Pronounced long-term trends in year-round diet composition of the European shag
 Phalacrocorax aristotelis

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13 Abstract (Word Count: 249)

Populations of marine top predators are exhibiting pronounced demographic changes due to 14 15 alterations in prey availability and quality. Changes in diet composition is a key potential 16 mechanism whereby alterations in prey availability can affect predator demography. Studies 17 of long-term trends in diet have focused on the breeding season. However, long-term changes 18 in non-breeding season diet is an important knowledge gap, since this is generally the most 19 critical period of the year for the demography of marine top predators. In this study, we analysed 495,239 otoliths from 5888 regurgitated pellets collected throughout the annual 20 cycle over three decades (1985–2014) from European shags Phalacrocorax aristotelis on the 21 22 Isle of May, Scotland (56°11'N, 02°33'W). We identified dramatic reductions in the frequency 23 of lesser sandeel Ammodytes marinus occurrence over the study, which was more pronounced during the non-breeding period (96% in 1988 to 45% in 2014), than the breeding 24 25 period (91% to 67%). The relative numerical abundance of sandeel per pellet also reduced 26 markedly (100% to 13%), with similar trends apparent during breeding and non-breeding 27 periods. In contrast, the frequencies of Gadidae, Cottidae, Pleuronectidae and Gobiidae all 28 increased, resulting in a doubling in annual prey richness from 6 prey types per year in 1988 29 to 12 in 2014. Our study demonstrates that the declining importance of the previously most prominent prey and marked increase in diet diversity is apparent throughout the annual cycle, 30 31 suggesting that substantial temporal changes in prey populations have occurred, which may 32 have important implications for seabird population dynamics.

33 Key words: seabird; long-term study; North Sea; winter diet; non-breeding diet; pellet
 34 analysis; lesser sandeel *Ammodytes marinus*.

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35 Introduction

Marine environments are changing rapidly across the globe due to a range of anthropogenic 36 37 activities, including pollution, overfishing and climate change (Halpern 2009, Poloczanska et 38 al. 2013). These effects have altered the abundance and distribution of lower trophic 39 organisms such as plankton, with consequences for mid-trophic level fish which are the 40 principal prey for a guild of marine top predators (Cury et al. 2000, Frederiksen et al. 2006). 41 Many marine top predator populations are declining markedly in association with these 42 changes in prey availability and quality (Paleczny et al. 2015, Sydeman et al. 2015). Altered 43 diet composition is a key potential mechanism whereby changes in prey availability can affect 44 marine top predators (Reid & Croxall 2001, Cury et al. 2011). Several studies have 45 demonstrated long-term changes in marine top predator diet, in particular seabirds (Miller & Sydeman 2004, Gaston & Elliott 2014). However, these studies have mainly been undertaken 46 during restricted periods of the annual cycle, because of logistical challenges of obtaining diet 47 48 data throughout the year. In seabirds, diet studies are usually conducted during the breeding season, from samples delivered by adults to offspring (Barrett et al. 2007). However, the non-49 50 breeding period is critically important for the population dynamics of seabirds, since most mortality occurs at this time (Weimerskirch 2002, Frederiksen et al. 2008). Thus, a key 51 question in understanding the link between changes in prey availability and seabird 52 53 population dynamics is the extent to which there have been long-term changes in non-54 breeding season diet, and whether these differ from those during the breeding season.

55 Our understanding of seabird diet outside the breeding period is largely based on indirect 56 methods such as stable isotopes and fatty acid analysis (Owen et al. 2013, Kowalczyk et al. 57 2014) or samples from shot/dead birds (Blake 1984, Harris et al. 2015). Such studies have 58 produced valuable insights into non-breeding diet, demonstrating marked differences from

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59 the breeding season, owing to a combination of altered prey availability (Kowalczyk et al. 60 2015), energetic constraints (Markones et al. 2010), habitat association (Ainley et al. 1996) and, in migratory species, altered locations (Ronconi et al. 2010). However, there is very 61 62 limited information on long-term changes in non-breeding diet. Green et al. (2015) examined 63 differences in breeding and non-breeding season diet in Cape gannets Morus capensis over a 64 thirty-year period. However, due to sporadic sampling, their trends analysis was restricted to the breeding period only. To our knowledge, no published studies have quantified long-term 65 66 trends in non-breeding season diet composition in seabirds, and compared these with trends in breeding season diet from the same population. 67

68 In this paper, we analysed three decades of year-round diet in the European shag Phalacrocorax aristotelis (hereafter shag) collected on the Isle of May, south-east Scotland. 69 The shag is a coastally distributed seabird that spends a large proportion of the day and every 70 night on land (Wanless & Harris 1997). Full-grown shags regularly regurgitate pellets 71 72 containing prey remains, which can be collected at accessible roosts, offering a rare 73 opportunity to quantify year-round diet (Barrett et al. 2007). Shags show a flexible foraging 74 strategy such that diet varies substantially across the species range. Lesser sandeel Ammodytes marinus (hereafter sandeel) is the dominant prey in many populations (Harris & 75 76 Wanless 1993, Velando & Freire 1999, Lilliendahl & Solmundsson 2006), but at others, 77 Gadoids (Gadidae), in particular saithe *Pollachius virens*, are the principal prey (Swann et al. 78 2008, Lorentsen et al. 2018). Seasonal variation in diet composition has been recorded in some populations in response to changes in prey availability (Velando & Freire 1999, 79 80 Lilliendahl & Solmundsson 2006). Previous studies of the Isle of May population demonstrated that, in the late 1980s and early 1990s, the diet of shags consisted mainly of 81 82 sandeels, with limited evidence of seasonal differences in diet composition (Harris and

