

# Bivariate Residual Plots with Simulation Polygons

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# Case study

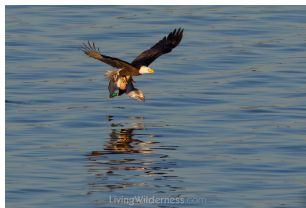
- Bald eagles and mallards



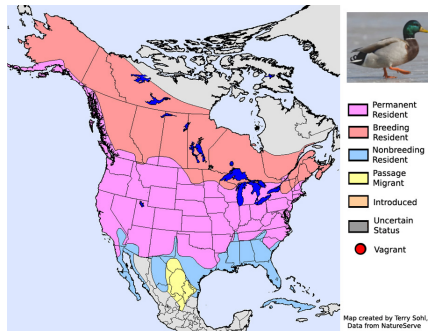
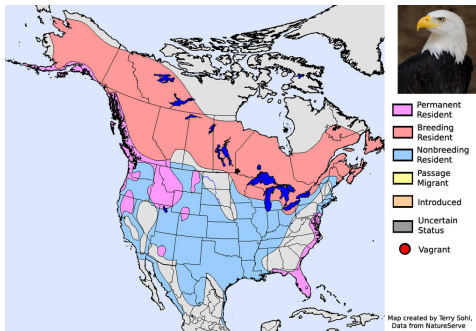


# Case study

- Bald eagles and mallards



# Case study

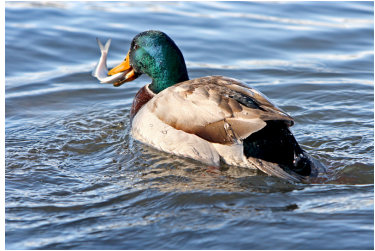


(Ridgely et al., 2003)



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- How should we assess goodness-of-fit for joint models?

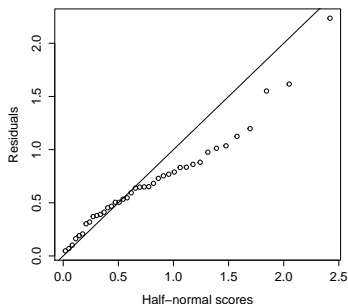
## For univariate models

“The ability of the human eye to find patterns in scatters of points is one strong reason for the use of graphical methods.” (A.C. Atkinson, 1985)

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- Half-normal plots with simulation envelopes
- Ordered absolute values of a model diagnostic vs. expected order statistics of a half-normal distribution  $\Phi^{-1}\left(\frac{i+n-\frac{1}{8}}{2n+\frac{1}{2}}\right)$

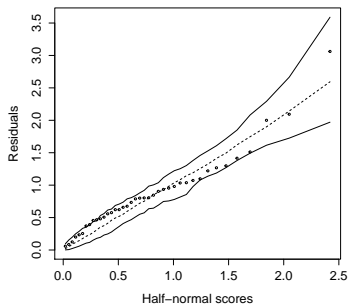


# Half-normal plots with simulation envelopes

- Fit model and obtain diagnostics in absolute value and in order
- Simulate 99 response variables using same model matrix, error distribution and fitted parameters
- Refit the model to each simulated sample and obtain the same diagnostics, again, sorted absolute values
- Compute desired percentiles (e.g. 2.5% and 97.5%) to form the envelope

# Half-normal plots with simulation envelopes

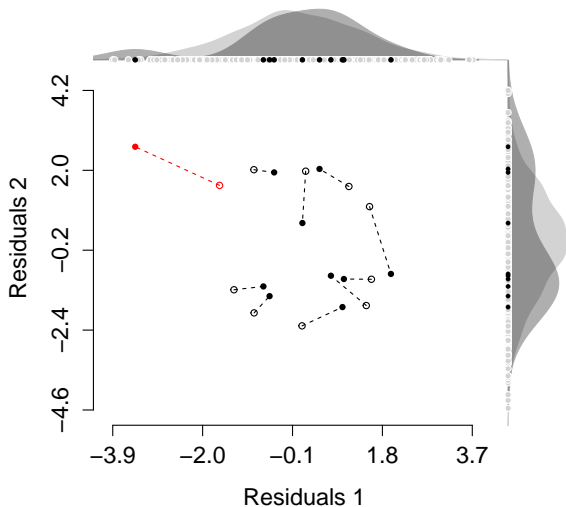
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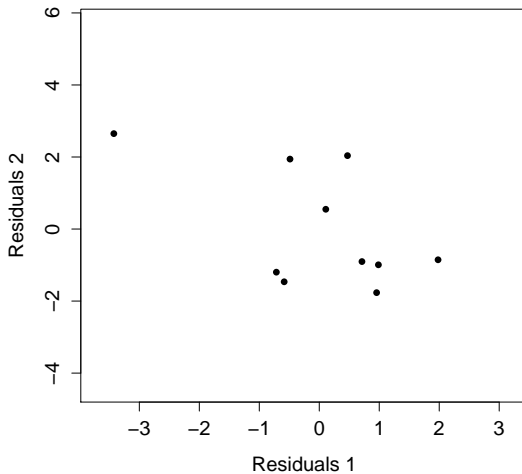


