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1 **Estimated PCDD/F TEQ and total TEQ concentrations in the serum of 7–10 year old Finnish**
2 **children**

3

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21

22 **Abstract**

23 Polychlorinated dibenzo-p-dioxins and furans (PCDD/Fs) and polychlorinated biphenyls (PCBs) are
24 persistent organic pollutants that have detrimental health effects. As people are exposed to them
25 mainly through the diet, EU has set maximum food dioxin and PCBs levels. EFSA CONTAM Panel
26 made new risk assessment in 2018 that lowered the tolerable weekly intake (TWI) from 14 pg-
27 TEQ/kg bw/week to 2 pg-TEQ/kg bw/week. Critical effect was decreased semen count at the age of
28 18-19 years if serum total TEQ at the age of 9 years exceeded the No Observed Adverse Effect
29 Level (NOAEL) of 7 pg/g lipid. However, it is largely unknown to what extent NOAEL is exceed
30 in European boys currently. We thus measured PCBs from small volume of serum in 184 Finnish
31 children 7-10 years of age. To estimate the TEQ levels of children from measured PCB levels, we
32 used our existing human milk PCDD/F and PCB concentrations to create a hierarchical Bayesian
33 regression model that was used to estimate TEQs from measured PCBs. For quality control (QC),
34 three pooled blood samples from 18 – 20 year old males were measured for PCDD/Fs and PCBs,
35 and estimated for TEQs. In QC samples measured and estimated TEQs agreed within 84% – 106%.
36 In our estimate for 7-10 year old children, PCDD/F TEQ exceeded NOAEL only in 0.5% and total
37 TEQ in 2.7% of subjects. Risk management following the decreased TWI proposed by the
38 CONTAM Panel should be carefully considered if total TEQ in children is already largely below
39 the NOAEL.

40

41

42 **Highlights:**

- 43 • Critical Russian study generated a NOAEL of 7 pg total TEQ/g lipid in 9 year old boys
- 44 • Nine major PCBs were measured in serum of 184 Finnish children 7–10 years of age
- 45 • Bayes model was used to estimate serum total TEQ concentrations with PCBs
- 46 • The model predicted that total TEQ was below the NOAEL in 97.3% of children
- 47 • Little health benefits are achieved with further reduction in food maximum levels

48

49 **Keywords**

50 PCDD/Fs, PCBs, impaired semen quality, tolerable weekly intake, European Food Safety Authority

51 (EFSA)

52 **1. Introduction**

53 Polychlorinated dibenzo-*p*-dioxins and furans (PCDD/Fs) and polychlorinated biphenyls (PCBs) are
54 persistent organic pollutants that accumulate in food chain. People are exposed to them mainly
55 through diet. Fish and seafood are typically the main contributors to dietary PCDD/F and PCB
56 exposure (EFSA, 2012). In addition, the intake of red and white meat, and lean and fatty fish, has
57 been linked to high levels of PCDD/Fs and dioxin-like compounds in maternal blood of European
58 pregnant women (Papadopoulou et al., 2014). In the latest Finnish intake assessment of PCDD/Fs
59 and PCBs conducted 20 years ago the contribution of fish to their intake was around 80% (Kiviranta
60 et al., 2004). In Finland, despite higher intake of PCDD/Fs and PCBs, high consumption of fish has
61 been linked to many beneficial health effects such as lower mortality from all causes and ischaemic
62 heart diseases (Turunen et al., 2008), hypotriglyceridemic and anti-inflammatory effects, improved
63 glucose-insulin metabolism and arterial elasticity (Turunen et al., 2013). A recent review indicates
64 that the benefits of diets providing moderate amounts of fish during pregnancy outweigh potential
65 detrimental effects of environmental contaminants in regards to offspring neurodevelopment
66 (Starling et al., 2015).

67 On the other hand, both compound groups have various detrimental health effects most of which are
68 mediated through the aryl hydrocarbon receptor (AhR). Toxic equivalency factors (TEFs) have
69 been set based on experimental evidence for 17 PCDD/F congeners and 12 dioxin-like PCB
70 congeners with respect to their potency to induce toxic or biological effects through the AhR. The
71 TEF value of the most toxic congener, TCDD, has been set to 1, and the TEF values of other
72 congeners to 0.00003–1. Multiplication of the measured amount of each congener by its respective
73 TEF value gives the amount that is toxicologically equivalent with TCDD. Summing up the TEF-
74 weighted amounts of all congeners in a mixture gives the approximate amount that is equivalent to
75 TCDD in toxicity (toxic equivalent sum, TEQ) (Van den Berg et al., 2006).

76 In 2001, the Scientific Committee on Food of the European Commission established a Tolerable
77 Weekly Intake (TWI) of 14 pg TEQ/kg bw/week for PCDD/Fs and PCBs based on detrimental
78 effects on male reproduction, immune system, cognitive recognition and development of
79 endometriosis in experimental animals (SCF, 2001). In November 2018 EFSA published a new
80 report of the Contaminant Panel (CONTAM Panel) “Risk for animal and human health related to
81 the presence of dioxins and dioxin-like PCBs in feed and food” (EFSA, 2018). In the critical
82 Russian study of EFSA risk assessment, increased serum TCDD and PCDD TEQ concentrations at
83 age of 8–9 years were associated with a decreased sperm concentration, total sperm count, and total
84 motile sperm count at the age of 18–19 years (Minguez-Alarcon et al., 2017). From this study, a No
85 Observed Adverse Effect Level (NOAEL) of 7.0 pg PCDD/F TEQ/g lipid in blood sampled at age
86 of 9 years was identified. This was based on the median serum PCDD/F TEQ level in the lowest
87 quartile of the critical Russian study. The CONTAM panel estimated by pharmacokinetic modelling
88 that boys breastfed for 12 months with 800 ml/day of milk containing 5.9 pg TEQ/g lipid followed
89 by weekly intake of 4 pg TEQ/kg bw/week for 8 years will result in the NOAEL serum level of 7.0
90 pg/g lipid by the age 9 years (EFSA, 2018).

91 Overall, the CONTAM Panel proposed an adult TWI of 2 pg TEQ/kg bw/week for the sum
92 PCDD/Fs and PCBs (EFSA, 2018). This is a drastic decrease from the previous TWI of 14 pg
93 TEQ/kg bw/week, and according to intake calculations in the EFSA report, a substantial portion of
94 the European population currently exceeds the proposed TWI. From a risk management perspective,
95 this might require lowering the maximum levels (European Commission, 2011) of PCDD/Fs and
96 PCBs especially in fish, which is the most important source of intake in most European countries.

97 Few data exists on the blood TEQ-levels in European pre-pubertal boys, and it is largely unknown
98 to what extent they currently exceed the NOAEL of 7.0 pg TEQ/g lipid. The required sample
99 volume for PCDD/F and non-ortho-PCB analysis is 50–100 ml of blood, and there are ethical and
100 practical issues that make it virtually impossible to obtain such samples. In Germany, in a pooled

101 blood sample of 10 year old boys (n=241) in 2002–2003, PCDD/F TEQ was 5.9 pg/g lipid and PCB
102 TEQ 5.6 pg/g lipid (Link et al., 2005). Levels in girls of the same age (pooled sample, n=223) were
103 almost identical. In the Netherlands, individual samples from 14 boys (median age 14.3 years)
104 collected in 2005 had lower median PCDD/F TEQ of 2.3 pg/g lipid (range 0.4–6.1 pg/g lipid) and
105 PCB TEQ of 1.5 pg/g lipid (range 0.04–7.8 pg/g lipid) (Leijts et al., 2008).

