# Geographical variation in the foraging behaviour of the pantropical red-footed booby

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ABSTRACT: While interspecific differences in foraging behaviour have attracted much 18 19 attention, less is known about how foraging behaviour differs between populations of a same species. Here we compared the foraging strategy of a pantropical seabird, the red-footed 20 booby (Sula sula), in five populations breeding in contrasted environmental conditions. The 21 foraging strategy strongly differed between sites, from short trips strictly diurnal in Europa 22 Island (Mozambique channel) to long trips including up to 5 nights at sea in Genovesa Island 23 (Galapagos archipelago). The Expectation Maximisation binary Clustering (EMbC) algorithm 24 was used to determine the different behaviours of individuals during their foraging trips 25 (travelling, intensive foraging, resting and relocating). During the day, the activity budget was 26 similar for all the breeding colonies. During the night, birds were primarily on the water, 27 drifting with currents. At all sites, birds similarly performed intensive foraging in zones of 28 Area-Restricted Search (ARS), although the size and duration of ARS zones differed 29 markedly. Red-footed boobies foraged over deep oceanic waters, with chlorophyll-a 30 concentrations varying between sites. Birds did not appear to target areas with higher 31 32 productivity. We suggest that range differences between populations may be linked to other factors such as intra- and interspecific competition. 33

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# **INTRODUCTION**

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The concept of species-typical behaviour assumes that behavioural traits are common among all members of a species (Greenberg & Haraway 1998). However, behavioural variation is commonly observed within a species (Lott 1991). While interspecific differences in foraging behaviour are well studied, less is known about how populations of the same species differ in their foraging behaviour.

Seabirds are 'central-place foragers' during the breeding period, since they nest on land and 46 47 forage at sea (Orians & Pearson 1979). Foraging strategies are usually linked to the local environmental conditions (e.g. Sims & Quayle 1998, Weimerskirch 1998, Burke & 48 Montevecchi 2009) and vary widely across seabird species (Shealer et al. 2002, Weimerskirch 49 2007). Some species search for unpredictable resources over wide areas covering large 50 distances during their foraging trips, while others specifically target oceanographic features 51 such as fronts, shelf edges or sea mounts to find prey (e.g. Schneider 1982, Haney 1986, 52 Weimerskirch 2007, Freeman et al. 2010). These oceanographic features play an essential role 53 54 in the dispersion and aggregation of nutrients and plankton, which attract both prev and predators. Moreover, it has been found that mesoscale and sub-mesoscale structures (e.g. 55 eddies and filaments) can increase primary productivity and consequently concentrate 56 associated predators such as seabirds (Nel et al. 2001, Weimerskirch et al. 2004, Tew-Kai et 57 58 al. 2009). In tropical oligotrophic waters, resources are scarcer and more heterogeneously distributed compared to temperate and polar waters (Longhurst & Pauly 1987, Ballance et al. 59 1997, Weimerskirch 2007). Several species of tropical seabirds feed in close association with 60 sub-surface predators, such as tuna and dolphins that bring prev to the surface in reach of 61 62 flying predators (Au & Pitman 1986, Hebshi et al. 2008).

The red-footed booby (Sula sula), hereafter RFB, is a non-migrant seabird species that lives 63 year-round in pantropical regions of the Atlantic, Pacific, and Indian Oceans (Nelson 1978). 64 65 During the breeding season, both partners of the pair take turns between nest-guarding and foraging trips. The RFB mainly feeds on flying fishes (Exocoetidae) and flying squids 66 (Ommastrephidae) (Nelson 1978, Schreiber et al. 1996). Since these prey occupy a low 67 68 trophic position, the chlorophyll-a concentration (a common proxy of the water productivity) could be an indicator of their spatial distribution. RFBs appear to target specific areas with 69 higher productivity in some sites (Ballance et al. 1997, Jaquemet et al. 2005, Weimerskirch et 70 71 al. 2005a) but not in others (Young et al. 2010).

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Besides local productivity, competition between individuals may also affect the distribution of the foraging zones around the colonies. Ashmole (1963) described the potential consequences of intraspecific competition on the fitness of central-place foragers like seabirds. He hypothesized that the more a colony hosts individuals, the more the surrounding waters can be locally depleted. This 'Ashmole's halo' (Birt et al. 1987) could lead birds from large colonies to travel further to find resources, inducing a decrease in reproductive success and thus a regulation of the colony size. RFBs breeding in the presence of other tropical booby species could reduce competition by partitioning resources, allowing for coexistence (Lack 1971). As
RFBs are known as the most pelagic booby species (Nelson 1978, Schreiber et al. 1996), we

- 82 expect them to have a longer foraging range in presence of other sulid species.
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84 Few studies have compared the foraging behaviour of a seabird species between different sites to better understand the factors affecting foraging strategies (e.g. Kappes et al. 2011, 85 Oppel et al. 2015). The wide distribution of RFB populations gives the opportunity to 86 examine the influence of contrasted biotic and abiotic conditions from different breeding sites 87 on foraging behaviour. The present study compares the foraging strategies of five different 88 populations of the RFB in the Indian and Pacific Oceans. Since the breeding sites have 89 contrasted local conditions, we predict that search strategies and foraging parameters will 90 differ between sites up to a certain level, constrained by the morphology and common habits 91 of the species. Knowing that the foraging strategy of the RFB varies substantially between the 92 stages of the breeding cycle (Mendez et al. 2016), we focused our study on the incubation 93 period only. 94

