| 1 | Patterns of variation in distylous traits and reproductive consequences in <i>Erythroxylum</i> |
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| 2 | species and populations |
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| 12 | Variation in distylous traits of <i>Erythroxylum</i> |
| 13 | |
| 14 | PREMISE OF THE STUDY: Distylous species possess two floral morphs with reciprocal |
| 15 | positioning of stigmas and anthers that is hypothesized to promote disassortative pollination. |
| 16 | Theoretical models predict equal morph frequencies, but many populations depart from the |
| 17 | expected 1:1 ratio, a pattern that often correlates with asymmetric mating between morphs |
| 18 | |
| | and/or presence of weak incompatibility system. Variation in reciprocity can also affect the |
| 19 | and/or presence of weak incompatibility system. Variation in reciprocity can also affect the likelihood of disassortative pollination, and hence reproductive fitness. |
| 19 20 | and/or presence of weak incompatibility system. Variation in reciprocity can also affect the likelihood of disassortative pollination, and hence reproductive fitness. METHODS: We described variation in incompatibility systems and morph ratio in four |
| 19 20 21 | and/or presence of weak incompatibility system. Variation in reciprocity can also affect the likelihood of disassortative pollination, and hence reproductive fitness. METHODS: We described variation in incompatibility systems and morph ratio in four <i>Erythroxylum</i> species to test if greater deviations from 1:1 ratios occur in populations of self- |
| 19 20 21 22 | and/or presence of weak incompatibility system. Variation in reciprocity can also affect the likelihood of disassortative pollination, and hence reproductive fitness. METHODS: We described variation in incompatibility systems and morph ratio in four <i>Erythroxylum</i> species to test if greater deviations from 1:1 ratios occur in populations of self- compatible species. We used adaptive inaccuracy and described upper and lower organ |

population means and coefficients of variation for fruit set to test if reciprocity could predict
female reproductive success.

26 KEY RESULTS: Morphs occurred in 1:1 ratios in most populations of three Erythroxylum species with distylous self-incompatibility. In self-compatible E. campestre populations 27 28 showed an excess of the long-styled morph, the short-styled morph, or were monomorphic for 29 the short-styled morph. We detected deviations from reciprocity, with total inaccuracy 30 ranging between 9.39% and 42.94%, and inaccuracy values were lowest in low organs. 31 Across populations, we found a positive relationship between inaccuracy and the coefficient 32 of variation of fruit set. 33 **CONCLUSIONS:** *Erythroxylum* species showed variation in the distylous syndrome, with 34 changes in the incompatibility system that corresponded with deviations from 1:1 morph

35 ratio, and variation in reciprocity that correlated with variation in female reproductive fitness.

36 **KEY WORDS:** breeding systems; Cerrado; disassortative mating; Erythroxylaceae;

37 heterostyly; isoplethy; monomorphism; morph bias; reciprocal herkogamy; sex

38 polymorphism.

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40 Heterostyly is a genetically controlled sex polymorphism described in 28 angiosperm 41 families, where plant populations possess two (distyly) or less commonly three (tristyly) floral 42 morphs (Barrett and Shore, 2008). In distylous species, the long-styled morph (L-morph) 43 presents the stigma above the anthers, while the stigma of the short-styled morph (S-morph) 44 appears below the anthers. Thus, floral morphs differ from one another in the positions in 45 which anthers and stigmas are reciprocally presented, an arrangement referred to as reciprocal 46 herkogamy (Barrett, 2002). Sex-organ reciprocity (L stigma-S anther and S stigma-L anther) 47 functions to promote cross-pollination between morphs (disassortative pollination) because

48 pollinators are likely to pick up pollen from the morphs on different parts of their bodies and 49 transfer pollen to stigmas at those same heights on flowers of the opposite morph (Darwin, 50 1877; Lloyd and Webb 1992a, b; Barrett, 2002). Together with reciprocal herkogamy, 51 heterostylous species commonly possess a heteromorphic self-incompatibility system 52 preventing self-fertilization and intra-morph cross-fertilization (Barrett and Richards, 1990), 53 and morph-specific ancillary characters (Ganders, 1979b; Dulberger, 1992). Hence, the 54 combination of herkogamy with reciprocally positioned high and low sexual organs should 55 reduce self-interference and pollen wastage on incompatible stigmas, and increase male 56 fitness (Barrett, 2002). At equilibrium, populations of distylous species are expected to show 57 a 1:1 ratio of style morphs (Pannell et al., 2005). Populations of many distylous species, however, show L- or S-biased morph ratios or populations fixed for a morph. These 58 59 conditions are associated with the breakdown of the distylous incompatibility system and with 60 random stochastic events (Eckert and Barrett, 1992; Zhou et al., 2012, 2017). In addition, the 61 presence of intra-morph compatibility and changes in the rates of disassortative pollen 62 transfer can result in deviations of the morph ratio (Barrett and Hodgins, 2006; Brys et al., 63 2008; Hodgins and Barrett, 2008; Pérez-Barrales and Arroyo, 2010; Consolaro et al., 2011; 64 Zhou et al., 2017).

65 An investigation of variability in the position of anthers and stigmas is an initial step 66 to interpret the ecological function and selection on the morphological features of the 67 distylous floral syndrome (Charlesworth and Charlesworth, 1979; Lloyd and Webb 1992a, b). 68 While reciprocity is based on the notion that reciprocal organs are placed at the same height, 69 heterostylous species display variation in high and low organ reciprocity, with anthers and 70 stigmas often deviating from the position of their "compatible" reciprocal organ (Pailler and 71 Thompson, 1997; Thompson and Dommée, 2000; Faivre and McDade, 2001; Pérez et al., 2004; Ferrero et al., 2009, 2011a, b; Keller et al., 2012; Sá et al., 2016; Armbruster et al., 72

73 2017). Deviations from perfect reciprocity (defined as exact similar position of reciprocal 74 high and low reproductive organs) can lower the probability of disassortative pollination, reduce fruit and seed production, or even facilitate a breakdown of the floral polymorphism 75 76 (Keller et al., 2014; Zhou et al., 2015; Jacquemyn et al., 2018; Wu et al., 2018; Brys and 77 Jacquemyn, 2019; but see Simón-Porcar et al., 2015). Under the assumption that the function 78 of heterostyly relies on the close matching between reciprocal organs (Darwin, 1877; 79 Jacquemyn et al., 2018; Brys and Jacquemyn, 2019), a morphological analysis of reciprocity 80 can help elucidate whether heterostylous species and populations are likely to have stable 81 reproductive systems. Armbruster et al. (2017) developed a metric to measure reciprocity 82 based on adaptive inaccuracy, with the aim of associating variation in reciprocity with 83 reproductive fitness. For distylous species, inaccuracy is based on the concept that the 84 adaptive optimum of an anther level is represented by the population mean of the reciprocal 85 stigma and vice versa, so that if all anthers and stigmas of a population are at the same height, 86 inaccuracy will be zero. Hence, inaccuracy is defined by the distance between the average 87 organ height with regards the population mean of its reciprocal organ, and the population 88 variance of each organ (Armbruster et al., 2009, 2017). Long and S individuals with stigma 89 heights similar to their compatible anther should experience higher disassortative pollen 90 transfer and hence higher seed or fruit production (Armbruster et al., 2017; Jacquemyn et al., 91 2018; Brys and Jacquemyn, 2019). Hence, populations of heterostylous species with lower 92 inaccuracy in reciprocity should have higher female reproductive success, as well as low 93 variability in female reproductive success. Therefore, a negative relationship between 94 inaccuracy and female fitness is expected, and a positive relationship between inaccuracy and 95 female fitness variation is predicted. To our knowledge, population level associations between 96 reproductive success and inaccuracy remain to be tested.

