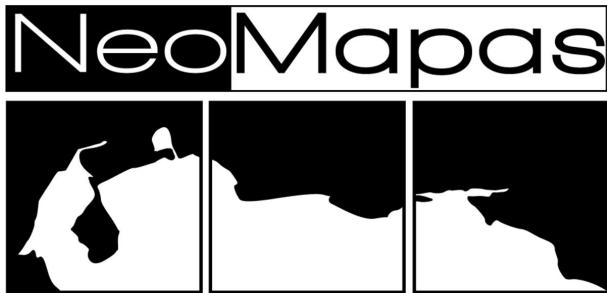


Monitoring Psittacidae in Venezuela: Distribution data and occupancy models for 2010



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1 Introduction

Venezuela has a high diversity of psittacids (Aves, Psittaciformes) but many species are threatened by increasing rate of land transformation (Oliveira-Miranda *et al.*, 2010) and illegal wildlife trade (Sánchez-Mercado *et al.*, 2017). The UICN reports declining regional trend for 34 of the 50 Psittacidae species occurring in the country, and six species are already under some threat category in the Venezuelan Red Data Book (Rodríguez & Rojas-Suárez, 2008).

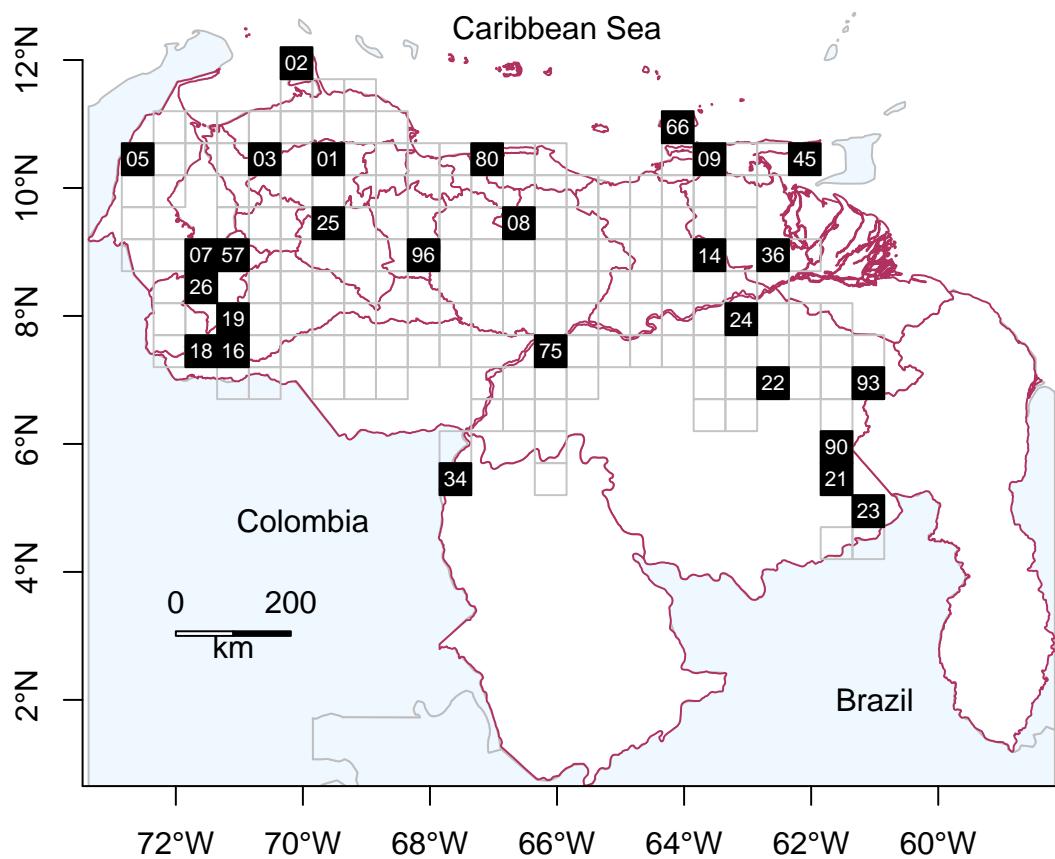
Here we provide an assessment of current distribution of Psittacid species in Venezuela using data from a national bird monitoring program carried out in 2010 as a component of the Neotropical Biodiversity Mapping Initiative (NeoMaps Ferrer-Paris *et al.*, 2013a; Rodríguez & Sharpe, 2002; Rodríguez *et al.*, 2012).

2 Methods

2.1 Distribution and monitoring data

Information on bird species present in Venezuela is taken from (Hilty, 2003). Range maps for all these species were obtained from BirdLife International (BirdLife International & Nature-Serve, 2014). Distribution records were retrieved from the Global Biodiversity Information Facility (GBIF.org, 2016).

NeoMaps bird survey was completed between March and April 2010 (Rodríguez *et al.*, 2012). Sampling universe consisted in 170 half-degree cells defined in the Venezuelan Biodiversity Grid, which cover over half of the country, but does not include the southern forest regions. Twenty seven cells were selected using a stratified sampling design based on environmental and biogeographical variables. Number in the following figure refer to NeoMaps transect codes (NM01 … NM93).



Standardized field sampling protocols for birds was implemented along a 40 km roadside transect within each cell. Two surveys were performed during two consecutive days in each transect: on the first day, 3-min point counts were performed at 50 stops, 800m apart. On the second day, cumulative species lists were recorded at a selection of 10 stops sampled for 9 min each, divided into three consecutive 3-min periods. Total sampling effort was 108 hours

of bird surveys (Rodríguez *et al.*, 2011; Rodríguez *et al.*, 2012)

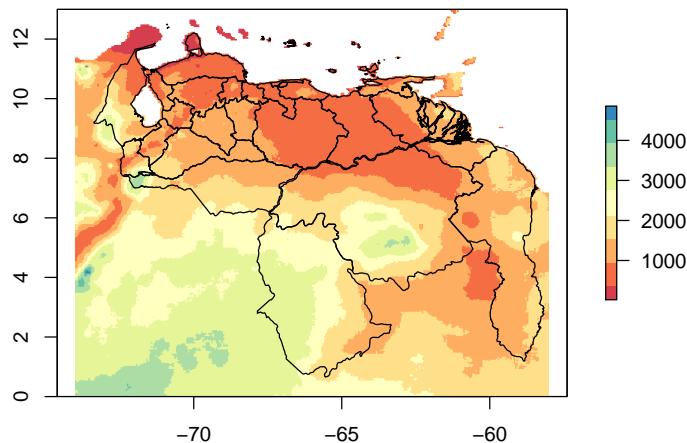
For this analysis we built detection histories for each psittacids species recorded by NeoMapas. We considered each stop as a “site” (i ; 1351 sites, 50 stops across 27 transects), and each timed survey period of 3 min as a “observation” (j), with duration $d = 3\text{min}$. For the first day survey, detections were recorded as “1” and lack of detections as “0”. For the cumulative list of the second day the detection history was filled with “0” until the first detection, and with null values (N) afterward. Thus valid detection histories for the second day are 1NN, 01N, 001 and 000, or NNN if the site was not visited on the second day (Ferrer-Paris *et al.*, 2013b). Time of the day was used as an observation covariate. Sites covariates were extracted from the spatial location of each site.

2.2 Sites covariates

In order to get representative data on climatic and vegetation condition at the time of the survey, we matched the location and date of each observations with time-series of environmental variables derived from the Moderate Resolution Radio Spectrometer (MODIS) sensors in Terra-Satellites and queried using the global MODIS Subsetting Tool (Land Processes Distributed Active Archive Center), and the Climate Hazards Group InfraRed Precipitation with Station data archive (CHIRPS version 2.0). We calculated the representative value of the variable for the year prior to the sampling time (approx. march 2009 - march 2010). We used the following variables:

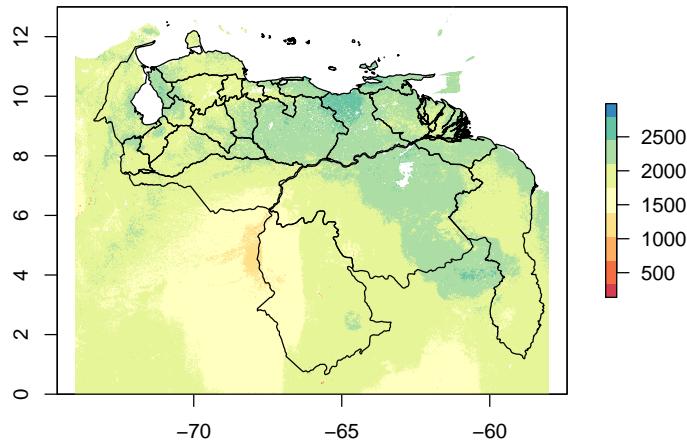
2.2.1 Total annual precipitation

Total precipitation of the year prior to NM sampling according to Chirps v 2.0(Funk *et al.*, 2015, pre01)



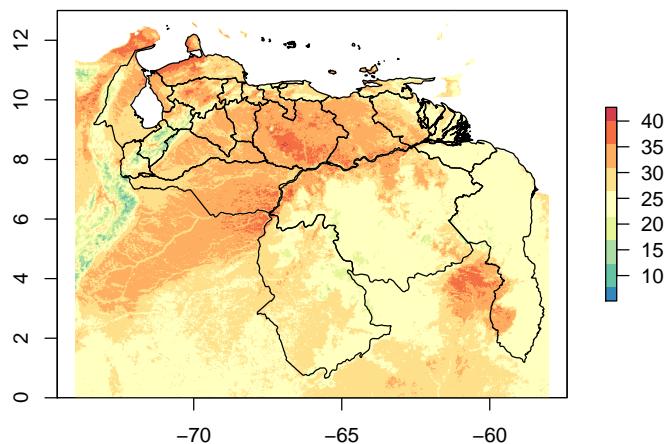
2.2.2 Potential Evapotranspiration

Total Potential Evapotranspiration of the year prior to NM (*Mu et al.*, 2011, pet01).



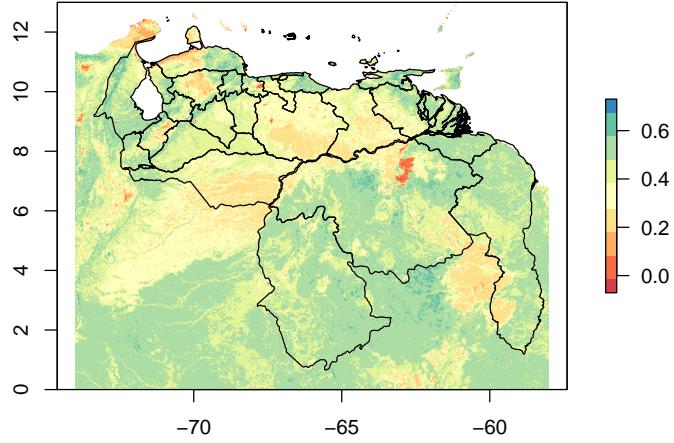
2.2.3 Land Surface Temperature

Mean LST of the year prior to NM (*Wan et al.*, 2004, dT01).



2.2.4 Enhanced Vegetation Index

Mean EVI of the year prior to NM sampling (*Huete et al.*, 2002, evi01), :



2.3 Occupancy models

We used a single-season occupancy model based on zero-inflated binomial models (MacKenzie *et al.*, 2006) to estimate a probability of occurrence for species detected in the surveys (Ψ). The occupancy state (z_i) of site i was modeled as $z_i \sim \text{Bernoulli}(\Psi_i)$, while the observation process was modeled as $y_{ij}|z_i \sim \text{Bernoulli}(z_i * p_{ij})$ in which p_{ij} represented site and occasion specific detection probability. Covariates of Ψ_i (site covariates) and p_{ij} (observation covariates) were modeled using the logit link.

We fitted eight models representing different combinations of covariates for probability of detection and probability of occurrence:

| Name | Detection | Presence |
|----------------|------------|---|
| nulo | $p \sim 1$ | $\Psi \sim 1$ |
| $p(h)Ps_i(.)$ | $p \sim h$ | $\Psi \sim 1$ |
| $p(.)Ps_i(V)$ | $p \sim 1$ | $\Psi \sim \text{evi01} + \text{evi01}^2$ |
| $p(h)Ps_i(V)$ | $p \sim h$ | $\Psi \sim \text{evi01} + \text{evi01}^2$ |
| $p(.)Ps_i(C)$ | $p \sim 1$ | $\Psi \sim dT01 + dT01^2 + pre01 + pre01^2 + pet01 + pet01^2$ |
| $p(h)Ps_i(C)$ | $p \sim h$ | $\Psi \sim dT01 + dT01^2 + pre01 + pre01^2 + pet01 + pet01^2$ |
| $p(.)Ps_i(VC)$ | $p \sim h$ | $\Psi \sim \text{evi01} + \text{evi01}^2 + dT01 + dT01^2 + pre01 + pre01^2 + pet01 + pet01^2$ |
| $p(h)Ps_i(VC)$ | $p \sim h$ | $\Psi \sim \text{evi01} + \text{evi01}^2 + dT01 + dT01^2 + pre01 + pre01^2 + pet01 + pet01^2$ |

We evaluated the individual performance of each model using the corrected Akaike Information Criterion (AICc). Then, we used the model with the best performance for each species to explain the lack of detections across the survey sites. For the sites without detections, we calculated the conditional probability of occurrence given that the species was not detected using empirical bayes methods (MacKenzie *et al.*, 2006). This probability (Ψ_{condl}) considers two components: whether sampling effort was enough to detect the species at least

once conditional on its presence ($p* = 1 - \Pi(1 - p)$), and the unconditional probability of occurrence given the values of the site covariates ($\hat{\Psi}$).

We used the unmarked, raster, and AICcmodavg packages of *R* to fit the models (Fiske & Chandler, 2011; Hijmans, 2017; Mazerolle, 2017; R Core Team, 2017)

3 Species accounts

3.1 Genus *Amazona*

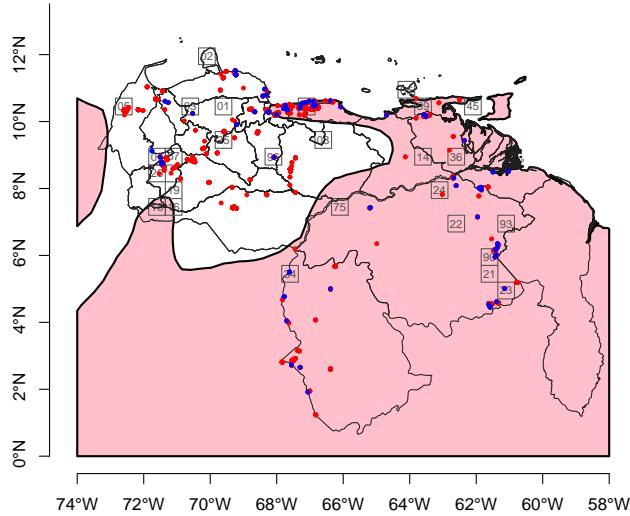
The genus *Amazona* is represented in Venezuela by eight species. The following table show species names and acronyms used in this text, with the number of distribution for each species according to the Global Biodiversity Information Facility (GBIF), and the number of records from the 2008-2012 period (GBIF.2010), and the number of detection in NeoMaps first day sampling (NM.M1), and the additional detections in the three sampling periods of the second day (NM.L1,NM.L2,NM.L3).

| | aspp | GBIF | GBIF.2010 | NM.M1 | NM.L1 | NM.L2 | NM.L3 |
|-----------------------------|-----------|------|-----------|-------|-------|-------|-------|
| <i>Amazona amazonica</i> | Amaz_amaz | 547 | 120 | 75 | 13 | 6 | 7 |
| <i>Amazona autumnalis</i> | Amaz_autu | 12 | 3 | NA | NA | NA | NA |
| <i>Amazona barbadensis</i> | Amaz_barb | 40 | 12 | 3 | 2 | 0 | 1 |
| <i>Amazona bodini</i> | Amaz_bodi | 10 | 0 | NA | NA | NA | NA |
| <i>Amazona dufresniana</i> | Amaz_dufr | 36 | 7 | NA | NA | NA | NA |
| <i>Amazona farinosa</i> | Amaz_fari | 212 | 46 | 12 | 3 | 1 | 1 |
| <i>Amazona mercenarius</i> | Amaz_merc | 13 | 1 | NA | NA | NA | NA |
| <i>Amazona ochrocephala</i> | Amaz_ochr | 719 | 264 | 108 | 33 | 4 | 5 |

Only four species were detected during NeoMaps surveys in 2010, but the four undetected species also had very few GBIF records in this time period. Please refer to Ferrer-Paris *et al.* (2013b,c) for alternative model parametrization and detailed data for this genus.

3.1.1 *Amazona amazonica*

Amazona amazonica is a widespread species. The following maps shows the expected distribution in Venezuela and surrounding countries according to BirdLife polygon maps (pink polygons) and curated distribution records from GBIF for the time period of 2008-2012. Blue dots represents detection of the species and red dots represent detection of other species of Psittacidae, but lack of detection of the target species.

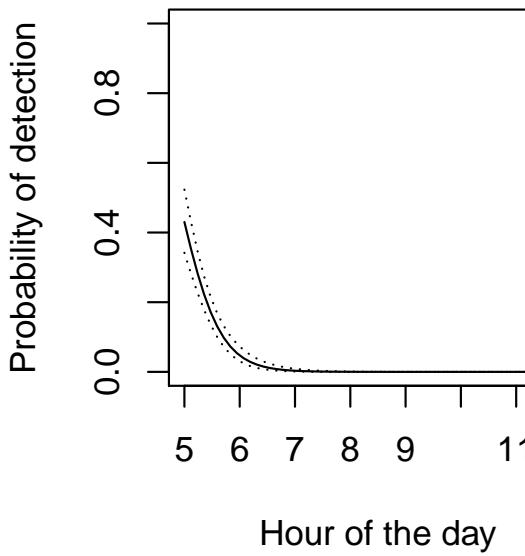


Some GBIF records are found outside the polygon of expected distribution. This can be an indication of incomplete representation of the known distribution in BirdLife maps, or taxonomic uncertainty in identification of GBIF records.

