Table 1 Results for the GM ingredients checks in 2024.

Food containing Maize (Corn)							
Sample No.	Sample description	Country of origin	Food chain stage	GM Event	Level	Result	Observations
1	Coarse cornmeal	Ireland	Retail	Bt-11, MON863, MON88017, MON89034, NK603 and TC1507	NQ	Compliant	(a)
2	Corn chips	Ireland	Retail	-	-	Compliant	-
3	Corn snack	Malaysia	Retail	-	-	Compliant	-
4	Cornflour	Ireland	Retail	-	-	Compliant	-
5	Cornflour	Ireland	Retail	-	-	Compliant	-
6	Cornflour	UK	Not stated	-	-	Compliant	-
7	Organic polenta	Italy	Retail	-	-	Compliant	-
8	Popcorn	Ireland	Retail	-	-	Compliant	-
9	Popcorn	Argentina	Not stated	MON89034	NQ	Compliant	(b)
10	Popcorn	South Africa	Manufacturer	-	-	Compliant	-
11	Popcorn	Brazil	Not stated	-	-	Compliant	-
12	Tortilla chips	Ireland	Manufacturer	MON810, MON89034 and TC1507	NQ	Compliant	(c)
13	Tortilla chips	Ireland	Manufacturer	-	-	Compliant	(d)

14	Tortilla chips	Ireland	Manufacturer	MON 89034 and TC1507	NQ	Compliant	(e)		
15	Tortilla chips	UK	Retail	-	-	Compliant	-		
16	Tortilla wraps	Ireland	Manufacturer	-	-	Compliant	-		
17	Tortilla wraps	Not stated	Packer	-	-	Compliant	-		
Food containing Rice									
Sample No.	Sample description	Country of origin	Food chain stage	GM Event	Level	Result	Observations		
1	Basmati rice	Ireland	Retail	-	-	Compliant	-		
2	Basmati rice	Ireland	Retail	-	-	Compliant	-		
3	Basmati rice	Not stated	Wholesale/Distribution	-	-	Compliant	-		
4	Brown rice drink	UK	Retail	-	-	Compliant	-		
5	Jasmine rice	Thailand	Not stated	-	-	Compliant	-		
6	Jasmine rice	Cambodia	Not stated	-	-	Compliant	-		
7	Long grain rice	Ireland	Retail	-	-	Compliant	-		
8	Long grain rice	Ireland	Retail	-	-	Compliant	-		
9	Long grain rice	United states	Not stated	-	-	Compliant	-		
10	Long grain rice	UK	Retail	-	-	Compliant	-		
11	Long grain rice	Portugal	Manufacturer	-	-	Compliant	-		
12	Long grain rice	Thailand	Retail	-	-	Compliant	-		

13	Parboiled rice	Uruguay	Wholesale/Distribution	-	-	Compliant	-	
14	Rice	India	Retail	-	-	Compliant	-	
15	Rice	UK	Retail	-	-	Compliant	-	
16	Rice & corn snack	UK	Retail	-	-	Compliant	-	
17	Rice & corn snack	UK	Not stated	-	-	Compliant	-	
18	Rice & corn snack	Ireland	Not stated	-	-	Compliant	-	
19	Rice cakes	UK	Not stated	-	-	Compliant	-	
20	Rice cakes	Ireland	Retail	-	-	Compliant	-	
21	Rice cakes	Ireland	Retail	-	-	Compliant	-	
22	Rice cakes	UK	Retail	-	-	Compliant	-	
23	Rice cakes	Ireland	Retail	-	-	Compliant	-	
24	Rice crackers	Taiwan	Not stated	-	-	Compliant	-	
25	Rice crackers	Ireland	Not stated	-	-	Compliant	-	
26	Rice-based cereals	UK	Not stated	-	-	Compliant	-	
27	Rice noodles	China	Not stated	-	-	Compliant	-	
	Food containing Soybean							
Sample No.	Sample description	Country of origin	Food chain stage	GM Event	Level	Result	Observations	

