

Chapter 2

Sample Preparation and Chromatographic Analysis



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Abstract Globally, mycotoxins in food and feedstuff have been considered public food safety concerns because they could seriously affect animal production and human health. Mycotoxins could also cause detrimental damages to various organs such as the spleen, liver, heart, and immune system, eventually resulting in a substantial reduction of the animals and accordingly economic losses. Furthermore, multiple mycotoxins have been found in a variety of different foods, which might lead to acute and chronic effects on humans. Thus, the authentic methods for accurate determination of multicomponent mycotoxins have gained great attention from researchers to policy-makers. We summarized the combined analytical procedure of sample preparation and mycotoxin determination in different mycotoxin-producing matrices such as rice, maize, cereals, feedstuff, milk, fruit, and vegetable, as well as biological samples. As expected, it will be valuable references to the routine analytical protocols for confirmational lab analysis.

Keywords Sample preparation · LC-MS/MS · HPLC · GC · Mycotoxin

2.1 Introduction

Mycotoxins are a group of secondary fungal metabolites and thus can contaminate a wide range of food and other commodity crops at each step in the supply chain (Bennett and Klich 2009). These natural contaminants may lead to acute and chronic effects, both in humans and animals, including immunotoxicity, hepatotoxic, nephrotoxicity, carcinogenic, and genotoxicity (Nicholson et al. 2004). The most frequently found mycotoxins in food and feed are generally produced by the genera *Aspergillus*, *Penicillium*, and *Fusarium*, including aflatoxins, deoxynivalenol, fumonisins, zearalenone, and ochratoxin A (Irena Kralj and Helena 2009; Andrade

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et al. 2017). These chemical compounds are extremely stable in nature, and thus, they can also be co-occurring in fully processed foods. Therefore, they represent a major concern for animal and human health. Many studies have shown that multi-mycotoxins had been found in different matrices such as cereals, maize, rice, fruit, and vegetable and biological samples (Al-Taher et al. 2017; da Luz et al. 2017; de Oliveira et al. 2017; Juan et al. 2017a; Fan et al. 2019). In order to determine mycotoxin residues in various food matrices to protect consumers' health, different analytical methods have been proposed with high sensitivity and accuracy. Recently, there is a growing interest in analytical methods for simplified sample preparation and techniques for the determination of single and multiple mycotoxins in foods.

2.2 Sample Preparation

In general, the most common methods applied to quantification on the levels of mycotoxins were based on first-step extraction with organic solvents of acetonitrile and methanol, followed by cleanup and dilution step (Almeida et al. 2012; Chia-Ding et al. 2013; Frenich et al. 2009). However, none of these methods could avoid matrix effects altogether. Matrix effect is a complex from samples and analyte-specific phenomenon observed in the analytical method for mycotoxins that can be compensated by matrix-matched calibration, isotope-labeled internal calibration, and sample dilution (Slobodchikova and Vuckovic 2018). The isotope-labeled internal standards could compensate for any changes in ionization caused by the presence of matrix interferences. Therefore, the best means is to cope with matrix effects and thus ensure reliable results. However, the generic application of isotopically labeled internal standards was terribly restricted by the high costs and limited availability.

In the field of mycotoxin analysis, regardless any chosen technique, cleanup after extraction is usually required, which is very crucial for reliable results finally obtained. This procedure required high precision and accuracy in order to isolate mycotoxins of interest from the matrix interferences. For matrix removal and analyte enrichment, the main sample preparation techniques that have been used to determine mycotoxin contaminations in various food matrices are as follows: solid phase extraction (SPE), immunoaffinity columns (IAC), liquid-liquid extraction (LLE), and quick, easy, cheap, effective, rugged and safe (QuEChERS) (García-Moraleja et al. 2015a, b; Liao et al. 2015; Natalia et al. 2013).

