



# **CONTENTS**



 We calculated the mean relative abundance and standard error for each compound class according to maturity status and compound origin (anthropogenic, natural, and unknown). To reduce the dimension of the data and to identify co-occurring compound classes, we ran a PCA on all 16 individuals with the logged sum of the relative abundance of all 26 compound classes (167 individual total congeners) as variables.

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#### *Identifying maturity-related HOC bioaccumulation*

 We used Mann-Whitney U tests to evaluate significant differences in overall number of compounds and total relative abundance between maturity status. We then created a Random Forest model to examine differences in compound class abundance between 51 mature and immature individuals using the randomForest R package<sup>1</sup>. Permutation tests 52 were conducted with the rfPermute package<sup>2</sup> to determine significance for each compound class. Compound class abundances were calculated only using compounds found in 10 or more individuals (76 compounds from 23 classes). Because Random Forest cannot be conducted with missing data, we estimated the relative abundance for the five excluded peaks from 3 different individuals by measuring the best available peak areas. These five measurements were only used in random forest analyses and were excluded from all other analyses and data summaries. A compound class was considered significant when permutation tests calculated a p-value of 0.05 or below.

 Differences in log testosterone levels due to maturity status were evaluated with t-tests. We then ran a linear regression between the first axis of principal component analysis (PC1; which represented variation in 15 anthropogenic and unknown compound classes), and blubber testosterone level to examine how endocrine health is related to HOC exposure. We regressed the negative value of PC1 (because PC1 was negatively related with prominent compound abundance) with age to control for age-related variation in HOC load and extracted the residuals from the model as a proxy of the variation not explained simply by age. Similarly, the log of blubber testosterone was regressed with average gonad mass. If both gonad mass were unavailable, we included only the right gonad mass. Residuals from each of the PC1-age and testosterone-gonad mass models were then used in a linear model to compare age-corrected PC1 (as the predictor variable) with gonad mass-corrected testosterone (the response variable). The logged sum of the congeners in the PC1 group were similarly age-corrected. The age correction was done by building a model with age and HOC load and extracting the residuals. The variation not explained simply by the age model was then used in the linear regression with testosterone (corrected by gonad mass) as above to confirm that this trend can be explained by the compound classes of interest. In mature individuals, the non-age corrected sum of the PC1 compounds were log transformed and regressed directly with gonad mass-corrected testosterone.

 We constructed two additional Random Forest regression models on mature individuals 82 to explore the effects of HOCs on testes endocrine function (classes: mtry  $= 7$ , ntree  $=$   $4,000$ , nrep = 2,000; individual compounds: mtry = 13, ntree = 8,000, nrep = 4,000). For both Random Forest analyses, we used only compounds found in 10 or more individuals (76 compounds from 23 classes) in mature individuals to exclude confounding factors due to bioaccumulation with age. The focus on mature individuals also reflects the greater likelihood that impaired or compromised testes function can be detected in mammals during near maximum testosterone production, i.e. when there is the greater physiological 89 demand to reproduce. The Random Forest models included gonad mass-corrected testosterone as the response variable and either compound class abundance or individual congener abundance as the predictor variable. Compound classes were considered significant when permutation tests calculated a p-value of 0.05 or below. In case of non- convergence (i.e. conflicting model results), we also assessed variable importance, or the mean decrease in accuracy when removed from the model (%IncMSE), where a higher variable importance represents a larger increase in predictive power with the inclusion of 96 the predictor variable.

# **TABLES AND FIGURES**



**Table S1:** The month and year of sample collection, maturity, and age for each of the individuals that were included in this research.

**Table S2:** A list of the compounds and corresponding group name detected across all 16 samples. For compound structure, identification, mass spectrum and chromatographic information, see Supplemental Material for Shaul et al<sup>4</sup>. The origin of each compound class is represented by the following: anthropogenic [A], natural [N], mixed [M], and unknown[U].

#### **Brominated/chlorinated diphenyl ether (B/CDE) [U]** PBCDE Br3Cl 3

**Brominated-anisole [M]** 2,4,6-tribromo anisole

**Brominated-indole [N]** 4,6-dibromoindole 5-bromoindole

#### **Chlordane-related [A]**

alpha chlordane chlordane related 2 chlordane related 4 chlordane related 6 chlordane related 7 chlordane related 9 chlordane related 10 chlordane related 13 chlordane related 14 cis nonachlor gamma chlordane oxychlordane trans nonachlor

### **Chlorinated-benzene [A]**

1,2,3,4-tetrachlorobenzene hexachlorobenzene pentachlorobenzene

### **Chlorinated-styrene [A]**

ethenyl benzene 3Cl polychlorinated styrene 7Cl **Chlorobenzaldehyde [U]** Chlorobenzaldehyde

## **Dichlorodiphenyltrichloroethanerelated (DDT-related) [A]** DDMU 1 DDMU 2 DDMU 3 DDMU 4 DDT related 2 DDT related 3 DDT related 12 DDT related 14 DDT related 16 DDT related 18 DDT related 20 DDT related 23 o,p'-DDD p,p'-DDD

