

1 Concentrations and profiles of organochlorine contaminants in North Pacific resident and transient killer
2 whale (*Orcinus orca*) populations

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24 **Abstract**

25 Organochlorine (OC) profiles have been used as chemical “fingerprints” to infer an animal’s foraging area.
26 North Pacific killer whale (*Orcinus orca*) populations are exposed to different levels and patterns of OCs based on
27 their prey, distribution, and amount of time spent in a particular area. To characterize concentrations and profiles of
28 OCs found in various populations of North Pacific killer whales, polychlorinated biphenyls (PCBs), including
29 dioxin-like congeners, DDTs, and hexachlorobenzene (HCB), were measured in biopsy blubber samples of photo-
30 identified resident (fish-eating) and transient (mammal-eating) killer whales collected from 1994 through 2002 from
31 Russian Far East waters to the waters of the west coast of the United States, representing 10 populations. We
32 compared blubber OC concentrations based on ecotype (resident vs. transient), sex and reproductive maturity, and
33 geographic area and over OC mixtures were examined to determine if we could detect segregated geographical areas
34 (foraging areas) among the six populations with sufficient sample sizes. Transients had significantly higher OC
35 concentrations than residents and adult male whales had consistently higher OC levels compared to adult females,
36 regardless of ecotype. Our OC profile findings indicate segregated foraging areas for the North Pacific killer
37 whales, consistent with observations of their geographic distributions. Several potential health risks have also been
38 associated with exposure to high levels of contaminants in top-level predators including reproductive impairment,
39 immune suppression, skeletal deformities, and carcinoma. The results of this baseline study provide information on
40 the geographic distribution of OCs found in North Pacific killer whales, results which are crucial for assessing the
41 potential health risks associated with OC exposure in this species.

42 **1.0 Introduction**

43 Killer whales (*Orcinus orca*) are widely distributed throughout the world's oceans, but are primarily found
44 at higher latitudes (Mitchell, 1975; Leatherwood and Dahlheim, 1978; Forney and Wade, 2006). In the North
45 Pacific, three different ecotypes of killer whale have been identified and named "residents," "transients," and
46 "offshores." These ecotypes differ genetically (Hoelzel et al., 1998, 2002; Barrett-Lennard, 2000; Parsons et al.,
47 2013; Moura et al., 2015) and in various aspects of their ecology, morphology, and behavior (Ford and Ellis, 1999;
48 Baird, 2000; Barrett-Lennard, 2000; Ford et al., 2000; Dahlheim et al., 2008; Emmons et al., 2018).

49 Resident killer whales from the North Pacific primarily consume fish with a preference toward salmonids
50 (*Oncorhynchus* spp.), but also consume Pacific herring (*Clupea pallasii*), Pacific halibut (*Hippoglossus stenolepis*),
51 lingcod (*Ophiodon elongates*), and rockfish (*Sebastes* spp.) (Ford et al., 1998; Saulitis et al., 2000; Ford and Ellis,
52 2006; Hanson et al., 2010; Ford et al., 2016). Because of their preference for salmon, seasonal movements of
53 resident killer whales are strongly tied to Pacific salmon migrations (Ford, 2009). In contrast, the preferred diet of
54 transient killer whales consists of marine mammals with movements linked to the presence of their prey in coastal
55 waters (Baird, 2000; Ford et al., 1998, 2000; Saulitis et al., 2000; Ford, 2009; Dahlheim and White, 2010).
56 Transient killer whales from the North Pacific target a wide variety of prey (e.g., Northern fur seals [*Callorhinus*
57 *ursinus*], harbor seals [*Phoca vitulina*], Steller sea lions [*Eumetopias jubatus*], gray whales [*Eschrichtius robustus*],
58 minke whales [*Balaenoptera acutorostrata*], Dall's porpoise [*Phocoenoides dalli*], and harbor porpoise [*Phocoena*
59 *phocoena*]) that likely changes seasonally (Jefferson et al., 1991; Dahlheim and White, 2010). Less information is
60 available on the prey consumed by offshore killer whales. This ecotype has been observed preying on fishes and
61 carcharinid sharks (Ford, 2009; Heise et al., 2003; Herman et al., 2005; Jones, 2006; Krahn et al., 2007a; Dahlheim
62 et al., 2008).

63 There are several resident killer whale populations in the eastern North Pacific. Two resident killer whale
64 populations, the Southern Resident killer whales and Northern Resident killer whales, inhabit the coastal waters of
65 Washington State, USA and southern British Columbia, Canada (WA/BC). The Southern Resident killer whales are
66 observed as far south as central California and as far north as the southern waters of southeast Alaska (Bigg et al.,
67 1990; Ford et al., 2000; Hanson et al., 2013; Carretta et al., 2017). Northern Resident killer whales primarily inhabit
68 the coastal and inland waters of British Columbia with travels south into Washington State waters and north into
69 southeast Alaska waters (Dahlheim et al., 1997; Ford et al., 2000). There are also several resident killer whale

70 populations found in southeast Alaska (SEAK), throughout the Gulf of Alaska (GOA), and westward to the eastern
71 and central Aleutian Islands (EAI and CAI, respectively) and Bering Sea (Dahlheim and Waite, 1993; Matkin et al.,
72 1999a; Matkin et al., 2007; Muto et al., 2016). Although the ranges of eastern North Pacific resident killer whales
73 overlap, each population appears to have a defined core area, an area the whale populations frequently inhabit
74 (Figure 1). Within each core area, whether it is offshore or more coastal, available prey species vary. Human
75 development and anthropogenic threats also vary within each core area. For example, the Southern Resident killer
76 whale's core area overlaps with high human population and urban development, whereas killer whales from more
77 remote core areas off Alaska are exposed to less industrial development and other human-related activities.

78 Similar to eastern North Pacific resident killer whales, several populations of transient killer whales occur
79 in marine waters from California through the GOA, Aleutian Islands, and Bering Sea (Matkin et al., 1999a; Matkin
80 et al., 2012; Muto et al., 2016). Most transient killer whales found in SEAK have been well documented (via
81 photographic matches) to frequently occur throughout the coastal waters of WA/BC (Dahlheim and White, 2010).
82 Conversely, no photographic matches have been found among SEAK transients and whales known from Prince
83 William Sound (Matkin et al., 1999b), or from the GOA, Aleutian Islands, and Bering Sea (Dahlheim, 1997).
84 Recent genotypic and observational data strongly suggest that a subdivision occurs between transients found in the
85 EAI with those that occur in the GOA (Matkin et al., 2007; Durban et al., 2010; Matkin et al., 2012; Parsons et al.,
86 2013). The genotypic data indicate that the eastern point of this subdivision between transients in the EAI and the
87 GOA likely occurs in and near the waters surrounding Kodiak Island. Amchitka Pass represents a division between
88 central and western Aleutian Islands (CAI and WAI) transients (Parsons et al., 2013). Subpopulations of transients
89 were also apparent in the eastern Aleutian Islands and Bering Sea (Parsons et al., 2013).

90 Organochlorines (OCs), such as DDTs and polychlorinated biphenyls (PCBs), are persistent, widespread
91 environmental contaminants that are resistant to metabolism and have been shown to bioaccumulate through marine
92 food webs (Fisk et al., 2001; Ruus et al., 2002; Hoekstra et al., 2003). However, some large marine mammal species,
93 notably polar bears (*Ursus maritimus*), have an excellent ability to metabolize some OCs (Muir et al., 1988; Letcher
94 et al., 1996, 1998). Accordingly, OCs have been used as intrinsic chemical tracers to infer sources of OCs, foraging
95 areas, and migration patterns of many marine species (Ramos and González-Solís, 2012), including Pacific herring
96 (West et al., 2008), Atlantic salmon (*Salmo salar*) (Svendsen et al., 2008; 2009), bluefish (*Pomatomus saltatrix*)

97 (Deshpande et al., 2016a), bluefin tuna (*Thunnus thynnus*) (Deshpande et al., 2016b), Greenland sharks (*Somniosus*
98 *microcephalus*) (Fisk et al., 2002), bottlenose dolphins (*Tursiops truncatus*) (Borrell et al., 2006; Fair et al., 2010),
99 humpback whales (*Megaptera novaeangliae*) (Elfes et al., 2010), and killer whales (Herman et al., 2005; Krahn et
100 al., 2007a). Marine environments have distinct OC patterns based on a variety of historical inputs (e.g., industrial
101 discharges, atmospheric deposition, current transport, etc.), and animals that do not readily metabolize these OCs
102 and forage for extended periods of time can accumulate OCs in proportion to their availability in those
103 environments. Species found in more coastal waters near high urban and human development regions are
104 potentially exposed to higher levels of OCs than species found in offshore waters and consequently will have higher
105 body burdens of these pollutants. However, remote, undeveloped offshore areas can receive some input of OCs
106 through various routes, such as long-range atmospheric transport from industrialized areas (Wania and Mackay,
107 1996).

108 Previous studies have reported concentrations of OCs in tissues of killer whales in the North Pacific.
109 Calambokidis et al. (1984) first reported high levels of PCBs and DDTs in blubber of an adult male transient killer
110 whale (PCBs: 250 $\mu\text{g g}^{-1}$, wet wt.; DDTs: 640 $\mu\text{g g}^{-1}$, wet wt.) and an adult male Southern Resident killer whale
111 (PCBs: 38 $\mu\text{g g}^{-1}$, wet wt.; DDTs: 59 $\mu\text{g g}^{-1}$, wet wt.) that stranded off the coasts of British Columbia and
112 Washington State, respectively, in the late 1970s. Since then, concentrations of environmental contaminants have
113 been measured in tissues of other stranded or incidentally caught killer whales from the North Pacific Ocean (Ono et
114 al., 1987; Jarman et al., 1996; Hayteas and Duffield, 2000; Krahn et al., 2004; Kajiwara et al., 2006). More recently,
115 concentrations of lipophilic contaminants have been determined in blubber biopsy samples of wild-ranging killer
116 whales (Ross et al., 2000; Ylitalo et al., 2001; Rayne et al., 2004; Herman et al., 2005; Krahn et al., 2007a, 2007b,
117 2008, 2009; McHugh et al., 2007; Wolkers et al., 2007; Noël et al., 2009; Jepson et al., 2016; Atkinson et al., 2019).
118 These studies, as well as others, have revealed large variability in OC levels across regions and within a species.

119 To increase our knowledge of baseline concentrations and profiles of OCs in various populations of North
120 Pacific killer whales and how these vary geographically over the North Pacific, PCBs (including dioxin-like
121 congeners) and OC pesticides were measured in biopsy blubber samples of 98 wild-ranging resident and transient
122 killer whales sampled from Russian Far East waters to the waters of Washington State, USA. We examined the
123 contribution of dioxin-like PCBs contributing to sum PCB total toxic equivalents ($\sum\text{PCB TEQ}$) values as well as the

124 individual DDT and PCB congeners contributing to their respective sum concentrations based on sex and
125 reproductive maturity category. Lastly, we examined the OC mixtures in whale blubber samples to assess whether
126 we could detect segregated foraging areas among six killer whale populations. We hypothesized that populations of
127 North Pacific killer whales would differ in their OC concentrations and profiles based on their ecotype (i.e., resident
128 or transient), foraging area (i.e., geographic distribution and location of core feeding area), and their sex and
129 reproductive maturity category (i.e., adult males, adult females, juveniles or unknown status).

130 **2.0 Materials and Methods**

131 2.1 Field Sample Collection

132 From 1994 through 2002, biopsy samples were collected from 98 free-ranging North Pacific killer whales.
133 Remote biopsy sampling techniques were similar to those described in Barrett-Lennard et al. (1996) and Ylitalo et
134 al. (2001). A small core containing skin and blubber (approximately 2.0 to 3.0 cm in length and 0.5 cm in diameter)
135 was obtained from each animal and the blubber portion of the biopsy sample was stored at -20° C until chemical
136 analysis.

137 In the current study, killer whales that were biopsy sampled were visually identified using photo-
138 identification catalogues (Dahlheim, 1997; Matkin et al., 1999a). Using the method of Bigg et al. (1986),
139 photographs were taken of the individual killer whales at the time they were biopsy sampled to confirm
140 identification. Sex assignment (i.e., male, female or unknown) was based on direct observations of whales in well-
141 studied populations (e.g., WA/BC, SEAK) or determined genetically for other populations. Reproductive maturity
142 class (i.e., juvenile, adult or unknown) of each whale was assessed in the field based on long-term sighting data or, if
143 not known, was assigned based on relative body size as described in Herman et al. (2005). We use sex and
144 reproductive maturity class assignments to create four sex/maturity categories (i.e., adult female, adult male,
145 juveniles, and unknowns). Killer whale populations sampled as part of this study are listed in Supplemental Table 1,
146 based on their ecotype (69 residents and 29 transients) and geographic sampling area (i.e., assumed foraging area).
147 Sex and reproductive maturity-category data (e.g., adult males, adult females, and juveniles) are known for the
148 majority of whales sampled and are provided in Supplemental Table 1. The population geographic area assignments
149 follow those described by Herman et al. (2005).

