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Biometric data for ten species from the families Malaconotidae, Platysteiridae and Laniidae based on museum specimens and ringing data

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Abstract

Biometric data is vital for species identification, taxonomic studies, ageing, and sexing. Museum collections and ringing data sets are important sources of biometric data for birds. We present biometric data for ten species from the families Malaconotidae, Platysteiridae and Laniidae based on 271 museum specimens and 184 ringing records. The species reported on are Bokmakierie, Brown-crowned Tchagra, Crimson-breasted Shrike, Brubru, Cape Batis, Chin-spot Batis, Pririt Batis, Magpie Shrike, Southern Fiscal, and Red-backed Shrike. The biometric data were compared between the sexes and to published data.

Keywords: sexual dimorphism, subspecies, citizen science, museum study skins, SAFRING, bird ringing.

Introduction

Biometric data are used for species identification, in taxonomic studies, investigating geographic size variation, ageing and sexing of individuals (Kühl and Burghardt 2013, Rowley et al. 2020). For southern African birds, Hockey et al. (2005) provide the most comprehensive summary of biometric data, but, in many instances, the data are incomplete and don't distinguish between subspecies or the sexes. For some species, no biometric data are reported. Many museums and research institutes hold collections of biological material that can provide valuable biometric data sets for taxonomic and systematic studies (Suarez and Tsutsui 2004). For example, the National Museum, Bloemfontein, has an extensive bird collection (>5,000 skins and >2,000 skeletons; https://nasmus. co.za/ornithology), each with corresponding collection data, e.g., location, date, and biometric data recorded upon collection. Additional biometric data are available from regional ringing initiatives using trained citizen scientists, such as southern Africa's SAFRING (de Beer et al. 2000). Rose et al. (2020) summarised biometric data from the SAFRING database, highlighting the importance of recording complete and accurate biometric information. Although Rose et al. (2020) reported sex-specific differences in species for which sufficient data were available, the authors did not distinguish between subspecies in their analyses. Comparisons of biometric data for subspecies and the sexes have been completed using regional ringing data, e.g., Karoo Prinia Prinia maculosa (de Swardt et al. 2018) and Namibian passerines (Paijmans and Bryson 2023), museum skins, e.g., Sabota Lark Calendulauda sabota (Marr et al. 2017), and combining museum and ringing data, e.g., Malachite Sunbirds Nectarinia famosa, Karoo Prinia, Drakensberg Prinia Prinia hypoxantha, and African Rock Pipit Anthus crenatus (de Swardt et al. 2003, de Swardt et al. 2018, de Swardt 2022). In this paper, we summarise selected biometric data for four species in the family Malaconotidae, three in the Platysteiridae, and three in the Laniidae based on specimens from the ornithology collection of the National Museum, Bloemfontein, and ringing data.

Methods

Representatives of the family Laniidae (Southern Fiscal *Lanius collaris*, Red-backed Shrike *Lanius collurio*, and Magpie Shrike *Lanius melanoleucus*), the Malaconoti-



dae (Crimson-breasted Shrike Laniarius atrococcineus, Brubru Nilaus afer, Brown-crowned Tchagra Tchagra australis, and Bokmakierie Telophorus zeylonus, and the Platysteiridae (Cape Batis Batis capensis, Chinspot Batis Batis molitor, and Pririt Batis Batis pririt) were selected for this study. Ringing surveys by DHDS from 1987 to 2023 were conducted at sites in the Free State. Northern Cape. Mpumalanga, and KwaZulu-Natal provinces as part of National Museum, Bloemfontein, research projects. Data were recorded as described by de Beer et al. (2000) and included culmen and tarsus lengths (± 0.1 mm) using Vernier callipers, wing and tail lengths (± 1 mm) using graduated rulers, and mass (± 0.1 g) using spring and electronic balances. Individuals were sexed in hand, where possible, based on size and plumage differences.

Two data sets were available for the museum specimens: dry skins collected pre-1995 and dry skins and skeletons prepared from fresh material post-1995 by DHDS. For the pre-1995 dry skins, hereafter Dry specimens, the mass as given on the collection card was used and the culmen, wing and tail lengths were taken following Eck et al. (2011). Tarsus measurements were excluded due to difficulty measuring the tarsus in study skins and shrinkage as the skins dry out (Kuczyński et al. 2002). For the specimens prepared from fresh material post-1995 (hereafter Fresh specimens), the sex, culmen, tarsus, wing, and tail lengths and mass were recorded for each specimen upon collection. For monomorphic species, e.g., tchagras, the bird was sexed during the preparation of the skin or skeletal specimens. Following the recommendation of Kuczyński et al. (2002), all measurements were recorded by one researcher (DHDS). For the statistical analyses, the post-1995 museum data were combined with the ringing data and analysed as Fresh specimens.

Biometrics of ten passerine species

All statistical analyses were done using the R statistical software (R Development Core Team 2023).

