

1 SI Methods

1.1 Chiropteran parturition data mining

Bat species that have been implicated in filovirus transmission either through RNA or antibody detection methods [1, 2] were systematically queried through Walker’s Bats of the World [3], Scopus, Web of Science, and Google Scholar. Generalized search terms including: “Birth*”; “Breed*”; “Reproduc*”; “Partur*” along with the Latin binomial and common species names. Bat species that had undergone taxonomic reassignment were queried under all names. Species that had been implicated, but did not reside in Africa, were queried at the genus level to identify representative sister species. Preliminary results suggested insufficient sample sizes to model parturition on a consistent per species, or per genus basis. Species were then aggregated into three taxonomic clusters: African fruit bats (family: *Pteropodidae*), molossid bats (family: *Molossidae*, and non-molossid microbats, according to Cumming and Bernard [4]. Publications were accessed by hand and the location and month of parturition events were extracted. Data points were restricted to those within mainland Africa. Studies that reported suckling neonates, lactating females, or attached pups were excluded as they did not provide explicate indication of the date, or location of parturition. Additional points for non-implicated species were added opportunistically as discovered and an African bat specialist (AM) was consulted to help fill holes in the dataset and to expand the data set to all mainland African bats. All gathered data was then compiled into a single, recirculating year for further analysis.

1.2 Covariate data

Covariate data was assembled for both the bat birthing models and EVD spillover models and is summarized in Table 2. Covariate layers were blocked into static variables and temporally dynamic variables depending on the data availability and hypotheses being tested. The static covariate layers used included those describing mammalian species biodiversity (both volant and non-volant) [5], land cover [6], and four of the 19 Bioclimatic variables representing 30 year average for precipitation seasonality, mean diurnal range, temperature seasonality, and temperature annual range [7]. Temporally dynamic covariate layers were used to describe mean monthly temperature, average monthly precipitation, potential evapo-transpiration [8] and an enhanced vegetative index [9].

Considering the vast size of Africa the study extent was restricted to locations in Africa receiving more than 500mL of precipitation annually as suggested by Schmidt *et al.* 2017 [10]. All covariate layers were cropped and masked to the precipitation layer, resolution was aggregated to 25km raster blocks and temporally dynamic covariates were normalized (mean zero and standard deviation of 1). All spatial handling was performed within R [11], using tools in the `sp` [12], `raster` [13], and `rgdal` [14] packages.

1.2.1 Mammalian diversity layers

Covariate layers describing non-volant mammalian, African fruit bat, molossid, and non-molossid microbats diversity were created using the IUCN red list terrestrial mammals distribution database [5]. Classification by taxonomic rank was performed by `taxize` package [15] and manually checked by hand (see `DiversityRasters.R`) and the number of overlapping distributions for each cell within the study area were counted to create the diversity raster layers.

1.2.2 Land cover

The global land cover database [6] was obtained for the year 2009 and custom aggregation was used to reduce the number of classes from 23 to five as described by Table 3.

1.2.3 Fragmentation index

A binary mask of the aggregated ‘forest’ class (as described above) from the Globcover 2009 dataset [6] was used to define forest patches using 4-connectivity. A fragmentation index was then calculated for each pixel by counting the number of unique patches within a 20x20 sliding window using tools in the `EBImage` package [16].

84 1.3 African bat birth pulse modeling

85 The birth pulses for each of the taxonomic clusters were modeled through a modified ecological
86 niche modeling (ENM) procedure devised for this analysis. Standard ENMs use occurrence (or
87 occurrence/absence) and covariate data to create regressive correlations between observed environ-
88 mental phenomena and observations. In this instance, we re-purposed these models to correlate
89 phenological environmental signals with bat birthing events within a temporally dynamic frame-
90 work.

91 Birthing events identified through the data mining were split by taxonomic grouping as de-
92 scribed above and observation/absence matrices were generated to a sliding two month window.
93 Records of a birth within the window were recorded as an occurrence within the 25x25km cell in
94 which it was observed. All locations for which there was a recorded birth for within the taxon
95 group, but no births were observed in those months were deemed “absence points”. The two month
96 sliding window was chosen to reflect the non-discrete nature of birthing within wild animals and
97 to help account for deficiencies in the source data. An additional ten, randomly generated, down-
98 weighted (0.001) points (five occurrence and five absence points) were added to all models. This
99 measure was taken to ensure the representation of all land cover categories in all models, and pro-
100 vide necessary pseudo-absence points in a small number of cases in which and insufficient number
101 of absence points existed, while having minimal impact on the model outcomes.

102 Temporal covariate data was selected using a wider six month sliding window centered on the
103 same two month window used for the occurrence/absences and included the two preceding, and two
104 following months. Due to the inherent correlation of the covariate layers (especially the temporally
105 dynamic covariate layers) and general non-independence of all selected environmental metrics, a
106 raster based principal components analysis (PCA) method was used to reduce the dimensionality
107 of all continuous predictors into six orthogonal principal components using the `rasterPCA` function
108 of the package `RStoobox` [17]. The categorical land cover variable was then added to the other
109 dimensionally reduced co-variates and occurrence/absence data were fed into the `biomod2` [18]
110 framework.

111 ENMs were procedurally generated for all taxonomic clusters and all pairs of consecutive months
112 where the number of combined observations was greater than 7. Ten iterations of the potential
113 eight preliminary models (generalized linear model, generalized boosted model, classification tree
114 analysis, artificial neural network, surface range envelope, flexible discriminate analysis, random
115 forest and maximum entropy) were computed using a standard 70:30 training:test ratio. Ensemble
116 models were then compiled based on the weighted mean of the receiver operating characteristic
117 (ROC score) to create a single consensus layer with the probability of event occurrence between 0
118 and 1.

119 1.4 qAIC

120 Models with and without bat associated terms were compared using a quasi-Akaike information
121 criteria (qAIC) scheme of the form:

$$\text{qAIC} = -2\log\text{Lik}/\hat{C} + 2k,$$

122 where \hat{C} is the overdispersion, and k represents the number of terms in the model.

