1	Evolution of nestling faeces removal in avian phylogeny
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19	Nest sanitation-related traits have often been explained at the intraspecific level as
20	reducing the probability of infection or detection by predators and parasites, but its
21	evolution within the avian phylogeny is still poorly understood. We compiled detailed
22	information of such traits for more than 400 bird species and, by means of modern

24 removing their offspring's faeces and the production of faecal sacs by nestlings.

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25 Furthermore, because the functional hypotheses used to explain nest sanitation

comparative methodologies, we reconstructed the evolution of adults' contribution to

behaviour assume potential effects of brood size, body mass, nestling period and diet, 26 we explored the association between these traits and those related to nest sanitation in a 27 phylogenetically controlled framework. Our results suggest that parental removal of 28 nestling faeces has driven the evolution of faecal sacs, while the ancestral states 29 involved birds with faecal sacs removed by parents. These results support the long-held 30 idea that faecal sacs facilitate the removal of faeces by parents. Moreover, we found that 31 animal diets and small body sizes have favoured the evolution of faecal sacs suggesting 32 the existence of some chemical and physical constraints in relation to the evolution of 33 the mucous covering. Our results highlight the importance of nest sanitation in the 34 evolution of birds and their life history characteristics. 35

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37 Keywords: birds, faeces, faecal sacs, nest sanitation, parent–offspring relationships

Nest sanitation behaviour is an important and widespread behaviour in birds 38 39 that, despite being known for a long time (Blair & Tucker, 1941; Herrick, 1900; Skutch, 1976; Thomson, 1934), is still poorly understood, particularly regarding its evolution 40 41 (Gow, Wiebe, & Musgrove, 2015; Guigueno & Sealy, 2012; Lang, Straight, & Gowaty, 2002). This is surprising because strategies and traits related to nest sanitation, or even 42 the effort devoted to such activities, have been suggested to help infer levels of selection 43 pressures acting within species-specific nest environments (Ibáñez-Álamo, Ruiz-44 Rodríguez, & Soler, 2014). Recent experimental studies have focused on investigating 45 adaptive values of removal by parents of nestling faeces, the most common form of nest 46 47 sanitation behaviour in birds (Guigueno & Sealy, 2012). Since faeces contain potentially pathogenic microorganisms, its removal would reduce the probability of 48 infection (Ibáñez-Álamo, Ruiz-Rodríguez, et al., 2014). In addition, parasites and nest 49 50 predators might use chemical (i.e. odours) and/or visual cues of nestling faeces to located active nests and, therefore, removing it from nests would reduce the probability 51 52 of nest predation (Ibáñez-Álamo, Ruiz-Raya, Roncalli, & Soler, 2014; Ibáñez-Álamo, Sanllorente, Arco, & Soler, 2013; Petit, Petit, & Petit, 1989; Weatherhead, 1984) and 53 parasitism (Ibáñez-Álamo, Ruiz-Raya, Rodríguez, & Soler, 2016). Nestlings of many 54 species encapsulate faeces within a mucous covering forming faecal sacs (Blair & 55 Tucker, 1941; Herrick, 1900; Thomson, 1934; Weatherhead, 1984). This covering acts 56 as a physical barrier to impede microbial infections of birds (Ibáñez-Álamo, Ruiz-57 Rodríguez, et al., 2014) and might facilitate manipulation and removal of faeces by 58 parents (White 1773, cited in Blair & Tucker, 1941; Blair & Tucker, 1941; Herrick, 59 1900; Pycraft, 1909; Thomson, 1934). Thus, the evolution of faecal sacs should be 60 associated with parental removal of faeces from nests, although this prediction has 61 never been tested. Either parental removal would more easily evolve in species with 62

nestlings producing faecal sacs, or the evolution of faecal sacs would be particularlybeneficial in species in which parents removed them.

