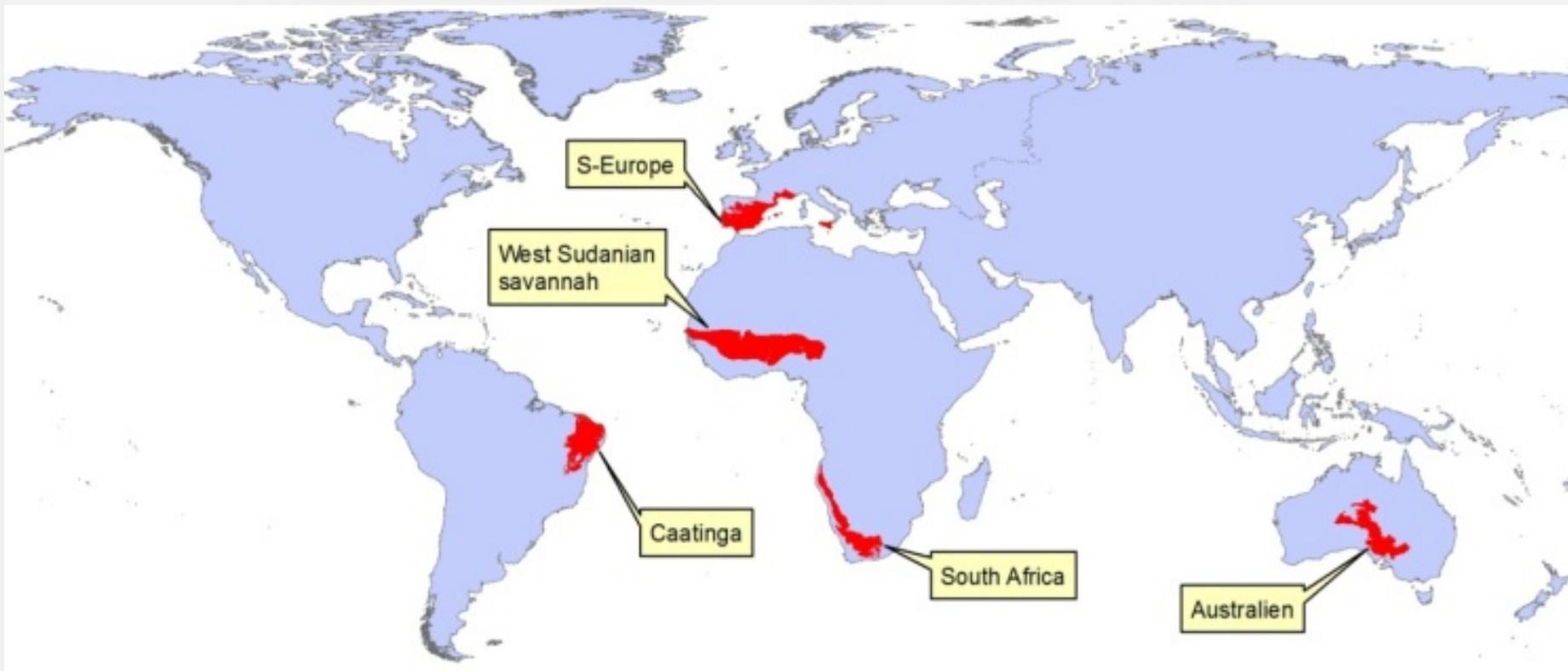


Contrasting EO indicators with biodiversity data: drylands

J.C. Campos, J.V. Leite, C.G. Vale and J.C. Brito

- 1 Test sites**
- 2 Validation of Indicators with Biodiversity Data - STATUS**
- 3 Validation of Indicators with Biodiversity Data - TRENDS**

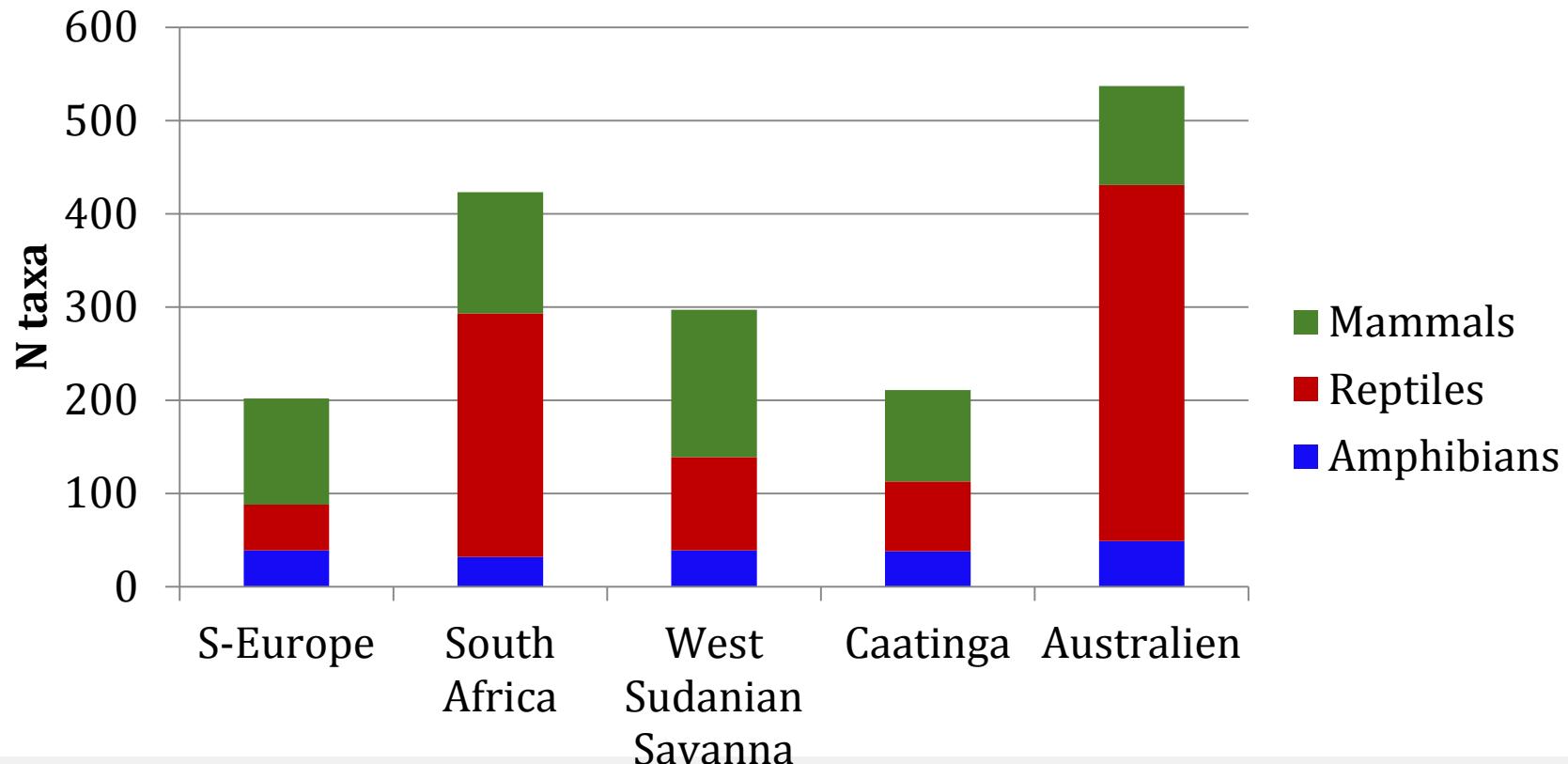
1. Test sites



2. Validation of Indicators with Biodiversity Data - STATUS

Biodiversity data

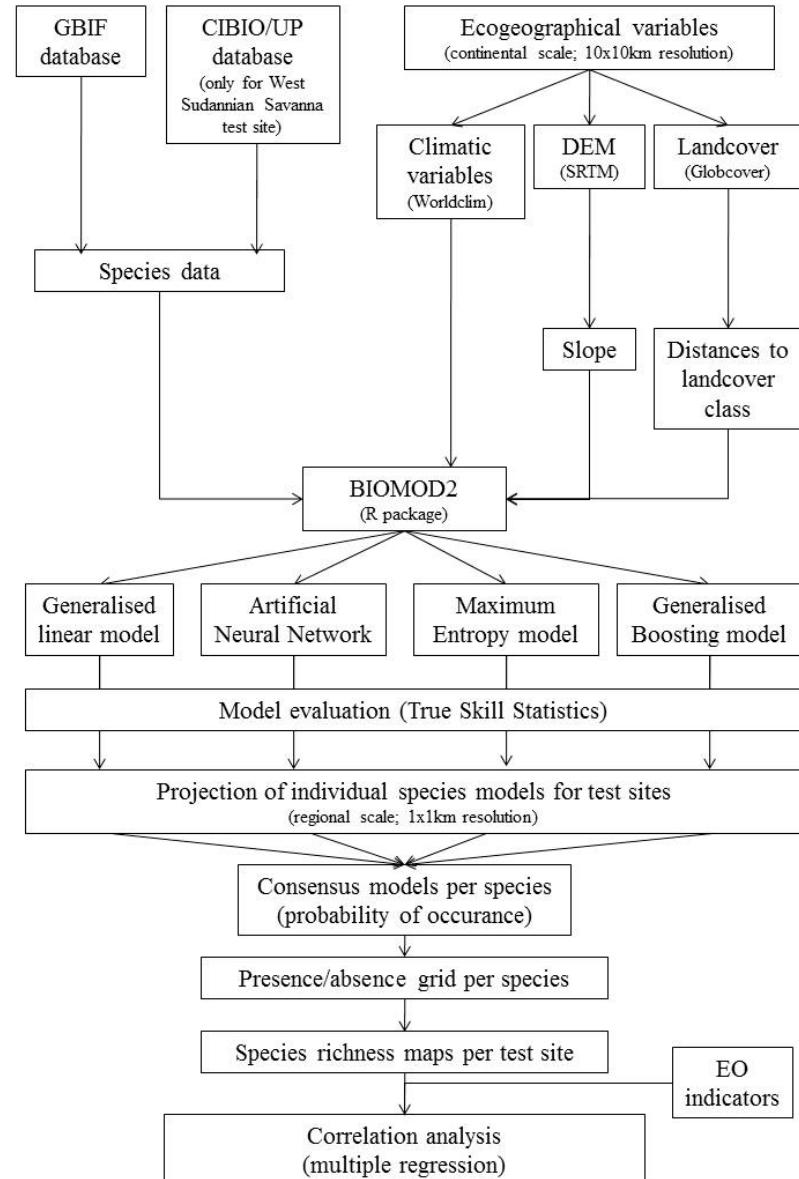
- Species richness of mammals, reptiles and amphibians (GBIF-only)