106 The main purpose of this study was to estimate the serum PCDD/F- and total TEQ (sum of PCDD/F
107 TEQ and PCB-TEQ) levels among 7–10 year old Finnish boys in two cohorts, and compare those to
108 the NOAEL of 7.0 pg TEQ/g lipid. To increase the sample size, girls of the same age from these
109 cohorts were also included in the dataset. As the direct analysis of PCDD/Fs and dioxin-like PCBs
110 (DL-PCBs) was not possible due to limited serum volume available, TEQ-levels were estimated
111 based on the levels of 9 most prevalent PCB congeners, which could be detected from a much
112 smaller serum volume (200 µl). For this estimation, we created a hierarchical Bayesian regression
113 model that was based on our previous PCDD/F and PCB data from 535 human milk samples
114 collected in 2000, 2005 and 2010 from 3 Finnish cities. For quality control of the selected approach,
115 we measured PCDD/Fs and PCBs from pooled blood samples of young male blood donors and
116 compared measured TEQs with pools' estimated TEQs.

117

118 **2. Materials and methods**

119 *2.1. Serum samples, human milk samples and pooled blood samples*

120 Serum samples (200 µl) of pre-pubertal boys and girls were collected from two cohorts. Details of
121 the samples are given in Table 1. In the MULTIEPIGEN study
122 (<https://cordis.europa.eu/project/id/742927>), serum samples were collected in April–August 2018,
123 from 12 boys and 24 girls 7–9 years old from comprehensive geographical coverage of Finland.
124 Triglycerides and total cholesterol were also determined from these samples, which enabled the
125 calculation of total serum lipids according to equation of (Phillips et al., 1989).

126 In the Extended Finnish PASTURE Cohort (LUKAS2) (Karvonen et al., 2009), serum samples
 127 were collected in September 2014–November 2015, from 78 boys and 70 girls 10 years of age from
 128 the city of Kuopio and surrounding countryside. As serum lipids were not determined in the
 129 LUKAS2 samples, the median of MULTIEPIGEN total serum lipids of (0.51%) was assumed for
 130 all EFPC samples.

131

132 **Table 1.** Background information for serum samples collected from different cohorts.

Cohort/City	Year of birth	Age at sampling	Nr. of samples
MULTIEPIGEN			
Helsinki			
Female	2009–2011	7–9	2
Kuopio			
Male	2011	7	3
Female	2009–2010	8–9	8
Oulu			
Male	2009–2010	8–9	2
Female	2009–2011	7–9	6
Tampere			
Male	2009–2011	7–9	5
Female	2009–2011	7–9	7
Turku			
Male	2009–2011	7–9	2
Female	2009	9	1
MULTIEPIGEN 7-9 years total			36
LUKAS2 (Kuopio and surrounding countryside)			

Male	2004	10	78
Female	2004	10	70

133

134 To create a regression model on the relationship of 9 PCBs to PCDD/F TEQ and total TEQ, we
 135 used our previous data on human milk samples collected in 2000 (n=113), 2005 (n=220) and 2010
 136 (n=202) from southern (Helsinki), central (Kuopio) and northern (Rovaniemi) Finland. Background
 137 information and (unpublished) summary of PCDD/F- and PCB-concentrations measured from these
 138 samples are given in Supplementary Information (SI) Table S1.

139 For quality control of the Bayesian estimation approach, blood samples of volunteer male donors
 140 aged 18–20 years from the cities of Helsinki (n=22), Tampere (n=20) and Turku (n=18) were
 141 collected by the Finnish Red Cross in February–March 2019. The pooled sample for each city was
 142 created by mixing 10 ml of blood from each donor and analysed for PCBs and PCDD/Fs from 100
 143 ml aliquot. We assumed congener profiles of PCDD/Fs and PCBs in these youngest possible
 144 volunteer male donors (18–20 years) to be the most representative available to those of young boys
 145 under scrutiny.

146

147 *2.2. Chemical analysis of serum, milk and whole blood samples*

148 From the 200 µl serum samples 9 PCB congeners were measured (PCB 74, 99, 118, 138, 153, 156,
 149 170, 180 and 187). PCBs 52, 101 and 183 were also measured but excluded from further analyses,
 150 because of high number of non-detects. Method consisted of liquid-liquid extraction, silica column
 151 clean-up and analysis by Agilent 7010 gas chromatograph triple quadrupole mass spectrometer
 152 (GC-MS/MS) (Koponen et al., 2013).

153 Standard extraction and clean up procedures used for PCDD/Fs and PCBs have been described
 154 earlier for mothers' milk (Kiviranta et al., 2004; Main et al., 2007) and pooled blood (Kiviranta et
 155 al., 2002) samples. Human milk samples were analysed with Agilent 6890 gas chromatograph

156 connected to Waters Autospec Ultima high resolution mass spectrometer (GC-HRMS) and pooled
157 blood samples (100 ml) with Agilent 7010 GC-MS/MS. Analysis of both mothers' milk and pooled
158 blood samples were conducted according to requirements laid down in the Commission Regulation
159 (EU) 2017/644 (EU, 2017). All methods used have been described in more detail in Supplementary
160 Information.

161
162 *2.3 Statistical testing of PCBs analysed from children's serum samples*

163 Basic statistical tests to compare measured concentrations 9 PCBs within and between
164 MULTIEPIGEN and LUKAS2 (mean, median, percentiles, t-test) were performed with IBM SPSS
165 Statistics 25. In the calculation of means, medians and percentiles of measured PCB concentrations
166 from MULTIEPIGEN and LUKAS2 cohorts, results below level of detection (LOD) were replaced
167 with LOD/2.

168
169 *2.4 Estimation of PCDD/F TEQ and total TEQ in serum samples using hierarchical Bayesian*
170 *regression model*

171 PCDD/F and PCB congener data from all human milk samples (n=535) and pooled blood donor
172 samples (n=3) were used to estimate the PCB congener coefficient distributions. These coefficient
173 distributions were then used to predict individual posterior distributions of PCDD/F TEQs and total
174 TEQs for each of the 184 children from MULTIEPIGEN and LUKAS2 studies.

175 To do this, TEQ values (separately for PCDD/F and PCB TEQ concentrations) were predicted using
176 linear regression analysis of log-transformed PCB concentrations. The regression formula was

177 $\ln y_i \sim N(\sum \beta_j \ln x_{i,j}, \tau),$

178 where y is the TEQ concentration, i is an index for individuals, j is an index for PCB congeners, N
179 is a normal distribution with parameters mean and precision, β is a congener-specific coefficient, x
180 is a congener concentration, and τ is the precision of the distribution. We used hierarchical Bayesian

181 regression with Gibbs sampling (Gelman et al., 2004) rather than frequentist linear regression,
182 because we were interested in the posterior TEQ distributions of individual people given their
183 personal PCB profiles.

184 Uninformative normal priors were used for β_j for each congener and uninformative gamma priors
185 were used for τ for PCDD/F TEQ and PCB TEQ concentrations. The total TEQ was calculated as
186 the sum of PCDD/F TEQ and PCB TEQ. The parameters β and τ were assumed to apply to all study
187 groups: children, blood donors, and mothers.

188 Blood donors and mothers were added to the modelled data set twice: first with the measured TEQ
189 value to estimate the parameter distributions, and then without the measured TEQ value to predict
190 individual posterior distributions about what the TEQ estimates would likely be if they were not
191 known.

192 Other independent variables than PCB concentrations were not used, because the explanatory power
193 of the model was already very good (R^2 ca. 0.9 for total TEQ, see supplementary information
194 Figure S1).

195 Regression analyses were performed using R statistical software (version 3.6.2, cran.r-project.org)
196 and JAGS Gibbs sampler (version 4.3.0).

197 The model code and synthetic data for testing is available at Github
198 (<https://github.com/jtuomist/dioxdisthuman>).