How can we do this for a bivariate model?

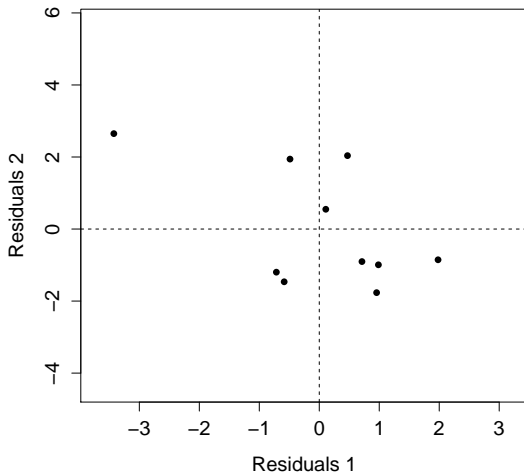
# How can we do this for a bivariate model?



# Bivariate residuals



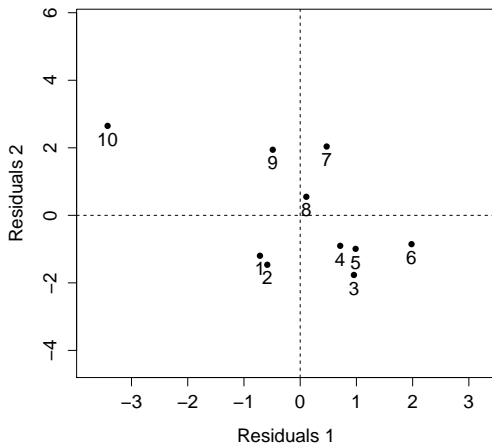
We need to order them



## Ordering by angles

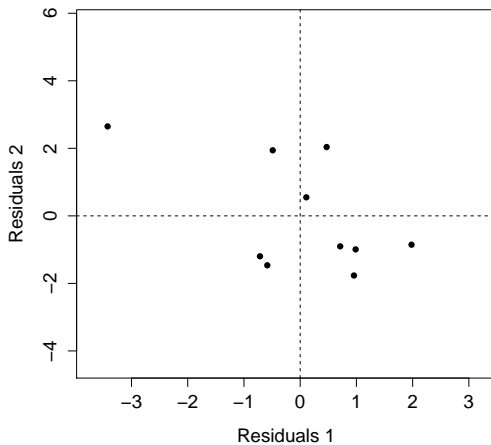
$$\alpha_i = \begin{cases} \tan^{-1} \left( \frac{y_i}{x_i} \right), & x > 0 \\ \tan^{-1} \left( \frac{y_i}{x_i} \right) + \pi, & x < 0 \text{ and } y \geq 0 \\ \tan^{-1} \left( \frac{y_i}{x_i} \right) + \pi, & x < 0 \text{ and } y < 0 \\ \pm \frac{\pi}{2}, & x = 0 \text{ and } y \neq 0 \\ \text{undefined}, & x = y = 0 \end{cases}$$

