	0.20	2 GS18	95	<0.05	<0.05		A58190 ER96ECS550 ESP000103
Spain 1996 (transgenic hybrid)	SL	0.80	0.27	2 GS18	95	< <u>0.05</u>	<0.05		A58190 ER96ECS550 ESP000104
Spain 1996 (transgenic hybrid)	SL	0.45	0.15	2 GS18	96	<0.05	<0.05		A58190 ER96ECS550 ESP000202

Country, Year,	Applica	ation			PHI days	Residues as mg/kg	glufosinate	free acid,	Ref
Variety	form	kg ai/ha	kg ai/hl	no ¹		glufosinate	MPP	NAG ²	
Spain 1996 (transgenic hybrid)	SL	0.60	0.20	2 GS18	96	<0.05	<0.05		A58190 ER96ECS550 ESP000203
Spain 1996 (transgenic hybrid)	SL	0.80	0.27	2 GS18	96	< <u>0.05</u>	<0.05		A58190 ER96ECS550 ESP000204
Spain 1996 (transgenic hybrid)	SL	0.45	0.15	2 GS18	113	<0.05	<0.05		A58190 ER96ECS550 ESP000302
Spain 1996 (transgenic hybrid)	SL	0.60	0.20	2 GS18	113	<0.05	<0.05		A58190 ER96ECS550 ESP000303
Spain 1996 (transgenic hybrid)	SL	0.80	0.27	2 GS18	113	< <u>0.05</u>	<0.05		A58190 ER96ECS550 ESP000304

¹. GS: growth stage at final treatment

GS18 8 leaves unfolded

GS19 9 or more leaves unfolded

GS23

5th leaf unfolded 6th leaf unfolded GS24

GS25 7th leaf unfolded

² included in glufosinate-ammonium result.

Table 45. Glufosinate residues in transgenic maize resulting from supervised trials in the USA and Argentina. Analyses of replicate samples are reported individually. Double-underlined residues are from treatments according to GAP and were used to estimate maximum residue levels.

Country, Year,	Applic	ation			PHI	Residues, mg/kg, as glufosinate free Ref acid ²				
Variety	form	kg ai/ha	kg ai/hl	No ¹	days	glufosinate- ammonium	MPP	NAG		
USA (NE) 1993 (LH59 X LH51) (LH119)(4) X T14)	SL	1.8 +2.6	1.9 +2.8	2 (60 cm)	96	NQ	NQ	NQ	ER-93-USA- 01-NE-02	
USA (IA) 1993 (LH216) (LH119) (4) X (T14))	SL	1.5	1.6	1 (60 cm)	100	NQ	NQ	NQ	ER-93-USA- 01-IA-02	
USA (IA) 1993 (LH216) X (LH119) (4) × T14)	SL	0.36		1 (30 cm)	118	NQ (3)	NQ (3)	NQ (3)	ER-93-USA- 01-IA-01	
USA (IA) 1993 (LH216) X (LH119) (4) × T14)	SL	0.50		1 (30 cm)	114	NQ (3)	NQ (3)	NQ (3)	ER-93-USA- 01-IA-01	
USA (IA) 1993 (LH216) X (LH119) (4) × T14)	SL	0.50		1 (60 cm)	114	<u>NQ</u> (3)	NQ (3)	NQ (3)	ER-93-USA- 01-IA-01	
USA (IL) 1993 (LH 59x LH51)(LH119) (4) X T14)	SL	0.36	0.39	1 (30 cm)	100	NQ (3)	NQ (3)	NQ (3)	ER-93-USA- 01-IL-01	
USA (IL) 1993 (LH 59x LH51)(LH119) (4) X T14)	SL	0.50	0.54	1 (30 cm)	100	NQ (3)	NQ (3)	NQ (3)	ER-93-USA- 01-IL-01	
USA (IL) 1993 (LH 59x LH51)(LH119) (4) X T14)	SL	0.50	0.54	1 (60 cm)	95	<u>NQ</u> (3)	NQ (3)	NQ (3)	ER-93-USA- 01-IL-01	

Country, Year,	Applic	ation			PHI	I Residues, mg/kg, as glufosinate free Ref acid ²				
Variety	form	kg ai/ha	kg ai/hl	No ¹	days	glufosinate- ammonium	MPP	NAG		
USA (IL) 1993 (LH 59x LH51)(LH119) (4) X T14)	SL	0.36 +0.50	0.39 +0.54	2 (60 cm)	95	<u>NQ</u> (3)	NQ (3)	NQ (3)	ER-93-USA- 01-IL-01	
USA (NE) 1993 (LH 59x LH51)(LH119) (4) X T14)	SL	0.36		1 (30 cm)	107	NQ (3)	NQ (3)	NQ (3)	ER-93-USA- 01-NE-01	
USA (NE) 1993 (LH 59x LH51)(LH119) (4) X T14)	SL	0.50	0.48	1 (30 cm)	107	NQ (3)	NQ (3)	NQ (3)	ER-93-USA- 01-NE-01	
USA (NE) 1993 (LH 59x LH51)(LH119) (4) X T14)	SL	0.50	0.48	1 (60 cm)	95	<u>NQ</u> (3)	NQ (3)	NQ (3)	ER-93-USA- 01-NE-01	
USA (IN) 1993 [LH119(4) X T14] X LH216)	SL	0.36	0.39	1 (30 cm)	115	NQ (3)	NQ (3)	NQ (3)	ER-93-USA- 01-IN-01	
USA (IN) 1993 [LH119(4) X T14] X LH216)	SL	0.50	0.54	1 (30 cm)	115	NQ (3)	NQ (3)	NQ (3)	ER-93-USA- 01-IN-01	
USA (IN) 1993 [LH119(4) X T14] X LH216)	SL	0.36 +0.50	0.39 +0.54	2 (60 cm)	106	NQ (3)	NQ (3)	NQ (3)	ER-93-USA- 01-IN-01	
USA (ND) 1993 (LH85 X LH160) (LH119)(4) X T14)	SL	0.36	0.39	1 (30 cm)	107	NQ (3)	NQ (3)	NQ (3)	ER-93-USA- 01-ND-01	
USA (ND) 1993 (LH85 X LH160) (LH119)(4) X T14)	SL	0.50	0.54	1 (30 cm)	107	NQ (3)	NQ (3)	NQ (3)	ER-93-USA- 01-ND-01	
USA (ND) 1993 (LH85 X LH160) (LH119)(4) X T14)	SL	0.36 +0.50	0.39 +0.54	2 (60 cm)	97	NQ (3)	NQ (3)	NQ (3)	ER-93-USA- 01-ND-01	
USA (ND) 1993 (LH85 X LH160) (LH119)(4) X T14)	SL	0.36	0.39	1 (30 cm)	117	NQ (3)	NQ (3)	NQ (3)	ER-93-USA- 01-ND-01	
USA (ND) 1993 (LH85 X LH160) (LH119)(4) X T14)	SL	0.50	0.54	1 (60 cm)	117	<u>NO</u> (3)	NQ (3)	NQ (3)	ER-93-USA- 01-ND-01	
USA (ND) 1993 (LH85 X LH160) (LH119)(4) X T14)	SL	0.36 +0.50	0.39 +0.54	2 (31 cm)	107	NQ (3)	NQ (3)	NQ (3)	ER-93-USA- 01-ND-01	
USA (MO) 1993 (LH216) X (LH119(4) X (T14))	SL	0.36	0.39	1 (30 cm)	97	NQ (3)	NQ (3)	NQ (3)	ER-93-USA- 01-MO-01	
USA (MO) 1993 (LH216) X (LH119(4) X (T14))	SL	0.36 +0.50	0.39 +0.54	2 (60 cm)	95	<u>NQ</u> (3)	NQ (3)	NQ (3)	ER-93-USA- 01-MO-01	
USA (CA) 1993 (LH59 X LH51) (LH119)(4) X T14)	SL	0.36	0.39	1 (30 cm)	119	NQ (3)	NQ (3)	NQ (3)	ER-93-USA- 01-CA-01	
USA (CA) 1993 (LH59 X LH51) (LH119)(4) X T14)	SL	0.36	0.39	1 (30 cm)	129	NQ (3)	NQ (3)	NQ (3)	ER-93-USA- 01-CA-01	
USA (CA) 1993 (LH59 X LH51) (LH119)(4) X T14)	SL	0.36 +0.50	0.39 +0.54	2 (60 cm)	106	<u>NQ</u> (3)	NQ (3)	NQ (3)	ER-93-USA- 01-CA-01	
USA (CA) 1993 (LH59 X LH51) (LH119)(4) X T14)	SL	0.36 +0.50	0.39 +0.54	2 (60 cm)	116	<u>NQ</u> (3)	NQ (3)	NQ (3)	ER-93-USA- 01-CA-01	

Country, Year,	Applic	ation			PHI	Residues, m acid ²	g/kg, as glu	ifosinate free	Ref
Variety	form	kg ai/ha	kg ai/hl	No ¹	days	glufosinate- ammonium	MPP	NAG	
USA (SD) 1993 (LH85 X LH160) (LH119)(4) X T14))	SL	0.36	0.71	1 (30 cm)	95	NQ (3)	NQ (3)	NQ (3)	ER-93-USA- 01-SD-01
USA (SD) 1993 (LH85 X LH160) (LH119)(4) X T14))	SL	0.50	0.99	1 (30 cm)	95	NQ (3)	NQ (3)	NQ (3)	ER-93-USA- 01-SD-01
USA (SD) 1993 (LH85 X LH160) (LH119)(4) X T14))	SL	0.36 +0.50	0.71 +0.99	2 (60 cm)	95 ¹	<u>NQ</u> (3)	NQ (3)	<u>0.07</u> (3)	ER-93-USA- 01-SD-01
USA (VA) 1993 (LH216) (LH119)(4) X T14)	SL	0.36	0.71	1 (30 cm)	106 ²	NQ	NQ	NQ	ER-93-USA- 01-VA-01
USA (VA) 1993 (LH216) (LH119)(4) X T14)	SL	0.50	0.99	1 (30 cm)	106 ⁶	NQ	NQ	NQ	ER-93-USA- 01-VA-01
USA (VA) 1993 (LH216) (LH119)(4) X T14)	SL	0.36 +0.50	0.71 +0.99	2 (60 cm)	97 ⁶	<u>NQ</u>	NQ	<u>0.07</u>	ER-93-USA- 01-VA-01
USA (MO) 1993 (LH59 X LH51) (LH119)(4) X (T14))	SL	0.50	0.53	1 (12-15 cm)	123	NQ (3)	NQ (3)	NQ (3)	ER-93-USA- 01-MO-02
USA (MO) 1993 (LH59 X LH51) (LH119)(4) X (T14))	SL	0.50	0.53	1 (25-30 cm)	118	NQ (3)	NQ (3)	NQ (3)	ER-93-USA- 01-MO-02
USA (MO) 1993 (LH59 X LH51) (LH119)(4) X (T14))	SL	0.50	0.53	1 (40-45 cm)	111	NQ (3)	NQ (3)	NQ (3)	ER-93-USA- 01-MO-02
USA (MO) 1993 (LH59 X LH51) (LH119)(4) X (T14))	SL	0.50	0.53	1 (55-66 cm)	101	<u>NQ</u> (3)	NQ (3)	NQ (3)	ER-93-USA- 01-MO-02
USA (MO) 1993 (LH59 X LH51) (LH119)(4) X (T14))	SL	0.50	0.53	1 (80-90 cm)	97	NQ (3)	NQ (3)	NQ (3)	ER-93-USA- 01-MO-02
USA (CA) 1993 (LH59 X LH51) (LH119)(4) X (T14))	SL	0.50	0.53	1 (12-15 cm)	127	NQ (3)	NQ (3)	NQ (3)	ER-93-USA- 01-CA-02
USA (CA) 1993 (LH59 X LH51) (LH119)(4) X (T14))	SL	0.50	0.53	1 (25-30 cm)	120	NQ (3)	NQ (3)	NQ (3)	ER-93-USA- 01-CA-02
USA (CA) 1993 (LH59 X LH51) (LH119)(4) X (T14))	SL	0.50	0.53	1 (40-45 cm)	114	NQ (3)	NQ (3)	NQ (3)	ER-93-USA- 01-CA-02
USA (CA) 1993 (LH59 X LH51) (LH119)(4) X (T14))	SL	0.50	0.53	1 (55-66 cm)	96	NQ (3)	NQ (3)	NQ (3)	ER-93-USA- 01-CA-02
USA (CA) 1993 (LH59 X LH51) (LH119)(4) X (T14))	SL	0.50	0.53	1 (80-90 cm)	96	<u>NQ</u> (3)	NQ (3)	NQ (3)	ER-93-USA- 01-CA-02
USA (FL) 1994 (LH51 X LH210) (LH119(4) X T14)	SL	0.40 +0.50	0.43 +0.54	2 (2-14 leaves)	69	NQ	NQ	glu	BK-94R-01- FL-01
USA (IA) 1994 (LH59 X LH51) (LH119(4) X T14))	SL	0.41 +0.53	0.44 +0.57	2 (60 cm)	122	NQ	NQ	glu	BK-94R-01- IA-01

¹ This crop suffered some phytotoxicity and the yield of grain was limited.

² Each of the treatments in the Virginia trials suffered from phytotoxicity and unusual weather conditions, resulting in limited collection of samples. CLICK HERE for continue

Country, Year,	Applica	ation			PHI	Residues, m	g/kg, as glu	ifosinate free	Ref
Variety	form	kg ai/ha	kg ai/hl	No ¹	days	glufosinate- ammonium	MPP	NAG	
USA (IL) 1994 (LH59 X LH51) (LH119(4) X T14))	SL	0.40 +0.50	0.43 +0.54	2 (60 cm)	118	<u>NQ</u>	NQ	glu	BK-94R-01- IL-01
USA (MN) 1994 (LH74) (LH82(3) X T14))	SL	0.38 +0.48	0.41 +0.52	2 (60 cm)	114	<u>0.05</u>	NQ	glu	BK-94R-01- MN-01
USA (MO) 1994 (LH59 X LH51) (LH119(4) X T14))	SL	0.40 +0.53	0.43 +0.57	2 (60 cm)	89	<u>NQ</u>	NQ	glu	BK-94R-01- MO-01
USA (NC) 1994 (LH51 X LH210) (LH119(4) X T14))	SL	0.38 +0.52	0.41 +0.56	2 (60 cm)	92	<u>NQ</u>	NQ	glu	BK-94R-01- NC-01
USA (NY) 1994 (LH74) (LH82(3) X T14))	SL	0.38 +0.53	0.41 +0.57	2 (60 cm)	88	<u>0.07</u>	NQ	glu	BK-94R-01- NY-01
USA (TX) 1994 (LH51 X LH210) (LH119(4) X T14))	SL	0.38 +0.46	0.41 +0.49	2 (60 cm)	97	<u>NQ</u>	NQ	glu	BK-94R-01- TX-01
USA (WA) 1994 (LH74) (LH82)3) X T14))	SL	0.40 +0.50	0.43 +0.54	2 (60 cm)	86	NQ	NQ	glu	BK-94R-01- WA-01
USA (WI) 1994 (LH85 X LH160) (LH119(4) X T14))	SL	0.38 +0.56	0.41 +0.60	2 (60 cm)	107	<u>NO</u>	NQ	glu	BK-94R-01- WI-01
USA (FL) 1994 (LH51 X LH210) (LH119(4) X T14)	SL200	0.40 +0.50	0.43 +0.54	2 (2-14 leaves)	69	NQ	NQ	glu	BK-94R-01- FL-01
USA (IA) 1994 (LH59 X LH51) (LH119(4) X T14))	SL	0.41 +0.53	0.44 +0.55	2 (60 cm)	122	NQ	NQ	glu	BK-94R-01- IA-01
USA (IL) 1994 (LH59 X LH51) (LH119(4) X T14))	SL	0.40 +0.50	0.43 +0.54	2 (60 cm)	118	NQ	NQ	glu	BK-94R-01- IL-01
USA (MN) 1994 (LH74) (LH82(3) X T14))	SL	0.40 +0.51	0.43 +0.55	2 (60 cm)	114	<u>0.07</u>	NQ	glu	BK-94R-01- MN-01
USA (MO) 1994 (LH59 X LH51) (LH119(4) X T14))	SL	0.40 +0.56	0.43 +0.60	2 (60 cm)	89	<u>0.05</u>	NQ	glu	BK-94R-01- MO-01
USA (NC) 1994 (LH51 X LH210) (LH119(4) X T14))	SL	0.40 +0.46	0.43 +0.49	2 (60 cm)	92	NQ	NQ	glu	BK-94R-01- NC-01
USA (NY) 1994 (LH74) (LH82(3) X T14))	SL	0.40 +0.53	0.43 +0.57	2 (60 cm)	88	<u>NQ</u>	NQ	glu	BK-94R-01- NY-01
USA (TX) 1994 (LH51 X LH210) (LH119(4) X T14))	SL	0.40 +0.49	0.43 +0.53	2 (60 cm)	97	<u>NQ</u>	NQ	glu	BK-94R-01- TX-01
USA (WA) 1994 (LH74) (LH82)3) X T14))	SL	0.40 +0.50	0.43 +0.54	2 (60 cm)	86	<u>NQ</u>	NQ	glu	BK-94R-01- WA-01
USA (WI) 1994 (LH85 X LH160) (LH119(4) X T14))	SL	0.38 +0.50	0.41 +0.54	2 (60 cm)	107	<u>NQ</u>	NQ	glu	BK-94R-01- WI-01
USA (MO) 1994 (LH59 X LH51) (LH119(4) X T14))	SL	2.0 +2.6	2.1 +2.5	2 (60 cm)	89	0.26 3.1 agf	0.06 0.75 agf	glu	BK-94R-03- MO-01

Country, Year,	Applic	cation	ation			Residues, m acid ²	Ref		
Variety	form	kg ai/ha	kg ai/hl	No ¹	days	glufosinate- ammonium	MPP	NAG	-
USA (MN) 1995 (LH202) (LH82 (4) x T25))	SL	0.39 +0.49	0.41 +0.53	2 (7-8 leaf)	100	NQ	NQ	glu	BK95R01- GGH-01
USA (KS) 1995 (B73(5)xT25) (LH51))	SL	0.39 +0.50	0.41 +0.53	2 (56-66 cm)	95	NQ	NQ	glu	BK95R01- MDA-02
USA (MI) 1995 (LH119 (9) x T14) (LH168))	SL	0.39 +0.50	0.41 +0.54	2 (61-77 cm)	111	NQ	NQ	glu	BK95R01- JRS-02
USA (MO) 1995 (B73 (5) x T25) (LH51))	SL	0.39 +0.52	0.44 +0.60	2 (60 cm)	90	<u>NQ</u>	NQ	glu	BK95R01- JLB-01
USA (NE) 1995 (LH119 (9) x T14) (LH51))	SL	0.39 +0.52	0.41 +0.55	2 (60 cm)	89	NQ	NQ	glu	BK95R01- MDA-01
USA (OH) 1995 (LH119 (9) x T14) (LH51))	SL	0.39 +0.52	0.43 +0.52	2 (60 cm)	117	NQ	NQ	glu	BK95R01- JRS-01
USA (TX) 1995 (LH216) (B73 (5) x T25))	SL	0.40 +0.50	0.42 +0.53	2 (53-66 cm)	84	NQ	NQ	glu	BK95R01- GLS-01
Argentina 1995)	SL	1.4	0.88	1	70	NQ	NQ		A57415 ER95ARG001 010102
Argentina 1996)	SL	1.4	0.88	1 GS17- 32)	104	NQ	NQ		A57414 ER96ARG001 010102

 1 Growth stage in the US trials is recorded as the height of the plant or, in some cases, as the number of leaves. 2 NQ - residues <LOD (0.05 mg/kg glufosinate free acid equivalents)

glu: included in glufosinate-ammonium result. agf: aspirated grain fraction from milling of the grain

Table 46. Glufosinate residues in transgenic maize resulting from supervised trials in Ontario, Canada. Double-underlined residues are from treatments according to GAP and were used to estimate maximum residue levels.

Year		Appli	cation		PHI	Re	sidues, mg/	kg	Ref
	form	kg ai/ha	kg ai/hl	no 1	days	glufosinate	MPP	NAG	
1993	SL	0.75		1 GS16	152	<0.05	<0.05	<0.05	A55207 93-IGN-C ON-1-2
1993	SL	1.5		1 GS16	152	<0.05	<0.05	<0.05	A55207 93-IGN-C ON-1-3
1993	SL	0.75		1 GS16	154	<0.05	<0.05	<0.05	A55207 93-IGN-C ON-2-2
1993	SL	1.5		1 GS16	154	<0.05	<0.05	<0.05	A55207 93-IGN-C ON-2-3
1994	SL	0.5		1 GS17-19	150	< <u>0.05</u>	<0.05	<0.05	A55207 94-IGN-C ON-1-2
1994	SL	1.0		1 GS17-19	150	<0.05	<0.05	<0.05	A55207 94-IGN-C ON-1-3

Year Application				PHI	Re	sidues, mg/	kg	Ref	
	form	kg ai/ha	kg ai/hl	no 1	days	glufosinate	MPP	NAG	
1994	SL	0.5		1 GS17-19	136	< <u>0.05</u>	<0.05	<0.05	A55207 94-IGN-C ON-2-2
1994	SL	1.0		1 GS17-19	136	< 0.05	<0.05	<0.05	A55207 94-IGN-C ON-2-3
1994	SL	0.5		1 GS17-19	136	< <u>0.05</u>	<0.05	<0.05	A55207 94-IGN-C ON-3-2
1994	SL	1.0		1 GS17-19	136	<0.05	<0.05	<0.05	A55207 94-IGN-C ON-3-3
1995	SL	0.6		1 GS18-19	138	< <u>0.05</u>	<0.05	<0.05	A55207 95-IGN-C ON-1-2
1995	SL	0.6		2 GS33	131	<0.05	<0.05	<0.05	A55207 95-IGN-C ON-1-3
1995	SL	1.2		1 GS40	126	<0.05	<0.05	<0.05	A55207 95-IGN-C ON-1-4
1995	SL	0.6		1 GS18	137	< <u>0.05</u>	<0.05	<0.05	A55207 95-IGN-C ON-2-2
1995	SL	0.6		2 GS33	128	<0.05	<0.05	<0.05	A55207 95-IGN-C ON-2-3
1995	SL	1.2		1 GS40	122	<0.05	<0.05	0.08	A55207 95-IGN-C ON-2-4
1995	SL	0.6		1 GS18-19	114	< <u>0.05</u>	<0.05	<0.05	A55207 95-IGN-C ON-3-2
1995	SL	0.6		2 GS33	107	<0.05	<0.05	<0.05	A55207 95-IGN-C ON-3-3
1995	SL	1.2		1 GS40	100	<0.05	<0.05	0.07	A55207 95-IGN-C ON-3-4
1995	SL	0.6		1 GS18	125	< <u>0.05</u>	<0.05	<0.05	A55207 95-IGN-C ON-4-2
1995	SL	0.6		2 GS33	117	<0.05	<0.05	<0.05	A55207 95-IGN-C ON-4-3
1995	SL	1.2		1 GS40	111	<0.05	<0.05	0.09	A55207 95-IGN-C ON-4-4

¹. GS: growth stage at final treatment GS16 6 leaves unfolded GS17 7 leaves unfolded

- 8 leaves unfolded
- GS18 GS19 9 or more leaves unfolded
- GS33 3 nodes detectable
- GS40 booting

Table 47. Glufosinate residues in soya beans resulting from supervised trials in the USA. Analyses of replicate samples from the treated plots are shown separately. Double-underlined residues are from treatments according to GAP and were used to estimate maximum residue levels.

Country,		App	lication		PHI	Residues, n	ng/kg, as g	lufosinate	Ref
(variety)	form	kg ai/ha	kg ai/hl	No ¹	days	glufosinate	MPP	NAG	
		0 -	0.71		~ ~	0.07 (0)	0.07 (0)	0.04.007	
USA (IA) 1993	SL	0.5	0.51	l trifaliata	85	< 0.05 (3)	<0.05 (3)	0.06 < 0.05	ER-93-USA-02-
(w98-7)	SI	0.5	0.51		65	12(3)	0 44 0 44	(2)	IA-01
	SL	0.5	0.51	full pod	05	1.2 (5)	0.39	5.0 5.4 4.7	
	SL	0.5	0.51	2	65	0.76 0.91	0.30 0.30	3.4 3.8 3.6	
				full pod		0.79	0.27		
USA (IL) 1993	SL	0.5	0.53	1	85	< 0.05 (3)	0.07 0.06	0.33 0.36	ER-93-USA-02-
(W98-7)	CI	0.5	0.52	trifoliate	65	0.06.1.0.000	0.06	0.32	IL-01
	SL	0.5	0.55	full pod	65	0.96 1.0 0.90	0.51 0.61	5.0 5.4 5.1	
	SL	0.5	0.53	2	65	0.95 0.95	0.62 0.68	5.1 5.5 5.4	
	~			full pod		0.93	0.65		
USA (MO) 1993	SL	0.5	0.53	Î	85	< 0.05 (3)	0.08 0.09	0.19 0.21	ER-93-USA-02-
(W98-7)				trifoliate			0.08	0.22	MO-01
	SL	0.5	0.53	1	65	1.4 (3)	0.89 0.61	6.7 6.7 7.2	
	CI	0.5	0.52	full pod	65	121011	0.79	505050	
	SL	0.5	0.55	2 full pod	65	1.2 1.0 1.1	0.74 0.75	5.9 5.8 5.9	
USA (IN) 1993	SL.	0.5	0.53	1 1	85	<0.05 (3)	0.08	0 12 0 10	ER-93-USA-02-
(W98-7)	SL	0.5	0.55	trifoliate	05	(0.05 (5)	< 0.05 (2)	0.08	IN-01
	SL	0.5	0.53	1	65	1.7 1.8 1.9	0.49 0.53	7.1 7.1 7.5	
				full pod			0.55		
	SL	0.5	0.53	2	65	1.6 1.8 1.8	0.56 0.64	5.9 6.1 7.1	
101 BD 1002	CT.	0.5	0.52	full pod	07	0.05 (0)	0.74	0.05 (0)	
USA (IN) 1993	SL	0.5	0.53	l trifoliata	85	<0.05 (3)	<0.05 (3)	<0.05 (3)	ER-93-USA-02-
(1198)	SL	0.5	0.53	1	65	0.93.0.91	0 37 0 33	383940	IIN-02
	5L	0.5	0.55	full pod	00	0.94	0.35	5.0 5.9 1.0	
	SL	0.5	0.53	1	75	0.80 0.84	0.29 0.32	3.1 3.6 3.7	
				full pod		0.81	0.36		
	SL	0.5	0.53	2	65	0.92 0.87	0.38 0.31	3.9 3.6 4.0	
	CI	0.5	0.52	full pod	75	0.88	0.35	212521	
	SL	0.5	0.55	full pod	15	0.68	0.30	5.4 5.5 5.4	
USA (AR) 1993	SL	0.5	1.1	1	85	< 0.05 (3)	< 0.05 (3)	0.07 0.09	ER-93-USA-02-
(W62)				trifoliate				0.08	AR-01
	SL	0.5	1.1	1	65	0.87 0.76	0.61 0.56	6.8 5.8 4.5	
				full pod		0.73	0.46		
	SL	0.5	1.1	2 full pod	65	0.89 0.88	0.58 0.49	7.6 6.6 7.0	
USA (MS) 1003	SI	0.5	1 1		85	0.81	0.38	0 10 0 10	
(W98)	SL	0.5	1.1	trifoliate	85	<0.05 (3)	<0.05 (3)	0.12	MS-01
(SL	0.5	1.1	1	65	0.47 0.52	0.44 0.41	5.3 4.9 4.7	
				full pod		0.48	0.27		
	SL	0.5	1.1	2	65	0.46 0.40	0.48 0.41	5.1 5.1 5.0	
	CT.	0.7	1.1	full pod	102	0.48	0.47	0.11	
USA (VA) 1993 (W62)	SL	0.5	1.1	l trifoliota	103	<0.05	<0.05	0.11	EK-95-USA-02-
(**02)	SL.	0.5	1.1	1	65	0.10	0.10	0.69	VA-01
	SL		1	full pod	55	0.10	0.10	0.07	
	SL	0.5	1.1	2	85	0.08	0.15	0.93	
				full pod					
USA (AR) 1994	SL	0.39	0.41	2	82	0.68 <u>0.83</u>	0.06 <u>0.06</u>	glu	BK-94R-02-AR-
(W62)		+0.50	+0.53	bloom					01

Country, Year		Appl	lication		PHI	Residues, n	ufosinate	Ref	
(variety)	form	kg ai/ha	kg ai/hl	No ¹	days	glufosinate	MPP	NAG	
USA (FL) 1994 (W62)	SL	0.39 +0.50	0.42 +0.54	2 bloom	93	0.58 <u>0.62</u>	0.09 <u>0.09</u>	glu	BK-94R-02-FL- 01
USA (IA) 1994 (W98)	SL	0.39 +0.50	0.42 +0.54	2 bloom	86	<u>1.1</u> 1.1	<u>0.14</u> 0.10	glu	BK-94R-02-IA-01
USA (IL) 1994 (transgenic group 3)	SL	0.39 +0.50	0.42 +0.54	2 bloom	69	<u>0.65</u> 0.52	<u>0.07</u> 0.06	glu	BK-94R-02-IL-01
USA (IN) 1994 (W98)	SL	0.39 +0.50	0.42 +0.54	2 bloom	77	<u>1.3</u> 0.89	<u>0.34</u> 0.15	glu	BK-94R-02-IN-01
USA (MO) 1994 (W98)	SL	0.39 +0.50	0.42 +0.54	2 bloom	77	<u>0.67</u> 0.41	<u>0.18</u> 0.17	glu	BK-94R-02-MO- 01
USA (NC) 1994 (W62)	SL	0.39 +0.50	0.42 +0.54	2 bloom	102	0.32 <u>0.43</u>	< 0.05 (2)	glu	BK-94R-02-NC- 01
USA (OH) 1994 (W98)	SL	0.39 +0.50	0.42 +0.54	2 bloom	76	<u>0.80</u> 0.71	<u>0.22</u> 0.18	glu	BK-94R-02-OH- 01
USA (PA) 1994 (W98)	SL	0.39 +0.50	0.42 +0.54	2 bloom	91	<u>0.73</u> 0.54	<u>0.08</u> 0.07	glu	BK-94R-02-PA- 01
USA (VA) 1994 (W62)	SL	0.39 +0.50	0.42 +0.54	2 bloom	82	<u>1.6</u> 1.3	<u>0.28</u> 0.36	glu	BK-94R-02-VA- 01
USA (NC) 1994 (transgenic Group 5 W62 A5403)	SL	0.4	0.43	1 3 rd node	136	<0.05 (2)	<0.05 (2)	glu	Project: BK-94R- 06 Trial#: NC-01
,	SL	0.4 +0.5	0.43 +0.54	2 6 th node	123	<0.05 (2)	<0.05 (2)	glu	
	SL	0.4 +0.5	0.43 +0.54	2 bloom	90	<u>0.79</u> 0.71	<u>0.13</u> 0.14	glu	
	SL	0.4 +0.5	0.43 +0.54	2 full pod	77	1.1 1.1	0.23 0.27	glu	
USA (IN) 1995 (MG III, W98 A3322 group 3)	SL	0.39 +0.50	0.39 +0.51	2 R2.5-R3	64	1.1 0.96	0.16 0.19	glu	BK-95R-04-DJL- 01
USA (IN) 1995 (MG III, W98 A3322 group 3)	SL	0.39 +0.50	0.39 +0.51	2 R2	62	<u>1.1</u> 0.90	<u>0.23</u> 0.24	glu	BK-95R-04-DJL- 02
USA (MO) 1995 (MG III W62 A5504 group 5)	SL	0.39 +0.50	0.41 +0.53	2 R1	71	0.62 <u>0.83</u>	0.11 <u>0.13</u>	glu	BK-95R-04-JLB- 01
USA (OH) 1995 (MG III, W98 A3322 group 3)	SL	0.39 +0.50	0.40 +0.52	2 R2	86	<u>0.32</u> 0.18	<u>0.07</u> 0.06	glu	BK-95R-04-JRS- 01
USA (MI) 1995 (MG III, W98 A3322 group 3)	SL	0.39 +0.50	0.41 +0.53	2 R2	81	<u>0.37</u> 0.33	<u>0.15</u> 0.10	glu	BK-95R-04-JRS- 02
USA (IA) 1995 (MG III, W98 A3322 group 3)	SL	0.39 +0.50	0.41 +0.53	2 R2	60	0.58 <u>0.68</u>	0.10 <u>0.10</u>	glu	BK-95R-04- GGH-01
USA (WI) 1995 (MG III, W98 A3322 group 3)	SL	0.38 +0.49	0.42 +0.54	2 R2.5	82	0.65 0.93	0.10 0.18	glu	BK-95R-04- GGH-02
USA (IA) 1995 (MG III, W98 A3322 group 3)	SL	0.39 +0.50	0.42 +0.53	2 R2	84	<u>0.32</u> 0.28	<0.05 (2)	glu	BK-95R-04- GGH-04
USA (NE) 1995 (MG III, W98 A3322 group 3)	SL	0.39 +0.50	0.41 +0.53	2 R3	80	0.12 0.62	0.05 0.06	glu	BK-95R-04- MDA-01
USA (KS) 1995 (MG III, W98 A3322 group 3)	SL	0.39 +0.50	0.41 +0.52	2 R2	78	<u>0.50</u> 0.42	<u>0.06</u> 0.05	glu	BK-95R-04- MDA-02

Country, Year		Appl	lication		PHI	Residues, mg/kg, as glufosinate free acid			Ref
(variety)	form	kg ai/ha	kg ai/hl	No ¹	days	glufosinate	MPP	NAG	
USA (LO) 1995 (MG III, W62 A5504 group 5)	SL	0.40 +0.54	0.42 +0.56	2 R2	70	<u>0.95</u> 0.64	<u>0.61</u> 0.84	glu	BK-95R-04-TLS- 01
USA (MS) 1995 (MG III, W62 A5504 group 5)	SL	0.39 +0.50	0.41 +0.53	2 R2	90	<u>1.1</u> 0.93	<u>0.16</u> 0.15	glu	BK-95R-04-TLS- 02
USA (IA) 1993 (transgenic W98- 7)	SL	2.5	2.7	1 trifoliate	95	<0.05	0.08	0.07	ER-93-USA-02- IA-02
USA (IN) 1994 (transgenic group 3 W98 A3322)	SL	2.0 +2.6	2.1 +2.7	2 bloom	77	3.4	1.1	glu	BK-94R-04-IN-01
USA (MO) 1994 (transgenic group 3 W98 A3322)	SL	2.0 +2.6	2.1 +2.7	2 bloom	77	1.5	1.8	glu	BK-94R-04-MO- 01

¹growth stage described or provided as "R" code

R1 beginning bloom

R2 full bloom

R3 beginning pod glu: included in glufosinate-ammonium result.