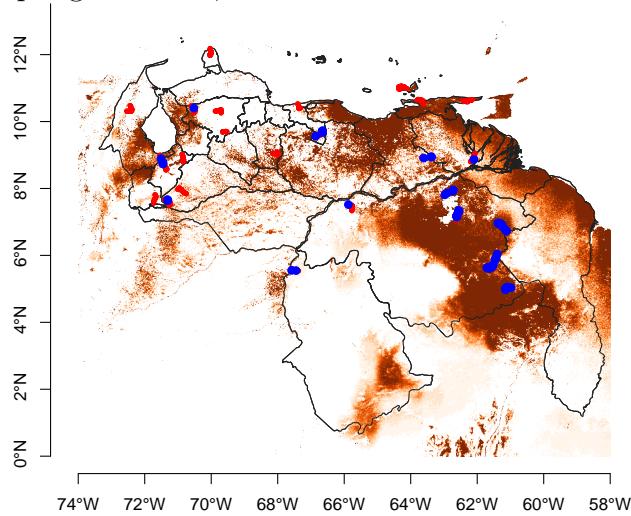
For this species we could fit several alternative occupancy models based on the large number of detections during the 2010 field survey data in Venezuela. We compare these models in the following table, where mod is the model description, n is the sample size (number of localities within its expected distribution or with evidence of presence), dtt is the number of detections, AICc is the corrected Akaike Information Criteria, Delta.AICc is the difference in AICc to the model with lowest AICc, AICw are the Akaike weights and LL is the log likelihood.

| | spp | mod | n | dtt | AICc | Delta.AICc | AICw | LL |
|---|-----------|-------------|-----|-----|----------|------------|-------|---------|
| 1 | Amaz_amaz | p(h)Psi(VC) | 900 | 97 | 553.7420 | 0.00 | 0.999 | -265.72 |
| 2 | Amaz_amaz | p(h)Psi(C) | 900 | 97 | 567.4946 | 13.75 | 0.001 | -274.65 |
| 3 | Amaz_amaz | p(.)Psi(VC) | 900 | 97 | 673.6545 | 119.91 | 0.000 | -326.70 |
| 4 | Amaz_amaz | p(.)Psi(C) | 900 | 97 | 696.9515 | 143.21 | 0.000 | -340.39 |
| 5 | Amaz_amaz | p(.)Psi(V) | 900 | 97 | 712.8495 | 159.11 | 0.000 | -352.40 |
| 6 | Amaz_amaz | nulo | 900 | 97 | 726.6031 | 172.86 | 0.000 | -361.29 |

For this species the p(h)Psi(VC) model had the highest support according to the AIC weights. This model corresponds to a time-dependent probability of detection and a effect of vegetation and climatic conditions on the probability of presence. The following figure shows a weighted estimate of probability of detection per hour of the day for the two models with $AICw > 0$.

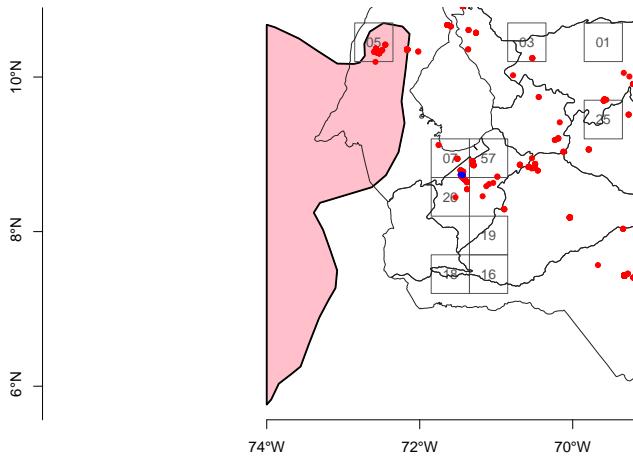


The following map shows the predicted (unconditional) probability of presence for the whole country based on the model with highest support and the values of the vegetation and climatic covariates. Darker colors indicate higher probabilities, dots represent NeoMaps sampling localities, blue dots indicate detections and red dots indicate lack of detections.



3.1.2 *Amazona autumnalis*

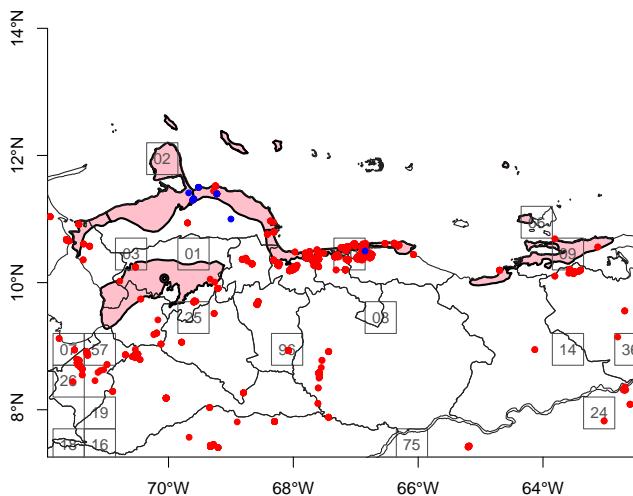
This species was expected in 50 sampling points from 'Rosario de Perijá', Zulia state (NeoMaps route NM05). However this species was not detected during the field work in 2010.



Known GBIF records are found outside the polygon of expected distribution. This can be an indication of incomplete representation of the known distribution in BirdLife maps, or taxonomic uncertainty in identification of GBIF records.

3.1.3 *Amazona barbadensis*

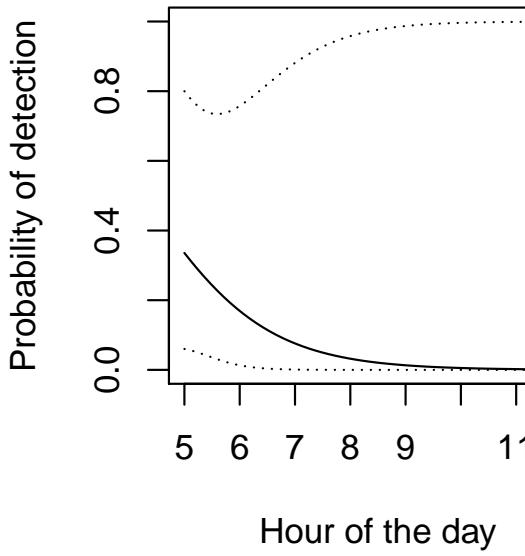
Amazona barbadensis is a species of restricted distribution in Venezuela and the Caribbean.



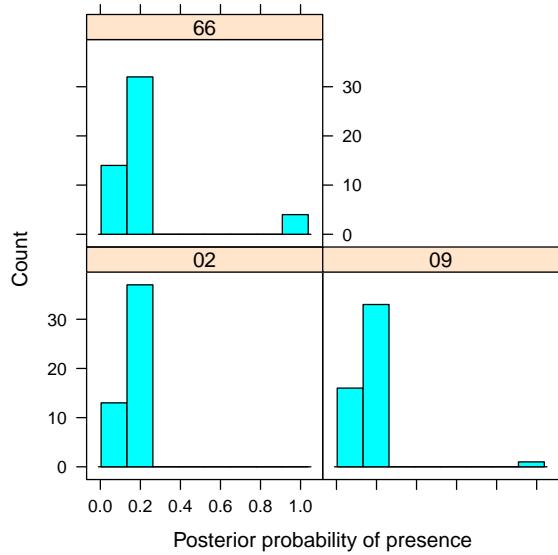
During NeoMaps field surveys in 2010, *Amazona barbadensis* was detected in few localities, and due to small sample size, only two models could be fitted, both have similar support according to AIC weights.

| | spp | mod | n | dtt | AICc | Delta.AICc | AICw | LL |
|---|-----------|------------|-----|-----|----------|------------|-------|--------|
| 7 | Amaz_barb | p(h)Psi(.) | 150 | 5 | 55.07084 | 0.00 | 0.571 | -24.45 |
| 8 | Amaz_barb | nulo | 150 | 5 | 55.64423 | 0.57 | 0.429 | -25.78 |

The following figure shows a weighted estimate of probability of detection per hour of the day for the two models with $AIC_w > 0$.



Unconditional probability of presence is relatively low (around 0, 2). For the three NeoMaps routes were the species was expected, presence was confirmed in very few localities (localities of detection), and true absence can be suspected in around 18% of the sampling localities were the sampling effort was enough to achieve low values of the posterior or conditional probability of presence ($\Psi_{post} < 0, 12$).

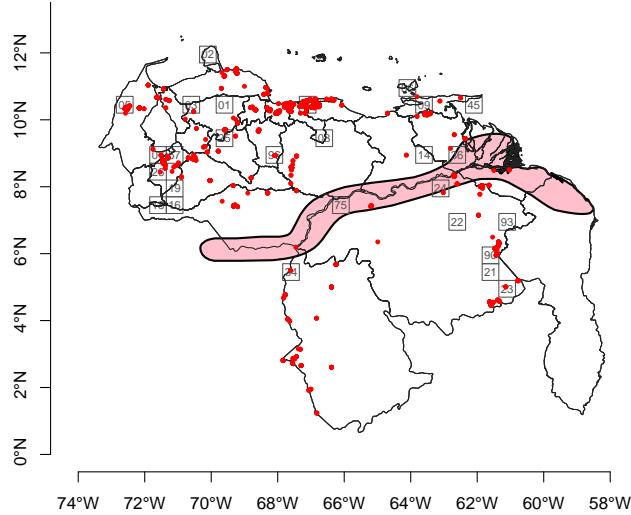


3.1.4 *Amazona bodini*

Amazona bodini is a species of restricted distribution in Venezuela. It was not detected by NeoMaps field work, and is not represented in GBIF record from the 2008-2012 period.

Amazona bodini was expected in 102 sampling points from 'Isla de Guara', Monagas-Delta

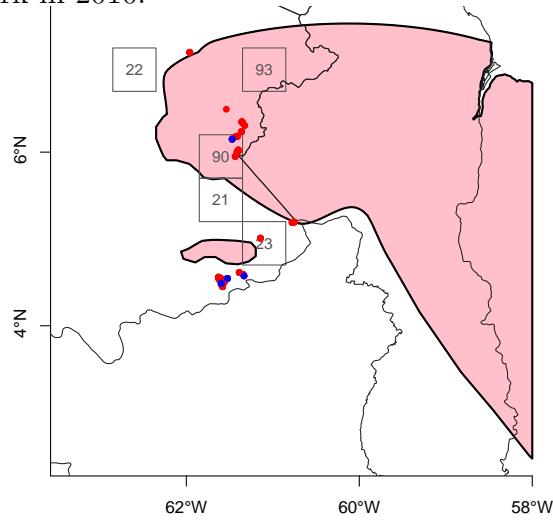
state (NeoMaps route NM36); 'Caicara del Orinoco', Bolívar state (NeoMaps route NM75). However this species was not detected during the field work in 2010.



3.1.5 *Amazona dufresniana*

Amazona dufresniana is a species of restricted distribution in Venezuela. It was not detected by NeoMaps field work, and has few GBIF record from the 2008-2012 period.

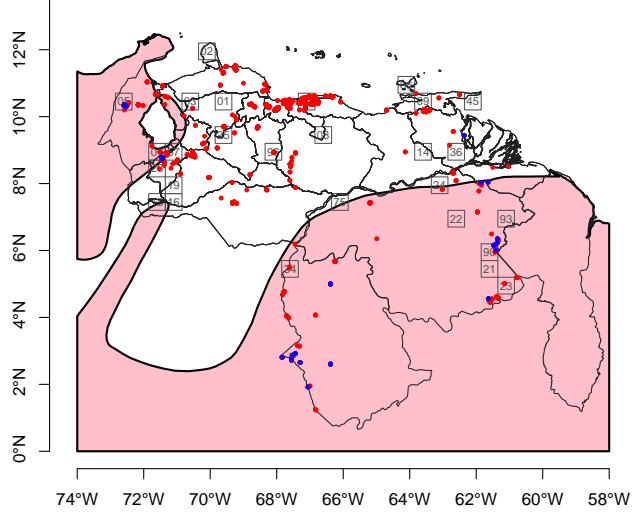
Amazona dufresniana was expected in 109 sampling points from 'Anacoco', Bolívar state (NeoMaps route NM93); 'La Escalera', Bolívar state (NeoMaps route NM90); 'Kavanayén', Bolívar state (NeoMaps route NM21). However this species was not detected during the field work in 2010.



3.1.6 *Amazona farinosa*

Amazona farinosa has a non-continuous distribution in Venezuela and was expected in 482 sampling points from 'Rosario de Perijá', Zulia state (NeoMaps route NM05); 'Sur del Lago', Zulia-Mérida state (NeoMaps route NM07); 'Guri', Bolívar state (NeoMaps route NM24); 'El Manteco', Bolívar state (NeoMaps route NM22); 'Caicara del Orinoco', Bolívar state

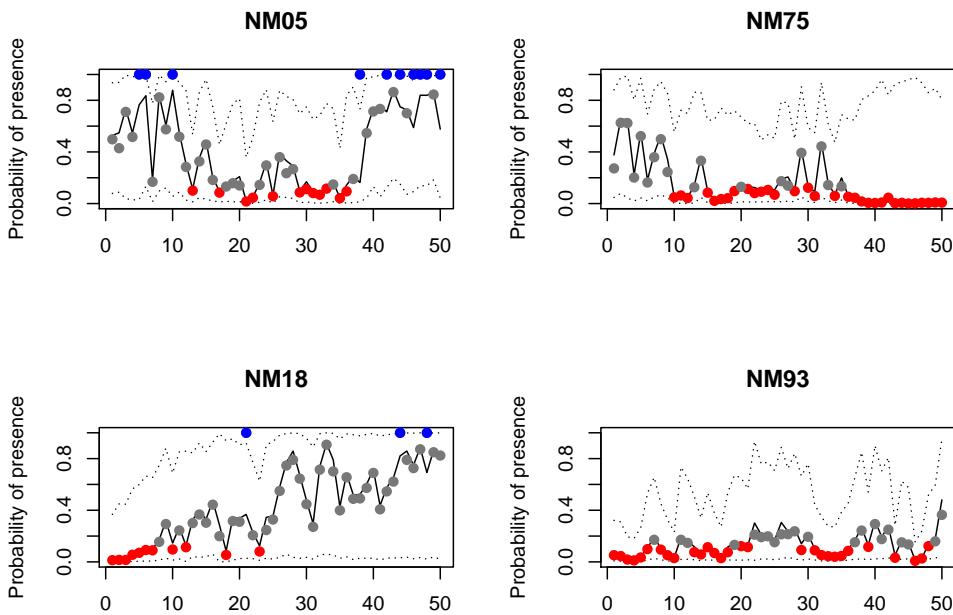
(NeoMaps route NM75); 'Gavilán', Amazonas state (NeoMaps route NM34); 'Anacoco', Bolívar state (NeoMaps route NM93); 'La Escalera', Bolívar state (NeoMaps route NM90); 'Kavanayén', Bolívar state (NeoMaps route NM21); 'Paraytepuy', Bolívar state (NeoMaps route NM23).



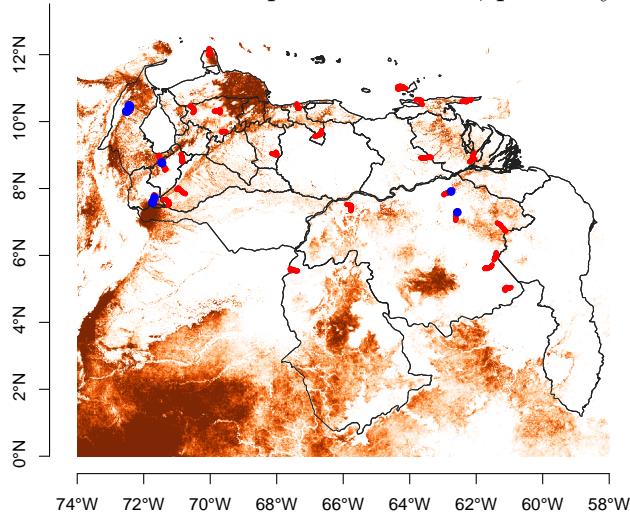
Available data allowed to fit models with constant probability of detection. The model with highest support according to AIC weights includes climatic and vegetation condition as covariates.

| | spp | mod | n | dtt | AICc | Delta.AICc | AICw | LL |
|----|-----------|-------------|-----|-----|----------|------------|-------|--------|
| 9 | Amaz_fari | p(.)Psi(VC) | 550 | 17 | 158.5461 | 0.00 | 0.940 | -69.07 |
| 10 | Amaz_fari | p(.)Psi(V) | 550 | 17 | 164.0841 | 5.54 | 0.059 | -78.01 |
| 11 | Amaz_fari | p(.)Psi(C) | 550 | 17 | 171.3484 | 12.80 | 0.002 | -77.54 |

According to the model with the best support the unconditional probability of presence varied along the NeoMaps routes depending on the value of the covariates. The figure show four examples of 40 km routes, ordered from beginning to end, two with detections (NM05 and NM18) and two without detections (NM75 and NM93). The lines show the predicted unconditional probability of presence (solid line: best estimate, dotted lines 95% confidence interval), and the dots represent the conditional probability given the observed detection history. Blue dots represent the localities of known occurrences, red dots are localities with very low posterior probabilities of presence ($\Psi_{post} < 0.125$), grey dots are localities with intermediate values.

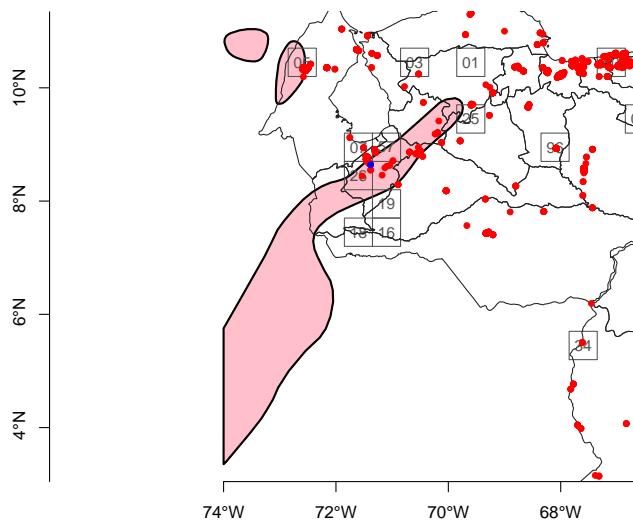


The predicted (unconditional) probability of presence for the whole country does reflect the expected distribution, but predicts high probability of presence in areas of Falcón and Lara state where the species is absent, probably due to biogeographic constrains.



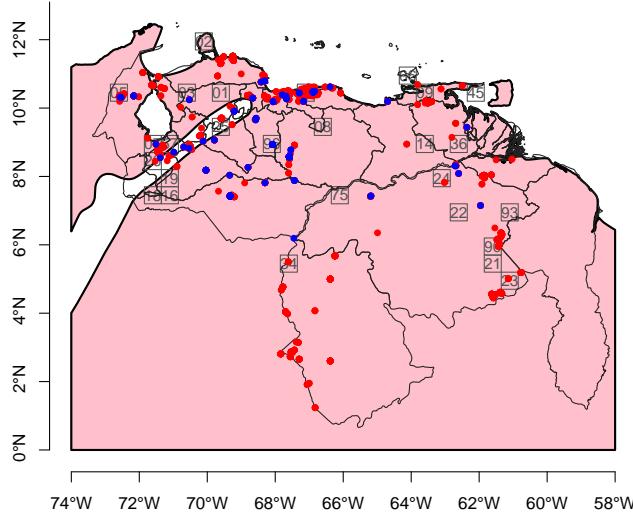
3.1.7 *Amazona mercenarius*

Amazona mercenarius was expected in 103 sampling points from 'Jají', Mérida state (NeoMaps route NM26); 'Piñango', Mérida state (NeoMaps route NM57). However this species was not detected during the field work in 2010.