# Ordering by angles



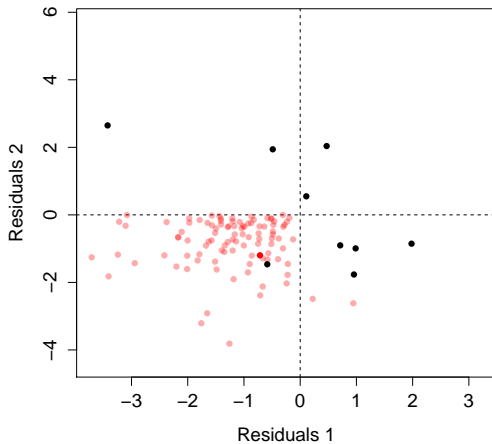
# Now we simulate

- Simulate 99 bivariate responses and refit model



# Now we simulate

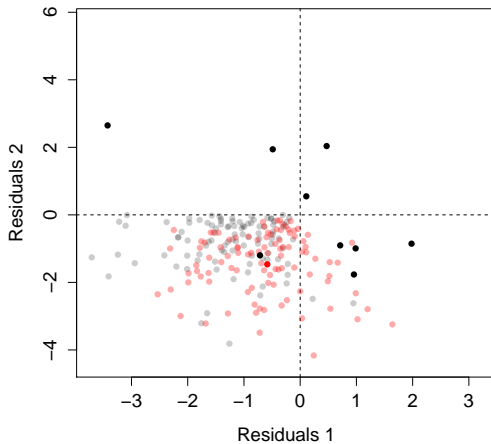
- Simulate 99 bivariate responses and refit model





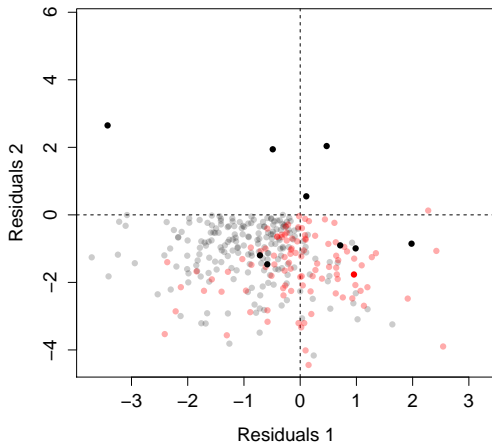
# Now we simulate

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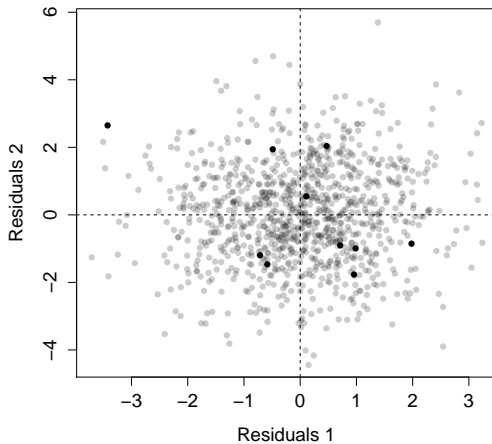
# Now we simulate

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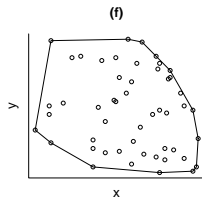
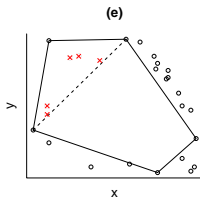
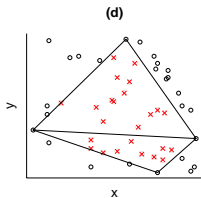
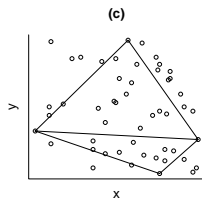
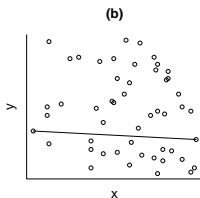
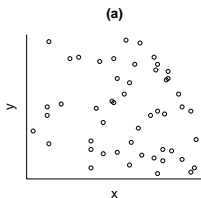
# Now we simulate

