1	On the Evolution of Extreme Structures: Static Scaling and the Function of Sexually
2	Selected Signals
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31 Introduction

32 Understanding how morphology scales with body size is one of the most 33 pervasive topics in organismal biology (Dial, Greene, & Irschick, 2008; Gould, 1966, 34 1974b, 1974a; J. Huxley, 1932; Schmidt-Nielsen, 1984; Templeton, Greene, & Davis, 35 2005; Thompson, 1917; Voje, 2016; West & Brown, 2005; West, Brown, & Enquist, 36 1997). The reason for this is simple - virtually every measurable aspect of an organism 37 scales with body size. Some relationships hold across hundreds of species, spanning 38 multiple orders of magnitude in overall size (e.g., Kleiber's Law (Kleiber, 1932); 39 Rubner's Surface Rule (Rubner, 1883; Von Bertalanffy, 1957); Cope's Rule (Stanley, 40 1973); Rensch's Rule (Abouheif & Fairbairn, 1997; Wolf U. Blanckenhorn, Meier, & 41 Teder, 2007; Fairbairn, 1997)). Others account for transformations in shape arising 42 during ontogeny (e.g., brain/body weight (Cock, 1966; Gould, 1974a, 1977); Dyar's Law 43 (Dyar, 1890)). Here we focus on 'static' allometry, scaling that occurs among individuals 44 of the same age sampled from within populations (Cheverud, 1982; sensu Cock, 1966; 45 Pélabon et al., 2013).

46 Perhaps the most striking pattern in the study of static scaling is the observation 47 that many extreme products of sexual selection – ornaments of choice and weapons of 48 intrasexual competition – scale steeply with body size (Bonduriansky & Day, 2003; 49 Eberhard, 1998; Egset et al., 2012; Emlen, 1996; Emlen & Allen, 2003; Fromhage & 50 Kokko, 2014; Gould, 1974b; Hongo, 2007; Kelly, 2005; Kodric-Brown, Sibly, & Brown, 51 2006; Miller & Emlen, 2010; Painting & Holwell, 2013; Shingleton, Frankino, Flatt, 52 Nijhout, & Emlen, 2007; Shingleton, Mirth, & Bates, 2008; L. W. Simmons & Tomkins, 53 1996; Stern & Emlen, 1999; Voje, 2016; Wilkinson, 1993). Specifically, when examined

on a log scale, the relationship between the size of these structures and body size is
greater than one ('positive allometry') (Gould, 1966; J. S. Huxley & Teissier, 1936;
Kerkhoff & Enquist, 2009; Shingleton & Frankino, 2013; Voje, 2016). These steep
scaling relationships cause ornaments and weapons to attain extraordinary proportions in
the largest individuals, inspiring descriptions such as 'extreme', 'exaggerated' (Darwin,
1871) and 'bizarre' (Gould, 1974b) (Fig. 1).

60 Early studies of static scaling often focused on the products of sexual selection, 61 including cervid antlers (Gould, 1973; J. Huxley, 1932; Thompson, 1917), fiddler crab 62 (Uca) chelae (J. Huxley, 1932), and beetle (Scarabaeidae) horns (Bateson & Brindley, 63 1892; Paulian, 1935). Since then, hundreds of sexually selected structures have been 64 examined, and the overwhelming majority scale steeply with body size (Emlen, 2008; 65 Emlen & Nijhout, 2000; Knell, Naish, Tomkins, & Hone, 2013b; Kodric-Brown et al., 66 2006; e.g., Otte & Stayman, 1979; Petrie, 1988, 1992; Voje, 2016). In fact, the link 67 between steep scaling and exaggerated ornaments and weapons is so widespread that 68 many consider the steepness of static allometry indicative of the intensity of sexual 69 selection acting on a structure (e.g., stalk-eyed fly (Diopsidae) eyestalks (Baker & 70 Wilkinson, 2001); frog (Anura) forelimbs (Schulte-Hostedde, Kuula, Martin, Schank, & 71 Lesbarrères, 2011); earwig forceps (L. W. Simmons & Tomkins, 1996)), and testing of 72 this 'positive allometry' hypothesis is frequently used to infer a sexual selection function 73 when natural observation is unattainable (e.g., trilobite spines (Knell & Fortey, 2005)). 74 The positive allometry hypothesis has, however, been met with resistance. 75 Bonduriansky (2007) noted that the near universality of this pattern may be an artefact of 76 the structures researchers elect to study. That is, when studies focus on morphological

77	scaling, scientists seek the extremes, so the literature is biased in favour of steep scaling
78	relationships (Emlen, 2008; Emlen & Nijhout, 2000; Kodric-Brown et al., 2006). Some
79	extreme structures known to function as sexually selected ornaments, such as elaborate
80	plumage in birds, do not scale positively with body size (José Javier Cuervo & Møller,
81	2001), nor do many genitalic traits, despite the fact that some experience strong selection
82	for increased size (Bertin & Fairbairn, 2007; W. U. Blanckenhorn, Kraushaar, Teuschl, &
83	Reim, 2004; Voje, 2016). Indeed, considering the full range of sexually selected
84	structures, including those that are not extreme in size, reveals that slopes are frequently
85	shallow or negative (Bonduriansky, 2007).
86	Furthermore, at least a few naturally selected structures, such as long bones in
87	large mammals (Bertram & Biewener, 1990; Christiansen, 1999) and cranial horns in
88	lizards (Bergmann & Berk, 2012), also scale positively with body size (Voje, 2016).
89	Clearly, sexual selection need not lead to the evolution of steep scaling, and other agents
90	of selection, such as locomotion and predator defence, occasionally lead to positive static
91	scaling. Where, then, does this leave the positive allometry hypothesis?
92	We argue that steep static scaling relationship slopes can be powerful clues to trait
93	function, particularly when combined with other morphological measures of among-
94	individual variation (e.g., trait-specific coefficients of variation; see below). In this
95	context, we suggest much of the controversy and inconsistency in the literature stems
96	from two sources. First, the positive allometry hypothesis has been applied to all sexually
97	selected structures, when, in fact, the logic holds only for a particular subset: sexually
98	selected signal structures where the size of the structure functions as an honest signal of
99	the body size or resource holding potential of their bearers. Second, tests of the positive

allometry hypothesis often rely on demonstrating a slope significantly greater than one.
While rich in historical precedent, this approach fails to incorporate the signalling
function of these structures. We propose future studies ask not whether the slope is
greater than one, but rather whether the slope is relatively steeper for the focal signal
structure than it is for other, more typically proportioned, non-signal related body parts. It
is the relative increase in slope that allows these structures to function effectively as
signals, and appropriate tests should incorporate this into their methods.