Table 1. Sample sizes for biometric data obtained from selected species in the families Malaconotidae, Platvsteiridae, and Laniidae, obtained from skin and skeleton specimens from the collection of the Ornithology Department, National Museum, Bloemfontein, and SAFRING data collected by DHDS from 1987 to 2023.

Species			Muse	eum			Ring	ging	(Combin	ed
	Dı	y	Fre	sh	Tot	al	Fre	sh	F	М	Total
	F	М	F	М	F	М	F	М			
Malaconotidae											
Bokmakierie	4	10	7	10	11	20			11	20	31
Brown-crowned Tchagra	6	6	7	21	13	27			13	27	40
Brubru	1	4	3	18	4	22			4	22	26
Crimson-breasted Shrike		6	1	10	1	16		2	1	18	19
Platysteiridae											
Cape Batis	5	2	1	4	6	6	12	7	18	13	31
Chinspot Batis	1	2	2	5	3	7		1	3	8	11
Pririt Batis	4	8	31	30	35	38	20	20	55	58	113
Laniidae											
Magpie Shrike	6	7	4	6	10	13	1		11	13	24
Red-backed Shrike	1	2	2	3	3	5	8	5	11	10	21
Southern Fiscal	12	25	4	6	16	31	55	53	71	84	155
	40	72	62	113	102	185	96	88	198	273	471

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Statistical significance was determined at p < 0.05. The than reported here (Paijmans and Bryson 2023), but Shapiro-Wilk test was used to evaluate the normality differences may be due to the skull line measurement of the data (shapiro.test). Differences in biometric being used for the Namibian study. parameters between the sexes were analysed using Analysis of Variance for normal data (aov) and the Brown-crowned Tchagra Tchagra australis damarensis Kruskal-Wallis test for non-parametric data (kruskal. Biometric data for 28 Fresh and 12 Drv specimens test). The data for Fresh and Dry specimens were of Tchagra australis damarensis are summarised in analysed separately, but the respective biometric Tables 2a.2 and 2b.2, respectively. We followed the parameters' ranges were compared to those for Fresh classification of Bowie (2005a) and Fry (2009). Dry male specimens. specimen males have significantly longer culmen

Results and Discussion

Biometric data for the four species in the Malaconotidae, three in the Platysteiridae, and three in the Laniidae included in this study were obtained from 287 museum specimens, 112 Dry and 175 Fresh specimens, and 184 ringed birds (Table 1). The sex of the specimens was strongly biased towards males for the museum specimens, while the ringing data was marginally biased towards females (Table 1). Recording of the five biometric parameters varied: only 44% of the records contained all five parameters, while only one parameter was missing for a further 21%. Mass was recorded for 95% of the records, wing length for 86%, tail length for 76%, culmen length for 67%, and tarsus length for 47%.

Malaconotidae

Bokmakierie Telophorus zeylonus zeylonus

Biometric data for 17 Fresh and 14 Dry specimens Biometric data for 21 Fresh and 5 Drv Nilaus afer of T. z. zeylonus are summarised in Tables 2a.1 and brubru specimens are summarised in Tables 2a.3 2b.1, respectively. The wing lengths of Fresh male and 2b.3, respectively. Only the mass for the Fresh specimens were significantly longer than the Fresh male specimens was significantly larger than that of females (Table 2a.1). No significant differences were the females (Table 2a.3); however, there were only found between the sexes for the Dry samples (Table three Fresh female specimens (for one of which 2b.1). The minimum wing length for the Dry males and only mass was recorded) and 18 Fresh male specifemale and Fresh females were outside the range of mens (Table 2a.3). There were insufficient specimens the Fresh males. Similarly, the minimum tarsus length for statistical analyses to be conducted for the Dry of the Dry females, maximum tarsus length of the specimens (Table 2b.3). The minimum mass and Fresh females, minimum tail length of the Dry males, minimum tarsus length for the Fresh females and maximum length of the Fresh females, and maximum the maximum tail length and maximum mass for the mass of the Dry males all exceeded the respective Dry males exceeded the range established by the ranges of the Fresh males. Fresh males.

The biometrics reported here are within the ranges The wing lengths and masses were similar to those and mean values of Oatley (2005) and Rose et al. given by Bowie (2005b), Rose et al. (2020), and Paij-(2020), although our maximum tail length is longer. mans and Bryson (2023). The remaining parameters The mean body mass of the Namibian subspecies *T*. were within the ranges presented by Bowie (2005b). z. thermophiles (Paijmans and Bryson 2023) is lower The mean tail lengths of males and females were than reported here for both sexes. Similarly, the culmen longer than those of Rose et al. (2020), although our sample size for females was low. The culmen lengths for the Namibian subspecies are also longer

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lengths than Dry females (Table 2b.2). The sample size for the Fresh males was three times greater than those for the Dry specimen males and both female groups, which might have influenced the results. The maximum tail length of the Fresh females exceeded that of the Fresh males.