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Wanless 1991, 1993). However, the North Sea has warmed substantially over the past three 83 84 decades (Høyer & Karagali 2016), which has resulted in changes in the distribution, abundance and diversity of many fish populations, including sandeel (Perry et al. 2005, Deurs, 85 Van et al. 2009, ter Hofstede et al. 2010). A recent analysis of diet based on prey delivered to 86 87 shag chicks on the Isle of May demonstrated a marked decline in the proportion of sandeel, from 0.99 (1985) to 0.51 (2014), over this period, along with a concurrent dietary 88 diversification (Howells et al. 2017). The authors attributed this dietary change to climate-89 90 mediated alterations in the availability of sandeels and alternative prey. Similarly, a 91 community-scale analysis of seabird breeding diet at this colony demonstrated a decline in 92 the importance of sandeels over the past three decades (Wanless et al. 2018). As local sandeel 93 populations are resident (Boulcott et al. 2007), it is probable that any effect of environmental 94 change on abundance or quality of these populations will affect both breeding and non-95 breeding diet of shags which over-winter on the Isle of May. Thus, we might predict a decline 96 in the importance of sandeel in the diet throughout the annual cycle. However, sandeel 97 availability varies among seasons since they are present in the water column during the spring and summer, but are buried in the sand during the winter, apart from a brief period when 98 99 they emerge to spawn (Wright & Bailey 1993). Furthermore, environmental conditions, 100 habitat use and energetic costs also vary between seasons (Daunt et al. 2014, Michelot et al. 2017). Thus, any changes in overall prey abundance or availability during the study might have 101 different effects on diet composition at different times of the year. However, whether long-102 103 term changes in diet composition outside the breeding season has matched trends observed 104 in diet during the breeding season (Howells et al 2017) is untested. Therefore, our specific 105 aims were to: a) quantify year round diet composition of shags over three decades; and b) 106 test whether dietary trends differ between the non-breeding and breeding period.

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107 Methods

108 Quantifying diet

109 The study was conducted between 1985 and 2014 at a European shag Phalacrocorax 110 aristotelis (hereafter shag) colony on the Isle of May National Nature Reserve, Firth of Forth, south-east Scotland (56°11'N, 02°33'W). Shags are present on the island throughout the year, 111 112 with a resident proportion of the breeding population joined in winter by migrants from other 113 locations (Grist et al. 2014), allowing for the collection of pellets throughout the year. Pellets 114 were collected opportunistically (mean number of sample days year⁻¹ ± SD: 23 ± 14; range: 3– 115 49) at roosts and breeding colonies using forceps, placed into a plastic bag and frozen. The breeding status and age of individuals that produced pellets was unknown. However, as chicks 116 117 do not produce pellets, all samples were from full-grown (i.e. fledged) birds (Russell et al. 118 1995).

119 Samples were submerged in a saturated solution of biological washing powder (Biotex[©]) and 120 heated at 40–50°C, until all soft tissue and mucus was digested. Residual hard parts (e.g. fish 121 otoliths, vertebrae and mouth parts, cephalapod beaks, mollusc shells and crustacea 122 exoskeletons) were then identified to the lowest possible taxon using keys in Härkönen (1986) 123 and Watt et al. (1997), allowing the presence/absence of each prey type to be recorded in 124 each pellet. Sandeels Ammodytes spp. (principally, lesser sandeels A. marinus; Harris and 125 Wanless 1991), the most frequent prey type recorded, have previously been classified in 126 dietary studies on the basis of age (Harris and Wanless 1991, Howells et al. 2017). However, 127 differentiating between sandeel age classes is generally not possible from otoliths obtained from pellets due to the effect of digestive erosion on otolith structure. Therefore, for the 128 129 purposes of this study, all sandeels were aggregated into a single prey category. The presence

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of sand was also noted, since it may arise from accidental ingestion when foraging in sandy habitats and therefore be an index of prey species that live in these habitats, notably sandeels (Winslade 1974, Holland et al. 2005). The number of otoliths of each prey type in each pellet was then counted. Each fish has two otoliths, but due to the large numbers that may be encountered in a pellet and the potential for otoliths within a pair to undergo differential digestion, it was not possible to accurately match otoliths from the same fish. Therefore, each otolith was treated as an individual sample within each pellet.

137 Pellet analysis has been used to quantify diet in a range of seabirds, including shags, 138 cormorants, skuas and terns (reviewed in Barrett et al. 2007). In appropriate study systems, 139 large sample sizes may be obtained in a non-intrusive way throughout the year. However, 140 quantifying diet from pellets involves two well-established limitations that must be 141 considered when interpreting the data. First, due to differential rates of erosion, small or soft prey may be completely absent or under-represented in pellets, with larger prey, or those 142 143 with more resilient body parts, more commonly retained (Barrett et al. 2007). For example, 144 Johnstone et al. (1990) showed that in captive shags the recovery of otoliths from Sprat 145 Sprattus sprattus, sandeel and Cod Gadus morhua was 17%, 20% and 52%, respectively. Accordingly, the most robust diet metric used to quantify prey in pellets is frequency of 146 147 occurrence, in which items are scored on the basis of presence or absence. This method does 148 not capture prey types that are completely digested, but accounts for any differential in 149 digestion rates among prey types that are recorded by giving equal weighting to prey types 150 irrespective of abundance in the sample. We also considered a second diet measure that is 151 typically quantified from pellets, the numerical abundance of different prey types. This 152 measure is more informative, but must be interpreted with care because it is more sensitive 153 to the effects of differential digestion rates (Barrett et al. 2007).

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A second limitation of quantifying diet from pellets is that the exact date when the prey were ingested is not known. However, the vast majority of pellets were fresh when collected, and they do not persist on rocks at our study colony because they disintegrate in rain or are consumed by herring gulls *Larus argentatus*, so we consider that pellets will have been produced within ca. two weeks of the sampling date.

159 Dietary response variables

For each pellet, we recorded the presence or absence of diagnostic remains (e.g. fish otolith, 160 vertebra, bone, mollusc shell, cephalopod beak) of each prey type. Frequency of occurrence 161 162 was then calculated as the percentage of pellets in which the prey type was found in each 163 period within each study year. We focused our analysis on frequency of occurrence of the top five most abundant fish prey: sandeel Ammodytes spp., Gadidae (Cod Fishes), Cottidae 164 (Cottids), Pleuronectidae (Flatfish) and Gobiidae (Gobies). All other prey types occurred in ≤ 165 166 10% of pellets and could thus not be analysed robustly, but due to their low prevalence in the 167 diet, we consider the omission of these prey unlikely to significantly affect our interpretation 168 of changes in diet composition.

Numerical abundance is typically quantified as the proportion of otoliths of a given fish prey type relative to all otoliths in the pellet. However, where the diet is dominated by a small number of prey types, as in this study (Sandeel 88% and Gadidae 7% of all otoliths), analysis of relative proportions leads to problems of interpretation, since a change in one prey type cannot be readily distinguished from a reciprocal change in the other. We therefore modelled number of sandeel otoliths relative to all prey otoliths and number of Gadidae otoliths relative to all non-sandeel prey otoliths. All other individual prey types occurred too infrequently for

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their relative abundance to be analysed. However, their summed contribution was < 5% of allotoliths.

