

1 **High trophic niche overlap in mixed-species colonies using**  
2 **artificial nests**

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Although successful at recovering endangered populations, conservation actions based on nest provisioning seldom consider how they shape the composition of communities and alter interspecific interactions. Specifically, the extent to which dietary overlap within these communities may affect the conservation of target species has rarely been assessed. In Southern Europe, large-scale nest-site provisioning aimed at recovering Lesser Kestrels *Falco naumanni* populations attracted non-target bird species, resulting in mixed breeding assemblages that might promote interspecific competition for resources during breeding. Here we used stable isotope analysis ( $\delta^{15}\text{N}$  and  $\delta^{13}\text{C}$ ) to assess inter- and intraspecific dietary segregation in these assemblages and investigate the mechanisms allowing species coexistence. We examined resource partitioning and trophic niche overlap among Lesser Kestrels, Common Kestrels *Falco tinnunculus*, European Rollers *Coracias garrulus*, Western Barn Owls *Tyto alba*, Little Owls *Athene noctua*, and Spotless Starlings *Sturnus unicolor*; and within species between parents and their offspring. Similar isotope ratios and highly overlapped niches, particularly among Lesser Kestrels, Rollers, and Starlings, suggest limited dietary segregation and use of similar prey. Within species, parent-offspring segregation was marked across all species. Our results indicate that species breeding in these assemblages occupy similar ecological niches, despite a potential increase in competition. High-resource availability in the area may permit coexistence but the viability of mixed-species groups may be compromised in areas with limited resources, which are predicted to expand with ongoing human and climate-induced changes. Conservation practices based on nest provisioning should consider the ecological niches of target and non-target species as well as their interactions.

**Keywords:** isotopic niche, dietary segregation, parent-offspring segregation, interspecific competition, mixed-species colonies, community



51 Human-induced changes in the environment shape the distribution of many species by  
52 restricting their access to resources such as food or nest-sites (Wiegand *et al.* 2005;  
53 Teckentrup *et al.* 2019). Ongoing biodiversity loss has prompted conservation projects  
54 including provisioning of artificial nest-sites to halt species decline in many taxa and regions  
55 (Harper *et al.* 2005; Catry *et al.* 2009; Olah *et al.* 2014; Sutherland *et al.* 2018), further re-  
56 shaping the nesting landscape and altering the composition and structure of communities  
57 (Duckworth *et al.* 2017; Catry & Catry 2019). Nest-site provisioning may modify  
58 interspecific interactions and trophic dynamics (Duckworth *et al.* 2017; Catry & Catry 2019),  
59 but their consequences can be subtle and potentially overlooked by researchers or  
60 conservation managers (Mainwaring *et al.* 2015).

61 According to niche theory and the competitive exclusion principle, species with similar  
62 niches, likely to compete with one another, are not expected to coexist (Hardin 1960; Pianka  
63 1981). If a new and artificial nesting landscape promotes the aggregation of potential  
64 competitor species, artificial nest-sites may turn into ecological traps by reducing long-term  
65 individual fitness through a reduction of resource availability or increased interspecific  
66 aggressive behaviours (Kappeler *et al.* 2015; Catry & Catry 2019).

67 To avoid competition, especially when resources are limited, coexisting species can segregate  
68 in at least one of three main axes: space, time, and diet (Pianka 1981). Segregation in spatio-  
69 temporal habitat utilisation occurs when species forage in different locations (e.g.  
70 insectivorous desert bats, Razgour *et al.* 2011), different periods of the day (e.g. African  
71 carnivore communities, Schuette *et al.* 2013), or when they have different breeding  
72 phenologies (e.g. sympatric penguin species, Clewlow *et al.* 2019). Dietary segregation  
73 occurs when species reduce their dietary niche overlap by foraging for alternative food  
74 resources or similar items with different sizes (Macarthur & Pianka 1966; Jenkins & Davoren  
75 2020; Mansor *et al.* 2021). Species may also contract their dietary niches to focus on certain

76 prey types that become temporarily super-abundant, which increases the dietary niche overlap  
 77 among species without increasing competition (Forero *et al.* 2004; Charter *et al.* 2018;  
 78 Denhard *et al.* 2020). Dietary niche segregation can also occur within species to reduce  
 79 intraspecific competition, through sexual or parent-offspring segregation, especially when the  
 80 potential for competitive interactions is high, such as for central place foragers during  
 81 reproduction (Orłowski *et al.* 2014; Catry *et al.* 2016a; Reisinger *et al.* 2020).

82 Here, we investigate dietary niche overlap in a multi-species bird assemblage created by  
 83 large-scale nest-site provisioning aimed at recovering the colonial Lesser Kestrel *Falco*  
 84 *naumanni* in southern Portugal (Catry *et al.* 2009). In a treeless landscape with few nesting  
 85 opportunities, many bird species took advantage of these artificial breeding structures – some  
 86 with more than 80 cavities, resulting in mixed-species colonies (Catry & Catry 2019). These  
 87 include Lesser Kestrels, Common Kestrels *Falco tinnunculus*, European Rollers *Coracias*  
 88 *garrulus*, Western Barn Owls *Tyto alba*, Little Owls *Athene noctua*, Western Jackdaws  
 89 *Corvus monedula*, Spotless Starling *Sturnus unicolor*, and Feral Pigeons *Columba livia*  
 90 (Catry & Catry 2019). Excluding Feral Pigeons and Western Jackdaws, the dietary habits of  
 91 these species, including mostly large arthropods and small rodents, suggest that their use of  
 92 trophic resources may overlap to a large extent (Jaksić *et al.* 1982; Motis *et al.* 1997; Tomé *et*  
 93 *al.* 2008; Catry *et al.* 2016a; Orihuela-Torres *et al.* 2017; Catry *et al.* 2019), especially during  
 94 the breeding season, when competition for food should peak as parents need to obtain  
 95 resources to fulfil their own energy requirements and those of their offspring (Orłowski *et al.*  
 96 2014; Antón-Tello *et al.* 2021). High densities of breeding pairs could lead to prey depletion  
 97 in the vicinity of the colonies, further increasing inter and intraspecific competition (*sensu*  
 98 Ashmole’s halo; Birt *et al.* 1987; Dehnhard *et al.* 2020; Jenkins and Davoren, 2020).

