

1 **Sexual dimorphism in adult Little Stints (*Calidris minuta*)**  
2 **revealed by DNA sexing and discriminant analysis**

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## Abstract:

**Background.** Sexual selection plays such an important role in a bird's life cycle that researchers must know the sex of an individual to interpret key aspects of its biology. Sexes of dimorphic species can be easily distinguished, but sexing monomorphic bird species often requires expensive and time-consuming molecular methods. The Little Stint (*Calidris minuta*) is a numerous species, monomorphic in plumage but showing a small degree of reversed sexual size dimorphism (RSSD). Females are larger than males but ranges of their measurements overlap, making Little Stints difficult to sex in the field. Our aim was to develop reliable sexing criteria for Little Stints in different stages of primary moult during their stay on the non-breeding grounds in South Africa using DNA-sexed individuals and discriminant function analysis.

**Methods.** We caught 348 adult Little Stints in 2008–2016 at their non-breeding grounds at Barberspan Bird Sanctuary (26°33'S, 25°36'E, Northwest Province, South Africa) using mist nets and walk-in traps. To identify the birds' sex we isolated DNA from blood samples collected in the field using PCR with P2/P8 primers. We used Storer's Dimorphism Index to assess the degree of sexual size dimorphism. Then we divided our sample into two groups: before and after primary moult. For each group we developed two functions: one using wing length only and the other a combination of morphometrics. Then we used a stepwise procedure to check which combination of measurements best discriminated sexes. To validate our result we used a jack-knife cross-validation procedure and Cohen-kappa statistics.

**Results.** DNA-sexed females were larger than males in all the morphometric features we measured. Birds with fresh primaries had on average 2.3 mm longer wings than those in worn plumage. A discriminant function using wing length ( $D_1$ ) correctly sexed 78.8% of individuals before moult, and a stepwise analysis revealed that a combination of wing length and tarsus ( $D_2$ ) correctly identified the sex of 82.7% of the birds. For birds with freshly moulted primaries a function using wing length ( $D_3$ ) correctly classified 83.4% of the individuals, and a stepwise analysis revealed that wing and total head length ( $D_4$ ) classified 84.7%.

**Discussion.** Sexual size differences in Little Stints are connected with their breeding biology. Females are bigger, which increases their fecundity; males are smaller, which increases their manoeuvrability during display flights and hence their mating success. Because of their extreme lack of breeding site fidelity we do not expect a geographical cline in their

**Comentado [ALS1]:** The observed SSD in the species is not due to sexual selection. Please review.

46 biometrics despite their wide breeding range. Available sexing criteria for Little Stints in the  
47 literature were developed using museum specimens, which often shrink, leading to  
48 misclassification of live birds. The sexing criteria we developed can be used for studies on  
49 Little **Stints** in the future and can be used on past data.

## 50 **Introduction**

51 An individual's sex is one of the most important factors shaping its biology. Male and female  
52 birds are subject to sex-specific selection pressures that entail differences in their biology,  
53 including migration strategies (Remisiewicz & Wennerberg, 2006; Jakubas et al., 2014),  
54 population structure (Nebel, 2006), foraging behavior (Mathot & Elner, 2004; Nebel, 2005),  
55 moult (Barshep et al., 2013) and physiology (Kulaszewicz, Wojczulanis-Jakubas & Jakubas,  
56 2015). Differences in the biology of males and females leads to diverging body sizes  
57 (Fairbairn 2007), and are expected to emerge if selection for a character is stronger in one sex  
58 than the other (Székely, Lislevand & Figuerola, 2007). Sexing monomorphic birds is difficult  
59 in the field, but can be done with molecular methods (Dubiec & Zagalska-Neubauer, 2006)  
60 using DNA isolated from blood samples (Owen, 2011), feathers (Bello, Francino & Sánchez,  
61 2001) or buccal swabs (Handel et al., 2006); however those methods are often stressful for  
62 the birds and expensive. For species in which males and females are monomorphic in  
63 plumage but show sexual size dimorphism (SSD) the sex can often be identified using  
64 morphometrics (Dechaume-Moncharmont, Monceau & Cézilly, 2011). In such cases the  
65 degree of SSD is crucial, because in some species measurements overlap between sexes,  
66 which might lead to misidentification (Jiménez, García-Lau & Gonzalez, 2015). Discriminant  
67 function analysis (DFA) enables observers to use a combination of morphological  
68 measurements to predict the sex of a bird with a certain probability (Tabachnick & Fidell,  
69 2007). However, the efficiency of this method depends on the accuracy of the measurements,  
70 which suffers if researchers are not calibrated with each other or if wing length is measured  
71 on feathers with different wear (Dechaume-Moncharmont, Monceau & Cézilly, 2011). In our  
72 study we focus on Little Stints, which are monomorphic in plumage but exhibit a small  
73 degree of **SSD**, with the females slightly larger than the males (del Hoyo, Elliott & Sargatal,  
74 1996). But the ranges of their measurements overlap, which impedes sexing them in the field  
75 (Prater, Marchant & Vuorinen, 2007). Sexing criteria for Little Stints in the literature are  
76 usually based on small samples or on museum specimens, which are known to shrink (Prater,  
77 Marchant & Vuorinen, 2007).