#### **MATERIALS AND METHODS**

#### Fieldwork

Data were collected from five sites: Europa Island (EU, Mozambique Channel), Christmas 100 Island (CI, Indian Ocean), Walpole Island (WA) and the Chesterfield Islands (CH, hereafter 101 'Chesterfield') off New Caledonia, and Genovesa Island (GEN) in the Galapagos Archipelago 102 103 (Fig. 1). All five sites host important breeding colonies of RFBs (Table 1). Our study examined the foraging behaviour of RFBs during the incubation period when male and female 104 alternate on the nest to incubate the egg (Nelson 1978). Timing of field work and numbers of 105 individuals studied on each breeding colony are given in Table 1. To study the movements of 106 birds at sea, incubating adults were selected randomly and fitted with 20 g (32 X 22 mm) 107 108 IGotU GPS loggers (Mobile Action Technology, Taipei). Depending on the site and date of deployment, locations were recorded every 10 s, 30 s, 60 s, 120 s or 300 s. GPS loggers were 109 attached to a group of three or four central tail feathers using Tesa tape (Wilson et al. 1997). 110 Birds were captured on nests that were previously identified and mapped. They were marked 111 on the tail or the breast with labile dye to identify the individual rapidly and from a distance. 112 Individuals were captured by hand or, for birds nesting higher in the trees, with a 6 m 113 telescopic fishing pole fitted with a nylon noose. In a few cases, both partners at the same nest 114 were fitted with GPS loggers. Birds were weighed in a bag with a spring balance, at both the 115 deployment and the retrieval of the GPS logger to determine gain or loss of weight. The study 116 birds were also measured (culmen height and length, wing length) at the recovery of the GPS 117 logger. Birds were sexed by their voice when possible (males have a higher pitched voice than 118 females; Nelson 1978) or by measurements (females are larger than males; Nelson 1978; 119 Weimerskirch et al. 2006). Blood samples were also collected on a sub-sample of 15 120

individuals in Europa in 2003 to verify the sex using molecular markers (Weimerskirch et al.2006).

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### Track parameters and behaviour labelling

A total of 199 tracks of birds leaving the island to go to the sea were analysed (Table 1). These tracks represented 1 to 8 successive foraging trips of 123 birds. Complete tracks were defined as trips for which GPS data were available from the departure of the bird from the nest to its return (90% of the dataset). Incomplete tracks, e.g. due to battery failure of the GPS device, were used only to estimate specific parameters when at least the beginning of the return phase of the foraging trip was present. Duration of foraging trip (h), total distance covered (km) and maximum range from the colony (km) were calculated for each track.

132 To determine the different behaviours of individuals during their foraging trips, we used the Expectation Maximisation binary Clustering (EMbC) algorithm (Garriga et al. 2016), a 133 variant of the maximum likelihood estimation of Gaussian mixture models (Redner & Walker 134 1984). The EMbC algorithm is a robust, non-supervised multi-variate clustering algorithm 135 that considers correlation and uncertainty of variables giving a meaningful local labelling 136 137 easily linked to biological interpretations. The annotation of behaviours was based on two input variables: the speed and the turning angle, obtained from successive locations. First, all 138 tracks were linearly interpolated with one location every 2 minutes and the maximum speed 139 was set to 90 km.h<sup>-1</sup> (Weimerskirch et al. 2005b). Each location was clustered by the 140 algorithm into four behaviour categories (Table 2): High velocity/Low turn (HL), High 141 velocity/High turn (HH), Low velocity/Low turn (LL), Low velocity/High turn (LH). A 142 behavioural mode was assigned to each cluster, as suggested by Louzao et al. (2014). The HL 143 and HH labels correspond respectively to travelling and relocating. Relocating reflects 144 important turns with a steady speed and can be interpreted as a displacement between 145 restricted areas of intensive foraging. The LL label corresponds to birds resting at the sea 146 surface, mostly sitting on water and drifting in a single direction induced by surface currents 147 (Weimerskirch et al. 2002). The LH label was interpreted as intensive foraging. A smoothing 148 procedure included in the package was applied to better accounts for the temporal 149 associations among behaviours. All trips from all breeding colonies were treated 150 simultaneously in the analysis conducted with the R package EMbC (Garriga et al. 2016). 151 Proportions of each behaviour along tracks were compared between sites and during daytime 152 or nighttime, i.e. when the sun was  $> 6^{\circ}$  below the horizon. All analyses were conducted in R 153 3.1.2 (R Development Core Team 2014). 154

Area-restricted search (ARS) was defined as at least 3 successive locations (i.e. 4 minutes) labelled as intensive foraging by the EMbC algorithm. To simplify the description of the different behaviour along the trajectory, we merged ARS zones when  $\leq$  4 locations labelled with other behaviours were observed between them (i.e. 10 minutes). The number of ARS zones per hour and their duration were calculated. The area covered was estimated through the minimum convex polygon around all the locations of a specific ARS zone. Each ARS was summarized in one central location by taking the median latitude and longitude.