150 Ninety eight killer whales from the North Pacific were sampled in this study from the following
151 geographical areas: 1) Northern Resident (n = 1) and Southern Resident (n = 3) killer whales from Washington
152 State/British Columbia (WA/BC); 2) resident (n = 19) and transient (n = 20) killer whales from southeast Alaska
153 (SEAK); 3) resident (n = 18) and transient (n = 2) killer whales from Gulf of Alaska/Prince William Sound (GOA);
154 4) resident (n = 20) and transient (n = 7) killer whales from eastern Aleutian Islands (EAI); and 5) resident killer
155 whales (n = 3) from Central Aleutian Islands (CAI). For the purposes of this paper, we also include data on resident
156 killer whales (n = 5) that occur in the western North Pacific (WNP) (Figure 1, Supplemental Table 1).

157 2.2 OC and Lipid Analyses

158 Killer whale biopsy blubber samples were analyzed at the National Marine Fisheries Service's Northwest
159 Fisheries Science Center in Seattle, WA for OCs using a high-performance liquid chromatography/photodiode array
160 detection (HPLC/PDA) method (Krahn et al., 1994). In this method, blubber (0.1 to 0.3g), sodium sulfate (5g),
161 hexane/pentane (1:1 v/v), and the surrogate standard (1,2,3,4-tetrachloro-p-dibenzodioxin; 250 ng) were
162 homogenized two times and the extracts combined. The analytes were separated from interfering compounds on a
163 gravity flow cleanup column that contained neutral, basic, and acidic silica gels with hexane/methylene chloride (1:1
164 v/v). Prior to the cleanup step, a 1-mL aliquot sample extract was removed for lipid quantitation by thin layer
165 chromatography with flame ionization detection (TLC/FID) (Ylitalo et al., 2005a). The remaining sample extract
166 was analyzed for eight dioxin-like PCB congeners (PCBs 77, 105, 118, 126, 156, 157, 169, 189), six other PCB
167 congener groups (PCB 138, PCB 180, PCBs 101/99/149/196, PCBs 128/123, PCBs 153/87, PCBs 170/194), *o,p'*-
168 DDD, *p,p'*-DDD, *p,p'*-DDE, *o,p'*-DDT, *p,p'*-DDT, and hexachlorobenzene (HCB) using a high-performance liquid
169 chromatography with photodiode array detection method (Krahn et al., 1994). The dioxin-like congeners were
170 resolved from other PCBs and OCs (listed above) by HPLC on two Cosmosil PYE analytical columns connected in
171 series and cooled to 16° C. The congeners were measured by an ultraviolet (UV) photodiode array detector and
172 were identified by comparing their UV spectra (collected from 200 to 310 nm) and retention times to those of
173 reference standards in a library. The purity of each analyte was confirmed by comparing spectra within a peak to the
174 apex spectrum. The lower limit of quantitation (LOQ) for PCBs and DDTs ranged from 0.37 to 4.2 ng g⁻¹, wet
175 weight (ww) and 1.2 to 5.6 ng g⁻¹, ww, respectively. The LOQ for HCB ranged from 0.34 to 1.9 ng g⁻¹, ww. The
176 ranges of these LOQ values are typical for the HPLC/PDA and are comparable to or lower than those reported in

177 other studies in which blubber samples were analyzed using the same analytical methods (Ylitalo et al., 2001;
178 Ylitalo et al., 2005; Greig et al., 2007).

179 Concentrations of summed PCBs (Σ PCBs) were calculated using HPLC/PDA analyte concentration data
180 and the following formula: Σ PCBs = Σ concentrations of 14 PCBs and PCB analyte groups listed above (based on
181 individual response factor) + Σ concentrations of “other PCBs” (calculated by summing areas of peaks identified as
182 PCBs and using an average PCB response factor). The response factors of the PCB congeners measured on the
183 HPLC/PDA system are similar (ranging from 0.65 to 0.75) regardless of degree of chlorination or other chemical
184 properties, using an average PCB response factor was warranted for the “other PCBs” measured in the samples.
185 Based on retention times and UV spectral data, the other PCBs included PCBs 31, 66, 70, 110, 182, 190, 200, and
186 202. Summed DDT (Σ DDTs) concentrations were calculated by adding the concentrations of *o,p'*-DDD, *p,p'*-DDD,
187 *p,p'*-DDE, *o,p'*-DDT, and *p,p'*-DDT. Summed PCBs, Σ DDTs, and HCB were reported as $\mu\text{g g}^{-1}$ lipid weight (lw).

188 In order to compare the PCB TEQ concentrations in the current study with those previously reported in
189 killer whales from Kenai Fjords/Prince William Sound, Alaska, we used the mammalian toxic equivalent factors
190 reported in Van den Berg et al. (1998). The PCB TEQs were calculated by multiplying the molar concentration of
191 each dioxin-like congener by the appropriate toxic equivalency factor (TEF) value for that compound. These TEQ
192 concentrations are conservative values as they are calculated solely on measurable concentrations of eight dioxin-
193 like PCBs (i.e., PCBs 77, 105, 118, 126, 156, 157, 169, 189) and did not include polychlorinated dibenzo-*p*-dioxins
194 (PCDDs) and polychlorinated dibenzo-*p*-furans (PCDFs). In addition, the LOQ values of our PDA are higher than
195 those of low and high resolution gas chromatography/mass spectrometry and thus may also contribute to
196 underestimated TEQ values. In previous studies, PCDDs and PCDFs, as well as dioxin-like PCB congeners, were
197 measured in the blubber of killer whales and TEQs were determined from the data (Kannan et al., 1988; Jarman et
198 al., 1996; Ross et al., 2000; Kajiwara et al., 2006; Noël et al., 2009). Although these studies reported wide ranges of
199 the TEQ values in these animals, dioxin-like PCBs contributed a much larger percentage (> 80%) to the TEQs than
200 PCDDs and PCDFs (< 20%).

201 Biopsy blubber samples of the killer whales were analyzed for lipid classes and percent lipid using a Mark
202 5 Iatroscan TLC/FID (Ylitalo et al., 2005a). Various lipid classes (i.e., sterol esters/wax esters, triglycerides, free
203 fatty acids, cholesterol, polar lipids) were separated on silica-based Chromarods (SIII) and developed in a solvent

204 system containing 60:10:0.02 hexane:diethyl ether:formic acid (v/v/v). The total percent lipid values were
205 calculated by adding the concentrations of these lipids classes and were reported as percent lipid.

206 2.3 Quality Assurance

207 Each sample batch contained 8 to 12 field samples, a method blank, and a National Institute of Standards
208 and Technology (NIST) blubber Standard Reference Material (SRM 1945) and were analyzed by HPLC-PDA
209 (Krahn et al., 1994; Ylitalo et al., 2001). The concentrations of $\geq 70\%$ of individual analytes that were measured in
210 the NIST SRM 1945 were within 35% of either end of the 95% confidence interval range of the published NIST
211 certified or reference OC concentration of that analyte. Method blanks contained no more than four analytes that
212 exceeded four times the LOQ, unless the analyte was not detected in the associated blubber samples in the set. The
213 percent recoveries of the surrogate standard for all field and quality assurance samples ranged from 71% to 99%.

214 2.4 Statistical Analyses

215 2.4.1 OC Concentrations and Ratios in North Pacific Killer Whales

216 Prior to these analyses, the OC concentrations (not the ratios) were lipid normalized to account for lipid
217 variation among samples (Balmer et al., 2019). These data were then log transformed to meet the criteria for normal
218 distribution and equal variances. We used a generalized linear modelling (GLM) approach to investigate the extent
219 to which variability in contaminant levels (i.e., dependent variables of HCB (ng/g, lw), \sum PCBs (ng/g, lw), \sum DDTs
220 (ng/g, lw), \sum PCB TEQs (pg/g, lw) and OC ratios (*p,p'*-DDT / \sum DDTs, and \sum DDTs / \sum PCBs) for the 98 killer
221 whales was explained by a suite of three independent, fixed effects, variables [i.e., ecotype, foraging area (habitat),
222 sex/maturity category] and one interaction term (ecotype*sex/maturity category). For each dependent variable, we
223 ran all possible model combinations of the four variables (i.e., 15 combinations). Collection year and whale
224 population were not included in our GLM analyses as these data were skewed for certain whale groups (e.g., SEAK
225 killer whales were sampled primarily from 1994 – 1997 whereas the other North Pacific whales were sampled
226 primarily in 2001 and 2002).

227 All model parameters were estimated by maximizing the likelihood function. To compare models, we
228 calculated four values for each model; Akaike's information criterion (AIC), delta AIC, relative likelihood and AIC
229 weight. Smaller AIC values indicate "better" models, and when comparing two models, we calculated the difference
230 in AIC values (delta AIC; Akaike, 1973; Burnham et al., 2011). A delta AIC of less than 2 indicates little difference
231 between competing models; a delta AIC of 2–10 indicates moderate support for a difference between the models,

232 and a delta AIC of greater than 10 indicates strong support (Burnham et al., 2011). Relative likelihood represents the
233 likelihood of a model given the data, whereas AIC weight is the discrete probability of each model (Burnham et
234 al.,2011). The best model was defined as having a delta AIC of 0.00, although preference was given to the simplest
235 model if two or more models had a delta AIC of less than 2. All statistical analyses were conducted in R version
236 3.5.2 (R Core Team 2018).

237 Analysis of variance (ANOVA) and the Tukey–Kramer Honestly Significant Difference (HSD) pairwise
238 post-hoc test were used to compare mean $\log(\sum\text{DDTs lw})$, $\log(\sum\text{PCBs lw})$, and $\log(\sum\text{PCB TEQs lw})$, and
239 untransformed mean OC ratios ($\sum\text{DDTs} / \sum\text{PCBs}$, and p,p' -DDE / $\sum\text{DDTs}$) between ecotypes, among geographical
240 areas and age/maturity categories, and the interaction (ecotype*sex/maturity category). For mean $\log(\text{HCB lw})$, we
241 compared mean values between ecotypes and sex/maturity categories. The Tukey–Kramer HSD test is one of a
242 number of post-hoc methods recommended to use to test differences between pairs of means among groups that
243 contain unequal sample sizes (Zar, 1999).

244 We also examined differences in mean levels and ratios of the OCs for adult resident males only ($n = 33$)
245 from four geographical areas (WNP, EAI, GOA, and SEAK). Data from adult males were used to avoid any
246 confounding issues associated with differences in OC levels and patterns based on maternal offloading to offspring.
247 Limited samples sizes precluded comparisons among resident adult male whales from other populations or among
248 other sex/maturity categories. The level of significance used for all statistical tests was $p \leq 0.05$. The univariate
249 analysis was completed using JMP Statistical Software (SAS Institute, Inc., Cary, NC).

250 *2.4.2 Assessing Segregated Diet and Foraging Areas among Killer Whale Populations*

251 Principal Component Analyses (PCA), as detailed in the software package Primer-E version 6 (Clark and
252 Warwick, 2006; Clark and Gorley, 2006) was used to further evaluate segregation in the relative abundance of OCs
253 in individual whales among resident and transient populations, potentially due to differences in diet and foraging
254 area among populations and/or differences in the composition of sex/maturity categories among populations. The
255 PCA was limited to a comparison among the six populations that had at least five biopsy samples (EAI residents,
256 GOA residents, SEAK residents, WNP residents, EAI transients, and SEAK transients), and included all
257 sex/maturity class samples for each population. Although OC mixtures in marine mammals can differ among sex
258 and reproductive maturity categories because females preferentially offload less hydrophobic contaminants to their

259 offspring via lactation (Ridgway and Reddy 1995, Pomeroy et al., 1996, Ylitalo et al., 2001, Debier et al., 2003,
260 Desforges et al., 2012), we hypothesized (and tested) that these differences would not be as pronounced as those
261 associated with variation in diet and foraging area among transient and resident populations. Dissimilar contaminant
262 mixtures (i.e., fingerprints) among populations of the same sex/maturity category would suggest inputs of specific
263 OCs associated with different sources (i.e., diet and foraging area of resident and transient populations).