We summarize literature on animal signalling to show why positive allometry is
likely when structures evolve as signals of body size, and why these structures are
predicted to scale more steeply with body size than other, non-signal structures measured
in the same individuals. By the same logic, we explain why other types of extreme
structures, such as those used in prey capture or locomotion, should not scale more

112 steeply than other body parts.

We test these predictions by comparing the slopes of a suite of extreme morphological structures (14 signal, 15 non-signal; Table 1) to slopes of more typically proportioned 'reference' structures within the same organism (rather than the traditional comparison to isometry, see below), and show that relatively steep slopes are common for structures that function as sexually selected signals but not for comparably extreme structures that function in other, non-signalling contexts.

119

120 Methods

121 Specimen/structure selection and morphological measures

122	All species with putatively 'extreme' structures - hereafter referred to as 'focal
123	structures' (see Appendix 1 for our classification of 'extreme') – and adequate sample
124	size (n ≈ 10) were surveyed from the Phillip L. Wright Zoological Museum at the
125	University of Montana (MT, USA), the Museum of Comparative Zoology at Harvard
126	(MA, USA), and the Emlen Lab Entomological Collection (MT, USA). Surveying all
127	species that met these criteria allowed for a relatively unbiased sample of both taxa and
128	structure type. However, since most sexually selected structures in insects are beetle
129	horns (reviewed in Emlen, 2008), the invertebrates surveyed here appear somewhat
130	Coleoptera-biased. Six additional datasets were sourced specifically for this analysis -
131	Jackson's chameleons (Triceros jacksonii) for the presence of both an extreme signal
132	(horns) and non-signal (tongue) structure, large bee flies (Bombylius major), sabre wasps
133	(Rhyssa persuasoria), and peacock moths (Saturnia pyri), for the presence of sexually
134	selected non-signal structures, and ceratopsids (Protoceratops and rewsi) and pterosaurs
135	(Rhamphorhynchus muensteri) to test the described methods on fossil datasets. Finally, it
136	should be noted that while the species/structures surveyed here were unbiased relative to
137	the sampled collections, the collections may have been biased either in taxa or in favour
138	of particularly exaggerated structures. If true, then the results presented here, and their
139	interpretation, may be limited to a particular subset of extreme morphology.
140	Focal structures of extant species were categorized as a 'sexually selected
141	signals'(i.e., structures used by potential mates or competitive rivals as visual signals of
142	the bearer's overall condition/quality (M. B. Andersson, 1994; Bradbury & Vehrencamp,
143	1998)) or 'non-signal' structures using relevant behavioural studies from the literature
144	(see Table 1). When literature on the focal species was unavailable, studies in closely

145	related species were used to infer trait function. Bill function in the American pelican
146	(Pelecanus erythrorphychos) was inferred from its sister species, P. occidentalis (Bels et
147	al., 2012; Kennedy, Taylor, Nádvorník, & Spencer, 2013; Orians, 1969; Schreiber,
148	Woolfenden, & Curtsinger, 1975). Lantern function in the Malagasy lantern bug (Zanna
149	madagascariensis) was inferred from several other Fulgoridae species with similar head
150	morphology (Hogue, 1984; Urban & Cryan, 2009). Snout function in the elephant shrew
151	(Elephantulus fuscus) was inferred from two species of the same genus with similar
152	rostral morphology, E. brachyrhynchus and E. myurus (Kingdon, 1974; Kratzing &
153	Woodall, 1988). Horn function in dung beetles (Sulcophanaeus menelas, Phanaeus
154	saphirinus, Othophagus lanista) was inferred from both a comprehensive review of horn
155	function in beetles (Eberhard, 1980) and empirical studies of dung beetle mating systems
156	(e.g., Emlen, Marangelo, Ball, & Cunningham, 2005; Moczek & Emlen, 2000). Hindleg
157	function in frog legged beetles (Sagra buqueti) was inferred from a closely related
158	species with similar leg morphology and mating behaviour (Katsuki, Yokoi, Funakoshi,
159	& Oota, 2014; O'Brien, Katsuki, & Emlen, 2017). Finally, the function of focal traits in
160	extinct species were inferred from key publications focused on 'bizarre' morphology in
161	the fossil record (Knell & Sampson, 2011; Knell, Naish, Tomkins, & Hone, 2013a; D. W.
162	Hone, Wood, & Knell, 2016; but see Padian & Horner, 2011, 2013, 2014).
163	Reference structures were then chosen for each species as structures that could be
164	consistently measured across all samples and lacked obvious functional connection with
165	the focal structure. These criteria appear adequate in choosing reference structures.
166	However, the authors recognize the limitation of using a single reference structure and
167	encourage the use of multiple reference structures per organism in future application of

the described methods. Doing so will better capture the scaling relationship of 'typical'
(i.e., non-signal) traits and help mitigate impact of choosing inappropriate reference
structures.

171	Measures of overall body size were based on established, taxon specific methods
172	for estimating body size. For species where established estimates of body size were
173	unavailable, methods were adopted from closely related taxa. A summary of study
174	species names, sample sizes, relevant morphological information (e.g., focal structure,
175	reference structure, body size measures), and literature used to establish sexually selected
176	signal/naturally selected non-signal function is provided in Table 1.
177	Dung beetles (Sulcophanaeus menelas), earwigs, mantidflies (Climaciella
178	brunnea), large bee flies, sabre wasps, and wildebeest (Connochaetes tourinus) were
179	measured using photographs (including scale bars) and ImageJ 1.50i software (NIH,
180	USA). S. menelas, earwigs, and mantidflies, large bee flies, and sabre wasps were
181	photographed using a 16.2 megapixel Nikon D5100 DSLR camera mounted on a
182	binocular stereo microscope (Leica S6D) set at a fixed distance. Wildebeest were
183	photographed using a 14.2 megapixel Nikon D3100 DSLR camera set at a fixed distance
184	designated to minimize perspective effects (i.e., approximating orthographic projection).
185	All other extant species were measured using digital callipers.
186	Measures of ceratopsians (Protoceratops and rewsi) and pterosaurs
187	(Rhamphorhynchus muensteri) were collected directly using digital callipers, from
188	photographs of specimens including scale bars, or from the literature when appropriate, to
189	maximize the number of available specimens (see Appendices 2 and 3).
190	