2.2.1 Solid Phase Extraction

Effective extraction of multiple mycotoxins from real complicated samples is the promise to perform either qualitative or quantitative analyses. Solid phase extraction (SPE) is by far sample preparation method used in routine analysis of

mycotoxins. This method can enrich the target analytes from different matrices. However, the drawback of SPE is time-consuming, complicated, and unsuitable for multiclass analysis (Xu et al. 2015; Arroyo-Manzanares et al. 2015). To overcome the limitations of the solid phase extraction, tremendous efforts have been made to improve this method. Recently, magnetic solid phase extraction (MSPE) has been developed to effectively separate mycotoxins (Hashemi et al. 2014; Cassandra et al. 2014). More recently, Zhao et al. have reported an analytical method based on magnetic solid phase extraction (MSPE) (Zhao et al. 2017). In their study, $\text{Fe}_3\text{O}_4@\text{nSiO}_2@\text{mSiO}_2$, a magnetic nanoparticle, was prepared and dispersed into the sample solution to extract mycotoxin, and the absorbent together with the mycotoxin can be conveniently separated from the solution when applying an external magnetic field. Under the optimized MSPE conditions, the limits of detection were obtained for mycotoxins at 0.210 $\mu\text{g}/\text{kg}$ for FB1, 0.0800 $\mu\text{g}/\text{kg}$ for OTA, and 1.03 $\mu\text{g}/\text{kg}$ for ZEA, respectively. The proposed MSPE procedure serves not only for sample cleanup but also for mycotoxin enrichment that greatly enhances the assay sensitivity.

Efforts have been made to improve this method by Du et al. (2018). In their research, microwave-assisted dispersive micro-solid phase extraction has been applied for the extraction to detect multiple mycotoxins in different matrices. The optimum sample extraction conditions include 5 mL of sample solution dispersant with 2.5 $\mu\text{g}/\text{mL}$ of nano zirconia. When shaking vigorously for 2 min, the obtained analytes were desorbed by 100 μL of chloroform. The optimized method for determination of multi-mycotoxins by MA-d- μ -SPE coupled with UHPLC-Q-TOF/MS showed the lower limits of detection (0.0036–0.033 $\mu\text{g}/\text{kg}$ for solid samples).

2.2.2 Quick, Easy, Cheap, Effective, Rugged, and Safe Method (QuEChERS)

In order to minimize the sample pretreatment but prevent more exposure to matrix effects, a quick, easy, cheap, effective, rugged, and safe (QuEChERS) method is a suitable alternative. It is an effective simple method applied for extracting mycotoxin from a different food, both in solid samples and in liquid samples (Miró-Abella et al. 2017; Arroyo-Manzanares et al. 2015; Zhou et al. 2018). Its extraction method was based on partitioning via salting-out between aqueous and organic layer, and its cleanup was based on dispersive SPE (dSPE) by combining MgSO_4 and different sorbents (i.e., C18, primary, secondary amine, etc.).

Miró-Abella et al. have developed for the simultaneous determination of 11 mycotoxins in plant-based beverage matrices, using a QuEChERS extraction (Miró-Abella et al. 2017). In their method, 10 mL of sample was added to a 50 mL centrifuge tube mixed with 10 mL ACN with 1% HCOOH and shaken for 3 min. It was then 4 g of MgSO_4 and 1 g of NaCl were added into the solution and shaken vigorously for 3 min. Method quantification limits were between 0.05 $\mu\text{g}/\text{L}$ (for aflatoxin G₁ and aflatoxin B₁) and 15 $\mu\text{g}/\text{L}$ (for deoxynivalenol and fumonisin B₂).

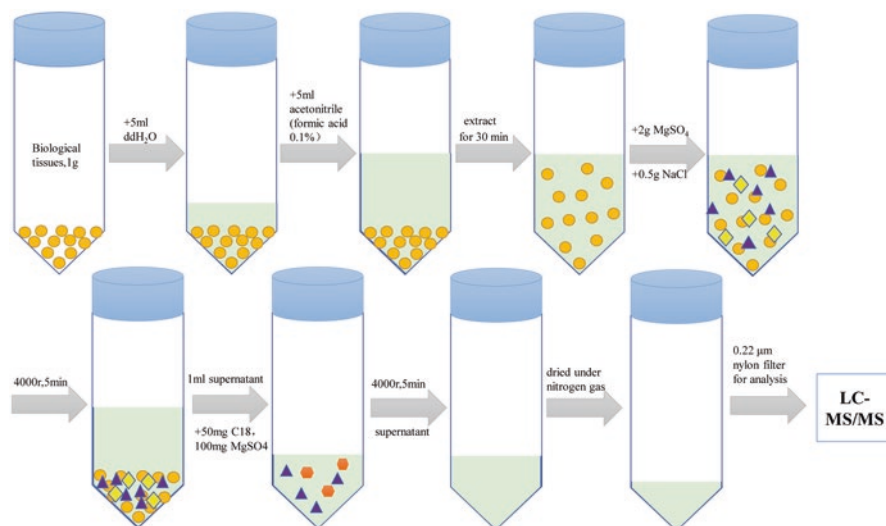


Fig. 2.1 Schematic diagrams of quick, easy, cheap, effective, rugged, and safe (QuEChERS) method for extracting mycotoxins