99 We used stable isotope analysis (SIA) to investigate inter and intraspecific overlap in the diet  
 100 of six bird species breeding in mixed colonies installed in artificial breeding structures to

101 better understand the mechanisms allowing their coexistence. Specifically, using stable  
102 isotope ratios of carbon ( $\delta^{13}\text{C}$ ) and nitrogen ( $\delta^{15}\text{N}$ ), we aimed to: (1) compare isotopic niche  
103 breadth and overlap among the species and (2) assess parent-offspring diet segregation within  
104 each species. We sampled feathers from nestlings and blood plasma from parent-offspring  
105 pairs during the breeding season, providing detailed insight into the dietary resources used by  
106 these species. We predict species with similar diets and similar trophic positions to show  
107 higher dietary overlap, while species at the top and bottom ends of the trophic spectrum to be  
108 more segregated from the rest of the assemblage. We consider the extent to which dietary  
109 overlap and niche segregation among potential competitors may affect the conservation of  
110 Lesser Kestrels.

111

## 112 **2. Methods:**

### 113 ***2.1 Study area and study species***

114 This study was carried out in the Castro Verde Special Protection Area (SPA, Natura 2000),  
115 Southeast Portugal (37°41'N, 8°05'W). At about 85,000 ha, this is the main area of agro-  
116 steppe in Portugal – a semi-natural habitat created by low intensity farming practices – and  
117 one of the main strongholds for several threatened farmland bird species in Western Europe  
118 (Moreira *et al.* 2007; BirdLife International 2021). Land use within the study area has  
119 remained relatively stable in recent decades, in part due to the implementation of agri-  
120 environmental policy schemes and funding mechanisms that ensure high-quality foraging  
121 habitat for many farmland birds (Catry *et al.*, 2013; Silva *et al.*, 2018). Although potential  
122 changes in prey availability were not assessed across the study period, habitat stability  
123 suggests no major changes in prey composition and abundance. Mixed-species colonies are  
124 mostly found in artificial nesting structures provided by LIFE conservation projects target at

125 Lesser Kestrels during the early 2000s (Catry *et al.* 2009), and to a lesser extent in abandoned  
126 farmland buildings, with nests inside wall cavities or under roof tiles. Colonies have been  
127 monitored annually and their composition (number of pairs of each species) has remained  
128 similar in recent years. Within these breeding assemblages, we focused on species that prey  
129 mainly on small vertebrates and arthropods, as these have higher potential for trophic niche  
130 overlap: Western Barn Owl, Little Owl, Common Kestrel, Lesser Kestrel, European Roller  
131 and Spotless Starling. Feral Pigeons and Western Jackdaws are mostly herbivorous and  
132 include a low proportion of animal prey in their diet (Murton & Westwood 1966; Högstedt  
133 1980; Soler *et al.* 1990), hence dietary overlap with the carnivorous or insectivorous species  
134 is expected to be very low. Furthermore, because of incomplete diet information, we would  
135 not be able to correctly interpret their isotopic niches and determine the diet overlap with the  
136 remaining species. Feral Pigeons and Western Jackdaws were, therefore, excluded from this  
137 study.

## 138 **2.2. Sample collection**

139 Feather sampling was carried out during the breeding seasons of 2014, 2016 and 2018 (Table  
140 1). Nestlings of Western Barn Owls (n=12), Little Owls (n=19), Common Kestrels (n=33),  
141 Lesser Kestrels (n=36), European Rollers (n=36), and Spotless Starlings (n=12) were  
142 sampled in their nests at the age of 3-4 weeks, when breast feathers are well developed.  
143 Approximately 3-5 breast feathers were plucked from one nestling per nest (keeping the  
144 feather base intact) and stored in separate plastic zip bags. Blood sampling of parent-  
145 offspring pairs – one parent and one nestling – of Little Owls (n=7), Common Kestrels (n=8),  
146 Lesser Kestrels (n=11), European Rollers (n=6), and Spotless Starlings (n=6) was carried out  
147 during the 2019 breeding season (Table 1). Parents and their offspring were sampled within a  
148 narrow temporal window ( $6.9 \pm 6.1$  days) at their nests. Approximately 150  $\mu$ L of blood was  
149 collected from the brachial vein and preserved in separate vials. The blood was centrifuged

for 10 min at 3400 rev/min within 3 hours of collection, to separate plasma from red blood cells, and plasma samples were frozen before being further prepared for stable isotope analysis (red blood cells were not analysed). We were unable to sample enough Western Barn Owl adults, and so parent-offspring segregation was not investigated for this species. Additionally, to aid the interpretation of stable isotope data, we sampled key prey types of each species in mixed-species colonies throughout the 2018 breeding season (Table 2). Prey species were identified and collected inside or near the nests and were kept frozen until processed for isotope analysis. All work involving bird handling and sample collection was approved by the Instituto de Conservação da Natureza e Florestas (ICNF, permits 547/2018, 427/2019).

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### 161 **2.3. Stable isotope analysis**

Stable isotope ratios have been widely used as dietary tracers to assess different habitats and dietary resources used by consumers, providing a proxy for species or individual ecological niches using minimally invasive methods (Forero *et al.* 2004; Catry *et al.* 2019; Dehnhard *et al.* 2020). Stable isotope analysis of nitrogen ( $\delta^{15}\text{N}$ ) and carbon ( $\delta^{13}\text{C}$ ) provides information on the trophic position of an organism in the food chain and on habitat use, respectively, allowing researchers to simultaneously examine dietary niche segregation/overlap of multiple species or individuals within assemblages and make inference about competition for resources (Inger & Bearhop 2008; Alonso *et al.* 2012; Catry *et al.* 2016b; Reisinger *et al.* 2020; Jenkins and Davoren, 2020). Stable isotope analysis also provides information on the dietary niche at multiple temporal scales depending on the tissue sampled, as isotope ratios reflect the diet during which that tissue was metabolically active.