The culmen, wing and tarsus lengths and the masses reported here are within the ranges of Bowie (2005a) and Rose et al. (2020); however, Bowie (2005a) reported unsexed biometric data for most of the biometric parameters. The mean tail lengths were longer than those of Bowie (2005a) and Rose et al. (2020) but similar to those of Paijmans and Bryson (2023). However, Namibian birds' body masses were lower than reported here (Paijmans and Bryson 2023).

Brubru Nilaus afer brubru

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Table 2a. Summary statistics of biometric data for selected species in the Malaconotidae based on SAFRING data collected by DHDS from 1987 to 2023 and samples curated after 1995 by DHDS at the National Museum, Bloemfontein. Sex-based differences in the biometric data were evaluated using Analysis of Variance except in cases where the data were not normally distributed, in which case the Kruskal-Wallis test was used (indicated by "KW" besides the p-value). The p-values in bold indicate significant differences in biometric parameters between the sexes. Where statistical analyses were not possible due to insufficient sample numbers, the p-value is replaced with #.

			F (n = 7)				I	M (n = 10)			р	Test
	n	mean ± SD	median	min	max	n	mean ± SD	median	min	max		
Culmen	6	23.2 ± 0.9	22.9	22.2	24.7	9	22.4 ± 2.4	21.9	19.5	26.3	0.489	
Tarsus	5	33.6 ± 2.5	32.5	30.8	37.0	7	33.1 ± 1.6	33.0	30.5	35.5	0.637	
Wing	7	97.4 ± 2.7	97.0	95.0	101.0	10	100.9 ± 3.3	100.5	97.0	107.0	0.037	
Tail	7	102.3 ± 9.5	100.0	88.0	118.0	9	100.1 ± 7.4	100.0	88.0	115.0	0.615	
Mass	4	66.8 ± 6.2	67.8	59.7	72.0	8	63.3 ± 7.6	64.8	47.4	72.5	0.447	

			F (n = 7)				I	M (n = 21)				
	n	mean ± SD	median	min	max	n	mean ± SD	median	min	max		
Culmen	6	17.3 ± 0.9	17.1	16.5	18.9	20	17.9 ± 1.3	18.1	15.1	20.1	0.222	
Tarsus	4	26.4 ± 0.3	26.3	26.2	26.8	15	25.9 ± 0.7	25.9	24.4	27.3	0.428	
Wing	6	77.2 ± 1.8	77.0	75.0	80.0	20	77.0 ± 3.1	77.5	72.0	83.0	0.910	
Tail	5	101.4 ± 6.7	100.0	95.0	112.0	20	96.2 ± 5.6	96.5	84.0	110.0	0.093	
Mass	7	36.6 ± 2.7	36.0	33.1	40.0	21	35.8 ± 4.5	36.1	25.0	43.0	0.727	

3. Brubru Nilaus afer brubru

			F (n = 3)					M (n = 18)				
	n	mean ± SD	median	min	max	n	mean ± SD	median	min	max		
Culmen	2	14.9 ± 0.1	14.9	14.8	14.9	16	15.2 ± 1.3	15.1	12.1	16.9	0.763	
Tarsus	2	21.7 ± 0.6	21.7	21.3	22.1	11	22.5 ± 0.4	22.6	21.5	22.9	0.057	KW
Wing	2	85.5 ± 2.1	85.5	84.0	87.0	18	85.2 ± 2.9	85.0	78.0	90.0	0.876	
Tail	2	57.5 ± 0.7	57.5	57.0	58.0	17	58.3 ± 2.5	59.0	54.0	62.0	0.667	
Mass	3	21.9 ± 1.8	22.3	20.0	23.5	18	25.2 ± 1.8	25.0	21.2	29.4	0.008	

4. Crimson-breasted Shrike Laniarius atrococcoineus

			F (n = 1)					M (n = 11)				
	n	mean ± SD	median	min	max	n	mean ± SD	median	min	max		
Culmen	1	20.0	20.0	20.0	20.0	9	20.8 ± 1.3	20.9	18.2	23.3	#	
Tarsus	1	31.1	31.1	31.1	31.1	8	32.9 ± 2.7	32.5	30.3	39.0	#	
Wing	1	94.0	94.0	94.0	94.0	11	99.2 ± 3.3	100.0	93.0	104.0	#	
Tail	1	98.0	98.0	98.0	98.0	10	101.1 ± 2.7	100.0	98.0	107.0	#	KW
Mass	1	42.0	42.0	42.0	42.0	11	47.4 ± 3.4	47.0	41.6	53.3	#	

Table 2b. Summary statistics of biometric data for selected species in the Malaconotidae based on museum skin specimens collected pre-1995 (Dry specimens) and held in the ornithological collection of the National Museum, Bloemfontein. Sex-based differences in the biometric data were evaluated using Analysis of Variance except where the data were not normally distributed, in which case the Kruskal-Wallis test was used (indicated by "KW" besides the p-value). The p-values in bold indicate significant differences in biometric parameters between the sexes. Where statistical analyses were not possible due to insufficient sample numbers, the p-value is replaced with #.