3.1.8 *Amazona ochrocephala*

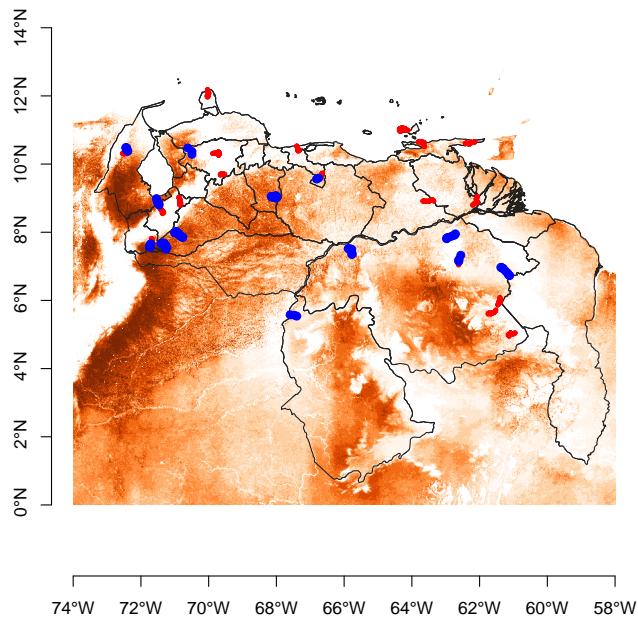
Amazona ochrocephala has a widespread distribution in Venezuela and was detected in several NeoMaps transects.



Available data allowed to fit several models with constant probability of detection. The model with highest support according to AIC weights includes climatic and vegetation condition as covariates.

| | spp | mod | n | dtt | AICc | Delta.AICc | AICw | LL |
|----|-----------|-------------|------|-----|-----------|------------|------|---------|
| 12 | Amaz_ochr | p(.)Psi(VC) | 1250 | 143 | 888.5061 | 0.00 | 1 | -434.16 |
| 13 | Amaz_ochr | p(.)Psi(C) | 1250 | 143 | 916.9785 | 28.47 | 0 | -450.43 |
| 14 | Amaz_ochr | p(.)Psi(V) | 1250 | 143 | 987.4805 | 98.97 | 0 | -489.72 |
| 15 | Amaz_ochr | nulo | 1250 | 143 | 1042.1294 | 153.62 | 0 | -519.06 |

The predicted (unconditional) probability of presence for the whole country does reflect the expected distribution.



3.2 Genus Ara

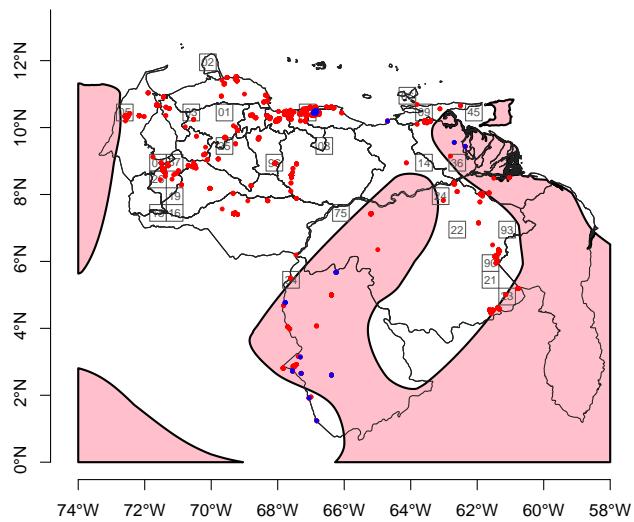
The genus Ara is represented in Venezuela by five species.

| | aspp | GBIF | GBIF.2010 | NM.M1 | NM.L1 | NM.L2 | NM.L3 |
|------------------|----------|------|-----------|-------|-------|-------|-------|
| Ara ararauna | Ara_arar | 140 | | 56 | NA | NA | NA |
| Ara chloropterus | Ara_chlo | 271 | | 52 | 4 | 4 | 4 |
| Ara macao | Ara_maca | 290 | | 117 | 6 | 1 | 1 |
| Ara militaris | Ara_mili | 80 | | 18 | 6 | 0 | 2 |
| Ara severus | Ara_seve | 457 | | 217 | 48 | 11 | 3 |

Four species were detected during NeoMaps surveys in 2010, the only exception was (*Ara ararauna*).

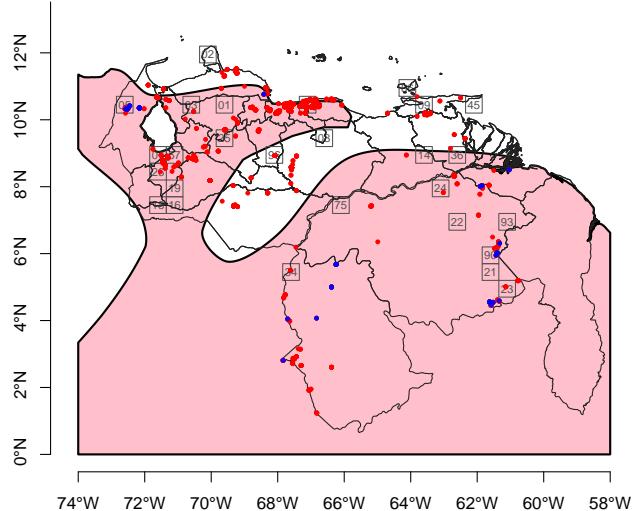
3.2.1 *Ara ararauna*

Ara ararauna was expected in 112 sampling points from 'Isla de Guara', Monagas-Delta state (NeoMaps route NM36); 'Gavilán', Amazonas state (NeoMaps route NM34); 'Paraytepuy', Bolívar state (NeoMaps route NM23). However this species was not detected during the field work in 2010.



3.2.2 *Ara chloropterus*

Ara chloropterus has a widespread distribution and was expected in 1045 sampling points from NeoMaps, but has few actual detections.



Available data allowed to fit models with constant probability of detection and variable probability of presence.

| | spp | mod | n | dtt | AICc | Delta.AICc | AICw | LL |
|----|----------|------------|------|-----|----------|------------|-------|--------|
| 16 | Ara_chlo | p(h)Psi(.) | 1050 | 13 | 149.4942 | 0.00 | 0.964 | -71.74 |
| 17 | Ara_chlo | nulo | 1050 | 13 | 156.0788 | 6.58 | 0.036 | -76.03 |

However, the model has a poor fit with a combination of high probability of occurrence and low probability of detection and too wide standard errors for the estimated parameters:

```
Call:  
occu(formula = ~hora ~ 1, data = UMF[os, ])
```

```

Occupancy (logit-scale):
  Estimate    SE      z P(>|z|)
  3.99 25.7 0.155  0.877

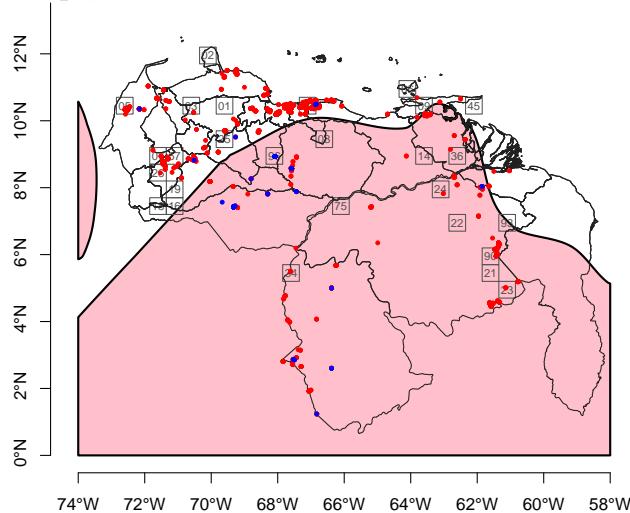
Detection (logit-scale):
  Estimate    SE      z P(>|z|)
(Intercept) -0.291 1.74 -0.168  0.8666
hora         -13.706 5.34 -2.568  0.0102

AIC: 149.4713
Number of sites: 1050
optim convergence code: 0
optim iterations: 63
Bootstrap iterations: 0

```

3.2.3 *Ara macao*

Ara macao has a widespread distribution and was expected in 607 sampling points from NeoMaps, but has few actual detections.



Available data allowed to fit several alternative models including covariates in probability of presence.

| | spp | mod | n | dtt | AICc | Delta.AICc | AICw | LL |
|----|----------|------------|-----|-----|-----------|------------|-------|--------|
| 18 | Ara_maca | p(.)Psi(C) | 600 | 8 | 79.22826 | 0.00 | 0.692 | -31.49 |
| 19 | Ara_maca | p(h)Psi(C) | 600 | 8 | 80.85336 | 1.63 | 0.307 | -31.27 |
| 20 | Ara_maca | p(.)Psi(V) | 600 | 8 | 93.83741 | 14.61 | 0.000 | -42.89 |
| 21 | Ara_maca | nulo | 600 | 8 | 100.99736 | 21.77 | 0.000 | -48.49 |
| 22 | Ara_maca | p(h)Psi(.) | 600 | 8 | 102.79301 | 23.56 | 0.000 | -48.38 |

The best model includes constant detection probability and effect on climatic covariates, but parameter values suggest a poor fit and high uncertainty.

```

Call:
occu(formula = ~1 ~ pet01 + I(pet01^2) + dT01 + I(dT01^2) + pre01 +
  I(pre01^2), data = UMF[os, ])

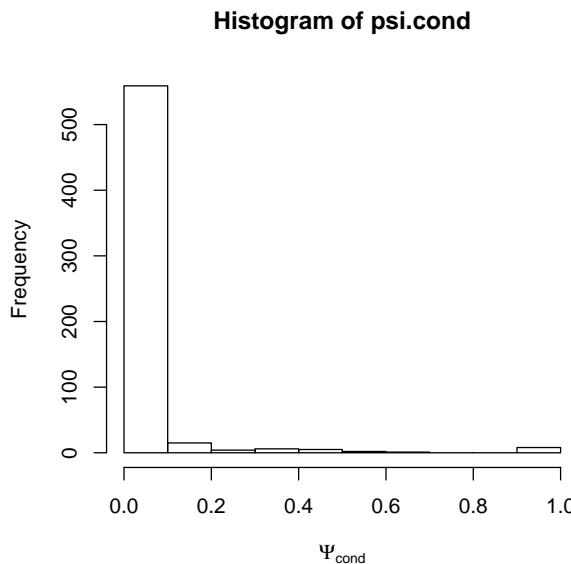
Occupancy (logit-scale):
      Estimate     SE      z P(>|z|)
(Intercept) -4.7783  2.11 -2.2634  0.0236
pet01        2.4933  2.23  1.1202  0.2626
I(pet01^2)   0.0817  1.13  0.0722  0.9424
dT01         6.0089  4.41  1.3627  0.1730
I(dT01^2)   -2.5151  2.85 -0.8835  0.3770
pre01        0.8769 11.11  0.0789  0.9371
I(pre01^2) -16.0437 24.46 -0.6560  0.5118

Detection (logit-scale):
      Estimate     SE      z P(>|z|)
-1.04 0.775 -1.35  0.179

AIC: 78.9846
Number of sites: 600
optim convergence code: 0
optim iterations: 74
Bootstrap iterations: 0

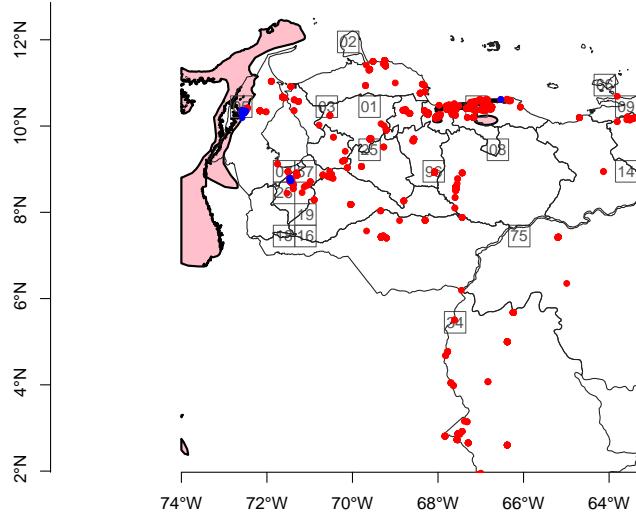
```

Unconditional probability of presence is relatively low in most of the sampling area, and presence was confirmed in very few localities (localities of detection).



3.2.4 *Ara militaris*

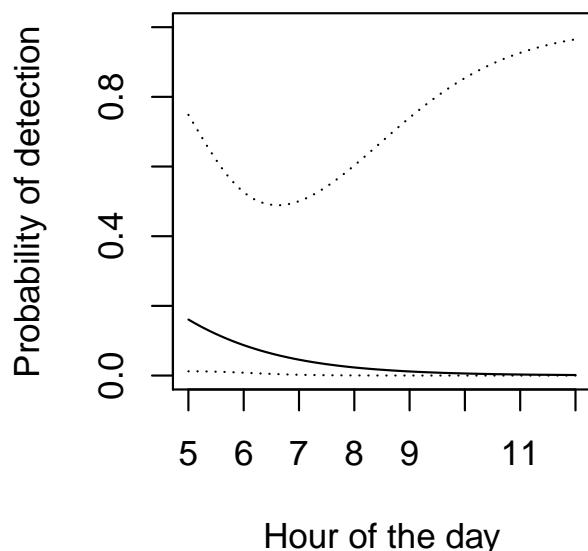
Ara militaris is found in the Andean and Coastal mountain ranges of Venezuela, and was expected in 35 sampling points from 'Rosario de Perijá', Zulia state (NeoMaps route NM05); 'Colonia Tovar', Aragua-Vargas state (NeoMaps route NM80).



Available data allowed to fit some alternative models with similar AICc values.

| | spp | mod | n | dtt | AICc | Delta.AICc | AICw | LL |
|----|----------|------------|-----|-----|----------|------------|-------|--------|
| 23 | Ara_mili | p(h)Psi(V) | 150 | 8 | 71.80744 | 0.00 | 0.365 | -30.70 |
| 24 | Ara_mili | p(.)Psi(V) | 150 | 8 | 72.38796 | 0.58 | 0.273 | -32.06 |
| 25 | Ara_mili | p(h)Psi(.) | 150 | 8 | 72.60602 | 0.80 | 0.245 | -33.22 |
| 26 | Ara_mili | nulo | 150 | 8 | 74.09608 | 2.29 | 0.116 | -35.01 |

The best model includes an effect of time of day on the detection probability.



Parameter estimates have high values and high uncertainty.

```

Call:
occu(formula = ~hora ~ evi01 + I(evi01^2), data = UMF[os, ])

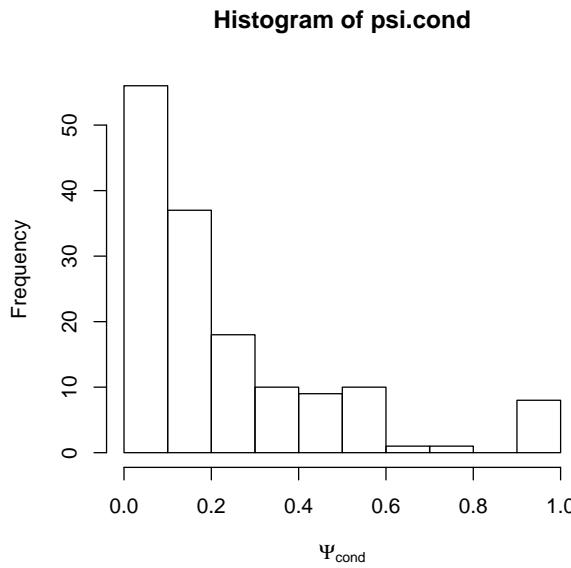
Occupancy (logit-scale):
            Estimate      SE      z P(>|z|)
(Intercept) -3.388 2.89 -1.173 0.241
evi01        0.996 5.48  0.182 0.856
I(evi01^2)   0.761 2.49  0.305 0.760

Detection (logit-scale):
            Estimate      SE      z P(>|z|)
(Intercept)  1.95 2.66  0.732 0.464
hora        -10.59 6.90 -1.534 0.125

AIC: 71.39078
Number of sites: 150
optim convergence code: 0
optim iterations: 41
Bootstrap iterations: 0

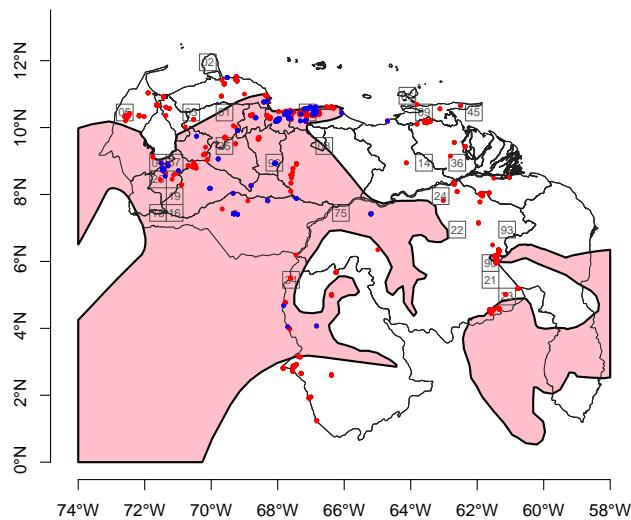
```

Unconditional probability of presence is relatively low in most of the sampling area, and presence was confirmed in very few localities (localities of detection).



3.2.5 *Ara severus*

Ara severus has a widespread but non continuous distribution and was expected in 642 sampling points from NeoMaps.



Available data allowed to fit models with constant probability of detection and probability of presence.

| | spp | mod | n | dtt | AICc | Delta.AICc | AICw | LL |
|----|----------|------------|-----|-----|----------|------------|------|---------|
| 27 | Ara_seve | p(.)Psi(V) | 700 | 61 | 450.5242 | 0.00 | 1 | -221.23 |
| 28 | Ara_seve | nulo | 700 | 61 | 482.0490 | 31.52 | 0 | -239.02 |

Parameter estimates have high values and high uncertainty.