199

200 **3. Results**

201 *3.1. Concentrations of PCBs measured in the serum of MULTIEPIGEN and LUKAS2 children*

202 Overall, the concentrations of PCBs in the serum samples from MULTIEPIGEN and LUKAS2
203 cohorts were very low (Table 2). For 9 PCB congeners (74, 99, 118, 138, 153, 156, 170, 180 and
204 187) mean and median concentrations were >LOD. These 9 congeners were included in the further

205 analysis. However, even of these 9 congeners, mean concentrations of PCBs 74, 99, 156, 118 and
 206 187 were below or around the LOQ of 5 pg/ml (Table 2).

207

208 **Table 2.** Concentrations of PCBs (pg/ml) measured in MULTIEPIGEN and LUKAS2 serum
 209 samples (n=184).*

Congener	% > LOD	Mean	Median	10th perc.	90th perc.
PCB52	1.8	<LOD	<LOD	<LOD	<LOD
PCB74	67	3.2	2.4	<LOD	5.3
PCB99	73	3.3	2.9	<LOD	5.6
PCB101	1.8	<LOD	<LOD	<LOD	<LOD
PCB118	95	5.3	4.5	2.7	8.8
PCB138	99	21	18	7.6	41
PCB153	100	38	31	12	72
PCB156	60	3.3	2.7	<LOD	7.1
PCB-170	96	10	8.4	2.7	21
PCB180	100	23	19	5.9	48
PCB183	34	<LOD	<LOD	<LOD	3.3
PCB187	85	5.6	4.5	<LOD	11

210 *LOD = 2 pg/ml and LOQ = 5 pg/ml for all congeners. In the calculation of means, medians and
 211 percentiles results <LOD were replaced with LOD/2.

212

213 Mean and median concentrations of \sum PCB-9 in MULTIEPIGEN and LUKAS2 cohorts by sex are
 214 given in SI Table S2. Due to 3 extreme samples, mean of \sum PCB-9 in MULTIEPIGEN boys (40
 215 ng/g lw) was 44% larger than in MULTIEPIGEN girls (28 ng/g lw), but difference in the median of
 216 \sum PCB-9 was only 18%. Overall, as no statistically significant within cohort difference in serum
 217 \sum PCB9 levels between boys and girls were found in 2-sided t-test in either MULTIEPIGEN or

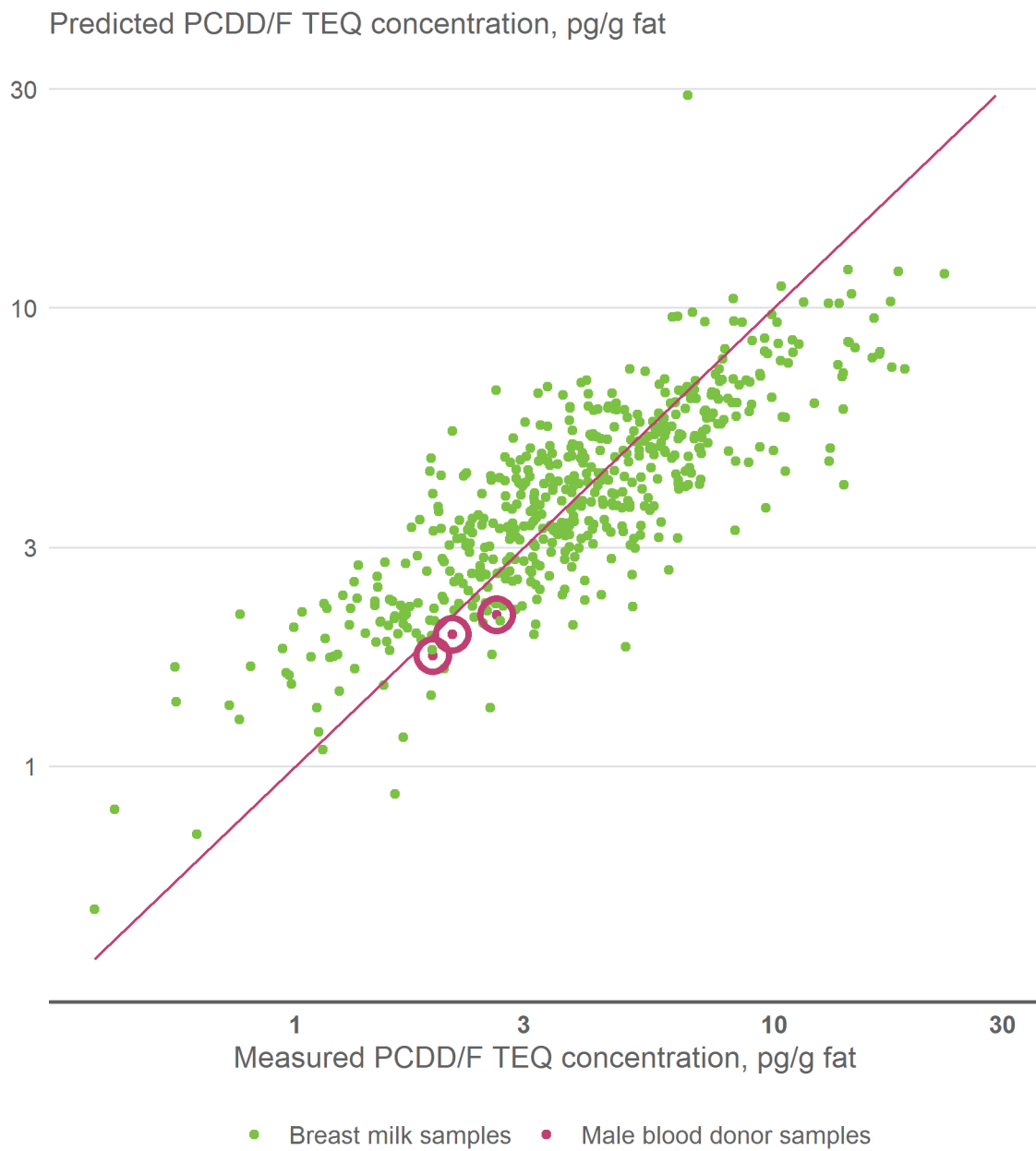
218 LUKAS2, we considered it appropriate to include also girls in the Bayesian analysis. However,
219 differences between cohorts were borderline or significant (SI Table S3).

