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1 The evolution and diversity of intra-male sperm translocation in

- 2 **Odonata: a unique behaviour in animals**
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16 **Abstract**

17 Behavioural diversity is a basic component of biodiversity, with implications in 18 ecological interactions at the intra- and interspecific levels. The reproductive 19 behaviour of Odonata (dragonflies and damselflies) is unique among insects 20 and conditioned by the anatomical separation between the male's reproductive 21 organs and the intromittent organ. Prior to mating, males must translocate 22 sperm from the genital pore in the 9th abdominal segment to the seminal vesicle 23 located ventrally in the 2nd abdominal segment. This behaviour, exclusive to 24 odonates, is known as intra-male sperm translocation (ST). Here, we review the 25 literature on ST, and use phylogenetic comparative analyses to investigate the 26 evolution of ST within the Odonata. Information on ST was compiled for 176 27 species, with the commonest variant being ST once per mating, after tandem 28 formation (66%). Other variants found were: ST involving precopulatory genital 29 touching (10%), ST by the male alone before tandem (16%) or after copulation 30 (5%), and repetition of ST during the same copulation (3%). The precopulatory 31 genital touching might have evolved to detect female receptivity. ST before 32 tandem formation might be favoured when mating opportunities are scarce and 33 copulations are brief. ST after mating might be favoured if males need to be 34 ready to copulate fast. Finally, repeated ST could have evolved through 35 postcopulatory sexual selection in males with limited sperm removal ability, as a 36 mean to improve their sperm competition. The most plausible scenario for the 37 evolution of ST is that the ancestors of the Odonata produced a spermatophore 38 and attached it to the body, leading towards the evolution of the secondary 39 genitalia in males. Our study emphasizes the role of behavioural diversity to 40 understand behavioural evolution.

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41 Significance statement

- 42 Unique behaviours are exclusive of a few individuals, populations and/or
- 43 species. The intra-male sperm translocation (ST) of dragonflies and damselflies
- 44 is a unique behaviour in animals: before mating, males need to transfer sperm
- 45 from the primary to the secondary genitalia, which are anatomically separated.
- 46 Thus, the viability and quality of sperm (*i.e.* fertility) will depend on the timing of
- 47 ST relative to copulation. Our literature review found a variety of ST variants,
- 48 being ST in tandem and before copulation the ancestral strategy. We discuss
- 49 putative evolutionary routes for all the variants found and emphasize the
- 50 importance of retrieving detailed observations of such unique behaviours in the
- 51 field, which could help to better understand behavioural evolution in this insect
- 52 group. Behavioural diversity is rarely addressed by conservation strategies,
- 53 despite unique behaviours being at a higher risk of extinction.
- 54
- 55 Keywords: dragonflies, damselflies, reproductive behaviour, ethodiversity

56 Introduction

57 Behavioural diversity or ethodiversity (Cordero-Rivera 2017a) is a fundamental 58 level of phenotypic variability that can promote population stability in changing 59 environments (e.g. Dingemanse et al. 2004), increase adaptability at the 60 intraspecific level (Berger-Tal and Saltz 2016) and inform about evolutionary 61 processes. Ethodiversity can thus play an important role in species resilience 62 and consequently in how we can manage species for conservation strategies. 63 Unfortunately, while many conservation strategies address the extinction of 64 species and its possible cascading effects across trophic levels (Pérez-Méndez 65 et al. 2016), the disappearance of behaviours is rarely considered (Caro and 66 Sherman 2012). Behaviours that are exclusive of a limited number of 67 individuals, populations or species, are relevant from a conservation 68 perspective, because they are in high risk of extinction (Caro and Sherman 69 2012). For instance, sexual conflict can be an engine of speciation, because it 70 may trigger antagonistic coevolution between the sexes, potentially leading to 71 rapid divergent evolution of the characters related to reproduction (e.g. Cordero-72 Rivera, 2017b). Therefore, the study of reproductive behaviours is relevant for 73 our understanding of diversification and speciation. 74 Insects from the order Odonata (dragonflies and damselflies) represent a clear

