



## Effects of Feeding Genetically Modified Crops to Domestic Animals: A Review<sup>#</sup>

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ARTICLE INFO	ABSTRACT
<p><sup>#</sup>This study was presented as an oral presentation at the 4th International Anatolian Agriculture, Food, Environment and Biology Congress (Afyonkarahisar, TARGİD 2019)</p> <p><i>Review Article</i></p> <p>Received : 26/06/2019 Accepted : 08/10/2019</p> <p><b>Keywords:</b> GM-crops Ruminant Poultry Performance Health</p>	<p>Genetically modified (GM) crops are being planted at large scale worldwide. In most of the countries, GM crops are processed into livestock feed. The land is used for cultivation of GM plants has been increased in recent years; in 2012 GM plants were grown on over 170 million hectares in 28 countries by 17.3 million farmers and extended to 185.1 million hectors in 2016 worldwide. GM plants have been used as feed for animals and the number of studies has proved their safety for animal and public health. This paper reviews the possible effects of GM crops on livestock, poultry, and aquatic animals by reviewing different type of studies, in which parameters such as performance, reproductive and health assessment were investigated. The most of peer-reviewed papers evaluating the effects of feeding animals with transgenic crops were based on GM plants with improved agronomic traits i.e. herbicide-tolerant plants and pets-tolerant plants; however, in some cases GM plants with boosted nutritional properties assessed. In most experiments, either Bt (<i>Bacillus thuringiensis</i>) maize, Roundup Ready (RR) soybean or both fed to animals. Measurable differences in various parameters were mostly observed in Bt maize and soybean fed separately or simultaneously to animals. In this review, scientific studies showing the effects of the use of GM products in the nutrition of domestic animals on performance, health and reproductive parameters are investigated.</p>

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### Introduction

Thousands of years ago, plants with favorable characteristics have been produced with the conventional breeding methods. Traits of desirable characteristics are selected, combined and propagated by sexual crossing method to produce new varieties, but it takes times. New technology has been introduced which not only overcome the hindrance of sexual incompatibility between variable plant species (Southgate et al., 1995) but also used to improve the nutritional contents of plants.

All GM crops are transgenic plants which have been genetically modified by using genetic engineering techniques (generally called Recombinant DNA technology) (Sticklen, 2005; Conard U, 2005; Ma JKC et al., 2003; James, 2011). Nowadays, large amounts of GM crops are being used as feed material in animals (Flachowsky et al., 2012).

The commonly grown transgenic crops are maize, cotton, soybean, rice, and canola (rapeseed) in which a gene has been introduced to plants by transgenesis either for insect resistance or herbicide tolerance (Flachowsky et al., 2012). GM maize is introduced mainly insect-resistant Bt (*Bacillus thuringiensis*) genes (James, 2011) and GM Roundup Ready (RR) soybeans are incorporated with herbicide tolerance genes (Sieradzki et al., 2006). The main focus of these studies to evaluate the safety of feeding GM crops to animal by considering animal and public health. In Europe, GM crop safety is assessed by the panel on GMO (Genetically Modified Organism) of the European Food Safety Authority (EFSA) based on comparative studies of their molecular, compositional, phenotypic, and agronomic traits and its near-isogenic equivalent crops (EFSA, 2011; EFSA, 2015). Hence, most studies evaluated the feeding GM crops effect on animal health and most on performance

parameters such as daily weight gain, feed conversion, dry matter intake, and reproduction (Aumaitre, 2004; Flachowsky et al., 2007). The effect of Bt maize and RR soybean on the performance of cattle, pigs, and poultry are most extensively studied in table 1-4. Bt maize contains the cry1Ab gene obtained from the *Bacillus thuringiensis* which showing resistance to a wide range of Lepidoptera insects. Mammals lack the intestinal receptors for the Cry1Ab protein produced by this gene and are not believed to be affected by this toxin (Schnepf et al., 1998).

