

## **COMMITTEE ON TOXICITY OF CHEMICALS IN FOOD, CONSUMER PRODUCTS AND THE ENVIRONMENT (COT)**

### **First draft statement on the potential risks of combined exposure to mycotoxins**

#### **Background**

1. The potential risks from combined exposure to mycotoxins was identified as a topic that the COT should consider during horizon scanning.
2. A preliminary scoping paper regarding the potential risks from combined dietary exposure to mycotoxins (TOX/2020/34)<sup>1</sup> was reviewed by the COT in July 2020. Following discussions, the Committee requested for the Secretariat to collate additional data regarding the availability of biomonitoring data for multiple mycotoxin exposures specific to the United Kingdom (UK) population and the inclusion of the toxic mode of action for the reviewed mycotoxins (as those in originally included in TOX/2017/30)<sup>2</sup>.  
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3. The additional data was presented in September 2020 (TOX/2020/44)<sup>3</sup>. It was discussed that testing platforms for the detection of multiple mycotoxins in food matrixes are still in development, as such no new data will be available in the near future. It was concluded that there is a lack of UK data on biomonitoring and co-occurrence of mycotoxins in food. Therefore, there was limited understanding of actual co-exposure to dietary mycotoxins.
4. The draft statement as presented in Annex A brings together the conclusions from these discussions and lists the research recommended by the COT.

#### **Questions on which the views of the Committee are sought**

5. Members are invited to consider the following questions and to raise any other matters that arise from the newly submitted data:
  - i). Are there any aspects that have been addressed during the COT review of combined exposure to mycotoxins that are not covered in the draft statement and which should be included?

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<sup>1</sup> TOX/2020/34 is available on the [COT website](#).

<sup>2</sup> TOX/2017/30 is available on the [COT website](#).

<sup>3</sup> TOX/2020/44 is available on the COT website consisting of the [cover page](#), [Annex A](#) and [Annex B](#).

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- ii). Do the conclusions accurately represent the views of the COT?
- iii). Does the Committee have any other comments on the structure and content of the draft statement?

**Secretariat**  
**October 2020**

## COMMITTEE ON TOXICITY OF CHEMICALS IN FOOD, CONSUMER PRODUCTS AND THE ENVIRONMENT (COT)

### First draft statement on the potential risks of combined exposure to mycotoxins

#### Background

1. The COT horizon scanning identified consideration of potential risks from combined exposure to mycotoxins as a priority.
2. Climate change could have a significant impact on the life cycle stages<sup>4</sup> and rates of the development of toxicogenic fungi which has the ability to modify host-resistance and host-pathogen interactions. In turn, this will influence the conditions for mycotoxin production that varies for each individual pathogen species and/or strain.
3. Furthermore, advances on analytical techniques have driven the simultaneous detection and quantification of multiple mycotoxins in both food and feed commodities (Krska *et al.*, 2007; De Santis *et al.*, 2017; Flores-Flores & González-Peñas, 2017; Bessaire *et al.*, 2019; Singh & Mehta, 2020; Agriopoulou *et al.*, 2020).
4. In light of this, new combinations of factors (mycotoxins/host plants and geographical location) will have to be considered when assessing the potential risks of combined exposure to mycotoxins.

#### Introduction

##### *Definitions*

5. Mycotoxins are toxic secondary metabolites produced by fungi and can cause adverse health effects in both humans and animals. Cereals are often the most severely affected crops; however, some nuts, fruits and spices can also be affected. Mycotoxins are stable low-molecular weight chemicals and are not often affected by food processing.
6. Mycotoxins of greatest concern to human health and livestock are produced by several fungal genera of filamentous fungi, namely *Aspergillus*, *Fusarium* and *Penicillium*, which produce aflatoxins (AFs), ochratoxin A

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<sup>4</sup> There are four main stages for the fungi life cycle; hyphal growth, spore formation, spore dispersal and spore germination.

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(OTA), patulin (PAT), fumonisins (FBs), zearalenone (ZEN), nivalenol (NIV), deoxynivalenol (DON), citrinin (CIT), T-2 and HT-2 toxins (WHO, 2018).

7. Exposure to dietary mycotoxins can lead to several adverse effects in humans, which include carcinogenic, teratogenic, hepatotoxic, nephrotoxic, and cytotoxic, immunological and haematological effects.

8. In addition to native mycotoxins, modified mycotoxins can also be produced by fungi or generated as part of the defence mechanism of the infected plant. They are described as metabolites that normally remain undetected during the testing for parent mycotoxins. It has been reported that some modified mycotoxins can be converted into the parent mycotoxin during digestion in humans and animals, and thus has the potential to lead to adverse health effects. Although, the toxicological data for this is scarce (Freire & Sant'Ana, 2018).

### **Prevalence and co-occurrence**

9. DON, FBs, and ZEN are the most prevalent mycotoxins in the world, with a prevalence of 66%, 56%, and 53%, respectively in cereals and cereal based products (Smith, 2016).

10. The production of mycotoxins can occur pre-, during or post-harvest. Several factors can influence the production of mycotoxins pre-harvest, such as the sowing time, plant density, soil conditions, irrigation, presence of weed and pest. During harvest, the influencing factors include drying, cleaning and sorting of the crops, whereas the post-harvest factors are mainly associated with storage and processing. Mycotoxin colonisation also depends on temperature, relative humidity, rainfall and water activity<sup>5</sup>. The presence of the fungi spores in crops does not always result in the production of mycotoxins, since optimal growth conditions are required for biosynthesis (Battilani *et al.*, 2020).

11. The natural co-occurrence of mycotoxins in food and feed is quite common and occurs for three main reasons; (i) some fungi can produce more than one mycotoxin (particularly *Fusarium* spp.), (ii) food commodities can be contaminated by several fungi and (iii) animal and human diets usually consist of multiple commodities.

12. At present, there is limited availability on data with regards to co-occurrence of mycotoxins in food commodities especially in Europe. In contrast, worldwide surveys for mycotoxins in feed are more common of which, the BIOMIN Mycotoxin Survey<sup>6</sup> is one of the most comprehensive.

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<sup>5</sup> Water activity is a measure of the availability in a substrate of water for microbial growth.

<sup>6</sup> The BIOMIN Mycotoxin Survey constitutes the longest running and most comprehensive survey of its kind. The survey results provide insights on the incidence of the six major mycotoxins (AFs, ZEN, DON, FBs, T-2 and OTA) in the agricultural commodities used for livestock feed in order to identify the potential risk posed to livestock animal production. Further information available on the [BIOMIN website](#).

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13. Updated results for January – June 2020 on the occurrence of mycotoxins in ~9,600 finished feed and raw commodity samples from 68 countries based on ~42,500 analyses were published in August 2020. It was observed that mycotoxins are rarely found alone; multiple mycotoxins co-contaminated feed materials (68% of all samples analysed for at least two mycotoxins). Results specific for Northern Europe shows high contamination of DON at an average of 531 ppb in straw (BIOMIN, 2020a).

14. Table 1 provides a mycotoxin prevalence breakdown for Central, Eastern, Northern and Southern Europe.

Table 1 - the BIOMIN Mycotoxin Survey Central, Eastern, Northern and Southern Europe results on prevalence of mycotoxins in animal feed (%) for January to March 2020 (reproduced from BIOMIN, 2020b).

<b>Mycotoxin</b>	<b>AF</b>	<b>ZEN</b>	<b>DON</b>	<b>T-2</b>	<b>FBs</b>	<b>OTA</b>
<b>Central Europe</b>	5	60	78	27	61	8
<b>Eastern Europe</b>	3	42	37	45	36	31
<b>Northern Europe</b>	0	31	59	20	21	5
<b>Southern Europe</b>	9	49	60	15	93	13

Abbreviations: AF = Aflatoxins; ZEN = Zearalenone; DON = Deoxynivalenol; T-2 = T-2 toxin; FBs = Fumonisin; OTA = Ochratoxin-A.

15. Some publications note that there is still limited knowledge on the presence and co-occurrence of multiple mycotoxins, both for native mycotoxins and their modified forms, in food and feed (Palumbo *et al.*, 2020), and that it is difficult to infer trends or recent developments regarding mycotoxin contamination in European feed from available data due to the influence of the respective cropping season's climate on the contamination levels which caused high year to year variation of results, as well as the differences in the applied analytical methods used to detect the contamination (Streit *et al.*, 2012).

### **Methods for sampling and measuring mixtures of mycotoxins in food matrices**

16. The main analytical methods to measure mycotoxins are enzyme-linked immunosorbent assay (ELISA), gas chromatography-mass spectrometry (GC-MS) and liquid chromatography-mass spectrometry (LC-MS). Currently, LC based methods are the most frequently used (Serrano *et al.*, 2012; Malachová *et al.*, 2018), with several mass spectrometric detectors such as fluorescence detector, single-quadrupole mass spectrometer, time-of-flight, triple-quadrupole, ion trap and orbital ion trap mass analysers, as well as hybrid systems that combine two types of analysers.