97 Erythroxylaceae is a heterostylous family with four tropical genera and ca. 260 species 98 of small trees and shrubs (White et al., 2019). The genus Erythroxylum P. Browne is the 99 largest of the family (ca. 250 species; White et al., 2019) and endemic to the neotropical 100 region. The main center of diversity is Brazil, with 114 species of which 74 are endemic 101 (Plowman and Hensold, 2004; Loiola et al., 2007), representing a monophyletic group within 102 the genus (White et al., 2019). Although knowledge of the reproductive biology of species is 103 scarce, *Erythroxylum* species show variation in breeding system, with heterostylous species 104 displaying variation in the strength of heteromorphic incompatibility system and population 105 morph ratio (Burck, 1895; Arroyo and Cabrera, 1978; Ganders, 1979a; Domínguez, 1990; 106 Berry et al., 1991; Domínguez et al., 1997; Barros, 1998; Pailler et al., 1998; Bianchi et al., 107 2000). A feature of Erythroxylum flowers that could affect reciprocity of anthers and stigmas 108 is the presence of two stamen whorls, which may or may not show similar length within the 109 flower.

110 In this study we investigated four species of *Erythroxylum* in Brazil to (i) describe the 111 incompatibility system of each species; (ii) describe variation in morph ratio among species 112 and populations; (iii) quantify the variation in the position of anthers and stigmas to 113 investigate inaccuracy of high and low reciprocal organs; and (iv) study the natural fruit set of 114 the morphs in different populations. These data were used to test the following predictions: 115 first, because previous research on *Erythroxylum* species indicated some degree of self-116 compatibility (Arroyo and Cabrera, 1978; Ganders, 1979a; Berry et al., 1991; Barros, 1998), 117 we predicted greater deviations from 1:1 morph ratio in self- and intra-morph compatible 118 species than in species with heteromorphic incompatibility system. Second, because 119 deviations from perfect reciprocity appear associated with lower disassortative pollen 120 deposition, which in turn should reduce fruit production (Jacquemyn et al., 2018; Brys and 121 Jacquemyn, 2019), we predicted a negative relationship between inaccuracy and average fruit

set at the population level. Also, we predicted a positive relationship between inaccuracy and the coefficient of variation of fruit set as increases in inaccuracy would be associated with higher uncertainty in disassortative pollen deposition.

125 MATERIAL AND METHODS

126 Species and study areas

127 The study species, E. campestre A.St.-Hil., E. deciduum A.St.-Hil., E. suberosum A.St.-Hil. 128 and E. tortuosum Mart., are found in the Brazilian Cerrado. In these species flowers are small 129 (ranging from 2,8 mm length in *E. campestre* to 4,5 mm in *E. suberosum*) with creamy white 130 petals (Fig. 1), and visited by various species of bees and wasps, or less frequently, by 131 butterflies and flies (Barros, 1998; R. Matias, University of Brasília, personal observation). At 132 the base, petals possess a nectariferous ligule appendage forming a small tube. Flowers have 133 ten stamens with filaments fused at the base, forming a short staminal tube around the 134 superior ovary. Stamens are organized in two whorls with five stamens each, placed opposite 135 to the sepals and petals. In the study species, the two stamen whorls in L flowers appear at 136 different heights but occur at similar heights in S flowers (Fig. 2; other Erythroxylum species 137 show variation in the two stamen whorls of L and S flowers; Pailler et al., 1998). Flowers of 138 both morphs are tricarpellate with only one fertile locule containing a single fertile ovule 139 (Loiola et al., 2007; Silva et al., 2016). In the present study, eleven Cerrado areas from 140 Central Brazil were selected for the population survey. Details on region, area name and 141 coordinates are included in Appendix S1 (see the Supplementary Data with this article). In 142 total, 26 populations were surveyed. Species and population surveys were completed during 143 the main flowering season between August and November of 2015 and 2016.

144 Incompatibility system

145 Using hand-pollinations, the incompatibility system was assessed in one population per 146 species in 2016 (populations 3, 10, 16 and 24; located in the same area). Between 16-19 147 flowering individuals per species were tagged, all of which were exposed to the following 148 treatments: (1) facilitated self-pollination using pollen from the same flower, (2) intra-morph 149 and (3) inter-morph cross-pollination by applying pollen from different plants of the same and 150 different morph respectively, (4) autonomous self-pollination by bagging unopened flowers, 151 (5) agamospermy by emasculating flowers and bagging them to prevent insect visitation, and 152 (6) control treatment as open pollination under natural pollination conditions. To describe the 153 incompatibility system a total of 2-7 flowers were used per treatment and on each individual 154 such that all plants used in the experiment received all treatments. Flowers were bagged 155 before and after the manipulations with small tulle bags to prevent uncontrolled pollen arrival, 156 except for the control treatment. Fruit development was assessed after two weeks. 157 Erythroxylum flowers produce a single seed so that fruit set corresponds to seed set. Data 158 were analysed using generalized linear mixed-effects models (hereafter GLMMs) to 159 investigate the effect of treatment, morph and interaction term on fruit set (number of fruits 160 formed/number of flowers used in each treatment per individual). Individual plants were 161 included as a random factor. Fruit production was modelled with a binomial distribution using 162 the probit function. The analyses were conducted using the glmer function in the lme4 163 package in R (Bates et al., 2015). To obtain the significance of each factor, a type-II analysis 164 of variance was conducted using the Anova function in the car package (Fox, 2015). The 165 Ismeans function with a Tukey adjustment from the package emmeans (Lenth, 2016) was 166 used as a post hoc test to detect significant differences between treatments. All statistical 167 analyses in this study were performed in the R statistical environment (R Core Team, 2019).

168 **Population morph ratios**

169 Population morph ratios were assessed during surveys conducted in 2015. In each area, all

170 plants intercepted along a 1000-2000m. transect were classified as L- or S-morph. Because

171 the areas investigated included at least two *Erythroxylum* species (see Appendix S1),

172 flowering individuals were verified to identify the species and floral morph. *Erythroxylum*

173 individuals do not appear to propagate clonally and the sampling of genets was unambiguous.

174 In all populations, chi-square (x^2) analysis with Yates correction was used to assess whether

175 population morph ratios departed from isoplethy (1:1), using $\alpha = 0.05$.

176 Floral morphology and reciprocal herkogamy

177 During the survey of population morph ratio, flowers were collected (n = 1-3 flowers per 178 individual plant; 18-49 individuals per population) and fixed in 70% ethanol for 179 measurements. Flowers were dissected, photographed and traits measured using ImageJ 1.45s 180 software (Rasband, 2011; http://imagej.nih.gov/ij/). The traits measured included stigma and 181 anther height, using as a reference point the top of the ovary (Fig. 2). Anther height of each 182 anther level was measured in the L-morph (Fig. 2). Differences between morphs and 183 populations in stigma and anther height per species were analysed with linear mixed-effects 184 models (hereafter LMMs) using the lmer function in the lme4 package (Bates et al., 2015). 185 Morph, population and interaction term were included in the models as fixed factors, and 186 individual plants as a random factor. Significance of each factor was tested with type-II 187 analysis of variance as described above. Residual analysis was used in order to ensure model 188 assumptions were met (homoscedasticity and normal distribution). Because the aim of the 189 analysis consists of detecting differences between morphs, we excluded monomorphic and 190 highly anisoplethic populations.