To date, considerable number of review articles on GM crops and their effects on livestock animals have been published up to 2014. In this article, literature are investigated on effects of feeding GM crops on domestic animals from the period 2014-2019 using the search engines like Scopus, Web of Sciences, Google Scholar and PubMed. Scientific publications reported on the experiments transgenic plants were fed to ruminants, pigs, poultry, or aquatic animals and their effect on health parameters were examined. The observed effects of these GM crops on various parameters in animals are summarized in table 1-4. For this purpose, only studies containing the control groups were selected. In published studies, this review evaluated the performance parameters (average daily weight gain, body weight, and feed conversion), reproductive traits (litter size, birth weight) and health parameters (body condition score, organ weight,

hematology, serum biochemistry, histopathology, clinical examination, immune response, and gastrointestinal microbiota).

### Effects of Feeding with GM Crops on Ruminants

Three studies with maize (MON 810) and soybean (Roundup Ready, MON 40-3-2) separately or simultaneously were conducted (Furgal Dierżuk et al., 2014; 2015; Tudisco et al., 2015) in ruminants (Table 1), significant differences between GM crops fed animals and control animals observed by (Tudisco et al., 2015). They evaluated the in vivo and post mortem carcass traits and immunoglobulin G concentration in kids born from goats fed GM and non-GM soybean. Furthermore, they also investigated the goat colostrum quality, in terms of chemical composition, immunoglobulin concentration, and the presence of feed DNA fragments. Birth weight significantly ( $P<0.05$ ) higher in the non-GM fed group while only carcass weights were significantly affected by the treatment resulting in GM soybean-fed group. Colostrum from GM-fed groups showed a significantly lower percentage of protein and fat, either serum or colostrum IgG concentration (mg/mL). Transgenic DNA sequences were detected in colostrum from the animals receiving GM-soybean meal but not observed in samples from the goats fed with non-GM soybean.

Table 1 Summary of studies testing the effects of GM-crops on ruminants

GM crop	Summary of studies
1	Country: Poland, Production stage: 10 days old bull Experimental design: 1 group of non-GM maize and soybean meal (n=10), 1 group of non-GM maize and GM soybean (n=10), 1 group of GM maize and non-GM soybean meal (n=10), 1 group of GM maize and GM soybean meal (n=10) Duration of experiment: 90 days Studied parameters: Performance parameters Observations: No adverse effects of GM components References: (Furgal Dierżuk et al., 2014)
2	Country: Poland, Production stage: Three weeks before parturition Experimental design: 1 group of non-GM maize and soybean meal (n=10), 1 group of non-GM maize and GM soybean (n=10), 1 group of GM maize and non-GM soybean meal (n=10), 1 group of GM maize and GM soybean meal (n=10) Duration of experiment: 305 <sup>th</sup> days of lactation Studied parameters: Performance parameters, milk composition, blood serum metabolite profiles Observations: No adverse effects of GM components References: (Furgal-Dierżuk et al., 2015)
3	Country: Italy, Production stage: 60 days before kidding Experimental design: 1 group of control (n=10), 1 group of treatment (n=10) Duration of experiment: Until 30 days of kids age Studied parameters: Goat colostrum quality, kid birth weight, and carcass weight Observations: Colostrum from GM-fed groups showed a significantly lower percentage of protein and fat and IgG concentration. Birth weight significantly higher in the non-GM fed the group while only carcass weights were significantly affected by the treatment resulting in the GM-fed group. References: (Tudisco et al., 2015)

1: Bt maize (MON 810) and soybean meal (Roundup Ready, MON 40-3-2), 2: Maize (MON 810) and soybean meal (Roundup Ready, MON 40-3-2), 3: GM soybean (MON40-3-2)

Furgal Dierżuk et al. (2014) performed an experimental study to determine the effects of GM maize (MON 810) and soybean meal (Roundup Ready, MON 40-3-2) on performance parameters, basal chemical composition of the *musculus thoracis* (MT), fatty acid composition of intramuscular fat, and transfer of transgenic DNA to calf

tissues, as well as the histological examination of organs and tissues of Polish Black and White Holstein Frisian calves. In this study, bulls aged 10 days were divided into four groups. The experiment was conducted for 90 days. There were no significant differences between all groups in final live weight, average daily weight gain, MT

chemical composition and fatty acid profile of intramuscular fat. The calf rumen fluid contained tDNA, but there was no tDNA in the intestinal content, blood, studied organs, or meat. Histological examination of the investigated organs and muscles found no differences among treatments.

