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1 **Short-term fasts increase levels of halogenated flame retardants in tissues of a wild**  
2 **incubating bird**

3

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16

17

18 **Abstract**

19 Many species are adapted for fasting during parts of their life cycle. For species  
20 undergoing extreme fasts, lipid stores are mobilized and accumulated contaminants can be  
21 released to exert toxicological effects. However, it is unknown if short-term fasting events may  
22 have a similar effect. The objective of this study was to determine if short successive fasts are  
23 related to contaminant levels in liver and plasma of birds. In ring-billed gulls (*Larus*  
24 *delawarensis*), both members of the pair alternate between incubating the nest for several hours  
25 (during which they fast) and foraging, making them a useful model for examining this question.  
26 Birds were equipped with miniature data loggers recording time and GPS position for two days  
27 to determine the proportion and duration of time birds spent in these two activities. Liver and  
28 plasma samples were collected, and halogenated flame retardants (HFRs) (PBDEs and  
29 dechlorane plus) and organochlorines (OCs) (PCBs, DDTs, and chlordane-related compounds)  
30 were determined. Most birds (79%) exhibited plasma lipid content below 1%, indicating a likely  
31 fasted state, and plasma lipid percent declined with the number of hours spent at the nest site.  
32 The more time birds spent at their nest site, the higher were their plasma and liver concentrations  
33 of HFRs. However, body condition indices were unrelated to either the amount of time birds  
34 fasted at the nest site or contaminant levels, suggesting that lipid mobilization might not have  
35 been severe enough to affect overall body condition of birds and to explain the relationship  
36 between fasting and HFR concentrations. A similar relationship between fasting and OC levels  
37 was not observed, suggesting that different factors are affecting short-term temporal variations in  
38 concentrations of these two classes of contaminants. This study demonstrates that short fasts can  
39 be related to increased internal contaminant exposure in birds and that this may be a confounding  
40 factor in research and monitoring involving tissue concentrations of HFRs in wild birds.

41

42 **Keywords:** PBDEs; PCBs; Dechlorane Plus; lipid mobilization; Fasting; birds; incubation

43

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48

49

50 **Research Approval**

51 All experimental protocols with the ring-billed gull were approved by the Institutional

52 Committee on Animal Care of the Université du Québec à Montréal.

53 **Introduction**

54 For many vertebrate species, chronic and short term fasting occurs regularly as part of their  
55 life cycle such as during hibernation, post-weaning, incubation, moulting, and migration, or can  
56 be part of regular seasonal restrictions in food availability or foraging frequency. During periods  
57 of fasting or starvation, animals mobilize lipid reserves to fuel their daily energy requirements.  
58 Depending on the species, body state and size of lipid stores, this metabolic state can last  
59 anywhere from several hours to months (Wang et al. 2006). Because many halogenated organic  
60 contaminants are predominantly lipid soluble and thus bioaccumulative in lipid-rich tissues, they  
61 can be released through lipid mobilization leading to increased circulating levels (Birnbaum  
62 1985). Studies to date on the effects of fasting on contaminant toxicodynamics in free-ranging  
63 animals have focused mainly on species that undergo dramatic fasting events where individuals  
64 loose a large proportion of their total body lipids such as polar bears (*Ursus maritimus*) during  
65 hibernation (Polischuk et al. 2002) and female eider ducks (*Somateria mollissima*) that fast for  
66 several weeks during incubation (Bustnes et al. 2010). In these situations, lipid mobilization  
67 during fasting can lead to increased circulating levels of contaminants as they are released from  
68 fat stores (Birnbaum 1985). Several studies have reported increased enzyme-mediated liver  
69 metabolism of contaminants following fasting, further confirming that increased organismal  
70 exposure occurs during these natural food deprivation phases (Helgason et al. 2010; Jorgensen et  
71 al. 1999; Routti et al. 2013; Vijayan et al. 2006).

72 The effects of less severe fasting events on contaminant concentrations in birds have received  
73 limited attention to date particularly for wildlife species undergoing mild fasting events. Many  
74 birds are not adapted for long-term fasting and thus they must forage frequently to cover their  
75 high daily energy requirements. In such species, even short-term or milder fasts have the

76 potential to enhance lipid mobilization, which may as a result impact internal contaminant  
77 concentrations (e.g. Routti et al. 2013). However, herring gull chicks (*Larus argentatus*)  
78 exposed to a mixture of environmental contaminants and subsequently fed a reduced (by 70%)  
79 but not fully restricted diet for one week, demonstrated a 10% loss in body lipids coupled with an  
80 increase in contaminants and their metabolites in liver, brain, and plasma (Routti et al. 2013).  
81 This study showed that even incomplete fasting can reduce body fat stores, increase lipid  
82 mobilization and associated contaminant release in birds that are not adapted for high lipid  
83 storage. Conversely, white crowned sparrows (*Zonotrichia leucophrys*) exposed to 1,1,1-  
84 trichloro-bis(4-chlorophenyl)ethane (*p,p'*-DDT) in the lab for 5 days and subsequently fasted did  
85 not show increases in *p,p'*-DDT or its metabolites, *p,p'*-DDD and *p,p'*-DDE, in various tissues  
86 following 20 minute, 4 or 9 hour fasting regimes despite the loss of up to 19% of their body  
87 mass (Scollon et al. 2012). Further research on different contaminants, in different species and  
88 under varying fasting regimes is clearly needed to resolve this question.

89 Once fat-soluble contaminants are released from their association with lipids, they enter  
90 circulation and may be redistributed into target organs and tissues to exert toxicity (e.g. Bigsby et  
91 al. 1997). This means that during periods of fasting, animals may be at an increased risk of  
92 adverse effects which may be compounded by other ecological and physiological challenges  
93 including or resulting from food shortage (Wingfield 1994), migration, reproduction, or disease  
94 (Hall et al. 2008; Keller et al. 2006). Which species are at increased risk of exposure from  
95 contaminant release from fat stores, and in what life cycle phases this may occur, are still not  
96 well understood nor are the toxicokinetics of different contaminant classes in this situation.

97 The objective of the present study was to determine if short successive fasts related to  
98 incubation bouts were linked to contaminant (chlorinated and brominated) concentrations in liver

99 and plasma of an omnivorous bird species, the ring-billed gull (*Larus delawarensis*). We used  
100 ring-billed gulls breeding near the metropolis of Montreal (QC, Canada) because they have been  
101 shown to accumulate elevated levels of halogenated flame retardants (HFRs) (Gentes et al.  
102 2012). Moreover, this species undergoes biparental incubation (Ryder 2012), where both parents  
103 alternate between bouts of incubation (and thus fasting) that may last several hours, and foraging  
104 trips (Marteinson et al. 2015). Because these gulls do not exhibit prolonged continuous fasts  
105 (e.g., several days), we measured their time-activity budget using high-resolution GPS-based  
106 tracking during the last 24 hours prior to tissue collection. The relationships between liver and  
107 plasma contaminant concentrations and the amount of time spent fasting at the nest site or  
108 foraging away from the colony were investigated. To evaluate whether or not birds were in a  
109 fasted state, the percent plasma lipids was assessed, and the body condition was examined. We  
110 hypothesized first that, *i*) if short term fasting was to elicit lipid mobilization and increased  
111 contaminant release from lipid stores that contaminant concentrations would increase in plasma  
112 and liver as a function of the duration of time birds spend fasting at the nest site, and *ii*) that this  
113 increase would be related to changes in whole body lipid stores during fasting. Alternatively, if  
114 contaminant concentrations in plasma were dependent largely on recent dietary intake of  
115 contaminated prey, HFR and OC concentrations in blood would be expected to decline with the  
116 number of hours spent at the nest site (and thus time since feeding), or increase as the proportion  
117 of time spent foraging increased, and be related to time spent foraging in habitats of varying  
118 contamination.

119

## 120 **Materials and Methods**

121 *Model species and fieldwork*

122           During the incubation period (May-June 2011), ring-billed gull males ( $n = 16$ ) and  
123 females ( $n = 13$ ) were sampled on Deslauriers Island ( $45^{\circ}42'45''\text{N}$ ,  $73^{\circ}26'25''\text{W}$ ) located in the  
124 St. Lawrence River downstream of Montreal (QC, Canada). Approximately 44,000 ring-billed  
125 gull pairs breed on this island annually (P. Brousseau, personal communication). These  
126 omnivorous gulls use the surrounding mosaic of agricultural, urban and suburban areas where  
127 they feed opportunistically (Patenaude Monette et al., 2014; Caron-Beaudoin et al., 2013). The  
128 most prominent HFRs previously determined in these ring-billed gulls were PBDEs, for which  
129 the sum of 45 congeners averaged ( $\pm$  SEM)  $205 \pm 32.0$  ng/g wet weight (ww) in liver and  $27 \pm$   
130  $4.05$  ng/g ww in plasma, which is the highest level recorded in gull tissues to date in Canada  
131 (Gentes et al. 2012). Dechlorane plus (DP), a suggested deca-BDE alternative, was also  
132 determined in all of the samples at low levels in liver (*anti*-DP:  $6.06 \pm 1.64$ ; *syn*-DP:  $2.38 \pm 0.67$ ;  
133 (Gentes et al. 2012).

134           Two-hundred ring-billed gull nests with one egg were georeferenced and monitored  
135 daily. Once clutches were completed (i.e., three eggs), males or females were initially captured  
136 at random on their nests using a radio-controlled noose trap and the number of incubation days  
137 completed was back calculated. Each bird was weighed ( $\pm 0.01$  g) and morphometric  
138 measurements (head, culmen, and tarsus length) were recorded using digital callipers ( $\pm 1$  mm).  
139 A miniature GPS data logger (GiPSy2, TechnoSmArt, Guidonia, Roma, Italy) was then attached  
140 on two central tail feathers (rectrices), and was recovered at the second capture two to three days  
141 later. GPS units weighed 14-15 g, thus representing 2-4% of the ring-billed gull's body mass  
142 (mean  $\pm$  SEM:  $478 \pm 7$  g), which was shown not to influence the daily energy expenditure (via  
143 field metabolic rate measurement) of ring-billed gulls for the same individuals as utilized in the  
144 present study (Marteinson et al. 2015). At the initial and second captures, both occurring during



145 the incubation phase, blood samples were obtained (3 mL and 8 mL for capture and recapture,  
146 respectively) using a heparinized 25-gauge needle and 10 mL-syringe. Because blood collection  
147 volume from the initial capture had to be restricted due to the body size of these birds and  
148 because their behaviour was monitored thereafter, contaminant analysis in blood was only  
149 performed on one blood sample set (the second). Following the second blood collection, birds  
150 were euthanized, sexed by gonadal examination, and the liver was collected. Adipose tissue was  
151 not harvested because it is not reliably present in large enough amounts in this species during the  
152 incubation phase (it is sparsely distributed), thus precluding chemical analysis in all individuals.  
153 In the field, blood samples were kept in amber plastic vials in a cooler, and were centrifuged in  
154 the laboratory within 10 hours to obtain plasma for chemical analysis (section 2.4). Liver  
155 samples were also kept in a cooler in the field, and stored at -20°C until chemical analysis  
156 (section 2.4). Approval for all handling and sampling procedures was obtained by the  
157 Institutional Committee on Animal Care of the Université du Québec à Montréal, which  
158 followed the Canadian Council on Animal Care guidelines.