Call:

```
occu(formula = ~1 ~ evi01 + I(evi01^2), data = UMF[os, ])
```

Occupancy (logit-scale):

| | Estimate | SE | z | P(> z) |
|-------------|----------|-------|--------|---------|
| (Intercept) | 4.22 | 6.21 | 0.679 | 0.497 |
| evi01 | 11.59 | 10.41 | 1.114 | 0.265 |
| I(evi01^2) | -7.84 | 8.13 | -0.964 | 0.335 |

Detection (logit-scale):

| Estimate | SE | z | P(> z) |
|----------|-------|-------|----------|
| -2.37 | 0.145 | -16.3 | 1.25e-59 |

AIC: 450.4667

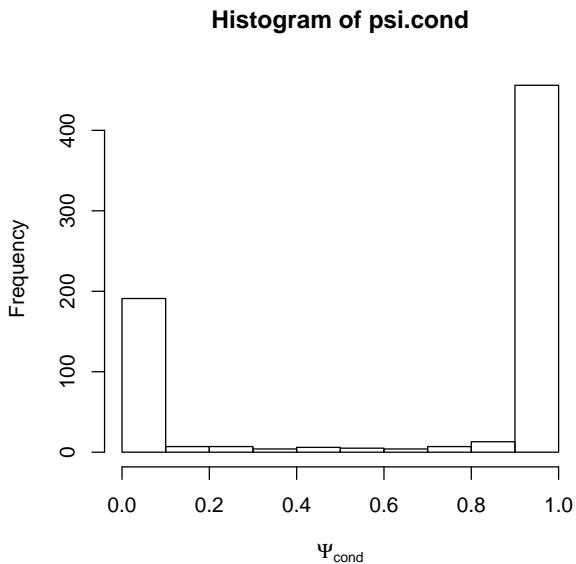
Number of sites: 700

optim convergence code: 0

optim iterations: 81

Bootstrap iterations: 0

The combination of parameter values results in extreme predictions of conditional probability of presence within the sampling area.



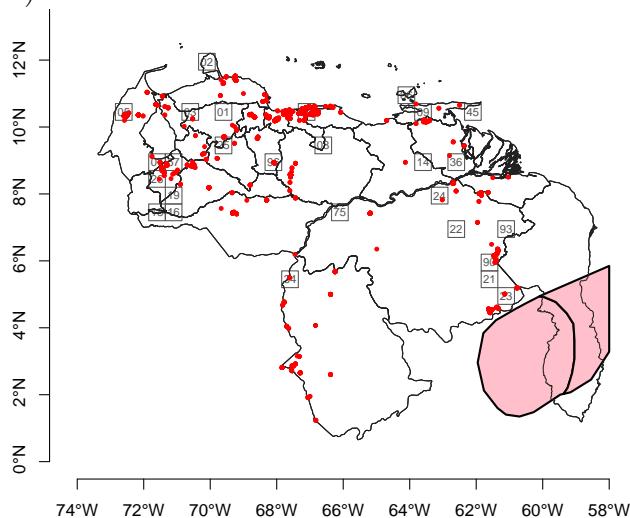
3.3 Genus Aratinga

Evidence for the presence of the genus *Aratinga* in Venezuela is scarce. One species is suspected to be present, but reliable records are missing.

| | aspp | GBIF | GBIF.2010 | NM.M1 | NM.L1 | NM.L2 | NM.L3 |
|------------------------------|-----------|------|-----------|-------|-------|-------|-------|
| <i>Aratinga solstitialis</i> | Arat_sols | 1 | 0 | NA | NA | NA | NA |

3.3.1 *Aratinga solstitialis*

The distribution of *Aratinga solstitialis* probably includes part of the Guayana Esequiba, but taxonomic and distribution information for this taxon needs further review (Silveira *et al.*, 2005).

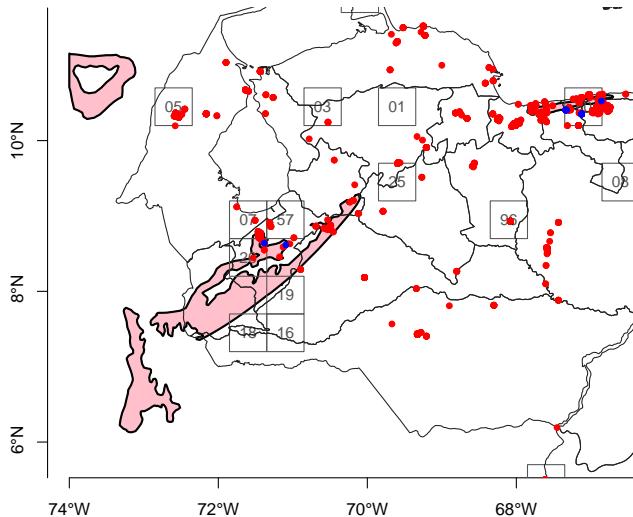


3.4 Genus *Bolborhynchus*, species *B. lineola*

The genus *Bolborhynchus* is represented in Venezuela by a single species.

| | aspp | GBIF | GBIF.2010 | NM.M1 | NM.L1 | NM.L2 | NM.L3 |
|------------------------------|-----------|------|-----------|-------|-------|-------|-------|
| <i>Bolborhynchus lineola</i> | Bolb_line | 30 | | 22 | NA | NA | NA |

Bolborhynchus lineola is a species of restricted distribution in Venezuela. *Bolborhynchus lineola* was expected in 96 sampling points from 'Sur del Lago', Zulia-Mérida state (NeoMaps route NM07); 'Jají', Mérida state (NeoMaps route NM26); 'San Joaquín de Navay', Táchira state (NeoMaps route NM18); 'Colonia Tovar', Aragua-Vargas state (NeoMaps route NM80). It was not detected by NeoMaps field work, and only has few GBIF record from the 2008-2012 period.



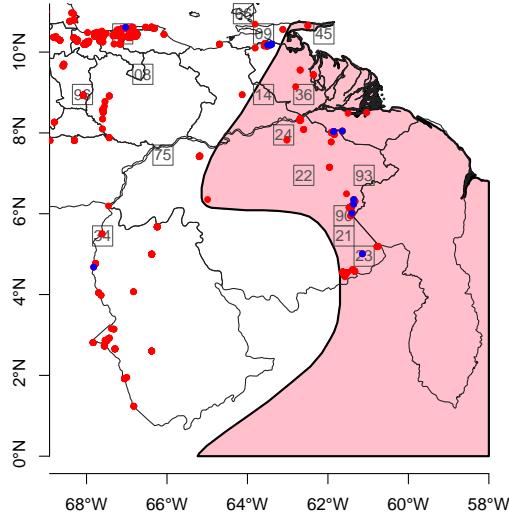
3.5 Genus *Brotogeris*

The genus *Brotogeris* is represented in Venezuela by three species, but only two were detected in NeoMaps surveys of 2010.

| | aspp | GBIF | GBIF.2010 | NM.M1 | NM.L1 | NM.L2 | NM.L3 | |
|-------------------------------|-----------|------|-----------|-------|-------|-------|-------|----|
| <i>Brotogeris chrysoptera</i> | Brot_chry | 51 | | 14 | 0 | 0 | 2 | 0 |
| <i>Brotogeris cyanoptera</i> | Brot_cyan | 27 | | 15 | NA | NA | NA | NA |
| <i>Brotogeris jugularis</i> | Brot_jugu | 412 | | 218 | 16 | 2 | 3 | 0 |

3.5.1 *Brotogeris chrysoptera*

Brotogeris chrysoptera is only found in eastern Venezuela, and was expected in 454 sampling points from 'Paria', Sucre state (NeoMaps route NM45); 'San Tomé', Anzoátegui-Monagas state (NeoMaps route NM14); 'Isla de Guara', Monagas-Delta state (NeoMaps route NM36); 'Guri', Bolívar state (NeoMaps route NM24); 'El Manteco', Bolívar state (NeoMaps route NM22); 'Anacoco', Bolívar state (NeoMaps route NM93); 'La Escalera', Bolívar state (NeoMaps route NM90); 'Kavanayén', Bolívar state (NeoMaps route NM21); 'Paraytepuy', Bolívar state (NeoMaps route NM23).



It was detected twice during the field work in 2010. Due to the scarce data available, only two models could be fitted. The null model had best support according to AIC weights, but a poor fit with a combination of high probability of occurrence and low probability of detection and too wide standard errors for the estimated parameters:

| | spp | mod | n | dtt | AICc | Delta.AICc | AICw | LL |
|----|-----------|------------|-----|-----|----------|------------|-------|--------|
| 40 | Brot_chry | nulo | 450 | 2 | 31.53931 | 0.00 | 0.734 | -13.76 |
| 41 | Brot_chry | p(h)Psi(.) | 450 | 2 | 33.56471 | 2.03 | 0.266 | -13.76 |

Call:
`occu(formula = ~1 ~ 1, data = UMF[os,])`

Occupancy (logit-scale):

| Estimate | SE | z | P(> z) |
|----------|----|--------|---------|
| 2.26 | 26 | 0.0869 | 0.931 |

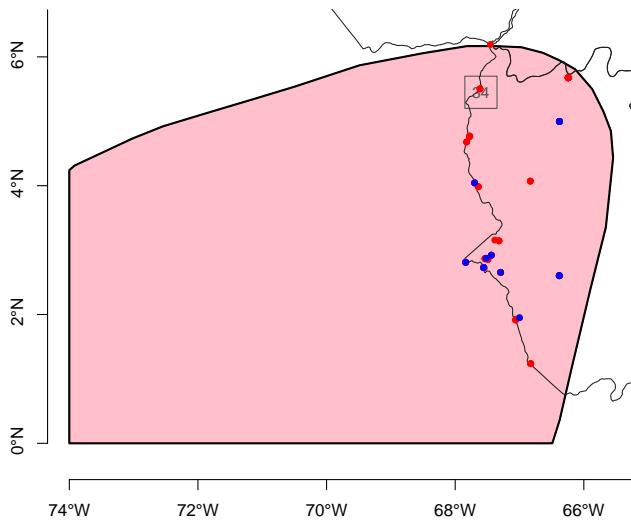
Detection (logit-scale):

| Estimate | SE | z | P(> z) |
|----------|------|-------|---------|
| -5.78 | 2.56 | -2.26 | 0.024 |

AIC: 31.51246
Number of sites: 450
optim convergence code: 0
optim iterations: 82
Bootstrap iterations: 0

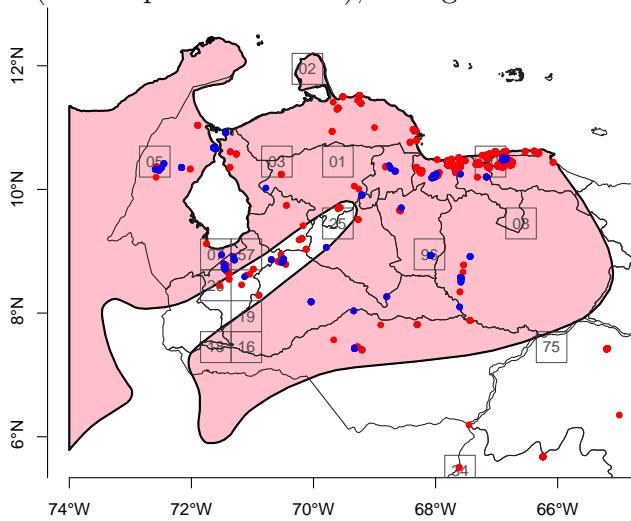
3.5.2 *Brotogeris cyanoptera*

Brotogeris cyanoptera is only found in souther Venezuela in Amazonas state, and was expected in 50 sampling points from 'Gavilán', Amazonas state (NeoMaps route NM34). It was not detected during the field work in 2010.



3.5.3 *Brotogeris jugularis*

Brotogeris jugularis is only found in western Venezuela, and was expected in 557 sampling points from 'Rosario de Perijá', Zulia state (NeoMaps route NM05); 'Quebrada Arriba', Lara-Falcón state (NeoMaps route NM03); 'Sur del Lago', Zulia-Mérida state (NeoMaps route NM07); 'Jají', Mérida state (NeoMaps route NM26); 'Piñango', Mérida state (NeoMaps route NM57); 'Capitanejo', Barinas state (NeoMaps route NM19); 'Otopún', Barinas state (NeoMaps route NM16); 'San Joaquín de Navay', Táchira state (NeoMaps route NM18); 'Paraguaná', Falcón state (NeoMaps route NM02); 'Río Tocuyo', Lara state (NeoMaps route NM01); 'Colonia Tovar', Aragua-Vargas state (NeoMaps route NM80); 'Corralito', Cojedes state (NeoMaps route NM96); 'Altagracia de Orituco', Aragua state (NeoMaps route NM08).

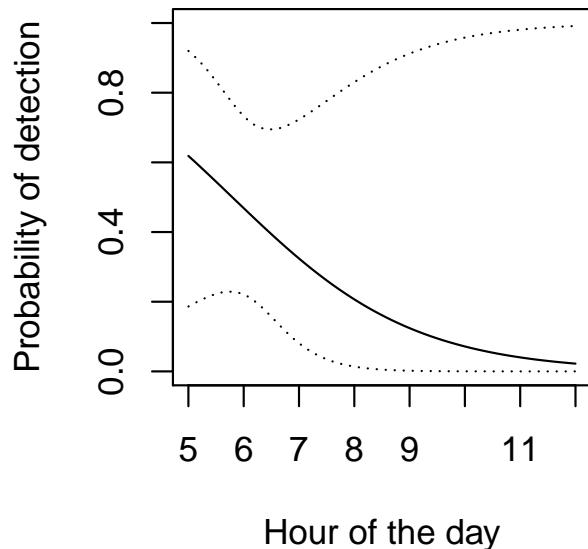


Available data allowed to fit several alternative models, and models with either climatic or vegetation covariates received considerable support according to AIC weights.

| | spp | mod | n | dtt | AICc | Delta.AICc | AICw | LL |
|----|-----------|------------|-----|-----|----------|------------|-------|--------|
| 42 | Brot_jugu | p(h)Psi(C) | 650 | 20 | 174.6283 | 0.00 | 0.551 | -78.17 |
| 43 | Brot_jugu | p(.)Psi(C) | 650 | 20 | 175.4298 | 0.80 | 0.369 | -79.60 |

| | | | | | | | | |
|----|-----------|---------------|-----|----|----------|-------|-------|--------|
| 44 | Brot_jugu | $p(.)\Psi(V)$ | 650 | 20 | 179.1705 | 4.54 | 0.057 | -85.55 |
| 45 | Brot_jugu | $p(h)\Psi(V)$ | 650 | 20 | 180.9018 | 6.27 | 0.024 | -85.40 |
| 46 | Brot_jugu | nulo | 650 | 20 | 199.9915 | 25.36 | 0.000 | -97.99 |
| 47 | Brot_jugu | $p(h)\Psi(.)$ | 650 | 20 | 200.7790 | 26.15 | 0.000 | -97.37 |

The best model includes an effect of time of day on the detection probability.



Call:

```
occu(formula = ~hora ~ pet01 + I(pet01^2) + dT01 + I(dT01^2) +
  pre01 + I(pre01^2), data = UMF[os, ])
```

Occupancy (logit-scale):

| | Estimate | SE | z | P(> z) |
|-------------|----------|-------|--------|---------|
| (Intercept) | -1.11 | 0.573 | -1.934 | 0.0531 |
| pet01 | 0.35 | 0.755 | 0.463 | 0.6433 |
| I(pet01^2) | -1.60 | 0.976 | -1.637 | 0.1016 |
| dT01 | -1.69 | 0.660 | -2.564 | 0.0103 |
| I(dT01^2) | -2.23 | 0.903 | -2.464 | 0.0137 |
| pre01 | 1.10 | 0.555 | 1.988 | 0.0468 |
| I(pre01^2) | -0.37 | 0.203 | -1.821 | 0.0685 |

Detection (logit-scale):

| | Estimate | SE | z | P(> z) |
|-------------|----------|------|-------|---------|
| (Intercept) | 3.51 | 2.42 | 1.45 | 0.147 |
| hora | -10.17 | 6.17 | -1.65 | 0.099 |

AIC: 174.3471

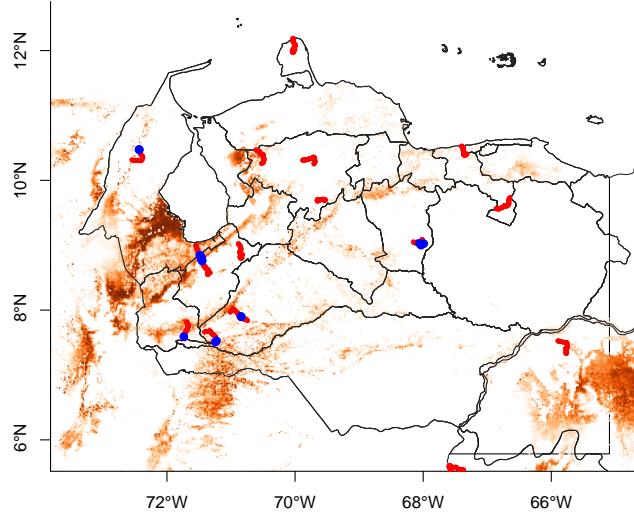
Number of sites: 650

```

optim convergence code: 0
optim iterations: 33
Bootstrap iterations: 0

```

The following map shows the predicted (unconditional) probability of presence for the whole country based on the model with highest support and the values of the climatic covariates.

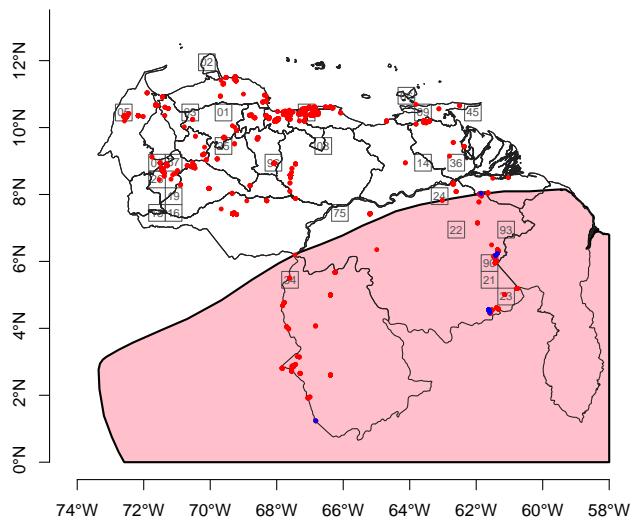


3.6 Genus *Deroptyus*, species *D. accipitrinus*

The genus *Deroptyus* is represented in Venezuela by a single species.

| | aspp | GBIF | GBIF.2010 | NM.M1 | NM.L1 | NM.L2 | NM.L3 |
|---|------|------|-----------|-------|-------|-------|-------|
| <i>Deroptyus accipitrinus</i> Dero_acci | | | 49 | 11 | NA | NA | NA |

Deroptyus accipitrinus is a species of restricted to South Venezuela, and was expected in 308 sampling points from 'Guri', Bolívar state (NeoMaps route NM24); 'El Manteco', Bolívar state (NeoMaps route NM22); 'Gavilán', Amazonas state (NeoMaps route NM34); 'Anacoco', Bolívar state (NeoMaps route NM93); 'La Escalera', Bolívar state (NeoMaps route NM90); 'Kavanayén', Bolívar state (NeoMaps route NM21); 'Paraytepuy', Bolívar state (NeoMaps route NM23). However this species was not detected during the field work in 2010, and only has few GBIF record from the 2008-2012 period.