264 The OC data selected for the PCA analyses excluded three of the 20 OCs (PCB 77, 126, and 169) because
265 they were detected in less than 30% of the samples. Fifteen of the samples had blank values for at least one analyte
266 because of interference by unidentified analytes preventing quantification. These interference-blank values were
267 randomly distributed among the sample-analyte combinations, except for one sample from a SEAK resident whale,
268 which had five blanks and was excluded from the analyses, resulting in 88 samples. The remaining missing values
269 were replaced with a value calculated by an expectation maximum likelihood algorithm (Primer-E, version 6). For
270 the remaining 17 analytes originally included in the PCA, undetectable values (131 of 1,496 sample analyte
271 combinations, 8.8% of the data set) were replaced with the average of the analyte-specific LOQ for this study. Prior
272 to analyzing with PCA, the OC data were pretreated by standardizing (i.e., computing the proportional contribution
273 of each OC compound concentration to the total OC concentration in each sample) and then transforming the data
274 by taking the square root to reduce the contribution of dominant compounds. The total number of OCs input to the
275 PCA procedure ($n = 17$) was reduced to five compounds that most efficiently described the OC patterns (HCB, PCB
276 180, *p,p'*-DDE, and *o,p'*-DDT, and *p,p'*-DDT,) by using the BEST/BVSTEP procedure (Primer-E, version 6) to
277 select OCs that contribute most to explaining the observed OC mixtures. The BEST/BVSTEP identifies the smallest
278 possible subset of a data (i.e., subset of OCs in this case) which, in combination, describes most of the pattern ($\rho >$
279 0.95) of the full data (i.e., all 17 OCs).

280 To test for significant difference in overall OC mixtures among populations (with all sex/maturity
281 categories included), pairwise comparisons of population mixtures were conducted with ANOSIM, using the R
282 statistic to identify the degree of segregation between-groups as detailed in the software package Primer-E version 6
283 (Clark and Warwick, 2006; Clark and Gorley, 2006). Values of the ANOSIM R statistic range from 0 (i.e., no
284 segregation, or complete similarity) to 1.0 (i.e., complete segregation, or no similarity) of a population. A p value of
285 < 0.05 was used as a guide for determining whether the measured segregation between populations (i.e., R statistic)
286 was statistically significant. Additionally, to evaluate difference in OC mixtures among populations due to diet and

287 foraging area (i.e., ecotype and geographic area), independent of the composition of sex/maturity categories among
288 populations, PC1 scores of whales of the same sex/maturity category were analyzed for significant difference among
289 populations using a t-test or ANOVA and Holm-Sidak post-hoc tests. The level of significance used for all statistical
290 tests was $p \leq 0.05$.

291 *2.4.3 Toxicological risks of OCs to North Pacific killer whales*

292 Concentrations of \sum PCBs and \sum PCB TEQs (see Supplemental Table 1) measured in individual North
293 Pacific killer whales in the current study were compared to threshold values associated with immune suppression
294 and vitamin A depression in harbor seals (PCBs 17 and TEQs 209 $\mu\text{g g}^{-1}$ lw), (De Swart et al., 1996; Ross et al.,
295 1996; Kannan et al., 2000) and reproductive dysfunction (PCBs 77 $\mu\text{g g}^{-1}$ lw) in ringed seals (Boon et al., 1987).

296 **3.0 Results**

297 *3.1 OC Concentrations and Ratios in North Pacific Killer Whales*

298 A wide range of OC concentrations and percent lipid values were determined in the blubber of killer whales
299 from the North Pacific (Supplemental Table 1). Among animals of known sex/maturity category, the lowest
300 concentrations of HCB (0.052 $\mu\text{g g}^{-1}$ lw), \sum DDTs (2.6 $\mu\text{g g}^{-1}$ lw), \sum PCBs (2.5 $\mu\text{g g}^{-1}$ lw), and \sum PCB TEQs (23 pg g^{-1}
301 lw) were measured in the blubber of adult female resident killer whales. The highest concentrations of \sum PCBs and
302 \sum PCB TEQs (920 $\mu\text{g g}^{-1}$ lw and 4,700 pg g^{-1} lw, respectively) were found in an adult female transient killer whale.
303 A juvenile transient killer whale had the highest concentration of \sum DDTs (1,700 $\mu\text{g g}^{-1}$ lw). The highest
304 concentration of HCB (11 $\mu\text{g g}^{-1}$ lw) was found in the blubber of both a juvenile and an adult male transient whale.
305 We also found that, regardless of ecotype, the rank order of OCs measured was \sum DDTs > \sum PCBs >> HCB.

306 The GLM results for the North Pacific killer whales indicated that most of the variability in the lipid-
307 normalized concentrations of HCB was best explained by ecotype*sex/maturity category interaction (Table 1;
308 Figure 2). Although the results of ANOVA revealed a significant interaction ($p < 0.001$) ecotype*sex/maturity
309 category for the $\log(\text{HCB lw})$, these differences could not be distinguished by Tukey-Kramer HSD post-hoc
310 pairwise tests. For \sum DDTs, \sum PCBs, and \sum PCB TEQs, the variability was best explained by the
311 ecotype*sex/maturity category interaction followed by geographical area (Table 1; Figures 3 through 5). ANOVA
312 results for the $\log(\sum\text{DDTs lw})$ and $\log(\sum\text{PCBs lw})$ showed significant differences between ecotypes ($p < 0.001$) and
313 among age/maturity categories ($p < 0.01$) but Tukey-Kramer HSD did not find differences among the sex/maturity

314 categories (Figures 3 and 4). Non-significant differences in $\log(\sum\text{DDTs lw})$ and $\log(\sum\text{PCBs lw})$ were found among
315 geographical areas or in the ecotype*sex/maturity category interaction. ANOVA tests revealed non-significant
316 differences in $\log(\sum\text{PCB TEQs lw})$ among geographical areas but found significant interaction of
317 ecotype*sex/maturity category ($p < 0.05$) (Figure 5). However, differences among the categories were not detected
318 using the Tukey-Kramer post-hoc tests.

319 For the other two dependent variables, ratios of p,p' -DDE / $\sum\text{DDTs}$ and $\sum\text{DDTs} / \sum\text{PCBs}$, the variability
320 was best explained by ecotype, sex/maturity category, and geographical area (Table 1; Figures 6 - 7). We found
321 significant differences in mean p,p' -DDE / $\sum\text{DDTs}$ ratios between ecotypes ($p < 0.001$), with transients having a
322 higher mean ratio than the resident mean ratio. Significant differences for this ratio were also found among
323 geographical areas ($p < 0.001$), with the CAI and EAI whales having lower mean ratios compared to the ratios
324 determined in SEAK and SRKW whales. Significant differences in mean ratios of p,p' -DDE / $\sum\text{DDTs}$ were found
325 among sex/maturity categories ($p < 0.05$) but the post-hoc pairwise tests did not denote differences among the
326 categories. Significant differences in the ratio of $\sum\text{DDTs} / \sum\text{PCBs}$ were found between ecotypes ($p < 0.01$) and
327 geographical area ($p < 0.0001$), with transients having higher mean ratios than residents and whales from EAI and
328 SEAK having higher mean $\sum\text{DDTs} / \sum\text{PCBs}$ ratios compared to the mean value determined for the GOA whales.
329 Among sex/maturity categories, no significant differences were found in either of these ratios.

330 To assess if we could detect any additional patterns within a similar sex/maturity category, ANOVA and
331 Tukey's HSD were used to compare mean concentrations of OCs and mean OC ratios among WNP, EAI, GOA, and
332 SEAK resident adult male killer whales. The mean (\pm SE) OCs (HCB, $\sum\text{DDTs}$, $\sum\text{PCBs}$), $\sum\text{PCB TEQs}$ and percent
333 lipid values varied among adult male residents from four populations (WNP, EAI, GOA, and SEAK; Table 2). The
334 percent lipid values in the blubber biopsy samples from adult male WNP, EAI, GOA, and SEAK resident killer
335 whales ranged from 5.0% to 38% (Table 2). Adult male WNP resident killer whales had statistically significant
336 lower lipid percent in blubber biopsy values compared to the EAI ($p = 0.0489$) and GOA ($p = 0.0455$) adult male
337 residents. SEAK adult male residents had significantly higher mean concentrations of $\sum\text{DDTs}$ ($p = 0.0120$), $\sum\text{PCBs}$
338 ($p = 0.0200$), and $\sum\text{PCB TEQs}$ ($p = 0.0143$) compared to EAI adult male residents. SEAK adult male residents also
339 had significantly higher mean concentrations of $\sum\text{DDTs}$ ($p < 0.0001$), $\sum\text{PCBs}$ ($p = 0.0014$), and $\sum\text{PCB TEQs}$ ($p =$
340 0.0003) compared to GOA adult male residents. The mean HCB concentration in the SEAK adult male residents
341 was significantly higher than those determined in the GOA ($p = 0.0286$) or WNP ($p = 0.0430$), but no significant

342 difference in mean HCB levels was found between EAI and SEAK adult males ($p = 0.3643$). Significant differences
343 in the mean $\sum\text{DDTs} / \sum\text{PCBs}$ were found among all populations (SEAK and GOA adult males: $p < 0.0001$; EAI and
344 GOA adult males: $p < 0.0001$; SEAK and WNP adult males: $p = 0.0310$; WNP and GOA adult males $p = 0.0227$;
345 SEAK and EAI adult males: $p = 0.0150$) except between WNP and EAI resident killer whales ($p = 0.9285$; Table 2).
346 The EAI resident killer whales had a significantly different mean p,p' -DDE / $\sum\text{DDTs}$ ratio than adult males from
347 GOA ($p = 0.0023$) and SEAK ($p = 0.0128$).

348 *3.2 Assessing Segregated Diet and Foraging areas among Killer Whale Populations*

349 A comparison of OC mixtures in killer whales indicated significant segregations in the six populations
350 analyzed: EAI residents, GOA residents, SEAK residents, WNP residents, EAI transients, and SEAK transients
351 (Figure 8, upper panel; Table 3 $R = 0.451$, $p = 0.001$), which included all adult males, adult females, juveniles and
352 unknowns in each population. Overall, PC1 accounted for 57.4% of the variation and illustrated larger difference in
353 OC mixtures of the transient vs. resident populations but also smaller difference among populations of the same
354 ecotype. Positive loadings for PC1 were highest for HCB, p,p' -DDT, and o,p' -DDT and were more dominant in
355 resident populations whereas negative loading were highest for p,p' -DDE and more dominant in transient
356 populations revealing higher accumulation of p,p' -DDE in the transient populations that feed at a higher trophic
357 level. In contrast, PC2 accounted for 24% of the variation and better illustrated differences in OC mixtures among
358 sex/maturity class. Positive loadings for PC2 were highest for highly lipophilic OCs PCB180 followed by p,p' -DDT,
359 a mixture observed more in adult females, whereas negative loadings of PC2 were highest o,p' -DDT, followed by
360 p,p' -DDE, mixture observed more in adults males, suggesting females are retaining these more lipophilic OCs.

361 Based on an examination of all individuals in each population (i.e., inclusion of all sex/maturity classes),
362 the greatest degree of segregation occurred between SEAK and EAI transients with each of the resident ecotypes
363 (Table 3, R ranges from 0.857 to 0.341 for all but one comparison). Transients (Figure 9, open symbols) generally
364 distributed further to the left on the PC1 axis (i.e., higher proportion p,p' -DDE and lower proportion of HCB, p,p' -
365 DDT, and o,p' -DDT), than residents (Figure 9, solid symbols). All comparisons of OC profiles between transients
366 and resident populations were significant ($p \leq 0.01$) except for the comparison between EAI transients and SEAK
367 residents ($p = 0.15$), which also had considerably less segregation ($R = 0.100$) than all other comparisons of
368 transients and residents ($R \geq 0.341$). For SEAK and EAI transients, the OC profiles were also significantly different
369 from each other, with SEAK transients having higher proportions of p,p' -DDE, but the degree of segregation was

370 less (Table 3, $R = 0.285$, $p = 0.004$). Overall, less segregation in OC mixtures was observed among resident
371 populations, however, the EAI, GOA, and SEAK populations are all significantly different from each other (Table 3
372 ($p < 0.001$ for all comparisons), with the greatest segregation between EAI and GOA ($R = 0.374$, followed by SEAK
373 and GOA ($R = 0.372$), and least between EAI and SEAK ($R = 0.289$). The WNP residents were significantly
374 segregated from the GOA resident killer whale population (Table 3, $R = 0.436$, $p = 0.016$) but not the SEAK and
375 EAI resident populations (Table 3, ($p \geq 0.35$); however, there were only five individuals within WNP population,
376 potentially limiting our ability to discriminate true differences that may exist.

377 Examination of a subset of PCA scores among populations of individuals of the same sex/maturity
378 demonstrate that variation in OC patterns among populations was primarily determined by ecotype, followed by
379 geographic feeding area (Figures 9 and 10) rather than the sex/maturity class of the whales. For example, the PCA
380 scores for female transient and resident killer whales from SEAK (Figure 9, upper panel) illustrate that transient
381 females have a much lower average PC1 score than the resident females (-1.1298 vs. 0.355 , t-test, $p < 0.001$)
382 Likewise, the PCA scores for male transient and resident killer whales from EAI (Figure 9, lower panel) illustrate
383 that transient females have a much lower average PC1 score than resident females (-0.646 vs. 0.844 , t-test, $p <$
384 0.001). Furthermore, examination of the PCA scores among resident males (Figure 10), illustrate significant
385 segregation among populations (ANOVA, $p < 0.001$) with average PC1 score for EAI males (0.844) significantly
386 higher of those from GOA males (0.214) and SEAK males (-0.316), which were not significantly different from
387 each other (Holm Sidak post-hoc test). The WNP resident males also had a significantly higher average PC1 score
388 (0.736) than the SEAK resident males.

