

# Low Geographic and Subspecific Variation in the Loud Call of the Widespread and Phenotypically Cryptic Northern Lesser Galago (*Galago senegalensis*) Suggests Taxonomic Uniformity

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## ABSTRACT

Like other nocturnal primates, many species of galago (Galagidae) are phenotypically cryptic, making their taxonomic status difficult to resolve. Recent taxonomic work has disentangled some of the confusion. This has resulted in an increase in the number of recognised galago species. The most widespread galago species, and indeed the most widespread nocturnal primate, is the northern lesser galago (*Galago senegalensis*) whose geographic range stretches >7,000 km across Africa. Based on morphology, 4 subspecies are currently recognised: *G. s. senegalensis*, *G. s. braccatus*, *G. s. sotikae* and *G. s. dunni*. We explore geographic and subspecific acoustic variation in *G. senegalensis*, testing three hypotheses: isolation by distance, genetic basis, and isolation by barrier. There is statistical support for isolation by distance for 2 of 4 call parameters (fundamental frequency and unit length). Geographic distance explains a moderate amount of the acoustic variation. Discriminant function analysis provides some degree of separation of geographic regions and subspecies, but the percentage of misdesignation is

high. Despite having (putative) parapatric geographic ranges, the most pronounced acoustic differences are between *G. s. senegalensis* and *G. s. durni*. The findings suggest that the Eastern Rift Valley and Niger River are significant barriers for *G. senegalensis*. The acoustic structures of the loud calls of 121 individuals from 28 widespread sites are not significantly different. Although this makes it unlikely that additional unrecognised species occur within *G. senegalensis* at the sites sampled, vast areas of the geographic range remain unsampled. We show that wide-ranging species do not necessarily exhibit large amounts of variation in their vocal repertoire. This pattern may also be present in nocturnal primates with smaller geographic ranges.

## INTRODUCTION

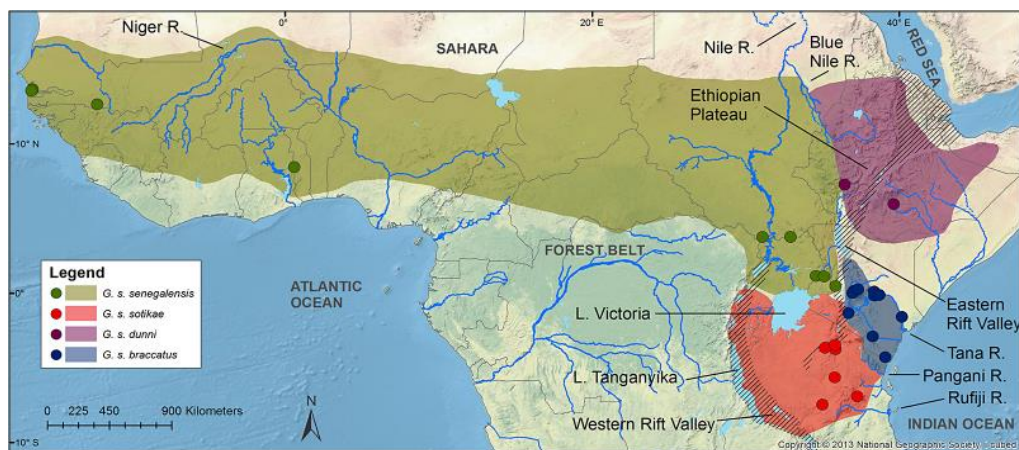
Geographic variation and dialects in vocalizations have been described within many species of vertebrates (e.g., Anura [Velásquez, 2014]; Psittaciformes [Brigham and Cebek, 1989]; Cetacea [Ford, 1991]; Chiroptera [Wright and Dahlin, 2018]; Rodentia [Slobodchikoff et al., 1998]). Intraspecies variation arises when differences in acoustic structure are larger among populations than within populations. While geographic variation relates to isolated populations, dialects arise in populations where gene flow is not restricted by distance or geographic barriers [Conner, 1982]. In primates, intraspecies geographic acoustic variation or dialects is documented for many taxa, including robust chimpanzees (*Pan troglodytes*) [Mitani et al., 1999], orang-utans (*Pongo* spp.) [Delgado et al., 2003], silvery gibbons (*Hylobates moloch*) [Dallmann and Geissmann, 2009], Japanese macaques (*Macaca fuscata*) [Green, 1975], vervet monkeys (*Chlorocebus pygerythrus*) [Price et al., 2014], Thomas's langurs (*Presbytis thomasi*) [Wich et al., 2008], red-chested moustached tamarins (*Saguinus labiatus*) [Maeda and Masataka, 1987], and small-eared greater galago (*Otolemur garnettii*) [Bettridge et al., in press]. Although there is ample evidence of intraspecies geographic variation in primate vocalizations, mechanisms that contribute to this variation are poorly understood [Dallmann and Geissmann, 2009]. Morphology, ecological selection, sexual selection, cultural transmission, and neutral processes (e.g.,

genetic drift) all potentially contribute to intraspecies acoustic variation. Indeed, some of these factors contribute simultaneously [Wilkins et al., 2013]. Studying intraspecies variation is important as it can significantly contribute to understanding the evolutionary processes that lead to variation in phenotypic traits [Clink et al., 2018]. Rapidly speciating lineages, such as swordtail crickets (Trigonidiinae), cicadas (Cicadoidea), and green lacewings (Chrysopidae), are predominantly identified by their calls [Wilkins et al., 2013]. As such, intraspecies divergence in acoustic signals may play a key role in speciation [Wilkins et al., 2013]. Moreover, vocalizations are a reliable tool for identifying cryptic (often nocturnal) species such as bats [Jones and Van Parijs, 1993], nightjars (Caprimulgidae) [Turner, 2011], frogs [Yuan and Ramli, 2013], and small-bodied primates [Zimmermann et al., 1988; Braune et al., 2008; Burton and Nietsch, 2010].