Furgal Dierżuk et al. (2015) conducted same study on 40 Polish Holstein-Friesian cows to evaluate the effects of GM maize (MON 810) and soybean meal (Roundup Ready, MON 40-3-2) on performance parameters, milk composition, blood serum metabolite profiles and transfer of tDNA into the milk of cows. This study was conducted from the 3<sup>rd</sup> week before parturition to the 305<sup>th</sup> day of lactation. There were no significant differences between transgenic and non-transgenic feeds in productivity, milk composition and blood metabolite profiles such as  $\beta$ -hydroxybutyric acid, free fatty acids, glucose, insulin, and progesterone. The transgenic DNA sequences of MON 810 and RR soybean meal were not detectable by PCR in milk.

### Effects of Feeding with GM Crops on Pigs and Rabbits

Many kinds of research were conducted to evaluate the safety of GM feeding in pigs and rabbits either by giving Bt rice (Liu et al., 2018a, Liu et al., 2018b, Liu et al., 2017) or Bt maize (Chen et al., 2016, Yalçın et al., 2018) presented in Table 2. Chen et al., (2016) conducted a study to investigate the chronic effect of transgenic maize comprise of Cry1Ab protein (1.64 mg/kg) to evaluate the growth performance, immune response and health of Wuzhishan pig after 196-day GM maize feeding. Long-term feeding Bt corn results had no adverse effects of GM maize on pigs. Later on, Liu et al., (2017) performed an experiment to evaluate the chronic effects of Bt rice carrying the Cry1Ab protein (1.64 mg/kg) on offspring of highly inbred Wuzhishan pigs. The result showed that only difference occurs in the average daily gain and feed conversion ratio (FCR) of the females in week 3 and males from weeks 1 to 10 were different between both groups and the body weight of the male pigs in week 2 was greater in the non-GM group than that of the Bt group ( $P < 0.05$ ). There was no effect on sex steroid level, hematology parameters, relative organ weights, or histopathology. Consequently, long-term intake of transgenic rice had no adverse effects on pig offspring. In addition, Liu et al., (2018b) observed the effects of GM rice containing Cry1Ab protein (1.64 mg/kg) on male and female pigs at least for 360 days. The result showed that Bt rice had no effect on growth indexes, reproductive performances, hematology, and organ- morphology after long-term feeding except that total protein and bilirubin was higher in the non-GM group compared to the Bt group ( $P < 0.05$ ) but total bilirubin difference did not exist in male pigs ( $P > 0.05$ ).

The effects of GM maize in New Zealand rabbits were evaluated by conducting an experiment to study effects on liver and kidney organ weights, malondialdehyde (MDA) and glutathione (GSH) levels of liver and kidney tissues. In this study, rabbits were divided into three groups: parental, first-generation and second-generation and further divided into subgroup i.e. control group and the treatment group, on which control group was fed with conventional maize (non-GM) and treatment groups were fed with GM maize.

The results indicated an increase in liver weights and decrease in kidney weights observed as generations passed in GM maize-fed group. Significant decrease in GSH and increase in MDA in all three generations fed with GM maize (Yalçın et al., 2018).

### Effects of Feeding with GM Crops on Poultry

In a series of experimental studies, transgenic soybean meal, maize, canola meal, and rice included separately or simultaneously in the diet did not adversely influence the health status, reproduction traits, and cellular immune response of broiler chickens, laying hens and quails (Korwin-Kossakowska et al., 2016; Sartowska et al., 2015; Halle and Flachowsky, 2014; Gao et al., 2014; Zhong et al., 2016; Liu et al., 2016; Lili et al., 2017; McNaughton et al., 2014). All the studies are illustrated in Table 3a and 3b.

Czerwinski et al., (2015a), Czerwinski et al., (2015b), Li et al., (2015) and Hameed et al., (2016) also found no negative effect of dietary supplementation of GM maize, rice, soybean meal and sugarcane on hematological, biochemical and histopathological parameters of broilers.

Lili et al. (2017), McNaughton et al. (2014), Halle and Flachowsky (2014) and Sartowska et al. (2015) did not observe any adverse effect of feeding transgenic maize, rice, soybean and canola meal on growth performance parameters, carcass traits, nutrient digestibility and intestinal microbiota of broilers, laying hens and quails.