**Comentado [ALS2]:** The authors should make it clear that the results should be used carefully because of the misclassification of sex (21.2 to 15.3%), according to the used FDA.

**Excluido:** sexual size dimorphism

We aim to develop reliable sexing criteria for Little Stints by discriminant functions that account for the stage of wear of their primaries using measurements of DNA-sexed individuals spending the non-breeding season in South Africa. We suggest that these discriminant functions serve as a useful tool, after adjustment to local measurement routines, for other researchers studying Little Stints at various stages of their life cycle.

## **Materials & methods**

### **Species and study area**

The Little Stint is a long-distance migrant wader, which breeds in the Eurasian tundra and spends the non-breeding season from Mediterranean coasts south throughout sub-Saharan Africa to South Africa in the south and southwestern Asia in the east (del Hoyo, Elliott & Sargatal, 1996; Tulp et al., 2002). Most Little Stints leave the non-breeding grounds and attempt breeding at the end of their first year; but a few stay in South Africa over the austral winter for a “gap year” (Underhill 1997; Underhill et al., 1999). Immature Little Stints can be distinguished from adults by the retained chestnut-fringed inner median coverts, which in adults are pale grey-brown tipped whitish (Prater, Marchant & Vuorinen, 2007). After arrival at the non-breeding grounds immature and adult birds undergo a complete moult, including the replacement of wing flight feathers. They complete this moult in January–March, after which the immatures become indistinguishable from the adults (Prater, Marchant & Vuorinen, 2007).

We caught Little Stints at Barberspan Bird Sanctuary (26°33'S, 25°36'E; North West Province, South Africa). This reserve is centered on a shallow alkaline lake whose area varies from 257 ha to 2000 ha, depending on rainfall (Milstein, 1975; Barnes, 1998). Lake Barberspan is fed by the Harts River. In the austral winter and during droughts it becomes the only permanent waterbody in the surrounding farmland, which makes it an important stopover, moulting and non-breeding destination for waterbirds, including Palearctic migrants (Barnes, 1998; Lipshutz et al., 2011; Remisiewicz & Avni, 2011). Barberspan Bird Sanctuary is a Wetland of International Importance in terms of the Ramsar Convention and an Important Bird Area according to BirdLife International (Cowan, 1995; Barnes, 1998; Remisiewicz & Avni 2011).