### 159 *Time-activity budget determination*

160 The same ring-billed gull individuals were used as those for which details on the activity  
161 budget and field metabolic rate have been previously reported (Marteinson et al. 2015). Briefly,  
162 the GPS data loggers recorded geographical positions ( $\pm$  5-10 m), velocity, date, and time at 4-  
163 minute intervals for two to three days (Caron-Beaudoin et al. 2013). Using combined  
164 information on velocity and position, each position was treated as an instantaneous behavioral  
165 sample for which the total time duration could be associated following Marteinson et al. (2015).  
166 Positions falling into the two activities of interest to address the present objective were defined  
167 as follows: 1) “Nest-site attendance” was assigned to all positions when birds were at the nest

168 site which included incubation as well as time spent near the nest (e.g., resting, guarding and/or  
169 preening), as the two behaviors could not be distinguished due to the spatial precision of the GPS  
170 data logger; and 2) “Foraging” was assigned to all positions when birds were away from the  
171 colony, but not flying. This coarse definition was used because specific activities such as  
172 walking, resting or preening similarly could not be distinguished (Marteinson et al. 2015).  
173 Though birds were GPS-tracked for 2-3 days, we assessed only positions from the last 24 hours  
174 prior to recapture. As a result, the first portion of each trajectory post-release (1-2 days) was not  
175 used in this experiment and any potentially confounding behavioural effects related to the initial  
176 capture (contention and handling) did not affect activity assessment. From the trajectories within  
177 the 24 hours prior to recapture, the proportions (%) of time spent in each of the two activities  
178 were calculated. Because birds were recaptured at the nest site, thus interrupting a bout of nest-  
179 site attendance, the number of hours that birds were attending the nest-site prior to recapture  
180 could also be determined. The proportion of time birds spent foraging was further categorized  
181 into habitat types utilized based on previous research in our laboratory (Caron-Beaudoin et al.  
182 2013; Gentes et al. 2015; Marteinson et al. 2015): 1) agricultural areas, 2) waterways (including  
183 the St. Lawrence River and other water bodies), and 3) urban areas (including urban, suburban,  
184 landfills, and wastewater treatment plant ponds).

### 185 *Body condition and nutritional state*

186 The body condition of ring-billed gulls was estimated via two methods. First, the  
187 condition index of individuals was determined for each sex separately by plotting the first  
188 principal component based on three morphometric measures (head, culmen, and tarsus length)  
189 against body mass, from which residuals were obtained and used as body condition index. This  
190 measure is useful in estimating body condition which will depend on how both muscle and fat

191 masses contribute to overall body mass. Second, body composition was estimated to determine  
192 the proportion of whole body lipids. To achieve this the determination of total body water  
193 percent by measuring the dilution of deuterium-labelled water in blood after a period of 1 hour  
194 following injection (described below) was conducted. Total body water percent has been shown  
195 to be a reliable non-destructive technique to estimate body condition in birds (McWilliams and  
196 Whitman 2013) and is strongly and negatively correlated with total body lipid content in birds  
197 and mammals (Farley and Robbins 1994; Mata et al. 2006) including gulls (glaucous gulls, *Larus*  
198 *hyperboreus*: (Shaffer et al. 2006); Verreault et al. 2007; Fig. S1). Briefly, as described in detail  
199 in Martinson et al. (2015), at the initial capture, 0.65 mL of water containing 37% deuterium  
200 (371,000 ppm) was injected into the pectoral muscle of the bird and after 1 hour of water  
201 equilibration in the body, a 3 mL blood sample was obtained. Duplicate 500  $\mu$ L aliquot samples  
202 of plasma were separated from whole blood and were assessed for  $^2\text{H}/^1\text{H}$  ratio using a  
203 Micromass Isoprime<sup>TM</sup> DI gas isotope ratio mass spectrometer (IRMS) coupled to a Aquaprep<sup>TM</sup>  
204 system using an equilibration method with the presence of hydrophobic platinum as a catalyst  
205 (Horita and Kendall 2004). Baseline levels of deuterium for this field site (colony) were  
206 determined from plasma of four control (uninjected) ring-billed gulls. The whole body water  
207 percent was calculated by dividing the dilution space for hydrogen (in grams of water) by body  
208 mass. The dilution space ( $N_d$ ) was calculated using the plateau approach as follows:  $N_d = \text{Mol}_{inj}$   
209  $(E_{mix} - E_{inj}) / (E_{wat} - E_{mix})$  where  $\text{Mol}_{inj}$  is the moles of  $^2\text{H}$  in the injectate,  $E_{mix}$  is the equilibrium  
210 enrichment of  $^2\text{H}$  in water (ppm of initial enrichment),  $E_{wat}$  is the background level of isotopes  
211 from control birds (in ppm) and  $E_{inj}$  is the enrichment of the injectate (in ppm) (eqn 17.11:  
212 Speakman 1997). The percent of whole body lipids was then calculated using the linear equation  
213 determined for the relationship between total body water mass and total body lipid mass in

214 glaucous gulls ( $y = -0.91x + 69.89$ ) which were inversely correlated ( $R^2 = 0.72$ ) (data from  
215 Shaffer et al. 2006 and Verreault et al. 2007; Fig. S1). The extractable percent lipid content was  
216 determined in plasma and liver as part of the chemical analysis (see section 2.4), and was used as  
217 a measure of recently acquired lipids through diet and as a proxy of fasting state. Plasma lipid  
218 content above 1% were used as a guideline to indicate whether birds had fed recently (K.  
219 Drouillard, unpublished data), whereas those below this threshold were likely to be in a fasted  
220 state, for example, fasted ring doves were found to have an average of 0.63% lipids in plasma  
221 while fasting (Drouillard and Norstrom 2000).

## 222 *Chemical analysis*

223 Ring-billed gull liver and plasma samples (both from the second capture) were analyzed  
224 for 37 PBDE congeners (BDE-1, -2, -3, -7, -10, -15, -17, -28, -49, -66, -71, -77, -85, -99, -100, -  
225 119, -126, 138, -139, -140, -153, -154/BB-153, -171, -180, -183/Dec-604, -184, -191, -196, -  
226 197/-204, -201, -203, -205, -206, -207, -208, and -209) as well as dechloranes (Dec) (Dec-602, -  
227 603, BDE-183/-604, and -604 CB), Chlordene Plus (CP) and Dechlorane Plus (*syn*- and *anti*-  
228 DP). Sample extraction and clean-up procedures were performed based on methods described  
229 previously in detail (Houde et al. 2014; Gentes et al 2012). Briefly, 1.0 g of liver or plasma was  
230 homogenized and ground with diatomaceous earth, and spiked with 200 ppb of an internal  
231 standard mixture (BDE-30, BDE-156,  $^{13}\text{C}$ -BDE-209,  $^{13}\text{C}$ -*syn*-DP, and  $^{13}\text{C}$ -*anti*-DP), and  
232 extracted with the solvent dichloromethane:*n*-hexane (50:50 volume ratio) using a pressurized  
233 liquid extraction system (Fluid Management Systems, Watertown, MA). The extractable lipid  
234 percentages in plasma and liver samples were determined gravimetrically. Samples were  
235 cleaned-up with a PBDE-free acid-basic-neutral column followed by a PBDE-free neutral  
236 alumina column (Fluid Management Systems). The identification and quantification of the

237 analytes was achieved using a gas chromatograph coupled to a mass spectrometer (GC-MS)  
238 (Agilent Technologies 5975C Series, Palo Alto, CA) in the electron capture negative ionization  
239 mode (ECNI). Quality assurance procedures included analysis of method blanks and standard  
240 reference material (SRM) (NIST 1947 Lake Michigan fish tissue) for each batch of ten samples.  
241 Background contamination of method blanks were low, however, blank correction was  
242 consistently performed for the following PBDEs: BDE-15, -17, -47, -49, -66, -71, -77, -85, -99, -  
243 100, -119, 126, -138, -139, -153, -154/BB-153, -171, -180, -183/Dec-604, -191, -196, -197/-204,  
244 -201, -203, and -209. The mean recovery of the internal standards in samples, blanks and SRMs  
245 were  $85.8 \pm 1.7\%$  for BDE-30,  $88.0 \pm 1.9\%$  for BDE-156,  $51.2 \pm 1.9\%$  for  $^{13}\text{C}$ -BDE-209,  $90.0 \pm$   
246  $2.2\%$  for  $^{13}\text{C}$ -*syn*-DP and  $90.2 \pm 2.2\%$  for  $^{13}\text{C}$ -*anti*-DP. Concentrations of PBDEs and other  
247 HFRs were quantified using an internal standard approach, and thus all analyte concentrations  
248 were inherently recovery-corrected. PBDE concentrations (seven congeners) determined in SRM  
249 showed less than 22% deviation from certified values.

250       Levels of OCs were reanalyzed in cleaned-up fractions obtained from liver tissue only  
251 (see sample preparation procedure above) because large enough volumes of plasma could not be  
252 collected from these mid-size birds. The compounds measured were: PCBs (CB-17, -18, -33, -  
253 44, -49, -52, -70, -74, -82, -87, -95, -99, -101, -110, -118, -128, -138, -149, -151, -153, -156, -  
254 158, -170, -171, -177, -180, -183, -187, -191, -194, -195, -206, -208, and -209),  
255 dichlorodiphenyltrichloroethane (*p,p'*-DDT) and its metabolites (*p,p'*-DDD, *p,p'*-DDE), Mirex,  
256 oxychlordan, *cis*- and *trans*-nonachlor, *cis*-chlordan, pentachlorobenzene (QCB),  
257 tetrachlorobenzene (1,2,3,4-TCB and 1,2,4,5-TCB), and octachlorostyrene (OCS). The  $^{13}\text{C}$ -*syn*-  
258 DP and  $^{13}\text{C}$ -*anti*-DP internal standards were used to recovery-correct OC concentrations as  
259 described above. Chemical analysis was performed using a Hewlett-Packard 5890 GC with 5973

260 mass selective detector (GC-MSD) operated in the electron impact (EI) mode, and using selected  
261 ion monitoring (SIM). For every batch of five samples, a MSD-PCB standard prepared from  
262 Aroclor 1242, 1254 and 1260 mixtures (AccuStandards, CT), two OC pesticide standards  
263 (Supelco, PA), a method blank and one in-house reference tissue (GLIER Detroit River fish  
264 homogenate) were also analyzed. PCB concentrations (32 congeners) determined in the in-house  
265 reference tissue showed less than 26% deviation from reference values.