Table 48. Glufosinate residues in rape seed resulting from supervised trials in France, Germany and the UK.

Country	Applic	ation			PHI	Residues	, mg/kg, exp	ressed as	Ref
Voor (vorioty)	form	ka si/ba	Ka ai/bl	No^1	dava	glufosinato	MDD	NAG^2	ł
Teal (vallety)	101111			INO O	111	giulosiliate	INIF F	NAU	4 5 4000
France 1994	SL	0.60	0.30	2	111	0.06	<0.05		A54228
(Falcon)		+0.45	+0.23	G\$35					ER93ECN553
									FRA000202
France 1994	SL	0.60	0.30	2	111	< 0.05	< 0.05		A54228
(Falcon)		+0.45	+0.23	GS35					ER93ECN553
									FRA000302
France 1994 (Gs	SL	0.60	0.30	2	121	0.07	< 0.05		A54228
401×90)		+0.45	+0.23	GS33					ER93ECN553
/									FRA000102
France 1995	SL	0.6	0.30	2	127	< 0.05	< 0.05		A56444
(Falcon B)				GS35					ER94ECN553
									FRA000203
France 1995	SL	0.67	0.32	2	133	< 0.05	< 0.05		A56444
(Falcon)	~ _	+0.60	+0.30	GS39					ER94ECN553
(1 410011)		10.00	10120	0.007					FRA000103
France 1996	SL	0.60	0.30	2	108	0.13	<0.05		A57955
(GMO hybrid)	52	0.00	+0.24	GS32	100	0110	(0100		ER96ECN553
· · · ·									FRA000103
France 1996	SL	0.60	0.30	2	108	0.12	< 0.05		A57955
(GMO hybrid)			+0.24	GS32					ER96ECN553
()									FRA000102

Country	Applic	ation			PHI	Residues	, mg/kg, exp	ressed as	Ref
Year (variety)	form	kg ai/ha	Kg ai/hl	No ¹	days	glufosinate	MPP	NAG ²	1
Germany 1995	SL	0.60	0.20	2	106	0.06	< 0.05		A56444
(Falcon G-S40/90)				GS35					ER94ECN553
									DEU030103
Germany 1995	SL	0.60	0.20	2	105	0.06	< 0.05		A56444
(Falcon)				GS35					ER94ECN553
									DEU040103
Germany 1995	SL	0.60	0.20	2	110	0.05	< 0.05		A56444
(Falcon)				GS35					ER94ECN553
G 1005	CT.	0.50	0.00		104	0.05	0.05		DEU040203
Germany 1995	SL	0.60	0.20	2	104	<0.05	<0.05		A56444
(Falcon)				G237					EK94ECN555
Cormony 1006	CI.	0.60	0.20	2	04	<0.05	<0.05		DE0050105
(Falcon)	SL	0.00	0.20	G\$32	94	<0.03	<0.05		A37933 FR96ECN553
(1 alcoll)				0552					DFU030102
Germany 1996	SL.	0.80	0.27	2	94	0.06	< 0.05		A57955
(Falcon)	52	0.00	0.27	GS32		0100	0102		ER96ECN553
(,									DEU030103
Germany 1996	SL	0.60	0.20	2	97	0.11	< 0.05		A57955
(Falcon)				GS32					ER96ECN553
									DEU050102
Germany 1996	SL	0.80	0.27	2	97	0.13	< 0.05		A57955
(Falcon)				GS32					ER96ECN553
G 1006	CT	0.60	0.00	2	07	0.14	0.05		DEU050103
Germany 1996	SL	0.60	0.20	2	97	0.14	<0.05		A5/955 EDOCECN552
(Falcon)				0337					EK90ECN555
Germany 1006	SI	0.80	0.27	2	07	0.12	<0.05		DE0040102 457955
s(Falcon)	SL	0.80	0.27	GS37)/	0.12	<0.05		FR96FCN553
s(r ulcoll)				0007					DEU040103
UK 1994	SL	0.60	0.30	1	275	< 0.05	< 0.05		A54227
				GS21					ER93ECN552
									GBR000102
UK 1994	SL	0.60	0.30	1	212	< 0.05 (2)	< 0.05 (2)		A54227
				GS21					ER93ECN552
THE 100.4	CT.	0.50	0.00	•	01	0.05	0.05		GBR000202
UK 1994	SL	0.60	0.30	2	91	<0.05	<0.05		A54228
		+0.45	+0.23	G\$35					ER93ECN553
LIV 1004 (Ealaon)	CI	0.60	0.20	2	116	<0.05	<0.05		GBR000202
UK 1994 (Faicoli)	SL	0.00	0.50	ے 32	110	<0.03	<0.05		A34220 ED03ECN553
		+0.45	+0.25	0333					GBR000102
UK 1996 (Falcon)	SL.	0.60	0.30	2	100	0.07	< 0.05		A57955
	52	0.00	0100	GS32	100	0107	10102		ER96ECN553
									GBR000103
UK 1996 (Falcon)	SL	0.60	0.30	2	92	0.1^{-3}	$< 0.05 \frac{3}{2}$		A57955
				GS32					ER96ECN553
									GBR000203
UK 1996 (Falcon)	SL	0.60	0.30	2	100	0.15	< 0.05		A57955
				GS32					ER96ECN553
UK 1006 (Eslage)	CI	0.80	0.40	2	100	0.07	<0.05		GBK000102
UK 1990 (Faicon)	പ	0.00	0.40	6532	100	0.07	<0.05		FR96ECN553
				0002					GBR000104
UK 1996 (Falcon)	SL	0.60	0.30	2	92	$0.1^{-\frac{3}{2}}$	< 0.05 3		A57955
				GS32					ER96ECN553
									GBR000202
UK 1996 (Falcon)	SL	0.80	0.40	2	92	0.12^{-3}	$< 0.05^{\frac{3}{2}}$		A57955
				GS32					ER96ECN553
									GBR000204

¹ : GS growth stage	
GS21	first side shoot detectable
GS32	2 visibly extended internodes
GS33	3 visibly extended internodes
GS35	5 visibly extended internodes
GS37	7 visibly extended internodes
GS80	beginning of ripening
GS83	30% of pods ripe
² included in glufosinate-am	monium result.

 $\frac{3}{2}$: Grain samples were harvested at GS80-83 and stored in paper bags for even drying and prepared to laboratory samples 10 days later.

Table 49. Glufosinate residues in canola resulting from supervised trials in Canada. Doubleunderlined residues are from treatments according to GAP and were used to estimate maximum residue levels.

		Appli	cation		PHI	Residues, mg/kg ²			Ref
Location,	form	kg ai/ha	kg ai/hl	No ¹	days	glufosinate	MPP	NAG	
Year, (variety)									
Canada (AB)	SL	0.75		1	63	< 0.05	< 0.05	< 0.05	XEN93-09-AB-
1992 (transgenic									018
HCN92)									A53770
Canada (AB)	SL	0.75		1	63	< 0.05	< 0.05	< 0.05	XEN93-09-
1992 (transgenic									AB019
HCN92)									A53770
Canada (AB)	SL	0.75		1	65	< 0.05	< 0.05	< 0.05	XEN93-09-
1992 (transgenic									AB020
HCN92)									A53770
Canada (SK)	SL	0.75		1	65	< 0.05	< 0.05	0.07	XEN93-09-
1992 (transgenic									SK024
HCN92)									A53770
Canada (SK)	SL	0.75		1	62	< 0.05	< 0.05	0.14	XEN93-09-
1992 (transgenic									SK026
HCN92)									A53770
Canada (MB)	SL	0.75		1	65	< 0.05 (2)	< 0.05 (2)	< 0.05 (2)	XEN93-09-
1992 (transgenic									MB015
HCN92)									A53770
Canada (MB)	SL	0.75		1	59	< 0.05	< 0.05	< 0.05	XEN93-09-
1992 (transgenic									MB016
HCN92)									A53770
Canada (MB)	SL	0.75		1	65	< 0.05	< 0.05	< 0.05	XEN93-09-
1992									MB022
									A53770
Canada (MB)	SL	0.75		1	65	< 0.05	< 0.05	0.08	XEN93-09-
1992									MB028
									A53770
Canada (AB)	SL	0.83		1	95	< 0.05	< 0.05	< 0.05	XEN93-29-AB-
1993 (transgenic				GS 3-6					TC-5
HCN92)				leaf					A56394
Canada (AB)	SL	1.5		1	95	< 0.05	< 0.05	< 0.05	XEN93-29-AB-
1993 (transgenic				GS 3-6					TC-5
HCN92)				leaf					A56394
Canada (AB)	SL	0.75	0.68	1	80	< 0.05	< 0.05	< 0.05	XEN93-29-AB-
1993 (transgenic				GS 3-5					TC-1
HCN92)				leaf					A56394
Canada (AB)	SL	1.5	1.36	1	80	< 0.05	< 0.05	< 0.05	XEN93-29-AB-
1993 (transgenic									TC-1
HCN92)									A56394
Canada (AB)	SL	0.5	0.45	1	75	< 0.05	< 0.05	< 0.05	XEN93-29-AB-
1993 (transgenic									TC-2
HCN92)									A56394

	Application				PHI	Res	Ref		
Location, Year, (variety)	form	kg ai/ha	kg ai/hl	No ¹	days	glufosinate	MPP	NAG	
Canada (AB) 1993 (transgenic HCN92)	SL	0.75	0.68	1	75	<0.05	<0.05	< 0.05	XEN93-29-AB- TC-2 A56394
Canada (AB) 1993 (transgenic)	SL	0.5	0.45	1 GS 4-5 leaf	75	<0.05	< 0.05	< 0.05	XEN93-29-AB- TC-3 A56394
Canada (AB) 1993 (transgenic)	SL	1.5		1 GS 4-5 leaf	75	<0.05	< 0.05	< 0.05	XEN93-29-AB- TC-3 A56394
Canada (AB) 1993 (transgenic HCN92)	SL	0.5	0.45	1 GS 3-5 leaf	83	<0.05	<0.05	<0.05	XEN93-29-AB- TC-4 A56394
Canada (AB) 1993 (transgenic HCN92)	SL	0.75	0.68	1 GS 3-5 leaf	83	<0.05	<0.05	<0.05	XEN93-29-AB- TC-4 A56394
Canada (MB) 1993 (transgenic HCN92)	SL	0.75	0.68	1 GS 4-6 leaf	69	0.12	<0.05	<0.05	XEN93-29-MB- TC-2 A56394
Canada (MB) 1993 (transgenic HCN92)	SL	0.5	0.45	1 GS 4-5 leaf	67	<0.05	<0.05	<0.05	XEN93-29-MB- TC-3 A56394
Canada (MB) 1993 (transgenic HCN92)	SL	0.75	0.68	1 GS 4-5 leaf	67	<0.05	<0.05	<0.05	XEN93-29-MB- TC-3 A56394
Canada (MB) 1993 (transgenic HCN92)	SL	0.5	0.45	1 GS 4-5 leaf	68	<0.05	<0.05	< 0.05	XEN93-29-MB- TC-4 A56394
Canada (MB) 1993 (transgenic HCN92)	SL	0.75	0.68	1 GS 4-5 leaf	68	<0.05	<0.05	< 0.05	XEN93-29-MB- TC-4 A56394
Canada (SK) 1993 (transgenic HCN92)	SL	1.0		1 GS 5 leaf	66	<0.05	<0.05	0.05	XEN93-29-SK- TC-1 A56394
Canada (SK) 1993 (transgenic HCN92)	SL	2.0		1 GS 5 leaf	66	<0.05	<0.05	0.10	XEN93-29-SK- TC-1 A56394
Canada (SK) 1993 (transgenic HCN92)	SL	0.5	0.45	1 GS 3-4 leaf	57	<0.05	< 0.05	<0.05	XEN93-29-SK- TC-3 A56394
Canada (SK) 1993 (transgenic HCN92)	SL	0.75	0.68	1 GS 3-4 leaf	57	<0.05	< 0.05	<0.05	XEN93-29-SK- TC-3 A56394
Canada (SK) 1993 (transgenic HCN92)	SL	0.5	0.45	1 GS 10 leaf	69	< <u>0.05</u>	<0.05	<0.05	XEN93-29-SK- TC-6 A56394
Canada (SK) 1993 (transgenic HCN92)	SL	0.75	0.68	1 GS 10 leaf	69	< <u>0.05</u>	<0.05	<0.05	XEN93-29-SK- TC-6 A56394
Canada (MT) 1994 (transgenic HCN92)	SL	0.4	0.36	1 GS 2 leaf	77	<0.05	<0.05	<0.05	A56392 Minto
Canada (MT) 1994 (transgenic HCN92)	SL	0.8	0.73	1 GS 2 leaf	77	<0.05	<0.05	<0.05	A56392 Minto
Canada (MT) 1994 (transgenic HCN92)	SL	1.2	1.1	1 GS 2 leaf	77	<0.05	<0.05	<0.05	A56392 Minto
Canada (MT) 1994 (transgenic HCN92)	SL	0.4	0.36	1 GS 5-6 leaf	70	<0.05	<0.05	<0.05	A56392 Minto

	Application				PHI	Residues, mg/kg ²			Ref
Location, Year, (variety)	form	kg ai/ha	kg ai/hl	No ¹	days	glufosinate	MPP	NAG	
Canada (MT) 1994 (transgenic HCN92)	SL	0.8	0.73	1 GS 5-6 leaf	70	<0.05	<0.05	<0.05	A56392 Minto
Canada (MT) 1994 (transgenic HCN92)	SL	1.2	1.1	1 GS 5-6 leaf	70	< 0.05	<0.05	0.06	A56392 Minto
Canada (SK) 1994 (transgenic HCN92)	SL	0.4	0.36	1 GS 2-3 leaf	73	<0.05	<0.05	<0.05	A56392 Indian Head
Canada (SK) 1994 (transgenic HCN92)	SL	0.8	0.73	1 GS 2-3 leaf	73	< 0.05	<0.05	<0.05	A56392 Indian Head
Canada (SK) 1994 (transgenic HCN92)	SL	1.2	1.1	1 GS 2-3 leaf	73	< 0.05	<0.05	<0.05	A56392 Indian Head
Canada (SK) 1994 (transgenic HCN92)	SL	0.4	0.36	1 GS 5-7 leaf	57	<0.05	<0.05	0.17	A56392 Indian Head
Canada (SK) 1994 (transgenic HCN92)	SL	0.8	0.73	1 GS 5-7 leaf	57	<0.05	<0.05	0.24	A56392 Indian Head
Canada (SK) 1994 (transgenic HCN92)	SL	1.2	1.1	1 GS 5-7 leaf	57	<0.05	<0.05	0.26	A56392 Indian Head
Canada (MT) 1994 (transgenic HCN92)	SL	0.4	0.36	1 GS 4-5 leaf	65	<0.05	<0.05	<0.05	A56392 Portage la Prairie
Canada (MT) 1994 (transgenic HCN92)	SL	0.8	0.73	1 GS 4-5 leaf	65	<0.05	<0.05	0.07	A56392 Portage la Prairie
Canada (MT) 1994 (transgenic HCN92)	SL	1.2	1.1	1 GS 4-5 leaf	65	<0.05	0.06	0.05	A56392 Portage la Prairie
Canada (AB) 1994 (transgenic HCN92)	SL	0.4	0.36	1 GS 2-4 leaf	77	<0.05	<0.05	<0.05	A56392 Vauxhall
Canada (AB) 1994 (transgenic HCN92)	SL	0.8	0.73	1 GS 2-4 leaf	77	< 0.05	<0.05	<0.05	A56392 Vauxhall
Canada (AB) 1994 (transgenic HCN92)	SL	1.2	1.1	1 GS 2-4 leaf	77	< 0.05	<0.05	<0.05	A56392 Vauxhall
Canada (AB) 1994 (transgenic HCN92)	SL	0.4	0.36	1 GS 4-6 leaf	67	<0.05	<0.05	0.08	A56392 Vauxhall
Canada (AB) 1994 (transgenic HCN92)	SL	0.8	0.73	1 GS 4-6 leaf	67	<0.05	<0.05	0.17	A56392 Vauxhall
Canada (AB) 1994 (transgenic HCN92)	SL	1.2	1.1	1 GS 4-6 leaf	67	0.05	< 0.05	0.24	A56392 Vauxhall

 1 GS: growth stage expressed in terms of leaf development. 2 The expression of residues is unclear. In reports A56392 and A56394 glufosinate residues may be expressed as free acid or ammonium salt. MPP and NAG may be expressed as themselves rather than as glufosinate free acid.

Table 50. Glufosinate residues in canola resulting from supervised trials in Australia.

		Арр	lication		PHI	Residues, mg/kg, as glufosinate free acid			Ref
Location,	form	kg ai/ha	kg ai/hl	No ¹	days	glufosinate	MPP^2	NAG ²	
Year,									
(variety)									
Australia (NSW)	SL	0.6	0.72	1	86	< 0.05 (2)			YH 96 AH
1996 (glufosinate				GS 3-5 leaf					AU NW 06
selective line)									
Australia (NSW)	SL	1.2	1.4	1	86	< 0.05 (2)			YH 96 AH
1996 (glufosinate				GS 3-5 leaf					AU NW 06
selective line)									
Australia (WA)	SL	0.6	0.83	1	124	<0.05 (2)			YH 96 AH
1996 (var ex				GS 1-2 leaf					AU WA 09
Canada)									
Australia (WA)	SL	1.2	1.7	1	124	<0.05 (2)			YH 96 AH
1996 (var ex				GS 1-2 leaf					AU WA 09
Canada)									
Australia (SA) 1996	SL	0.6	0.75	1	95	<0.05 (2)			YH 96 AH
(transgenic				GS 4-6 leaf					AU SA 17
Canadian line)									
Australia (SA) 1996	SL	1.2	1.5	1	95	< 0.05 (2)			YH 96 AH
(transgenic				GS 4-6 leaf					AU SA 17
Canadian line)									
Australia (Vic) 1997	SL	0.6	0.6	1	105	< 0.05 (2)			YH 96 AH
(Canadian tolerant)				GS 2 leaf					AU VN 12
Australia (Vic) 1997	SL	1.2	1.2	1	105	<0.05 (2)			YH 96 AH
(Canadian tolerant)				GS 2 leaf					AU VN 12

¹GS: growth stage expressed in terms of leaf development. ²included in glufosinate-ammonium result.

Table 51. Glufosinate residues in tolerant sugar beet root resulting from foliar application in supervised trials in France, Germany and the UK.

Country y (variety	year		Appl	lication		PHI	Residues, mg/	kg, as glufe acid	osinate free	Ref
	, ,	form	kg ai/ha	kg ai/hl	no ¹	days	glufosinate	MPP	NAG	
France (transgenic hybrid)	1995	SL	0.60	0.30	2 GS31	85	0.87	<0.05	2	A56446 ER95ECN555 FRA000102
France (transgenic hybrid)	1996	SL	0.80	0.32	2 GS19	96	0.05	<0.05	2	A57574 ER96ECN551 FRA000102
France (transgenic hybrid)	1996	SL	0.80	0.32	2 GS19	96 124	0.12 0.06	<0.05 <0.05	2	A57574 ER96ECN551 FRA000202
France (transgenic hybrid)	1996	SL	0.80	0.32	2 GS19	91	0.48	<0.05	2	A57574 ER96ECN551 FRA000302
Germany (Sä-Nr. 951 36)	1995 959 –	SL	0.80	0.27	2 GS31	96 116	0.39 0.3	<0.05 0.05	2 2	A57574 ER96ECN551 DEU010102
UK (transgenic hybrid)	1995	SL	0.60	0.30	2 GS19	83	0.79	<0.05	2	A56446 ER95ECN555 GBR000102
UK (transgenic hybrid)	1995	SL	0.60	0.60	2 GS31	94	0.32	<0.05	2	A56446 ER95ECN555 GBR000202

Country y (variety	year)		Appl	ication		PHI	Residues, mg	/kg, as glufo acid	osinate free	Ref
		form	kg ai/ha	kg ai/hl	no ¹	days	glufosinate	MPP	NAG	
UK (Roberta)	1996	SL	0.80	0.80	2 GS31	111	0.88 0.99	0.06 <0.05	2	A57574 ER96ECN551 GBR000102
UK (transgenic hybrid)	1996	SL	0.80	0.80	2 GS19	105	0.52	<0.05	2	A57574 ER96ECN551 GBR000202

¹: GS growth stage GS14 GS15

4 leaves (2nd pair unfolded) 5 leaves unfolded

GS19

9 or more leaves unfolded

GS31 beginning of crop cover formation ² included in glufosinate-ammonium result.

Table 52. Glufosinate residues in tolerant sugar beet root resulting from foliar application in supervised trials in the USA.

Location,		Appl	lication		PHI Residues, mg/kg, as glufosinate fre			sinate free	Ref
Year,	£	1	1-2-2:4-1	N ₂ 1	1	alufacianta	acid	NAC^2	
(variety)	SI			2	120	$\leq 0.05(2)$	$\frac{\text{WIPP}}{0.14(2)}$	NAG	DV 05D 06
(Liberty Link)	SL	10.21	0.21	S 8 leaf	139	<0.03 (2)	0.14 (2)		DK-95K-00
(Liberty Link)		+0.20	+0.20 +0.20	0-leal					(Regimen B)
USA (CA) 1995	SL	0.43	0.43	3	139	<0.05 (2)	0 30 0 32		BK-95R-06
(Liberty Link)	51	+0.40	+0.39	6-leaf	107	(0.05 (2)	0.50 0.52		CA-016
(+0.40	+0.40						(Regimen C)
USA (CA) 1995	SL	0.61	0.59	1	139	<0.05 0.05	0.27 0.31		BK-95R-06
(Liberty Link)				6-leaf					CA-016
									(Regimen D)
USA (MN) 1995	SL	0.19	0.21	3	95	< 0.05 (2)	< 0.05 (2)		BK-95R-06
(Liberty Link)		+0.20		8-leaf					KBT-01
		+0.20							(Regimen B)
USA (MN) 1995	SL	0.40	0.43	3	95	0.09 (2)	< 0.05 (2)		BK-95R-06
(Liberty Link)		+0.41		8-leaf					KBT-01
110 A 0 D 1005	CT.	+0.41	0.64	-	07	0.10.0.10	0.05 (0)		(Regimen C)
USA (MN) 1995	SL	0.61	0.64	2	95	0.12 0.10	<0.05 (2)		BK-95R-06
(Liberty Link)		+0.62		8-leaf					KB1-01
UGA (ID) 1005	CI	0.01	0.01	2	41	0.06 -0.05	-0.05 (2)		(Regimen D)
USA (ID) 1995 (Liberte Link)	SL	0.21	0.21	3 9 1aaf	41	0.06 < 0.05	<0.05 (3)		BK-95K-00
(Liberty Link)		+0.20		8-leal		<0.05)			TWM-01
USA (ID) 1005	SI	+0.21	0.42	2	41	0 16 0 15	<0.05 (2)		Regimen B)
(Liberty Link)	SL	0.41	0.42	S 8 loof	41	0.10 0.15	<0.03 (2)		DK-95K-00 TWM 01
(LIOCITY LIIK)		+0.43		0-icai					Regimen C)
USA (ID) 1995	SL	0.63	0.64	2	41	0.21	0.06		BK-95R-06
(Liberty Link)	51	+0.61	0.01	8-leaf		0.21	0.00		TWM-01
(Regimen D)
USA (ND) 1995	SL	0.21	0.21	3	103	0.08 0.06	< 0.05 (3)		BK-95R-06
(Liberty Link)		+0.21	+0.22	11-12-leaf		0.08			PGM-01
		+0.22	+0.21						(Regimen B)
USA (ND) 1995	SL	0.43	0.43	3	104	0.14 (2)	< 0.05 (2)		BK-95R-06
(Liberty Link)		+0.43		11-12-leaf					PGM-01
		+0.46							(Regimen C)
USA (ND) 1995	SL	0.86	0.85	2	104	0.15 0.12	< 0.05 (2)		BK-95R-06
(Liberty Link)		+0.64	+0.63	11-12-leaf					PGM-01
									(Regimen D)
USA (MI) 1996	SL	0.60	0.61	2	109	0.12 0.13	0.05 0.06		BK-96R-01
(Liberty Link)			+0.63	8-leaf					R05-02
									(Regimen B)

Location,		Appl	ication		PHI Residues, mg/kg, as glufosinate f			sinate free	Ref
Year,		1	1				acid		
(variety)	form	kg ai/ha	kg ai/hl	No ¹	days	glufosinate	MPP	NAG ²	
USA (MI) 1996	SL	0.60	0.64	3	106	0.24 0.21	0.05 < 0.05		BK-96R-01
(Liberty Link)		+0.39	+0.40	10-leaf					R05-02
	CT.	+0.60	+0.66	2	02	0.07.0.10	0.05 (0)		(Regimen C)
USA (OH) 1996 (Liberty Link)	SL	0.60	0.63	2 8 loof	83	0.27 0.12	<0.05 (2)		BK-96K-01
(Liberty Link)			+0.07	8-leal					(Rogimon B)
USA (OH) 1996	SL	0.60	0.61	3	77	0 56 0 78	<0.05 (2)		REGIMEND)
(Liberty Link)	SL	+0.39	+0.41	10-leaf	,,	0.50 0.70	<0.05 (2)		R05-03
(Licerty Linit)		+0.60	+0.66	10 104					(Regimen C)
USA (ND) 1996	SL	0.60	0.63	2	67	0.17 0.16	< 0.05 (2)		BK-96R-01
(Liberty Link)				7.5-8.5-leaf					R05-04
									(Regimen B)
USA (ND) 1996	SL	0.60	0.63	3	62	0.54 0.70	< 0.05 (2)		BK-96R-01
(Liberty Link)		+0.39	+0.41	9.5-11-leaf					R05-04
		+0.60	+0.63						(Regimen C)
USA (NE) 1996	SL	0.60	0.63	2	115	< 0.05 (2)	< 0.05 (2)		BK-96R-01
(Liberty Link)				8-leaf					R07-01
	CT.	0.00	0.61	2	100	0.07.0.05	0.05 (2)		Regimen B)
USA (NE) 1996	SL	0.60	0.61	3	108	0.07 0.05	<0.05 (2)		BK-96R-01
(Liberty Link)		+0.39	+0.41	10-leaf					R07-01
USA (ND) 1006	C1	+0.00	+0.00	2	72	0.12.0.14	<0.05 (2)		Regimen C)
(Liberty Link)	SL	0.00	0.05	Z 8 loof	15	0.12 0.14	<0.03 (2)		DK-90K-01
(Liberty Link)				o-leal					(Regimen B)
USA (ND) 1996	SL	0.60	0.63	3	66	0.28.0.33	0.07.0.11		BK-96R-01
(Liberty Link)	5L	+0.39	+0.41	10-leaf	00	0.20 0.33	0.07 0.11		R07-02
(Excerty Emily)		+0.60	+0.63	10 Ioui					(Regimen C)
USA (CO) 1996	SL	0.60	0.59	2	80	< 0.05 (2)	< 0.05 (2)		BK-96R-01
(Liberty Link)				8-leaf					R08-01
									(Regimen B)
USA (CO) 1996	SL	0.60	0.63	3	68	0.53 0.55	< 0.05 (2)		BK-96R-01
(Liberty Link)		+0.39	+0.40	10-leaf					R08-01
		+0.60	+0.58						(Regimen C)
USA (CO) 1996	SL	0.60	0.61	2	86	0.11 (2)	< 0.05 (2)		BK-96R-01
(Liberty Link)			+0.59	8-leaf					R09-01
	~ -								(Regimen B)
USA (CO) 1996	SL	0.60	0.64	3	81	0.27 0.30	<0.05 (2)		BK-96R-01
(Liberty Link)		+0.39	+0.41	10-leaf					(Decimon C)
	CI	+0.00	+0.01	2	122	0.06.0.08	0.07.0.06		(Regimen C)
(Liberty Link)	SL	0.00	0.05 ± 0.62	Z 8-leaf	152	0.00 0.08	0.07 0.00		BK-90K-01 B10-01
(LIDEITY LIIK)			+0.02	0-leal					(Regimen B)
USA (CA) 1996	SL.	0.60	0.61	3	122	0 37 0 36	0.06.0.07		BK-96R-01
(Liberty Link)	51	+0.39	+0.42	10-leaf	122	0.57 0.50	0.00 0.07		R10-01
(+0.60	+0.63						(Regimen C)
USA (ID) 1996	SL	0.60	0.62	2	128	0.07 0.06	< 0.05 (2)		BK-96R-01
(Liberty Link)				8-10-leaf					R11-01
· · · ·									(Regimen B)
USA (ID) 1996	SL	0.60	0.61	3	121	0.19 0.22	< 0.05 (2)		BK-96R-01
(Liberty Link)		+0.39	+0.41	10-leaf					R11-01
		+0.60	+0.63						(Regimen C)
USA (CA) 1996	SL	2.8	3.0	3	136	0.23	0.93		BK-96-R05
(Liberty Link)		+3.0	+3.2	8-leaf					
		+3.0	+3.2						