173 Here, we use feathers from nestlings of the six study species to examine isotopic breadth and  
174 overlap among species, as they offer information for the period during which the feather was  
175 being formed, adequately characterising the diet of the nestlings during the breeding season  
176 for the feather growing period (Inger & Bearhop 2008; Weiss *et al.* 2009). Feathers were  
177 washed in double baths of 0.25N sodium hydroxide solution alternated with baths of double  
178 distilled water to remove any surface contaminants, and were then air dried and then cut into  
179 small fragments (Catry *et al.* 2008). To investigate parent-offspring dietary segregation, we  
180 used blood plasma. Blood plasma has a higher turnover rate than feathers, reflecting the  
181 averaged diet over shorter periods (days), and thus allows for a temporally aligned  
182 comparison between parent and offspring signatures (Inger & Bearhop 2008; Reisinger *et al.*  
183 2020). Blood plasma from adult birds was additionally used to infer niche overlap between  
184 species in addition to nestling feathers. Information from nestling plasma was only used to  
185 examine parent-offspring segregation, as it does not add information to that obtained with  
186 nestling feathers (sample size was smaller and restricted to one year, and plasma reflects  
187 information on the birds' diet over a much shorter period). Plasma samples of birds and soft  
188 tissues (muscle) of prey were dried in an oven at 60°C for 48h and then homogenised to a  
189 powder. Prey samples were then processed for lipid extraction by immersion in a 2:1  
190 chloroform/methanol solution with a solvent volume three to five times larger than the  
191 sample volume (Logan *et al.* 2008). Samples were then mixed for 30s, left undisturbed for ca.  
192 30 min, further centrifuged for 10 min at 3400 rev/min, and the supernatant containing  
193 solvent and lipids was removed. This process was repeated at least three times (until the  
194 solvent was clear) and samples were then re-dried at 60°C for 24h to remove any remaining  
195 solvent. Between 0.8 and 1.2 mg of each sample (feathers, plasma, and prey) were weighted  
196 and stored in tin cups for stable carbon and nitrogen isotope assays. Isotopic ratios were  
197 determined by continuous-flow isotope ratio mass spectrometry at the Stable Isotopes and

Instrumental Analysis facility of the faculty of Sciences, University of Lisbon. Results are presented conventionally as  $\delta$  values in parts per thousand (‰) relative to the IAEA CH7 and Glucose BCR for  $\delta^{13}\text{C}$ , and USGS25 and USGS35 for  $\delta^{15}\text{N}$ . The precision of the isotope ratio analysis, calculated using values from six to nine replicates of laboratory standard material (casein) interspersed among samples in every bath analysis, was 0.04-0.05‰ and 0.02-0.04‰ for  $\delta^{13}\text{C}$  and  $\delta^{15}\text{N}$ , respectively.

#### **2.4 Data analysis**

Feather isotopic signatures for each species from all years were pooled together as the relative isotopic position of each species remained similar (Table 1; Fig. S1).

Differences in isotopic signatures among species were examined using pairwise multivariate analyses of variance performed on nitrogen and carbon simultaneously. Then analyses of variance were performed on carbon and nitrogen separately, followed by pairwise comparisons between each pair of species. Each set of analyses was performed for both nestling feathers and adult plasma. Non-parametric tests were used for nestlings (PERMANOVA, Kruskal-Wallis, and Wilcoxon rank sum tests), and parametric tests for adults (MANOVA, ANOVA, and Tukey tests), after assessing data normality.

Prey signatures were only used to broadly characterise the birds' diet and help interpret differences and similarities in the isotopic niches of the different consumer species, i.e., without quantitatively assessing the contribution of each prey type (Jackson & Parnell 2020).

Prey items with similar isotopic signals were pooled within each taxonomic order (SIA group, Table 2), and then plotted in the isotopic biplot graph to aid the interpretation of consumer signatures using trophic enrichment factors (TEF). Due to lack of reported discrimination factors for all consumer species, we used the mean ( $\pm$  SD) value of trophic discrimination factors for feather and plasma samples of other bird species reported in

published studies reviewed by Caut *et al.* (2009; feathers:  $\delta^{13}\text{C} = 2.16 \pm 1.53$ ,  $\delta^{15}\text{N} = 2.84 \pm 1.14$ ; plasma:  $\delta^{13}\text{C} = -0.08 \pm 0.85$ ,  $\delta^{15}\text{N} = 2.82 \pm 0.31$ ).

To measure isotopic niche breadth of each species and estimate the degree of niche overlap between species, corrected and Bayesian standard ellipse areas ( $\text{SEA}_\text{C}$  and  $\text{SEA}_\text{B}$ , respectively) were calculated using the SIBER package for R (Jackson & Parnell 2020).  $\text{SEA}_\text{B}$  were used to quantitatively compare niche breadth among species, using 1 000 000 iterations, three chains, a burn-in of 1000 and thinning of 10, using a vague normal prior (Jackson & Parnell 2020). Overlap among  $\text{SEA}_\text{C}$  for all pairs of species and for parents and offspring within each species were estimated to reflect the degree of trophic resources shared. For each species (i) in one pair (i,j), a value of overlap ( $\text{Ov}_{[i]}$ ) was calculated as the ratio between the area of overlap between the two  $\text{SEA}_\text{C}$  ( $A_{[i,j]}$ ) and its own  $\text{SEA}_{[i]}$  ( $A_{[i]}$ ), expressed as a proportion ( $\text{Ov}_{[i]} = A_{[i,j]}/A_{[i]}$ ; Catry *et al.* 2016b). Average overlap per species was estimated as the average of all overlaps involving that species. Additionally, to evaluate parent-offspring segregation within each species, differences in Euclidean distances between the centroids of parents and offspring were calculated and tested using a residual permutation procedure based on Turner *et al.* (2010). Afterward, differences in  $\delta^{15}\text{N}$  and  $\delta^{13}\text{C}$  between parents and offspring were compared with t-tests or Mann-Whitney U tests for parametric and non-parametric data, respectively. Assumptions of each test were met prior analysis. All analyses were conducted in the R statistical environment (R Development Core Team, 2016).