Galagos (Galagidae) are small-bodied (50–1,700 g) nocturnal primates endemic to sub-Saharan Africa where they occur in forest, woodland, and bushland [Bearder and Masters, 2013; Butynski et al., 2013; Nekaris, 2013]. The taxonomy of the Galagidae has long been contentious, primarily because many members of this family are phenotypically cryptic. With the help of molecular techniques, comparative morphology, and behavioural comparisons, including bioacoustics, the taxonomy of this family is now better understood. There has been an increase in the number of species recognised within Galagidae: Schwarz [1931] recognised 5 species, Hill [1953] 6, Jenkins [1987] 7, Olson [1979] 11, Bearder et al. [1995] and Honess [1996] 17, Kingdon [1997] 18, Groves [2001] 23, Grubb et al. [2003] 24, Butynski et al. [2013] 18, Nekaris [2013] 18, and Rowe and Myers [2016] 20.

The galago, and indeed the nocturnal primate, with the widest (>7,000 km) and largest geographic range, the northern lesser galago (*Galago senegalensis*), occurs from the Atlantic Ocean in West Africa to the Red Sea and Indian Ocean in eastern Africa (Fig. 1) [Nash et al., 2013; Nekaris, 2013]. Based on morphological, behavioural, genetic, and acoustic data, 7 species have been split from *G. senegalensis*: southern lesser galago (*Galago moholi*), Somali lesser galago (*Galago gallarum*), spectacled lesser galago (*Galago matschiei*), Malawi dwarf galago (*Paragalago nyasae*), Zanzibar dwarf galago (*Paragalago zanzibaricus*), Mozambique dwarf galago (*Paragalago granti*), and Kenya coast dwarf

galago (*Paragalago cocos*) [Zimmermann et al., 1988; Nash et al., 1989; Zimmermann, 1990; Anderson, 1999, 2001; Bearder, 1999; Anderson et al., 2000; DelPero et al., 2000; Masters and Bragg, 2000; Groves, 2001, 2005; Grubb et al., 2003; Butynski and De Jong, 2004; Roos et al., 2004; Masters et al., 2007; Nekaris and Bearder, 2011; Nash et al., 2013].



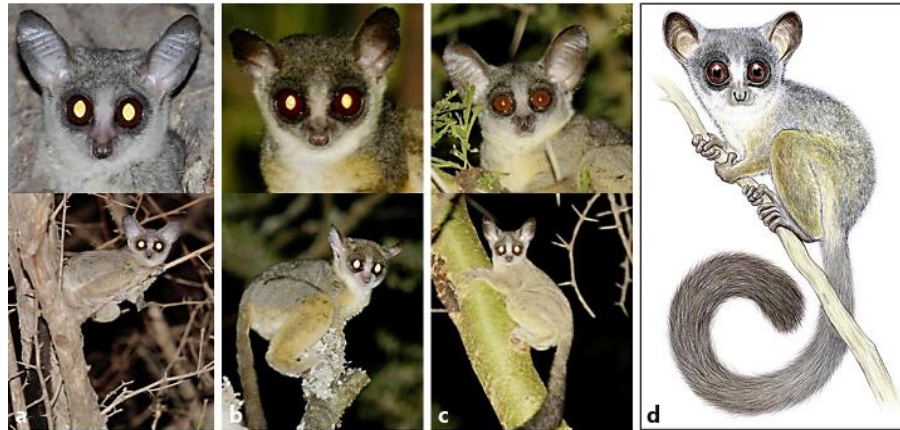
**Fig. 1.** Geographic ranges of the 4 subspecies of northern lesser galago (*Galago senegalensis*). Also shown are the sites at which honk calls were recorded and the more likely barriers to gene flow.

Species identification based on morphology is appropriate for animals that recognise one another by sight (as is typically the case for diurnal primates), but potentially problematic for nocturnal animals that mostly rely on scent and sound. Many nocturnal primates, including galagos, emit loud calls (also referred to as advertising calls or long-range calls) in the context of attracting companions and repelling rivals [Bearder et al., 1995]. The loud calls of galagos are highly species-specific and, therefore, are excellent tools for species identification [Masters, 1993; Bearder et al., 1995]. For example, formerly conspecific *G. moholi* and *G. senegalensis* were elevated to species level due to differences in their homologous loud calls and vocal repertoires [Zimmermann et al., 1988; Zimmermann, 1990; Bearder et al., 1995; Anderson et al., 2000]. Genetic data later confirmed this taxonomy [Pozzi et al., 2015]. The distinctive loud call of *G. gallarum*, previously considered a subspecies of *G. senegalensis*, confirmed its species status [Butynski and De Jong, 2004, 2013]. Recently, the Angolan dwarf galago (*Galagoides kumbirensis*) was recognised partly based on loud call and vocal repertoire [Svensson et al., 2017].

Although there is extensive information on species specificity of loud calls in galagos, there is little knowledge of intraspecies variation of these calls and mechanisms that contribute to this variation.

*Galago senegalensis* has an extensive vocal repertoire with at least 18 spectrographically different call types [Zimmermann, 1985; Zimmermann et al., 1988; Bearder et al., 1995]. *Galago senegalensis* belongs to the galago group of ‘repetitive callers’. Its loud call, also referred to as the ‘honk’ or ‘woo’, is arranged at regular intervals into ‘bouts’. Bouts may comprise hundreds of repeated honks [Bearder et al., 1995]. Vocal profile of *G. senegalensis* available at: <http://www.wildsolutions.nl/vocal-profiles/galago/senegalensis/>. Honk loud calls (hereafter referred to as honk call) show harmonic spectra with a fundamental frequency of about 400 Hz and slow frequency modulation. Honk calls can be heard by the human ear at >200 m under most conditions [Y. de Jong and T. Butynski, pers. observation] and are, therefore, suitable for long-distance communications among conspecifics [Zimmermann, 1985, 1995; Schneiderová et al., 2016]. Honk calls are emitted in gathering, contact, and display situations [Zimmermann, 1985; Bearder et al., 1995; Anderson et al., 2000; Svensson and Bearder, 2013]. Both sexes emit honk calls when leaving sleeping sites at dusk and when returning to sleeping sites at dawn, as well as throughout the night [Bearder et al., 2003]. Honk calls are particularly common during mating and birth periods [Bearder et al., 1995], and when population density is relatively high [Y. de Jong and T. Butynski, pers. observation]. *Galago senegalensis* is a particularly suitable species for investigating geographic and intraspecific vocal variation as it is polytypic, locally common, lives in a wide range of habitats, has a large geographic range with many potential barriers to gene flow, and has a honk call that is given in long bouts that are relatively easy to record and analyse.