1. Bokmaki	erie 7	Telephorus zeylo	1. Bokmakierie Telephorus zeylonus														
			F (n = 4)					M (n = 10)			р	Test					
	n	mean ± SD	median	min	max	n	mean ± SD	median	min	max							
Culmen	3	22.4 ± 1.4	23.2	20.8	23.2	4	21.9 ± 1.1	21.6	21.1	23.4	0.631						
Tarsus	3	31.2 ± 1.6	30.8	29.8	32.9	5	32.7 ± 1.3	32.5	31.2	34.0	0.437						
Wing	3	96.0 ± 3.6	95.0	93.0	100.0	6	98.2 ± 2.0	98.5	96.0	100.0	0.275						
Tail	3	96.7 ± 5.7	95.0	92.0	103.0	6	97.3 ± 8.8	96.5	87.0	112.0	0.909						
Mass	3	64.2 ± 4.2	65.0	59.7	68.0	10	69.4 ± 10.3	69.7	54.5	84.0	0.429						
2. Brown-c	rown	ed Tchagra Tcha	agra australis	damarens	is												
			F (n = 6)					M (n = 6)									
	n	mean ± SD	median	min	max	n	mean ± SD	median	min	max							
Culmen	5	16.7 ± 1.1	16.7	15.3	17.9	4	18.5 ± 0.8	18.2	17.8	19.6	0.038						
Tarsus	4	27.0 ± 2.6	26.6	24.8	30.0	3	25.6 ± 1.2	25.1	24.7	26.9	0.882	KW					
Wing	5	76.0 ± 1.9	76.0	73.0	78.0	4	76.8 ± 1.7	76.5	75.0	79.0	0.555						
Tail	5	95.2 ± 6.4	95.0	88.0	103.0	4	91.8 ± 4.6	92.5	86.0	96.0	0.397						

I. DUKIIIAK		lelephonus zeyh	511u3 2091011u3	•									
			F (n = 4)					M (n = 10)			р	Test	
	n	mean ± SD	median	min	max	n	mean ± SD	median	min	max			
Culmen	3	22.4 ± 1.4	23.2	20.8	23.2	4	21.9 ± 1.1	21.6	21.1	23.4	0.631		
Tarsus	3	31.2 ± 1.6	30.8	29.8	32.9	5	32.7 ± 1.3	32.5	31.2	34.0	0.437		
Wing	3	96.0 ± 3.6	95.0	93.0	100.0	6	98.2 ± 2.0	98.5	96.0	100.0	0.275		
Tail	3	96.7 ± 5.7	95.0	92.0	103.0	6	97.3 ± 8.8	96.5	87.0	112.0	0.909		
Mass	3	64.2 ± 4.2	65.0	59.7	68.0	10	69.4 ± 10.3	69.7	54.5	84.0	0.429		
2. Brown-crowned Tchagra Tchagra australis damarensis													
			F (n = 6)					M (n = 6)					
	n	mean ± SD	median	min	max	n	mean ± SD	median	min	max			
Culmen	5	16.7 ± 1.1	16.7	15.3	17.9	4	18.5 ± 0.8	18.2	17.8	19.6	0.038		
Tarsus	4	27.0 ± 2.6	26.6	24.8	30.0	3	25.6 ± 1.2	25.1	24.7	26.9	0.882	KW	
Wing	5	76.0 ± 1.9	76.0	73.0	78.0	4	76.8 ± 1.7	76.5	75.0	79.0	0.555		
Tail	5	95.2 ± 6.4	95.0	88.0	103.0	4	91.8 ± 4.6	92.5	86.0	96.0	0.397		
Mass	6	35.4 ± 1.8	35.0	33.5	38.5	6	34.9 ± 4.0	35.0	29.5	40.6	0.801		

3. Brubru Nilaus afer brubru

			F (n = 1)					M (n = 4)				
	n	mean ± SD	median	min	max	n	mean ± SD	median	min	max		_
Culmen						3	14.7 ± 0.6	14.8	14.1	15.2	#	_
Tarsus						1	21.5	21.5	21.5	21.5	#	
Wing						3	87.3 ± 1.5	87.0	86.0	89.0	#	
Tail						3	61 ± 1.7	60.0	60.0	63.0	#	
Mass	1	26.1	26.1	26.1	26.1	4	26.2 ± 3.7	26.5	21.5	30.1	#	

4. Crimson-breasted Shrike Laniarius atrococcoineus

			F (n = 0)					M (n = 6)				
	n	mean ± SD	median	min	max	n	mean ± SD	median	min	max		
Culmen						5	21.3 ± 0.8	21.5	20.0	21.9	#	
Tarsus						1	33.1	33.1	33.1	33.1	#	
Wing						5	102.8 ± 4.5	101.0	98.0	110.0	#	
Tail						5	108.6 ± 5.2	110.0	100.0	113.0	#	
Mass						6	52.1 ± 6.7	53.0	40.2	60.0	#	





lengths of Namibian birds are longer than reported here, but this may be due to Paijmans and Bryson (2023) using the skull line measurement.

