Manuscript STOTEN-D-20-13879 Jedynak et al.

Response to the Reviewer

We are very grateful for the opportunity to submit a revised version of our manuscript entitled "Prenatal exposure to a wide range of environmental chemicals and child behaviour between 3 and 7 years of age - an exposome-based approach in 5 European cohorts" for publication in the Science of the Total Environment Journal. We appreciate the time and effort that the Reviewer dedicated to providing feedback on our manuscript and are grateful for the insightful comments and valuable improvements to our work. We have incorporated most of the suggestions made by the Reviewer. Please see below for a point-by-point response to the reviewer's comments and concerns.

Reviewer: The manuscript by Jedynak et al. addresses a relevant study question during an important window of development such as pregnancy in 5 European cohorts. Indeed, this is one of the few studies examining the role of chemical mixtures on children's behavior. The authors studied 708 mother-child pairs across Europe and found that higher prenatal urinary BPA and MnBP concentrations were associated with more externalizing problems, which is in line with both the epidemiological and toxicological evidence available. On the contrary, other exposures including Copper and persistent organic pollutants (POPs) showed inverse associations, which in the case of POPs could be influenced by adipose tissue kinetics. Overall, the study objective is very relevant, the design appropriate and the manuscript is concise and very well-written, so I feel this will be a nice contribution to the field. However, I believe the following comments will help to improve the manuscript:

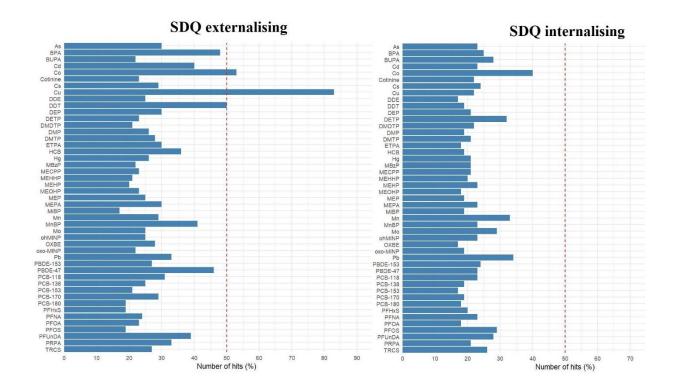
Major comments:

Reviewer's comment 1: In this study you decided to implement LASSO for variable selection. In a previous study you used Elastic-net (ENET) models (Philippat et al., 2019 doi: 10.1289/EHP3523). When comparing results from LASSO and ExWAS, it appears that LASSO was too restrictive. Would it be possible to additionally implement ENET and compare performance with LASSO?

Authors' Response: In this study, we did not implement the ENET procedure due to the lack of ready-to-use tools (neither in R or STATA) to apply ENET on multiply imputed data where the outcome variables (in our case SDQ scores) follow a negative binomial distribution. Obtaining the meaningful estimates for multiply imputed data directly from the penalized regression model would require a method to aggregate the coefficients of selected variables for each imputed dataset and to assign a standard error for the estimates of variables that were not selected. These operations are not straightforward and, to our best knowledge, no tools for performing them on data with negative binomial outcome exist. We did not

encounter such issues for the paper cited by the Reviewer (Philippat et al., 2019), since analyses were based on the non-imputed data and normally distributed outcome.

While, as described above, obtaining meaningful ENET coefficients is not straightforward when relying on multiple imputed datasets and non-gaussian outcome, ENET can still be used to select variables that can then be fitted into a new, non-penalized multiple regression model which produces the final estimates. For multiply imputed data this can be achieved by applying the same method as we used for LASSO, i.e. by fitting the ENET on each of the 100 imputed datasets and retaining an exposure only if it was selected in at least 50% of runs. Below we present the results of the variable selection using ENET, with the optimal α chosen by maximizing the prediction log-likelihood using 10-fold cross-validation and the corresponding λ +1SE. In line with the result obtained for the LASSO, the ENET algorithm did not retain any biomarker for the SDQ internalising score (Figure 1). For the externalising score, the ENET selected copper, a biomarker also selected by the LASSO, as well as two other biomarkers (cobalt and dichlorodiphenyltrichloroethane) that have not been identified by LASSO.



Revision Figure 1: Percentage of times each biomarker was selected by the ENET algorithm within 100 runs (one per imputed dataset). Red dashed line represents the threshold of 50% of runs. Models were adjusted for cohort, season of conception, child sex and age at SDQ assessment, parity, and maternal factors: education level, work status, age, pre- pregnancy BMI, and prenatal active smoking status.

Effect estimates obtained by fitting variables selected by ENET to a non-penalized multiple regression model are shown in Table 1 below. In this model, only copper was significantly associated with the externalising behavioural score and its estimates and CIs were almost identical as those obtained for the ExWAS. Since the results obtained with ENET were very similar to those for LASSO and ExWAS, we decided not to include this analysis in the manuscript.

Revision Table 1: Adjusted associations between prenatal exposure to environmental contaminants and SDQ externalising score selected by ENET (n = 708).

Behavioural		Main ExW	45	Multiple non-penalized		
		Main Ex VV	10	regression model		
outcome	Exposures included in the model	IRR (95%CI)	p value	IRR (95%CI)	p value	
SDQ externalising score	Cobalt (Co)	1.10 (0.98; 1.24)	0.112	1.11 (0.98; 1.25)	0.094	
	Copper (Cu)	0.90 (0.82; 0.98)	0.021	0.90 (0.82; 0.99)	0.024	
	Dichlorodiphenyltrichloroethane (DDT)	0.92 (0.84; 1.00)	0.045	0.93 (0.85; 1.01)	0.075	

The multiple regression model was adjusted for cohort, season of conception, child sex and age at SDQ assessment, parity, and maternal factors: education level, work status, age, pre- pregnancy BMI, and prenatal active smoking status. Abbreviations: BMI = body mass index. CI = confidence interval of the IRR estimate. ExWAS = exposome-wide association study. IRR = incident rate ratio. SDQ = Strengths and Difficulties Questionnaire.

