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Eliza H.K. Leat, Sophie Bourgeon, Sveinn A. Hanssen, Aevar Petersen, Hallvard Strøm, Tor Harry Bjørn, Geir W. Gabrielsen, Jan Ove Bustnes, Robert W. Furness, Ane Haarr, Katrine Borgå. The effect of long-range transport, trophic position and diet specialization on legacy contaminant occurrence in great skuas, Stercorarius skua, breeding across the Northeast Atlantic. Environmental Pollution. 244, 2019, 55-65, ISSN 0269-7491.

The article has been published in final form by Elsevier at http://dx.doi.org/10.1016/j.envpol.2018.10.005

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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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ABSTRACT

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

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

INTRODUCTION

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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 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 food web with increasing HOC concentrations with trophic position (Borgå et al. 2001; Ruus et al. 2002; Borgå et al. 2004). As great skuas are migratory, a central question is whether the 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 up to 22% of the variation in HOC concentrations in some of the breeding colonies, leaving much unexplained variance in HOC occurrence to other factors such as local sources during the breeding period (Leat et al. 2013). Local HOC sources, both the baseline exposure and ecological factors like trophic position and diet specialization, may influence the HOC occurrence, as they affect the trophic magnification in the food web (Borgå et al. 2012). A principal source of spatial variation in contaminants in the abiotic environment is the location relative to primary emission sources. Primary emission sources of HOCs have been concentrated in industrialized- and agricultural areas mostly in the mid-latitudes of the northern hemisphere (Breivik et al. 2004). However, long-range transport of HOCs in the atmosphere has distributed these pollutants to remote environments, including the Arctic (Muir and de Wit 2010). Although long-range transport is considered the main source of contaminants to the Arctic, local sources such as settlements and military sites may contribute as well (Brown et al. 2009). One of the physical processes governing long-range atmospheric transport is cold condensation; that is, semi-volatile HOCs shift from gas phase to condensed

phases with decreasing temperature (Wania and Mackay 1993). Differences in chemical's volatility lead to compositional shifts in pollutant pattern along latitudinal and altitudinal temperature gradients, a process often referred to as global distillation (Wania and Mackay 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 than a temperature gradient, that leads to environmental gradients of pollutants (von Waldow et al. 2010a, von Waldow et al. 2010b). Comparisons of HOCs in wildlife between eastern and western Greenland at similar latitude appear to support the remoteness hypothesis, with higher HOC concentrations in eastern Greenland closest to European emission sources (Vorkamp et al. 2004). However, observational data show that latitudinal gradients often coincide with remoteness from emission sources, particularly in Europe (von Waldow et al. 2010a, von Waldow et al. 2010b).

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).

The effect of diet on biomagnification of HOCs is well known and established for food webs using stable isotopes of nitrogen ($\delta^{15}N$) as a measure of relative trophic position (Fisk et al. 2001, Borga et al. 2012). Even within a seabird colony, a positive association between trophic 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.

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

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 long-range 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

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

MATERIALS and METHODS

Sampling

Blood was sampled using heparinized syringes from 204 incubating adult great skuas in five colonies in 2008; Ny-Ålesund 78°55'N, n=5; Bjørnøya 74°29'N, n=15; Hjelmsøya 71°03'N, n=4; South-East Iceland 63°52'N, n=21; and Runde 62°23'N, n=5 (listed with decreasing latitude), and in three colonies in 2009; Bjørnøya n=51; Iceland n=56, and Shetland 60°09'N, n=47 (Table 1, Table S1). Plasma and red blood cells were separated by centrifuging and frozen at -20°C. Birds were sexed by molecular methods using DNA extracted from red blood cells following a modified protocol (Griffiths et al. 1998). Whilst all birds sampled in 2009 were successfully sexed, some of the individuals sampled in 2008 were not due to insufficient quantity of red blood cells for analyses.