- Simulate 99 bivariate responses and refit model

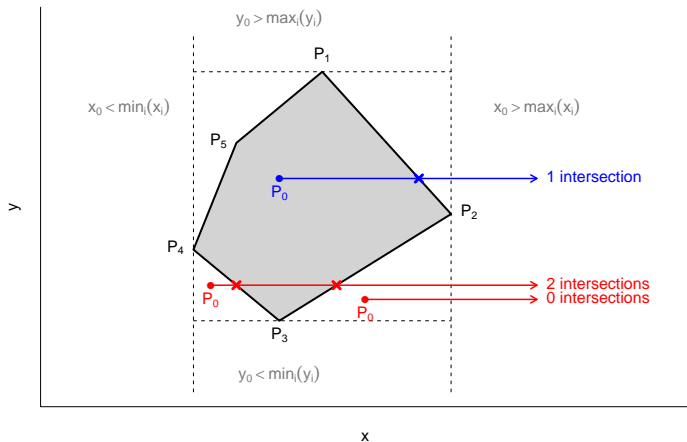


# Now we must obtain our “envelope”

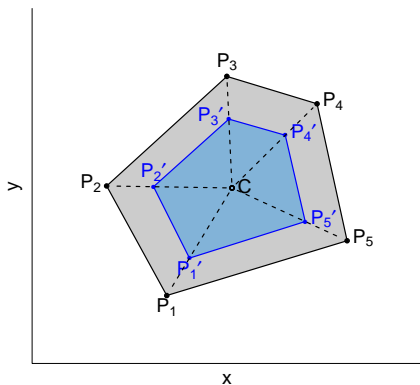
## ■ Convex Hull



# Is the point inside?

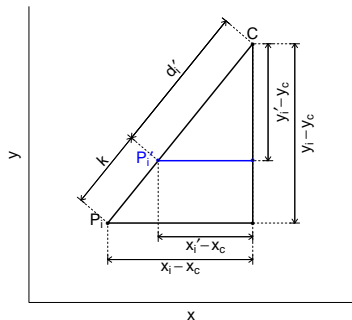


# 95% of a polygon



$$A_{\mathbf{P}} = \frac{1}{2} \left| \sum_{i=1}^{n-1} (x_i y_{i+1} - x_{i+1} y_i) + (x_n y_1 - x_1 y_n) \right| \quad A_{\mathbf{P}'} = \gamma A_{\mathbf{P}}, \quad 0 < \gamma < 1$$

# 95% of a polygon



$$\frac{x'_i - x_C}{x_i - x_C} = \frac{y'_i - y_C}{y_i - y_C} = \frac{d'_i}{d_i}$$

$$x'_i = \frac{d'_i}{d_i}(x_i - x_C) + x_C = \frac{d_i x_i - k \tilde{x}_i}{d_i}$$

$$y'_i = \frac{d'_i}{d_i}(y_i - y_C) + y_C = \frac{d_i y_i - k \tilde{y}_i}{d_i}$$

$$\tilde{x}_i = x_i - x_C \text{ and } \tilde{y}_i = y_i - y_C$$

## 95% of a polygon

$$ak^2 + bk + c = 0$$

$$a = \sum_{i=1}^{n-1} \frac{\tilde{x}_i \tilde{y}_{i+1} - \tilde{x}_{i+1} \tilde{y}_i}{d_i d_{i+1}} + \frac{\tilde{x}_n \tilde{y}_1 - \tilde{x}_1 \tilde{y}_n}{d_n d_1}$$

$$b = \sum_{i=1}^{n-1} \left\{ \frac{d_i(\tilde{x}_{i+1} y_i - x_i \tilde{y}_{i+1}) + d_{i+1}(x_{i+1} \tilde{y}_i - \tilde{x}_i y_{i+1})}{d_i d_{i+1}} \right. \\ \left. + \frac{d_n(\tilde{x}_1 y_n - x_n \tilde{y}_1) + d_1(x_1 \tilde{y}_n - \tilde{x}_n y_1)}{d_n d_1} \right\}$$

$$c = 2(A \pm \gamma A_{\mathbf{P}})$$

$$\hat{k} = \min_i \{k_i \in \mathbb{R}\}$$



# 95% of a polygon

- Another possibility is to scale the distances  $d_i$  from the centroid to the vertices to  $\sqrt{\alpha} \times d_i$  and the resulting coordinates are:

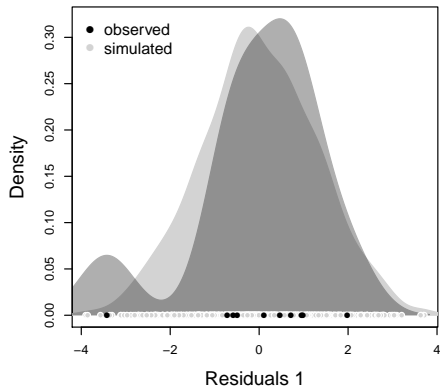
$$x_i^* = \sqrt{\alpha}(x_i - x_C) + x_C$$