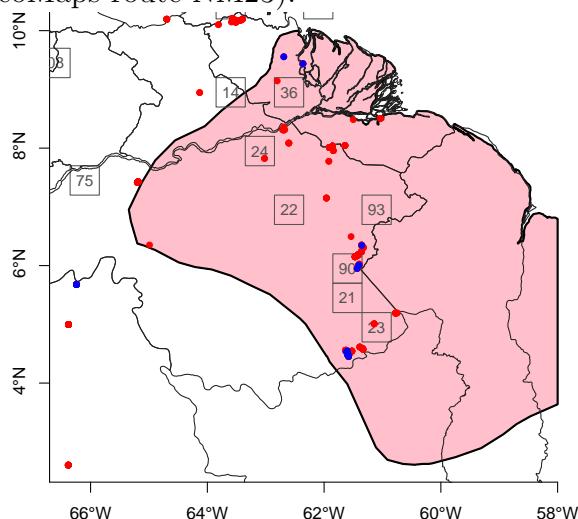


3.7 Genus *Diopsittaca*, species *D. nobilis*

This genus is represented in Venezuela by a single species.

| | aspp | GBIF | GBIF.2010 | NM.M1 | NM.L1 | NM.L2 | NM.L3 |
|--------------------------------------|------|------|-----------|-------|-------|-------|-------|
| <i>Diopsittaca nobilis</i> Diop_nobi | | 58 | | 20 | 1 | 1 | 0 |

In Venezuela *Diopsittaca nobilis* is restricted to the Guyana region south of the Orinoco river, and was expected in 361 sampling points from 'San Tomé', Anzoátegui-Monagas state (NeoMaps route NM14); 'Isla de Guara', Monagas-Delta state (NeoMaps route NM36); 'Guri', Bolívar state (NeoMaps route NM24); 'El Manteco', Bolívar state (NeoMaps route NM22); 'Anacoco', Bolívar state (NeoMaps route NM93); 'La Escalera', Bolívar state (NeoMaps route NM90); 'Kavanayén', Bolívar state (NeoMaps route NM21); 'Paraytepuy', Bolívar state (NeoMaps route NM23).



It was detected twice during the field work in 2010, and only has few GBIF record from the 2008-2012 period.

Due to the scarce data available, the model has a poor fit with a combination of high probability of occurrence and low probability of detection and too wide standard errors for the estimated parameters:

```

Call:
occu(formula = ~1 ~ 1, data = UMF[os, ])

Occupancy (logit-scale):
Estimate    SE      z P(>|z|)
 2.82 48.7 0.058  0.954

Detection (logit-scale):
Estimate    SE      z P(>|z|)
 -5.82 2.83 -2.06  0.0396

AIC: 31.52855
Number of sites: 450
optim convergence code: 0
optim iterations: 111
Bootstrap iterations: 0

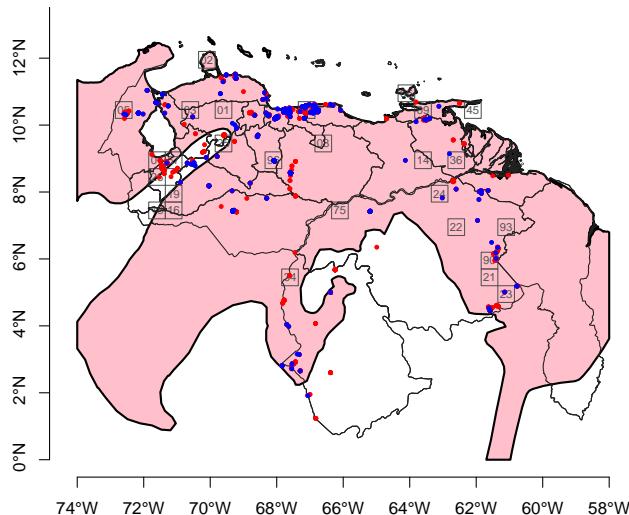
```

3.8 Genus *Eupsittula*, species *Eupsittula pertinax*

The genus *Eupsittula* contains one widespread and common species.

| aspp | GBIF | GBIF.2010 | NM.M1 | NM.L1 | NM.L2 | NM.L3 |
|---------------------|-----------|-----------|-------|-------|-------|-------|
| Eupsittula pertinax | Arat_pert | 1102 | 482 | 155 | 27 | 11 |

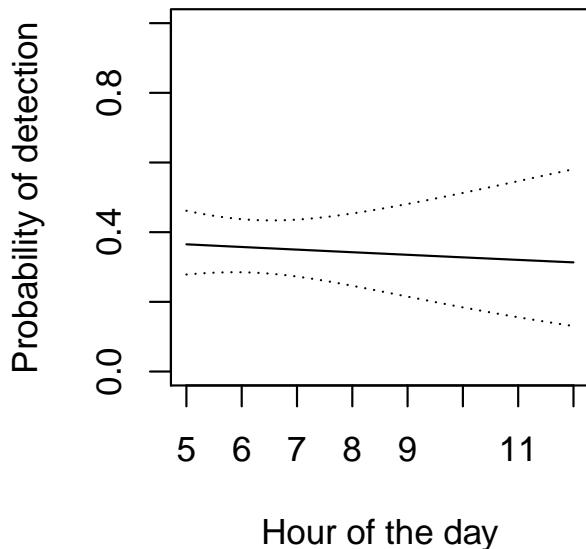
Eupsittula pertinax is widely distributed in Venezuela, and was expected in 1180 sampling points.



Available data allowed to fit several alternative models, and models with both climatic and vegetation covariates received most support according to AIC weights.

| | spp | mod | n | dtt | AICc | Delta.AICc | AICw | LL |
|----|-----------|-------------|------|-----|----------|------------|-------|---------|
| 31 | Arat_pert | p(.)Psi(VC) | 1300 | 190 | 1177.645 | 0.00 | 0.665 | -578.74 |
| 32 | Arat_pert | p(h)Psi(VC) | 1300 | 190 | 1179.135 | 1.49 | 0.316 | -578.46 |
| 33 | Arat_pert | p(.)Psi(C) | 1300 | 190 | 1185.397 | 7.75 | 0.014 | -584.64 |
| 34 | Arat_pert | p(h)Psi(C) | 1300 | 190 | 1187.145 | 9.50 | 0.006 | -584.50 |
| 35 | Arat_pert | p(.)Psi(V) | 1300 | 190 | 1284.377 | 106.73 | 0.000 | -638.17 |
| 36 | Arat_pert | p(h)Psi(V) | 1300 | 190 | 1286.388 | 108.74 | 0.000 | -638.17 |
| 37 | Arat_pert | nulo | 1300 | 190 | 1293.649 | 116.00 | 0.000 | -644.82 |
| 38 | Arat_pert | p(h)Psi(.) | 1300 | 190 | 1295.622 | 117.98 | 0.000 | -644.80 |

There seems to be a slight effect of the hour of the day on the probability of detection for this species.



However, the best supported model includes a constant p .

Call:

```
occu(formula = ~1 ~ evi01 + I(evi01^2) + pet01 + I(pet01^2) +
  dT01 + I(dT01^2) + pre01 + I(pre01^2), data = UMF[os, ])
```

Occupancy (logit-scale):

| | Estimate | SE | z | P(> z) |
|-------------|----------|--------|--------|----------|
| (Intercept) | -1.5567 | 0.2604 | -5.979 | 2.25e-09 |
| evi01 | 0.5476 | 0.1828 | 2.996 | 2.74e-03 |
| I(evi01^2) | 0.2696 | 0.1410 | 1.912 | 5.58e-02 |
| pet01 | 0.0308 | 0.1473 | 0.209 | 8.34e-01 |
| I(pet01^2) | -0.1484 | 0.0713 | -2.082 | 3.74e-02 |
| dT01 | 1.8196 | 0.2364 | 7.697 | 1.39e-14 |
| I(dT01^2) | 0.1373 | 0.1712 | 0.802 | 4.23e-01 |
| pre01 | 0.0623 | 0.1910 | 0.326 | 7.44e-01 |

```
I(pre01^2)      0.0602 0.0872 0.691 4.89e-01
```

Detection (logit-scale):

| Estimate | SE | z | P(> z) |
|----------|------|-------|----------|
| -0.592 | 0.17 | -3.49 | 0.000485 |

AIC: 1177.474

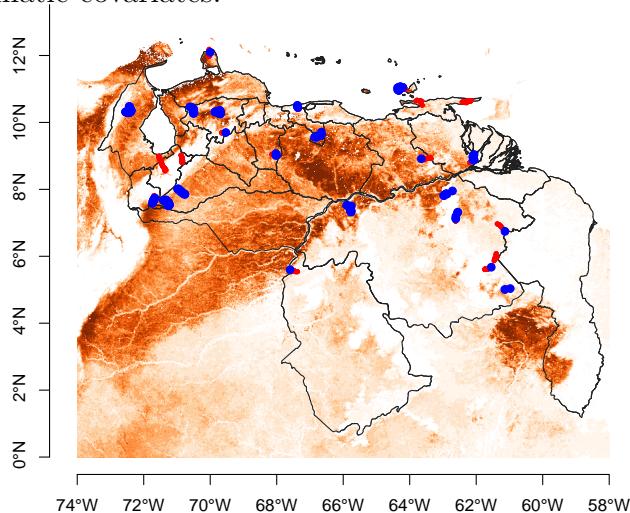
Number of sites: 1300

optim convergence code: 0

optim iterations: 51

Bootstrap iterations: 0

The following map shows the predicted (unconditional) probability of presence for the whole country based on the model with highest support and the values of the vegetation and climatic covariates.



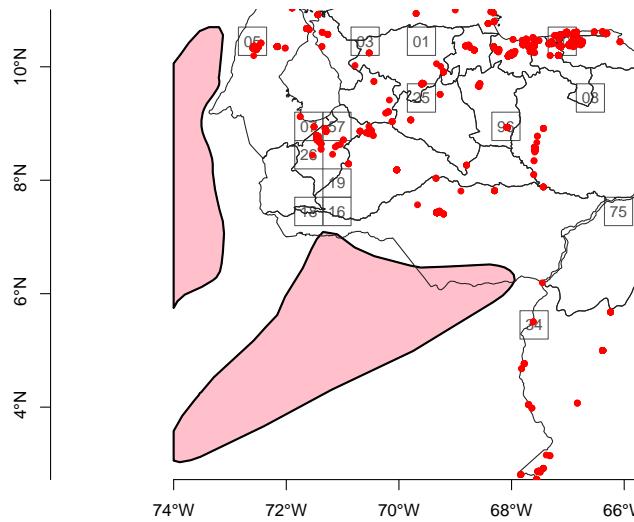
3.9 Genus *Forpus*

The genus *Forpus* is represented in Venezuela by three species, but only one species was detected in NeoMaps surveys of 2010.

| | aspp | GBIF | GBIF.2010 | NM.M1 | NM.L1 | NM.L2 | NM.L3 |
|------------------------------|-----------|------|-----------|-------|-------|-------|-------|
| <i>Forpus conspicillatus</i> | Forp_cons | 1 | 0 | NA | NA | NA | NA |
| <i>Forpus modestus</i> | Forp_mode | 9 | 0 | NA | NA | NA | NA |
| <i>Forpus passerinus</i> | Forp_pass | 991 | 443 | 31 | 8 | 5 | 5 |

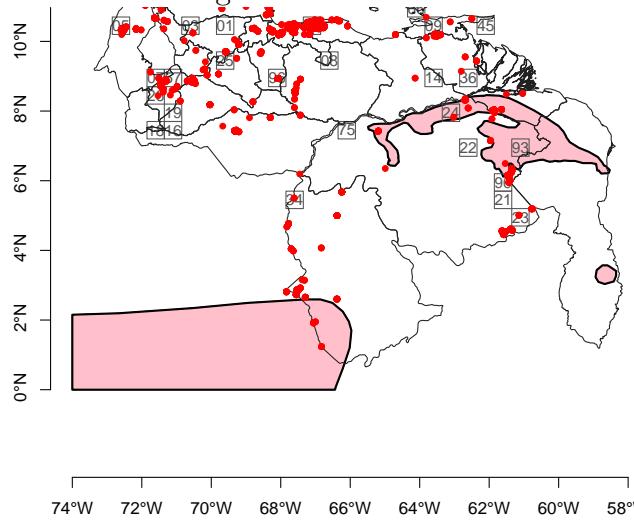
3.9.1 *Forpus conspicillatus*

Forpus conspicillatus is only found in south-western Venezuela, and it was not expected in NeoMaps sampling region.



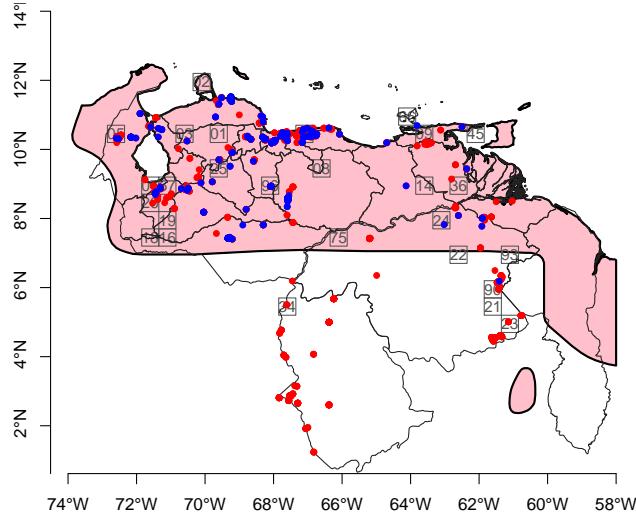
3.9.2 *Forpus modestus*

Forpus modestus was expected in 102 sampling points from 'Guri', Bolívar state (NeoMaps route NM24); 'Anacoco', Bolívar state (NeoMaps route NM93). However this species was not detected during the field work in 2010.



3.9.3 *Forpus passerinus*

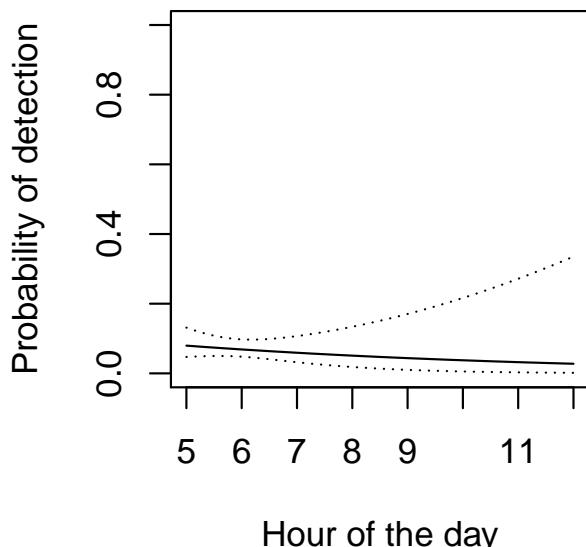
Forpus passerinus has a widespread distribution in northern Venezuela. It was expected in 1060 sampling points from NeoMaps sampling localities..



Available data allowed to fit several alternative models, but the model with most support included climatic covariates.

| | spp | mod | n | dtt | AICc | Delta.AICc | AICw | LL |
|----|-----------|------------|------|-----|----------|------------|------|---------|
| 49 | Forp_pass | p(h)Psi(C) | 1050 | 47 | 399.3574 | 0.00 | 1 | -190.59 |
| 50 | Forp_pass | nulo | 1050 | 47 | 441.3997 | 42.04 | 0 | -218.69 |
| 51 | Forp_pass | p(h)Psi(.) | 1050 | 47 | 442.0069 | 42.65 | 0 | -217.99 |
| 52 | Forp_pass | p(.)Psi(V) | 1050 | 47 | 443.3331 | 43.98 | 0 | -217.65 |
| 53 | Forp_pass | p(h)Psi(V) | 1050 | 47 | 444.0884 | 44.73 | 0 | -217.02 |

The best model includes an effect of time of day on the detection probability.



The fitted model had high estimated parameter values.

Call:

```
occu(formula = ~hora ~ pet01 + I(pet01^2) + dT01 + I(dT01^2) +
```

```
pre01 + I(pre01^2), data = UMF[os, ])
```

Occupancy (logit-scale):

| | Estimate | SE | z | P(> z) |
|-------------|----------|-------|--------|---------|
| (Intercept) | 1.314 | 1.256 | 1.046 | 0.2954 |
| pet01 | -3.515 | 2.046 | -1.718 | 0.0858 |
| I(pet01^2) | 6.256 | 2.765 | 2.262 | 0.0237 |
| dT01 | -0.652 | 0.964 | -0.677 | 0.4985 |
| I(dT01^2) | -4.043 | 1.750 | -2.310 | 0.0209 |
| pre01 | -4.137 | 2.156 | -1.919 | 0.0550 |
| I(pre01^2) | -5.171 | 2.361 | -2.190 | 0.0285 |

Detection (logit-scale):

| | Estimate | SE | z | P(> z) |
|-------------|----------|-------|--------|---------|
| (Intercept) | -2.08 | 0.789 | -2.634 | 0.00845 |
| hora | -1.49 | 2.220 | -0.669 | 0.50339 |

AIC: 399.1843

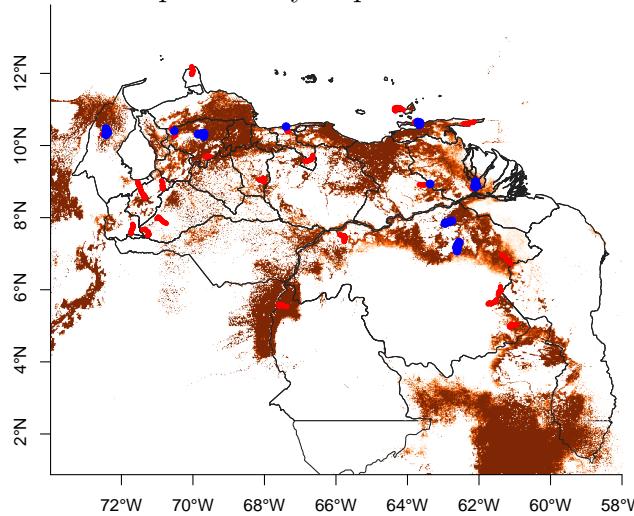
Number of sites: 1050

optim convergence code: 0

optim iterations: 63

Bootstrap iterations: 0

However, the combination of coefficients resulted in predictions with high contrast in unconditional probability of presence.

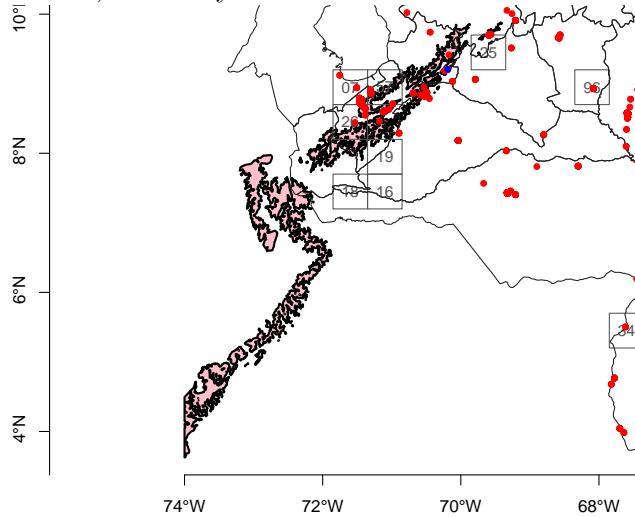


3.10 Genus Hapalopsittaca, species *H. amazonina*

The genus Hapalopsittaca is represented in Venezuela by a single species.

| aspp | GBIF | GBIF.2010 | NM.M1 | NM.L1 | NM.L2 | NM.L3 |
|----------------|-----------|-----------|-------|-------|-------|-------|
| Hapalopsittaca | amazonina | Hapa_amaz | 12 | 1 | NA | NA |

Hapalopsittaca amazonina is a species of restricted distribution in Venezuela. *Hapalopsittaca amazonina* was expected in 43 sampling points from 'Jají', Mérida state (NeoMaps route NM26); 'Piñango', Mérida state (NeoMaps route NM57). It was not detected by NeoMaps field work, and only has few GBIF record from the 2008-2012 period.