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## Foraging behaviour and environmental drivers

Kernel estimation (Worton 1989) was used to determine the utilization distribution (UD) 164 probability based on the locations of individuals. Kernel density estimates offer the 165 advantages of being widely used to identify population-level core habitat areas. We used the 166 function kernelUD implemented in the R package adehabitatHR (Calenge 2006) using the 167 reference bandwidth which produces contiguous cores without over-smoothing. Choosing a 168 secant projection and a narrow zone minimizes the distortions in a map generated from 169 projection. To estimate the size of general (95%) and core (50%) foraging areas, we used the 170 function getverticeshr with adapted local projections (Europa: Moznet / UTM zone 37S; 171 Genovesa: WGS 84 / UTM zone 16S; Christmas: RGNC91-93 / Lambert New Caledonia; 172 Chesterfield and Walpole: RGNC 1991 / Lambert New Caledonia). Depth was obtained from 173 the one arc-minute resolution GEBCO bathymetric dataset using the R package marmap 174 175 (Pante & Simon-Bouhet 2013). Monthly composites of chlorophyll-a concentration (chl-a, in mg.m<sup>-3</sup>) were obtained at a spatial resolution of 4 km from Aqua MODIS satellite using the R 176 package *rerddap* (Chamberlain 2016). At a finer time-scale, we used a self-written script to 177 obtain 11-days composites of chl-a concentration at 4 km resolution (GlobColour, merged 178 sensor type and GSM algorithm) using the software GNA Octave (Eaton et al. 2014). 179

For each site, the accessible area was defined by a circle around the colony with a radius 180 corresponding to the maximum range recorded by GPS tracking. The foraging area was 181 delimited by the minimum convex polygon that included all ARS zones of all birds. The 182 accessible but not prospected area was defined as the accessible area to which the prospected 183 area was subtracted. Monthly chl-a concentration was compared between prospected areas 184 and non-prospected areas. Comparisons between travelling and ARS locations were made 185 using 11-day chl-a concentration. Prior to data analysis, travelling locations were resampled 186 with one location every 10 minutes to reduce autocorrelation and have a number of locations 187 in the same order of magnitude than the number of ARS zones. 188

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# Statistical analysis

As some individuals were tracked during several trips, linear mixed-effects models with 191 'individual' as random factor were applied to avoid pseudoreplication. We used the function 192 *lmer* from the R package *lme4* (Bates et al. 2015) to test for differences in trip parameters 193 between breeding colonies. Tukey's HSD test was used to calculate post-hoc comparisons on 194 each factor in the model using the function *glht* from the R package *multcomp* (Hothorn et al. 195 2008). When residuals were not normally distributed, variables were square-root transformed. 196 When the data still did not meet the assumptions, we used a Kruskal-Wallis rank sum test and 197 198 Tukey and Kramer (Nemenyi) test for pairwise comparisons with Tukey-Dist approximation for independent samples from the R package PMCMR (Pohlert 2014). Data from Europa (in 199 years 2003 and 2013; Table 1) were pooled since no significant differences between years 200 were observed. Data from Genovesa (in years 2009 and 2014; Table 1) were analysed 201

separately to take into account the inter-annual effects. Males and females were pooled in all 202 analyses since no significant effect of the sex was observed when doing comparisons of track 203 parameters (p > 0.05). Generalized linear mixed model (GLMM) with binomial family and 204 205 logit link were applied to compare environmental parameters between ARS and travelling with 'individual' and 'track' as random factors using the function glmer from the R package 206 *lme4* (Bates et al. 2015). Values of the dependent variables are given as mean  $\pm$  standard 207 208 deviation. The Marascuilo procedure (1966) was used to compare the pairwise proportions of the behaviours defined according to the EMbC algorithm (Garriga et al. 2016) among 209 breeding colonies. 210

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# Trip parameters

RESULTS

216 Foraging parameters varied extensively between sites. Individuals from Europa undertook short foraging trips exclusively, lasting on average less than 7 h, with a maximum range of 50 217 km, and never spent the night at sea (Fig. 2 and Fig. 3). At Walpole, apart from two trips that 218 lasted 60 h including three nights at sea, trips were only slightly longer than those of Europa 219 (mean duration 8 h, mean range 80 km). In Chesterfield, some trips were longer than 24 hours 220 221 and nights spent at sea were frequently observed, yielding a mean trip duration of 21 h and a maximum range of 125 km. In contrast, overnight trips were common in Genovesa, with a 222 mean duration of 22 h and range of 122 km in 2009, and higher values in 2014 (37 h and 176 223 km). Four trips included 4 nights at sea and one trip included 5 nights at sea. The furthest 224 location was recorded 472 km away from the colony. Birds from Christmas Island made 225 significantly longer trips in duration than in the other sites (45 h on average), but the 226 maximum range recorded (164 km on average) was not greater. Four trips included 4 nights at 227 sea (Fig. 2 and Fig. 3). 228