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17. An important and critical step in the analytical process is sample preparation and clean-up with techniques including solid phase extraction, matrix solid-phase dispersion, liquid–liquid and solid–liquid partitioning, accelerated solvent extraction, multifunctional columns and immunoaffinity columns (Serrano et al., 2012; Agriopoulou *et al.*, 2020).

18. The main analytical method used for detecting and measuring co-occurrence of very low concentrations of mycotoxins is liquid chromatography tandem mass spectrometry (LC–MS/MS) (Serrano *et al.*, 2012; Gambacorta *et al.*, 2018; Malachová *et al.*, 2018; Shi *et al.*, 2019; Battliani *et al.*, 2020; Palumbo *et al.*, 2020;) however, multi-mycotoxins analyses are not widely performed due to their associated high cost.

### **Current state of authoritative assessment and research**

19. The COT have previously reviewed the European Food Safety Authority (EFSA) external report on mycotoxin mixtures in food and feed (Battiliani *et al.*, 2020), several opinions by authoritative groups on some mycotoxin combinations, the work produced by the Mycotoxin mixtures (MYCOMIX) project led by Paula Alvito and her colleagues, as well as being aware of the Mycotoxin and Toxigenic Moulds (MYTOX) research group co-ordinated by Sarah De Saeger; as presented in TOX/2020/34<sup>7</sup>. Brief summaries are provided below.

#### *European Food Safety Authority (EFSA) external scientific report on mycotoxin mixtures*

20. In this EFSA report titled “*Mycotoxin Mixtures in Food and Feed: Holistic, Innovative, Flexible Risk Assessment Modelling Approach*” (MYCHIF) Battiliani *et al.*, (2020) performed an extensive literature review out across four topics relating to the investigation of mycotoxin mixtures present in food and feed. These topics were:

- i). Ecology and interaction with host plants of mycotoxin producing fungi, mycotoxin production, recent developments in mitigation actions of mycotoxins in crop chains;
- ii). Analytical methods for native, modified and co-occurring mycotoxins;
- iii). Toxicity, toxicokinetics, toxicodynamics and biomarkers relevant to humans and animals and;
- iv). Modelling approaches, and key reference values for exposure, hazard and risk modelling.

21. The data collected from these were then stored in the MYCHIF platform hosted by EFSA. The main objective of which was to develop an integrated method supported by modelling, for the risk assessment of

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mycotoxin mixtures in food and feed. Each topic will be summarised in the following paragraphs.

22. It was observed that *Aspergillus* spp., *Fusarium* spp., and *Penicillium* spp., were the most relevant mycotoxins worldwide, this also extends to *Alternaria* spp. and *Claviceps* spp. to a minor extent. The production of mycotoxigenic fungi are not commonly host specific, since their occurrence is mainly associated with a specific crop depending on its region of growth and meteorological conditions. A fungus can produce different types of mycotoxins (e.g. sterigmatocystin which is a precursor for AFs), as such contamination of food and feed stuffs can occur concurrently. Modified mycotoxins may also co-occur with native varieties as a result of fungi-host plant interaction or during processing.

23. The MYCHIF report deemed that temperature, relative humidity, rainfall, are the most important ecological factors that influence fungal colonisation of substrates. Additionally, each species would have its own ecological needs and requirements. In general, mycotoxins are stable compounds and can accumulate over time (both during crop growth and post-harvest). Therefore, mitigation of contamination requires both good practices at all production stages.

24. In terms of methodologies used for mycotoxin analysis, these are split into two categories. Firstly, screening tests provide qualitative or semi-quantitative results. They are generally based on antibody recognition and these methods are often relatively straightforward to carry out. The other category is confirmatory analysis which provides confirmation of fungi species identity and quantitative results. The most widely used quantification method is High Performance Liquid Chromatography (HPLC). LC-MS is also used to identify and quantify mycotoxins. A number of high- or ultra-chromatography coupled to mass spectrometry systems have the ability to measure both regulated mycotoxins and other lesser tested for mycotoxins with analytical standards available (e.g. CIT, sterigmatocystin etc) together in different feed food commodities, however, there are some limitations in these systems including: cost, sensitivity to include lower limits of quantification and detection of *in vivo* metabolites, and a harmonised fit for purpose methodology characterised by the ability to measure multiple mycotoxins. Furthermore, what remains important is the meaningful interpretation of the mycotoxin mixture data to their related effects in human and animal health.

25. Toxicity, toxicokinetics and toxicodynamic parameters for humans and animals were collected from the literature to build three databases in the MYCHIF platform. These are: *in vitro* toxicokinetic data, *in vivo* toxicokinetic data, and *in vivo* toxicity data. A limited number of articles on mixtures were observed in comparison to those only exploring effects of single compounds. The information available only covers a limited number of combinations of mycotoxins. Toxicokinetic data were mainly reported in pigs and chickens, and rats. Toxicodynamic modelling of mycotoxin mixtures could not be performed using the currently available data.

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26. The mycotoxin dose, exposure pathway, interspecies and intraspecies differences were identified to be the most important parameters that may influence the toxicokinetic mixtures.

27. As part of the MYCHIF project, it was highlighted that testing of all mycotoxin mixture combinations is unfeasible, as such, focus should be given on the prioritisation of mycotoxin mixtures, the creation of harmonised methods for generating *in vitro* and if required additional *in vivo* toxicokinetic data, and utilising predictive kinetic modelling that includes uncertainty, and inter- and intraspecies variability analysis.

28. In terms of the exposure assessment, human biomonitoring data was collected from the literature; 66/176 articles that focused on biomarker studies of multi-mycotoxins were selected for further analysis. A multi-biomarker study was defined whereby both the parent and one or more metabolite was measured. Regarding biomonitoring in humans, AFs is the most widely studied mycotoxin followed by OTA, DON, FBs, ZEN and other emerging mycotoxins such as alternaria (ALT), tenuazonic acid, fusarenon-X (Fus-X,) neosolaniol, CIT, NIV, T-2, 4,15-diacetoxyscirpenol (4,15-DAS), and enniatins (ENNs) in a very few studies. The most common sample matrix was urine, followed by serum, plasma, blood, breast milk, colostrum and amniotic fluid.

29. The simultaneous determination of more than one mycotoxin in human biological fluids presents as a new challenge in mycotoxin biomonitoring. There are several constraints; in an analytical context, there is a lack of method standardisation and the unavailability of commercial reference standards (especially glucuronides). Their use in exposure assessments may be premature and cannot be fully exploited since there is a lack of: knowledge of effects of different combinations on the bioavailability of each individual compounds, the excretion rate, and a consensus of a validated biomarker to be used in context to a multi-mycotoxin analysis.

30. Overall, there is still a lack of harmonisation in the experimental settings (e.g. for the use and validation of analytical methods) and design of biomonitoring studies (e.g. the selection of a candidate biomarker), in the data collection and in the definition of performance of fit for purpose analytical methods. These highlighted issues made it not possible for Battilani *et al.*, (2020) to exploit the biomonitoring dataset for exposure assessment goals. They recommend that international study guidelines should be prepared to support the production of data.

31. A human case study was presented to risk assess two mycotoxin mixtures which can occur and co-occur in cereal based food products (DON, FBs and ZEN, and T2/HT-2 toxin, DON and NIV) using the component-based approach (CBA)<sup>8</sup> and provisional daily intake modelling methodologies (further detailed in the Exposure assessment section). A problem formulation, exposure assessment, hazard assessment and risk characterisation were

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<sup>8</sup> Component based approaches are used to estimate the hazard or risk of combined exposures based on information on exposure and hazard for each individual component.



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completed. A Margin of Exposure (MOE)<sup>9</sup> value of 100 was chosen as a reference/cut off since the considered mycotoxins were neither genotoxic nor carcinogenic. In brief, due to data gaps and limitations a robust risk assessment could not be performed, however, results from the CBA modelling approach calculated MOE values <100 for both mixtures, indicating that there is a need to refine the risk assessment or a potential health risk.

32. A hierarchy map (based on EU Member states) was made possible when considering exposure to T2/HT-2 toxin, DON and NIV for adults. The maps provided a visual representation of higher risk exposure groups. With reference to the co-occurrence and occurrence data collected for the United Kingdom (UK); T2/HT-2 toxin, DON and NIV adult exposure levels based on the mean co-occurrence data and mean consumption value were >0.8 ,0.6-1.2 and 0.24-0.3 µg/kg bw/day, respectively.

33. To put into context the above exposure values, their health-based guidance values are provided. The acute reference dose (ARfD) for T2/HT-2 toxin is 0.3 µg/kg bw (EFSA, 2017a). DON and its acetylated forms have an ARfD of 8 µg/kg bw or a tolerable daily intake (TDI) of 1 µg/kg bw (EFSA, 2017b). Whilst, NIV has a TDI of 1.2 µg/kg bw (EFSA, 2013).