### 3. Results

#### 3.1 Stable isotope signatures

Feathers were collected from 148 nestlings, and plasma samples were taken from 38 parent-offspring pairs of six different species; mean ( $\pm$  SD) stable isotope signatures ( $\delta^{13}\text{C}$ ,  $\delta^{15}\text{N}$ ) for each species are presented in Table 1 and Figure 1. A PERMANOVA on  $\delta^{13}\text{C}$  and  $\delta^{15}\text{N}$  of

246 nestling feathers revealed differences among species ( $F_{(5,142)} = 23.618$ ,  $P = 0.001$ ), both in  
 247  $\delta^{13}\text{C}$  and in  $\delta^{15}\text{N}$  (Kruskal-Wallis tests  $\delta^{13}\text{C}$ :  $\chi^2_{(5,142)} = 15.948$ ,  $P = 0.007$ ;  $\delta^{15}\text{N}$ :  $\chi^2_{(5,142)} =$   
 248  $87.323$ ,  $P < 0.001$ ). Post-hoc tests indicated that  $\delta^{13}\text{C}$  only differed between Lesser Kestrels  
 249 and European Rollers ( $P = 0.02$ ). Western Barn Owls and Little Owls fed at the highest  
 250 trophic level (most enriched  $\delta^{15}\text{N}$  values) of the community, significantly higher than all  
 251 other species (all  $P < 0.005$ ). Common Kestrels had more enriched  $\delta^{15}\text{N}$  values than Lesser  
 252 Kestrels and European Rollers, (both  $P < 0.001$ ), but similar values to Spotless Starlings ( $P =$   
 253  $0.257$ ). Spotless Starlings also had more enriched  $\delta^{15}\text{N}$  than Lesser Kestrels ( $P = 0.016$ ) and  
 254 European Rollers ( $P < 0.001$ ) and Lesser Kestrels had higher values than European Rollers ( $P$   
 255  $= 0.012$ ). Regarding the adult community, blood isotopic signatures were only different  
 256 among species for  $\delta^{15}\text{N}$  values (MANOVA:  $F_{(4,33)} = 4.97$ ,  $P < 0.001$ ; ANOVA  $\delta^{13}\text{C}$ :  $F_{(4,33)} =$   
 257  $2.038$ ,  $P = 0.112$ ;  $\delta^{15}\text{N}$ :  $F_{(4,33)} = 10$ ,  $P < 0.001$ ). The trophic rank of adults was similar to that  
 258 of nestlings. Little Owls and Common Kestrels had the highest  $\delta^{15}\text{N}$  values (adult Western  
 259 Barn Owls were not sampled), with Little Owls being in a significantly higher position than  
 260 Lesser Kestrels, European Rollers, and Spotless Starlings (all  $P < 0.001$ ), while Common  
 261 Kestrels had only significantly higher  $\delta^{15}\text{N}$  values than Spotless Starlings ( $P$ -values: Lesser  
 262 Kestrel = 0.061, European Roller = 0.068, Spotless Starling = 0.030). Adult Lesser Kestrels,  
 263 European Rollers and Spotless Starlings had similar  $\delta^{15}\text{N}$  values (all  $P > 0.900$ ; Fig. 1B). In  
 264 short, trophic position (according to  $\delta^{15}\text{N}$  values) of nestlings and adults in the community  
 265 was Western Barn Owl = Little Owl > Common Kestrel = Spotless Starling > Lesser Kestrel  
 266 > European Roller; and Little Owl = Common Kestrel > Spotless Starling = Lesser Kestrel =  
 267 European Roller; respectively (Fig.1 and 2).

268 Isotopic signatures of the main prey items (corrected by TEF) are given in Figure 1. The 10  
 269 potential prey groups comprise a broad isotopic space, with mean  $\delta^{13}\text{C}$  values ranging from -  
 270  $27.3 \pm 0.6$  to  $23.2 \pm 0.3$  and mean  $\delta^{15}\text{N}$  from  $3.4 \pm 1.5$  to  $10.9 \pm 3.0$  (Table 2; Fig. 1). The

relative trophic position of the six species and their relation to prey suggest that Common Kestrels, Lesser Kestrels, European Rollers, and Spotless Starlings feed their offspring mostly with Orthoptera and to a lesser extent with Coleoptera and vertebrates, while the opposite occurs in Western Barn Owls and Little Owls (Fig. 1A). Regarding adults, Little Owls and Common Kestrels had closer signatures to vertebrates, while Lesser Kestrels, European Rollers, and Spotless Starlings maintained a lower trophic position.

### 3.2 Interspecific isotopic niche overlap

As observed for the consumers' isotopic space described above, differences in trophic niches were found mainly between species likely including vertebrates in their diet, i.e., Western Barn Owls and Little Owls, and those preying mostly on arthropods, such as Lesser Kestrels, European Rollers, and Spotless Starlings; with Common Kestrels in an intermediate position (Fig. 2). Amongst nestlings, Little Owls had the widest niche breadth ( $SEA_B = 4.352$ ; Table 3), only slightly overlapping with Common Kestrels and Spotless Starlings (23% and 12%, respectively; Table S1) but fully overlapping with Western Barn Owls ( $SEA_B = 0.755$ ; Table 3; Fig. 2A). Common Kestrels and Spotless Starlings also showed a high overlap with each other (63% and 75%) as well as with Little Owls (59% and 37% respectively; Table S1). Lesser Kestrel nestlings had the smallest niche breadth ( $SEA_B = 0.505$ ; Table 3) which highly overlapped with the dietary niche of European Rollers, Common Kestrels, and Spotless Starlings (60%, 40% and 60%, respectively; Table S1). Amongst adults, Lesser Kestrels, European Rollers, and Spotless Starlings showed high niche overlap, while Common Kestrels and Little Owls were clearly segregated (Fig. 2B; Table S1).