In this study we examined variation in the honk call of *G. senegalensis* across the species’ geographic range, across subspecies, and across potential geographic barriers. The taxonomic arrangement applied here for *G. senegalensis* is that of Groves [2001], Grubb et al. [2003], Butynski et al. [2013], and Nekaris [2013], who recognise 4 subspecies (Fig. 2): Senegal lesser galago (*G. s. senegalensis*), Kenya lesser galago (*G. s. braccatus*), Tanzania lesser galago (*G. s. sotikae*), and Ethiopia lesser galago (*G. s. dunni*).



**Fig. 2.** Comparison of pelage colour and pattern, and ear length, of the 4 subspecies of northern lesser galago (*Galago senegalensis*): *G. s. senegalensis*, The Gambia (a); *G. s. braccatus*, central Kenya (b); *G. s. sotikae*, southwestern Kenya (c); *G. s. dunni* (d). Photographs by S.K. Bearder (a), Y.A. de Jong and T.M. Butynski (b and c). Drawing by Stephen Nash.

We tested three hypotheses, all of them based on the assumption that variation occurs within the honk call of *G. senegalensis* and that neutral processes contributed to this acoustic variation.

1 The isolation by distance hypothesis is based on the fact that differentiation among distant populations leads to evolutionary change [Wright, 1943]; a subspecies with a large geographic range is expected to have more within subspecies honk call variation than a subspecies with a small geographic range. In addition, a relationship between geographic distance and acoustic differences is expected throughout the range [e.g., Irwin et al., 2008; Campbell et al., 2010].

2 According to the genetic basis hypothesis, differences among the honk calls of the 4 subspecies have a genetic basis; honk call variation is expected to correspond to the 4 subspecies [e.g., Hill et al., 2013].

3 According to the isolation by barrier hypothesis, populations are isolated by geographic barriers (e.g., deep valleys, big rivers) [e.g., Baker et al., 2003]; geographic barriers are expected to explain honk call variation.

We also investigated the intraspecies variation of honk calls with respect to the potential existence of additional species and subspecies within *G. senegalensis*.

## METHODS

### *Study Species and Regions*

*Galago senegalensis*, a medium-sized galago (adult body weight approx. 200 g) [Jenkins, 1987; Groves, 2001; Nash et al., 2013], forages and sleeps alone or in small groups. Solitary individuals are, however, part of social networks [Bearder et al., 2003; Nash et al., 2013; Nekaris, 2013]. Although the 4 *G. senegalensis* subspecies are phenotypically similar, with an understanding of the diagnostic traits (Fig. 2; Table 1) and experience, they can be visually distinguished in the field [De Jong and Butynski, 2018a]. Several geographic barriers appear to isolate, or partly separate, these 4 subspecies (Fig. 1):

1 *G. s. senegalensis* (type locality Senegal): limited by Atlantic Ocean in the west; north by Sahara Desert; east by western edge of Ethiopian Plateau, Blue Nile River, Eastern (Gregory) Rift Valley; south by northern edge of Masai Mara Plains, northern edge of Lake Victoria, forest belt.

2 *G. s. braccatus* (type locality Tsavo, Kenya): limited by Eastern Rift Valley in the west; north by arid region (including Chalbi Desert); east by Tana River, Indian Ocean; south by Pangani River, northern lower slopes of Mount Kilimanjaro/Mount Meru.

3 *G. s. sotikae* (type locality Telek River, Sotik, Kenya): limited by Western (Albertine) Rift Valley (particularly Lake Tanganyika) in the west; north by Pangani River, southern lower slopes of Mount Kilimanjaro/Mount Meru, southern lower slope of Mau Escarpment, southern edge of Lake Victoria, forest belt; east by Indian Ocean; south by Rufiji River, Udzungwa Mountains, Southern Highlands.

4 *G. s. dunni* (type locality Fafan River, Somalia): limited by western edge of Ethiopian Plateau, Blue Nile River, Eastern Rift Valley in the west; north by northern edge of Ethiopian Plateau; east by eastern edge of Ethiopian Plateau, arid region; south by arid region (including Chalbi Desert); southwest by Eastern Rift Valley. Competition with *G. gallarum* might be a limitation to the east and south.

Within the geographic range of *G. senegalensis* we recognise 4 regions: (1) west of the Niger River (only *G. s. senegalensis*); (2) between the Niger River and the Nile River (only *G. s. senegalensis*);

(3) east of the Nile River and west of the Eastern Rift Valley (*G. s. senegalensis*, *G. s. sotikae*, *G. s. dunni*); (4) east of the Eastern Rift Valley and/or on the Ethiopian Plateau (*G. s. braccatus*, *G. s. sotikae*, *G. s. dunni*).

**Table 1.** Phenotypic characters which, together, distinguish the 4 subspecies of northern lesser galago (*Galago senegalensis*)

	<i>G. s. senegalensis</i>	<i>G. s. braccatus</i>	<i>G. s. sotikae</i>	<i>G. s. dunni</i>
Relative ear length	Medium	Short	Long	Short
Forehead	Grey, sometimes with buff wash	Grey with russet wash	Grey; yellow wash in southern part of range	Grey or greyish-brown
Dorsum	Grey	Grey to brownish-grey	Brownish-grey with yellow wash	Grey
Flanks	Grey with yellow wash	Grey with intense russet wash	Greyish-brown with yellow wash	Grey with yellow wash
Outer lower hindlegs	Grey, sometimes with yellow wash	Intense russet or bright buff	Grey, sometimes with yellow wash	Grey with yellow wash
Delineation of colours between upper and lower outer hindlegs	Weak or none	Sharp	None	Weak
Base of tail	Dark grey to greyish-brown	Brown with russet wash	Greyish-brown with yellow wash	Grey
Tip of tail	Blackish or brown	Blackish or brown	Dark grey or brown	Dark grey or brown

Based on Jenkins [1987], Groves [2001], De Jong and Butynski [2009, 2018a], Nash et al. [2013], and Svensson and Bearder [2013].