Population mean and variance in anther and stigma height were used to investigate
adaptive inaccuracy in reciprocity as described by Armbruster et al. (2017). In heterostylous
species, the phenotypic optimum of L stigmas corresponds to the population mean of S anther

194 height, and the phenotypic optimum of S anthers corresponds to the population mean of L 195 stigma height. Similarly, the phenotypic optimum of S stigmas corresponds to the population 196 mean of L anther height, and the phenotypic optimum of L anthers corresponds to the 197 population mean of S stigma height (Darwin, 1877; Barrett 2002; Armbruster et al., 2017). 198 Inaccuracy in reciprocity is then estimated as the contribution of bias (differences in mean 199 height of reciprocal organs) and imprecision (variance in organ height) to departures from 200 close matching between reciprocal organs (Armbruster et al., 2009, 2017). We used the 201 population mean and variance of each organ type to calculate inaccuracy values for high (L 202 stigmas [S] and S anthers [A]; Eqn 1) and low organs (L anthers [a] and S stigmas [s]; Eqn 2). 203 Because L flowers have two stamen whorls of different lengths (Fig. 2), the low organ level 204 inaccuracy was estimated in two ways. The height measures of the two anther levels (aA and 205 aa for upper and lower anther level, respectively) were analyzed together in the same equation 206 (Eqn 2) by using the population mean and variance of the L anthers without distinction of 207 anther position. In addition, the contribution of each anther level to inaccuracy in reciprocity 208 was estimated by using upper (Eqn 3; aA) and lower anthers (Eqn 4; aa) separately. We 209 generated these two inaccuracy estimates because it is unknown whether one anther level 210 contributes more to the pollination of S stigmas, or whether both contribute equally. Hence, 211 the comparison of inaccuracy values obtained without distinction between the upper and 212 lower anther (a), and those estimated with the upper (aA) and lower (aa) L anther can be 213 helpful to infer the potential role of the two anther levels in disassortative pollen transfer 214 (Thompson et al., 2012). In the equations presented below, letters with bars correspond to the 215 population mean of organs whereas V corresponds to the variance of the organs (Armbruster 216 et al., 2017).

217 Inaccuracy_{high organs} =
$$(\overline{A} - \overline{S})^2 + V_A + V_S$$
 Eqn 1

218 Inaccuracy_{low organs with both L anthers} = $(\overline{a} - \overline{s})^2 + V_a + V_s$ Eqn 2

- 219 Inaccuracy_{low organs with upper L anthers} = $(\overline{aA} \overline{s})^2 + V_{aA} + V_s$ Eqn 3
- 220

Inaccuracy_{low organs with lower L anthers} = $(\bar{aa} - \bar{s})^2 + V_{aa} + V_s$ Eqn 4

221 The unit of inaccuracy is trait units squared, and the results of high (Eqn 1) and low 222 organ inaccuracies (Eqn 2) were summed to provide a total inaccuracy value per population. 223 Equations 3 and 4 represent inaccuracy estimates for low organs using the upper (aA) and 224 lower (aa) anther of L flowers. To compare levels of inaccuracy between populations and 225 species, values were standardized by the squared mean of all anther and stigma heights 226 recorded for each population and adjusted to a proportional scale (Armbruster et al., 2017). 227 Low values of inaccuracy correspond to high levels of reciprocity, e.g., close matching 228 between reciprocal organs with low dispersal around the optimal values.

229 Population variation of natural fruit set and relationship with inaccuracy

230 Natural fruit production was studied in 12 of the 26 populations (2-4 populations per species) 231 in 2015. Ten to 27 individual plants were randomly selected in each population, and 4-102 232 flowers on each plant were tagged and left exposed to pollinators. Fruit development was 233 inspected 3-4 weeks later. Differences among populations in fruit set (number of fruits 234 formed/number of flowers marked) were analysed with a GLMM using the glmer function 235 with a binomial distribution and probit function. This analysis included the terms population, 236 morph and the interaction effect; individual plant was included as random factor. Significance 237 of each factor was tested with type-II analysis of variance; when a factor was statistically 238 signifficant, post hoc tests were conducted as described above. Sequential Bonferroni 239 correction was used to account for multiple tests.

Using the standardized inaccuracy values, LMMs were used to evaluate the relationship between high (Eqn1) and low (Eqn 2) organ inaccuracy with female reproductive success. Three populations of the self-compatible species were excluded from the analyses 243 because fruit set can result from either self- or cross-pollination (see results below). 244 Specifically, two analyses were conducted using inaccuracy values as the independent 245 variable and the mean and the coefficient of variation of fruit set for each population and 246 morph as dependent variables. In these analyses, species and population nested within species 247 were included as random effects in the models. To obtain the statistical significance of the 248 models, we performed a type-II analysis of variance. Because the sample size was small (only 249 nine populations and a total sample size of 18, including two morphs per population), the term 250 morph was not included in the analyses (a model incorporating morph showed that this term 251 was not statistically significant in the analyses, results not shown).

252 **RESULTS**

253 Incompatibility system

254 In E. campestre autonomous self-pollination and all hand pollination treatments rendered 255 similar fruit set (P> 0.875 for all comparisons), while agamospermy and open pollination 256 treatments produced lower fruit set (P < 0.006 for all comparisons; Table 1 and Fig. 3). In the 257 rest of species, fruit set after the inter-morph cross-pollination treatment was much higher 258 than all the other treatments (P<0.006 for all comparisons; Table 1 and Fig. 3), with the 259 exception of the open pollination treatment in E. deciduum, suggesting absence of pollen 260 limitation for this species. In all cases, there were no statistically significant differences 261 between morphs (Table 1). Taken together, these results indicate that E. campestre is a self-262 compatible species, whereas E. deciduum, E. suberosum and E. tortuosum present the typical 263 heteromorphic self-incompatibility of distylous species.

264 **Population morph ratios**

Of the eight populations of *E. campestre*, three were isoplethic (had equal numbers of L and S
plants) and five showed variation in the morph ratio, with one population fixed for the S-

morph, two populations biased for the L-morph and two populations biased for the S-morph
(Table 2; Fig. 4). Populations of the remaining species displayed 1:1 morph ratio, except for
one L-biased population of *E. suberosum* and one L-biased population of *E. tortuosum* (Table
270 2; Fig. 4).

271 Floral morphology and reciprocal herkogamy

272 All species showed statistically significant differences in the position of anthers and stigmas 273 between morphs (Tables 3, 4), and all species other than E. deciduum showed differences 274 between populations. For stigma height, the interaction term morph * population was 275 statistically significant for all species except for *E. deciduum* (Table 4). The interaction term 276 for the upper anther of the L-morph was statistically significant and marginally significant for 277 E. campestre and E. deciduum respectively, while the interaction effect using the lower L 278 anther was significant in all species (Table 4). These results suggest that populations of 279 species harbor substantial variation in anther and stigma height.