Reviewer's comment 2: Results section 3.2.2. Internalizing score. In ExWAS, the authors selected a threshold of p<0.05 to select associations. While I understand this approach, the authors face one ironical finding in their analysis: For MnBP-externalizing, the p-value was 0.048, and the association was selected. For DETP-internalizing, the p-value was 0.053, and this association has not even mentioned in the manuscript. If one checks if this potential association is driven by a unique cohort, then finds that there is a consistent pattern towards more internalizing problems in four out of the five cohorts examined: BiB 2.04 (1.33;3.12); EDEN 1.12 (0.91;1.36); INMA 0.98 (0.81;1.18); KANC 1.15 (0.78;1.67) and RHEA 1.11 (0.92;1.33). Therefore, I think the DETP association should be acknowledged in both the results section and the abstract, and highlight that future studies need to confirm or ruled out this association.

Authors' Response: As suggested by the Reviewer, the association between DETP and SDQ internalizing score is now mentioned in the abstract, results, discussion and Table 2.

Reviewer's comment 3: Following the previous example, I think the authors could maintain the p-threshold at <0.05, but also add that the direction and pattern of associations across the five cohorts was also accounted in the selection of the most relevant associations. Thus, this study identified patterns of

associations that, although not significant at the conventional level, would be worth it to highlight as candidate associations requiring further confirmation. Examples are: cadmium and cobalt for externalizing problems; and cobalt, manganese, BUPA and TRCS for internalizing problems.

Authors' Response: We agree that the consistency of the results across the cohorts is an important criterion to select the most relevant associations between prenatal exposures and SDQ scores. However, we do not find it feasible to rely on the direction and pattern of association across the cohorts to select the associations to be mentioned in the result section. Effect estimates obtained for each cohort for the biomarkers mentioned by the Reviewer had wide confidence intervals that most often included 1, which made it difficult to define the direction and pattern of those associations (see Table 2 below). Moreover, apart from the mentioned issues, the association between Co and the SDQ internalising score mentioned by the Reviewer showed a moderate heterogeneity across the cohorts ($I^2 = 0.42$) pointing rather towards the inconsistency of the results rather than any common pattern. For these reasons we decided to not expand the list of associations mentioned in the results section.

Revision Table 2: Adjusted associations between prenatal exposure to environmental contaminants suggested by the Reviewer and SDQ externalising (upper panel) and internalising scores (lower panel), extracted from the Appendix Table 6 and 7 (n = 708).

	ExWAS			ExWAS with cohort-exposure interaction ^a				
Exposure	IRR (95%CI)							\mathbf{I}^2
	IRR (95%CI)	p value	BiB	EDEN	INMA	KANC	RHEA	
Cadmium (Cd)	1.08 (0.97; 1.20)	0.182	1.17 (0.76;1.80)	1.11 (0.92;1.33)	1.03 (0.88;1.22)	1.09 (0.92;1.29)	1.12 (0.97;1.29)	< 0.001
Cobalt (Co)	1.10 (0.98; 1.24)	0.112	1.21 (0.85;1.73)	1.21 (0.99;1.47)	1.04 (0.86;1.26)	0.99 (0.75;1.30)	1.09 (0.92;1.30)	< 0.001

	ExWAS			ExWAS with cohort-exposure interaction ^a				
Exposure					IRR (95%CI)			\mathbf{I}^2
	IRR (95%CI)	p value	BiB	EDEN	INMA	KANC	RHEA	
Cobalt (Co)	1.10 (0.96; 1.27)	0.179	1.50 (1.02;2.21)	1.06 (0.85;1.33)	1.08 (0.84;1.38)	0.77 (0.54;1.09)	1.23 (0.98;1.55)	0.420
Manganese (Mn)	1.11 (0.98; 1.26)	0.099	1.07 (0.70;1.63)	1.07 (0.87;1.32)	1.12 (0.87;1.44)	1.03 (0.76;1.40)	1.19 (0.98;1.44)	< 0.001
N-butyl paraben (BUPA)	1.09 (0.98; 1.21)	0.121	1.24 (0.89;1.72)	1.03 (0.83;1.27)	1.24 (1.02;1.51)	0.99 (0.67;1.46)	1.00 (0.82;1.21)	0.111
Triclosan (TRCS)	1.08 (0.96; 1.21)	0.204	1.83 (1.15;2.90)	1.12 (0.91;1.39)	1.00 (0.81;1.25)	1.04 (0.69;1.55)	1.02 (0.82;1.25)	< 0.001

The regression models were adjusted for cohort, season of conception, child sex and age at SDQ assessment, parity, and maternal factors: education level, work status, age, pre- pregnancy BMI, and prenatal active smoking status. Abbreviations: BMI = body mass index. CI = confidence interval of the IRR estimate. ExWAS = exposome-wide association study. IRR = incident rate ratio. SDQ = Strengths and Difficulties Questionnaire. BiB = Born in Bradford. EDEN = Étude des Déterminants Pré et Postnatals du Développement et de la Santé de l'Enfant. INMA = Infancia y Medio Ambiente. KANC = Kaunas Cohort. RHEA = Mother-Child Cohort in Crete.