Crimson-breasted Shrike Laniarius atrococcineus

Biometric data for 12 Fresh and 6 Dry Laniarius atrococcineus specimens are summarised in Tables 2a.4 and 2b.4, respectively. No statistical comparisons between the sexes were possible because, for females, only one Fresh specimen was measured.

The maximum tail length and maximum mass for Dry males exceeded those of Fresh males.

Biometrics of ten passerine species

The tarsus, wing, and tail lengths and mass reported here are within the ranges recorded by Bowie (2005c), Rose et al. (2020), and Paijmans and Bryson (2023). However, the mean body mass for males, mainly from Free State Kalahari thornveld, was lower than those of Rose et al. (2020) but similar to Paijmans and Bryson's (2023) value. However,

Table 3a. Summary statistics of biometric data for selected species in the Platysteiridae based on SAFRING data collected by DHDS from 1987 to 2023 and samples curated after 1995 by DHDS at the National Museum, Bloemfontein. Sex-based differences in the biometric data were evaluated using Analysis of Variance except in cases where the data were not normally distributed, in which case the Kruskal-Wallis test was used (indicated by "KW" besides the p-value). The p-values in bold indicate significant differences in biometric parameters between the sexes. Where statistical analyses were not possible due to insufficient sample numbers, the p-value is replaced with #.

1. Cape Bat	is Batis	capensis hollio	layi									
		F	⁼ (n = 13)				Ν	/I (n = 11)			р	Test
	n	mean ± SD	median	min	max	n	mean ± SD	median	min	max		
Culmen	6	11.6 ± 0.7	11.6	10.8	12.6	9	11.9 ± 0.7	11.9	10.8	12.9	0.155	
Tarsus	5	22.0 ± 0.5	21.9	21.6	22.8	7	21.2 ± 1.0	21.1	20.1	23.0	0.881	
Wing	13	64.0 ± 3.4	63.0	58.0	70.0	10	64.7 ± 2.8	65.5	60.0	68.0	0.949	
Tail	8	48.4 ± 4.1	49.0	40.0	53.0	10	51.3 ± 4.7	52.5	45.0	60.0	0.264	
Mass	12	12.3 ± 1.1	12.5	9.9	13.9	9	13.4 ± 1.2	13.5	10.9	14.5	0.046	

2. Chinspot Batis Batis molitor palliditergum

			F (n = 2)				I	M (n = 6)			
-	n	mean ± SD	median	min	max	n	mean ± SD	median	min	max	
Culmen	2	12.5 ± 0.1	12.5	12.4	12.5	6	12.6 ± 1	12.5	11.2	14.2	0.884
Tarsus	2	18.4 ± 0.6	18.4	17.9	18.8	5	18.3 ± 1.1	18.0	16.8	19.8	0.940
Wing	2	60.0 ± 2.8	60.0	58.0	62.0	6	61.5 ± 1.6	61.0	60.0	64.0	0.369
Tail	2	48.5 ± 0.7	48.5	48.0	49.0	6	47.3 ± 2.3	47.0	45.0	50.0	0.532
Mass	2	11.0 ± 1.2	11.0	10.1	11.8	6	11.5 ± 0.8	11.4	10.4	12.5	0.468

3. Pririt Batis Batis pririt pririt

		F	⁼ (n = 51)				Ν	/l (n = 50)				
-	n	mean ± SD	median	min	max	n	mean ± SD	median	min	max		
Culmen	37	10.4 ± 0.8	10.3	8.3	11.6	39	11.1 ± 0.8	10.9	9.8	12.9	0.001	
Tarsus	33	17.7 ± 1.0	17.8	16.2	20.4	28	17.5 ± 1.4	17.5	13.8	21.3	0.648	KW
Wing	46	55.9 ± 1.9	56.0	52.0	65.0	48	56.9 ± 2.4	57.0	48.0	68.0	0.001	KW
Tail	41	46.1 ± 2.9	46.0	40.0	55.0	44	46.0 ± 2.8	45.0	40.0	51.0	0.776	KW
Mass	51	9.1 ± 1.2	8.9	6.9	12.5	50	9.1 ± 1	8.9	7.5	12.0	0.788	KW

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the culmen lengths of Namibian birds were longer male samples for statistical analyses. The following than reported here (Paijmans and Bryson 2023). measurements were outside the ranges for the Fresh male specimens: maximum culmen length Platysteiridae for the Dry females, minimum tarsus length for the Cape Batis Batis capensis hollidayi Dry males, and both minimum and maximum wing Biometric data for 24 Fresh and 7 Dry Batis capenlengths, minimum tail length and minimum mass sis hollidayi specimens are summarised in Tables for the Fresh females. The data for B. c. hollidayi 3a.1 and 3b.1, respectively. Only the mass of the represents newly published biometric data for this Fresh male and female specimens differed signifsubspecies, as Smith (2005a) only listed biometric icantly (Table 3a.1). There were insufficient Dry data of B. c. capensis.