123

124 2 Tables

Table 1: African bat birth pulse data mining results

<i>Species</i>	Annual Births	Locations	References
<i>Pteropodidae</i>			
<i>Casinycteris ophiodon</i>	1	1	[19]
<i>Eidolon helvum</i>	1	3	[20, 21, 22, 23]
<i>Epomophorus gambianus</i>	1-2	3	[24, 25, 26]
<i>Epomophorus labiatus</i>	2	1	[27]
<i>Epomophorus wahlbergi</i>	1-2	2	[28, 29]
<i>Epomopos buettikoferi</i>	2	2	[26, 30]
<i>Epomopos franqueti</i>	2	3	[31, 32, 27]
<i>Hypsignathus monstrosus</i>	2	3	[33, 34, 19]
<i>Micropteropus pusillus</i>	1-2	6	[35, 36, 37, 25, 38, 26]
<i>Myonycteris angolensis</i>	1	2	[24, 19]
<i>Myonycteris leptodon</i>	1-2	3	[21, 19]
<i>Myonycteris torquata</i>	2	1	[35]
<i>Nanonycteris veldkampi</i>	1	2	[19, 25]
<i>Rousettus aegyptiacus</i>	1-2	12	[19, 25, 39, 40, 41, 34, 36, 24, 42, 43]
<i>Molossidae</i>			
<i>Chaerephon pumilus</i>	2-3	4	[44, 45, 46, 47]
<i>Chaerephon pumilus</i>	1	1	[48]
<i>Mops condylurus</i>	2	5	[49, 45, 50, 46]
<i>Mops midas</i>	1	1	[51]
<i>Otomops harrisoni</i>	1	2	[46]
<i>Tadarida aegyptiaca</i>	1	1	[52]
<i>Tadarida fulminans</i>	2	1	[53]
Non- <i>Molossidae</i> Microbats			
<i>Cardioderma cor</i>	2	1	[54]
<i>Coleura afra</i>	2	1	[47]
<i>Eptesicus hottentotus</i>	1	1	[55]
<i>Hipposideros abae</i>	1	1	[37]
<i>Hipposideros beatus</i>	1	1	[56]
<i>Hipposideros caffer</i>	1	3	[57, 58, 59]
<i>Hipposideros caffer rubber</i>	1	2	[37, 19]
<i>Hipposideros cyclops</i>	1	1	[37]
<i>Hipposideros gigas</i>	1	2	[60, 61]
<i>Hipposideros rubber</i>	1	1	[62]
<i>Hipposideros vittatus</i>	1	1	[63]
<i>Lavia frons</i>	1	2	[37, 64]
<i>Miniopterus fraterculus</i>	1	1	[65]
<i>Miniopterus inflatus</i>	1	1	[61]
<i>Miniopterus minor</i>	1	1	[66]
<i>Miniopterus natalensis</i>	1	4	[67, 65, 68, 69]
<i>Myotis bocagei</i>	2	1	[56]
<i>Myotis tricolor</i>	1	1	[70]
<i>Neoromicia capensis</i>	1	1	[71]
<i>Neoromicia nanus</i>	1	4	[72, 73, 74, 50]
<i>Neoromicia somalicus</i>	1	1	[50]
<i>Neoromicia zuluensis</i>	1	1	[75]
<i>Nycteris grandis</i>	1	1	[76]
<i>Nycteris hispidar</i>	2	1	[37]
<i>Nycteris nana</i>	1	1	[37]
<i>Nycteris thebaica</i>	1	1	[57]
<i>Nycticeinops schlieffeni</i>	1	3	[77, 50, 75]
<i>Pipistellus rusticus</i>	1	1	[78]
Continued on next page			

Table 1 – continued from previous page

<i>Species</i>	Annual Births	Locations	References
<i>Rhinolophus basii</i>	1	1	[75]
<i>Rhinolophus capensis</i>	1	1	[79]
<i>Rhinolophus clivossis</i>	1	1	[79]
<i>Rhinolophus darlingi</i>	1	1	[80]
<i>Rhinolophus fumigatus</i>	1	1	[75]
<i>Rhinolophus landeri</i>	1	1	[59]
<i>Rhinolophus mossambicus</i>	1	1	[81]
<i>Rhinolophus simulator</i>	1	2	[81, 50]
<i>Scotoecus hirundo</i>	1	1	[50]
<i>Scrotophilus borbonicus</i>	1	1	[77]
<i>Scrotophilus dinganii</i>	1-2	3	[50, 72, 36]
<i>Scrotophilus leucigaster</i>	1	1	[82]
<i>Scrotophilus vividis</i>	1	1	[77]
<i>Taphozous hildegardeae</i>	2	1	[66]
<i>Taphozous mauritanus</i>	2	2	[75, 50]

Table 2: Covariate layers for modelling

Description	Analysis	Variable Type	References
Non-volant mammalian biodiversity	Both	Static	[5]
Land cover (modified)	Both	Static	[6]
Precipitation seasonality	ENM	Static	[7]
Mean diurnal range	ENM	Static	[7]
Temperature seasonality	ENM	Static	[7]
Temperature annual range	ENM	Static	[7]
Mean monthly temperature	ENM	Temporal	[7]
Mean monthly precipitation	ENM	Temporal	[7]
Potential evapo-transpiration	ENM	Temporal	[8]
Enhanced vegetative index	ENM	Temporal	[9]
Vegetation fragmentation index	spatGLM	Static	This study
Human population density	spatGLM	Static	[83]

Table 3: Reclassification of land cover variables

Value	Label	Class
11	Post-flooding or irrigated croplands (or aquatic)	crop
14	Rainfed croplands	crop
20	Mosaic cropland (50-70%) / vegetation (grassland/shrubland/forest) (20-50%)	crop
30	Mosaic vegetation (grassland/shrubland/forest) (50-70%) / cropland (20-50%)	grass
40	Closed to open (>15%) broadleaved evergreen or semi-deciduous forest (>5m)	forest
50	Closed (>40%) broadleaved deciduous forest (>5m)	forest
60	Open (15-40%) broadleaved deciduous forest/woodland (>5m)	forest
70	Closed (>40%) needleleaved evergreen forest (>5m)	forest
90	Open (15-40%) needleleaved deciduous or evergreen forest (>5m)	forest
100	Closed to open (>15%) mixed broadleaved and needleleaved forest (>5m)	forest
110	Mosaic forest or shrubland (50-70%) / grassland (20-50%)	forest
120	Mosaic grassland (50-70%) / forest or shrubland (20-50%)	grass
130	Closed to open (>15%) (broadleaved or needleleaved; evergreen or deciduous) shrubland (<5m)	grass
140	Closed to open (>15%) herbaceous vegetation (grassland; savannas or lichens/mosses)	grass
150	Sparse (<15%) vegetation	grass
160	Closed to open (>15%) broadleaved forest regularly flooded (semi-permanently or temporarily)	forest
170	Closed (>40%) broadleaved forest or shrubland permanently flooded - Saline or brackish water	forest
180	Closed to open (>15%) grassland or woody vegetation on regularly flooded or waterlogged soi	grass
190	Artificial surfaces and associated areas (Urban areas >50%)	urban
200	Bare areas	other
210	Water bodies	other
220	Permanent snow and ice	other
230	No data (burnt areas; clouds;...)	NA

Table 4: The abbreviations abr, mic, and mol, represent African fruit bats, non-molossid microbats and molossid bats respectively. The numbers included within the model names are the calendar months over which the model was produced. No models were created for January-February, and July-August for the non-molossid microbats.