In 2018, Zheng Yan et al. developed and validated a liquid chromatography-tandem mass spectrometry (LC-MS/MS) method for the simultaneous quantification of ZEN family mycotoxins including ZEN and most relevant key metabolites in different tissues (Wang et al. 2018). In this study, the QuEChERS method was used to extract the analytes, and the procedure was shown at Fig. 2.1.

2.3 Major Techniques for the Determination of Mycotoxins

Accurate quantitation of mycotoxins at trace levels in real samples is comprehensively important for subsequent studies on confirmation legal judgments and exposure assessment. Therefore, it is very extremely important to improve and guarantee the performance of the applied analytical methodologies. To date, numerous methods and quantitative techniques for mycotoxins have been well established to investigate the occurrence of multiple mycotoxins in different matrices, to evaluate their toxicokinetics and toxicological dynamics in cell models and experimental animals. Previously, enzyme-linked immunosorbent assay (ELISA), high-performance liquid chromatography (HPLC), gas chromatography (GC), and thin-layer chromatography (TLC) are commonly used for quantitation of mycotoxins (Gutzwiller et al. 2014; Tamura et al. 2015; Rodríguez-Carrasco et al. 2014). These analytical methods are sensitive and selective but unsuitable for on-site monitoring because of bulky instrumentation in lab. In addition, these methods are relatively expensive and require well-trained, skilled persons for the operation and laborious sample preparation and cleanup procedures. At the same time, due to the case that several groups

of mycotoxins are frequently co-occurring even in the same sample, effective and reliable analytical methods are strictly required to detect multicomponent mycotoxins. Recently, with the fast developments in mass spectrometry (MS) technology, it enables to have the higher selectivity and sensitivity and provided the capacity to distinguish multiple contaminants from co-extractives in one individual sample. As a tendency, instrumental analysis on quantification of different mycotoxins requires the minimal sample loading volume via the utmost simplified green procedure, without much chemical solutions involved. Universally, the triple-quadrupole tandem MS (MS/MS) is considered the gold standard for quantitative analysis of multiple compounds (Koesukwiwat et al. 2014). Therefore, LC-MS/MS has been widely developed and employed for the analysis of single mycotoxin or group of mycotoxins. Previous studies demonstrated that LC-MS/MS is capable of simultaneously detecting multiple mycotoxins in different feedstuffs and foods, such as cereals (wheat, maize and rice), wheat-derived products, animal-borne feed, milk, and plant-based beverage (Oliveira et al. 2017; Suzane et al. 2017; Zhao et al. 2015; Jiang et al. 2018).

2.3.1 Cereals and Grains

Cereals are main staple foods in diets worldwide, which are highly valued as a source of nutrients, vitamins, minerals, and complex carbohydrates. One of the major challenges of the quality and safety of cereals is the natural contamination by mycotoxin-producing fungi post invasion into the host plants. This contamination can feasibly occur in the crops and worsen during harvesting, transportation, drying, processing, and/or further developing during storage of cereals and its derived products (Irena Kralj and Helena 2009). *Fusarium* head blight (FHB) mainly caused by *Fusarium graminearum* spp. was the common fungal disease in wheat and cereals (Prandini et al. 2009). The most common mycotoxins causing FHB were considered to be fumonisins, zearalenone, and deoxynivalenol (Krska et al. 2003; Ramirez et al. 2004). In addition to the parental mycotoxin forms, some food could also contaminate derivative/transformed compounds during food processing or as results of plant/animal metabolisms (Rychlik et al. 2014). Thus from now, it should be possible to suggest that accurate and reliable methods for the determination of mycotoxins and their modified forms in cereals are totally required in order to ensure the quality of these products and at most preserve consumer health.