### *Study Sites and Honk Call Recordings*

*Galago senegalensis* honk calls included in this study were recorded during 1994–2018 from wild individuals at 28 sites (Fig. 1; Table 2). Of 131 recordings, 121 were suitable for further analyses. Galagos were detected by their calls or eye shine and observed at close proximity when feasible, for as long as possible. Some of the galagos were habituated or became accustomed to researcher(s) within hours [e.g., Svensson and Bearder, 2013]. Since the honk calls were recorded by various researchers over a 24-year period, the recording equipment varied from high-quality cassette recorders to digital recorders. Some recordings were obtained by an audio recorder (AudioMoth by Open Acoustic Devices) [Hill et al., 2018]. As quality and completeness of recordings varied, there was a limit to the number of acoustic parameters that could be reliably measured. For instance, for many recordings it was not possible to



assess whether the beginning and the end of the honk call were included. Hence, bout length, a call parameter commonly used in animal vocalization studies, was not considered in this study.

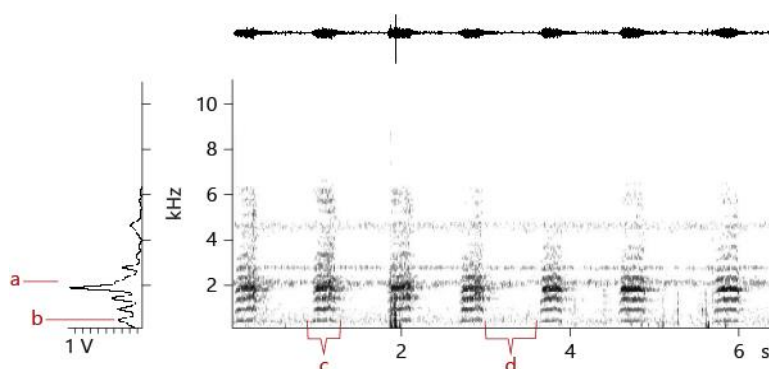
**Table 2.** Sites at which honk calls of northern lesser galago (*Galago senegalensis*) used in this study were obtained

Country	Site	Subspecies	Habitat	Altitude, m a.s.l.	Mean annual rainfall <sup>1</sup> , mm	Mean annual temperature <sup>1</sup> , °C	Recorded by	Call recordings, <i>n</i>
Kenya	Kora NP	b	Bushland, woodland	400	47	27	Y.J./T.B.	2
	Tsavo West NP	b	Bushland, woodland, grassland	850	64	25	Y.J./T.B.	1
	Tana River Primate Nat R	b	Forest, grassland, woodland, bushland	40	49	28	T.B./Y.J.	3
	Meru NP	b	Forest, woodland, bushland, grassland	370	99	26	Y.J./T.B.	10
	Borana Ranch, Laikipia	b	Woodland, bushland, grassland	1,900	97	18	T.B./C.D.S.	3
	Kithima Kia Mukuu NR, Isiolo	b	Bushland, woodland	1,550	99	21	Y.J./T.B.	1
	North Kilimangodo, Kwale	b	Bushland surrounded by agriculture	250	74	26	Y.J./T.B.	1
	Ololula Forest, Nairobi	b	Forest	1,800	76	19	T.B.	1
	Chebseon, Southwest Mau FR	so	Forest	2,350	95	19	T.B.	1
	Oi Pejeta Conservancy, Laikipia	b	Woodland, bushland, grassland	1,800	114	16	Y.J./T.B.	2
	Saiwa Swamp NP	se	Forest	1,850	94	18	Y.J./T.B.	2
	Mwingi Nat R	b	Forest, bushland, woodland	510	62	26	Y.J./T.B.	5
	Tumbili Estate, Laikipia	b	Woodland, bushland	1,800	56	25	Y.J./T.B.	5
Tanzania	Lake Manyara NP	so	Grassland, woodland, bushland	970	44	22	Y.J./T.B.	1
	Ruaha NP	so	Woodland	830	50	24	P.H.	2
	Lake Eyasi	so	Woodland	1,040	51	22	A.P.	31
	Meia Meia	so	Woodland, bushland	1,300	47	22	Y.J./T.B.	10
	Melela Nzuri, Morogoro	so	Woodland	510	93	25	Y.J./T.B.	5
	Kwakachinja	so	Woodland	985	48	22	C.B.	3
Ethiopia	Bidree Village	d	Woodland	1,700	64	21	T.B.	4
	Kafa Biosphere Reserve	d	Forest, bamboo	1,600	130	19	K.S.	1
Uganda	Mt Elgon NP	se	Forest, grassland, bamboo	2,800	98	18	L.A.	2
	Agoro-Agu FR	se	Forest, woodland	1,400	99	23	Y.J./T.B.	6
	Otzi FR	se	Forest, woodland	1,290	90	26	Y.J./T.B.	3
Ghana	Kyabobo NP	se	Woodland, forest	180	108	26	S.B./J.T./R.P.	5
Gambia	Niumi NP	se	Woodland, grassland	2	73	28	S.B./M.S.	3
Senegal	Fathala WR	se	Woodland, grassland	35	68	28	I.S.	6
	Fongoli	se	Woodland	200	95	29	G.E.	2

NP, National Park; NR, Nature Reserve; Nat R, National Reserve; FR, Forest Reserve; WR, Wildlife Reserve. b, *G. s. braccatus*; d, *G. s. dummi*; se, *G. s. senegalensis*; so, *G. s. sotikae*. A.P., Andrew Perkin; C.B., Caroline Bettridge; C.D.S., C. D. Schaaf; G.E., Grace Ellison; I.S., Irena Schneiderová; J.T., Jessica Tombs; K.S., Karina Schell; L.A., Lesley Ambrose; M.S., Magdalena Svensson; P.H., Paul Honess; R.P., Roland Plesker; S.B., Simon Bearder; T.B., Thomas Butynski; Y.J., Yvonne de Jong. <sup>1</sup> Mean annual rainfall and mean annual temperature 1991–2015 ([www.worldbank.org/climateportal](http://www.worldbank.org/climateportal)).

## Acoustic Analyses

Honk calls were digitalised (if recorded by a cassette recorder) at a sample rate of 22.05 kHz and 16-bit size and analysed using Avisoft SASLab Pro software (R. Specht, Berlin; version 5.2). Each recording was converted into a spectrogram, ensuring that only high-quality recordings were included in the analyses. Spectrogram parameters were set as follows: FFT length 512; 50% overlap; hamming window, yielding time, and frequency measurement precision of 5.3 ms and 94 Hz. Acoustic parameters measured from spectrograms of honk calls (Fig. 3) were: (a) mean dominant frequency (measured in Hz); (b) mean fundamental frequency (first harmonic, measured in Hz); (c) mean unit length (basic element of a call that is represented as a continuous tracing along the temporal axis of the spectrogram, measured in seconds); and (d) mean interunit interval (time gap between 2 consecutive units along the temporal axis of a spectrogram, measured in seconds).