Table 3b.Summary statistics of biometric data for selected species in the Platysteiridae based on museum skin specimens collected pre-1995 (Dry specimens) and held in the ornithological collection of the National Museum, Bloemfontein. Sex-based differences in the biometric data were evaluated using Analysis of Variance except where the data were not normally distributed, in which case the Kruskal-Wallis test was used (indicated by "KW" besides the p-value). The p-values in bold indicate significant differences in biometric parameters between the sexes. Where statistical analyses were not possible due to insufficient sample numbers, the p-value is replaced with #.

			F (n = 5)				1	M (n = 2)			р	Tes
	n	mean ± SD	median	min	max	n	mean ± SD	median	min	max	P	
Culmen	2	13.1 ± 0.1	13.1	13.0	13.2	1	11.3	11.3	11.3	11.3	#	
Tarsus	-	23.0	23.0	23.0	23.0	. 1	19.7	19.7	19.7	19.7	" #	
Wing	2	63 ± 2.0	63.0	61.0	65.0	1	67.0	67.0	67.0	67.0	" #	
Tail	2	50 ± 2.0	50.0	50.0	50.0	1	51.0	51.0	51.0	51.0	#	
Mass	5	12.4 ± 0.9	12.5	11.1	13.3	2	13.1 ± 1.5	13.2	12.1	14.2	" 0.410	
		Batis molitor pal										
			F (n = 1)				l					
	n	mean ± SD	median	min	max	n	mean ± SD	median	min	max		
Culmen	1	12.2	12.2	12.2	12.2						#	
Tarsus											#	
Wing	1	55.0	55.0	55.0	55.0						#	
Tail	1	46.0	46.0	46.0	46.0						#	
Mass	1	9.0	9.0	9.0	9.0	2	11.7 ± 0.9	11.7	10.8	12.6	#	
3. Pririt Bat	is Batis	pririt pririt										
			F (n = 4)									
	n	mean ± SD	median	min	max	n	mean ± SD	median	min	max		
Culmen	4	11.4 ± 0.7	11.7	10.3	11.9	8	11.6 ± 0.8	11.7	10.5	12.8	0.620	
Tarsus	1	15.3	15.3	15.3	15.3	3	17.7 ± 0.1	17.8	17.6	17.8	0.157	K١
Wing	4	56 ± 1.8	56.0	54.0	58.0	8	56.5 ± 1.4	57.0	54.0	58.0	0.610	
Tail	4	46.8 ± 4.1	46.0	43.0	52.0	7	47.1 ± 2.9	47.0	42.0	50.0	0.855	
Mass	4	9.4 ± 0.9	9.8	8.0	10.0	7	9.8 ± 1.1	10.0	8.4	11.0	0.539	



The wing and tail lengths reported here were longer than those of Smith (2005a) and Rose et al. (2020), while the culmen lengths were shorter. Smith (2005a) only presented mass data for B. c. erythrophthalma from the eastern highlands of Zimbabwe. The mass data of Rose et al. (2020) (for all subspecies) was similar to the values reported here.

Chinspot Batis Batis molitor palliditergum

Biometric data for 8 Fresh and 3 Dry Batis molitor *palliditergum* specimens are summarised in Tables 3a.2 and 3b.2, respectively. There were insufficient Dry samples for statistical analyses. No significant differences between the sexes were found for the Fresh specimens (Table 3a.2); however, only two female specimens were included in the analyses. The minimum wing length and minimum mass for the Fresh and Dry female specimens exceeded the range of the Fresh male specimens.

The biometric data reported here for *B. m. palliditergum* are similar to those of Smith (2005b) and Rose et al. (2020). However, Smith (2005b) only reported on B. *m. molitor*, and Rose et al. (2020) did not distinguish between subspecies.

Pririt Batis Batis pririt pririt

Biometric data for 101 Fresh and 12 Dry Batis pririt pririt specimens are summarised in Tables 3a.3 and 3b.3. respectively. No significant differences were found for the Dry specimens (Table 3b.3). The culmen length and wing length of the Fresh males were significantly larger than those of the Fresh females (Table 3b.3). The maximum tail length of both the Fresh and Dry female specimens and the minimum and maximum mass for the Fresh female specimens exceeded the range of the Fresh male specimens.

Tarsus, wing and tail lengths of *B*. *p*. *pririt* in this study were similar to those of Spottiswoode (2005) and Rose et al. (2020), but the culmen lengths were shorter than those of Spottiswoode (2005). The Namibian subspecies B. p. affinis have similar ranges for all biometrics except mass, which was lower than reported here for B. p. pririt (Paijmans and Bryson 2023).

