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1	The effect of long-range transport, trophic position and diet
2	specialization on legacy contaminant occurrence in great skuas,
3	Stercorarius skua, breeding across the Northeast Atlantic
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5	Eliza H.K Leat ¹ Sophie Bourgeon ² Sveinn A. Hanssen ² Aevar Petersen ³ Hallvard
6	Strøm ⁴ Tor Harry Bjørn ⁵ Geir W. Gabrielsen ⁴ Jan Ove Bustnes ² Robert W. Furness ¹
7	Ane Haarr ⁶ Katrine Borgå ⁶⁷ *
8	
9	
10	¹ College of Medical Veterinary and Life Sciences Graham Kerr Building University of
11	Glasgow Glasgow G12 8QQ UK
12	² The Arctic University of Norway (UiT) Department of Arctic and Marine Biology
13	Tromsø Norway
14	³ Brautarland 2 108 Reykjavik Iceland
15	⁴ Norwegian Polar Institute FRAM Centre Postbox 6606 Langnes NO- 9296 Tromsø
16	Norway
17	⁵ Bio-Bjørn Kirkenesveien 4 9602 Hammerfest Norway
18	⁶ Department of Biosciences University of Oslo Pb 1066 Blindern 0316 Oslo Norway
19	⁷ Norwegian Institute for Water Research Gaustadalleén 21 0349 Oslo Norway
20	
21	*Corresponding author:
22	Katrine Borgå
23	Department of Biosciences University of Oslo
24	Pb 1066 Blindern

- 25 0316 Oslo
- 26 Norway
- 27
- 28 <u>katrine.borga@ibv.uio.no</u> Phone: +47 915 888 92

29 ABSTRACT

30 High levels of halogenated organic contaminants (HOCs) have been found in the marine 31 predatory seabird great skua (Stercorarius skua) from breeding colonies in the Northeastern 32 Atlantic, with large unexplained inter-colony variation. The present study aimed at analyzing 33 if the HOCs occurrence in breeding great skuas in remote colonies was explained by local 34 baseline food web exposure determined by long-range transport, or by ecological factors such 35 as diet specialization and relative trophic position in the breeding area. The occurrence of 36 organochlorine pesticides (OCPs), polychlorinated biphenyls (PCBs), and polybrominated diphenyl ethers (PBDEs) was analyzed in plasma of 204 adult great skuas collected over two 37 38 years (2008 and 2009) and 5 colonies across the North-Atlantic from Shetland to Svalbard. 39 The Σ HOCs levels in plasma ranged across two orders of magnitude, from 40 to 7600 ng/g 40 (wet weight) and differed significantly across the great skua colonies. The variation in 41 contaminant occurrence among colonies did not reflect long-range transport through a 42 latitudinal or remoteness gradient, as the second northernmost colony (Bjørnøya), had the 43 highest contaminant concentrations. No latitudinal or remoteness gradient was evident in the 44 contaminant pattern among the colonies. The contaminant levels increased significantly with increasing δ^{15} N values, and regurgitated pellets of undigested prev suggested that great skuas 45 with higher δ^{15} N values had a higher proportion of bird prey in their diet, mostly seabirds. In 46 contrast, great skuas from colonies with lower δ^{15} N and lower contaminant level fed mostly 47 on fish. The enrichment of δ^{13} C increased with decreasing δ^{15} N and lower contaminant levels. 48 49 Therefore, individual behavior of great skuas, such as migration strategies and diet 50 specialization, rather than long-range transport and thus baseline food web exposure, explain 51 among and within colony variance in contaminant occurrence.

- 53 Keywords: Great skua, *Stercorarius skua*, long-range transport, Arctic, global fractionation,
 54 remoteness index
- 55
- 56 Capsule: Trophic level and diet specialization, not long-range transport, explains the large-
- 57 scale variation in contaminant occurrence in the avian predator great skuas breeding in the
- 58 Northeastern Atlantic.

59 INTRODUCTION

60 High levels of halogenated organic contaminants (HOCs) have been found in the marine top predator great skua (Stercorarius skua) from remote breeding colonies in the Northeastern 61 62 Atlantic with large unexplained inter-colony variation (Leat et al. 2013). Adult birds are exposed to HOC through the diet, and lipid soluble recalcitrant HOCs biomagnify through the 63 64 food web with increasing HOC concentrations with trophic position (Borgå et al. 2001; Ruus 65 et al. 2002; Borgå et al. 2004). As great skuas are migratory, a central question is whether the 66 source of HOCs is predominantly from the local food web near the breeding colonies or from their wintering areas. The great skuas' wintering areas have previously been shown to explain 67 68 up to 22% of the variation in HOC concentrations in some of the breeding colonies, leaving 69 much unexplained variance in HOC occurrence to other factors such as local sources during 70 the breeding period (Leat et al. 2013). Local HOC sources, both the baseline exposure and 71 ecological factors like trophic position and diet specialization, may influence the HOC 72 occurrence, as they affect the trophic magnification in the food web (Borgå et al. 2012). 73

74 A principal source of spatial variation in contaminants in the abiotic environment is the 75 location relative to primary emission sources. Primary emission sources of HOCs have been 76 concentrated in industrialized- and agricultural areas mostly in the mid-latitudes of the 77 northern hemisphere (Breivik et al. 2004). However, long-range transport of HOCs in the 78 atmosphere has distributed these pollutants to remote environments, including the Arctic 79 (Muir and de Wit 2010). Although long-range transport is considered the main source of 80 contaminants to the Arctic, local sources such as settlements and military sites may contribute 81 as well (Brown et al. 2009). One of the physical processes governing long-range atmospheric 82 transport is cold condensation; that is, semi-volatile HOCs shift from gas phase to condensed

83 phases with decreasing temperature (Wania and Mackay 1993). Differences in chemical's 84 volatility lead to compositional shifts in pollutant pattern along latitudinal and altitudinal 85 temperature gradients, a process often referred to as global distillation (Wania and Mackay 86 1993). As an alternative to global distillation, the remoteness hypothesis suggests that it is the differential removal of pollutants from the air with distance from emission sources, rather 87 88 than a temperature gradient, that leads to environmental gradients of pollutants (von Waldow 89 et al. 2010a, von Waldow et al. 2010b). Comparisons of HOCs in wildlife between eastern 90 and western Greenland at similar latitude appear to support the remoteness hypothesis, with 91 higher HOC concentrations in eastern Greenland closest to European emission sources 92 (Vorkamp et al. 2004). However, observational data show that latitudinal gradients often 93 coincide with remoteness from emission sources, particularly in Europe (von Waldow et al. 94 2010a, von Waldow et al. 2010b).