Benefits associated with nest sanitation behaviour may depend on ecological and 65 life history characteristics. It is, for instance, possible that the costs associated with the 66 presence of nestling faeces in the nest depend on the volume and contents of faeces. 67 Everything else being equal, negative impacts of small faeces in terms of attracting nest 68 69 predators or vectors for potentially dangerous microorganisms/parasites would be lower than those of larger faeces (Ibáñez-Álamo et al., 2016; Petit et al., 1989). Moreover, if 70 faeces are not removed and accumulate in the nest during nestling development, species 71 72 with longer nestling periods and larger broods would differentially suffer higher costs than species with the opposite life history traits. Therefore, we expect species with 73 longer nestling periods or larger broods to be those with faeces removal. 74

75 Species-specific diet is another element that might have influenced the evolution of nest sanitation-related traits (Guigueno & Sealy, 2012). Diet affects the prevalence of 76 77 potentially pathogenic microorganisms in chicken faeces (Ryu, Park, Bang, Kang, & Hwangbo, 2016) and the presence of animal components in the food of livestock 78 produces faeces with more intense odours than those feeding only from plants (e.g. 79 Mackie, Stroot, & Varel, 1998). Consequently, there are good reasons to think that diet 80 characteristics could affect the costs of not removing faeces from nests in terms of 81 microbial infection or detectability by predators. We explored this possibility by 82 analysing the relationship between diet (i.e. animals or plants) and nest sanitation-83 84 related traits (faeces removal and faecal sac production).

Previous studies on the subject have been focused on exploring within-species
variation in a small number of bird species (e.g. Herrick, 1900; Ibáñez-Álamo et al.,
2016, 2013; Ibáñez-Álamo, Ruiz-Raya, et al., 2014; Ibáñez-Álamo, Ruiz-Rodríguez, et

al., 2014; Petit et al., 1989; Quan, Li, Wang, & Goodale, 2015; Thomson, 1934;
Weatherhead, 1984), but exploring the interspecific associations among traits related to
nest sanitation and ecological and life history characteristics is essential to understand
the evolution of nest sanitation in birds (Gow et al., 2015). Trying to fill this gap, we
investigated these scenarios potentially affecting the evolution of nest sanitation using
information collected from the literature for more than 400 bird species (19 Orders).

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#### 95 Methods

# 96 Data collection

After checking reviews on the topic (Blair & Tucker, 1941; Guigueno & Sealy, 2012; 97 Thomson, 1934), we searched for related articles in the Web of Science and Google 98 99 Scholar by using the following keywords: 'sanitation', 'nest sanitation', 'nest cleaning' and 'f(a)ecal sacs'. We also checked the Handbook of the Birds of the World (Del Hoyo, 100 Elliott, Sargatal, Christie, & de Juana, 2016) for information on nest sanitation-related 101 traits. The literature used for each species is listed in the Supplementary Material (Table 102 103 S1). For each species, we gathered information on (1) whether faeces are removed from 104 nests; we considered a species to show removal when adults and/or offspring in some 105 way avoided its accumulation in the nest. We noted (2) the presence of faecal sacs and 106 whether (3) parents and (4) offspring remove nestling faeces at least during part of the 107 nestling period. If parents removed nestling faeces, we also noted (5) which sex (male, 108 female or both) was responsible. For seven species (Accipiter nisus, Ardea cinerea, Buteo buteo, Chiroxiphia caudata, Falco peregrinus, Haliaeetus albicilla, Ocyceros 109 110 birostris), we completed information for these variables using video recordings 111 available on the Internet (ARKive.org). We did not include precocial species in our data set given that we were interested in the evolution of removing nestling faeces fromnests.

We also collected data on (6) body mass, (7) brood size, (8) nestling period 114 115 duration, (9) diet (animals versus plants) and (10) nesting habits (hole, semihole and open nesters) using the Handbook of the Birds of the World (Del Hoyo et al., 2016). We 116 considered the diet of a species to be of animal origin when the nestlings were fed 117 118 mainly with animals such as invertebrates (i.e. insects), vertebrates (i.e. birds) or carrion. Those species whose nestlings were mainly fed with seeds, fruits or nectar were 119 classified as having plant diets. For those species without specific information on the 120 121 nestling diet, we used the main feeding source for the species. All collected information is reported in Table S2. 122

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# 124 *Statistical analyses*

125 Because the expected interspecific associations may have a strong phylogenetic 126 component (Harvey & Pagel, 1991) we used the phylogenetic association among the bird species considered in our analyses. Phylogenetic relationships were estimated in 127 the Mesquite environment (Maddison & Maddison, 2015) as the consensus (i.e. 128 129 majority rules consensus) tree of 1000 phylogenetic trees downloaded from http://birdtree.org/ (Jetz, Thomas, Joy, Hartmann, & Mooers, 2012; Table S3). The 130 131 predicted associations were subsequently explored with phylogenetically controlled analyses. 132