Kim (2019) conducted a study on two-day-old broilers for 32 days by using GM maize and soybean simultaneously. They evaluated the growth performance, nutrient digestibility, carcass weight and meat quality of broilers. They found that total lysine, methionine, and threonine of non-GM grains were lower than that of GM grains. The protein content of GM soybean meal was higher than that of non-GM soybean meal. Feed intake and FCR were greater ( $P < 0.05$ ) in broilers provided with non-GM diet than that of the GM group from day 17 to 32. A decrease in FCR has observed in birds fed the GM diet through the entire experiment ( $P < 0.05$ ). No significant impacts on blood profile, meat quality, and nutrient digestibility were found in response to dietary treatments.

Papineni et al., (2017) conducted a 42-day broiler study with diets containing toasted DAS-44406-6 soybean meal to evaluate nutritional wholesomeness and safety compared with non-transgenic, near-isoline soybean and non-transgenic references varieties as conventional comparators (Dairyland 99915, Porter 75148, and Williams 82). On measurements, only thigh weight was numerically different between birds fed DAS-44406-6 soybean meal and those fed isoline soybean meal. However, no significant differences to thigh weight were observed between birds fed DAS-44406-6 soybean and any of the non-transgenic reference varieties. Results indicate that DAS-44406-6 soybeans are nutritionally equivalent to conventional varieties.

Czerwiński et al., (2017) checked the effects of GM soybean meal and maize on the diversity and activity of microbiota inhabiting terminal gut segments in broiler chickens. In the ileum and caecum of all groups, members representing the orders *Clostridiales*, *Lactobacillales*, and *Selenomonadales* were present, accompanied by *Bifidobacteriales* in the caecum.

Table 2 Summary of studies testing the effects of GM-crops on pigs and rabbits

GM crop	Summary of studies
1	Country: China, Production stage: 3-month-old pigs Experimental design: Non-GM (female N= 7, male n = 7), GM rice (female n= 7, male n= 7) Duration of experiment: 75 days Studied parameters: Growth, reproductive performance, hematology, Histopathology organ morphology Observations: No effects of GM feeding observed References: (Liu et al., 2018a)
2	Country: China, Production stage: 3-month-old pigs Experimental design: Non-GM (female n = 6, male n = 5), Gm-rice (female n = 11, male n = 5) Duration of experiment: 360 days Studied parameters: Growth, hematology, organ weights and histopathology Observations: The average daily gain and feed conversion ratio of the female pigs in week 3 and the male pigs in weeks 1 were significantly different (P<0.05) between the Bt and isogenic groups. References: (Liu et al., 2017)
3	Country: China, Production stage: 3-month-old pigs Experimental design: Non-GM group (n= 26), GM group (n= 27) Duration of experiment: 360 and 420 days (Two generations) Studied parameters: Gut microbiota, histopathology Observations: No adverse effects of GM feeding observed References: (Liu et al., 2018b)
4	Country: China, Production stage: 40 days old pigs Experimental design: Isogenic corn–soybean meal-control diet, Bt corn–soybean meal-control diet Duration of experiment: 196 days Studied parameters: Growth, immune response, and health Observations: No adverse effects of GM feeding observed References: (Chen et al., 2016)
5	Country: Turkey, Production stage: Rabbit Experimental design: A control group (3 female, 3 male), GM maize treated group (3 female, 3 male) Duration of experiment: parental, first-generation and second-generation Studied parameters: Liver and kidney organ weights, malondialdehyde (MDA) and glutathione (GSH) levels of liver and kidney tissues Observations: Increase in liver weights and decrease in kidney weights observed as generations passed. Significant decrease in GSH and an increase in MDA in all three generations fed with GM maize. References: (Yalçın et al., 2018)

1: Bt rice (Cry1Ab protein), 2: Bt rice (Cry1Ab protein), 3: Bt rice (Cry1Ab protein), 4: Bt Maize (MON 810), 5: Bt Maize (MON 810)

The diversity of the order *Lactobacillales* in the ileum and caecum of birds fed GM maize was reduced, while that of *Lactobacillales* in the ileum and *Bifidobacteriales* in the caecum of birds fed GM soybean was higher compared with conventional maize and soybean. The use of GM and conventional maize and soybean did not affect the activity of microbiota measured as bacterial enzyme activity and the concentration of short-chain fatty acids in the ileal and caecal digesta. The GM maize did not change the resistance of *E. coli* or *Clostridium* against antibiotics, while GM soybean slightly increased the resistance of *Clostridium* from the ileum against kanamycin and those from caecum against kanamycin and erythromycin compared with conventional feedstuffs. In conclusion, the use of GM soybean and maize MON810 in diets did not affect the broiler intestinal ecosystem.