**Comentado [ALS3]:** It would be better represented on a map of the country with a study area

## 108 Data collection

109 During September–April in 2008–2016 we captured 348 adult Little Stints using walk-in  
110 traps (Busse & Meissner, 2015) and mist nets. We ringed and measured each bird.  
111 Measurements included: bill length (from the feather line to the bill tip), total head length  
112 (from the back of the skull to the bill tip) and tarsus length (from the tarsal joint to the distal  
113 end of the tarso-metatarsus), taken with callipers of 0.1 mm accuracy, and wing length  
114 (maximum flattened chord, as in Evans (1986)), tarsus-plus-toe (Piersma, 1984) taken to the  
115 nearest 1 mm with a ruler, and body mass to 1 g or 0.1 g in different years (weighed with an  
116 electronic scale). We used only morphological measurements taken by MR (SAFRING  
117 ringing permit 1454), as recommended in Henry et al. (2015) because measurements taken by  
118 different ringers can reduce the accuracy of sex discrimination. We also took blood samples  
119 from a brachial vein (Owen, 2011) from all ringed Little Stints and preserved the samples in  
120 96% ethanol for DNA sexing. Only trained, experienced team members handled the birds and  
121 took the blood samples to ensure safety standards. All the procedures were approved by the  
122 management of Barberspan Bird Sanctuary, under permits from SAFRING and the  
123 Department of Rural, Environmental and Agricultural Development, North West Provincial  
124 Government, South Africa.

## 125 Molecular sex identification

126 We isolated DNA from blood samples using the Blood Mini kit (A&A Biotechnology,  
127 Poland). The next step was PCR with the pair of P2 and P8 primers (Griffiths et al., 1998)  
128 using a modified protocol. Total volume of PCR was 20 µl, the reaction mix included: 7.5 µl  
129 REDTaq® ReadyMix™ (Sigma Aldrich), 3.5 µl of water, 1 µl MgCl<sub>2</sub>, 1 µl P2 primer (10  
130 µM), 1 µl P8 primer (10 µM) and 2 µl of the DNA sample. For PCR amplifications we used  
131 an Eppendorf Mastercycler with this thermic profile: an initialisation at 94°C for 2 minutes,  
132 40 cycles of denaturation at 94°C for 30 seconds, 40 cycles of annealing at 50°C for 30  
133 seconds and 40 cycles of elongation at 72°C for 2 minutes, ending with a final elongation at  
134 72°C for 5 minutes. This method is based on the amplification of chromo-helicase-DNA-  
135 binding (CHD) genes found on avian sex chromosomes. The CHD-Z gene (ca 370 bp) is  
136 located on the Z chromosome, therefore it is present in both sexes. The CHD-W gene (ca 390  
137 bp) occurs only on chromosome W, therefore it is unique to females (Griffiths et al., 1998).  
138 We separated the products of PCR with electrophoresis on 3.5% agarose gel (75V, 120 min)  
139 stained with ethidium bromide (samples from 2008–2013) and Midori Green Advanced DNA

**Comentado [ALS4]:** low accuracy in body mass measurement

140 Stain (NIPPON Genetics, samples from 2013–2016). Products were visualised in UV light,  
141 one band was visible for males, which indicated ZZ chromosomes, and two bands for females  
142 (ZW chromosomes). The method enabled us to sex 100% of the birds we caught.

### 143 Statistical analysis

144 For each morphometric measurement we used Storer's dimorphism index ( $DI_{\text{Storer's}}$ ) to assess  
145 the degree of SSD in Little Stints (Storer, 1966), using the formula:

146 
$$DI_{\text{Storer's}} = \frac{\text{mean } \varphi - \text{mean } \sigma}{\text{mean } \sigma + \text{mean } \varphi} * 100$$

147 We compared all the morphometrics we had measured of males and females using the two-  
148 sample *t*-test. Then we divided the birds into two groups: those before and those after primary  
149 moult, and compared the measurements of birds from these two groups. Birds caught in  
150 active moult were classified as “before moult” because they moult their outermost primary  
151 last. However, we did not measure wing length of any bird whose outermost primary was  
152 heavily damaged or if it was growing. For each group we used Discriminant Function  
153 Analysis to determine the best set of measurements for sexing Little Stints with a two-fold  
154 approach. First, we used wing length alone as a discriminant factor because this measurement  
155 differed the most between the sexes. The second approach used a stepwise method including  
156 other measurements after conducting pairwise correlation of all the measurements. We used  
157 only one of a pair of correlated measurements at a time in the stepwise procedure to avoid  
158 multicollinearity. We did not include the body mass of Little Stints, because it changes during  
159 the non-breeding season during pre-migratory fuelling. The aim of producing two different  
160 discriminant functions for each group was to make these functions applicable for different  
161 data sets, because wing length is the most commonly taken measurement, in contrast to tarsus  
162 and total head lengths. All the assumptions of discriminant function analysis were met  
163 (Tabachnick & Fidell, 2007), including the homogeneity of covariances (Box's M test), the  
164 homogeneity of variance (Levene's test) and the normal distributions of the measurements for  
165 males and females separately in each of the two groups, and we confirmed no  
166 multicollinearity of the selected measurements ( $r < 0.50$  for all pairwise correlations). We  
167 computed prior classification probabilities from the group sizes because of the unequal  
168 number of males and females in our sample (Tabachnick & Fidell, 2007). To validate our  
169 models we used a jack-knife procedure to assess the percentage of correctly sexed individuals  
170 by discriminant function analysis (Dechaume-Moncharmont et al. 2011). This cross-