#### 266 *Data analysis*

267 The complete data sets (including contaminant concentrations, time-activity budget, body  
268 condition indices, and percent lipids) were obtained for 16 males and 13 females for plasma  
269 contaminant concentrations and 14 males and 11 females for liver contaminant concentrations.  
270 All statistical analyses were conducted using wet weight (ww)-based concentrations of these  
271 lipophilic contaminants did not increase in proportion to tissue lipids (Hebert and Keenleyside  
272 1995). More specifically, for liver samples, HFR concentrations were unrelated to liver lipid  
273 percents (Pearson's Correlation Analysis;  $0.982 > p > 0.140$ ), which was consistent with OC  
274 concentrations (Spearman's Correlation Analyses;  $0.810 > p > 0.071$ ). Plasma HFR  
275 concentrations, were correlated with plasma lipid percents (see results below), although  
276 negatively, and thus opposite to what is predicted for lipophilic contaminants according to  
277 Hebert and Keenleyside (1995). Regardless, a parallel set of relationships between lipid percents  
278 and plasma HFR concentrations based on lipid-normalized data are presented in the  
279 Supplementary Information (Fig. S5) which demonstrates similar results to those based on ww  
280 concentrations. For HFRs, PBDE congeners that were determined (i.e. above the detection limit)  
281 in both liver and plasma of all individuals as well as the sum of all analyzed congeners  
282 ( $\Sigma_{37}$ PBDEs) were used in analyses. The  $\Sigma_{37}$ PBDEs were calculated by summing the

283 concentrations of all congeners above their respective detection limits. The sum of dechloranes  
284 was also assessed which, depending on their detection, for plasma included levels of *syn*- and  
285 *anti*-DP, and for liver it additionally included Dec-602, -603, and -604 CB as well as CP. For  
286 OCS, PCB congeners that were quantified in liver of all individuals were also analyzed as were  
287  $\Sigma_{38}$ PCBs, *p,p'*-DDE, *p,p'*-DDT, and  $\Sigma$  nonachlors (sum of *cis*- and *trans*-nonachlor).

288 Relationships between contaminant concentrations (plasma or liver) and the following  
289 variables were assessed using Spearman's Rank Correlation analyses: percent lipids in either  
290 liver or plasma, body condition measures (body condition index, whole body water and lipid  
291 percents), activity measures (time spent incubating or foraging and number of hours spent at the  
292 nest-site prior to recapture); correlation analyses between body condition and activity measures  
293 or the number of incubation days completed were also assessed. Spearman's Rank Correlation  
294 analyses were used for both sexes combined due to low sample size and to control for the effect  
295 of any outliers due to the ranking of the data. The contaminants were listed in order of  
296 decreasing *r* value to rank the strength of the relationships because different congeners will have  
297 varying halogenation and lipophilicity (expected to show differing behaviour during fasting).  
298 Data were additionally split into two groups defined as follows, and compared for contaminant  
299 concentrations and condition measures using t-tests: 1) birds in the lower (25%) and upper  
300 (75%) quartile groups based on percent plasma lipids and 2) birds in the lower (25%) and upper  
301 (75%) quartile groups based on hours spent in nest-site attendance.

302 To determine and rank which factors influenced the sum concentrations of the contaminant  
303 classes, a series of Generalized Linear Models (GLM) were conducted and ranked using Akaike's  
304 Information Criterion corrected for small sample sizes (AICc) (Burnham and Anderson 2002;  
305 Burnham et al. 2011). Variables that may affect contaminant levels were assessed including

306 temporal variables (capture date, Julian egg laying date), sex, body mass, % plasma lipids, and  
307 activity and body condition measures (body condition index, whole body water and lipid  
308 percents). Only models with one or two predictor variables as well as the intercept were  
309 conducted due to the low sample size. Related variables were not included in the same model.  
310 For comparison, the null model was assessed, and only models with  $AIC_c$  values lower than that  
311 of the null model were retained. Additionally, models for which predictor variables for which the  
312 parameter estimate 95% confidence intervals included zero were not considered. For each model,  
313 the  $\Delta AIC_c$  was calculated as well as the weight ( $w$ ) (Burnham and Anderson 2002). Simple  
314 linear regressions were also conducted with the variables for each model to generate adjusted  $R^2$   
315 values. Statistical analyses were conducted using IBM SPSS 20, and a 0.05 significance level  
316 was employed where applicable. Means are reported with standard error ranges.

## 317 **Results**

### 318 *Time-activity budget*

319 The proportions of time ring-billed gulls spent in nest-site attendance and foraging were  
320 consistent with those previously reported for a larger set of individuals including the present  
321 birds (Marteinson et al. 2015). Birds spent 47-90% of the 24-hour tracking period prior to  
322 recapture in nest-site attendance (mean:  $70 \pm 0.02\%$ ). The amount of time that birds had spent in  
323 this activity prior to their recapture at the nest ranged from 0.2 to 18.9 hours (mean:  $5.4 \pm 0.2$  h).  
324 Birds spent from 0 to 30% of the tracking period in foraging activities (mean:  $11 \pm 0.02\%$ ). The  
325 proportions of time ring-billed gulls spent foraging in different sites were consistent with those  
326 previously reported for a larger set of individuals, including the present birds (Gentes et al.  
327 2015). Gulls in the present sub-sample spent  $70 \pm 0.1\%$  of time foraging in agricultural areas, 29



328  $\pm 0.06$  % in urban areas and  $1 \pm 0.01$ % of their time on waterways in the last 24 hours prior to  
329 capture. Activities were unrelated to body condition measures (condition index, whole body  
330 water or lipid percent).

### 331 *Contaminant concentrations and profiles in liver and plasma*

332 Concentrations of HFRs (PBDEs and DP isomers) in the present ring-billed gull plasma  
333 and liver samples collected in 2011 were comparable to those previously reported for individuals  
334 sampled in 2010 (Gentes et al. 2012). The mean  $\Sigma_{37}$ PBDE concentrations in the present ring-  
335 billed gulls were  $138 \pm 22.2$  ng/g ww in liver (Fig. 1) and  $32.7 \pm 4.7$  ng/g ww in plasma. Among  
336 these, 13 PBDE congeners were quantifiable in all individuals in both liver and plasma including  
337 tetra-BDE (BDE-47), penta-BDEs (BDE-99 and -100), hexa-BDEs (BDE-138, -153, and -  
338 154/BB-153), hepta-BDE (BDE-183), octa-BDEs (BDE-196, -197, and -201), nona-BDEs  
339 (BDE-207 and -208), and deca-BDE (BDE-209). This congener profile was dominated by BDE-  
340 209, followed by the major penta-mixture congeners BDE-99, -47, -153, and -100 in descending  
341 order in both liver and plasma (Fig. S2). The percent contributions of these five major congeners  
342 to  $\Sigma_{37}$ PBDE concentrations in liver samples (with almost identical proportions in plasma) were  
343  $26 \pm 2\%$ ,  $24 \pm 1\%$ ,  $18 \pm 1\%$ ,  $7 \pm 0.4\%$  and  $6 \pm 0.4\%$ , respectively. The  $\Sigma$ dechlorane  
344 concentrations were  $6.6 \pm 1.3$  ng/g ww in liver (Fig. 1) and  $0.5 \pm 0.1$  ng/g ww in plasma.

345 Several OCs were detected in ring-billed gull liver samples. The mean  $\Sigma_{38}$ PCB  
346 concentrations in the gull livers were  $138 \pm 16.1$  ng/g ww (Fig. 1). A total of 19 PCB congeners  
347 were detected in the liver of all individuals, which included penta-CBs (CB-99, -118, -105, and -  
348 128), hexa-CBs (CB-138, -153, -158, and -156), hepta-CBs (CB-170, -171, -180, -183, and -  
349 191), octa-CBs (-194, -195, and -199) and the nona-CB-206. The congeners present in the

350 highest concentrations were CB-153, -138, -180, -187 and -118 in descending order (Fig. S3).  
351 Several other OCs were quantified in liver of all individuals (Fig. 1): *trans*-nonachlor (mean:  
352  $10.8 \pm 1.7$  ng/g ww), *cis*-nonachlor ( $0.7 \pm 0.2$  ng/g ww), *p,p'*-DDE ( $107 \pm 12.3$  ng/g ww), and  
353 *p,p'*-DDT ( $33.9 \pm 4.7$  ng/g ww) (Fig. 1). A few other OCs were detected in some of the  
354 individuals including QCB ( $n = 8$ ), *cis*-chlordane ( $n = 7$ ), *p,p'*-DDD ( $n = 5$ ), Mirex ( $n = 16$ ), and  
355 *trans*-chlordane ( $n = 1$ ), whereas 1,2,4,5-TCB, 1,2,3,4-TCB, OCS, and oxychlordane were not  
356 detected in any individuals. Mean concentrations of  $\Sigma_{37}$ PBDEs in liver were similar to  $\Sigma$   
357  $_{38}$ PCBs in this tissue (Fig. 1).

#### 358 *Liver, plasma and whole body lipid contents and contaminant concentrations*

359 The mean extractable plasma lipid percent of ring-billed gulls was  $0.8 \pm 0.04\%$  (range:  
360  $0.07 - 1.4\%$ ). The majority of birds (79%;  $n = 23$ ) exhibited plasma lipid content below 1%,  
361 with six individuals having percent lipids above that fasting state guideline level (Fig. 2).  
362 However, the plasma lipid percent declined significantly with the number of hours birds had  
363 spent on their nest site prior to recapture ( $r = -0.40$ ,  $p = 0.033$ ), which corresponded to a 0.02%  
364 decline in lipid percentage per hour of nest-site attendance (Fig. 2). Based on the patterns  
365 observed, plasma lipid contents between 0.5 and 0.8% were the most commonly encountered and  
366 no birds had plasma lipids exceeding 0.8% after their nest-site attendance lasted  $\geq 8$  hours.

367 The mean liver lipid percent of ring-billed gulls was  $6.0 \pm 0.2\%$  (range: 4.1-8.5%). Liver  
368 lipid percent was unrelated to the number of hours birds had fasted prior to recapture. Whole  
369 body lipid content estimated by the deuterium labelled water method was  $23.6 \pm 2.4\%$  (range:  
370 17.4-28.3%). Similar to what was observed for liver lipids, whole body lipids did not show any  
371 significant relationship with the number of hours birds had fasted prior to recapture.