75 example of rare behavioural diversity, regarding their copulatory behaviour. In 76 most insects, copulation is performed with both partners oriented in opposite 77 directions (Figure 1A). However, the reproductive behaviour of odonates is 78 unique because during copulation male and female remain attached by two 79 points, forming the so-called "copulatory wheel" (Figure 1B). In this position, the 80 male anal appendages located at the tip of the abdomen grasp the female 81 thorax (or head), and the female genitalia located at the distal part of the 82 abdomen, contacts with the male secondary genitalia, positioned in the proximal 83 part of the abdomen. Such copulatory position occurs because the male 84 intromittent organ is situated in the second and third abdominal segments 85 (secondary genitalia), while the male primary genitalia are located in the ninth 86 abdominal segment. Therefore, before insemination, the male must translocate 87 sperm from the primary to the secondary genitalia, a behaviour called intra-male 88 sperm translocation (hereafter, ST) (Figure 1C-F). The ST behaviour is not

found in any other insect (Shuker and Simmons 2014; Córdoba-Aguilar et al.
2018), although indirect insemination is also found in octopuses and spiders.
The ST it is thus a clear example of diversity at the behavioural level with
important evolutionary implications, because the viability and quality of the
sperm, and hence fertility, will depend on the timing of ST relative to copulation
(Rivas-Torres et al. unpublished).

95 However, little is known about its evolution, its relation with the evolution of the 96 copulatory wheel, and particularly its diversity within the order. Our main aim 97 was therefore to explore the variation of ST behaviour from an evolutionary 98 perspective and discuss its possible connection with the copulatory wheel. We 99 performed a literature review on the diversity of ST across Odonata, and we 100 used phylogenetic comparative analyses to investigate the evolution of the ST 101 behaviour within this insect order. Our literature review revealed differences in 102 ST behaviour within and between species, and the comparative phylogenetic 103 analyses suggested ST in tandem and before copulation, as the ancestral 104 behaviour in Odonata. We discuss potential evolutionary routes for all the ST 105 behaviours found, which could help to better understand behavioural evolution 106 in this insect group, as well as certain aspects of the evolution of sexual 107 behaviours and divergence in other animals.

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109 Material and Methods

110 Literature review of ST behaviour

111 Published data on ST in odonates were searched by querying Google scholar 112 (http://scholar.google.com) for "sperm translocation" and "odonat*" or "intra-113 male sperm transfer" and "odonat*". Searches were carried out in December 114 2016. Additional searchers were completed in November 2018 looking for ST 115 behaviour in papers written in Spanish, French, German and Italian and manually looking in old English references, where the behaviour could be 116 117 described in different terms (e.g. "filling of the sperm vesicle"). Japanese 118 references were also included thanks to the help of colleagues. We also 119 screened manually the papers published in Martinia (from 1991 to 2016) and in 120 the Journal of the British Dragonfly Society (from 1995 to 2016), which are two 121 non-digitized odonatological journals). We collated all available observations of

ST, including how this behaviour was performed, and its duration (with sample 122 123 size and standard error). We compiled information for a total of 176 species 124 (see Results). Full details of the literature reviewed, variables considered, and 125 species included in the analyses are given in Supplementary Materials 1 and 2. 126 For the purposes of our study, we have categorized ST in four characters: (i) 127 before or after copula, (ii) alone or in tandem, (iii) non-repeated or repeated, 128 and (iv) with or without precopulatory genital touching (see Table 1 and 129 Results). Since the description of the precopulatory genital touching might not 130 be stated explicitly in the revised papers, we established that when something 131 similar to this behaviour was described just before the ST, we considered it as 132 precopulatory genital touching.

133

134 Phylogenetic tree reconstruction

135 We aimed at understanding the evolution of ST using phylogenetic comparative

136 analyses (Harvey and Pagel 1991). Therefore, we first constructed a

137 phylogenetic tree including the odonate species for which data on ST were

138 available in the literature. We searched GenBank (Clark et al. 2016,

139 <u>https://www.ncbi.nlm.nih.gov/genbank/</u>) for sequences of the mitochondrial

140 genes COI and 16S, and the nuclear genes 18S and 28S. In total, we retrieved

sequence data for 129 species out of the 176 for which we had data on ST. We

also retrieved sequences of the same genes from 31 species without

143 information on ST, to increase the resolution of the tree, and for six species of