2. Validation of Indicators with Biodiversity Data - STATUS

Validation methods

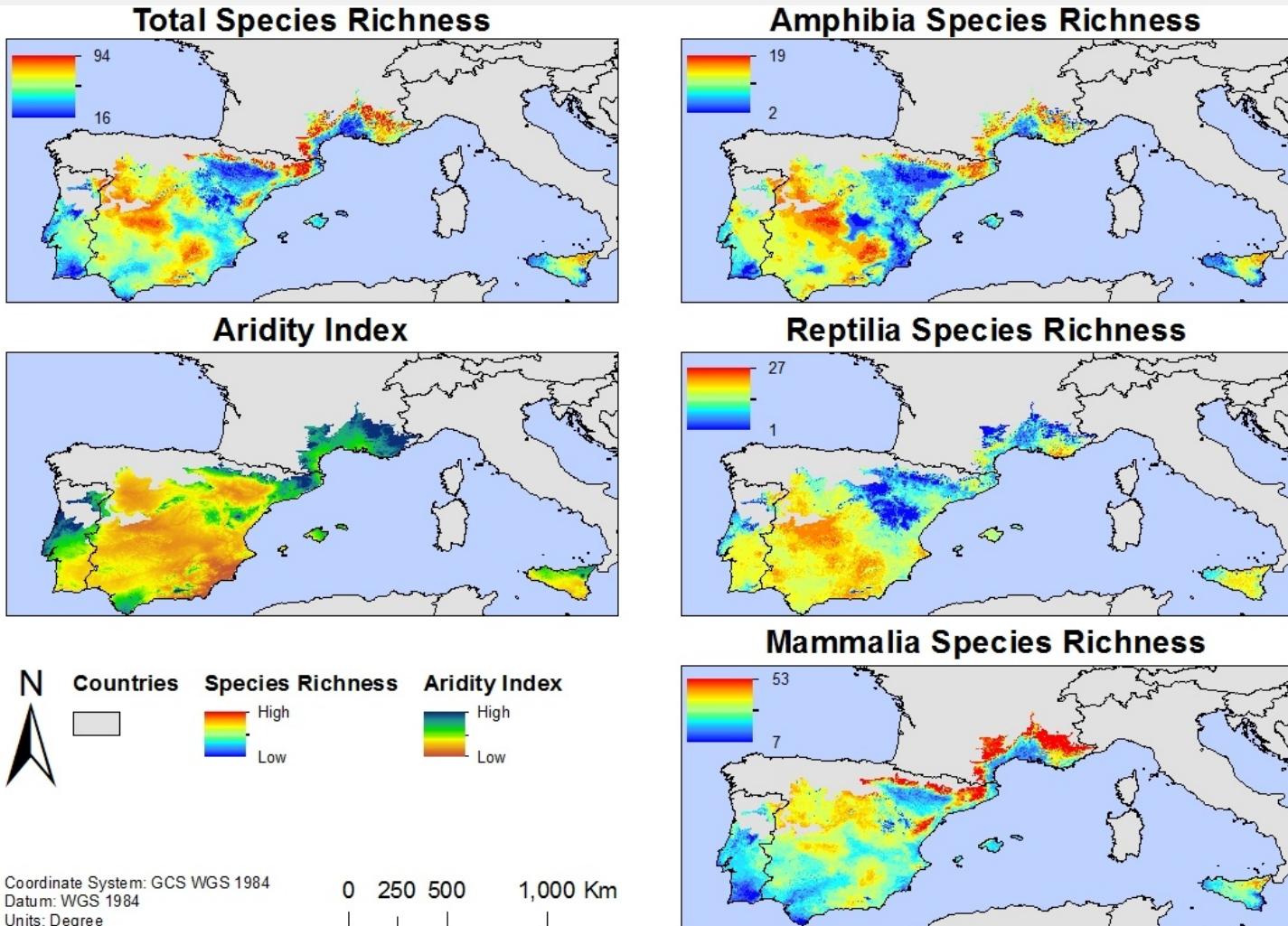
- Biodiversity distribution data
- Climatic, topographical and habitat data for each test site
- Development of distribution models for each species
- Combination of models to derive predictions of species richness
- Correlation of indicators with predictions of species richness distribution



2. Validation of Indicators with Biodiversity Data - STATUS

Validation results

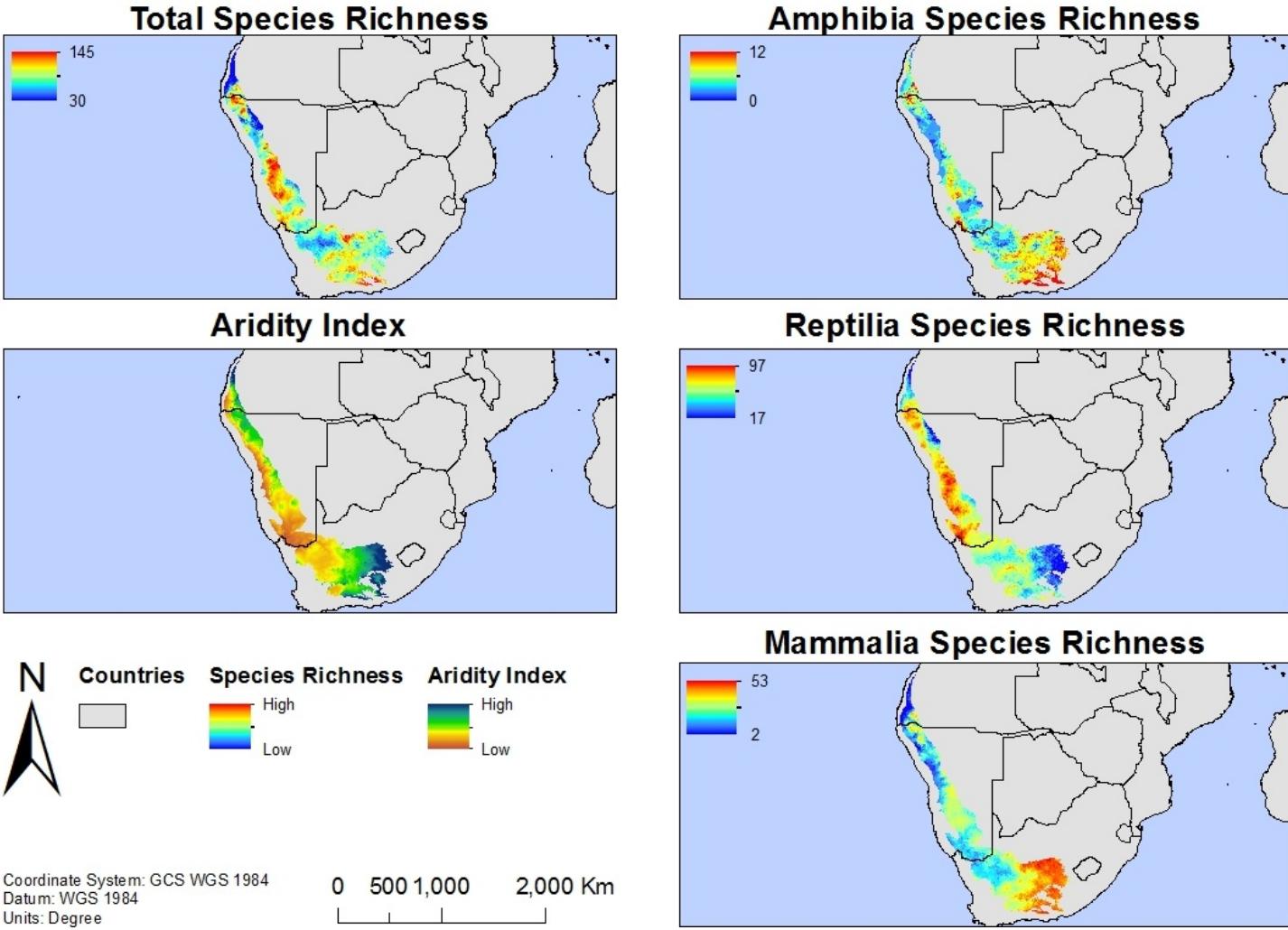
- Species richness distribution (total/by group)
- S-Europe
- Hotspots in Central Iberia, NE Spain, S France, and Sicily island



2. Validation of Indicators with Biodiversity Data - STATUS

Validation results

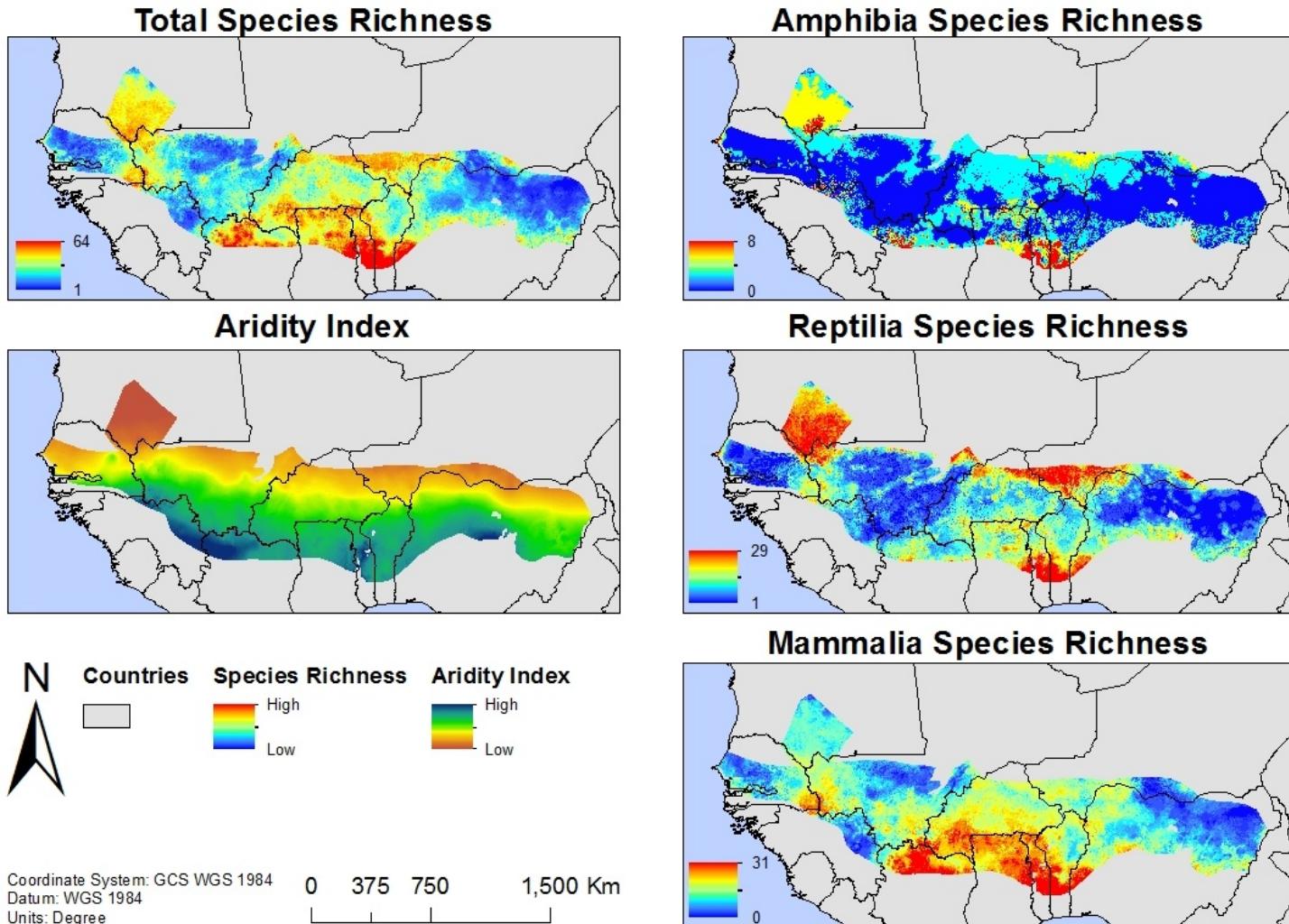
- Southern Africa
- Test site with highest species richness
- Reptile hotspots: Central-N regions
- Amphibian and mammal hotspots: S regions



