

Supplementary Material

Functional trait filtering of aerial insectivorous forest bats in Amazonian cattle-dominated landscapes

This supplementary material contains:

- **Figure A1** Estimates of wing morphology values
- **Figure A2** Environmental variables values
- **Figure A3** Ordination plots
- **Table A1** Trait values
- **Table A2** Model output tables
- Landscape variable calculations

Figure A1. Estimating wing morphology values

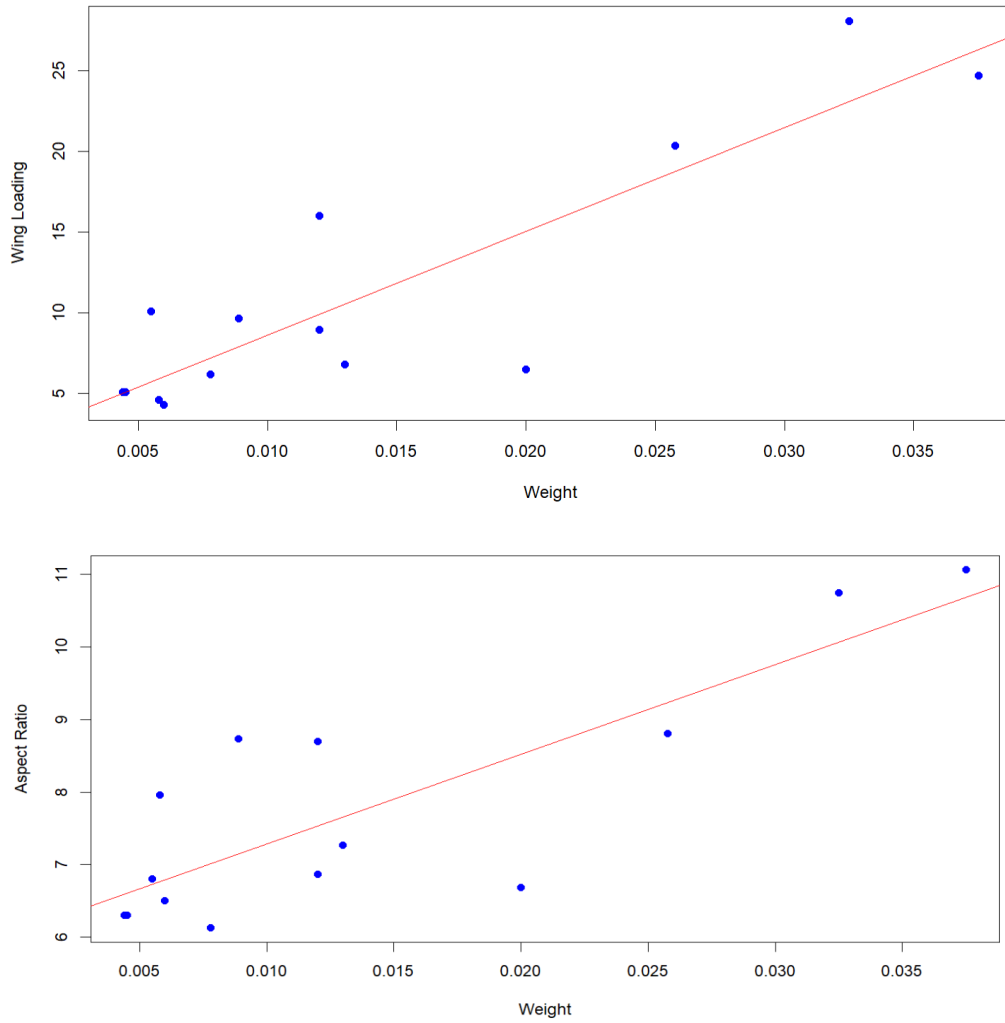


Figure A1. Linear regression demonstrating the relationship between body mass and (a) relative wing loading (rWL) and (b) aspect ratio (AR) of the AIBs included in the analysis. Equations of the regression were used to estimate values of rWL and AR for *Noctilio albus*, *Saccopteryx canescens*, and *Peropteryx trinitatis*. $rWL = 642.732 * Weight + 2.187$, adjusted $R^2 = 0.7558$; $AR = 123.4593 * Weight + 6.0485$, adjusted $R^2 = 0.6503$.