280 Values of standardized total inaccuracy calculated without distinction between the L 281 anthers whorls ranged between 9.39% and 42.94% (Table 5). Erythroxylum suberosum and E. 282 tortuosum displayed the lowest inaccuracy values per species, ranging between 9.39% and 283 20.01% in E. suberosum and between 13.93% and 21.64% in E. tortuosum (Table 5). 284 Inaccuracy values in E. deciduum were the largest of all species, ranging between 38.07% and 285 42.94%. Erythroxylum campestre showed substantial variation in inaccuracy (Table 5), with 286 populations showing low values (e.g., 13.48% in population 8) and high values (e.g., 41.16% 287 in population 5), covering the range displayed by all species. Low organ inaccuracy estimated 288 with the upper and lower L anther considered separatelly revealed different patterns. In E. 289 campestre, E. suberosum and E. tortuosum inaccuracy was smaller when S stigmas were analyzed against the upper anthers of L flower (s-aA), with values ranging from 0.70% to 290 291 9.08%. For these species, inaccuracy values estimated with the upper anther level ranged

between 4.01% and 22.51% (Table 5). In contrast, *E. deciduum* showed the opposite pattern,

with larger inaccuracy values for the upper anthers ranging between 53.28% and 71.52%, and

between 1.59% and 3.54% for the lower anthers. For *E. campestre*, *E. suberosum* and *E.*

295 tortuosum high organs displayed larger inaccuracy values than low organs, whereas low organ

inaccuracy in *E. deciduum* was larger than the high value. However, this appeared to result

from the large contribution to inaccuracy of the upper L anther (Table 5).

298 **Population variation of natural fruit set and relationship with inaccuracy**

299 Populations of *E. campestre* (the only self-compatible species) and *E. tortuosum* showed

300 similar patterns of fruit set, with no differences between morphs, populations or the

301 interaction effect (Table 1; Fig. 5). Erythroxylum deciduum displayed substantial variation in

302 fruit set among populations and for the morph * population term, with similar fruit set

303 between morphs in populations 9 and 12, and larger fruit set for the L-morph in population 10

304 (Table 1; Fig. 5). Fruit set in *E. suberosum* differed among populations, but no differences

305 were detected between morphs (the interaction term was not statistically signifficant; Table 1;

306 Fig. 5).

The relationship between mean fruit set of L and S plants and organ inaccuracy (using the low organ inaccuracy value without distinction between the upper and lower L anthers) was not significant ($x^2 = 0.253$; df = 1; P = 0.615). In contrast, there was a positive and significant relationship between the coefficient of variation of fruit set and measures of inaccuracy ($x^2 = 7.961$; df = 1; P = 0.005; Fig. 6), suggesting that greater inaccuracy reciprocity values correlate with greater variation in fruit set.

313 **DISCUSSION**

314 In the present study, we described the incompatibility system, variation in morph ratio and 315 upper and lower organ reciprocity in four distylous species of *Erythroxylum*, and used those 316 data to investigate associations between inaccuracy in sex-organ reciprocity and the 317 population mean and coefficient of variation in fruit set among populations. We predicted a 318 negative relationship between inaccuracy and fruit production, because low inaccuracy values 319 correspond to high reciprocity, which in turn should increase disassortative pollen transfer 320 and female fitness (Jacquemyn et al., 2018; Brys and Jacquemyn, 2019). While we did not 321 detect a negative relationship of inaccuracy with mean fitness as measured by fruit 322 production, we found a positive relationship between inaccuracy in reciprocity and the 323 coefficient of variation of fruit set. Although this result was based on a limited number of 324 populations, it suggests that the morphological variation of anthers and stigmas, and the 325 magnitude of overlap between reciprocal organs affects the function of distyly in 326 Erythroxylum.

327 We found that in most of the populations of the species with the typical heteromorphic 328 incompatibility system, namely E. deciduum, E. suberosum and E. tortuosum, L and S plants 329 occurred at equal frequency. In contrast, populations of *E. campestre* displayed variation in 330 morph ratio, including isoplethic, L-biased, S-biased and S-monomorphic populations. Unlike 331 other Erythroxylum species (Ganders, 1979a; Domínguez, 1990; Pailler et al., 1998; Bianchi 332 et al., 2000; Silva et al., 2007), E. campestre is self-compatible and all crosses formed seeds. 333 The 1:1 morph ratio in populations of distylous species is often associated with the presence 334 of a heteromorphic incompatibility system (Pailler and Thompson, 1997; Pailler et al., 1998; 335 Sá et al., 2016). In self-compatible species or intra-morph compatible species, promotion of 336 legitimate pollination and disassortative mating is also an important mechanism for the 337 maintenance of the polymorphism, and changes in mating patterns can drive changes in 338 morph ratio, and the fixation of a morph (Barrett and Hodgins, 2006; Pérez-Barrales and 339 Arroyo, 2010, Simón-Porcar et al., 2014; Zhou et al., 2015; Ferrero et al., 2017). 340 Unfortunately, we were unable to obtain pollinator observations or estimate pollen transfer

341 rates to understand if departures of the 1:1 ratio correlated with rates of legitimate pollination 342 and the function of distyly (but see discussion below), particularly in the self-compatible 343 Erythroxylum species. Variation in morph ratio could also be explained in the context of 344 stochastic population events, such as habitat fragmentation, reduction of population size or 345 founder events, which are often accompanied by changes in the incompatibility system and 346 selection of autonomous self-pollination (Eckert and Barrett, 1992; Brys et al., 2008; Costa et 347 al., 2016; Zhou et al., 2012, 2017). However, our population data per species is limited and 348 larger population surveys would be necessary to model the relationship between random 349 population events, breakdown of the incompatibility system and the loss of the polymorphism 350 in *Ervthroxylum*.

351 We used adaptive inaccuracy to quantify reciprocity. Adaptive inaccuracy uses the mean and the variance to interpret the adaptive significance of the position of anthers and 352 353 stigmas in relation to pollen pick-up and delivery (Armbruster et al., 2009, 2017), which is 354 critical for the function of heterostyly (Armbruster et al., 2006). In our analysis, when 355 averaged across multiple populations, the values at species level were higher than those 356 obtained in species of Primula L. (Primulaceae; Armbruster et al., 2017) and Pulmonaria L. 357 (Boraginaceae; Jacquemyn et al., 2018), suggesting that reciprocity in distylous Erythroxylum 358 species is lower and less accurate. Our results showed that high organs had larger inaccuracy 359 values than low organs. In *Primula* and *Pulmonaria*, high organs contributed more strongly to 360 inaccuracy, a pattern attributed to developmental variation, which is often greater in large 361 organs (Armbruster et al., 2017; Jacquemyn et al., 2018; Brys and Jacquemyn, 2019). In 362 flowers with long tubes and epipetalous stamens, like those in Primula and Pulmonaria, 363 anther height is usually influenced by floral development and the correlation between the stamens and the floral tube, which might then affect the placement of anthers and stigmas, 364 365 and therefore reciprocity, with larger effects for high organs if S anthers express more

366 variability in their height (Faivre, 2000; Thompson and Dommée, 2000; Faivre and McDade, 367 2001; Pérez-Barrales et al., 2014; Santos-Gally et al., 2015). Erythroxylum species present 368 open and relatively small corollas. Filaments are relatively free and only fused at the base of 369 the ovary and not to the corolla. While our results agreed with an apparent tendency of larger 370 inaccuracy values in high organs, as detected in both Primula and Pulmonaria, the 371 morphology of *Erythroxylum* flowers probably imposes low developmental restrictions, and 372 allows expression of more variability in organ height position compared to epipetalous 373 flowers. This might explain the relatively larger inaccuracy values compared to those in 374 Primula and Pulmonaria (Armbruster et al., 2017; Jacquemyn et al., 2018; Brys and 375 Jacquemyn, 2019). Future morphological studies of floral development will be useful to gain 376 insights into the importance of floral development in the expression of reciprocity (Faivre, 377 2000; Faivre and McDade, 2001).