2. Validation of Indicators with Biodiversity Data - STATUS

Validation results

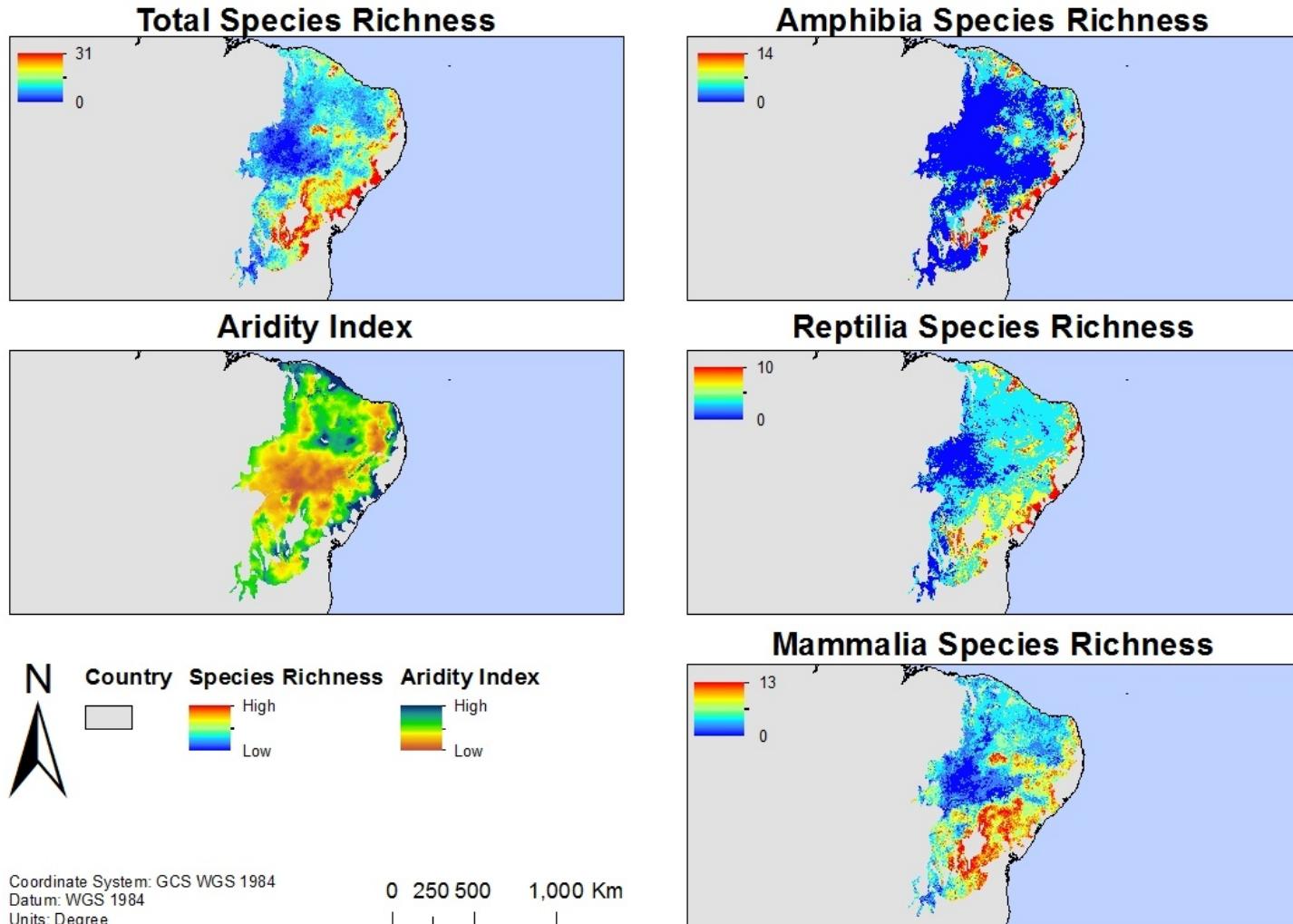
- West Sudanian savanna
- Hotspots in southern and northern limits, probably related to biome transition



2. Validation of Indicators with Biodiversity Data - STATUS

Validation results

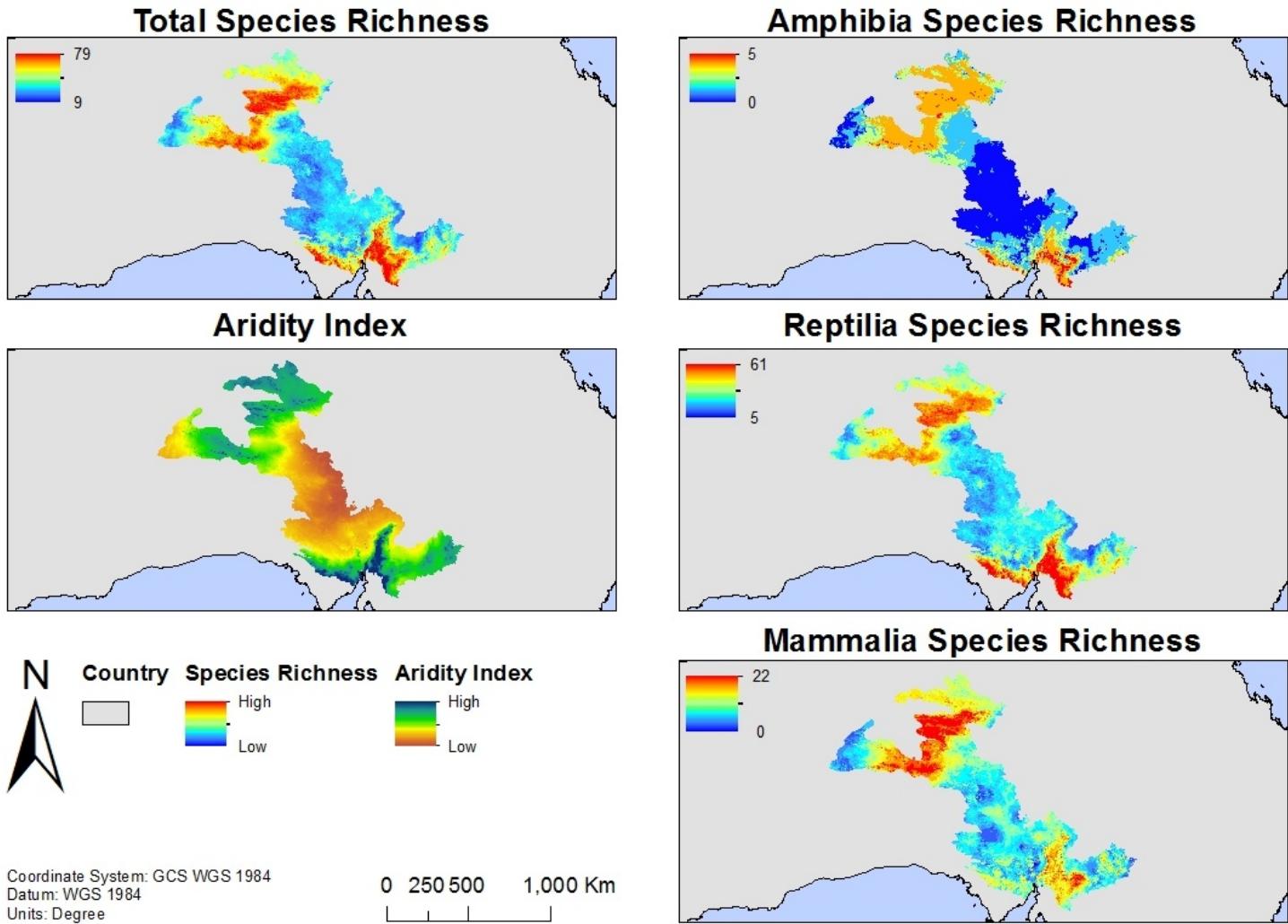
- Caatinga
- Relatively low sample size of species data
- Models predict relatively low species richness per grid cell
- Hotspots in the E-SE region, close to Cerrado and Atlantic forest



2. Validation of Indicators with Biodiversity Data - STATUS

Validation results

- Australien
- Hotspots in N and S regions
- Highly localized hotspots in central region, associated to desert water springs



2. Validation of Indicators with Biodiversity Data - STATUS

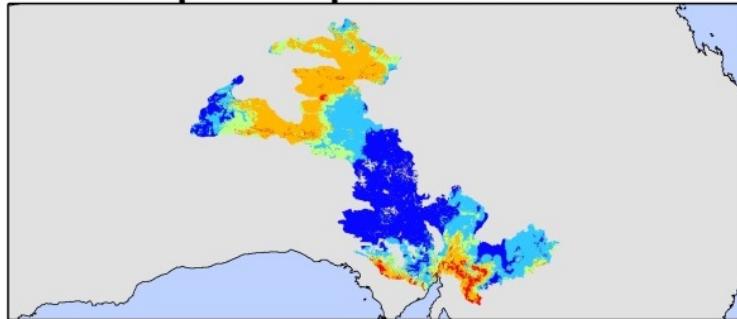
- Correlations between species richness and indicators

Test Sites	Indicators	fAPAR						Rain Use Efficiency						TRMM	
		Cyclic Fraction		Dry Season		Vegetation Year		Cyclic Fraction		Dry Season		Vegetation Year		Vegetation Year	
		Species Richness	Mean	Var	Mean	Var	Mean	Var	Mean	Var	Mean	Var	Mean	Var	Mean
Caatinga	Total	0.001	0.183	0.502	0.343	0.304	-0.054	0.361	0.254	0.446	0.316	0.250	0.107	-0.205	0.101
	Amphibia	-0.011	0.147	0.458	0.207	0.266	-0.095	0.209	0.223	0.279	0.231	0.124	0.089	-0.073	0.146
	Reptilia	0.012	0.212	0.421	0.322	0.251	0.002	0.369	0.284	0.392	0.323	0.223	0.183	-0.197	0.146
	Mammalia	0.002	0.121	0.379	0.309	0.240	-0.036	0.316	0.163	0.408	0.246	0.248	0.040	-0.221	0.011
	Total	0.094	-0.088	0.104	-0.124	-0.081	-0.086	-0.160	0.039	-0.185	0.130	0.009	-0.157	0.281	-0.104
South Europe	Amphibia	0.171	-0.083	0.180	-0.206	0.237	-0.117	-0.063	-0.093	0.092	-0.193	0.089	-0.011	0.328	0.081
	Reptilia	0.089	0.112	-0.150	0.052	-0.085	0.160	0.086	0.066	-0.072	0.075	0.059	0.257	-0.244	0.405
	Mammalia	0.011	-0.165	0.168	-0.127	0.144	-0.169	-0.155	-0.240	0.061	-0.241	-0.063	-0.393	0.413	-0.426
	Total	-0.148	-0.148	-0.148	-0.148	-0.148	-0.148	-0.148	-0.148	-0.148	-0.148	-0.148	-0.148	-0.148	-0.148
West Sudanian	Amphibia	-0.443	-0.443	-0.157	0.321	-0.357	0.119	-0.424	0.392	-0.216	0.314	-0.404	0.382	-0.072	-0.029
	Reptilia	-0.435	0.371	-0.182	0.377	-0.358	0.371	-0.441	0.386	-0.262	0.373	-0.424	0.394	-0.129	-0.036
	Mammalia	0.268	-0.154	0.370	-0.173	0.329	-0.185	0.041	-0.164	0.299	-0.258	0.183	-0.214	0.488	-0.457
	Total	-0.007	-0.007	-0.007	-0.007	-0.007	-0.007	-0.007	-0.007	-0.007	-0.007	-0.007	-0.007	-0.007	-0.007
South African	Amphibia	0.243	-0.352	0.357	-0.291	0.308	-0.372	0.198	-0.321	0.412	-0.283	0.378	-0.362	0.295	-0.378
	Reptilia	-0.291	0.362	-0.373	0.436	-0.370	0.376	0.079	0.422	-0.373	0.405	-0.284	0.446	-0.469	0.389
	Mammalia	0.323	-0.510	0.322	-0.345	0.357	-0.524	0.104	-0.395	0.384	-0.303	0.405	-0.443	0.418	-0.417
	Total	0.569	-0.514	0.461	-0.459	0.617	-0.535	0.225	-0.418	0.472	-0.296	0.583	-0.476	0.534	-0.218
Australia n	Amphibia	0.602	-0.541	0.453	-0.478	0.628	-0.560	0.194	-0.426	0.457	-0.258	0.574	-0.480	0.631	-0.161
	Reptilia	0.589	-0.522	0.481	-0.478	0.638	-0.542	0.271	-0.445	0.496	-0.339	0.617	-0.496	0.500	-0.283
	Mammalia	0.349	-0.342	0.274	-0.266	0.388	-0.360	0.051	-0.231	0.274	-0.104	0.336	-0.290	0.447	-0.006