### Effects of Feeding with GM Crops on Aquatic Animals

Data on the effects of GM crops feeding on aquatic animals have mostly concerned Bt maize in which most of the scientific publications observed no adverse influence of GM supplementations except (Gu et al., 2014). All the studies are explained in Table 4. They observed that mortality, growth performance and body composition were

similar in fish fed the GM and non-GM maize varieties. The Bt maize-fed fish, however, displayed minor but significantly decreased digestive enzyme activities of leucine aminopeptidase and maltase, as well as the decreased concentration of gut bile salts, but significantly increased amylase activity at some sampling points. Histomorphological, radiographic and mRNA expression evaluations did not reveal any biologically relevant effects of Bt maize in the gastrointestinal tract, liver or skeleton. Cry1Ab protein or other compositional differences in GM Bt maize may cause minor alterations in intestinal responses in juvenile salmon, but without affecting overall survival, growth performance, development or health.

To check the biosafety of transgenic crops in aquatic animals, Zhu et al. (2015) conducted a study to evaluate the effects of Bt rice on frogs. They carried out the study for 90 days with 30% Bt rice, 30% parental rice or no rice as a control. On the assessment of biological, clinical, pathological and growth parameters, non-significant differences were found in body weight, body length, animal behavior, visceral organ weight, liver and kidney function, or the microstructure of some tissues between the froglets fed on the Bt rice containing diet and those fed on the parental rice or control diets.

Table 3a Summary of studies testing the effects of GM-crops on poultry

GM crop	Summary of studies
1	Country: South Korea, Production stage: Two-day-old male Ross 308 broilers Experimental design: 1 GM maize-soybean meal based Diet, 2 Non-GM maize-soybean meal based Diet Duration of experiment: 32 days Studied parameters: Growth performance, nutrient digestibility, carcass weight, and meat quality Observations: Feed intake and FCR were greater ( $P<0.05$ ) in broilers provided with non-GMO diet than that of the GMO group from d 17 to 32. A decrease in FCR has observed in birds fed the GMO diet through the entire experiment ( $P<0.05$ ). References: (Kim, 2019)
2	Country: USA, Production stage: At the day of the hatch (Ross 708) Experimental design: 7 dietary treatments Duration of experiment: 32 days Studied parameters: Performance and carcass yields Observations: No statistically significant differences were observed between treatments and control References: (McNaughton et al., 2014)
3	Country: China, Production stage: Day-old female Arbor Acres broilers Experimental design: 1 group fed with GM rice, 1 group fed with the parental line Duration of experiment: 42 days Studied parameters: Intestinal microbiota Observations: No adverse effects on the broiler intestinal microbiota References: (Lili et al., 2017)
4	Country: China, Production stage: Day old female Arbor Acres chicks Broiler Experimental design: 1 group fed with GM rice, 1 group fed with the parental line Duration of experiment: 42 days Studied parameters: Immunological assessment Observations: No significant differences were observed References: (Liu et al., 2016)
5	Country: China, Production stage: 55 weeks age of laying hen Experimental design: 1 control group (non-transgenic near-isoline corn), 1 group of GM maize diet, 1 group of reference corn diet Duration of experiment: Week 1 to 4, week 5 to 8 and week 9 to 12 Studied parameters: Growth, egg quality, organ health indicators Observations: No differences were observed in birds fed with treatments and control diet References: (Zhong et al., 2016)
6	Country: China, Production stage: 50-week-old laying hens Experimental design: 1 control group (n=72), 1 GM group (n=72) Duration of experiment: 16 weeks Studied parameters: Organ weight, serum biochemical parameters, and nutrient digestibility Observations: No adverse effects were found References: (Gao et al., 2014)
7	Country: Germany, Production stage: 31 week age of laying hens Experimental design: 1 control group (30 hens each and 3 cockerels), 1 GM group (30 hens each and 3 cockerels) Duration of experiment: 4 generations of laying hen Studied parameters: Animal health, laying performance, feed intake, feed efficiency and hatchability of chickens Observations: No significant influence was observed References: (Halle and Flachowsky, 2014)
8	Country: Poland, Production stage: Japanese quail Experimental design: 1 group of GM soya (Roundup Ready) and non-GM Maize, 1 group of GM maize and non-GM soya, 1 group of non-GM soya and maize. Duration of experiment: 10 generations of Japanese quail Studied parameters: Reproduction, survival rate, growth, egg-laying performance Observations: No effect between treatments and control References: (Sartowska et al., 2015)
9	Country: Poland, Production stage: Japanese quail Experimental design: 1 group of GM soya (Roundup Ready) and non-GM, Maize 1 group of GM maize and non-GM soya, 1 group of non-GM soya and maize. Duration of experiment: 10 generations of Japanese quail Studied parameters: Health status, necropsy findings, and organ histopathology Observations: No, any effect was observed References: (Korwin-Kossakowska et al., 2016)