144 Ephemeroptera, to be used as outgroups in the analysis. The final dataset

145 included a total of 160 Odonata species (including 31 without data for ST),

146 which represented 14 families of Zygoptera and 11 families of Anisoptera (see

147 Supplementary Material Table S2).

148 All sequences were imported into Geneious version 9.1.7

149 (<u>http://www.geneious.com</u>, Kearse et al. 2012) for visual inspection before

alignment. Sequences were aligned using the ClustalW algorithm (Thompson et

al. 1994), as implemented in Geneious. We used BEAST version 2.4.8.0

152 (Bouckaert et al. 2014) to reconstruct the phylogenetic relationships among the

153 study species. The phylogenetic reconstruction was performed with the

154 nucleotide substitution models selected for each gene by the bModelTest

155 package version 1.0.4 (Bouckaert and Drummond 2017), and a strict clock 156 model and the Yule speciation model as priors. The analysis was run for 10 157 million generations and sampled every 1,000 generations. The output was 158 examined with Tracer version 1.6 (Rambaut et al. 2014) to assess convergence 159 of the Markov-chain Monte Carlo onto a stationary distribution through the 160 analysis of trace plots and effective sample sizes (ESS) of the model 161 parameters (ESS > 200 was considered acceptable). TreeAnnotator version 162 2.4.8, included in the BEAST package, was used to build a maximum clade 163 credibility tree after discarding 10% of sampled trees as burn-in. The consensus 164 tree was visualized and edited with FigTree version 1.4.2 (available at: 165 http://tree.bio.ed.ac.uk/software/figtree/). This tree was pruned to contain only 166 those species with available ST information using the R package ape (Paradis 167 et al. 2004), and the pruned tree was subsequently used for the phylogenetic 168 comparative analyses.

169

170 *Phylogenetic comparative analyses*

171 We estimated the ancestral states of our study discrete characters related to 172 sperm transfer using the function ace in the R package ape (Paradis et al. 173 2004). Our character states were ST performed (i) before vs. after copula; (ii) in 174 tandem vs. alone vs. both in tandem and alone; (iii) with repetition vs. with no 175 repetition vs. with and without repetition and (iv) with precopulatory genital 176 touching vs. without precopulatory genital touching vs. with and without 177 precopulatory genital touching (Robertson and Tennessen 1984). We note that 178 the combined states (i.e. both in tandem or alone) correspond to intra- or 179 interpopulation variation of the ST behaviour. We used maximum likelihood 180 estimation with equal rates of transition (Pagel 1994) and the likelihood of the 181 ancestral states was computed using a joint estimation procedure (see Pupko et 182 al. 2000). Finally, we also investigated possible correlated evolution between 183 pairs of characters. For this analyses, we first recoded the study characters to 184 binary. We coded the combined states to the least common variant of ST for 185 each character. Our rationale behind this decision is that we wanted to give 186 more weight to the least common behaviours, which are already under-187 represented in our dataset. Even though species with more than one type of ST

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- 188 have not stably evolved only one ST behaviour, they could offer hints on the
- evolution of ST across odonates. We used Pagel's (1994) function to detect
- 190 correlated evolution as implemented in the R package phytools (Revell 2012),
- 191 with the function fitDiscrete from the package geiger (Harmon et al. 2008).
- 192