All the foraging areas of RFB were over oceanic waters but their size clearly differed between 229 sites (Fig. 4). Europa had the smallest foraging area evenly distributed around the island (95% 230 and 50% kernels: 22 243 km<sup>2</sup> and 3 863 km<sup>2</sup> respectively; Fig. 4). The four other sites showed 231 directionality in foraging area. Birds foraged principally to the north-east of Walpole (54 988 232 km², 12 420 km²), to the west of Chesterfield (57 992 km², 14 422 km²), to the east of 233 Genovesa (2009: 60 438 km<sup>2</sup>, 12 497 km<sup>2</sup>; 2014: 132 784 km<sup>2</sup>, 28 206 km<sup>2</sup>) and to the east of 234 235 Christmas (111 900 km<sup>2</sup>, 18 388 km<sup>2</sup>). The surface area covered by birds from Genovesa in 2014 was approximately six times larger when compared to birds from Europa. The direction 236 of all foraging areas was not related to the main wind direction (Table 1 and Fig. 4). 237

The multiple pairwise comparisons (Marascuilo procedure) showed that the percentages of the different behaviours during the foraging trips were not significantly different between sites during the day (Fig. 5). During the night, the high proportion of resting behaviour in Christmas was significantly different from all the other sites. The proportion of resting behaviour in Genovesa differed also from Europa and Chesterfield. The proportion of relocating behaviour at Christmas was significantly different from Europa and Chesterfield. After the sunset, individuals from Europa were mainly travelling for short periods until they reached the colony. Foraging bouts occurred occasionally, and birds never rested on the sea surface. In the other sites, the more the birds tended to spend entire nights at sea, the more a resting behaviour was observed.

Only 2.5% of the dataset (5 tracks from 4 birds) did not contain ARS. The number of ARS 248 zones per hour was slightly different between breeding colonies ( $F_{4.91} = 2.81$ , p = 0.03), with 249 0.5-0.7 ARS.h<sup>-1</sup> on average (Fig. 6). Only Europa and Christmas differed significantly 250 (Tukey's HSD test, p = 0.016), with the highest values observed in Europa (up to 1.67 ARS.h<sup>-</sup> 251 <sup>1</sup>). The mean duration of ARS differed between sites ( $F_{4.91} = 5.91$ , p < 0.001). ARS lasted on 252 average between 16 min and 28 min (Fig. 6). ARS from birds in Europa and Chesterfield, 253 doing relatively short trips, were significantly longer than in Genovesa (Tukey's HSD test, p 254 = 0.01 and p = 0.04 respectively) and Christmas (Tukey's HSD test, p = 0.01 and p = 0.03255 respectively). Walpole was intermediate (Tukey's HSD test, p > 0.05). Long ARS lasting 256 more than one hour were occasionally observed at Europa but were rare at other sites. Mean 257 ARS surface ranged between 0.45 km<sup>2</sup> and 1.86 km<sup>2</sup> (Fig. 6), with often larger values for 258 Europa and Chesterfield which were statistically different from Christmas (Tukey's HSD test, 259 p = 0.01 and p = 0.04 respectively). No inter-annual effect was observed in Genovesa for the 260 three descriptive parameters (Tukey's HSD test, p > 0.05). 261

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# Foraging areas and oceanographic conditions

For Europa, Genovesa and Walpole, the incubation period occurred 2-3 months after the 264 annual peak of chl-a in the waters around each island, and 2-3 months before the peak at 265 Chesterfield and Christmas (Fig. 7). We observed a high variability in the concentration of 266 chl-a between study sites. Inside the foraging areas (Fig. 8), waters around Europa and 267 Walpole showed a particularly low concentration (mean 0.07 mg.m<sup>-3</sup>), significantly different 268 from the three other sites (Tukey's HSD test, p < 0.05). The chl-a concentration was 269 considerably higher in the foraging areas of birds from Genovesa and Christmas (more than 270 0.15 mg.m<sup>-3</sup> on average). Chesterfield was intermediate (0.11 mg.m<sup>-3</sup> on average) but not 271 significantly different from Genovesa and Christmas (Tukey's HSD test, p > 0.05). 272

Birds from Europa foraged in all directions with no specific orientation (Fig. 8). The foraging 273 areas of the four other sites were clearly oriented towards specific directions. RFB did not 274 seem to especially favour areas of higher chl-a concentration (Table 3). A slightly higher 275 276 monthly chl-a concentration in the prospected area was observed in Europa and was more pronounced in Christmas. In the three other sites, the mean chl-a concentration was similar or 277 slightly but not significantly higher in the non-prospected area. Regarding the bathymetry, 278 birds foraged over relatively shallow oceanic waters at Europa, Chesterfield, Genovesa and 279 Walpole, with average depths ranging between 2 000 - 3 000 m (Table 3). Most birds from 280 Christmas moved over a deep oceanic trench during their foraging trips, leading to an average 281

depth of c. 5 000 m in the foraging area. Depending on the site, the bathymetry wasalternatively higher in the prospected or the non-prospected area (Table 3).