178 Diet diversity was quantified by calculating sample-level prey richness, which was the number 179 of prey types recorded in each pellet. Due to the effects of digestion on prey items, it was not 180 generally possible to identify all body parts to species level, but to a higher taxonomic level 181 which varied with prey type (fish: family; Crustacea and Mollusca: subphylum; Polychaeta: 182 class). As prey richness is a count, the aggregate, annual prey richness (pooling all pellets in 183 each year) was systematically higher than the sample average (sample-level prey richness: median: 5; range: 0–9; annual prey richness: median: 12; range: 6–14). However, as annual 184 185 prey richness is a measure of the total number of prey types exploited each year, we included 186 it in our analysis.

187 Defining breeding and non-breeding periods

For the purpose of this study, a study year commenced at the onset of breeding in one 188 189 calendar year and ended at the commencement of breeding in the subsequent calendar year. 190 To determine the timing of onset of breeding in each study year we calculated the month in 191 which the population median egg laying date occurred, estimated from weekly observations 192 at long-term monitoring plots (1985–2014: median day of year: range: 101–181; Newell et al. 193 2015; updated). In shags, average incubation duration of a clutch of three eggs, the modal 194 clutch size in this population, is 36 days (Potts et al. 1980), with fledging occurring at a mean 195 of 53 days after hatching (range: 48–58, n = 35; Potts et al. 1980). Therefore, we defined each 196 breeding season as the month of median egg laying date plus the following three months. 197 This four month period was longer than the breeding period of individual pairs (~3 months), but was designed to capture the spread of laying that occurs in each year (Daunt et al. 2007). 198

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We found that 97% of all observations of breeding activity (defined as observations of incubating eggs or brooding chicks; n = 29,075) at our long-term monitoring plots occurred in this four month time window, confirming that it was a robust representation of the breeding period. The non-breeding period commenced in the first month after the breeding period until the last month before the month of median laying date in the following year (range of months: breeding: April–September; non-breeding: August–May; Supplementary Material Table S1).

206 Statistical analysis

207 All statistical analyses were conducted using the R programming software (version 3.4.0, R 208 Development Core Team 2016). To test for temporal trends and effects of period (breeding 209 vs non-breeding) on sample-level presence, relative numerical abundance and prey richness, we fitted Generalised Linear Mixed Models (hereafter GLMMs), using the 'glmer' function in 210 211 the 'lme4' package (Bates et al. 2015). Binomial models with a logit-link function were fitted 212 for presence and relative numerical abundance, and Poisson models with a log-link function 213 for sample-level prey richness. For each of the sample-level dietary components we fitted a 214 global model containing fixed effects of year, period and a year by period interaction. This 215 framework allowed us to test for temporal trends, the differences between periods, and differing temporal trends between breeding and non-breeding periods in each of the dietary 216 217 components. Within each model, we also included random effects for month, year and month 218 nested within year, to account for residual temporal autocorrelation. To account for 219 overdispersion, we also included an individual, sample-level random effect in models of 220 sandeel otoliths relative to all prey and Gadidae relative to non-sandeel prey (Harrison 2015). 221 We did not consider sample date as an explanatory variable, since this variable had no clear

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biological relevance, due to the variable time elapsed between pellet production andcollection.

To identify trends in annual prey richness, where there was just a single value per year, we 224 fitted a Poisson GLMM with a log-link function. We subtracted 6 (the minimum annual prey 225 226 richness value over the study) from each value, so that the data are consistent with the 227 distributional properties of the Poisson distribution. However, we present the results and 228 plots on the original, unadjusted scale. This step was not necessary with the sample-level prey 229 richness data, as the minimum value was zero i.e. pellets where no species were identified. 230 Visual inspection indicated that the annual prey richness may be exhibiting non-linear trends. 231 To test this, a global model containing both a linear and quadratic numeric fixed effect of year 232 was fitted, along with a categorical, annual level random effect of year to account for 233 overdispersion (Harrison 2015). We weighted each annual prey richness value by the number of pellets per year and included a fixed (offset) effect of log(number of pellets year⁻¹) to 234 235 account for any systematic change in annual prey richness with annual sample size.

236 In order to compare models with different fixed effects but the same random structure we 237 used maximum likelihood in all models (Zuur et al. 2009). In each analysis, the fixed effect of 238 year was centred on zero (by subtracting mean year from each value) and rescaled (by dividing the centred value by the standard deviation of year). The inclusion of all years in the analysis 239 240 led to difficulties with model convergence. Preliminary analyses confirmed that this was 241 caused by the inclusion of years where samples were not collected in both the breeding and 242 non-breeding periods, so these were excluded from the modelling process (707 samples in 7 243 years; 1985-87, 1994, 1998-99, 2008).

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Model selection was performed on the four models (null, year, period, and year by period 244 245 interaction) for each variable using Akaike's Information Criterion corrected for small sample 246 sizes (AICc), where the best-supported model was considered to have the lowest AICc value 247 compared to alternative models. Models within two AICc (Δ AICc < 2) of the top model were 248 deemed as having similar levels of support (Burnham & Anderson 2002), unless they 249 contained an additional parameter, in which case they were considered uninformative (Arnold 2010). Analysis was conducted according to an established protocol (Zuur et al. 2010), 250 251 with the 'MuMIn' (Bartoń 2016) package used to obtain model selection outputs (see 252 Supplementary Material for full details of model selection). Due to the large number of models, we only report those within 10 AICc points of the best model in the main text. 253

254 For figures and tables, annual means were calculated by pooling all samples in each period 255 within a year. For presence, each mean value was calculated as the frequency of occurrence 256 i.e. the percentage of samples in which the prey class was present. For numerical abundance, 257 each mean value was calculated as the proportion of all otoliths of a given prey type relative 258 to all otoliths. To aid comparison with frequency of occurrence, we converted numerical 259 abundance proportions into percentages. Study years commenced at the onset of breeding, 260 so each spanned two calendar years. All study years were retained in figures of annual mean data (1985–2014), with model plots presented over the range of years included in the analysis 261 262 (1988–2014).