**Excluído:** sexual size dimorphism

**Comentado [ALS5]:** ( \* 0,5)

**Comentado [ALS6]:** Results in table 1 are not accordig this formula.

**Comentado [ALS7]:** But how, if bill lenght and Weight have a larger Storer index than wing lenght. So, the largest SSD was in bill and weight (I think body mass is not a good choice)

validation technique predicts the sex of each individual using a discriminant function calculated for all the birds except the individual being classified (Hair et al., 1995). We had unequal samples of males and females, so we assessed the effectiveness of our proposed functions by calculating Cohen's kappa statistic (Titus & Mosher, 1984). It estimates the improvement made by the results of discriminant analysis over random chance: 0 = no improvement over chance, 1 = full compliance (Titus & Mosher, 1984). The optimal cutting score was calculated as a weighted average of the group centroids (Hair et al., 1995). Statistical analyses were performed in IBM SPSS Statistics for Windows, version 22.0 (IBM Corp., Armonk, N.Y., USA). All tests were two-tailed and the accepted level of significance was  $P < 0.05$ .

## Results

### Morphological differences between the sexes

We identified 187 males and 165 females using DNA sexing. Analysis of morphometrics and Storer's dimorphism index ( $DI_{\text{Storer's}}$ ) also revealed sexual differences. On average females were bigger than males in all morphological measurements (Table 1). In all our measurements, birds before and after wing moult, differed only in wing length ( $t_{311} = 7.69$ ,  $p < 0.001$ ), which was on average 2.3 mm longer in those with fresh primaries after moult than in those before moult with worn primaries. We therefore conducted the discriminant analyses separately for these two groups.

Some morphological measurements were correlated (correlation coefficients for males and females between tarsus and tarsus-plus-toe were  $r=0.73$  and  $r=0.79$  and for culmen and total head length  $r=0.81$  and  $r=0.83$ ). We chose wing length, total head length (because it is less prone to errors than bill length measured to the feather line), and tarsus length as the best factors for discriminant analysis.

### Discriminant functions for adult Little Stints before primary moult

Using measurements of 156 adult Little Stints (70 females and 86 males) taken before they had moulted their primaries, and using only wing length as a discriminant factor, we obtained the equation:

$$D_1 = -47.496 + 0.484 (\text{wing}),$$

**Comentado [ALS8]:** 187 + 165 = 352  
Methods and Data collection = 348 birds. Where 4 birds not evaluated the morphometrics?

201 which allowed us to correctly classify 78.8% of the birds. A jack-knife cross-validation  
202 procedure yielded the same success rate, and our random chance-corrected procedure showed  
203 that our proposed classification was 56.8% better than chance ( $\kappa = 0.568 \pm 0.082$  SE,  $P <$   
204  $0.001$ ). If  $D_1 > 0.17$ , the bird was classified as a female, and if  $D_1 < 0.17$  as a male (Fig. 1)

205 In the stepwise procedure including three selected measurements (wing, tarsus and  
206 total head length), only the combination of wing and tarsus length was a significant  
207 discriminant factor. The best discriminant function we obtained was

208 
$$D_2 = -50.428 + 0.421 (\text{wing}) + 0.420 (\text{tarsus}).$$

209 The percentage of correctly classified birds was 82.7%, more accurate than the previous  
210 equation. The cross-validation procedure correctly classified 80.8% of individuals and the  
211 classification was 65.1% better than chance ( $\kappa = 0.651 \pm 0.081$  SE,  $P < 0.001$ ). If  $D_2 >$   
212  $0.18$  then the individual was a female and if  $D_2 < 0.18$  it was a male (Fig. 2).