$$y_i^* = \sqrt{\alpha}(y_i - y_C) + y_C$$

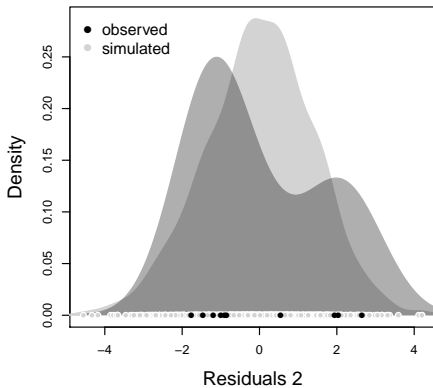
for the new polygon  $\mathbf{P}^* = \overline{P_1^* \dots P_v^*}$ , with  $P_i^* = (x_i^*, y_i^*)$ .

# Adding density plots

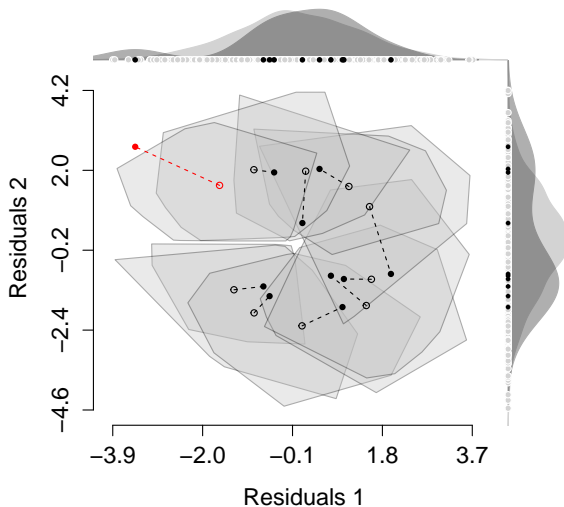
(a)



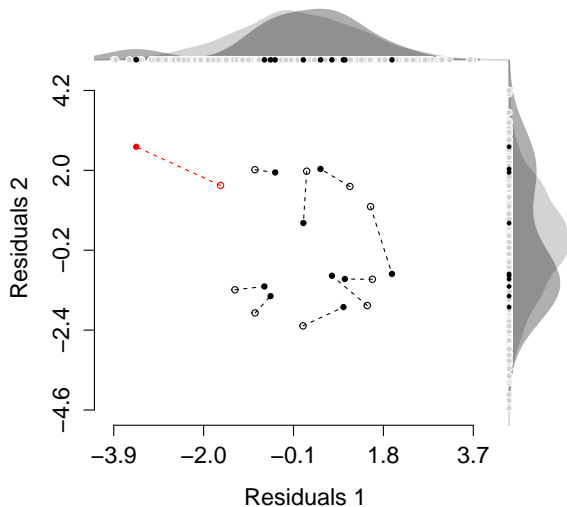
(b)



# Final display

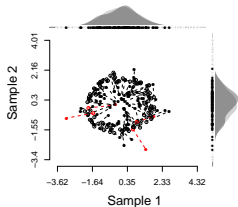


# Final display

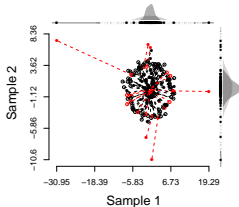


# Expected shapes

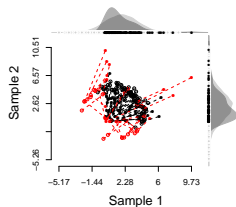
(a) Normal



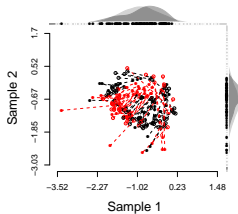
(b) Heavy-tailed



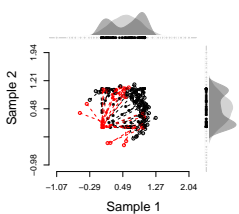
(c) Right-skewed



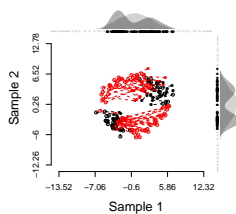
(d) Left-skewed



(e) Bimodal (beta)

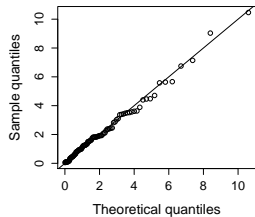


(f) Bimodal (mixture)