Reviewer's comment 4: Discussion lines 274-277: Good point with the residual confounding advantage of pooling the five cohorts. Just after this sentence, or in any other part, you should also acknowledge that the variability among cohorts and procedures also presents limitations that would probably tend to introduce some random noise and bias toward the null. Could you briefly elaborate this part? For example, if BPA or MnBP was measured by different labs, without an interlaboratory comparison, this will tend to increase variability in exposure-SDQ associations. If urine samples were collected at different trimesters in each cohort, this will also add variability due to normal dilution effects, etc.

Authors' Response: The Reviewer is right. Since the cohorts were recruited before the start of the HELIX project, collection of biological samples during pregnancy was not harmonized leading to different timings (i.e., different trimester) for exposure assessment across cohorts. It is also true that, for some cohorts, the same biomarker was sometimes assessed in different laboratories. However, the results of interlaboratory comparisons performed in the framework of the HELIX protocol show a high correlation between assessments performed by different laboratories (see Table 3 below). For instance, phenol metabolites were assessed by both the Norwegian Institute of Public Health (NIPH) and the Centers for Disease Control and Prevention (CDC) in 35 urine samples of the EDEN cohort. Correlation coefficients between these two measurements were high (≥ 0.90, see Table 3 below). Results for all interlaboratory comparisons performed within the Helix framework can be found in the Supplementary material of (Tamayo-Uria et al. 2019), pp. 16-19.

<u>Revision Table 3</u>: Spearman correlation coefficients between 12 phenol urinary maternal samples analysed in the NIPH and CDC laboratories.

	NIPH						
CDC	MEPA	ETPA	PRPA	BUPA	BPA	OXBE	TRCS
MEPA	1.0	-	-	-	-	-	-
ETPA	-	0.98	-	-	-	-	-
PRPA	-	-	0.90	-	-	-	-
BUPA	-	-	-	1.0	-	-	-
BPA	-	-	-	-	0.90	-	-
OXBE	-	-	-	-	-	0.91	-
TRCS	-	-	-	-	-	-	0.99

Abbreviations: BPA = bisphenol A. BUPA = n-butyl paraben. ETPA = ethyl paraben. MEPA = methyl paraben. OXBE = oxybenzone. PRPA = propyl paraben. TRCS = triclosan. CDC = Centers for Disease Control and Prevention. NIPH = Norwegian Institute of Public Health.

All these issues are now mentioned in the discussion section: "On the other hand, since the cohorts were recruited before the start of the HELIX project, collection of biological samples during pregnancy was not harmonized leading to different timings (i.e., different trimester) for exposure assessment across

cohorts. Additionally, for some cohorts the same exposure biomarker was sometimes assessed by different laboratories (see Appendix Table 2), which may partly explain the between-cohort heterogeneity of the results observed for some exposures. This should not have a strong impact on our results since the interlaboratory comparisons performed in the framework of the HELIX protocol suggested a high correlation between assessments performed in different laboratories. For instance, correlation coefficients between phenol urinary concentrations measured by the Norwegian Institute of Public Health and Centers for Disease Control and Prevention in 12 maternal samples of the EDEN cohort were ≥ 0.90 (Supplementary of (Tamayo-Uria et al. 2019)). ".

General comments:

Reviewer's comment: Please check the manuscript for blank spaces throughout the text.

Authors' Response: We have revised the manuscript for blank spaces. As they are not present in the .docx document, we assume these were artefacts of PDF conversion. We will make sure they will not appear in the revised version of the manuscript.

Reviewer's comment: Check Appendix Tables 5 and 6, and complete estimates, so all appear with two decimals, even if it is for example 1.00 instead of 1, or 0.90 instead of 0.9, to be consistent throughout the table.

Authors' Response: We have corrected the tables accordingly.

Reviewer's comment: Highlights: Copper and POPs findings are not mentioned.

Authors' Response: We have mentioned the association between copper and SDQ score in the highlights: "Copper was associated with decreased behavioural scores". However, we decided to not mention the POPs findings because they were mainly seen among women with insufficient weight gain during pregnancy and were not consistent with the literature.

Reviewer's comment: Abstract: Line 22 "The association with PFOS" instead of "An".

Authors' Response: We have corrected the sentence.

Reviewer's comment: Introduction: I like the introduction, concise and informative.

Authors' Response: Thank you.

Reviewer's comment: Line 69: In a previous STOTEN paper, Kim et al., analyzed different chemical families in relation to child neurodevelopment:

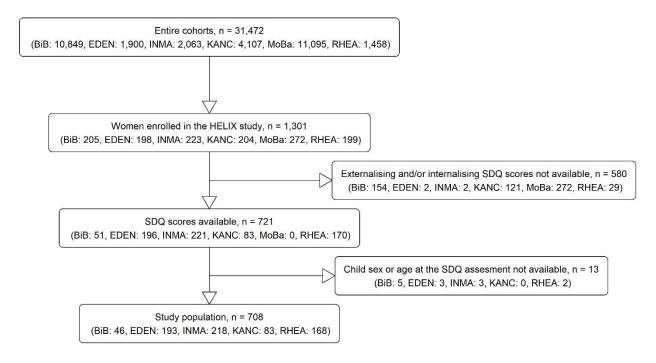
Kim et al., 2018. Association between maternal exposure to major phthalates, heavy metals, and persistent organic pollutants, and the neurodevelopmental performances of their children at 1 to 2 years of age-CHECK cohort study. Sci Total Environ. doi: 10.1016/j.scitotenv.2017.12.058.

Authors' Response: We have updated the reference.

Methods

Reviewer's comment: 2.1 Study design and population: A flow-chart explaining the inclusion of mother-child couples from each cohort would be helpful.

Authors' Response: We have added a flow-chart to the appendix (Appendix Figure 1). See also below.