Most of the characters we considered are of binary nature, including the contribution of adults and nestlings to removing faeces from nests. In most species (349 of 370) the removal of nestling faeces is only performed by parents, but in some, nestlings also contribute (11 species) or even are the only ones in charge of this task by

directly disposing of their own faeces (10 species). Because of the limited number of 137 138 species that do not remove faeces from the nests, or with nestlings participating in this task, we did not include more than one categorical independent factor in our model. As 139 140 we were mainly interested in exploring the evolution of nest sanitation behaviour of adults in relation to nestling traits, we considered the binomial information of whether 141 142 adults remove nestling faeces without the help of their offspring. Information on 143 nestling diet was also compiled as binomial information (mainly animals versus mainly plants). In addition to binomial variables, some statistical models also include 144 continuous independent factors, namely, body mass, nestling period and brood size. 145 146 Thus, given that all dependent factors were of binomial distribution and the need to control phylogenetic relationships, we used a phylogenetic generalized linear mixed 147 model for binary data (binaryPGLMM; Ives & Garland, 2014; Ives & Helmus, 2011) as 148 149 implemented in the R (version 3.2.3. R Core Team, 2016) statistical environment with the appropriate libraries ('ape'; Paradis, Claude, & Strimmer, 2004), 'MASS' (Venables 150 151 & Ripley, 2002) and 'mvtnorm' (Genz & Bretz, 2009). The binaryPGLMM package 152 performs linear regressions for binary phylogenetic data, estimating regression coefficients (hereafter 'estimate') with approximate standard error. At the same time, it 153 estimates the strength of the phylogenetic signal in the residuals (hereafter 's') and gives 154 155 an approximate conditional likelihood ratio test for the hypothesis that there is no signal (Ives & Garland, 2014; Ives & Helmus, 2011). 156

Additionally, we explored possible associations in character evolution and the direction of changes along the phylogenetic tree of several pairs of traits: (1) between parental contribution to removing faeces from nests (alone or not) and nestling production of faecal sacs, (2) between nestling diet (animals or plants) and adult contribution to removing faeces, and (3) between nestling diet and production of faecal

sacs. Briefly, we used Pagel's discrete method to test models of independent and 162 163 dependent evolution (Harvey & Pagel, 1991). This method compares the ratio of likelihood of two models: one of the models where the rates of change in each character 164 165 are independent of the state, and a second model where rates of change depend on the state of the other trait. Since likelihoods associated with each of the eight possibilities of 166 167 transition are estimated, this approach provides a good method to study evolutionary 168 pathways through estimations of transition rates between pairs of binary character states (i.e. test for any effects in Mesquite 3.04). In addition, we also tested for the possibility 169 that character X depended on character Y, or that character Y depended on X. We 170 171 performed these analyses as implemented in Mesquite (Maddison & Maddison, 2015) with 50 ML replicates over 1000 repeated simulations. 172

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## 174 **Results**

## 175 *Faeces removal by adults and presence of faecal sacs*

We collected information on nest sanitation-related characters of 417 species of birds 176 from 19 different Orders. Within the 396 species that remove nestling faeces from their 177 178 nests (94.96% of all species analysed), reliable information on production of faecal sacs 179 and on adult and nestling participation in faeces removal was available for 370 species. 180 Most of these species (96.76%) produce faecal sacs, and in most of them (97.49%) only 181 adults remove faeces from nests. In nests of the 12 species with no faecal sacs, faeces 182 removal is mainly done by nestlings alone (83.33%) although adults contribute in two of them. Another interesting result is that, when adults participate in faeces removal, 183 184 both male and female perform this behaviour in the majority of species (91.9%) and just in a few of them either the female (6.8%) or the male (1.3%) do it exclusively. 185

The study of evolutionary associations among these traits along the phylogenetic 186 187 tree suggests that the evolution of parental contribution to faeces removal from nests depends on the presence of faecal sacs. Nodes with no faecal sacs and only adults 188 189 removing nestling faeces, as well as those with faecal sacs and nestlings able to dispose of their own faeces, evolved to nodes with faecal sacs and only adults removing faeces 190 more than expected by random (Fig. 1). Therefore, the evolutionary acquisition of 191 192 faecal sacs was more likely in nodes with adult removal and no nestling help (difference 193 in log-likelihood = 2.92, P = 0.029), while the trait that only adults remove faeces depends on whether the ancestral species had faecal sacs (difference in log-likelihood = 194 195 17.92, *P* = 0.047; Fig. 1).