**Fig. 3.** Northern lesser galago (*Galago senegalensis*) honk call spectrogram showing the 4 call parameters included in this study: dominant frequency (**a**); fundamental frequency (**b**); unit length (**c**); interunit interval (**d**).

Intraindividual variation in unit length and interunit interval is considerably smaller than interindividual variation. More specifically, when measuring multiple unit lengths per call, the standard deviation was 7.78% of the mean value within individuals ( $n = 100$ ) but 19.62% among individuals ( $n = 121$ ). Similarly, when measuring multiple interunit intervals per call, the standard deviation is 9.57% of the mean value within individuals ( $n = 99$ ) but 21.52% among individuals ( $n = 121$ ). Based on this, and to avoid non-independence when including multiple measurements from single individuals, we applied, for

all 4 call parameters, the mean for individuals rather than the range. To help ensure that only 1 honk call bout per individual was analysed, we selected only 1 call per hour from the same location. While it is unlikely, we are, nevertheless, not absolutely certain that no individual provided more than 1 honk call bout to this study.

### *Statistical Analyses*

To test the isolation by distance hypothesis, 2 high-quality calls were selected from each of 21 sites (Fig. 1) and compared against one another. Only 1 high-quality call was available from the remaining 7 sites. This resulted in a total of 49 recordings. Geographic distances of the 28 sites were measured in ArcMAP 10.4 and rounded up to the nearest 5 km. Within-site calls were assumed to have been recorded at <5 km. For all 4 call parameters we measured differences among calls by subtracting the smallest value from the largest value and then divided that value by the largest difference measured (thus, expressing differences as a proportion of the maximum recorded difference). To approach a normal distribution, we log-transformed geographic distances and differences in call parameters. We calculated Pearson's correlation coefficient for the relationship between geographic distance and difference for the 4 call parameters separately and for their mean. Because data from each site were used in comparisons with all other sites, samples are not independent. To overcome this, we regrouped the 670 comparisons in 49 distance classes (i.e., equal to the original sample size), each containing 10–15 comparisons.

To test whether the observed variation can be explained by genetic basis (genetic basis hypothesis) or by separated geographic regions (isolation by barrier hypothesis), we applied multivariate analysis of variance. We compared the absolute measurements of the 4 call parameters using the geographic ranges of the 4 subspecies. In addition, call parameter measurements were taken of the 4 geographic regions. All useable calls ( $n = 121$ ) were included in this analysis. This was followed by post hoc testing (Tukey) to determine which differences among subspecies and geographic regions were statistically significant.

In testing subspecies differences and the effects of geographic regions, we applied a canonical discriminant function analysis (DFA), applying all usable calls ( $n = 121$ ). Four a priori recognised groups were used for testing subspecies differences and for testing the effects of geographic regions (groups as

defined above). For both DFA analyses we extracted the first 2 functions. These explained 65.9 and 22.2% of the variance for subspecies, and 77.3 and 21.7% of the variance for geographic regions. In all analyses we accepted significance when  $p$  was  $<0.05$  in a two-tailed test.

## RESULTS

### *Isolation by Distance Hypothesis*

Apart from unit length and dominant frequency, which are weakly but significantly correlated ( $R = 0.335$ ,  $n = 49$ ,  $p = 0.023$ ) (Table 3), the 4 call parameters are not correlated with each other (all Pearson's  $R < 0.208$ ,  $n = 49$ ,  $p > 0.165$ ). Variation in call parameters, expressed as standard deviation divided by the mean or the range, shows no relationship with geographic range size. Therefore, *G. s. senegalensis*, the subspecies with the largest geographic range, does not have greater honk call variation than *G. s. braccatus*, the subspecies with the smallest range.

There is no significant relationship between geographic distance (log-transformed) and the difference in interunit interval (log-transformed) ( $R = 0.208$ ,  $n = 49$ ,  $p = 0.152$ , Fig. 4c) or dominant frequency (log-transformed) ( $R = 0.119$ ,  $n = 49$ ,  $p = 0.413$ , Fig. 4d). There is a highly significant positive relationship between geographic distance (log-transformed) and the difference in fundamental frequency (log-transformed) (Pearson's  $R = 0.507$ ,  $n = 49$ ,  $p = 0.0002$ , Fig. 4a), unit length (log-transformed) ( $R = 0.297$ ,  $n = 49$ ,  $p = 0.0384$ , Fig. 4b), and the 4 parameters combined ( $R = 0.511$ ,  $n = 49$ ,  $p < 0.0002$ , Fig. 4e). The coefficient of determination is moderate to low ( $R^2 = 0.257$ ,  $R^2 = 0.088$ , and  $R^2 = 0.261$ , respectively). As such, geographic distance explains a moderate amount of the observed differences in fundamental frequency, unit length, and overall acoustic variation.

**Table 3.** Means and standard deviations for 5 honk call parameters for each of the 4 subspecies of northern lesser galago (*Galago senegalensis*). Ranges in parentheses.

Subspecies, <i>n</i>	Maximum duration, s	Dominant frequency, Hz	Fundamental frequency, Hz	Unit length, s	Interunit interval, s
<i>G. s. senegalensis</i> (29)	124	1,296±284 (670–1,790)	641±258 (300–1,215)	0.29±0.06 (0.20–0.42)	0.68±0.14 (0.46–0.89)
<i>G. s. braccatus</i> (35)	124	1,038±348 (350–1,630)	580±206 (320–1,060)	0.24±0.04 (0.18–0.33)	0.80±0.17 (0.51–1.17)
<i>G. s. sotikae</i> (52)	94	1,305±362 (410–1,980)	634±229 (320–1,140)	0.28±0.05 (0.18–0.41)	0.81±0.17 (0.45–1.24)
<i>G. s. dunni</i> (5)	28	1,186±396 (980–1,890)	424±134 (320–640)	0.21±0.07 (0.17–0.33)	0.72±0.12 (0.65–0.93)