At a finer scale, the 11-day composite chl-a concentration and the bathymetry were compared 284 between ARS and travelling segments of a trip (Table 4). We found no significant differences 285 in chl-a for Europa (GLMM, p = 0.50) and Genovesa (GLMM, p = 0.08 and p = 0.62 in 2009 286 and 2014, respectively). Higher values were observed inside ARS than during travelling for 287 Christmas (GLMM, p < 0.001) and lower values for Chesterfield (GLMM, p = 0.03) and 288 Walpole (GLMM, p=0.05). No significant differences in bathymetry between ARS and 289 290 travelling were observed in Europa, Walpole and Chesterfield. ARS occurred in significantly deeper waters than travelling in Genovesa (GLMM, p = 0.01 and p < 0.0001 in 2009 and 291 2014, respectively) and Christmas (GLMM, p < 0.0001). 292

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#### **DISCUSSION**

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This study is the first to compare the foraging behaviour and its relationship with 298 oceanographic conditions for a seabird species during a same breeding stage across breeding 299 colonies over a large extent of the species' pantropical range. We found significant inter-300 301 colony differences in foraging behaviour, especially extensive differences in foraging duration and range between sites. These differences were not directly explained by chl-a 302 concentration, a proxy of marine productivity. However, some similarities common to all sites 303 were observed at a fine spatio-temporal scale, as the proportion of the different behaviours 304 305 during the day and the main characteristics of area-restricted search (ARS) zones. Beyond environmental conditions, we suggest that intra- and interspecific competition within a colony 306 and with adjacent colonies can explain the large diversity of foraging strategies used by the 307 308 red-footed booby.

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# Intraspecific differences in foraging behaviour

Our study confirms that RFBs are oceanic foragers throughout their range but also indicates 312 that the foraging behaviour differs substantially among colonies. Birds nesting on Europa 313 undertook relatively short and exclusively diurnal foraging trips. The foraging trips of the 314 315 birds from Walpole were similar to those from Europa, except for two trips including nights at sea. The duration of trips was respectively higher in Chesterfield, Genovesa and Christmas, 316 where trips lasting more than a day were frequently observed. The longest durations and 317 ranges were observed in Genovesa but birds from Christmas made the longest trips on 318 average. Until the present study, red-footed boobies were thought to undertake diurnal 319 320 foraging trips exclusively based on preliminary results from GPS tracking (Weimerskirch et

al. 2005a, Young et al. 2010). The only locality where it was suggested from observation that 321 trips can last more than one day was in the Galapagos (Nelson 1978; Schreiber et al. 1996). 322 Here we confirmed the previous visual observations in the Galapagos, reporting birds leaving 323 324 the colony of Genovesa up to five days, and we showed that during these long trips birds can forage at up to 472 km from the colony. At night, the percentages of the different behaviours 325 varied extensively across the five breeding colonies of RFB. At Europa, birds travelled 326 327 rapidly in order to return to the colony and rest on land. For the other breeding colonies, slow and linear trajectories suggested that the birds floated on the water during the night, being 328 drifted by surface currents. Foraging activity was rare, occurring presumably during dawn and 329 dusk. Since RFBs are visual foragers with likely crepuscular vision, nocturnal foraging is 330 constrained by the lack of ambient light (Ashmole & Ashmole 1967). Weimerskirch et al. 331 (2005a) suggested that predation may be a reason for the RFBs from Europa to stay on land 332 during the night. RFBs from Genovesa, Chesterfield, Walpole and Christmas frequently 333 drifted on the sea surface at night, susceptible to attacks from below. Observations of foot 334 damage to Nazca Boobies Sula granti in the Galapagos indicated possible attacks from 335 toothed sub-surface predators (Zavalaga et al. 2012). Sharks are known to attack seabirds 336 (Johnson et al. 2006, Meyer et al. 2010) but since they are potentially present at all five 337 studied sites, predation risk may not be the main factor explaining the different foraging 338 behaviour observed. Sharks may rely on vision to detect seabirds on the surface, inducing that 339 340 resting at night may not be a high-risk behaviour. Birds could then afford nocturnal predation risk for example in case of low prey encounter during the previous day (Zavalaga et al. 2012). 341

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#### Similarities in diurnal foraging

Individuals of a species possess similar behaviours, even if discrete populations do not mix. 345 This 'Species-Typical Behaviour' is influenced by species genes and social-natural 346 environment (Haraway & Maples 1998). Many similarities appeared in the way RFBs used 347 their environment during the day. Generally, they left from and returned to the colony in a 348 straight trajectory. During the middle sections of the foraging trip, birds simultaneously 349 reduced their speed and increased their sinuosity, suggesting that they found a patch of prev 350 (Weimerskirch et al. 2005a). ARS frequency, size and duration showed large variability 351 within sites. This variability may reflect a fine adjustment of the time spent in a patch of food 352 according to its prey abundance and distribution, before moving to another. In order to 353 optimize their foraging trip, birds should minimise the travelling time between foraging areas 354 and their colonies (Charnov 1976). ARS were more variable at Europa, where trips were short 355 and strictly diurnal, with higher occurrence, larger sizes and longer durations than for the 356 other breeding colonies. Furthermore, the EMbC behaviour analysis of birds from Europa 357 showed a higher proportion of intensive foraging behaviour during trips, reflecting an 358 optimization of the daily trip. Overall similarity in foraging strategy might be related to the 359 fact that tropical seabirds generally feed in association with subsurface predators like tuna and 360 dolphins that make the preys available at the surface (Au & Pitman 1986). However, a study 361

reported that RFBs from Hawaii did not associate with any subsurface predator in greater proportion than what would be expected by chance (Hebshi et al. 2008). Further research is still needed to better understand the foraging strategies of tropical seabirds in oligotrophic waters.