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## 197 *Nest sanitation, body mass, brood size and nestling period*

198 Nestling faeces removal was not significantly affected by body mass, brood size or nestling period (all binary PGLMM: P > 0.16). In contrast, faecal sacs were more 199 200 frequent in species of lower body mass (binary PGLMM: estimate (SE) = -3.48 (1.19), Z = 2.93, P = 0.003; phylogenetic signal s2: 6.52, P = 0.001) and shorter nestling period 201 (binary PGLMM: estimate (SE) = -9.69 (3.80), Z = 2.55, P = 0.011: phylogenetic signal 202 s2: 7.12, P = 0.0003). Brood size did not predict the existence of faecal sacs (binary 203 204 PGLMM: P = 0.5221) and, when considering all three life history traits together, body mass (binary PGLMM: P = 0.027), but not the duration of the nestling period (binary 205 PGLMM: P = 0.160) or brood size (binary PGLMM: P = 0.408), reached statistical 206 207 significance explaining faecal sac evolution. When body mass and parental contribution 208 to removing nestling faeces from nests (i.e. with no help from nestlings) were 209 considered together, the presence of faecal sacs tended to be more common in species in which only adults remove faeces (binary PGLMM: P = 0.07), while the effect of body 210

mass reached statistical significance (binary PGLMM: P = 0.025). Moreover, nestlings help parents to remove faeces from nests more often in large species (binary PGLMM: estimate (SE) = 2.11 (0.62), Z = 3.40, P = 0.0007: phylogenetic signal s2: 4.64, P =0.0001). Species that do not remove nestling faeces are similarly distributed among hole (6.3%), semihole (1.9%) and open nesters (4.9%) suggesting that is unlikely that nesting habits affected the evolution of this behaviour.

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#### 218 *Nest sanitation and nestling diet*

The behaviour of removing nestling faeces from the nest was not associated with 219 offspring diet (88.0% of the 50 species with plant diets removed faeces while 95.9% of 220 the 363 species with animal diets did so; binary PGLMM: estimate (SE) = 0.01 (0.83), Z 221 = 0.01, P = 0.99: phylogenetic signal s2: 3.56, P < 0.0001). Nestlings of species feeding 222 223 on plants more often produced faeces within a mucous covering than those eating 224 mainly animals (100% of the 32 species with plant diets produced faecal sacs while 225 96.2% of the 313 species with animal diets did so; binary PGLMM: estimate (SE) = -226 7.56 (3.38), Z = 2.24, P = 0.025), even after controlling for the effect of body mass (binary PGLMM: estimate (SE) = -4.31 (0.97), Z = 4.45, P < 0.0001; phylogenetic 227 signal of the model:  $s_2 = 2.29$ , P < 0.0001). In contrast, species-specific nestling diets 228 229 were not associated with parental removal of faeces (binary PGLMM: estimate (SE) = 1.88 (1.72), Z = 1.09, P = 0.27), but were associated with body mass (binary PGLMM: 230 estimate (SE) = 2.25 (0.65), Z = 3.44, P = 0.0005; phylogenetic signal of the model: s2 231 = 5.013, P < 0.0001). Results from correlations of the evolution of the characters 232 considered showed that the production of faecal sacs (Fig. 2), but not parental 233 contribution to remove faeces from nests (differences in log-likelihood = 2.62, P =234 0.23), tended to be associated with nestling diet. We found evidence supporting the 235

influence of nestling faecal sacs on diets (differences in log-likelihood = 5.76, P < 0.0001) and of diets on faecal sacs (differences in log-likelihood = 4.31, P = 0.04). Thus, diet is likely to be responsible for the evolution of faecal sacs but probably not for the role that adults play in nest sanitation behaviour.