34. To conclude, the following data gaps and research recommendations were observed and/or suggested. There is limited knowledge on the presence and co-occurrence of multiple mycotoxins, both for native mycotoxins and their modified forms, in food and feed. Available analytical methods have limitations for the routine monitoring of modified and multi-mycotoxins in food and feed. In the context of multi-mycotoxin analysis and the use of LC-MS; there remains an urgent need for the following: lower costs, fit for purpose methods characterised by the ability to measure multiple mycotoxins (with lower limits of quantification for all co-occurring mycotoxins; including their metabolites investigated *in vivo*), and availability of commercial reference materials for providing reliable quantitative results.

35. In terms of toxicity data, a limited number of articles on mixtures were observed in comparison to those only exploring effects of single compounds. The available studies only cover a very limited combination of mycotoxins and the available toxicokinetic data is mainly in livestock species (pigs and chickens), as well as rats. The modelling of toxicodynamic features of mycotoxin mixtures could not be performed based on the limited number of data available. The development of prioritisation criteria for mycotoxin mixtures to be tested was suggested as a further research priority. In addition to this, consistent methodologies and harmonised guidelines for generating *in vitro* and *in vivo* toxicokinetic (TK) and toxicodynamic (TD) data are needed to provide consistent data for pharmacologically based toxicokinetics and benchmark dose modelling of mycotoxin mixtures. Current analytical methods should have the capability to detect and analyse real world samples. Finally,

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<sup>9</sup> The margin of exposure (MOE) is the ratio of the point of departure (typically the benchmark dose – lower confidence limit for a tumorigenic response in experimental animals), to the estimated human exposure for a genotoxic carcinogen. MOE values that are ≥10,000 have been considered to indicate low concern.

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the utilisation of TK and TD modelling should be further considered and explored.

36. With reference to the use of biomarkers for exposure assessment purposes; there is a need to derive qualitative and quantitative correlations between the mycotoxin intake from food and from other possible routes of exposure like dermal or inhalation (which may be important to consider as part of occupational hazard assessments, however, it is unclear how realistic it would be to achieve this level of granularity from such a population based study).

#### *Opinions by authoritative groups on some mycotoxin combinations*

37. Assessments of some binary mycotoxin combinations have been carried out by EFSA, Joint Food and Agriculture Organization and World Health Organisation Expert Committee on Food Additives (JECFA) and the Scientific Committee on Food (SCF). These are summarised in the next following paragraphs.

#### EFSA reviews

38. During their review of 4,15-DAS in 2018, the EFSA Panel on Contaminants in the Food Chain (CONTAM) considered the combined effects and interactions of 4,15-DAS with T-2 and HT-2 toxins, AFs, OTA and FBs. Following the analysis of the available database describing possible effects of combined exposure to 4,15-DAS and other mycotoxins, the EFSA CONTAM Panel concluded that the data was weak and inconclusive (EFSA, 2018).

39. The EFSA CONTAM Panel also examined available publications addressing interactions of CIT with PAT, AFs and OTA, particularly on the subject of synergism in 2012. It was concluded by the EFSA CONTAM Panel that the available evidence indicated that CIT at low doses does not exacerbate the toxic effects of other mycotoxins and that the combined effect of CIT and OTA is, at most, additive (EFSA, 2012a).

#### JECFA reviews

40. The JECFA have previously considered the toxicology associated with concurrent exposure to FBs and other mycotoxin agents in 2011. The reviewed *in vitro* and *in vivo* studies were found to only involve single doses of each individual mycotoxins, and as such the JECFA Committee concluded that none of the studies were adequate for quantitative assessment of interactions (JECFA, 2011).

41. Although studies by Carlson *et al.*, (2001) and Gelderbloom *et al.*, (2002) were noted; these documented the ability of FB1 to promote AFB1 hepatocarcinogenicity in trout and orally dosed pure FB1 in rats induced precancerous lesions, respectively. The treatment regime and dose at which these effects were observed in trout was through exposure of 100 ppb of AFB1 and then exposure to  $\geq 23$  FB1 ppm for 42 weeks (Carlson *et al.*, 2001).

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Whilst in male Fisher rats (n=5-8/group) it was AFB1 at 17 µg/kg bw per day (via oral gavage) for 14 days, and exposure to FB1 at 250 mg/kg diet (post recovery period of 21 days in between) (Gelderbloom *et al.*, 2002). The potential interaction between DNA-reactive AFB1 and FB1, with its potential to induce regenerative proliferation was noted as a concern by the JECFA Committee (JECFA, 2011).

42. The topic was revisited in the JECFA 2018 evaluation (Riley *et al.*, 2018). Synergistic effects were reported by Qian *et al.*, (2016) for the development of preneoplastic lesions (e.g. increased number of apoptotic cells and placental form of glutathione-S-transferase), where male F344 rats (n=13) were exposed to pure AFB1 (equivalent to 15 µg/kg bw per day for 14 days) and pure FB1 (equivalent to 25 mg/kg bw per day for 21 days), alone or sequentially (the rats were treated with AFB1 and then FB1, with a recovery period of 21 days in between).

43. JECFA noted that, even though there are additive or synergistic effects observed from FB1 and AFB1 co-exposure in laboratory animals in inducing the development of preneoplastic lesions and hepatocellular carcinoma (as discussed above), there was currently no data available on such effects in humans. Furthermore, two prospective epidemiological studies (Magoha *et al.*, 2016; Shirima *et al.*, 2015), do not support the hypothesis of an interaction between AFB1 and FB1 in childhood stunting. JECFA concluded that there were few data available to support co-exposure as a contributory factor in human disease. However, the interaction between AFB1 (genotoxic), and FBs, which have the potential to induce regenerative cell proliferation (particularly at exposures above the provisional maximum tolerable daily intake), remained a concern.

44. It was recommended that exposures to both compounds should be reduced and that emphasis on human studies should be on biomarker-based approaches.

45. JECFA further assessed the combined toxicity of FBs and DON in their 2018 evaluation. They concluded that some of the effects from co-exposure were suggestive of being additive or more than additive, however, the effect is dependent on the endpoints measured (JECFA, 2018).

#### SCF review

46. The SCF have provided basic summary for the relative potency and dose additivity of trichothecenes in 2002 (SCF, 2002). Although different types of trichothecenes appear to cause similar effects (T-2 toxin, HT-2 toxin, deoxynivalenol and nivalenol) at the biochemical and cellular level and there are similarities in toxic effects, there are also considerable differences in the spectrum of toxic effects *in vivo*. Large, non-systematic potency differences between these toxins are seen when different endpoints are being considered. Additionally, there are only few studies that address the combined effects of these toxins.

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47. In *in vitro* studies, dose additivity as well as antagonism has been observed for T-2 toxin, DON and NIV, whilst in *in vivo* studies only antagonism was observed. At the time of review, the SCF were not aware of any other NIV with other trichothecene combinations examined *in vivo*. As such with only *in vitro* studies suggesting dose additivity, the establishment of the nature of combined effects or relative potencies of trichothecenes was not further explored. In conjunction to this, the SCF did not support the establishment of a group TDI for all trichothecenes evaluated as synergism was not observed.

#### MYCOMIX

48. The MYCOMIX project carried out in 2013-2015 aimed to contribute and fill the gap concerning the risk assessment of children to multiple mycotoxins in infant foods (Alvito *et al.*, 2015). Three questions were postulated. Firstly, are children exposed daily to one or several mycotoxins via the diet. Secondly, can this co-exposure affect children's health, and lastly are there interactive effects in toxicity of mixtures of mycotoxins.

49. The final study report was published in 2018 (Assunção *et al.*, 2018). Analysis of 52 different cereal-based products revealed a co-occurrence of mycotoxins in 75% of the analysed samples, with two or more mycotoxins occurring simultaneously. The highest number of mycotoxins detected simultaneously was seven and the combinations of two (OTA and DON; OTA and FBs) and four (AFs, OTA and ZEN) mycotoxins were the most commonly detected, with a percentage of occurrence of 6% for each combination.

50. Food diary analysis revealed that ~92% of the children (n=75; 18 males and 20 females (13-24 months), 9 males and 9 females (25-36 months), 7 males and 12 females (36-47 months)) consumed one or more cereal-based products, and at least once in three days. 42%, 65% and 65% consumed breakfast cereals, infant cereals and biscuits, respectively. The mean daily consumption of these food groups, for all children (both non-consumers and consumers), were 5.6 g (breakfast cereals), 25.3 g (infant cereals) and 8.7 g (biscuits). For the only consumers group, the values increase to 15.4 g, 38.7 g, and 13.4 g, for the same food groups respectively.