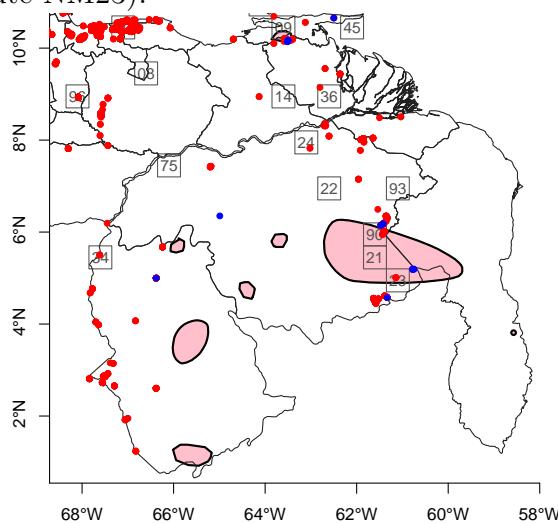


3.11 Genus Nannopsittaca, species *N. panychlora*

The genus Nannopsittaca is represented in Venezuela by a single species.

| | aspp | GBIF | GBIF.2010 | NM.M1 | NM.L1 | NM.L2 | NM.L3 |
|--------------------------|-----------|------|-----------|-------|-------|-------|-------|
| Nannopsittaca panychlora | Nann_pany | 42 | | 15 | 1 | 0 | 0 |

Nannopsittaca panychlora is a species of restricted distribution in Venezuela, it was expected in 146 sampling points from 'La Escalera', Bolívar state (NeoMaps route NM90); 'Kavanayén', Bolívar state (NeoMaps route NM21); 'Paraytepuy', Bolívar state (NeoMaps route NM23).



It was only detected once during NeoMaps field work. The fitted model had large estimates and large uncertainty in both parameters.

```

Call:
occu(formula = ~1 ~ 1, data = UMF[os, ])

Occupancy (logit-scale):
Estimate    SE      z P(>|z|)
4.41 93.2 0.0473 0.962

Detection (logit-scale):
Estimate    SE      z P(>|z|)
-5.46 1.5 -3.64 0.000268

AIC: 16.95734
Number of sites: 150
optim convergence code: 0
optim iterations: 73
Bootstrap iterations: 0

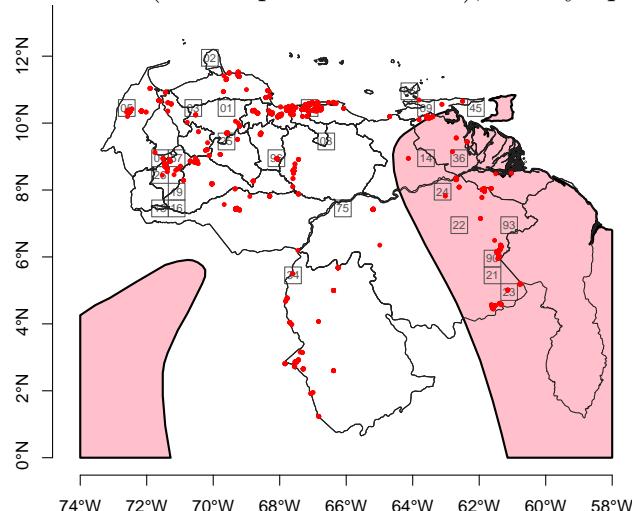
```

3.12 Genus Orthopsittaca, species *O. manilatus*

The genus Orthopsittaca is represented in Venezuela by a single species.

| | aspp | GBIF | GBIF.2010 | NM.M1 | NM.L1 | NM.L2 | NM.L3 |
|-----------------------------------|------|------|-----------|-------|-------|-------|-------|
| Orthopsittaca manilatus Orth_mani | | | | 1 | 0 | 1 | 1 |

Orthopsittaca manilatus is a species of restricted distribution in Venezuela, it was expected in 407 sampling points from 'San Tomé', Anzoátegui-Monagas state (NeoMaps route NM14); 'Isla de Guara', Monagas-Delta state (NeoMaps route NM36); 'Guri', Bolívar state (NeoMaps route NM24); 'El Manteco', Bolívar state (NeoMaps route NM22); 'Anacoco', Bolívar state (NeoMaps route NM93); 'La Escalera', Bolívar state (NeoMaps route NM90); 'Kavanayén', Bolívar state (NeoMaps route NM21); 'Paraytepuy', Bolívar state (NeoMaps route NM23).



The small number of detections during NeoMaps field work allowed to fit two alternative models with similar AICc support.

| | spp | mod | n | dtt | AICc | Delta.AICc | AICw | LL |
|----|-----------|------------|-----|-----|----------|------------|-------|-------|
| 55 | Orth_mani | p(h)Psi(.) | 450 | 5 | 63.46122 | 0.00 | 0.522 | -28.7 |
| 56 | Orth_mani | nulo | 450 | 5 | 63.63516 | 0.17 | 0.478 | -29.8 |

The fitted model had large estimates and large uncertainty in both parameters and is thus non-informative for inferences or predictions.

```

Call:
occu(formula = ~hora ~ 1, data = UMF[os, ])

Occupancy (logit-scale):
Estimate    SE      z P(>|z|)
 3.25 40.6 0.08  0.936

Detection (logit-scale):
Estimate    SE      z P(>|z|)
(Intercept) -1.02 3.26 -0.313  0.754
hora        -12.42 9.37 -1.326  0.185

AIC: 63.40741
Number of sites: 450
optim convergence code: 0
optim iterations: 68
Bootstrap iterations: 0

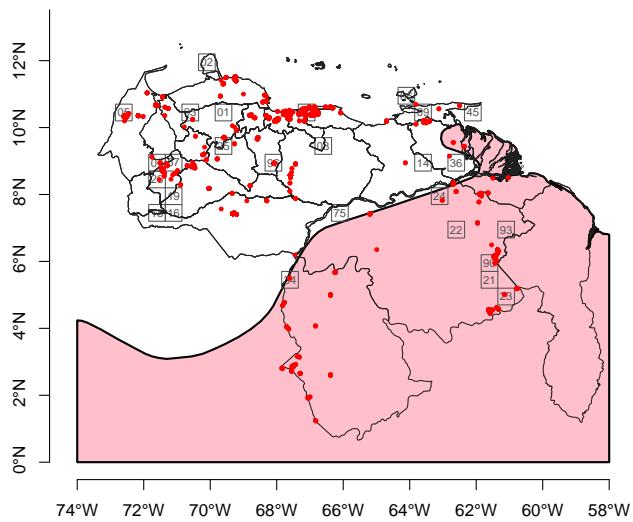
```

3.13 Genus Pionites, species *P. melanocephala*

The genus Bolborhynchus is represented in Venezuela by a single species.

| | aspp | GBIF | GBIF.2010 | NM.M1 | NM.L1 | NM.L2 | NM.L3 |
|------------------------|-----------|------|-----------|-------|-------|-------|-------|
| Pionites melanocephala | Pion_mela | 5 | 0 | 3 | 1 | 0 | 1 |

Pionites melanocephala is found in southern Venezuela. It was expected in 0 sampling points from .



With the few number of detections, only one model could be fitted for this species. The fitted model had high estimates and high uncertainty in detectability parameters, and its predictions are not informative.

```

Call:
occu(formula = ~hora ~ 1, data = UMF[os, ])

Occupancy (logit-scale):
Estimate      SE      z P(>|z|)
-2.52 2.12 -1.19  0.234

Detection (logit-scale):
            Estimate      SE      z P(>|z|)
(Intercept) 2.96 5.29 0.559  0.576
hora        -15.40 11.96 -1.288  0.198

AIC: 60.56913
Number of sites: 350
optim convergence code: 0
optim iterations: 83
Bootstrap iterations: 0

```

3.14 Genus Pionus

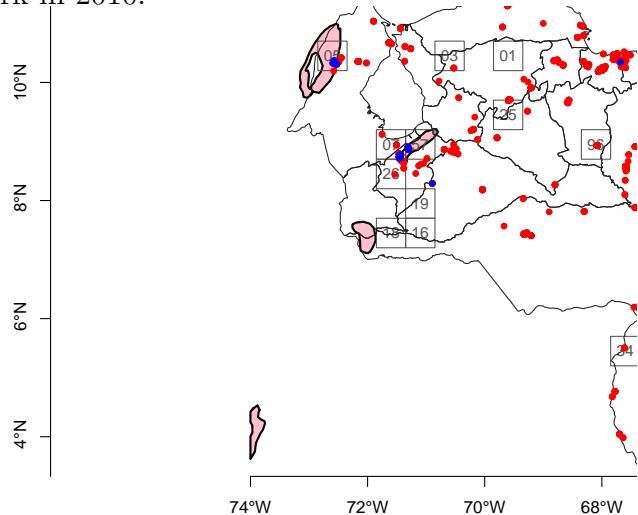
The genus *Pionus* is represented in Venezuela by five species, but only one species was detected in NeoMaps surveys of 2010.

| | aspp | GBIF | GBIF.2010 | NM.M1 | NM.L1 | NM.L2 | NM.L3 |
|----------------------------|------------------|------|-----------|-------|-------|-------|-------|
| <i>Pionus chalcopterus</i> | <i>Pion_chal</i> | 44 | 24 | NA | NA | NA | NA |
| <i>Pionus fuscus</i> | <i>Pion_fusc</i> | 46 | 4 | NA | NA | NA | NA |
| <i>Pionus menstruus</i> | <i>Pion_mens</i> | 356 | 86 | 28 | 11 | 7 | 3 |

| | | | | | | | |
|-------------------|-----------|-----|----|----|----|----|----|
| Pionus seniloides | Pion_seni | 0 | 0 | NA | NA | NA | NA |
| Pionus sordidus | Pion_sord | 142 | 38 | NA | NA | NA | NA |

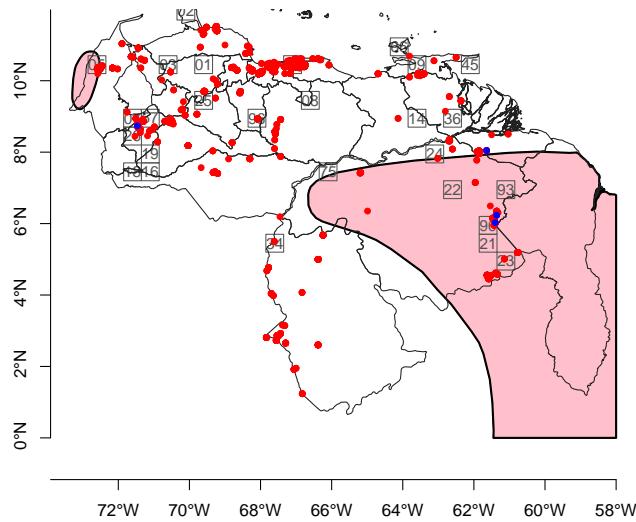
3.14.1 *Pionus chalcopterus*

Pionus chalcopterus was expected in 38 sampling points from 'Rosario de Perijá', Zulia state (NeoMaps route NM05); 'Sur del Lago', Zulia-Mérida state (NeoMaps route NM07); 'Jají', Mérida state (NeoMaps route NM26). However this species was not detected during the field work in 2010.



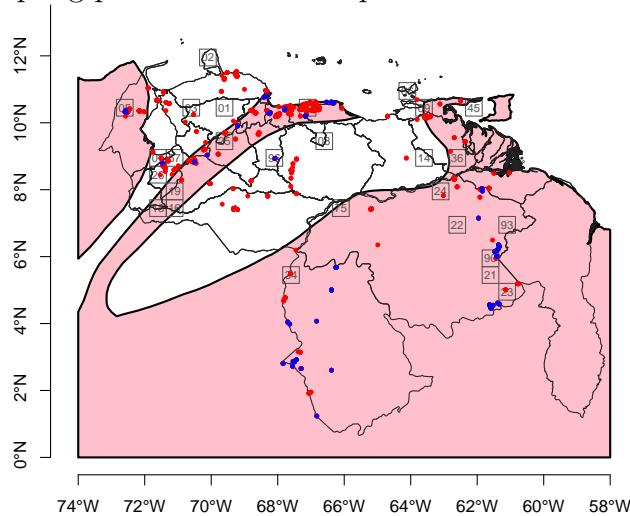
3.14.2 *Pionus fuscus*

Pionus fuscus was expected in 283 sampling points from 'Rosario de Perijá', Zulia state (NeoMaps route NM05); 'Guri', Bolívar state (NeoMaps route NM24); 'El Manteco', Bolívar state (NeoMaps route NM22); 'Caicara del Orinoco', Bolívar state (NeoMaps route NM75); 'Anacoco', Bolívar state (NeoMaps route NM93); 'La Escalera', Bolívar state (NeoMaps route NM90); 'Kavanayén', Bolívar state (NeoMaps route NM21); 'Paraytepuy', Bolívar state (NeoMaps route NM23). However this species was not detected during the field work in 2010.



3.14.3 *Pionus menstruus*

Pionus menstruus has a widespread but non continuous distribution and was expected in 798 sampling points from NeoMaps.



Available data allowed to fit several alternative models.

| | spp | mod | n | dtt | AICc | Delta.AICc | AICw | LL |
|----|-----------|-------------|-----|-----|----------|------------|-------|---------|
| 60 | Pion_mens | p(h)Psi(VC) | 850 | 47 | 361.1700 | 0.00 | 0.783 | -169.43 |
| 61 | Pion_mens | p(h)Psi(.) | 850 | 47 | 364.3834 | 3.21 | 0.157 | -179.18 |
| 62 | Pion_mens | p(h)Psi(V) | 850 | 47 | 366.2987 | 5.13 | 0.060 | -178.11 |
| 63 | Pion_mens | p(.)Psi(VC) | 850 | 47 | 401.6749 | 40.50 | 0.000 | -190.71 |
| 64 | Pion_mens | p(.)Psi(V) | 850 | 47 | 415.4491 | 54.28 | 0.000 | -203.70 |
| 65 | Pion_mens | nulo | 850 | 47 | 419.0509 | 57.88 | 0.000 | -207.52 |

The model with most supports include a effect of time of the day on probability of detection and effect of climatic and vegetation covariates on probability of presence.

```

Call:
occu(formula = ~hora ~ evi01 + I(evi01^2) + pet01 + I(pet01^2) +
    dT01 + I(dT01^2) + pre01 + I(pre01^2), data = UMF[os, ])

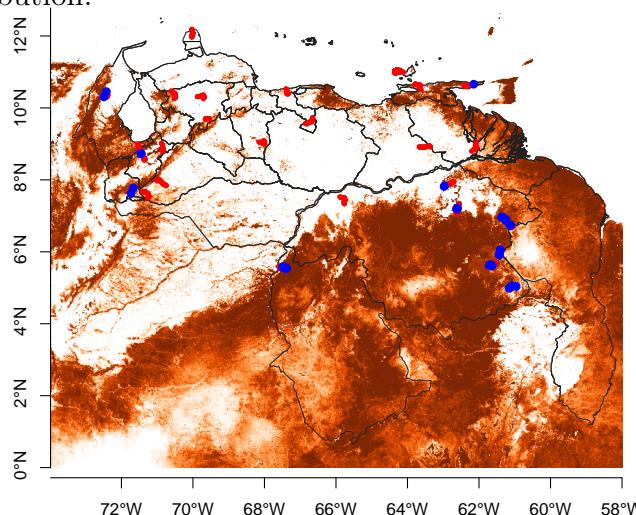
Occupancy (logit-scale):
            Estimate      SE      z P(>|z|)
(Intercept) -1.600 1.261 -1.27  0.204
evi01       -1.226 1.038 -1.18  0.238
I(evi01^2)   1.873 1.586  1.18  0.238
pet01        1.463 1.409  1.04  0.299
I(pet01^2)   0.589 0.585  1.01  0.315
dT01        -2.077 1.587 -1.31  0.191
I(dT01^2)   -1.102 1.033 -1.07  0.286
pre01        2.618 1.614  1.62  0.105
I(pre01^2)  -0.584 0.436 -1.34  0.181

Detection (logit-scale):
            Estimate      SE      z P(>|z|)
(Intercept)  3.82  1.14  3.36 7.87e-04
hora        -19.76 3.63 -5.45 5.04e-08

AIC: 360.8549
Number of sites: 850
optim convergence code: 0
optim iterations: 105
Bootstrap iterations: 0

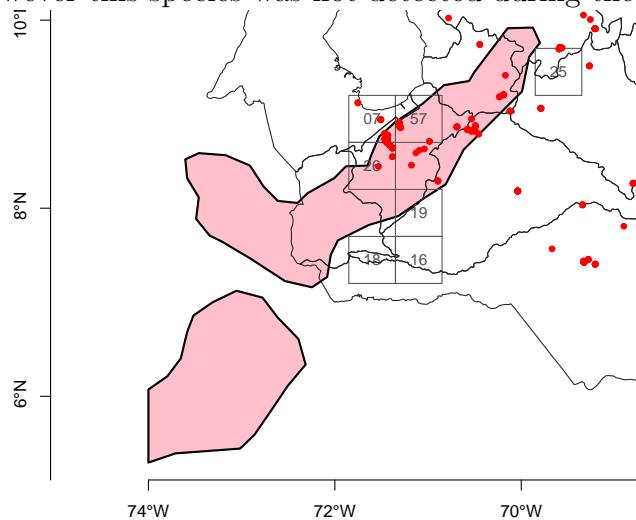
```

The predicted unconditional probability of presence matches very well the expected distribution.