### 3.3 Parent-offspring segregation

Differences between adults and nestlings were found in  $\delta^{15}\text{N}$  values for every species except Little Owls (Little Owl:  $w_{(12)} = 38$ ,  $P = 0.097$ ; Common Kestrel:  $w_{(14)} = 68$ ,  $P = 0.001$ ; Lesser Kestrel  $w_{(20)} = 107$ ,  $P = 0.001$ ; European Roller:  $t_{(10)} = 2.753$ ,  $P = 0.025$ ; Spotless Starling:  $w_{(10)} = 34$ ,  $P = 0.009$ ). No differences in  $\delta^{13}\text{C}$  values were found for any species (Little Owl:  $t_{(12)} = -1.961$ ,  $P = 0.075$ ; Common Kestrel:  $t_{(14)} = 0.381$ ,  $P = 0.709$ ; Lesser Kestrel:  $t_{(20)} = 1.256$ ,  $P = 0.228$ ; European Roller:  $t_{(10)} = 0.115$ ,  $P = 0.115$ ; Spotless Starling:  $t_{(10)} = -0.083$ ,  $P = 0.936$ ). Despite this, Euclidean distance between centroid location of adults and nestlings was significantly different in all species (all  $P < 0.01$ ; Table S2), indicating a high parent-offspring segregation in all of them (Fig. 3).

## 4. Discussion

Identifying the dietary niche of species within assemblages is crucial to understand the mechanisms allowing the coexistence of potential competitors in a specific area and design adequate conservation management actions for mixed-species colonies. Here, we used carbon and nitrogen stable isotope ratios from nestlings and from parent-offspring pairs to provide a detailed assessment of the dietary resources used by a group of six bird species breeding in dense mixed-species colonies established as the result of an extensive nest-provisioning program (Catry *et al.* 2009; Catry & Catry, 2019). Overall, our results show a high intraspecific (parent-offspring) but low interspecific dietary segregation in these assemblages, suggesting a high potential for interspecific competition for resources during the breeding season.

### 4.1 Interspecific overlap in isotopic niches

317 Identical isotopic signatures and highly overlapping niches suggest that species breeding in  
318 these assemblages rely on similar prey to feed their offspring. As reported in previous studies,  
319 the low  $\delta^{15}\text{N}$  isotopic signatures of Lesser Kestrel and European Roller nestlings likely reflect  
320 a diet based on Orthoptera such as grasshoppers and bush-crickets (> 50% and 70% of all  
321 prey delivered to Lesser Kestrel and European Roller nestlings in the study area, respectively;  
322 Catry *et al.* 2016a, 2019). Thus, the high overlap observed between the two species was  
323 expected. Little Owl nestlings had the largest isotopic niche breadth in our study, overlapping  
324 with Common Kestrels but segregating from Lesser Kestrels and European Rollers. Common  
325 Kestrels and Little Owls have more generalist diets that include vertebrates but may expand  
326 their niche to consume more abundant prey such as beetles and grasshoppers, especially  
327 during chick rearing (Tomé *et al.* 2008; Orihuela-Torres *et al.* 2017). Remains of birds,  
328 amphibians, reptiles, and small mammals, but also of Orthoptera, were frequently found in  
329 Little Owl and Common Kestrel nests in the study area during chick rearing, confirming their  
330 eclectic diet. Nestlings of Western Barn Owls had the most enriched nitrogen signatures of  
331 the assemblage, but their dietary niche breadth suggests a more specialised, vertebrated-based  
332 diet (Jaksic *et al.* 1982; Riegert *et al.* 2021; authors pers.obs), segregating from all other  
333 species except Little Owls. Although there is no previous information on Spotless Starling  
334 diet in our study area, in general they rely mostly on small insects such as hoverflies, ants,  
335 and larvae of Coleoptera and Lepidoptera (Motis *et al.* 1997), which should give them a  
336 distinct isotope signature. However, we often observed Spotless Starlings feeding their  
337 offspring with grasshoppers, suggesting a similar diet and potentially explaining the observed  
338 niche overlap with the remaining community.

339 Isotopic characterisation of the adult community indicated a similar trophic rank, although  
340 the information retrieved from adult plasma is probably not representative of the whole  
341 breeding season, and results are based on few individuals in a single year. Adult Lesser

342 Kestrels and European Rollers showed a high overlap with Spotless Starlings, suggesting  
343 similar diets, while Common Kestrels and Little Owls had more enriched  $\delta^{15}\text{N}$  values, likely  
344 reflecting the consumption of vertebrates.

#### 345 ***4.2 Parent-offspring dietary segregation***

346 Despite the lack of significant differences when comparing  $\delta^{15}\text{N}$  and  $\delta^{13}\text{C}$  values between  
347 Little Owl adults and nestlings separately, differences in Euclidean distance suggests that all  
348 species studied in these mixed colonies (including Little Owls) exhibited clear parent-  
349 offspring dietary segregation during the chick-rearing period. During a season of high energy  
350 demand, parents are expected to feed their offspring with high energy items, keeping less  
351 profitable resources for themselves (Orłowski *et al.* 2014). While this was already reported  
352 for Lesser Kestrels and European Rollers (Catry *et al.* 2016a, 2019), we could not find studies  
353 addressing parent-offspring segregation in the other studied species. This study reveals that  
354 Little Owls, Common Kestrels and Spotless Starlings also exhibit marked parent-offspring  
355 segregation in trophic resources, reinforcing that this may be a common evolutionary strategy  
356 to avoid intraspecific competition.