95

Most studies of spatial distribution and long-range transport of pollutants focus on abiotic
compartments such as soils water and air (Agrell et al. 1999, Meijer et al. 2003, Gioia et al.
2006, Gioia et al. 2008, Lohmann et al. 2009). Few studies have been carried out in biota, but
a study of great black-backed gull (*Larus marinus*) along the Norwegian coast recognized the
pattern of contaminant occurrence predicted by global distillation (Steffen et al. 2006). Also,
studies of eagles suggested that at remote sites without local point sources, long-range
transport of contaminants is important (Elliott et al. 2009, Fort et al. 2014).

104 The effect of diet on biomagnification of HOCs is well known and established for food webs 105 using stable isotopes of nitrogen (δ^{15} N) as a measure of relative trophic position (Fisk et al. 106 2001, Borga et al. 2012). Even within a seabird colony, a positive association between trophic 107 position and contaminant concentrations may be significant (Sagerup et al. 2002), although

such intraspecific relationship is rarely documented in wildlife, probably due to restrictions in
sample sizes and lack of sufficient dietary variance among individuals. Whilst most great skua
pairs are generalist predators, some specialize on specific prey types and in some cases on
individual species (Furness 1987, Phillips et al. 1997). Therefore, some of the variation in
HOC occurrence in top predators such as great skuas may be explained by variation in diet,
either among or within colonies.

114

115 The distribution of great skua breeding colonies across the Northeast Atlantic cover a large 116 spatial scale (60°N to 80°N), and thus provides the opportunity to analyze if the spatial 117 variance of HOCs reflects the pattern caused by long range transport, or if trophic level and 118 diet specialization overrides this food web baseline signal. Concentrating on a single species 119 controls for some aspects of biological variability, which can affect HOC concentrations such 120 as species-specific biotransformation rates, and physiological adaptations, for example 121 metabolic rates. The great skua feed opportunistically on a wide range of prey, including 122 adults and chicks of seabirds, terrestrial mammals, pelagic fish, demersal fish discarded from 123 fishing boats, prey gained through kleptoparasitism of other seabirds, and some caught in 124 surface waters (Furness 1987). Indigestible prey remains are regurgitated in the form of 125 pellets, from which prey can be identified to taxon or in some cases species level. The diet of 126 individual skua pairs can be studied as they are highly territorial, and regurgitated pellets are 127 thus highly likely to be produced by the birds occupying that territory.

128

The aim of the present study was to unravel if the occurrence of HOCs in great skuas across the North Atlantic was explained by local baseline food web exposure determined by longrange transport, or by relative trophic positions and local diet habits. We hypothesized that if long-range atmospheric transport is the dominant process affecting HOC concentrations in

133 great skuas, plasma concentrations would decrease either with increasing latitude or 134 remoteness from emission sources. In addition, the pattern of HOCs was predicted to show 135 increasing proportion of less halogenated (thus more volatile) compounds in the northern 136 colonies or with increasing remoteness. If the HOC levels were predominantly described by 137 the great skuas trophic position and diet specialization we expected the HOC levels to be related to the stable isotope signatures of nitrogen ($\delta^{15}N$) and carbon ($\delta^{13}C$), and to prev 138 139 remains in regurgitated pellets. None of the breeding colonies included in this study are close 140 to large urban areas, therefore it is presumed that sources of HOCs are dominated by long-141 range atmospheric transport rather than local emission sources.

142

143 MATERIALS and METHODS

144 Sampling

145 Blood was sampled using heparinized syringes from 204 incubating adult great skuas in five

146 colonies in 2008; Ny-Ålesund 78°55'N, n=5; Bjørnøya 74°29'N, n=15; Hjelmsøya 71°03'N,

147 n=4; South-East Iceland 63°52'N, n=21; and Runde 62°23'N, n=5 (listed with decreasing

148 latitude), and in three colonies in 2009; Bjørnøya n=51; Iceland n=56, and Shetland 60°09'N,

149 n=47 (Table 1, Table S1). Plasma and red blood cells were separated by centrifuging and

150 frozen at -20°C. Birds were sexed by molecular methods using DNA extracted from red blood

- 151 cells following a modified protocol (Griffiths et al. 1998). Whilst all birds sampled in 2009
- 152 were successfully sexed, some of the individuals sampled in 2008 were not due to insufficient
- 153 quantity of red blood cells for analyses.

154 *Remoteness indices*

155 Remoteness of the colonies was estimated using remoteness indices with night-time light

156 emissions and cropland areas as proxies for HOC emissions from industrial and agricultural

157 activities respectively (von Waldow et al. 2010b) (see supporting information for details). For

the two emission scenarios, the colonies increased in remoteness as follows; Shetland and
Runde < Hjelmsøya and Iceland < Bjørnøya < Ny-Ålesund using night-time light emissions
and Shetland and Runde < Hjelmsøya < Iceland < Bjørnøya < Ny-Ålesund using cropland
area. Recently quantification of remoteness was modulated to take into account the
contaminants' physicochemical properties (Göktaş and MacLeod 2016). As these results were
highly correlated with von Waldow's remoteness index, the colonies were ranked according
to the original remoteness index.

165

166 Contaminant analyses

167 OCs (HCB, Dieldrin, Mirex, QCB, OCS, β-HCH, trans-nonachlor, oxychlordane, cis-

168 nonachlor, *cis*-chlordane, *pp* '-DDT, *pp* '-DDE, *pp* '-DDD, and PCB congeners: 31/28, 52, 49,

169 74, 70, 101, 99, 110, 149, 118, 153, 105/132, 138, 158, 187, 183, 128, 177, 156/171, 180,

170 191, 170, 201, 195/208, 194, 205, 206, 209) and PBDEs (congeners 47, 99, 100, 153, 154)

171 were extracted and measured in a total of 204 plasma samples at the Great Lakes Institute for

172 Environmental Research (GLIER) at the University of Windsor Ontario Canada by gas

173 chromatography electron capture detection (GC-ECD) (Lazar et al. 1992). Lipid content

174 (extractable organic content) in plasma was gravimetrically determined. The details of the

analyses are described in Leat et al. (2013). Recoveries of the internal recovery standard PCB-

176 30 were 77.4 +10% (mean + SD) in 2008 and 69.5 + 9.6 % in 2009. The data were recovery

177 corrected. Method detection limit was 0.08 µg/kg for PCBs and 0.05 µg/kg for all other OCs.

178 Method detection limits for PBDEs varied by congener; PBDE-47 0.373 µg/kg, PBDE-99

179 0.361 μg/kg, PBDE-100 0.471 μg/kg, PBDE-153 0.420 μg/kg, PBDE-154 0.488 μg/kg.