Model	ROC	Sensitivity	Specificity
mic2.mic3	0.986	96.296	96.97
mic3.mic4	0.979	97.959	87.013
mic4.mic5	0.985	97.561	89.412
mic5.mic6	0.993	97.222	98.889
mic6.mic7	0.995	100	98.131
mic8.mic9	0.997	100	97.368
mic10.mic11	0.994	96.154	100
mic11.mic12	0.993	92.683	100
mic12.mic1	0.985	98.551	91.228
mol1.mol2	0.969	100	82.222
mol2.mol3	0.993	100	90.625
mol3.mol4	0.993	100	93.75
mol4.mol5	0.983	96.296	93.939
mol5.mol6	0.952	95.238	84.615
mol6.mol7	0.96	95.238	84.615
mol7.mol8	0.962	90.476	89.744
mol8.mol9	0.976	84.211	100
mol9.mol10	0.972	84.211	100
mol10.mol11	0.952	92.308	88.235
mol11.mol12	0.961	90.244	100
mol12.mol1	0.969	93.478	92.857
ptr1.ptr2	0.987	97.015	91.589
ptr2.ptr3	0.988	99.01	93.151
ptr3.ptr4	0.991	96.124	95.556
ptr4.ptr5	0.994	97.248	95.385
ptr5.ptr6	0.992	94.643	96.61
ptr6.ptr7	0.988	96.364	94.958
ptr7.ptr8	0.983	94.444	94.118
ptr8.ptr9	0.982	92.553	96.25
ptr9.ptr10	0.991	93.75	100
ptr10.ptr11	0.994	95.918	100
ptr11.ptr12	0.991	94	98.387
ptr12.ptr1	0.989	95.349	96.183

Table 5: Model for all alternative animal spatGLM models. Terms not included in that model are denoted by “-”

Model:	Null		D_{tot}		D_{tax}		P_{tot}		$D_{tot} + P_{tot}$		$D_{tot} \times P_{tot}$		$D_{tax} + P_{tax}$		$D_{tax} \times P_{tax}$	
	Estimate	P value	Estimate	P value	Estimate	P value	Estimate	P value	Estimate	P value	Estimate	P value	Estimate	P value	Estimate	P value
Intercept)	-32.81	<0.001	-30.02	<0.001	-27.56	<0.001	-38.2	0	-36.39	<0.001	-29.63	<0.001	-30.61	<0.001	-31.34	<0.001
P_{afb}	-	-	-	-	-	-	-	-	-	-	-	-	2	0.07	-	-
λ_{afb}	-	-	-	-	-	-	-	-	-	-	-	-	-	1.08	0.01	-
P_{mic}	-	-	-	-	-	-	-	-	-	-	-	-	-0.03	0.98	-	-
λ_{mic}	-	-	-	-	-	-	-	-	-	-	-	-	-	0.06	0.79	-
P_{mol}	-	-	-	-	-	-	-	-	-	-	-	-	-0.28	0.81	-	-
λ_{mol}	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-0.18	0.64
P_{total}	-	-	-	-	-	-	2.11	0.24	1.74	0.31	-	-	-	-	-	-
λ_{total}	-	-	-	-	-	-	-	-	-	-	0.23	0.67	-	-	-	-
P_{afb2}	-	-	-	-	-	-	-	-	-	-	-	-	1.27	0.25	-	-
λ_{afb2}	-	-	-	-	-	-	-	-	-	-	-	-	-	0.79	0.04	-
P_{mic2}	-	-	-	-	-	-	-	-	-	-	-	-	1.67	0.08	-	-
λ_{mic2}	-	-	-	-	-	-	-	-	-	-	-	-	-	0.2	0.3	-
P_{mol2}	-	-	-	-	-	-	-	-	-	-	-	-	0.75	0.56	-	-
λ_{mol2}	-	-	-	-	-	-	-	-	-	-	-	-	-	0.47	0.31	-
P_{total2}	-	-	-	-	-	-	6.17	0.02	5.65	0.02	-	-	-	-	-	-
λ_{total2}	-	-	-	-	-	-	-	-	-	-	1.51	0.03	-	-	-	-
P_{afb4}	-	-	-	-	-	-	-	-	-	-	-	-	-0.72	0.49	-	-
λ_{afb4}	-	-	-	-	-	-	-	-	-	-	-	-	-	0.46	0.19	-
P_{mic4}	-	-	-	-	-	-	-	-	-	-	-	-	-1.09	0.28	-	-
λ_{mic4}	-	-	-	-	-	-	-	-	-	-	-	-	-	-0.32	0.09	-
P_{mol4}	-	-	-	-	-	-	-	-	-	-	-	-	-0.69	0.55	-	-
λ_{mol4}	-	-	-	-	-	-	-	-	-	-	-	-	-	0.02	0.95	-
P_{total4}	-	-	-	-	-	-	0.96	0.54	0.57	0.71	-	-	-	-	-	-
λ_{total4}	-	-	-	-	-	-	-	-	-	-	0	1	-	-	-	-
P_{afb6}	-	-	-	-	-	-	-	-	-	-	-	-	0.9	0.42	-	-
λ_{afb6}	-	-	-	-	-	-	-	-	-	-	-	-	-	-	0.6	0.16
P_{mic6}	-	-	-	-	-	-	-	-	-	-	-	-	-1.69	0.09	-	-
λ_{mic6}	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	0.12
P_{mol6}	-	-	-	-	-	-	-	-	-	-	-	-	-1.15	0.31	-	-
λ_{mol6}	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-0.3	0.44
P_{total6}	-	-	-	-	-	-	2.45	0.17	2.08	0.24	-	-	-	-	-	-
λ_{total6}	-	-	-	-	-	-	-	-	-	-	0.46	0.4	-	-	-	-
D_{afb}	-	-	-	-	5.52	<0.001	-	-	-	-	-	-	5.93	<0.001	-	-
D_{mic}	-	-	-	-	-2	0.34	-	-	-	-	-	-	-1.54	0.33	-	-
D_{mol}	-	-	-	-	1.5	0.26	-	-	-	-	-	-	1.47	0.13	-	-
D_{bat}	-	-	2.21	0.01	-	-	-	-	0.91	0.36	-	-	-	-	-	-
PopDen	-1.7	<0.001	-1.79	<0.001	-2.03	<0.001	-1.84	<0.001	-1.84	<0.001	-1.84	<0.001	-2.01	<0.001	-1.87	<0.001
fragIndex	0.54	0.15	0.53	0.15	0.03	0.97	0.44	0.34	0.44	0.31	0.46	0.25	0.1	0.84	0.32	0.48
BVD	4.96	<0.001	4.73	<0.001	4.27	<0.001	4.62	<0.001	4.57	<0.001	4.55	<0.001	4.06	<0.001	4.13	<0.001
D_{MNB}	6.34	<0.001	3.97	<0.001	3.74	0.09	6.08	<0.001	5.16	<0.001	4.05	<0.001	3.67	0.04	5.36	<0.001
qAIC	325.99		271.56		481.71		358.26		320.38		272.43		280.96		340.56	
\hat{c}	0.82		0.69		1.32		0.92		0.81		0.69		0.71		0.87	

Table 6: Model for all alternative animal spatGLM models with modified African birth pulse terms. Terms not included in that model are denoted by “-”