389 *3.3 Toxicological risks of OCs to North Pacific killer whales*

390 Of the 69 resident killer whales analyzed in the current study (Supplemental Table 1), 62% (43/69) of these
391 animals had blubber Σ PCB levels that exceeded the threshold for immune suppression and vitamin A depression
392 whereas only 4% (3/69) of these whales (2 adult males from SRKW population and 1 juvenile from SEAK) had
393 blubber PCB concentrations above the ringed seal reproductive dysfunction threshold. All of the transient whales
394 except one adult female from SEAK (Σ PCBs: $58 \mu\text{g g}^{-1}$ lw) had blubber Σ PCB levels that exceeded both of these
395 threshold values. With regard to blubber PCB TEQ concentrations, all transient (29/29) and 57% (39/69) of the
396 resident whales had values that exceeded the 209 pg g^{-1} lw threshold value.

397

398 **4.0 Discussion**

399 *4.1 OC Concentrations and Ratios in North Pacific Killer Whales*

400 Variability in OC concentrations measured in the blubber of marine mammals is largely due to several
401 factors including diet, age, sex or reproductive history, birth order, body composition, and nutritive condition
402 (Borrell, 1993; de Swart et al., 1994; Aguilar et al., 1999; Ross et al., 2000; Ylitalo et al., 2001; Krahn et al., 2007a;
403 Krahn et al., 2009; Mongillo et al., 2016). In the current study, baseline concentrations and profiles of OCs in
404 various populations of North Pacific killer whales were characterized from biopsy blubber samples of 98 wild-
405 ranging resident and transient killer whales sampled from across the North Pacific, ranging from the Russian Far
406 East to Washington State, USA. We hypothesized that populations of North Pacific killer whales would differ in
407 their OC concentrations and profiles based on their ecotype, geographical (foraging) area, and by sex/maturity
408 category. In addition, our study provides information on the levels of Σ PCB TEQs for killer whale populations
409 across this region.

410 Transient killer whales across the expanse of the North Pacific had significantly higher mean blubber
411 concentrations of OCs than those measured in residents in the same sex/maturity categories (See Figures 2 – 5).
412 These findings were expected because previous feeding ecology and observational field studies have shown that
413 North Pacific transient killer whales feed primarily at a higher trophic level (i.e., marine mammals) than residents,
414 whose diet is composed primarily of fish (Ford et al., 1998, 2000; Ford and Ellis, 2006; Saulitis et al., 2000; Herman
415 et al., 2005; Krahn et al., 2007a; Matkin et al., 2007; Dahlheim and White, 2010). Because OCs are generally
416 resistant to metabolism and environmental degradation, and the majority of OC accumulation in adult killer whales
417 is from their diet, we would expect killer whales that feed at higher trophic levels to have a higher exposure to these
418 pollutants. Moreover, our results corroborate findings reported in previous studies in which OC levels in transient
419 killer whales were substantially higher than in resident killer whales (e.g., Ross et al., 2000; Ylitalo et al., 2001;
420 Herman et al., 2005; Krahn et al., 2007a).

421 In both transient and resident killer whales, adult females had lower OC concentrations than that observed
422 in juvenile or adult male killer whales (See Figures 2-5; Supplemental Table 1). Numerous studies have found
423 significant differences in OC concentrations between adult male and adult female killer whales (Ross et al., 2000;
424 Ylitalo et al., 2001; Krahn et al., 2009). This is because reproductive females are able to transfer a substantial

425 amount of their OC body burdens to their calves, especially through lactation (Fukushima and Kawai, 1981; Tanabe
426 et al., 1981; Tanabe et al., 1982; Borrell et al., 1995). Unlike adult whales that receive the majority of OCs from
427 their diet, calves receive the majority of their pollutant load from their mothers through transplacental transfer and
428 nursing (Tanabe et al., 1982; Aguilar and Borrell, 1994; Borrell et al., 1995; Ridgway and Reddy, 1995; Desforges
429 et al., 2012; Cadieux et al., 2016).

430 Examination of OC ratios between ecotypes revealed that transient killer whales had significantly higher
431 $\sum\text{DDTs} / \sum\text{PCBs}$ ratios than resident killer whales (Figure 7). Krahn et al. (2007a) also found higher $\sum\text{DDTs} /$
432 $\sum\text{PCBs}$ ratios in adult male West Coast transients compared to those determined in adult male residents. Borrell
433 (1993) examined the variations in $\sum\text{DDTs} / \sum\text{PCBs}$ ratios in long-finned pilot whales (*Globicephala melas*),
434 Atlantic white-sided dolphins (*Lagenorhynchus acutus*), harbor porpoises, fin whales (*Balaenoptera physalus*), sei
435 whales (*B. borealis*), and sperm whales (*Physeter macrocephalus*) from the northeastern North Atlantic and found
436 an increasing ratio with body size and a decreasing ratio with trophic level. In contrast, Pinzone et al. (2015) found
437 that long-finned pilot whales, fin whales, and sperm whales from the Mediterranean Sea that feed at different trophic
438 levels did not have significant differences in $\sum\text{DDTs} / \sum\text{PCBs}$, further suggesting that other factors may also
439 influence this ratio in cetaceans.

440 It has also been previously suggested that geographical foraging areas influence the $\sum\text{DDTs} / \sum\text{PCBs}$ ratio.
441 For example, among the three Southern Resident killer whale pods (J, K, and L pods), the $\sum\text{DDTs} / \sum\text{PCBs}$ ratios
442 were significantly higher in K and L pods than the ratios determined for the J pod (Krahn et al., 2007b, 2009).
443 Relatively high levels of DDTs are found in the marine environment off California, creating what has been called a
444 “California signature” where $\sum\text{DDTs}$ are high relative to $\sum\text{PCBs}$ (Calambokidis and Barlow, 1991; Jarman et al.,
445 1996; Krahn et al., 2007b). Consequently, salmon stocks that originate from the more urban southern waters
446 generally have higher DDTs than stocks that originate from British Columbia or Alaska (Mongillo et al., 2016), as
447 this OC was primarily used as an insecticide in agricultural regions (EPA, 1972; Eganhouse et al., 2000; Blasius and
448 Goodmanlowe, 2008). Supported by field observations of K and L pods off California waters and no observations
449 of J pod, the authors suggested this difference in the chemical fingerprint was likely due to the pods feeding in
450 spatially distinct areas during certain times of the year. Krahn et al. (2007a) also reported that GOA resident killer
451 whales had statistically lower $\sum\text{DDTs} / \sum\text{PCBs}$ ratios and statistically higher $\sum\text{PBDEs} / \sum\text{PCBs}$ than the EAI or

452 CAI whales. Although the primary driver for the variations in this ratio through the food web is not well known, it
453 is likely that the geographic proximity to regional OC sources is an important driver.

454 Some OC ratios can be used to estimate how recently the pollutant loading into a local ecosystem occurred.
455 For example, *p,p'*-DDE, a stable and persistent isomer that originates from the breakdown of DDT, is more
456 abundant over time and through the food web (Aguilar, 1984; Borrell, 1993) than the parent *p,p'*-DDT. In pinnipeds
457 and odontocetes, it has been estimated that the conversion of DDT to DDE can take years and continue after the
458 DDT exposure has ceased (Aguilar, 1984). A relatively higher ratio of *p,p'*-DDE / \sum DDTs may suggest that DDT
459 has been in the environment for a long time, whereas a relatively smaller ratio may indicate exposure to a 'fresher'
460 source of this OC pesticide. This ratio can also be influenced by the concentration of \sum DDT in the individual
461 (higher body burdens of \sum DDT can result in higher rates of dehydrochlorination and result in higher levels of DDE)
462 (Borrell and Aguilar 1987, Martineau et al., 1987). In the current study, transient killer whales in all sex/maturity
463 categories had significantly higher *p,p'*-DDE / \sum DDTs than resident killer whales in the corresponding class,
464 reflecting its persistence through the food web and may also reflect the intensification of dehydrochlorination given
465 their relatively higher body burdens than resident whales. The mean *p,p'*-DDE / \sum DDTs values measured in
466 resident and transient North Pacific killer whales was above 0.8 and 0.9, respectively (Figure 6). Similarly,
467 McHugh et al. (2007) sampled blubber from killer whales from British and Irish waters and reported that the *p,p'*-
468 DDE / \sum DDTs ratio ranged from 0.7 to 0.96, suggesting a more historical contamination source of DDTs.

469 *4.2 Assessing Segregated Diet and Foraging areas among Killer Whale Populations*

470 Distinct chemical fingerprints, suggesting distinct foraging areas and/or diets, were measured among the
471 four resident and two transient populations (Figure 8), based on the PCA analyses of the relative abundance of five
472 OC analytes: HCB, non-dioxin-like PCB 180, and DDT metabolites *p,p'*-DDE, *p,p'*-DDT, and *o,p'*-DDT. The
473 greatest distinction in fingerprints occurred between transient populations and resident populations, consistent with
474 the differences in OC concentrations and ratios observed for these ecotypes. As discussed above, transient killer
475 whales in all sex/maturity categories had significantly higher mean *p,p'*-DDE / \sum DDTs ratios than resident killer
476 whales in the corresponding sex/majority category, likely associated with persistence of *p,p'*-DDE through the food
477 web. However, we also observed differences in fingerprints of EAI and SEAK transients, likely associated with
478 their foraging areas. The SEAK transients have been well documented (via photographic matches) to frequently
479 occur throughout the coastal waters of British Columbia and Washington State (Dahlheim and White, 2010).

480 Conversely, SEAK transients have not been observed in Prince William Sound (Matkin et al., 1999b), or the Gulf of
481 Alaska, Aleutian Islands, and Bering Sea (Dahlheim, 1997).

482 We also observed significant differences in chemical fingerprints among resident populations, consistent
483 with differences in p,p' -DDE / \sum DDTs among adult males from resident populations. The EAI resident killer
484 whales had a significantly different mean p,p' -DDE / \sum DDTs ratio than adult males from GOA and SEAK. SEAK
485 residents, which are known to inhabit waters off southeast Alaska, had significantly different profiles than EAI and
486 GOA residents (Table 2). It is likely that SEAK resident killer whales consume salmon with a more southern range
487 for part of the year including those from British Columbia, Washington, Oregon and California given these
488 populations are known to consume Chinook salmon; the Chinook salmon in SEAK are dominated by southern
489 stocks; and strong synchronized demographic rates were found between SEAK resident killer whales and SRKWs,
490 which suggests a common environmental driver (Ward et al., 2016). These southern stocks of adult Chinook salmon
491 (e.g. stocks from British Columbia, Washington, Oregon, and California) tend to have higher OC concentrations
492 than Chinook populations from Alaska (Mongillo et al., 2016). Supporting this hypothesis, we found significantly
493 higher OC concentrations in blubber of adult male SEAK resident killer whales than in adult males from the WNP,
494 EAI, and GOA that likely consume different salmon stocks and other fish species. We also found significant
495 differences in mean blubber percent lipid values among these adult resident males based on geographic areas, but
496 the reasons for these differences are not known. It should be noted that the SEAK adult males were sampled
497 approximately four to seven years prior to adult males from the three other geographical areas. The effects of
498 different sampling years on the killer whale blubber OC concentrations, however, are likely to be negligible as the
499 declines in concentrations of PCBs, DDTs, and HCB in marine biota have been estimated to range from 2 – 8% per
500 year since their attaining maximum concentrations in the 1980s (AMAP 2015, West et al., 2017; Bolton et al.,
501 2020). It is more likely that the differences in OC profiles among resident adult male killer whales are largely driven
502 by their segregated feeding locations. Future effort should include additional collection of biopsy samples of these
503 populations to compare and assess temporal trends to this baseline study to confirm if concentrations have continued
504 to decline since the early 2000s.

505 *4.3 Toxicological risks of OCs to North Pacific killer whales*

506 A number of potential health risks have been associated with exposure to high levels of contaminants in
507 top-level predators such as killer whales. For example, Hall et al. (2006a) reported that the risk of infection in

508 harbor porpoise from the United Kingdom doubled when blubber PCB concentrations exceeded 45 $\mu\text{g g}^{-1}$ lipid.
509 Thus, OC exposure may indirectly affect populations of marine mammals by increasing susceptibility to
510 opportunistic pathogens at lower exposure levels than are necessary to observe direct toxicity or death. In addition,
511 high levels of OCs in other marine mammal species have been linked to reproductive impairment, immune
512 suppression, anemia, endocrine disruption, skeletal deformities, and carcinoma (Reijnders, 1986; Subramanian, et
513 al., 1987; de Swart et al., 1995; Ross et al., 1996; Beckmen et al., 2003; Schwacke et al., 2012; Ylitalo et al., 2005b;
514 Desforges et al., 2016).