372 The plasma lipid percent was negatively associated with the plasma concentrations of  
373 several individual PBDEs (in order of decreasing  $r$ : BDE-154/BB-153, -209, -47, -196, -197, -  
374 201, -208, and -207;  $-0.31 < r < -0.37$ ;  $0.014 < p < 0.050$ ),  $\Sigma_{37}$ PBDEs ( $r = -0.33$ ,  $p = 0.031$ ) (Fig.  
375 3B) as well as  $\Sigma$ dechloranes ( $r = -0.39$ ,  $p = 0.022$ ). Similarly, liver concentrations of several  
376 individual PBDE congeners (in order of decreasing  $r$ : BDE-183/Dec-604, -153, -154/BB-153, -  
377 196, -197/204, -100, -47, -201, -209, and -138;  $-0.33 < r < -0.43$ ;  $0.007 < p < 0.046$ ).  $\Sigma_{37}$ PBDEs  
378 ( $r = -0.40$ ,  $p = 0.013$ ) (Fig. 3A) and  $\Sigma$ dechloranes ( $r = -0.35$ ,  $p = 0.03$ ) were also negatively  
379 associated with plasma lipid content. Despite the variation in bromine content of the various  
380 PBDE congeners (from tetra- to deca-brominated) that were related to plasma lipid percent,  
381 correlations all showed a very similar  $r$ , and thus only the relationships between plasma lipid  
382 percents and  $\Sigma_{37}$ PBDEs in liver or plasma are displayed graphically for brevity (Fig. 3). Birds  
383 with low plasma lipid percents (25% quartile;  $n = 13$ ) ranged from 0.4-0.7% and those with high  
384 plasma lipid percents (75% quartile,  $n = 11$ ) had mean levels above the 1% threshold (0.93-  
385 1.43%). The mean  $\Sigma_{37}$ PBDEs in liver and plasma of low plasma lipid birds was  $74.4 \pm 19.4$  ng/g  
386 ww and  $41.0 \pm 6.7$  ng/g ww which were 2.3 and 2.1 times higher, respectively, compared to high  
387 lipid plasma birds (mean liver:  $33.1 \pm 10.7$  ng/g ww; plasma:  $19.4 \pm 4.6$  ng/g ww). The  
388 difference between low and high plasma lipid birds was significant for  $\Sigma_{37}$ PBDEs in plasma ( $t_{22}$   
389 = 2.55,  $p = 0.018$ ) with a similar, but non-significant trend for liver concentrations ( $t_{18} = 1.86$ ,  $p$   
390 = 0.079).

391 PCB and OC pesticide concentrations in liver were not significantly related to the lipid  
392 percents in plasma (Fig 3). For these compounds, birds grouped into low plasma lipid categories  
393 showed mean  $\Sigma_{38}$ PCB concentration of  $161 \pm 51.5$  ng/g ww in liver compared to  $134 \pm 32.0$  ng/g

394 for birds in the high plasma lipid category, representing a 1.2 times decrease which did not differ  
395 significantly.

396

397 *Relationships between contaminant concentrations and time-activity budget*

398 With respect to the HFRs, as the proportion of time that birds spent in nest-site  
399 attendance increased, so did their concentrations of  $\Sigma_{37}$ PBDEs in both liver ( $r = 0.49$ ,  $p = 0.013$ )  
400 and plasma ( $r = 0.44$ ,  $p = 0.020$ ). Several of the major congeners in liver (in order of decreasing  
401  $r$ : BDE-99, -197/204, -201, -209, -153, -47, -154/BB-153, -196, -183/Dec-604, -138, -208, -100,  
402 and -207:  $0.49 < r < 0.57$ ;  $0.003 < p < 0.033$ ) and plasma (in order of decreasing  $r$ : BDE-100, -  
403 153, -154/BB-153, -47, -209, -153, and -99:  $0.39 < r < 0.43$ ;  $0.022 < p < 0.042$ ) also significantly  
404 increased with the proportion of time birds spent in nest-site attendance. The positive correlation  
405 between nest-site attendance and  $\Sigma_{37}$ PBDE concentrations was stronger when the number of  
406 hours at the nest site before recapture was considered for both liver ( $r = 0.56$ ,  $p = 0.004$ ; Fig. 4A)  
407 and plasma concentrations ( $r = 0.57$ ,  $p = 0.002$ ; Fig. 4B, with a similar figure for lipid-corrected  
408 values in plasma: Fig. S5).  $\Sigma_{37}$ PBDE concentrations in this sub-sample of the colony increased  
409 on average by 18.0 ng/g ww per hour spent on the nest site prior to capture in liver and 2.5 ng/g  
410 ww on average per hour in plasma (Fig. 5). Most of the major congeners in liver (in order of  
411 decreasing  $r$ : BDE-153, -154/BB-153, -138, -99, -100, -197/-204, -209, -201, -183/Dec-604, -  
412 196, -207, and -47:  $0.44 < r < 0.60$ ;  $0.001 < p < 0.033$ ) and plasma (in order of decreasing  $r$ :  
413 BDE-209, -154/BB-153, -153, -99, -153, -100, -47, -196, -197, -207, -208, and -201:  $0.41 < r <$   
414  $0.57$ ;  $0.002 < p < 0.032$ ) also increased with the number of hours spent at the nest site before  
415 recapture. Additionally,  $\Sigma$ dechlorane concentrations in liver were positively related to the  
416 proportion of time spent at the nest-site ( $r = 0.40$ ,  $p = 0.050$ ). Birds with low nest site attendance

417 (25% quartile,  $n = 7$ ) were present at the nest site for 1.5 hours or less (mean:  $0.8 \pm 0.4$ ) and  
418 those with high nest-site attendance (75% quartile,  $n = 7$ ) were present at the nest for 7.3 hours or  
419 more (mean:  $13.2 \pm 1.6$  h). Low nest-site-attendance birds exhibited a mean  $\Sigma_{37}$ PBDE  
420 concentrations of  $14.7 \pm 3.0$  ng/g ww in plasma and  $52.7 \pm 13.2$  ng/g ww in liver which were 3.8  
421 and 5.2 times lower, respectively, compared to those in the high nest-site attendance group  
422 (mean plasma:  $56.2 \pm 15.4$  ng/g ww; mean liver:  $275 \pm 105$  ng/g ww). Concentrations of OCs  
423 were unrelated to nest-site attendance measures or the proportion of time spent foraging. Similar  
424 to what was reported by Gentes et al. (2015) for a larger set of ring-billed gulls (which included  
425 the present individuals), the proportion of time spent foraging in agricultural and urban areas as  
426 well as in waterways were not related to HFR concentrations. Similarly, OC concentrations in  
427 liver were unrelated to the proportion of time spent in these three different foraging habitats.

#### 428 *Relationships with body condition measures*

429 Body condition index (mean:  $-0.71 \pm 3.74$  g), total body water percent (mean:  $51 \pm$   
430  $0.42\%$ ) and total body lipid percent (mean:  $23.61 \pm 0.38\%$ ) were not related to nest-site  
431 attendance measures, proportion of time spent foraging, number of incubation days completed or  
432 contaminant concentrations, nor were they related to one another.

433

#### 434 *Model selection for $\Sigma$ PBDEs in liver and plasma*

435 AIC analysis was conducted exclusively for PBDEs because only this contaminant class  
436 showed relationships with predictor variables as described above (Table 1). The variation in  
437 liver concentrations of  $\Sigma_{37}$ PBDEs were best explained by the model including the number of  
438 hours spent at the nest-site prior to recapture in combination with the capture date ( $w = 0.43$ ),

439 which explained 38% of the variation. The remaining models that showed significant effects all  
440 had considerably lower AIC weights in comparison to the top models ( $w = 0.12-0.02$ )  
441 demonstrating their minimal effect on liver  $\Sigma_{37}$ PBDEs in models containing plasma lipid % ( $w$   
442  $= 0.11-0.05$ ) and the sex ( $w = 0.02$ ). The variations in  $\Sigma_{37}$ PBDEs concentrations in plasma  
443 were best explained by the proportion of time birds had spent in nest-site attendance ( $w = 0.29$ )  
444 which accounted for 18% of the variation. The second best model closely followed and was the  
445 same as the top model for  $\Sigma_{37}$ PBDEs in liver- i.e., the number of hours spent at the nest-site in  
446 combination with capture date ( $w = 0.22$ ). After these two top models, AIC weights ranged from  
447 0.15 to 0.07, thus comparatively having a minimal effect on plasma  $\Sigma_{37}$ PBDE concentrations.  
448 Sex explained only a small proportion of the variations in plasma  $\Sigma_{37}$ PBDEs ( $w = 0.15$ ). Julian  
449 lay date, body mass and measures of body condition did not explain any variation in  $\Sigma_{37}$ PBDE  
450 concentrations in both plasma and liver. Plasma % lipids did not explain any variation in plasma  
451  $\Sigma_{37}$ PBDEs.

## 452 **Discussion**

453 This study demonstrates that time-activity budgets of wild birds can influence their  
454 contaminant concentrations in liver and plasma. More specifically, the percentage of time spent  
455 in nest-site attendance, rather than the time engaged in foraging activities, was significantly  
456 related to increased plasma and liver concentrations of PBDEs and DP isomers in ring-billed  
457 gulls during the incubation period. This suggests that short nest-site attendance bouts in birds  
458 have a greater and previously underestimated influence on within-tissue concentration variations  
459 compared to foraging, at least with respect to HFRs. However, unexpectedly, a similar pattern of  
460 enrichment of OCs in liver as a function of time spent in nest-site attendance did not occur

461 despite the fact that log  $K_{ow}$  values of the contaminant classes are largely overlapping: e.g., 4.6-  
462 6.9 for PCBs (Han et al. 2006), 5.75 for DDT (Veith et al. 1979), and 5.7-8.3 for PBDEs  
463 (Braekevelt et al. 2003; Han et al. 2006). This may imply different causal factors contributing to  
464 short-term temporal variations in tissue-specific contaminant levels between present suite of OCs  
465 and HFRs. Plasma lipids decreased with time spent fasting as would be predicted, and HFR  
466 concentrations increased with time spent fasting which agrees in part with our first hypothesis.  
467 However, this did not occur concurrently with the required changes in body condition during  
468 fasting in these gulls, that is, there was no relationship between whole body lipids and time spent  
469 fasting (incubating) or HFR concentrations. Therefore, we must reject our first hypothesis that  
470 short-term fasting caused whole body lipid mobilization and contaminant release in these  
471 incubating gulls. The lack of relationship between time spent foraging or in different habitats,  
472 which should have varying contamination (e.g. urban vs. agricultural), with plasma or liver HFR  
473 or OC concentrations implies that we must also reject our second hypothesis predicting that  
474 recent intake of a contaminated diet may be at the basis for the relationships between HFRs and  
475 time spent fasting in these birds. These results suggest that other underlying mechanisms may  
476 explain the relationships between HFR concentrations and time spent fasting (incubating) in  
477 ring-billed gulls. These other possible mechanisms may be related to the effect of time of capture  
478 during the incubation phase or differing depuration rates related to metabolic biotransformation  
479 of individual PBDE congeners in the gulls. Further study on these avenues and others is  
480 warranted and larger sample sizes are needed.