Figure A2. Environmental variables values

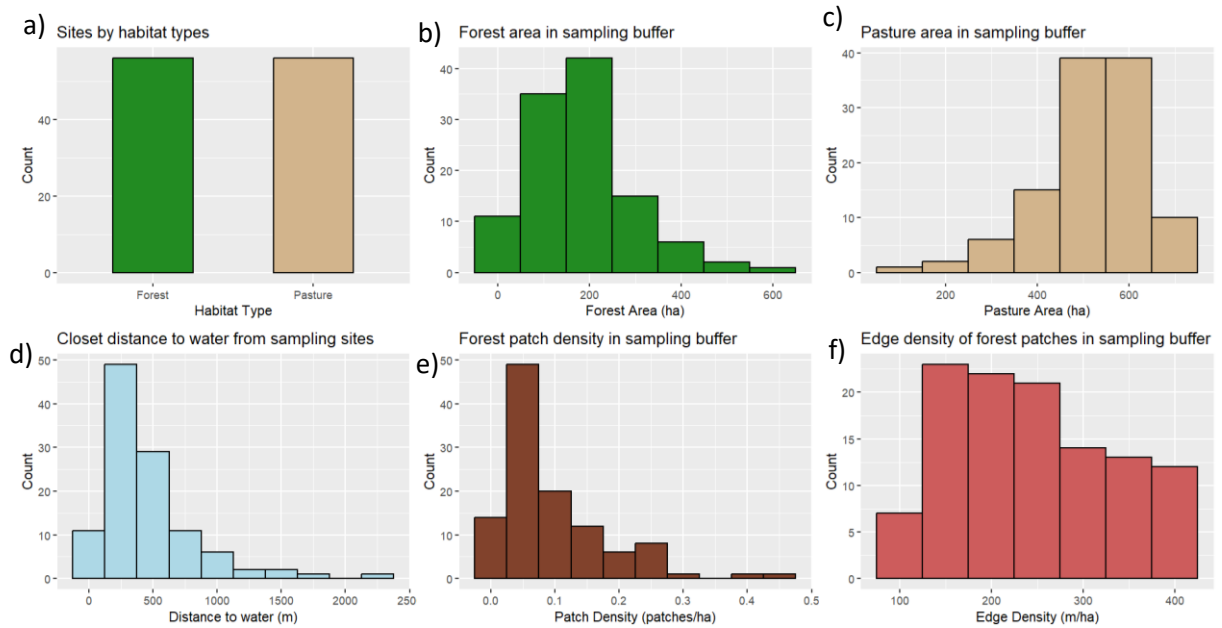


Figure A2. Values of the six environmental variables included in the analysis (a) Habitat type (either forest or pasture), (b) Forest area in the sampling buffer (ha), (c) Pasture area in the sampling buffer (ha), (d) The closest distance to water from each of the sampling sites, (e) the patch density of forest in the sampling buffer, and (f) the edge density of forest patches in the sampling buffer