¹: growth stage is described in terms of number of leaves. ² included in glufosinate-ammonium result. CLICK HERE for continue

Country,	Applic	ation			PHI	Residues, as g	lufosinate free	acid, mg/kg ²	Ref
Year, (variety)	form	kg ai/ha	kg ai/hl	No ¹	days	Glufosinate	MPP	NAG	
France 1993 (LH163×LH164- ×F2×T14)	SL	0.45	0.23	2 GS25	0 39 119	20 shoot 0.10 shoot <0.05 shoot	0.17 shoot <0.05 shoot <0.05 shoot	13 shoot 0.44 shoot 0.06 shoot	A54225 ER93ECN550 FRA000402
					119	<0.05 cob	<0.05 cob	glu	
France 1993 (LH206×LH74- ×F2×T14)	SL	0.45	0.23	2 GS24	0 43 119	6.2 shoot <0.05 shoot <0.05 shoot	0.15 shoot <0.05 shoot <0.05 shoot	8.6 shoot <0.05 shoot <0.05 shoot	A54225 ER93ECN550 FRA000502
					119	<0.05 cob	<0.05 cob	glu	
France 1993 (MLC 2101/T14)	SL	0.45	0.23	2 GS24	0 48 84	27 shoot 1.2 shoot <0.05 shoot	0.11 shoot <0.05 shoot <0.05 shoot	19 shoot 0.76 shoot <0.05 shoot	A54225 ER93ECN550 FRA000102
					84	<0.05 cob	<0.05 cob	glu	
France 1993 (MLC 2101/T14)	SL	0.45	0.23	2 GS24	0 51 92	24 shoot 0.24 shoot <0.05 shoot	0.47 shoot 0.06 shoot 0.12 shoot	7.0 shoot <0.05 shoot <0.05 shoot	A54225 ER93ECN550 FRA000202
					92	<0.05 cob	0.23 cob	glu	
France 1993 (MLC 2101/T14)	SL	0.45	0.23	2 GS24	0 39 106	26 shoot 0.09 shoot <0.05 shoot	0.22 shoot <0.05 shoot <0.05 shoot	17 shoot <0.05 shoot 0.05 shoot	A54225 ER93ECN550 FRA000302
					106	<0.05 cob	<0.05 cob	glu	
France 1994 ((LH82(4)T25)sf (- 2)xSH298	SL	0.45	0.23	2 GS25	0 35 90 90	24 shoot 0.1 shoot 0.07 shoot <0.05 cob	<0.05 shoot <0.05 shoot <0.05 shoot <0.05 cob	glu glu glu glu	A54229 ER94ECN550 FRA000202
France 1994 (F6×M1)	SL	0.45	0.23	2 GS25	0 35 77	9.9 shoot <0.05 shoot <0.05 shoot	0.13 shoot <0.05 shoot <0.05 shoot	6.5 shoot 0.11 shoot <0.05 shoot	A54226 ER93ECS550 FRA000102
					77	<0.05 cob	<0.05 cob	glu	
France 1994 (LH824xT25)AF4x- B73)	SL	0.45	0.15	2 GS25	0 67 67	43 shoot <0.05 shoot <0.05 cob	0.27 shoot <0.05 shoot <0.05 cob	glu glu glu	A54230 ER94ECS550 ER 4000102
France 1994 (SH298(LH82(4)T2 5)sf(2))	SL	0.45	0.23	2 GS25	0 27 69 69	24 shoot 0.12 shoot 0.36 shoot <0.05 cob	<0.05 shoot <0.05 shoot <0.05 shoot <0.05 shoot <0.05 cob	glu glu glu glu	A54229 ER94ECN550 FRA000102
France 1995 (F2xLH82-T25)	SL	0.45	0.23	2 GS18	0 44 72 72	25 shoot 0.51 shoot 0.47 shoot <0.05 cob	0.26 shoot <0.05 shoot <0.05 shoot <0.05 cob	glu glu glu glu	A56445 ER95ECN550 FRA000202
France 1995 (F2xLH82-T25)	SL	0.45	0.23	2 GS19	0 44 72 72	25 shoot 0.83 shoot 0.82 shoot <0.05 cob	0.29 shoot <0.05 shoot <0.05 shoot <0.05 cob	glu glu glu glu	A56445 ER95ECN550 FRA000203
France 1995 (tolerant 039866)	SL	0.45	0.23	2 GS18	0 55 73 73	20 shoot 0.12 shoot 0.12 shoot <0.05 cob	0.2 shoot <0.05 shoot <0.05 shoot <0.05 cob	glu glu glu glu	A56445 ER95ECN550 FRA000103
France 1995 (tolerant 039866)	SL	0.45	0.23	2 GS18	0 55 73 73	15 shoot 0.1 shoot <0.05 cob	0.15 shoot <0.05 shoot <0.05 shoot <0.05 cob	glu glu glu glu	A56445 ER95ECN550 FRA000102

Table 53. Glufosinate residues in maize forage and fodder resulting from supervised trials in France, Germany, Italy, The Netherlands and Spain.

Country,	Applica	ation			PHI	Residues, as g	lufosinate free	acid, mg/kg ²	Ref
Year, (variety)	form	kg ai/ha	kg ai/hl	No ¹	days	Glufosinate	MPP	NAG	
France 1995 (Transgenic hybrid)	SL	0.53	0.27	2 GS18	0 49 65 65	7.9 shoot 0.43 shoot 0.16 shoot <0.05 cob	0.08 shoot 0.06 shoot 0.06 shoot <0.05 cob	4.2 shoot 0.28 shoot 0.37 shoot <0.05 cob	A54760 ER95ECS550 FRA000103
France 1995 (Transgenic hybrid)	SL	0.53	0.27	2 GS18	0 49 65 65	8.3 shoot 0.14 shoot <0.05 shoot <0.05 cob	0.05 shoot <0.05 shoot <0.05 shoot <0.05 cob	4.2 shoot 0.54 shoot 0.21 shoot <0.05 cob	A54760 ER95ECS550 FRA000102
France 1995 (Transgenic hybrid)	SL	0.45	0.23	2 GS19	0 49 60 60	8.9 shoot <0.05 shoot 0.05 shoot <0.05 cob	0.13 shoot 0.05 shoot 0.07 shoot <0.05 cob	5.1 shoot 0.24 shoot 0.37 shoot <0.05 cob	A54760 ER95ECS550 FRA000203
France 1995 (Transgenic hybrid)	SL	0.45	0.23	2 GS19	0 49 60 60	14 shoot <0.05 shoot <0.05 shoot <0.05 cob	0.14 shoot <0.05 shoot <0.05 shoot <0.05 cob	6.5 shoot 0.13 shoot 0.14 shoot <0.05 cob	A54760 ER95ECS550 FRA000202
France 1996 (transgenic hybrid)	SL	0.60	0.24	2 GS18	0 56 75 75	9.1 shoot 0.14 shoot 0.19 shoot <0.05 cob	0.05 shoot <0.05 shoot <0.05 shoot <0.05 cob	glu glu glu glu	A58191 ER96ECN550 FRA000102
France 1996 (transgenic hybrid)	SL	0.80	0.32	2 GS18	0 56 75 75	18 shoot 0.28 shoot 0.65 shoot <0.05 cob	0.08 shoot <0.05 shoot <0.05 shoot <0.05 cob	glu glu glu glu	A58191 ER96ECN550 FRA000103
France 1996 (transgenic hybrid)	SL	0.45	0.18	2 GS18	0 56 75 75	16 shoot 0.4 shoot 0.31 shoot <0.05 cob	0.1 shoot <0.05 shoot <0.05 shoot <0.05 cob	glu glu glu glu	A58191 ER96ECN550 FRA000104
France 1996 (transgenic hybrid)	SL	0.60	0.24	2 GS18	0 57 76 76	26 shoot 0.24 shoot 0.57 shoot <0.05 cob	0.09 shoot <0.05 shoot <0.05 shoot <0.05 cob	glu glu glu glu	A58191 ER96ECN550 FRA000302
France 1996 (transgenic hybrid)	SL	0.80	0.32	2 GS18	0 57 76 76	24 shoot 0.48 shoot 0.54 shoot <0.05 cob	0.09 shoot <0.05 shoot <0.05 shoot <0.05 cob	glu glu glu glu	A58191 ER96ECN550 FRA000303
France 1996 (transgenic hybrid)	SL	0.45	0.18	2 GS18	0 57 76 76	14 shoot 0.2 shoot 0.23 shoot <0.05 cob	<0.05 shoot <0.05 shoot <0.05 shoot <0.05 cob	glu glu glu glu	A58191 ER96ECN550 FRA000304
France 1996 (transgenic hybrid)	SL	0.60	0.24	2 GS18	0 62 76 76	16 shoot 0.13 shoot 0.22 shoot <0.05 cob	0.07 shoot <0.05 shoot <0.05 shoot <0.05 cob	glu glu glu glu	A58191 ER96ECN550 FRA000402
France 1996 (transgenic hybrid)	SL	0.80	0.32	2 GS18	0 62 76 76	25 shoot 0.2 shoot 0.27 shoot <0.05 cob	0.08 shoot <0.05 shoot <0.05 shoot <0.05 cob	glu glu glu glu	A58191 ER96ECN550 FRA000403
France 1996 (transgenic hybrid)	SL	0.45	0.18	2 GS18	0 62 76 76	20 shoot <0.05 shoot 0.15 shoot <0.05 cob	<0.05 shoot <0.05 shoot <0.05 shoot <0.05 cob	glu glu glu glu	A58191 ER96ECN550 FRA000404
France 1996 (transgenic hybrid)	SL	0.45	0.18	2 GS18	0 46 75 75	16 shoot 0.31 shoot 0.37 shoot <0.05 cob	0.05 shoot <0.05 shoot <0.05 shoot <0.05 cob	glu glu glu glu	A58190 ER96ECS550 FRA000302
France 1996 (transgenic hybrid)	SL	0.60	0.24	2 GS18	0 46 75 75	19 shoot 0.33 shoot 0.64 shoot <0.05 cob	0.06 shoot <0.05 shoot <0.05 shoot <0.05 cob	giu glu glu glu	A58190 ER96ECS550 FRA000303

Year, (variety) form kg aiha	Country,	Applic	ation			PHI	Residues, as g	lufosinate free	acid, mg/kg ²	Ref
France 1996 (masgenic hybrid) SL (masgenic hybrid)	Year, (variety)	form	kg ai/ha	kg ai/hl	No ¹	days	Glufosinate	MPP	NAG	
(fransgenic hybrid) Finace 1996 SL 0.60 0.24 2.0 0.47 shoot -0.05 shoot elia FRA000304 (fransgenic hybrid) SL 0.60 0.24 2 0 18 shoot 0.05 shoot elia AS8191 (fransgenic hybrid) SL 0.80 0.22 2 0 23 shoot 0.05 shoot elia AS8191 (fransgenic hybrid) SL 0.80 0.22 2 0 23 shoot 0.06 shoot elia AS8191 (fransgenic hybrid) SL 0.45 0.18 2 0 23 shoot 0.05 shoot elia AS8191 (fransgenic hybrid) SL 0.45 0.18 2 0 12 shoot 0.05 shoot elia ER96ECS50 (fransgenic hybrid) SL 0.45 0.18 2 0 31 shoot 0.05 shoot elia ER96ECS50 (fransgenic hybrid) SL 0.45 0.18 2 0 31 shoot 0.05 shoot	France 1996	SL	0.80	0.32	2	0	34 shoot	0.09 shoot	glu	A58190
Prance 1996 SL 0.60 0.24 2 0 18 shoot 0.05 solo 0.05 solo Pin FA00030 (transgenic hybrid) L 0.60 0.24 2 0 18 shoot 0.07 shoot -0.05 shoot Pin FRA00030 Pin FRA00020 Pin FRA000200 Pin FRA000200 Pin FRA000201 Pin FRA000102 Pin FRA000102 Pin FRA000102 Pin FRA000102 Pin FRA000102 Pin FRA000102 <td< td=""><td>(transgenic hybrid)</td><td></td><td></td><td></td><td>GS18</td><td>46</td><td>0.44 shoot</td><td><0.05 shoot</td><td>giu</td><td>ER96ECS550</td></td<>	(transgenic hybrid)				GS18	46	0.44 shoot	<0.05 shoot	giu	ER96ECS550
$ \begin{array}{ c c c c c c c c c c c c c c c c c c c$						75	0.47 shoot	<0.05 shoot	glu	FRA000304
$ \begin{split} & \mbox{France 1996} & \mbox{SL} & 0.60 & 0.24 & 2 & 0 & 18 shoot & 0.07 shoot & e^{10} & ER96ECN50 \\ & \mbox{France 1996} & \mbox{SL} & 0.80 & 0.32 & 2 & 0 & 23 shoot & 0.05 shoot & e^{10} & ER96ECN50 \\ & \mbox{France 1996} & \mbox{SL} & 0.45 & 0.18 & 2 & 0 & 23 shoot & 0.05 shoot & e^{10} & ER96ECN50 \\ & \mbox{France 1996} & \mbox{SL} & 0.45 & 0.18 & 2 & 0 & 23 shoot & 0.05 shoot & e^{10} & ER96ECN50 \\ & \mbox{France 1996} & \mbox{SL} & 0.45 & 0.18 & 2 & 0 & 23 shoot & 0.05 shoot & e^{10} & ER96ECN50 \\ & \mbox{France 1996} & \mbox{SL} & 0.45 & 0.18 & 2 & 0 & 23 shoot & 0.05 shoot & e^{10} & ER96ECN50 \\ & \mbox{France 1996} & \mbox{SL} & 0.45 & 0.18 & 2 & 0 & 33 shoot & 0.05 shoot & e^{10} & ER96ECN50 \\ & \mbox{France 1996} & \mbox{SL} & 0.60 & 0.18 & 2 & 0 & 31 shoot & 0.05 shoot & e^{10} & ER96ECN50 \\ & \mbox{France 1996} & \mbox{SL} & 0.60 & 0.24 & 2 & 0 & 36 shoot & 0.05 shoot & e^{10} & ER96ECS50 \\ & \mbox{France 1996} & \mbox{SL} & 0.60 & 0.24 & 2 & 0 & 36 shoot & 0.05 shoot & e^{10} & ER96ECS50 \\ & \mbox{France 1996} & \\mbox{SL} & 0.80 & 0.32 & 2 & 0 & 36 shoot & 0.05 shoot & e^{10} & ER96ECS50 \\ & \mbox{France 1996} & \\mbox{SL} & 0.45 & 0.18 & 2 & 0 & 36 shoot & 0.05 shoot & e^{10} & ER96ECS50 \\ & \mbox{France 1996} & \\mbox{SL} & 0.45 & 0.18 & 2 & 0 & 36 shoot & 0.05 shoot & e^{10} & ER96ECS50 \\ & \mbox{France 1996} & \\mbox{SL} & 0.45 & 0.18 & 2 & 0 & 0.5 shoot & 0.05 shoot & e^{10} & ER96ECS50 \\ & \mbox{France 1996} & \\mbox{SL} & 0.45 & 0.18 & 2 & 0 & 0.5 shoot & 0.05 shoot & e^{10} & ER96ECS50 \\ & \mbox{France 1996} & \\mbox{SL} & 0.45 & 0.18 & 2 & 0 & 0.5 shoot & 0.05 shoot & e^{10} & ER96ECS50 \\ & \mbox{France 1996} & \\mbox{SL} & 0.45 & 0.18 & 2 & 0 & 0.5 shoot & 0.05 shoot & e^{10} & ER96ECS50 \\ & \mbox{France 1996} & \\mbox{SL} & 0.45 & 0.18 & 2 & 0 & 0.5 shoot & 0.05 shoot & e^{10} & ER96ECS50 \\ & \mbox{France 1996} & \\mbox{SL} & 0.45 & 0.18 & 2 & 0 & 0.5 shoot & 0.05 shoot & e^{10} & ER96ECS50 \\ & \mbox{France 1996} & \\mbox{SL} & 0.45 & 0.15 & 0.25 & 0.05 shoot & 0.05 shoot & 0.05 $						75	<0.05 cob	<0.05 cob	glu	
	France 1996	SL	0.60	0.24	2	0	18 shoot	0.07 shoot	glu	A58191
Prace 1996 SL 0.80 0.32 2 0 2.3 bnot 0.08 bnot 20 AS8191 France 1996 SL 0.80 0.32 2 0 2.3 bnot 0.08 shoot 20 AS8191 France 1996 SL 0.45 0.44 shoot 0.05 shoot 20 AS8191 France 1996 SL 0.45 0.18 2 0 2.2 shoot 0.09 shoot 20 A58191 France 1996 SL 0.45 0.18 2 0 31 shoot 0.05 shoot 20 76 -0.05 shoot 20 76 -0.05 shoot 20 76 -0.05 shoot 20 78 -0.05 shoot 20 78 -0.05 shoot 20 78 -0.05 shoot 20 78 -0.05 shoot 20 -0.05 shoot -0.05 shoot -	(transgenic hybrid)				GS19	56	0.27 shoot	<0.05 shoot	glu	ER96ECN550
$ \begin{array}{ c c c c c c c c c c c c c c c c c c c$						76	0.75 shoot	<0.05 shoot	glu	FRA000202
France 1996 SL 0.80 0.32 2 0 25 shoot 0.08 shoot 2 mask of the shoot	F 400.6	<i>a</i> .	0.00	0.00	-	76	<0.05 cob	<0.05 cob	glu	1.50101
(Hansgenic hybrid) SL (G19) 56 (J.44 shoot) (J.05 shoot) Jack (RA000203) France 1996 SL 0.45 0.18 2 0.2 shoot 0.09 shoot Shi FRA000204 France 1996 SL 0.45 0.18 2 0 31 shoot 0.14 shoot Shoot Shi FRA000204 France 1996 SL 0.45 0.18 2 0 31 shoot 0.14 shoot Shoot Shoot FRA000102 France 1996 SL 0.60 0.24 2 0 36 shoot 0.24 shoot 4.05 shoot 2.05 sob Shoot FRA000102 France 1996 SL 0.60 0.24 2 0 36 shoot 0.24 shoot 4.05 shoot 2.05 sob Shoot FRA000103 France 1996 SL 0.80 0.32 2.0 0 6.5 shoot 0.05 shoot Shoot FRA000104 France 1996 SL 0.80 0.31 shoot 0.03 shoot 0.05 shoo	France 1996	SL	0.80	0.32	2	0	23 shoot	0.08 shoot	glu	A58191
	(transgenic hybrid)				GS19	56	0.44 shoot	<0.05 shoot	glu	ER96ECN550
						76 76	0.46 shoot	< 0.05 shoot	glu	FRA000203
	E	CT	0.45	0.10		/0	<0.05 COD	<0.05 COD	glu	4.59101
(Hamsgenic hybrid) SL 0.45 0.18 2 0 0.17 shoot 76 0.03 shoot 0.05 shoot 91 2000 shoot 91 2000 shoot 0.05 shoot 91 2000 shoot 91 2000 shoot 91 2000 shoot 0.05 shoot 91 2000 shoot 91	France 1996	SL	0.45	0.18	2	0	22 shoot	0.09 shoot	glu	A58191
France 1996 (transgenic hybrid) SL ERGE/ESS50 0.45 (uransgenic hybrid) SL ERGE/ESS50 0.45 (uransgenic hybrid) SL ERGE/ESS50 0.60 (uransgenic hybrid) SL ERGE/ESS50 0.18 (uransgenic hybrid) SL ERGE/ESS50 0.24 (uransgenic hybrid) SL ERGE/ESS50 0.38 (GETMAM) 0.38 (noot (uransgenic hybrid) 0.45 (uransgenic hybrid) SL ERGE/ESS50 0.24 (uransgenic hybrid) SL ERGE/ESS50 0.24 (uransgenic hybrid) SL ERGE/ESS50 0.38 (noot (GETMAM) 0.38 (noot (uransgenic hybrid) 0.45 (uransgenic hybrid) 0.45 (uransgenic hybrid) 0.45 (uransgenic	(transgenic hybrid)				GS19	50 76	0.17 shoot	< 0.05 shoot	glu	EK96ECN550
Prance 1996 (transgenic hybrid) SL (transgenic hybrid) 0.45 (transgenic hybrid) 0.60 (transgenic hybrid) 0.81 (transgenic hybrid) 0.81 (transgenic hybrid) 0.82 (transgenic hybrid) 0.82 (transgenic hybrid) 0.82 (transgenic hybrid) 0.43 (transgenic hybrid) 0.45 (transgenic hybrid) 0.45 (transgenic hybrid) 0.45 (transgenic hybrid) 0.45 (transgenic hybrid) 0.84 (transgenic hybrid) 0.						70 76	< 0.05 should < 0.05 sob	< 0.05 shoot	glu	FKA000204
Prince 1996 SL 0.45 0.18 2 0 15 1001 0.14 shout Prince Prince 1996 SL 0.45 CS19 64 0.05 shoot 0.05 shoot 0.05 shoot 0.05 shoot 0.05 shoot 0.07 shoot 0.07 shoot Prince 1996 SL 0.60 0.24 2 0 35 shoot 0.05 shoot 0.02 shoot Prince 1996 SL 0.80 0.24 2 0 36 shoot 0.05 shoot Prince 1996 SL 0.80 0.32 2 0 50 shoot 0.02 shoot Prince 1996 SL 0.80 0.32 2 0 50 shoot 0.02 shoot Prince 1996 SL 0.80 0.32 2 0 65 shoot 0.05 shoot Prince 1996 SL 0.45 0.18 2 0 65 shoot 0.05 shoot Prince 1996 SL 0.45 0.18 2 0 65 shoot 0.05 shoot Prin A58190 France 1996 SL 0.60 0.24 2 0	Emma 1006	CI	0.45	0.10	2	70	<0.05 C00	<0.05 C00	glu	4.59100
(framsgenic hybrid) SL (transgenic hybrid) SL (transgenic hybrid) SL (transgenic hybrid) SL (transgenic hybrid) SL (transgenic hybrid) 0.60 (transgenic hybrid) SL (transgenic hybrid) 0.80 (transgenic hybrid) SL (transgenic hybrid) 0.80 (transgenic hybrid) SL (transgenic hybrid) 0.80 (transgenic hybrid) SL (transgenic hybrid) 0.80 (transgenic hybrid) SL (transgenic hybrid) 0.81 (transgenic hybrid) SL (transgenic hybrid) 0.81 (transgenic hybrid) SL (transgenic hybrid) 0.82 (transgenic hybrid) 0.60 (transgenic hybrid) 0.82 (transgenic hybrid) 0.60 (transgenic hybrid) 0.82 (transgenic hybrid) 0.60 (transgenic hybrid)	France 1996	SL	0.45	0.18	2	0	31 shoot	0.14 shoot	glu	A58190
France 1996 (transgenic hybrid) SL SL SL SL SL SL SL SL SL SL SL SL SL S	(transgenic hybrid)				0319	04	<0.05 shoot	< 0.05 shoot	glu	ER90EC5550
France 1996 (transgenic hybrid) SL E 0.60 0.60 0.24 0.24 2 C S1 2 0 S1 0 S1 00 0.08 shoot 91 20.05 cob 0.08 shoot 0.05 shoot 91 20.05 cob 0.05 shoot 0.05 shoot 91 4 0.05 shoot 0.05 shoot 91 A58190 clos shoot 91 A58190 clos shoot 91 France 1996 (transgenic hybrid) SL 2 0.80 0.32 2 C S1 0 GS19 50 shoot 91 0.24 shoot 0.12 shoot 91 20.05 cob 0.05 shoot 91 20.05 shoot 0.05 shoot 91 20.05 shoot 0.05 shoot 91 20.05 shoot 91 20.05 shoot 91 20.05 shoot 91 20.05 shoot 0.05 shoot 91 ER96ECS550 91 France 1996 (transgenic hybrid) SL 2 0.45 0.18 2 C S1 0 S1 6.51 shoot 86 -0.05 shoot 91 Ch05 shoot 91 A58190 France 1996 (transgenic hybrid) SL 2 0.60 0.24 2 C S1 0 S1 8.9 shoot -0.05 cob -0.05 shoot 91 ER96ECS550 91 France 1996 (transgenic hybrid) SL 2 0.80 0.32 2 C S1 0 S1 7.2 shoot -0.05 shoot 86 -0.05 shoot 91 A58190 France 1996 (transgenic hybrid) SL 2 0.45 0.15 2 C S1 0 S1 <s1< td=""> 0.15</s1<>						91	<0.05 snoot	<0.05 shoot	glu	FRA000102
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	E	CT	0.00	0.24	-	91	<0.05 COD	<0.05 COD	glu	4.59100
(fransgenic hybrid) SL 0.80 0.32 2 0 50 shoot c.0.05 shoot glu FRA000103 France 1996 SL 0.80 0.32 2 0 50 shoot 0.24 shoot glu A58190 France 1996 SL 0.45 0.18 2 0 50 shoot -0.05 shoot glu Bfa96CS550 glu FRA000103 France 1996 SL 0.45 0.18 2 0 65 shoot -0.05 shoot glu FRA00202 ftransgenic hybrid) SL 0.45 0.18 2 0 6.5 shoot -0.05 shoot glu FRA00202 ftransgenic hybrid) SL 0.60 0.24 2 0 8.9 shoot -0.05 shoot glu FRA00202 ftransgenic hybrid) SL 0.60 0.24 2 0 8.9 shoot -0.05 shoot glu FRA00203 ftransgenic hybrid) SL 0.60 0.32 2 0 7.2 shoot -0.05 sho	France 1996	SL	0.60	0.24	2	0	36 shoot	0.2 shoot	glu	A58190
France 1996 (transgenic hybrid) SL (transgenic hybrid) 0.80 (solution) 0.85 (solution)	(transgenic hybrid)				GS19	04 01	0.06 shoot	<0.05 shoot	glu	EK96EC5550
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $						91	0.08 shoot	<0.05 shoot	glu	FRA000103
$ \begin{array}{c} \mbox{France 1996} \\ \mbox{(transgenic hybrid)} \\ \mbox{(transgenic hybrid)} \\ \mbox{France 1996} \\ \mbox{(transgenic hybrid)} \\ \mbox{SL} \\ \mbox{0.8L} \\ \mbox{0.8L} \\ \mbox{0.8L} \\ \mbox{0.4S} \\ \mbox{0.3L} \\ \mbox{0.4S} \\ \$	E 1007	CT.	0.00	0.22	2	91	<0.05 cob	<0.05 cob	glu	4.50100
(fransgenic hybrid) SL 0.45 0.15 2 00 2 8000 91 <0.05 shoot 91 <0.05 sho	France 1996	SL	0.80	0.32	2	0	50 shoot	0.24 shoot	glu	A58190
France 1996 (transgenic hybrid) SL SL Status 0.45 (unsgenic hybrid) 0.45 SL SL SL SL SL SL SL SL SL 0.45 (1.45) 0.18 (1.45) 2 (1.45) 0 (1.45) 0 (1.45) 0 (1.45) 0 (1.45) 1.45 (1.45) 0 (1.45) 1.45 (1.45)	(transgenic hybrid)				GS19	64 01	0.12 shoot	<0.05 shoot	glu	ER96ECS550
France 1996 (transgenic hybrid) SL 0.45 0.18 2 0 6.5 5000 2003 boot en A58190 France 1996 (transgenic hybrid) SL 0.65 0.18 2 0 6.5 5000 <0.05 shoot						91 01	0.13 shoot	< 0.05 shoot	glu	FRA000104
Prance 1996 SL 0.45 0.18 2 0 5.8 moother c.005 shoother c.005 s	Emmas 1006	CI	0.45	0.10	2	91	<0.03 C00	<0.05 cob	glu	4.59100
	(trance 1990	SL	0.45	0.18	2 CS10	0 57	0.5 shoot	< 0.05 shoot	glu	A38190
$ \begin{array}{ c c c c c c c c c c c c c c c c c c c$	(transgenic hybrid)				0319	57	0.25 shoot	< 0.05 shoot	glu	ER90EC5550
France 1996 (transgenic hybrid)SL0.600.242 GS1908.9 shoot GS19<0.05 shoot S0 <0.05 shoot glu <0.0						00 86	< 0.05 should < 0.05 sob	< 0.05 shoot	glu	FKA000202
	Emmas 1006	CI	0.60	0.24	2	00	<0.03 C00	<0.05 cob	glu	4.59100
(Iransgenic hybrid) SL 0.80 0.32 2 0 7.2 shoot <0.05 shoot glu FRA000203 France 1996 SL 0.80 0.32 2 0 7.2 shoot <0.05 shoot	(transce 1990	SL	0.00	0.24	2 CS10	0 57	8.9 snoot	< 0.05 shoot	glu	A38190
$ \begin{array}{ c c c c c c c c c c c c c c c c c c c$	(transgenic hybrid)				0319	57	0.27 shoot	< 0.05 shoot	glu	ER90EC3330
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $						86	< 0.05 coh	< 0.05 shoot	glu	TRA000205
	Eranaa 1006	CT.	0.80	0.22	2	0	<0.05 C00	<0.05 cbb	glu	458100
$ \begin{array}{ c c c c c c c c c c c c c c c c c c c$	(transgenic hybrid)	SL	0.80	0.52	2 GS10	0 57	0.11 shoot	< 0.05 shoot	glu	FR06FCS550
Germany 1994 (F2xLH82- 4)T25sf(2)SL 0.45 0.15 (LH82(4)T25sf'4F2) 0.15 2 ($1482^{-4}T25sf'4F2)$ 0.45 0.15 2 (2 (148	(transgenie nybrid)				0517	86	0.11 shoot	< 0.05 shoot	glu	ER/02C3550
Germany 1994SL 0.45 0.15 2 GS24 0 23 shout 0.65 shout g^{lu} g^{lu} A54229 ER94ECN550 DEU040202Germany 1994SL 0.45 0.15 2 GS24 0 23 shout 0.65 shout $(0.05 \text{ shout}$ g^{lu} g^{lu} g^{lu} A54229 ER94ECN550 DEU040202Germany 1994 (F2xLH82- 4)T25sf(2)SL 0.45 0.15 2 2 0 2 39 shout 0.53 shout 6.05 cob $d55 \text{ shout}$ $d55 \text{ shout}$ g^{lu} $d52 \text{ cob}$ Germany 1994 (LH82(4)T25st2xS- H298)SL 0.45 0.15 2 2 0 44 29 shout $d.05 \text{ shout}$ $d.05 \text{ shout}$ $d53 \text{ shout}$ $d53 \text{ shout}$ Germany 1995 (LH82^4T25sf^4F2)SL 0.45 0.15 2 2 0 38 shout 29 shout $d.05 \text{ shout}$ g^{lu} g^{lu} $d.54229$ g^{lu} Germany 1995 (LH82^4T25sf^4F2)SL 0.45 0.15 2 2 0 38 shout $d.05 \text{ shout}$ $d.05 \text{ shout}$ g^{lu} g^{lu} $d.54229$ g^{lu} Germany 1995 (LH82^4T25sf^4F2)SL 0.45 0.15 2 2 0 38 shout 0.17 shout g^{lu} g^{lu} $d.56445$ g^{lu} Germany 1995 (LH82^4T25sf^4F2)SL 0.45 0.15 2 2 0 78 0.97 shout 0.17 shout g^{lu} g^{lu} $d.56445$ g^{lu} <td></td> <td></td> <td></td> <td></td> <td></td> <td>86</td> <td>< 0.05 cob</td> <td>< 0.05 shoot</td> <td>glu</td> <td>TRA000204</td>						86	< 0.05 cob	< 0.05 shoot	glu	TRA000204
Germany 1994 SL 0.45 0.15 2 0 37 0.74 shoot c0.05 shoot glu glu <thglu< th=""> glu glu</thglu<>	Germany 1994	SI	0.45	0.15	2	0	23 shoot	0.65 shoot	glu	Δ 54229
Germany 1994 (F2xLH82- 4)T25sf(2) SL 0.45 0.15 2 0.15 0 2 0 0 39 shoot (SS shoot) 0.05 shoot (SS shoot) glu glu A54229 (ER94ECN550 DEU040202 Germany 1994 (F2xLH82- 4)T25sf(2) SL 0.45 0.15 2 0 0 2 39 shoot (SS shoot) 0.53 shoot (SS shoot) glu glu A54229 (ER94ECN550 DEU050102 Germany 1994 (LH82(4)T25st2xS- H298) SL 0.45 0.15 2 0.15 0 2 2 0 0 29 shoot <0.05 shoot (SS shoot) glu glu A54229 (ER94ECN550 DEU050102 Germany 1994 (LH82(4)T25st2xS- H298) SL 0.45 0.15 2 0 0 2 2 0 0 29 shoot <0.05 shoot (SS shoot) glu glu A54229 (ER94ECN550 DEU050102 Germany 1995 (LH82^4T25sf^4F2) SL 0.45 0.15 2 0 0 30 shoot <0.05 shoot (SS shoot) glu glu A56445 ER95ECN550 Germany 1995 (LH82^4T25sf^4F2) SL 0.45 0.15 2 0 0 30 shoot 0.17 shoot (0.05 shoot) glu glu A56445 ER95ECN550 Guruany 1995 (LH82^4T25sf^4F2) SL 0.45 0.15 2 0.15 0 30 shoot (78 0.09 cob 0.05 shoot (0.05 shoot) glu glu	Germany 1774	SL	0.45	0.15	2 GS24	37	0.74 shoot	< 0.05 shoot	glu	FR94ECN550
$ \begin{array}{ c c c c c c c c c c c c c c c c c c c$					0524	55	0.74 shoot 0.67 shoot	<0.05 shoot	glu	DEU040202
$ \begin{array}{ c c c c c c c c c c c c c c c c c c c$						55	< 0.05 cob	<0.05 cob	glu	DEC010202
$ \begin{array}{c ccccccccccccccccccccccccccccccccccc$						76	<0.05 cob	<0.05 cob	glu	
$ \begin{array}{c ccccccccccccccccccccccccccccccccccc$	Germany 1994	SL	0.45	0.15	2	0	39 shoot	0.53 shoot	glu	A 54229
$\begin{array}{c c c c c c c c c c c c c c c c c c c $	(F2xLH82-	SE	0.15	0.15	- GS23	55	< 0.05 shoot	< 0.05 shoot	glu	ER94ECN550
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	(12.12.102) 4)T25sf(2)				0020	87	< 0.05 shoot	< 0.05 shoot	glu	DEU050102
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $.)(_)					87	<0.05 cob	<0.05 cob	glu	
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	Germany 1994	SL	0.45	0.15	2	0	29 shoot	<0.05 shoot	glu	A54229
$ \begin{array}{c ccccccccccccccccccccccccccccccccccc$	(LH82(4)T25st2xS-	22	01.10	0.10	- GS25	31	0.78 shoot	< 0.05 shoot	glu	ER94ECN550
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	H298)					51	0.19 shoot	<0.05 shoot	glu	DEU030102
Germany 1995 (LH82^4T25sf^4F2) SL 0.45 0.15 2 0 30 shoot 0.17 shoot glu Germany 1995 SL 0.45 0.15 2 0 30 shoot 0.17 shoot glu A56445 GS18 38 1.5 shoot <0.05 shoot	/					84	0.26 shoot	<0.05 shoot	glu	
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $						84	<0.05 cob	<0.05 cob	glu	
$\begin{array}{c ccccccccccccccccccccccccccccccccccc$	Germany 1995	SL	0.45	0.15	2	0	30 shoot	0.17 shoot	glu	A56445
$ \begin{array}{c ccccccccccccccccccccccccccccccccccc$	(LH82^4T25sf^4F2)			5.10	- GS18	38	1.5 shoot	< 0.05 shoot	glu	ER95ECN550
78 0.09 cob <0.05 cob glu	(• •	78	0.79 shoot	< 0.05 shoot	glu	DEU010202
						78	0.09 cob	<0.05 cob	glu	