Remoteness indices

Remoteness of the colonies was estimated using remoteness indices with night-time light emissions and cropland areas as proxies for HOC emissions from industrial and agricultural 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. Contaminant analyses OCs (HCB, Dieldrin, Mirex, QCB, OCS, \u03b3-HCH, trans-nonachlor, oxychlordane, cisnonachlor, cis-chlordane, pp'-DDT, pp'-DDE, pp'-DDD, and PCB congeners: 31/28, 52, 49, 74, 70, 101, 99, 110, 149, 118, 153, 105/132, 138, 158, 187, 183, 128, 177, 156/171, 180, 191, 170, 201, 195/208, 194, 205, 206, 209) and PBDEs (congeners 47, 99, 100, 153, 154) were extracted and measured in a total of 204 plasma samples at the Great Lakes Institute for Environmental Research (GLIER) at the University of Windsor Ontario Canada by gas chromatography electron capture detection (GC-ECD) (Lazar et al. 1992). Lipid content (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-30 were 77.4 +10% (mean + SD) in 2008 and 69.5 + 9.6 % in 2009. The data were recovery corrected. Method detection limit was 0.08 µg/kg for PCBs and 0.05 µg/kg for all other OCs. Method detection limits for PBDEs varied by congener; PBDE-47 0.373 µg/kg, PBDE-99 0.361 μg/kg, PBDE-100 0.471 μg/kg, PBDE-153 0.420 μg/kg, PBDE-154 0.488 μg/kg.

181 Dietary descriptors

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Stable isotopes of nitrogen and carbon (δ^{15} N and δ^{13} C respectively) were measured in red blood cells as described in detail elsewhere (Leat et al. 2013). The analytical precision based 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 laboratory fish muscle standard was included for 2009 data for which the standard deviations were 0.22% and 0.08% for $\delta^{15}N$ and $\delta^{13}C$, respectively. The variation in spatial baseline isotopic signature among remote sites (Green et al. 2016) is assumed to be negligible compared to signature variation caused by differences in diet in the great skuas. In 2009, breeding territories of blood-sampled birds in each colony were searched for regurgitated pellets every 2-3 days, until the nest failed or fieldwork ceased. In 2008 and 2009, additional sampling of regurgitated pellets from great skuas breeding territories also took place on an ad hoc basis in the Bjørnøya colony. Pellets were examined and classified by prey type; bird, fish, mammal or other, and identified to species level whenever possible. Carcasses of large birds, e.g. ~20 days old great skua chicks and rabbits (Shetland) were counted as equivalent to 1 pellet. Sagittal otoliths recovered from the regurgitated pellets were used to identify the fish species consumed (Härkönen 1986). A minority of all pellets contained more than one prey type and were divided proportionally between the identified groups of prey i.e. 0.5 fish, 0.5 bird. Vegetation in pellets was excluded from the dietary data analysis as it was considered to have been ingested incidentally along with animal dietary items. Territories with > 5 regurgitated pellets were classified as specialists if > 70% of pellets came from one dietary prey group (bird fish or mammal) (Votier et al. 2004). Otherwise, they were classified as generalists. Regurgitated pellets were destroyed once sampled, and identified on site to prevent double counting, or they were removed for further examination. Food regurgitated whilst handling the birds was not included in calculations of diet composition, but was used to identify prey items from the pellets.

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Data analysis

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

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

In all analyses, PCBs were grouped according to homologue group as volatility and longrange transport depends upon the degree of halogenation. Detection rates of the five PBDEs analyzed (PBDE-47, 99, 100, 153, and 154) varied between congeners, colonies and years (Figure S2). PBDE 47 was detected in 97% of the samples and in samples where all five PBDEs were above detection limit (n=89, four individuals sampled in both years were excluded from the 2009 dataset), PBDE 47 was positively correlated with the other PBDEs (Spearman's rank correlations: rho = 0.81, S = 22236, rho = 0.84, S = 18349, rho = 0.61, S = 46348, rho = 0.67, S = 38931, p < 0.001 in all cases) (Figure S3). Therefore PBDE 47 was used as a proxy for commercial penta-BDEs in analysis of spatial variation in PBDE concentration.

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.