Appendix Figure 1: Study flow-chart. Abbreviations: BiB = Born in Bradford. EDEN = Étude des Déterminants Pré et Postnatals du Développement et de la Santé de l'Enfant. INMA = Infancia y Medio Ambiente. KANC = Kaunas Cohort. MoBa = Norwegian Mother, Father and Child Cohort Study. RHEA = Mother-Child Cohort in Crete. SDQ = Strengths and Difficulties Questionnaire. Adapted from (Maitre et al. 2018).

Reviewer's comment: Line 85: Strenght[s] and Difficulties Questionnaire [delete "s"].

Authors' Response: We have corrected the sentence.

Reviewer's comment: Lines 84-85: you mention child twice in the same sentence.

Authors' Response: We have removed the repeated word.

Reviewer's comment: 2.2 Exposure assessment: Many different chemicals were measured. In lines 96 and 97 the authors cite two papers, but this seems insufficient to described the methodologies, laboratories and quality controls followed for all chemicals and metabolites. Was each chemical family assessed by the same lab in all the cohorts? Interlaboratory comparisons were available? In the supplementary material, a table providing brief but relevant information for each chemical compound would be helpful. For example, it would be nice to know the laboratory in which the sample was analyzed in each cohort, the method (HPLC, GCMS...), the exact biological sample used in each cohort (e.g. for PFAS, while serum/plasma can be compared, PFAS measured in whole blood would probably lead to a reduction in PFAS concentrations). This information can help to decide to which extent cohorts and measurements are comparable, and to perform a more informative interpretation of study findings in light of the variability observed among cohorts.

Authors' Response: We have added a table with the requested information to the appendix (Appendix Table 2). See also below.

Appendix Table 1: Exposure assessment – biological matrices, timing of sample collection, laboratories, methodologies and quality controls (adapted from (Haug et al. 2018)).

T. II. 6	W-170	nn	Cohort	NV 5 :	DVIII :
Family of exposures	BiB	EDEN	KANC	INMA	RHEA
OCs					
Biological matrix	Serum/plasma	Serum	-	Serum	Serum
Timing of sample collection, mean GW (SD)	26.6 (1.4)	26.1 (1.2)	13.7 (2.0)	39.4 (1.3)	14.1 (3.7)
Laboratory	NIPH (Norway)	NIPH (Norway)	NIPH (Norway)	LSPG (Spain)	National Institute for Health and Welfare, Chemical Exposure Unit, Kuopio (Finland)
Analytical method	GC-MS/MS	GC-MS/MS	GC-MS/MS	GC-MS	GC-MS/MS
Interlaboratory comparison	samples are spiked, and most ar		ation range than our samples usual accept for PCB-153, and for PCB-17	ly are. Despite these factors Z- so 70 in 1 sample.	cores below 2 were obtained for
Standard reference material	the non-fortified sample (195	from NIST, Organic Contaminants in 37) were 2-18%. The deviation from the m. The RSDs of the 6 different injection from the certifie	he certified values was 2-20%, exc	cept for HCB (50%) and PCB-153) were 2-20%, and for most analy	3 (76%), where a small blank
Reference for the analytic method	Modified (Caspersen et al. 2016) Modified (Caspersen et al. 2016)	Modified (Caspersen et al. 2016)	(Goñi et al. 2007)	(Koponen et al. 2013)
PBDEs					
Biological matrix	Serum/plasma	Serum	-	Serum	Serum
Timing of sample collection, mean GW (SD)	26.6 (1.4)	26.1 (1.2)	13.7 (2.0)	39.4 (1.3)	14.1 (3.7)
Laboratory	NIPH (Norway)	NIPH (Norway)	NIPH (Norway)	NIPH (Norway)	National Institute for Health and Welfare, Chemical Exposure Unit, Kuopio (Finland)
Analytical method	GC-MS/MS	GC-MS/MS	GC-MS/MS	GC-MS	GC-MS/MS
Interlaboratory comparison		AMAP interlaboratory comparison s nalytes are in a much higher concentra			
Standard reference material		from NIST, Organic Contaminants in) were 2-18%. The deviation from the were 2-20%, and for most analytes	e certified values was 2-20%. The l	RSDs of the 6 different injections	
Reference for the analytic method	Modified (Caspersen et al. 2016) Modified (Caspersen et al. 2016)	Modified (Caspersen et al. 2016)	Modified (Caspersen et al. 2016)	(Koponen et al. 2013)
PFASs					
Biological matrix	Serum/plasma	Serum	Whole blood	Plasma	Serum
Timing of sample collection, mean GW (SD)	26.6 (1.4)	26.1 (1.2)	13.7 (2.0)	39.4 (1.3)	14.1 (3.7)