191 Statistical analyses

192 Statistical analyses were performed in R 3.3.2 (R Core Development Team 2016). 193 Measurements were \log_{10} transformed and mean standardized prior to analysis. Ordinary 194 least squares (OLS) regression was used to assess scaling relationship slope (Kilmer & 195 Rodríguez, 2016; Smith, 2009; Warton, Duursma, Falster, & Taskinen, 2012; Warton, 196 Wright, Falster, & Westoby, 2006). For every species, focal structure size and reference 197 structure size were regressed on body size in separate models. Analyses of covariance 198 (ANCOVA) were then used to compare regression slopes of focal structure size on body 199 size (β_{focal}) to regression slopes of reference structure size on body size ($\beta_{\text{reference}}$) within 200 the same species (i.e., to determine whether or not there was a significant interaction 201 between body size and trait group (focal/reference) in explaining trait size). (Differences 202 in intercept were not analysed, since all data were mean-standardized prior to analysis.) 203 In addition, slope estimates (β_{focal} and $\beta_{\text{reference}}$) were collected from each model and 95% 204 confidence intervals constructed. These 95% confidence intervals were then compared 205 between focal and reference structures within the same species. 206 Mean β_{focal} was calculated for sexually selected signal structures and compared to

206 Mean β_{focal} was calculated for sexually selected signal structures and compared to 207 mean β_{focal} calculated for non-signal structures using Welch's t test. Mean $\beta_{reference}$ was 208 calculated for species with sexually selected signal structures and compared to mean 209 $\beta_{reference}$ for species with exaggerated non-signal structures using Welch's t-test. 95% 210 confidence intervals were constructed around mean $\beta_{reference}$ for species with sexually 211 selected signal structures and mean $\beta_{reference}$ for species with non-signal structures and 212 compared. The difference between β_{focal} and $\beta_{reference}$ ($\Delta\beta_{focal-reference}$) was calculated for 213 each species. Mean $\Delta\beta_{focal-reference}$ for species with sexually selected signal structures was

214 compared to mean $\Delta\beta_{\text{focal-reference}}$ for species with non-signal structures using Welch's t-215 test. 95% confidence intervals were constructed around mean $\Delta\beta_{\text{focal-reference}}$ for sexually 216 selected signal structures and mean $\Delta\beta_{\text{focal-reference}}$ for non-signal structures and compared. 217 Coefficients of variation were calculated for every structure. Mean coefficient of 218 variation was calculated across all signal structures and compared to the mean coefficient 219 of variation compared across all non-signal structures using 95% confidence intervals and 220 Welch's t test.

221

222 **Results**

223 Results of species-level analyses are summarized in Table 1, including slope estimates (β_{focal} and $\beta_{\text{reference}}$) and adjusted R² values for all models, differences between 224 β_{focal} and $\beta_{reference}$ ($\Delta\beta_{focal-reference}$), ANCOVA results, 95% confidence intervals 225 surrounding β_{focal} , $\beta_{\text{reference}}$, and $\Delta\beta_{\text{focal-reference}}$, and coefficients of variation. For the 226 227 majority of species with sexually selected signal structures, β_{focal} was significantly greater 228 than $\beta_{\text{reference}}$ (Table 1; Appendix 4). For two of these species, whitetail deer and 229 wildebeest, β_{focal} was greater than $\beta_{\text{reference}}$, but 95% confidence intervals surrounding 230 these estimates were overlapping and the ANCOVA showed no significant difference 231 between β_{focal} and $\beta_{\text{reference}}$. In pronghorn antelope, 95% confidence intervals surrounding 232 β_{focal} and $\beta_{\text{reference}}$ were overlapping, but ANCOVA showed a (slightly) significant 233 difference between β_{focal} and $\beta_{\text{reference}}$. Earwigs, on the other hand, displayed non-234 overlapping 95% confidence intervals surrounding β_{focal} and $\beta_{\text{reference}}$, but the ANCOVA showed no significant difference between β_{focal} and $\beta_{\text{reference}}$. For all species with 235

236	exaggerated, non-signal structures, β_{focal} and $\beta_{reference}$ were either not significantly
237	different, or $\beta_{reference}$ was significantly higher than β_{focal} (Table 1; Appendix 5). Unlike
238	extreme sexually selected signal structures, extreme non-signal structures appear to scale
239	similarly to reference structures within the same organism. Mean slope (β_{focal}) of all
240	exaggerated sexually selected signal structures was greater than the mean slope (β_{focal}) of
241	all non-signal structures ($t_{13.543} = -3.835$, $p < 0.01$) and 95% confidence intervals were
242	non-overlapping (95% CI mean β_{focal} for sexually selected signal structures [1.709, 4.56];
243	95% CI mean β_{focal} for non-signal structures [0.374, 0.783]). Mean $\Delta\beta_{\text{focal-reference}}$ for
244	sexually selected signal structures was greater than mean $\Delta\beta_{focal-reference}$ for non-signal
245	structures ($t_{14.164} = 4.079$, $p = 0.001$; Appendix 6) and 95% confidence intervals did not
246	overlap (95% CI mean $\Delta\beta_{\text{focal-reference}}$ for sexually selected signal structures [1.072, 3.831];
247	95% CI mean $\Delta\beta_{\text{focal-reference}}$ for non-signal structures [-0.501, 0.078]).
248	Coefficients of variation were significantly higher for extreme, sexually selected
249	signal structures (mean = 15.444, 95% CI [9.325, 21.562]) than for non-signal structures
250	(mean = 5.351, 95% CI [3.263, 7.438]) ($t_{16.043}$ = 3.37, $p < 0.01$; Appendix 7).
251	
252	Discussion