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#### Foraging behaviour and productivity

Previous studies assumed that RFBs may forage in more productive areas (Ballance et al. 368 1997, Jaquemet et al. 2005, Weimerskirch et al. 2005a). However, Young et al. (2010) did not 369 370 find any major variation of chl-a concentration around a site that harbours a large RFB colony, the Palmyra Atoll (Northern Pacific), that would support this hypothesis. After 371 examining the chl-a concentration inside prospected and non-prospected areas, we found that 372 birds from Christmas Island targeted productive areas with deep sea bed, but not birds from 373 the other colonies. At all the remaining sites, birds would have been able to reach more 374 productive waters within their range if they had flown in another direction. Top marine 375 376 predators such as cetaceans and seabirds target productive waters to increase their encounter rate with prev patches in restricted areas (Jaquemet et al. 2005). At Europa, the feeding of 377 great frigatebirds Fregata minor is positively linked with dynamical fronts at the edge of 378 eddies (Weimerskirch et al 2004, Tew-Kai et al. 2009, De Monte et al. 2012, Jaquemet et al. 379 2014). However, the distribution of frigatebirds is negatively influenced by chl-a 380 381 concentration, suggesting that frigatebirds do not seem to directly target high primary productivity (Thiers et al. 2014). Similarly, the productivity found in the foraging area of 382 masked boobies Sula dactylatra in the eastern tropical Pacific is not significantly different 383 from the non-prospected area within the foraging range of the population (Weimerskirch et al. 384 2008). As tropical waters are characterised by an overall lower productivity compared to 385 temperate or polar waters (Longhurst & Pauly 1987), the distribution and abundance of prev 386 is believed to be more unpredictable than in colder waters (Ashmole 1971). Here we see that 387 the chl-a concentration is not a good indicator of foraging areas of RFB in tropical 388 389 environments.

390 Time lags, physical forcings and food web processes can thwart the link between primary productivity and the distribution of predators. Because seabirds do not feed directly on 391 primary producers, a natural delay due to energy transfer between phytoplankton, fish or 392 squid occurs. For example, in the Benguela current system, this phenomenon takes up to 393 several weeks (Grémillet et al. 2008). Moreover, seabird prey seems to be less uniformly 394 distributed than plankton (Piontkovski & Williams 1995). For top marine predators, long 395 time-series of chl-a may be better indicators of productive habitats than finer temporal-scale 396 measurements (Survan et al. 2012). Static non-biological features, such as water depth and 397 distance to shore, can be better explanatory variables than chl-a (Nur et al. 2011). Since we 398 did not find a direct effect of the bathymetry nor the chl-a in four of the five sites, other 399 factors may account for the differences of foraging ranges observed between breeding 400 colonies. 401

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#### **Resource partitioning**

405 Resource competition may lead to adaptations that reduce niche overlap (Gause 1934) and thus explain differences in seabird foraging area and behaviour (Rome & Ellis 2004, Lance et 406 al. 2005). In mixed colonies, seabirds may have to cope with interspecific and intraspecific 407 competition. Birds from larger colonies could also have to forage further than birds from 408 smaller colonies because individuals foraging close to the colony may cause local prev 409 depletion (Ashmole 1963, Furness & Birkhead 1984, Jovani et al. 2015). The sizes of RFB 410 colonies differ extensively between study sites. The small population at Europa (2 800 - 3 800 411 412 pairs, Le Corre & Jouventin 1997) had the shortest foraging range while the large population 413 at Genovesa (140 000 pairs, Nelson 1978) had the longest foraging range, suggesting that intraspecific competition may partly explain the differences in foraging range between 414 breeding colonies. In Genovesa, high intraspecific competition may lead birds to travel 415 several days including nights at sea and thus reach great distances. Grémillet et al. (2004) 416 studied two close colonies of Cape gannets Morus capensis in South Africa and found that 417 birds from the larger colony did longer foraging trips in duration and range. Similarly, mean 418 foraging trip duration of the northern gannet Morus bassanus from colonies in Britain and 419 Ireland was positively correlated with colony size (Lewis et al. 2001). In tropical ecosystems, 420 tracking of masked boobies from two islands differing in colony size, surrounded by similar 421 422 oligotrophic waters, was also consistent with Ashmole's hypothesis (Oppel et al. 2015). In Clipperton (Pacific Ocean), masked boobies showed a particularly long foraging range 423 (average range of 103 km, maximum 242 km; Weimerskirch et al. 2008) and the huge colony 424 425 size (> 100 000 individuals) might explain that range.

426 Present or even previous competition could produce interspecific variation in foraging behaviour (Trivelpiece et al. 1987). The RFB is the only booby species present in Europa, 427 while the four other sites host one or two other booby species. A little or no interspecific 428 429 competition could explain why foraging trips were almost evenly distributed in a short range around Europa, and only during the day. The RFB is the smallest booby species and may fly 430 further in presence of other booby species because of lower flight costs. Incubating RFBs in 431 Johnston Atoll (central Pacific) made diurnal trips significantly longer than those made by 432 brown boobies Sula leucogaster, with a mean duration of 14 h and 6.7 h respectively (Lewis 433 et al. 2004). RFBs and masked boobies from Palmyra Atoll showed strong differences in their 434 foraging behaviour and ranges, with RFBs being again more pelagic than masked boobies 435 (Young et al. 2010). The two same species in Tromelin Island (Indian Ocean), surrounded by 436 more oligotrophic waters, demonstrated interspecific segregation at the level of core foraging 437 areas but not at the scale of the whole foraging region (Kappes et al. 2011). However, intra-438 and interspecific competition did not explain the higher maximum ranges observed in Tern 439 Island (3 000 - 5000 RFB pairs) compared to Palmyra Atoll that hosts 25 000 RFB pairs and 440 one more tropical booby species (Young et al. 2015). However, the two islands have different 441 oceanographic contexts thus potential environmental effects may overshadow the competition 442 effect. 443