263 Results

264 Pellet composition

A total of 5888 pellets were collected between 1985 and 2014 (n = 23 years; mean ± SD pellets
year⁻¹: 256 ± 212; range: 31–973), with 5,668 (96%) containing at least one identifiable prey

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267 type. The data set comprised 3140 pellets from the breeding period (mean ± SD pellets year⁻ 268 ¹: 136 ± 112; range: 0–342) and 2748 from the non-breeding period (mean ± SD pellets year⁻ 269 ¹: 119 ± 132; range: 0–538; Supplementary Material Table S2). 270 Fifteen individual prey types were identified using all prey remains (Table 1). Fish were the 271 dominant prey, with sandeel the most frequently encountered, occurring in 79% of pellets (Table 1; Figure 1a). The next most frequently encountered prey was Gadidae (Cod Fishes) 272 273 occurring in 41% of pellets (Figure 1b), followed by Cottidae (Cottids; 20%; Figure 1c), 274 Pleuronectidae (Righteye Flounders; 19%; Figure 1d) and Gobiidae (Gobies; 19%; Figure 1e). 275 All other prey occurred in ≤ 10% of pellets (Table 1). Sand occurred in 52% of pellets (Figure 276 1e; Table 1). The median sample-level and annual prey richness was 5 (range: 0–9) and 12

277 (range: 6–14), respectively.

We recorded 495,239 otoliths belonging to 11 fish prey types, with 4913 (83%) pellets containing at least one identifiable otolith (Table 2; mean \pm SD otoliths pellet⁻¹: 84 \pm 117; range 0–1,048). Otoliths were dominated by sandeel (88%; mean \pm SD relative numerical abundance per sample: 70 \pm 40%). Gadidae were the second most common (7%; mean \pm SD: 46 \pm 41% numerical abundance relative to all non-sandeel otoliths). All other fish prey comprised < 5% of otoliths (Table 2).

284 Temporal and seasonal changes in pellet composition

The best-supported model for sandeel presence contained an effect of year, period and a year by period interaction (Table 3; full model selection table presented in Supplementary Material Table S3). Overall, sandeel frequency of occurrence decreased markedly in both the breeding and non-breeding periods. However, the decline was more pronounced during the nonbreeding period, from 96% in 1988 to 45% in 2014, compared to 91% to 67% during the

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290 breeding season (data values: Figure 1a; predicted values from model: Figure 2a). The best-291 supported model for both Gadidae and Cottidae presence contained an effect of year only 292 (Table 3; Table S3). Gadidae frequency of occurrence increased from 22% in 1988 to 66% in 293 2014 (data values: Figure 1b; predicted values from model: Figure 2b), whereas Cottidae 294 frequency of occurrence increased from 5% in 1988 to 45% in 2014 (data values: Figure 1c; 295 predicted values from model: Figure 2c; Table 3; Table S3). Overall, there was an increase in 296 Pleuronectidae presence over the study, driven predominantly by the breeding period, when 297 frequency of occurrence increased from 7% (1988) to 23% (2014), with frequency during the 298 non-breeding period remaining relatively constant at 15% in 1988 and 14% in 2014 (data 299 values: Figure 1d; predicted values from model: Figure 2d; Table 3; Table S3). Gobiidae 300 presence increased overall between 1988 and 2014, but there was a significant interaction 301 between year and period such that presence was higher during the non-breeding period at 302 the start of the study (breeding 2%; non-breeding 6%), while by the end of the study the 303 frequency was the same in both periods (breeding 21%; non-breeding 21%; data values: 304 Figure 1e; predicted values from model: Figure 2e; Table 3; Table S3). Presence of sand 305 displayed a substantial decline over the study, with a significant year by period interaction 306 such that frequency reduced from 44% to 19% during breeding and 92% to 16% in the non-307 breeding period (data values: Figure 1f; predicted values from model: Figure 2f; Table 3; Table 308 S3).

Sandeel numerical abundance relative to all otoliths decreased from 100% in 1988 to 13% in 2014, but there was no evidence of a difference between the breeding and non-breeding periods (data values: Figure 3a; predicted values from model: Figure 4a; Table 4; full model selection table presented in Supplementary Material Table S4). The decline was less marked at the start of the study, but accelerated from the early 2000s. Gadidae numerical abundance

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314 relative to all non-sandeel otoliths reduced overall, but was consistently higher during 315 breeding (data values: Figure 3b; predicted values from model: Figure 4b; Table 4; Table S4). 316 The magnitude of change was similar in the two seasons, from 68% (1988) to 48% (2014) in 317 the breeding period, and from 54% (1988) to 34% (2014) in the non-breeding period. 318 Sample-level prey richness increased over the study, but with a more marked increase during breeding (from 1.16 prey types pellet⁻¹ in 1988 to 3.36 in 2014) than non-breeding (1.67 prey 319 320 types pellet⁻¹ in 1988 to 2.69 in 2014; data values: Figure 5a; predicted values from model: 321 Figure 6a; Table 5; full model selection table presented in Supplementary Material Table S5). 322 Annual prey richness displayed a quadratic trend over the study, increasing from 6.27 prey 323 types year⁻¹ in 1988 to 12.31 in 2014, with a peak of 15.80 in 2007 (data values: Figure 5b; 324 predicted values from model: Figure 6b; Table 5; Table S5). However, a model containing a linear effect of year received similar support, providing strong evidence for an increasing 325 trend in annual prey richness. 326

327 Discussion

We identified dramatic changes in the diet composition of full-grown European shags 328 329 Phalacrocorax aristotelis (hereafter shag) on the Isle of May over the past three decades both 330 during and outside the breeding season. The dominance of lesser sandeels Ammodytes 331 marinus (hereafter sandeel) decreased, with the decline in sandeel occurrence more marked 332 during the non-breeding period. In contrast, the frequency of Gadidae, Cottidae, 333 Pleuronectidae and Gobiidae increased. Prey richness also increased over the course of the 334 study, in particular during the breeding period. These marked changes highlight the importance of monitoring changes in diet composition throughout the annual cycle. 335

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336 Dietary change

337 Our findings of an overall decline in the dietary contribution of sandeel throughout the annual 338 cycle, support our general prediction that changes in the importance of sandeels over time 339 would be similar in breeding and non-breeding diets, since local sandeel populations are 340 resident (Boulcott et al. 2007). One explanation for this year-round reduction is climate-341 mediated alterations in the abundance, availability or profitability of sandeels associated with 342 rising temperatures in the North Sea (Arnott & Ruxton 2002, Deurs, Van et al. 2009). Similar 343 dietary changes have been observed in other seabird populations in response to changes in 344 prey availability (Miller & Sydeman 2004, Gaston & Elliott 2014, Green et al. 2015). Howells 345 et al. (2017) also recorded a reduction in the length of sandeels fed to nestling shags at this 346 colony over the past three decades, which, due to the negative, non-linear relationship 347 between calorific content and sandeel size (Hislop et al. 1991, Wanless et al. 2005), may be linked to the decreasing prevalence in shag diet. However, due to substantial digestive 348 349 erosion of sandeel otoliths in pellets (Johnstone et al. 1990), it was not possible to use otolith 350 length-fish length relationships to infer changes in sandeel length in this study. With flexible 351 foraging behaviours, as evidenced by the wide range of prey types exploited throughout their 352 range, shags may be able to adjust their diet in response to availability and quality of alternative prey. Such flexibility may be a key mechanism underpinning the dietary trends 353 354 observed in this study, such that sandeel may have become scarcer or lessened in profitability 355 compared to alternative prey, which may themselves have become more abundant or 356 profitable. Data suggests that the energy density of alternative prey is similar to sandeels 357 (Spitz et al. 2010). However, in the absence of estimates of prey availability or capture rates, 358 it is not possible to fully establish the causes underpinning these temporal patterns in diet 359 composition. Industrial fisheries may also reduce the availability of sandeels, with knock-on