Laniidae

Southern Fiscal Lanius collaris (mixed subspecies L. c. collaris and L. c. subcoronatus)

Biometric data for 118 Fresh and 37 Dry Lanius collaris specimens are summarised in Tables 4a.1 and 4b.1,

respectively. The data set was not recorded on the subspecies level because Fuchs et al. (2011) found extensive gene flow among southern African populations and limited molecular divergence between L. collaris subspecies characterised by eyebrow markings. The white eyebrow is a phenotypic adaptation for foraging in arid environments (Fuchs et al. 2011). Significant differences between the sexes were found in wing length for both the Dry and Fresh specimens (Tables 4a.1 and 4b.1). In addition, significant differences were found in mass for the Fresh specimens (Table 4a.1) and tail length for the Dry specimens (Table 4b.1). The minimum culmen length of the females and Dry males exceeded the range of the Fresh male specimens (Tables 4a.1 and 4b.1).

The wing and tail lengths and masses reported here have similar ranges and means to those of Rose et al. (2020). Dean (2005a) did not report means for wing, tail, tarsus, and culmen lengths but reported mass data for both sexes lower than those reported here. Biometric data for the Namibian subspecies L. c. aridicolus reported similar ranges for all parameters except the masses, which were also lower than those reported here (Paijmans and Bryson, 2023).

Red-backed Shrike Lanius collurio

Biometric data for 18 Fresh and 3 Dry Lanius collurio specimens are summarised in Tables 4a.2 and 4b.2, respectively. No statistical comparisons were possible for the Dry specimens due to the low sample numbers (Table 4b.2). No significant differences between the sexes were found for the Fresh specimens (Table 4a.2). Only the minimum tail length for the Dry male specimens exceeded the range for the Fresh males.

Herremans (2005) listed biometric data for mixed Palaearctic subspecies of Lanius collurio similar to those in this study. The wing and tail lengths and mass reported here fall within the range of values reported by Rose et al. (2020) and Paijmans and Bryson (2023).

Magpie Shrike Lanius melanoleucus melanoleucus

Biometric data for 11 Fresh and 13 Dry Lanius melanoleucus melanoleucus specimens are summarised in Tables 4a.3 and 4b.3, respectively. No significant differences were recorded between biometric data for the Fresh males and females (Table 4a.2), but the culmen lengths for the Dry specimens were significantly different (Table 4b.2). The following parameters exceeded the

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Table 4a. Summary statistics of biometric data for selected species in the Laniidae based on SAFRING data collected by DHDS from 1987 to 2023 and samples curated after 1995 by DHDS at the National Museum, Bloemfontein. Sex-based differences in the biometric data were evaluated using Analysis of Variance except in cases where the data were not normally distributed, in which case the Kruskal-Wallis test was used (indicated by "KW" besides the p-value). The p-values in bold indicate significant differences in biometric parameters between the sexes. Where statistical analyses were not possible due to insufficient sample numbers, the p-value is replaced with #.

1. Southern Fiscal Lanius collaris (mixed subspecies L. c. collaris and L. c. subcoronatus)													
		F	⁼ (n = 59)				Ν	р	Test				
-	n	mean ± SD	median	min	max	n	mean ± SD	median	min	max			
Culmen	34	16.1 ± 1.8	15.8	13.3	20.3	37	16.7 ± 1.2	16.5	15.1	20.5	0.137		
Tarsus	24	26.1 ± 2.3	25.8	22.2	31.0	26	27.4 ± 2.2	27.3	22.1	31.5	0.078		
Wing	57	97.2 ± 2.6	98.0	89.0	103.0	64	100.2 ± 5.7	99.0	88.0	130.0	< 0.001	KW	
Tail	44	109.5 ± 6.6	110.0	91.0	121.0	52	111.5 ± 10.9	112.0	77.0	130.0	0.109	KW	
Mass	57	41.6 ± 4.8	41.5	30.2	60.0	64	43.9 ± 4.5	43.5	30.0	55.5	0.001	KW	

2. Red-backed Shrike Lanius collurio

	F (n = 10)						l				
	n	mean ± SD	median	min	max	n	mean ± SD	median	min	max	
Culmen	6	14.3 ± 1.1	14.7	12.3	15.3	6	14.5 ± 1.8	13.9	13.0	17.8	0.808
Tarsus	4	22.5 ± 6.3	23.9	13.8	28.5	5	23.3 ± 2.1	22.3	22.1	27.0	0.796
Wing	10	91.3 ± 3.2	90.5	87.0	96.0	8	90.0 ± 3.3	89.0	86.0	95.0	0.428
Tail	6	77.0 ± 2.8	78.0	72.0	80.0	6	76.7 ± 3.2	76.5	72.0	82.0	0.851
Mass	10	27.3 ± 2.3	28.5	23.0	29.1	8	28.7 ± 2.2	29.1	25.5	31.0	0.206