Figure A3. Ordination plots

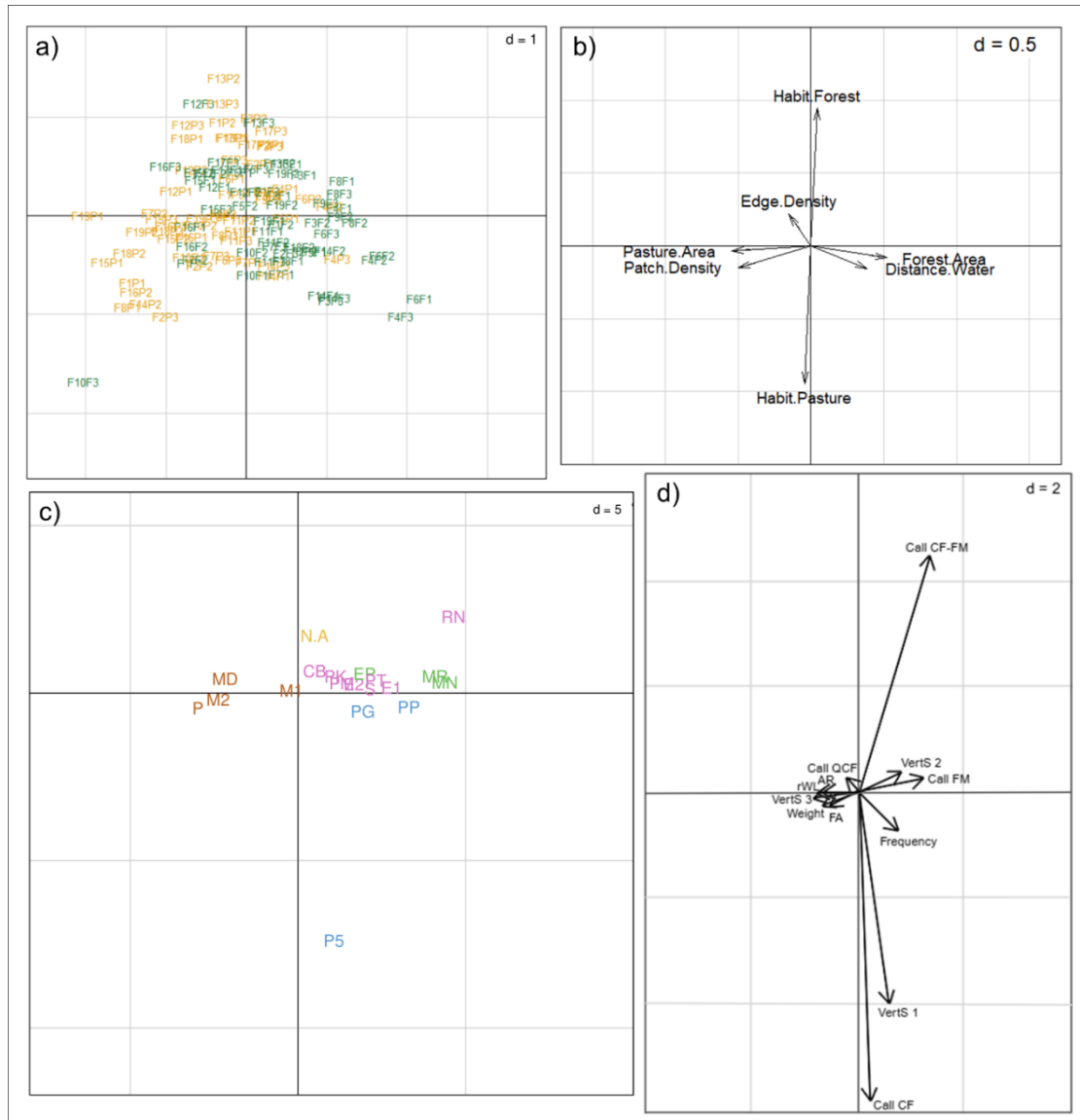


Figure A3: The first two axes from the RLQ analysis showing the distribution in ordination space of (a) Sites: Yellow = Pasture, Green = Forest (b) Environmental variables (c) Sonotypes: Orange = Molossidae, Purple = Emballonuridae, Green = Vespertilionidae, Yellow = Noctilionidae, Blue = Mormoopidae (See Table 1 for sonotype abbreviations) and (d) Traits (see Table 2 for trait abbreviations).

Table A1. Trait values

Table A1. Values of the functional traits for each of the 19 species/sonotypes included in the analysis. Relative wing loading = Wing loading (adult mass multiplied by gravitational acceleration divided by wingspan) divided by mass raised to a third: $rWL = (Mg/S)/mass^{1/3}$; Aspect ratio = The square of the adult wingspan, divided by the area of the wing: $AR = B^2/S$.