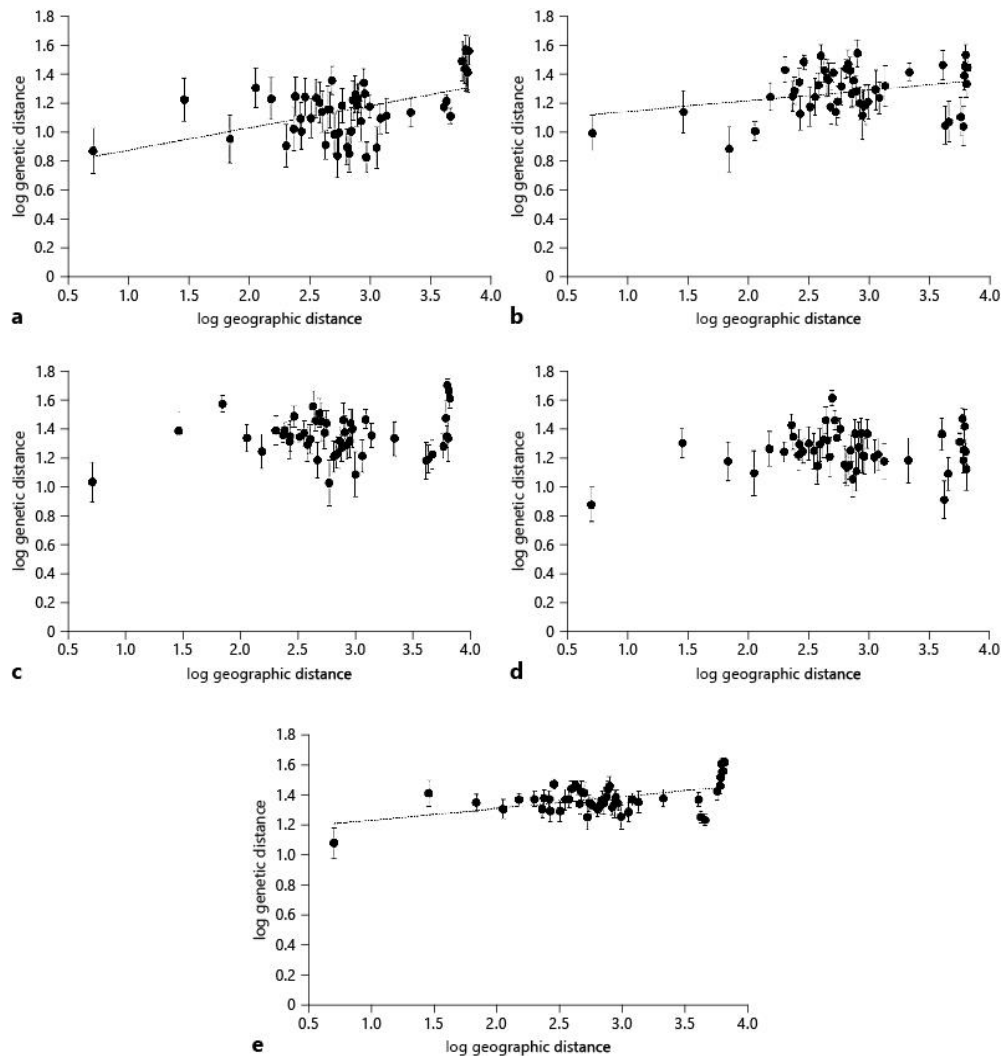


Fig. 4. Difference among 4 parameters of the honk call of individual northern lesser galagos (*Galago senegalensis*) in relation to geographic distance: fundamental frequency (a); unit length (b); interunit intervals (c); dominant frequency (d); combined (e). Values are means  $\pm$  SEM. Note that for a, b, and e, the relationship is statistically significant, but not for c and d. As such, no trendline is presented for c and d.

## Genetic Basis Hypothesis

There are significant acoustic differences among the honk calls of all 4 *G. senegalensis* subspecies ( $F_{12, 302} = 4.326$ ,  $p < 0.001$ ; Wilk's  $\lambda = 0.657$ , partial  $\eta^2 = 0.131$ ), in unit length ( $F_{3, 117} = 8.828$ ,  $p < 0.001$ ), interunit interval ( $F_{3, 117} = 3.993$ ,  $p = 0.010$ ), and dominant frequency ( $F_{3, 117} = 6.221$ ,  $p = 0.001$ ), but not in fundamental frequency (Table 3). A Tukey post hoc test indicates that unit length is significantly different between *G. s. senegalensis* and *G. s. braccatus* ( $p < 0.001$ ), between *G. s. senegalensis* and *G. s. dunni* ( $p = 0.007$ ), between *G. s. braccatus* and *G. s. sotikae* ( $p = 0.002$ ), and between *G. s. sotikae* and *G. s. dunni* ( $p = 0.022$ ). For interunit interval, *G. s. senegalensis* differs significantly from *G. s. braccatus* ( $p = 0.045$ ) and *G. s. sotikae* ( $p = 0.008$ ). Finally, for dominant frequency, *G. s. senegalensis* differs significantly from *G. s. braccatus* ( $p = 0.012$ ), and *G. s. braccatus* from *G. s. sotikae* ( $p < 0.001$ ; Table 3).

The DFA designated, on average, 52.1% of the individual honk calls to the correct subspecies, whereas given that there are 4 subspecies, the prior probability for each subspecies is 25%. Correct subspecies designation is highest for *G. s. braccatus* (65.7%) and lowest for *G. s. sotikae* (44.2%; Table 4). *G. alago s. braccatus* and *G. s. dunni* are separated from *G. s. sotikae* and *G. s. senegalensis* mostly on DFA function 1 (which correlates most strongly with unit length and dominant frequency). *Galago s. braccatus* and *G. s. sotikae* are separated from *G. s. dunni* and *G. s. senegalensis* mostly on DFA function 2 (which correlates most strongly with interunit length).

**Table 4.** Designation of individual northern lesser galago (*Galago senegalensis*) to each of the 4 subspecies and to each of 4 geographic regions based on canonical discriminant function analysis of the honk call. Percent in parentheses.