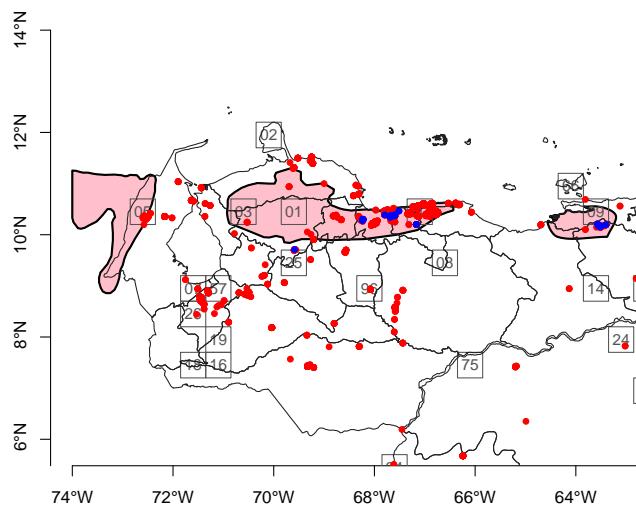
3.14.4 *Pionus seniloides*

Pionus seniloides was expected in 128 sampling points from 'Sur del Lago', Zulia-Mérida state (NeoMaps route NM07); 'Jaí', Mérida state (NeoMaps route NM26); 'Piñango', Mérida state (NeoMaps route NM57); 'San Joaquín de Navay', Táchira state (NeoMaps route NM18). However this species was not detected during the field work in 2010.



3.14.5 *Pionus sordidus*

Pionus sordidus was expected in 162 sampling points from 'Rosario de Perijá', Zulia state (NeoMaps route NM05); 'Quebrada Arriba', Lara-Falcón state (NeoMaps route NM03); 'Araya', Sucre state (NeoMaps route NM09); 'Río Tocuyo', Lara state (NeoMaps route NM01); 'Colonia Tovar', Aragua-Vargas state (NeoMaps route NM80). However this species was not detected during the field work in 2010.

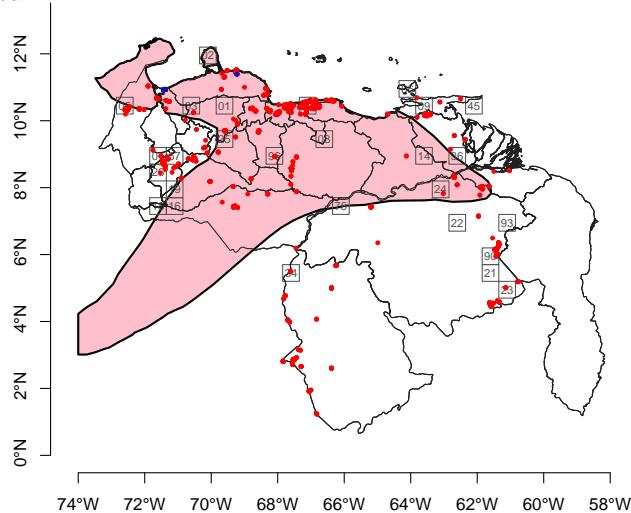


3.15 Genus *Thectocercus*, species *T. acuticaudatus*

The genus *Thectocercus* is represented in Venezuela by a single species.

| | aspp | GBIF | GBIF.2010 | NM.M1 | NM.L1 | NM.L2 | NM.L3 |
|--------------------------------------|------|------|-----------|-------|-------|-------|-------|
| Thectocercus acuticaudatus Arat_acut | 38 | | | 5 | 3 | 0 | 0 |

Thectocercus acuticaudatus was expected in 575 sampling points. Although the species was detected during NeoMaps surveys of 2010, no model could be fitted with the available data.



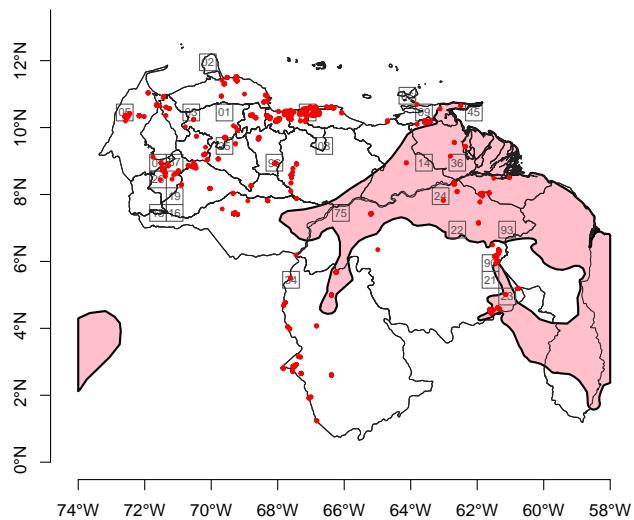
3.16 Genus Psittacara

The genus *Psittacara* is represented in Venezuela by two species, both were detected in NeoMaps surveys.

| | aspp | GBIF | GBIF.2010 | NM.M1 | NM.L1 | NM.L2 | NM.L3 |
|-------------------------------------|------|------|-----------|-------|-------|-------|-------|
| Psittacara leucophthalmus Arat_leuc | 0 | | | 0 | 5 | 2 | 0 |
| Psittacara wagleri Arat_wagl | 1190 | | | 101 | 4 | 0 | 0 |

3.16.1 *Psittacara leucophthalmus*

Psittacara leucophthalmus is only found in eastern Venezuela, and was expected in 411 sampling points from 'Paria', Sucre state (NeoMaps route NM45); 'San Tomé', Anzoátegui-Monagas state (NeoMaps route NM14); 'Isla de Guara', Monagas-Delta state (NeoMaps route NM36); 'Guri', Bolívar state (NeoMaps route NM24); 'El Manteco', Bolívar state (NeoMaps route NM22); 'Caicara del Orinoco', Bolívar state (NeoMaps route NM75); 'Anacoco', Bolívar state (NeoMaps route NM93); 'La Escalera', Bolívar state (NeoMaps route NM90); 'Paraytepuy', Bolívar state (NeoMaps route NM23).



This species was only detected in few NeoMaps localities. Due to the scarce data available, only two models could be fitted. These models had a poor fit with large estimates and large uncertainty for both parameters.

```

spp          mod   n dtt      AICc Delta.AICc  AICw    LL
29 Arat_leuc p(h)Psi(.) 450    7 80.87169      0.00 0.712 -37.41
30 Arat_leuc      nulo 450    7 82.68527      1.81 0.288 -39.33

Call:
occu(formula = ~hora ~ 1, data = UMF[os, ])

Occupancy (logit-scale):
Estimate   SE     z P(>|z|)
-2.75  1.09 -2.51  0.012

Detection (logit-scale):
Estimate   SE     z P(>|z|)
(Intercept) 3.87  4.04  0.958  0.338
hora       -16.50 10.27 -1.607  0.108

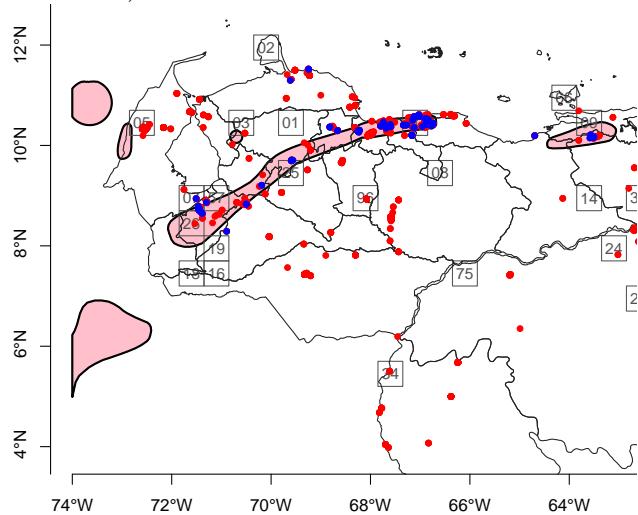
AIC: 80.81787
Number of sites: 450
optim convergence code: 0
optim iterations: 44
Bootstrap iterations: 0

```

3.16.2 *Psittacara wagleri*

Psittacara wagleri is only found in northern mountain ranges of Venezuela, and was expected in 215 sampling points from 'Sur del Lago', Zulia-Mérida state (NeoMaps route NM07); 'Jají',

Mérida state (NeoMaps route NM26); 'Piñango', Mérida state (NeoMaps route NM57); 'Yacambú', Lara state (NeoMaps route NM25); 'Colonia Tovar', Aragua-Vargas state (NeoMaps route NM80).



This species was only detected in few NeoMaps localities. Due to the scarce data available only one model could be fitted. This model had a poor fit with a combination of low probability of occurrence and high probability of detection and too wide standard errors for the estimated parameters:

```

spp          mod    n dtt      AICc Delta.AICc AICw      LL
39 Arat_wagl p(h)Psi(.) 250     4 47.11455           0     1 -20.51

Call:
occu(formula = ~hora ~ 1, data = UMF[os, ])

Occupancy (logit-scale):
Estimate      SE      z P(>|z|)
-4.12 0.504 -8.17 3.04e-16

Detection (logit-scale):
Estimate      SE      z P(>|z|)
(Intercept) 12.24 452 0.0271  0.978
hora        -1.41 NaN   NaN     NaN

AIC: 47.01699
Number of sites: 250
optim convergence code: 0
optim iterations: 23
Bootstrap iterations: 0

```

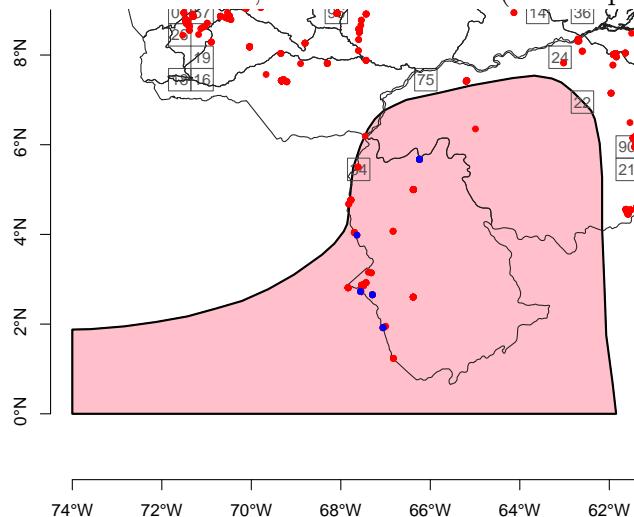
3.17 Genus Pyrilia

The genus Pyrilia is represented in Venezuela by three species, only one species was detected in NeoMaps surveys.

| | aspp | GBIF | GBIF.2010 | NM.M1 | NM.L1 | NM.L2 | NM.L3 |
|--------------------|-----------|------|-----------|-------|-------|-------|-------|
| Pyrilia barrabandi | Pion_barr | 36 | | 11 | 0 | 2 | 0 |
| Pyrilia caica | Pyri_caic | 26 | | 4 | NA | NA | NA |
| Pyrilia pyrilia | Pyri_pyi | 20 | | 12 | NA | NA | NA |

3.17.1 *Pyrilia barrabandi*

Pyrilia barrabandi is only found in southern Venezuela, and was expected in 50 sampling points from 'Gavilán', Amazonas state (NeoMaps route NM34).



It was detected twice during the field work in 2010. Due to the scarce data available, only two models could be fitted. The null model had best support according to AIC weights, but a poor fit with a combination of high probability of occurrence and low probability of detection and too wide standard errors for the estimated parameters:

| spp | mod | n | dtt | AICc | Delta.AICc | AICw | LL |
|--------------|------------|----|-----|----------|------------|------|-------------|
| 57 Pion_barr | nulo | 50 | 2 | 22.75325 | | 0.00 | 0.744 -9.25 |
| 58 Pion_barr | p(h)Psi(.) | 50 | 2 | 24.89061 | | 2.14 | 0.256 -9.18 |

```
Call:  
occu(formula = ~1 ~ 1, data = UMF[os, ])
```

```
Occupancy (logit-scale):  
Estimate SE z P(>|z|)  
3.98 53.9 0.0738 0.941
```

```
Detection (logit-scale):  
Estimate SE z P(>|z|)
```

-3.59 1.25 -2.86 0.00417

AIC: 22.49793

Number of sites: 50

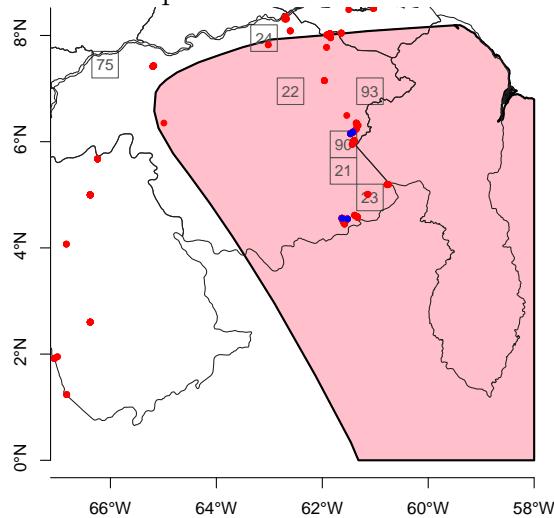
optim convergence code: 0

optim iterations: 69

Bootstrap iterations: 0

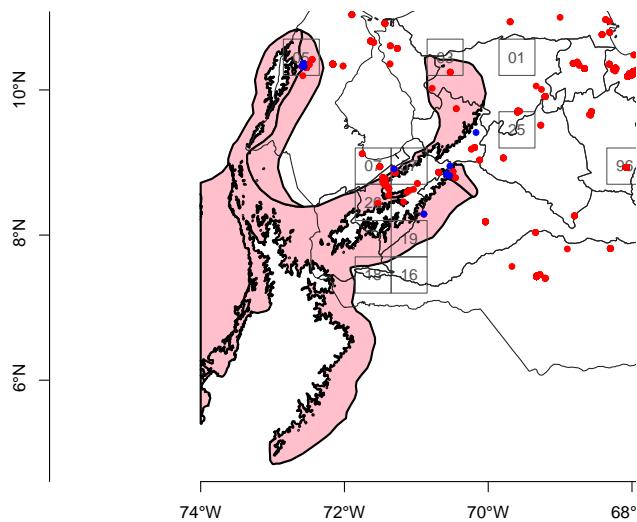
3.17.2 *Pyrilia caica*

Pyrilia caica was expected in 299 sampling points from 'Guri', Bolívar state (NeoMaps route NM24); 'El Manteco', Bolívar state (NeoMaps route NM22); 'Anacoco', Bolívar state (NeoMaps route NM93); 'La Escalera', Bolívar state (NeoMaps route NM90); 'Kavanayén', Bolívar state (NeoMaps route NM21); 'Paraytepuy', Bolívar state (NeoMaps route NM23). However this species was not detected during the field work in 2010.



3.17.3 *Pyrilia pyrilia*

Pyrilia pyrilia was expected in 150 sampling points from 'Quebrada Arriba', Lara-Falcón state (NeoMaps route NM03); 'Sur del Lago', Zulia-Mérida state (NeoMaps route NM07); 'Jají', Mérida state (NeoMaps route NM26); 'Capitanejo', Barinas state (NeoMaps route NM19); 'Otopún', Barinas state (NeoMaps route NM16); 'San Joaquín de Navay', Táchira state (NeoMaps route NM18). However this species was not detected during the field work in 2010.



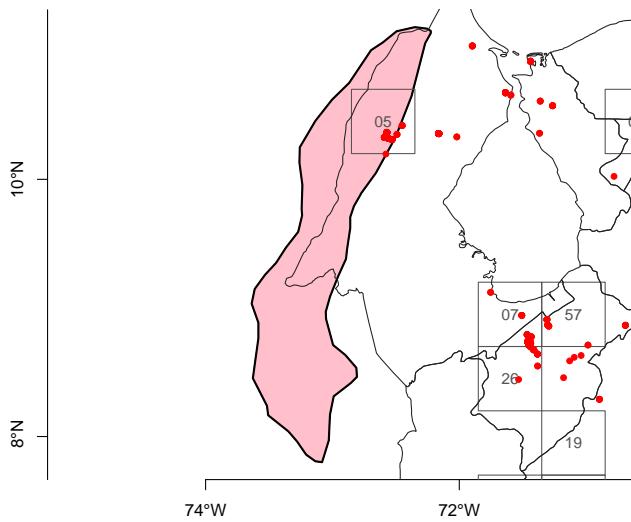
3.18 Genus Pyrrhura

The genus Pyrrhura is represented in Venezuela by seven species and five were detected in NeoMaps surveys.

| | aspp | GBIF | GBIF.2010 | NM.M1 | NM.L1 | NM.L2 | NM.L3 |
|-----------------------|-----------|------|-----------|-------|-------|-------|-------|
| Pyrrhura caeruleiceps | Pyrr_caer | 37 | 2 | NA | NA | NA | NA |
| Pyrrhura egregia | Pyrr_egre | 60 | 9 | 3 | 1 | 0 | 0 |
| Pyrrhura emma | Pyrr_emma | 37 | 2 | NA | NA | NA | NA |
| Pyrrhura hematotis | Pyrr_hoem | 234 | 71 | 1 | 0 | 0 | 0 |
| Pyrrhura melanura | Pyrr_mela | 10 | 3 | 1 | 1 | 0 | 1 |
| Pyrrhura picta | Pyrr_pict | 130 | 31 | 6 | 2 | 2 | 0 |
| Pyrrhura rhodocephala | Pyrr_rhod | 96 | 23 | 1 | 0 | 0 | 0 |

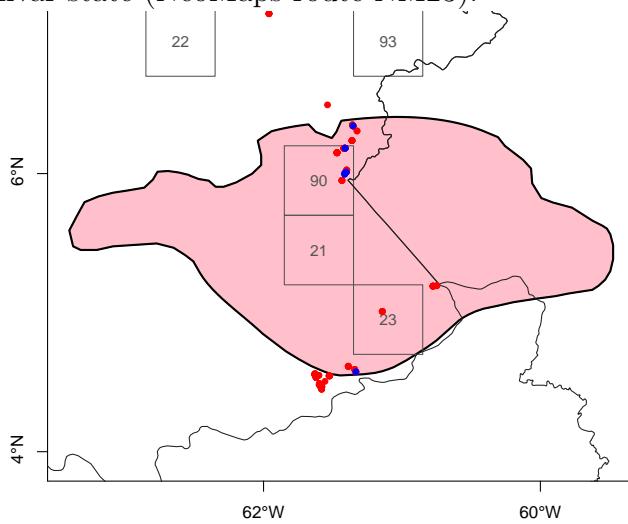
3.18.1 *Pyrrhura caeruleiceps*

Pyrrhura caeruleiceps was expected in 7 sampling points from 'Rosario de Perijá', Zulia state (NeoMaps route NM05). However this species was not detected during the field work in 2010.



3.18.2 *Pyrrhura egregia*

Pyrrhura egregia was expected in 152 sampling points from 'La Escalera', Bolívar state (NeoMaps route NM90); 'Kavanayén', Bolívar state (NeoMaps route NM21); 'Paraytepuy', Bolívar state (NeoMaps route NM23).