193 **Results**

194 Variability of ST behaviour

- 195 We found a total of 123 papers that provided information on ST from 176
- 196 species of odonates, belonging to 22 families (Table 1, Supplementary
- 197 Materials 1 and 2). The ST is a behaviour that usually lasts for some seconds
- but shows high variability (mean \pm SE: 16.7 \pm 2.7 sec, range: 0.2 to 150 sec, N
- 199 = 82 species) (Table 1).
- 200 At least five different variants of ST were described across species: (i) male
- alone, before the formation of precopulatory tandem; (ii) in precopulatory
- tandem, without genital touching, only once per copula; (iii) in precopulatory
- 203 tandem after genital touching, only once per copula; (iv) in tandem, repeated
- 204 during copulation; and (v) male alone, after copula (see Table 1). However, in
- some species [e.g. Libellula quadrimaculata (L., 1758) or Erythemis
- simplicicollis (Say, 1839)] we found a combination of the ST variants due to
- 207 intra- or interpopulation variability (Figure 2).
- 208 Overall, the majority of species (66%) perform ST in precopulatory tandem
- 209 (Table 1), hence before copulation and only once per copula. Anisoptera males
- 210 perform ST alone, before tandem formation in a higher proportion (31%)
- 211 compared with Zygoptera (9%). The ST in tandem and repeated was found for
- 212 4% of species of Zygoptera, but no in Anisoptera (Table1). Finally, ST alone,
- after copula was found for 6% of the species of Zygoptera and 3% of Anisoptera
- 214 (Table 1).
- 215 The families Coenagrionidae and Libellulidae, are the most speciose and also
- 216 display more variants of ST behaviour, including all five in the first family. The
- 217 majority of coenagrionids studied performed ST in precopulatory tandem, and
- 218 only once per copula (58%; Table 1). The second most common variant was the
- 219 ST in tandem, but only after precopulatory genital touch (32%; Table 1). In
- libellulids, ST in tandem, only once per copula, was also the commonest variant

(61%; Table 1), and the second commonest ST was male alone, but in this
case, before tandem formation (33%; Table 1). ST by the male alone, after
copulation is a rare behaviour, but is the dominant behaviour in the Euphaeidae
(57%) and Polythoridae (67%) and it is also performed by the only living

225 representative of Pseudolestidae.

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227 Evolution of ST behaviour

DNA sequences were available for 129 out of the 176 species for which
information on ST behaviour was available in the literature (Supplementary
Material Table S2). The consensus phylogenetic tree was congruent with the
currently accepted and relatively well-established phylogeny of the order (*e.g.*Lorenzo-Carballa and Cordero-Rivera, 2014, but see Dijkstra et al. 2013), and
the support for the main clades was high in most cases (Supplementary
Material Figure S1).

- In our dataset, the majority of species perform ST before copula (Figure 2). Six
 species perform ST after copula (four Zygoptera and two Anisoptera).
- 237 According to the ancestral reconstruction for this trait, the variant of ST after
- 238 copula has evolved independently in all cases except for the group
- 239 Euphaea/Anisophaea (Zygoptera). Regarding ST in tandem and/or alone, most
- 240 species perform it in tandem, while 22 species do it alone (seven Zygoptera and
- 241 15 Anisoptera), and nine species use both strategies (three Zygoptera and six
- 242 Anisoptera). These variants have independently evolved in all cases except for
- 243 the group Euphaea/Anisophaea (Zygoptera), and the clade
- 244 Nannothemis/Pachydiplax/Erythemis and for the species of Leucorrhinia
- 245 (Anisoptera). In the case of ST performed repeatedly or not, we found that the
- 246 majority of species perform it just once. Only within Zygoptera, we found three
- 247 species that repeat sperm transfer and two that perform both variants
- 248 (corresponding to five independent origins). Regarding ST performed with or
- 249 without genital touching, only five species have ST with genital touching and
- three perform both variants (all of them within Zygoptera). These cases
- 251 correspond to independent origins except for the group of Ischnura ramburii, I.
- 252 graellsii and I. elegans. In summary, the ancestral state of ST would be before
- 253 copula, in precopulatory tandem, non-repeated and without genital touching.

Finally, we did not find any significant correlated evolution between the different variants of ST (P > 0.05).

256

257 Discussion

Our review of the literature indicates that there are at least five variants of ST
behaviour among Odonata. Our work also highlights the lack of basic
information on the ST behaviour for the vast majority of species within this
insect group: the total number of odonate species is estimated around 6000
(Lorenzo-Carballa and Cordero-Rivera, 2014), however we only found data on
ST behaviour for 176 species.

264 Variation of ST behaviours

In most of the odonate species studied, males perform ST after having grasped
the female in precopulatory tandem and once per copula (variant ii). This variant
of ST is the ancestral state, according to our comparative phylogenetic
analyses. Variant (i), the completion of ST by the male alone, before finding a
female (Table 1), could be advantageous in species where females are rarely
encountered and copulations are brief. Males that performed ST alone would be
ready to copulate as soon as they grasp a female in tandem.

272 Variant (iii), involves performing ST only after precopulatory genital touching

273 (Table 1), a behaviour that presumably signals female receptivity to the male

274 (Robertson and Tennessen, 1984). This behaviour might derive from the basal

behaviour (*i.e.*variant ii), if males commonly encounter unreceptive females.