515 Although the health effects from mixture interactions is largely unknown, neglecting to consider potential
516 interactive effects may underestimate risk to individual killer whales or their population (Mongillo et al., 2016). For
517 example, certain mixtures of pollutants can interact additively or synergistically and enhance toxicity (e.g. Eriksson
518 et al., 2006, Gao et al., 2009), whereas some may have antagonistic interactions and reduce toxicity (e.g. Yordy et
519 al., 2010). Desforges et al. (2017) used contaminant mixtures from the blubber of polar bears and killer whales for
520 in-vitro experiments with immune cells of multiple species and observed lower effect levels from the mixtures
521 relative to the single compounds. Pellacani et al. (2014) investigated the cytotoxic effects of several combinations of
522 persistent pollutants and observed antagonistic, additive, and synergistic interactions from one mixture. Because real
523 world exposures of mixtures can contain a range of interactions, it makes predicting effects extremely difficult.

524 There have been several attempts to model relative impacts of OCs on marine mammal populations. Hall et
525 al. (2006b) estimated the effects of PCB accumulation rates on potential population growth rates in a bottlenose
526 dolphin population and found that the current PCB accumulation rates may be depressing the population growth
527 rate. More recently, Desforges et al. (2018) modeled relative PCB effects on killer whale reproduction and immune
528 function to assess the potential risk on long-term viability and population size. The model predicted the high risk of
529 population crashes for many killer whale populations (e.g., those found in more industrialized urban areas and those
530 that eat at higher trophic levels). Their predictions have since been the center of an active and ongoing debate in the
531 online (eLetter) forum of the journal *Science* about how PCBs could cause population declines.

532 In the current study, transients appear to be at higher risk of health effects compared to residents as their
533 blubber PCBs and PCB TEQs (Supplemental Table 1) exceeded threshold levels established for marine mammals.
534 As previously noted, we found that 97% of transient North Pacific killer whales had blubber $\sum\text{PCBs}$ that exceeded
535 the reproductive dysfunction threshold value established for ringed seals (Boon et al., 1987) while only 4% of the

536 resident whales had blubber concentrations that exceeded this value. Regardless of geographical area, adult males
537 comprised the largest fraction of animals that had blubber PCB and PCB TEQ levels that exceeded immune
538 suppression and vitamin A thresholds for marine mammals, also placing them at higher risk of deleterious effects.
539 Therefore, examining differences in OC concentrations based on ecotype, as well as population and sex/maturity
540 class is important when assessing the relative risk of potential health impacts from OC exposure in North Pacific
541 killer whales.

542 *4.4 Conclusion*

543 North Pacific killer whales are long-lived, high trophic level predators that are exposed to multiple
544 stressors. Some of these populations are in close proximity to human activities and development and thus exposed
545 to relatively high levels of OCs, making them particularly vulnerable to adverse health effects. Our study results
546 indicate high variability in OC concentrations in North Pacific killer whales and that segregation in foraging areas
547 appears to be a primary driver of variability in OC profiles we observed among the populations within the ecotypes.
548 Current concentrations and profiles of organochlorine contaminants in North Pacific resident and transient killer the
549 levels have likely reduced over the 20-year period since samples were collected. The results of this baseline study
550 are crucial for examining time trend changes in the region and assessing the potential health risks associated with
551 OC exposure in this species.

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1 Figures

2 Figure 1. Map of the North Pacific Ocean detailing core foraging areas inhabited by killer whale populations from
3 which biopsy samples were collected. Ninety eight killer whales from the North Pacific were sampled in the
4 following geographical areas: 1) resident whales from Washington State/British Columbia (WA/BC); 2) resident and
5 transient whales from southeast Alaska (SEAK); 3) resident and transient whales from Gulf of Alaska/Prince
6 William Sound (GOA/PWS); 4) resident and transient whales from eastern Aleutian Islands (EAI); 5) resident
7 whales from Central Aleutian Islands (CAI), and resident whales that occur in the western North Pacific (WNP).
8

9 Figure 2. Plot of $\log(\text{HCB lw})$ with respect to sex/maturity class and ecotype. Resident and transient ecotypes are
10 colored blue and red, respectively. Results of analysis of variance (ANOVA) revealed a significant interaction
11 ecotype*sex/maturity class ($F_{2,9} = 4.7$; $p < 0.05$); however, Tukey-Kramer honestly significant difference post-hoc
12 pairwise tests indicated non-significant differences ($p > 0.05$) among sex/maturity classes.
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14 Figure 3: Plot of $\log(\sum \text{DDTs lw})$ with respect to a) geographical area, and b) sex/maturity class and ecotype.
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17 differences in the interaction of ecotype*sex/maturity class. However, significant differences were observed
18 between ecotypes ($F_{1,90} = 182$; $p < 0.001$) and among age and sex classes ($F_{3,95} = 6.9$; $p < 0.01$). Tukey-Kramer
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25 differences in the interaction ecotype*sex/maturity class. Significant differences were observed between ecotypes
26 ($F_{1,90} = 230$; $p < 0.001$) and among sex/maturity classes ($F_{3,95} = 6.9$; $p < 0.01$). Tukey-Kramer honestly significant
27 difference post-hoc pairwise tests indicated non-significant differences ($p > 0.05$) among sex/maturity classes.
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29 Figure 5: Plot of $\log(\sum \text{PCB TEQs lw})$ with respect to a) geographical area and b) sex/maturity class and ecotype.
30 Resident and transient ecotypes are colored blue and red, respectively. Results of analysis of variance (ANOVA)
31 revealed non-significant differences in $\log(\sum \text{PCBs TEQs lw})$ among geographical areas, yet a significant interaction
32 ecotype*sex/maturity class was observed ($F_{2,90} = 3.2$; $p < 0.05$). Tukey-Kramer honestly significant difference post-
33 hoc pairwise tests indicated non-significant differences ($p > 0.05$) for the interaction ecotype*sex/maturity class.
34

35 Figure 6: The ratio of p,p' -DDE/ $\sum \text{DDTs}$ with respect to a) ecotype, b) sex/maturity class, and c) geographical area.
36 ANOVA tests revealed significant differences in the ratio of p,p' -DDE/ $\sum \text{DDTs}$ between ecotypes ($F_{1,95} = 59$; $p <$
37 0.001), among sex/maturity classes ($F_{3,93} = 2.9$; $p < 0.05$), and geographical areas ($F_{5,91} = 7.3$; $p < 0.001$). Tukey-
38 Kramer honestly significant difference post-hoc pairwise tests indicated significant differences ($p < 0.05$) between
39 geographical areas, which are denoted by different letters, but did not detect differences among sex/maturity classes
40 ($p > 0.05$).
41

42 Figure 7: The ratio of $\sum \text{DDTs}/\sum \text{PCBs}$ with respect to a) ecotype, b) sex/maturity class, and c) geographical area.
43 ANOVA tests revealed significant differences in the ratio of $\sum \text{DDTs}/\sum \text{PCBs}$ between ecotypes ($F_{1,95} = 9.8$; $p < 0.01$)
44 and among areas ($F_{5,91} = 7.3$; $p < 0.001$), but not among sex/maturity classes. Tukey-Kramer honestly significant
45 difference post-hoc pairwise tests indicated significant differences ($p < 0.05$) between areas (denoted by different
46 letters).
47

48 Figure 8. Plot of the first two principal components based on the Principal Component Analysis (PCA) of
49 proportions of 5 OCs (HCB, PCB 180, p,p' -DDE, p,p' -DDT, and o,p' -DDT) measured in blubber samples of killer
50 whales populations from the North Pacific Ocean revealing the degree of segregation on OC mixtures among
51 populations (upper panel) and among sex/maturity classes (lower pane). Collectively, both PCAs explain 81.5% of
52 the variation, with PC1 accounting for 57.4%, showing higher proportions of HCB, p,p' -DDT, and o,p' -DDT in
53 resident whale populations higher proportions of p,p' -DDE in transient populations. PC2 accounts for 24% of the
54 variation, with higher proportions of PCB180 in adults females and higher proportion of p,p' -DDE in adult males

55 (upper panel).EAI = eastern Aleutian Islands; GOA = Gulf of Alaska; SEAK =southeast Alaska; and WNP =
56 western North Pacific.

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58 Figure 9. Plot of the PC1 and PC2 scores for resident (solid symbols) and transient (open symbols) populations of
59 SEAK adult females (upper plot) and EAI adult males (lower plot), revealing the segregation in PC1 scores for OC
60 mixtures between resident and transient populations within a geographical (foraging) area is not influenced by
61 sex/maturity of the whales. Plotted scores are of subset scores from the PCA of proportions of 5 OCs measured in
62 blubber samples of killer whale populations from the North Pacific Ocean plotted in Figure 8). SEAK =southeast
63 Alaska EAI = eastern Aleutian Islands, SEAK =southeast Alaska.

64
65 Figure 10. Plot of the PC1 and PC2 scores for adult males from SEAK (circles symbols), GOA (upward triangles
66 symbols), WNP (downward trianglesymbols), and EAI (square symbols) resident populations, suggesting that
67 segregation in PC1 scores for organochlorine (OC) mixtures among resident population is primarily determined by
68 diet and geographical (foraging) area rather than sex/maturity of whales. Plotted scores are a subset of scores from
69 the PCA of proportions of 5 OCs measured in blubber samples of killer whale populations from the North Pacific
70 Ocean plotted in Figure 8. EAI = eastern Aleutian Island; SEAK =southeast Alaska, GOA = Gulf of Alaska, and
71 WPN = Western North Pacific.

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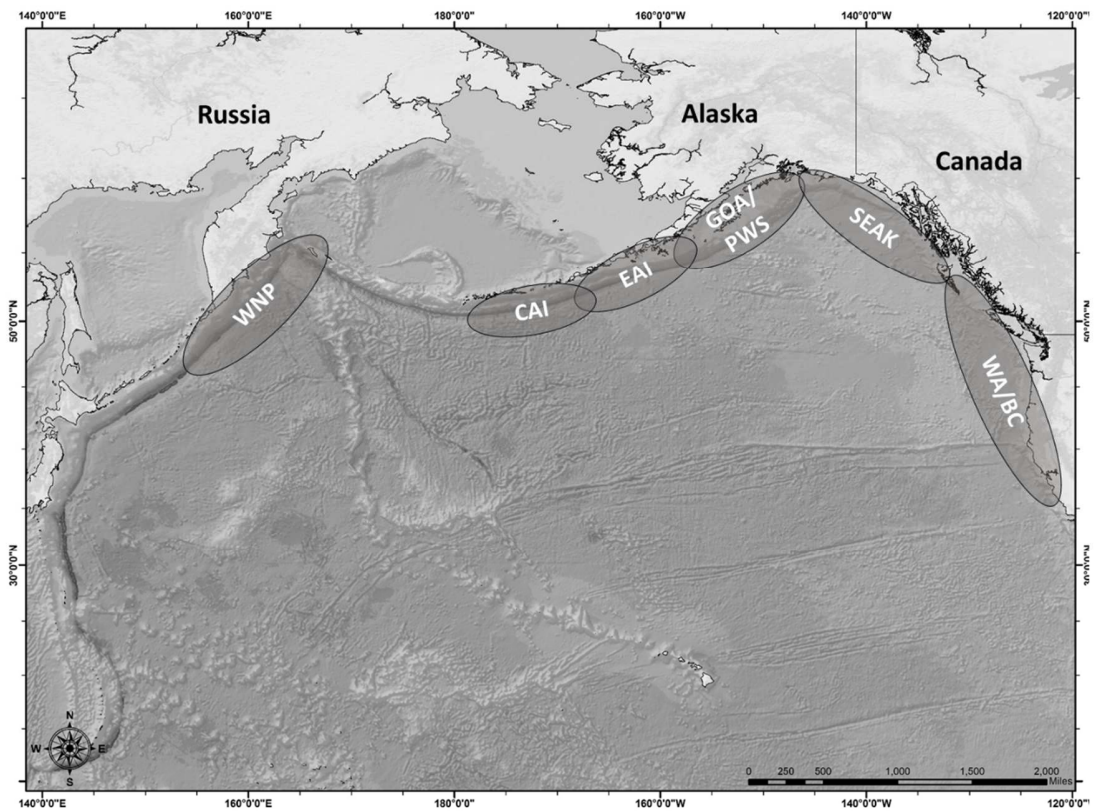