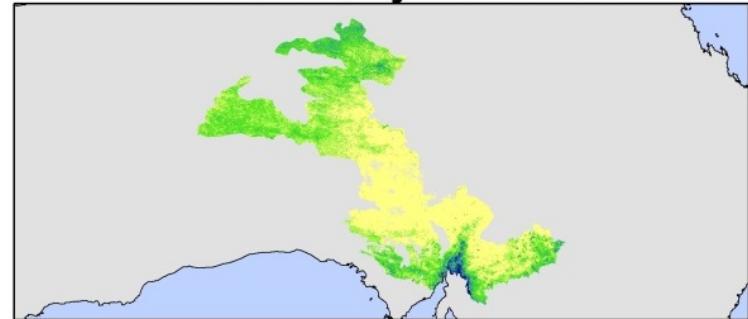
2. Validation of Indicators with Biodiversity Data - STATUS

- Australien
- Amphibian species richness and the three most highly correlated indicators

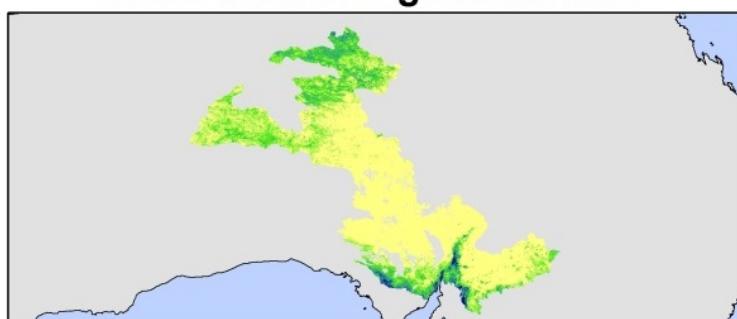
Amphibia Species Richness



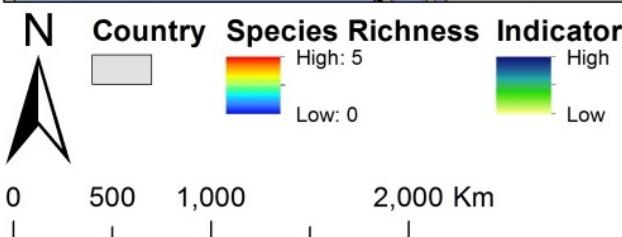
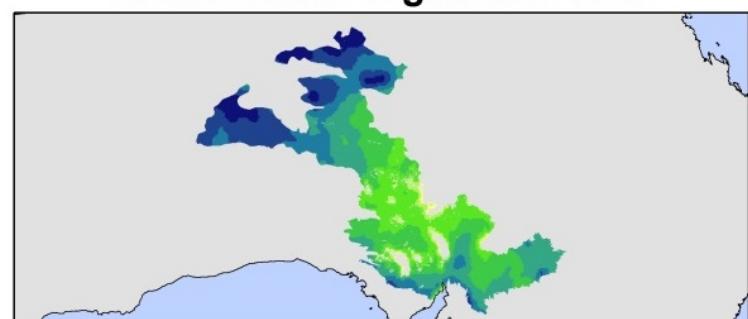
Mean fAPAR Cyclic Fraction



Mean fAPAR Vegetation Year

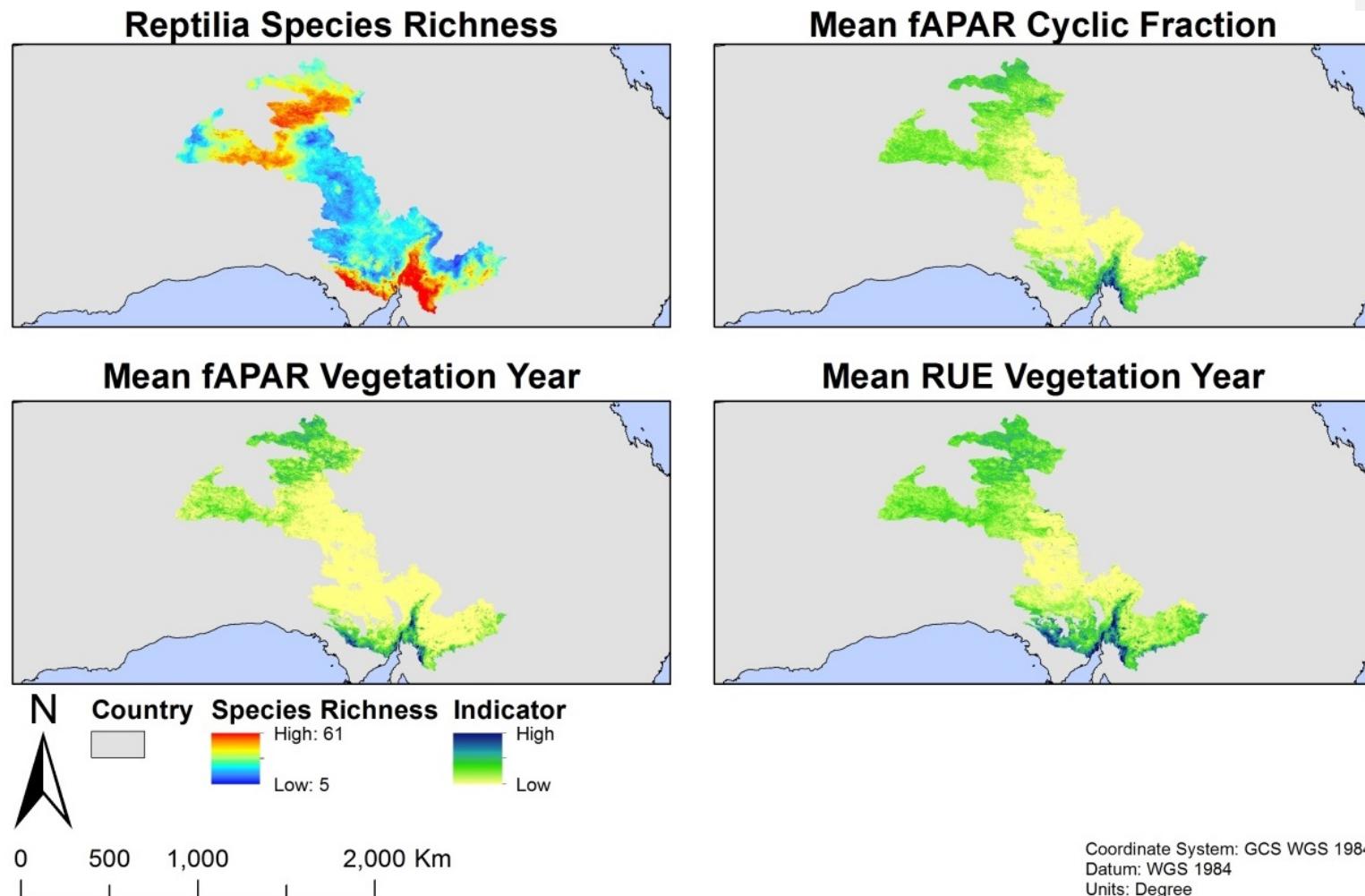


Mean TRMM Vegetation Year



2. Validation of Indicators with Biodiversity Data - STATUS

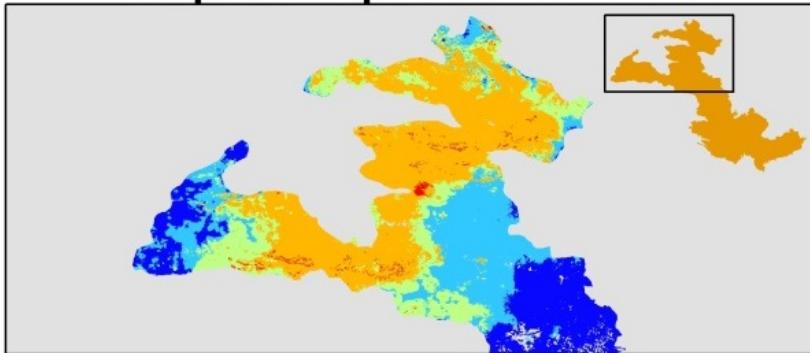
- Australien
- Reptilia species richness and the three most highly correlated indicators



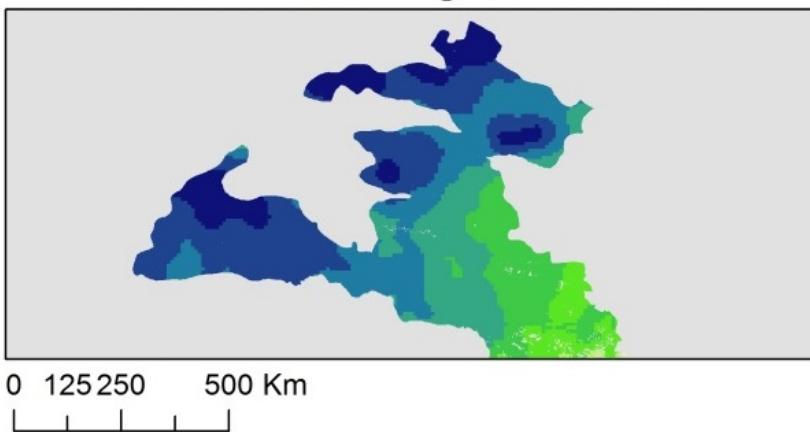
2. Validation of Indicators with Biodiversity Data - STATUS

- Australien

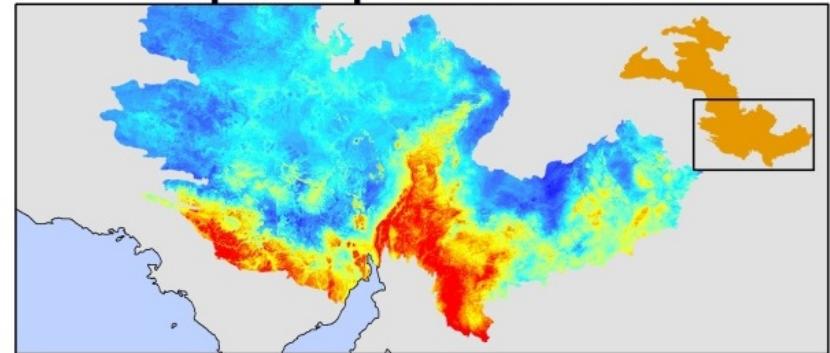
Amphibia Species Richness



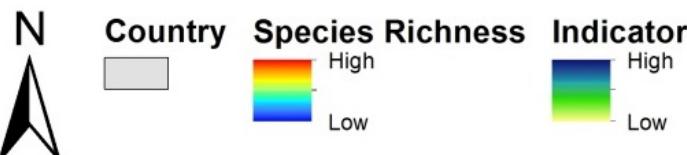
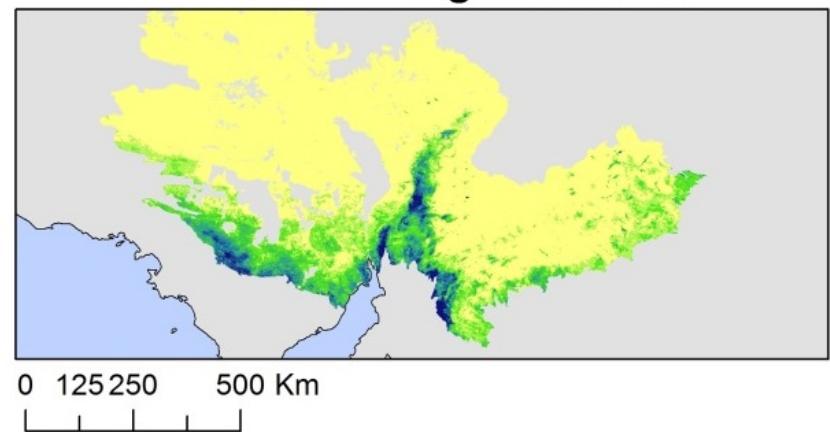
Mean TRMM Vegetation Year