378 In our study, inaccuracy values of low organs were influenced by the presence of two 379 anther whorls in the L-morph. Of all species, E. deciduum showed the greatest differences 380 between the two inaccuracy values for low organs (s-aA and s-aa), and consequently great 381 imprecision between S stigmas and the upper anthers of L flower (s-aA). We believe that high 382 inaccuracy values can decrease the amount of disassortative pollination, but pollination 383 studies are necessary to test this hypothesis in Erythroxylum species. In Narcissus assoanus 384 Dufour ex Schult. f., a style dimorphic species with intra-morph compatibility and with two 385 anther whorls in L and S flowers, female fertility of S flowers increased when pollen donors 386 were the low L anthers rather than the low S anthers, a fertility pattern explained by the closer 387 match of S stigmas with the low L anthers than the low anther level of S flowers (Thompson 388 et al., 2012).

The data presented here add additional evidence that heterostylous species often depart
from theoretical expectations of high reciprocity (Pailler and Thompson, 1997; Thompson and

391 Dommée, 2000; Faivre and McDade, 2001; Keller et al., 2012; Armbruster et al., 2017; 392 Jacquemyn et al., 2018). The ultimate goal of understanding the evolutionary significance of 393 heterostyly is to establish a relationship between variation in reciprocity, disassortative pollen 394 transfer and reproductive success. We found a positive and significant relationship between 395 inaccuracy and the coefficient of variation in fruit set. These findings agree with recent 396 experiments evaluating how inaccuracy in reciprocity influences disassortative pollination in 397 distylous Pulmonaria (Jacquemyn et al., 2018) and Primula veris L. (Brys and Jacquemyn, 398 2019). In these studies, disassortative pollen transfer increased in L and S flowers with greater 399 reciprocity (e.g. low inaccuracy). Consistent with pollen deposition patterns, seed set in P. 400 *veris* declined with increasing inaccuracy, indicating a strong link between disassortative 401 pollen deposition and seed set (Brys and Jacquemyn, 2019). It is therefore possible that larger 402 inaccuracy in *Erythroxylum* populations decreased legitimate pollen deposition, making 403 disassortative mating unpredictable, which in turn would result in greater fruit set variation. 404 This result shows that variation in reciprocity could potentially have far-reaching effects on 405 patterns of fruit production, and may even initiate or facilitate the breakdown of this floral 406 polymorphism, especially if these changes are accompanied by modifications in the 407 heteromorphic incompatibility system (Barrett and Shore, 2008).

408 We did not detect a relationship between inaccuracy and mean fruit set. Erythroxylum 409 species produce only one seed per flower, and in the presence of a heteromorphic 410 incompatibility system, a small amount of compatible pollen may be sufficient to fertilize a 411 single ovule. This contrasts with species with a larger ovule number, like Primula veris, 412 where disassortative pollen transfer correlated with seed set (Brys and Jacquemyn, 2019). 413 Alternatively, variation in fruit set could be explained in the context of pollen limitation 414 (Knight et al., 2005). For example, in the experiment to describe the incompatibility system, 415 fruit set after open pollination was smaller than the legitimate hand-pollination pollination

treatment (Fig. 3). The low fruit set could be explained under low visitation rate or inefficient
pollination service. Fruit production under natural open pollination was substantially different
among populations, which probably reflects geographic variation in the pollinator service.
Future research in *Erythroxylum* should incorporate pollinator observations to understand how
pollinator interactions contribute to disassortative pollen transfer. Ultimately, data on
pollinator behavior and efficiency can help to establish a clear relationship between
inaccuracy, pollen flow dynamics and the function of distyly in *Erythroxylum*.

423 In this study, we described the variation of the distylous syndrome in Erythroxylum 424 species and populations, and how this variation is partitioned between morphs. We conclude 425 that the most important features of typical distyly, namely the morphological syndrome and 426 heteromorphic incompatibility system, seem to be conserved in E. deciduum, E. suberosum 427 and E. tortuosum. The morphological syndrome is present in E. campestre, but this species is 428 self-compatible and populations displayed great variation in morph ration, including S-429 monomorphic populations. Variation in morph ratio and the presence of self-compatibility may indicate failure in the function of the polymorphism. Monomorphism might be a derived 430 431 condition, as reported in other distylous species and populations (Mast et al., 2006; Zhou et 432 al., 2012; Ruiz-Martín et al., 2018). As different clades of Erythroxylum generally occur in 433 distinct geographic areas (White et al., 2019), the presence of monomorphism in species that 434 occur in the Brazilian Cerrado and Northern Venezuela (E. undulatum; Berry et al., 1991) 435 possibly supports the hypothesis of multiple independent origins of monomorphism from 436 distylous ancestors. Taken together, our findings suggest that *Erythroxylum* is a promising 437 genus to further investigate the selection, function and evolution of heterostyly.

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447 AUTHOR CONTRIBUTIONS

448 All authors have made substantial contributions to conception, design and interpretation of

- 449 data. R.M. collected and analyzed the data. R.M. and R.P.B. led the writing. H.C. helped with
- 450 discussion and text revision.

451 DATA AVAILABILITY STATEMENT

All data needed to evaluate the conclusions in the paper are present in the paper and/or the
Supplementary Materials. Additional data related to this paper may be requested from the
authors.

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631 SUPPORTING INFORMATION

- 632 Additional Supporting Information may be found online in the supporting information section
- 633 at the end of the article.
- 634 APPENDIX S1. Locations of the 26 populations studied. DF= Federal District; GO= Goiás;
- 635 MG= Minas Gerais.

636 TABLES

- 637 **TABLE 1.** Results of the analysis of variance for the comparisons of the fruit set to describe
- 638 the incompatibility system and natural fruit set of four *Erythroxylum* species.

| Source of variation | E. campestre | | | <i>E. d</i> | E. deciduum | | | E. suberosum | | | E. tortuosum | | |
|---|----------------------------------|------------------|----------------------------------|-----------------------------------|------------------|-----------------------------------|-----------------------------------|------------------|-----------------------------------|----------------------------------|------------------|----------------------------------|--|
| Source of variation | x ² | df | Р | x ² | df | Р | x ² | df | Р | x ² | df | Р | |
| Incompatibility system | | | | | | | | | | | | | |
| Morph | 0.424 | 1 | 0.514 | 0.091 | 1 | 0.763 | 1.165 | 1 | 0.559 | 3.627 | 1 | 0.057 | |
| Treatment | 46.440 | 5 | < 0.001 | 55.617 | 5 | < 0.001 | 45.984 | 5 | < 0.001 | 16.518 | 5 | < 0.01 | |
| Morph * Treatment | 6.053 | 5 | 0.301 | 8.493 | 5 | 0.131 | 0.354 | 5 | 0.997 | 0.000 | 5 | 1.000 | |
| Natural fruit set | | | | | | | | | | | | | |
| Morph | 0.837 | 1 | 0.360 | 0.274 | 1 | 0.601 | 0.000 | 1 | 0.996 | 0.481 | 1 | 0.488 | |
| Population | 4.053 | 2 | 0.132 | 30.852 | 2 | < 0.001 | 23.464 | 3 | < 0.001 | 0.267 | 1 | 0.605 | |
| Morph * Population | 0.177 | 2 | 0.915 | 7.901 | 2 | 0.019 | 2.869 | 3 | 0.412 | 0.000 | 1 | 0.978 | |
| Morph * Treatment Natural fruit set Morph Population Morph * Population | 6.053 0.837 4.053 0.177 | 5 1 2 2 | 0.301 0.360 0.132 0.915 | 8.493 0.274 30.852 7.901 | 5 1 2 2 | 0.131 0.601 <0.001 0.019 | 0.354 0.000 23.464 2.869 | 5 1 3 3 | 0.997 0.996 <0.001 0.412 | 0.000 0.481 0.267 0.000 | 5 1 1 1 | 1.000 0.488 0.605 0.978 | |