180

181 Dietary descriptors

Stable isotopes of nitrogen and carbon (δ^{15} N and δ^{13} C respectively) were measured in red 182 183 blood cells as described in detail elsewhere (Leat et al. 2013). The analytical precision based 184 on the standard deviation of a NIST 8414 standard (bovine muscle) for 2008 and 2009 data was 0.07 and 0.18‰ for δ^{15} N and 0.08 and 0.09‰ for δ^{13} C, respectively. An internal 185 186 laboratory fish muscle standard was included for 2009 data for which the standard deviations were 0.22‰ and 0.08‰ for δ^{15} N and δ^{13} C, respectively. The variation in spatial baseline 187 188 isotopic signature among remote sites (Green et al. 2016) is assumed to be negligible 189 compared to signature variation caused by differences in diet in the great skuas.

190 In 2009, breeding territories of blood-sampled birds in each colony were searched for 191 regurgitated pellets every 2-3 days, until the nest failed or fieldwork ceased. In 2008 and 192 2009, additional sampling of regurgitated pellets from great skuas breeding territories also 193 took place on an *ad hoc* basis in the Bjørnøya colony. Pellets were examined and classified by 194 prey type; bird, fish, mammal or other, and identified to species level whenever possible. 195 Carcasses of large birds, e.g. ~20 days old great skua chicks and rabbits (Shetland) were 196 counted as equivalent to 1 pellet. Sagittal otoliths recovered from the regurgitated pellets were 197 used to identify the fish species consumed (Härkönen 1986). A minority of all pellets 198 contained more than one prey type and were divided proportionally between the identified 199 groups of prey i.e. 0.5 fish, 0.5 bird. Vegetation in pellets was excluded from the dietary data 200 analysis as it was considered to have been ingested incidentally along with animal dietary 201 items. Territories with > 5 regurgitated pellets were classified as specialists if > 70% of pellets 202 came from one dietary prey group (bird fish or mammal) (Votier et al. 2004). Otherwise, they 203 were classified as generalists. Regurgitated pellets were destroyed once sampled, and 204 identified on site to prevent double counting, or they were removed for further examination. 205 Food regurgitated whilst handling the birds was not included in calculations of diet 206 composition, but was used to identify prey items from the pellets.

207 Data analysis

208 Data from 2008 and 2009 were analyzed separately, as not all colonies were sampled, as not 209 all variables were included in both years. That is, whilst the sex of all individuals sampled in 210 2009 was identified using DNA, the majority of individuals sampled in 2008 were not sexed 211 due to insufficient material for DNA analysis. Treating the years separately allows sex to be 212 included as an explanatory variable in the analysis of the 2009 data. Twelve individuals, three 213 from Bjørnøya and nine from Iceland, sampled in both 2008 and 2009 were included in the 214 analysis. Since males do the majority of foraging during incubation and often regurgitate food 215 to the female at the nest (Catry and Furness 1999, Caldow and Furness 2000) the registered 216 data (HOCs, stable isotopes, regurgitated pellets) in males and females of the same nest (pair) 217 may not be independent. To ensure independence of data, only the first bird sampled per 218 territory/nest was included in the analysis.

219

220 OCs were included in the data analysis if concentrations were above method detection limits 221 in \geq 70% across the colonies. In total 40 organochlorines were included in the analysis: 28 222 PCBs ranging from tri- to deca- homologues, and 12 organochlorine pesticides including 223 HCB, OCS, β-HCH, Mirex, trans-nonachlor, cis-nonachlor, oxychlordane, pp'-DDT, pp'-224 DDD, and pp'-DDE. Non-detects in the included contaminants were replaced by modelled 225 values below the detection limit, using the distribution of concentrations above the detection 226 limit of an individual HOC as described in Leat et al. (2013) and in supplementary 227 information (Helsel 2006). Some of the 2009 data are published in Bourgeon et al. (2012) and 228 Leat et al. (2013).

229

230 In all analyses, PCBs were grouped according to homologue group as volatility and long-

231 range transport depends upon the degree of halogenation. Detection rates of the five PBDEs

232 analyzed (PBDE-47, 99, 100, 153, and 154) varied between congeners, colonies and years 233 (Figure S2). PBDE 47 was detected in 97% of the samples and in samples where all five 234 PBDEs were above detection limit (n=89, four individuals sampled in both years were 235 excluded from the 2009 dataset), PBDE 47 was positively correlated with the other PBDEs 236 (Spearman's rank correlations: rho = 0.81, S = 22236, rho = 0.84, S = 18349, rho = 0.61, S = 237 46348, rho = 0.67, S = 38931, p < 0.001 in all cases) (Figure S3). Therefore PBDE 47 was 238 used as a proxy for commercial penta-BDEs in analysis of spatial variation in PBDE 239 concentration.

240

Lipid content in plasma did not differ significantly between colonies (2008: F_{447} = 0.39, p = 0.81; 2009: F_{249} = 2.4, p = 0.098, two low outliers were removed to meet assumption of normality of residuals) and all HOC concentrations are presented on a wet weight basis.

245 Contaminant concentrations and patterns among colonies were investigated using the 246 multivariate methods redundancy analysis (RDA) and principal component analysis (PCA) using the *vegan* package in R (R Development Core Team 2015). Colony, $\delta^{15}N$, $\delta^{13}C$ and 247 248 lipid content were included as explanatory variables (active in RDA, passive in PCA) for both 249 years. The percentage of variation attributable to each explanatory variable was assessed by 250 RDA. This was done by comparing the residual variance of the full model with models 251 constrained to each explanatory variable in turn, separately (by conditioning out the other 252 explanatory variables). Sex was included as explanatory variable in 2009, only. HOC 253 concentrations were logarithmically transformed to reduce variance heterogeneity and 254 skewness, whilst to investigate HOC pattern the data were standardized by norm to analyze 255 the relative contribution of each contaminant to the total.