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360 effects on seabird diet composition. However, the sandeel fishery off eastern Scotland did not 361 overlap spatially with the foraging distribution of this shag population (Bogdanova et al. 362 2014). Furthermore, the fishery was only operational between 1990 and 1999 (Daunt et al. 2008). As such, we would have expected a stepped reduction in sandeel occurrence in the 363 364 diet over this period, which was not what we found. Similarly, Wanless et al. (2018) did not 365 record a reduction in sandeel occurrence in the diet of the seabird community breeding at 366 the colony during the 1990s. We therefore consider it unlikely that top-down fishing pressure 367 was driving the observed trends in sandeel dietary contribution.

368 The steeper decline in sandeel frequency of occurrence during the non-breeding period may 369 be linked to reduced foraging capacity at this time of the year, as a result of shortened day 370 length, adverse weather and absence of sandeels in the water column, apart from a brief 371 period during spawning (Wright & Bailey 1993, Frederiksen et al. 2008, Daunt et al. 2014). Accordingly, any changes in overall prey availability over the course of the study might have 372 373 had a more pronounced effect on diet composition at this time of year than during the 374 breeding season. However, no seasonal difference in the rate of change was apparent in 375 sandeel numerical abundance. This disparity with sandeel occurrence may arise because 376 numerical abundance is quantified as the proportion relative to other prey, which themselves 377 may have shown seasonal differences in trends. However, we could not test this since we 378 could not distinguish changes in sandeels from reciprocal changes in other prey. Whatever 379 the mechanism, the lack of difference between breeding and non-breeding periods in the 380 trend in numerical abundance of sandeels relative to other prey suggests that this species has 381 shown similar declines throughout the year in terms of biomass consumed. The overall reduction in frequency of sand is in line with these conclusions. Sand ingestion likely reflects 382 383 accidental ingestion when foraging for sandeels, since shags generally extract sandeels

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directly from within the sand sediment (Watanuki et al. 2008), whereas other prey species
that live in these habitats, such as Pleuronectidae and Callionymidae, are more likely captured
on the sea floor.

The increase in dietary frequency of Gadidae accords with recent evidence of a distributional 387 388 shift into Scottish waters of some Gadiformes in recent years (Cormon et al. 2014), including 389 saithe Pollachius virens, the principle prey of shags is some populations. Pleuronectidae 390 frequency also increased in the diet over the last thirty years, so shags may have continued 391 to forage in sandy areas through the course of the study, but increasingly targeted 392 Pleuronectidae, and other prey associated with sandeel habitats, such as Callionymidae, 393 rather than sandeels. Gobiidae also increased, but this prey class is predominantly associated 394 with rocky areas, which accords with past work on this population demonstrating the use of multiple habitats (Watanuki et al. 2008). Gadidae otoliths relative to other non-sandeel prev 395 396 reduced over the study, suggesting that other non-sandeel prey have increased more rapidly 397 than Gadidae. However, there was strong evidence that Gadidae numerical abundance 398 relative to other non-sandeel prey was consistently higher during breeding. This is in contrast 399 to Lilliendahl and Solmundsson (2006) who observed a higher prevalence of Gadidae in 400 Icelandic shag pellets during winter. One possible explanation is that many Gadidae species use inshore waters as nursery grounds, with immatures moving into shallow, coastal feeding 401 402 areas in the Firth of Forth during summer (Bergstad et al. 1987, Heessen et al. 2015).

One consequence of these dietary changes is that both sample-level and annual prey richness
increased over the study, with the latter peaking in 2007. Long-term dietary diversification
has also been observed in other seabird species in response to changes in prey availability
(Gaston & Elliott 2014). The parallel increase in diversity at the single pellet and whole year

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407 scale suggests that, on average, the population is now exhibiting an individual 408 generalist/population generalist structure of resource use (Bolnick et al. 2003). Seasonal 409 patterns of sample-level prey richness changed over the study, such that the increase was 410 more pronounced during breeding, in line with seasonal differences in the pattern of change 411 among Pleuronectidae and Gobiidae frequency of occurrence. Climate-mediated changes in 412 fish populations have been widely reported in the North Sea, including changes in the abundance and distribution of many species (Perry et al. 2005, Dulvy et al. 2008). Thus, the 413 414 dietary trends observed in our study population may be indicative of reductions in the 415 abundance and availability of sandeel, increases in non-sandeel prey or a combination of both. These changes may vary among seasons, but without independent data on any 416 abundance of these prey types it is currently not possible to distinguish these alternatives. 417

418 Limitations

419 It is important to recognise the limitations of estimating year-round diet from pellets when interpreting our results. The most important limitation of pellet analysis is the potential for 420 421 underrepresentation of soft-bodied or easily digestible prey (Barrett et al. 2007). For example, 422 Pholidae and Callionymidae (the otoliths of which are poorly sampled by pellet analysis) can 423 form a substantial proportion of chick diet in this population (Howells et al. 2017), but were recorded infrequently in pellets. One important consequence of this is patterns of long-term 424 change over time might have been different had we been able to detect all prey types. In 425 426 particular, the increase in diversity over the course of the study may be greater than we could 427 demonstrate if more digestible prey than sandeels have become more common in the diet 428 throughout the year, as indicated from our diet data obtained from regurgitates (Howells et 429 al. 2017). A further limitation of our study is that we had to pool all sandeel age-classes. As a

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430 result, we could not examine temporal and seasonal patterns in the relative contribution of 431 different age classes, in contrast to our recent analysis of diet from regurgitations (Howells et 432 al. 2017). Another consideration is that due to substantial differences in detection rates with sandeel size (i.e. larger fish are better represented in pellets; Johnstone et al. 1990), some of 433 434 the observed reduction in sandeel relative numerical abundance may have been exacerbated by changes in detectability, since average sandeel length declined over the course of the study 435 (Howells et al. 2017). However, given the dramatic trends observed in this study and the 436 437 comparatively small decrease in sandeel size observed in chick diet (from in 104.5 mm 1988 438 to 92.0 in 2014), we consider our observation of a decline in sandeel abundance to be robust to this limitation. Finally, uncertainty in the date of pellet production could also have affected 439 440 our results, for example by assigning pellets to the wrong period. However, given the length 441 of non-breeding and breeding periods (several months) compared with the maximum likely 442 duration between pellet production and collection (ca. two weeks), and the fitting of month 443 as a random term in our models, we do not consider that this error would have had a strong 444 impact on our results.