Model:	P_{tot}^*		$D_{tot} + P_{tot}^*$		$D_{tot} \times P_{tot}^*$		$D_{tox} + P_{tox}^*$		$D_{tox} \times P_{tox}^*$	
	Estimate	P value	Estimate	P value	Estimate	P value	Estimate	P value	Estimate	P value
Intercept)	-25.46	<0.001	-27.81	<0.001	-31.44	<0.001	-29.16	<0.001	-33.39	<0.001
P_{afb}^*	-	-	-	-	-	-	2.2	0.05	-	-
λ_{afb}^*	-	-	-	-	-	-	-0.13	0.89	2.32	0.06
P_{mic}^*	-	-	-	-	-	-	-	-	-	1
λ_{mic}^*	-	-	-	-	-	-	-0.79	0.5	-	-
P_{mol}^*	-	-	-	-	-	-	-	-	0.13	0.78
λ_{mol}^*	-	-	-	-	-	-	-	-	-	-
P_{total}^*	2.41	0.42	3.78	0.17	-	-	-	-	-	-
λ_{total}^*	-	-	-	-	2.8	0.04	-	-	-	-
P_{afb}^{*2}	-	-	-	-	-	-	0.99	0.38	-	-
λ_{afb}^{*2}	-	-	-	-	-	-	-	-	1.7	0.17
P_{mic}^{*2}	-	-	-	-	-	-	1.58	0.12	-	-
λ_{mic}^{*2}	-	-	-	-	-	-	-	-	0.19	0.41
P_{mol}^{*2}	-	-	-	-	-	-	0.34	0.8	-	-
λ_{mol}^{*2}	-	-	-	-	-	-	-	-	0.49	0.36
P_{total}^{*2}	-6.27	0.13	-5.04	0.18	-	-	-	-	-	-
λ_{total}^{*2}	-	-	-	-	-1.6	0.32	-	-	-	-
P_{afb}^{*4}	-	-	-	-	-	-	-1.57	0.2	-	-
λ_{afb}^{*4}	-	-	-	-	-	-	-	-	-0.79	0.51
P_{mic}^{*4}	-	-	-	-	-	-	-0.94	0.38	-	-
λ_{mic}^{*4}	-	-	-	-	-	-	-	-	-0.41	0.07
P_{mol}^{*4}	-	-	-	-	-	-	-0.59	0.63	-	-
λ_{mol}^{*4}	-	-	-	-	-	-	-	-	-0.12	0.81
P_{total}^{*4}	-6.03	0.15	-4.81	0.2	-	-	-	-	-	-
λ_{total}^{*4}	-	-	-	-	-1.55	0.35	-	-	-	-
P_{afb}^{*6}	-	-	-	-	-	-	0.78	0.51	-	-
λ_{afb}^{*6}	-	-	-	-	-	-	-	-	1.44	0.28
P_{mic}^{*6}	-	-	-	-	-	-	-1.38	0.18	-	-
λ_{mic}^{*6}	-	-	-	-	-	-	-	-	-0.3	0.16
P_{mol}^{*6}	-	-	-	-	-	-	-1.07	0.36	-	-
λ_{mol}^{*6}	-	-	-	-	-	-	-	-	-0.25	0.58
P_{total}^{*6}	2.18	0.48	3.49	0.22	-	-	-	-	-	-
λ_{total}^{*6}	-	-	-	-	2.68	0.05	-	-	-	-
D_{afb}	-	-	-	-	-	-	5.97	<0.001	-	-
D_{mic}	-	-	-	-	-	-	-1.65	0.31	-	-
D_{mol}	-	-	-	-	-	-	1.6	0.1	-	-
D_{bat}	-	-	2.21	0.01	-	-	-	-	-	-
PopDen	-1.72	<0.001	-1.79	<0.001	-1.78	<0.001	-2.01	<0.001	-1.91	<0.001
fragIndex	0.54	0.15	0.54	0.13	0.54	0.11	0.07	0.88	0.13	0.82
BVD	4.93	<0.001	4.75	<0.001	4.79	<0.001	3.96	<0.001	3.8	<0.001
D_{MNB}	5.97	<0.001	3.89	<0.001	4.1	<0.001	3.54	0.05	4.26	0.02
qAIC	315.64		253.83		252.21		295.64		390.43	
\hat{c}	0.79		0.63		0.63		0.75		1.05	

Table 7: Model for all alternative human spatGLM models. Terms not included in that model are denoted by “-”

Model:	Null	D_{tot}		D_{tax}		B_{tot}		$D + B_{tot}$		$D \times B_{tot}$		$D + B_{tax}$		$D \times B_{tax}$		
		Estimate	P value	Estimate	P value	Estimate	P value	Estimate	P value	Estimate	P value	Estimate	P value	Estimate	P value	
Intercept)	-29.82	<0.001	-29.03	<0.001	-29.66	<0.001	-28.73	<0.001	-32.76	<0.001	-28.73	<0.001	-31.62	<0.001	-28.62	<0.001
P_{afb}	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
λ_{afb}	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
P_{mic}	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
λ_{mic}	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
P_{mol}	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
λ_{mol}	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
P_{total}	-	-	-	-	-	-	0.79	0.22	1.88	0.27	0.79	0.22	-	-	-	-
λ_{total}	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
P_{afb2}	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
λ_{afb2}	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
P_{mic2}	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
λ_{mic2}	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
P_{mol2}	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
λ_{mol2}	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
P_{total2}	-	-	-	-	-	-	1.94	0.01	4.5	0.04	1.94	0.01	-	-	-	-
λ_{total2}	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
P_{afb4}	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
λ_{afb4}	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
P_{mic4}	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
λ_{mic4}	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
P_{mol4}	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
λ_{mol4}	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
P_{total4}	-	-	-	-	-	-	-0.13	0.78	-0.5	0.71	-0.13	0.78	-	-	-	-
λ_{total4}	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
P_{afb6}	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
λ_{afb6}	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
P_{mic6}	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
λ_{mic6}	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
P_{mol6}	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
λ_{mol6}	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
P_{total6}	-	-	-	-	-	-	0.09	0.87	0.29	0.85	0.09	0.87	-	-	-	-
λ_{total6}	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
D_{afb}	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
D_{mic}	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
D_{mol}	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
D_{bat}	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
PopDen	-0.23	0.24	-0.29	0.05	-0.25	0.07	-0.28	0.04	-0.28	0.05	-0.28	0.04	-0.23	0.08	-0.27	0.08
fragIndex	-0.21	0.54	-0.26	0.34	-0.38	0.17	-0.31	0.25	-0.3	0.26	-0.31	0.25	-0.36	0.22	-0.36	0.22
BVD	-11.04	0.99	-11.29	0.98	-11.21	0.98	-10.94	0.98	-11.01	0.98	-10.94	0.98	-11.43	0.98	-11.37	0.98
D_{MNB}	5.61	<0.001	2.88	0.02	2.92	0.01	3.37	<0.001	3.36	0.01	3.37	<0.001	3.36	<0.001	4.72	<0.001
OB_{an}	5.17	<0.001	4.96	<0.001	4.47	<0.001	4.74	<0.001	4.74	<0.001	4.74	<0.001	4.5	<0.001	4.73	<0.001
OB_{panj-1}	2.02	0.28	1.92	0.17	2.03	0.1	1.91	0.14	1.93	0.14	1.91	0.14	2.71	0.02	2.62	0.06
qAIC	1076.48	613.37	512.88	533.35	677.95	522.57	533.35	630.23	457.58	630.23	457.58	630.23	457.58	630.23	457.58	630.23
\hat{c}	1.87	1.08	0.91	0.93	1.18	0.92	0.93	1.10	0.79	1.10	0.79	1.10	0.79	1.10	0.79	1.10