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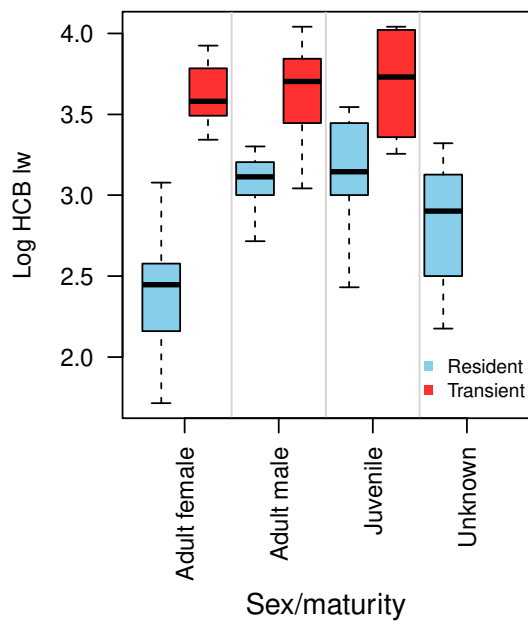


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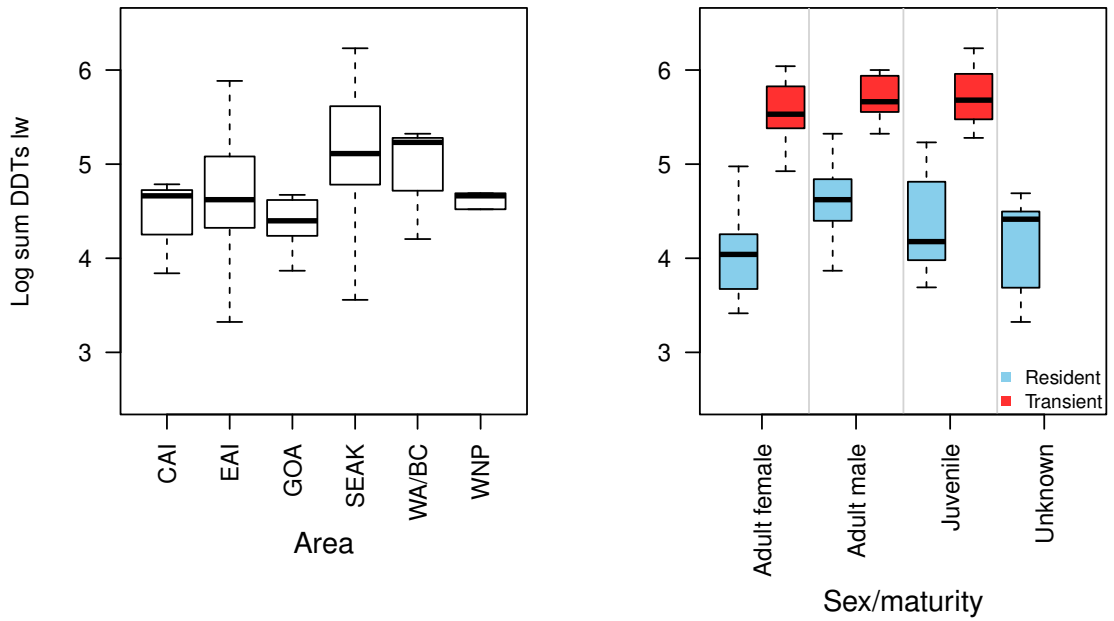


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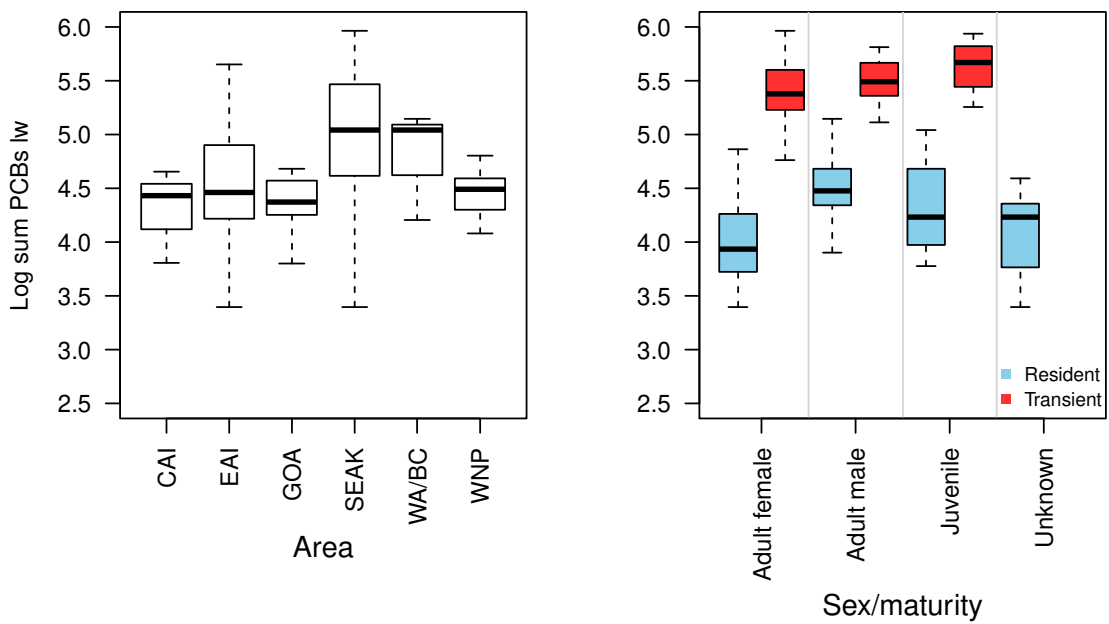


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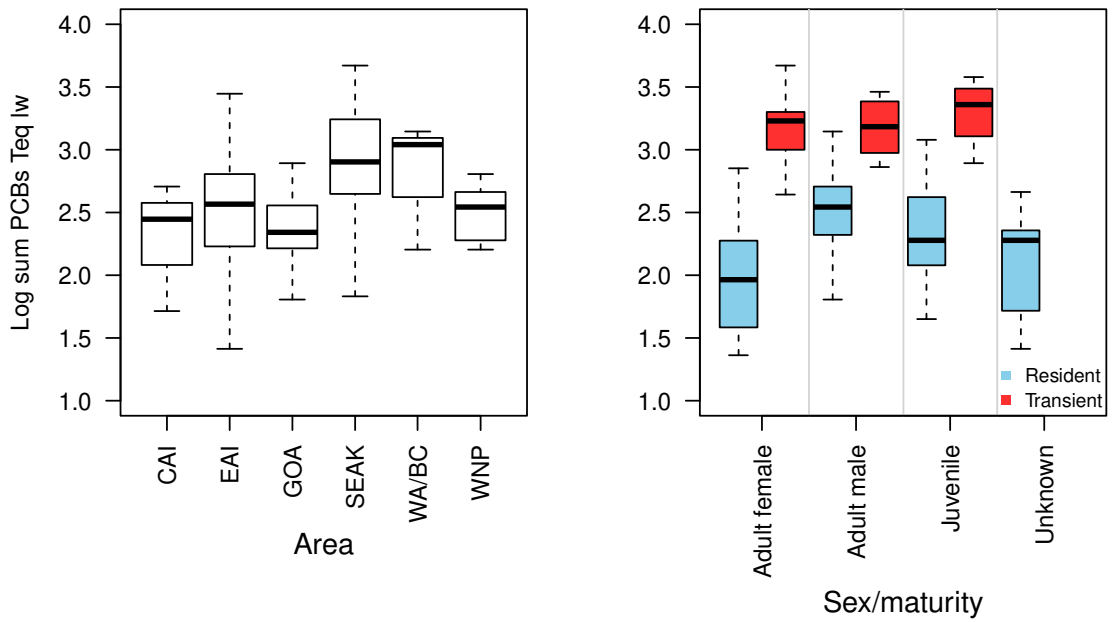


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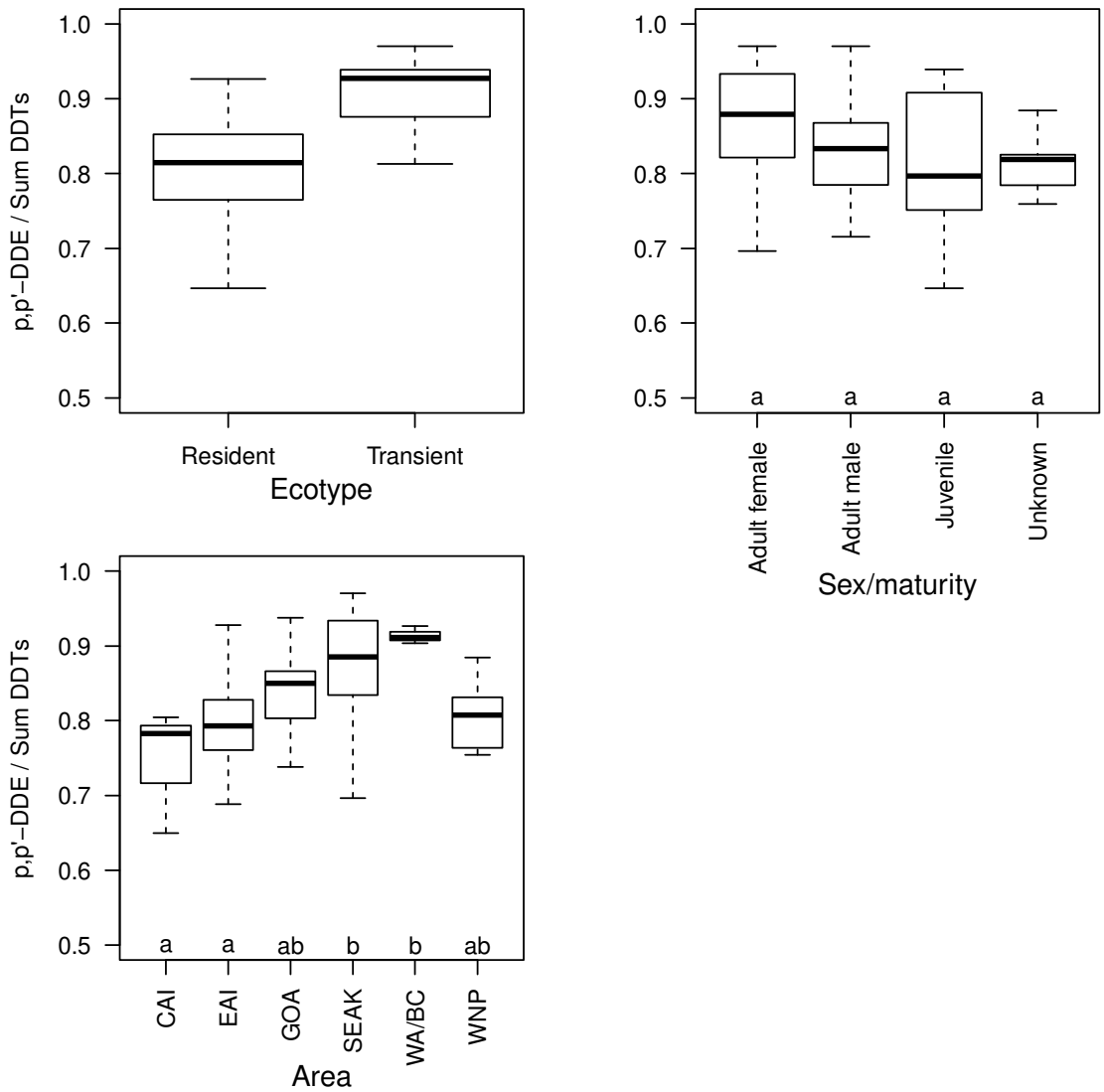