#### 213 **Discriminant functions for adult Little Stints after primary moult**

214 The second group we analysed was 159 adult Little Stints (82 females and 77 males) after  
215 their complete post-breeding moult, with all new primaries. The function obtained using only  
216 wing length was

217 
$$D_3 = -52.184 + 0.520 (\text{wing}).$$

218 This function correctly classified 83.4% of the individual birds, which was confirmed by the  
219 cross-validation procedure. The proposed classification was 66.9% better than chance ( $\kappa =$   
220  $0.669 \pm 0.080$  SE,  $P < 0.001$ ). When  $D_3 > -0.038$  the individual was classified as a female,  
221 when  $D_3 < -0.038$  as male (Fig. 1).

222 The stepwise analysis revealed that the best discriminating combination of  
223 measurements was wing and total head length (THL), according to the equation

224 
$$D_4 = -59.310 + 0.445 (\text{wing}) + 0.377 (\text{THL}).$$

225 This function correctly classified 84.7% of the individuals in the sample, which was the  
226 highest proportion of all the equations we present. The cross-validation procedure showed  
227 that the proposed equation was 84.1% correct and 69.4% better than chance ( $\kappa = 0.694 \pm$   
228  $0.080$  SE,  $P < 0.001$ ). If  $D_4 > -0.041$  a Little Stint was classified as a female and if  $D_4 < -$   
229  $0.041$  as a male (Fig. 2).



## 230 Discussion

231 We showed that the female Little Stints were larger than the males, not only in wing length,  
232 as described earlier (Cramp & Simmons, 1983; Prater, Marchant & Vuorinen, 2007), but also  
233 in other body features. Thus we recommend a combination of several measurements in  
234 discriminant functions as a useful tool in studies of sexual differences of this species. Such  
235 size difference might benefit each sex in different ways, the females through increased  
236 fecundity and the males through higher mating success with smaller size, which we discuss.

237 **Wing length was the most dimorphic morphometric**, with females having longer  
238 wings than males. Mean wing lengths based on the sexing criteria in Prater et al. (2007) and  
239 established by Tree (1974) is 95.9 mm for adult males and 99.5 mm for adult females, 1.5  
240 mm and 1.7 mm shorter than our results (Table 1). The sexing criteria presented in Prater et  
241 al. (2007) **was** developed using museum specimens, thus subjected to shrinkage (Jenni &  
242 Winkler, 1994). We compared our differences with the one obtained using a regression  
243 equation in Engelmoer et al. (1983), where: shrinkage =  $0.006 \times \text{fresh wing-length [mm]} +$   
244  $0.976$ , and predicted shrinkage for both sexes was 1.6 mm, similar to what we observe in our  
245 sample. Moulting and feather wear are important considerations when taking feathered  
246 measurements like wing length (Meissner, 2005; Jiménez, García-Lau & Gonzalez, 2015),  
247 thus we propose different discriminant functions for birds before and after moulting. During the  
248 non-breeding season all Little Stints should be carefully examined for moulting because their  
249 outermost primary might still be growing. Measuring wing lengths of birds moulting P10  
250 would underestimate wing length and could even cause **misclassifications**.

251 Sexual differences in the size of Little Stints might be closely related to their breeding  
252 biology. Male mating success is a strong form of **sexual selection** that affects their  
253 morphometrics. During the breeding season male Little Stints perform display flights,  
254 favouring smaller size to increase manoeuvrability (Figuerola, 1999; Székely et al., 2000).  
255 The females' bigger size might be connected with increasing fecundity, because Little Stints  
256 exhibit successive bigamy of both sexes where females lay two clutches in a short period  
257 (Cramp & Simmons, 1983; Hildén, 1983) and a larger body size allows her to compensate  
258 better for the increased energy expenditure of egg production (Jönsson & Alerstam, 1990).  
259 Sexual differences in morphometrics cannot be explained by a division in parental care,  
260 because each clutch and brood is cared for by a single parent of either sex (Tulp et al., 2002).  
261 Small calidridine sandpipers, such as Western Sandpiper (Nebel, 2005) and Least Sandpiper

**Comentado [ALS9]:** That is not true. If bill length and Weight have a larger Storer index than wing length. So, the largest SSD was in bill length and body mass. The Storer's index indicates how much SSD you have in each measure.