445 Demographic and conservation implications

The year-round reduction in the importance of sandeels in shag diet and associated dietary diversification may have important demographic consequences. In shags, the majority of mortality occurs in winter (Aebischer 1986, Harris & Wanless 1996, Frederiksen et al. 2008), linked to foraging capacity in more challenging environmental conditions (Daunt et al. 2006, 2014, Lewis et al. 2015). Such changes may also be important during pre-breeding, when diet composition can be a key determinant of subsequent reproductive success (Sorensen et al. 2009). Prey availability during the breeding season is also a key determinant of breeding

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success (Daunt et al. 2001, Frederiksen et al. 2007). Crucially, effects on fitness are likely to 453 depend on the relative profitability of different prey types throughout the annual cycle (Hislop 454 455 et al. 1991, Litzow et al. 2004). Due to the difference in habitat associations between prey types, the dietary change observed may also have important implications for shag foraging 456 457 distributions (Bogdanova et al. 2014, Michelot et al. 2017). The increase in proportion of nonsandeels in the diet could alter interactions with anthropogenic activities, such as offshore 458 renewable developments or recreation. Shags in this population are partial migrants, 459 460 whereby a proportion of individuals remain resident throughout the year while the remainder migrate (Grist et al. 2014). Studies that estimate diet composition during the non-breeding 461 period throughout the population range would deliver a more complete picture of the 462 463 potential implications for population dynamics and conservation management.

464 In summary, we identified substantial alterations in diet composition of a population of shags throughout the annual cycle over a thirty-year period. Our results accord with recent climate-465 466 mediated changes in the distribution and abundance of many ecologically and commercially 467 important fish species in the North Sea, most notably sandeel. To our knowledge, this study 468 is the first to have quantified long-term trends in seabird diet outside the breeding season. 469 The similarities and differences observed in these long-term trends compared with those during the breeding season highlight the importance of considering the diet of seabirds 470 471 throughout the annual cycle in assessments of long-term dietary change. That the decline in 472 sandeel frequency and abundance is apparent both during and outside the breeding season 473 suggests that substantial temporal changes in prey populations have occurred, and may have 474 important implications for seabird population dynamics in the region.

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482 Compliance with Ethical Standards

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680 Tables

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Table 1 Summary table of frequency of occurrence of each prey type and sand between 1985-

683 86 and 2014-15, including the % for all pellets combined, mean of annual % and range of

- 684 annual %. Prey that could not be identified to any taxonomic level are referred to as
- 685 Unidentified.

Prey	Pellets (%)	Annual Mean ± SD	Annual Range
Sandeel	4668 (79%)	77% ± 17	47–96%
Gadidae	2409 (41%)	46% ± 17	22–75%
Cottidae	1149 (20%)	21% ± 16	2–47%
Pleuronectidae	1145 (19%)	19% ± 9	4–33%
Gobiidae	1126 (19%)	22% ± 16	2–56%
Crustacea	585 (10%)	13% ± 7	2–27%
Callionymidae	414 (7%)	9% ± 9	0–25%
Pholidae	364 (6%)	7% ± 10	0–32%
Mollusca	354 (6%)	7% ± 8	0–26%
Zoarcidae	346 (6%)	8% ± 9	0–40%
Labridae	210 (4%)	5% ± 5	0–16%
Polychaeta	181 (3%)	4% ± 3	0–7%
Syngnathinae	50 (1%)	1% ± 3	0–15%
Clupeidae	23 (< 1%)	< 1% ± < 1	0–2%
Agonidae	7 (< 1%)	< 1% ± < 1	0–2%
Unidentified	366 (6%)	6% ± 7	0–30%
Sand	3070 (52%)	48% ± 22	12-84%

- Table 2 Summary table of otolith numerical abundance for each fish prey type between 1985-86 and 2014-15, including total number of otoliths
- 688 (and %), annual mean ± SD number of otoliths pellet⁻¹, and range of annual number of otoliths.

Prey	Otolith	Annual Mean ± SD	Annual Range
Sandeel	434,629 (88%)	62.63 ± 36.8	1081–97,665
Gadidae	33,897 (7%)	6.93 ± 5.60	139–5044
Gobiidae	9830 (2%)	1.17 ± 1.24	3–1098
Cottidae	6558 (1%)	2.05 ± 1.80	11–1500
Pleuronectidae	6,291 (1%)	1.11 ± 0.86	2–737
Pholidae	1787 (< 1%)	0.29 ± 0.56	0–691
Zoarcidae	1031 (< 1%)	0.25 ± 0.40	0–262
Callionymidae	805 (< 1%)	0.17 ± 0.30	0-214
Labridae	335 (< 1%)	0.08 ± 0.09	0–64
Clupeidae	64 (< 1%)	0.01 ± 0.05	0–25
Agonidae	12 (< 1%)	< 0.01 ± < 0.01	0–5

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- 690 Table 3 Model selection table for Generalised Linear Mixed Models testing for effects of year, period and a year by period interaction (*) on
- 691 presence of each prey type. Periods are reported as non-breeding (NB) relative to breeding. Table shows model rank compared to other models,
- 692 model structure, fixed effect estimates, standard errors, z ratios, number of parameters (k), difference in AICc between top model and selected
- 693 model (Δ AICc) and Akaike weight relative to other models (ω_i). Due to the large number of prey types and models, we only report those models
- 694 within 10 AICc points of the top model, which is shown in bold (for full model selection tables see Table S3).