Subspecies	<i>senegalensis</i>	<i>braccatus</i>	<i>sotikae</i>	<i>dunni</i>	Total
<i>G. s. senegalensis</i>	14 (48.3)	9 (31.0)	4 (13.8)	2 (6.9)	29
<i>G. s. braccatus</i>	3 (8.6)	23 (65.7)	5 (14.3)	4 (11.4)	35
<i>G. s. sotikae</i>	12 (23.1)	10 (19.2)	23 (44.2)	7 (13.5)	52
<i>G. s. dunni</i>	1 (20.0)	1 (20.0)	0 (0.0)	3 (60.0)	5
Geographic region	West of Niger R.	Between Niger R. and Nile R.	West of Eastern Rift Valley	East of Eastern Rift Valley	Total
West of Niger R.	6 (40.0)	2 (13.3)	3 (20.0)	4 (26.7)	15
Between Niger R. and Nile R.	0 (0.0)	1 (33.3)	0 (0.0)	2 (66.7)	3
West of Eastern Rift Valley	2 (16.7)	2 (16.7)	5 (41.7)	3 (25.0)	12
East of Eastern Rift Valley	15 (16.5)	20 (22.0)	17 (18.7)	39 (42.9)	91

### *Isolation by Barrier Hypothesis*

There are significant differences in 2 of the 4 honk call parameters among geographic regions ( $F_{12, 302} = 2.246$ ,  $p = 0.010$ ; Wilk's  $\lambda = 0.798$ , partial  $\eta^2 = 0.73$ ): unit length ( $F_{3, 117} = 3.715$ ,  $p = 0.014$ ), and interunit interval ( $F_{3, 117} = 4.628$ ,  $p = 0.004$ ), but not in dominant frequency or fundamental frequency. A Tukey post hoc test revealed that unit length in populations east of the Eastern Rift Valley is significantly different from populations west of the Eastern Rift Valley ( $p = 0.014$ ). For interunit interval, *G. senegalensis* in the western-most part of the range (i.e., west of the Niger River) differs significantly from *G. senegalensis* in the eastern-most part of the range (i.e., east of the Eastern Rift Valley;  $p = 0.002$ ).

The DFA designated, on average, 42.1% of all honk calls to the correct geographic regions. Given that there are 4 regions, the probabilities for each is 25%. Correct subspecies designation is highest for individuals east of the Eastern Rift Valley (42.9%). Correct designation for individuals west of the Eastern Rift Valley (41.7%) is better than by chance (Table 4). Individuals in the western-most part of the species' range and those in the eastern-most part of the species' range can be separated on DFA function 1 (which correlates most strongly with interunit length).

## **DISCUSSION**

Comparison of acoustic features among and within taxa can provide a better understanding of their taxonomic arrangement, phylogenetic relationships, and ecological interactions [Hartmann et al., 2002]. This study is the first to describe and analyse intraspecies variation in the structure of the honk call of *G. senegalensis*, and its 4 subspecies, over its large geographic range. Three hypotheses were tested: isolation by distance hypothesis, genetic basis hypothesis, and isolation by barrier hypothesis. We found low to moderate levels of support for all three hypotheses.

The isolation by distance hypothesis predicts that subspecies with larger geographic ranges have more variation in their honk call than subspecies with small geographic ranges. Statistically significant support for the isolation by distance hypothesis was found for 2 of the 4 call parameters, as well as for all 4 parameters combined. Distance, however, explains only a moderate amount of the acoustic variation.

The subspecies with the largest geographic range, *G. s. senegalensis*, shows no greater honk call variation than does *G. s. braccatus*, the subspecies with the smallest geographic range.

The genetic basis hypothesis predicts that honk call variation corresponds to the 4 subspecies recognised for *G. senegalensis*. While the absolute differences among these taxa are small and unmarked, we did find statistically significant acoustic differences in honk calls among the 4 subspecies for 3 of the 4 call parameters. These differences are most pronounced between *G. s. senegalensis* and *G. s. dunni*. It appears that temporal parameters, and to a lesser extent frequency parameters, are responsible for subspecies-level differences. This is in agreement with earlier studies. For example, Zimmermann [1990] measured interunit intervals for captive *G. s. senegalensis* and *G. s. braccatus*, and in accordance with our results, found higher interunit intervals for *G. s. braccatus*. Species-level differences, on the other hand, are often strongly reflected in differences in fundamental frequency and its modulation, in addition to differences in overall temporal patterns [Zimmermann et al., 1988; Zimmermann, 1990].

The isolation by barrier hypothesis predicts greater honk call variation between regions divided by putative geographic barriers. Sample sizes available to test this hypothesis are, unfortunately, small and unequal, potentially limiting the explanatory power of our analysis. Significant differences are, however, found in 2 of the 4 call parameters among the 4 geographic regions. Unit length of honk calls east of the Eastern Rift Valley is significantly different from populations west of the Eastern Rift Valley. This suggests that the Eastern Rift Valley serves, or has served, as a barrier for *G. senegalensis*, as it has for other primate species [Butynski and De Jong, 2007]. Likewise, the Niger River seems to act, or has acted, as a significant barrier.

Due to considerable morphological variation within *G. senegalensis*, Masters and Bragg [2000] postulate that there may be unrecognised taxa within this widespread species. However, the acoustic structures of the honk calls from the 28 sites (7 countries) included in this study are not noticeably different. This suggests that all 121 individuals sampled are *G. senegalensis*. This is not to say that unrecognised taxa within *G. senegalensis* do not occur – as vast geographic regions remain unsampled



(e.g., the Sahel, the sub-Sahel, Ethiopia, Somalia, northern Kenya, Rwanda, Burundi, and western Tanzania; Fig. 1).

In addition to distance and isolation, divergence in acoustic signals among populations can be driven by various adaptive mechanisms. Variation in the acoustic structure of loud calls might be attributed to differences in morphology, including body size [Fitch and Hauser, 2003; Dávalos et al., 2018]. For example, small bats emit higher frequency calls than do larger bats [Jones, 1996]. In mammals, energy required for call production is generated in the lungs and calls are produced by vocal folds, subsequently modified in the vocal tract [Fitch and Hauser, 2003; Ey et al., 2007]. Larger mammals, therefore, can produce calls with lower fundamental and dominant frequencies than small mammals [Inoue, 1988; Fitch and Hauser, 1995]. This applies to some mammals, such as grasshopper mice (*Onychomys* spp.) [Hafner and Hafner, 1979], red deer (*Cervus elaphus*) [Reby et al., 2005], koalas (*Phascolarctos cinereus*) [Charlton et al., 2011], but not to others, such as ground squirrels (*Spermophilus* spp.) [Matrosova et al., 2007]. This study found that the lowest fundamental frequency and lowest dominant frequency occur in *G. s. dunnii*, the largest subspecies (mean head-body length of 188 mm,  $n = 5$ ) compared to the smaller *G. s. senegalensis* (mean head-body length 164 mm,  $n = 55$ ) and *G. s. braccatus* (mean head-body length 164 mm,  $n = 10$ ) [Nash et al., 2013]. It is, however, unlikely that honk call variation can be explained by the small differences in body size among these subspecies.