**Excluido:** ere

**Comentado [ALS10]:** Bill length shows the largest Storer's index of all measurements. Please discuss why it did not prove to be a good measure for building a DFA.

Supporting literature:

- Nebel et al. 2011. The evolution of sexual bill-size dimorphism in shorebirds: a morphometric test of the resource partitioning hypothesis. *Evolutionary Ecology Research* 13:35–44.

- Scherer et al. 2014. Sexual dimorphism and body condition of wintering Whiterumped Sandpipers in southern Brazil.

**Comentado [ALS11]:** Nebel and Thompson (2011) evaluated the evolution of sexual bill-size dimorphism in 151 shorebird species and found that SSD is more pronounced in bill length as a resource-exploiting trophic character than nontrophic traits such as tarsus and wing length and body mass. The authors suggest that the comparative approach reveals that these patterns are not strictly the result of contemporary natural or sexual selection but rather reflect the shared ancestry of extant species.

(Nebel, 2006), exhibit substantial sexual differences in bill length, which is not the most dimorphic morphometric in Little Stints (Table 1) and suggests that they do not exhibit sexual differences in their foraging niche.

**Comentado [ALS12]:** Excluding body mass, the largest SSD is in Bill length (4,48)! Please, revise our conclusions.

| Measurement                 | Females |                 |             | Males |                |             | t     | p       | D <sub>storer's</sub> |
|-----------------------------|---------|-----------------|-------------|-------|----------------|-------------|-------|---------|-----------------------|
|                             | N       | Mean (SD)       | Range       | N     | Mean (SD)      | Range       |       |         |                       |
| Wing length (mm)            | 152     | 101.19 (± 2.34) | 95 – 106    | 164   | 97.42 (± 2.19) | 92 – 102    | 14.81 | < 0.001 | 3.80                  |
| Total head length (mm)      | 163     | 39.30 (± 1.06)  | 36.0 – 42.3 | 185   | 38.26 (± 0.98) | 35.8 – 41.1 | 9.62  | < 0.001 | 2.69                  |
| Bill length (mm)            | 163     | 18.11 (± 0.92)  | 16.3 – 20.5 | 185   | 17.32 (± 0.81) | 15.1 – 20.1 | 8.55  | < 0.001 | 4.48                  |
| Tarsus length (mm)          | 162     | 22.09 (± 0.86)  | 19.7 – 24.2 | 185   | 21.55 (± 0.78) | 19.4 – 24.4 | 6.16  | < 0.001 | 2.47                  |
| Tarsus-plus-toe length (mm) | 163     | 40.94 (± 1.27)  | 37 – 44     | 185   | 40.15 (± 1.34) | 37 – 46     | 5.69  | < 0.001 | 1.96                  |
| Weight (g)                  | 162     | 24.48 (± 3.66)  | 19 – 42     | 185   | 21.98 (± 2.43) | 17 – 33     | 7.42  | < 0.001 | 10.78                 |

A wide breeding range, such as that of the Little Stint, often results in a geographical variation in biometrics (Zwarts et al., 1996; Dmitrenok et al., 2007), which might distort a DFA. We do not expect such differences in our study because Little Stints are an opportunistic species that show no natal philopatry or breeding-site fidelity and breed wherever they find favourable environmental conditions (Hildén 1983; Underhill et al. 1993; Tomkovich & Soloviev 1994). Their polygamous breeding system where males and females both frequently have two partners enhances gene-flow in the population, limiting geographical clines in the morphometrics of Little Stints.