Laboratory	NIPH (Norway)	NIPH (Norway)	NIPH (Norway)	Institute for Occupational Medicine, RWTH Aachen University (Germany)	NIPH (Norway)					
Analytical method	GC-MS	GC-MS	GC-MS	Online column-switching LC– MS–MS analysis	GC-MS					
Interlaboratory comparison	We participated 3 times in the AMAP interlaboratory comparison study during the period where HELIX samples were analysed. Each round included 3 samples samples are spiked in a wide concentration range. Z- scores below 2 were obtained for all contaminants except for PFUnDA in 2 samples.									
Standard reference material	In total 15 samples from 4 roun	In total 15 samples from 4 rounds of the AMAP interlaboratory comparison study were analysed during the period where HELIX samples were analysed, and the mean deviation from the assigned value varied between 8 and 17%.								
Reference for the analytic method	(Haug et al. 2009)	(Haug et al. 2009)	(Poothong et al. 2017)	(Manzano-Salgado et al. 2015)	(Haug et al. 2009)					
Metals and semi-metals										
Biological matrix	Whole blood	Whole blood	Whole blood	Cord whole blood	Whole blood					
Timing of sample collection, mean GW (SD)	26.6 (1.4)	26.1 (1.2)	13.7 (2.0)	39.4 (1.3)	14.1 (3.7)					
Laboratory	ALS Scandinavia (Sweden)	ALS Scandinavia (Sweden)	ALS Scandinavia (Sweden)	Hg; LSPA (Spain)	ALS Scandinavia (Sweden)					
Analytical method	ICP-SFMS	ICP-SFMS	ICP-SFMS	AAS	ICP-SFMS					
Standard reference material		rm whole blood reference materials were blinded for the laboratory. The								
Reference for the analytic method	(Rodushkin et al. 2000)	(Rodushkin et al. 2000)	(Rodushkin et al. 2000)	(Ramon et al. 2011)	(Rodushkin et al. 2000)					
Phthalate metabolites										
Phthalate metabolites Biological matrix	Urine	Urine	-	Urine	Urine					
	Urine 26.6 (1.4)	Urine 26.1 (1.2)	34.2 (1.3)	Urine NA	Urine 14.1 (3.7)					
Biological matrix Timing of sample collection,			- 34.2 (1.3) NIPH (Norway)							
Biological matrix Timing of sample collection, mean GW (SD)	26.6 (1.4)	26.1 (1.2)	. ,	NA Bioanalysis Research Group at the Hospital del Mar Medical	14.1 (3.7)					
Biological matrix Timing of sample collection, mean GW (SD) Laboratory	26.6 (1.4) NIPH (Norway) LC-MS/MS	26.1 (1.2) NIPH (Norway) LC-MS/MS reference material SRM 3673 were ar	NIPH (Norway) LC-MS/MS	NA Bioanalysis Research Group at the Hospital del Mar Medical Research Institute (Spain) HPLC-MS HELIX samples were analysed, and	14.1 (3.7) NIPH (Norway) LC-MS/MS					
Biological matrix Timing of sample collection, mean GW (SD) Laboratory Analytical method	26.6 (1.4) NIPH (Norway) LC-MS/MS	26.1 (1.2) NIPH (Norway) LC-MS/MS reference material SRM 3673 were ar	NIPH (Norway) LC-MS/MS nalysed during the period where I	NA Bioanalysis Research Group at the Hospital del Mar Medical Research Institute (Spain) HPLC-MS HELIX samples were analysed, and	14.1 (3.7) NIPH (Norway) LC-MS/MS					
Biological matrix Timing of sample collection, mean GW (SD) Laboratory Analytical method Standard reference material Reference for the analytic method Phenols	26.6 (1.4) NIPH (Norway) LC-MS/MS In total 42 samples of NIST r (Sabaredzovic et al. 2015)	26.1 (1.2) NIPH (Norway) LC-MS/MS reference material SRM 3673 were ar assign (Sabaredzovic et al. 2015)	NIPH (Norway) LC-MS/MS nalysed during the period where I ned value varied between 1 and 2	NA Bioanalysis Research Group at the Hospital del Mar Medical Research Institute (Spain) HPLC-MS HELIX samples were analysed, and 5%. (Valvi et al. 2015)	14.1 (3.7) NIPH (Norway) LC-MS/MS the mean deviation from the (Sabaredzovic et al. 2015)					
Biological matrix Timing of sample collection, mean GW (SD) Laboratory Analytical method Standard reference material Reference for the analytic method	26.6 (1.4) NIPH (Norway) LC-MS/MS In total 42 samples of NIST r	26.1 (1.2) NIPH (Norway) LC-MS/MS reference material SRM 3673 were ar assign	NIPH (Norway) LC-MS/MS nalysed during the period where I ned value varied between 1 and 2	NA Bioanalysis Research Group at the Hospital del Mar Medical Research Institute (Spain) HPLC-MS HELIX samples were analysed, and 5%.	14.1 (3.7) NIPH (Norway) LC-MS/MS the mean deviation from the					
Biological matrix Timing of sample collection, mean GW (SD) Laboratory Analytical method Standard reference material Reference for the analytic method Phenols	26.6 (1.4) NIPH (Norway) LC-MS/MS In total 42 samples of NIST r (Sabaredzovic et al. 2015)	26.1 (1.2) NIPH (Norway) LC-MS/MS reference material SRM 3673 were ar assign (Sabaredzovic et al. 2015)	NIPH (Norway) LC-MS/MS nalysed during the period where I ned value varied between 1 and 2	NA Bioanalysis Research Group at the Hospital del Mar Medical Research Institute (Spain) HPLC-MS HELIX samples were analysed, and 5%. (Valvi et al. 2015)	14.1 (3.7) NIPH (Norway) LC-MS/MS the mean deviation from the (Sabaredzovic et al. 2015)					
Biological matrix Timing of sample collection, mean GW (SD) Laboratory Analytical method Standard reference material Reference for the analytic method Phenols Biological matrix Timing of sample collection,	26.6 (1.4) NIPH (Norway) LC-MS/MS In total 42 samples of NIST r (Sabaredzovic et al. 2015) Urine	26.1 (1.2) NIPH (Norway) LC-MS/MS reference material SRM 3673 were ar assign (Sabaredzovic et al. 2015) Urine	NIPH (Norway) LC-MS/MS nalysed during the period where I ned value varied between 1 and 2 (Sabaredzovic et al. 2015)	NA Bioanalysis Research Group at the Hospital del Mar Medical Research Institute (Spain) HPLC-MS HELIX samples were analysed, and 5%. (Valvi et al. 2015) Urine	14.1 (3.7) NIPH (Norway) LC-MS/MS the mean deviation from the (Sabaredzovic et al. 2015) Urine					