256

257	The sum of the OCs, and PBDE 47, were used to examine the overall relationships between
258	diet, mainly carbon source (δ^{13} C), and relative trophic level (δ^{15} N), and HOCs. Relationships
259	between stable isotopes and HOCs concentrations were investigated using general linear
260	models with backwards stepwise selection to remove non-significant variables. HOC
261	concentration data were log transformed to reduce heterogeneity and skewness (Full model
262	for 2009 in Equation 1).
263	
264	$\log \Sigma OC \text{ or } \log \Sigma PBDEs = \delta^{15}N + \delta^{13}C + \% \text{ lipid} + \text{sex} + \epsilon \qquad (Eq. 1)$
265	
266	The relationship between diet measured by regurgitated pellets and stable isotopes was
267	investigated by linear models whereas the relationship between pellets and HOCs was
268	examined visually due to small sample sizes of pellets.
269	
270	RESULTS AND DISCUSSION
271	Overall, the data show that great skuas feeding at higher trophic levels and specializing on
272	bird prey (inferred from stable isotopes and regurgitated pellets) have higher concentrations of
273	pollutants, and that this is the main explanation for variation in HOC occurrence among
274	colonies, rather than global distillation and remoteness (colonies spanning nearly 19 degrees
275	latitude) to primary sources and wintering sites (Leat et al. 2013). Large sample sizes were
276	obtained from Bjørnøya, Runde, Iceland and Shetland (15 – 56 individuals), whereas small
277	sample sizes from Hjelmsøya, Runde and Ny-Ålesund $(4 - 5 \text{ individuals})$ limit the
278	interpretation of results from these colonies. Contaminant concentrations decreased in the
279	following order: Bjørnøya (2009) >Bjørnøya (2008) > Hjelmsøya (2008) > Runde (2008) >
280	Iceland (2009) > Ny Ålesund (2008) > Iceland (2008) > Shetland (2009). <i>pp</i> '-DDE and PCB
281	153 were the contaminants of highest concentrations in all colonies for both years, followed

282 by oxychlordane, mirex, and trans-nonachlor (Table 1).

283

284 HOCs occurrence in great skua plasma and long-range transport

285 To meet the predictions of global distillation or remoteness hypotheses, Runde/Shetland and 286 Ny-Ålesund should be extremes, representing least remote/lowest latitude and most remote/highest latitude, respectively. However, great skuas from Ny-Ålesund and Runde had 287 288 intermediate contaminant concentrations compared to those of the other colonies (Table 1). 289 Great skuas from Shetland had lowest contaminant concentrations, despite being the least 290 remote/lowest latitude colony. The HOC concentrations were higher in the remote and high 291 latitude colony Bjørnøya, compared to the other colonies both in 2008 and 2009. Skuas from 292 Bjørnøya had twice as high mean ΣOC concentrations as the second most contaminated 293 colony in 2008 (Hjelmsøya), and more than 3.5 times the mean concentrations from Iceland in 294 both years (Table 1).

295

296 Lipophilic HOC concentrations in the blood of birds are known to increase with decreasing 297 body mass and lipid reserves (Bustnes et al. 2012b, Bustnes et al. 2017). Therefore, 298 differences among colonies in the average body condition of great skuas offer a potential 299 explanation for spatial differences in HOC concentrations. However, the present study is 300 biased towards the selection of breeding birds. Great skuas are long-lived and may offset 301 breeding seasons if conditions are poor (Catry et al. 1998). Hence, birds in poor condition are 302 unlikely to have been sampled. There was, however, no significant difference in body mass or 303 wing length of great skuas among colonies sampled in 2009 (Bourgeon et al. 2012). Thus, 304 body condition was not considered an important explanatory factor for variation in HOC 305 occurrence in the present study, and therefore not included in the statistical analyses.

307 PCA plots of HOC concentrations and patterns are presented in supporting information 308 (Figure S1). No clear latitudinal gradient was found (selected representative HOCs are 309 presented in Table 1). The direct multivariate redundancy analyses (RDA) accounted for 32% 310 and 60% of the variance in HOC concentrations in 2008 and 2009, respectively (Table 2). 311 Significant explanatory variables accounting for most of the variance in concentrations were, in decreasing order: colony, lipid%, δ^{13} C and δ^{15} N in 2008; and colony, δ^{15} N, δ^{13} C and lipid 312 313 in 2009 (Figure S1, Table 2). The direct multivariate redundancy analyses accounted for 52% 314 and 18% of the variance in HOC patterns in 2008 and 2009, respectively (Table 2). 315 Significant explanatory variables accounting for most of the variance in patterns were, in decreasing order: colony, $\delta^{15}N$ and $\delta^{13}C$ in 2008; and colony, sex, $\delta^{15}N$ and $\delta^{13}C$ in 2009 316 317 (Figure S1, Table 2). Male great skuas had higher HOC concentrations compared to females, 318 which is in accordance with findings in other avian wildlife (Bustnes et al. 2003, Bustnes et 319 al. 2005, Bustnes et al. 2007, Bustnes et al. 2017). However, neither the lipid content or sex 320 were found to be major contributors to the inter-colony difference in HOC occurrence in great 321 skuas. There was no clear latitudinal trend in the relative pattern of HOCs in general, or in 322 PCB homologue patterns in either year. The contaminants with the largest variance in relative 323 contribution among the samples were penta- to nona-CB homologue groups, pp'-DDE, pp'-324 DDT, oxychlordane and Mirex. If long-range transport would explain the contaminant pattern 325 found in great skuas, the pattern in the Bjørnøya colony should have had proportionally more 326 volatile HOCs, and the pattern in the Iceland colony would have been intermediate to 327 Shetland and Bjørnøya, which is not the case. 328

329 The lack of a latitudinal or remoteness gradient in concentrations and pattern of HOCs in

330 great skuas is in contrast to abiotic studies (Agrell et al. 1999, Meijer et al. 2003, Gioia et al.

331 2006) and the few previous studies of geographic distribution of HOCs in biota (Ter Schure et

332 al. 2002, Vorkamp et al. 2004, Steffen et al. 2006, Bourgeon et al. 2012, Bustnes et al. 2012a). 333 Most of the species in which latitudinal or remoteness gradients in HOCs have been found are 334 cold blooded species occupying lower trophic levels compared to the great skua (Ter Schure 335 et al. 2002, Vorkamp et al. 2004, Bourgeon et al. 2012, Bustnes et al. 2012a). The HOC 336 pattern in the skuas across the colonies is dominated by recalcitrant and biomagnifying HOCs. 337 Therefore, biotransformation of HOCs in the trophic levels preceding that of the great skua, 338 and in the great skua itself, seem to obscure long-range abiotic transport patterns. In general, 339 biotransformation is efficient in warm blooded species compared to cold blooded lower 340 trophic level species where the contaminant pattern to a higher degree resembles that of the 341 abiotic environment (Borga et al. 2004). The dominance of highly chlorinated PCBs 342 congeners in the pattern of HOCs in Bjørnøya birds in 2008 is indicative of a pattern 343 dominated by biomagnification, rather than long-range transport (Borgå et al. 2001).