253 Within species, sexually selected signal structures scaled steeply with body size 254 (Table 1; Appendix 4). In the majority of sexually selected species surveyed here, the 255 scaling relationship of the signal (β_{focal}) was significantly steeper than that of the 256 reference structure ($\beta_{reference}$). Surprisingly, this pattern did not hold for whitetail deer 257 (*Odocoileus virginianus*) or wildebeest. In these species, β_{focal} was greater than $\beta_{reference}$, 258 but there was no significant difference between β_{focal} and $\beta_{reference}$. Similarly, for earwigs, 259 the ANCOVA showed no significant difference between β_{focal} and $\beta_{reference}$, but β_{focal} was 260 greater than Breference and 95% confidence intervals surrounding these estimates were non-261 overlapping (Table 1; Appendix 4). These results may be an artefact of relatively small 262 sample size (e.g., n < 18 for whitetail deer) and/or biased sampling (e.g., hunters 263 favouring largest antlered males in sampled populations), since previous work has shown 264 positive allometry and/or strong selection for these, and similar, weapons (e.g., Kruuk et 265 al., 2002; Melnycky, Weladji, Holand, & Nieminen, 2013; Lundrigan, 1996; L. W. 266 Simmons & Tomkins, 1996). Alternatively, these structures may function strictly as 267 weapons (i.e., tools) of intrasexual competition, not as visual signals of quality. If true, 268 then steep scaling between weapon and body size is not expected (McCullough, Miller, & 269 Emlen, 2016, see below). Overall, our results for sexually selected signal structures are 270 consistent with previous work showing that these types of extreme structures tend to be 271 positively allometric (Bonduriansky & Day, 2003; Emlen, 2008; Green, 1992; Kodric-272 Brown & Brown, 1984; Kodric-Brown et al., 2006; Petrie, 1988, 1992; L. W. Simmons 273 & Tomkins, 1996; Voje, 2016).

Every exaggerated non-signal structure measured scaled with a slope that was either less than, or not significantly different from, that of the reference structure (Table 1; Appendix 5). In addition, across species, the scaling relationship (β_{focal}) of sexually selected signal structures was significantly steeper than that of non-signal structures ($t_{11.902} = -3.23$, p < 0.01). Even within the same organism, non-signal structures scaled at a shallower rate than sexually selected signals. In Jackson's chameleon, for example, where both an extreme sexually selected signal, horn length, and an extreme non-signal

281 prey capture structure, tongue length, were surveyed, horn size scaled at a much steeper

rate compared to the reference structure than did tongue size (Table 1; Fig. 2).

283

284 Why signals should scale more steeply than other body parts

285 Many studies have considered what makes a good signal (reviewed in Bradbury & 286 Vehrencamp, 1998; Maynard Smith & Harper, 2003; Searcy & Nowicki, 2006). In the 287 context of sexual selection, receivers are often females who use variation in signal 288 expression as a basis for mate choice, or males who use these signals to determine the 289 resource holding potential (i.e., fighting ability) of rival males (M. B. Andersson, 1994; 290 Bradbury & Vehrencamp, 1998; Hardy & Briffa, 2013). In both cases, information 291 encoded in the signal pertains to the overall genetic quality and/or condition of the bearer 292 (reviewed in Neff & Pitcher, 2005).

293 Although any phenotype could, in principle, be used as a signal (provided it is 294 detectable and variable across individuals), some make more effective signals than 295 others. The best signals are conspicuous - bigger or brighter than other body parts 296 (Bradbury & Vehrencamp, 1998). However, it is not just the structure that must be 297 conspicuous. Variation in the expression of that structure is key to mate and rival 298 assessment, and the more pronounced the differences, the better. For this reason, signal 299 structures are often selected to be more variable in their expression than other, 300 surrounding, non-signal structures (Alatalo, Höglund, & Lundberg, 1988; José Javier 301 Cuervo & Møller, 2001; Emlen, Warren, Johns, Dworkin, & Lavine, 2012; Fitzpatrick, 302 1997; Petrie, 1992; Pomiankowski & Moller, 1995; Rowe & Houle, 1996; L. W. 303 Simmons & Tomkins, 1996; Tazzyman, Iwasa, & Pomiankowski, 2014; Wallace, 1987). Hypervariability in trait size amplifies associated variation in male quality, making these
otherwise subtle differences easier to see (Hasson, 1991; Tazzyman et al., 2014; Wallace,
1987).

307	Effective signals must also be honest. If poor quality males can cheat by
308	producing effective signals, then reliability of the signal plummets and receivers should
309	focus on other traits. One form of honesty arises when the growth of signal traits is
310	condition-sensitive (Biernaskie, Grafen, & Perry, 2014; Bonduriansky, 2006;
311	Bonduriansky & Day, 2003; Grafen, 1990; Iwasa, Pomiankowski, & Nee, 1991;
312	Johnstone, 1997; Kodric-Brown et al., 2006; Nur & Hasson, 1984; Pomiankowski, 1987;
313	Zeh & Zeh, 1988). Condition-sensitive growth of signal structures may 'capture' genetic
314	or environmental variation underlying overall quality, making these signals virtually
315	impossible to fake (Miller & Moore, 2007; Rowe & Houle, 1996; Wilkinson & Taper,
316	1999). Indeed, sexually selected signal structures are notoriously sensitive to stress,
317	parasite load, and nutrition (Cotton, Fowler, & Pomiankowski, 2004; Ezenwa & Jolles,
318	2008; Gosden & Chenoweth, 2011; Hamilton & Zuk, 1982; Izzo & Tibbetts, 2015; Knell
319	& Simmons, 2010; Kruuk et al., 2002; Skarstein & Folstad, 1996).
320	Hypervariability through heightened condition sensitivity causes structures to be
321	reliable and informative as signals of quality (M. B. Andersson, 1994; M. Andersson &
322	Iwasa, 1996; M. Andersson & Simmons, 2006; Bradbury & Vehrencamp, 1998), and
323	these basic characteristics are shared by a wealth of sexually selected signals (reviewed in
324	Bradbury & Vehrencamp, 1998). When information contained in a sexually selected
325	signal involves individual differences in the size of a structure, and when among-
326	individual variation in condition or genetic quality manifests as differences in overall

327 body size, then selection for increasingly effective signals should lead to the evolution of 328 not just higher trait-specific coefficients of variation, but also to a relatively steeper 329 scaling relationship slope (Biernaskie et al., 2014; Green, 1992; Kodric-Brown & Brown, 330 1984; Kodric-Brown et al., 2006; Petrie, 1988). The steeper the slope, the more variable 331 the focal structure will be relative to surrounding body parts. Mechanistically, when 332 variation in condition is driven by differential access to nutrition, then the evolution of 333 heightened condition-sensitive growth in a particular structure, relative to others, will 334 also manifest as an increase in the steepness of the slope for that structure (Emlen et al., 335 2012; Lavine, Gotoh, Brent, Dworkin, & Emlen, 2015; Mirth, Frankino, & Shingleton, 336 2016; Shingleton & Frankino, 2013). Thus, for this particular subset of signal structures, 337 the positive allometry hypothesis should hold. Indeed, the steeper the scaling relationship 338 slope, the better the signal will be, leading to the evolution of larger and larger structures 339 with steeper and steeper patterns of static scaling.