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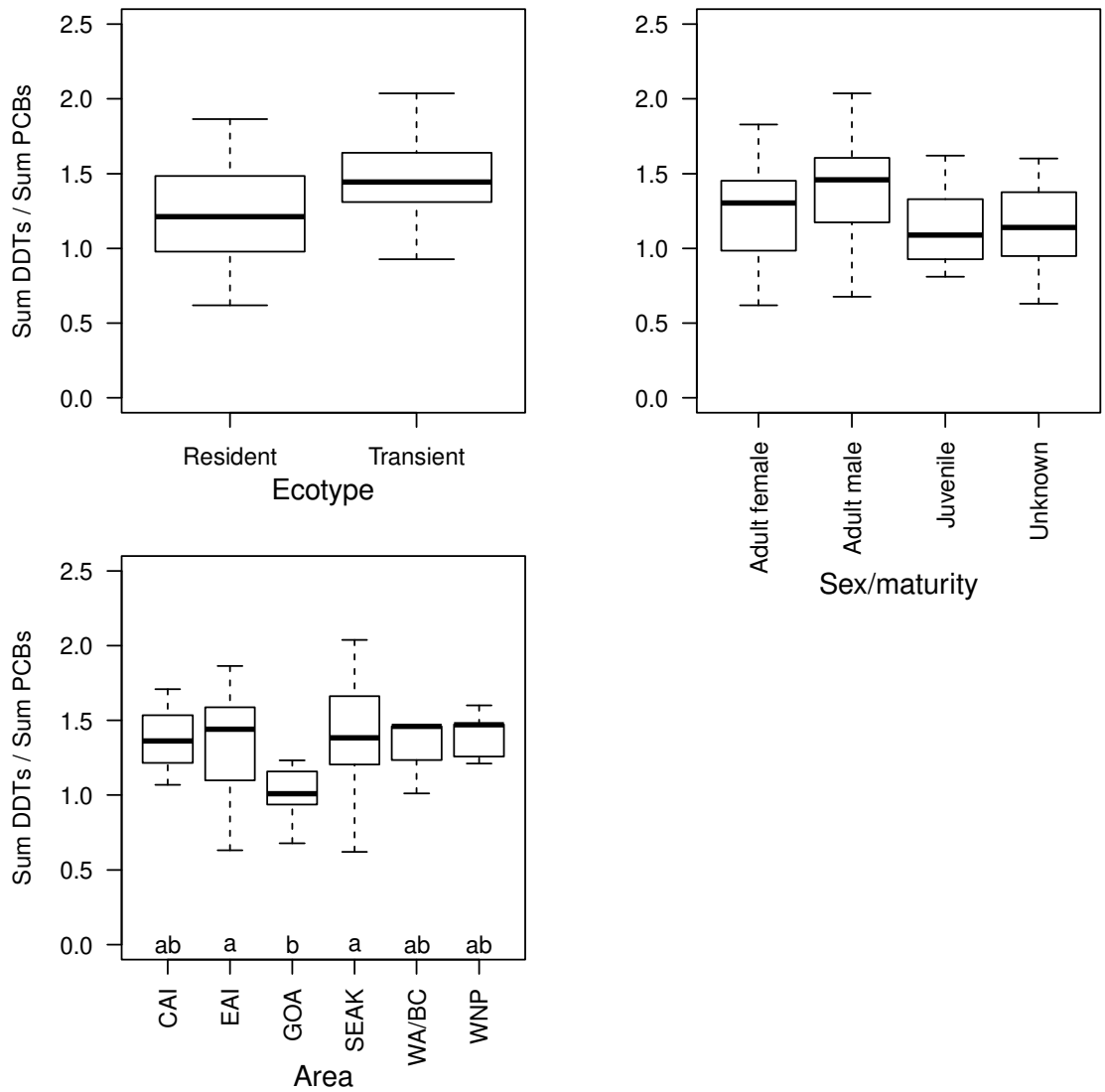


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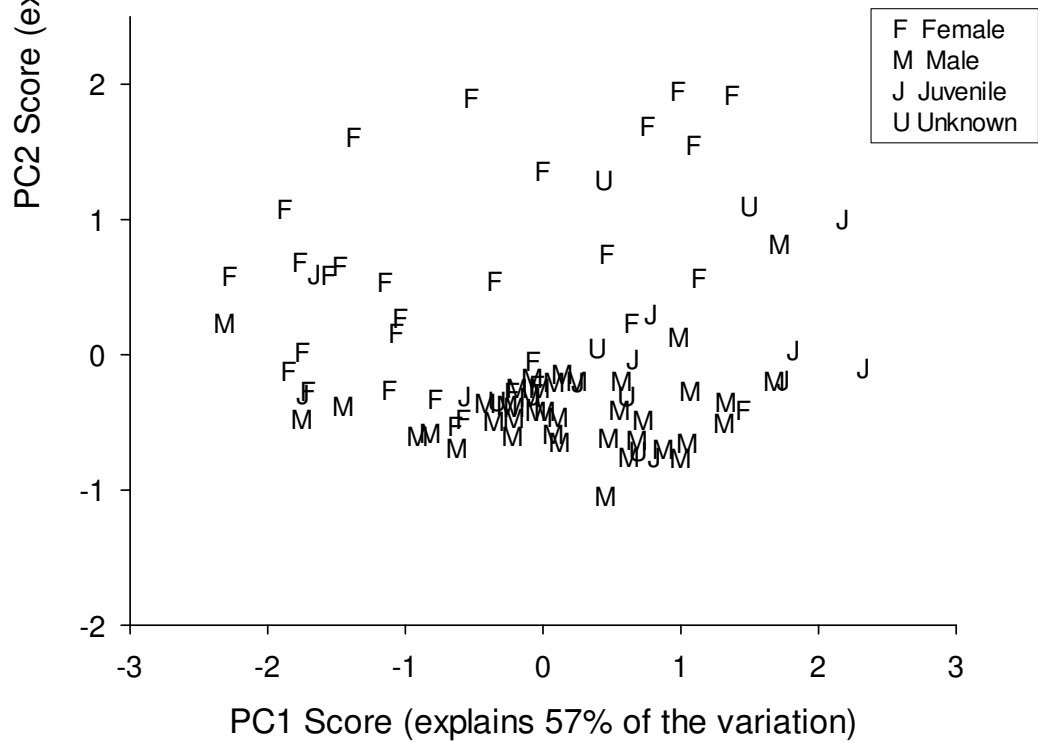
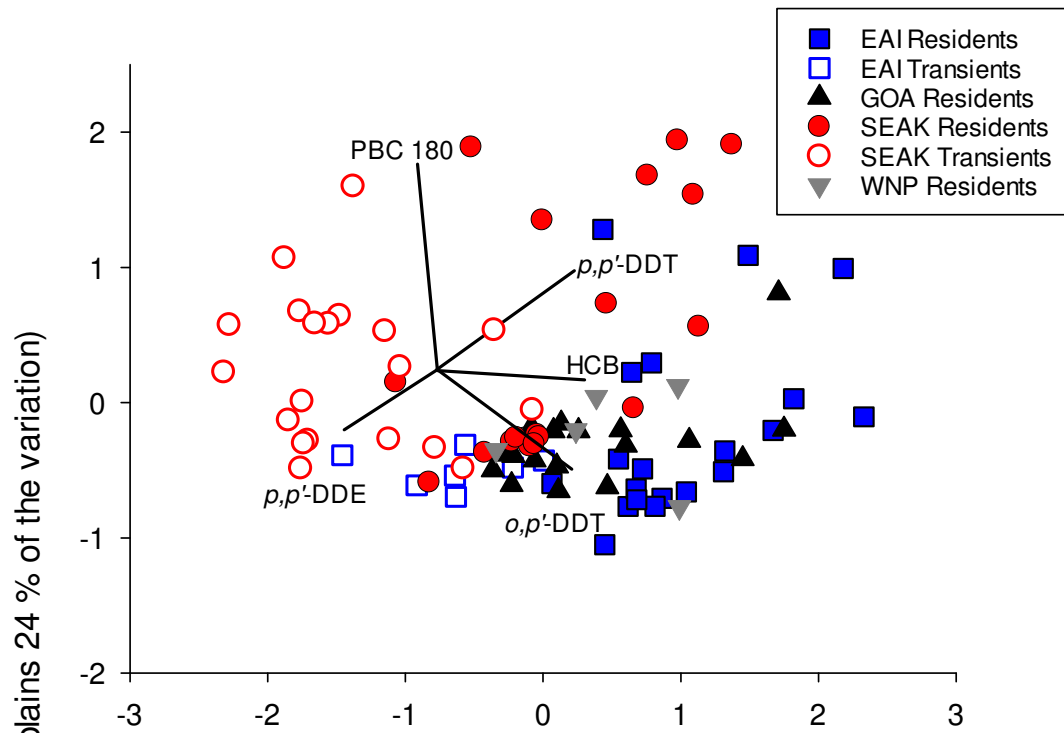


Figure 9. Plot of the PC1 and PC2 scores for resident (solid symbols) and transient (open symbols) populations of SEAK adult females (upper plot) and EAI adult males (lower plot), revealing the segregation in PC1 scores for OC mixtures between resident and transient populations within a geographical (foraging) area is not influenced by sex/maturity of the whales. Plotted scores are of subset scores from the PCA of proportions of 5 OCs measured in blubber samples of killer whale populations from the North Pacific Ocean plotted in Figure 8). SEAK =southeast Alaska EAI = eastern Aleutian Islands, SEAK =southeast Alaska.

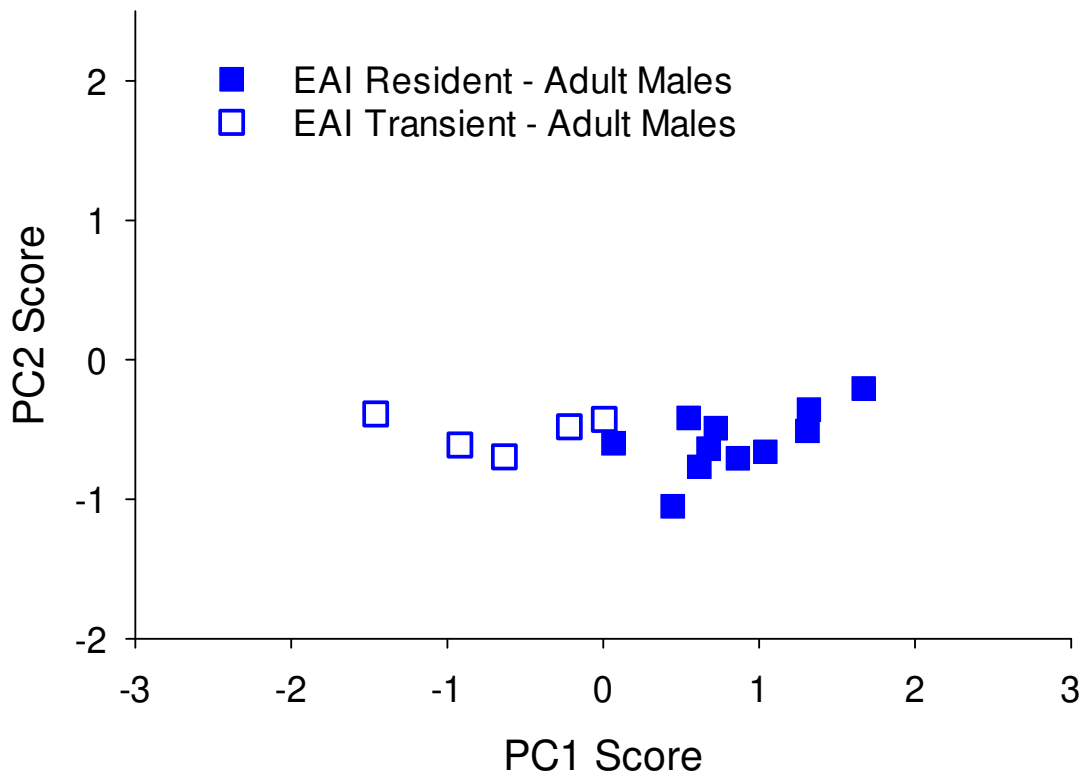
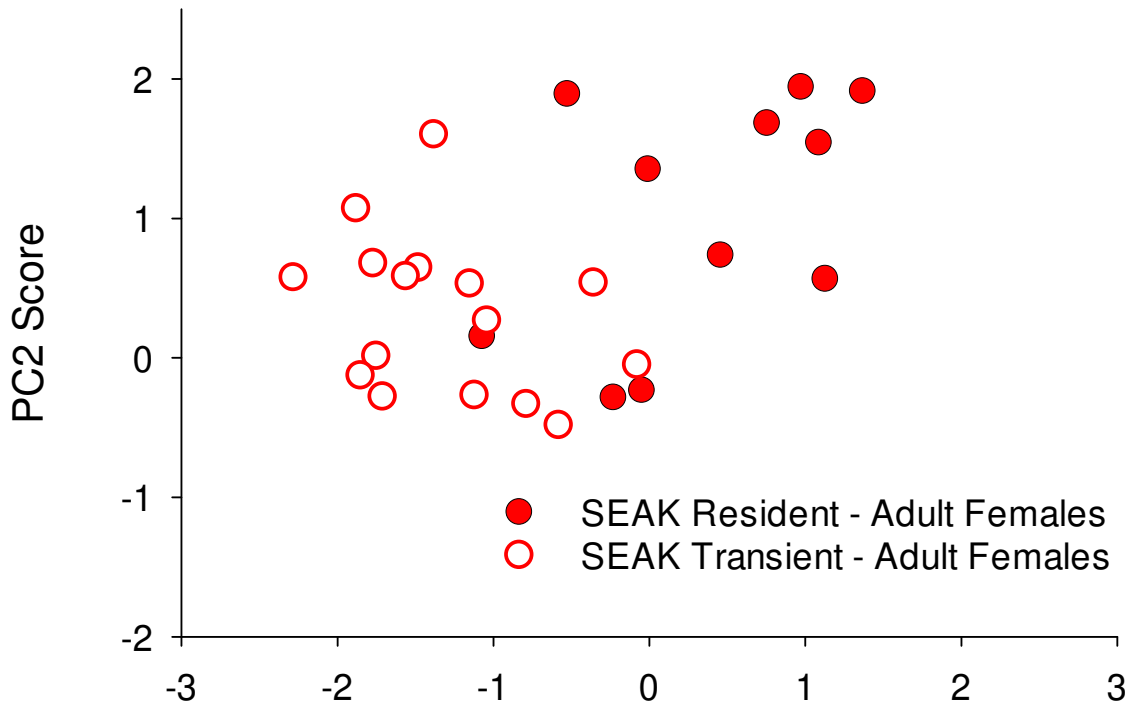
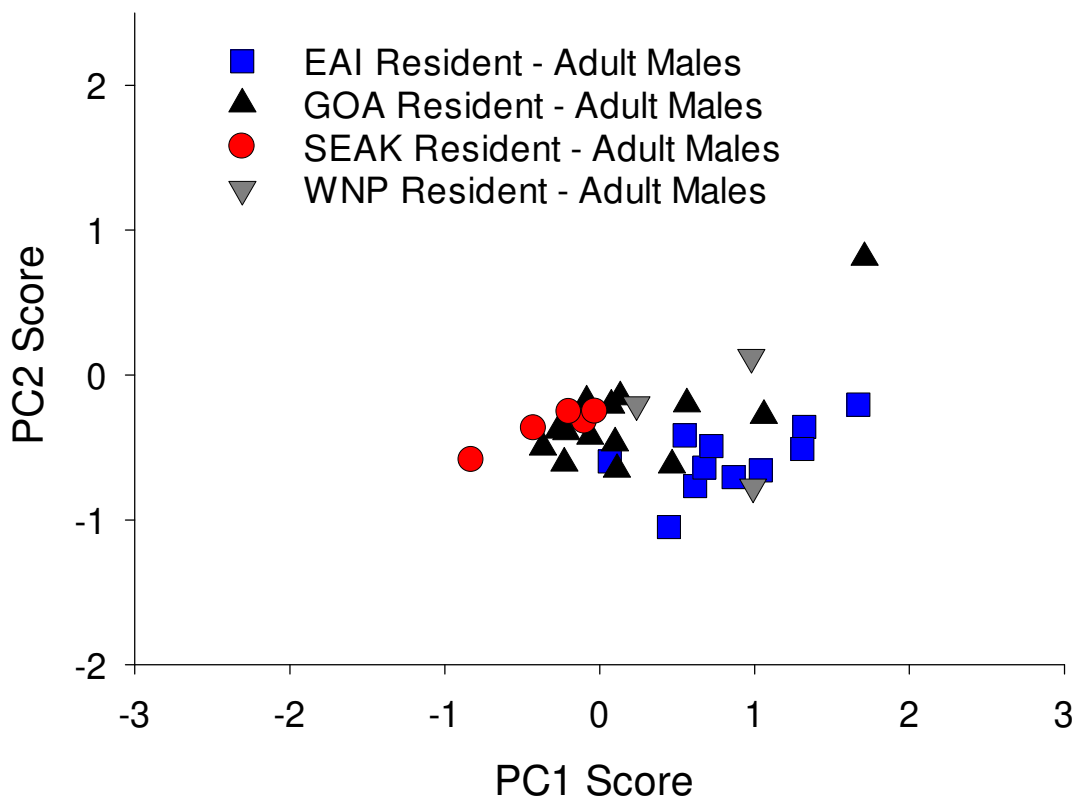