3. Magpie Shrike Lanius melanoleucus melanoleucus

	F (n = 5)											
	n	mean ± SD	median	min	max	n	mean ± SD	median	min	max		
Culmen	4	19.6 ± 0.4	19.7	19.1	20.0	6	19.1 ± 2	19.2	16.1	21.2	0.617	
Tarsus	3	33.7 ± 2.0	33.1	32.1	35.9	5	35.3 ± 1.6	35.4	33.2	36.8	0.251	
Wing	5	135.4 ± 2.6	136.0	132.0	139.0	6	138.7 ± 3.2	140.0	133.0	142.0	0.101	
Tail	4	269.0 ± 20.0	272.5	245.0	286.0	5	280.4 ± 33.9	293.0	220.0	300.0	0.142	KW
Mass	5	84.8 ± 8.0	83.0	76.0	98.0	6	95.3 ± 8.2	95.0	83.4	106.2	0.062	

range of the Fresh male specimens: minimum tarsus et al. (2020), although higher mean body masses are length for both Fresh and Dry females, minimum wing recorded for males and females here. Dean (2005b) length for Fresh females and Dry males, minimum tail listed unsexed data for culmen and tarsus lengths. length for Dry females, maximum tail length for both Dry males and females, minimum mass for Fresh females, Conclusions and maximum mass for Dry females.

This paper demonstrates the use of museum and ringing data to generate biometric data comparable to those The tail lengths found in the study were considerably of other studies. The study emphasises the importance shorter than the >340.0 mm maximum tail lengths reof compiling comprehensive biometric data for birds ported by Dean (2005b). Wing and tail biometrics were ringed or sacrificed for museum collections. We concur within the ranges reported by Dean (2005b) and Rose with Rose et al. (2020) that a ringed or sacrificed "bird



8



Biometrics of ten passerine species

Table 4b. Summary statistics of biometric data for selected species in the Laniidae based on museum skin specimens collected pre-1995 (Dry specimens) and held in the ornithological collection of the National Museum, Bloemfontein. Sex-based differences in the biometric data were evaluated using Analysis of Variance except where the data were not normally distributed, in which case the Kruskal-Wallis test was used (indicated by "KW" besides the p-value). The p-values in bold indicate significant differences in biometric parameters between the sexes. Where statistical analyses were not possible due to insufficient sample numbers, the p-value is replaced with #.

		F	⁼ (n = 12)				Ν		р	Test		
	n	mean ± SD	median	min	max	n	mean ± SD	median	min	max		
Culmen	5	14.6 ± 1.0	15.2	13.1	15.4	6	15.8 ± 0.6	15.7	14.9	16.8	0.048	
Tarsus	1	26.2	26.2	26.2	26.2						#	
Wing	5	95.6 ± 4.7	97.0	89.0	101.0	6	101.3 ± 3.2	100.0	98.0	107.0	0.039	KW
Tail	5	105.2 ± 4.5	103.0	100.0	110.0	6	111.5 ± 2.9	110.5	108.0	115.0	0.021	KW
Mass	12	41.6 ± 6.2	40.0	34.7	52.0	25	40.7 ± 6.1	40.9	27.2	50.0	0.676	KW

	F (n = 1)						I				
	n	mean ± SD	median	min	max	n	mean ± SD	median	min	max	
Culmen	1	13.8	13.8	13.8	13.8	2	15.1 ± 0.1	15.1	15.0	15.1	#
Tarsus						2	26.2 ± 1.2	26.2	25.3	27.0	#
Wing	1	90.0	90.0	90.0	90.0	2	91.0 ± 1.4	91.0	90.0	92.0	#
Tail	1	75.0	75.0	75.0	75.0	2	74.0 ± 8.5	74.0	68.0	80.0	#
Mass						2	31.9 ± 4.5	31.9	28.7	35.0	#

3. Magpie Shrike Lanius melanoleucus melanoleucus

	F (n = 6)						I					
	n	mean ± SD	median	min	max	n	mean ± SD	median	min	max		
Culmen	3	18.7 ± 0.8	19.1	17.8	19.1	5	20.1 ± 0.6	20.0	19.2	20.8	0.029	
Tarsus	3	34.2 ± 1.4	34.8	32.6	35.1	4	35.4 ± 0.8	35.5	34.5	36.1	0.196	
Wing	3	137.7 ± 1.5	138.0	136.0	139.0	5	134.4 ± 5.6	135.0	125.0	140.0	0.294	KW
Tail	3	271.3 ± 53.8	285.0	212.0	317.0	5	293.6 ± 9.5	295.0	283.0	305.0	0.764	KW
Mass	6	100.2 ± 14.8	95.5	85.0	125.0	7	95.7 ± 4.1	95.5	89.0	102.0	0.886	KW

with no associated data has limited scientific use", highlighting the responsibility of researchers and bird ringers to record all reasonable biometric parameters as comprehensively and as accurately as possible.

Even though there are very low sample numbers for some of the species included in this study, the data presented represents the first published biometric data records for several subspecies in the Malaconotidae, Platysteiridae, and Laniidae. The biometric data for these subspecies could be improved by including specimens from other museum collections and incorporating ringing data from other bird ringers operating in the study area. The data could be supplemented with SAFRING data if the sex of the bird is specified in the biometric data and a subspecies can be confidently assigned to the record.

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