Country,	Applica	ation			PHI	Residues, as g	lufosinate free	acid, mg/kg ²	Ref
Year,	form	kg ai/ha	kg ai/hl	No ¹	days	Glufosinate	MPP	NAG	
(variety)								- 1	
Germany 1995	SL	0.45	0.15	2	0	18 shoot	0.1 shoot	glu	A56445
(LH82^4125sf^4SH				GS18	59	<0.05 shoot	<0.05 shoot	glu	ER95ECN550
-298)					77	0.05 shoot	<0.05 shoot	glu	DEU050102
G 1005	CI.	0.45	0.15	2	77	<0.05 cob	<0.05 cob	olu	156115
Germany 1995	SL	0.45	0.15	2	0	26 shoot	0.08 shoot	glu	A56445
$(LH82^{4}I25sf^{4}xF)$				GS18	48	0.34 shoot	< 0.05 shoot	glu	ER95ECN550
2)					90	0.25 shoot	< 0.05 shoot	glu	DE0010102
Commony 1006 (E2	C1	0.60	0.20	2	90	<0.05 C00	<0.05 c00	glu	A 59101
bybride IH 82)	SL	0.00	0.20	2 GS18	0 55	20 shoot	< 0.05 shoot	glu	AJ0191 ED06ECN550
ilyonue - L11-62)				0310	103	0.12 shoot	< 0.05 shoot	glu	DFU050102
					103	<0.05 cob	<0.05 cob	glu	DEC030102
Germany 1996 (F2	SL	0.80	0.27	2	0	39 shoot	0.09 shoot	glu	A58191
hybride - LH-82)	SE	0.00	0.27	- GS18	55	0.16 shoot	< 0.05 shoot	glu	ER96ECN550
				0.510	103	0.28 shoot	<0.05 shoot	glu	DEU050103
					103	<0.05 cob	<0.05 cob	glu	
Germany 1996 (F2	SL	0.45	0.15	2	0	23 shoot	0.05 shoot	glu	A58191
hybride - LH-82)				GS18	55	0.14 shoot	<0.05 shoot	glu	ER96ECN550
•					103	0.15 shoot	<0.05 shoot	glu	DEU050104
					103	<0.05 cob	<0.05 cob	glu	
Germany 1996	SL	0.60	0.20	2	0	24 shoot	0.07 shoot	glu	A58191
(Facet - transgen)				GS18	40	0.15 shoot	<0.05 shoot	glu	ER96ECN550
					69	0.17 shoot	<0.05 shoot	glu	DEU060102
					69	<0.05 cob	<0.05 cob	glu	
Germany 1996	SL	0.80	0.27	2	0	40 shoot	0.1 shoot	glu	A58191
(Facet - transgen)				GS18	40	0.18 shoot	<0.05 shoot	glu	ER96ECN550
					69	0.52 shoot	<0.05 shoot	glu	DEU060103
					69	<0.05 cob	<0.05 cob	glu	
Germany 1996	SL	0.45	0.15	2	0	31 shoot	0.08 shoot	glu	A58191
(Facet - transgen)				GS18	40	0.08 shoot	<0.05 shoot	glu	ER96ECN550
					69	0.15 shoot	<0.05 shoot	glu	DEU060104
a 100 f	a .	0.70	0.00		69	<0.05 cob	<0.05 cob	olu	
Germany 1996	SL	0.60	0.20	2	0	32 shoot	0.1 shoot	glu	A58191
(Facet - transgen)				G218	38 70	0.23 shoot	< 0.05 shoot	glu	EK96ECN550
					70	<0.05 cob	< 0.05 shoot	glu	DE0000202
Germany 1006	SI	0.80	0.27	2	0	<0.05 C00	<0.05 cob	glu	A 58101
(Facet - transgen)	SL	0.80	0.27	2 GS18	38	0.55 shoot	< 0.05 shoot	glu	FR96FCN550
(1 deet - transgen)				0510	50 70	0.33 shoot	<0.05 shoot	glu	DFU060203
					70	< 0.05 coh	<0.05 shoot	glu	DEC000205
Germany 1996	SL	0.45	0.15	2	0	23 shoot	0.08 shoot	glu	A58191
(Facet - transgen)	~=	01.10	0.12	- GS18	38	0.29 shoot	< 0.05 shoot	glu	ER96ECN550D
(70	0.9 shoot	<0.05 shoot	glu	EU060204
					70	<0.05 cob	<0.05 cob	glu	
Germany 1996	SL	0.60	0.20	2	0	34 shoot	0.17 shoot	glu	A58191
(LH82 T25 SF x				GS19	48	0.12 shoot	<0.05 shoot	glu	ER96ECN550
F2)					106	0.09 shoot	<0.05 shoot	glu	DEU010202
					106	<0.05 cob	<0.05 cob	glu	
Germany 1996	SL	0.80	0.27	2	0	45 shoot	0.17 shoot	glu	A58191
(LH82 T25 SF x				GS19	48	0.16 shoot	<0.05 shoot	glu	ER96ECN550
F2)					106	0.13 shoot	<0.05 shoot	glu	DEU010203
	~-				106	<0.05 cob	<0.05 cob	5 ^{ru}	
Germany 1996	SL	0.45	0.15	2	0	27 shoot	0.12 shoot	gh	A58191
(LH82 T25 SF x				GS19	48	0.08 shoot	<0.05 shoot	oh	ER96ECN550
F2)					106	0.06 shoot	<0.05 shoot	glu	DEU010204
a 100.6	GT	0.00	0.00		106	<0.05 cob	<0.05 cob	olu	4 50101
Germany 1996	SL	0.60	0.20	2	0	28 shoot	0.13 shoot	glu	A58191
(LH82 125 SF x Eq)				0819	50 112	U.1 shoot	<0.05 shoot	glu	EK90ECN550
re)					113	< 0.00 shoot	< 0.05 shoot	glu	DEU010102
			l	I	113	~0.0J COD	NUU COU		

Country,	Applic	ation			PHI	Residues, as g	lufosinate free	acid, mg/kg ²	Ref
Year, (variety)	form	kg ai/ha	kg ai/hl	No ¹	days	Glufosinate	MPP	NAG	
Germany 1996 (LH82 T25 SF x Fe)	SL	0.80	0.27	2 GS19	0 50 113 113	27 shoot 0.21 shoot 0.15 shoot <0.05 cob	0.14 shoot <0.05 shoot <0.05 shoot <0.05 cob	glu glu glu glu	A58191 ER96ECN550 DEU010103
Germany 1996 (LH82 T25 SF x Fe)	SL	0.45	0.15	2 GS19	0 50 113 113	21 shoot 0.07 shoot 0.05 shoot <0.05 cob	0.11 shoot <0.05 shoot <0.05 shoot <0.05 cob	glu glu glu glu	A58191 ER96ECN550 DEU010104
Italy 1994 ([B73(4)xT14]sfx - LM82)	SL	0.45	0.15	2 GS23	0 40 67 67	42 shoot 0.75 shoot 0.68 shoot <0.05 cob	0.17 shoot <0.05 shoot <0.05 shoot <0.05 cob	glu glu glu glu	A54230 ER94ECS550 ITA000102
Italy 1995 (Transgenic hybrid)	SL	0.45	0.15	2 GS18	0 45 77 77	10 shoot <0.05 shoot <0.05 shoot <0.05 cob	0.08 shoot <0.05 shoot <0.05 shoot <0.05 cob	4.3 shoot 0.22 shoot 0.16 shoot <0.05 cob	A54760 ER95ECS550 ITA000104
Italy 1995 (Transgenic hybrid)	SL	0.45	0.15	2 GS18	0 48 78 78	16 shoot <0.05 shoot <0.05 shoot <0.05 cob	0.12 shoot <0.05 shoot <0.05 shoot <0.05 cob	4.8 shoot 0.14 shoot 0.1 shoot <0.05 cob	A54760 ER95ECS550 ITA000204
Italy 1995 (Transgenic hybrid)	SL	0.45	0.15	2 GS18	0 45 77 77	7.8 shoot <0.05 shoot 0.07 shoot <0.05 cob	0.05 shoot <0.05 shoot <0.05 shoot <0.05 cob	4.1 shoot 0.21 shoot 0.45 shoot <0.05 cob	A54760 ER95ECS550 ITA000103
Italy 1995 (Transgenic hybrid)	SL	0.45	0.15	2 GS18	0 48 78 78	19 shoot <0.05 shoot <0.05 shoot <0.05 cob	0.13 shoot <0.05 shoot <0.05 shoot <0.05 cob	5.2 shoot 0.08 shoot 0.15 shoot <0.05 cob	A54760 ER95ECS550 ITA000203
Italy 1995 (Transgenic hybrid)	SL	0.45	0.15	2 GS18	0 45 77 77	11 shoot <0.05 shoot <0.05 shoot <0.05 cob	0.11 shoot <0.05 shoot <0.05 shoot <0.05 cob	3.4 shoot 0.11 shoot 0.12 shoot <0.05 cob	A54760 ER95ECS550 ITA000102
Italy 1995 (Transgenic hybrid)	SL	0.45	0.15	2 GS18	0 48 78 78	15 shoot <0.05 shoot <0.05 shoot <0.05 cob	0.17 shoot <0.05 shoot <0.05 shoot <0.05 cob	3.7 shoot 0.06 shoot 0.12 shoot <0.05 cob	A54760 ER95ECS550 ITA000202
Italy 1996 (Transgenic hybrid)	SL	0.80	0.27	2 GS18	0 38 64 64	46 shoot 0.76 shoot 1.0 shoot <0.05 cob	0.12 shoot <0.05 shoot <0.05 shoot <0.05 cob	glu glu glu glu	A58190 ER96ECS550 ITA000105
Italy 1996 (Transgenic hybrid)	SL	0.80	0.27	2 GS18	0 39 67 67	20 shoot 0.35 shoot 0.46 shoot <0.05 cob	0.05 shoot <0.05 shoot <0.05 shoot <0.05 cob	glu glu glu glu	A58190 ER96ECS550 ITA000205
Italy 1996 (transgenic hybrid)	SL	0.60	0.20	2 GS18	0 38 64 64	25 shoot 0.3 shoot 0.57 shoot <0.05 cob	0.09 shoot <0.05 shoot <0.05 shoot <0.05 cob	glu glu glu glu	A58190 ER96ECS550 ITA000103
Italy 1996 (transgenic hybrid)	SL	0.80	0.27	2 GS18	0 38 64 64	37 shoot 0.47 shoot 1.1 shoot <0.05 cob	0.11 shoot <0.05 shoot <0.05 shoot <0.05 cob	glu glu glu glu	A58190 ER96ECS550 ITA000104
Italy 1996 (transgenic hybrid)	SL	0.60	0.20	2 GS18	0 39 67 67	15 shoot 0.21 shoot 0.3 shoot <0.05 cob	0.08 shoot <0.05 shoot <0.05 shoot <0.05 cob	glu glu glu glu	A58190 ER96ECS550 ITA000203
Italy 1996 (transgenic hybrid)	SL	0.80	0.27	2 GS18	0 39 67 67	22 shoot 0.36 shoot 0.71 shoot <0.05 cob	0.1 shoot <0.05 shoot <0.05 shoot <0.05 cob	glu glu glu glu	A58190 ER96ECS550 ITA000204

Country,	Applica	ation			PHI	Residues, as g	lufosinate free	acid, mg/kg ²	Ref
Year,	form	kg ai/ha	kg ai/hl	No ¹	days	Glufosinate	MPP	NAG	
(variety)									
Italy 1996	SL	0.80	0.27	2	0	21 shoot	0.05 shoot	glu	A58190
(Transgenic hybrid)				GS19	38	0.18 shoot	<0.05 shoot	glu	ER96ECS550
					54	0.79 shoot	<0.05 shoot	glu	ITA000305
					54	<0.05 cob	<0.05 cob	glu	
Italy 1996	SL	0.60	0.20	2	0	19 shoot	0.09 shoot	giu	A58190
(transgenic hybrid)				GS19	38	0.58 shoot	<0.05 shoot	glu	ER96ECS550
					54	0.56 shoot	<0.05 shoot	glu	ITA000303
				-	54	<0.05 cob	<0.05 cob	glu	
Italy 1996	SL	0.80	0.27	2	0	20 shoot	0.09 shoot	glu	A58190
(transgenic hybrid)				GS19	38	0.43 shoot	<0.05 shoot	glu	ER96ECS550
					54	0.7 shoot	<0.05 shoot	glu	ITA000304
	a.		0.15.0.00	-	54	<0.05 cob	<0.05 cob	olu	- ID
Netherlands 1995	SL	0.9	0.45-0.23		71	0.41 shoot	<0.05 shoot	glu	6 HMA 52-509
(Hudson)	a.		0.15.0.00	(6 leaf)	/1	<0.05 cob	<0.05 cob	olu	- ID
Netherlands 1995	SL	0.9	0.45-0.23	1	85	0.81 shoot	0.05 shoot	glu	6 HMA 52-510
(Scarlet)	~~			(5 leaf)	85	<0.05 cob	<0.05 cob	glu	
Netherlands 1995	SL	0.9	0.45-0.23	1	72	1.7 shoot	0.24 shoot	olu	6 HMA 52-264
(Scarlet)	~~			(9 leaf)	72	0.06 cob	0.07 cob	glu	
Spain 1994 (LH119-	SL	0.45	0.15	2	0	27 shoot	0.28 shoot	olu	A54230
6xT14/LH82)				GS25	40	0.14 shoot	<0.05 shoot	glu	ER94ECS550
					70 70	0.05 shoot	<0.05 shoot	glu	ESP000102
g : 1005	CI.	0.45	0.15	2	/0	<0.05 cob	<0.05 cob	401	154760
Spain 1995	SL	0.45	0.15	2	0	24 shoot	0.46 shoot	4.2 shoot	A54/60
(Transgenic hybrid)				GS18	46 57	< 0.05 shoot	<0.05 shoot	0.08 shoot	EK95EC5550
					57	<0.07 shoot	< 0.05 shoot	< 0.25 shoot	ESP000105
Spain 1005	C1	0.45	0.15	2	57	<0.03 C00	< 0.03 cou	< 0.05 cob	151760
(Transgenic hybrid)	SL	0.45	0.15	∠ G\$18	0 46	0.05 shoot	< 0.39 shoot	0.15 shoot	A34700 FR95EC\$550
(Transgenie nyond)				0510	40 57	0.03 shoot	< 0.05 shoot	0.13 shoot	EK75EC5550 ESP000102
					57	< 0.05 cob	< 0.05 shou	< 0.05 coh	L51 000102
Spain 1996	SL.	0.45	0.15	2	0	14 shoot	0.06 shoot	glu	A 58190
(transgenic hybrid)	5L	0.15	0.15	- GS18	36	0.25 shoot	<0.05 shoot	glu	ER96ECS550
())					54	0.28 shoot	<0.05 shoot	glu	ESP000102
					54	<0.05 cob	<0.05 cob	glu	
Spain 1996	SL	0.60	0.20	2	0	22 shoot	0.07 shoot	glu	A58190
(transgenic hybrid)				GS18	36	0.62 shoot	<0.05 shoot	glu	ER96ECS550
(****8***)***)					54	0.31 shoot	<0.05 shoot	glu	ESP000103
					54	<0.05 cob	<0.05 cob	glu	
Spain 1996	SL	0.80	0.27	2	0	19 shoot	0.07 shoot	glu	A58190
(transgenic hybrid)				GS18	36	0.9 shoot	<0.05 shoot	glu	ER96ECS550
					54	0.54 shoot	<0.05 shoot	glu	ESP000104
					54	<0.05 cob	<0.05 cob	glu	
Spain 1996	SL	0.45	0.15	2	0	12 shoot	0.06 shoot	glu	A58190
(transgenic hybrid)				GS18	35	0.66 shoot	<0.05 shoot	glu	ER96ECS550
					55	0.54 shoot	<0.05 shoot	glu	ESP000202
					55	<0.05 cob	<0.05 cob	glu	
Spain 1996	SL	0.60	0.20	2	0	17 shoot	0.06 shoot	glu	A58190
(transgenic hybrid)				GS18	35	1.2 shoot	<0.05 shoot	glu	ER96ECS550
					55	0.72 shoot	<0.05 shoot	giu	ESP000203
				_	55	<0.05 cob	<0.05 cob	glu	
Spain 1996	SL	0.80	0.27	2	0	24 shoot	0.08 shoot	giu	A58190
(transgenic hybrid)				GS18	35	0.88 shoot	<0.05 shoot	giu glu	ER96ECS550
					55	0.48 shoot	<0.05 shoot	oh	ESP000204
	~-				55	<0.05 cob	<0.05 cob	o-"	
Spain 1996	SL	0.45	0.15	2	0	25 shoot	0.18 shoot	oh	A58190
(transgenic hybrid)				GS18	46	0.33 shoot	0.08 shoot	glu	ER96ECS550
					59 50	0.47 shoot	0.05 shoot	glu	ESP000302
					39	<0.05 cob	<0.05 cob	0 -	

Country, Application					PHI	Residues, as g	acid, mg/kg ²	Ref	
Year, (variety)	form	kg ai/ha	kg ai/hl	No ¹	days	Glufosinate	MPP	NAG	
Spain 1996 (transgenic hybrid)	SL	0.60	0.20	2 GS18	0 46 59 59	36 shoot 0.66 shoot 0.59 shoot <0.05 cob	0.25 shoot 0.05 shoot <0.05 shoot <0.05 cob	glu glu glu glu	A58190 ER96ECS550 ESP000303
Spain 1996 (transgenic hybrid)	SL	0.80	0.27	2 GS18	0 46 59 59	28 shoot 0.8 shoot 1.0 shoot <0.05 cob	0.23 shoot 0.05 shoot 0.07 shoot <0.05 cob	glu glu glu glu	A58190 ER96ECS550 ESP000304

¹. GS: growth stage at final treatment

GS18 8 leaves unfolded GS19

9 or more leaves unfolded GS23

GS24

5th leaf unfolded 6th leaf unfolded 7th leaf unfolded GS25

 2 shoot: stems + leaves.

glu: included in glufosinate-ammonium result.

Table 54. Glufosinate residues in maize forage and fodder resulting from supervised trials in the USA and Argentina. Analyses on replicate samples are reported individually. Double-underlined residues are from treatments according to GAP and were used to estimate maximum residue levels.

Location,	Application				PHI Residues, mg/kg, as glufosinate free acid ³						Ref
Year,	form	kg	kg ai/hl	No ¹	days	Commo	glufosinate-	MPP	NAG		
(variety)		ai/ha	-			dity ²	ammonium				
USA (IA) 1993	SL	0.36		1	30	Forage	0.05 0.08 0.09	NQ (3)	0.36 0.49	0.45	ER-93-
(transgenic (LH216) X				(30 cm)	60	silage	NQ (3)	NQ (3)	0.18 0.19	0.17	USA-01-IA-
(LH119) (4) × T14)					118	fodder	NQ (3)	NQ (3)	0.15 0.19 0.	.19	01
USA (IA) 1993	SL	0.50		1	30	forage	$0.07\ 0.08\ 0.18$	NQ (3)	0.36 0.49	0.45	ER-93-
(transgenic (LH216) X				(30 cm)	60	silage	NQ (3)	NQ (3)	0.14 0.12	0.16	USA-01-IA-
(LH119) (4) × T14)					118	fodder	NQ (2) 0.05	NQ (3)	0.20 (3)		01
USA (IA) 1993	SL	0.50		1	30	forage	0.12 0.12 0.07	NQ (3)	0.80 0.75	0.51	ER-93-
(transgenic (LH216) X				(60 cm)	62	silage	0.12 <u>0.13</u> 0.08	NQ (3)	0.71 <u>0.94</u>	0.47	USA-01-IA-
(LH119) (4) × T14)					114	fodder	<u>0.13</u> 0.15 0.12	NQ 0.05 NQ	<u>0.81</u> 0.64 0.	.48	01
USA (IL) 1993	SL	0.36	0.39	1	30	forage	NQ (3)	NQ (3)	0.10 0.06	NQ	ER-93-
(transgenic (LH 59x				(30 cm)	60	silage	NQ (3)	NQ (3)	NQ	(3)	USA-01-IL-
LH51)(LH119) (4) X					100	fodder	NQ (3)	NQ (3)	NQ (3)		01
T14)											
USA (IL) 1993	SL	0.50	0.54	1	30	forage	NQ (3)	NQ (3)	0.06 0.18	0.06	ER-93-
(transgenic (LH 59x				(30 cm)	60	silage	NQ (3)	NQ (3)	NQ	(3)	USA-01-IL-
LH51)(LH119) (4) X					100	fodder	NQ (3)	NQ (3)	NQ (3)		01
T14)											
USA (IL) 1993	SL	0.50	0.54	1	30	forage	NQ 0.06 0.08	NQ (3)	0.19 0.48	0.47	ER-93-
(transgenic (LH 59x				(60 cm)	61	silage	<u>NQ</u> (3)	NQ (3)	<u>0.20</u> 0.17	0.17	USA-01-IL-
LH51)(LH119) (4) X					95	fodder	<u>NQ</u> (3)	NQ (3)	0.21 0.16 <u>0</u> .	25	01
T14)											
USA (IL) 1993	SL	0.36	0.39	2	30	forage	0.13 0.14 0.13	NQ (3)	0.64 0.76	0.51	ER-93-
(transgenic (LH 59x		+0.50	+0.54	(60 cm)	6195	silage	0.07 <u>0.09</u> 0.07	NQ (3)	0.47 <u>0.69</u>	0.51	USA-01-IL-
LH51)(LH119) (4) X						fodder	<u>NQ</u> (3)	NQ (3)	0.13 0.12 <u>0.</u>	15	01
T14)											
USA (NE) 1993	SL	0.36		1	30	forage	NQ (3)	NQ (3)	NQ	(3)	ER-93-
(transgenic (LH 59x				(30 cm)	72	silage	NQ (3)	NQ (3)	NQ	(3)	USA-01-
LH51)(LH119) (4) X					107	fodder	NQ (3)	NQ (3)	NQ (3)		NE-01
T14)											
USA (NE) 1993	SL	0.50	0.48	1	30	forage	NQ (3)	NQ (3)	0.09 0.05	0.14	ER-93-
(transgenic (LH 59x				(30 cm)	72	silage	NQ (3)	NQ (3)	NQ	(3)	USA-01-
LH51)(LH119) (4) X					107	fodder	NQ (3)	NQ (3)	NQ (3)		NE-01
T14)											