SPS	Species	Frequency	Call_structure	Weight	AR	WL	VertS	FA_smallest*
E2	<i>Centronycteris maximiliani/centralis</i>	40	QCF	5.8	7.9575	4.61	2	41.5
CB	<i>Cormura brevirostris</i>	28	QCF	8.9	8.73	9.66	2	41
EP	<i>Eptesicus brasiliensis</i>	32	FM	12	6.8648	8.94	2	40
M1	<i>Molossus molossus</i>	38.25	QCF	12	8.7	16	3	37
MN	<i>Myotis nigricans</i>	60	FM	4.5	6.3	5.1	2	30
MR	<i>Myotis riparius</i>	50	FM	4.4	6.3	5.1	2	31
PK	<i>Peropteryx kappleri</i>	31	QCF	8.1	7.3268	6.66	2	43
PM	<i>Peropteryx macrotis</i>	38	QCF	5.5	6.8	10.1	2	43
P	<i>Promops</i> sp.	32.25	QCF	37.5	11.0626	24.7	3	48
PG	<i>Pteronotus gymnonotus</i>	55	CF-FM	13	6.4883	6.46	1	50
PG	<i>Pteronotus gymnonotus</i>	55	CF-FM	13	8.0428	7.1	1	50
P5	<i>Pteronotus parnellii</i> 55	55	CF	20	6.68	6.48	1	50
PP	<i>Pteronotus personatus</i>	68.5	CF-FM	7.5	6.7	5.97	1	40
S	<i>Saccopteryx bilineata</i>	43.5	QCF	9.3	4.5373	4.62	2	40.5
S	<i>Saccopteryx bilineata</i>	43.5	QCF	9.3	8.1719	7.66	2	40.5
S	<i>Saccopteryx leptura</i>	51.5	QCF	4.8	5.6774	6.25	2	40.5
MD	<i>Eumops hansae</i>	20	QCF	16.5	8.6961	22.49	3	44.2
MD	<i>Eumops auripendulus</i>	20	QCF	32	10.6799	19.89	3	44.2
MD	<i>Eumops maurus</i>	20	QCF	22	8.4748	13.13	3	44.2
MD	<i>Cynomops</i> sp	20	QCF	32.55	7.3659	25.96	3	44.2
M2	M2	32.5	QCF	32.5	10.0647	22.79	3	41.5
M2	M2	32.5	QCF	32.5	11.0626	38.01	3	41.5
M2	<i>Molossus ater</i>	32.5	QCF	32.5	11.1	23.4	3	41.5
RN	<i>Rhynchonycteris naso</i>	98.88	CF-FM	6	6.5	4.3	2	35
N.A	<i>Noctilio albus</i>	73.06	CF-FM	25.7	9.22	18.71	2	54
E1	<i>Saccopteryx canescens</i>	54	QCF	5	6.67	5.4	2	35
PT	<i>Peropteryx trinitatis</i>	43.63	QCF	6	6.79	6.04	2	36

* Lowest values from the range of FA values were used from Lopez-Baucells et al (2016)

Table A2. Model output tables

Rao's Q				
	Estimate	Std Error	Z value	P-value
Intercept	-1.17013	0.06558	-17.842	<2e-16
Habitat (pasture)	-0.05385	0.07407	-0.727	0.467

Simpson's				
	Estimate	Std Error	Z value	P-value
Intercept	0.81791	0.07955	10.282	<2e-16
Habitat (pasture)	-0.05303	0.08651	-0.613	0.54

Uniqueness				
	Estimate	Std Error	Z value	P-value
Intercept	-0.66581	0.05301	-12.560	<2e-16
Habitat (pasture)	-0.05283	0.05970	-0.885	0.376

Landscape variable calculations

Landscape variables were calculated in QGIS using land-use data from MapBiomass Brazil (Land cover and use, Collection 9).

Forest and pasture cover was calculated as follows:

$$\text{Land-use cover} = \text{Total area of land-use pixel} / \text{Buffer area}$$

Forest patch density was calculated as follows:

$$\text{Forest patch density (patches per ha)} = (\text{number of forest patches} / \text{total area of forest in buffer (m}^2\text{)}) * 10,000$$

Edge density was calculated as follows:

$$\text{Edge density (meters per ha)} = (\text{total length of forest patch edges (m)} / \text{buffer area (m}^2\text{)}) * 10,000$$