| (10) | 1 . • | 6.6 | T .1 | 1 | • |
|------|-------------|----------|----------|------|----------|
| 647 | nonulations | of four | Ervthrox | whim | snecies |
| 012 | populations | 01 10 01 | шуштол | yuum | species. |

| Species | Number of plants | df | x ² | Р | Characterization of |
|--------------|------------------|----|-----------------------|---------|----------------------------|
| Population | (S:L-morph) | ui | | Ŧ | populations |
| E. campestre | | | | | |
| 1 | 40 (21:19) | 1 | 0.03 | 0.874 | Isoplethy |
| 2 | 56 (17:39) | 1 | 7.88 | < 0.001 | Anisoplethy (†L) |
| 3 | 39 (21:18) | 1 | 0.10 | 0.749 | Isoplethy |
| 4 | 39 (32:7) | 1 | 14.77 | < 0.001 | Anisoplethy (†S) |
| 5 | 37 (1:36) | 1 | 31.24 | < 0.001 | Anisoplethy (<i>\L</i>) |
| 6 | 39 (19:20) | 1 | 0.00 | 1.000 | Isoplethy |
| 7 | 43 (43:0) | 1 | 41.02 | < 0.001 | Monomorphic |
| 8 | 41 (39:2) | 1 | 31.61 | < 0.001 | Anisoplethy $(\uparrow S)$ |
| E. deciduum | | | | | |
| 9 | 41 (18:23) | 1 | 0.39 | 0.532 | Isoplethy |
| 10 | 40 (18:22) | 1 | 0.23 | 0.635 | Isoplethy |
| 11 | 39 (15:24) | 1 | 1.64 | 0.200 | Isoplethy |
| 12 | 34 (13:21) | 1 | 1.44 | 0.230 | Isoplethy |
| E. suberosum | × , | | | | 1 2 |
| 13 | 32 (12:20) | 1 | 1.53 | 0.216 | Isoplethy |
| 14 | 34 (14:20) | 1 | 0.74 | 0.391 | Isoplethy |
| 15 | 48 (22:26) | 1 | 0.19 | 0.665 | Isoplethy |
| 16 | 40 (26:14) | 1 | 3.03 | 0.082 | Isoplethy |
| 17 | 42 (21:21) | 1 | 0.00 | 1.000 | Isoplethy |
| 18 | 35 (16:19) | 1 | 0.11 | 0.735 | Isoplethy |
| 19 | 30 (7:23) | 1 | 7.50 | 0.006 | Anisoplethy (1L) |
| 20 | 72 (40:32) | 1 | 0.68 | 0.409 | Isoplethy |
| E. tortuosum | | | | | 1 2 |
| 21 | 42 (27:15) | 1 | 2.88 | 0.090 | Isoplethy |
| 22 | 49 (23:26) | 1 | 0.08 | 0.775 | Isoplethy |
| 23 | 31 (9.22) | 1 | 4 65 | 0.031 | Anisoplethy $(\uparrow L)$ |
| 24 | 39(22.17) | 1 | 0.41 | 0.522 | Isoplethy |
| 25 | 37 (22:15) | 1 | 0.97 | 0.324 | Isoplethy |
| 26 | 42 (19:23) | 1 | 0.21 | 0.643 | Isoplethy |
| 20 | 12 (17.23) | 1 | 0.21 | 0.015 | isopicity |

- **TABLE 3.** Flower morphometrics (in mm) of four *Erythroxylum* species: sample size for the
- 652 two floral morphs, mean (\pm variance) of the height of each organ type, and mean (\pm variance)

| Species | n I. | n S- | I stiames | Santhers | S stiamos | I anthers - | Two anther leve | l of the L-morph |
|--------------|-------|-------|-------------|-------------|-------------|--------------|-----------------|------------------|
| Dopulation | n L- | n S- | L suginas | (Λ) | 5 stigillas | L allulers - | Upper anthers | Lower anthers |
| Fopulation | morph | morph | (3) | (A) | (8) | (a) | (aA) | (aa) |
| E. campestre | | | | | | | | |
| 1 | 18 | 17 | 3.11 (0.52) | 3.25 (0.54) | 1.47 (0.20) | 0.88 (0.11) | 1.11 (0.10) | 0.64 (0.02) |
| 2 | 58 | 25 | 2.49 (0.22) | 2.13 (0.10) | 1.40 (0.03) | 0.84 (0.08) | 1.06 (0.04) | 0.62 (0.02) |
| 3 | 36 | 36 | 3.50 (0.34) | 2.99 (0.09) | 1.44 (0.02) | 1.06 (0.12) | 1.31 (0.08) | 0.81 (0.03) |
| 4 | 12 | 62 | 3.10 (0.14) | 3.51 (0.13) | 1.65 (0.05) | 1.01 (0.12) | 1.25 (0.07) | 0.76 (0.04) |
| 5 | 42 | 2 | 2.40 (0.57) | 1.95 (0.01) | 1.02 (0.04) | 0.84 (0.15) | 1.13 (0.09) | 0.55 (0.05) |
| 6 | 28 | 30 | 3.73 (0.51) | 3.90 (0.16) | 1.79 (0.05) | 1.12 (0.25) | 1.47 (0.18) | 0.76 (0.07) |
| 7 | 0 | 19 | - | 3.47 (0.45) | 1.77 (0.10) | - | - | - |
| 8 | 1 | 33 | 2.61 | 2.23 (0.12) | 1.17 (0.02) | 1.02 (0.11) | 1.26 | 0.78 |
| Average | | | 2.99 (0.38) | 2.93 (0.20) | 1.46 (0.06) | 0.97 (0.13) | 1.23 (0.09) | 0.70 (0.04) |
| E. deciduum | | | . , | | . , | . , | | |
| 9 | 40 | 30 | 3.58 (0.18) | 4.21 (0.61) | 1.51 (0.08) | 2.61 (1.27) | 3.64 (0.27) | 1.58 (0.13) |
| 10 | 44 | 39 | 3.77 (0.30) | 3.85 (0.17) | 1.49 (0.06) | 2.84 (1.47) | 3.87 (0.62) | 1.80 (0.16) |
| 11 | 39 | 30 | 3.73 (0.19) | 4.22 (0.53) | 1.43 (0.09) | 2.57 (1.21) | 3.61 (0.17) | 1.52 (0.04) |
| 12 | 33 | 30 | 3.54 (0.21) | 3.80 (0.51) | 1.49 (0.05) | 2.62 (0.99) | 3.52 (0.17) | 1.72 (0.16) |
| Average | | | 3.66 (0.22) | 4.02 (0.46) | 1.48 (0.07) | 2.66 (1.24) | 3.66 (0.31) | 1.66 (0.12) |
| E. suberosum | | | | | | | . , | |
| 13 | 25 | 6 | 4.29 (0.31) | 4.35 (0.11) | 2.12 (0.03) | 1.51 (0.35) | 2.03 (0.12) | 1.00 (0.06) |
| 14 | 30 | 20 | 4.53 (0.29) | 4.47 (0.37) | 2.03 (0.04) | 1.59 (0.66) | 2.18 (0.46) | 1.00 (0.16) |
| 15 | 45 | 38 | 4.57 (0.58) | 4.48 (0.26) | 2.16 (0.07) | 1.73 (0.54) | 2.31 (0.32) | 1.16 (0.09) |
| 16 | 18 | 38 | 4.88 (0.56) | 4.81 (0.51) | 2.26 (0.10) | 2.06 (0.54) | 2.65 (0.21) | 1.46 (0.15) |
| 17 | 29 | 31 | 5.83 (0.53) | 4.95 (0.52) | 2.28 (0.06) | 2.30 (0.70) | 2.99 (0.32) | 1.61 (0.12) |
| 18 | 49 | 39 | 4.90 (0.19) | 4.54 (0.22) | 2.07 (0.05) | 1.82 (0.38) | 2.36 (0.11) | 1.27 (0.06) |
| 19 | 40 | 14 | 5.14 (0.27) | 4.30 (0.81) | 2.05 (0.05) | 2.04 (0.45) | 2.60 (0.18) | 1.47 (0.07) |
| 20 | 27 | 55 | 4.19 (0.41) | 3.71 (0.29) | 1.75 (0.06) | 1.38 (0.34) | 1.85 (0.15) | 0.91 (0.08) |
| Average | | | 4.79 (0.39) | 4.45 (0.39) | 2.09 (0.06) | 1.80 (0.50) | 2.37 (0.23) | 1.24 (0.10) |
| E. tortuosum | | | . , | | . , | . , | | |
| 21 | 15 | 47 | 3.91 (0.32) | 3.91 (0.27) | 1.61 (0.06) | 1.27 (0.44) | 1.85 (0.16) | 0.69 (0.03) |
| 22 | 42 | 37 | 3.87 (0.34) | 3.99 (0.36) | 1.61 (0.04) | 1.28 (0.40) | 1.80 (0.21) | 0.76 (0.03) |
| 23 | 36 | 19 | 3.88 (0.30) | 3.51 (0.10) | 1.49 (0.05) | 1.13 (0.31) | 1.60 (0.13) | 0.65 (0.03) |
| 24 | 42 | 50 | 4.48 (0.27) | 4.12 (0.33) | 1.64 (0.05) | 1.40 (0.34) | 1.92 (0.09) | 0.88 (0.04) |
| 25 | 31 | 47 | 4.64 (0.22) | 4.18 (0.44) | 1.64 (0.10) | 1.50 (0.60) | 2.21 (0.13) | 0.79 (0.05) |
| 26 | 45 | 15 | 3.70 (0.31) | 3.40 (0.42) | 1.49 (0.05) | 1.01 (0.15) | 1.34 (0.07) | 0.69 (0.02) |
| Average | | | 4.08 (0.29) | 3.85 (0.32) | 1.58 (0.06) | 1.27 (0.37) | 1.79 (0.13) | 0.74 (0.03) |