276 High densities of unreceptive females would be predicted in populations/species

whose females remain near the reproductive site in the maturation phase, like

278 many *Ischnura* (Cordero et al., 1998). When females give no refusal signs,

279 males can remain in tandem for very long periods, even for a full day in the

280 laboratory (Cordero et al. 1992). Genital touching would therefore be adaptive

for males, avoiding wasting sperm with unreceptive females, but also for

females, because they would be released faster if signalling their

283 unreceptiveness. In odonates, it has been assumed that females cannot be

forced to copulate (e.g. Fincke 1997), although forced matings are possible in

285 populations with high male densities and females ovipositing unguarded

286 (Cordero-Rivera and Andrés 2002).

287 Only two species, *Coenagrion scitulum* (Rambur, 1842) and *Megaloprepus* 288 *caerulatus* (Drury, 1782), are known to routinely perform variant (iv), i.e. 289 repeated ST in one copulation (Table 1, Figure 1D). This behaviour has been 290 also recorded occasionally in three other zygopteran species: Lestes barbarus 291 (Fabricius, 1798) (Lestidae), *Ischnura aurora* (Brauer, 1865) (Coenagrionidae) 292 and Perissolestes remotus (Williamson & Williamson, 1924) (Perilestidae) (see 293 Supplementary Material 1). In the case of C. scitulum, the repetition of ST 294 during the copulation has been interpreted as a mechanism evolved in the 295 context of sperm competition (Córdoba-Aguilar and Cordero-Rivera 2008). In 296 this species, males are apparently unable to remove a significant portion of 297 sperm from rivals using their genital ligula (Cordero et al. 1995). Therefore, by 298 repeating ST and insemination, they might over-compete rival sperm. However, 299 other species of odonates [e.g. Hypolestes trinitatis (Gundlach, 1888)] are also 300 known to have limited ability to remove sperm, but ST is not repeated (Torres-301 Cambas and Cordero-Rivera 2011), and hence the link between both 302 phenomena is not straightforward.

303 The most intriguing variant for ST is variant (v), *i.e.* the translocation of sperm 304 by the male alone, after copulation (Figure 1F, Table 1). This behaviour was 305 first reported for the coenagrionid Mortonagrion hirosei Asahina, 1972, from 306 Japan (Naraoka 2014), and then observed in nine additional species, including 307 four Euphaeidae, and Pseudolestes mirabilis Kirby, 1900 (Cordero-Rivera and 308 Zhang 2018). Translocating sperm after copulation could be explained if these 309 species have evolved physiological mechanisms to maintain sperm alive for 310 long periods of time (until next mating). However, at least for the first copula, 311 males have to fill their sperm vesicle before copulation. Males could routinely 312 perform ST each morning to be prepared for the next mating, but this has never 313 been observed in *P. mirabilis*, the only species studied in detail (Cordero-Rivera 314 and Zhang 2018).

- 315 In some species, we found several of the ST variants due to intra- or
- 316 interpopulation variability. For example, males of *Hetaerina americana*
- 317 Fabricius, 1798 were reported performing ST alone before copula in one
- population, in tandem without precopulatory touch in another population and in
- tandem with precopulatory touch at a third locality (Supplementary material

320 Table 1). Two ST variants were also reported for different populations of *I*. 321 aurora and Libellula quadrimaculata Linnaeus, 1758, and between males of a 322 single population of Aeshna cyanea Muller, 1764 (Supplementary material 323 Table 1). Intraspecific variability seems rare, and unlikely to be detected in 324 short-term studies. For instance, only 5% of the 137 ST observed in two 325 Enallagma species occurred before tandem formation (Logan, 1971; cited by 326 Corbet, 1999). The reasons behind this intraspecific diversity of ST are 327 unknown, but these species are excellent candidates for further investigation of 328 the diversification of ST strategies within odonates.