51. Worst-case exposure for the summed daily intake of mycotoxins present in cereal-based products (breakfast cereal, infant cereal and biscuits) are presented in

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52. *Table 2.*

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Table 2 - Sum of worst-case children's daily intake of mycotoxins present in cereal-based products (breakfast cereal, infant cereal and biscuits) based on a deterministic approach (reproduced from Assunção *et al.*, 2018).

	<b>Consumers and non-consumers</b>	<b>Only consumers</b>
<b>Toxins</b>	<b>Sum of daily intake (ng/kg bw/day)</b>	
<b>AFM<sub>1</sub></b>	0.069	0.116
<b>AFB<sub>1</sub></b>	0.012	0.028
<b>AFB<sub>2</sub></b>	0.003	0.006
<b>AFG<sub>1</sub></b>	0.016	0.028
<b>OTA</b>	0.131	0.227
<b>FB1</b>	6.4	14.0
<b>FB2</b>	1.0	2.6
<b>DON</b>	57.22	112.78
<b>NIV</b>	2.68	6.60
<b>ZEN</b>	0.86	1.64

Abbreviations: AFM<sub>1</sub>; Aflatoxin M1, AFB<sub>1</sub>; Aflatoxin B1, AFB<sub>2</sub>; Aflatoxin B2, AFG<sub>1</sub>; Aflatoxin G1, OTA; Ochratoxin A, FB<sub>1</sub>; Fumonisin B1, FB<sub>2</sub>; Fumonisin B2, DON; Deoxynivalenol, NIV; Nivalenol, ZEN; Zearalenone.

53. Breakfast cereals were the highest contributor for the estimated daily intake of mycotoxins by Portuguese children under three years old, revealing the highest values for FBs, trichothecenes, ZEN and AFB<sub>1</sub>. On the other hand, processed cereal-based foods (flours) presented the highest contribution for the estimated daily intakes of AFM<sub>1</sub>, AFB<sub>2</sub>, AFG<sub>1</sub>, and OTA.

54. The risks derived from exposure of children to mixtures of mycotoxins in breakfast cereals were assessed, and the authors concluded that there was a potential health concern in this population group.

55. The authors recommended that a national monitoring program was carried out with the aim of establishing protective values in legislation. Additionally, further research should be conducted to obtain toxicological data, including health consequences resulting from early-life exposures to multiple mycotoxins. Good farming and food production processes should aim to reduce the generation of mycotoxins in crops, and greater consideration of decontamination of foods destined for children consumption.

## MYTOX

56. MYTOX<sup>10</sup> is a multi-disciplinary research group, which deals with issues involving toxigenic moulds, mycotoxins, mycotoxins and human health, and mycotoxins and animal health.

57. The mycotoxins and human health research unit includes all research projects in relation to the occurrence of mycotoxins and its effects on human health including epidemiological studies, risk assessment studies and

<sup>10</sup> Further information can be found at the [MYTOX website](#).

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scenario analyses. It is, however, unspecified whether there are ongoing projects on the potential risks of aggregate dietary exposure to mycotoxins.

## Toxicokinetics

58. The Committee previously reviewed the available toxicokinetic data relating to combined exposures to mycotoxins (Warth *et al.*, 2013; Battilani *et al.*, 2020), and concluded that there is a limited number of studies in order to fully understand the toxicokinetic profiles of varied mycotoxin mixtures *in vivo* in humans.

59. Battilani *et al.*, (2020) further highlighted the complexity of studying the toxicokinetic of mycotoxin mixtures, suggesting that it needed to be addressed on a case-by-case approach. Mycotoxin dosage, exposure pathway, interspecies and intraspecies differences were identified among the most important parameters that may influence the toxicokinetics of mixtures.

## Toxicology

60. This section is presented in three parts. Firstly, the Committee's review on the toxicity of single mycotoxins. Secondly, the review of the available data on relative potencies for mycotoxin groups, and lastly, the review of the toxicity of common binary mixtures found in literature.

61. The COT has previously reviewed the toxicity of single mycotoxins; especially in the diet of infants and children aged 0-12 months and 1-5 years, respectively (TOX/2017/30)<sup>11</sup>. Annex B of TOX/2020/44<sup>12</sup>, presented a detailed yet succinct overview of all mycotoxins previously covered in the scope of TOX/2017/30. The collated information includes; their associated mycotoxins, the species of fungus that produces them, their mode of action (MOA), key toxicological endpoints, as well as their recommended health-based guidance values as set by authoritative bodies such as EFSA, JECFA, SCF *etc.*

62. In mixtures toxicology, there are three main different categories of interactions between mycotoxins. These are:

- i). Additive – where the observed effect of the mycotoxin combination is the sum of the individual effects of the two studied toxins;
- ii). Antagonistic – where the observed effect of the mycotoxin combination is less than expected from the sum of the individual effects of the studied mycotoxins and;

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<sup>11</sup> The first scoping paper [TOX/2017/30](#) and the resulting [addendum to the 0-5 years Overarching Statement](#) are available on the COT website.

<sup>12</sup> Annex B of TOX/2020/44 is available on the [COT website](#).

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- iii). Synergistic – where the observed effect of the mycotoxin combination is greater than what is expected in comparison to the sum of the individual effects of the two studied mycotoxins.

63. A paper by Speijers & Speijers (2004) on combined toxic effects was identified to be one of the first reviews to assess the topic. It concluded that, at the time tools were not fully developed to establish the type of interaction or whether there is any interaction at all (with regards to trichothecenes). More recent reviews have also been published by Grenier and Oswald (2011), De Ruyck *et al.*, (2015), Alassane-Kpembi *et al.*, (2017), Lee & Ryu (2017), and Battilani *et al.*, (2020).

64. Authoritative groups such as EFSA, JECFA, and SCF have provided opinions on some binary mycotoxin mixtures (refer to paragraphs 20-47).

65. The COT reviewed literature on binary mixtures as the co-occurrence of two mycotoxins in food commodities is the more commonly reported. A summary of OTA, AFB1 and *Fusarium* spp. mycotoxins and their interactions with other mycotoxins is presented in *Table 3* *Table 5*, respectively.

66. The majority of studies testing for the combined effects of mycotoxins are *in vitro*, with cell viability endpoints (e.g. apoptosis, necrosis, DNA damage, oxidative damage and immunotoxicity) being the most commonly assessed. Available *in vivo* data reported potential adverse effects on the liver, kidneys and teratogenicity.

67. Other considerations for the potential adverse effects on the microbiota (Baines *et al.*, 2013; Liew & Mohf-Redzwan, 2018) and endocrine system (Demaegdt *et al.*, 2016) were also reviewed by the Committee.

68. The Committee observed that there were a number of mycotoxins with MOAs involving ribosomal protein synthesis inhibition; however, there was a lack of information on possible additive toxicity. Additionally, there was a large amount of variability in the methodology utilised since there is currently no harmonisation on combinative testing strategies for each toxicological endpoint for each plausible mycotoxin combination.



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Table 3 - Provides a highlight summary of the observed combinative effects of ochratoxin A with fumonisin B1, zearalenone, and citrinin in different *in vitro* methods.

<b>Mycotoxin mixture</b>	<b>Method</b>	<b>Cellular or animal model</b>	<b>Endpoint</b>	<b>Combination effect</b>	<b>Reference</b>
<b>OTA + FB<sub>1</sub></b>	<i>In vitro</i>	Rat C6 glioma, Vero monkey and human Caco-2 cells	Cytotoxicity	Synergistic	Creppy <i>et al.</i> , (2004)
	<i>In vitro</i>	PK-15 cells	Cytotoxicity	Additive	Šegvić Klarić <i>et al.</i> , (2007)
	<i>In vitro</i>	human and pig lymphocytes	Cytotoxicity	Synergistic	Mwanza <i>et al.</i> , (2009)
	<i>In vitro</i>	Male Wistar rats	Genotoxicity	Synergistic	Domijan <i>et al.</i> , (2006)
<b>OTA + ZEN</b>	<i>In vitro</i>	hHepG2 cells	Cytotoxicity	Antagonistic	Wang <i>et al.</i> , (2014)
	<i>In vitro</i>	hHepG2 and KK-1 cells	Cytotoxicity	Additive	Li <i>et al.</i> , (2014)
<b>OTA + CIT</b>	<i>In vitro</i>	Piglets lymphocytes	Immunotoxicity	Synergistic	Bernhoft <i>et al.</i> , (2004)
	<i>In vitro</i>	Monkey kidney vero cells	Cytotoxicity	Synergistic	Bouslimi <i>et al.</i> , (2008b)
	<i>In vitro</i>	PK-15 cells	Cytotoxicity	Antagonistic	Šegvić Klarić <i>et al.</i> , (2012)

Abbreviations: OTA = Ochratoxin A; FB1 = Fumonisin B1; ZEN = Zearalenone; CIT = Citrinin; PK-15 = Porcine kidney 15 epithelial cells; hHepG2 = human hepatoma cells G2; KK-1 = murine ovarian granular cells.