Reptilia Species Richness



Mean fAPAR Vegetation Year



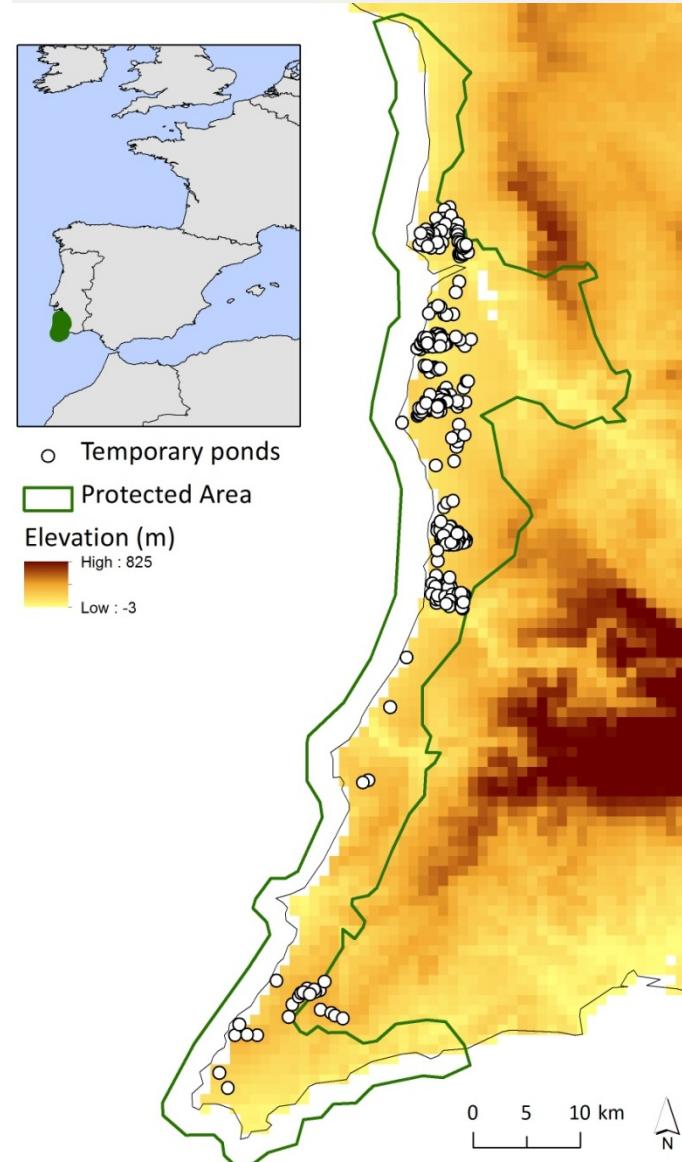
3. Validation of Indicators with Biodiversity Data - TRENDS

- Trend data

Dryland	Contact	Institution	Reference
S-Europe	Pedro Beja	CIBIO, Portugal	Ferreira & Beja 2013
South Africa	Rita Covas	CIBIO, Portugal	Covas et al. 2008
West Sudanian savannah	Adam Konecky Patrick Triplet	Institute of Vertebrate Biology, Academy of Sciences of the Czech Republic OMPO, Oiseaux migrateurs du Paléarctique occidental, Paris	Konecky et al. 2009 Triplet et al. 2010
Caatinga	Júlio Ferreira Izabel Silva	Universidade Estadual Da Paraíba. Centro De Ciências Biológicas E Da Saúde (CCBS) Universidade Federal de Sergipe	Ferreira 2011 Silva 2013
Australien	Simon Ferrier	CSIRO, Ecosystem Sciences, Australia	Waiting contact

3. Validation of Indicators with Biodiversity Data - TRENDS

- S-Europe
- 50 temporary ponds (sampled 2002, 2009, 2010 and 2011)
- Amphibian richness across the breeding season (February-June)
 - N Taxa: nº of taxa detected (max. 13 species)
 - Abundance: average nº of individuals (max. 9 species)
- Temporal overlap between indicators and biodiversity data: 8 ponds retained
- Temporal sample size and null variance: 3 ponds retained



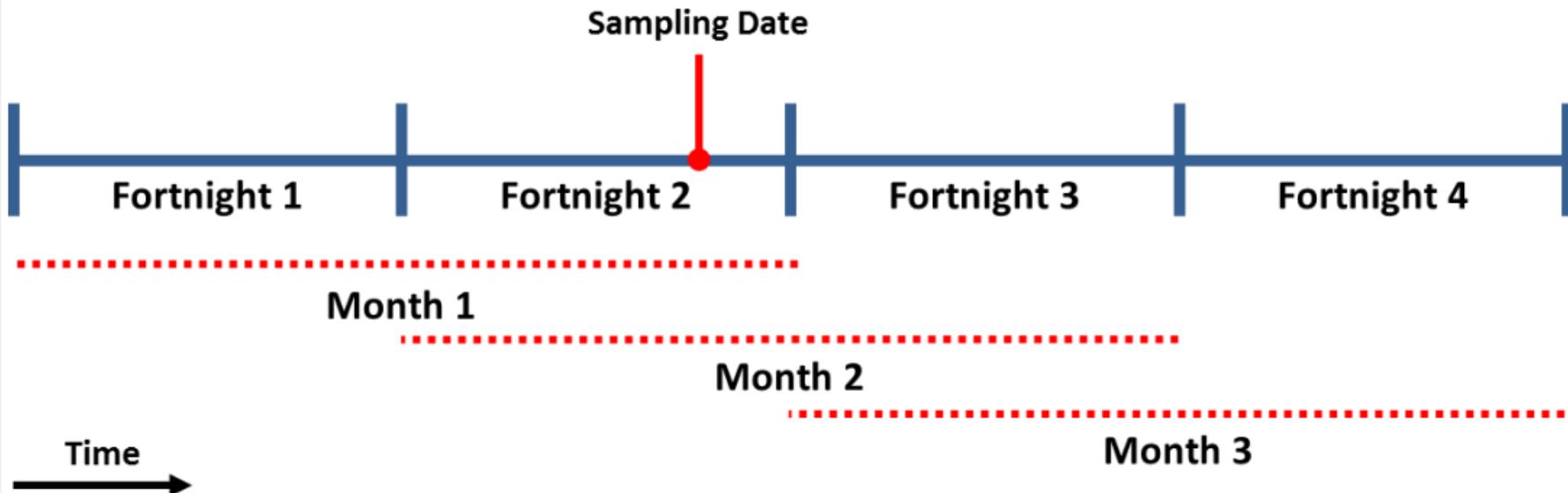
[Biological Conservation 165 \(2013\) 179–186](#)

Mediterranean amphibians and the loss of temporary ponds:
Are there alternative breeding habitats?

Mário Ferreira, Pedro Beja *

3. Validation of Indicators with Biodiversity Data - TRENDS

- S-Europe
- Biodiversity data: day / Indicator data (fAPAR): 15-days
- 4 aggregation schemes of biodiversity data essayed: Fortnights 1, 2, 3 and 4
- 4 aggregation schemes of biodiversity/indicator data essayed: Month 1, 2, and 3, and Maximum across all fortnights (not depicted)



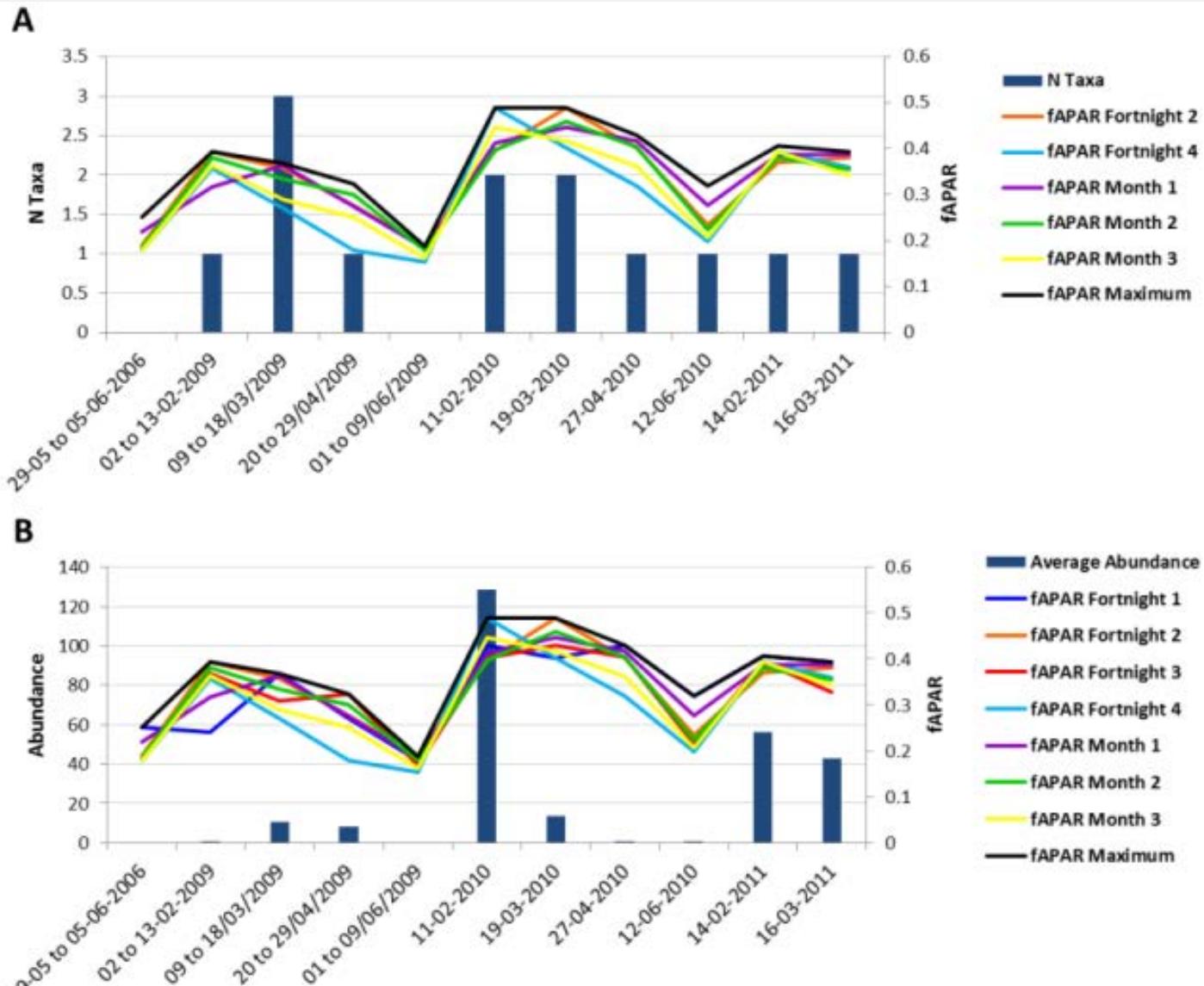
3. Validation of Indicators with Biodiversity Data - TRENDS

- S-Europe
- Significant positive correlations between indicator (fAPAR) and biodiversity data (N taxa and Abundance)
- No single aggregation scheme essayed significant across all test-ponds

Indicators (fAPAR)	N Taxa			Abundance		
	A10	C30	G03	A10	C30	G03
Fortnight 1	0.56	0.16	0.13	0.66 (0.028)	0.06	0.36
Fortnight 2	0.62 (0.041)	-0.51	0.36	0.61 (0.048)	-0.14	0.3
Fortnight 3	0.57	0.7 (0.016)	0.66 (0.027)	0.65 (0.03)	0.24	0.41
Fortnight 4	0.62 (0.041)	0.35	0.46	0.83 (0.002)	-0.03	0.77 (0.006)
Month 1	0.68 (0.022)	-0.29	0.3	0.68 (0.022)	-0.13	0.46
Month 2	0.62 (0.041)	0.65 (0.032)	0.4	0.66 (0.027)	0.25	0.36
Month 3	0.66 (0.026)	0.69 (0.019)	0.51	0.82 (0.002)	0.3	0.71 (0.014)
Maximum	0.66 (0.026)	0.39	0.48	0.78 (0.005)	-0.09	0.57