329 Why has ST evolved?

330 The evolution of ST behaviour is likely related to the atypical copulation position 331 in the Odonata. One plausible scenario for the evolution of ST behaviour is that 332 ancestors of modern odonates produced a spermatophore, and deposited it on 333 the substrate, a behaviour currently observed in arachnids, myriapods and 334 wingless hexapods (Proctor 1998). The thick cerci of Namurotypus Brauckmann 335 and Zessin, 1989 males (ancestor of the carboniferous Odonata) could have 336 been used to firmly grasp the female behind her compound eyes. The male 337 could then have directed the female over the spermatophore (Bechly et al. 338 2001), in a way similar to the 'drag off' behaviour observed in whipspiders 339 (Amblypygi) (Weygoldt 1969). Attaching the spermatophore to the male body is 340 clearly more efficient, and could be the selective pressure required for the 341 evolution of secondary genitalia in odonates (reviewed in Cordero-Rivera and 342 Córdoba-Aguilar 2010), and hence, the ST behaviour.

An alternative, yet more speculative hypothesis for the evolution of ST could be related to sexual cannibalism, since females of several zygopterans are known to sometimes attack and eat mature conspecific males (Cordero 1992). We are aware of only one case of sexual cannibalism described during copulation:

Robertson (1985) observed a female of *Ischnura ramburi* (Selys, 1850) that was
repeatedly chased and grasped by a male until she finally attacked him and ate
out his thorax, but the male succeeded initiating copulation before dying in
copula. Other predatory animals, like spiders or octopuses, have a specialised
appendix used as an intromittent organ to introduce the spermatophore in the

352 female reproductive organs. These two groups also show sexual cannibalism

353 (Ibáñez and Keyl 2010; Li et al. 2012). Therefore, it could be hypothesised that 354 the tandem and wheel position of odonates during copulation, along with the 355 intra-male ST, might allow males to avoid sexual cannibalism (Chapman et al. 356 2003; Schneider 2014). From this idea, we propose the hypothesis that when 357 the risk of sexual cannibalism is high, selection favours the evolution of 358 secondary mechanisms to inseminate females safely for the males. This could 359 be tested with a review of the insemination behaviours across different animal 360 taxa and within a phylogenetic context.

361 Conclusions

362 Some of the ST variants seem to have evolved several times, but this behaviour 363 needs further investigation in a larger number of species. No other insect group 364 shows a behaviour equivalent to the ST of odonates (although some similarities 365 can be found with the pedipalps of spiders and the hectocotylus of octopuses), 366 and consequently we lack comparative evidence to understand the evolution of 367 this enigmatic behaviour. ST is sometimes performed very fast and therefore 368 careful video recording of this behaviour, particularly in Anisoptera, is needed to 369 avoid confusing movements to clean the abdomen with true ST. We currently

370 lack information for a number of families of both Anisoptera (e.g.

371 Austropetaliidae, Neopetallidae and Macromiidae) and Zygoptera (e.g.

372 Chlorocypgidae, Amphipterygidae and Lestoideidae), and most of the species

373 with information on ST are from temperate zones and fewer from tropical areas.

374 This bias can be explained by the scarcity of field studies in tropical regions,

and also due to the low sexual activity observed in tropical species (e.g.

376 Sanmartín-Villar and Cordero-Rivera, 2016).

377 Our study emphasizes also the relevance (and scarcity) of detailed natural

378 history observations for most species. We expect that this review will encourage

the scientific community towards more research in diversity of reproductive

380 behaviours, with a special focus on tropical species. This applies not only to

odonates, but to other animal taxa. This information, combined with modern

- 382 molecular techniques and phylogenetic hypotheses, is fundamental to
- 383 understand relevant questions about behavioural evolution as well as
- 384 behavioural diversity (ethodiversity). This is a necessary step to increase

- awareness on the importance of conserving not only species, but alsobehaviours (Cordero-Rivera, 2017a).
- 387

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Figure captions

Fig. 1: Insects usually mate with both sexes facing opposite directions, like in butterflies (**A**: *Hipparchia* sp. Fabricius, 1807; Satyridae). However, in odonates (**B**; *Orthetrum brunneun* (Fonscolombe, 1837), Libellulidae), the sexes are orientated to the same direction, allowing flying in copula. The intra-male sperm translocation can take place after the male has grasped the female in tandem, like in (**C**) *Neurobasis chinensis* (Linnaeus, 1758) (Calopterygidae), with or without precopulatory genital touching. In some species, this behaviour is repeated up to seven times in the same copulation, like in (**D**) *Coenagrion scitulum* (Rambur, 1842) (Coenagrionidae). Finally, in some species males translocate sperm alone before copulation, like in (**E**) *Diphlebia lestoides* (Selys, 1853), or after copulation, very close to the female as in (**F**) *Euphaea masoni* Selys, 1879 (Euphaeidae). Pictures: A. Cordero-Rivera, except (E), courtesy of Reiner Richter.