Interlaboratory comparison	We participated twice in the External Quality Assessment Scheme for organic substances in urine for BPA and TRCS. Each round included 2 samples. Z- scores below 2 were obtained for both contaminants in all samples.								
Standard reference material	In total 145 samples of NIST r	reference material SRM 3673 were a assigned value varie	analysed during the period where led between 1 and 19%, except BPA		d the mean deviation from the				
Reference for the analytic method	(Sakhi et al. 2018)	(Philippat et al. 2012)	(Sakhi et al. 2018)	(Sakhi et al. 2018)	(Sakhi et al. 2018)				
OP pesticide metabolites									
Biological matrix	Urine	Urine	-	Urine	Urine				
Timing of sample collection, mean GW (SD)	26.6 (1.4)	26.1 (1.2)	34.2 (1.3)	NA	14.1 (3.7)				
Laboratory	NIPH (Norway)	NIPH (Norway)	NIPH (Norway)	NIPH (Norway)	NIPH (Norway)				
Analytical method	UPLC-TOF	UPLC-TOF	UPLC-TOF	UPLC-TOF	UPLC-TOF				
Reference	(Cequier et al. 2016)	(Cequier et al. 2016)	(Cequier et al. 2016)	(Cequier et al. 2016)	(Cequier et al. 2016)				
Cotinine	-				_				
Biological matrix	Urine	Urine	-	Urine	Urine				
Timing of sample collection, mean GW (SD)	26.6 (1.4)	26.1 (1.2)	34.2 (1.3)	NA	14.1 (3.7)				
Laboratory	Fürst Medical Analysis Laboratory (Norway)	Fürst Medical Analysis Laboratory (Norway)	Fürst Medical Analysis Laboratory (Norway)	Public Health Laboratory of Bilbao - LSPPV (Spain)	Fürst Medical Analysis Laboratory (Norway)				
Analytical method	The Immulite® 2000 Nicotine Metabolite (Cotinine) 600 Test on an Immulite 2000 XPi from Siemens Healthineers	The Immulite® 2000 Nicotine Metabolite (Cotinine) 600 Test on an Immulite 2000 XPi from Siemens Healthineers	The Immulite® 2000 Nicotine Metabolite (Cotinine) 600 Test on an Immulite 2000 XPi from Siemens Healthineers	LC-MS	The Immulite® 2000 Nicotine Metabolite (Cotinine) 600 Test on an Immulite 2000 XPi from Siemens Healthineers				
Reference for the analytic method	-	-	-	(Aurrekoetxea et al. 2013)	-				
Creatinine									
Biological matrix	Urine	Urine	-	Urine	Urine				
Timing of sample collection, mean GW (SD)	26.6 (1.4)	26.1 (1.2)	34.2 (1.3)	NA	14.1 (3.7)				
Laboratory	Fürst Medical Analysis Laboratory (Norway)	National Center for Environmental Health laboratory at the CDC (US)	Fürst Medical Analysis Laboratory (Norway)	Public Health Echevarne Laboratory of Barcelona (Spain)	Fürst Medical Analysis Laboratory (Norway)				
Analytical method	AU680 Chemistry System form Beckman Coulter using DRI® Creatinine-Detect® Test	Enzymatic reaction using a Roche Hitachi 912 chemistry analyzer (Roche Hitachi, Basel, Switzerland)	AU680 Chemistry System form Beckman Coulter using DRI® Creatinine-Detect® Test	Jaffé method - Beckman Coulter© AU5400	AU680 Chemistry System form Beckman Coulter using DRI® Creatinine-Detect® Test				
Reference for the analytic method	-	-		-					
Lipids									
Biological matrix	Serum/plasma	Serum	-	Serum	Serum				
Timing of sample collection, mean GW (SD)	26.6 (1.4)	26.1 (1.2)	13.7 (2.0)	39.4 (1.3)	14.1 (3.7)				

Laboratory	Laboratory (Norway)	Laboratory (Norway)	Laboratory (Norway)	Bizkaia (Spain)	(Greece)
	ADVIA® Chemistry XPT	ADVIA® Chemistry XPT	ADVIA® Chemistry XPT	Cobas Mira self-analyzer (Roche	
Analystical mathod	System, and the FS kit from	System, and the FS kit from	System, and the FS kit from	Diagnostic, Basel, Switzerland)	Standard enzymatic method
Analytical method	DiaSys was used to measure	DiaSys was used to measure	DiaSys was used to measure	using an Enzymatic-Colorimetric	Standard enzymatic method
	concentrations of phospholipids	concentrations of phospholipids	concentrations of phospholipids	method with Spinreact reagents	
Reference for the analytic method	-	-	-	-	-

Abbreviations: GW = gestational week. NA = not available. SD = standard deviation. BPA = bisphenol A. DDT = dichlorodiphenyltrichloroethane. HCB = hexachlorobenzene. OC = organochlorine compound. OP = organophosphate. PBDE = polybrominated diphenyl ether. PCB = polychlorinated biphenyl. PFAS = per- and polyfluoroalkyl substance. PFUnDA = perfluoroundecanoate. TRCS = triclosan. RSD = relative standard deviation. SRM = Standard reference material. AAS = thermal decomposition, amalgamation and atomic absorption spectrometry. GC-MS/MS = gas chromatography coupled with tandem mass spectrometry. GC-MS = gas chromatography mass spectrometry. GC-MS-NICI = gas chromatography-negative-ion chemical ionization mass spectrometry. HPLC-MS = ultra-performance liquid chromatography coupled to tandem mass spectrometry. ICP-SFMS = conductively coupled plasma-sector field mass spectrometry. LC-MS/MS = liquid chromatography-mass spectrometry. Q-ICP-MS = inductively coupled plasma quadruple mass spectrometry. SPE-HPLC-MS/MS = on-line solid-phase extraction coupled to isotope dilution high performance liquid chromatography-tandem mass spectrometry. UPLC-MS/MS = ultra-performance liquid chromatography-tandem mass spectrometry. UPLC-MS/MS = ultra-performance liquid chromatography-tandem mass spectrometry. AMAP = Arctic Monitoring and Assessment Program "Ring Test for Persistent Organic Pollutants in Human Serum". CDC = Centers for Disease Control and Prevention. LSPA = Laboratorio de Salud Pública de Alava. LSPG = Laboratorio de Salud Pública de Guipúzcoa. NIPH = Norwegian Institute of Public Health. NIST = National Institute of Standards and Technology, USA. BiB = Born in Bradford. EDEN = Étude des Déterminants Pré et Postnatals du Développement et de la Santé de l'Enfant. INMA = Infancia y Medio Ambiente. KANC = Kaunas Cohort. RHEA = Mother-Child Cohort in Crete.