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Table 4 - Provides a highlight summary of the observed combinative effects of aflatoxin B<sub>1</sub> with aflatoxin B<sub>2</sub>, fumonisin B<sub>1</sub>, deoxynivalenol, zearalenone, T-2 toxin, and ochratoxin A in different *in vitro* and *in vivo* models.

<b>Mycotoxin mixture</b>	<b>Method</b>	<b>Cellular or animal model</b>	<b>Toxicity endpoint</b>	<b>Combination effect</b>	<b>Reference</b>
<b>AFB<sub>1</sub> + AFB<sub>2</sub></b>	<i>In vitro</i>	hUVEC	Cytotoxicity	Synergistic	Braicu <i>et al.</i> , (2010)
	<i>In vitro</i>	hLF and A 2780	Cytotoxicity	Additive	
	<i>In vitro</i>	Rat liver slices	Toxicity	Antagonistic	Friedman <i>et al.</i> , (1997)
<b>AFB<sub>1</sub> + FB<sub>1</sub></b>	<i>In vitro</i>	Spleen mononuclear cells	Cytotoxicity	Synergistic	Mary <i>et al.</i> , (2012)
	<i>In vitro</i>	hHep-G2 and hBEAS-2B cells	Cytotoxicity	Additive and antagonistic	McKean <i>et al.</i> , (2006b)
	<i>In vivo</i>	Fischer 344 rats	Acute	Synergistic	
	<i>In vivo</i>	Male Fischer rats	Hepatotoxicity	Synergistic	Gelderblom <i>et al.</i> , (2002)
	<i>In vivo</i>	White rabbits	Hepatotoxicity	Synergistic	Orsi <i>et al.</i> , (2007)
	<i>In vivo</i>	Male Wistar rats	Hepatotoxicity	Synergistic	Theumer <i>et al.</i> , (2008)
<b>AFB<sub>1</sub> + DON</b>	<i>In vitro</i>	PK-15 cells	Cytotoxicity	Synergistic	Lei <i>et al.</i> , (2013)
	<i>In vitro</i>	BRL 3A cells	Cytotoxicity	Synergistic	Sun <i>et al.</i> , (2015)
	<i>In vitro</i>	Ames test	Mutagenicity	Synergistic	Šmerák <i>et al.</i> , (2001)
<b>AFB<sub>1</sub> + ZEN</b>	<i>In vitro</i>	PK-15 cells	Cytotoxicity	Synergistic	Lei <i>et al.</i> , (2013)
	<i>In vitro</i>	BRL 3A cells	Cytotoxicity	Synergistic	Sun <i>et al.</i> , (2015)
<b>AFB<sub>1</sub> + T-2</b>	<i>In vitro</i>	Ames test	Mutagenicity	Synergistic	Šmerák <i>et al.</i> , (2001)
	<i>In vitro</i>	hBEAS-2B cells	Cytotoxicity	Additive and synergistic	McKean <i>et al.</i> , (2006a)
	<i>In vivo</i>	Fischer 344 rats	Acute	Additive	
<b>AFB<sub>1</sub> + OTA</b>	<i>In vitro</i>	Ames test	Mutagenicity	Synergistic	Sedmíková <i>et al.</i> , (2001)
	<i>In vitro</i>	Monkey kidney vero cells	Cyto and genotoxicity	Additive	Golli-Bennour <i>et al.</i> , (2010)
	<i>In vitro</i>	hHep-G2 cells	Cytotoxicity	Additive	Corcuera <i>et al.</i> , (2011)
	<i>In vitro</i>	hHep-G2 cells	Genotoxicity	Antagonistic	
	<i>In vivo</i>	Wistar rat dams	Teratogenicity	Antagonistic	Wangikar <i>et al.</i> , (2004)
	<i>In vivo</i>	Male Sprague Dawley rats	Hepatotoxicity and nephrotoxicity	Synergistic	Abdel-Wahhab <i>et al.</i> , (2015)

Abbreviations: AFB<sub>1</sub> = Aflatoxin B<sub>1</sub>; AFB<sub>2</sub> = Aflatoxin B<sub>2</sub>; FB<sub>1</sub> = Fumonisin B<sub>1</sub>; DON = Deoxynivalenol; ZEN = Zearalenone; T-2 = T-2 toxin; OTA = Ochratoxin A; HUVEC = Human umbilical vein endothelial cells; HLF= Human lung fibroblasts; hHep-G2 = Human hepatoma G2 cells; hBEAS-2B = Human bronchial epithelial cells; PK-15 = Porcine kidney 15 epithelial cells; BRL 3A = Buffalo rat liver cells.

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Table 5 - Provides a highlight summary of the observed combinative effects of various combined *Fusarium* mycotoxins including: zearalenone and fumonisin B<sub>1</sub>; zearalenone and T-2 toxin; deoxynivalenol and zearalenone; deoxynivalenol and T-2 toxin; deoxynivalenol and 15-acetyldeoxynivalenol; deoxynivalenol and nivalenol; and deoxynivalenol and fumonisin B<sub>1</sub> in different *in vitro* and *in vivo* methods.

<b>Mycotoxin mixture</b>	<b>Method</b>	<b>Cellular or animal model</b>	<b>Endpoint</b>	<b>Combination effect</b>	<b>Reference</b>
<b>ZEN + FB<sub>1</sub></b>	<i>In vitro</i>	Human Caco-2 cells	Cytotoxicity	Antagonistic	Kouadio <i>et al.</i> , (2007)
			Lipid peroxidation	Synergistic	
			Inhibition of DNA synthesis	Antagonistic	
			DNA fragmentation	Synergistic	
<b>ZEN + T-2</b>	<i>In vitro</i>	hCFU-GM cells	Myelotoxicity	Additive	Ficheux <i>et al.</i> , (2012)
<b>DON + ZEN</b>	<i>In vitro</i>	hCFU-GM cells	Myelotoxicity	Additive	Bensassi <i>et al.</i> , (2014)
	<i>In vitro</i>	HCT116 cells	Cytotoxicity	Antagonistic	
<b>DON + T-2</b>	<i>In vitro</i>	hCFU-GM cells	Myelotoxicity	Additive or synergistic	Ficheux <i>et al.</i> , (2012)
<b>DON + 15-AcDON</b>	<i>In vitro</i>	hGES-1 cells	Cytotoxicity	Synergistic	Yang <i>et al.</i> , (2017)
<b>DON + NIV</b>	<i>In vitro</i>	Human Caco-2 cells	Cytotoxicity	Synergistic and additive	Alassane-Kpembi <i>et al.</i> , (2013)
	<i>In vitro</i>	hGES-1 cells	Cytotoxicity	Synergistic	Yang <i>et al.</i> , (2017)
<b>DON + FB<sub>1</sub></b>	<i>In vitro</i>	Human Caco-2 cells	Cytotoxicity	Synergistic	Kouadio <i>et al.</i> , (2007)
	<i>In vitro</i>	hCFU-GM cells	Myelotoxicity	Antagonistic	Ficheux <i>et al.</i> , (2012)
	<i>In vivo</i>	Male crossbred castrated piglets	Morphological changes	Antagonistic	Bracarense <i>et al.</i> , (2012)
			Immunological changes	Synergistic-Antagonistic	

Abbreviations: ZEN = Zearalenone; T-2 = T-2 toxin; DON = Deoxynivalenol; 15-AcDON = 15-acetyldeoxynivalenol; NIV = Nivalenol; FB<sub>1</sub> = Fumonisin B<sub>1</sub>; hCFU-GM = Human colony forming unit-granulocyte and macrophage cells; HCT116 = human colon carcinoma cell line; GES-1 = Human gastric epithelial cells.

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## **Exposure assessment**

69. As stated previously, the co-occurrence of mycotoxins in food and feed is possible since some fungi species are able to produce more than one mycotoxin (for example the *Fusarium* spp.), food commodities can also be contaminated by several fungi species. A combined exposure is therefore highly likely in humans due to varied diets and/or one food commodity may be contaminated with more than one mycotoxin.

70. The completion of an exposure assessment is challenging when limited information is available. Little to no UK relevant data could be obtained from the literature where all age groups were considered. It would be necessary to assess all age groups to determine those who would be of greater risk. Additionally, different methodologies have been observed in the literature to assess the levels of exposure (e.g. food diaries, biomarker analyses etc), as such performing data comparison may not be accurate. The development and application of multi-analyte methods has been advancing as detailed earlier; however, this has not yet been internationally applied as a gold standard for assessing the presence of multiple mycotoxins in food commodities. The use of current methodologies for mycotoxin analysis (e.g. HPLC) still presents an issue in terms of management of left-censored data.