This species was detected in few localities during the field work in 2010. Several alternative models were fitted.

| | spp | mod | n | dtt | AICc | Delta.AICc | AICw | LL |
|----|-----------|------------|-----|-----|----------|------------|-------|--------|
| 66 | Pyrr_egre | p(.)Psi(V) | 150 | 3 | 32.51359 | 0.00 | 0.460 | -12.12 |
| 67 | Pyrr_egre | nulo | 150 | 3 | 33.49337 | 0.98 | 0.282 | -14.71 |
| 68 | Pyrr_egre | p(h)Psi(V) | 150 | 3 | 34.65420 | 2.14 | 0.158 | -12.12 |
| 69 | Pyrr_egre | p(h)Psi(.) | 150 | 3 | 35.57612 | 3.06 | 0.100 | -14.71 |

The fitted models with extreme values in parameters and high uncertainty in estimates are not informative.

```

Call:
occu(formula = ~1 ~ evi01 + I(evi01^2), data = UMF[os, ])

Occupancy (logit-scale):
    Estimate      SE      z P(>|z|)
(Intercept) -3.40 0.899 -3.784 0.000154
evi01        2.07 1.828  1.131 0.258179
I(evi01^2)  -1.44 2.180 -0.661 0.508699

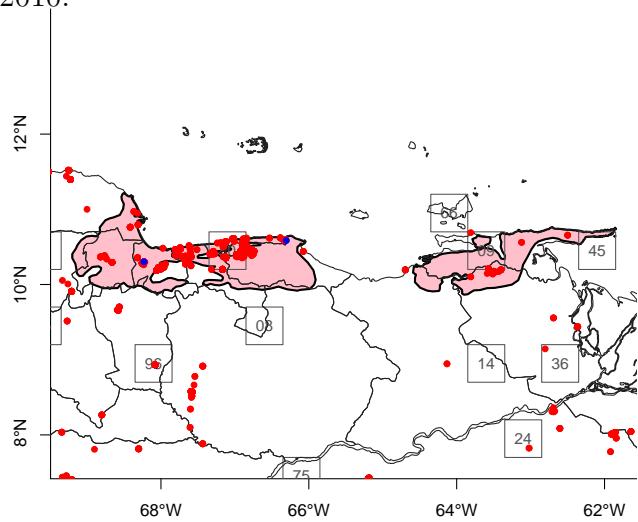
Detection (logit-scale):
    Estimate      SE      z P(>|z|)
               9.44 85.4 0.11  0.912

AIC: 32.23772
Number of sites: 150
optim convergence code: 0
optim iterations: 63
Bootstrap iterations: 0

```

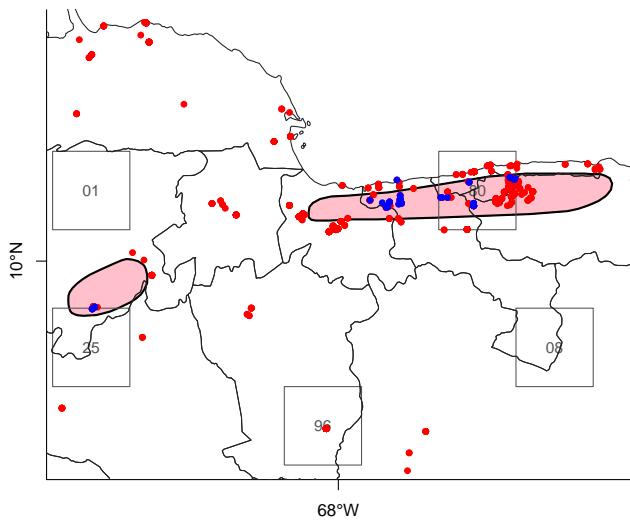
3.18.3 *Pyrrhura emma*

Pyrrhura emma was expected in 107 sampling points from 'Araya', Sucre state (NeoMaps route NM09); 'Paria', Sucre state (NeoMaps route NM45); 'Colonia Tovar', Aragua-Vargas state (NeoMaps route NM80). However this species was not detected during the field work in 2010.



3.18.4 *Pyrrhura hoematotis*

Pyrrhura hoematotis was expected in 85 sampling points from 'Yacambú', Lara state (NeoMaps route NM25); 'Colonia Tovar', Aragua-Vargas state (NeoMaps route NM80).



This species was only detected once during the field work in 2010.

| spp | mod | n | dtt | AICc | Delta.AICc | AICw | LL |
|-------------------|-----|-----|-----|----------|------------|------|--------|
| 70 Pyrr_hoem nulo | | 100 | 1 | 15.32454 | | 0 | 1 -5.6 |

The fitted models with extreme values in parameters and high uncertainty in estimates are not informative.

```

Call:
occu(formula = ~1 ~ 1, data = UMF[os, ])

Occupancy (logit-scale):
Estimate SE z P(>|z|)
-4.59 1.01 -4.56 5.07e-06

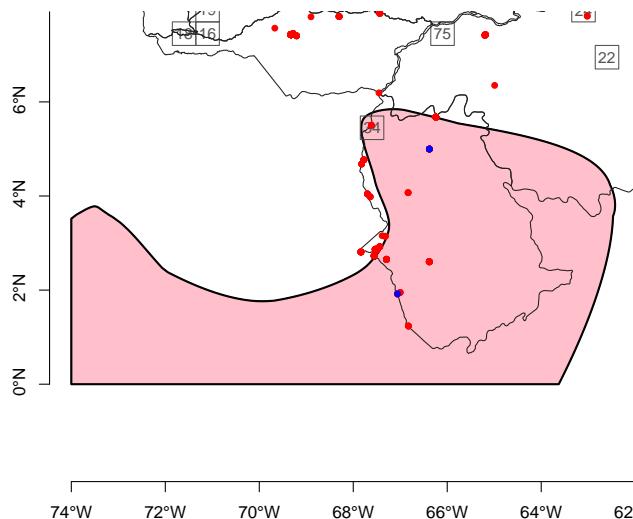
Detection (logit-scale):
Estimate SE z P(>|z|)
 6.66 62 0.107 0.915

AIC: 15.20083
Number of sites: 100
optim convergence code: 0
optim iterations: 66
Bootstrap iterations: 0

```

3.18.5 *Pyrrhura melanura*

Pyrrhura melanura was expected in 50 sampling points from 'Gavilán', Amazonas state (NeoMaps route NM34).



This species was only detected in few occasions during the field work in 2010.

| | spp | mod | n | dtt | AICc | Delta.AICc | AICw | LL | |
|----|-----------|------|--------|-----|------|------------|------|----|--------|
| 71 | Pyrr_mela | p(h) | Psi(.) | 50 | 3 | 31.23506 | 0 | 1 | -12.36 |

The fitted model with extreme values in parameters and high uncertainty in estimates are not informative.

```

Call:
occu(formula = ~hora ~ 1, data = UMF[os, ])

Occupancy (logit-scale):
Estimate      SE      z P(>|z|)
 6.16  91.5  0.0673  0.946

Detection (logit-scale):
Estimate      SE      z P(>|z|)
(Intercept) -0.602  3.19 -0.189  0.850
hora        -8.094 10.12 -0.800  0.424

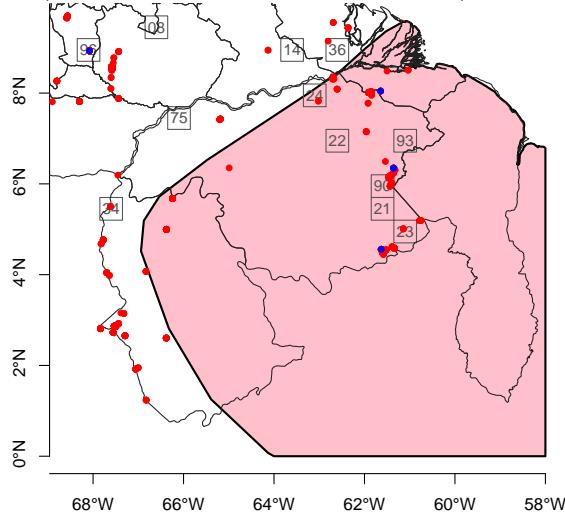
AIC: 30.71332
Number of sites: 50
optim convergence code: 0
optim iterations: 40
Bootstrap iterations: 0

```

3.18.6 *Pyrrhura picta*

Pyrrhura picta was expected in 317 sampling points from 'Isla de Guara', Monagas-Delta state (NeoMaps route NM36); 'Guri', Bolívar state (NeoMaps route NM24); 'El Manteco', Bolívar state (NeoMaps route NM22); 'Anacoco', Bolívar state (NeoMaps route NM93); 'La

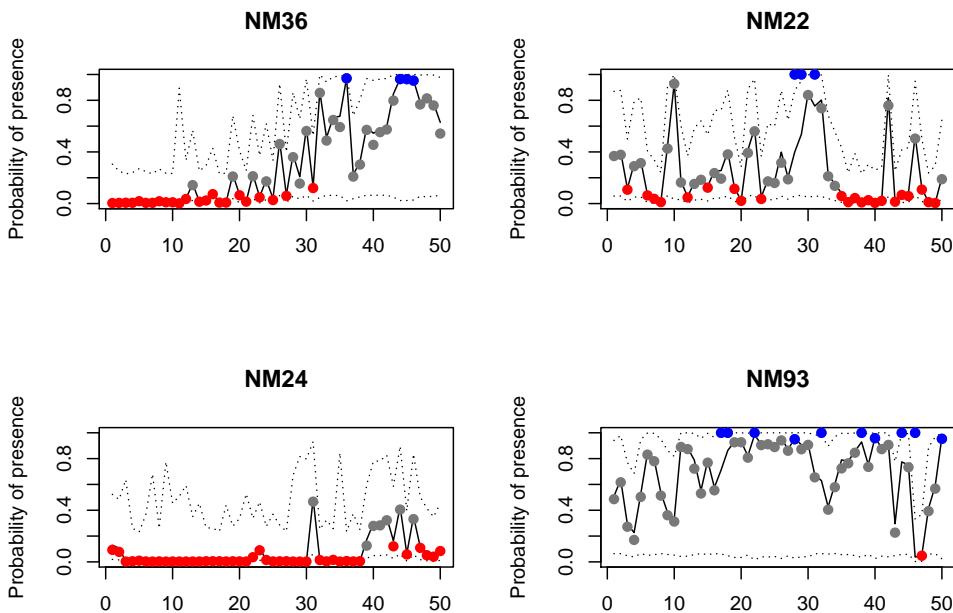
'Escalera', Bolívar state (NeoMaps route NM90); 'Kavanayén', Bolívar state (NeoMaps route NM21); 'Paraytepuy', Bolívar state (NeoMaps route NM23).



Available data allowed to fit several alternative models.

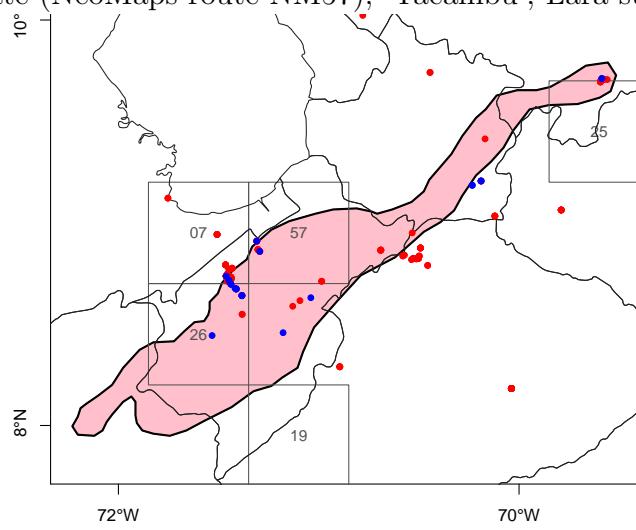
| | spp | mod | n | dtt | AICc | Delta.AICc | AICw | LL |
|----|-----------|------------|-----|-----|-----------|------------|-------|--------|
| 72 | Pyrr_pict | p(.)Psi(V) | 350 | 10 | 89.88223 | 0.00 | 0.724 | -40.88 |
| 73 | Pyrr_pict | p(h)Psi(V) | 350 | 10 | 91.82109 | 1.94 | 0.275 | -40.82 |
| 74 | Pyrr_pict | nulo | 350 | 10 | 104.14575 | 14.26 | 0.001 | -50.06 |
| 75 | Pyrr_pict | p(h)Psi(.) | 350 | 10 | 106.01953 | 16.14 | 0.000 | -49.98 |

According to the model with the best support the unconditional probability of presence varied along the NeoMaps routes depending on the value of the covariates. The figure show four examples of 40 km routes, ordered from beginning to end, three with detections (NM22, NM36 and NM93) and one without detections (NM24). The lines show the predicted unconditional probability of presence (solid line: best estimate, dotted lines 95% confidence interval), and the dots represent the conditional probability given the observed detection history. Blue dots represent the localities of known occurrences, red dots are localities with very low posterior probabilities of presence ($\Psi_{post} < 0.125$), grey dots are localities with intermediate values.



3.18.7 *Pyrrhura rhodocephala*

Pyrrhura rhodocephala was expected in 160 sampling points from 'Sur del Lago', Zulia-Mérida state (NeoMaps route NM07); 'Jají', Mérida state (NeoMaps route NM26); 'Piñango', Mérida state (NeoMaps route NM57); 'Yacambú', Lara state (NeoMaps route NM25).



This species was only detected once during the field work in 2010.

| | spp | mod | n | dtt | AICc | Delta.AICc | AICw | LL |
|----|-----------|------------|------|-----|------------|------------|------|------------|
| 76 | Pyrr_rhod | | nulo | 200 | 1 16.65263 | | 0.00 | 0.737 -6.3 |
| 77 | Pyrr_rhod | p(h)Psi(.) | 200 | 1 | 18.71410 | | 2.06 | 0.263 -6.3 |

The fitted models with extreme values in parameters and high uncertainty in estimates are not informative.

```
Call:  
occu(formula = ~1 ~ 1, data = UMF[os, ])  
  
Occupancy (logit-scale):  
Estimate SE z P(>|z|)  
-5.29 1 -5.28 1.31e-07  
  
Detection (logit-scale):  
Estimate SE z P(>|z|)  
8.41 152 0.0555 0.956  
  
AIC: 16.59171  
Number of sites: 200  
optim convergence code: 0  
optim iterations: 15  
Bootstrap iterations: 0
```

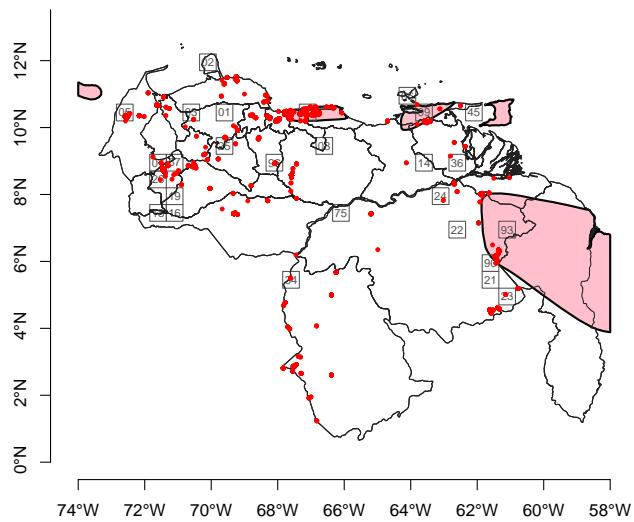
3.19 Genus Touit

No species of the genus Touit was detected by NM surveys. There are also few records in GBIF.

| | aspp | GBIF | GBIF.2010 | NM.M1 | NM.L1 | NM.L2 | NM.L3 |
|--------------------|-----------|------|-----------|-------|-------|-------|-------|
| Touit batavica | Toui_bata | 1 | 0 | NA | NA | NA | NA |
| Touit dilectissima | Toui_dile | 0 | 0 | NA | NA | NA | NA |
| Touit huetii | Toui_huet | 4 | 2 | NA | NA | NA | NA |
| Touit purpurata | Toui_purp | 0 | 0 | NA | NA | NA | NA |

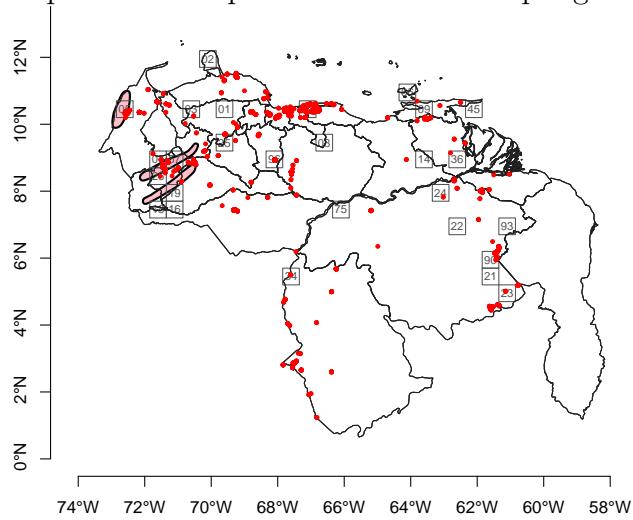
3.19.1 *Touit batavica*

This species was expected in 183 NM sampling localities.



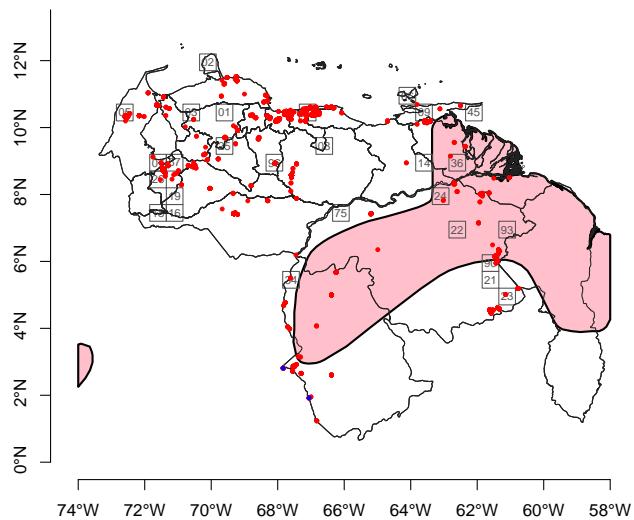
3.19.2 *Touit dilectissima*

This species was expected in 87 NM sampling localities.



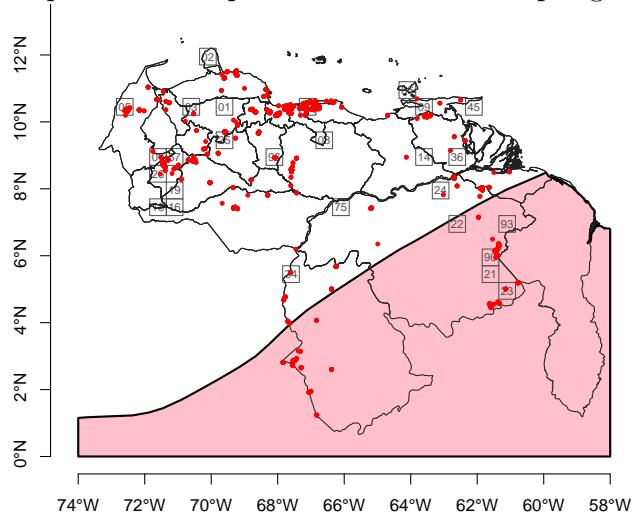
3.19.3 *Touit huetii*

This species was expected in 213 NM sampling localities.



3.19.4 *Touit purpurata*

This species was expected in 222 NM sampling localities.



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