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

The sum of the OCs, and PBDE 47, were used to examine the overall relationships between diet, mainly carbon source (δ^{13} C), and relative trophic level (δ^{15} N), and HOCs. Relationships between stable isotopes and HOCs concentrations were investigated using general linear models with backwards stepwise selection to remove non-significant variables. HOC concentration data were log transformed to reduce heterogeneity and skewness (Full model for 2009 in Equation 1).

$$\log \Sigma OC \text{ or } \log \Sigma PBDEs = \delta^{15}N + \delta^{13}C + \% \text{ lipid} + \text{sex} + \epsilon$$
 (Eq. 1)

The relationship between diet measured by regurgitated pellets and stable isotopes was investigated by linear models whereas the relationship between pellets and HOCs was examined visually due to small sample sizes of pellets.

RESULTS AND DISCUSSION

Overall, the data show that great skuas feeding at higher trophic levels and specializing on bird prey (inferred from stable isotopes and regurgitated pellets) have higher concentrations of pollutants, and that this is the main explanation for variation in HOC occurrence among colonies, rather than global distillation and remoteness (colonies spanning nearly 19 degrees latitude) to primary sources and wintering sites (Leat et al. 2013). Large sample sizes were obtained from Bjørnøya, Runde, Iceland and Shetland (15 – 56 individuals), whereas small sample sizes from Hjelmsøya, Runde and Ny-Ålesund (4 – 5 individuals) limit the interpretation of results from these colonies. Contaminant concentrations decreased in the following order: Bjørnøya (2009) >Bjørnøya (2008) > Hjelmsøya (2008) > Runde (2008) > Iceland (2009) > Ny Ålesund (2008) > Iceland (2008) > Shetland (2009). pp'-DDE and PCB 153 were the contaminants of highest concentrations in all colonies for both years, followed

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

HOCs occurrence in great skua plasma and long-range transport

To meet the predictions of global distillation or remoteness hypotheses, Runde/Shetland and 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 intermediate contaminant concentrations compared to those of the other colonies (Table 1). Great skuas from Shetland had lowest contaminant concentrations, despite being the least remote/lowest latitude colony. The HOC concentrations were higher in the remote and high latitude colony Bjørnøya, compared to the other colonies both in 2008 and 2009. Skuas from Bjørnøya had twice as high mean Σ OC concentrations as the second most contaminated colony in 2008 (Hjelmsøya), and more than 3.5 times the mean concentrations from Iceland in both years (Table 1).

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

PCA plots of HOC concentrations and patterns are presented in supporting information (Figure S1). No clear latitudinal gradient was found (selected representative HOCs are presented in Table 1). The direct multivariate redundancy analyses (RDA) accounted for 32% and 60% of the variance in HOC concentrations in 2008 and 2009, respectively (Table 2). 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 in 2009 (Figure S1, Table 2). The direct multivariate redundancy analyses accounted for 52% and 18% of the variance in HOC patterns in 2008 and 2009, respectively (Table 2). 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 (Figure S1, Table 2). Male great skuas had higher HOC concentrations compared to females, which is in accordance with findings in other avian wildlife (Bustnes et al. 2003, Bustnes et al. 2005, Bustnes et al. 2007, Bustnes et al. 2017). However, neither the lipid content or sex were found to be major contributors to the inter-colony difference in HOC occurrence in great skuas. There was no clear latitudinal trend in the relative pattern of HOCs in general, or in PCB homologue patterns in either year. The contaminants with the largest variance in relative contribution among the samples were penta- to nona-CB homologue groups, pp'-DDE, pp'-DDT, oxychlordane and Mirex. If long-range transport would explain the contaminant pattern found in great skuas, the pattern in the Bjørnøya colony should have had proportionally more volatile HOCs, and the pattern in the Iceland colony would have been intermediate to Shetland and Bjørnøya, which is not the case. The lack of a latitudinal or remoteness gradient in concentrations and pattern of HOCs in great skuas is in contrast to abiotic studies (Agrell et al. 1999, Meijer et al. 2003, Gioia et al.