### *Stepwise approach*

71. A stepwise approach to the exposure assessment was considered by the Committee (summarised below).

72. Firstly, mycotoxins should be categorised based on toxicological similarities where an endpoint is defined. This will then determine how occurrence data for the considered mycotoxins should be grouped together to calculate total residues for each mycotoxin group by summation in the exposure assessment (either in one food or multiple foods). An opportunity to note any missing data can be recorded throughout this step.

73. The exposure should be then calculated deterministically, and if major exceedances are observed in relation to the toxicological endpoint a probabilistic calculation should be considered. The estimated exposure can then be compared against the health-based guidance value to determine the MOE. Depending on the endpoint, an MOE value that is  $\leq 100$  (for non-genotoxic and non-carcinogenic compounds) or 10,000 (for genotoxic carcinogens) would indicate a level of risk whilst values that are  $\geq 100$  or 10,000 indicates no appreciable cause of concern (EFSA, 2012b). The MOE value aids in putting exceedances into perspective.

74. Lastly, if probabilistic modelling was carried out; a sensitivity analysis should be considered for assessing the impact of different variables.

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### *Potential data sources*

75. Three potential data sources to use in the exposure assessment were proposed and summarised in the following paragraphs.

#### *FSA – Mycotoxin Total Diet Study*

76. The FSA has previously carried out a Total Diet Study (TDS)<sup>13</sup> that included mycotoxin analysis in 2011 (FS102081), where a total number of 3,312 food samples were analysed for the presence of mycotoxins. The main aim of the study was to calculate background exposure to various mycotoxins from the whole diet and to compare exposure to those calculated by other sources (Stratton *et al.*, 2017).

77. Co-occurrences were observed in the TDS dataset. For example, sample S14-042859 (a wholemeal bread) contained DON, some ergot alkaloids and also a low level of OTA.

78. The following limitations were observed with the TDS dataset including; that it was limited to a small number of food groups, some recovery rates were poor, food samples were collected from 2009 and as such may not be reflective of the current levels of mycotoxins detected in foods. Finally, as mentioned multi-mycotoxin analysis was not consistently used for each food sample.

79. Further information received from the project manager; Susan MacDonald (personal communication, 2020) has confirmed that a method for multi-mycotoxin analysis (*i.e.* different classes/families) was not performed for the TDS. This was due to the different chemistries and properties of the mycotoxins themselves rather than the different food matrices. Therefore, samples were analysed by several methods to obtain the full suite of analyte results with the lowest reporting limits achievable. Although not strictly a multi-analyte method, mycotoxins from the same family (*e.g.* ergot alkaloids and trichothecenes) were detected using one methodology. The possibility and availability of a multi-mycotoxin method were also discussed with the project manager. It was confirmed that a methodology is both possible and available although this means that compromises have to be made in order to make it suitable for all tested mycotoxins. The compromises include the lack of dedicated extraction techniques, sample clean-up and analyte enrichment. This results in higher reporting limits, which in turn can affect the estimate of intake (*i.e.* overestimation), as well as potentially requiring additional resources in terms of sample re-analyses to avoid false positives.

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<sup>13</sup> The Food Standards Agency Total Diet Study on Mycotoxins can be found on the [FSA website](#).

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### EFSA – MYCHIF Platform

80. It was noted that UK co-occurrence data was presented in the scientific report for MYCHIF, therefore the MYCHIF platform (Battilani *et al.*, 2020) was included as a potential data source.

81. A human case study was presented as part of the MYCHIF report (Chapter 3.8.2, pp. 88) where an aggregate chronic exposure assessment for two mycotoxin mixtures in cereal food sources (1: DON, FBs and ZEN and 2: T-2/HT-2 toxin, DON and NIV) was carried out using two modelling methodologies: CBA and provisional daily intake approach.

82. In the CBA, the co-occurrence of mycotoxins and consumption data of cereal-food based products (as an example) for each single mycotoxin were combined to obtain an individual mycotoxin exposure, this was then summed up to obtain the total exposure, under the dose addition assumption. The main uncertainty identified was the adopted deterministic approach of the input modelling data for mycotoxin concentrations. Due to the scarcity of concentration data for many countries, a probabilistic approach was applied at an EU level only. It was assumed that the maximum exposure limits (*i.e.* the lower and upper bound highest 95th percentile chronic exposure) were the most conservative values. The risk decision was based on the calculated MOE values, based on the methodology and assumptions in the MYCHIF case study the MOE values were <100 for all age groups (adolescent, adult and elderly).

83. The provisional daily intake (expressed in  $\mu\text{g}/\text{kg bw}/\text{day}$ ) models the internal dose with the available human biomarker data to derive exposure to the mixture. This was estimated by combining the mycotoxin concentration in the urine, the available excretion rate for each of the mycotoxin in the mixture, the human body weight and the daily urine excretion volume ( $\mu\text{g}/\text{L}$  mycotoxin, L urine in 24 hrs, % excretion rate, kg bw, respectively). Values were calculated for single mycotoxins present in the mixture and for the mixture. A hazard exposure index<sup>14</sup> was used to estimate the risk, if the value is  $\leq 1$  the combined risk as deemed acceptable, whereas when it is  $> 1$  a potential concern is possible. The identified uncertainties included the default body weight of 70 kg, the excretion rate where values were derived from a single study or from correlation approximations, urine volumes where the urine was not corrected for dilution factors, and data representativeness. Overall, a hazard exposure index could not be quantified due to the uncertainties for single mycotoxins described above, since these would also need to be integrated into the analysis of mixtures where additional variables should be considered for unknown toxicokinetics and toxicodynamics, as well as unknown synergistic/additive effects.

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<sup>14</sup> A hazard exposure index is a risk-assessment tool, which can indicate whether further investigations are required for mixtures. It is based on dose addition assumptions; it is the sum of the hazard quotients of the chemicals in the mixture.

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84. The MYCHIF data was located and the file downloaded. Extraction of relevant UK data was attempted; however, the datasets are extremely complex and were not accompanied with straightforward guidance. The Secretariat has made the relevant contact in order to gain further guidance. The following benefits to using this data have been observed; the use of multi-mycotoxin analysis, the more recent data collection of mycotoxin co-occurrences in cereal commodities, and the integration of singular and multi-biomarker mycotoxin analysis data.

85. Nevertheless, at present it has been difficult to determine potential limitations due to the complexity of the datasets. Uncertainties for both modelling methodologies were also identified.

#### *Exposure data derived from the literature*

##### *Non-UK data*

86. Co-exposures have been reported in the literature for various age groups such as those observed from the MYCOMIX Portuguese studies in children as seen in paragraph 48.

87. For the exposure assessment, non-UK co-occurrence data and consumption data for each single mycotoxin could be used to obtain an individual mycotoxin exposure. These could then be totalled for each co-occurrence type and also for each mycotoxin family to obtain the total exposure within each food or food group.

88. Obvious limitations include the use of non-UK data which may not be applicable to consumers in the UK, however, the use of non-UK data may expedite an exposure assessment to reveal common mycotoxin combinations as well as the most affected food groups in Europe.

##### *Human biomonitoring data*

89. The use of human biomonitoring (HBM) data was explored as part of TOX/2020/44<sup>15</sup>, outputs from this exercise are summarised below.

90. Biological monitoring utilises biomarkers<sup>16</sup> to represent or estimate the internal exposure as a result of inhalation, ingestion or dermal exposure to a chemical, and as such, biomarkers are indicators of exposure, effect, and/or susceptibility. Typically, exposure assessments to any dietary contaminant is based on intakes from food (or feed), otherwise known as the external

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<sup>15</sup> TOX/2020/44 is available on the [COT website](#).

<sup>16</sup> A biomarker is a naturally occurring molecule, gene, or characteristic by which a particular pathophysiological or physical process, disease etc. can be identified.

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exposure or oral dose. However, the bioaccessibility<sup>17</sup> and bioavailability<sup>18</sup> of the contaminant determines the internal exposure.

91. Mycotoxins can be classified as short-lived chemicals that can only be effectively measured if the individual is undergoing continuous or continual exposures or if the timing of exposure(s) is known. Mycotoxin biomarkers have been defined as the compounds themselves (e.g. parent compounds and/or a metabolite) or as a result of interaction with target molecules (e.g. DNA or protein adducts) (Marín *et al.*, 2018). Urinary excretion mainly represents recent mycotoxin intake, whereas measurements in plasma/serum are more likely to represent long-term exposure.

92. The main analytical methods employed to perform biomarker analyses are based on either chromatography (e.g. LC) or immunochemistry (e.g. ELISA).