Fig. 2: The variants of intra-male sperm translocation behaviour, mapped on a phylogeny of the studied Odonata. The ancestral behaviour of sperm translocation was estimated to be before copula, in precopulatory tandem, non-repeated and without genital touching. Family names have been added to the main branches of the tree.

Figure 1







The evolution and diversity of intra-male sperm translocation in Odonata: a unique behaviour in animals

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Author for correspondence: Anais Rivas-Torres, email: <u>arivasto@gmail.com</u>, address: ECOEVO Lab, Universidade de Vigo, Escola de Enxeñaría Forestal, Campus A Xunqueira, 36005 Pontevedra, Galiza, Spain, telephone: +34657885598 Table 1. Summary of the diversity of sperm translocation behaviour in the Odonata (N: number of species; %: percentage of variants

of ST behaviour).

	(i) Male alone, before precopulatory tandem		(ii) In precopulatory tandem, only once per copula		(iii) In precopulatory tandem, after genital touching, only once per		(iv) In tandem, repeated during copula		(v) Male alone, after copula		Total	Duration (sec)
Suborder and	N	0/	N	0/	copula	0/	N	0/	N	0/	N	Moon SE (N)
	IN	70	IN	70	IN CO	70	N E	70	N O	70	N 100	
Zygoptera	11	8.5	85	65.9	20	15.5	5	3.9	8	6.2	129	19.4±3.1 (67)
Calopterygidae	3	12.5	20	83.3	1	4.2	0	0	0	0	24	6.1±0.9 (13)
Coenagrionidae	2	3.3	35	58.3	19	31.7	3	5.0	1	1.7	60	20.7±4.8 (35)
Diphlebiidae	1	100	0	0	0	0	0	0	0	0	1	
Euphaeidae	2	28.6	1	14.3	0	0	0	0	4	57.1	7	
Hemiphlebiidae	0	0	1	100	0	0	0	0	0	0	1	3.5 (1)
Lestidae	0	0	14	93.3	0	0	1	6.7	0	0	15	39.7±9.0 (10)
Megapodagrionidae	1	14.3	6	85.7	0	0	0	0	0	0	7	13.3±2.2 (4)
Perilestidae	0	0	1	50.0	0	0	1	50.0	0	0	2	
Platycnemididae	2	40.0	3	60.0	0	0	0	0	0	0	5	
Platystictidae	0	0	1	100	0	0	0	0	0	0	1	1.3 (1)
Polythoridae	0	0	1	33.3	0	0	0	0	2	66.7	3	2.6 (1)
Protoneuridae	0	0	1	100	0	0	0	0	0	0	1	18.8 (1)
Pseudolestidae	0	0	0	0	0	0	0	0	1	100.0	1	
Synlestidae	0	0	1	100	0	0	0	0	0	0	1	6.0 (1)
Anisoptera	20	31.3	42	65.6	0	0.0	0	0.0	2	3.1	64	5.0±3.9 (15)
Aeshnidae	2	14.3	12	85.7	0	0	0	0	0	0	14	0.2 (1)
Chlorogomphidae	1	100	0	0	0	0	0	0	0	0	1	
Cordulegastridae	0	0	2	100	0	0	0	0	0	0	2	
Corduliidae	2	33.3	4	66.7	0	0	0	0	0	0	6	

Gomphidae	2	66.7	1	33.3	0	0	0	0	0	0	3	5.0 (1)
Libellulidae	12	33.3	22	61.1	0	0	0	0	2	5.6	36	5.3±4.6 (13)
Petaluridae	1	50.0	1	50.0	0	0	0	0	0	0	2	
Anisozygoptera	0	0.0	1	100.0	0	0.0	0	0.0	0	0.0	1	
							-					
Epiophlebiidae	0	0	1	100	0	0	0	0	0	0	1	