2.3 Behavioral outcomes

Reviewer's comment: Lines 101-105: Apart from this rationale, it should be included that this approach was also selected to reduce the number of multiple comparisons performed. Subscales also provide interesting and predictive information, but in this case, investigating 47 exposures and all subscales, would lead to a multiple comparison issue difficult to disentangle, and a huge amount of data difficult to present to the reader.

Authors' Response: We have updated the text accordingly: "Moreover, given our limited sample size and the large number of studied exposures, combining the SDQ sub-scales limited the number of performed tests."

2.4. Statistical analysis

Reviewer's comment: Lines 107-108: reference for this sentence?

Authors' Response: We have added a relevant reference.

Reviewer's comment: Lines 109-110: "haemal lipophilic biomarker concentrations by total fat percentage" It reads as if you normalized POPs by total body fat percentage. Probably you want to say that lipophilic biomarker concentrations were standardized by serum lipids and expressed as ng/g of lipid.

Authors' Response: We clarified: "Haemal lipophilic biomarker concentrations were standardized and expressed in ng/g of total lipids in serum or plasma."

Reviewer's comment: Why season of conception was included in the models? For specific biomarkers such as BP-3, season of sample collection would be relevant, but I do not totally see its role as a relevant confounder of the associations tested.

Authors' Response: Exposure to several environmental contaminants considered in our study is dependent on behaviours regulated by season, e.g. on time spent outdoors. That included exposure to some heavy metals (e.g. cadmium, lead and mercury that are common air pollutants (Tchounwou et al. 2012)) or substances present in sunscreens (e.g. oxybenzone also known as BP-3 (American Association of Pharmaceutical Scientists 2018)) or in other seasonal cosmetics (e.g. parabens). On the other hand, there is evidence on season of birth being associated with emotional and behavioural regulation in children (Asano et al. 2016) or risk of neurodevelopmental disorders (e.g. (Zerbo et al. 2011). For these reasons we decided to include the season of conception in the models.

Results

13

Reviewer's comment: Lines 162-165: Median externalizing score for BiB children was 0.5, in striking contrast with the rest of cohorts, in which the median was between 5 and 6. Based on this huge difference, what is your criteria to not exclude this cohort from the analysis at least in a sensitivity?

Authors' Response: We have provided an additional sensitivity analysis for the SDQ externalizing scores where we excluded the BiB cohort and compared the results with the main ExWAS. Results of this additional analysis are now included in the main text and in Table 2, see also below. Exclusion of the BiB cohort did not strongly affect our results for the SDQ externalising score, except of widening of the confidence interval and increase of p value for DDT.

<u>Revision Table 4</u>: Adjusted associations between the prenatal exposure to environmental contaminants and SDQ externalising scores, extracted from the Table 2.

Behavioural			ExWAS			ExWAS after exclusion of the BiB cohort		
outcome	Exposure	Exposure family			FWER			
			IRR (95%CI)	p value	p value	IRR (95%CI)	p value	n
	BPA	Phenol	1.06 (1.01; 1.12)	0.028	0.842	1.06 (1.01; 1.12)	0.026	662
	Cu	Essential element	0.90 (0.82; 0.98)	0.021	0.631	0.91 (0.83; 1.00)	0.042	662
SDQ	DDT	Organochlorine compound	0.92 (0.84; 1.00)	0.045	1	0.94 (0.86; 1.03)	0.174	662
externalising	MnBP	Phthalate	1.06 (1.00; 1.13)	0.048	1	1.06 (1.00; 1.13)	0.041	662
	PCB-138	Organochlorine compound	0.88 (0.79; 0.99)	0.035	1	0.88 (0.79; 0.99)	0.031	662
	PFUnDA	Perfluoroalkyl substance	0.92 (0.84; 0.99)	0.034	1	0.90 (0.82; 0.98)	0.013	662

The regression model was adjusted for cohort, season of conception, child sex and age at SDQ assessment, parity, and maternal factors: education level, work status, age, pre- pregnancy BMI, and prenatal active smoking status. Abbreviations: BiB = Born in Bradford. BMI = body mass index. CI = confidence interval of the IRR estimate. ExWAS = exposome-wide association study. FWER = family wise error rate adjustment. IRR = incident rate ratio. SDQ = Strengths and Difficulties. BPA = bisphenol A. Cu = copper. DDT = dichlorodiphenyltrichloroethane. MnBP = mono-n-butyl phthalate. PCB = polychlorinated biphenyl. PFUnDA = perfluoroundecanoate.

Reviewer's comment: Lines 195-197: Appendix Figure 1 is informative. It could be included in the main manuscript. In that case change "Addequate to Adequate".

Authors' Response: The figure was moved to the main manuscript with corrected legend.