481 Few studies have been conducted on the effects of fasting on contaminant concentrations in  
482 birds. A few examples demonstrate that concentrations of contaminants increase during  
483 prolonged periods of food restriction. For example, in common eiders, where the female fasts

484 continuously for four weeks during the incubation, levels of PCB-153, *p,p'*-DDE and  
485 hexachlorobenzene increased between day five and day 20 in the incubation period (Bustnes et  
486 al. 2010). Moreover, in overwintering greater scaup (*Aythya marila*), which experience reduced  
487 availability and simultaneous reductions in body fat stores, individuals captured later in the  
488 winter had higher wet weight-based levels of  $\Sigma$ PCBs and  $\Sigma$ DDTs (including DDD, DDE) than  
489 birds caught early in the winter (Perkins and Barclay 1997). This effect has been confirmed even  
490 in species that are not adapted for extreme fasts. For instance, herring gull chicks exposed to a  
491 mixture of contaminants (OCs and PBDEs) in the laboratory and subsequently fasted for one  
492 week, exhibited contaminant concentrations that were three times higher in liver, plasma, and  
493 brain compared to a non-fasted exposed group (Routti et al. 2013). Changes in feeding rates and  
494 body mass throughout the breeding season have similarly been associated to contaminant  
495 concentrations in some seabirds, both those adapted to long fasts or not, including kittiwakes  
496 (*Rissa trydactyla*) (Henriksen et al. 1996), and Adélie penguins (*Pygoscelis adeliae*) and  
497 southern fulmars (*Fulmarus galacialoides*) (van den Brink et al. 1998). More specifically, in the  
498 penguin and fulmar study, lower body mass during times of fast (egg-laying or incubation) or  
499 times of higher physical exertion (chick-provisioning) were associated with higher levels of  
500 several OCs in blood and uropygial oil compared to periods where birds were able to replenish  
501 fat stores (van den Brink et al. 1998). Similarly for the kittiwakes during breeding, levels of  
502  $\Sigma$ PCBs in the brain were four-fold higher at the end of chick-rearing period during which they  
503 lost 20% of their body mass compared to the pre-breeding period (Henriksen et al. 1996).  
504 Collectively, these studies demonstrate that relationships exist between fasting and contaminant  
505 concentrations, but the processes behind these remain difficult to explain.



506 Contaminant mobilization during tissue lipid metabolism is commonly invoked as a  
507 mechanism describing the increase in tissue contaminant concentrations post-fasting (Daley et al.  
508 2014). When animals cannot obtain energy from food, they must draw upon stored energy  
509 reserves. The loss of lipids due to metabolism of triglycerides stored in adipose tissue, which  
510 reflects the main proximate component contributing to the partitioning capacity of organisms for  
511 hydrophobic contaminants, results in a fugacity gradient between adipose tissue and blood. This  
512 favors net diffusion of chemicals from adipose tissue until a new inter-tissue equilibrium is  
513 achieved. The process of lipid mobilization during fasting typically follows a defined sequence  
514 of events. In the first metabolic phase of fasting, stored liver glycogen and fatty acids from fat  
515 stores are used to supply glucose to tissues (Wang et al. 2006). However, these sources of  
516 energy are rapidly depleted (from hours to days) after which the organism will enter the second  
517 phase of food deprivation in which glycerol is released from adipose tissue and used for the  
518 production of glucose (Wang et al. 2006). This second phase is longer and, depending on the  
519 species and size of lipid stores, can last weeks to months (Wang et al. 2006). Fasting birds  
520 typically enter phase two after a few days of fasting - two days in red-legged partridges  
521 (*Alectoris rufai*) (Rodriguez et al. 2005) which are not well-adapted to fasting condition, four  
522 days in gentoo penguins (*Pygoscelis papua*) which are adapted to short-term fasts, and seven  
523 days in King penguins (*Aptenodytes patagonica*) which are adapted to several months of fasting  
524 (Cherel et al. 1993). Gulls (Larids) fall into the middle of this spectrum, entering phase two after  
525 approximately four to five days, as documented for example in herring (Totzke et al. 1999) and  
526 yellow-legged gulls (*Larus cachinnans*) (Alonso-Alvarez and Ferrer 2001).

527 During lipid loss, contaminant levels become concentrated in adipose tissue as lipids are  
528 depleted leading to an elevation in chemical fugacity. This creates a fugacity gradient between

529 adipose and blood that favors chemical diffusion and mobilization from the major storage  
530 compartment (fat) to other tissues in the body (Daley et al. 2014). For most bioaccumulative  
531 contaminants, inter-tissue distribution kinetics are considered rapid compared to whole body  
532 elimination. Therefore, the kinetics associated with tissue-to-tissue transfer are commonly  
533 ignored when formulating simple, non-physiologically based bioaccumulation models (Selck et  
534 al. 2013). For example, in the ring dove (*Streptopelia risoria*), rapidly perfused organs such as  
535 brain, liver and gonads achieved 90% steady state with blood PCB-153 in less than three days,  
536 while fat and carcass required ten days to achieve steady state with blood (Daley et al. 2013). In  
537 contrast, whole body elimination rates for PCB-153 in the same species was reported to be on the  
538 order of 2.9 to 3.7 years (Drouillard and Norstrom 2003). The rate at which contaminants are  
539 released from fat stores during fasting and how this differs between different chemicals remains  
540 largely unknown. Some evidence indicates that not all contaminant classes follow the same  
541 dynamics of release during lipolysis. For example, in fasting polar bears, some OCs increased in  
542 concentrations in adipose tissue during fasting including  $\Sigma$ chlordanes and PCBs, suggesting that  
543 they are less readily released from this compartment, whereas others declined such as  $\Sigma$ DDTs  
544 and  $\Sigma$  hexachlorocyclohexanes, suggesting that they were released more easily into circulation  
545 and/or were potentially eliminated from the body at a higher rate through metabolic  
546 biotransformation (Polischuk et al. 2002). There is some evidence to suggest that contaminants  
547 with lower log  $K_{ow}$  (i.e., are less lipophilic) may be released more efficiently from blubber in  
548 grey seals (*Halichoerus grypus*) (Vanden Berghe et al. 2012).

549         However, in the present study, there is no evidence to support mobilization of lipid stores  
550 as the mechanism behind the relationships between HFR concentrations and time spent fasting.  
551 For these birds, the maximum time ring-billed gulls spent on their nesting site was in the order of

552 20 hours, which can be enough to weight loss in birds that are not adapted for prolonged fasting  
553 (e.g. Scollon et al. 2012). However, even though individual ring-billed gulls would have  
554 underwent several such relatively short fasting events over their 28 days incubation period, there  
555 was no relationship between body condition expressed as whole body lipids and time spent at the  
556 nesting site, body condition and number of incubation days nor between body condition  
557 measures and contaminant concentrations in liver or plasma. This points to a lack of condition  
558 deterioration over time during this phase and suggests that these birds remained within the first  
559 metabolic phase of fasting, and were relying more heavily on fatty acids and glycogen reserves  
560 compared to stored triglycerides. Loss of fatty acids from adipose tissue would also decrease the  
561 partitioning capacity of the tissue, but to a lesser extent than triglycerides (Dulfer and Govers  
562 1995; Dulfer et al. 1996). Ring-billed gulls forage between short-term incubation bouts, and tend  
563 to exploit areas of high and predictable food availability (e.g., anthropogenic-related habitats and  
564 agricultural fields) and quality (fat and protein content) (Caron-Beaudoin et al. 2013; Patenaude-  
565 Monette et al. 2014). This may have allowed them to replenish labile tissue energy stores as they  
566 were used up on a daily basis. This is in line with other seabird species exhibiting biparental care  
567 that maintain body mass during the incubation period (e.g., Moreno 1989) despite prolonged  
568 bouts of intermittent fasting at the nest. Given that whole body lipids in birds of the present study  
569 did not change significantly in relation to time devoted to nest-site attendance, fugacity gradients  
570 generated as a result of adipose tissue depletion cannot explain the increasing trend of plasma  
571 and liver HFR concentrations with time spent at the nest site. This conclusion is reinforced by  
572 the fact that OCs showed no evidence of enriched liver concentrations. Mobilization of  
573 contaminants from lipid-rich tissues during weight loss would be expected to increase all  
574 hydrophobic contaminant concentrations (including OCs and HFRs in present birds) in tissues to

575 a similar extent. For example, Routti and colleagues (2013) exposed herring gull chicks to a  
576 mixture of environmentally relevant contaminants for 45 days and subsequently fasted (70%  
577 food reduction) treatment birds for seven days to compare chemical enrichment in tissues  
578 relative to non-fasted individuals (controls). The above study demonstrated that all contaminant  
579 classes including  $\Sigma$ PCBs,  $\Sigma$ DDTs,  $\Sigma$ CHLORs and  $\Sigma$ PBDEs increased in liver, brain, and plasma  
580 over those of the non-fasted group to a similar extent.

581 It is more likely that the relationships between HFR concentrations and time spent fasting  
582 in ring-billed gulls at the nest site was related to recent contaminant exposure which may have  
583 changed as the spring season progressed. Three alternative explanations would be that ring-  
584 billed gulls had been: i) exposed to a “diluted meal” during foraging activities rather than the  
585 contaminated diet we predicted from the habitats they use, ii) exposed to a non-dietary source of  
586 HFRs at the nesting site itself or iii) exposure to PBDEs may have altered the incubation  
587 behavior of these gulls. In the diluted meal hypothesis, birds returning from foraging and which  
588 have fed on a relatively clean meal (low HFR concentrations) would be expected to experience  
589 dilution of plasma contamination due to an influx of clean dietary lipids (e.g., portomicrons)  
590 and/or production of *de novo* lipids generated from assimilated nutrients (Drouillard and  
591 Norstrom 2000; Drouillard and Norstrom 2001). In this scenario, differences in the behavior of  
592 different contaminant classes might be explained by the fact that the diluted meal was less  
593 contaminated with HFRs, but not OCs. Based on Fig. 2, peak plasma lipid contents were  
594 observed in birds had been incubating for less than a few hours prior to capture, followed by  
595 depletion of plasma lipids to apparent fasting levels after eight to ten hours. This profile is very  
596 similar to what was observed in ring doves after achieving  $C_{max}$  of plasma lipids following a  
597 controlled feeding study that demonstrated an exponential return of plasma lipids approaching

598 fasting levels after approximately seven hours (Drouillard and Norstrom 2000). This would be  
599 surprising because these birds do not rely heavily on aquatic habitats for feeding which act as  
600 sinks for PCBs (reviewed in: Beyer and Biziuk 2009), but they do frequently utilize  
601 anthropogenic-related habitats to forage (e.g., Caron-Beaudoin et al., 2013) which can be highly  
602 contaminated with HFRs (Venier and Hites 2008). As further evidence, recent (last 24 hours)  
603 use of landfills and wastewater treatment plant ponds has been linked to increased concentrations  
604 of BDE-209 in these gulls nesting in the same colony as utilized in the present study (Gentes et  
605 al. 2015).

606         However, the contaminant patterns in liver of ring-billed gulls do not support this  
607 hypothesis. Dilution of liver lipids post-feeding is also possible given that liver is capable of  
608 generating lipids from absorbed carbohydrates and proteins. Indeed, lipid-normalized  
609 egg/maternal tissue concentration ratios in herring gulls and other bird species have been  
610 hypothesized to occur as a result of dilution of the liver lipid pool during yolk production  
611 resulting from *de novo* lipid production in the liver (Braune and Norstrom 1989; Drouillard and  
612 Norstrom 2001; Norstrom et al. 1986). In the case of herring gulls, a mean lipid-normalized  
613 egg/maternal tissue ratio of 0.7 was apparent, suggesting a 1.4-fold dilution factor of liver OC  
614 levels during yolk production (Braune and Norstrom 1989). A much higher egg/maternal tissue  
615 dilution factor of 2.9-fold was observed in ring doves which approaches the 5.2-fold differences  
616 in liver HFR residues observed in ring-billed gulls from the present study. However, the period  
617 of yolk formation is considered an extreme case where *de novo* lipid production in the liver is  
618 expected to be maximized and does not likely reflect the situation present for incubating ring-  
619 billed gulls. Second, there was no evidence for a change in liver lipid content with time at the  
620 nesting site, which would be expected to be observed under a situation of lipid dilution in liver.