#### 357 ***4.3 Niche theory applied to mixed-species colonies in an artificial nesting landscape***

358 If on one hand the competitive pressure among species breeding together in dense  
359 aggregations can be alleviated through parent-offspring partitioning in their food resources,  
360 on the other hand the low dietary segregation among species can lead to an increase of  
361 interspecific competitive interactions (Catry & Catry 2019).

362 These results raise questions as to the mechanisms allowing species coexistence. According  
363 to ecological niche theory, species with high dietary overlap may still partition their niche  
364 and avoid interspecific competition through spatial or temporal segregation in habitat  
365 utilisation (Pianka 1981; Reisinger *et al.* 2020). In the mixed-species colonies studied, there



366 is no evidence for temporal or spatial segregation, as the breeding phenology and chick  
367 rearing period of all species coincided, and as central-place foragers, all species hunted in the  
368 vicinity of the colonies (Casagrande *et al.* 2008; Catry *et al.* 2012; Šalek & Lövy 2012; Catry  
369 *et al.* 2017a). However, different species could still select different habitats or have hunting  
370 strategies that would foster niche segregation, an aspect that should be further investigated.

371 Alternatively, high niche overlap among sympatric species may reflect high availability of  
372 resources (Pianka 1981; Charter *et al.* 2018; Jenkins & Davoren 2020;). In the Castro Verde  
373 SPA, traditional farming practices associated with agro-steppes are maintained through  
374 specific agri-environmental policy schemes, ensuring high-quality foraging habitats for many  
375 farmland species (Catry *et al.* 2013, 2017a; Silva *et al.* 2018). Large areas of low-intensity  
376 grazed fallows support a high abundance of Orthoptera, and the chick-rearing period of  
377 Lesser Kestrels and European Rollers (and likely of the whole species assemblage except  
378 Western Barn Owls) coincides with the peak abundance of grasshoppers (Catry *et al.* 2016a,  
379 2017b, 2019). From mid-May, the abrupt increase in grasshopper availability (Catry *et al.*  
380 2017b), may allow species to contract their dietary niche to focus on temporarily super-  
381 abundant prey, increasing their overlap without necessarily increasing interspecific  
382 competition (Forero *et al.* 2004; Jenkins & Davoren 2020). The lack of differences in  
383 productivity between European Rollers breeding solitary and in mixed-species colonies, or  
384 between Lesser Kestrels in small or large colonies (Table S3), suggests that prey depletion  
385 around colonies in our study area is unlikely and that food availability is high enough to  
386 allow species with similar niches to coexist. This also suggest the study area might be able to  
387 support additional colonies, but this could imply other conservation implications, such as  
388 increasing these species' reliance on artificial nests and the financial costs associated with it  
389 (Gameiro *et al.* 2020).

390 While we found high trophic overlap among species within the studied community, we lack  
391 evidence to understand whether this may constrain the coexistence of these mixed-species  
392 colonies under some circumstances, an issue that is of particular importance for species of  
393 conservation concern, such as Lesser Kestrel and European Roller. Future studies could  
394 address how species coexist in areas with low resource availability, which are predicted to  
395 increase with ongoing human and climate-induced habitat changes (Tschardtke *et al.* 2005;  
396 Marcelino *et al.* 2020; Raven & Wagner 2021). Our study highlights the need to consider the  
397 potential niche overlap between target and sympatric species, especially when adopting  
398 conservation measures that may alter the composition of communities, such as those based on  
399 nest-site provisioning.

400

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421

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585 **Tables**

586 **Table 1:** Carbon ( $\delta^{13}\text{C}$ ) and nitrogen ( $\delta^{15}\text{N}$ ) stable isotope signatures (mean  $\pm$  SD) of six bird  
 587 species breeding in mixed colonies in Castro Verde, Southern Portugal. Nestling feathers and  
 588 blood (plasma) from parent-offspring pairs were collected during the breeding seasons of  
 589 2014-2018 (pooled) and 2019, respectively. n = sample size.

	Feathers (nestlings)				Blood		
	$\delta^{13}\text{C}$	$\delta^{15}\text{N}$	n		$\delta^{13}\text{C}$	$\delta^{15}\text{N}$	n
Western Barn Owl	$-23.42 \pm 0.58$	$10.64 \pm 0.45$	12	-	-	-	-
Little Owl	$-23.81 \pm 1.07$	$10.08 \pm 1.37$	19	Nestling	$-25.21 \pm 0.45$	$10.03 \pm 1.11$	7
				Adult	$-25.63 \pm 0.34$	$11.15 \pm 0.74$	7
Common Kestrel	$-23.64 \pm 0.55$	$8.88 \pm 1.05$	33	Nestling	$-25.28 \pm 0.34$	$8.77 \pm 0.63$	8
				Adult	$-25.20 \pm 0.44$	$10.07 \pm 0.47$	8
Lesser Kestrel	$-23.78 \pm 0.25$	$7.76 \pm 0.66$	36	Nestling	$-25.55 \pm 0.19$	$7.93 \pm 0.76$	11
				Adult	$-25.38 \pm 0.40$	$8.94 \pm 0.91$	11
European Roller	$-23.52 \pm 0.42$	$7.34 \pm 0.69$	36	Nestling	$-25.42 \pm 0.26$	$7.30 \pm 0.67$	6

Spotless Starling	$-23.84 \pm 0.52$	$8.54 \pm 0.89$	12	Adult	$-25.17 \pm 0.23$	$8.78 \pm 1.14$	6
				Nestling	$-25.48 \pm 0.26$	$6.62 \pm 1.03$	6
				Adult	$-25.49 \pm 0.20$	$8.61 \pm 1.02$	6

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591 **Table 2:** Carbon ( $\delta^{13}\text{C}$ ) and nitrogen ( $\delta^{15}\text{N}$ ) stable isotope signatures (mean  $\pm$  SD) of main  
592 prey sampled in Castro Verde, Southern Portugal. Taxa within coleoptera and orthoptera  
593 were grouped into SIA groups according to their similarity in isotopic signatures. n = sample  
594 size.