Response	Rank	Model	Estimate	SE	z value	k	ΔAICc	ω
Sandeel	1	i + year + period + year*period				4	0	1.00
		year	-0.54	0.22	-2.42			
		period (NB)	0.24	0.16	1.52			
		year* period (NB)	-0.64	0.15	-4.35			
Gadidae	1	i + year				2	0	0.43
		year	0.67	0.11	6.35			
Gadidae	2	i + year + period + year*period				4	0.12	0.41
		year	0.83	0.13	6.36			
		period (NB)	0.02	0.18	0.1			
		year*period (NB)	-0.28	0.14	-1.99			
Gadidae	3	i + year + period (NB)				3	1.95	0.16
		year	0.67	0.11	6.27			
		period (NB)	0.04	0.19	0.24			
Gobiidae	1	i + year + period + year*period				4	0	0.78
		year	0.91	0.19	4.75			
		period (NB)	0.7	0.26	2.69			
		year*period (NB)	-0.41	0.19	-2.16			
Gobiidae	2	i + year + period (NB)				3	2.57	0.22
		year	0.72	0.16	4.45			
		period (NB)	0.84	0.25	3.33			
Pleuronectidae	1	i + year + period + year*period				4	0	0.98
		year	0.46	0.12	3.96			
		period (NB)	0.23	0.21	1.1			
		year*period (NB)	-0.47	0.14	-3.33			
Pleuronectidae	2	i + year + period (NB)				3	8.96	0.01
		year	0.23	0.1	2.36			
		period (NB)	0.37	0.2	1.84			
Cottidae	1	i + year				2	0	0.64

		year	0.92	0.14	6.62			
Cottidae	2	i + year + period				3	2	0.24
		year	0.92	0.14	6.62			
		period (NB)	-0.01	0.19	-0.06			
Cottidae	3	i + year + period + year*period				4	3.3	0.12
		year	0.98	0.16	6.27			
		period (NB)	0	0.19	-0.01			
		year*period (NB)	-0.11	0.14	-0.84			
Sand	1	i + year + period + year*period				4	0	1.00
		year	-0.41	0.25	-1.66			
		period (NB)	1.62	0.32	5.1			
		year*period (NB)	-1.01	0.25	-4.04			

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Table 4 Model selection table for Generalised Linear Mixed Models testing for effects of year, period and a year by period interaction (*) on

697 numerical abundance of sandeel (relative to all otoliths) and Gadidae (relative to all non-sandeel otoliths). Periods are reported as non-breeding

698 (NB) relative to breeding. Table shows model rank compared to other models, model structure, fixed effect estimates, standard errors, z ratios,

699 number of parameters (k), difference in AICc between top model and top model (Δ AICc) and Akaike weight relative to other models (ω_i). Due to

the large number of prey types and models, we only report those models within 10 AICc points of the top model, which is shown in bold (for full

701 model selection tables see Table S4).

Response	Rank	Model	Estimate	SE	z value	k	ΔΑΙϹϲ	ωί
Sandeel	1	i + year				2	0	0.65
		year	-2.84	0.31	-9.03			
Sandeel	2	i + year + period (NB)				3	1.89	0.25
		year	-2.83	0.32	-8.87			
		period (NB)	-0.14	0.39	-0.35			
Gadidae	1	i + year + period				3	0	0.92
		year	2.07	0.12	16.61			
		period (NB)	-0.65	0.23	-2.79			

Running head: Trends in year round shag diet

Table 5 Model selection table for Generalised Linear Mixed Models testing for effects of year, period and a year by period interaction (*) on sample-level prey richness, and linear and quadratic trends in annual prey richness. Periods are reported as non-breeding (NB) relative to breeding. Table shows model rank compared to other models, model structure, fixed effect estimates, standard errors, z ratios, number of parameters (k), difference in AICc between top model and top model (Δ AICc) and Akaike weight relative to other models (ω_i). Due to the large number of prey types and models, we only report those models within 10 AICc points of the top model, which is shown in bold (for full model selection tables see Table S5). Models with similar levels of support as the top model indicated with ⁺.

Response	Rank	Model	Estimate	SE	z value	k	ΔAICc	ω
Sample-level prey richness	1	i + year + season (NB) + year*period (NB)			4	0	1.00
		year	0.37	0.04	8.95			
		period (NB)	0.14	0.06	2.31			
		year*period (NB)	-0.2	0.04	-5.23			
Annual prey richness	1	i + year + year²				3	0	0.66
		year	0.93	0.26	3.62			
		year ²	-0.86	0.37	-2.31			
	2†	i + year				2	1.33	0.34
		year	1.23	0.27	4.63			

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Figure 2. Fitted lines and 95% confidence intervals for frequency of occurrence between 1988-89 and 2014-15 for: a) Sandeel; b) Gadidae; c) Cottidae; d) Pleuronectidae; e) Gobiidae; and f) sand. Plots with a single line indicate a year effect only. Plots with two fitted lines indicate differences in trends between periods (breeding period: solid line; non-breeding period: dashed line).

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Figure 3 Interannual variation in breeding and non-breeding percentage numerical abundance, expressed as the mean across pellets between 1985-86 and 2014-15, for: a) sandeels (relative to all prey), and b) non-sandeels (relative to all non-sandeel prey). Blank years are those in which no pellets were collected. Values presented as percentages to aid comparison with frequency of occurrence.

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prey), and b). Gadidae (relative to all non-sandeel prey) between 1988-89 and 2014-15. Plots with a single line indicate year effect only. Plots with two fitted lines indicate differences between periods (breeding period: solid line; non-breeding period: dashed line). Values presented as percentages to aid comparison with frequency of occurrence.





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Figure 6 Fitted lines and 95% confidence intervals for modelled a) sample-level prey richness and b) annual prey richness between 1988-89 and 2014-15. Plots with two fitted lines indicate differences in trends between periods (breeding period: solid line; non-breeding period: dashed line). The linear and quadratic terms in the sample and annual prey richness plots appear as quadratic and cubic terms, respectively, due to the Poisson distribution of the data.

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- 712 Supplementary Material
- Pronounced long-term trends in year-round diet composition of the European shag *Phalacrocorax aristotelis*

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- Table S1 Start month, end month and length (in months) of breeding and non-breeding
- periods in each study year, together with the total number of months contributing to each
- 526 study year.