621 Finally, *de novo* lipid production by the liver would dilute all hydrophobic chemicals to the same  
622 extent, but only HFRs and not OCs showed changes in liver concentration with time on the  
623 nesting activity in ring-billed gulls. Thus, the “diluted meal” hypothesis could be considered  
624 consistent with the observed trends in plasma concentrations of HFRs and lack of trends for OCs,  
625 although not consistent with the patterns observed for liver.

626 The second hypothesis implies that ring-billed gulls were exposed to elevated HFR and  
627 baseline OC concentrations while at the nesting site. The exposure source was apparently not  
628 related to dietary exposure at the nesting site because the change in plasma lipid content with  
629 time at the nest is consistent with birds reverting to a fasting state during this activity.  
630 Alternative exposure routes could be as a result of inhaled HFR-contaminated dusts/particles or  
631 air at the local nesting site. Assimilated material from the respiratory tract would be expected to  
632 show up rapidly in perfused tissues including blood and liver, followed by slower redistribution  
633 to adipose tissues (Daley et al. 2013). HFRs can be present in the atmosphere in their volatilized  
634 form which is more common for lower-brominated congeners, or in association with  
635 dust/particles which is more common for higher-brominated congeners including BDE-209  
636 (reviewed in: Hale et al. 2006). Several HFRs including PBDEs have been measured in outdoor  
637 air globally (reviewed in: Hale et al. 2006) with the highest concentrations recorded outside  
638 point-source areas in urbanized environments (e.g., Venier and Hites 2008). Recently, PBDE  
639 concentrations in outdoor air have begun to exceed concentrations of PCBs in several reports  
640 (reviewed in: Hale et al. 2006). This is consistent with the finding that the present ring-billed  
641 gulls demonstrated similar mean  $\Sigma$ PBDEs concentrations to  $\Sigma$ PCBs, the latter of which have  
642 previously dominated in bird tissue. Few studies have measured HFRs in Canadian city air  
643 though, recently in Toronto, air contained 38 pg/m<sup>3</sup>  $\Sigma$ PBDEs in combined gas and particle

644 phases, as well as a number of emerging HFRs including *syn*- and *anti*-DP (Shoeib et al. 2015).  
645 Concentrations of contaminants in the atmosphere can be affected by temperature and weather as  
646 well as other atmospheric variables (e.g. (Melymuk et al. 2012) and references therein) and since  
647 these change throughout the spring season when incubation takes place in the present location,  
648 this may play a role in the identified effect of capture date on PBDE concentrations in these  
649 gulls. However, the diet of these birds also changes as the spring progresses (Patenaude-Monette  
650 et al. 2014), and further research is required to determine how the capture date may be related to  
651 contaminant exposure in these birds. The contribution of atmospheric exposure to HFR  
652 contamination in wildlife is largely unknown. However, it has been confirmed as an important  
653 source of exposure to humans via indoor air and house dust (reviewed in: (Hale et al. 2006)), and  
654 thus wildlife living in urban areas may be receiving significant (and underestimated) exposure to  
655 HFRs via air and associated particles and via ingestion of these contaminated particles during  
656 preening. Further research on this avenue of exposure to HFRs in ring-billed gulls is currently  
657 ongoing in our laboratory to determine how much this route of exposure may be contributing to  
658 their body burdens.

659         Finally, the possibility that exposure to PBDEs may have altered the incubation or nest-  
660 site attentiveness of these gulls should be explored as they have been shown to disrupt related  
661 endpoints. In laboratory studies, exposure to PBDEs has caused reduced nest-related behavior  
662 during courtship and brood rearing in American kestrels (*Falco sparverius*) (Ferne et al. 2008,  
663 Martinson et al. 2010), as well as reduced nest temperatures during incubation (Sullivan et al.  
664 2013). Additionally, in wild male glaucous gulls, greater plasma sum PBDE concentrations were  
665 associated with reductions in the hormone prolactin which regulates incubation behavior  
666 (Verreault et al., 2008, and references therein). As such to date, exposure to PBDEs appears to

667 have a negative impact on incubation and prolactin levels in birds, suggesting that this may not  
668 explain the positive correlation between PBDEs and time spent at the nest in the present ring-  
669 billed gulls, however further research on this avenue may be warranted.

670

## 671 **Conclusions**

672 This study demonstrates increased HFR concentrations in ring-billed gull plasma and  
673 liver tissues during the incubation period while birds were present at the nesting site. More  
674 specifically, bouts of nest-site attendance in ring-billed gulls lasting only a few hours were  
675 associated with significant increase in levels of PBDEs and DP isomers (liver and plasma), but  
676 not OCs (liver). Conversely, the amount of time that birds were engaged in foraging activities,  
677 body condition and percent body fat were unrelated to contaminant concentrations. The changes  
678 in plasma HFR concentrations may be consistent with the hypothesis that birds were exposed to  
679 a diluted meal with respect to HFR levels, although this would not apply for OCs. However, the  
680 patterns in liver do not support this explanation because all hydrophobic organic contaminants  
681 should increase in liver as fasting duration increases. Alternatively, these gulls may have been  
682 significantly exposed to non-dietary sources of HFRs at the nest site (i.e., via inhalation of air  
683 and HFR-laden particles or ingestion via preening of particles adsorbed to their plumage).  
684 Ultimately, further study is needed to understand the increased liver and plasma HFR  
685 concentrations in these birds while they spend time fasting at the nest site. However, regardless  
686 of the underlying mechanisms and exposure routes, the results presented herein highlight the fact  
687 that short-term fasting events related to, for example nest-site attendance, may significantly  
688 influence levels of HFRs in birds, and hence the potential toxicity associated with these  
689 compounds. As such, the amount of time spent in this activity just prior to capture may become



690 a confounding factor in monitoring and research on contaminants, including biological effect  
691 studies in wild birds. This may also apply to a wide variety of other situations where short-term  
692 fasting occurs including other reproductive phases, migration, or even daily overnight fasting.  
693 Further research on how contaminant concentrations relate to these life cycle stages is warranted.

694

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702 authors thank M. Patenaude-Monette and M.-L. Gentes.

703

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863



864 **Table 1:** Generalized linear models ranked by Akaike's Information Criterion for small sample sizes ( $AIC_c$ ) and linear regression  
865 adjusted  $R^2$  for variables that explain  $\Sigma_{37}$ PBDEs determined in plasma and liver of ring-billed gulls (*Larus delawarensis*) breeding  
866 near Montreal (QC, Canada).

Response variable	Model	Parameter estimate	$\Delta AIC_c$	$w$	$R^2$
		B (CI)			
$\Sigma_{37}$ PBDEs liver ( $n = 25$ )	# hours at nest-site + capture date	1.0(0.3 – 1.7); -8.8 (-14 – -2.7)	0	0.43	0.38
	% nest-site attendance + capture date	499.5 ( 33.0 – 966.0); -7.5 (-14.3 – -0.7)	2.49	0.12	0.36
	plasma lipid % + date	-7.7 (-14.5 – -0.9); -200.4 ( -395.5 – -5.3)	2.79	0.11	0.35
	plasma lipid % + % nest-site attendance	-201.3 (-397.9 – -4.64)513.5 (42.3 – 984.8)	3.07	0.09	0.35
	capture date	-10.1 (-17.0 – -3.2)	3.69	0.07	0.25
	% nest-site attendance	682.4 (237.8 – 1158.1)	3.95	0.06	0.24
	# hours at nest-site	1.2 (0.4 – 2.0)	3.96	0.06	0.24
	% plasma lipids	-2.8 (-476.6 – -76.0)	4.41	0.05	0.23
	sex	-147.3 (-270.0 – -24.5)	5.82	0.02	0.18
	null	n/a	8.22	0.01	n/a
$\Sigma_{37}$ PBDEs plasma ( $n = 29$ )	% nest-site attendance	117.0 (21.7 – 212.2)	0	0.29	0.18
	# hours at nest-site + date	0.1 (0.0 – 0.3); -1.5 ( -2.8 – -0.1)	0.51	0.22	0.24
	sex	-24.5 (-47.6 – -1.4)	1.24	0.15	0.14
	date	-1.5 (-3.0 – - 0.4)	1.48	0.14	0.13
	# h nest-site	0.2 ( 0.0 – -0.1)	1.7	0.12	0.12
	null	n/a	2.70	0.07	n/a

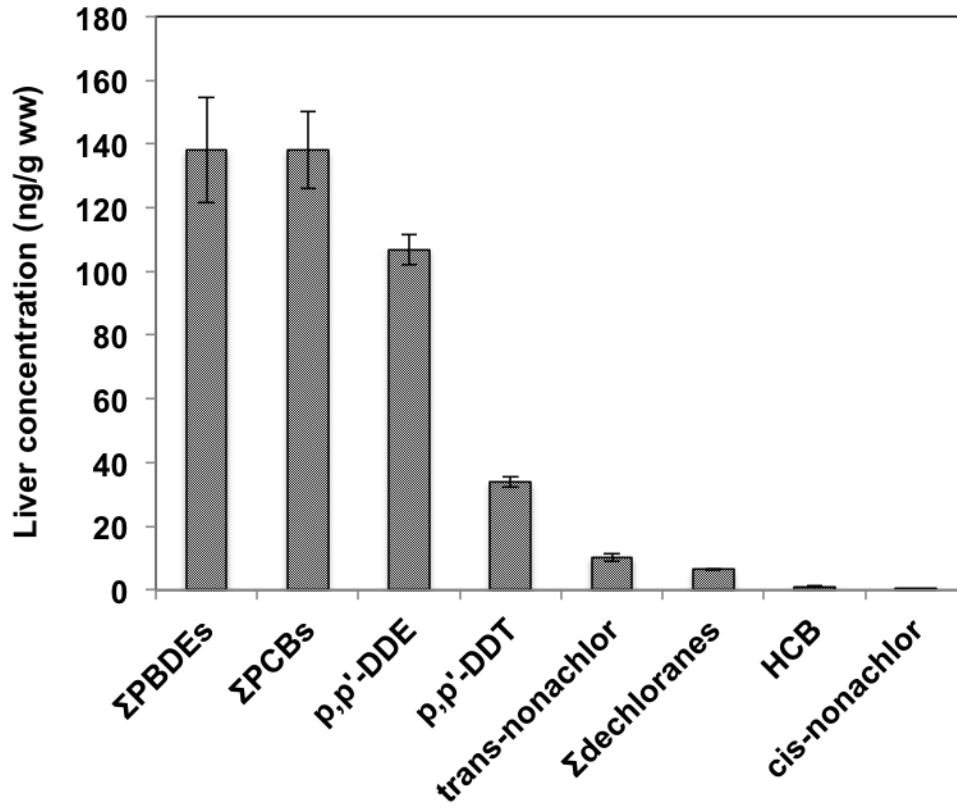
867 B = parameter estimates. CI = 95% confidence intervals.  $\Delta AIC_c$  = the difference in Akaike's Information Criterion (corrected for  
868 small sample sizes value) between a model and the top model.  $w$  = model weight. The  $R^2$  refers to the adjusted coefficient of  
869 determination generated from linear regression analysis. Variables with  $AIC_c$  values above the null model (thus explaining no  
870 variation in the response variable) included body mass, condition index, Julian lay date, proportion of time spent foraging, proportion

871 of time spent in different habitat types (urban, agricultural, waterway), percent body water or fat and for plasma  $\Sigma_{37}$ PBDEs the %  
872 plasma lipids.