	$\delta^{13}\text{C}$	$\delta^{15}\text{N}$	n	SIA group
<b>Small mammals</b>				
<i>Apodemus sp.</i>	-25.6 $\pm$ 0.8	7.0 $\pm$ 1.9	7	
<i>Mus sp.</i>	-25.3 $\pm$ 0.4	8.1 $\pm$ 0.9	3	
Soricidae	-24.9 $\pm$ 0.3	8.2 $\pm$ 1.6	2	
<b>Birds</b>				
<i>Coturnix sp.</i>	-26.1 $\pm$ 0.3	7.2 $\pm$ 0.5	2	
<i>Sturnus unicolor</i>	-25.3 $\pm$ 0.6	8.5 $\pm$ 1.2	3	
<b>Reptiles</b>				
<i>Chalcides striatus</i>	-25.3 $\pm$ 0.5	8.2 $\pm$ 0.8	8	
<b>Chilopoda</b>				
<i>Scolopendra sp.</i>	-25.9 $\pm$ 0.8	9.0 $\pm$ 0.7	8	
<b>Coleoptera</b>				
Carabidae und.	-25.5 $\pm$ 0.5	10.9 $\pm$ 3.0	3	1
<i>Silpha sp.</i>	-25.5 $\pm$ 0.7	10.8 $\pm$ 3.4	3	1
Tenebrionide und.	-26.3 $\pm$ 0.3	9.7 $\pm$ 0.2	2	1
<i>Amphimallon nigrum</i>	-26.6 $\pm$ 0.8	6.6 $\pm$ 0.9	10	2
<i>Carabus rugosus</i>	-26.7 $\pm$ 0.8	5.9 $\pm$ 1.0	2	2
<i>Chrysolina bankii</i>	-27.1 $\pm$ 0.2	7.0 $\pm$ 1.0	3	2
<i>Pterostichus ebenus</i>	-25.9 $\pm$ 0.1	7.0 $\pm$ 0.2	2	2
<i>Sepidium sp.</i>	-26.8 $\pm$ 0.8	6.5 $\pm$ 0.7	8	2
<i>Bubas bison</i>	-27.3 $\pm$ 0.6	8.9 $\pm$ 0.4	5	3
<i>Netocia sp.</i>	-23.2 $\pm$ 0.3	5.6 $\pm$ 3.7	2	4
Protaetia sp.	-23.9	7.4	1	4
<b>Orthoptera</b>				
Acrididae	-26.4 $\pm$ 0.7	3.8 $\pm$ 1.0	12	5
Tettigonidae	-25.3 $\pm$ 0.7	3.4 $\pm$ 1.5	14	5
<i>Gryllotalpa sp.</i>	-27.2 $\pm$ 0.7	6.4 $\pm$ 1.0	7	6

595

596

597 **Table 3:** Isotopic niche breadth of nestlings (feathers) and adults (plasma) from bird species  
598 breeding in mixed colonies. Bayesian standard ellipse areas (SEAb, with 95% credible  
599 intervals) and sample size-corrected standard ellipse areas (SEAc) are shown. *n* = sample  
600 size.

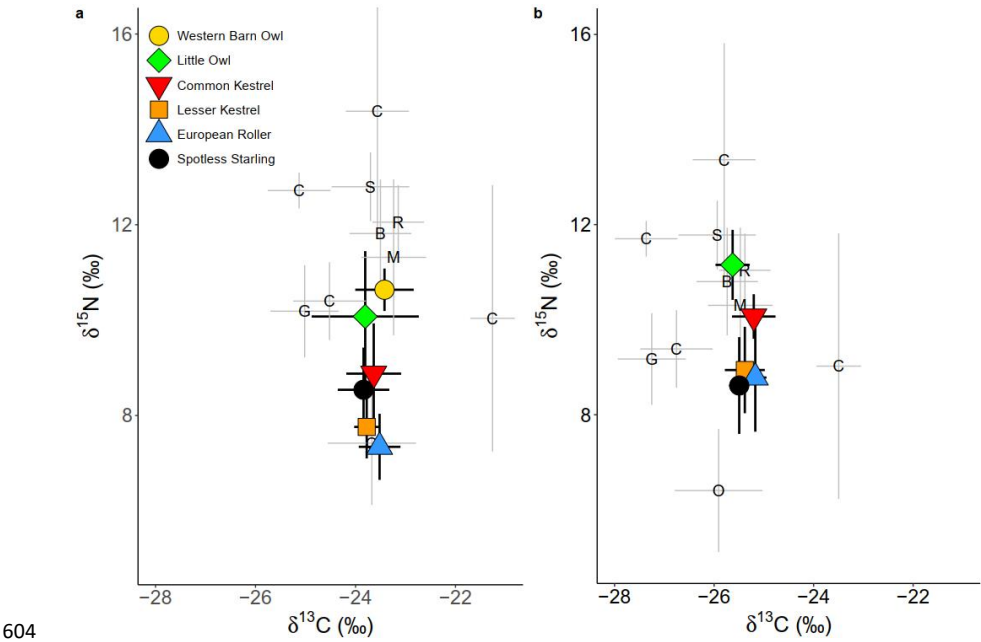
Consumers	n	SEAb (95% credible interval)	SEAc
<b>Nestlings (feathers)</b>			
Western Barn Owl	12	0.755 (0.439-1.451)	0.899
Little Owl	19	4.352 (2.824-7.175)	4.848
Common Kestrel	33	1.763 (1.265-2.543)	1.872
Lesser Kestrel	36	0.505 (0.369-0.718)	0.535
European Roller	36	0.830 (0.602-1.173)	0.871
Spotless Starling	12	1.320 (0.770-2.540)	1.575
<b>Adults (plasma)</b>			
Little Owl	7	0.688 (0.333-1.682)	0.922
Common Kestrel	8	0.558 (0.290-1.297)	0.736
Lesser Kestrel	11	1.058 (0.590-2.064)	1.254
European Roller	6	0.578 (0.274-1.592)	0.785
Spotless Starling	6	0.532 (0.248-1.446)	0.764