Reviewer's comment: Table 1: Maternal work status. Numbers in the "overall distribution" column are interchanged between "unemployed" and "employed".

Authors' Response: We have corrected the numbers accordingly.

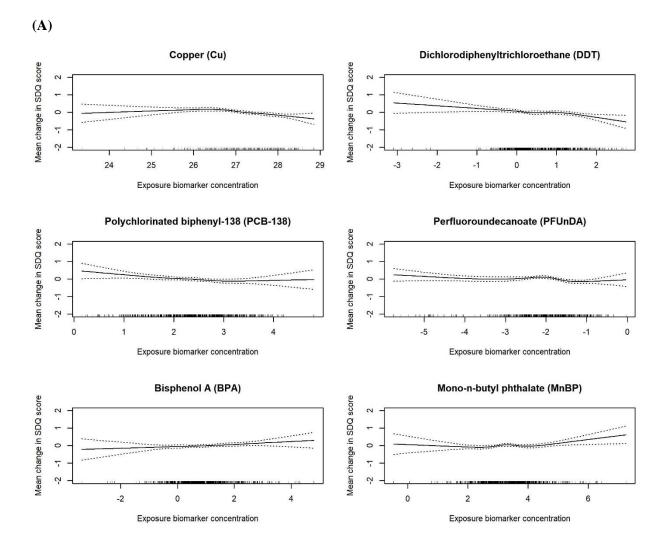
Reviewer's comment: Figure 1 footnote: Could you include the sample size for each of the cohorts? Or in any other place?

Authors' Response: We have added the requested information in the Figure 1 footnote.

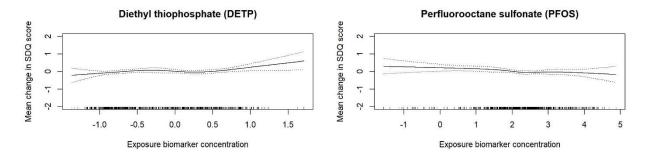
Discussion

Reviewer's comment: Lines 213-214: Have you checked the functional shape of the Cu association using for example LOWESS or GAMs? For essential metals, as you have stated, too much or too little would cause problems. Thus, a monotonic dose-response is not expected.

Authors' Response: We now use the GAM with the restricted cubic splines function fitted on the biomarker concentrations that were detected in the LASSO and ExWAS as associated with the SDQ scores. Results are presented in the Appendix Figure 2, see also below. All detected exposure-SDQ score associations were linear.



(B)



Appendix Figure 2: GAMs with restricted cubic splines function fitted on the log₂ and IQR transformed prenatal concentrations of exposures selected by the LASSO and ExWAS as associated with the externalising (A) and internalising (B) SDQ scores. Solid line represents the fit and dashed line the standard error of the fit. The model was adjusted for cohort, season of conception, child sex and age at SDQ assessment, parity, and maternal factors: education level, work status, age, pre- pregnancy BMI, and prenatal active smoking status. Abbreviations: GAM = generalized additive model. ExWAS = exposomewide association study. IQR = inter-quartile range. LASSO = least absolute shrinkage and selection operator. SDQ = Strengths and Difficulties Questionnaire.

Reviewer's comment: Lines 225-239: I agree with this paragraph. However, I think the authors should first mention that prenatal BPA has been associated with both externalizing and internalizing behaviors in previous studies (updated review Table 1 Mustieles and Fernández, 2020). Indeed, results from this study support that BPA may be related to both internalizing and externalizing problems, although externalizing problems were more evident. In Appendix Table 6, for internalizing problems, BPA showed the following overall estimate: 1.04 (0.98, 1.12) p=0.209. Additionally, four out of five cohorts showed positive estimators: BiB 1.14 (0.83;1.55); EDEN 1.16 (0.98;1.36); INMA 1.02 (0.93;1.13); KANC 0.98 (0.78;1.23) and RHEA 1.04 (0.92;1.19). Given that the BPA-behavior relationship is one of the most studied, it would be worth it to mention this trend.

Authors' Response: In the discussion section we have now acknowledged the previous studies that had reported associations with the SDQ internalizing score and we have mentioned that the effect estimate for this association was also positive in our study population: "Previous studies also reported higher scores on the internalizing behaviour sub-scale in association with the prenatal exposure to bisphenol A (Braun et al. 2011, 2017; Evans et al. 2014; Grohs et al. 2019; Harley et al. 2013; Li et al. 2020; Perera et al. 2012, 2016; Philippat et al. 2017; Roen et al. 2015). While not significant (p value = 0.21), effect estimate for our study population also suggested a positive association between BPA and internalizing SDQ score (IRR: 1.04 (95%CI: 0.98, 1.12))."

Reviewer's comment: Lines 237-239: The experimental and epidemiologic data for BPA and behavior has been updated in Mustieles and Fernández 2020. Additionally, you can also cite Nesan et al., 2018 for

experimental studies.

Mustieles V, Fernández MF. Bisphenol A shapes children's brain and behavior: towards an integrated neurotoxicity assessment including human data. Environ Health. 2020 Jun 9;19(1):66. doi: 10.1186/s12940-020-00620-y.

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Authors' Response: We have updated the references according to the Reviewer's request.

Conclusions

Reviewer's comment: Lines 307-308: "These negative associations must be interpreted with caution as they were not reported in previous studies, [and could even result from changes in body composition during pregnancy]."

Authors' Response: We have corrected the sentence accordingly: "Cu, DDT, PCB-138, PFOS and PFUnDA were negatively associated with child behaviour scores, suggesting lower risk of behavioural problems. These negative associations were not reported in previous studies and for lipophilic compounds (DDT and PCB-138) could even result from changes in body composition during pregnancy."

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