344

345 Relative trophic position and HOC concentrations in great skua

346 The stable isotope signatures differed among colonies (Figure 1, Table 1, MANOVA 2008: 347 Pillai = 0.87 F_{448} = 10 p < 0.001; 2009: Pillai = 0.96 F_{2150} = 70 p < 0.001) and there were no differences between sexes (2009: Pillai = $0.018 \text{ F}_{1150} = 1 \text{ p} = 0.26$). In 2008, great skuas from 348 Bjørnøya had higher δ^{15} N than Iceland, Ny-Ålesund and Runde skuas (r² = 0.42, F₄₄₈ = 10.5, 349 350 p <0.001, Tukey p <0.005). Hjelmsøya, with similar δ^{15} N to Bjørnøya, had higher δ^{15} N values than Iceland (p = 0.042). In 2009, skuas from Bjørnøya had higher δ^{15} N values than Iceland 351 and Shetland skuas (Kruskal-Wallis $\chi^2 = 121$, df = 2, p < 0.001, post hoc comparisons 352 Wilcoxon Signed rank tests p<0.001), and Iceland skuas had higher $\delta^{15}N$ than Shetland (p 353 354 <0.001). As stable isotopes and colony could not be judged as independent variables, colony 355 was excluded from models investigating the relationship between HOC concentrations and 356 stable isotopes.

358	Across colonies there was a positive linear relationship between $\delta^{15}N$ and ΣHOC
359	concentrations (2008: R ² = 0.11, slope=-0.55, p=0.007, 2009: R ² = 0.59, slope=0.93, p<0.0001,
360	Figure 2). Although lipophilic recalcitrant HOCs show increasing concentrations with trophic
361	position in the food web (Borgå et al. 2001, Fisk et al. 2001, Ruus et al. 2002, Lavoie et al.
362	2010), such relationships are rarely shown within one species. However, $\delta^{15}N$ levels were
363	found to be positively related to OC concentrations in both bald eagles (Haliaeetus
364	leucocephalus) (Elliott et al. 2009) and white-tailed eagles in Norway (H. albicilla) (Bustnes
365	et al. 2013), and to HCB, DDTs and PCBs (from muscle) in Bjørnøya glaucous gulls (Larus
366	<i>hyperboreus</i>) (Sagerup et al 2002.), and weak positive relationships were found between $\delta^{15}N$
367	and HOCs in liver of northern fulmars (Fulmarus glacialis) (Knudsen et al. 2007). Even
368	though there are no measurements of baseline $\delta^{15}N$ from the study area, we assume that $\delta^{15}N$
369	is an indicator of relative trophic level. Green et al. (2016) reported that blue mussels (Mytilus
370	edulis) occupy the same trophic level along the whole Norwegian coast (range: 5-8‰), with
371	some stations appearing to be outliers (particularly low or high $\delta^{15}N$). However, the $\delta^{15}N$
372	levels in these stations was attributed to anthropogenic influence, i.e. increased anthropogenic
373	input. Thus, we assume that the difference in isotopic signal across the remote colonies in the
374	present study, reflects differences in diet specializations rather than differences in baseline
375	isotopic signal. In the present study, positive linear relationships between $\delta^{15}N$ and log sum
376	HOCs within the colonies were found for the Iceland colony in 2009 but not 2008 (R^2 =0.07,
377	$p=0.05$, and $R^2=0.17$, $p=0.5$, respectively), in Shetland in 2009 ($R^2=0.10$ $p=0.03$), and the
378	Bjørnøya colony in 2008 but not 2009 ($R^2=0.26$, p=0.03, and $R^2=0.004$, p=0.67,
379	respectively). These results suggest variation in individual diet specialization or prey taxon
380	availability between breeding seasons.
381	

382 The regurgitated pellets indicated differences in diet and occurrence of specialist and 383 generalist predators among colonies, with trophic position increasing from Shetland to Iceland to Bjørnøya ($\chi^2 = 296$, df = 4, p < 0.001, respectively). Great skuas from Bjørnøya had a 384 higher proportion of bird in their diet (2008: $\chi^2 = 229$, df = 1, p < 0.001, 2009: $\chi^2 = 132$, df = 1, 385 p <0.001), whilst great skuas in Iceland and Shetland had a higher proportion of fish ($\chi^2 = 86$ 386 and $\chi^2 = 95$, p <0.001) (Table 4). Great skuas from Shetland also had a higher proportion of 387 388 other diet items. Of the regurgitated pellets in Shetland, 19% (n = 71) contained mammal 389 remains, mostly rabbit (Oryctolagus cuniculus), whilst only one regurgitated mammal pellet 390 was recorded in Iceland, At Bjørnøya, the arctic fox (Vulpes lagopus) is the only mammal 391 present. The regurgitated mammal pellets from Shetland were concentrated in 14 territories 392 out of 42, of which only 7 had more than one regurgitated mammal pellet. Of the territories at 393 Bjørnøya where 5 or more pellets were collected (n = 56, a mean of 13 pellets per nest), the 394 majority were bird specialists (2008: 67%, 2009: 77%, Table 4). In contrast, 47% of great 395 skua territories from Shetland were occupied by fish specialists and 43% by generalists. In 396 Iceland, only three territories had more than 5 pellets, and recorded two fish specialists and a 397 generalist. Details of identified bird and prey species can be found in Supplementary 398 Information.

399

400 Carbon source and HOC occurrence

401 Bjørnøya, Hjelmsøya and Runde were depleted in δ^{13} C compared to Iceland and Ny-Ålesund

402 in 2008 ($r^2 = 0.74$, $F_{448} = 38$, p < 0.001, Tukey p < 0.005) (Table 1). Likewise, in 2009 skuas

403 from Bjørnøya were depleted in δ^{13} C compared to Iceland and Shetland (r² = 0.52, F₂₁₅₁ = 83,

404 p <0.001, Tukey p <0.001) (Table 1). Across all colonies, there was a negative linear

405 relationship between carbon source (δ^{13} C) and Σ HOC concentrations (2008: R²= 0.19, slope=-

406 0.73, p=0.0004, 2009: R^2 = 0.24, slope=-1.34, p<0.0001, Figure 2). Most reported studies have

found few or no significant relationships between δ^{13} C values and HOCs in avian tissues 407 408 (Ruus et al. 2002, Sagerup et al. 2002, Knudsen et al. 2007), however, Lavoie et al. (2010) found sum brominated flame retardants (BFRs) to increase with δ^{13} C. Studies of coastal 409 410 breeding eagles report that PCBs, DDE and PBDEs in chicks increase with marine input and increasing δ^{13} C levels (less negative values) (Elliott et al., 2009), and thus is opposite to the 411 412 findings from the present study. The study by Elliott et al., (2009) also included some 413 individual eagles breeding in estuaries or further up river and therefore reflect more riverine 414 and freshwater systems. For white-tailed eagles in Norway, there was a negative relationship between δ^{13} C levels and POPs (Bustnes et al. 2013) similar to this study. δ^{13} C measurements 415 416 from the present study varied only from -19.8 to -17.7‰, which is at the end of scale 417 compared to the study by Elliott et al. (2009), which might explain the contrasting findings. 418