340 A few exceptions should be noted, however. First, body size is not always 341 correlated with overall genetic quality or condition, as is the case for many fishes (Bolger 342 & Connolly, 1989) and birds (José J. Cuervo & Møller, 2009). In these species, signals 343 are still expected to be condition-sensitive and hypervariable. However, because 344 condition is not correlated with body size, differences in the relative sizes of signal 345 structures may not covary with body size (e.g., Bonduriansky & Day, 2003; José J. 346 Cuervo & Møller, 2009; Fitzpatrick, 1997; Pomfret & Knell, 2006). (This was true for 347 several focal non-signal traits, and several reference traits surveyed here (indicated by low adjusted R^2 values; Table 1). Indeed, future analyses may benefit from choosing 348 349 reference structures that more tightly covary with body size.) Similarly, signals that vary

350 in other ways besides size (e.g., colour, behaviour, chemical signals) are also not 351 expected to scale with body size. Finally, sexually selected traits that do not function as 352 signals (e.g., peacock moth antennae, measured here; Table 1; Appendix 5), are not 353 predicted to scale steeper than reference structures, since hypervariation and/or condition 354 sensitivity may actually decrease performance. This includes sexually selected weapons 355 that function only as tools of battle and not as signals of quality, condition, or resource 356 holding potential (McCullough et al., 2016). For these structures, trait expression should 357 be proportional across the entire population, even when selection favours large relative 358 trait sizes. Large structures may display especially high scaling relationship intercepts 359 compared to other traits in the body, but since there is no hypervariation and/or 360 heightened condition sensitivity, the slope should not differ from that of a reference 361 structure. Consequently, we suggest much of the confusion regarding the link between 362 positive allometry and sexual selection can be resolved by recognizing that the positive 363 allometry hypothesis applies only to those structures that act as visual signals of among-364 individual variation in condition or genetic quality and, in fact, it applies only to a subset 365 of these, signals whose information involves differences in signal size in species where 366 quality is approximated by variation in overall size. For these structures, sexual selection 367 is predicted to drive the evolution of extreme trait size and unusually steep scaling.

368

369 Testing the positive allometry hypothesis against reference structures, rather than
370 isometry

We suggest three reasons for testing the positive allometry hypothesis incomparison with reference structures, rather than with isometry. First, inferring signal

373 function for a structure that scales steeply only makes sense if that structure scales more 374 steeply than other body parts. Steep scaling relationship slopes are relevant because they 375 cause structures to be better signals than other, surrounding body parts. The properties 376 that make them effective signals are relative: they are more variable and more condition-377 sensitive in their growth than other body parts. Sexual selection favours receivers who 378 pay attention to these structures because, by doing so, individuals make more informed 379 decisions than they would if they focused on other body parts. Consequently, the pattern 380 that matters for inferring a sexually selected signal function is the difference in slope 381 between the putative signal and other, non-signal, structures.

382 Second, detecting hyperallometry in a focal structure without comparing the slope 383 to a control can be misleading. It is possible for non-signal structures to scale steeply. 384 Indeed, in our sample of non-signal exaggerated structures, gaboon viper (*Bitis gaboncia*) 385 fangs, elephant shrew (Elephantulus fuscus) snouts, and mantidfly forelegs all scaled 386 with relatively slopes (i.e., $\beta > 1$), but the reference structures were hyperallometric too 387 (Appendix 5; Table 1). Had we focused only on the absolute value of the scaling 388 relationship slope we would have erroneously inferred a signal function for these 389 structures when, in fact, their scaling relationship slopes were no different from those of 390 surrounding body parts. These structures lack the critical properties of an informative 391 signal despite being hyperallometric.

Finally, comparing measured slopes with isometry places undue emphasis on the estimated slope *per se*. Isometry may be intuitive in principle, but actually detecting it, or rejecting it, depends a lot on the particular landmarks selected, the units of measurement involved, and the chosen measure of body size (Bookstein, 1989; Jungers, Falsetti, &

396 Wall, 1995; e.g., Mosimann & James, 1979). For this reason, focusing tests of the 397 positive allometry hypothesis exclusively on rejection of a slope of one may be 398 misleading, especially in the context of interspecific comparisons where landmarks and 399 measures of body size/condition often differ (e.g., Bolger & Connolly, 1989; Jakob, 400 Marshall, & Uetz, 1996, p. d; Peig & Green, 2010). Focusing instead on the slopes of 401 focal structures compared to those of reference structures delivers an internally controlled 402 assay for the properties of a structure's expression that matter. Significant increases in the 403 slope of a focal structure relative to other body parts means that the focal structure has the 404 predicted properties of a signal, and we suggest this constitutes evidence in favour of a 405 function for that structure as a sexually selected signal.

406

407 Diversity of exaggerated morphology

408 Not all sexually selected structures are signals, but many experience strong 409 selection for increased size. In arthropods with low population density, for example, 410 males search for receptive females and selection can lead to the evolution of elaborate 411 antennae and/or enlarged eyes (e.g., peacock moth antennae, measured here; Table 1). 412 This results in pronounced sexual dimorphism in relative trait size and, in some species, 413 exaggerated male sensory structures (M. B. Andersson, 1994; Bertin & Cezilly, 2003; 414 Lefebvre, 2000; Thornhill, 1981). Similarly, antagonistic coevolutionary arms races 415 arising from conflict between males and females can drive rapid evolution of genitalia 416 (Arnqvist & Rowe, 2002, 2005; Brennan, Clark, & Prum, 2009; Parker, 1979; Leigh W. 417 Simmons, 2014). In both contexts, sexual selection drives the evolution of extreme size, 418 but these structures do not function as signals. There is little covariance between trait

419 *variation* and fitness and, thus, no benefit in traits being hypervariable or extra condition

420 sensitive. For these traits, steep scaling slopes are not expected (e.g., Eberhard, 1998,

421 2010; Hosken & Stockley, 2004).