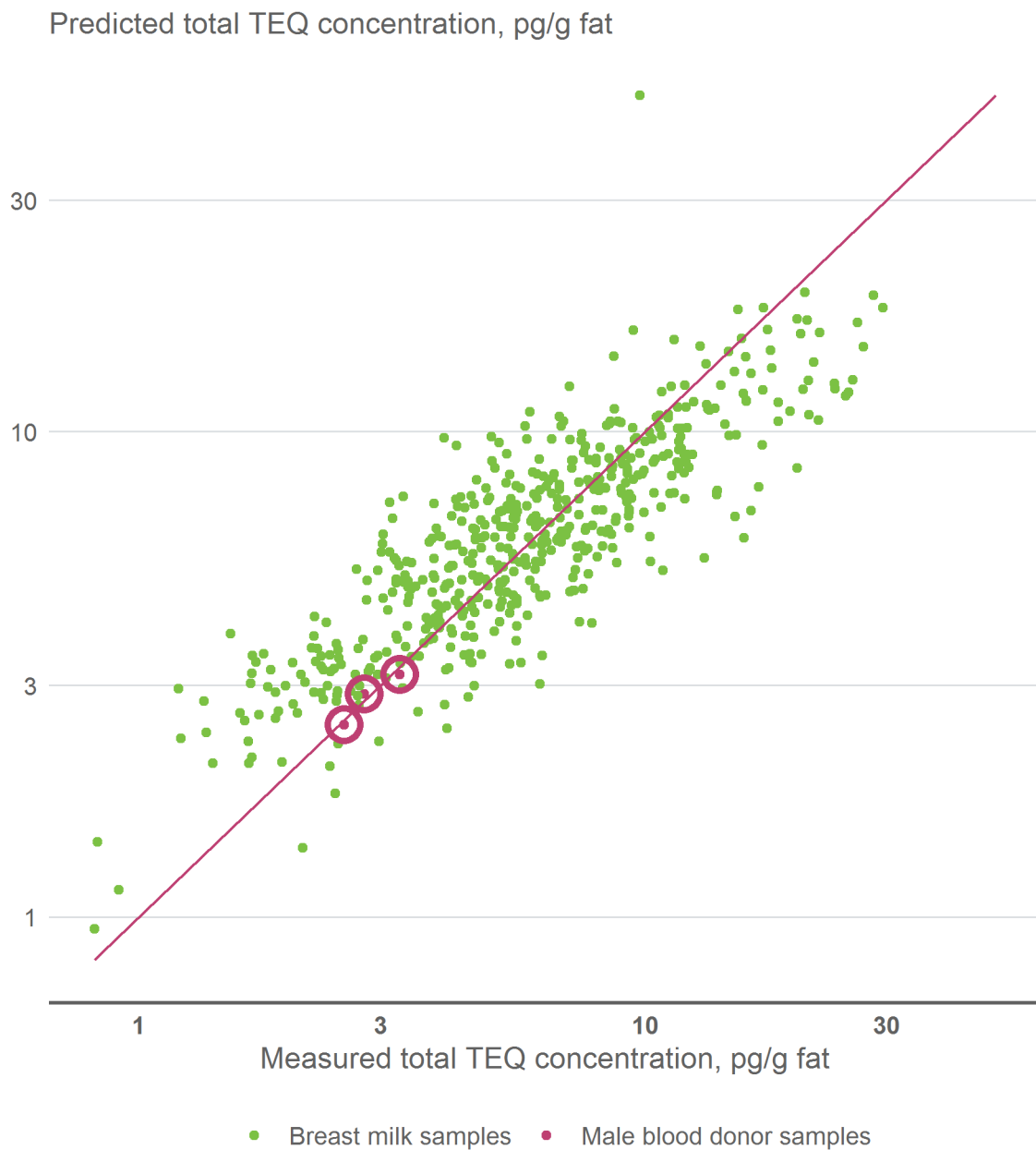
220

221 *3.2. TEQ-prediction power of PCBs for human milk and pooled blood samples*

222 The power of 9 PCBs to predict TEQs was tested in 3 different ways using human milk samples.
223 Results are shown in SI Figure S1. For PCDD/F TEQ, PCB-TEQ and total TEQ, the best
224 correlation is obtained when congener-specific concentration data of all 9 PCBs contribute to the
225 model. Obviously the fit for PCB-TEQ was better than for PCDD/F TEQ, but slightly surprisingly
226 the fit for total TEQ was similar to or even better than for PCB TEQ. Overall, information rich
227 models (congener-specific data of all 9 PCBs as predictors) are superior for all estimated TEQs
228 compared to single congener or sum of congeners.

229 Figure 1 shows measured versus predicted PCDD/F TEQ and total TEQ using 9 PCBs as predictors
230 for human milk and pooled blood samples. For human milk samples, models appear to slightly
231 underestimate both TEQs at low concentrations and overestimate at high concentrations. However,
232 between 3 and 10 pg/g that encloses the NOAEL of 7 pg total TEQ/g lipid, the fit is satisfactory.
233 Pooled blood samples used for quality control are relatively close to one-to-one line in Figure 1.
234 Measured and predicted TEQ-concentrations are shown numerically in SI Table S4. Compared to
235 measured lower bound concentrations, predicted concentrations of PCDD/F TEQs range from 84%
236 to 94% and those of total TEQs from 99% to 106%, respectively.





238

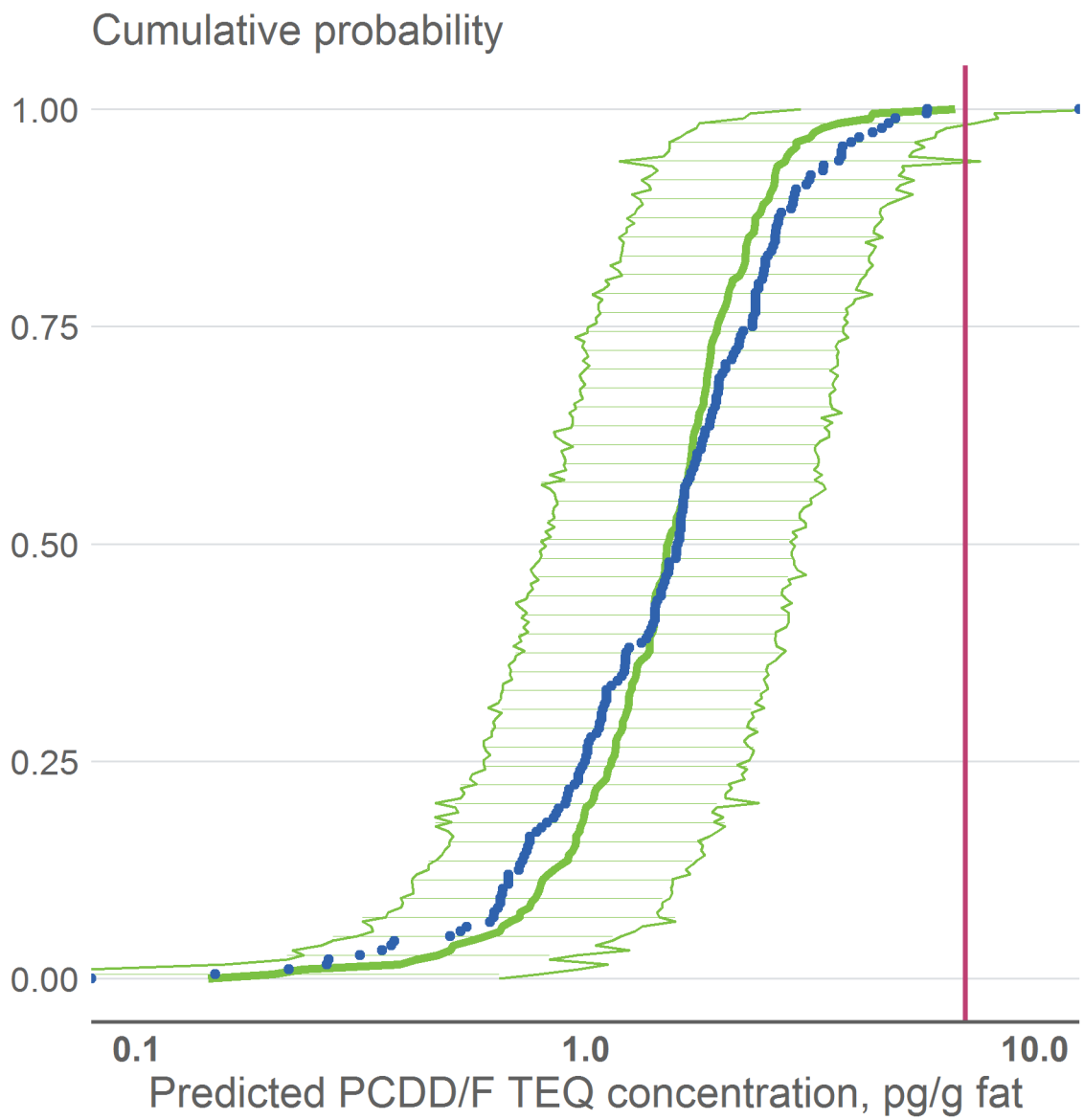
B)

239 Figure 1. Prediction accuracy of 9 PCBs for A) PCDD/F TEQ and B) total TEQs with Bayesian
 240 hierarchical regression model for human milk (green dots) and pooled blood (red circles) samples.