Spatial patterns in δ^{13} C are well known in the Southern Ocean with δ^{13} C decreasing with 419 420 increasing latitude (Quillfeldt et al. 2010), but less is known from the northern hemisphere. In the present study, δ^{13} C in great skuas does not simply decrease with increasing latitude, as 421 422 Ny-Ålesund δ^{13} C is comparable to Iceland rather than Bjørnøya. Runde skuas are grouped with Bjørnøya and Hjelmsøya in δ^{13} C values. We hypothesized that the Shetland skuas feed to 423 a greater degree on the terrestrial (less seabird prey, more mammal prey) system as reflected 424 425 in the regurgitated pellets, and that this further contributes to lower HOC exposure, as HOC 426 levels in the terrestrial system are generally lower than the marine system for the North Atlantic and Arctic region (AMAP 2004). If so, Ny-Ålesund and Iceland would also reflect a 427 428 more enriched δ^{13} C accompanied with lower HOC levels compared to e.g. Bjørnøya. Remains of mammals were found in regurgitated pellets from Iceland, and also the Ny-Ålesund skuas 429 430 are known predators on eggs and chicks of terrestrial feeding geese (Bustnes, Personal communication). However, none of these colonies had a distinct terrestrial δ^{13} C signal. 431

Bjørnøya was the colony with the lowest δ^{13} C values indicating a marine signal, and the highest levels of HOCs, which support the general assumption that marine food webs are more contaminated than terrestrial food webs. However, further research is needed on the baseline levels of δ^{13} C along a northern-Arctic, latitudinal gradient, as has been done for southern-Antarctic areas.

437

438 *Measuring diet – combining stable isotopes and pellet data*

439 When combined with regurgitated pellets, the indication for colony differences in HOCs 440 being mostly driven by the trophic position at breeding site is even stronger. Across both years, birds from Bjørnøya had the highest HOC concentrations and the highest $\delta^{15}N$ values, 441 442 which probably resulted from Bjørnøya skuas having the highest proportion of bird prey in 443 their diet. Bjørnøya had the highest proportion of individuals specializing on bird prey, and 444 these skuas had consistently higher concentrations of both OCs and PBDEs compared to 445 generalist skuas in Bjørnøya. This has also been found for glaucous gulls (*Larus hyperboreus*) 446 at Bjørnøva (Bustnes et al. 2000). Shetland at the other extreme had a more mixed diet 447 dominated by fish, with only one bird prey specialist, several skuas feeding on terrestrial mammals (rabbit), and the lowest δ^{15} N values and HOC concentrations across colonies. 448 449 Iceland had intermediate HOC concentrations and $\delta^{15}N$ values, and a higher proportion of bird 450 prey in their diet, than birds at the Shetland colony. No pellets were collected from the three 451 smaller colonies, Ny-Ålesund, Hjelmsøya and Runde, holding less than 60 pairs at each 452 location (around 10, 5 and 50 pairs for both years, respectively). These three colonies are 453 located near to large colonies of other species including geese and eider ducks, providing 454 potential prey either through kleptoparasitism or by preying directly on eggs, chicks or adult 455 birds. Small great skua colonies have been shown to have a higher incidence of seabird

456 predation (Votier et al. 2007). However, HOC concentrations in Ny-Ålesund, Hjelmsøya and
457 Runde were lower than birds from Bjørnøya, where seabird prey dominate.

458

459 Stable isotopes and regurgitated pellets both provided information on the diet of great skuas in 460 the breeding season, and can be used to analyze the relationship between diet and HOC 461 concentrations. One obvious difference is that the regurgitated pellets only reflect the 462 momentary diet and give specific information on prey items, whereas the stable isotopes 463 integrate the carbon source and relative positioning in the food web over time. In the present 464 study, red blood cells were analyzed for stable isotopes, which integrates the stable isotopes 465 over the past 2-3 weeks (Bearhop et al. 2002). Pellets tend to over represent the proportion of 466 mammal and bird in the diet due to their amount of indigestible material, and the longevity of 467 pellets composed of fur or feathers in the environment (Furness and Hislop 1981, Votier et al. 468 2001). Regurgitates may also be biased in the other direction as a bird diet may be more 469 difficult to regurgitate than a fish diet (Furness 1987). However, presuming that the bias in 470 pellet data towards birds and mammals is constant across colonies, pellets still reflect 471 differences in diet among colonies.

472

473 Contaminant levels compared to other studies

The organochlorine concentrations in great skua plasma were amongst the highest found in plasma of seabirds in contemporary studies. Glaucous gulls (*Larus hyperboreus*), particularly those breeding in Bjørnøya, are known to have high concentrations of HOCs associated to negative effects (Verreault et al. 2010). Great skua from Bjørnøya in 2009 had twice the DDT concentrations found in glaucous gulls from Bjørnøya in 2002 and 2004 (Verreault et al. 2005). Higher OC concentrations in great skua compared to seabirds of a similar ecological niche is not confined to Bjørnøya. Also, in Ny-Ålesund on Svalbard, OC concentrations in

great skuas exceeded those of glaucous gulls (Verreault et al. 2006), although both great skuas
and glaucous gulls have lower levels in Ny-Ålesund than Bjørnøya (Verreault et al. 2005,
Verreault et al. 2006). OC concentrations in great skuas from Hjelmsøya are higher than in
whole blood of great black-backed gulls at Hornøya in northern Norway (Bustnes et al. 2005).
There are no close comparisons across species available for the least contaminated colony
Shetland, however, plasma concentrations there exceeded those of great black-backed gulls
whole blood concentrations in Norway (Bustnes et al. 2005).

488

489 As in the present study of great skuas, PBDEs are generally found at low concentrations 490 compared to many of the most persistent OCs in the above-mentioned avian top predators 491 (Verreault et al. 2006, Chen and Hale 2010). As with OC, PBDE concentrations in great skuas 492 exceeded those of glaucous gulls (Verreault et al. 2007). Although the dietary niches of 493 glaucous gulls, great black-backed gulls, and great skuas overlap (Cramp and Simmons 1983) 494 the exact composition of their diets may differ both during the breeding and non-breeding season. Differences in migration route and wintering areas may also contribute to both inter-495 496 and intra-species differences (Steffen et al. 2006, Magnusdottir et al. 2012, Leat et al. 2013). 497

498 The HOCs that dominated the great skua pattern at all colonies (pp'-DDE, PCB-153,

499 oxychlordane, mirex, trans-nonachlor), also dominate the pattern in other seabird species

500 (Bustnes et al. 2005, Helberg et al. 2005, Borgå et al. 2007, Lavoie et al. 2010), due to their

high recalcitrance and lipid solubility. Trophic level and diet specialization are the most likely
explanation for the higher levels in great skuas than other species, in addition to physiological
differences such as metabolic rate, longevity, or biotransformation ability.