Table 3b Summary of studies testing the effects of GM-crops on poultry

10	Country: Poland, Production stage: Day old broilers Experimental design: GM-maize groups (n = 24), 2 GM-soybean groups (n = 24), 2 GM-maize and GM-soybean groups (n = 24), 2 control groups (n = 24) Duration of experiment: 28 days Studied parameters: Hematology, histopathology, organ weight Observations: The lower proportion of T helper and T cytotoxic cells within lymphocytes in all GM-fed animals; higher spleen weight in GM-maize fed animals; increased the count of apoptotic cells in villi tips in GM-maize fed animals; higher width of tunica mucosa in the jejunum in GM-maize fed animals References: (Czerwinski et al., 2015a; Czerwinski et al., 2015b)
11	Country: Pakistan, Production stage: 2-week old broilers Experimental design: 3 GM-sugarcane groups (n = 10) Duration of experiment: 120 days Studied parameters: Serum biochemistry, histopathology, organ weight Observations: No effect of GM feed observed References: (Hameed et al., 2016)
12	Country: China, Production stage: Day old broilers Experimental design: 1 GM-rice group (n = 90), 1 control group (n = 90) Duration of experiment: 6 weeks Studied parameters: Serum biochemistry, histopathology, organ weight Observations: No effect of GM feed observed References: (Li et al., 2015)

1: Bt maize (MON 810) and soybean meal (Roundup Ready), 2: GM canola meal from event DP-Ø73496-4 glyphosate acetyltransferase (*gat4621*), 3: Bt rice (*Cry1Ac/Cry1Ab*), 4: Bt rice (*cry1Ab/cry1Ac*), 5: Bt maize (*mCry1Ac*), 6: Phytase transgenic maize, 7: Bt maize, 8: Bt maize (MON 810) and soybean meal (Roundup Ready), 9: Bt maize (MON 810) and soybean meal (Roundup Ready), 10: Bt maize (MON 810) and soybean meal (Roundup Ready), 11: Bt sugarcane (*Cry1Ac*), 12: Bt rice (*Cry1Ab/Ac*)

Table 4 Summary of studies testing the effects of GM-crops on aquatic animals

1	Country: Norway, Production stage: Juveniles Experimental design: 1 pair was a fish meal based, 1 pair was soybean meal based, (Each diet contained 20% Maize either Bt-maize or non-GM) Duration of experiment: 99 days Studied parameters: Survival, growth performance, digestive function, morphology, immune and stress response parameters Observations: The Bt-maize fed fish, however, displayed minor but significantly decreased digestive enzyme activities as well as decreased concentration of gut bile salts, but significantly increased amylase activity at some sampling points. References: (Gu et al., 2014)
2	Country: China, Production stage: Adult Zebrafish Experimental design: 1 group of 0.1 mg/L of proteins, 1 group of 1 mg/L of proteins, 1 group of 10 mg/L of proteins 1 group of Chlorpyrifos toxicant as a positive control Duration of experiment: 132-hour post fertilization Studied parameters: Developmental, biochemical, and molecular parameters Observations: No significant differences after Cry1C or Cry2A exposure. References: (Gao et al., 2018)
3	Country: China, Production stage: Froglets Experimental design: 1 group of 30% Bt rice, 1 group of non-GM rice, 1 control group with no rice, 1 group of Chlorpyrifos toxicant as a positive control Duration of experiment: 90 days Studied parameters: Growth, biological, clinical, and pathological assessments Observations: No significant differences were observed References: (Zhu et al., 2015)
4	Country: United States, Production stage: Adult zebrafish Experimental design: 1 control male group (given non-GM feed), 1 control female group (given non-GM feed), 1 experimental male group (given GM corn feed), and 1 experimental female group (given GM corn feed), Each diet contained 20% maize either Bt-maize or non-GM Duration of experiment: 8 weeks Studied parameters: Embryonic development, histology and transgenic DNA transfer Observations: No significant differences were observed between treatment and control groups. References: (Rayan et al., 2015)