Different analytical techniques have been proposed to monitor the large range of mycotoxins in cereals. Table 2.1 has summarized recent studies of method for determination of multi-mycotoxins in cereals and maize by LC-MS/MS. A multi-component method to determine aflatoxins, deoxynivalenol, fumonisins, zearalenone, ochratoxin A, and some metabolites in cereals was developed by Andrade et al. (2017). In this study, acidified ACN used as the extraction solvent followed by LC-MS/MS analysis proved to be suitable, rapid, and cost-effective for the multi-mycotoxin determination in wheat, maize, and rice products. This method

Table 2.1 Representative studies on analytical methods and limit of quantification (LOQ) in cereals and grains

Matrix	Limit of quantification ($\mu\text{g}/\text{kg}$)													Detection method	References	
	AFB ₁	AFB ₂	AFG ₁	AFG ₂	DON	15-AcDON	3-AcDON	D3G	FB ₁	FB ₂	FB ₃	OTA	ZEN			Sample preparation
Maize	1.2	0.7	0.7	0.5	39.0	121.0	77.0	84.0	19.0	8.0	32.0	4	24	SLE-LTP	LC-MS/MS	Andrade et al. (2017)
Grain	0.3	4.0	1.0	3.8	18.4				14.7	16.0	1.1	0.9		IAC	LC-MS/MS	Rodríguez-Carrasco et al. (2016)
Maize									12.5	12.5	12.5			SPE	LC-MS/MS	Wang and Li (2015)
Wheat					125	125	125		63	63	63		125	SPE	LC-MS/MS	Blesa et al. (2014)
Sorghum	5	5	10	10	25	62.5	31		25	50	50	2.5	6	SPE	LC-MS/MS	Ediage et al. (2015)
Wheat					3.9		11.1	6.6					5.7		LC-MS/MS	Berthiller et al. (2005)
Rice	0.5	1.2	1.0	1.6	40	72	48	84	21	12	24	3	16	SLE-LTP	LC-MS/MS	Andrade et al. (2017)
Cereal					75.2				2.89	2.36	0.65		38.6	QuEChERS	LC-MS/MS	Arroyo-Manzanares et al. (2015)

was satisfactorily validated for the simultaneous quantitation of multi-mycotoxins, with LOQs ranging from 0.5 to 121 $\mu\text{g}/\text{kg}$ listed in Table 2.1. Natalia Arroyo-Manzanares et al. proposed an UHPLC-MS/MS method combined with a sample treatment based on QuEChERS (4 g MgSO_4 , 1 g NaCl, 1 g sodium citrate and 0.5 g disodium hydrogen citrate sesquihydrate) for the determination of multi-mycotoxins in cereals (Arroyo-Manzanares et al. 2015). The analytical method for detecting fusarenon-X, T-2 and HT-2 toxin, and citrinin was also developed in this study with low LODs and LOQs. A UPLC-ESI-MS/MS method for the simultaneous determination of multi-mycotoxins including zearalenone, aflatoxins, and deoxynivalenol in feedstuffs was developed by David Romera et al. (2018). In their study, acetonitrile-water-formic acid (80:19:1, v/v/v) was carried out to extract target and nontarget mycotoxins in feedstuff and then detected by combining QTOF-MS and MS/MS detectors with UPLC. In 2019, E.J. Llorent-Martínez et al. developed an alternative flow-through luminescence optosensor for the quantification of ZEA (Llorent-Martínez et al. 2019). With this purpose, a QuEChERS procedure was carried out for the extraction of ZEA from different cereal samples and feedstuff materials. Solid-phase spectroscopy (SPS) in a multi-commutated system was implemented by making use of three-way solenoid valves, which provides an enhancement in the sensitivity and selectivity of the method. An ultra-performance liquid chromatography-electrospray tandem mass spectrometry (UPLC-ESI+/-MS/MS) method for the simultaneous analysis of citrinin (CIT) and ochratoxin A (OTA) in feed was first developed by Celine Meerpoel et al. (2018). In their study, the involved mycotoxins were extracted from these matrices using a QuEChERS-based extraction method without any further cleanup step. Furthermore, they detected citrinin (CIT) and ochratoxin A (OTA) in the selected matrices including pig feed and wheat flour with low LOQs. A study proposed a liquid chromatography-tandem mass spectrometry method using the selected reaction monitoring mode for determining fumonisins B1 and B2 (Oliveira et al. 2017). They modified the extraction procedure based on a matrix solid-phase dispersion approach, using 1 g of silica gel for dispersion and elution with 70% of ammonium formate aqueous buffer (50 mmol/L, pH 9), demonstrating a simple, cheap, and chemically friendly sample preparation procedure.