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

al. 2002, Vorkamp et al. 2004, Steffen et al. 2006, Bourgeon et al. 2012, Bustnes et al. 2012a). Most of the species in which latitudinal or remoteness gradients in HOCs have been found are cold blooded species occupying lower trophic levels compared to the great skua (Ter Schure et al. 2002, Vorkamp et al. 2004, Bourgeon et al. 2012, Bustnes et al. 2012a). The HOC pattern in the skuas across the colonies is dominated by recalcitrant and biomagnifying HOCs. Therefore, biotransformation of HOCs in the trophic levels preceding that of the great skua, and in the great skua itself, seem to obscure long-range abiotic transport patterns. In general, biotransformation is efficient in warm blooded species compared to cold blooded lower trophic level species where the contaminant pattern to a higher degree resembles that of the abiotic environment (Borga et al. 2004). The dominance of highly chlorinated PCBs congeners in the pattern of HOCs in Bjørnøya birds in 2008 is indicative of a pattern dominated by biomagnification, rather than long-range transport (Borgå et al. 2001). Relative trophic position and HOC concentrations in great skua The stable isotope signatures differed among colonies (Figure 1, Table 1, MANOVA 2008: Pillai = $0.87 \text{ F}_{448} = 10 \text{ p} < 0.001$; 2009: Pillai = $0.96 \text{ F}_{2150} = 70 \text{ p} < 0.001$) and there were no differences between sexes (2009: Pillai = $0.018 F_{1150} = 1 p = 0.26$). In 2008, great skuas from Bjørnøya had higher δ^{15} N than Iceland, Ny-Ålesund and Runde skuas ($r^2 = 0.42$, $F_{448} = 10.5$, 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 and Shetland skuas (Kruskal-Wallis $\chi^2 = 121$, df = 2, p <0.001, post hoc comparisons Wilcoxon Signed rank tests p<0.001), and Iceland skuas had higher $\delta^{15}N$ than Shetland (p <0.001). As stable isotopes and colony could not be judged as independent variables, colony was excluded from models investigating the relationship between HOC concentrations and stable isotopes.

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Across colonies there was a positive linear relationship between $\delta^{15}N$ and ΣHOC concentrations (2008: R^2 = 0.11, slope=-0.55, p=0.007, 2009: R^2 = 0.59, slope=0.93, p<0.0001, Figure 2). Although lipophilic recalcitrant HOCs show increasing concentrations with trophic position in the food web (Borgå et al. 2001, Fisk et al. 2001, Ruus et al. 2002, Lavoie et al. 2010), such relationships are rarely shown within one species. However, $\delta^{15}N$ levels were found to be positively related to OC concentrations in both bald eagles (Haliaeetus leucocephalus) (Elliott et al. 2009) and white-tailed eagles in Norway (H. albicilla) (Bustnes et al. 2013), and to HCB, DDTs and PCBs (from muscle) in Bjørnøya glaucous gulls (Larus hyperboreus) (Sagerup et al 2002.), and weak positive relationships were found between $\delta^{15}N$ and HOCs in liver of northern fulmars (Fulmarus glacialis) (Knudsen et al. 2007). Even though there are no measurements of baseline $\delta^{15}N$ from the study area, we assume that $\delta^{15}N$ is an indicator of relative trophic level. Green et al. (2016) reported that blue mussels (Mytilus edulis) occupy the same trophic level along the whole Norwegian coast (range: 5-8%), with some stations appearing to be outliers (particularly low or high δ^{15} N). However, the δ^{15} N levels in these stations was attributed to anthropogenic influence, i.e. increased anthropogenic input. Thus, we assume that the difference in isotopic signal across the remote colonies in the present study, reflects differences in diet specializations rather than differences in baseline isotopic signal. In the present study, positive linear relationships between $\delta^{15}N$ and log sum HOCs within the colonies were found for the Iceland colony in 2009 but not 2008 (R²=0.07, 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 Bjørnøya colony in 2008 but not 2009 ($R^2=0.26$, p=0.03, and $R^2=0.004$, p=0.67, respectively). These results suggest variation in individual diet specialization or prey taxon availability between breeding seasons.