### **Dichlorobenzophenone [A]**

4,4'-dichlorobenzophenone

# **Dimethyl bipyrroles (DMBPs) [N]**

DMBP Br2Cl2 1 DMBP Br2Cl3 DMBP Br2Cl4 DMBP Br3Cl2 1 DMBP Br3Cl2 4 DMBP Br3Cl3 DMBP Br4Cl 1 DMBP Br4Cl 2 DMBP Br4Cl2 DMBP Br5Cl DMBP 4Br 1

# **Dimethyl bipyrroles (DMBPs) [N] (cont'd)**

DMBP 4Br 2 DMBP 4Br 3 DMBP 5Br 1 DMBP 5Br 2 DMBP 6Br DMBP 6Cl

# **Hexachlorocyclohexane-related**

**(HCH-related) [A]** alpha BHC beta BHC cyclohexene 4Cl 1

# **Heptachlor-related [A]**

heptachlor epoxide heptachlor related 1 heptachlor related 2 heptachlor related 3

# **Methyl bipyrroles (MBPs) [N]**

MBP 4Br 1 MBP 4Br 2 MBP 4Br 3 MBP 4Br 4 MBP 6Cl 1 MBP 6Cl 2 MBP 6Cl 3 MBP 7Cl

# **Methoxy brominated/chlorinated**

**diphenyl ether (MeO-B/CDE) [N]** MeOBCDE Br3Cl

#### **Methoxy brominated diphenyl ether (MeO-BDE) [N]**

2'-MeOBDE-68 6-MeOBDE-47 MeOBDE 3Br MeOBDE 4Br

## **Methoxy polybrominated byphenyl (MeO-PBB) [N]** di-MeOPBB-80

**Methylenebistrichloroanisole [A]** Methylenebistrichloroanisole

#### **Methylsulfonyl polychlorinated biphenyl (Methylsulfonyl-PCB) [A]**

methylsulfonylPCB 5Cl 1 methylsulfonylPCB 5Cl 2 methylsulfonylPCB-101

## **Mirex-related [A]**

mirex mirex 1Cl mirex 2Cl 1 mirex 2Cl 2 mirex related 1

## **Polybrominated biphenyl (PBB) [A]**

BB-49 BB-52 BB-101 BB-153 BB 4Br 1 BB 5Br 2

# **Polybrominated diphenyl ethers**

**(PBDEs) [A]** BDE-17/25 BDE-28/33 BDE-47 BDE-49 BDE-66 BDE-75 BDE-99 BDE-100 BDE-116 BDE-153 BDE-154 BDE-155 BDE 5Br 1

# **Polybrominated hexadroxanthene**

**derivative (PBHD) [N]** PBHD 3Br 1 PBHD 3Br 2 PBHD 3Br 3 PBHD 4Br

#### **Pyrrolidinecarbonyl-chloride [A]**

pyrrolidinecarbonyl chloride

# **Tris(chlorophenyl)methane**

**(TCPM) [A]** TCPM TCPM 1 TCPM 2 TCPM 3 TCPM 4 TCPM 4Cl 1 TCPM 4Cl 2 TCPM 4Cl 3 TCPM 5 TCPM 6

# **Tris(chlorophenyl)methanol (TCPMOH) [A]**

TCPME

## **Unknown [U]**

unknown-11 unknown-13 unknown-14 unknown-15 unknown-17 unknown-21 unknown-22 unknown-27 unknown-32 unknown-34 unknown-35 unknown-36 unknown-38 unknown-39 unknown-41 unknown-43 unknown-44 unknown-46 unknown-47

unknown-49

# **Unknown [U] (cont'd)**

unknown-55 unknown-56

## **Unknown-2 [U]**

unknown-2-2 unknown-2-3 unknown-2-4 unknown-2-5 unknown-2-8 unknown-2-10

### **Unknown-3 [U]**

unknown-3-1 unknown-3-2 unknown-3-3

## **Unknown-4 [U]**

unknown-4-2 unknown-4-3 unknown-4-5 unknown-4-8 unknown-4-11 unknown-4-12 unknown-4-13 unknown-4-14 unknown-4-16 unknown-4-17

# **Unknown-5 [U]**

unknown-5-2

### **Unknown-6 [U]**

unknown-6-1 unknown-6-3

#### **Unknown-7 [U]**

unknown-7-1 unknown-7-2

# **Unknown-8 [U]**

unknown-8-2



Figure S1: A principal components analysis (PCA) plot of the samples according to maturity (mature/M and immature/I).



**Figure S2:** Regressions examining the relationship between PC1 (adjusted for age) and blubber testosterone (adjusted for gonad mass). Circles and triangles represent data from immature and mature animal samples, respectively.



Figure S3: The mean decrease in accuracy when removed from the model (%IncMSE) for all compound classes. Red bars represent significant variables with  $p < 0.05$ .



**Figure S4:** The mean decrease in accuracy when removed from the model (%IncMSE) for all compounds. Red bars represent significant variables with  $p < 0.05$ .



Figure S5: The relative abundance of individual compounds that significantly correlated with gonad mass adjusted testosterone concentration and had a variable importance greater than five. Variable importance in %IncMSE (I) and a p-value (p) is provided for each compound. Color denotes the origin of the compound (anthropogenic, unknown, or natural).

#### **REFERENCES**

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