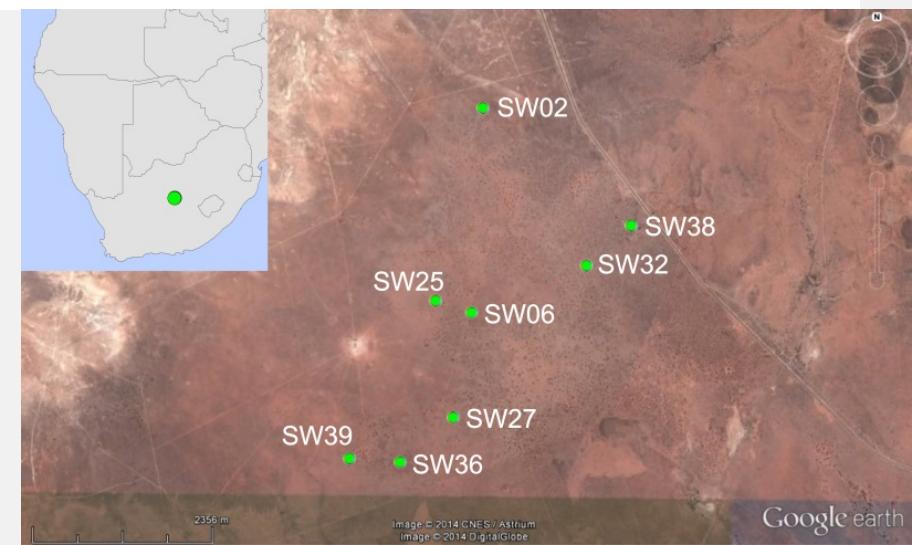
3. Validation of Indicators with Biodiversity Data - TRENDS

- S-Europe
- Pond A10
- Significant correlations:
 - almost all N taxa data aggregation schemes
 - all abundance aggregation schemes
- Sampling gaps in abundance



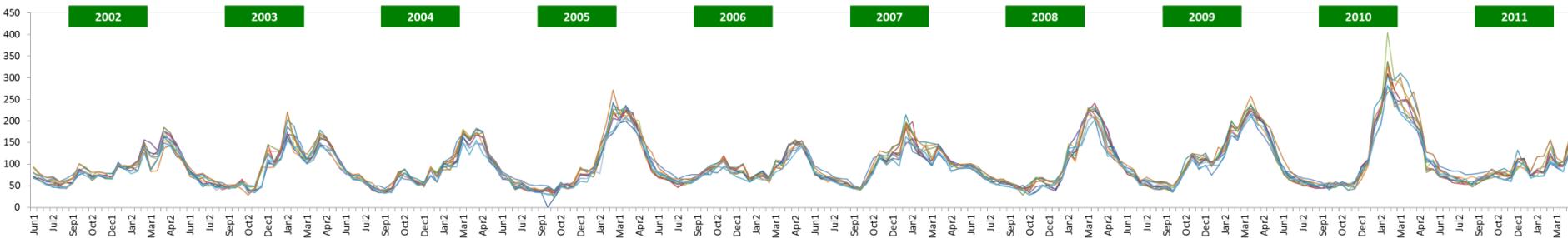
3. Validation of Indicators with Biodiversity Data - TRENDS

- **South Africa**
- **Benfontein Game Farm, Northern Cape Province, South Africa**
- **Data from various bird colonies in 2010 and 2012 on:**
 - **number of eggs (clutch size)**
 - **number of nests used (as a proxy to the number of breeding birds within a colony)**
 - **breeding season length**
- **Eight colonies selected (max. data possible)**



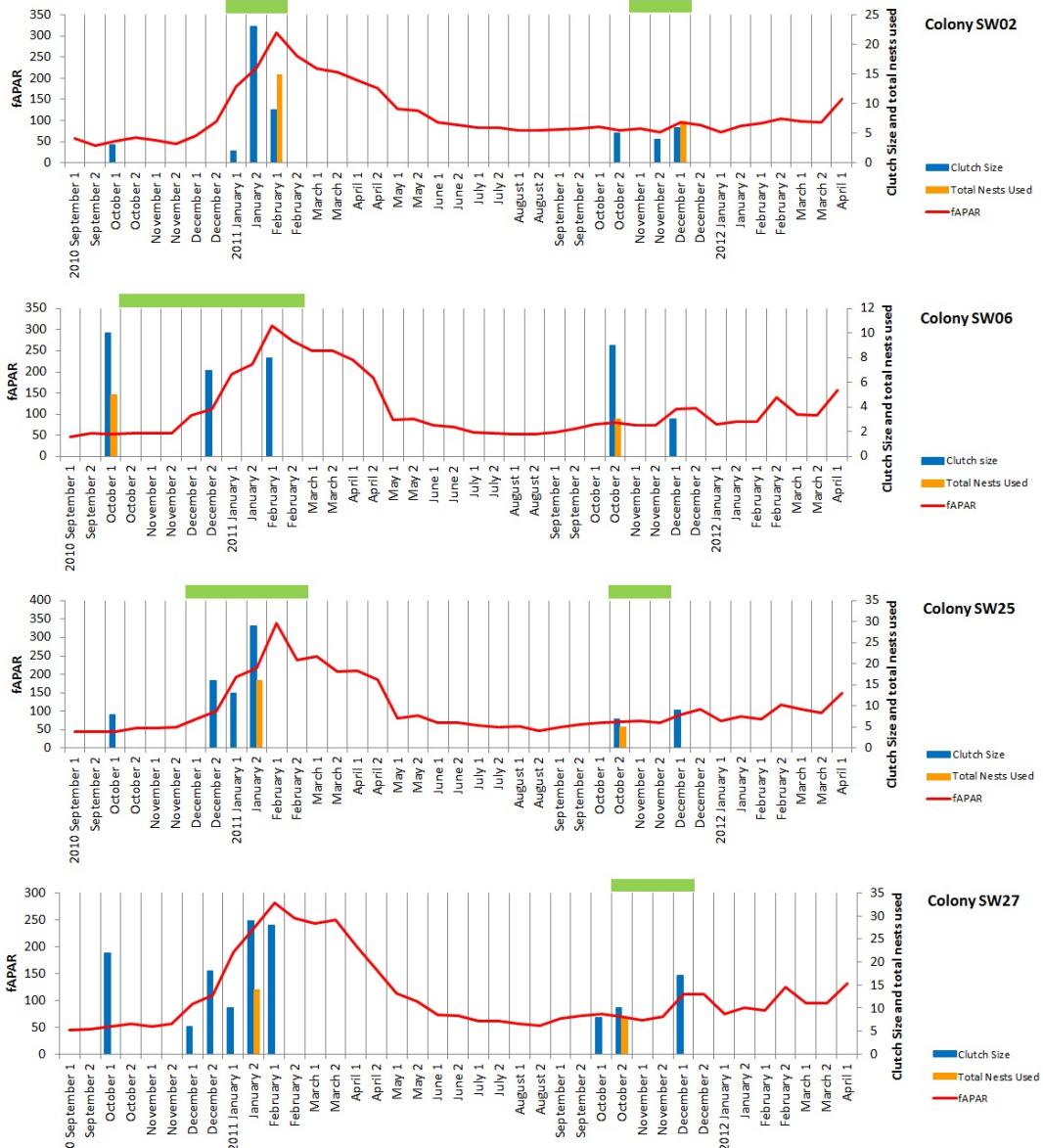
3. Validation of Indicators with Biodiversity Data - TRENDS

- South Africa
- Fortnight fAPAR values for the eight selected colonies along with theoretical reproductive season of sociable weavers between June 2002 and April 2012
- Apparent positive relationship between breeding and indicators



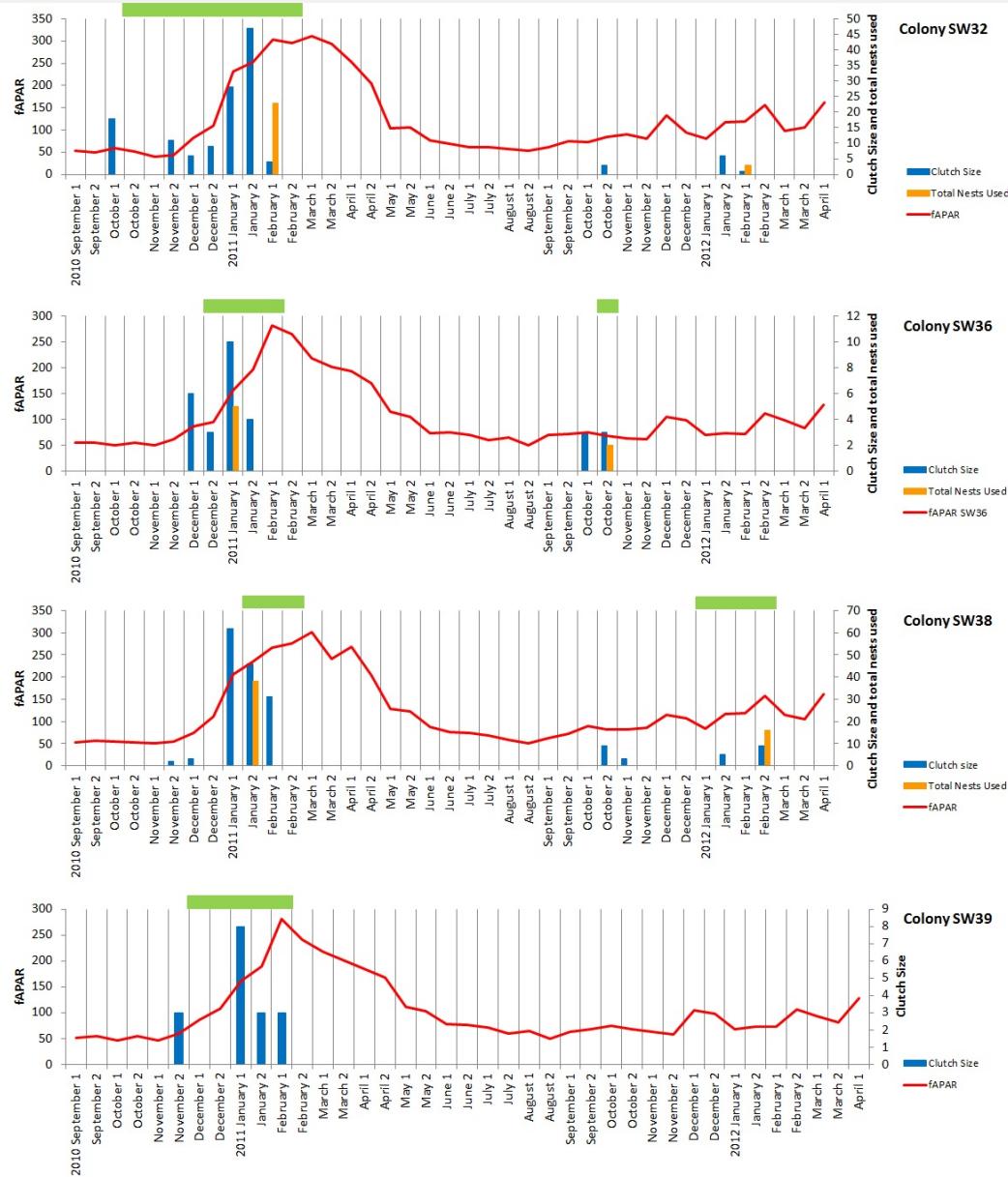
3. Validation of Indicators with Biodiversity Data - TRENDS

- South Africa
- Fortnight fAPAR values along with clutch size, total nests used, and breeding season length for four colonies
- Clutch sizes larger and egg laying events more frequent during 2nd fortnight of Dec. and 1st fortnight of Feb. – match with fAPAR peaks
- N eggs, the frequency at which they are laid, and N of nests used are much higher in 2010 than in 2011, as well as fAPAR