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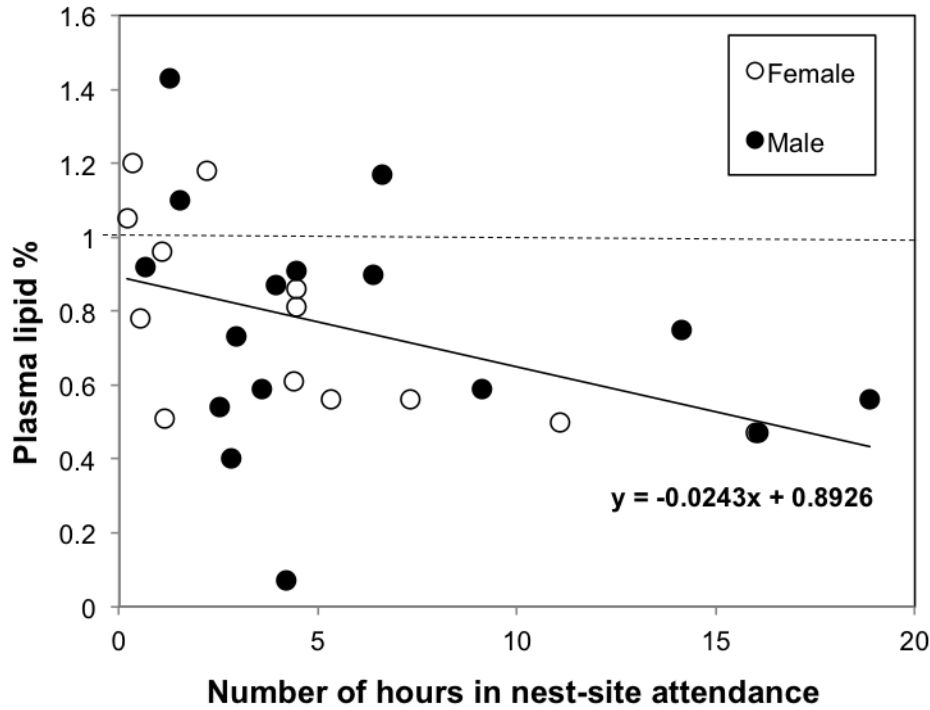
875 **Figure 1:** Mean concentrations ( $\pm$  SEM) of organochlorines and halogenated flame retardants in  
876 liver of ring-billed gulls (*Larus delawarensis*) collected during the incubation period near  
877 Montreal (QC, Canada).



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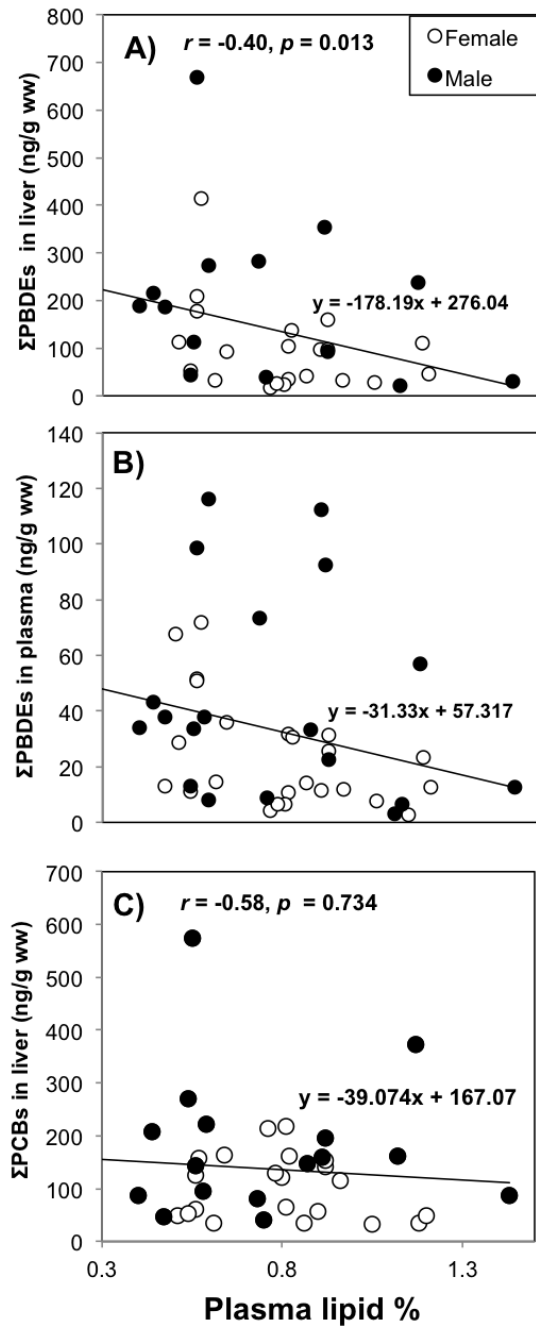
880 **Figure 2:** Relationship (Spearman's Rank Correlation) between plasma lipid percent and  
881 number of hours spent at the nest site in the last 24 hours prior to recapture in incubating ring-  
882 billed gulls (*Larus delawarensis*) near Montreal (QC, Canada). Birds with less than 1% plasma  
883 lipids (dashed line) were assumed to be in a fasted state (K. Drouillard, unpublished data).



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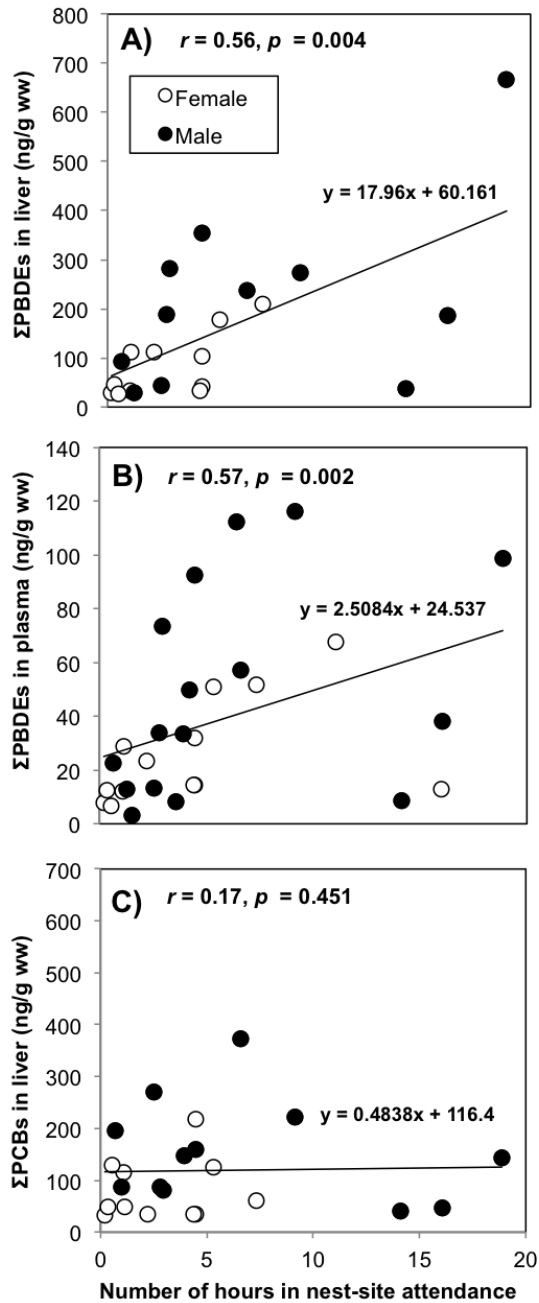
885

886 **Figure 3:** Relationships (Spearman's Rank Correlation) between the concentrations of  $\Sigma_{37}$ PBDE  
887 in liver (A) and plasma (B) and (C)  $\Sigma_{38}$ PCBs in liver and the percent lipids determined in plasma  
888 of incubating ring-billed gulls (*Larus delawarensis*) near Montreal (QC, Canada).



889

890 **Figure 4:** Relationships (Spearman's Rank Correlation) between the number of hours birds spent  
891 attending the nest-site in the last 24 hours prior to recapture and  $\Sigma_{37}$ PBDE concentrations in A)  
892 plasma and B) liver, and C)  $\Sigma_{38}$ PCBs in liver of incubating ring-billed gulls (*Larus delawarensis*)  
893 near Montreal (QC, Canada).



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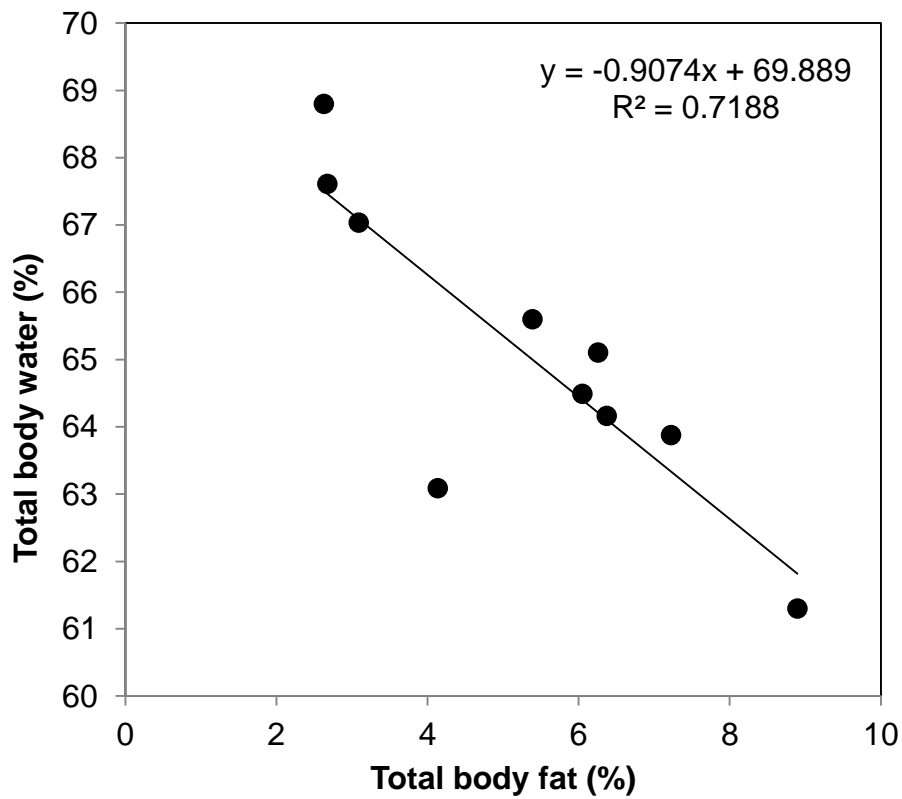
895 **Supplementary information**

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896

897 **Methods**

898 **Figure S1:** Relationship between total body water and total body fat of glaucous Gulls (*Larus*  
899 *hyperboreus*) from Svalbard, Norway. Data from Shaffer et al. (2006) and Verreault et al.  
900 (2007).