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

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Carbon source and HOC occurrence

Bjørnøya, Hjelmsøya and Runde were depleted in δ^{13} C compared to Iceland and Ny-Ålesund in 2008 ($r^2 = 0.74$, $F_{448} = 38$, p < 0.001, Tukey p < 0.005) (Table 1). Likewise, in 2009 skuas from Bjørnøya were depleted in δ^{13} C compared to Iceland and Shetland ($r^2 = 0.52$, $F_{2151} = 83$, p < 0.001, Tukey p < 0.001) (Table 1). Across all colonies, there was a negative linear relationship between carbon source (δ^{13} C) and ΣHOC concentrations (2008: $R^2 = 0.19$, slope=-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 (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 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 findings from the present study. The study by Elliott et al., (2009) also included some individual eagles breeding in estuaries or further up river and therefore reflect more riverine 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 from the present study varied only from -19.8 to -17.7‰, which is at the end of scale compared to the study by Elliott et al. (2009), which might explain the contrasting findings. Spatial patterns in δ^{13} C are well known in the Southern Ocean with δ^{13} C decreasing with 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 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 a greater degree on the terrestrial (less seabird prey, more mammal prey) system as reflected in the regurgitated pellets, and that this further contributes to lower HOC exposure, as HOC 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 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 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.

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Bjørnøya was the colony with the lowest $\delta^{13}C$ 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 $\delta^{13}C$ along a northern-Arctic, latitudinal gradient, as has been done for southern-Antarctic areas.

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Measuring diet – combining stable isotopes and pellet data

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

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

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

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). As in the present study of great skuas, PBDEs are generally found at low concentrations compared to many of the most persistent OCs in the above-mentioned avian top predators (Verreault et al. 2006, Chen and Hale 2010). As with OC, PBDE concentrations in great skuas exceeded those of glaucous gulls (Verreault et al. 2007). Although the dietary niches of glaucous gulls, great black-backed gulls, and great skuas overlap (Cramp and Simmons 1983) 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 interand intra-species differences (Steffen et al. 2006, Magnusdottir et al. 2012, Leat et al. 2013). The HOCs that dominated the great skua pattern at all colonies (pp'-DDE, PCB-153, oxychlordane, mirex, trans-nonachlor), also dominate the pattern in other seabird species (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

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differences such as metabolic rate, longevity, or biotransformation ability.

A strength of the present study was the large sample size covering the great skua breeding 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 levels within the same species. These results show that large scale spatial variations in HOC occurrence in great skua is dominated by ecological factors such as local diet and trophic position. In combination with previous studies it can be concluded that these factors are more important than both wintering area and long-range transport of contaminants from primary sources. Condition, lipid content and sex differences are less important factors in the broader spatial perspective. As such, for remote regions, long-range transport and predictions of global abiotic distribution of contaminants is not sufficient to predict risk of contaminant exposure in wildlife.

ACKNOWLEDGEMENTS

This work was supported by the Research Council of Norway (project 184830). Fieldwork in Shetland was carried out under license from the Home Office and bird trapping and ringing permits from the British Trust for Ornithology. We thank Scottish Natural Heritage and the Holbourn family for permission to work in Foula. Fieldwork in Bjørnøya was carried out under permit from the Governor of Svalbard, Stavanger Museum, and the Directorate for Nature Management, with financial support from the SEAPOP-program (www.seapop.no). Aaron T. Fisk at the Great Lakes Institute of Environmental Research University of Windsor Canada is acknowledged for contaminant and stable isotope analyses. We thank all those who assisted with fieldwork.

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Table 1. Concentrations (μ g/kg ww) of selected organochlorines (OCs) and polybrominated diphenylethers (PBDEs) in plasma and stable isotope signatures of nitrogen (δ^{15} N) and carbon (δ^{13} C) in red blood cells (‰) of adult great skuas by breeding colony and year (arithmetic mean± SD, min-max of concentrations).