2.3.2 Milk and Beverage

Milk is an important source that provides nutrition for the growth, development, and maintenance of human health, particularly to younger children and infants. For an instance, breast milk is considered to be the ideal food for infants since it provides the best original health nutrition for their development and growth. However, milk can also be a carrier of mycotoxins and their metabolites, causing various physiological problems in individuals. Additionally, mycotoxins present in milk could transform into the blood and tissues of breeding mammals and thus brought to the forefront for milk safety (Degen et al. 2017). The most studied mycotoxins in milk

are aflatoxins and ochratoxin A and to a lesser extent, trichothecenes and zearalenone (Flores-Flores and González-Peñas 2015; Huang et al. 2014). Previous studies have shown considerable amount of AFM1 in milk and dairy products. As confirmed, aflatoxin M1 is produced by lactating animals and mothers when the cows consumed the feeds or foods contaminated with aflatoxin B1 (AFB1) through cytochrome P450-associated enzymes in the liver. Then, AFM1 was secreted through urine and milk (Dashti et al. 2009). In the last decades, several validated analytical methods have been developed for multi-mycotoxin determination in milk, which allow studying the presence of mycotoxins in milk. Table 2.2 has listed recent studies of method for determination of multi-mycotoxins in milk and milk product by LC-MS/MS. A reliable and universal QuEChERS-based UPLC-MS/MS method was newly developed for the determination of multi-mycotoxins in milk (Zhou et al. 2018). As for the choice of the mobile elution phase, MeOH and ACN were considered as candidates. The results showed that compared with MeOH, ACN provided sharper peak profiles and facilitated the elution of analytes.

Since milk samples always present complex matrix with multiple components, complex interaction should exist among the detected mycotoxins and matrix, as well as other functional nutrients. Herein, several methods have been proposed to alleviate matrix effects. For example, an m- μ dSPE method using Fe₃O₄@pDA NPs has been applied for the extraction of six mycotoxins before their LC-MS analysis from different milk and yogurt samples in the study of Javier González-Sálamo et al. (2017). Moreover, a highly sensitive and specific multiclass method by combining multi-mycotoxin IAC enrichment procedure and UHPLC-Q-Orbitrap was developed and validated to be capable of quantifying 14 mycotoxins in raw milk (Mao et al. 2017). Interestingly, multi-mycotoxin IAC cleanup utilized there could produce the extract-alleviating matrix effects, while the addition of acetonitrile in milk was found to be beneficial for the extraction. As proved, the developed method has been successfully applied to routine analyses for quantitation of mycotoxins in raw milk.

2.3.3 Fruits and Vegetables

Fruits and vegetables are highly susceptible to be infested by pathogenic fungi due to their high water content and abundance of nutrients. Among all fruits, berry fruit (i.e., small berries such as strawberries, blueberries, and red currants) is especially susceptible to be contaminated by fungi considering that their soft and fragile skin is susceptible to small lesions, which allow the growth of spoilage fungi. They may decompose during any processing stage of growth, pre- and post harvests, as well as storage and transportation. Therefore, several classes of mycotoxins can simultaneously contaminate fruits and vegetable (Sanzani et al. 2016). Mycotoxins associated with fruits and vegetables mainly include *Alternaria* toxins, ochratoxin A (OTA), patulin (PAT), and trichothecenes. Although consumers are easily tend to cut out the visible rotten areas of the infected fruits and vegetables before consumption, several

Table 2.2 Representative studies on analytical methods and limit of quantification (LOQ) in milk product