1: Bt-maize (MON810), 2: Cry1C and Cry2A proteins, 3: Bt rice (*Cry1Ab/1Ac*), 4: Bt maize (MON89034× MON88017)

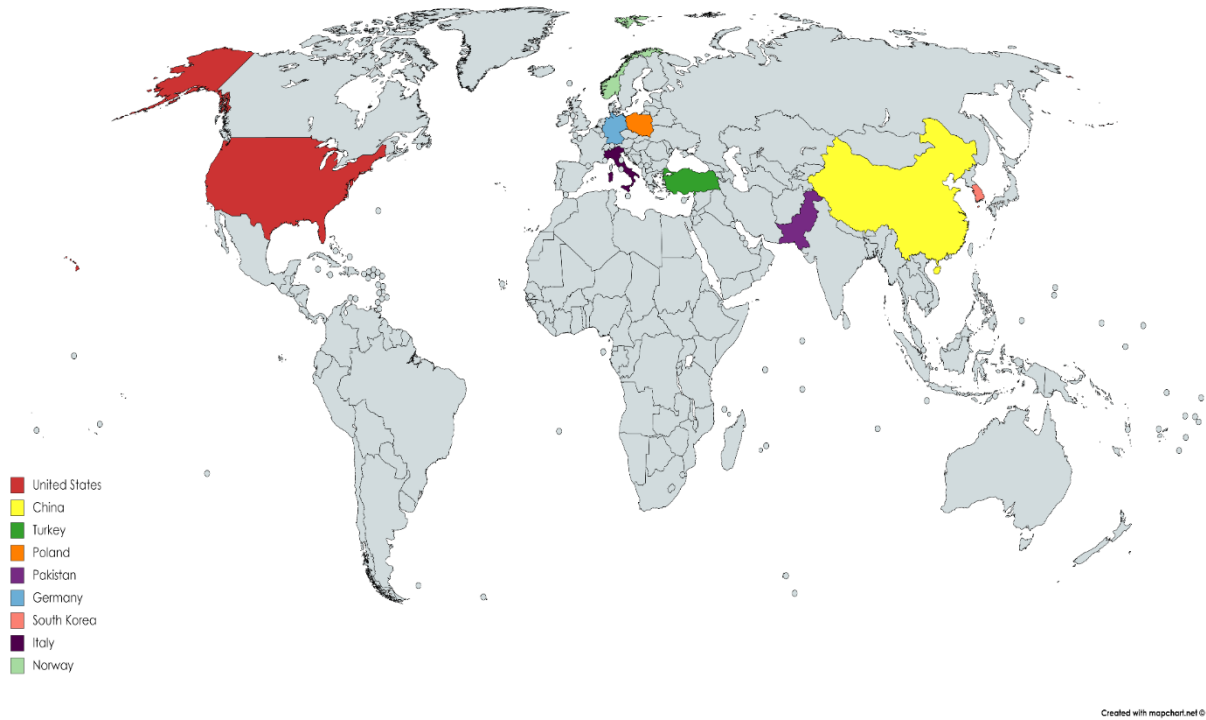


Fig 1 Country where the majority of studies investigated the effects of feeding transgenic crops to animals have been conducted during 2014-2019