653 of upper and lower anthers in the L-morph.

661 TABLE 4. Results of the analysis of variance for the comparisons between floral morphs and 662 populations of stigma height and anther height of four *Erythroxylum* species. High anther 663 height corresponds to the comparison between anthers of the S-morph and upper anthers of 664 the L-morph. Low anther height corresponds to the comparison between anthers of the S-

665 morph and lower anthers of the L-morph.

| | Source of variation | E. campestre | | E. deciduum | | | E. suberosum | | | E. tortuosum | | | |
|------------------|---------------------|----------------|----|-------------|----------------|----|--------------|----------------|----|--------------|----------------|----|---------|
| | Source of variation | x ² | df | Р | x ² | df | Р | x ² | df | Р | x ² | df | Р |
| | Stigma height | | | | | | | | | | | | |
| | Morph | 622.7 | 1 | < 0.001 | 1083.9 | 1 | < 0.001 | 1955.4 | 1 | < 0.001 | 3146.1 | 1 | < 0.001 |
| | Population | 105.1 | 4 | < 0.001 | 1.3 | 3 | 0.717 | 118.7 | 6 | < 0.001 | 49.7 | 5 | < 0.001 |
| | Morph * Population | 35.4 | 4 | < 0.001 | 1.2 | 3 | 0.762 | 46.9 | 6 | < 0.001 | 42.5 | 5 | < 0.001 |
| | High anther height | | | | | | | | | | | | |
| | Morph | 1165.8 | 1 | < 0.001 | 14.7 | 1 | < 0.001 | 870.6 | 1 | < 0.001 | 1877.0 | 1 | < 0.001 |
| | Population | 189.0 | 4 | < 0.001 | 3.9 | 3 | 0.271 | 131.9 | 6 | < 0.001 | 76.0 | 5 | < 0.001 |
| | Morph * Population | 89.5 | 4 | < 0.001 | 7.8 | 3 | 0.051 | 6.0 | 6 | 0.424 | 7.0 | 5 | 0.221 |
| | Low anther height | | | | | | | | | | | | |
| | Morph | 2482.4 | 1 | < 0.001 | 907.7 | 1 | < 0.001 | 2780.1 | 1 | < 0.001 | 5606.1 | 1 | < 0.001 |
| | Population | 180.5 | 4 | < 0.001 | 0.9 | 3 | 0.835 | 136.5 | 6 | < 0.001 | 35.7 | 5 | < 0.001 |
| | Morph * Population | 154.0 | 4 | < 0.001 | 17.3 | 3 | < 0.001 | 14.7 | 6 | 0.022 | 27.2 | 5 | < 0.001 |
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| 677 | TABLE 5. Estimates of inaccuracy in reciprocity in species and populations of <i>Erythroxylum</i> |
|-----|--|
| 678 | (standardized inaccuracy in parentheses). The values of inaccuracy were standardized by the |
| 679 | squared mean of all anther and stigma height recorded for each population. Total inaccuracy |
| 680 | for each population (column 4) is the sum of the inaccuracy values of high and low organs |
| 681 | (columns 2 and 3). Columns 5 and 6 include inaccuracy (absolute value in units of mm^2 and |
| 682 | percentage standardized) of the low organs (S stigmas and L anthers) considering separately |
| 683 | the upper (aA) and lower (aa) anthers of the L-morph. |