Matrix	Limit of quantification ($\mu\text{g}/\text{kg}$)										Sample preparation	Detection method	References
	AFM ₁	AFM ₂	AFG ₁	AFG ₂	AFB ₁	AFB ₂	ZEN	ZAN	OTA	OTB			
Cow milk							2.15	1.57			m- μdSPE	LC-MS	González-Sálamo et al. (2017)
Liquid milk	1					0.1				2	SPE	LC-MS/MS	Wang and Li (2015)
Milk powder	5					0.5				10	SPE	LC-MS/MS	Wang and Li (2015)
Milk	0.01	0.02	0.01	0.02	0.01	0.01	0.2	0.1	0.01	0.01	QuEChERS	UPLC-MS/MS	Zhou et al. (2018)
Milk	0.001	0.001	0.003	0.003	0.003	0.003	0.025	0.01	0.001	0.001	IAC	LC/Q-Orbitrap	Mao et al. (2017)
Milk	0.02					0.03	0.04	0.15	0.02		SPE	LC-MS/MS	Jiang et al. (2018)
Milk		0.1	0.02	0.02	0.02	0.02	1		0.1	0.05	MSPE	LC-MS/MS	Zhao et al. (2015)

mycotoxins, especially those mentioned above, could still be transferred into the remaining parts indicating that mycotoxins could be present in fresh fruit and healthy processed products with low concentrations (Monbaliu et al. 2010). The potential adverse effects on human health could be caused by the presence of mycotoxins, considering the concurrent intake of different contaminated food and drinks by the population. Thus, sensitive, robust, and accurate analytical methods for specific analysis of multiple mycotoxins in fruits and vegetables are indeed necessary. Recently, high-performance liquid chromatography-tandem mass spectrometry (HPLC-MS/MS) has been increasingly accepted for mycotoxin analyses owing to its high selection, sensitivity, and anti-interference ability. Table 2.3 has summarized several studies of method for determination of multi-mycotoxins in fruits and vegetables by LC-MS/MS. A rapid and accurate analytical method for the determination of three *Alternaria* mycotoxins in fruit and juice was developed and validated by Charalampos K. Myresiotis et al. (Myresiotis et al. 2015). The QuEChERS extraction technique coupled with HPLC-DAD was used for the simultaneous determination of three *Alternaria* mycotoxins. Hao Dong et al. have developed a rapid and robust UHPLC-MS/MS method for the determination of AOH, AME, TEA, TEN, DON, PAT, and OTA in fresh fruit and vegetable (Dong et al. 2019). QuEChERS technique was finally applied for effective purification, and extraction was performed with acetonitrile under acidic conditions. In this method, a C18 column by gradient elution was applied to separate mycotoxins. Mass spectrometry analysis was scanned by ESI+ and ESI− dynamic switching using multiple reaction monitoring (MRM). In 2016, Meng Wang et al. have developed a highly sensitive, rapid, and reliable method for the simultaneous determination of eight mycotoxins in fruits by ultra-high-performance liquid chromatography-tandem mass spectrometry based on homemade “MCX + NH₂” SPE cleanup (Wang et al. 2016). Before SPE cleanup, acidified aqueous acetonitrile and an additional salt-out step using NaCl were employed. In their study, ESI positive mode was selected to analyze four mycotoxins including TEA, ALT, OTA, and CIT, and ESI negative mode was selected to analyze the other four mycotoxins including AOH, AME, TEN, and PAT. In addition, the CORTECS C18 column was carried out in this method. In 2018, Sara De Berardisa et al. have developed a liquid chromatography-tandem mass spectrometry method for the determination of four *Alternaria* toxins, i.e., AOH, AME, TEN, and TEA in tomato-based and fruit-based products (De et al. 2018). The use of an organic solvent (acetonitrile) as an extraction solvent was particularly appropriate for TeA extraction in this method. QuEChERS extraction method was performed and liquid chromatography coupled to tandem mass spectrometry (LC-MS/MS) was carried out in the study of Juan et al. (2017b). The simultaneous quantification of OTA and *Alternaria* toxins in berry by-products, jam and juice, was successfully proposed and validated, with good sensitivity to detect these mycotoxins.

Table 2.3 Representative studies on analytical methods and limit of quantification (LOQ) in fruit and vegetable