504

505 A strength of the present study was the large sample size covering the great skua breeding 506 population of the north-eastern Atlantic which allowed a broader spatial perspective. Large sample sizes are needed to test relationships between dietary descriptors and contaminant 507 508 levels within the same species. These results show that large scale spatial variations in HOC 509 occurrence in great skua is dominated by ecological factors such as local diet and trophic 510 position. In combination with previous studies it can be concluded that these factors are more 511 important than both wintering area and long-range transport of contaminants from primary 512 sources. Condition, lipid content and sex differences are less important factors in the broader 513 spatial perspective. As such, for remote regions, long-range transport and predictions of 514 global abiotic distribution of contaminants is not sufficient to predict risk of contaminant 515 exposure in wildlife.

516

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527

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- 686

	2008	NORTH		SOUTH		2009		SOUTH
	Ny-Alesund (n=5)	Bjørnøya (n=15 ^a)	Hjelmsøya (n=4)	Iceland (n=21)	Runde (n=5)	Bjørnøya (n=51)	Iceland (n=56)	Shetland (n=47)
% Lipids	1 ±0.4	0.84 ±0.42	1.04 ±0.12	0.90 ±0.3	0.90 ±0.12	0.92 ±0.18	0.88 ±0.21	0.87 ±0.17
	0.5-1.6	0.33-1.73	0.96-1.22	0.08-1.5	0.74-1.05	0.30-1.2	0.11-1.4	0.5-1.2
$\delta^{15}N$	13.2 ±0.3	13.9 ±0.4	13.7 ±0.4	13.1 ±0.4	13.3 ±0.3	14.6 ±0.4	13.3 ±0.4	12.7 ±0.4
$\delta^{13}C$	12.8-13.7	13.4 -14.9	13.3-14.2	12.5 - 13.6	13.0-13.7	13.7-15.3	12.0-14.0	11.0-13.6
	-18.2 ±0.2	-19.1±0.3	-19.3 ±0.4	-18.2 ±0.3	-18.8 ±0.2	-19.1 ±0.3	-18.5 ±0.3	-18.5 ± 0.3
	-18.5 to -17.9	-19.8 to -18.8	-19.7 to -18.9	-18.6 to -17.7	-19.0 to -18.6	-19.7 to -18.5	-19.3 to -17.9	-19.2 to -17.9
β-НСН	1.9 ± 1.4	0.497 ±0.443	1.03 ± 1.35	0.532 ±0.628	0.383 ±0.227	2.4 ±2	1.1 ±1	0.192 ±0.192
	0.134-3.8	ND-1.8	0.205-3	ND-2.2	ND-0.56	0.381-11	ND-4.2	ND-1.2
Oxy-Chlordane	14 ± 11	63 ±39	21 ±30	12 ± 11	15 ±22	98 ±69	17 ±16	3.9 ± 7.6
HCB	0.979-35	12-142	4.1-65	ND-44	2.9-54	7.7-335	1.8-92	0.424-51
	12 ±7.4	10 ±8	11 ±9.3	4.7 ±3.1	6.5 ±4.5	23 ±11	4.6 ±2.4	1.9 ±1.8
NC.	1.3-20	2.6-28	2.9-25	0.972-14	2.6-14	4.5-57	0.741-11	0.6-12.5
Mirex	14 ±8.4	18 ±12	9.9 ±8.7	12 ±18	7.7 ±8.3	51 ±34	12 ±11	4 ± 4.5
	1.5-28	ND-39	3-22	ND-88	2.5-22	7.3-134	2-64	0.628-26
pp'-DDE	163 ± 108	196 ±130	179 ±188	175 ±148	160 ± 173	526 ±307	164 ± 158	55 ±60
PCB 153	10-325	40-508	51-458	31-544	50-461	59-1286	19-782	7.7-347
	127 ±69	383 ±181	143 ±102	113 ±106	122 ±91	485 ±301	120 ±102	61 ±55
500	8.9-222	93-676	64-284	19-515	40-269	107-1660	22-509	9-279
ΣOCs	496 ±248	1657 ±820	825 ±600	445 ±458	679 ±519	2410 ± 1452	641 ±507	316 ±249
	39-757)	404-3073	375-1694	81-2270	269-1557	532-7619	141-2754	81-1376
PBDE 47	11 ±7.6	13 ±9.3	12 ± 8.8	3.9 ±4	15 ± 21	17 ±25	7.1 ±4.6	3.9 ± 2
	ND-25	3.4-39	4.9-23	ND-16	2.7-52	ND-127	ND-21	0.935-10

Table 1. Concentrations (µg/kg ww) of selected organochlorines (OCs) and polybrominated diphenylethers (PBDEs) in plasma and stable

isotope signatures of nitrogen ($\delta^{15}N$) and carbon ($\delta^{13}C$) in red blood cells (‰) of adult great skuas by breeding colony and year (arithmetic

689 mean± SD, min-max of concentrations).

687

688

a n=13 for % Lipids

Table 2 Multivariate redundancy analysis of HOC concentrations and patterns in great skua

93 94		Explanatory variable	Variation explained (%)	p-value)	Total variation explained by RDA (%)
95	Concentratio	<u>on</u>			
96	2008 a	Colony	34%	0.001	32%
97		Lipid %	28%	0.003	
98		$\delta^{15}N$	5.3%	0.001	
99		$\delta^{13}C$	6%	0.004	
00					
01	2009	Colony	9.8%	0.001	61%
02		Lipid %	0.6%	0.04	
03		$\delta^{15}N$	3.1%	0.001	
04		$\delta^{13}C$	0.3%	0.001	
05		Sex ^a	3.2%	0.15	
ص ⁰⁶	Patterns				
07	2008 a	Colony	23%	0.001	48%
08		Lipid %	4.5%	0.12	
)9		$\delta^{15}N$	0.6%	0.002	
10		$\delta^{13}C$	0.3%	0.001	
11					
12	2009	Colony	53%	0.001	18%
13		Lipid %	1.6%	0.98	
14		$\delta^{15}N$	7.9%	0.001	
15		$\delta^{13}C$	4.8%	0.001	
16		Sex ^a	12%	0.04	
7	a Sex was no	ot included in th	e 2008 models	as the sex of	f the majority of individuals was

692 plasma collected in 2008 (5 colonies) and 2009 (3 colonies).

718 unknown

Table 3. Diet composition of breeding great skuas from three breeding colonies in 2008 and

- 720 2009 estimated from regurgitated pellets. Number and type of diet specialists (≥70% of pellets
- belonging to one dietary group, Votier et al. 2004), and generalists from study territories
- where more than 5 pellets were recorded.