601

602 **Figure legends**

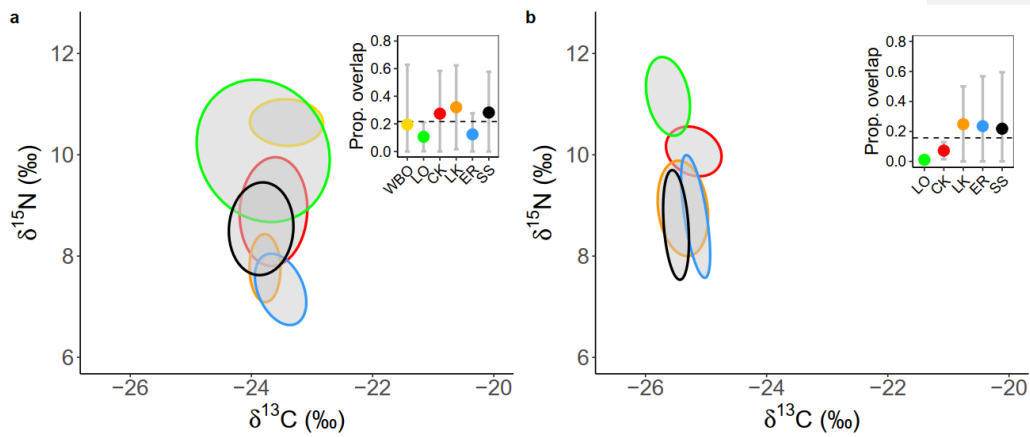
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**Commented [RF1]:** update legend to full species names, and capitalise each word

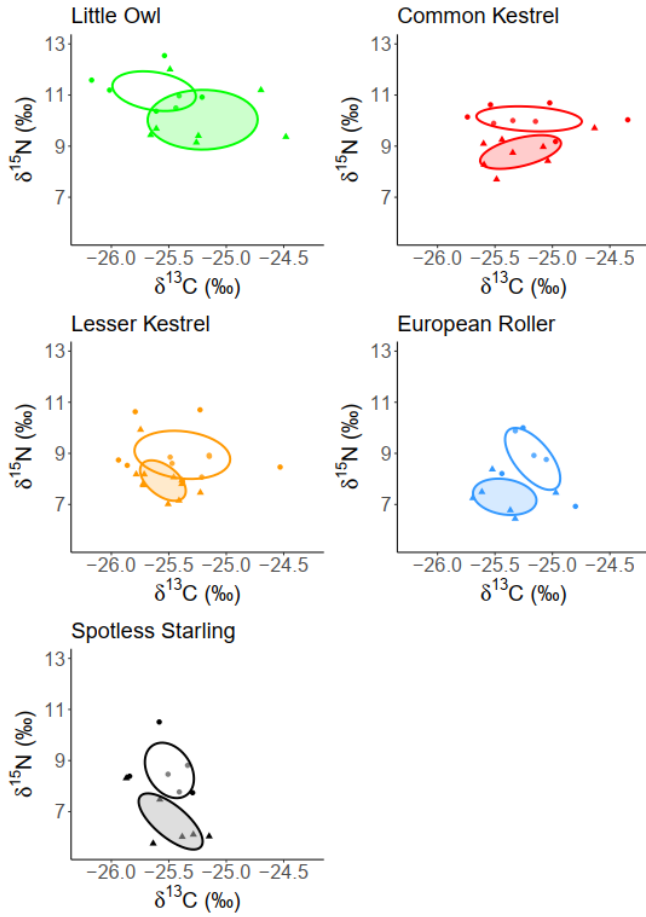


604

605 **Fig. 1:** Carbon ( $\delta^{13}\text{C}$ ) and nitrogen ( $\delta^{15}\text{N}$ ) stable isotope signatures (mean  $\pm$  SD) of nestlings  
606 (A) and adults (B) from six bird species breeding in mixed colonies established in an  
607 artificial nesting landscape in Southern Portugal. Isotope signatures of nestlings and adults  
608 were obtained from feathers (collected in 2014, 2016, and 2018, pooled) and blood plasma  
609 (collected in 2019), respectively. Signatures of main prey items are shown. M, small  
610 mammals; B, birds; R, reptiles; S, scolopendra; C, coleoptera; O, orthoptera; G,  
611 *Gryllotalpa*. Stable isotope ratios of prey items were obtained from muscle/soft tissue and  
612 corrected for the trophic discrimination factor (based on Caut et al. 2009, see methods).  
613 Because different tissues were used for nestlings and adults, direct comparisons in trophic  
614 position between the two age groups should be avoided.



**Fig. 2:** Overlap in isotopic niches (corrected standard ellipses; SEAc) among nestlings (A) and adults (B) from six bird species breeding in mixed colonies established in an artificial breeding landscape in Southern Portugal. WBO = Western Barn Owl, LO = Little Owl, CK = Common Kestrel, LK = Lesser Kestrel, ER = European Roller, SS = Spotless Starling. Isotope signatures of nestlings and adults were obtained from feathers and blood plasma, respectively. Inset plot (upper right corner) shows the mean ( $\pm$ SD) niche overlap for each species and the mean overlap of the community (horizontal dashed line).



**Fig. 3** Overlap in isotopic niches (corrected standard ellipses; SEAc) between parents (clear) and their offspring (filled) of bird species breeding in mixed colonies established in an artificial nesting landscape in Southern Portugal. Stable isotope ratios ( $\delta^{13}\text{C}$  and  $\delta^{15}\text{N}$ ) were measured using blood plasma from 7 Little Owl, 8 Common Kestrel, 11 Lesser Kestrel, 6 Roller, and 6 Starling parent-offspring pairs (one parent and one offspring in each pair). Circles and triangles show individual signatures of parents and nestlings, respectively.