The exclusion by adjacent colonies is also known to potentially influence the directionality of 445 the foraging trips (Wakefield et al. 2013). The small foraging range observed in Europa may 446 be caused by a small competition since the island hosts a relatively small RFB colony, with 447 no other tropical boobies and no other island in the vicinity. Genovesa is one of the 448 Northeastern islands of the Galapagos archipelago. Since the foraging range was clearly 449 oriented in the east direction, birds may avoid competition with colonies from other species 450 that have shorter ranges (Anderson 1991). Lastly, no island is present in the vicinity of 451 452 Christmas Island, where the foraging area towards Java Island seems to be mainly driven by the environment. Although resource partitioning between distant colonies is difficult to 453 evaluate, our data suggest that resource partitioning may also have an influence on the 454 foraging behaviour observed at the colony-scale. 455

To conclude, the environmental context and competition may affect the foraging behaviour of the RFB, a central-place forager in tropical oligotrophic waters. To better understand the patterns observed in infra-specific studies, multi-species studies and information about the local environment seem essential to assess the impact of each effect resulting in the foraging behaviour.

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670	LEGEND TO TABLES
671	
672 673 674 675 676	Table 1: Study sites and data collected on incubating red-footed boobies fitted with GPS loggers. EU = Europa, WA = Walpole, CH = Chesterfield, GEN = Genovesa, CI = Christmas. S= South, E = East. RFB = Red-footed booby <i>Sula sula</i> , BB = Brown booby <i>Sula</i> <i>leucogaster</i> , MB = Masked booby <i>Sula dactylatra</i> , AB = Abbott's booby <i>Papasula abbotti</i> , NB = Nazca booby <i>Sula granti</i>
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678 679	Table 2: Values of the delimiters of speed and turning angle of the four behaviours assigned by the EMbC algorithm
680	
681 682 683 684 685	Table 3: Comparison of the chlorophyll-a concentration (chl-a, in mg.m <sup>-3</sup> ) and the bathymetry (bathy, in m) between the foraging area and the accessible but non-prospected area of red-footed boobies <i>Sula sula</i> from five different breeding colonies. Values expressed as mean $\pm$ SD and significantly higher means are in bold for pairwise comparisons. EU = Europa, WA = Walpole, CH = Chesterfield, GEN = Genovesa, CI = Christmas
686	
687 688 689 690 691	Table 4: Comparison of chlorophyll-a concentration (chl-a, in mg.m <sup>-3</sup> ) and bathymetry (bathy, in m) between travelling and area-restricted search (ARS) zones of red-footed boobies <i>Sula sula</i> from five different breeding colonies. Significantly higher values (mean $\pm$ SD) generated from GLMM models are in bold. EU = Europa, WA = Walpole, CH = Chesterfield, GEN09 = Genovesa (2009), GEN14 = Genovesa (2014), CI = Christmas
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Site	Island size	Colony location	Main wind direction	Study period	Number of RFB tracked	Number of tracks	RFB population size (pairs)	Other booby species (pairs)
<b>F</b> U	28	40.3°E	0E	2003: 09/08 - 09/23	9	9	2 200 2 200 8	1
EU	km²	22.3°S	5E	2013: 09/23 - 10/16	13	34	2 800-3 800 "	/
WA	2 km²	158.4°E 9.9°S	SE	2012: 05/27 - 06/02	17	25	c.1000 <sup>f</sup>	BB (100s) <sup>b</sup>
СН	< 10 km²	168.9°E 22.6°S	SE	2014: 09/20 - 09/24	7	13	7 200-7 300 <sup>c</sup>	MB (280-500 <sup>c</sup> ) BB (3 800-5 800 <sup>c</sup> )
GE N	14 km²	105.6°E 10.5°S	S	2014: 07/29 - 08/22	15	39	140 000 <sup>e</sup>	NB (> 1000 <sup>f</sup> )
CI	135 km²	-89.9°E 0.3°S	SE	2009: 11/13 - 11/25 2014: 11/10 - 11/25	26 36	37 42	12 000 <sup>d</sup>	AB (2 500 <sup>d</sup> ) BB (5 000 <sup>d</sup> )
	Km²	0.3°S		2014: 11/10 - 11/25	30	42		BB (5 000 °)

<sup>a</sup> Le Corre & Jouventin 1997, <sup>b</sup> Spaggiari et al. 2007, <sup>c</sup> Borsa et al. 2010, <sup>d</sup> James & McAllan 2014, <sup>e</sup> Nelson 1978, <sup>f</sup> H. Weimerskirch (pers. comm.)