240

#### 241 **Discussion**

Adults of most altricial bird species remove their nestling faeces from their nests, and 242 these faeces are usually encapsulated in a mucous covering that reduces the probability 243 244 of bacterial infection (Ibáñez-Álamo, Ruiz-Rodríguez, et al., 2014) and, apparently, facilitates faecal handling by parents (White 1773, cited in Blair & Tucker, 1941; Blair 245 & Tucker, 1941; Herrick, 1900; Pycraft, 1909; Thomson, 1934). Furthermore, there 246 247 seems to be no differences between males and females as both sexes perform this behaviour in most species in which adults participate in the removal of nestling faeces. 248 Our results show that these traits (adult removal and faecal sacs) are almost completely 249 fixed in altricial birds and much more widespread than previously thought (Blair & 250 Tucker, 1941; Skutch, 1976; Thomson, 1934), which suggests that parental participation 251 252 in cleaning the nest is critical for offspring development. Nest sanitation is an essential component of parental behaviour, but its functionality has only been studied 253 254 intraspecifically (Carere & Alleva, 1998; Dell'Omo, Alleva, & Carere, 1998; Gow et al., 2015; Lang et al., 2002; Thomson, 1934). To our knowledge, this is the first 255 256 interspecific study on the topic and the only one looking at the evolution of nest sanitation within the avian phylogeny. Our main findings are (1) the existence of 257 258 correlated evolution between adult removal of faeces and the production of faecal sacs by nestlings and (2) the detection of significant associations between life history 259 characteristics (i.e. nestling period and body mass), the evolution of faecal sacs and 260

parental contribution to removing faeces from nests. Finally, we also found (3) evidence suggesting a role of nestling diet on the evolution of faecal sacs. All these results considered together suggest that nest sanitation behaviour of adults, together with nestling diet, have influenced the evolution of the structure of nestling faeces, particularly in relation to the presence of the mucous covering (faecal sacs). Below we discuss possible evolutionary implications of our findings.

Altriciality is considered the derived character of precocial ancestors (Starck & 267 Ricklefs, 1998) and selection pressures favouring the evolution of nest sanitation would 268 be higher in altricial species whose nestlings develop within the nest environment 269 270 (Guigueno & Sealy, 2012). Because we were interested on the evolution of removing nestling faeces from nests, we did not include precocial species in our data set (apart 271 from one semiprecocial species, Antigone antigone) and thus cannot discuss the 272 273 possibility of correlated evolution between nest sanitation and altriciality. However, 274 since adults removing nestling faeces and nestlings producing faecal sacs were the 275 ancestral states (Fig. 1), we speculate that these traits had already evolved in precocial 276 bird species. Nest sanitation behaviours occur not only during the nestling phase, but also during the egg incubation stage (Guigueno & Sealy, 2012), for instance renovating 277 green materials (Petit, Hossaert-McKey, Perret, Blondel, & Lambrechts, 2002) or 278 279 removing broken eggs or eggshells from the nest (Soler et al., 2011; Tinbergen et al., 1962). Associated benefits of nest sanitation should be higher during the nestling period 280 and thus these traits would rapidly become fixed in altricial bird populations. However, 281 282 that the presence of faecal sacs is the ancestral state and that we only found them in altricial species in our literature search also suggest that the mucous covering might 283 have evolved in relation to altriciality. 284

We found evidence of correlated evolution between parental removal of faeces 285 286 and nestling production of faecal sacs. Faecal sacs mainly evolved in species in which adults remove faeces from their nests without help of their nestlings, while faeces 287 288 removal exclusively by adults evolved mainly in species with faecal sacs (Fig. 1). These associations would be explained by faecal sacs facilitating faeces removal by parents 289 (White 1773, cited in Blair & Tucker, 1941), an idea already mentioned in the 18th 290 century and now widely accepted by the scientific community (Blair & Tucker, 1941; 291 292 Herrick, 1900; Pycraft, 1909; Thomson, 1934) even though it has not previously been demonstrated. Our results offer the first demonstration of this hypothesis and we 293 294 encourage future studies to investigate whether predictions of the evolutionary pattern found here fit the ecological context by, for example, exploring intraspecific covariation 295 in faecal sac resistance and the efficacy of parents removing nestling faeces. 296