	2008	NORTH			SOUTH		NORTH	SOUTH
	Ny-Alesund	Bjørnøya	Hjelmsøya	Iceland	Runde	Bjørnøya	Iceland	Shetland
	(n=5)	$(n=15^a)$	(n=4)	(n=21)	(n=5)	(n=51)	(n=56)	(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
	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
δ^{13} C	-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
	0.979-35	12-142	4.1-65	ND-44	2.9-54	7.7-335	1.8-92	0.424-51
HCB	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
	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
	10-325	40-508	51-458	31-544	50-461	59-1286	19-782	7.7-347
PCB 153	127 ±69	383 ± 181	143 ± 102	113 ±106	122 ±91	485 ±301	120 ± 102	61 ±55
	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

a = 13 for % Lipids

Table 2 Multivariate redundancy analysis of HOC concentrations and patterns in great skua plasma collected in 2008 (5 colonies) and 2009 (3 colonies).

693 694		Explanatory variable	Variation explained (%)	p-value	Total variation explained by RDA (%)
695	Concentration	<u>1</u>			
696	2008 a	Colony	34%	0.001	32%
697		Lipid %	28%	0.003	
698		$\delta^{15}N$	5.3%	0.001	
699		δ^{13} C	6%	0.004	
700					
701	2009	Colony	9.8%	0.001	61%
702		Lipid %	0.6%	0.04	
703		$\delta^{15}N$	3.1%	0.001	
704		δ^{13} C	0.3%	0.001	
705		Sex ^a	3.2%	0.15	
706 ₀	<u>Patterns</u>				
707	2008 a	Colony	23%	0.001	48%
708		Lipid %	4.5%	0.12	
709		$\delta^{15}N$	0.6%	0.002	
710		δ^{13} C	0.3%	0.001	
711					
712	2009	Colony	53%	0.001	18%
713		Lipid %	1.6%	0.98	
714		$\delta^{15}N$	7.9%	0.001	
715		δ^{13} C	4.8%	0.001	
716		Sex ^a	12%	0.04	

a Sex was not included in the 2008 models as the sex of the majority of individuals was

⁷¹⁸ unknown

Table 3. Diet composition of breeding great skuas from three breeding colonies in 2008 and
 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 $\delta^{15}N$ and $\delta^{13}C$
727	Figure 2. Relationship between isotopes and contaminant concentrations
728	Figure 3. Relationship between isotopes and diet inferred from pellet type
729	Figure 4. Contaminant concentrations between different specialists (diet inferred from pellets)
730	
731	

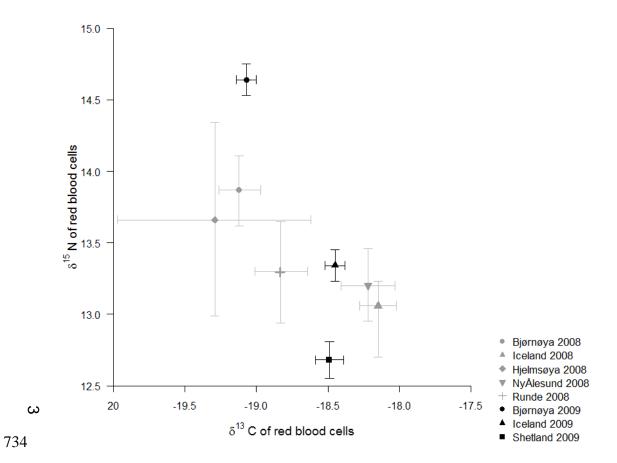


Figure 1. Great skua stable isotope values (‰) of nitrogen ($\delta^{15}N$) and carbon ($\delta^{13}C$) in red blood cell (n = 214) by colony and year. Mean with 95% confidence intervals.