241 *3.3 PCDD/F TEQ and total TEQ concentrations in the serum of MULTIEPIGEN and*
242 *LUKAS2 children, estimated by Bayesian hierarchical regression model*

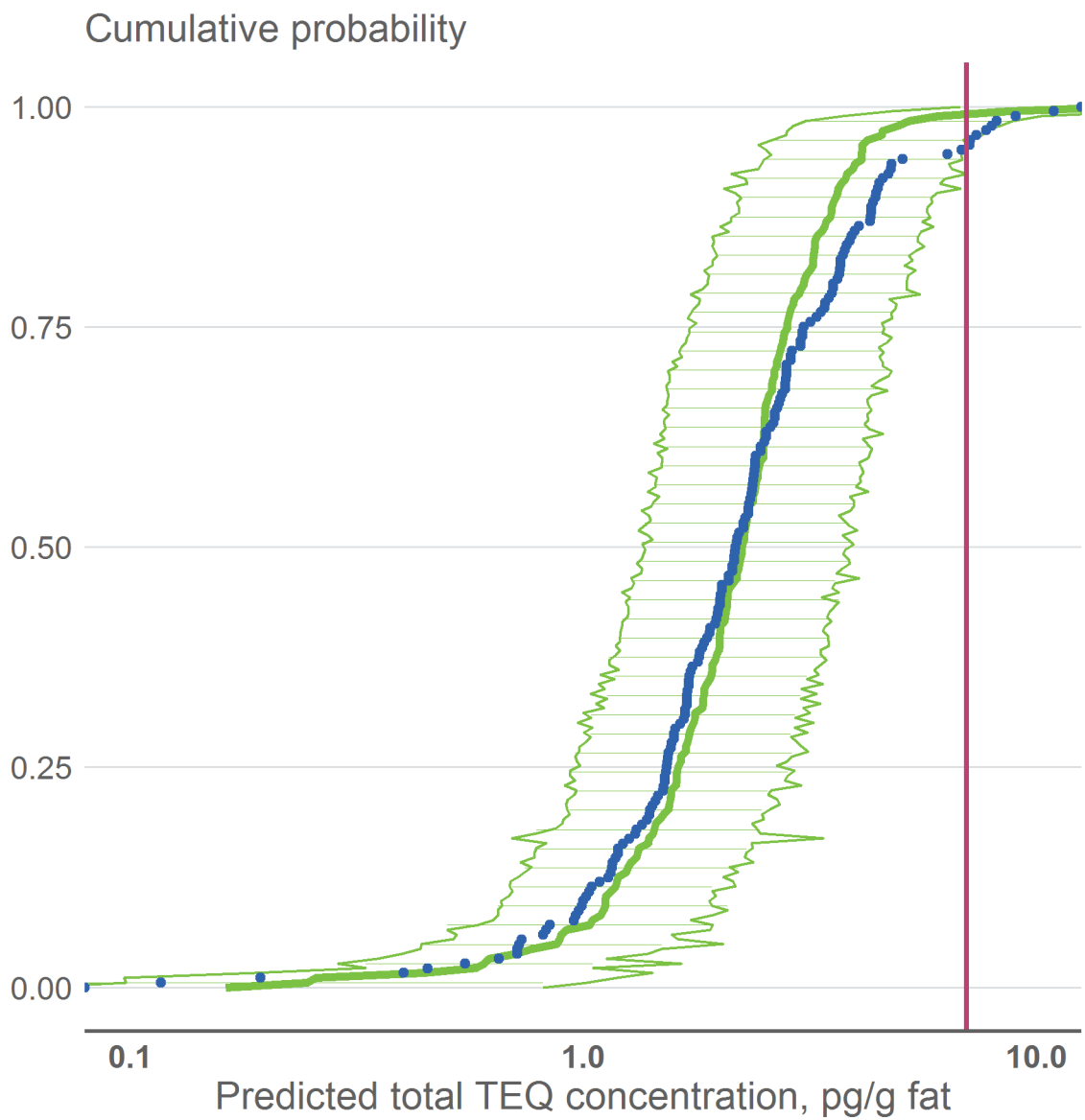
243 Cumulative probability distribution of predicted PCDD/F TEQs and total TEQs in children's
244 serum from MULTIEPIGEN and LUKAS2 combined compared to NOAEL of 7.0 pg TEQ/g
245 lipid are shown in Figure 2. All expected values for PCDD/F TEQ and 98,9% of expected
246 total TEQ values were below NOAEL. However, random draws from each child's probability
247 distribution is a more realistic approach to evaluate NOAEL exceedance. For 0.5% and 2.7 %
248 of the children PCDD/F TEQ and total TEQ was exceeded, respectively.

249



+ Expected value and 5-95% CI
 + Random draw
 + NOAEL 7 pg/g

250
251 A)



+ Expected value and 5-95% CI
 + Random draw
 + NOEL 7 pg/g fat

252
253 B)

254 Figure 2. Cumulative probability distribution of predicted A) PCDD/F TEQs and B) total
 255 TEQs in children's serum from MULTIEPIGEN and LUKAS2 combined. Green line =
 256 Expected value for each child; Green "saw-edged" line =90% CI (5-95%) individual

257 distributions of expected value; Blue line = Random draw from the probability distributions
258 of each child. Random draws points are independently arranged in increasing order, and they
259 do not refer to the same individual as expected value and its CI on the same row.

260

261 **4. Discussion**

262 The aim of this study was to estimate whether the NOAEL of 7.0 pg/g lipid identified from
263 the Russian study is exceeded for PCDD/F or total TEQ among 7–10 year old Finnish
264 children. When the estimated TEQ-concentration for each child was randomly drawn from
265 individual probability distributions, PCDD/F TEQ exceeded NOAEL in 0.5% and total TEQ
266 in 2.7% of subjects. These results and the overall decrease of human exposure imply that risk
267 management actions following the drastically decreased TWI proposed by the CONTAM
268 Panel should be carefully considered. If relatively remote risks define the dietary
269 recommendations, indirect consequences on total dietary risks may be great. For example,
270 exclusion of fish from the market in the situation where PCDD/F levels in the serum of boys
271 potentially affected are already largely below the NOAEL would only lead to losses of fish
272 related health benefits for consumers.

273 The approach selected to estimate the levels of PCDD/F TEQ and total TEQ in the blood of
274 boys around 9 years of age has to be seen in the context where serum available per individual
275 is barely enough for the measurement of major PCBs with a highly sensitive analytical
276 method. Total volume of serum available from all 184 samples received for this study could
277 have enabled congener specific analysis of PCDD/Fs and mono-ortho-PCBs from a single
278 pooled sample, but individual variation in concentrations would have been lost. Thus, we

279 utilized our extensive dataset from mothers' milk to model the relationship between 9 PCBs
280 measurable from small volume of serum and TEQ concentrations.

281 In the construction of hierarchical Bayesian regression model we used mothers' milk results
282 from years 2000, 2005 and 2010. These birth years match closely the birth years of the blood
283 donors (1999-2001), and children from LUKAS2 (2004) and MULTIEPIGEN (2009-2011).
284 There is some variation in the PCDD/F- and PCB-congener profiles between serum/blood and
285 breast milk. For example, in our milk samples median contribution of PCB-126 to total TEQ
286 was 22–25% (10th–90th percentiles 16–33%) and median contribution of PCB-TEQ to total
287 TEQ was 30–34% (10th–90th percentiles 24–43%) depending on the year. In pooled blood
288 samples PCB-126 contributed only 12–16% to the lower bound of total TEQ, and PCB-TEQ
289 contributed 20–24% to total TEQ. Lower contribution of PCB 126 to total TEQ in blood
290 compared to breast milk is in agreement with results from Japanese paired samples (Todaka et
291 al., 2010).