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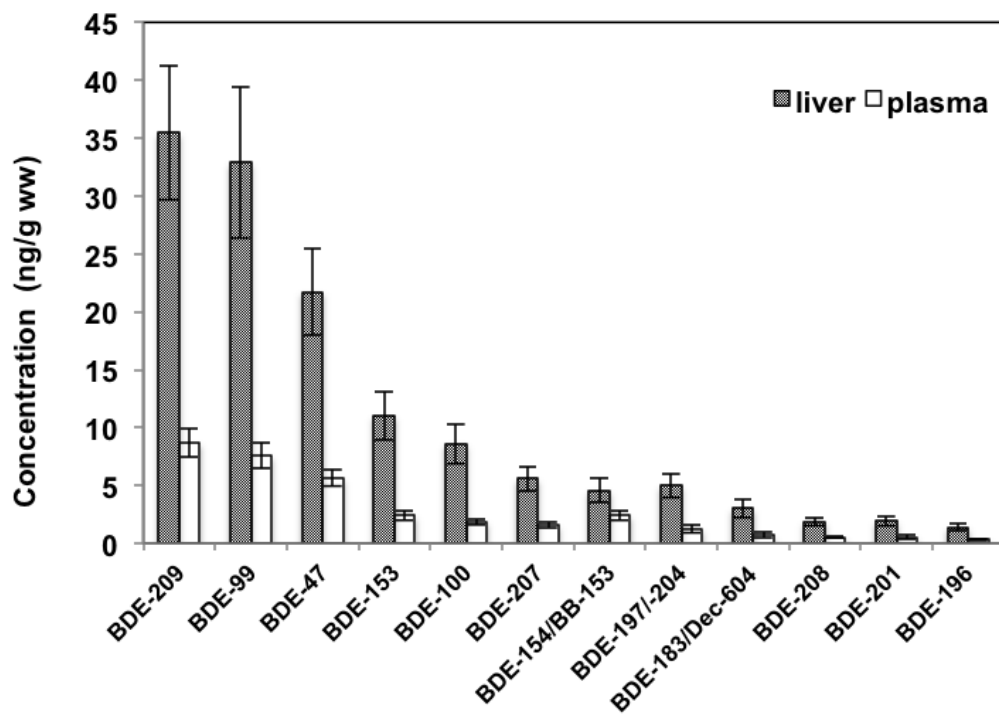
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906 **Results**

907 **Figure S2:** Concentrations of major PBDE congeners determined in liver and plasma of ring-  
908 billed gulls (*Larus delawarensis*) during the incubation phase near Montreal (QC, Canada).

909



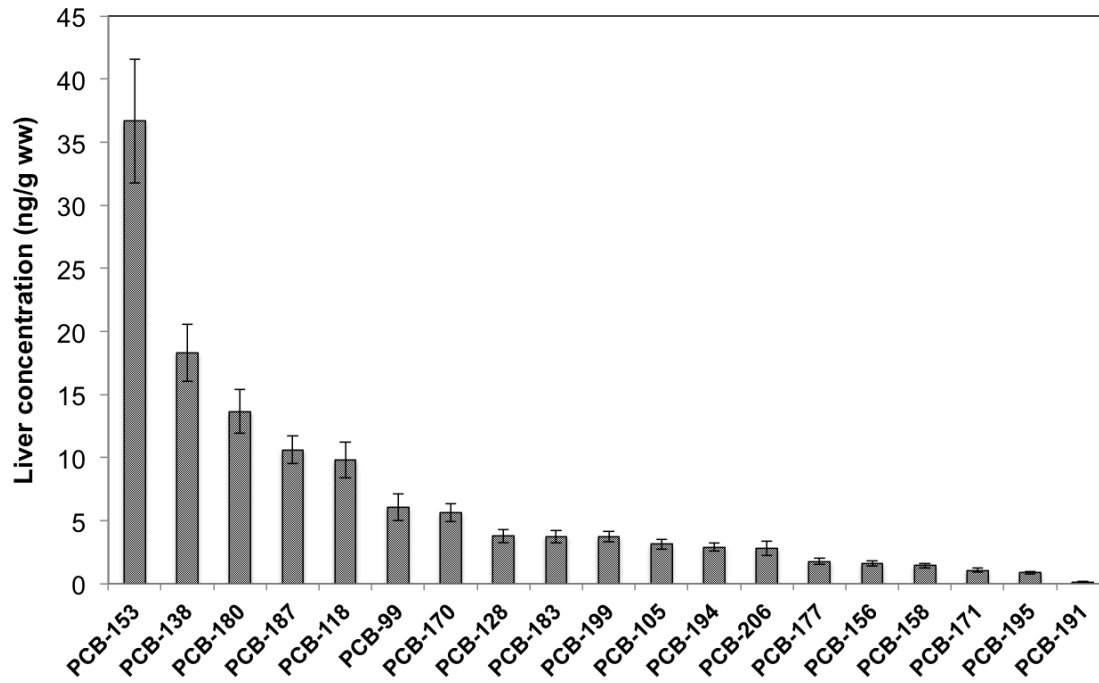
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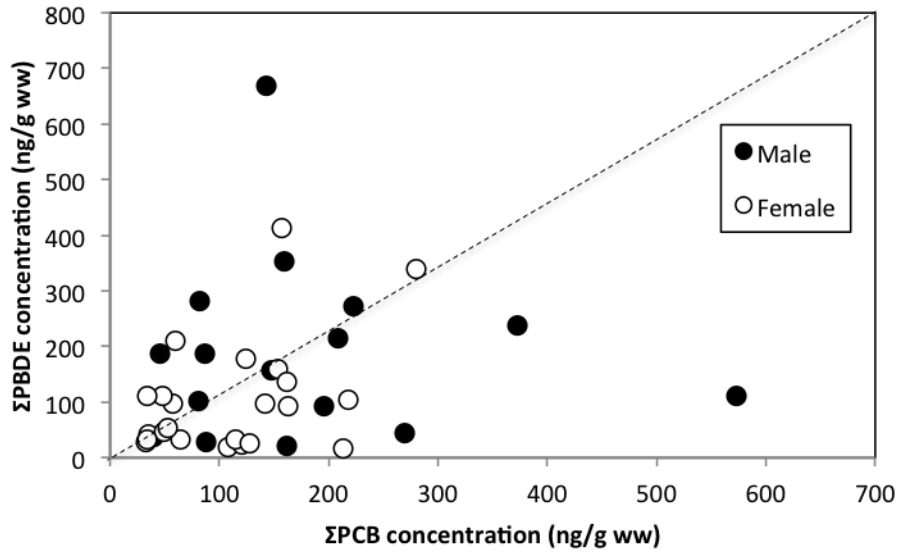


913 **Figure S3:** Concentrations of major PCB congeners determined in liver of ring-billed gulls  
914 (*Larus delawarensis*) during the incubation phase near Montreal (QC, Canada).



915  
916  
917

918 **Figure S4:** Concentrations of  $\Sigma_{37}$ PBDE vs.  $\Sigma_{38}$ PCBs in liver of ring-billed gulls (*Larus*  
919 *delawarensis*) nesting near Montreal (QC, Canada). Points above the dashed line represent birds  
920 for which  $\Sigma$ PBDE exceed  $\Sigma$ PCB concentrations and vice versa for points below this line.



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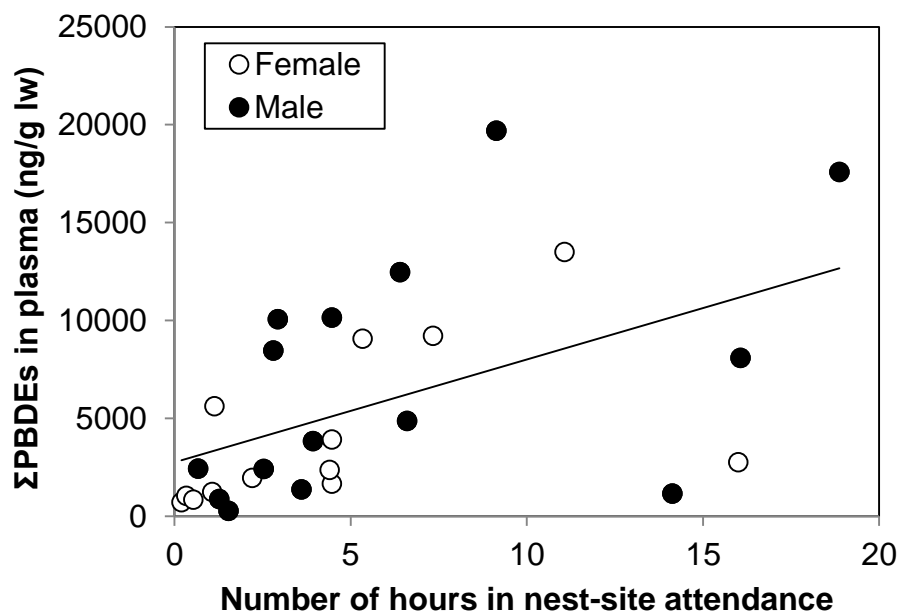
924 Relationships between lipid-corrected plasma HFR concentrations and time-activity budget

925 As the proportion of time that birds spent in nest-site attendance increased, so did their  
926 lipid-corrected concentrations of  $\Sigma_{37}$ PBDEs plasma ( $r = 0.46, p = 0.012$ ) as well as several of the  
927 individual PBDE congeners (in order of decreasing  $r$ : BDE-154/BB-153, -47, -99, -100, -153, -  
928 201, -197, -183:  $0.39 < r < 0.43, 0.007 < p < 0.003$ ). The positive relationship between nest-site  
929 attendance and PBDEs was stronger when the number of hours at the nest site before recapture  
930 was considered; all lipid-corrected congener concentrations were positively related to this  
931 activity measure ( $0.046 > p > 0.001$ ) as were  $\Sigma_{37}$ PBDEs ( $r = 0.61, p = 0.001$ ; Fig. S5) and  
932  $\Sigma$ dichloranes ( $r = 0.39, p = 0.043$ ).

933

934 **Figure S5:** Relationship (Spearman's Rank Correlation) between the number of hours birds were  
935 attending the nest-site in the last 24 hours prior to recapture and lipid corrected  $\Sigma_{37}$ PBDE  
936 concentrations in plasma of incubating ring-billed gulls (*Larus delawarensis*) near Montreal  
937 (QC, Canada).

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