Apart from morphological cues, such as body size, differences in habitat structure drives natural selection for variation in mammal vocalizations, as in Gunnison's prairie dogs (*Cynomys gunnisoni*) [Perla and Slobodchikoff, 2002] and whistling rats (*Parotomys* spp.) [Le Roux et al., 2002]. Generally, taxa which prefer dense vegetation have vocalization with lower frequencies, narrower bandwidths, fewer frequency modulations, longer notes, and longer internote intervals [Morton, 1975]. Slower modulated elements are favoured in denser vegetation whereas longer calls with short, rapidly repeated elements perform better in open habitats [Wiley and Richards, 1978]. *Galago senegalensis* lives in a wide variety of habitats (woodland, bushland, riverine forest, moist forest) over a wide altitudinal range (0–2,800 m a.s.l.) [L. Ambrose, Y. de Jong, and T. Butynski, pers. observations]. Honk call diversity within *G.*

*senegalensis* might, therefore, be partly explained by habitat structure. Unfortunately, systematic data on habitat structure were not always collected when this study's honk call recordings were made.

Some diurnal primates, such as ring-tailed lemur (*Lemur catta*), pygmy marmoset (*Cebuella pygmaea*), Japanese macaque, and Geoffroy's spider monkey (*Ateles geoffroyi*), adapt the acoustic structure of their loud call depending on the physical distance of the potential receiver [Snowdon and Hodun, 1981; Oda, 1996; Sugiura, 2007; Ordóñez-Gómez et al., 2017]; calls have a higher frequency as distance increases. This might be a mechanism to avoid environmental background noise and to increase localizability [Snowdon and Hodun, 1981; Oda, 1996; Sugiura, 2007]. This study did not assess whether the frequency of the honk call of *G. senegalensis* is influenced by intragroup dispersal. It is likely, however, that nocturnal primates which occur in small, dispersed groups adapt their loud call frequencies in response to the often ample environmental background noises.

As indicated earlier, geographic limits of all 4 *G. senegalensis* subspecies are, in most cases, poorly understood (Fig. 1). For example, although Hill [1953] suggests that the Blue Nile River is the western limit for *G. s. dunni* and the eastern limit for *G. s. senegalensis*, this remains to be validated. Similarly, the southern limit for *G. s. dunni* and the northern limit of *G. s. braccatus* are uncertain. It appears that *G. s. sotikae* meets *G. s. senegalensis* in the vicinity of Lake Edward, Lake George, and Katonga River, and again off the northeastern corner of Lake Victoria, but this is based on limited data.

Many primate taxa in Africa occur along phenotypic clines [Mayr, 1956]. In eastern Africa, this includes baboons (*Papio* spp.), savanna monkeys (*Chlorocebus* spp.), patas monkeys (*Erythrocebus patas* spp.), and gentle monkeys (*Cercopithecus mitis* spp.), as well as *G. senegalensis* [Jolly, 1993; Frost et al., 2003; De Jong and Butynski, 2009; Butynski and De Jong, 2012; Cardini et al., 2012; Y. de Jong and T. Butynski, pers. observations]. In Kenya, *G. s. braccatus* occurs east of the Eastern Rift Valley and *G. s. sotikae* occurs west of the Eastern Rift Valley as well as within the Rift Valley (Fig. 1). Butynski and De Jong [2012] recognise an east-west cline within *G. senegalensis* that extends from at least the Laikipia Plateau in central Kenya (*G. s. braccatus*), across the Eastern Rift Valley at Lake Naivasha, to at least Lake Manyara in central northern Tanzania (*G. s. sotikae*). The more obvious phenotypical changes along

this cline are ear length and colour of the dorsum, outer hindlegs, and tail. Although this cline can be viewed in photographs available on the ‘Galago Photographic Map’ [De Jong and Butynski, 2018b], additional photographs from many more sites along this cline are required before the phenotypic changes can be described in detail. Although this cline calls into question the validity of *G. s. braccatus* and *G. s. sotikae*, this study found that honk call unit length and dominant frequency of individuals east of the Eastern Rift Valley are significantly different from those of individuals west of the Eastern Rift Valley. It may be that additional subspecies occur within *G. senegalensis*, and/or that more phenotypic clines will be detected. These should be sought as their presence and location will contribute to our understanding of the evolutionary history, biogeography, and taxonomy of *G. senegalensis*, and are likely to have important implications for the conservation of genetic, ecological, and behavioural diversity within this species.

The study of communication systems, and in particular vocalizations, in nocturnal primates, including night monkeys (*Aotus* spp.), tarsiers (Tarsiidae), lemurs (Lemuriformes), galagos, and lorises (Lorisinae), has revealed novel insights into the evolutionary history of these often cryptic animals. In some taxa, big differences in vocal behaviour and call types strongly suggest presence of multiple species. While variation in vocalizations is expected to be largest in wide-ranging species, *G. senegalensis*, the nocturnal primate with the largest geographic range, displays only minor differences in its calls. Other nocturnal species, when studied over large geographic scales, may exhibit similar uniformity or may markedly differ. Exploring this in more detail is an exciting and important avenue for further research.

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The authors have no ethical conflicts to disclose.

## **DISCLOSURE STATEMENT**

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## **AUTHOR CONTRIBUTIONS**

This project was conceived by M.S. Svensson and S.K. Bearder. M.S. Svensson, T.M. Butynski, Y.A. de Jong, S.K. Bearder, and I. Schneiderová collected field recordings of *G. senegalensis*. V. Nijman and M.S. Svensson analysed the data, and M.S. Svensson, T.M. Butynski, Y.A. de Jong, S.K. Bearder, I. Schneiderová, and V. Nijman wrote the paper. All authors approved the final manuscript.

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