| | | | | Inaccuracy of the low organs | | | | |
|--------------|---------------|---------------|----------------|----------------------------------|---------------|--|--|--|
| Species | Inaccuracy b | y organ type | Total incommon | considering the different height | | | | |
| Population | | | | of anthers in the L-morph | | | | |
| | High (S-A) | Low (s-a) | - | (s-aA) | (s-aa) | | | |
| E. campestre | | | | | | | | |
| 1 | 1.08 (22.81%) | 0.66 (14.04%) | 1.74 (36.85%) | 0.43 (9.08%) | 0.90 (19.01%) | | | |
| 2 | 0.44 (15.07%) | 0.42 (14.35%) | 0.87 (29.42%) | 0.18 (6.18%) | 0.66 (22.51%) | | | |
| 3 | 0.69 (13.64%) | 0.28 (5.49%) | 0.96 (19.13%) | 0.12 (2.29%) | 0.44 (8.68%) | | | |
| 4 | 0.44 (8.24%) | 0.58 (10.93%) | 1.03 (19.17%) | 0.28 (5.23%) | 0.89 (16.61%) | | | |
| 5 | 0.77 (32.00%) | 0.22 (9.16%) | 0.99 (41.16%) | 0.14 (5.67%) | 0.30 (12.62%) | | | |
| 6 | 0.70 (10.02%) | 0.75 (10.86%) | 1.45 (20.89%) | 0.33 (4.82%) | 1.17 (16.90%) | | | |
| 8 | 0.26 (8.55%) | 0.15 (4.93%) | 0.42 (13.48%) | 0.02 (0.70%) | 0.17 (5.52%) | | | |
| Average | 0.63 (15.76%) | 0.44 (9.96%) | 1.07 (25.73%) | 0.21 (4.85%) | 0.65 (14.55%) | | | |
| E. deciduum | | | | | | | | |
| 9 | 1.20 (13.49%) | 2.56 (28.88%) | 3.76 (42.37%) | 4.89 (55.12%) | 0.21 (2.40%) | | | |
| 10 | 0.47 (5.32%) | 3.35 (37.62%) | 3.82 (42.94%) | 6.37 (71.52%) | 0.32 (3.54%) | | | |
| 11 | 0.96 (10.74%) | 2.58 (28.94%) | 3.54 (39.67%) | 4.99 (56.00%) | 0.14 (1.59%) | | | |
| 12 | 0.79 (9.70%) | 2.32 (28.36%) | 3.12 (38.07%) | 4.36 (53.28%) | 0.26 (3.20%) | | | |
| Average | 0.86 (9.81%) | 2.70 (30.95%) | 3.56 (40.76%) | 5.15 (58.98%) | 0.23 (2.68%) | | | |
| E. suberosum | | | | | | | | |
| 13 | 0.42 (4.48%) | 0.76 (8.05%) | 1.18 (12.53%) | 0.16 (1.74%) | 1.34 (14.28%) | | | |
| 14 | 0.67 (6.71%) | 0.89 (8.93%) | 1.56 (15.64%) | 0.52 (5.23%) | 1.25 (12.60%) | | | |
| 15 | 0.85 (8.08%) | 0.80 (7.61%) | 1.64 (15.69%) | 0.42 (4.00%) | 1.17 (11.20%) | | | |
| 16 | 1.07 (8.70%) | 0.68 (5.55%) | 1.75 (14.25%) | 0.47 (3.80%) | 0.89 (7.22%) | | | |
| 17 | 1.81 (12.30%) | 0.76 (5.15%) | 2.57 (17.46%) | 0.88 (5.96%) | 0.63 (4.28%) | | | |
| 18 | 0.54 (4.89%) | 0.50 (4.50%) | 1.04 (9.39%) | 0.24 (2.20%) | 0.75 (6.75%) | | | |
| 19 | 1.78 (15.62%) | 0.50 (4.39%) | 2.29 (20.01%) | 0.54 (4.74%) | 0.46 (4.01%) | | | |
| 20 | 0.93 (12.23%) | 0.53 (7.02%) | 1.46 (19.25%) | 0.22 (2.84%) | 0.85 (11.15%) | | | |
| Average | 1.01 (9.13%) | 0.68 (6.40%) | 1.69 (15.53%) | 0.43 (3.81%) | 0.92 (8.94%) | | | |
| E. tortuosum | | | | | | | | |
| 21 | 0.59 (8.20%) | 0.61 (8.53%) | 1.20 (16.73%) | 0.28 (3.87%) | 0.93 (12.97%) | | | |
| 22 | 0.71 (9.80%) | 0.55 (7.65%) | 1.26 (17.45%) | 0.29 (4.04%) | 0.81 (11.21%) | | | |
| 23 | 0.54 (8.68%) | 0.50 (7.94%) | 1.04 (16.61%) | 0.20 (3.13%) | 0.79 (12.68%) | | | |
| 24 | 0.73 (8.61%) | 0.45 (5.32%) | 1.18 (13.93%) | 0.23 (2.67%) | 0.67 (7.91%) | | | |
| 25 | 0.86 (9.64%) | 0.72 (8.04%) | 1.58 (17.68%) | 0.55 (6.11%) | 0.88 (9.81%) | | | |
| 26 | 0.82 (14.24%) | 0.43 (7.40%) | 1.25 (21.64%) | 0.14 (2.50%) | 0.71 (12.28%) | | | |
| Average | 0.71 (9.86%) | 0.54 (7.48%) | 1.25 (17.34%) | 0.28 (3.72%) | 0.80 (11.14%) | | | |

686 FIGURE LEGENDS

687 **FIGURE 1.** Flowers of *Erythroxylum* species studied in the Cerrado of Central Brazil,

688 showing L- and S-morph of E. campestre (A, B), E. deciduum (C, D), E. suberosum (E, F)

689 and *E. tortuosum* (G, H).

690 FIGURE 2. Schematic representation of the relative position of stigmas and anthers in the S

and L-morph of *Erythroxylum* species studied and the flower traits measured. Letters in

692 parentheses correspond to the initials used for each type of organ in the analysis of

693 reciprocity. For clarity, only four of 10 stamens per flower are drawn (two per whorl).

694 **FIGURE 3.** Mean ± standard error of fruit set after hand-pollination treatments in individuals

695 of four *Erythroxylum* species. Pollination treatments were: agamospermy (Agam.),

696 autonomous self-pollination (Self-aut.), facilitated self-pollination (Self-fac.), intra-morph

697 cross-pollination (Intra) and inter-morph cross-pollination (Inter). Results from flowers

698 exposed to natural pollinations are also shown (control). As there were no statistical

699 differences between morphs within each species (results not shown), the results for both

700 morphs are plotted together. Treatments with different letters were significantly different from

701 each other at the 0.05 level in post-hoc tests.

702 **FIGURE 4.** Geographic distribution and variation in morph ratio in 26 populations of four

703 *Erythroxylum* species studied in Cerrado areas of Central Brazil. Black and grey segments

indicate the proportions of the L- and S-morph, respectively. Numbers correspond to the code

of each population. Details on sample size per population are provided in Table 2.

706 FIGURE 5. Population mean ± standard error of natural fruit set in S (black bars) and L

707 plants (grey bars) in four *Erythroxylum* species. Location of sampled populations are shown

in Figure 4. Populations with different letters were significantly different from each other at

the 0.05 level in post-hoc tests. Asterisks indicate statistically significant differences between

710 morph within populations or in overall, considering the 0.05 level in post-hoc tests. Sequential

711 Bonferroni correction was used to account for multiple tests. Except for *E. campestre*, natural

712 fruit set of these populations (mean and coefficient of variation) was used to investigate

associations with high and low inaccuracy (see Material and Methods for details).

714 FIGURE 6. Relationship between inaccuracy and the coefficient of variation in fruit set for S

715 (grey points) and L plants (black points) in nine populations of three different Erythroxylum

516 species. Dashed line shows predicted relationship and the shaded area indicates the 95%

717 confidence interval.







Treatment







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APPENDIX S1. Locations of the 26 populations studied. DF= Federal District; GO= Goiás;

MG= Minas Gerais.

| Region and name of areas | Population code | Coordinates |
|--|-----------------|------------------|
| Brasília (DF) - Parque Nacional de Brasília | 13, 21 | 15°43'S, 47°57'W |
| Brasília (DF) - APA do Paranoá | 1, 14 | 15°45'S, 47°51'W |
| Brasília (DF) - RECOR | 2, 15, 22 | 15°56'S, 47°55'W |
| Catalão (GO) - Pasto do Pedrinho | 9, 23 | 18°09'S, 47°56'W |
| Catalão (GO) - Fazenda Pé do Morro | 3, 10, 16, 24 | 18°06'S, 47°59'W |
| Catalão (GO) - Pires Belo | 4 | 17°52'S, 47°42'W |
| Caldas Novas (GO) - PESCAN | 5, 11, 17 | 17°46'S, 48°39'W |
| Uberlândia (MG) - Clube Caça & Pesca Itororó | 12, 18 | 19°00'S, 48°18'W |
| Uberlândia (MG) - Estação Ecológica do Panga | 6, 19, 25 | 19°10'S, 48°23'W |
| Diamantina (MG) - Ribeirão dos Infernos | 7 | 18°16'S, 43°32'W |
| Diamantina (MG) - Pedra da Baleia | 8, 20, 26 | 18°24'S, 43°60'W |