3. Validation of Indicators with Biodiversity Data - TRENDS

- South Africa
- Fortnight fAPAR values along with clutch size, total nests used, and breeding season length for four colonies



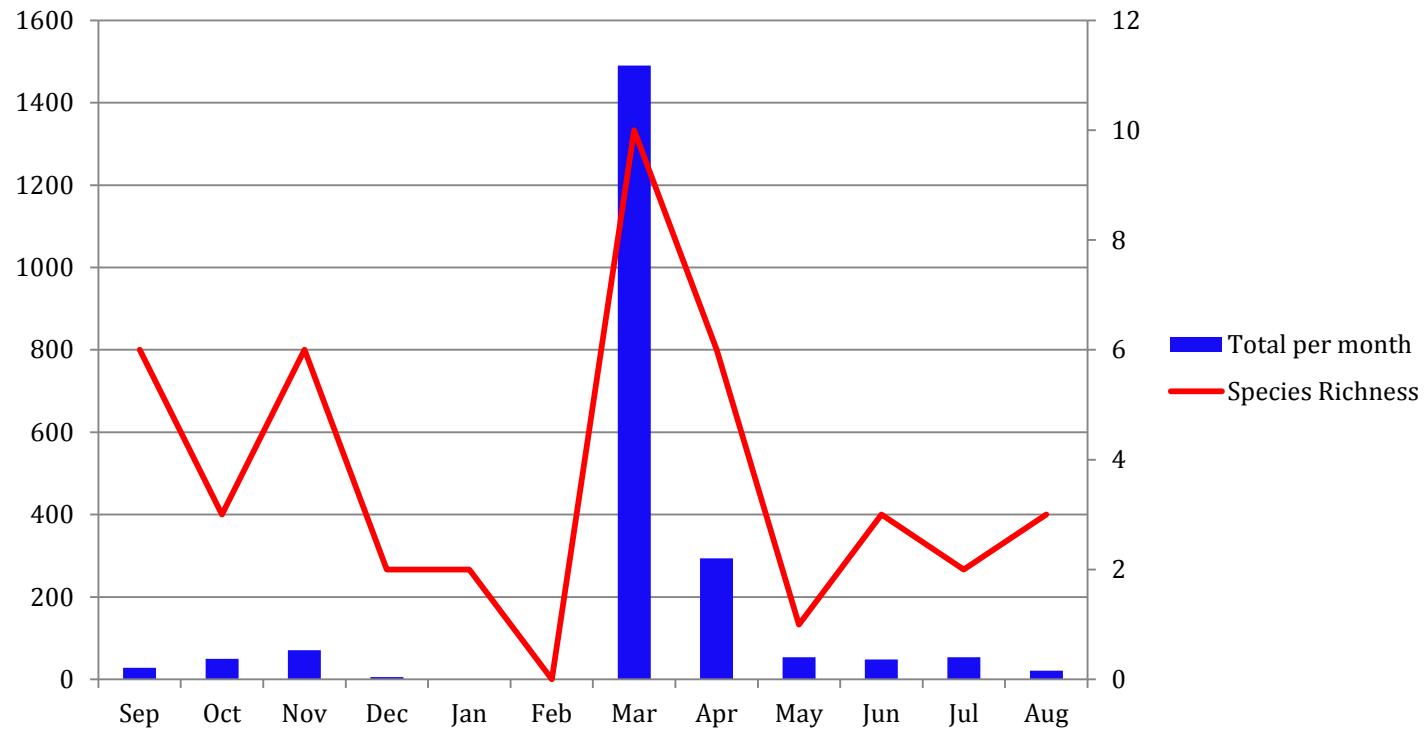
3. Validation of Indicators with Biodiversity Data - TRENDS

- Caatinga
- Monumento Natural Grota de Sergipe
- Tadpole abundance during September 2011 and August 2012



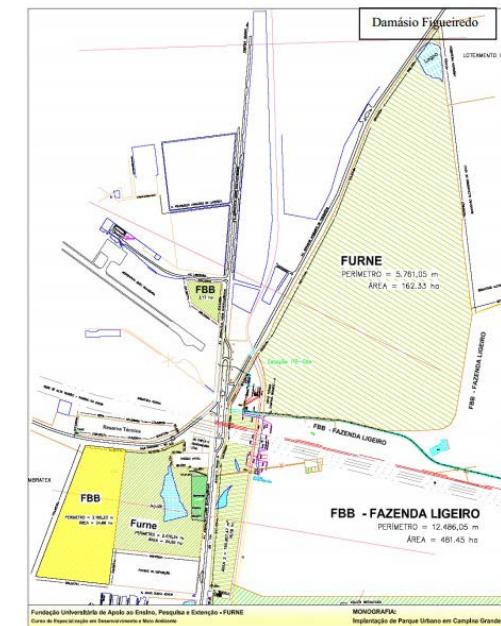
3. Validation of Indicators with Biodiversity Data - TRENDS

- Caatinga



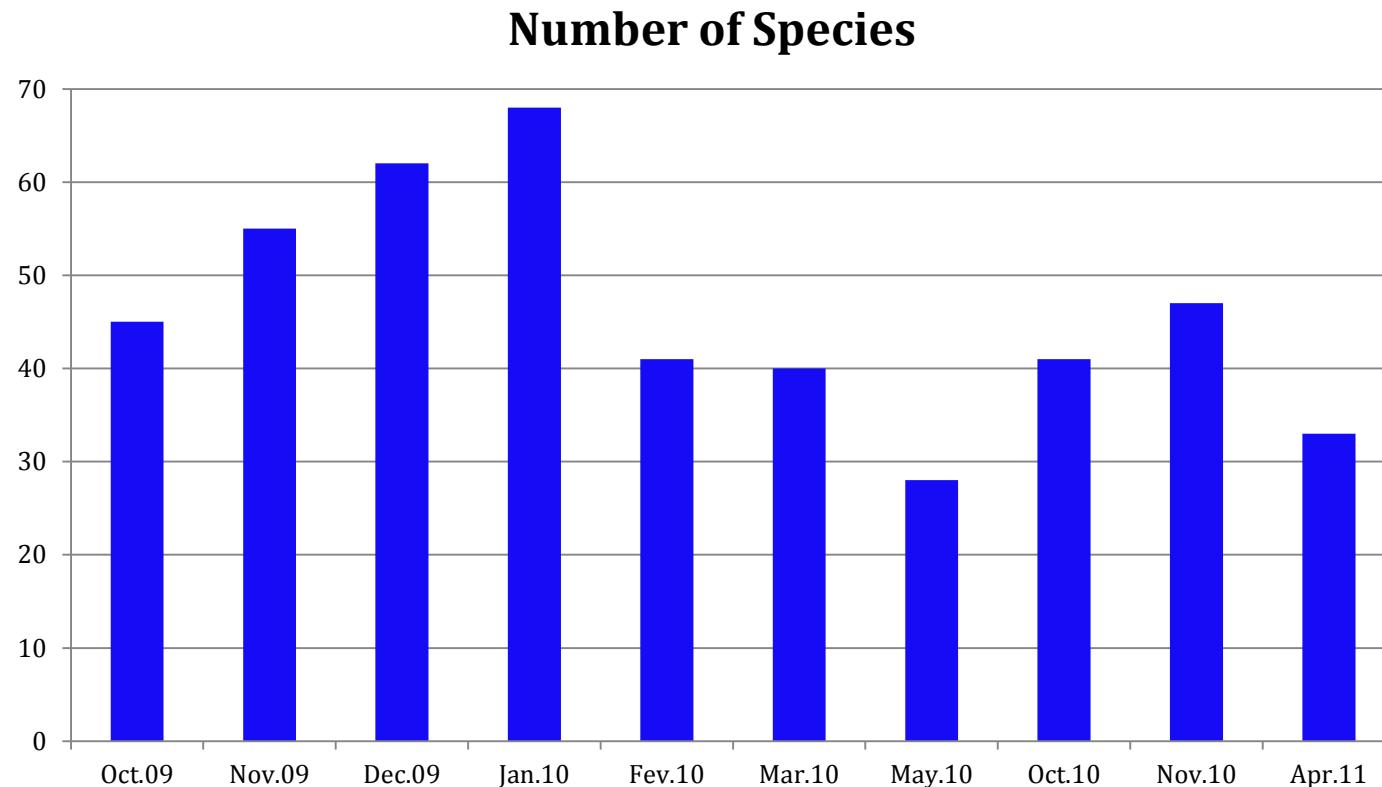
3. Validation of Indicators with Biodiversity Data - TRENDS

- Caatinga
- Complexo Aluizio Campos, Campina Grande
- Bird species richness between September 2009 and April 2011



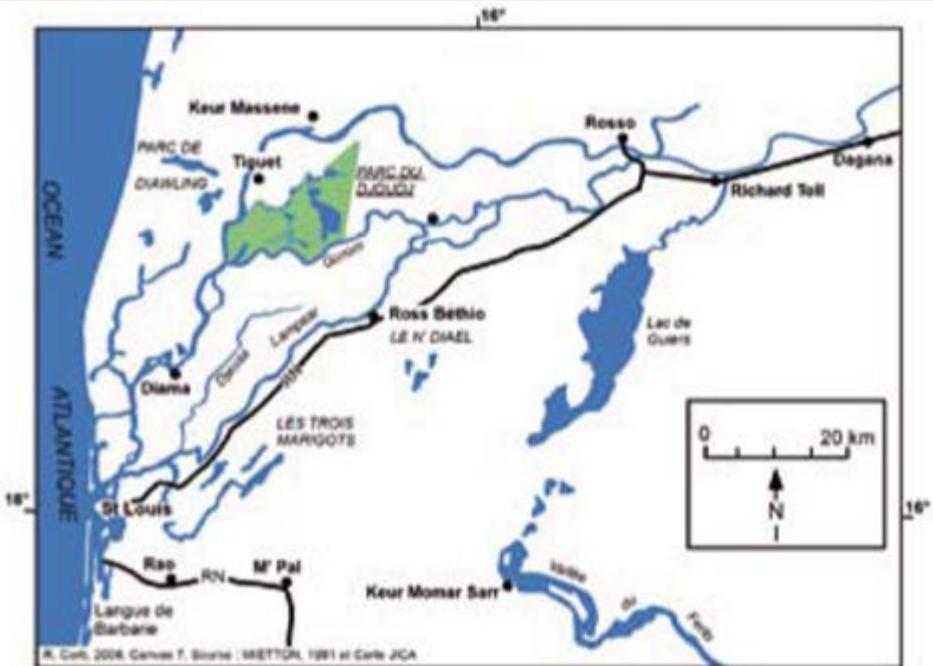
3. Validation of Indicators with Biodiversity Data - TRENDS

- Caatinga



3. Validation of Indicators with Biodiversity Data - TRENDS

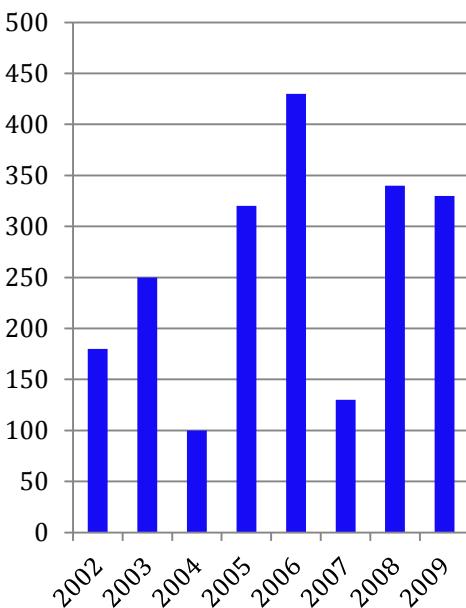
- West Sudanian Savannah
- National Park (Parc national des oiseaux du Djoudj)
- Data between 2002-2010
- Annual bird population counts