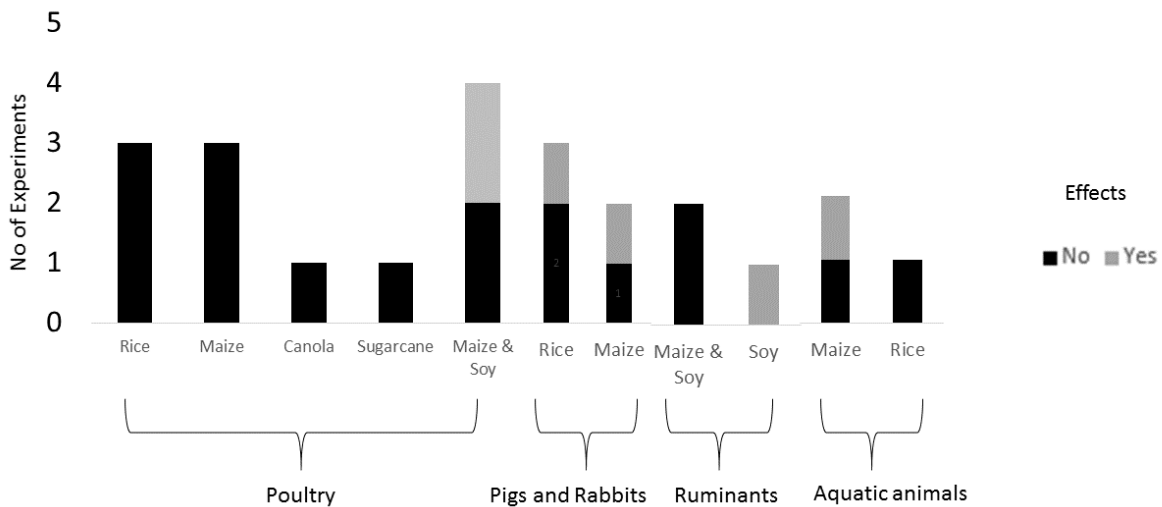


Fig 2 Number of experiments per animal species and GM crop in which various parameters were studied. Black bars indicate the experiments in which no health effects were observed. Grey bars indicate the experiments in which effects were observed (i.e. a significant difference ( $P < 0.05$ ) between GM-fed animals and control animals for at least one health parameter).

Rayan et al. (2015) assessed the effect of feeding stacked GM corn event (MON89034 × MON88017) on embryonic development, adult tissue histology and assessed the possibility of transgenic DNA transfer, using zebrafish as an animal model. No changes or abnormalities were observed in embryonic morphology nor in any of the

tissues examined histologically. In addition, no plant DNA reference genes or transgenic DNA were found in any of the analyzed samples. Results of these studies indicate that the safety and nutrition of GM crops are similar to non-GM crops and growth, health and development were not adversely affected by dietary intake of GM-crops.

## Conclusion

There are several studies that show different effects of feeding with GM feed sources in different animals. In other relevant studies of feeding GM crops, not only production parameters but also the metabolic status of animals were analyzed and only a few minor treatment differences have been found, with less or no biological relevance for poultry and livestock. The majority of studies investigating the effects of feeding transgenic crops to animals have been conducted in Europe and Asia with most experiments being performed in Poland and China. While few studies were performed in Germany, Norway, Italy, Pakistan, South Korea, and the USA (Fig. 1). Almost half of the experimental studies investigated hematology and histopathology, whereas about a quarter of the studies investigated organ weight, serum biochemistry, and the immune response. The significant differences in health, immune and production parameters between GM-fed animals and control animals were most often observed when GM maize and soybean were fed either separately or simultaneously, especially when fed to poultry (Fig. 2). Despite the presence of international guidelines for feeding trials with GM crops in animals, many published studies undergo from serious weaknesses with respect to experimental design, statistical analysis, and the use of non-GM feed for comparison purposes. The majority of the reviewed studies indeed fed the control animals with the near-isogenic counterpart of the GM crop that was evaluated, although the growing conditions of the crops were not always provided. A well-described protocol must be designed to evaluate the effects of GM feed on animal health particularly to evaluate the health effects of second-generation GM crops. In our review, only experiments with first-generation GM crops were included. Health effects of feed based on the second generation GM crops have not been evaluated yet in livestock experiments. For these GM crops feeding studies in target animals are a prerequisite to evaluate their effect on health parameters because their composition is substantially modified and no near-isogenic counterpart may be available. Based on this review, we concluded that there is no clear evidence that GM feed has adverse effects on animal production. The result presented in many studies shows that commercialized transgenic crops can be safely included in feed for animals and it can affect positively on the production as well as on health status of the animals so it could be safe to feed GM crops to livestock, poultry, and aquatic animals.

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