		Breeding	S		Non-breeding	g	Total
Year	Start	End	Length (months)	Start	End	Length (months)	Length
1985-86	May	August	4	September	March	7	11
1987-88	April	July	4	August	March	8	12
1988-89	April	July	4	September	April	8	12
1989-90	May	August	4	September	April	8	13
1990-91	June	September	4	October	May	8	12
1991-92	June	September	4	October	April	7	11
1992-93	May	August	4	September	May	9	13
1993-94	June	September	4	October	April	7	11
1994-95	May	August	4	September	March	7	12
1998-99	May	August	4	September	April	8	12
1999-00	May	August	4	September	April	8	12
2003-04	April	July	4	August	April	9	13
2004-05	May	August	4	September	April	8	12
2005-06	May	August	4	September	April	8	12
2006-07	May	August	4	September	April	8	12
2007-08	May	August	4	September	April	8	12
2008-09	May	August	4	September	April	8	12
2009-10	April	July	4	August	March	8	12
2010-11	April	July	4	August	March	8	12
2011-12	April	July	4	August	March	8	12
2012-13	April	July	4	August	March	8	12
2013-14	May	August	4	September	April	8	12
2014-15	April	July	4	August	March	8	12

Running head: Trends in year round shag diet

Table S2 Sample size of pellets in each month between 1985-86 and 2014-15. Bold values indicate breeding period in each year. The non-breeding period in each year starts in the month after the breeding period and ends in the month preceding the commencement of the breeding period in the following calendar year.

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						Мо	nth						
Year	Jan	Feb	Mar	Apr	May	Jun	Jul	Aug	Sep	Oct	Nov	Dec	Total
1985	-	-	-	-	0	0	0	0	0	53	0	0	53
1987	0	0	0	0	0	0	0	0	0	30	0	0	30
1988	0	0	40	39	31	10	30	24	0	0	29	0	203
1989	0	0	38	27	65	83	67	40	42	157	30	0	549
1990	29	36	40	93	111	80	80	99	80	75	0	0	723
1991	0	0	69	119	102	70	91	55	60	61	0	0	627
1992	0	0	80	93	65	56	110	78	19	30	0	0	531
1993	0	41	43	40	76	79	40	0	0	0	0	0	319
1994	0	0	9	29	28	43	31	21	0	0	0	0	161
1998	0	0	0	0	0	0	49	0	0	0	0	0	49
1999	0	0	0	0	0	153	189	0	0	0	0	0	342
2003	0	0	0	0	0	10	72	20	0	0	0	0	102
2004	0	0	0	0	26	83	97	16	0	0	0	0	222
2005	0	0	12	101	104	54	54	22	0	9	0	27	383
2006	23	17	38	51	91	83	24	27	0	28	0	0	382
2007	15	0	29	22	8	29	59	16	0	0	0	0	178
2008	0	0	0	25	0	21	10	0	0	0	0	0	56
2009	0	0	0	24	30	29	19	0	10	0	0	0	112
2010	18	0	26	0	8	6	37	15	31	0	0	18	159
2011	0	18	4	0	5	17	4	63	35	0	16	11	173
2012	28	23	33	0	1	36	41	18	52	30	17	16	295
2013	42	11	0	0	0	9	19	7	0	15	0	33	136
2014	20	28	0	0	1	20	14	20	0	0	0	0	103
2015	0	0	0	-	-	-	-	-	-	-	-	-	-

Table S3 Full model selection table for Generalised Linear Mixed Models testing for effects of year, period and a year by period interaction (*)on presence of each prey type. Periods are reported as non-breeding (NB) relative to breeding. Table shows model rank compared to other models, model structure, number of parameters (k), difference in AICc between top model and top model (Δ AICc) and Akaike weight relative to other models (ω_i). Top models are shown in bold.

Response	Rank	Model	k	Δ AICc	ω _i
Sandeel	1	i + year + period + year*period	4	0	1.00
	2	i + year	2	15.38	<0.01
	3	i + year + period	3	16.76	<0.01
	4	i	1	25.75	<0.01
	5	i + period	2	27.12	<0.01
Gadidae	1	i + year	2	0	0.43
	2	i + year + period + year*period	4	0.12	0.41
	3	i + year + period	3	1.95	0.16
	4	i	1	17.44	<0.01
	5	i + period	2	18.9	<0.01
Gobiidae	1	i + year + period + year*period	4	0	0.78
	2	i + year + period	3	2.57	0.22
	3	i + year	2	12.35	<0.01
	4	i + period	2	12.63	<0.01
	5	i	1	22.66	<0.01
Pleuronectidae	1	i + year + period + year*period	4	0	0.98
	2	i + year + period	3	8.96	0.01
	3	i + year	2	10.34	0.01
	4	i + period	2	11.82	<0.01
	5	i	1	13.37	<0.01
Cottidae	1	i + year	2	0	0.64
	2	i + year + period	3	2	0.24
	3	i + year + period + year*period	4	3.3	0.12
	4	i	1	19.1	<0.01
	5	i + period	2	21.07	<0.01
Sand	1	i + year + period + year*period	4	0	1.00
	2	i + year + period	3	13.44	<0.01
	3	i + period	2	23.41	<0.01
	4	i + year	2	93.67	<0.01
	5	i	1	102.36	<0.01

Table S4 Full model selection table for Generalised Linear Mixed Models testing for effects of year, period and a year by period interaction (*) on numerical abundance of sandeel (relative to all otoliths) and Gadidae (relative to all non-sandeel otoliths). Periods are reported as non-breeding (NB) relative to breeding. Table shows model rank compared to other models, model structure, number of parameters (k), difference in AICc between top model and top model (Δ AICc) and Akaike weight relative to other models (ω_i). Top models are shown in bold.

Response	Rank	Model	k	Δ AICc	ω
Sandeel	1	i + year	2	0	0.65
	2	i + year + period	3	1.89	0.25
	3	i + year + period + year*period	4	3.89	0.09
	4	i	1	25.24	<0.01
	5	i + period	2	26.5	<0.01
Gadidae	1	i + year + period	3	0	0.92
	2	i + year	2	5.21	0.07
	3	i + year + period + year*period	4	8.84	0.01
	4	i	1	79.63	<0.01
	5	i + period	2	81.67	<0.01

Table S5 Full model selection table for Generalised Linear Mixed Models testing for effects of year, period and a year by period interaction (*) on sample-level prey richness, and linear and quadratic trends in annual prey richness. Periods are reported as non-breeding (NB) relative to breeding. Table shows model rank compared to other models, model structure, number of parameters (k), difference in AICc between top model and top model (Δ AICc) and Akaike weight relative to other models (ω_i). Top models are shown in bold. Models with similar levels of support indicated with [†]

Response	Rank	Model	k	Δ AICc	ω _i
Sample prey richness	1	i + year + period + year*period	4	0	1
	2	i + year + period	3	23.84	<0.01
	3	i + year	2	32.04	<0.01
	4	i + period	2	45.01	<0.01
	5	i	1	54.95	<0.01
Annual prey richness	1	i + year + year²	3	0	0.66
	2 ⁺	i + year	2	1.33	0.34
	3	i	1	12.14	<0.01

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