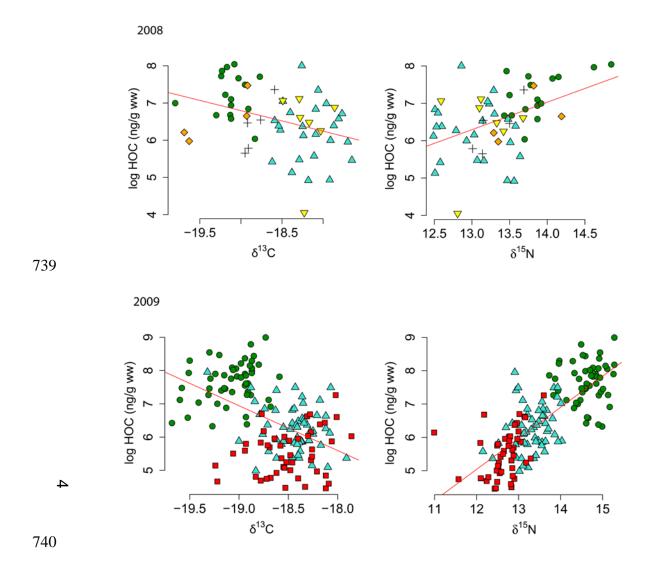


Figure 2. Relationships between the sum of halogenated organic contaminants (HOCs), concentrations in plasma, and stable isotopes of nitrogen ($\delta^{15}N$) and carbon ($\delta^{13}C$) in red blood cells of great skuas by colony and year (top: 2008, bottom: 2009).

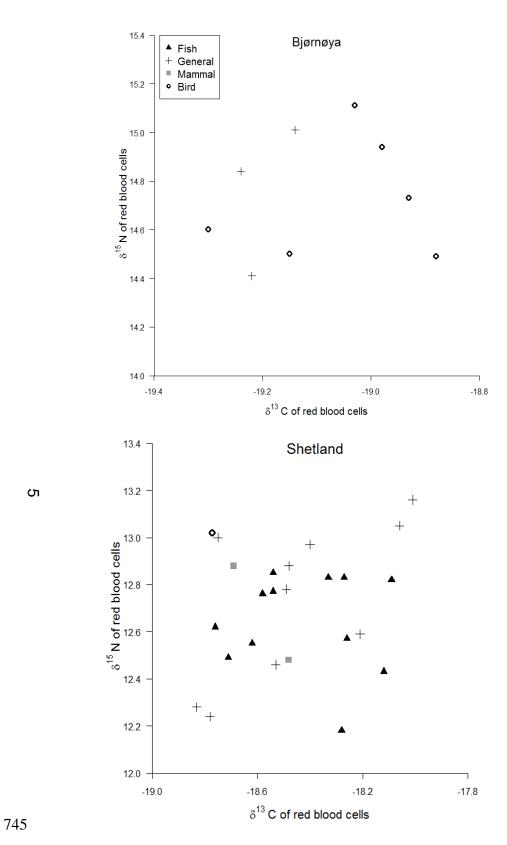


Figure 3. Stable isotope values (‰) of red blood cells of great skuas from Bjørnøya (top) and
 Shetland (below), classified from pellet type as specialists or generalists.

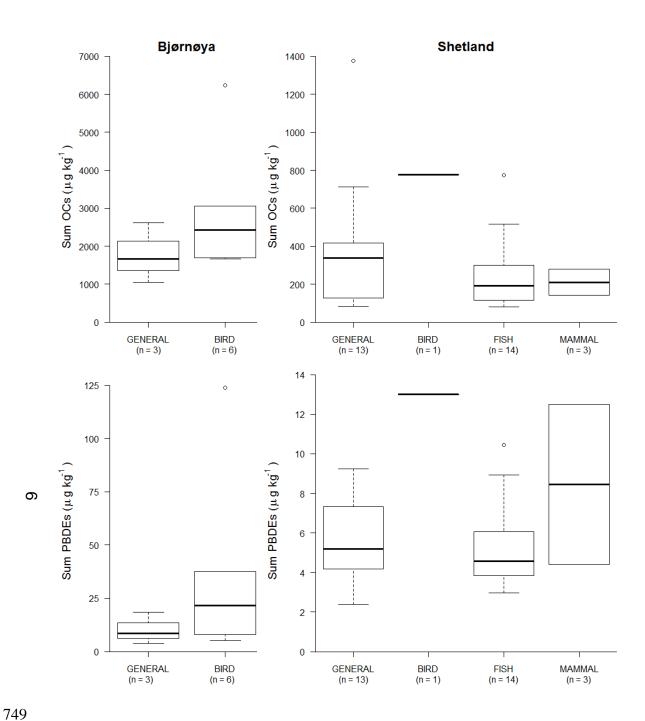


Figure 4. Sums of organochlorine and PBDE concentrations (µg/kg, ww) by territorial diet specialisms classified by pellet type.