422 Exaggerated size can also arise through natural selection as, for example, in some 423 locomotor, prey capture, and feeding structures (reviewed in Lavine et al., 2015). 424 Appendages such as praying mantis forelimbs and antlion mandibles function like levers, 425 snapping closed to grasp prey. For these species, longer forelimbs or mandibles perform better than shorter ones both because they move faster at their tips, and because they 426 427 sweep through a larger 'kill zone' (Loxton & Nicholls, 1979; Maldonado, Levin, & Pita, 428 1967). However, like sensory and genitalic structures of sexual selection, large size in 429 these naturally selected structures is not related to a signal function. There is no benefit to 430 hypervariability or heightened condition sensitivity, and steep scaling relationship slopes 431 are not expected.

432 Here, we provide measures of static allometry for 15 extreme non-signalling 433 structures (Table 1; Appendix 5). None are sexually dimorphic, and none scaled more 434 steeply than other, typically proportioned, body parts. Jackson's chameleons provide 435 perhaps the best example of all, since males in this species have both types of extreme 436 structure: three horns on the head that function as a signal of competitive ability (Bustard, 437 1958), and an elongated tongue used to capture prey. Even though the tongue is relatively 438 larger than the horns, tongues scaled with a slope that was shallower than the reference 439 structure. Horns, in contrast, scaled disproportionately steeply (Fig. 2). Clearly, the 440 evolution of extreme structures need not entail relative increases in static allometry slope,

and steep slopes, when they occur, can provide valuable clues to a sexually selectedsignal function.

443

444 Inferring function for extreme structures in extinct taxa

445 Unlike most organisms described above, the behaviour of extinct taxa cannot be 446 observed. Even so, lines of evidence can be drawn from static, morphological data to 447 provide testable hypotheses of behaviour (D. W. E. Hone & Faulkes, 2014). For example, 448 hypotheses surrounding mechanical function, such as those involving anchors for 449 musculature or levers that increase moment arms, can be assessed (and potentially 450 rejected) using data from fossils (e.g., D. W. Hone, Naish, & Cuthill, 2012; Knell & 451 Fortey, 2005). Similarly, we maintain the use of static scaling relationship slopes and 452 coefficients of variation may provide a means for inferring a sexually selected signal 453 function for extreme morphology in the fossil record. 454 Static scaling relationships have been used already to infer function in the fossil 455 record (Gould, 1973; D. W. Hone et al., 2016; Knell & Fortey, 2005). However, such 456 inferences remain controversial (e.g., Padian & Horner, 2011, 2013, 2014; Knell & 457 Sampson, 2011; Knell et al., 2013a; D. W. Hone & Mallon, 2017; Mallon, 2017). One 458 issue is that collecting multiple individuals from the same fossil locality and horizon (i.e., 459 a single population) is difficult. Sample sizes are often small or gathered from animals 460 separated in space and/or time, and animals are rarely sexed (e.g., D. W. Hone & Mallon, 461 2017). As a result, detection of even fundamental patterns in morphology, such as sexual 462 dimorphism, remains elusive (Mallon, 2017; but see Sengupta, Ezcurra, & 463 Bandyopadhyay, 2017). Another issue is that distinguishing between different signal

464 functions is often difficult. Social dominance and sexually selected signals, for example,
465 are often confluent and distinguishing between them is complex. In addition, the cooption
466 of extreme structures to multiple functions, thereby exposing them to multiple patterns of
467 selection, may further confound these data (e.g., dugong tusks; Anderson, 1979;

468 Domning & Beatty, 2007).

469 Despite these limitations, we suggest behaviour can be inferred from the fossil 470 record using the methods and logic described above. We predict that when focal 471 structures act as signals of overall body size, both the slope of the static scaling 472 relationship and the coefficient of variation will be steeper/greater in the putative signal 473 structure than in reference structures used as controls. As 'proof of concept' for this 474 approach, we included two putative sexually selected signal structures from the fossil 475 record in our analyses, the enlarged cephalic frill of the ceratopsian dinosaur 476 Protoceratops and rewsi (adapted and expanded from D. W. Hone et al., 2016), and the 477 tail vane of the pterosaur, *Rhamphorhynchus*. In both cases, the focal structure scaled 478 more steeply with body size and had a higher coefficient of variation than reference 479 structures measured in the same individual (Fig. 3; Table 1), implying a signalling 480 function.

Overall, we believe this method useful for inferring extreme structure function in the fossil record (perhaps even more useful when analysed in conjunction with other patterns in morphology - e.g., changes in complexity during ontogeny, high variation in trait shape and size between species lineages). Both morphological scaling relationships and coefficients of variation can be reliably measured in fossil specimens, even when sample size is small. We recommend the use of these methods in subsequent analyses of

487	extreme or 'bizarre' morphology in the fossil record, and are hopeful that they might
488	provide insight into the ongoing debate regarding sexual selection in non-avian dinosaurs.
489	
490	Overall, we suggest that when applied specifically and exclusively to
491	disproportionately large animal structures that function as signals of overall body size,
492	and when assessed through comparison with surrounding, non-signal structures rather
493	than through detection of an estimated slope greater than 1, the positive allometry
494	hypothesis holds. Sexually selected signal structures are predicted to – and, in fact,
495	appear to – scale more steeply with body size than non-signal structures. For this reason,
496	we suggest that relative patterns of trait scaling offer powerful clues to trait function,
497	particularly when combined with other measures of trait expression such as trait specific
498	coefficients of variation.
499	
500 501	Data archive
502	Datasets supporting this article will be uploaded to Dryad
503	
504	

- **Competing interests**
- 506 The authors have no competing interests to report

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- 946 Figure 1: Extreme non-signal (ns) and sexually selected (ss) signal structures. Clockwise
- 947 from top right; bighorn sheep horns (O. canadensis; ss), Jackson's chameleon horns (T.
- 948 *jacsonii*; ss), praying mantis forelimbs (Mantodea; ns), ichneumon wasp ovipositor
- 949 (Ichneumonoidea; ss non-signal), gaboon viper fangs (*B. gaboncia*; ns), and dung beetle
- 950 horns (Scarabaeidae, ss). Photos credited in Acknowledgments.





952

954	Figure 2: Static scaling relationships for an extreme sexually selected signal structure
955	(horns; red; left; $n = 40$) and an extreme, non-signal naturally selected structure (tongue;
956	blue; right; $n = 25$) in Jackson's chameleons (<i>T. jacksonii</i>). Red and blue indicate focal
957	structures. Grey indicates the reference structures. Lines represent ordinary least squares
958	regression of standardized log_{10} structure size on standardized log_{10} body size. In
959	Jackson's chameleon, the extreme sexually selected signal (horn length) scales at a
960	significantly steeper rate than the reference structure (hindfoot length). The extreme non-
961	signal structure (tongue length) does not. 95% CI for horn length [3.358, 5.159], tongue
962	length [0.251, 0.949], and hindlimb length [1.13, 1.979].