Table 8: Model for all alternative human spatGLM models with modified African birth terms. Terms not included in that model are denoted by “-”

Model:	P_{tot}^*		$D_{tot} + P_{tot}^*$		$D_{tot} \times P_{tot}^*$		$D_{tax} + P_{tax}^*$		$D_{tax} \times P_{tax}^*$	
	Estimate	P value	Estimate	P value	Estimate	P value	Estimate	P value	Estimate	P value
Intercept)	-3.81	<0.001	-9.4	0.29	-29.74	<0.001	-30.63	<0.001	-29.93	<0.001
P_{afb}^*	-	-	-	-	-	-	-2.41	0.05	-	-
λ_{afb}^*	-	-	-	-	-	-	1.83	0.89	-1.83	0.06
P_{mic}	-	-	-	-	-	-	1.83	0.89	-	-
λ_{mic}	-	-	-	-	-	-	-0.19	0.5	0.26	1
P_{mol}	-	-	-	-	-	-	-	-	-	-
λ_{mol}	-	-	-	-	-	-	-	-	-0.15	0.78
P_{total}^*	-17.59	0.42	-15.29	0.01	-	-	-	-	-	-
λ_{total}^*	-	-	-	-	-2.23	0.21	-	-	-	-
P_{afb2}^*	-	-	-	-	-	-	0.91	0.38	-	-
λ_{afb2}^*	-	-	-	-	-	-	-0.06	0.12	1.89	0.17
P_{mic2}	-	-	-	-	-	-	-	-	-	-
λ_{mic2}	-	-	-	-	-	-	0.29	0.8	-0.18	0.41
P_{mol2}	-	-	-	-	-	-	-	-	-	-
λ_{mol2}	-	-	-	-	-	-	-	-	-0.09	0.36
$P_{total*2}$	-0.29	0.13	0.71	0.84	-	-	-	-	-	-
$\lambda_{total*2}$	-	-	-	-	2.65	0.09	-	-	-	-
P_{afb*4}	-	-	-	-	-	-	-0.39	0.2	-	-
λ_{afb*4}	-	-	-	-	-	-	-0.78	0.38	0.53	0.51
P_{mic4}	-	-	-	-	-	-	-	-	-	-
λ_{mic4}	-	-	-	-	-	-	0.2	0.63	-0.24	0.07
P_{mol4}	-	-	-	-	-	-	-	-	-	-
λ_{mol4}	-	-	-	-	-	-	-	-	0.25	0.81
$P_{total*4}$	-2.46	0.15	-0.91	0.8	-	-	-	-	-	-
$\lambda_{total*4}$	-	-	-	-	2.14	0.17	-	-	-	-
P_{afb*6}	-	-	-	-	-	-	1.6	0.51	-	-
λ_{afb*6}	-	-	-	-	-	-	-0.27	0.18	2.34	0.28
P_{mic6}	-	-	-	-	-	-	-	-	-	-
λ_{mic6}	-	-	-	-	-	-	1.34	0.36	-0.04	0.16
P_{mol6}	-	-	-	-	-	-	-	-	-	-
λ_{mol6}	-	-	-	-	-	-	-	-	0.44	0.58
$P_{total*6}$	-9.41	0.48	-7.28	0.09	-	-	-	-	-	-
$\lambda_{total*6}$	-	-	-	-	0.07	0.96	-	-	-	-
D_{afb}	-	-	-	-	-	-	2.87	<0.001	-	-
D_{mic}	-	-	-	-	-	-	1.86	0.19	-	-
D_{mol}	-	-	-	-	-	-	-0.59	0.46	-	-
D_{bat}	-	-	2.1	0.04	-	-	-	-	-	-
PopDen	-0.21	0.2	-0.25	0.09	-0.29	0.07	-2.01	0.07	-0.25	0.08
fragIndex	-0.25	0.43	-0.28	0.33	-0.23	0.41	0.07	0.12	-0.38	0.17
BVD	-11.93	0.98	-11.88	0.98	-11.22	0.98	3.96	0.98	-11.53	0.98
D _{MNB}	4.73	<0.001	3.15	0.01	3.42	0.01	3.54	0.01	4.09	<0.001
OB _{an}	5.14	<0.001	5	<0.001	5.09	<0.001	4.56	<0.001	4.58	<0.001
OB _{ant-1}	2.25	0.14	2.19	0.1	2.1	0.15	2.89	0.01	2.89	0.02
qAIC	757.58		595.88		664.63		454.93		542.84	
\hat{c}	1.35		1.06		1.16		0.78		0.96	