## Conclusions

DNA sexing remains the most reliable method for monomorphic species, but discriminant functions are useful when researchers cannot collect or process DNA samples, get permits or when sex identification is needed in field studies. The functions we developed can be applied on measurements collected in the past and can strengthen the analysis when individual birds' sex must be known. To make our functions more applicable for researchers we present equations that can be used with different types of morphological measurements that might be available and equations that consider the wear of primary feathers. Our results are based on relatively big samples and all of the measurement were taken by one ringer, but ringing teams

284 should regularly calibrate all people taking measurements (Dechaume-Moncharmont,  
285 Monceau & Cézilly, 2011), which improves the accuracy of sexing by discriminant functions.  
286 The functions we suggest should be applied cautiously to data from other researchers, and  
287 should probably first be adjusted to each study, considering possible differences in measuring  
288 routines with those of our team. We suggest application of these or similar discriminant  
289 functions as a useful tool facilitating studies of differences in the biology of the sexes at  
290 different stages of the life cycle, in Little Stints and in other species of small sexual size  
291 dimorphism.

## 292 **Acknowledgements**

293 We thank the staff at Barberspan Bird Sanctuary for their help and for making us feel  
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**Comentado [ALS13]:** The SSD and the Storer's index of the species was not discussed. Studies have reported that sexual size dimorphism decreases with size when females are the larger sex (Rensch's rule), and is determined by a combination of sexual competition and male agility.

313 Dmitrenok M, Puglisi L, Demongin L, Gilbert G, Polak M, Bretagnolle V. 2007.  
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425 Morphological features of adult male and female Little Stints.

426 t, p – results of t-test comparing the sexes. DI<sub>storer's</sub> – Storer's dimorphism index.

| <u>Measurement</u>                 | <u>Females</u> |                        |                    | <u>Males</u> |                       |                    |             |                   |              |
|------------------------------------|----------------|------------------------|--------------------|--------------|-----------------------|--------------------|-------------|-------------------|--------------|
|                                    | <u>N</u>       | <u>Mean (SD)</u>       | <u>Range</u>       | <u>N</u>     | <u>Mean (SD)</u>      | <u>Range</u>       |             |                   |              |
| <u>Wing length (mm)</u>            | <u>152</u>     | <u>101.19 (± 2.34)</u> | <u>95 – 106</u>    | <u>164</u>   | <u>97.42 (± 2.19)</u> | <u>92 – 106</u>    |             |                   |              |
| <u>Total head length (mm)</u>      | <u>163</u>     | <u>39.30 (± 1.06)</u>  | <u>36.0 – 42.3</u> | <u>185</u>   | <u>38.26 (± 0.98)</u> | <u>35.0 – 41.0</u> |             |                   |              |
| <u>Bill length (mm)</u>            | <u>163</u>     | <u>18.11 (± 0.92)</u>  | <u>16.3 – 20.5</u> | <u>185</u>   | <u>17.32 (± 0.81)</u> | <u>15.1 – 20.1</u> | <u>8.55</u> | <u>&lt; 0.001</u> | <u>4.48</u>  |
| <u>Tarsus length (mm)</u>          | <u>162</u>     | <u>22.09 (± 0.86)</u>  | <u>19.7 – 24.2</u> | <u>185</u>   | <u>21.55 (± 0.78)</u> | <u>19.4 – 24.4</u> | <u>6.16</u> | <u>&lt; 0.001</u> | <u>2.47</u>  |
| <u>Tarsus-plus-toe length (mm)</u> | <u>163</u>     | <u>40.94 (± 1.27)</u>  | <u>37 – 44</u>     | <u>185</u>   | <u>40.15 (± 1.34)</u> | <u>37 – 46</u>     | <u>5.69</u> | <u>&lt; 0.001</u> | <u>1.96</u>  |
| <u>Weight (g)</u>                  | <u>162</u>     | <u>24.48 (± 3.66)</u>  | <u>19 – 42</u>     | <u>185</u>   | <u>21.98 (± 2.43)</u> | <u>17 – 33</u>     | <u>7.42</u> | <u>&lt; 0.001</u> | <u>10.78</u> |

**Comentado [ALS14]:** The results are not according the Storer's index informed in the methods  
Wing = (101,19 – 97,42)/ (101,19 + 97,42) \*100  
Wing = 1,89  
DI<sub>storer's</sub> =  $\frac{\text{mean } \varphi - \text{mean } \sigma}{\text{mean } \sigma + \text{mean } \varphi} * 100$