1	Edamame beans	Ireland	Retail	-	-	-	-
2	Fermented soya	Ireland	Retail	MON89788	NQ	Compliant	(f)
3	Miso instant soup	Ireland	Retail	-	-	Compliant	(g)
4	Soy protein chunks	Ireland	Retail	-	-	Compliant	-
5	Soy protein mince	Ireland	Not stated	-	-	Compliant	-
6	Soya ball	Ireland	Retail	-	-	Compliant	-
7	Soya beans	Canada	Retail	MON89788	NQ	Compliant	(h)
8	Soya beans	Ireland	Retail	MON89788	NQ	Compliant	(h)
9	Soya beans	Ireland	Retail	-	-	Compliant	-
10	Soya chunks	India	Retail	-	-	Compliant	-
11	Soya drink	Ireland	Retail	-	-	Compliant	-
12	Soya drink	Ireland	Retail	-	-	Compliant	-
13	Soya drink	Ireland	Retail	-	-	Compliant	-
14	Soya drink	Ireland	Retail	-	-	Compliant	-
15	Soya protein	Ireland	Retail	-	-	Compliant	-
16	Tofu	Not stated	Not stated	GTS 40-3-2	NQ	Compliant	(i)
17	Tofu	Germany	Retail	-	-	Compliant	-

- (a) Detection of the genetic elements: the rice actin 1 intron sequence (I-rActI), the 35s promoter (p-35S), the FMV promoter (p-FMV), the NOS terminator (t-NOS), the E9 terminator (t-E9), the *cry1Ab/Ac*, *cry3Bb1*, *pat*, and *cp4-epsps* genes, and the *ctp2-cp4epsps* construct. The GM maize events Bt-11, MON863, MON88017, MON89034, NK603 and TC1507 were detected; however, no quantification was performed due to too high Cycle Threshold (Ct) values observed during the PCR amplification process.
- (b) Detection of the genetic elements: the 35s promoter (p-35S), the FMV promoter (p-FMV) and the NOS terminator (t-NOS). The GM maize event MON89034 was detected; however, no quantification was performed due to too high Ct values observed during the PCR amplification process.
- (c) Detection of the genetic elements: the rice actin 1 intron sequence (I-rActI), the 35s promoter (p-35S), the NOS terminator (t-NOS), the *pat* gene and the *ctp2-cp4epsps* construct. The genetic markers for both endogenous maize and rice were detected. The GM maize events MON810, MON89034 and TC1507 were detected; however, no quantification was performed due to too high Ct values observed during the PCR amplification process.
- (d) Detection of the genetic elements: the 35s promoter (p-35S) and the ctp2-cp4epsps construct. No GM maize events were detected.
- (e) Detection of the genetic elements: the rice actin 1 intron sequence (I-rActI), the 35s promoter (p-35S), the FMV promoter (p-FMV), the NOS terminator (t-NOS), the *pat* gene and the *ctp2-cp4epsps* construct. The GM maize events MON89034 and TC1507 were detected; however, no quantification was performed due to too high Ct values observed during the PCR amplification process.
- (f) Detection of the genetic elements: the E9 terminator (t-E9) and the *ctp2-cp4epsps* construct. The GM soya event MON89788 was detected; however, no quantification was performed due to too high Ct values observed during the PCR amplification process.
- (g) The genetic markers for both endogenous rice and soyabean were detected.
- (h) Detection of the genetic elements: the E9 terminator (t-E9), the FMV promoter (p-FMV) and the *ctp2-cp4epsps* construct. The GM soya event MON89788 was detected; however, no quantification was performed due to too high Ct values observed during the PCR amplification process.
- (i) Detection of the genetic elements: the 35s promoter (p-35S), the NOS terminator (t-NOS) and the E9 terminator (t-E9). The GM soya event GTS 40-3-2 was detected; however, no quantification was performed due to too high Ct values observed during the PCR amplification process. NQ, not quantified.