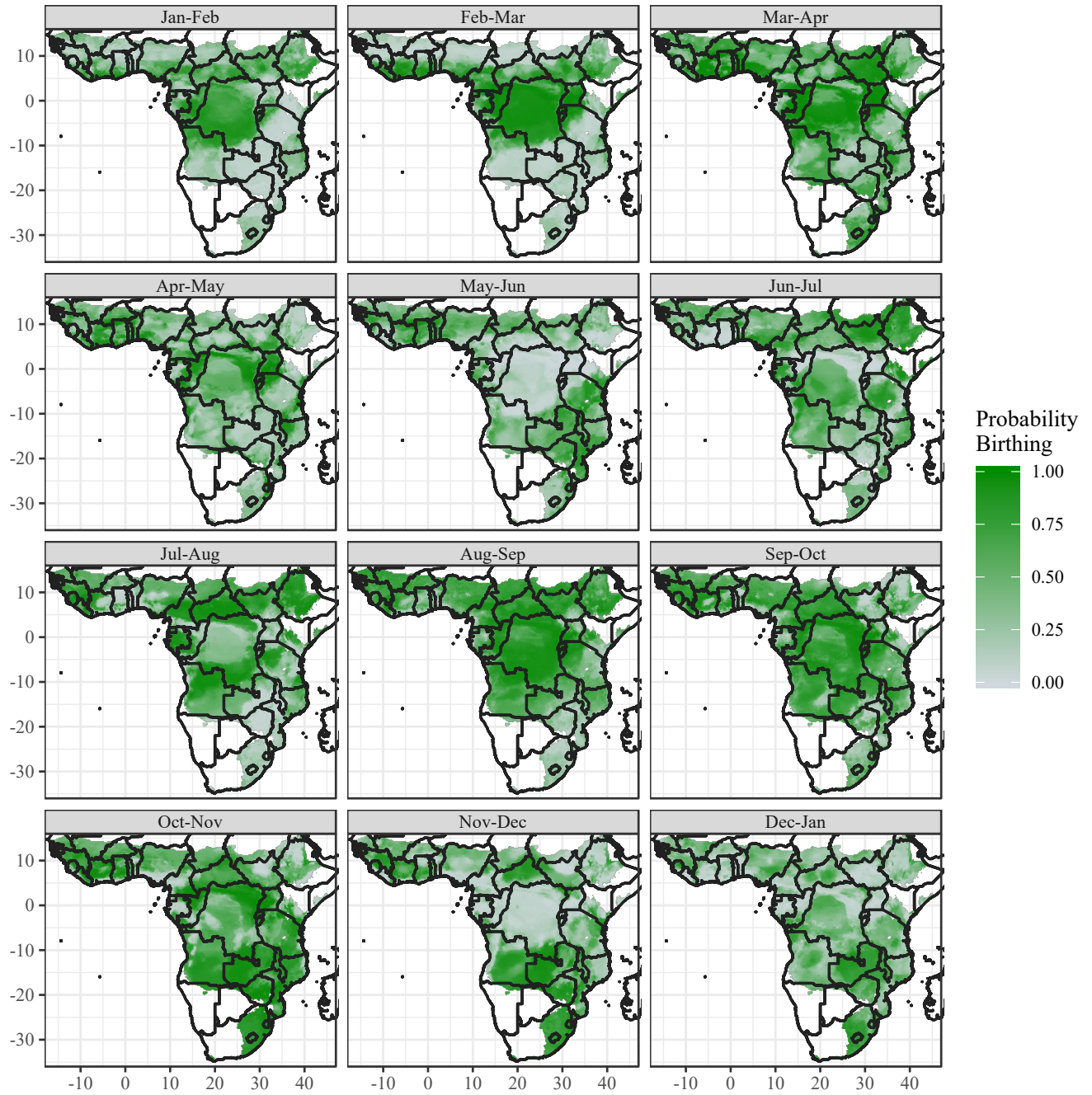


Figure 1: Monthly probability of birthing for African fruit bats

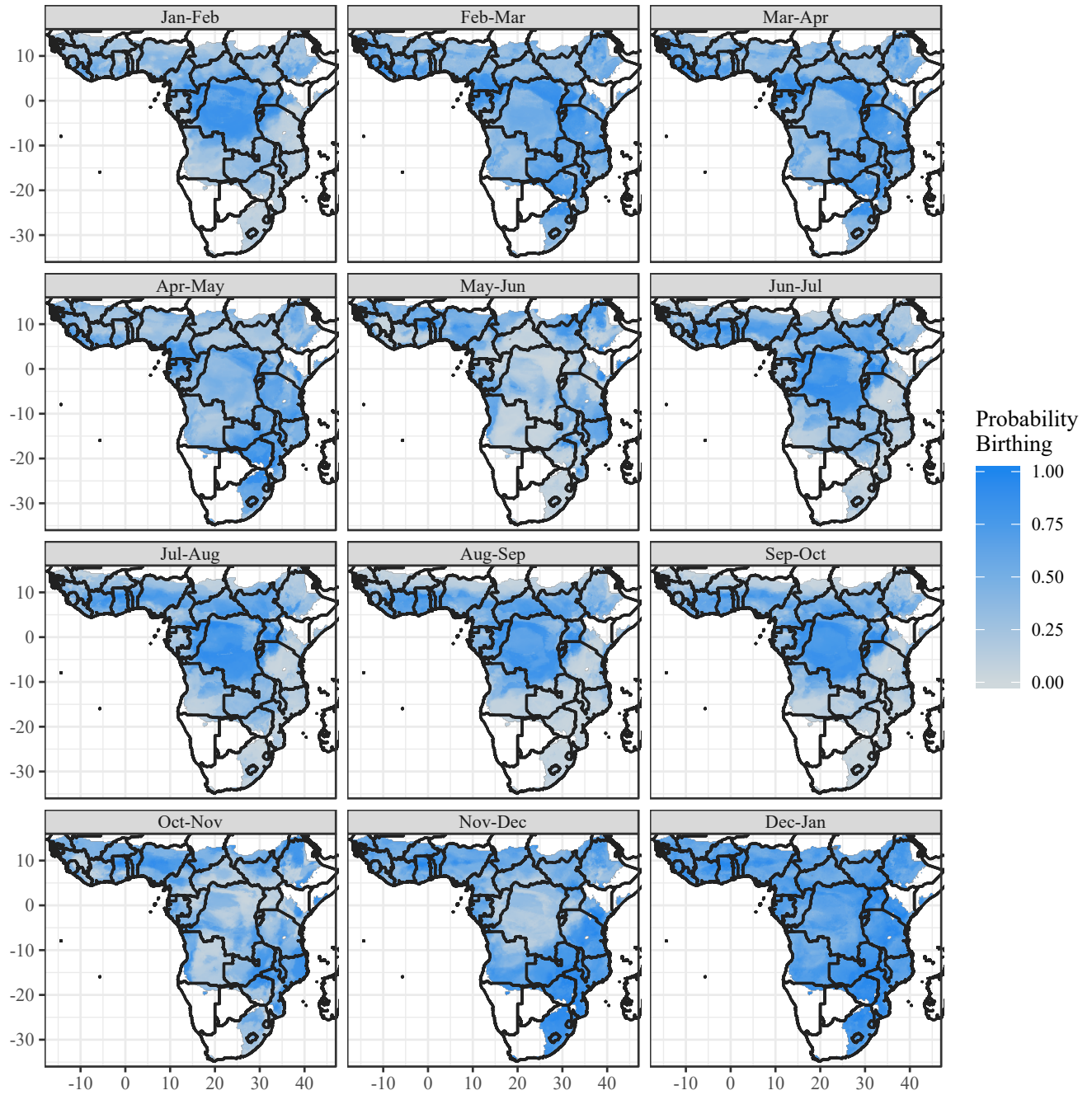


Figure 2: Monthly probability of birthing for molossid bats

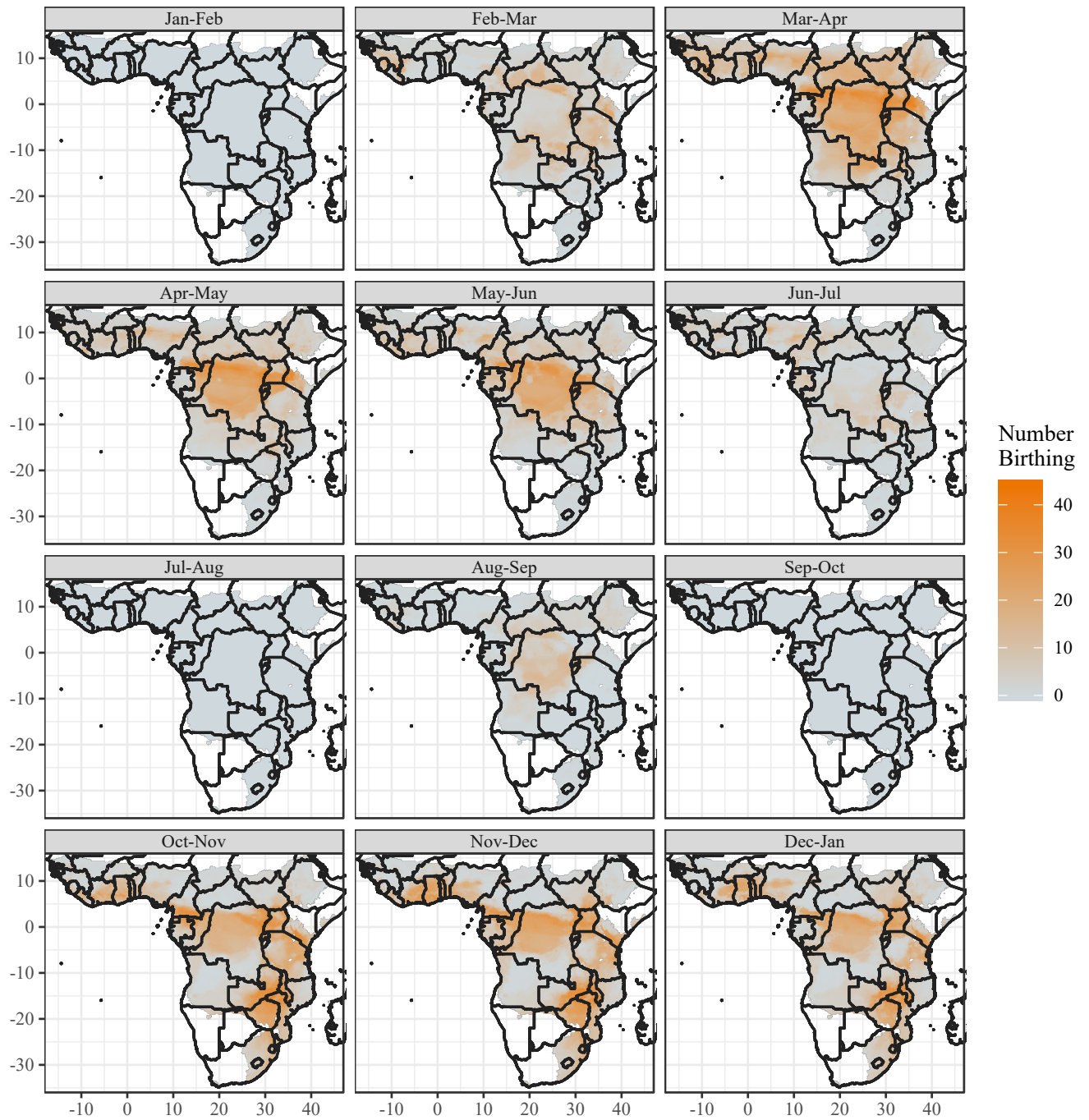