93. European human biomonitoring initiatives such as the Consortium to Perform Human Biomonitoring on a European Scale (COPHES)<sup>19</sup>, Human Early-Life Exposome (HELIX)<sup>20</sup>, and European Union Human Biomonitoring (HBM4EU)<sup>21</sup>, as well as typical literature databases (PubMed, Science Direct, Google Scholar, Scopus and Zenodo) and the Information Platform for Chemical Monitoring (IPChem) platform – were mined for any relevant UK biomonitoring data on combined exposures to mycotoxins.

94. Both COPHES and HELIX did not include exposure to mycotoxins in the scope of their work, however, in the HBM4EU initiative it was. In brief, Alvito *et al.*, (2019)<sup>22</sup> conclude that there are numerous factors that need to be considered when attempting to integrate biomarker data for exposure assessment – and thus the following risk assessment. These factors include: the validation and harmonisation of analytical methods to assess mycotoxin exposure biomarkers, a greater understanding of the current exposure levels of the European population to multiple mycotoxins and whether this differs for each Member State *etc.*

95. There are currently no UK government led HBM initiatives relating to mycotoxins, however, scientific interest for this has and continues to grow which has led to several publications. These publications were previously reviewed by the COT. It was observed that the available literature seems to focus on estimating DON exposures from using total DON (free DON and DON-glucuronides) in urinary samples as biomarkers in the UK population. Only one other study for OTA exposure was reported by Gilbert *et al.*, (2001).

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<sup>17</sup> Bioaccessibility describes events that take place during food digestion for transformation into potentially bio-accessible material, the absorption/assimilation through epithelial tissue and pre-systemic metabolism.

<sup>18</sup> Bioavailability describes the fraction of bio-accessible material which is likely to reach the systemic circulation.

<sup>19</sup> The COPHES final report is available on the [EU HBM website](#), a brief [technical report](#) is also available.

<sup>20</sup> Further information on the HELIX project is available on the [CORDIS EUROPA website](#).

<sup>21</sup> A pdf file for a brief informative guide for HBM4EU is available on the [HBM4EU website](#).

<sup>22</sup> The HBM4EU mycotoxin paper by Alvito *et al.*, (2019) is available on the [HBM4EU website](#).



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96. The presence of DOM-1, a metabolite that is formed by microbiota metabolism in urine is rarely reported suggesting that it may not be a suitable as an exposure biomarker for DON (Turner *et al.*, 2008a, 2008b, 2009, 2010, 2011; Hepworth *et al.*, 2012; Wells *et al.*, 2016; Papageorgiou *et al.*, 2018a, 2018b). From the reviewed studies, a proportion of the UK population (adults, children, adolescents, pregnant women, elderly and vegetarians) exceeded the TDI for DON of 1 µg/kg bw (EFSA, 2017b).

97. At the time of review, only one report by Gratz *et al.*, (2020) performed multi-mycotoxin biomarker analyses. In this pilot study, UK children (n=21) were estimated to frequently exceed the TDI for 52% of DON and 95% of OTA cases.

#### *Summary of exposure assessment*

98. As discussed, there is little to no relevant UK co-occurrence data for mycotoxins. Available data either from food surveys, total diet studies and other databases have their own associated limitations.

99. Although the advancement and availability of detection techniques and equipment has progressed the development of biomarkers of exposure to mycotoxins are limited mainly to AFs, OTA, DON, FBs, ZEN and to a lesser extent – emerging mycotoxins such as Fus-X, CIT, NIV, T-2 toxin, 4,15-DAS, ENNs, ALT and tenuazonic acid.

100. Understanding the toxicokinetics of mycotoxin metabolites and their availability in different biological samples (e.g. OTA has the potential to be transferred to breastmilk) and how they may correlate to the exposure still needs further investigation. Additionally, there is a lack of harmonisation in the experimental settings and design, with particular reference to data collection and in the definition of performance criteria of fit for purpose analytical methods. Oversights on sampling strategy were also noted, for example the lack of knowledge regarding the stability of the biomarker, the defined time of sampling, and detailed information regarding the way of sample collection and storage. Furthermore, there is a need for the development of biomarkers of exposure for the detection of masked mycotoxins

101. As such it was concluded by the COT that further research is required to enable the inclusion of HBM data for a robust exposure assessment.

#### **Risk assessment**

102. It is generally accepted that the effects of a combination of mycotoxins cannot be predicted based solely on their individual effects and that, in addition to additivities and synergies, there can also be antagonisms observed.

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103. The combined toxic effects that are observed will greatly depend on the experimental design. Factors such as the type of experimental cells or animal models, the duration of exposure, the dosage and relation between mycotoxins (*i.e.* the ratio of each mycotoxin in the mixture), the tested endpoint and methodology used; including any statistical aspects used for modelling scenarios.

104. Potential uncertainties could arise when comparing between toxicity studies utilising “*natural*” contaminated test samples and purified extracts. For example, in livestock studies where feed is naturally contaminated with DON, a higher toxicity was observed when compared to exposure groups treated purified DON. This result was attributed to the presence of additional fungal metabolites, where low concentrations of ZEN or the 3- or 15-AcDON precursors were found in some cases.

105. The main challenges in hazard assessment of multiple mycotoxins include; the lack of accurate information regarding the toxicokinetic profiles of mycotoxin mixtures and their bioaccessibility (*i.e.* the actual percentage of mycotoxins that can be absorbed in the small intestine) that would enable a more accurate risk assessment. Additionally, the variability within mycotoxin bioaccessibility values depends on the compound, food product, contamination level and the nature of contamination (spiked or naturally contaminated). Furthermore, the breadth of *in vitro* digestion models used to assess the bioaccessibility of mycotoxins constitutes another important challenge. The suitability and reliability of *in vitro* models for *in vivo* extrapolation are also required to be comparable to the situation *in vivo*.

106. The mycotoxin absorption constitutes as another challenge considering that toxins could reach the intestine as the parent compound or as metabolites formed during digestion; the available methods for mycotoxin metabolites are also still in its infant stages. Studies on combined genotoxic effects of mycotoxin mixtures should also be further developed.

107. As for the exposure assessment, there is little to no relevant UK co-occurrence data for mycotoxins and HBM data. Available data either from food surveys, total diet studies and other databases have their own associated limitations and at this time an exposure assessment could not be performed.

108. In terms of risk assessment, one of the main challenges posed to risk characterisation is the absence of toxicological data. A deeper understanding of the interactions between multiple mycotoxins at a molecular level will assist in drawing real life conclusions on the health impact of human exposure to mycotoxin mixtures.

109. Based on the limitations discussed above and the absence of an exposure assessment, a full risk assessment on the potential adverse effects of aggregate dietary exposure to mycotoxins could not be carried out presently.

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## Further considerations

110. Concerns over the effect of climate change on fungi and host specific interactions were mentioned previously in this document. Moretti *et al.*, (2019) provided a narrative review on the potential emerging mycotoxin risks under a climate change scenario in Europe. It was hypothesised that the contamination risk of aflatoxin (produced by *Aspergillus flavus*) in maize in South and Central-Europe will extend to new regions in the next 30 years. The *Fusarium* spp. species profile on wheat are also hypothesised to change in Northern, Central and Southern-Europe. It is unknown whether these changes will be similar across these regions. As a result, new combinations of mycotoxins/host plants/geographical areas may arise. It was recommended by Moretti *et al.*, (2019) that developments of new diagnostic tools, a deeper knowledge of both biology, and genetics of toxigenic fungi may be required.

111. Co-exposure to mycotoxins from breast milk and infant formula may also need to be considered in infants and young children. Several publications have already considered this as a potential risk. For example, Ortiz *et al.*, (2018) have investigated the multiple mycotoxin exposure of infants and young children (0-23 months) via breastfeeding and complementary/weaning foods consumption in Ecuadorian highlands and Braun *et al.*, (2020) whom performed a longitudinal assessment of mycotoxin co-exposures in exclusively breastfed infants in Austria.

## Summary

112. Mycotoxins are toxic secondary metabolites produced by fungi and is capable of causing adverse health effects in both humans and animals. Those of greatest concern to human health are produced by several fungal genera of filamentous fungi, namely *Aspergillus*, *Fusarium* and *Penicillium* spp.

113. DON, FBs, and ZEN are the most prevalent mycotoxins in the world with regards to cereals and cereal based products, with a prevalence of 66%, 56%, and 53%, respectively. Regulation limits for these compounds are based on considerations for the toxicity of single exposures. There are several reports where co-exposure to multiple mycotoxins are observed in both humans and animals. As such, the literature was reviewed to investigate the potential risks of aggregate dietary exposure from mycotoxins.

114. An external EFSA report by Battilani *et al.*, (2020) was recently published; the group carried out an extensive literature review whereby a platform was built named MYCHIF. The database is comprised of four topics including: the ecological background of mycotoxins and their interactions with host plants, the available analytical methods to detect the co-occurrence of mycotoxins, the toxicological and biomarkers data relevant to humans and animals and modelling approaches in order to perform risk modelling.