Colony	Bjørnøya	Bjørnøya	Iceland	Shetland			
Year	2008	2009	2009	2009			
Date	07/07-19/07	12/07-18/08	04/06-11/07	09/06-28/07			
No. of pellets	146	215	115	368			
% Composition							
Fish	23	17	71	57			
Bird	76	82	27	19			
Mammal	N/A	0	1	19			
Other	1	1	1	5			
No of territories with more than 5 pellets							
Specialists	11	13	2	30			
Fish	0	0	2	14			
Bird	8	10	0	1			
Mammal	0	0	0	2			
Generalists	3	3	1	13			

725 FIGURE CAPTIONS

- 726 Figure 1. Relationship between δ^{15} N and δ^{13} C
- Figure 2. Relationship between isotopes and contaminant concentrations
- Figure 3. Relationship between isotopes and diet inferred from pellet type
- Figure 4. Contaminant concentrations between different specialists (diet inferred from pellets)

730

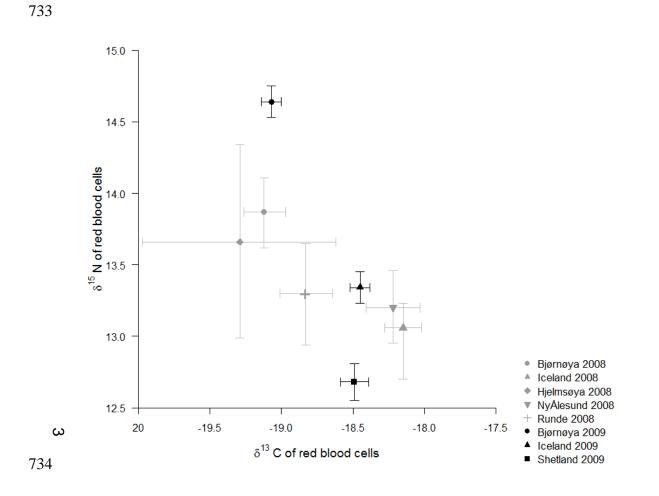


Figure 1. Great skua stable isotope values (‰) of nitrogen ($\delta^{15}N$) and carbon ($\delta^{13}C$) in red blood cell (n = 214) by colony and year. Mean with 95% confidence intervals.

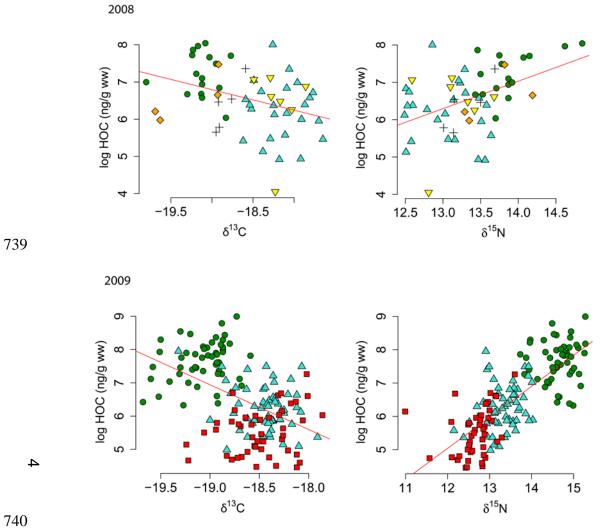


Figure 2. Relationships between the sum of halogenated organic contaminants (HOCs),

concentrations in plasma, and stable isotopes of nitrogen ($\delta^{15}N$) and carbon ($\delta^{13}C$) in red

blood cells of great skuas by colony and year (top: 2008, bottom: 2009).

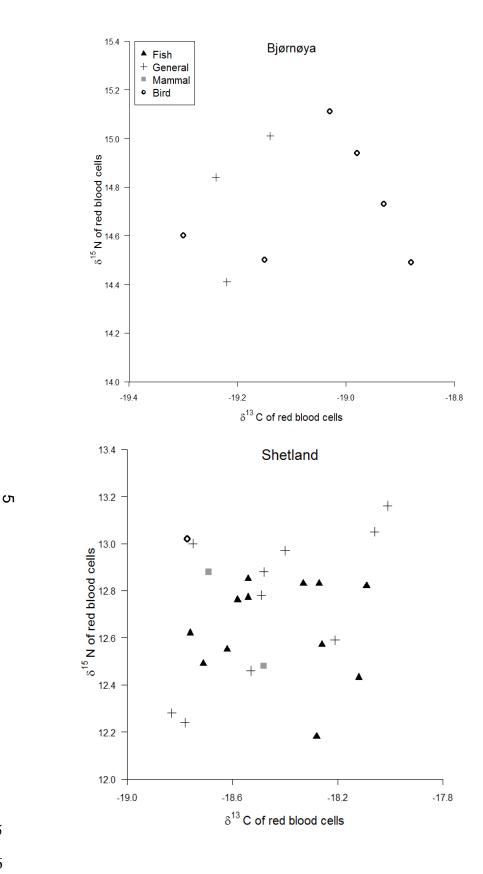
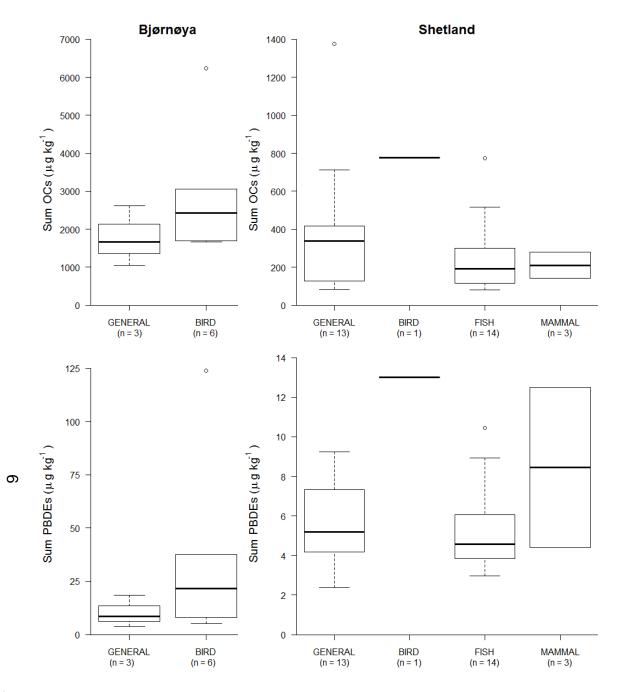


Figure 3. Stable isotope values (‰) of red blood cells of great skuas from Bjørnøya (top) and
Shetland (below), classified from pellet type as specialists or generalists.





750 Figure 4. Sums of organochlorine and PBDE concentrations ($\mu g/kg$, ww) by territorial diet

751 specialisms classified by pellet type.

752