# 707 Table 2:

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	Delimiter				
Behaviour	Speed min (km.h <sup>-1</sup> )	Speed max (km.h <sup>-1</sup> )	Turn min (radians)	Turn max (radians)	
Resting	0	4	0	0.30	
Intense foraging	0	14	0.30	3.14	
Travelling	4	90	0	0.31	
Relocating	14	90	0.31	3.14	

711 Table 3:
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Breeding	Donomotor	Area categ	ory
colony	Parameter -	Non-prospected	Foraging
EU	chl-a	$0.149\pm0.014$	$0.153 \pm 0.018$
EU	bathy	$-2975 \pm 554$	$-3023\pm391$
XXZ A	chl-a	$0.129 \pm 0.033$	$0.111\pm0.017$
WA	bathy	$-2290 \pm 1481$	$-2634 \pm 1383$
CU	chl-a	$0.111\pm0.033$	$0.111\pm0.027$
CH	bathy	$-2176\pm1092$	$\textbf{-2058} \pm \textbf{797}$
CENOO	chl-a	$\textbf{0.207} \pm \textbf{0.050}$	$0.190\pm0.036$
GEIN09	bathy	$-2726\pm709$	$-2268\pm503$
CEN14	chl-a	$0.220\pm0.049$	$0.203\pm0.039$
GEN14	bathy	$-2763\pm679$	$\textbf{-2203} \pm \textbf{512}$
CI	chl	$0.182\pm0.039$	$\textbf{0.207} \pm \textbf{0.043}$
CI	bathy	$-4905 \pm 1055$	$-5096 \pm 1283$

715	Table 4.
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Breeding	Parameter	Beha	viour	Test	
colony		Travelling	ARS	Z	p-value
EU	chl-a	$0.076\pm0.059$	$0.069\pm0.053$	-0.669	0.5036
	bathy	$-2828\pm703$	$-2890 \pm 631$	-1.045	0.2961
	chl-a	$\boldsymbol{0.077 \pm 0.015}$	$0.069\pm0.017$	-1.961	0.0499
WA	bathy	$-2369 \pm 1636$	$-3153 \pm 1806$	-0.869	0.3851
GU	chl-a	$\textbf{0.119} \pm \textbf{0.028}$	$0.112\pm0.022$	-2.179	0.0294
СН	bathy	$-1894 \pm 864$	$-1969 \pm 853$	-1.588	0.1122
GENIAA	chl-a	$0.111 \pm 0.055$	$0.103\pm0.038$	-1.747	0.0806
GEN09	bathy	$-1924\pm507$	$-2029 \pm 496$	-3.195	0.014
GEN14	chl-a	$0.190\pm0.067$	$0.184\pm0.061$	-0.500	0.617
	bathy	$-2154\pm508$	$-2288 \pm 463$	-4.239	< 0.001
CI	chl-a	$0.146\pm0.063$	$\textbf{0.156} \pm \textbf{0.069}$	3.708	< 0.001
CI	bathy	$-5297 \pm 1221$	$-5503 \pm 1104$	-3.423	< 0.001

720	LEGEND TO FIGURES
721	
722 723	Figure 1: Locations of the five breeding colonies of red-footed boobies <i>Sula sula</i> studied during incubation
724 725 726 727	Figure 2: Boxplots of trip duration (h) and maximum range (km) for red-footed boobies <i>Sula sula</i> from five different breeding colonies. Different letters above boxes indicate significant differences (Tukey's HSD test). EU = Europa, WA = Walpole, CH = Chesterfield, GEN09 = Genovesa (2009), GEN14 = Genovesa (2014), CI = Christmas
728 729 730	Figure 3: Distribution of trip duration (h) and maximum range (km) for red-footed boobies <i>Sula sula</i> from five different breeding colonies. $EU = Europa$ , $WA = Walpole$ , $CH = Chesterfield$ , $GEN09 = Genovesa$ (2009), $GEN14 = Genovesa$ (2014), $CI = Christmas$
731 732 733	Figure 4: General (95% kernel density estimation, <i>light</i> ) and core (50% kernel density estimation, <i>dark</i> ) foraging areas of red-footed boobies <i>Sula sula</i> from five different breeding colonies superimposed on bathymetric maps. Colony sites are indicated by a yellow star
734 735 736 737	Figure 5: Mean proportion of each behaviour for red-footed boobies <i>Sula sula</i> from five different breeding colonies. Behaviour was determined along tracks according to EMbC analysis. Results are displayed in the form of pie charts according to the site and the period of the day
738 739 740 741	Figure 6: Number per hour, mean duration (min) and mean surface ( $km^2$ ) of area-restricted searches (ARS) zones for red-footed boobies <i>Sula sula</i> from five different breeding colonies. Different letters above boxes indicate significant differences (Tukey's HSD test). EU = Europa, WA = Walpole, CH = Chesterfield, GEN = Genovesa, CI = Christmas
742 743 744	Figure 7: Time series of the monthly chlorophyll-a concentration in the accessible area of red- footed boobies <i>Sula sula</i> from five different breeding colonies. Grey boxes indicate incubation periods
745 746 747 748	Figure 8: Foraging areas (red polygons) and accessible areas (black circles) of red-footed boobies <i>Sula sula</i> from five different breeding colonies, superimposed on monthly chlorophyll-a concentration maps. Colony sites are indicated by a yellow star and grey dots represent the centroid of each area-restricted search (ARS) zone.
749	
750	

Figure 1:



Figure 2:









Figure 5:











Figure 7:





0.3

0.2