292 These differences in congener profiles are possibly reflected in the slight underestimation of
293 hierarchical Bayesian regression model predicted PCDD/F TEQs (84–99%) from pooled
294 blood samples. However, the prediction accuracy was considered satisfactory and proved the
295 ability of hierarchical Bayesian regression model to predict PCDD/F TEQ also from
296 MULTIEPIGEN and LUKAS2. As Red Cross donors were 10 years older than the boys of
297 critical age, our underlying assumption is that the half-life of 9 PCB congeners are close to
298 those PCDD/F congeners that make up most of the PCDD/F TEQ. Congeners that contributed
299 80% to PCDD/F TEQ were 1,2,3,7,8-PeCDD, 1,2,3,6,7,8-HxCDD and 2,3,4,7,8-PeCDF. The
300 elimination half-life of PCDD TEQ has been estimated at 8.7 years (EFSA 2018) and that of

301 2,3,4,7,8-PeCDF 7–20 years depending on the study (Flesch-Janys et al., 1996; Rohde et al.,
302 1999; Ogura, 2004; Milbrath et al., 2009). The estimates of intrinsic adult elimination half-
303 lives at background levels for PCB congeners 118, 138, 153, 170, 180, 187 range from 9 to 15
304 years (Ritter et al., 2009). Furthermore, (Grandjean et al., 2008) found no indication that
305 intrinsic elimination half-lives of PCBs depend on age or are shorter in children than in adults
306 after correcting for the effect of growth. Thus, half-lives of PCDD/F TEQ and 9 PCBs appear
307 reasonably similar to justify our assumption.

308 For random draw, the proportion of children exceeding the NOAEL was increased from 0.5%
309 for PCDD/F TEQ to 2.7% for total TEQ. Percentages remain small, but relative increase from
310 PCDD/F TEQ to total TEQ was large. However, the interpretation of total TEQ is somewhat
311 complicated. In the EFSA risk assessment there was also a strong recommendation to
312 reevaluate the TEF-value of 0.1 assigned for PCB-126. Observations from *in vitro* studies
313 with human cells show that PCB-126 is much less potent in humans than suggested by the
314 WHO2005-TEF of 0.1.. This was seen by EFSA to be reflected in the critical study on
315 Russian boys, where PCDD/F TEQ (p=0.04) but not total TEQ (p=0.61) was associated with
316 decreased sperm concentration (EFSA 2018).

317 Small percentage of children exceeding the NOAEL may on part depend on the age of
318 children in our cohorts. We combined MULTIEPIGEN and LUKAS2 children in the Bayes
319 model due to limited number of samples available from MULTIEPIGEN (n=36) even if mean
320 Σ PCB9 (ng/g lipid) in MULTIEPIGEN children was significantly (p=0.001) higher than in
321 LUKAS2. This difference can largely be explained by growth dilution as the median Σ PCB9
322 concentration in MULTIEPIGEN subjects converges with increasing age from 7 years (40

323 ng/g lipid, n=7) to 8 years (27 ng/g lipid, n=9) and to 9 years (23 ng/g lipid, n=19) towards to
324 the median Σ PCB9 concentration (17 ng/g lipid, n=148) in LUKAS2 subjects all 10 years of
325 age. However, at the age of 9 years only a slightly larger fraction of LUKAS2 children would
326 have exceeded the NOAEL of 7.0 pg/g lipid based on the shape of PCDD/F and total TEQ
327 distributions (Figure 2A and 2B).

328 The TWI of 2 pg TEQ/kg bw for the sum PCDD/Fs and PCBs proposed by the CONTAM
329 Panel was based on the accumulation models used (EFSA, 2018). Regarding the model used
330 for boys in the EFSA risk assessment (milk 5.9 pg TEQ/g for 1 year + intake 4 pg
331 TEQ/kg/week for 8 years results in serum level of 7 pg TEQ/g lipid), our data on PCDD/F
332 basis supports it relatively well. Median PCDD/Fs in breast milk around 2005–2010 were 2–4
333 pg TEQ/g lipid (Table S1). Based on the Finnish intake study (Kiviranta et al., 2001), our fish
334 data (Airaksinen et al., 2018) and recent Swedish intake data (Livsmedelverket, 2017), adult
335 dietary PCDD/F intake around the years 2010–2018 could be estimated at 2–4 pg TEQ/kg
336 bw/week that would imply an intake of 4–8 pg TEQ/kg bw/week for children. However,
337 considering that this intake is likely overestimated due to lower consumption of Baltic Sea
338 fatty fish among children compared to the whole population estimate almost 20 years ago
339 (Kiviranta et al., 2004), our estimated median serum PCDD/F TEQ levels of 2–3 pg/g lipid
340 around 10 years of age is in good agreement with the EFSA model. On the contrary, the long
341 term maternal accumulation model of EFSA (milk 5.9 pg TEQ/g lipid for 1 year + intake 2 pg
342 TEQ/kg/week for 34 years results in milk level of 5.9 pg TEQ/g lipid) seems largely
343 overestimated. Our unpublished median milk PCDD/F concentration in primipara mothers in
344 1987 was 23 pg TEQ/g lipid, calculated intake around 2000 was 5–6 pg TEQ/ kg bw/week

345 (Kiviranta et al., 2001; Kiviranta et al., 2004) and an estimate of PCDD/Fs in the milk of a 33
346 year old primipara mother in year 2020 is ≤ 2.5 pg TEQ/g lipid extrapolated from 2010 milk
347 data assuming 4–5% annual decrease. Thus, even if our intake estimate from 2000 would be
348 50% overestimated for women entering the reproductive age, the maternal accumulation
349 model of EFSA would still be clearly overprotective compared to our measured levels in
350 breast milk from 1987 through 2010 and estimated levels in 2020. This also calls for caution
351 in risk management options considering the neurodevelopmental benefits of maternal fish
352 consumption during pregnancy and lactation.

353 This study has some limitations. Very small amount of serum was available from each child
354 and it was thus not possible to measure PCDD/F TEQ and total TEQ from pooled serum
355 samples of children. It would have been useful to compare the pool concentrations with the
356 mean of hierarchical Bayesian regression estimates. Children in LUKAS2 were 1–2 years
357 older than in the critical Russian study that could have resulted in slight underestimation in
358 the proportion of children exceeding the NOAEL of 7.0 pg/g lipid. The number of boys
359 especially in MULTIEPIGEN was limited and we merged the cohorts as pre-pubertal
360 PCDD/F and PCB levels within cohorts were not significantly different between sexes.
361 Finally, we did not have serum lipids measured from LUKAS2, but used the median of
362 MULTIEPIGEN total lipids for all LUKAS2 subjects. However, this was assumed to have
363 only a minor impact on results as the range of total lipids in MULTIEPIGEN (and supposedly
364 in LUKAS2 also) was narrow (5th–95th percentile 0.43–0.64%) and was expected to be similar
365 also in LUKAS2.