968	Figure 3:	Static scaling	g relationshi	ps for extreme	putative sexual	lly selected	signal
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- 969 structures in ceratopsians (*Protoceratops andrewsi*; left; n = 38) and pterosaurs
- 970 (*Rhamphorhynchus muensteri*; right; n = 10). Red indicates putative signal structures.
- 971 Grey indicates reference structure. Lines represent the ordinary least squares regression
- 972 of standardized log_{10} structure size on standardized log_{10} body size. In both species, the
- scaling relationship of the putative signal trait is steeper than that of the reference trait (*P*.
- andrewsi: 95% CI for slope of focal structure [1.173, 1.353], 95% CI for slope of
- 975 reference structure [0.925,1.039]; *R. muensteri*: 95% CI for slope of focal structure
- 976 [1.332, 2.930], 95% CI for slope of reference structure [0.871, 1.262]), consistent with a
- history of selection for a hypervariable sexually selected signal. Inlaid photographs
- 978 display study species with focal structures highlighted in red. Photos credited in
- 979 Acknowledgments.
- 980



Standardized log_{10} body size (head length)

Fig.%3



983 Table 1: Summary of study species and results.

985 INCLUDED SEPARATELY AS EXCEL TABLE

987	Table 1 footnotes
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988 ((f) = focal trait,	$(\mathbf{r}) = \mathbf{reference trait},$	$\mathbf{C}\mathbf{V} = \mathbf{c}$	coefficient of	f variation,	β = slope	of scaling
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relationship between trait size and body size, $\Delta\beta$ = difference between $\beta_{(f)}$ and $\beta_{(r)}$, \ddagger

990 extinct species, $R^{2}_{(f)}$ = adjusted R^{2} of scaling relationship between focal trait size and

- body size, $R^{2}_{(r)}$ = adjusted R^{2} of scaling relationship between reference trait and body
- size, * = sexual dimorphism may be impossible to detect (see D. W. Hone & Mallon,
- 993 2017). F and *p* value from ANCOVA comparing $\beta_{(f)}$ and $\beta_{(r)}$ within the same species.

1006 Appendix 1: Identifying extreme morphology

1007 Many extreme structures appear self-evident. Some, like beetle horns, are massive 1008 in absolute and relative size and few would contest their designation as extreme. Others 1009 are more ambiguous. Butterfly wings, for example, rarely earn the title of extreme yet, 1010 their ontogenetic growth and relative size are more akin to that of beetle horns than other 1011 insect wings (Nijhout & Emlen, 1998). Examples like this highlight the ambiguity 1012 surrounding extreme morphology and the subjective nature of categorizing structures as 1013 extreme. This uncertainty, in part, stems from the lack of established criteria for 1014 designating a structure as extreme. For over a century, researches have explored the 1015 evolution of extreme morphology (M. B. Andersson, 1994; reviewed in Darwin, 1871; 1016 Emlen, 2008). Yet, to our knowledge, not once has the term 'extreme' been defined. 1017 Recognizing and limiting bias is a vital component of biological research and, 1018 given the large body of work dedicated toward putatively extreme structures, we believe 1019 a consistent method for identifying these structures is needed. Here we suggest three 1020 (potentially overlapping) categories of extreme - ontogenetically, statically, and 1021 evolutionarily extreme – and provide guidelines for assigning structures to each category. 1022 1023 **Ontogenetically Extreme:** Ontogenetically extreme structures are those displaying rates 1024 of growth, often occurring in bursts close to reproductive maturity, that outpace other 1025 surrounding structures. Examples include the horns of beetles and the wings of 1026 lepidopterans, both of which grow to drastic proportions during the same timeframe as 1027 other, more typically proportioned structures (Nijhout & Emlen, 1998). Ontogenetically

extreme should be distinguished by rates of growth that are faster than those of referencestructures within the same organism.

1030

1031 Statically Extreme: Statically extreme structures are disproportionately larger than other 1032 structures when sampled across same stage (generally adult) individuals within a 1033 population. Relative size of a focal trait can be assessed by comparing the size of the 1034 focal trait to other, analogous traits in the same sex (e.g., harlequin beetle (Acrocinus 1035 *longimanus*) forelegs are relatively larger than midlegs or hindlegs (Zeh, Zeh, & 1036 Tavakilian, 1992)) or by comparing the size of the same trait across sexes (e.g., harlequin 1037 beetle forelegs are disproportionately larger in males than they are in females (Zeh et al., 1038 1992)). Statically extreme structures should be distinguished by comparing slopes and/or 1039 intercepts of the static scaling relationships (trait size versus body size) of the focal and 1040 reference traits.

1041

1042 **Evolutionarily Extreme:** Evolutionarily extreme structures are extreme when compared 1043 with homologous structures in closely related organisms. Examples include the hindlegs 1044 of jerboas, which are relatively longer than the hindlegs of their quadrupedal ancestors 1045 (Miljutin, 2008; Dipodidae; Wu et al., 2014) and the raptorial forelimbs of mantidflies 1046 (mantispidae; Ohl, Barkalov, & Xin-Yue, 2004). Evolutionarily extreme structures can be 1047 distinguished by a) comparing static scaling relationships (slopes and/or intercepts) of 1048 individuals sampled from populations of ancestral and derived species; b) comparing 1049 mean relative trait size of ancestral and derived species (e.g., Wu et al., 2014); and/or c)

- 1050 by mapping changes in trait size onto a phylogeny and testing for lineage specific
- 1051 changes in relative trait size (Wu et al., 2014).