Figure 3: Monthly probability of birthing for non-molossid microbats

No models were generated for January-February or July-August.

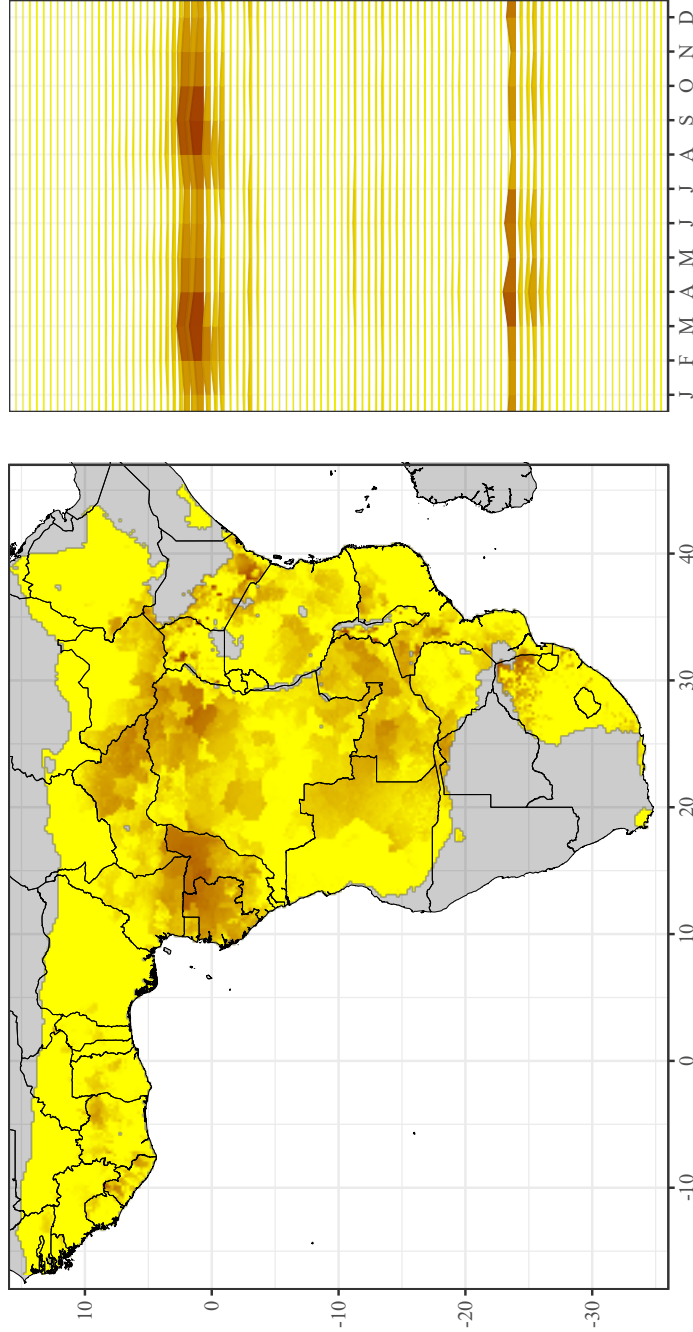


Figure 4: *Left:* Mean risk of EVD spillover to non-human reservoir hosts. *Right:* Risk of EVD spillover aggregated by latitude and month.