Year, Variety Form (a) Ref Ref Ref Ref Ref Ref USA (ND 1993) (LISA) (LII1) (LII1) (A) K. 0.50 0.44 0.50 0.44 0.54 0.56 0.58 0.57 0.58 <th>Location,</th> <th colspan="3">Application</th> <th></th> <th>PHI</th> <th></th> <th>Residues, mg/kg</th> <th>Ref</th>	Location,	Application				PHI		Residues, mg/kg	Ref		
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	Year,	form	kg	kg ai/hl	No^1	days	Commo	glufosinate-	MPP	NAG	
USA (NS) 1993 SL 0.50 0.48 1 0.60	(variety)	~~	ai/ha	0.40			dity ²	ammonium			
Har Big (L11) 9(A) X No. (3) OS (2)	USA (NE) 1993 (transgonia (LH 50y	SL	0.50	0.48	1	30 60	forage	$0.16\ 0.20\ 0.15$	NQ (3)	1.4 1.3 1.2	ER-93-
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	(LH 39X)				(60 cm)	92	fodder	$0.03 \frac{0.03}{0.03} 0.00$	1100000000000000000000000000000000000	0.30 <u>0.48</u> 0.40 0.68 0.58 0.57	USA-01- NE-01
$ \begin{array}{c} \begin{tabular}{ c c c c c c c c c c c c c c c c c c c$	T14)					95	Touter	<u>0.10</u> 0.08 0.14	<u>0.11</u> 0.09 0.09	<u>0.08</u> 0.38 0.37	NL-01
$ \begin{array}{c} (armsenic [LH1]9(4) \\ VTH3 KLIP1(5) \\ (armsenic [LH1]9(4) \\ VTH3 KLIP1(5) \\ (armsenic [LH1]9(4) \\ XTH3 KLIP1(5) \\ (armsenic [LH1]9(4) \\ XTH3 KLIP1(5) \\ (armsenic [LH1]9(4) \\ VTH3 KLIP1(5) \\ (armsenic [LH1]9(4) \\ (armsenic [LH1]9(4) \\ (b, ATH3 KLIP1(5) \\ (armsenic [LH1]9(4) \\ (a$	USA (IN) 1993	SL	0.36	0.39	1	30	forage	NO 0.07 0.05	NO (3)	0.34 0.46 0.40	ER-93-
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	(transgenic [LH119(4)	~-			(30 cm)	69	silage	NQ (2) 0.06	NQ (3)	0.18 0.18 0.30	USA-01-
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	X T14] X LH216)					115	fodder	0.11 NQ 0.05	0.06 NQ (2)	0.25 0.12 0.12	IN-01
$ \begin{array}{c} (30 \ cm) \ [6p] \ silage \\ (30 \ cm) \ [6p] \ silage \\ (30 \ cm) \ [6p] \ silage \\ (31 \ cm) \ [6p] \ silage \\ (31 \ cm) \ [6p] \ (31 \ c$	USA (IN) 1993	SL	0.50	0.54	1	30	forage	0.06 0.05 0.05	NQ (3)	0.42 0.40 0.46	ER-93-
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	(transgenic [LH119(4)				(30 cm)	69	silage	NQ 0.09 NQ	NQ (3)	0.22 0.14 0.20	USA-01-
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	X T14] X LH216)				_	115	fodder	0.05 NQ 0.05	NQ (2) 0.06	0.13 0.11 0.15	IN-01
$ \begin{array}{c} \mbox{dist} (L1119(4)) & (-1.3)$	USA (IN) 1993 (transgania [L 1110(4)	SL	0.36	0.39	2 (60 am)	30 60	forage	0.26 0.25 0.23	0.06 NQ (2)	1.6 1.5 1.4	ER-93-
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	(Tallsgellic [LH119(4)] X T14] X LH216)		+0.50	+0.54	(00 cm)	106	fodder	$0.22\ 0.18\ 0.21$ $0\ 32\ 0\ 31\ 0\ 45$	0.09 NQ(2) 0.18.0.13.0.16	1.4 1.5 1.9 1 2 1 4 1 7	USA-01- IN-01
$ \begin{array}{c} (ransgenic (LH85 X \\ LH160) (LH19)(4) X \\ T14) \\ USA (ND) 1993 \\ UTA (SD) 1993 \\ USA (ND) 1993 \\ USA ($	USA (ND) 1993	SL	0.36	0.39	1	30	forage	NO (3)	NO (3)	0.19 0.20 0.23	ER-93-
$ \begin{array}{ c c c c c c c c c c c c c c c c c c c$	(transgenic (LH85 X				(30 cm)	80	silage	NQ (3)	NQ (3)	0.06 NQ NQ	USA-01-
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	LH160) (LH119)(4) X					107	fodder	NQ (3)	NQ (3)	NQ (3)	ND-01
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	T14)										
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	USA (ND) 1993	SL	0.50	0.54	1	30	forage	0.07 0.06 NQ	NQ (3)	0.35 0.29 0.24	ER-93-
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	(transgenic (LH85 X)				(30 cm)	80	silage	NQ (3)	NQ (3)	NQ 0.05 NQ	USA-01-
$ \begin{array}{c} 1.123 (ND) 1993 \\ (Itransgenic (LH85 X) \\ (LH160) (LH19)(4) X \\ T14) \\ (Itransgenic (LH216) X \\ (LH19(4) X (T14)) \\ (Itransgenic (LH216) X \\ (Itransgenic (LH216) X$	$L \Pi 100) (L \Pi 119)(4) \Lambda$ T14)					107	Todder	NQ (3)	NQ (3)	NQ (3)	ND-01
	USA (ND) 1993	SL	0.36	0.39	2	30	forage	NO (3)	NO (3)	0.29 0.22 0.20	ER-93-
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	(transgenic (LH85 X	52	+0.50	+0.54	(60 cm)	72	silage	<u>NO</u> (3)	NQ (3)	0.12 0.12 0.10	USA-01-
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	LH160) (LH119)(4) X					97	fodder	<u>NQ</u> (3)	NQ (3)	0.07 <u>0.11</u> 0.08	ND-01
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	T14)										
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	USA (MO) 1993	SL	0.36	0.39	1	30	forage	0.07 NQ NQ	NQ (3)	0.22 0.16 0.14	ER-93-
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	(transgenic(LH216) X				(30 cm)	64 07	silage	NQ (3)	NQ (3)	0.06 NQ NQ	USA-01-
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	$\frac{(LH119(4) X (114))}{USA (MO) 1002}$	CI	0.26	0.20	2	97	fodder	NQ (3)	NQ (3)	NQ(3)	MO-01
$\begin{array}{c c c c c c c c c c c c c c c c c c c $	(transgenic(I H216) X	SL	± 0.50	0.39 ±0.54	(60 cm)	50 60	silage	$0.33\ 0.20\ 0.17$ $0.26\ 0.11\ 0.08$	NQ(3)	0.90 1.1 0.90 1 5 0.89 0.70	EK-93- USA-01-
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	(LH119(4) X (T14))		10.50	10.54	(00 cm)	95	fodder	$0.10\ 0.10\ 0.12$	NO (3)	$0.46\ 0.52\ 0.60$	MO-01
	USA (CA) 1993	SL	0.36	0.39	1	30	forage	NQ (3)	NQ (3)	0.13 0.11 0.08	ER-93-
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	(transgenic (LH59 X				(30 cm)	73	silage	NQ (3)	NQ (3)	NQ (3)	USA-01-
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	LH51) (LH119)(4) X					119	fodder	NQ (3)	NQ (3)	NQ (3)	CA-01
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	T14)	CT.	0.26	0.20	2	20	c	0.12.0.12	0.00.0.11	0.52 0.54	ED 02
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	USA (CA) 1993 (transgenic (LH59 X	SL	0.36 ⊥0.50	0.39 ±0.54	2 (60 cm)	30 71	ilage	0.12 0.13 NO (3)	0.08 0.11 NO (3)	0.52 $0.540.10 0.26 0.16$	ER-93- USA-01-
$\begin{array}{c c c c c c c c c c c c c c c c c c c $	(Hansgeine (E115) X LH51) (LH119)(4) X		10.50	10.54	(00 cm)	106	fodder	0.09010009	0.13.0.10.0.10	0.55 0.60 0.59	CA-01
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	T14)					100	rouder	<u></u> 0.07	0110 <u></u> 0110	0.00	0.1.01
	USA (SD) 1993	SL	0.36	0.71	1	30	forage	0.12 NQ NQ	NQ (3)	0.14 0.15 0.13	ER-93-
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	(transgenic (LH85 X				(30 cm)	71	silage	NQ (3)	NQ (3)	NQ (3)	USA-01-
$\begin{array}{c c c c c c c c c c c c c c c c c c c $	LH160) (LH119)(4) X					95	fodder	NQ (3)	NQ (3)	0.07 0.08 0.08	SD-01
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	114))	SI	0.50	0.00	1	20	forage	0.05.0.07.0.06	NO (2)	0.21 0.44 0.25	ED 02
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	(transgenic (LH85 X	SL	0.50	0.99	(30 cm)	30 71	silage	NO (3)	NQ (3)	$0.21 \ 0.44 \ 0.23$ $0.09 \ 0.09 \ 0.10$	USA-01-
$\begin{array}{c ccccccccccccccccccccccccccccccccccc$	LH160) (LH119)(4) X				(2000)	95	fodder	NO NO 0.05	NQ (3)	0.20 0.14 0.22	SD-01
$ \begin{array}{c} \text{USA (SD) 1993} \\ (\text{transgenic (LH85 X)} \\ \text{LH160) (LH119)(4) X} \\ \text{USA (VA) 1993} \\ (\text{transgenic (LH216)} \\ (\text{LH119)(4) X T14)} \end{array} \\ \begin{array}{c} \text{SL} \\ \text{usage} \\ \text{obs} \\ o$	T14))								~~~		
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	USA (SD) 1993	SL	0.36	0.71	2	30	forage	0.29 0.23 0.19	NQ (3)	0.89 1.4 1.2	ER-93-
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	(transgenic (LH85 X		+0.50	+0.99	(60 cm)	60	silage	<u>0.08</u> 0.07 NQ	NQ (3)	<u>0.09</u> 0.09 0.10	USA-01-
114)) USA (VA) 1993 SL 0.36 0.71 1 33 forage 0.05 NQ 0.06 NQ (3) 0.41 0.22 0.33 ER-93- (LH119)(4) X T14) (30 cm) 60 silage NQ (3) NQ (3) 0.46 0.29 0.35 ER-93- (LH119)(4) X T14) (30 cm) 60 silage NQ NQ NQ NQ (3) 0.46 0.29 0.35 ER-93- (transgenic (LH216) (LH119)(4) X T14) 0.50 0.99 1 33 forage 0.07 NQ NQ NQ (3) 0.46 0.29 0.35 ER-93- (LH119)(4) X T14) 0.50 0.99 1 33 forage 0.07 NQ NQ NQ (3) 0.30 0.12 0.24 USA-01- USA (VA) 1993 SL 0.36 0.71 2 30 forage 0.31 0.36 0.41 0.22 0.35 ER-93- (transgenic (LH216) 10.50 +0.59 (60 cm) 61 silage 0.49 0.64 0.17 0.30 0.12 0.24 USA-01- (LH119)(4) X T14) 0	LH160) (LH119)(4) X T14))					95	fodder	<u>0.20</u> 0.09 0.20	0.06 NQ 0.07	<u>1.7</u> 0.79 1.6	SD-01
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	114))	SI	0.26	0.71	1	22	forage	0.05 NO.0.06	NO (2)	0 41 0 22 0 22	ED 02
$\begin{array}{c c c c c c c c c c c c c c c c c c c $	(transgenic (LH216)	SL	0.30	0.71	(30 cm)	55 60	silage	NO (3)	NQ (3)	$0.41 \ 0.22 \ 0.33$ $0.15 \ 0.11 \ 0.08$	USA-01-
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	(LH119)(4) X T14)				(00 011)	00	Siluge			0.110 0.111 0.000	VA-01
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	USA (VA) 1993	SL	0.50	0.99	1	33	forage	0.07 NQ NQ	NQ (3)	0.46 0.29 0.35	ER-93-
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	(transgenic (LH216)				(30 cm)	60	silage	NQ NQ 0.06	NQ (3)	0.30 0.12 0.24	USA-01-
$ \begin{array}{c ccccccccccccccccccccccccccccccccccc$	(LH119)(4) X T14)	~~			-	106	fodder	NQ NQ	0.11 0.07	0.10 0.07	VA-01
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	USA (VA) 1993	SL	0.36	0.71	2 (60)	30 61	torage	0.31 0.36 0.44	NQ (3) 0 17 0 24 0 20	1.9 2.4 3.1	ER-93-
$\begin{array}{c ccccccccccccccccccccccccccccccccccc$	(LH_{16}) (LH119)(4) X T14)		+0.50	+0.99	(00 cm)	01 97	fodder	0.49 <u>0.64</u> 0.44	0.17 0.54 0.20 0.41 0.43	2.4 <u>2.3</u> 2.1	USA-01- VA-01
$\begin{array}{c ccccccccccccccccccccccccccccccccccc$	USA (MN) 1993	SL	0.52	1.1	1	30	forage	0.34 0.27 0.25	NO (3)	1.5 1.5 1.2	ER-93-
(LH119)(4) X T14) MN-01	((LH85 X LH160)		5.02		(76 cm)	66	silage	0.20 0.20 0.28	NQ (3)	1.3 0.99 1.4	USA-01-
	(LH119)(4) X T14)						_				MN-01

Location,	Application			PHI		Residues, mg/kg, as glufosinate free acid ³				
Year,	form	kg	kg ai/hl	No ¹	days	Commo	glufosinate-	MPP	NAG	
(variety)	CI	ai/ha	0.52	1	20	dity	ammonium	NO (2)	0.00 0.12 0.11	ED 02
(transgenic (LH59 X	SL	0.50	0.55	(12 cm)	20 30	forage	NQ (3)	NQ (3)	0.09 0.12 0.11 NO (3)	EK-95- USA-01-
LH51) (LH119)(4) X				()	40	forage	NQ (3)	NQ (3)	NQ (3)	MO-02
(T14))					60	silage	NQ (3)	NQ (3)	NQ (3)	
	CI	0.50	0.52	1	123	fodder	NQ (3)	NQ (3)	NQ (3)	ED 02
USA (MO) 1993 (transgenic (LH59 X	SL	0.50	0.53	1 (28 cm)	20 30	forage	NQ (3) NO (3)	NQ (3) NO (3)	0.19 0.14 0.06 0.06 0.19 NO	ER-93- USA-01-
LH51) (LH119)(4) X				(20 011)	40	forage	NQ (3)	NQ (3)	NQ (3)	MO-02
(T14))					60	silage	NQ (3)	NQ (3)	NQ (3)	
	CT.	0.50	0.52	1	118	fodder	NQ (3)	NQ (3)	NQ (3)	ED 02
USA (MO) 1993 (transgenic (LH59 X	SL	0.50	0.53	1 (46 cm)	20 30	forage	0.09 0.08 0.07 0.06 0.05 NO	NQ (3) NO (3)	$0.68 \ 0.55 \ 0.46$ $0.29 \ 0.23 \ 0.16$	ER-93- USA-01-
LH51) (LH119)(4) X				(40 cm)	40	forage	NQ (3)	NQ (3)	$0.29 \ 0.23 \ 0.10$ $0.06 \ 0.12 \ 0.11$	MO-02
(T14))					60	silage	NQ (3)	NQ (3)	0.13 NQ NQ	
					111	fodder	NQ (3)	NQ (3)	NQ (3)	
USA (MO) 1993 (transgonia (LH50 V	SL	0.50	0.53	1 (61 am)	20	forage	$0.41\ 0.40\ 0.40$	NQ (3) 0.05 NO NO	2.7 3.3 1.9	ER-93-
(11139 X) LH51) (LH119)(4) X				(01 cm)	30 40	forage	$0.27 \ 0.37 \ 0.33$ $0.50 \ 0.22 \ 0.44$	NO (3)	2.5 1.2 2.2	MO-02
(T14))					60	silage	<u>0.14</u> 0.14 0.14	<u>0.31</u> 0.45 0.34	<u>1.0</u> 0.63 0.74	
					101	fodder	<u>0.23</u> 0.20 0.18	<u>0.05</u> NQ NQ	<u>0.91</u> 0.71 0.79	
USA (MO) 1993	SL	0.50	0.53	1	20	forage	0.17 0.11 0.14	NQ (3)	0.77 0.67 0.82	ER-93-
(transgenic (LH59 X) LH51) (LH119)(4) X				(86 cm)	30 40	forage	$0.11\ 0.14\ 0.06$ $0\ 11\ 0\ 11\ 0\ 08$	NQ (3) NO (3)	0.74 0.90 $0.520.73 0.87 0.57$	USA-01- MO-02
(T14))					40 61	silage	NO 0.05 NO	NQ (3)	$0.31 \ 0.45 \ 0.34$	10-02
< <i>"</i>					97	fodder	0.13 0.06 0.08	NQ (3)	0.31 0.27 0.30	
USA (CA) 1993	SL	0.50	0.53	1	20	forage	NQ (3)	NQ (3)	NQ (3)	ER-93-
(transgenic (LH59 X				(13-15	30	forage	NQ (3)	NQ (3)	NQ (3)	USA-01-
LH51) (LH119)(4) X (T14))				cm)	40 60	forage	NQ (3) NO (3)	NQ (3) NO (3)	NQ (3) NO (3)	CA-02
(114))					127	fodder	NQ (3)	NQ (3)	NQ (3)	
USA (CA) 1993	SL	0.50	0.53	1	20	forage	0.25 0.20 0.24	0.12 0.14 0.11	1.3 0.87 1.2	ER-93-
(transgenic (LH59 X				(25-33	30	forage	0.06 NQ 0.05	NQ 0.05 NQ	0.25 0.23 0.35	USA-01-
LH51) (LH119)(4) X				cm)	40	forage	NQ (3)	NQ (3)	$0.11 \ 0.14 \ 0.13$	CA-02
(114))					120	fodder	NQ (3)	NQ (3)	NQ (3) 0.07 NO 0.06	
USA (CA) 1993	SL	0.50	0.53	1	20	forage	0.24 0.29 0.18	0.08 0.08 0.06	1.2 1.4 1.3	ER-93-
(transgenic (LH59 X				(33-51	30	forage	0.09 0.16 0.13	NQ NQ 0.05	0.48 0.62 0.61	USA-01-
LH51) (LH119)(4) X				cm)	40	forage	NQ 0.09 0.08	NQ (3)	0.31 0.43 0.34	CA-02
(114))					$60 \\ 114$	silage	NQ (3) NO (3)	NQ (3) NO NO 0 07	0.09 0.06 0.08	
USA (CA) 1993	SL	0.50	0.53	1	20	forage	0.46042051	0.11.0.10.0.09	18 16 14	ER-93-
(transgenic (LH59 X		0.00	0.00	(41-81	30	forage	0.44 0.31 0.33	0.11 0.10 0.08	1.5 1.4 1.6	USA-01-
LH51) (LH119)(4) X				cm)	40	forage	0.16 0.16 0.18	0.07 0.07 0.08	0.77 0.65 0.86	CA-02
(T14))					60 06	silage	NQ 0.09 <u>0.17</u>	0.06 NQ <u>0.07</u> 0.12 0.10 0.00	$0.21 \ 0.30 \ 0.55$	
USA (CA) 1993	SI	0.50	0.53	1	20	forage	<u>0.23</u> 0.26 0.09 0 29 0 36 0 34	0.06.0.07.0.06	<u>1.4</u> 0.88 0.00 0.88 0.96 0.95	FR-93-
(transgenic (LH59 X	5L	0.50	0.55	1 66-117	30	forage	0.21 0.34 0.19	0.07 0.06 0.05	$0.84 \ 0.89 \ 0.54$	USA-01-
LH51) (LH119)(4) X				cm)	40	forage	0.11 0.14 0.22	NQ (3)	0.60 0.49 0.67	CA-02
(T14))					60	silage	0.08 <u>0.09</u> NQ	NQ (3)	$0.33 \ \underline{0.59} \ 0.20$	
USA (EL) 1004	CI.	0.40	0.42	2	96	fodder	<u>0.18</u> 0.18 0.23	<u>0.07</u> 0.06 0.07	<u>2.4</u> 1.5 1.5 ^{glu}	DV 04D
(transgenic (LH51 X	SL	+0.50	+0.54	2 (2-14	29 60	silage	0.70	NO	glu	DK-94K- 01-FL-01
LH210) (LH119(4) X				leaves)	69	fodder	1.0	0.21	glu	
T14)									-1-	
USA (IA) 1994	SL	0.41	0.44	2	31	forage	0.66	NQ	giu elu	BK-94R-
(transgenic (LH59 X) (LH51) (LH119(4) X)		+0.55	+0.57	(61 cm)	100	forage	0.30	NQ	glu	01-IA-01
T14))		L			122	fodder	0.25	NQ	glu	
USA (IL) 1994	SL	0.40	0.43	2	31	forage	0.26	NQ	glu	BK-94R-
(transgenic (LH59 X		+0.50	+0.54	(61 cm)	60	silage	0.11	NQ	giu ơlu	01-IL-01
LH51) (LH119(4) X					76 110	forage	0.12	NQ	glu	
114)) USA (MN) 1994	SI	0.38	0.41	2	31	forage	<u>9.12</u> 1.1	NO	glu	BK-94P-
(transgenic (LH74)	50	+0.48	+0.52	- (61 cm)	60	silage	1.1	NQ	glu	01-MN-01
(LH82(3) X T14))					93	forage	0.91	NQ	glu	
					114	fodder	0.99	0.14	giù	

Location,	Application			PHI		Ref				
Year,	form	kg	kg ai/hl	No ¹	days	Commo	glufosinate-	MPP	NAG	
(variety)		ai/ha	-		•	dity ²	ammonium			
USA (MO) 1994	SL	0.40	0.43	2	30	forage	1.6	NQ	glu	BK-94R-
(transgenic (LH59 X		+0.53	+0.57	(61 cm)	60	silage	1.2	NQ	glu	01-MO-01
LH51) (LH119(4) X					69	forage	0.70	0.08	giu giu	
T14))	~~			-	89	fodder	<u>1.5</u>	0.26	ghu	
USA (NC) 1994 (transgenie (LU51 V	SL	0.38	0.41	2 (61 am)	30 50	forage	0.21	NQ	glu	BK-94R-
(transgenic (LH51 X)		+0.52	+0.56	(61 cm)	28 60	forage	0.10	NQ	glu	01-NC-01
T(4) X = T(4)					92	fodder	0.07	NO	glu	
USA (NY) 1994	SL	0.38	0.41	2	30	forage	4.0	NO	glu	BK-94R-
(transgenic (LH74)	22	+0.53	+0.57	- (61 cm)	69	silage	2.1	NO	glu	01-NY-01
(LH82(3) X T14))				` '	77	forage	2.9	NQ	glu	
					88	fodder	5.3	0.1	glu	
USA (TX) 1994	SL	0.38	0.41	2	30	forage	0.44	NQ	glu	BK-94R-
(transgenic (LH51 X		+0.46	+0.49	(61 cm)	60	silage	0.32	NQ	glu	01-TX-01
LH210) (LH119(4) X					74	forage	0.27	NQ	giu giu	
T14))	~~			-	97	fodder	0.26	<u>0.07</u>	ghu	
USA (WA) 1994	SL	0.40	0.43	$\frac{2}{(1 \text{ and})}$	30	forage	0.77	0.09 NO	glu	BK-94R-
(transgenic(LH/4))		+0.50	+0.54	(61 cm)	00 76	forage	0.24	NQ 0.07	glu	01-WA-01
$(L1102)3) \times 114))$					86	fodder	0.56	0.12	glu	
USA (WI) 1994	SL.	0.38	0.41	2	30	forage	0.61	NO	glu	BK-94R-
(transgenic (LH85 X	5L	+0.56	+0.60	(61 cm)	60	silage	0.36	NO	glu	01-WI-01
LH160) (LH119(4) X				()	92	forage	0.33	NQ	glu	
T14))					107	fodder	0.43	NQ	glu	
USA (FL) 1994	SL200	0.40	0.43	2	29	forage	1.8	NQ	glu	BK-94R-
(transgenic (LH51 X		+0.50	+0.54	(2-14	60	silage	0.84	NQ	glu	01-FL-01
LH210) (LH119(4) X				leaves)	69	fodder	1.2	0.29	glu	
T14)									du	
USA (IA) 1994	SL	0.41	0.44	2	31	forage	0.53	NQ	giu	BK-94R-
(transgenic (LH59 X		+0.51	+0.55	(61 cm)	60 100	silage	0.40	NQ	glu	01-IA-01
LH51) (LH119(4) X T14))					100	fodder	0.17	NQ	glu	
114))	SI	0.40	0.43	2	31	forage	0.22	NO	glu	BK 04P
(transgenic (LH59 X	SL	+0.50	+0.54	(61 cm)	60	silage	0.15	NO	glu	01-IL-01
LH51) (LH119(4) X		. 0.00		(01 011)	76	forage	0.11	NO	glu	01 12 01
T14))					118	fodder	0.11	NQ	glu	
USA (MN) 1994	SL	0.40	0.43	2	31	forage	2.4	NQ	glu	BK-94R-
(transgenic (LH74)		+0.51	+0.55	(61 cm)	60	silage	<u>1.7</u>	NQ	glu	01-MN-01
(LH82(3) X T14))					93	forage	1.4	NQ	giu glu	
				-	114	fodder	2.6	0.23	glu	
USA (MO) 1994	SL	0.40	0.43	$\frac{2}{\sqrt{1}}$	30	forage	1.5	NQ	glu	BK-94R-
(transgenic (LH59 X)		+0.56	+0.60	(61 cm)	60 60	forage	<u>1.2</u> 0.86	NQ 0.08	glu	01-MO-01
T(4) (LIII 19(4) X					89	fodder	1.2	0.08	glu	
USA (NC) 1994	SL.	0.40	0.43	2	30	forage	0.19	NO	glu	BK-94R-
(transgenic (LH51 X	5L	+0.46	+0.49	(61 cm)	58	silage	0.13	NO	glu	01-NC-01
LH210) (LH119(4) X				` '	69	forage	0.09	NQ	glu	
T14))					92	fodder	0.08	NQ	glu	
USA (NY) 1994	SL	0.40	0.43	2	30	forage	1.2	NQ	glu	BK-94R-
(transgenic (LH74)		+0.53	+0.57	(61 cm)	60	silage	<u>0.74</u>	NQ	giu	01-NY-01
(LH82(3) X T14))					77	forage	0.35	NQ	giu elu	
	CT.	0.40	0.42	2	88	fodder	<u>1.5</u>	NQ	glu	DV 04D
USA (TX) 1994 (transgenie (LU51 V	SL	0.40	0.43	2 (61 am)	30 60	forage	0.52	NQ	glu	BK-94R-
(LH) I H210) (I H110(A) V		+0.49	+0.55	(01 cm)	74	forage	0.30	NO	glu	01-1A-01
T(4)					97	fodder	0.25	0.07	glu	
USA (WA) 1994	SL	0.40	0.43	2	30	forage	0.70	0.09	glu	BK-94R-
(transgenic (LH74)		+0.50	+0.54	(61 cm)	60	silage	0.24	0.10	glu	01-WA-01
(LH82)3) X T14))				ľ í	76	forage	0.37	0.11	glu	
					86	fodder	0.43	0.15	glu	
USA (WI) 1994	SL	0.38	0.41	2	30	forage	0.49	NQ	glu	BK-94R-
(transgenic (LH85 X		+0.50	+0.54	(61 cm)	60	silage	0.19	NQ	giu	01-WI-01
LH160) (LH119(4) X					92	forage	0.28	NQ	elu	
114))					107	Iodder	<u>0.50</u>	NQ	· · ·	

Location,	ocation, Application				PHI		Residues, mg/kg	Ref		
Year,	form	kg	kg ai/hl	No ¹	days	Commo	glufosinate-	MPP	NAG	
(variety)		ai/ha	-		-	dity ²	ammonium			
USA (MN) 1995	SL	0.39	0.41	2	30	forage	2.4	0.06	glu	BK95R01-
(transgenic (LH202)		+0.49	+0.53	(7-8	60	forage	1.5	0.08	glu	GGH-01
(LH82 (4) x T25))				leaf)	100	fodder	1.2	0.10	glu	
USA (KS) 1995	SL	0.39	0.41	2	32	forage	1.3	NO	glu	BK95R01-
(transgenic		+0.50	+0.53	(56-66	62	forage	0.97	NO	glu	MDA-02
(B73(5)xT25) (LH51))				cm)	77	forage	1.1	0.09	glu	
					95	fodder	1.2	0.14	glu	
USA (MI) 1995	SL	0.39	0.41	2	30	forage	0.73	0.08	glu	BK95R01-
(transgenic (LH119 (9)		+0.50	+0.54	(61-77	61	forage	0.46	0.07	glu	JRS-02
x T14) (LH168))				cm)	98	forage	0.42	0.11	glu	
					111	fodder	0.50	0.15	glu	
USA (MO) 1995	SL	0.39	0.44	2	10	forage	4.1	0.07	glu	BK95R01-
(transgenic (B73 (5) x		+0.52	+0.60	(61 cm)	20	forage	1.5	0.05	glu	JLB-01
T25) (LH51))				Ì	29	forage	1.0	NQ	glu	
					40	forage	0.82	NQ	glu	
					60	forage	0.49	0.05	glu	
					81	forage	0.17	0.05	glu	
					90	fodder	0.51	0.18	glu	
USA (NE) 1995	SL	0.39	0.41	2	20	forage	0.41	NQ	glu	BK95R01-
(transgenic (LH119 (9))	+0.52	+0.55	(61 cm)	60	forage	0.33	NQ	glu	MDA-01
x T14) (LH51))					80	forage	0.30	NQ	glu	
					89	fodder	0.53	NQ	glu	
USA (OH) 1995	SL	0.39	0.43	2	31	forage	1.9	NQ	glu	BK95R01-
(transgenic (LH119 (9)		+0.52	+0.52	(61 cm)	60	forage	1.6	0.07	glu	JRS-01
x T14) (LH51))					88	forage	0.63	0.17	glu	
					117	fodder	1.1	0.22	glu	
USA (TX) 1995	SL	0.40	0.42	2	10	forage	3.2	0.07	glu	BK95R01-
(transgenic (LH216)		+0.50	+0.53	(53-66	20	forage	1.6	NQ	glu	GLS-01
(B73 (5) x T25))				cm)	30	forage	0.88	NQ	glu	
					40	forage	0.85	NQ	glu	
					61	forage	0.72	NQ	glu	
					71	forage	0.83	0.07	glu	
					84	fodder	1.6	0.14	glu	
Argentina 1996	SL	1.4	0.88	1	1	shoot	54	0.34		A57414
(transgenic)				GS17-	34	shoot	0.50	NQ		ER96ARG0
-				32)	54	shoot	1.9	NQ		01010102
					54	cob	0.08	NQ		

¹. growth stage at final treatment is given as the height of the plant. ² commodity descriptions: forage immature whole plant

silage mature whole plant collected prior to the dent stage

fodder mature plant without ears, collected at grain harvest

³ NQ - residues <LOD (0.05 mg/kg glufosinate free acid equivalents) glu: included in glufosinate-ammonium result.