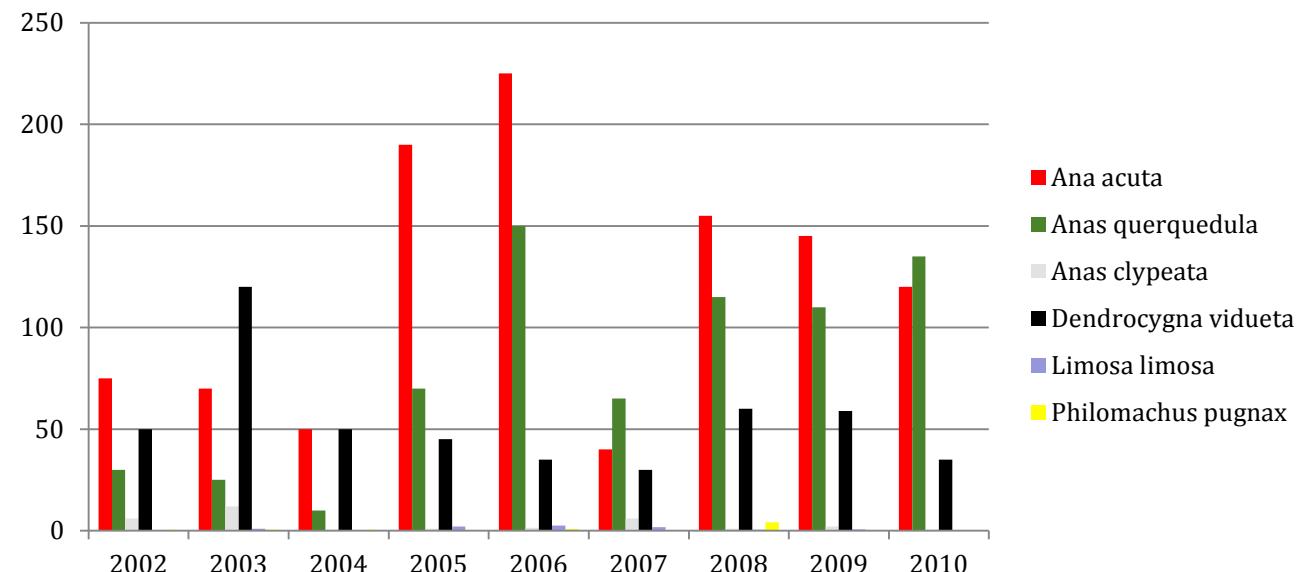
3. Validation of Indicators with Biodiversity Data - TRENDS

- West Sudanian Savannah

Total

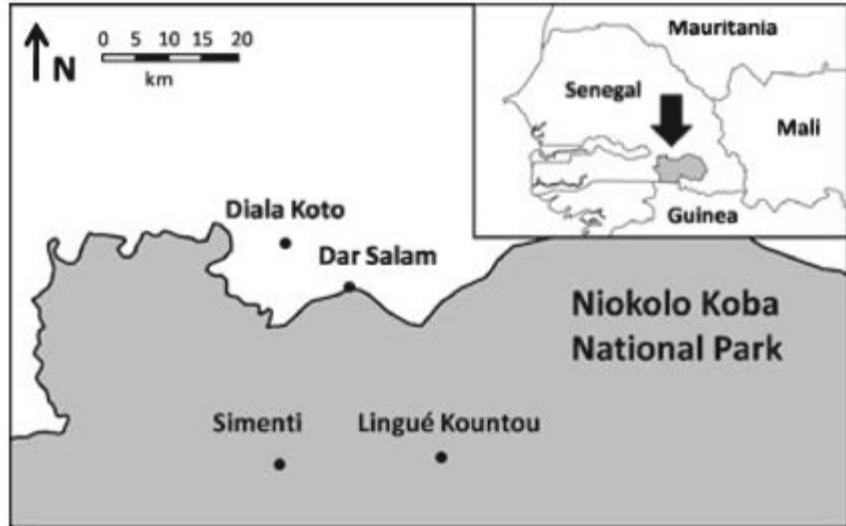


Species



3. Validation of Indicators with Biodiversity Data - TRENDS

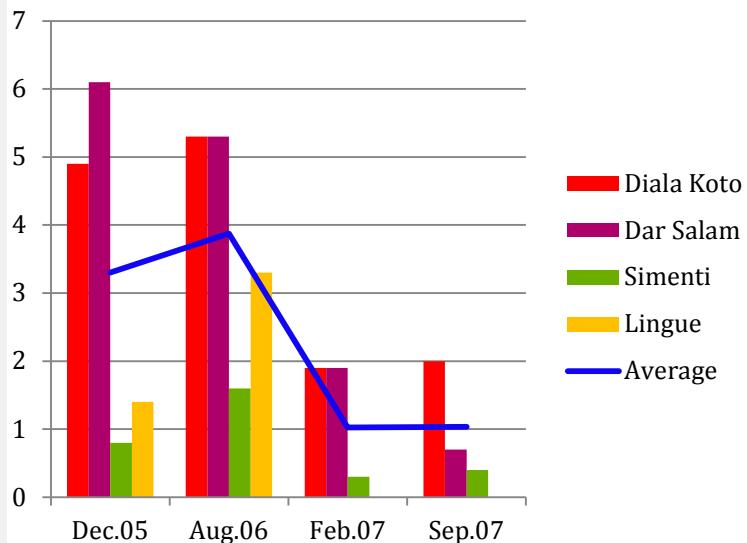
- West Sudanian Savannah
- National Park (Parc national du Niokolo Koba)
- Data between 2005-2007
- Abundance and species richness of small mammals



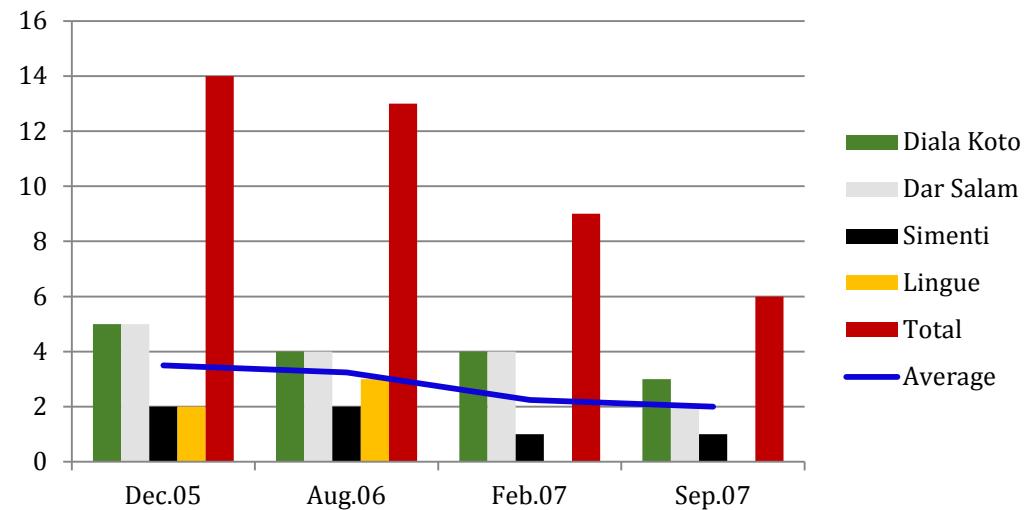
3. Validation of Indicators with Biodiversity Data - TRENDS

- West Sudanian Savannah

Abundance



Species richness



Thank you!!

- **Environmental Data**
 - **6 factors representing averages and extremes in precipitation and temperature**
 - **WorldClim database**
 - **1 factor representing slope derived from DEM (SRTM)**

CODE	Factor
BIO1	Annual Mean Temperature
BIO5	Max Temperature of Warmest Month
BIO6	Min Temperature of Coldest Month
BIO12	Annual Precipitation
BIO13	Precipitation of Wettest Month
BIO14	Precipitation of Driest Month
SLOPE	Slope

- **Environmental Data**
- **Land-cover categories per continent**
- **Distances to land cover categories**
- **Globcover 2006**

Continent	Code	Description
Africa	D03	Mosaic vegetation (50-70%) / cropland (20-50%)
	D07	Open (15-40%) broadleaved deciduous forest/woodland (>5m)
	D09	Closed to open (>15%) (broadleaved or needleleaved, evergreen or deciduous) shrubland (<5m)
	D10	Closed to open (>15%) herbaceous vegetation (grassland, savannahs or lichens/mosses)
	D14	Bare areas
	D15	Consolidated bare areas (hardpans, gravels, bare rock, stones, boulders)
Australia	D16	Non-consolidated bare areas (sandy desert)
	D03	Closed to open (>15%) broadleaved evergreen or semi-deciduous forest (>5m)
	D05	Mosaic forest or shrubland (50-70%) / grassland (20-50%)
	D06	Mosaic grassland (50-70%) / forest or shrubland (20-50%)
	D01	Rainfed croplands
	D07	Closed to open (>15%) (broadleaved or needleleaved, evergreen or deciduous) shrubland (<5m)
Europe	D08	Closed to open (>15%) herbaceous vegetation (grassland, savannahs or lichens/mosses)
	D09	Sparse (<15%) vegetation
	D01	Rainfed croplands / herbaceous crops / shrub or tree crops
	D02	Mosaic cropland (50-70%) / vegetation (grassland/shrubland/forest) (20-50%)
	D04	Closed (>40%) broadleaved deciduous forest (>5m)
	D06	Open (15-40%) needleleaved deciduous or evergreen forest (>5m)
South America	D08	Closed to open (>15%) mixed broadleaved and needleleaved forest (>5m)
	D12	Sparse (<15%) vegetation (grassland or shrubland)
	D01	Rainfed croplands and shrub or tree crops
	D02	Mosaic cropland (50-70%) / grassland/shrubland/forest (20-50%)
	D04	Closed to open (>15%) broadleaved evergreen or semi-deciduous forest (>5m)
	D03	Mosaic vegetation (grassland/shrubland/forest) (50-70%) / cropland (20-50%)
	D10	Closed to open (>15%) (broadleaved or needleleaved, evergreen or deciduous) shrubland (<5m)
	D05	Closed (>40%) broadleaved evergreen and/or semi-deciduous forest (>5m)

- **Species distribution models**
- **4 modeling techniques applied:**
 - Maximum Entropy model (MXT);
 - Generalised Linear models (GLM);
 - Artificial Neural Networks (ANN);
 - Generalised Boosting models (GBM);
- **Individual model quality evaluated by True Skill Statistics (threshold 0.7)**
- **Projection of continental models to each test site**
- **Ensemble prediction of probability of species occurrence (per taxonomic group)**
- **Estimates of species richness distribution in each test site**

```
 esa@geneplan: ~  
+  
+ ### Initialisation (C: pseudoabs-->random)  
+ myBiomodData <- BIOMOD_FormattingData(resp.var = myResp,  
+ expl.var = myExpl,  
+ resp.xy = myRespCoord,  
+ resp.name = myRespName,  
+ PA.nb.rep = 2,  
+ PA.nb.absences = 10000,  
+ na.rm=TRUE,  
+ PA.strategy = 'random')  
+  
+ myBiomodData  
+  
+ plot(myBiomodData)  
+  
+  
+ ### Options definition  
+ myBiomodOption <- BIOMOD_ModelingOptions()  
+ ### Modelling  
+ myBiomodModelOut <- BIOMOD_Modeling(  
+ myBiomodData,  
+ models = c('GLM', 'GBM', 'ANN', 'MAXENT'),  
+ models.options = myBiomodOption,  
+ NbRunEval=1,  
+ DataSplit=80,  
+ Yweights=NULL,  
+ VarImport=2,  
+ models.eval.meth = c('KAPPA', 'ROC', 'TSS'),  
+ SaveObj = TRUE,  
+ rescal.all.models = TRUE,  
+ do.full.models = FALSE)  
+  
+  
+ ### save models evaluation scores and variables importance on hard drive  
+ capture.output(getModelsEvaluations(myBiomodModelOut),  
+ file=file.path(sp.n,  
+ "models_evaluation.txt"))
```