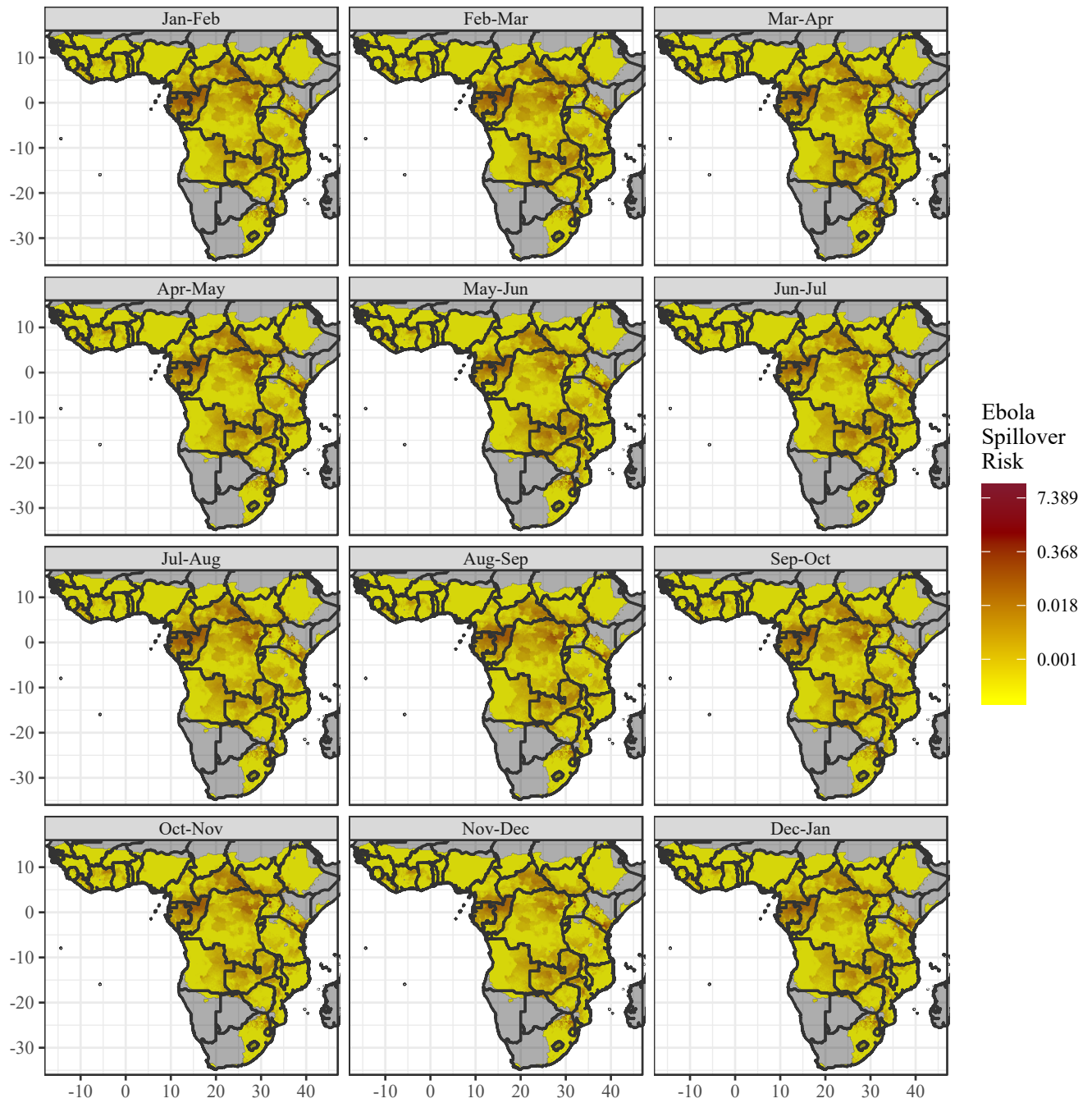


Figure 5: Monthly risk of ebolavirus spillover to non-human spillover hosts

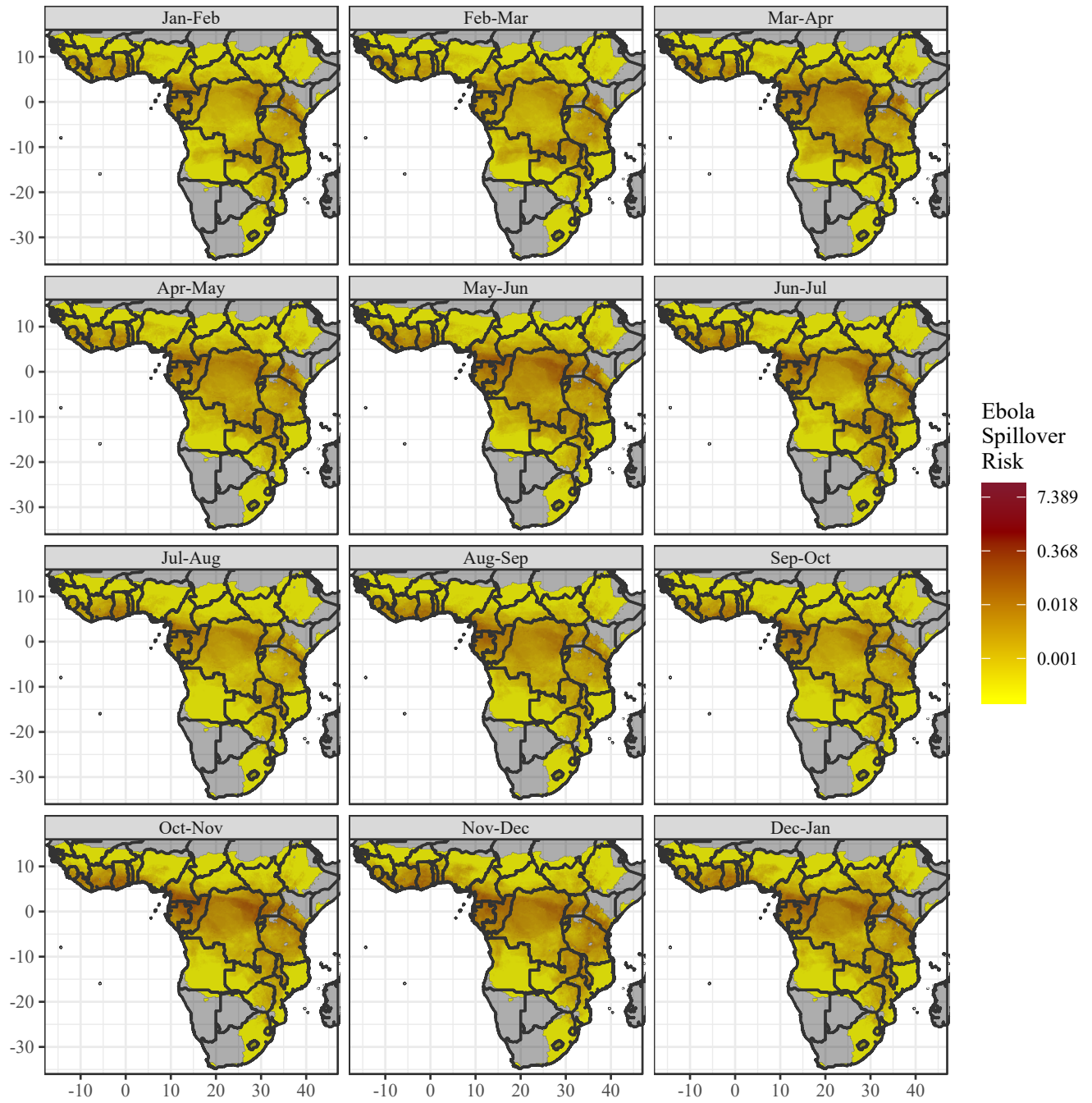


Figure 6: Monthly risk ebolavirus spillover into humans

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