Matrix	Limit of quantification ($\mu\text{g}/\text{kg}$)										Detection method	References
	AOH	AME	TEA	TEN	DON	PAT	OTA	ALT	CIT	Sample preparation		
Tomato	3.5	3.5		1.75						LLE	LC-MS/MS	Rodríguez-Carrasco et al. (2016)
Sweet cherries	1.0	1.0	1.0	1.0		5.0	1.0	1.0	1.0	SPE	UPLC-MS/MS	Wang et al. (2016)
Apples	1.0	1.0	1.0	1.0		5.0	1.0	1.0	1.0	SPE	UPLC-MS/MS	Wang et al. (2016)
Tomato	1.0	1.0	1.0	1.0		5.0	1.0	2.0	1.0	SPE	UPLC-MS/MS	Wang et al. (2016)
Orange	1.0	1.0	1.0	1.0		5.0	1.0	1.0	2.0	SPE	UPLC-MS/MS	Wang et al. (2016)
Tomato	16	20	4	160						QuEChERS	LC-MS/MS	De et al. (2018)
Jam berries	7	15		5			3.5			QuEChERS	LC-MS/MS	Juan et al. (2017b)
Juice berries	5	9		2			5			QuEChERS	LC-MS/MS	Juan et al. (2017b)
Vegetable	1.0	0.2	1.0	0.4	1.5	10.0	5.0			QuEChERS	LC-MS/MS	Dong et al. (2019)

2.3.4 *Biological Samples*

In general, external exposure assessment with the scope of risk assessment is used to food contamination and consumption data to estimate human exposure to different mycotoxins for different human populations such as infants and children. However, food and feedstuffs cannot accurately estimate the real disease risk to the human beings. Therefore, determination of mycotoxins and its derivatives in biological samples such as plasma and urine can monitor the mycotoxin levels directly and promote effective exposure assessment, which is crucial for the establishment of regulatory limits. Several findings of direct biomonitoring showed that a higher prevalence of some mycotoxins was observed (Gerding et al. 2015; Huybrechts et al. 2015). These studies clearly emphasized the importance of direct biomonitoring of mycotoxins in biological samples.

For example, a liquid chromatography-tandem mass spectrometry method for multi-mycotoxins in fish plasma was developed (Tolosa et al. 2016). However, LOQs of this method ranged from 1 to 17 ng/ml and are not suitable for the determination of mycotoxins in human plasma. Irina et al. developed a sensitive liquid chromatography-mass spectrometry method for 17 mycotoxins in human plasma (Slobodchikova and Vuckovic 2018). In their study, they studied the effect of a different method of sample preparation (solvent precipitation, SPE, and LLE) and LC separation on simultaneous analysis of mycotoxins. Finally, in order to eliminate the need for immunoaffinity extraction and minimize matrix effects, they developed the method based on three-step liquid-liquid extraction with ethyl acetate. In this method, 15 of these mycotoxins met the accuracy and precision of 80–120% and $\leq 20\%$ RSD at different concentrations tested. The obtained LOQs of all mycotoxins were in the range of 0.1–0.5 ng/ml except NIV (3 ng/ml).

2.4 Conclusion

Till now, some effective and reliable analytical methods have been well established to determine mycotoxins contaminated in food and feedstuff at legislated levels. For more emphasis, proficiency testing (PT) is an effective procedure for quality assurance and performance verification in different laboratories from different countries at the global level, ensuring that laboratory validation and inter- or intra-laboratory procedures are working satisfactorily (Girolamo et al. 2016). To be specific, the maize samples contaminated with multiple mycotoxins, including DON, FB1, FB2, ZEA, T-2, HT-2, OTA, AFB1, AFG1, AFB2, and AFG2, and the wheat samples contaminated with DON, ZEA, T-2, HT-2, and OTA were selected as real materials. As resulted, the majority of laboratory participants had the ability to provide acceptable results for the simultaneous analysis of DON, fFB1 and B2, OTA, ZEN, T-2 and HT-2, and AFB1 and G1 in maize and DON, OTA, ZEN, and T-2 and HT-2 in wheat, respectively. In addition, acidified ACN-water as extraction dilution and

injection without cleanup extract were of the best performance. However, only a few laboratories were capable to analyze AFB₂ and AFG₂ and FB₁ together with other mycotoxins in maize and T-2 in wheat. More importantly, some modified or derived forms of different mycotoxins were reported and proved to have identical toxicities to the parental molecule, such as D3G and 3- and 15-ADON, respectively. To be specific, via the currently used LC-MS/MS method, it is too tough for absolute separation and identification of 3- and 15-ADON. Also, it is very difficult to obtain the standards of the masked forms of the key mycotoxins (D3G, Z14G, and Z14 S), which have aroused as great challenges regarding the typed chromatographic analyses.

In one word, further improvements on sample preparations and optimization of equipment parameters of LC-MS/MS methodologies to gain higher sensitivity, fewer matrix effects, and more targets involved are still required for ongoing multi-mycotoxin analysis.

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