Year	Appli	cation		PHI	Residues, mg/		Ref	
	form	kg ai/ha kg ai/hl	No^1	days	glufosinate	MPP	NAG	
1993	SL	0.75	1	106	<0.05 shoot	<0.05 shoot	<0.05 shoot	A55207
			GS16	152	<0.05 shoot	<0.05 shoot	<0.05 shoot	93-IGN-C
				152	<0.05 cob	<0.05 cob	<0.05 cob	ON-1-2
1993	SL	1.5	1	106	<0.05 shoot	<0.05 shoot	<0.05 shoot	A55207
			GS16	152	<0.05 shoot	< 0.05 shoot	<0.05 shoot	93-IGN-C
1002	CI	0.75	1	152	< 0.05 cob	<0.05 cob	<0.05 cob	ON-1-3
1995	SL	0.75	I GS16	107	< 0.03 shoot	< 0.03 shoot	< 0.03 shoot	A33207 93 IGN C
			0510	154	< 0.05 should < 0.05 cob	< 0.05 should < 0.05 cob	< 0.05 shout < 0.05 cob	ON-2-2
1993	SL.	1.5	1	107	<0.05 cos	<0.05 cos	<0.05 shoot	A55207
	~ -		GS16	154	<0.05 shoot	<0.05 shoot	<0.05 shoot	93-IGN-C
				154	<0.05 cob	<0.05 cob	<0.05 cob	ON-2-3
1994	SL	0.5	1	99	<0.05 shoot	<0.05 shoot	<0.05 shoot	A55207
			GS17-	137	<0.05 cob	<0.05 cob	<0.05 cob	94-IGN-C
			19					ON-1-2
1994	SL	1.0	1	99	<0.05 shoot	<0.05 shoot	<0.05 shoot	A55207
			GS17-	137	<0.05 cob	<0.05 cob	<0.05 cob	94-IGN-C
100.4	CT.	0.5	19	02	0.05 1	0.05 1	0.24	ON-1-3
1994	SL	0.5		93	< 0.05 shoot	< 0.05 shoot	0.34 shoot	A55207
			10	114	<0.03 000	<0.03 000	<0.03 COD	94-IGN-C
1994	SI	1.0	1	93	<0.05 shoot	<0.05 shoot	0.20 shoot	A 55207
1774	SL	1.0	GS17-	114	< 0.05 should < 0.05 cob	< 0.05 should < 0.05 cob	<0.20 shoot	94-IGN-C
			19		0.05 000	0.05 000	0.05 000	ON-2-3
1994	SL	0.5	1	93	<0.05 shoot	<0.05 shoot	0.32 shoot	A55207
			GS17-	114	<0.05 cob	<0.05 cob	<0.05 cob	94-IGN-C
			19					ON-3-2
1994	SL	1.0	1	93	<0.05 shoot	<0.05 shoot	0.15 shoot	A55207
			GS17-	114	<0.05 cob	<0.05 cob	<0.05 cob	94-IGN-C
		-	19	_				ON-3-3
1995	SL	0.6	1	68	<0.05 shoot	<0.05 shoot	<0.05 shoot	A55207
			GS18- 10	104	<0.05 cob	<0.05 cob	<0.05 cob	95-IGN-C
1005	SI	0.6	2	61	<0.05 shoot	<0.05 shoot	0.28 shoot	A 55207
1775	SL	0.0	2 GS33	97	< 0.05 should < 0.05 cob	< 0.05 should < 0.05 cob	<0.28 shoot	95-IGN-C
			0000	21	0.05 000	10.05 000	0.05 000	ON-1-3
1995	SL	1.2	1	56	0.12 shoot	<0.05 shoot	0.86 shoot	A55207
			GS40	92	<0.05 cob	<0.05 cob	<0.05 cob	95-IGN-C
								ON-1-4
1995	SL	0.6	1	71	<0.05 shoot	<0.05 shoot	0.11 shoot	A55207
			GS18	105	<0.05 cob	<0.05 cob	<0.05 cob	95-IGN-C
100 7	a .			- 0				ON-2-2
1995	SL	0.6	2	62 06	0.05 shoot	<0.05 shoot	0.33 shoot	A55207
			0333	90	<0.05 COD	<0.05 COD	<0.05 COD	95-IGN-C
1005	SI	1.2	1	56	0.50 shoot	<0.05 shoot	3.8 shoot	A 55207
1775	SL	1.2	GS40	90	<0.05 cob	< 0.05 should < 0.05 cob	<0.05 cob	95-IGN-C
								ON-2-4
1995	SL	0.6	1	56	<0.05 shoot	<0.05 shoot	<0.05 shoot	A55207
			GS18-	103	<0.05 cob	<0.05 cob	<0.05 cob	95-IGN-C
			19			ļ		ON-3-2
1995	SL	0.6	2	64	<0.05 shoot	<0.05 shoot	<0.05 shoot	A55207
			GS33	96	<0.05 cob	<0.05 cob	<0.05 cob	95-IGN-C
1005	GT		1		0.42	0.05	2.7 .	ON-3-3
1995	SL	1.2		57	0.43 shoot	<0.05 shoot	3.7 shoot	A55207
			0540	89	<0.05 cob	<0.05 cob		95-IGN-C
	1							UN-3-4

Table 55. Glufosinate residues in transgenic maize forage and fodder resulting from supervised trials in Ontario, Canada.

Year	Application				PHI	Residues, mg/		Ref	
	form	kg ai/ha	kg ai/hl	No^1	days	glufosinate	MPP	NAG	
1995	SL	0.6		1	70	<0.05 shoot	<0.05 shoot	<0.05 shoot	A55207
				GS18	104	<0.05 cob	<0.05 cob	<0.05 cob	95-IGN-C
									ON-4-2
1995	SL	0.6		2	63	<0.05 shoot	<0.05 shoot	0.09 shoot	A55207
				GS33	96	<0.05 cob	<0.05 cob	<0.05 cob	95-IGN-C
									ON-4-3
1995	SL	1.2		1	57	0.40 shoot	<0.05 shoot	2.7 shoot	A55207
				GS40	90	<0.05 cob	<0.05 cob	0.10 cob	95-IGN-C
									ON-4-4

¹GS: growth stage at final treatment GS16 6 leaves unfolded GS17 7 leaves unfolded GS18 8 leaves unfolded GS19 9 or more leaves unfolded GS33 3 nodes detectable

GS40 booting

Table 56. Glufosinate residues in rape seed forage and fodder resulting from supervised trials in France, Germany, and the UK.

Country Year		Applic	ation		PHI	Residues, mg/	kg, as glufosinat	e free acid	Ref
(variety)	form	kg ai/ha	Kg ai/hl	No ¹	days	glufosinate	MPP	NAG ²	
France 1994	SL	0.60	0.30	1 at	0	65 shoot	0.49 shoot		A54228
(Falcon)				GS23	116	0.06 shoot	<0.05 shoot		ER93ECN553
		+0.45	+0.23	2 at	0	20 shoot	0.16 shoot		FRA000202
				GS35	10	7.0 shoot	<0.05 shoot		
					73	0.89 shoot	<0.05 shoot		
					73	0.06 pod	<0.05 pod		
					111	1.6 straw	<0.05 straw		
France 1994	SL	0.60	0.30	1 at	0	58 shoot	0.17 shoot		A54228
(Falcon)				GS23	116	0.05 shoot	0.05 shoot		ER93ECN553
		+0.45	+0.23	2 at	0	24 shoot	0.19 shoot		FRA000302
				GS35	10	7.2 shoot	<0.05 shoot		
					73	0.82 shoot	<0.05 shoot		
					73	0.1 pod	<0.05 pod		
					111	1.3 straw	<0.05 straw		
France 1994 (Gs	SL	0.60	0.30	1 at	0	6.5 shoot	0.05 shoot		A54228
401×90)				GS21	73	0.12 shoot	<0.05 shoot		ER93ECN553
		+0.45	+0.23	2 at	0	45 shoot	0.34 shoot		FRA000102
				GS33	38	0.16 shoot	<0.05 shoot		
					99	0.11 shoot	<0.05 shoot		
					99	0.24 pod	<0.05 pod		
					121	0.69 straw	0.13 straw		
France 1995	SL	0.6	0.30	2	0	7.9 shoot	0.08 shoot		A56444
(Falcon B)				GS35	16	2.5 shoot	<0.05 shoot		ER94ECN553
					69	0.39 shoot	<0.05 shoot		FRA000203
					69	0.12 pod	<0.05 pod		
					127	0.75 straw	<0.05 straw		
France 1995	SL	0.67	0.32	2	0	19 shoot	0.07 shoot		A56444
(Falcon)		+0.60	+0.30	GS39	28	2.5 shoot	<0.05 shoot		ER94ECN553
					78	1.0 shoot	<0.05 shoot		FRA000103
					78	0.15 pod	<0.05 pod		
					133	1.2 straw	0.11 straw		
Country Year		Applic	ation		PHI	Residues, mg/	Residues, mg/kg, as glufosinate free act		
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(variety)	form	kg ai/ha	Kg ai/hl	No ¹	days	glufosinate	MPP	NAG ²	
France 1996	SL	0.60	0.30	2	0	25 shoot	0.3 shoot		A57955
(GMO hybrid)			+0.24	GS32	17	6.0 shoot	0.08 shoot		ER96ECN553
					58	1.8 shoot	<0.05 shoot		FRA000103
					58	0.57 pod	<0.05 pod		
F 100.0		0.50	0.00		108	5.5 straw	0.22 straw		
France 1996	SL	0.60	0.30	2	0	13 shoot	0.11 shoot		A57955
(GMO hybrid)			+0.24	G832	1/	2.9 shoot	<0.05 shoot		ER96ECN553
					28 59	0.95 shoot	<0.05 shoot		FRA000102
					28 108	0.4 pod	<0.05 pod		
Germany 1005	SI	0.60	0.20	2	0	14 shoot	0.13 shoet		156111
(Falcon G-	SL	0.00	0.20	GS35	7	9.3 shoot	0.28 shoot		FR94FCN553
(1 alcoli 0- S40/90)				0555	53	1.6 shoot	< 0.05 shoot		DFU030103
5-10/202					53	0.32 pod	<0.05 shoot		DECOSOTOS
					106	1.0 straw	0.11 straw		
Germany 1995	SL	0.60	0.20	2	0	14 shoot	0.15 shoot		A56444
(Falcon)	~-			GS35	20	2.7 shoot	<0.05 shoot		ER94ECN553
` '					55	2.4 shoot	<0.05 shoot		DEU040103
					55	0.38 pod	<0.05 pod		
					105	2.7 straw	0.18 straw		
Germany 1995	SL	0.60	0.20	2	0	12 shoot	0.14 shoot		A56444
(Falcon)				GS35	12	4.0 shoot	<0.05 shoot		ER94ECN553
					47	1.6 shoot	<0.05 shoot		DEU040203
					47	0.52 pod	<0.05 pod		
					110	2.3 straw	0.12 straw		
Germany 1995	SL	0.60	0.20	2	0	19 shoot	0.08 shoot		A56444
(Falcon)				GS37	21	1.0 shoot	<0.05 shoot		ER94ECN553
					61	0.15 shoot	<0.05 shoot		DEU050103
					61	0.1 pod	<0.05 pod		
a 100 f	<i>a</i> .	0.50	0.00		104	0.17 straw	<0.05 straw		
Germany 1996	SL	0.60	0.20	2	0	21 shoot	0.32 shoot		A57955
(Falcon)				6332	0 51	4.5 shoot	< 0.05 shoot		EK96ECN555
					51	1.1 shoot	<0.05 shoot		DE0050102
					94	0.25 pou 1 7 straw	< 0.05 pou		
Germany 1996	SI	0.80	0.27	2	0	28 shoot	0.00 shaw		A 57955
(Falcon)	SL	0.00	0.27	G\$32	6	5.8 shoot	0.43 shoot 0.07 shoot		ER96ECN553
(I alcoli)				0032	51	1.1 shoot	< 0.05 shoot		DEU030103
					51	0.22 pod	<0.05 pod		220000100
					94	4.3 straw	0.19 straw		
Germany 1996	SL	0.60	0.20	2	0	20 shoot	0.19 shoot		A57955
(Falcon)				GS32	12	4.5 shoot	0.08 shoot		ER96ECN553
					54	0.43 shoot	<0.05 shoot		DEU050102
					54	0.13 pod	<0.05 pod		
					97	1.8 straw	0.09 straw		
Germany 1996	SL	0.80	0.27	2	0	26 shoot	0.28 shoot		A57955
(Falcon)				GS32	12	4.7 shoot	0.06 shoot		ER96ECN553
					54	1.1 shoot	<0.05 shoot		DEU050103
					54	0.35 pod	<0.05 pod		
	CT.	0.60	0.20	-	9/	2.3 straw	0.08 straw		157055
(Felger)	SL	0.60	0.20	2	0	10 shoot 5.2 at -1	0.2 shoot		AJ/YJJ EDOGEONIES
(raicon)				0331	17	J.∠ shoot	0.07 shoot		DELIDADIO2
					+/ 47	0.19 pod	< 0.05 shoot		DE0040102
					97	3.0 straw	0.14 straw		
Germany 1006	SI	0.80	0.27	2	0	18 shoot	0.2 shoot		A 57955
(Falcon)		0.00	5.27	G\$37	7	6.7 shoot	0.07 shoot		ER96ECN553
(i dicon)				0007	, 47	1.7 shoot	<0.05 shoot		DEU040103
					47	0.51 pod	<0.05 pod		
					97	3.2 straw	0.17 straw		

Item kg ai/ha Kg ai/ha Kg ai/ha No ¹ days glufosinate MPP NAG ² UK 1994 SL 0.60 0.30 1 at 0 37 shoot 0.49 shoot A5422 UK 1994 SL 0.60 0.30 1 at 0 37 shoot 0.05 shoot ER93E UK 1994 SL 0.60 0.30 1 at 0 224 <0.05 shoot <0.05 shoot ER93E UK 1994 SL 0.60 0.30 1 at 0 224 <0.05 shoot <0.05 shoot GBR0 UK 1994 SL 0.60 0.30 1 at 0 57 46 shoot 0.67 0.60 shoot A5422 UK 1994 SL 0.60 0.30 1 at 0 57 46 shoot 0.67 0.60 shoot A5422 UK 1994 SL 0.60 0.30 1 at 0 57 46 shoot 0.67 0.60 shoot ER93E UK 1994 SL 0.60 0.30 1 at 0	7 3CN552 00102 7 3CN552 00202 8 3CN553 00202
UK 1994 SL 0.60 0.30 1 at GS21 0.59 (37 shoot) 37 shoot (0.05 shoot) 0.49 shoot (0.05 shoot) A5422 (0.05 shoot) UK 1994 SL 0.60 0.30 1 at (0.30) 0 37 shoot) 0.49 shoot) A5422 UK 1994 SL 0.60 0.30 1 at (0.30) 0 37 shoot) (0.05 shoot)	7 3CN552 00102 7 3CN552 00202 8 3CN553 00202
OK 1994 SL 0.60 0.30 1 at 0 57 short 0.05 short ER931 GS21 159 <0.05 shoot	CN552 00102 7 3CN552 00202 8 3CN553 00202
UK 1994 SL 0.60 0.30 1 at 0 57 46 shoot <0.05 shoot <0.05 shoot GBR0 UK 1994 SL 0.60 0.30 1 at 0 57 46 shoot <0.05 shoot	00102 7 3CN552 00202 8 3CN553 00202
UK 1994 SL 0.60 0.30 1 at 0 0 57 46 shoot 275 <0.05 shoot <0.05 pod 275 <0.60 shoot <0.05 straw A5422 UK 1994 SL 0.60 0.30 1 at 1 at 1 at 1 at 0 57 46 shoot <0.05 shoot	7 3CN552 00202 8 3CN553 00202
UK 1994 SL 0.60 0.30 1 at GS21 0.32 57 46 shoot (3275) 0.67 0.60 shoot (3275) A5422 (305 straw) UK 1994 SL 0.60 0.30 1 at 1 at 1 at 1 at 1 at 1 at 1 at 1 at	7 3CN552 00202 8 3CN553 00202
UK 1994 SL 0.60 0.30 1 at GS21 0 57 46 shoot 0.67 0.60 shoot A5422 UK 1994 SL 0.60 0.30 1 at I at 0 57 46 shoot 0.67 0.60 shoot A5422 UK 1994 SL 0.60 0.30 1 at I at 0 57 46 shoot 0.67 0.60 shoot ER93E I g2 <0.05 shoot	7 3CN552 00202 8 3CN553 00202
UK 1994 SL 0.60 0.30 1 at GS21 0 57 46 shoot 0.67 0.60 shoot A5422 UK 1994 SL 0.60 0.30 1 at GS21 0 57 46 shoot 0.67 0.60 shoot ER93E UK 1994 SL 0.30 1 at I32 0.11 shoot <0.05 shoot	7 3CN552 00202 8 3CN553 00202
GS21 132 0.11 shoot <0.05 shoot	8 00202 8 3 3 3 5 5 3 5 3 5 3 5 3 5 3 5 3 5 3 5
192 <0.05 shoot <0.05 shoot GBR0 192 <0.05 (2) pod	00202 8 3CN553 00202
192 <0.05 (2) pod <0.05 (2) pod	8 3CN553 00202
	8 2CN553 20202
212 <0.05 straw <0.05 straw	8 2CN553 30202
UK 1994 SL 0.60 0.30 1 at 0 52 shoot 0.62 shoot A5422	ECN553 00202
GS21 121 <0.05 shoot <0.05 shoot ER93E	00202
+0.45 +0.23 2 at 0 16 shoot 0.08 shoot GBR0	
GS35 11 2.2 shoot <<0.05 shoot	
71 0.18 shoot <0.05 shoot	
71 0.07 pod <0.05 pod	
91 0.5 straw <0.05 straw	
UK 1994 (Falcon) SL 0.60 0.30 1 at 0 13 shoot 0.42 shoot A5422	8
GS21 159 0.13 shoot <0.05 shoot ER93E	CN553
+0.45 +0.23 2 at 0 0.11 shoot <0.05 shoot GBR0	00102
GS35 25 1.2 shoot <0.05 shoot	
65 0.31 shoot <0.05 shoot	
65 0.09 pod <0.05 pod	
116 0.47 straw <0.05 straw	
UK 1996 (Falcon) SL 0.60 0.30 2 0 16 shoot 0.31 shoot A5795	5
GS32 10 4.8 shoot 0.07 shoot ER96E	CN553
61 1.3 shoot <0.05 shoot GBR0	00103
61 0.17 pod <0.05 pod	
100 1.3 straw 0.06 straw	
UK 1996 (Falcon) SL 0.60 0.30 2 0 26 shoot 0.17 shoot A5795	5
GS32 9 6.7 shoot 0.11 shoot ER96E	CN553
58 1.8 shoot <0.05 shoot GBR0	30203
58 0.31 pod <0.05 pod	
92 5.7 straw 0.11 straw	~
UK 1996 (Falcon) SL $[0.60]$ $[0.30]$ 2 $[0]$ 14 shoot $[0.28$ shoot [A3/95]	J ICNEE2
$\begin{array}{c ccccccccccccccccccccccccccccccccccc$	UN333
61 0 20 rod 0 05 rod	J0102
100 15 straw	
IW 1996 (Falcon) SI 0.80 0.40 2 0 20 shoot 0.45 shoot A 5705	5
$GS32 \ 10 \ 69 \ shoot \ 0.45 \ shoot \ FP06E$	J ICN553
61 12 short 0.0 short GBP0	00104
61 0 19 nod <0.05 nod	50104
$100 14 ext{ straw} 0.05 ext{ straw}$	
$\frac{100}{1100} \frac{1100}{1100} \frac{1000}{0000} \frac{1000}{000} 1000$	5
GS32 9 52 short 0.00 short FR96E	CN553
58 1.5 shoot <0.05 shoot GBR0	00202
58 0.22 pod <0.05 pod	
92 3.3 straw^3 0.09 straw ³	
UK 1996 (Falcon) SL 0.80 0.40 2 0 30 shoot 0.09 shoot A5795	5
GS32 9 7.7 shoot 0.1 shoot ER96F	CN553
58 1.9 shoot <0.05 shoot GBR0	00204
58 0.34 pod <0.05 pod	-
92 4.5 straw ³ 0.12 straw ³	

¹ : GS growth stage	
GS21	first side shoot detectable
GS32	2 visibly extended internodes
GS33	3 visibly extended internodes
GS35	5 visibly extended internodes
GS37	7 visibly extended internodes
GS80	beginning of ripening
GS83	30% of pods ripe
2 included in glufosinate-amm	onium result.

³: Straw samples were harvested at GS80-83 and stored in paper bags for even drying and prepared to laboratory samples 10 days later.

Table 57. Glufosinate residues in canola forage and fodder resulting from supervised trials in Australia.

Location,		App	lication		PHI	Residues, mg/kg,	as gluf	osinate	Ref
Year					-	free a	cid		
(variety)	form	kg ai/ha	kg ai/hl	No ¹	days	glufosinate	MPP^2	NAG ²	
Australia (NSW)	SL	0.6	0.72	1	86	<0.1 straw			YH 96 AH
1996 (glufosinate				GS 3-5 leaf					AU NW 06
selective line)									
Australia (NSW)	SL	1.2	1.4	1	86	0.12 straw			YH 96 AH
1996 (glufosinate				GS 3-5 leaf					AU NW 06
selective line)									
Australia (WA)	SL	0.6	0.83	1	29	1.0 2.4 forage dw			YH 96 AH
1996 (var ex				GS 1-2 leaf	56	<0.05 forage dw			AU WA 09
Canada)					124	<0.1 (2) straw			
Australia (WA)	SL	1.2	1.7	1	29	2.5 2.3 forage dw			YH 96 AH
1996 (var ex				GS 1-2 leaf	56	<0.05 forage dw			AU WA 09
Canada)					124	<0.1 (2) straw			
Australia (SA)	SL	0.6	0.75	1	42	0.8 0.6 forage dw			YH 96 AH
1996 (transgenic				GS 4-6 leaf	95	0.13 0.12 straw			AU SA 17
Canadian line)									
Australia (SA)	SL	1.2	1.5	1	42	2.0 1.6 forage dw			YH 96 AH
1996 (transgenic				GS 4-6 leaf	95	0.14 0.12 straw			AU SA 17
Canadian line)									
Australia (Vic)	SL	0.6	0.6	1	42	<0.05 (2) forage			YH 96 AH
1997 (Canadian				GS 2 leaf	105	<0.1 (2) straw			AU VN 12
tolerant)									
Australia (Vic)	SL	1.2	1.2	1	42	<0.05 (2) forage			YH 96 AH
1997 (Canadian				GS 2 leaf	105	<0.1 (2) straw			AU VN 12
tolerant)									

¹: GS growth stage expressed in terms of leaf development ² included in glufosinate-ammonium result.

Table 58. Glufosinate residues in soya bean forage and fodder resulting from supervised trials in the USA.

Location, Year		App	olication		PHI		Residues, mg	/kg, expressed a	as glufosinate	Ref
(variety)	form	kg ai/ha	kg ai/hl	No ¹	days		glufosinate	MPP	NAG	
USA (IA)	SL	0.5	0.51	1	87	hay	0.22	0.25 0.23	0.72 0.56	ER-93-USA-
1993 (W98-				trifoliate						02-IA-01
/)	SL	0.5	0.51	1	65	hay	2.8	1.3	6.7	
				full pod		2				
	SL	0.5	0.51	2 full pod	65	hay	2.1	1.1	5.1	
USA (IL)	SL	0.5	0.53	1 1	85	hay	0.15	0.32	0.88	ER-93-USA-
1993 (W98-				trifoliate		2				02-IL-01
7)	SI	0.5	0.53	1	65	how	13	13	58	
	SL	0.5	0.55	full pod	05	пау	1.5	1.5	5.0	
	SL	0.5	0.53	2	65	hay	1.8	2.2	8.1	
	C1	0.5	0.52	full pod	07	hor	0.00	0.22	0.22	
1993 (W98-	SL	0.5	0.55	1 trifoliate	0/	пау	0.09	0.22	0.52	02-MO-01
7)										
	SL	0.5	0.53	1 full pod	65	hay	1.8	1.4	6.9	
	SL	0.5	0.53	2	65	hay	2.0	1.6	7.4	
				full pod						
USA (IN)	SL	0.5	0.53	1 trifolioto	87	hay	0.16	0.41	0.47	ER-93-USA-
1993 (w98- 7)				unonate						02-119-01
, ,	SL	0.5	0.53	1	65	hay	2.5	2.5	5.6	
	SI	0.5	0.53	full pod	65	how	3.2	3.0	73	
	SL	0.5	0.55	full pod	05	пау	3.2	5.0	7.5	
USA (IN)	SL	0.5	0.53	1	87	hay	0.06 0.05 0.06	0.16 0.17 0.18	0.15 0.14 0.16	ER-93-USA-
1993 (W98)	SI	0.5	0.53	trifoliate	65	hav	151627	131216	463071	02-IN-02
	SL	0.5	0.55	full pod	05	nay	1.5 1.0 2.7	1.5 1.2 1.0	4.0 3.7 7.1	
	SL	0.5	0.53	2	65	hay	2.3 3.6 1.9	1.9 2.5 1.8	6.8 9.7 5.8	
USA (AR)	SI	0.5	11	full pod	112	hav	<0.05	<0.05	0.12	FR-93-USA-
1993 (W62)	SL	0.5	1.1	trifoliate	112	nay	<0.05	<0.05	0.12	02-AR-01
	SL	0.5	1.1	1	65	hay	1.1	0.93	5.0	
	SL	0.5	11	full pod	65	hav	14	12	63	
		0.5		full pod	00	ildy	1.1	1.2	0.5	
USA (MS)	SL	0.5	1.1	1	86	hay	0.10	0.08	0.29	ER-93-USA-
1993 (W98)	SL	0.5	11	trifoliate	65	hav	0.58	0.58	47	02-MS-01
	22	0.0		full pod	00	iiuj	0100	0.00		
	SL	0.5	1.1	2	65	hay	0.80	0.63	4.6	
USA (VA)	SL	0.5	1.1	Tuli pod	103	hav	0.43	0.31	1.3	ER-93-USA-
1993 (W62)	SE	0.5		trifoliate	105	ilay	0.15	0.01	1.5	02-VA-01
	SL	0.5	1.1	1	64	hay	0.31	0.24	1.2	
	SL	0.5	1.1	$\frac{1}{2}$	64	hav	0.31	0.33	1.5	
				full pod						
USA (NC)	SL	0.4	0.43	1 2 rd 1	21	forage	0.20	< 0.05	glu glu	Project: BK-
(transgenic				3 node	42 136	forage hav	<0.05 <0.05	<0.05 <0.05	glu	94R-06 Trial#: NC-01
Group 5										
W62 A5403)	CT	0.4	0.42	2	21	for	0.26.0.22	0.20.0.24	glu	
	SL	+0.5	+0.54	6^{tn} node	123 ²¹	hav	<0.05 (2)	<0.29 0.34	glu	
	SL	0.4	0.43	2	21	forage	2.9 3.2	0.27 0.36	glu	
		+0.5	+0.54	bloom	90	hay	0.61 0.49	0.31 0.29	giu	

Location,	Application				PHI		Residues, mg	kg, expressed a	as glufosinate	Ref
Year,										
(variety)	form	kg ai/ha	kg ai/hl	No ¹	days		glufosinate	MPP	NAG	
	SL	0.4	0.43	2	11	forage	2.6 (2)	0.31 0.32	glu	
		+0.5	+0.54	full pod	77	hay	1.2 1.0 0.86	0.65 0.58 0.51	glu	

¹ growth stage described glu: included in glufosinate-ammonium result.

Table 59. Glufosinate residues in tolerant sugar beet tops and leaves resulting from foliar application in supervised trials in France, Germany and the UK.

Country, Year		App	lication		PHI	Residues, n	ng/kg, as glufosi acid	nate free	Ref
(variety)	form	kg ai/ha	kg ai/hl	No ¹	davs	glufosinate	MPP	NAG ²	-
France 1995 (transgenic hybrid)	SL	0.60	0.30	2 GS31	0 12 16	10 plan 4.1 plan 4.1 plan	t0.1 plant <0.05 plant <0.05 plant		A56446 ER95ECN555 FRA000102
					23 85	2.8 plan 1.1 leaves	<0.05 plant <0.05 leaves		
France 1996 (transgenic hybrid)	SL	0.80	0.32	2 GS19	0 13 20 55 96	31 plan 3.3 plan 1.6 plan 0.34 plan <0.05	t0.18 plant t<0.05 plant t<0.05 plant t<0.05 plant t<0.05 leaves		A57574 ER96ECN551 FRA000102
France 1996 (transgenic hybrid)	SL	0.80	0.32	2 GS19	0 13 20 54 96	26 plan 2.4 plan 1.8 plan 0.68 plan <0.05 leaves	0.11 plant <0.05 plant <0.05 plant <0.05 plant <0.05 plant <0.05 plant		A57574 ER96ECN551 FRA000202
France 1996 (transgenic hybrid)	SL	0.80	0.32	2 GS19	0 11 16 51 91	20 plan 4.0 plan 3.6 plan 1.6 plan 0.44 leaves	0.14 plant 0.07 plant <0.05 plant <0.05 plant <0.05 plant <0.05 leaves		A57574 ER96ECN551 FRA000302
Germany 1995 (Sä-Nr. 951 959 - 36)	SL	0.80	0.27	2 GS31	0 23 39 58 96	24 plan 4.0 plan 1.9 plan 1.2 plan 0.53 leaves	0.09 plant <0.05 plant <0.05 plant 0.05 plant 0.06 leaves		A57574 ER96ECN551 DEU010102
UK 1995 (transgenic hybrid)	SL	0.60	0.30	2 GS19	0 4 12 27 83	9.3 plan 6.9 plan 4.7 plan 2.7 plan 1.3 leaves plan	t <0.05 plant t 0.05 plant t <0.05 plant t <0.05 plant <0.05 plant <0.05 leaves		A56446 ER95ECN555 GBR000102
UK 1995 (transgenic hybrid)	SL	0.60	0.60	2 GS31	0 8 21 38 94	6.5 plan 1.9 plan 1.8 plan 1.1 plan 0.65 leaves	t <0.05 plant <0.05 plant t <0.05 plant t <0.05 plant <0.05 plant <0.05 leaves		A56446 ER95ECN555 GBR000202
UK 1996 (Roberta)	SL	0.80	0.80	2 GS31	0 14 43 71 111	11 plan 6.1 plan 2.4 plan 2.2 plan 1.4 leaves	0.08 plant 0.07 plant 0.06 plant 0.06 plant 0.07 leaves		A57574 ER96ECN551 GBR000102
UK 1996 (transgenic hybrid)	SL	0.80	0.80	2 GS19	0 14 40 64 105	12 plan 5.2 plan 2.0 plan 1.3 plan 0.83 leaves	0.15 plant 0.09 plant 0.05 plant 0.05 plant <0.05 plant <0.05 leaves		A57574 ER96ECN551 GBR000202

¹: GS growth stage GS14 4 leaves (2nd pair unfolded) GS15 5 leaves unfolded GS19 9 or more leaves unfolded GS31 beginning of crop cover formation ² included in glufosinate-ammonium result.

Table 60. Glufosinate residues in tolerant sugar beet tops resulting from foliar application in supervised trials in the USA.