366

367 **5. Conclusions**

368 We used a hierarchical Bayesian regression model to estimate PCDD/F TEQs and total TEQs
369 in 7–10 year old Finnish children’s serum from measured serum levels of major PCBs. More
370 than 99% of children were estimated to have serum PCDD/F TEQ concentration below the
371 NOAEL of 7.0 pg/g lipid identified from the critical Russian study for impaired semen quality
372 later in life. For total TEQ, the respective estimate was more than 97%. Due to cohorts used,
373 the mean age of our children was slightly less than 10 years compared to 8–9 years in the
374 critical Russian study. Our estimate is thus likely a slight underestimate of the true
375 concentrations in 8–9 year old Finnish boys. However, our estimate is accurate enough to
376 conclude that possible risk management actions following the drastically decreased TWI of 2
377 pg TEQ/kg bw/week established by the CONTAM Panel should be carefully considered
378 before acceptance. There are little if any potential health risks that can be avoided with
379 decreased maximum levels in foods, but a plenty of health benefits to lose if fish consumption
380 is decreased as a result lowered maximum levels. We recommend more biomonitoring studies
381 especially on the levels of PCDD/Fs and PCBs in breast milk as such studies are the easiest to
382 conduct. Also analysis of PCDD/Fs and PCBs in the serum of young boys either from pooled
383 samples or estimated using regression models as presented here is warranted. Risk
384 management actions should also rely on measurements of sensitive target population
385 concentrations, not solely on modelling.

386

387 **Declaration of interest**

388 None

389

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392

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1 **Supplementary information**

2

3 **Estimated PCDD/F TEQ and total TEQ concentrations in the serum of 7–10 year old Finnish**
4 **children**

5

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26

27 **Analysis of PCBs from serum**

28 From the 200 µl serum samples (Young Finns Study and Extended Finnish Pasture Cohort) 9 PCB
29 congeners (74, 99, 118, 138, 153, 156, 170, 180 and 187) were measured. Sum of these PCBs is
30 denoted as \sum PCB-9 in tables below. The method has been described earlier in detail (Koponen et al.
31 2013). In brief, ethanol and ¹³C-labelled internal standards of each compound in toluene were
32 added to samples (200 µl) in test tubes and mixed to precipitate the proteins and equilibrate internal
33 standards. Dichloromethane-hexane (1:4) was added for extraction followed by activated silica to
34 bind the sample water, ethanol and precipitate. Samples were mixed, and layers were allowed to
35 separate. The upper dichloromethane-hexane layer was poured to a solid phase extraction cartridge
36 (SPE cartridge) containing from bottom to top 10% AgNO₃ impregnated silica and a mixture of
37 Na₂SO₄ and silica. The lower layer in the test tube was extracted again with dichloromethane-
38 hexane that was also poured to SPE-cartridge. Elution of SPE-cartridges was continued with
39 dichloromethane-hexane and the eluate was concentrated to 15-20 µl for gas chromatography - high
40 triple quadrupole mass spectrometry (GC-MS/MS) analysis. The instrument used was Agilent 7010
41 GC-MS/MS system (Wilmington, DE, USA) GC column was DB-5MS UI (J&W Scientific, 20m,
42 ID 0.18 mm, 0.18 µm film).

43 As control serum sample NIST SRM 1958 with known concentrations of PCBs was analysed in
44 each batch (n=8) of samples. In addition, due to generally low levels of POPs in YFS and EFPC
45 samples, also SRM 1958 diluted 1/10 with New Born Calf Serum (Dil-SRM 1958) was analysed in
46 each sample batch. Mean accuracy (91-110% of SRM 1958 certified/reference concentrations and
47 85-102% of calculated concentrations in diluted SRM 1958) and precision (co-efficient of variation
48 (CV) ≤ 4.1% for SRM 1958 and ≤ 5.4% for Dil-SRM 1958) were acceptable for measured PCBs in
49 control samples. Limit of quantitation (LOQ) was 5 pg/ml and limit of detection (LODs) 2 pg/ml
50 for 9 PCBs.

51

52 **Analysis of PCDD/Fs and PCBs mothers' milk and pooled blood samples**

53 From the mothers' milk and pooled blood samples (100 ml), 17 dioxin congeners and 23 PCBs
54 (including 12 dioxin like PCBs and 11 other PCBs including the above 9 PCBs) were analysed.
55 Sample pre-treatment included liquid-liquid extraction, gravimetric determination of lipid content,
56 multilayer silica column cleanup and fractionation by activated carbon column as previously
57 described for milk (Kiviranta et al., 2004; Main et al., 2007) and blood (Kiviranta et al. 2002).
58 Cleaned milk samples were analysed with Waters Autospec Ultima high resolution mass
59 spectrometer (GC-HRMS) and pooled blood samples with Agilent 7010 GC-MS/MS. GC column

60 was DB-5MS UI (J&W Scientific, 60m, ID 0.25 mm, 0.25 µm film) in both cases. ¹³C-labelled
61 internal standards of each congener were used for quantification. LOQs of each congener were
62 calculated using the “signal/noise ratio of 3:1” –approach for mothers’ milk samples and using the
63 “lowest concentration point on a calibration curve” –approach for pooled blood samples as laid
64 down in the Commission Regulation (EU) 2017/644 (EU 2017) due to instrumental techniques used
65 in each matrix, respectively.

66 Regarding the quality control, during the years of milk sample analysis (2001-2011), laboratory
67 participated every year successfully to Interlaboratory Comparison of Persistent Organic Pollutants
68 in food (ILC POPs) since the first round in 2000 (<https://www.fhi.no/en/studies/ilc-pop/>). For the
69 recent analysis of pooled blood samples, an in-house spiked quality control sample was extracted
70 and analysed with GC-MS/MS that had a recovery of 89% for PCDD/F TEQ, 111% for PCB-TEQ
71 and 88% for the sum of 23 PCBs as compared to long term in-house average, respectively.

72 Laboratory of chemistry at the Finnish Institute for Health and Welfare is an accredited testing
73 laboratory T077 by Finnish Accreditation Services (FINAS) since 1996. Scope of accreditation
74 includes all of the methods used within this study.

75

Table S1. Median (10th - 90th percentiles) concentrations of Σ PCB-9 (ng/g lipid), PCDD/F-TEQ and Total-TEQ (pg/g lipid) in mothers' milk samples collected from the cities of Helsinki, Kuopio and Rovaniemi in 2000, 2005 and 2010.

Collection year	City	n	% of 1st-2nd-3rd birth**	Age (range)	Σ PCB-9 (ng/g lipid)	PCDD/F-TEQ (pg/g lipid)	Total TEQ (pg/g lipid)
2000							
	Helsinki	56	52 - 34 - 11	31 (26 - 37)	131 (60 - 245)	8.8 (4.2 - 16)	12 (6.1 - 24)
	Kuopio	57	39 - 32 - 25	31 (24 - 36)	93 (47 - 164)	6.3 (3.1 - 13)	9.2 (4.6 - 17)
2005							
	Helsinki	94	41 - 47 - 9	30 (25 - 37)	73 (40 - 140)	4.1 (2.1 - 8.2)	6.8 (3.3 - 14)
	Kuopio	83	48 - 28 - 16	29 (22 - 36)	53 (29 - 101)	3.9 (2.1 - 6.6)	5.5 (3.0 - 9.7)
	Rovaniemi	43	23 - 47 - 14	30 (24 - 39)	61 (33 - 113)	3.4 (2.1 - 6.9)	5.4 (3.4 - 10.4)
2010							
	Helsinki	77	53 - 36 - 8	31 (27 - 37)	57 (28 - 122)	3.3 (1.4 - 5.9)	5.1 (2.3 - 9.1)
	Kuopio	71	32 - 31 - 20	30 (24 - 36)	39 (16 - 76)	2.1 (1.1 - 4.4)	3.1 (1.6 - 6.2)
	Rovaniemi	54	39 - 35 - 17	30 (25 - 37)	48 (23 - 107)	2.6 (1.2 - 5.7)	4.0 (1.9 - 8.1)

* Parity above 3 is what is missing from 100 %

Table S2. Mean and median (10th and 90th percentiles) concentrations of measured Σ PCB-9 (ng/g lipid) by cohort and sex.