Figure 10. Plot of the PC1 and PC2 scores for adult males from SEAK (circles symbols), GOA (upward triangles symbols), WNP (downward triangles symbols), and EAI (square symbols) resident populations, suggesting that segregation in PC1 scores for organochlorine (OC) mixtures among resident population is primarily determined by diet and geographical (foraging) area rather than sex/maturity of whales. Plotted scores are a subset of scores from the PCA of proportions of 5 OCs measured in blubber samples of killer whale populations from the North Pacific Ocean plotted in Figure 8. EAI = eastern Aleutian Island; SEAK =southeast Alaska, GOA = Gulf of Alaska, and WPN = Western North Pacific.



1 Tables

2 Table 1. Results of the generalized linear modelling approach that assessed the independent variables (ecotype, that
3 explained variability in each of the six dependent variables [$\log(\text{HCB lw})$, $\log(\sum\text{PCBs lw})$, $\log(\sum\text{DDTs lw})$,
4 $\log(\sum\text{PCB TEQs lw})$, ratio p,p' -DDE / $\sum\text{DDTs}$, ratio $\sum\text{DDTs} / \sum\text{PCBs}$]. Relative likelihood (rel.like) is the
5 likelihood of a model given the data, and AIC weight (aic.wt) is the discrete probability of each model. Only models
6 that are indistinguishable (i.e., delta AIC of ≤ 2.0) are displayed.

7
8 Table 2. Mean (\pm SE) and concentration range of organochlorines ($\mu\text{g g}^{-1}$, lipid weight, lw, for HCB, $\sum\text{DDTs}$,
9 $\sum\text{PCBs}$ and pg g^{-1} lw for $\sum\text{PCB TEQs}$), percent lipid, and mean organochlorine ratios in blubber biopsy samples
10 from adult male western North Pacific (WNP), eastern Aleutian Island (EAI), Gulf of Alaska (GOA), and southeast
11 Alaska (SEAK) resident killer whales.

12
13 Table 3. ANOSIM statistical results for pair-wise comparison of organochlorine (OC) patterns among killer whale
14 populations among. R varies between 0 and 1, although small negative values close to zero are possible. R values
15 closer to 1 signify a higher degree of separation. Statistically significant differences are noted with an *. SEAK =
16 Southeast Alaska, EAI = Eastern Aleutian Island, GOA = Gulf of Alaska, WNP = Western North Pacific.

17

Table 1. Results of the generalized linear modelling approach that assessed the independent variables (ecotype, that explained variability in each of the six dependent variables [$\log(\text{HCB } 1w)$, $\log(\sum \text{PCBs } 1w)$, $\log(\sum \text{DDTs } 1w)$, $\log(\sum \text{PCB TEQs } 1w)$, ratio p,p' -DDE / $\sum \text{DDTs}$, ratio $\sum \text{DDTs} / \sum \text{PCBs}$]. Relative likelihood (rel.like) is the likelihood of a model given the data, and AIC weight (aic.wt) is the discrete probability of each model. Only models that are indistinguishable (i.e., delta AIC of ≤ 2.0) are displayed.

Model	AIC	delta.aic	rel.like	aic.wt
Log(HCB)~ecotype*sex/maturity	63.6	0.0	1.0	0.2
Log(HCB)~ecotype+ecotype*sex/maturity	63.6	0.0	1.0	0.2
Log(HCB)~ sex/maturity+ecotype*sex/maturity	63.6	0.0	1.0	0.2
Log(HCB)~ecotype+sex/maturity+ecotype*sex/maturity	63.6	0.0	1.0	0.2
Log(\sumPCBs)~area+ecotype*sex/maturity	66.4	0.0	1.0	0.2
Log(\sum PCBs)~ecotype+area+ecotype*sex/maturity	66.4	0.0	1.0	0.2
Log(\sum PCBs)~area+sex/maturity+ecotype*sex/maturity	66.4	0.0	1.0	0.2
Log(\sum PCBs)~ecotype+area+sex/maturity+ecotype*sex/maturity	66.4	0.0	1.0	0.2
Log(\sum PCBs)~ecotype+area+sex/maturity	67.7	1.3	0.5	0.1
Log(\sumDDTs)~area+ecotype*sex/maturity	97.3	0.0	1.0	0.2
Log(\sum DDTs)~ecotype+area+ecotype*sex/maturity	97.3	0.0	1.0	0.2
Log(\sum DDTs)~area+sex/maturity+ecotype*sex/maturity	97.3	0.0	1.0	0.2
Log(\sum DDTs)~ecotype+area+sex/maturity+ecotype*sex/maturity	97.3	0.0	1.0	0.2
Log(\sum DDTs)~ecotype+area+sex/maturity	98.0	0.7	0.7	0.2
Log(\sumPCB TEQs)~area+ecotype*sex/maturity	79.0	0.0	1.0	0.2
Log(\sum PCB TEQs)~ecotype+area+ecotype*sex/maturity	79.0	0.0	1.0	0.2
Log(\sum PCB TEQs)~area+ sex/maturity+ecotype*sex/maturity	79.0	0.0	1.0	0.2
Log(\sum PCB TEQs)~ecotype+area+ sex/maturity+ecotype* sex/maturity	79.0	0.0	1.0	0.2
Ratio p,p'-DDE/\sumDDTs~ecotype+area+sex/maturity	-303.3	0.0	1.0	0.5
Ratio \sumDDTs / \sumPCBs ~ ecotype+area+sex/maturity	22.2	0.0	1.0	0.5

Best model for each dependent variable shown in bolded text.

Table 2. Mean (\pm SE) and concentration range of organochlorines ($\mu\text{g g}^{-1}$, lipid weight, lw, for HCB, Σ DDTs, Σ PCBs and pg g^{-1} lw for Σ PCB TEQs), percent lipid, and mean organochlorine ratios in blubber biopsy samples from adult male western North Pacific (WNP), eastern Aleutian Island (EAI), Gulf of Alaska (GOA), and southeast Alaska (SEAK) resident killer whales.

Population	<i>n</i>	Percent lipid	$\mu\text{g g}^{-1}$, lw			pg g^{-1} , lw	OC Ratios (\pm SE)	
			HCB	Σ DDTs	Σ PCBs	Σ PCB TEQs	Σ DDTs/ Σ PCBs	<i>p,p'</i> - DDE/ Σ DDTs
WNP resident (range) Significance ¹	3	6.3 \pm 1.1 (5.0 - 8.6) B	1.1 \pm 0.30 (0.56 - 1.6) B	52 \pm 23 (15 - 94) A, B	36 \pm 15 (12 - 64) A, B	380 \pm 140 (160 - 640) A, B	1.385 \pm 0.087 B	0.775 \pm 0.016 A, B
EAI resident (range) Significance ¹	11	19 \pm 2.2 (6.7 - 31) A	1.4 \pm 0.11 (0.87 - 2.0) A, B	44 \pm 5.5 (21 - 69) B	30 \pm 3.6 (16 - 49) B	330 \pm 38 (140 - 550) B	1.455 \pm 0.070 B	0.778 \pm 0.011 B
GOA resident (range) Significance ¹	14	19 \pm 2.2 (6.0 - 38) A	1.2 \pm 0.09 (0.52 - 1.7) B	27 \pm 3.7 (4.3 - 47) B	25 \pm 3.3 (6.3 - 48) B	250 \pm 35 (64 - 500) B	1.040 \pm 0.041 C	0.838 \pm 0.011 A
SEAK resident (range) Significance ¹	5	9.5 \pm 1.2 (7.0 - 13) A, B	1.8 \pm 0.09 (1.6 - 2.0) A	120 \pm 5.2 (100 - 130) A	67 \pm 2.8 (60 - 71) A	770 \pm 54 (620 - 950) A	1.763 \pm 0.018 A	0.845 \pm 0.012 A

¹ Unlike letters indicate significant differences, Tukey-Kramer Honestly Significant Difference (HSD) test, $p < 0.05$.

Table 3. ANOSIM statistical results for pair-wise comparison of organochlorine (OC) patterns among killer whale populations among. R varies between 0 and 1, although small negative values close to zero are possible. R values closer to 1 signify a higher degree of separation. Statistically significant differences are noted with an *. SEAK = Southeast Alaska, EAI = Eastern Aleutian Island, GOA = Gulf of Alaska, WNP = Western North Pacific.

Populations	R	p	
SEAK Transient vs. EAI Resident	0.857	0.001	*
SEAK Transient vs. WNP Resident	0.737	0.001	*
SEAK Transient vs. GOA Resident	0.687	0.001	*
SEAK Transient vs. SEAK Resident	0.486	0.001	*
EAI Transient vs. WNP Resident	0.563	0.004	*
EAI Transient vs. EAI Resident	0.470	0.002	*
EAI Transient vs. GOA Resident	0.341	0.013	*
EAI Transient vs. SEAK Resident	0.100	0.154	
EAI Transient vs. SEAK Transient	0.285	0.004	*
GOA Resident vs. WNP Resident	0.436	0.016	*
EAI Resident vs. GOA Resident	0.374	0.001	*
SEAK Resident vs. GOA Resident	0.372	0.001	*
EAI Resident vs. SEAK Resident	0.289	0.001	*
EAI Resident vs. WNP Resident	0.037	0.350	
SEAK Resident vs. WNP Resident	-0.062	0.631	