Location,		Appli	cation		PHI	Residues, mg/	kg, as glufosinat	e free acid	Ref
Year,	C	1	1	NL 1	1	Cl. Calinata	MDD	NAC^2	
(variety)	Iorm	kg ai/na	kg ai/ni	NO	days	Glufosinate	MPP	NAG	DV 05D 06
USA (CA) 1995	SL	0.21	0.21	3	10	0.19 0.23	< 0.05 (2)		BK-95R-06
(Liberty Link)		+0.20	+0.20	8-leaf	15	0.31 0.29	0.14 0.17		CA-016
		+0.20	+0.20		30	0.23 0.28	0.53 0.54		(Regimen B)
					60 120	0.13 0.12	0.370.33		
110A (CA) 1005	CT	0.42	0.42	2	139	<0.05(3)	0.08 0.06 0.12		DV OFD OC
USA (CA) 1995	SL	0.43	0.43	3	10	0.39 0.46	<0.05 (2)		BK-95K-06
(Liberty Link)		+0.40	+0.39	6-leaf	15	1.04 1.1 1.2	0.51 0.37 0.48		CA-016
		+0.40	+0.40		50 60	0.03 0.70	1.2 1.1		(Regimen C)
					120	0.39 0.32	0.88 0.78		
USA (CA) 1005	CI.	0.61	0.50	1	159	< 0.03(2)	0.21 0.23		DV 05D 06
(Liberty Link)	SL	0.01	0.39	I 6 loof	10	5.0 5.5 2 5 2 8 2 0	0.23 0.22		DK-93K-00
(Liberty Link)				0-leal	15	2.5 2.6 2.0	$0.36\ 0.44\ 0.42$		(Dagiman D)
					50 60	1.2 1.4	1.2 1.5		(Regimen D)
					130	0.48 0.00 0.43	$0.82 \ 0.70 \ 0.81$		
					139	<pre>0.05 0.08</pre>	0.29 0.22 0.21		
USA (MN) 1005	SI	0.10	0.21	3	05	<0.05 (2)	<0.05 (2)		BK 05P 06
(Liberty Link)	SL	10.19	0.21	J 8 loof	95	<0.03 (2)	<0.03 (2)		DK-95K-00 KRT 01
(LIDEITY LIIK)		± 0.20		0-10ai					(Regimen B)
USA (MN) 1005	SI	0.20	0.43	3	95	<0.05 (2)	<0.05 (2)		(Reginen D) BK-95R-06
(Liberty Link)	5L	± 0.40	0.43	8-leaf)5	<0.05 (2)	<0.05 (2)		KRT_01
(Enterty Entry)		+0.41		0 lea					(Regimen C)
USA (MN) 1995	SL	0.61	0.64	2	95	0 10 0 09	<0.05 (2)		RK-95R-06
(Liberty Link)	5L	+0.62	0.04	2 8-leaf	,,,	0.10 0.09	(0.05 (2)		KBT-01
(Exectly Ellik)		10.02		0 Ioui					(Regimen D)
USA (ID) 1995	SL	0.21	0.21	3	49	0.08 0.09	<0.05 (2)		BK-95R-06
(Liberty Link)	22	+0.20	0.21	8-leaf		0100 0102	(0100 (2)		TWM-01
(+0.21							Regimen B)
USA (ID) 1995	SL	0.41	0.42	3	49	0.22 0.23	< 0.05 (2)		BK-95R-06
(Liberty Link)		+0.43		8-leaf					TWM-01
× , ,		+0.40							Regimen C)
USA (ID) 1995	SL	0.63	0.64	2	49	0.31	0.05		BK-95R-06
(Liberty Link)		+0.61		8-leaf					TWM-01
									Regimen D)
USA (ND) 1995	SL	0.21	0.21	3	103	0.05 0.09 0.05	< 0.05 (3)		BK-95R-06
(Liberty Link)		+0.21	+0.22	11-12-					PGM-01
		+0.22	+0.21	leaf					(Regimen B)
USA (ND) 1995	SL	0.43	0.43	3	104	0.11 0.07 0.11	< 0.05 (3)		BK-95R-06
(Liberty Link)		+0.43		11-12-					PGM-01
		+0.46		leaf					(Regimen C)
USA (ND) 1995	SL	0.86	0.85	2	104	0.07 0.08	< 0.05 (2)		BK-95R-06
(Liberty Link)		+0.64	+0.63	11-12-					PGM-01
				leaf					(Regimen D)
USA (MI) 1996	SL	0.60	0.61	2	109	0.14 0.16	0.05 < 0.05		BK-96R-01
(Liberty Link)			+0.63	8-leaf					R05-02 (Regimen
									B)
USA (MI) 1996	SL	0.60	0.64	3	106	0.30 (2)	<0.05 (2)		BK-96R-01
(Liberty Link)		+0.39	+0.40	10-leaf					R05-02 (Regimen
		+0.60	+0.66						C)

Location,		Appli	cation		PHI	Residues, mg	Residues, mg/kg, as glufosinate free acid		
(variety)	form	ko ai/ha	kg ai/hl	No ¹	davs	Glufosinate	MPP	NAG^2	
USA (OH) 1996	SL	0.60	0.63	2	83	0.16(2)	<0.05 (2)	1010	BK-96R-01
(Liberty Link)	SL	0.00	+0.67	2 8-leaf	05	0.10 (2)	<0.05 (2)		R05-03 (Regimen
(Licenty Linit)				0 100					B)
USA (OH) 1996	SL	0.60	0.61	3	77	0.46 (2)	< 0.05 (2)		BK-96R-01
(Liberty Link)		+0.39	+0.41	10-leaf					R05-03 (Regimen
		+0.60	+0.66						C)
USA (ND) 1996	SL	0.60	0.63	2	67	0.25 0.24	< 0.05 (2)		BK-96R-01
(Liberty Link)				7.5-8.5-					R05-04 (Regimen
		0.00	0.49	leaf			0.07 (0)		B)
USA (ND) 1996	SL	0.60	0.63	3	62	0.65 0.53	<0.05 (2)		BK-96R-01
(Liberty Link)		+0.39	+0.41	9.5-11-					R05-04 (Regimen
	CI	+0.60	+0.03		115	(0.05.(0)	(0.05.(2))		C)
(Liberty Link)	SL	0.00	0.05	∠ 8-leaf	115	<0.03 (2)	<0.03 (2)		DK-90K-01 R07-01 Regimen
(Liberty Link)				0-leal					R07-01 Regimen
USA (NE) 1996	SL	0.60	0.61	3	108	<0.05 (2)	<0.05 (2)		BK-96R-01
(Liberty Link)	22	+0.39	+0.41	10-leaf	100	(0100 (2)			R07-01 Regimen
		+0.60	+0.66						C)
USA (ND) 1996	SL	0.60	0.63	2	73	0.13 0.16	< 0.05 (2)		BK-96R-01
(Liberty Link)				8-leaf					R07-02 (Regimen
									B)
USA (ND) 1996	SL	0.60	0.63	3	66	0.23 0.24	0.06 0.08		BK-96R-01
(Liberty Link)		+0.39	+0.41	10-leaf					R07-02 (Regimen
		+0.60	+0.63			0.07 (0)	0.07 (0)		C)
USA (CO) 1996	SL	0.60	0.59	2	80	<0.05 (2)	<0.05 (2)		BK-96R-01
(Liberty Link)				8-lear					R08-01 (Regimen
USA (CO) 1996	SL	0.60	0.63	3	68	0.38(2)	<0.05 (2)		BK-96R-01
(Liberty Link)	SL	+0.39	+0.40	10-leaf	00	0.50 (2)	(0.05 (2)		R08-01 (Regimen
(Licenty Linit)		+0.60	+0.58	10 100					C)
USA (CO) 1996	SL	0.60	0.61	2	86	0.06 (2)	< 0.05 (2)		BK-96R-01
(Liberty Link)			+0.59	8-leaf					R09-01 (Regimen
									B)
USA (CO) 1996	SL	0.60	0.64	3	81	0.22 0.24	< 0.05 (2)		BK-96R-01
(Liberty Link)		+0.39	+0.41	10-leaf					R09-01 (Regimen
	~-	+0.60	+0.61	-					C)
USA (CA) 1996	SL	0.60	0.65	2	132	<0.05 0.07	<0.05 (2)		BK-96R-01
(Liberty Link)			+0.62	8-leaf					R10-01 (Regimen
USA (CA) 1006	CI.	0.60	0.61	2	122	0 10 0 26	0.06.0.08		В) РИ 06Р 01
(Liberty Link)	SL	0.00 ⊥0.39	± 0.01	J 10-leaf	122	0.19 0.20	0.00 0.08		BR-90R-01 R10-01 (Regimen
(Liberty Link)		+0.60	+0.63	10-icai					C)
USA (ID) 1996	SL	0.60	0.62	2	128	0.10.0.07	<0.05 (2)		BK-96R-01
(Liberty Link)	22	0.00	0.02	8-10-		0.10 0.07			R11-01 (Regimen
				leaf					B)
USA (ID) 1996	SL	0.60	0.61	3	121	0.32 0.30	0.06 0.05		BK-96R-01
(Liberty Link)		+0.39	+0.41	10-leaf					R11-01 (Regimen
		+0.60	+0.63						C)

¹ growth stage is described in terms of number of leaves. ² included in glufosinate-ammonium result.

Table 61. Glufosinate residues in almond hulls resulting from supervised trials in California, USA, 1985 Analyses of samples from replicate plots are shown separately. Double-underlined residues are from treatments according to GAP and were used to estimate maximum residue levels. CLICK HERE

(Variety)	Application				PHI, days	Residues, mg/kg free	g, as glufosinate acid	Ref
	Form	kg ai/ha	kg ai/hl	no.		glufosinate	MPP	
(Carmel)	SL	1.7		3	15	< <u>0.5</u> (3)	<0.5 (3)	A48446 07-CA-85-018
(Carmel)	SL	3.4		3	15	<0.5 (3)	<0.5 (3)	A48446 07-CA-85-018
(Special)	SL	1.7		3	15	< <u>0.5</u> (3)	<0.5 (3)	A48446 07-CA-85-018
(Special)	SL	3.4		3	15	<0.5 (3)	<0.5 (3)	A48446 07-CA-85-018
(Non-Peril)	SL	1.7		3	14	< <u>0.5</u> (3)	<0.5 (3)	A48446 07-CA-85-037
(Non-Peril)	SL	3.4		3	14	<0.5 (3)	<0.5 (3)	A48446 07-CA-85-037

Farm animal feeding studies

Studies on cows and laying hens were reported.

Groups of 3 lactating Holstein dairy cows (each weighing 430-610 kg) were dosed with glufosinate-ammonium + NAG (15+85) in gelatin capsules at total nominal levels equivalent to 9.1, 27 and 91 ppm glufosinate free acid in the diet for 28 consecutive days (Czarnecki and Brady, 1995b). Doses were administered after the morning milking in four separate capsules, one containing glufosinate-ammonium and three containing NAG. Milk was collected twice each day and pooled for analysis. On day 29 the animals from each group were slaughtered. The cows consumed a nominal 21 kg (average 16.8-25.4 kg) feed each per day.

The dosing mixture (glufosinate-ammonium 15 parts + NAG 85 parts) was chosen to represent the typical terminal residue composition in glufosinate-resistant crops that might be fed to animals.

Residues were not detected in milk from the lowest dosing level, detected in a few samples from the middle level, and consistently detected the highest dose group where a plateau was reached on day 3. Two milk samples from the same cow (on days 16 and 18) had unusually high residues, which corresponded with a period of low feed consumption (a constant dose representing a higher feed concentration) and low milk production. Residues in the tissues were below the LODs at the lower doses. At the highest dose they were detected in the kidneys and liver, but not in muscle or fat.

Table 62. Residues in the tissues of lactating dairy cows dosed with glufosinate-ammonium + NAG (15+85) in gelatin capsules at levels equivalent to 9.1, 27 and 91 ppm glufosinate free acid in the diet for 28 consecutive days (Czarnecki and Brady, 1995b).

Tissue	Residues, mg/kg, expressed as glufosinate free acid								
	9.1 pj	pm	27]	opm	91 ppm				
	glufosinate + MPP		glufosinate +	MPP	Glufosinate +	MPP			
	NAG		NAG		NAG				
Muscle	<0.05 (3)	< 0.05 (3)	<0.05 (3)	< 0.05 (3)	< 0.05 (3)	< 0.05 (3)			
Fat,	< 0.05 (3)	< 0.05 (3)	< 0.05 (3)	< 0.05 (3)	< 0.05 (3)	< 0.05 (3)			
perirenal									
Liver	<0.1 (3)	<0.1 (3)	<0.1 (3)	<0.1 (3)	< 0.1 (3)	0.28 0.29 0.25			
Kidneys	<0.1 (3)	<0.1 (3)	<0.1 (3)	<0.1 (3)	0.10 0.14 0.15	<0.1 (2) 0.13			

Day	Residues, mg/kg, expressed as glufosinate free acid										
	9.1	ppm	27 ppm	1	91	opm					
	glufosinate +	MPP	glufosinate +	MPP	glufosinate + NAG	MPP					
	NAG		NAG		-						
1	< 0.02 (3)	< 0.02 (3)	< 0.02 (3)	< 0.02 (3)	< 0.02 (3)	< 0.02 (3)					
3	< 0.02 (3)	< 0.02 (3)	< 0.02 (3)	< 0.02 (3)	0.05 0.04 0.05	< 0.02 (3)					
5	<0.02 (3)	< 0.02 (3)	< 0.02 (3)	< 0.02 (3)	0.05 0.04 0.03	< 0.02 (3)					
7	< 0.02 (3)	< 0.02 (3)	0.02 < 0.02 (2)	< 0.02 (3)	0.04 0.05 0.02	< 0.02 (3)					
9	< 0.02 (3)	< 0.02 (3)	< 0.02 (3)	< 0.02 (3)	0.03 0.03 0.03	< 0.02 (3)					
11	< 0.02 (3)	< 0.02 (3)	0.02 < 0.02 (2)	< 0.02 (3)	0.04 0.03 0.03	< 0.02 (3)					
14	< 0.02 (3)	< 0.02 (3)	< 0.02 (3)	< 0.02 (3)	0.03 0.03 0.03	< 0.02 (3)					
16	< 0.02 (3)	< 0.02 (3)	<0.02 (2) 0.02	< 0.02 (3)	0.05 0.23 0.03	<0.02 0.03 <0.02					
18	< 0.02 (3)	< 0.02 (3)	< 0.02 (3)	< 0.02 (3)	0.04 0.14 0.03	<0.02 0.03 <0.02					
21	< 0.02 (3)	< 0.02 (3)	0.03 < 0.02 (2)	< 0.02 (3)	0.05 0.05 0.04	< 0.02 (3)					
23	< 0.02 (3)	< 0.02 (3)	0.02 < 0.02 0.03	< 0.02 (3)	0.064 0.03 0.02	< 0.02 (3)					
25	< 0.02 (3)	<0.02 (3)	0.02 < 0.02 (2)	< 0.02 (3)	0.072 0.02 0.03	<0.02 (3)					
28	< 0.02 (3)	< 0.02 (3)	0.02 < 0.02 (2)	< 0.02 (3)	0.05 0.04 0.03	<0.02 (3)					

Table 63. Residues in the milk of lactating dairy cows dosed with glufosinate-ammonium + NAG (15+85) in gelatin capsules at levels equivalent to 9.1, 27 and 91 ppm glufosinate free acid in the diet for 28 consecutive days (Czarnecki and Brady, 1995b).

Groups of 20 white leghorn laying hens (each bird weighing 1.4-1.5 kg) were dosed with glufosinate-ammonium + NAG L-isomer, 15+85 in gelatin capsules at nominal levels equivalent to 0.36, 1.1 and 3.6 ppm glufosinate free acid in the diet for 28 days (Czarnecki and Brady, 1995a; Helsten, 1995; Crotts and McKinney, 1995). On day 29 twelve hens from each group were slaughtered. The remaining hens from each group were placed on a residue-free diet and killed on days 35 and 42. Eggs were collected daily.

The dosing levels were chosen on the assumption of feed consumption of 180 g/bird/day, but the birds consumed 133-160 g feed each per day. The actual dosing levels would be about 20% higher than intended.

No residues of glufosinate + NAG or MPP were found above the LOD in any of the tissues or eggs (Table 64).

Table 64. Residues in the tissues and eggs of hens dosed with glufosinate + NAG, (15+85) at levels equivalent to 0.36, 1.1 and 3.6 ppm in the feed for 28 days (Czarnecki and Brady, 1995a).

Commodity	Residues expressed as glufosinate free acid, mg/kg						
	0.36 ppm		1.1	ppm	3.6 ppm		
	glufosinate +	MPP	glufosinate +	MPP	glufosinate +	MPP	
	NAG		NAG		NAG		
Skin	< 0.05	< 0.05	< 0.05	< 0.05	< 0.05	< 0.05	
Muscle	< 0.05	< 0.05	< 0.05	< 0.05	< 0.05	< 0.05	
Liver	< 0.1	<0.1	< 0.1	<0.1	< 0.1	<0.1	
Fat	< 0.05	< 0.05	< 0.05	< 0.05	< 0.05	< 0.05	
Eggs ¹	< 0.05	< 0.05	< 0.05	< 0.05	< 0.05	< 0.05	

¹Eggs from days 1, 3, 5, 7, 9, 11, 14, 16, 18, 21, 23, 25 and 28 were analysed.

FATE OF RESIDUES IN STORAGE AND PROCESSING

The effects of commercial processing on residues in sugar beet, maize, soya bean seed and canola seed were reported.

Zietz and Simpson (1997)processed tolerant sugar beet roots (80-100 kg in each trial) from three glufosinate-ammonium residue trials in France, Germany and the UK (Hees and Werner, 1997d) and measured the residues in the processed fractions. The crops were each treated twice with glufosinate-ammonium at 0.80 kg ai/ha with the final treatment at GS19 (9 or more leaves unfolded) or GS31 (beginning of crop cover formation) and harvest 111-124 days later. The process (Figure 6) simulated a commercial operation.

The results are shown in Table 65.

Residues generally tended to remain in the juice and ultimately to appear in the molasses. Residues were not detected (<0.05 mg/kg) in the raw sugar.

Processing factors were calculated only for the glufosinate + NAG residues because the MPP residues in the sugar beet were too low to be useful. Processing factors for raw sugar could not be determined because residues were not found in the sugar in any of the trials. The processing factors for molasses were 5.0, 11.5 and 3.8 (mean 6.8) showing that glufosinate residues in the roots become concentrated in the molasses after removal of water.



Figure 6. Sugar beet processing.

Table 65. Glufosinate residues in sugar beet roots and processing fractions arising from glufosinateammonium treatment of transgenic sugar beet in trials in Germany (DEU010102), France (FRA000202) and the UK (GBR000102) (Hees and Werner 1997d, Zietz and Simpson 1997).

Commodity	Residues, mg/kg as glufosinate free acid						
	DEU010102		FRA0	FRA000202		000102	
	glufosinate + MPP		glufosinate +	MPP	glufosinate +	MPP	
	NAG		NAG		NAG		
Root	0.3	0.05	0.06	< 0.05	0.99	< 0.05	
Pressed pulp	< 0.05	< 0.05	< 0.05	< 0.05	0.13	< 0.05	
Raw juice	0.25	< 0.05	0.11	< 0.05	0.77	0.05	
Clarified juice	0.19	< 0.05	0.09	< 0.05	0.82	0.05	
Concentrated	0.72	0.14	0.55	0.05	2.5	0.17	
juice							

Commodity	Residues, mg/kg as glufosinate free acid						
	DEU010102		FRA000202		GBR000102		
	glufosinate + NAG	MPP	glufosinate + NAG	MPP	glufosinate + NAG	MPP	
Raw sugar	< 0.05	< 0.05	< 0.05	< 0.05	< 0.05	< 0.05	
Molasses	1.5	0.23	0.69	0.06	3.8	< 0.05	

Brady and Bertrand (1997) treated transgenic sugar beet, variety T-4A, with glufosinateammonium three times at 3.0 kg ai/ha, about 5 times the label rate and harvested the crop 136 days after the final treatment at the eighth leaf stage. The laboratory processing of 141 kg sugar beet roots was designed to simulate a commercial process (Englar, 1997) and was similar to that in Figure 6. The results are shown in Table 66.

In this trial, in contrast to the Zietz and Simpson trial, MPP was the main component of the residue. The total residue was used to calculate processing factors of 0 (<0.08) for refined sugar, and 6.4 for molasses.

Table 66. Glufosinate residues in sugar beet root and processing fractions arising from glufosinateammonium treatment of transgenic sugar beet in trials in the USA at 3.0 kg ai/ha and harvest 136 days after the last of three treatments (Brady and Bertrand, 1997).

Commodity	Residues, mg/kg as glufosinate free acid				
	glufosinate + NAG	MPP			
Root	0.27 0.24	1.0 0.96			
Dried pulp	0.14 0.15	0.56 0.63			
Refined sugar	<0.05 (2)	<0.05 (2)			
Molasses	1.7 1.4	6.5 6.1			

Czarnecki (1996) showed that the residues in grain dust were about 10 times as high as in the grain from glufosinate-ammonium-treated tolerant maize. The grain dust contained 0.62 mg/kg of MPP and 3.0 mg/kg glufosinate + NAG from grain containing MPP and glufosinate + NAG at 0.062 and 0.31 mg/kg respectively, all expressed as glufosinate free acid. Residue levels increased as particle size decreased, with levels in the <425 μ m fraction 2.5-3 times those in the >2450 μ m fraction.

To provide samples for a processing study in the USA Brady *et al.* (1994) treated glufosinate-tolerant maize (60 cm stage) with glufosinate-ammonium at 0.50 kg ai/ha (3 times the proposed label rate) in a trial in Iowa. In a second trial in Nebraska glufosinate-ammonium was applied at 0.36 kg ai/ha to 30 cm maize with a second treatment at 0.50 kg ai/ha at the 60 cm growth stage. The maize was treated by wet and dry milling processes (Figure 7). The results are shown in Table 67.

The residues were all below the LOD (0.05 mg/kg) in the Iowa trial. NAG was present at the LOD in the meal, flour and grits and at 0.1 mg/kg in the hulls in the Nebraska trial. The best estimates for processing factors are corn meal, corn flour and grits 1, hulls 2.

Table 67. Residues in maize and processed fractions resulting from dry and wet milling of glufosinate-tolerant maize treated at excessive rates with glufosinate-ammonium (Brady *et al.* 1994).

Commodity	Iowa trial			Nebraska trial		
		F	Residues, mg/	kg as glufosinate		
	Glufosinate	MPP	NAG	Glufosinate	MPP	NAG
Maize	< 0.05	< 0.05	< 0.05	< 0.05	< 0.05	0.054
Corn meal (dry milled)	< 0.05	< 0.05	< 0.05	< 0.05	< 0.05	0.051
Corn flour (dry milled)	< 0.05	< 0.05	< 0.05	< 0.05	< 0.05	0.053
Corn hulls (dry milled)	< 0.05	$<\!\!0.05$	< 0.05	< 0.05	< 0.05	0.10
Corn grits (dry milled)	< 0.05	$<\!\!0.05$	< 0.05	< 0.05	< 0.05	0.05
Corn crude oil (dry milled)	< 0.05	$<\!\!0.05$	< 0.05	< 0.05	< 0.05	< 0.05
Corn refined oil (dry milled)	< 0.05	< 0.05	< 0.05	< 0.05	< 0.05	< 0.05
Corn hulls (wet milled)	< 0.05	< 0.05	< 0.05	< 0.05	< 0.05	< 0.05
Corn starch (wet milled)	< 0.05	< 0.05	< 0.05	< 0.05	< 0.05	< 0.05
Corn crude oil (wet milled)	< 0.05	< 0.05	< 0.05	< 0.05	< 0.05	< 0.05
Corn refined oil (wet milled)	< 0.05	< 0.05	< 0.05	< 0.05	< 0.05	< 0.05

Figure 7. Wet and dry milling of maize (Brady et al., 1994).



Maize, wet milling

Maize, dry milling

Brady (1995e) processed seed from transgenic soya bean crops treated at excessive rates with glufosinate-ammonium and measured the residues, expressed as glufosinate free acid, in

		soya bea	n seeds	grain dust		
	Trial	glufosinate	MPP	glufosinate	MPP	
		+ NAG		+ NAG		
, I	IN-01	3.4 mg/kg	1.1 mg/kg	31mg/kg	8.7 mg/kg	
l	MO-01	1.5 mg/kg	1.8 mg/kg	4.3mg/kg	4.7 mg/kg	
		-	-	-	-	

the aspirated grain fractions (grain dust). In the two trials 32 kg of soya bean seed produced 78 and 228 g of grain dust. Residues in the grain dust were 8.6 and 2.8 times the levels in the grain in the two trials.

Czarnecki *et al.* (1994a) treated a transgenic soya bean crop in the USA (Iowa) with glufosinate-ammonium at 2.6 kg ai/ha (5 times the normal rate) at the six-trifoliate growth stage and harvested the soya beans for processing 96 days later. After processing 31 kg of the soya beans (Figure 8) the hulls contained the highest residues with lower levels in the meal. Residues of



Figure 8. Simulated commercial processing of soya beans

glufosinate, NAG and MPP were not detected (<0.05 mg/kg) in the refined or crude oil. The results are shown in Table 68.

The calculated processing factors for residues in the meal and crude or refined oil are 1.3 and 0 (<0.3) respectively.

Table 68. Residues of glufosinate, NAG and MPP in seeds and processed commodities from glufosinate-treated transgenic soya beans (Czarnecki *et al.*, 1994a).

Commodity	Residues as glufosinate free acid, mg/kg					
	glufosinate	NAG	MPP			
Seeds	< 0.05	0.07	0.08			
Hulls	0.07	0.27	0.30			
Meal	< 0.05	0.08	0.12			
Refined oil	< 0.05	< 0.05	< 0.05			
Crude oil	< 0.05	< 0.05	< 0.05			

MacDonald (1996a) treated a transgenic canola crop in Canada (Saskatchewan) with single applications of glufosinate-ammonium at 0.75, 1.5 and 3.8 kg ai/ha (up to 4 times the normal rate) at the 4-6 leaf growth stage and harvested the seed (3.5-5 kg) for processing 70 days later.

Processing was on a small scale but was intended to simulate commercial practice. The canola seed was dried and cleaned and, after conditioning, crude oil was produced first by pressing and then by hexane extraction of the cake. The cake was toasted. The crude oil was refined, bleached and deodorised. The results are shown in Table 69.

Glufosinate itself was not detected in the seed or any processed fraction. The main residue was NAG. No residues were detected in the oils. The residues of NAG remain with the meal and

change little during toasting. The calculated processing factor for untoasted meal is 3.1 (2.7, 3.1, 3.7) and for toasted meal 3.7 (3.1, 3.3, 4.8).

Table 69. Residues in seed and processed fractions from transgenic canola treated with glufosinate at three rates and harvested 70 days later in Canada (MacDonald 1996a).

Commodity	Residues, mg/kg as glufosinate free acid									
	0.75	kh ai/ha		1.5	1.5 kg ai/ha			3.8 kg ai/ha		
	glufosinate	MPP	NAG	Glufosinate	MPP	NAG	glufosinate	MPP	NAG	
Seed	< 0.05	< 0.05	0.063	< 0.05	< 0.05	0.060	< 0.05	< 0.05	0.21	
Meal, untoasted	< 0.05	< 0.05	0.17	< 0.05	< 0.05	0.22	< 0.05	0.11	0.64	
Meal, toasted	< 0.05	< 0.05	0.21	< 0.05	0.05	0.29	< 0.05	0.11	0.64	
Oil, crude	< 0.05	< 0.05	< 0.05	< 0.05	< 0.05	< 0.05	< 0.05	< 0.05	< 0.05	
Oil, refined	< 0.05	< 0.05	< 0.05	< 0.05	< 0.05	< 0.05	< 0.05	< 0.05	< 0.05	
Oil, bleached	< 0.05	< 0.05	< 0.05	< 0.05	< 0.05	< 0.05	< 0.05	< 0.05	< 0.05	
Oil, deodorized	< 0.05	< 0.05	< 0.05	< 0.05	< 0.05	< 0.05	< 0.05	< 0.05	< 0.05	
Soapstock	< 0.05	< 0.05	< 0.05	< 0.05	< 0.05	< 0.05	< 0.05	< 0.05	0.083	

Residues in the edible portion of food commodities

The only information was the from processing studies on soya bean, canola and sugar beet.

The processing factors for the residues in soya bean meal and crude or refined oil were 1.3 and 0 (<0.3) respectively. Residues were not detected (<0.05 mg/kg) in the crude or refined oil.

The processing factor for residues in untoasted canola meal was 3.1 and for toasted meal 3.7. Residues were not detected (<0.05 mg/kg) in the crude, refined, bleached or deodorized oil.

The processing factors for raw sugar and molasses in one trial were 0 (<0.35) and 6.8 respectively. In another trial the processing factors for refined sugar and molasses were 0 (<0.08) and 6.4 respectively. No residues were detected (<0.05 mg/kg) in the raw or refined sugar.

RESIDUES IN FOOD IN COMMERCE OR AT CONSUMPTION

No information.

NATIONAL MAXIMUM RESIDUE LIMITS

The Meeting was informed of the following national MRLs.