Cohort	Year of birth	Age	n	Mean	Median (10 th - 90 th perc.)
YFS					
Boy	2009-2011	7-9	12	40.3	30.9 (16.7 - 90.5)
Girl	2009-2011	7-9	24	27.9	26.1 (3.61 - 53.1)
EPFC					
Boy	2004	10	78	20.3	16.1 (7.00 - 41.1)
Girl	2004	10	70	17.9	17.2 (8.2 - 30.4)

Table S3. Difference in the Σ PCB9 (ng/g lipid) in boys and girls within and between cohort groups by t-test (p-values).

Cohort	Group	YFS		EFPC*	
		B 7-9 y	G 7-9 y	B 10 y	G 10 y
YFS	B 7-9 y		0.152	0.023	0.012
	G 7-9 y	0.152		0.053	0.011

*No statistically significant difference in Σ PCB9 between boys and girls within EFPC cohort by t-test (p=0.267).

Table S4. Measured concentrations of PCDD/F and PCBs congeners, PCDD/F-TEQs and Total-TEQs in pooled blood samples[#], and estimated PCDD/F-TEQ and Total-TEQ using measured concentrations of 9 PCBs with Bayesian hierarchical regression model.

City	Helsinki	Tampere	Turku
PCDD/Fs (pg/g lipid)			
2,3,7,8-TCDD	<0.56	<0.53	<0.49
1,2,3,7,8-PeCDD	0.82	0.60	0.62
1,2,3,4,7,8-HxCDD	<0.57	<0.50	<0.47
1,2,3,6,7,8-HxCDD	3.79	3.81	3.73
1,2,3,7,8,9-HxCDD	<0.55	<0.49	<0.46
1,2,3,4,6,7,8-HpCDD	8.74	6.20	7.36
OCDD	110	77.9	88.2
2,3,7,8-TCDF	<1.6	<1.5	<1.4
1,2,3,7,8-PeCDF	<1.0	<0.99	<0.91
2,3,4,7,8-PeCDF	2.76	1.76	2.14
1,2,3,4,7,8-HxCDF	1.70	1.23	1.20
1,2,3,6,7,8-HxCDF	1.59	0.88	1.32
1,2,3,7,8,9-HxCDF	<0.34	<0.32	<0.30
2,3,4,6,7,8-HxCDF	<0.48	<0.44	<0.41
1,2,3,4,6,7,8-HpCDF	6.13	4.15	5.30
1,2,3,4,7,8,9-HpCDF	<0.38	<0.34	<0.32
OCDF	<1.4	<1.3	<1.2
non-ortho-PCB (pg/g lipid)			
PCB-77	32.6	29.2	23.5
PCB-81	<1.3	<1.4	<1.1
PCB-126	3.87	3.57	4.42
PCB-169	6.05	6.15	5.33
mono-ortho-PCB (ng/g lipid)			
PCB-105	0.27	0.22	0.22
PCB-114	<0.15	<0.14	<0.13
PCB-118*	1.30	1.07	1.07
PCB-123	<0.18	<0.17	<0.16
PCB-156*	0.67	0.70	0.51
PCB-157	<0.22	<0.21	<0.19
PCB-167	<0.29	<0.27	<0.26
PCB-189	<0.15	<0.13	<0.12
di-ortho-PCB (ng/g lipid)			
PCB-28	0.72	0.73	0.83
PCB-52	0.25	0.26	0.23
PCB-74*	0.54	0.47	0.48

PCB-99*	0.81	0.63	0.70
PCB-101	0.40	0.37	0.32
PCB-138*	5.27	4.41	3.98
PCB-153*	8.41	7.91	6.52
PCB-170*	2.29	2.38	1.73
PCB-180*	4.87	5.24	3.62
PCB-183	0.55	0.48	0.42
PCB-187*	1.54	1.42	1.11
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Congener based PCDD/F-TEQ (pg/g lipid)			
lower bound	2.54	1.85	2.04
medium bound	3.01	2.29	2.45
upper bound	3.49	2.74	2.86
Congener based Total-TEQ (pg/g lipid)			
lower bound	3.18	2.45	2.70
medium bound	3.67	2.91	3.12
upper bound	4.15	3.37	3.55
<hr/>			
Bayes estimated PCDD/F-TEQ (pg/g lipid)			
	2.14	1.73	1.91
Bayes estimated Total-TEQ (pg/g lipid)			
	3.15	2.52	2.86

Samples from 18-20 year old males from Helsinki (22 donors), Tampere (20 donors) and Turku (18 donors).

*Congener included to 9 PCBs used in modelling.

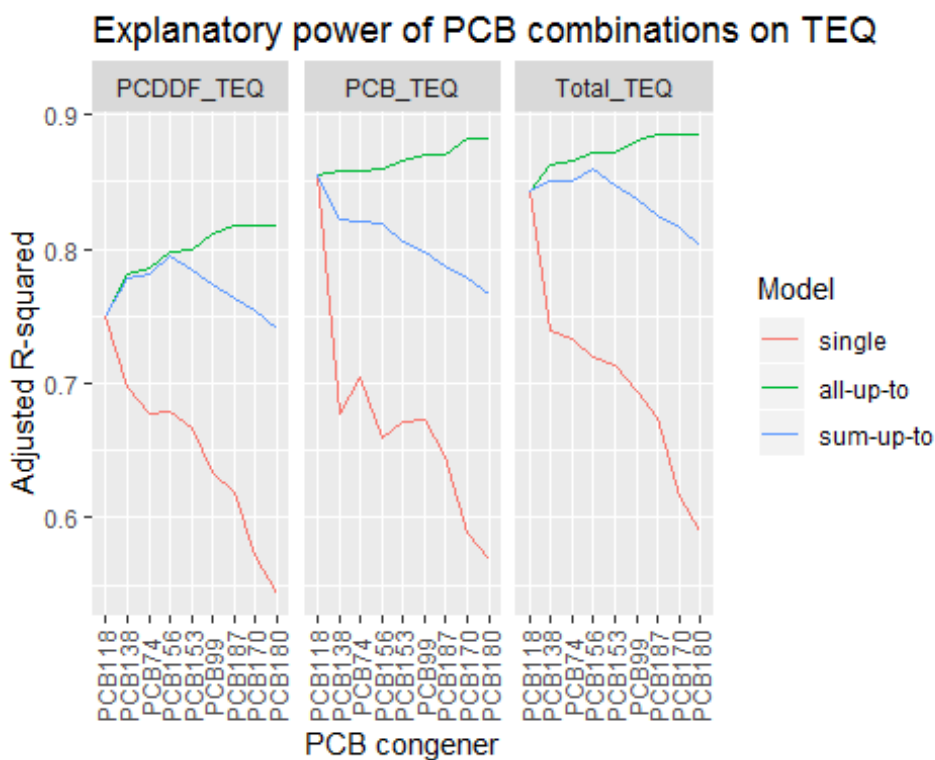


Figure S1. Explanatory power of different combinations of PCBs for TEQ-prediction. Measured PCDD/F and PCB concentrations from human milk (n=535) and pooled blood donor samples (n=3) were used to construct the hierarchical Bayesian regression model that predicted TEQ concentrations using measured PCBs. The congeners on x axis are ordered according to decreasing R^2 -value for Total-TEQ prediction by single PCB congener. Single = only one PCB congener concentration as predictor (1 predictor value); all-up-to: congeners added one-by-one to model, but congener-specific concentration data of all added congeners remain in the model (1-9 predictor values); sum-up-to: congener concentrations summed one-by-one on the top of previous congeners in the model (1 predictor value).

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