- 1073 Appendix 2: Sources for *Protoceratops andrewsi* data. AMNH = American Museum of
- 1074 Natural History (New York, USA); MPC = Mongolian Palaeontological Centre
- (Ulaanbaatar, MN); IVPP = Institute of Vertebrate Palaeontology and 1075
- 1076 Palaeoanthropology (Beijing, CN); ZPAL = Zoological Institute of Paleobiology, Polish
- 1077 Academy of Sciences (Warsaw, PL); CMNH/CM = Carnegie Museum of Natural History
- 1078 (Pittsburgh, USA); NHM = Natural History Museum (London, UK).
- 1079

1079	Protoceratops andrewsi			
	Source	Specimen	Number	
1080	Dodson (1976)	AMNH	6419	
1000	Dodson (1976)	AMNH	6434	
	Dodson (1976)	AMNH	6430	
1081	Dodson (1976)	AMNH	6251	
1001	Dodson (1976)	AMNH	6431	
	Dodson (1976)	AMNH	6486	
1082	Dodson (1976)	AMNH	6432	
1002	Dodson (1976)	AMNH	6428	
	Dodson (1976)	AMNH	6409	
1083	Dodson (1976)	AMNH	6480	
1005	Dodson (1976)	AMNH	6444	
	Dodson (1976)	AMNH	6485	
1084	Dodson (1976)	AMNH	6408	
1004	Dodson (1976)	AMNH	6433	
	Dodson (1976)	AMNH	6429	
1085	Dodson (1976)	AMNH	6439	
1005	Dodson (1976)	AMNH	6441	
	Dodson (1976)	AMNH	6477	
1086	Dodson (1976)	AMNH	6417	
1000	Dodson (1976)	AMNH	6425	
	Dodson (1976)	AMNH	6413	
1087	Dodson (1976)	AMNH	6414	
1007	Dodson (1976)	AMNH	6438	
	Dodson (1976)	AMNH	6466	
1088	Dodson (1976)	AMNH	6467	
1000	Handa et al. (2012)	MPC	100/539	
	Hone et al. (2014)	MPC	100/534	
1089	Hone et al. (2014)	MPC	100/526 B	
1007	Hone et al. (2014)	MPC	100/526 C	
	Fastovsky et al. (2011)	MPC	100/530 A	
1090	Fastovsky et al. (2011)	MPC	100/530 B	
	Fastovsky et al. (2011)	MPC	100/530 C	
	Fastovsky et al. (2011)	MPC	100/530 D	
1091	Fastovsky et al. (2011)	MPC	100/530 E	
	Fastovsky et al. (2011)	MPC	100/530 F	
	Fastovsky et al. (2011)	MPC	100/530 G	
1092	Fastovsky et al. (2011)	MPC	100/530 H	
	Unpublished photos	IVPP	23899	
	Unpublished photos	IVPP	Unnumbered Medium	
1093	Unpublished photos	IVPP	Unnumbered Small	
	Unpublished photos	ZPAL	MgD-II/2b	
	Unpublished photos	ZPAL	MgD-II/5	
1094	Unpublished photos	CMNH	9185	
	Unpublished photos	NHM	5134	
	Unpublished photos	NHM	6442	
1095	Unpublished photos	NHM	6440	
	Unpublished photos	AMNH	6418	
	Unpublished photos	AMNH	6637	
	Unpublished photos	AMNH	6422	
	Unpublished photos	AMNH	6485	

1096	Appendix 3: Sources for	Rhamphorhvnchus muensteri data. BSP = Palaeontolo	gical
	TT		0

- 1097 Museum, Munich (Munich, DE); YPM = Yale Peabody Museum (CT, USA) ;SMF =
- 1098 Forschungsinstitut und Naturmuseum Senckenberg (Frankfourt, DE); CMNH/CM =
- 1099 Carnegie Museum of Natural History (Pittsburgh, USA); SOS = Jura Museum (Eichstätt
- 1100 DE); NHM = Natural History Museum (London, UK); TMP = Royal Tyrell Museum of
- 1101 Palaeontology (Alberta, CA); MBR = Museo Argention de Ciencias Naturales (Buenos
- 1102 Aires, AR); BMNS = Brazoport Museum of Natural Science (TX, USA); NMS =
- 1103 National Museums of Scotland (Edinburgh, UK); TPI = Thanksgiving Point Institute
- 1104 (North American Museum of Ancient Life, UT, USA).
- 1105

Rhamphorhynchus muensteri

1107	Specimen	Specimen	Number	Wellnhofer number
1107	Wellnhofer (1975)	BSP	1960 .I. 470	9
1100	Wellnhofer (1975)	BSP	1938 .I .503	11
1108	Wellnhofer (1975)	Eichstaett		28
	Wellnhofer (1975)	YPM	1778	33
1109	Wellnhofer (1975)	SMF	R 4128	43
	Wellnhofer (1975)	СМ	11429	53
1110	Wellnhofer (1975)	BSP	1907 37	60
•	Wellnhofer (1975)	SOS	3558	77
1111	Wellnhofer (1975)	NA	NA	102
1111	Hone (2012)	NHM	W1198z0077/0001	
	Direct Measurement	TMP	2008.041.0001	
1112	Direct Measurement	MBR	3650.3	
	Direct Measurement	BMNS	21	
1113	Measured from photo	NMS	G.1994.13.1.	
	Measured from Photo	TPI	1012	
1114				
1115				
1115				
1117				
1116				
1117				
1118				
-				

Appendix 4: Scaling relationships for extreme sexually selected signal structures.
Lines represent ordinary least squares regression of log₁₀ standardized structure size on
log₁₀ standardized body size (slope estimates and sample sizes reported in Table 1). Red
points and lines represent focal traits. Grey points and lines represent reference traits.



1130 Appendix 5: Scaling relationships for extreme naturally selected/non-signal

structures. Lines represent ordinary least squares regression of log₁₀ standardized
structure size on log₁₀ standardized body size (slope estimates and sample sizes reported
in Table 1). Blue points and lines represent focal traits. Grey points and lines represent
reference traits.

1135



Standardized log₁₀ body size

- 1137 Appendix 6: Comparison of Δβ_{focal-reference} (difference between the scaling
- 1138 relationship slope of focal traits and reference traits) between extreme sexually
- 1139 selected signal traits (n = 14) and extreme non-signal selected traits (n = 15). $\Delta\beta_{\text{focal-}}$

1140 reference of extreme sexually selected signal structures is significantly greater than $\Delta\beta_{\text{focal-}}$

- 1141 reference of extreme non-signal structures ($t_{15.616} = 4.153 p < 0.001$).
- 1142
- 1143



- 1147 Appendix 7: Comparison of coefficients of variation (CV) between extreme sexually
- selected signal traits (n = 14) and extreme non-signal selected traits (n = 15). CVs of
- 1149 extreme sexually selected signal structures is significantly greater than CVs of extreme
- 1150 non-signal structures ($t_{16.043} = 3.37, p < 0.01$).
- 1151