297 We found no evidence supporting the expected associations between nestling 298 faeces removal and life history characteristics (see above), but body size and duration of 299 the nestling period were negatively associated with the presence of faecal sacs. We 300 predicted the opposite direction, however, because nest sanitation tasks should be more beneficial for larger species (i.e. higher production of faeces) and for those with longer 301 nestling periods (higher accumulation of faeces). When considering both variables in 302 303 the same model, only body mass reached statistical significance. Explanations of these 304 results might be related to physical constraints of the mucous covering: above a certain weight it may not guarantee the integrity of faecal sacs during parental removal. New 305 306 data are necessary to test this and other possible scenarios explaining the detected association between body mass and faecal sac production. Independently of the reason, 307 308 our results show that nestlings participated in nest sanitation tasks more often in large species (see Results), which may be related to the difficulty adults have removing faeces 309

with a weak, or no, mucous covering. Thus, because of the adaptive value of nest
sanitation (Kölliker, Royle, & Smiseth, 2012), nestlings of large species should directly
dispose of their own faeces from nests more often than those of small species.

313 We also argued that nestling diet should have affected the evolution of nest sanitation behaviour and we found partial support for this hypothesis because faecal 314 315 sacs are more likely to evolve in species with nestlings feeding mainly on vegetal 316 material (Fig. 2). Diet, however, does not seem to have affected the evolution of parental sanitation behaviour. Moreover, faecal sacs appeared less often in species with 317 nestlings mainly fed with animals. This is the most common source of food in birds 318 (72.4%; calculated from Burin, Kissling, Guimarães, Şekercioğlu, & Quental, 2016), 319 and, above, we proposed various scenarios in which the mucous covering could provide 320 fitness advantages for species with animal diets. We found the opposite result, however, 321 322 since faecal sacs appear more often in species with vegetal diets. We could speculate 323 that animal diets favour faeces with some specific components or in such concentrations 324 (i.e. uric acids) that might negatively interact with the mucous covering, but our results 325 do not allow us to discuss this or other scenarios (i.e. detectability by predators and/or parasites). Thus, although our results emphasize the importance of nestling diet for the 326 evolution of nest sanitation behaviour (i.e. faecal sacs), further investigation is 327 necessary to detect the underlying causes. 328

Summarizing, our comparative analyses support the hypothetical correlated evolution of parent and offspring traits related to nest sanitation, and emphasize the importance of nest sanitation in the evolution of birds in general and their life history characteristics in particular. We hope that our findings encourage further research directed to explore functional hypotheses of the detected evolutionary patterns.

334

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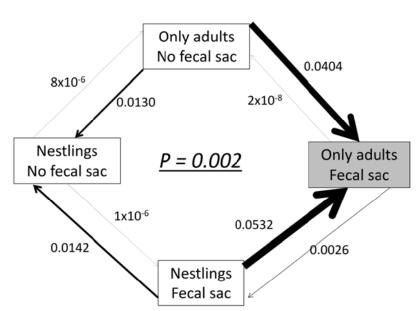
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# 462 Figure legends

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471 Fig 1: Flow diagram showing correlated evolution between the production of faecal sacs by nestlings (faecal sac versus no faecal sac) and faeces removal by adults (only adults 472 versus nestlings). The number associated with each arrow represents the maximum 473 474 likelihood value and, together with arrow thickness, indicates which transition is expected to be the most common. Probability (P) in the centre of the diagram indicates 475 476 the probability of correlated evolution between the binary traits after 10 ML replicates and 1000 repeated simulations. Parsimony ancestral states of characters are shown in 477 the grey box. 478

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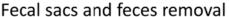
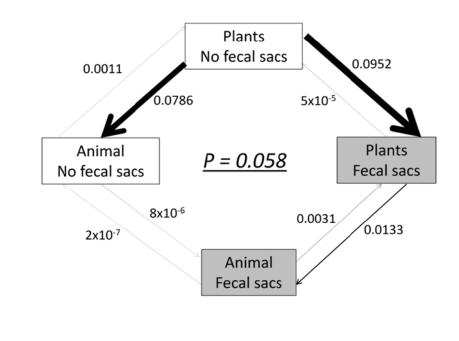


Fig 2: Flow diagram showing correlated evolution between the production of faecal sacs 480 by nestlings (faecal sac versus no faecal sac) and diet (animals versus plants). The 481 number associated with each arrow represents the maximum likelihood value and, 482 together with arrow thickness, indicates which transition is expected to be the most 483 common. Probability (P) in the centre of the diagram indicates probability of correlated 484 evolution between the binary traits after 10 ML replicates and 1000 repeated 485 simulations. Parsimony ancestral states of characters are shown in grey boxes. 486 481

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Fecal sacs and diets

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