115. Using the gathered information, a case study was carried out for two mycotoxin mixtures (1: DON, FBs and ZEN and 2: T-2/HT-2 toxin, DON and

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NIV). Biomarker data was utilised as the basis for the exposure assessment which was carried out probabilistically either with a component-based and provisional daily intake approaches.

116. For the CBA an MOE of <100 for all age groups (adolescent, adult and elderly) was calculated, where as a hazard index exposure index could not be calculated for the provisional daily intake approach. A hierarchy map (based on EU member states) for adults was compiled and provided a visual representation of higher risked exposure groups to T2/HT-2 toxin, DON and NIV.

117. In terms of data gaps, there is still limited knowledge on the presence and co-occurrence of multiple mycotoxins, both for native mycotoxins and their modified forms, in food and feed since current analytical methods have limitations. Furthermore, there is a limited number of toxicity data and there is a lack of consensus on methodologies and guidelines for generating *in vitro* and *in vivo* TK and TD. Model definitions are also required for the utilisation of biomarkers for exposure assessments.

118. Other opinions by authoritative groups on some mycotoxin mixtures were also summarised. The EFSA CONTAM Panel concluded that the available data for interactions between 4,15-DAS and other mycotoxins (T-2 and HT-2 toxins, AFs, OTA and FBs) is weak and inconclusive (EFSA, 2018). In contrast, the EFSA CONTAM Panel concluded that the combined effect of CIT and OTA is at most additive (EFSA, 2012). The JECFA Committee concluded that even though there are additive or synergistic effects observed from FB1 and AFB1 co-exposure in laboratory animals in inducing the development of preneoplastic lesions or hepatocellular carcinoma (Torres *et al.*, 2015; Carlson *et al.*, 2001; Gelderbloom *et al.*, 2002), there was currently no data available on such effects in humans. The combined toxicity of FBs and DON were suggestive of being additive or more than additive, however, the observed effect is dependent on the endpoints measured (JECFA, 2018).

119. The main analytical methods to measure mycotoxins are: ELISA, GC and LC-MS, whilst LC-MS/MS is the main analytical method for detecting and measuring co-occurrence of very low concentrations of mycotoxins, however, these advanced multi-mycotoxin techniques are not yet commonly applied in routine screening analyses due to their associated high cost.

120. The co-occurrence of mycotoxins in food and feed is quite common since some fungi can produce more than one mycotoxin (particularly *Fusarium* spp.), food commodities can be contaminated by several fungi, and animal and human diets usually consist of multiple commodities.

121. In terms of toxicokinetic data, only one human study was identified to have investigated and analysed the combined exposure to DON and ZEN (Warth *et al.*, 2013). This study, however, had its limitations which was mainly due to the number of volunteers (n=1 male) and in effect, does not cover inter-individual variations. It is hypothesised that the toxicokinetics of mycotoxin mixtures may need to be addressed on a case-by-case approach, however, it

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is recognised that the mycotoxin dose, exposure pathway, interspecies and intraspecies differences are identified as the most influential parameters that may affect observations.

122. The toxic effects of some binary mycotoxins were discussed in this paper (e.g. AFB1 and FB1, OTA and DON etc). The availability of *in vivo* data directly relevant for humans is scarce with most studies only covering a limited number of mycotoxin combinations and more generally focused on animal models of agricultural importance *i.e.* pigs and chickens.

123. It seems that the toxicity of combinations cannot be predicted based on the toxicity of individual mycotoxins. Furthermore, there is a large amount of variability between each methodology carried out for studies since there is currently no harmonisation on combinative testing strategies for each toxicological endpoint.

124. In terms of exposure assessments, the use of biomarkers data was explored in the MYCHIF report (see paragraphs 28-32 and 80-85). It was concluded that the use of biomarkers may be premature due since there is a lack of: knowledge on the human bioavailability of the toxin combination, and the excretion rate. There are also several limitations associated with multi-biomarker monitoring:

- i). Biological fluids contain extremely low analyte concentrations following dietary exposure, as such sample preparation is crucial to obtain acceptable limit of detection;
- ii). There is a great chemical diversity of analytes and makes this clean-up methodologies challenging (e.g. polar compounds like glucuronides);
- iii). Careful optimisation needs to be carried out to overcome matrix effects and interfering matrix peaks, eluents, the chromatographic gradient, and the dilution factor;
- iv). The co-elution of matrix components is said to have a negative influence the accuracy of quantitative methods through ion suppression or enhancement in the ion source and;
- v). In general, there is a lack of authentic reference standards and certified reference materials and a consensus of a validated biomarker to be used in context to a multi-mycotoxin analysis.

125. A suggested major research gap was the investigation of potential concurrent exposure of mycotoxins with other environmental chemicals that may exhibit some interactive activity and/or exert a biological function converging in the same molecular pathways.

126. A stepwise approach for the exposure assessment was detailed (see paragraphs 71-74) in this paper, these were based on the used the on use of deterministic and if necessary, probabilistic approaches. Three data sources were also identified to use in the exposure assessment, these were the FSA TDS Mycotoxin Study (Stratton et al., 2017), the MYCHIF platform (Battilani *et al.*, 2020), exposure data derived from literature.

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127. Based on the limitations presented throughout this document and the absence of an exposure assessment, a full risk assessment on the potential adverse effects of combined dietary exposure to mycotoxins could not be carried out presently.

### **COT conclusions and recommended research**

128. As noted, a risk assessment could not be carried out on the potential risks to combined exposure of mycotoxins for several reasons. There is a lack of harmonisation of approaches/methodologies and data analysis/modelling for toxicological investigations. Additionally, the underlying mechanisms of interactions between each mycotoxin combination is yet to be fully elucidated and understood. Further considerations for risk assessment include the potential toxic effects of mycotoxin mixtures on the gut microbiota and the endocrine system. Co-exposures from breastmilk and weaning foods must also be considered for infants and young children.

129. Furthermore, the availability of food consumption data is scarce, and the development of multi-analyte methods is still not yet fully applied as standard. Lastly, the management of left-censored data, the use of probabilistic models and a multi-biomarker approach should be consistent and have a well-defined approach.

130. The Committee noted that there was a lack of UK data, particularly in biomonitoring; however, there are a number of studies ongoing. Although the UK will not be collecting new data for mycotoxins under the HBM4EU initiative. However, in the future, more data could be obtained through Health Protection Research Units. Such research was considered to be a priority by the COT.

131. Members recommended as a pragmatic first step that a review should be carried out of the compounds which appeared to show a common effect on protein synthesis, assuming dose additivity, to determine whether there was any potential concern from co-exposure to these mycotoxins.

132. Research is needed on mycotoxins affecting ribosomal protein synthesis to determine whether they do exhibit dose additivity in their effects, to help develop a reliable basis for their cumulative risk assessment.

**COT**  
**October 2020**

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## Abbreviations

15-AcDON	15-acetyldeoxynivalenol
3-AcDON	3-acetyldeoxynivalenol
4,15-DAS	4,15-diacetoxyscirpenol
AFB1	Aflatoxin B1
AFB2	Aflatoxin B2
AFM1	Aflatoxin M1
AFs	Aflatoxins
ARfD	Acute reference dose
CBA	Component-based approach
CIT	Citrinin
CONTAM	Panel on Contaminants in the Food Chain
COPHES	Consortium to Perform Human Biomonitoring on a European Scale
COT	Committee on Toxicity of Chemicals in Food, Consumer Products and the Environment
DNA	Deoxyribonucleic acid
DOM-1	De-epoxy deoxynivalenol
DON	Deoxynivalenol
EFSA	European Food Safety Authority
ELISA	Enzyme-linked immunosorbent assay
ENN	Enniatins
FB1	Fumonisin B1
FBs	Fumonisin
Fus-X	Fusarenon-X
GC-MS	Gas chromatography-mass spectroscopy
HBM	Human biomonitoring
HBM4EU	European Union Human Biomonitoring
HELIX	Human Early-Life Exposome
HPLC	High-performance liquid chromatography
JECFA	Joint Food and Agriculture Organization and World Health Organisation Expert Committee on Food Additives
LC	Liquid chromatography
LC-MS	Liquid chromatography-mass spectroscopy
LC-MS/MS	Liquid chromatography tandem mass spectrometry
MOA	Mode of action
MOE	Margin of exposure
MYCHIF	Mycotoxin Mixtures in Food and Feed: Holistic, Innovative, Flexible Risk Assessment Modelling Approach
MYCOMIX	Mycotoxin mixtures
MYTOX	Mycotoxin and Toxigenic Moulds
NIV	Nivalenol
OTA	Ochratoxin A
PAT	Patulin
SCF	Scientific Committee on Food
TD	Toxicodynamic
TDI	Tolerable daily intake
TDS	Total diet study
TK	Toxicokinetic

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UK  
ZEN

United Kingdom  
Zearalenone



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