Country	MRL,	Commodity
-	mg/kg	
Argentina 1/	0.05	corn grain, corn forage, corn fodder
	<u>2</u> /	top fruits, stone fruits, citrus, grapes
Australia 3/	15	mixed pasture (legume/grasses)
	5	edible offal (mammalian)
	0.2	assorted tropical and sub-tropical fruits – inedible peel
	0.1	berries and other small fruits, citrus fruits, meat (mammalian), tree nuts, pome fruits
	0.05	milks, stone fruits
Austria <u>4</u> /	0.1	all crops
Belgium <u>5</u> /	2	potatoes

Country	MRL,	Commodity
	mg/kg	
	0.05	all other crops
Brazil <u>6</u> /	0.05	lettuce, cotton, banana, citrus, potatoes, coffee, apple, maize, peach, Savoy-cabbage,
		soya beans, wheat, grape, bean
Canada <u>7</u> /	6	lentils
	3	rape seed (canola)
	1	liver and kidneys of cattle, goats, hogs, poultry and sheep
	0.5	dry white beans
	0.4	potatoes
	0.2	corn
Colombia	<u>8/</u>	cotton, banana, soya bean, papaya, citrus, grape, strawberry, passion-flower, African palm, potatoes, coffee
Denmark		2/
Finland		2/
France <u>10</u> /	0.5	citrus, small berries, bananas, cherries, stone fruits, pome fruits, hazel nuts, tree nuts, olives, carrots, chicory, cabbage, spinach, beans, lettuce, lamb's lettuce, sweet maize, onions, potatoes
Germany 11/	3	pulses, sunflower seeds with hulls
	1	potatoes, rape seed
	0.5	currants, kiwifruit
	0.2	bananas, citrus
	0.1	other food of plant origin
Greece		2/
Ireland		<u>9</u> /
Italy	0.1	citrus, small berries and small fruits, stone fruits, pome fruits, tree nuts, hazel nuts,
-		kiwis, onion, cabbage, lettuce, beans, carrots, kohlrabi, radish, asparagus, potatoes, soya beans
Japan	0.5	potato, rice, tea
	0.3	mandarin, other citrus, large fruits, small fruits, oil seeds, nuts
	0.2	wheat, other cereals, vegetables, beans (immature + pods), leafy vegetables, rhizomes,
	0.1	bulb vegetables, mushrooms
T 1 5/	0.1	soya bean
Luxemburg <u>5</u> /	0.2	potatoes
	0.05	inuits and vegetables
Netherlands <u>12</u> /	0.5	
	0.05	fruit, vegetables
N	0-0.05 <u>13</u> /	other food commodities
Norway		
Poland	3	legume vegetables
D . 114/	0.2	fruits, vegetables (except legume vegetables)
Portugal <u>14</u> /	3	barley
	1	wheat
	0.5	potatoes
	0.1	cıtrus, berries, brassicae
	0.1 <u>15</u> /	tea, hop
	0.05 <u>15</u> /	pome fruits, stone fruits, other fruits, roots and tubercles, bulbs, fruits and vegetables, hard shelled fruits, leaf vegetables and fresh spice plants, fresh leguminosae, stalks,
		fungi, leguminosae (dried) grains, oil seeds, other cereals
Spain <u>16</u> /	0.05	fruits, vegetables, potatoes, cereals
	0.01 <u>17</u> /	legumes, oily seeds, tea and other infusions, dried hops, spices, tobacco, sugar beet, sugarcane, forages and hays, dried products (raisins, plums, etc)
South Africa <u>18</u> /	2	soya beans
	0.2	canola, maize
Sweden		<u>9</u> /
Switzerland	0.5	potatoes
	0.05	fruits and vegetables
UK <u>1</u> 9/	5.0	barley
	3.0	peas

Country	MRL,	Commodity
-	mg/kg	
	1.0	potatoes, wheat, linseed, field beans
	0.5	rape seed
USA	25	aspirated grain fractions (grain dust) 20/
	6	corn stover <u>20</u> /
	5	soya bean hulls <u>20</u> /
	4	corn forage <u>20</u> /
	2	soya beans <u>20</u> /
	0.5	almond hulls <u>21</u> /
	0.3	banana (0.2 in pulp) <u>22</u> /
	0.2	corn grain <u>20</u> /
	0.1	meat by-products such as kidneys and liver of cattle, goats, hogs, horses, poultry and sheep $\underline{21}/$
	0.1	tree nuts <u>21</u> /
	0.05 <u>21</u> /	apples, cattle fat, cattle meat, eggs, goats fat, goats meat, grapes, hogs fat, hogs meat,
		horses fat, horses meat, poultry fat, poultry meat, sheep fat, sheep meat
	0.02	milk <u>21</u> /

1/ Definition of the residue: glufosinate-ammonium, expressed as glufosinate free acid

- 2/ For these crops Argentina follows MRLs established in Brazil or Germany.
- 3/ Definition of the residue: sum of glufosinate-ammonium and 3-[hydroxy(methyl)phosphinoyl]propionic acid, calculated as glufosinate (free acid).
- 4/ Definition of the residue: DL-homoalanin-4-yl-(methyl)phosphinic acid: in total calculated as glufosinate
- 5/ Definition of the residue: DL-homoalanin-4-yl-(methyl)phosphinic acid and 3-methylphosphinicopropionic acid calculated as glufosinate free acid.
- 6/ Definition of the residue: glufosinate-ammonium, expressed as glufosinate free acid
- <u>7</u>/ Definition of the residue: 4-(hydroxy-(methyl)-phosphinoyl)-DL-homoalaninate ammonium salt, including the metabolite 3-methyl-phosphinicopropionic acid.
- 8/ Definition of the residue: the authorities accept the MRLs established by Codex and EPA
- 9/ No MRLs have been established by Denmark, Finland, Greece, Ireland, Norway or Sweden. Codex MRLs or those established in other EU countries are accepted
- 10/ Definition of the residue: 4-(hydroxy(methyl)phosphinoyl)-D,L-homoalanine
- 11/ Definition of the residue: DL-homoalanin-4-yl-(methyl)phosphinic acid and 3-methylphosphinicopropionic acid calculated as glufosinate
- 12/ Definition of the residue: glufosinate-ammonium, expressed as glufosinate.
- $\underline{13}$ / Range of 0 0.05 mg/kg because the LOD is 0.05 mg/kg
- 14/ Definition of the residue: glufosinate-ammonium and its metabolites 2-acetamido-4-methylphosphinico-butanoic acid and 3-methylphosphinicopropionic acid, expressed as glufosinate
- $\underline{15}$ / At the limit of determination
- 16/ Definition of the residue: glufosinate and its ammonium salt, expressed as glufosinate;
- 17/0.01 mg/kg is based on an earlier lower limit of quantification but will be adjusted to 0.05 mg/kg in near future for harmonisation purposes
- 18/ Definition of the residue: Glufosinate-ammonium and its metabolites 2-acetamido-4-methylphosphinico-butanoic acid and 3-methylphosphinicopropionic acid, expressed a glufosinate free acid equivalents. MRLs are temporary until 1999 and will be re-evaluated in the light of JMPR evaluation and Codex Alimentarius standards.
- 19/ Definition of the residue: DL-homoalanin-4-yl-(methyl)phosphinic acid and 3-methylphosphinicopropionic acid calculated as glufosinate free acid.
- <u>20</u>/ Definition of the residue: glufosinate-ammonium and its metabolites 2-acetamido-4-methylphosphinico-butanoic acid and 3-methylphosphinicopropionic acid. Time-limited tolerance expiring 13 July 1999.
- <u>21</u>/ Definition of the residue: glufosinate and its metabolite 3-methylphosphinico-propionic acid. Time-limited tolerance expiring 13 July 1999.
- 22/ Definition of the residue: combined residues of glufosinate ammonium and 3-methylphosphinicopropionic acid expressed as glufosinate equivalents. Time-limited tolerance expiring 18 Jan 2000.

APPRAISAL

Glufosinate-ammonium was first evaluated for residues and toxicology by the 1991 JMPR and subsequently for residues in 1994. Glufosinate-tolerant crops have now been developed with new GAP and different residue patterns, requiring new MRLs.

The Meeting received information on metabolism and environmental fate, registered uses and supervised residue trials on tropical fruits, tree nuts and various genetically-modified field crops. Feeding and processing studies were also reported.

Studies on the metabolism of glufosinate-ammonium in genetically modified rape seed (canola), sugar beet, maize, soya beans and tomatoes showed that the tolerant crops rapidly converted the active glufosinate isomer (L-glufosinate) to *N*-acetyl-glufosinate (NAG). The main components of the residue in tolerant plants are the L-isomer of NAG and D-isomer of glufosinate.

The Meeting received information on animal metabolism studies on rats, lactating goats and laying hens.

When <u>rats</u> were dosed orally with L– $[3,4-^{14}C]$ NAG some de-acetylation to glufosinate occurred and some glufosinate was bioavailable, but unchanged NAG was the main component (85-89%) of the TRR in the faecal extracts. Almost all of the administered ¹⁴C was excreted in the faeces within 4 days.

When rats were dosed orally with $[3,4-^{14}C]$ glufosinate-ammonium, 75-89% of the ^{14}C was excreted in the faeces and 8-11% in the urine within 48 hours. The main components in the faecal extracts were glufosinate (77% of administered ^{14}C), NAG (7.5%), 4-methylphosphinico-2-hydroxybutanoic acid or MHB (4.3%) and 3-methylphosphinicopropionic acid or MPP (1.3%). The main components in the urine were glufosinate (4.3%) and MPP (0.8%).

Very small amounts of the administered ¹⁴C were found in the tissues (<0.1%) and milk (<0.02%) of a lactating goat dosed orally for 4 days with [3,4-¹⁴C]glufosinate. Levels of ¹⁴C reached a plateau in the milk by day 2. Levels of ¹⁴C were higher in the kidneys and liver than in other tissues. Glufosinate and MPP constituted about 50% and 30% respectively of the residue in both kidneys and liver. Glufosinate accounted for 50% of the ¹⁴C in the milk. When a lactating goat was dosed orally with L-[3,4-¹⁴C]NAG the disposition of ¹⁴C in the tissues and milk was similar to that after dosing with glufosinate. Glufosinate was the main residue in the kidneys, liver and milk with NAG and MPP forming a substantial part of the residue in the kidneys and liver.

Less than 0.02% of the administered ¹⁴C was present in the edible tissues when <u>laying hens</u> were dosed orally for 14 days with $[3,4-^{14}C]$ glufosinate-ammonium. MPP and glufosinate were the main residues identified in the liver and eggs respectively. After dosing orally for 14 days with L- $[3,4-^{14}C]$ NAG less than 0.1% of the administered dose was present in the edible tissues and blood. NAG was the main residue identified in liver and egg yolks while glufosinate was the main residue in egg whites.

The Meeting received information on metabolism studies in rape seed, canola, sugar beet, maize, soya beans and tomatoes.

Genetically modified <u>rape</u> plants rapidly acetylated glufosinate. Cut rape plants were placed in a nutrient solution containing $[3,4-{}^{14}C]$ glufosinate-ammonium for 6 days, by which time 57% of the ${}^{14}C$ in the plants was associated with NAG and 36% with glufosinate.

In glufosinate-tolerant <u>canola</u> plant tissues sampled 1 hour after treatment with $[^{14}C]$ glufosinate, 73% and 18% of the ^{14}C was present as glufosinate and NAG respectively,

demonstrating very rapid acetylation of glufosinate. After 21 days 60%, 21% and 7% of the ¹⁴C corresponded to NAG, glufosinate and MPP respectively.

When glufosinate-tolerant <u>sugar beet</u> plants were sprayed with $[3,4-^{14}C]$ glufosinateammonium the racemic isomer composition in the surface residue was unchanged, but in the absorbed residue L-glufosinate was metabolised to L-NAG.

Glufosinate was generally a minor component of the residue in treated tolerant <u>maize</u>. NAG was the main residue in the forage, silage and fodder, while MPP was the major component in grain, cobs and husks. The GLC enforcement analytical method for glufosinate, MPP and NAG was in reasonable agreement with a radiolabel method for the residues in the forage.

NAG was the main residue in the forage, straw, pods and beans of treated tolerant <u>soya bean</u> plants. MPP levels exceeded glufosinate levels in the pods and beans. The GLC enforcement method and an HPLC radiolabel method were in reasonable agreement in analyses of forage, straw, pods and beans at the higher residue levels, but at low levels the result from the enforcement method was less than from the ¹⁴C measurement.

The translocation of [¹⁴C]glufosinate-ammonium from treated leaves to shoots, other leaves and roots was approximately 4 times as fast in glufosinate-resistant <u>tomato</u> plants as in susceptible tomatoes. Most of the surface residue was glufosinate itself, but this was very rapidly converted to NAG once absorbed into the leaves. NAG constituted essentially all the residue in the ripe fruit harvested 60 or 74 days after the plants were treated.

The Meeting received information on the degradation and dissipation of glufosinateammonium in soil, residues in rotational crops and fate in water-sediment systems.

Glufosinate disappeared with a half-life of about 3-6 days during aerobic incubation with a sandy loam soil. MPP, the major product, reached its maximum after about 14 days incubation. MPA, 2-methylphosphinicoacetic acid, became the main residue after long intervals. Glufosinate suffered 62% and 31% mineralization during 120 days incubation with and without incorporation of plant material into the soil respectively.

In a dissipation study glufosinate-ammonium, applied 3 times to bare ground, dissipated quickly with calculated half-lives of 15, 7.2 and 2.7 days, the increased rates probably being related to increased soil moisture and temperature. The estimated half-lives for MPP after its residues peaked were 38, 14 and 16 days, and for MPA 25, 19 and 7 days. No residues were detected below a 45-60 cm depth section, but glufosinate and MPP reached a depth of 30-45 cm on several occasions during the study. Glufosinate and its degradation products have some mobility in soil but their rapid dissipation ensures that travel down the soil profile is limited.

When L-[3,4-¹⁴C]NAG was incubated in a sandy loam soil it was very rapidly converted to L-glufosinate, which was then itself broken down and mineralized. The comparable rate of mineralization and production of unextractable residues suggests that the degradation pathway for NAG is through glufosinate. Further experiments showed that the half-life of NAG was only hours.

No important degradation products other than CO_2 were identified when $[2-^{14}C]MPA$ was incubated in a sandy loam under aerobic conditions. Estimated decline and mineralization half-lives were 24 and 74 days respectively. Degradation in a loamy sand was much slower. When MPP was incubated under aerobic conditions in a sandy loam, MPA was the only significant product after 120 days.

After 3 days incubation of a sandy loam soil with tolerant tomato leaves containing residues, mainly of glufosinate and NAG, most of the residue had been converted to MPP, which itself was degraded more slowly to MPA and ultimately to CO_2 .

Residues of degradation products of glufosinate should be undetectable or at very low levels in rotational crops. When radishes, lettuce and wheat were sown in a confined rotational crop study 28 days after glufosinate treatment of bare ground (to simulate re-sowing a failed crop) residues of MPP and MPA were present at low levels in the crops, demonstrating possible uptake of these compounds. The pattern of residues was similar in the three crops. When sowing was 119 days after treatment (to simulate a following crop) residues were not detectable in lettuce or radishes but MPP and MPA were detected by ¹⁴C methods at very low levels in the wheat grain and straw.

The photolytic breakdown of glufosinate in natural waters was very slow.

MPP became the major component of the residue within a few days when [3,4-¹⁴C]glufosinate-ammonium was incubated in a <u>water-sediment</u> system at 20°C. Glufosinate itself disappeared with a half-life of 3 days, but only 25% mineralization occurred during the 361 days of the study. In other experiments the rates of degradation were shown to be affected by the source of the water and the residue level of glufosinate (with faster disappearance at lower levels). In all cases most of the residue was in the water phase.

Methods of residue analysis

The main components of the residue in genetically modified tolerant crops are glufosinate, NAG and MPP. Analytical methods have been designed to measure the three components separately or, because glufosinate and NAG produce the same derivative in the analytical procedure, to measure glufosinate and NAG combined and MPP separately. Residues are extracted from the finely ground sample with water, and the extract is cleaned up on an anion exchange resin column. After solvent exchange, NAG and MPP are separated from glufosinate on a cation exchange column. The residues are taken up in glacial acetic acid and methylated, and glufosinate acetylated, with trimethyl orthoacetate in refluxing acetic acid. After solvent exchange and final clean-up on a silica gel cartridge the derivatized residues are determined by GLC with flame-photometric detection.

Modifications of the extraction and initial clean-up are needed for samples such as maize oil, fats and milk. A variation of the method dispenses with the cation exchange separation and determines glufosinate and NAG as a combined GLC peak because both compounds produce the same analytical derivative. The LOD for crop samples is typically 0.05 mg/kg for each analyte. Analytical recoveries have been extensively tested and found satisfactory on many substrates.

Analysts should be aware that transgenic glufosinate-tolerant soya beans plants can convert L-glufosinate to NAG very rapidly, giving apparently low analytical recoveries in spiked samples.

Glufosinate, NAG and MPP were shown to be stable during frozen storage for intervals of 12, 15 or 24 months in the following substrates: genetically modified maize and soya beans and their processed commodities, cow and chicken tissues, milk, eggs, susceptible maize grain, and transgenic rape seed and sugar beet roots. The 1994 JMPR reported that residues of glufosinate and MPP (described as Hoe 061517) in apples, oranges, kiwifruit, maize, soya beans and almonds were stable during frozen storage.

Some samples from the supervised trials were stored for 2 years, but the storage stability studies have demonstrated that the residues were still stable.

Definition of the residue

The current definition includes glufosinate and MPP and is based on the residues occurring in conventional crops. When glufosinate is used on glufosinate-tolerant crops NAG is produced. It should be included in the residue definition for enforcement because NAG is generally the main component of the residue and because the same derivative is produced in the analytical method from glufosinate itself and NAG, and in the simplified method both appear in the GLC peak from their common derivative. The revised residue definition is also suitable for commodities from conventional crops because if NAG is absent it will not contribute to the analytical result and if present at low levels it is necessarily included in the analytical result.

A suitable revised residue definition would be *Sum of glufosinate-ammonium, 3-(hydroxy(methyl)phosphinoyl)propionic acid and N-acetyl-glufosinate expressed as glufosinate (free acid)*, but the Meeting could not consider the adoption of this definition until the toxicological evaluation of NAG had been completed.

The residue reported in the supervised trials consists of three components, but is often reported with the glufosinate and NAG residue combined. The metabolism studies show that residues of the main component constitute 65-75% of the combined residue when all three components are at measurable levels. It follows that if all three components are below the LOD a reasonable assumption is that the combined residue is also below or close to the LOD. When one component is above and the others are below the LOD, the combined residue is assumed to be equal to the residue of the main component.

The method of calculating the total residue for various situations is illustrated by the following example.

Glufosinate	MPP	NAG	Total
< 0.05	< 0.05	< 0.05	< 0.05
< 0.05	< 0.05	0.06	0.06
0.05	< 0.05	0.09	0.14

Information was made available on uses of glufosinate around fruit trees and nut trees. Limited information was provided on GAP for use of glufosinate on transgenic crops.

Glufosinate-ammonium is registered for use in Australia as a directed spray for weed control around avocados, bananas, feijoa, guava, kiwifruit, litchis, mangoes, papaya, passion fruit, pineapples and rambutans at 0.20-1.0 kg ai/ha. Malaysia has similar registered uses for glufosinate-ammonium as a directed herbicide spray around bananas, carambola, durians, guava, jack fruit and mangoes at 0.3-0.5 kg ai/ha. Residues in the fruit would generally not be expected from this type of use. Glufosinate itself is not taken up by roots but MPP, the main degradation product in soil, can be absorbed by the roots and translocated through the crop.

Supervised trials

Supervised trials were reported on tropical fruits, nut trees, maize, soya beans, rape, canola and sugar beet.

Residues were not detected in <u>avocados</u> in 3 Australian trials where glufosinate was used at 1.0, 1.2 and 2.0 kg ai/ha. In one trial at 1.2 kg ai/ha, glufosinate was detected at 0.06 mg/kg (presumably direct contamination during application) in 1 sample on day 0, but in no other samples. Residues were not detected in the fruit from 3 Australian trials on <u>mangoes</u> where the application rates were 1.0, 1.2 and 2.0 kg ai/ha or 1 trial on <u>papayas</u> at 1.2 kg ai/ha.

Residues were not detected in the fruit from Malaysian trials on <u>carambola</u> and <u>guavas</u> (2 trials at 0.5 kg ai/ha and 2 at 1 kg ai/ha on each fruit).

The Meeting agreed to consider these fruits together as "tropical fruits with inedible peel" but noted that a CXL of 0.2 mg/kg had already been established for banana, which would have to be excluded from the group. The residues in rank order from the 16 trials were <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05, <0.05,

In view of the type of use and generally unquantifiable residues at 3-5 sampling intervals in all of the trials the Meeting estimated a maximum residue level of 0.05* mg/kg and an STMR of 0.05 mg/kg for glufosinate in Assorted tropical and sub-tropical fruits – inedible peel (except banana).

The US registered use for glufosinate around <u>almond</u>, <u>pecan</u> and <u>walnut</u> trees permits a directed application at 1.7 kg ai/ha, with no more than 5.1 kg ai/ha total per year, and a 14-day PHI. MPP residues were 0.07 mg/kg in almonds from a US trial at the label rate and up to 0.22 mg/kg in a trial at twice the label rate. MPP is the main degradation product in soil, so its presence demonstrates the possibility of root uptake rather than contamination of foliage by spray. Residues were not detected in almonds from 3 other trials at the label rate or 3 trials at the double rate. The trials were in California in 1985.

Residues were not detected (<0.05 mg/kg) in nuts from 3 trials on pecans (USA, 1985) with glufosinate applied at 1.7 kg ai/ha and 3 at twice that rate. In 2 trials nuts were harvested 21 days after application rather than 14 days, but the use pattern is sufficiently close to GAP. Residues were not detected (<0.05 mg/kg) in nuts from 6 walnut trials at 1.7 kg ai/ha, 5 trials 3.4 kg/ha or 3 at other rates higher than GAP in the USA in 1985.

The Australian registration for glufosinate-ammonium permits 0.2-1.0 kg ai/ha as a directed spray around nut trees. Residues were not detected (<0.1, <0.05 mg/kg) in <u>macadamia</u> nuts in Australian trials from 1992 and 1995 at the label rate (2 trials) and double rate (2 trials).

Italian registration permits a directed application of 0.5-1.6 kg ai/ha (maximum 2.5 kg ai/ha per year) for weed control around <u>hazelnuts</u>. Residues were not detected (<0.05 mg/kg) in nuts from 5 hazelnut trials in Italy in 1985 according to GAP.

The Meeting considered the 4 almond, 6 pecan, 14 walnut, 4 macadamia and 5 hazelnut trials as a group. The residues in the 33 trials were <0.05 (30), 0.07 and <0.1 (2) mg/kg. The Meeting estimated a maximum residue level of 0.1 mg/kg and an STMR level of 0.05 mg/kg for glufosinate in tree nuts.

Glufosinate-ammonium is registered in Canada for use on transgenic <u>maize</u> at 0.30-0.50 kg ai/ha with final application at the 8-10 leaf stage. In Canadian trials in 1994-5 no residues (<0.05 mg/kg) were detected in 7 trials at 0.5 or 0.6 kg ai/ha or 3 trials at 1.0 kg/ha applied at growth stage GS 17-19 (7-9 leaves), or in 4 trials at 0.6 kg ai/ha at a later growth stage (GS 33, 3 nodes detectable).

Glufosinate-ammonium is registered in Portugal for use on transgenic maize at 0.40-0.80 kg ai/ha with the final application when the plant has no more than 8-10 leaves. In 7 Italian trials and 3 Spanish trials in 1996 according to Portuguese GAP no residues (<0.05 mg/kg) were detected in the harvested maize.

The German registration for glufosinate-ammonium on tolerant maize allows a single application of 0.9 kg ai/ha during the 3-8 leaf growth stage or 2 applications 6 weeks apart at 0.45 kg ai/ha with the final application at the 8-leaf stage. Residues were below the LOD (0.05 mg/kg) in 13 trials in Germany where the trial conditions (2 applications of 0.45-0.60 kg ai/ha, the second at GS 18

or GS 19, i.e. 8 or 9 leaves) were considered to comply with the registered use, 22 trials in France in accord with German GAP, and in 5 trials in Germany and 7 in France where glufosinate-ammonium was used at excessive rates (2 applications of 0.80 kg ai/ha).

Glufosinate-ammonium is registered for use in the USA on tolerant maize with 2 applications at 0.23-0.41 kg ai/ha, the second at a plant height of 60 cm with harvest 70 days later. Supervised trials on maize in the USA at rates of 0.40-0.50 kg ai/ha (1 or 2 applications) at the nominated growth stage were accepted as equivalent to the maximum GAP. The PHI (70 days) was considered secondary to the growth stage in deciding the timing of the applications. In many cases the grain was harvested 90-120 days after the final treatment. The residues (glufosinate + NAG + MPP expressed as glufosinate) in the 35 trials according to GAP were <0.05 (29), 0.05 (2) and 0.07 (4) mg/kg.

In summary, the residues of glufosinate + NAG + MPP expressed as glufosinate in tolerant maize were Canada <0.05 (14) mg/kg, Italy <0.05 (7) mg/kg, Spain <0.05 (3) mg/kg, Germany <0.05 (18) mg/kg, France <0.05 (29) mg/kg, the USA <0.05 (29), 0.05 (2) and 0.07 (4) mg/kg. The residues in rank order were <0.05 (100), 0.05 (2) and 0.07 (4) mg/kg.

The Meeting estimated a maximum residue level of 0.1 mg/kg for glufosinate in maize and noted that this was equivalent to the existing CXL. The major residue in the glufosinate-tolerant crop is NAG, which is not included in the current residue definition. NAG was included in the reported residues where the derivatization GLC procedure without an ion-exchange separation step was used, and was below the LOD when determined separately.

Glufosinate is registered for use on transgenic glufosinate-tolerant <u>canola</u> in Canada with 1 application at 0.60 kg ai/ha or 2 applications of 0.30-0.50 kg ai/ha, the final application at the early bolting growth stage. The label carries the instruction not to graze the treated crop or cut for hay. The Meeting was informed that the 10 leaves stage is very close to bolting. Only two supervised trials in Canada met the condition of final treatment at the 10-leaf stage, each with 1 application of 0.5 or 0.75 kg ai/ha. Residues were not detected (<0.05 mg/kg) in either of the trials, but 2 trials were insufficient to support a recommendation.

Glufosinate-ammonium is registered for use in the USA on tolerant soya beans with 2 applications at 0.23-0.41 kg ai/ha, the second application at bloom with harvest 70 days later. Supervised trials on soya beans in the USA with 2 applications of 0.40-0.50 kg ai/ha at the nominated growth stage were accepted as equivalent to the maximum GAP. The PHI (70 days) was considered as secondary to the growth stage in deciding the timing of the applications. In the trials the grain was harvested 62-102 days after the final treatment. The residues of glufosinate + NAG + MPP expressed as glufosinate in the 20 trials according to GAP in rank order (median underlined) were 0.32, 0.39, 0.43, 0.52, 0.56, 0.71, 0.72, 0.78, 0.81, <u>0.85</u>, <u>0.89</u>, 0.92, 0.96, 1.02, 1.24, 1.26, 1.33, 1.56, 1.64 and 1.88 mg/kg.

The Meeting estimated a maximum residue level of 2 mg/kg for soya beans but could not recommend it as suitable for use as an MRL until the toxicological evaluation of NAG had been completed.

Residues in almond hulls from 3 trials on almonds at the label rate and 3 at twice that rate (see above) were all <0.5 mg/kg. Unfortunately, residues in the hulls were not determined in the 2 trials where MPP was detected in the almonds. However, the Meeting estimated a maximum residue level of 0.5 mg/kg for almond hulls, which is recommended for use as an MRL.

The registered use of glufosinate-ammonium on tolerant maize in the USA (see above) specifies pre-harvest intervals of 60 and 70 days for maize forage and fodder respectively. The supervised trials on maize in the USA described above, which complied with GAP, showed residues

of glufosinate + NAG + MPP expressed as glufosinate in the forage or silage harvested 60-92 days after treatment (the residue is reasonably persistent) in rank order, median underlined, of 0.09, 0.12, 0.12, 0.13, 0.17, 0.2, 0.26, 0.28, 0.3, 0.32, 0.33, 0.36, 0.4, 0.48, 0.53, 0.53, 0.54, 0.54, 0.68, 0.74, 0.78, 0.78, 0.79, 0.9, 1.07, 1.1, 1.19, 1.2, 1.45, 1.67, 1.7, 1.71, 1.76, 2.9 and 3.48 mg/kg.

The residues of glufosinate + NAG + MPP expressed as glufosinate in maize fodder harvested 84-122 days after treatment were 0.07, 0.08, 0.11, 0.12, 0.15, 0.22, 0.25, 0.25, 0.32, 0.33, 0.43, 0.5, 0.53, 0.58, 0.65, 0.68, 0.69, <u>0.72</u>, 0.8, 0.94, 0.95, 1.13, 1.19, 1.32, 1.42, 1.5, 1.69, 1.74, 1.76, 1.78, 1.96, 2.31, 2.65, 2.83 and 5.4 mg/kg.

The Meeting estimated maximum residue levels of 5 mg/kg in maize forage and 10 mg/kg in maize fodder but could not recommend them as suitable for use as MRLs until the toxicological evaluation of NAG had been completed.

Feeding studies on lactating dairy cows and laying hens were reported.

Lactating dairy cows were dosed with glufosinate-ammonium + NAG (15 + 85% to simulate the typical terminal residue) at rates equivalent to 9, 27 and 91 ppm glufosinate free acid equivalents in the diet for 28 days. Residues were not detected in the milk at the 9 ppm feeding level, but reached a plateau on day 3 at the 91 ppm level. Residues (glufosinate, NAG and MPP) in the tissues were below the LODs at the lower feeding levels; at 91 ppm they were detected in the kidneys and liver, but not in muscle or fat. MPP was the major residue in the liver.

<u>Laying hens</u> were dosed with glufosinate-ammonium + NAG (15 + 85) at levels equivalent to 0.36, 1.1 and 3.6 ppm glufosinate free acid equivalents in the diet for 28 days. Residues (glufosinate, NAG and MPP) were not detected in the tissues or eggs.

Studies on the fate of residues during the commercial food processing of sugar beet, maize, soya beans and canola were reported.

When tolerant <u>sugar beet</u> were treated with glufosinate-ammonium and processed, residues of glufosinate, NAG and MPP tended to remain in the juice and ultimately appear in the molasses. Residues were not detected (<0.05 mg/kg) in the raw sugar. The processing factors for glufosinate + NAG for raw sugar in 3 trials were nominally 0 (<0.17), 0 (<0.83) and 0 (<0.05) on the basis of the residues in the roots. The mean processing factor for molasses was 6.8, showing that glufosinate residues are concentrated in the molasses on evaporation of the water. In another study on sugar beet with fivefold application rates MPP was the main residue component and again no residues were detected in the raw sugar. The calculated processing factors were 0 (<0.08) for refined sugar and 6.4 for molasses.

In 2 processing trials (wet and dry milling) of glufosinate-tolerant maize treated with excessive rates of glufosinate-ammonium, only NAG was quantifiable in the grain from only one trial. In this trial no residues (<0.05 mg/kg) were detected in crude or refined oil from either process or in the starch and hulls from the wet milling process. The estimated processing factors were meal 1, grits 1, and hulls 2, all from dry milling.

Residues of glufosinate, NAG and MPP were not detected (<0.05 mg/kg) in refined or crude oil in a processing study with a transgenic <u>soya bean</u> crop treated with glufosinate-ammonium at a fivefold rate. The calculated factors for processing seed to meal and seed to crude or refined oil are 1.3 and 0 (<0.3) respectively.

Glufosinate itself was not detected in the seed or any processed fraction of a transgenic canola crop treated at a fourfold rate with glufosinate-ammonium. The main residue was NAG. The

calculated processing factor for untoasted meal was 3.1 and for toasted meal 3.7. No residues (<0.05 mg/kg) were detected in the oils.

Information was made available to the Meeting on national MRLs. Governments have adopted a variety of residue definitions.

RECOMMENDATIONS

On the basis of data from supervised trials the Meeting estimated the maximum residue levels and STMRs listed below. The maximum residue levels are recommended for use as MRLs.

Definition of the residue for compliance with MRLs and for the estimation dietary intake: sum of glufosinate-ammonium and 3-[hydroxy(methyl)phosphinoyl]propionic acid calculated as glufosinate (free acid)

Commodity		Recommended MRL, mg/kg		STMR
CCN	Name	New	Previous	
FI 0030	Assorted tropical and sub- tropical fruits - inedible peel ¹	0.05*	-	0.05
FI 0341	Kiwifruit	W^2	0.05*	
TN 0085	Tree nuts	0.1	-	0.05
AM 0660	Almond hulls	0.5	-	

¹Except Banana

²Replaced by recommendation for group MRL

*At or about the limit of determination.

DIETARY RISK ASSESSMENT

Estimated STMRs for glufosinate-ammonium on tropical fruit and tree nuts have been added to the previous list of MRLs (21) for other commodities. The estimated dietary intakes of glufosinate-ammonium expressed as glufosinate for the 5 GEMS/Food regional diets were in the range of 3 to 10% of the ADI. The Meeting concluded that the intake of residues of glufosinate-ammonium resulting from its uses that have been considered by the JMPR is unlikely to present a public health concern.

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