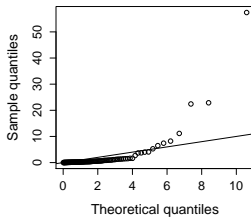


# Expected shapes

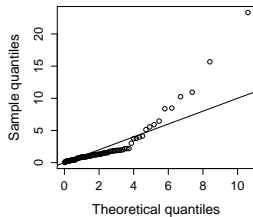
(a) Normal



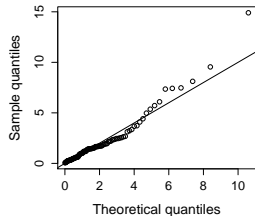
(b) Heavy-tailed



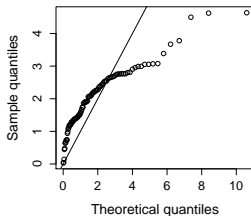
(c) Right-skewed



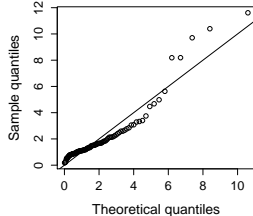
(d) Left-skewed



(e) Bimodal (beta)



(f) Bimodal (mixture)



## An example using simulated data

$$\mathbf{Y}_i = \begin{bmatrix} Y_{1i} \\ Y_{2i} \end{bmatrix} \sim N_2 \left( \begin{bmatrix} \mu_{1i} \\ \mu_{2i} \end{bmatrix}, \begin{bmatrix} \sigma_1^2 & \sigma_{12} \\ \sigma_{12} & \sigma_2^2 \end{bmatrix} \right),$$

Marginally

$$Y_{ji} \sim N(\mu_{ji}, \sigma_j^2), \quad j = 1, 2,$$

$$\text{Cov}(Y_{1i}, Y_{2i}) = \sigma_{12}$$

# Model fitting

$$\mu_{ji} = \beta_{j0} + \beta_{j1}x_i$$

$$L(\boldsymbol{\beta}, \boldsymbol{\Sigma}) = \prod_{i=1}^n (2\pi)^{-1} |\boldsymbol{\Sigma}|^{-\frac{1}{2}} \exp \left\{ -\frac{1}{2} (\mathbf{y}_i - \boldsymbol{\mu}_i)^\top \boldsymbol{\Sigma}^{-1} (\mathbf{y}_i - \boldsymbol{\mu}_i) \right\}$$



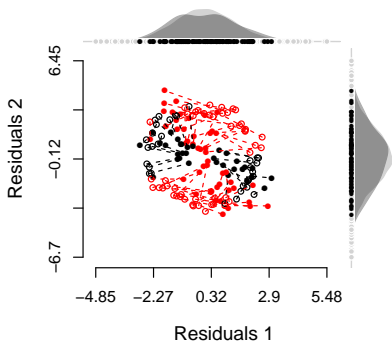
# Estimates

Table 1: Parameter estimates (standard errors) for both models fitted to the simulated correlated bivariate normal data and true values

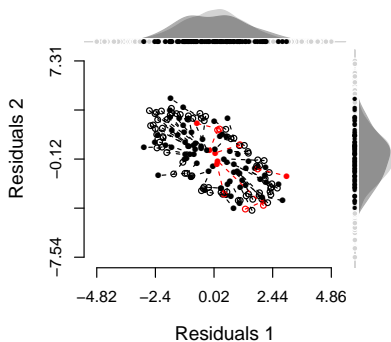
Parameter	Assuming independence	Estimating covariance	True value
$\beta_{10}$	1.50 (0.35)	1.50 (0.35)	2.00
$\beta_{11}$	0.43 (0.06)	0.43 (0.06)	0.40
$\beta_{20}$	0.78 (0.48)	0.78 (0.48)	0.20
$\beta_{21}$	0.18 (0.08)	0.18 (0.08)	0.20
$\sigma_1^2$	1.79 (0.28)	1.79 (0.28)	2.00
$\sigma_2^2$	3.48 (0.55)	3.49 (0.55)	3.00
$\sigma_{12}$	0.00 (—)	-1.61 (0.33)	-1.70
$-2 \times \loglik$	600.42	557.10	—

# Bivariate residual plots with simulation polygons

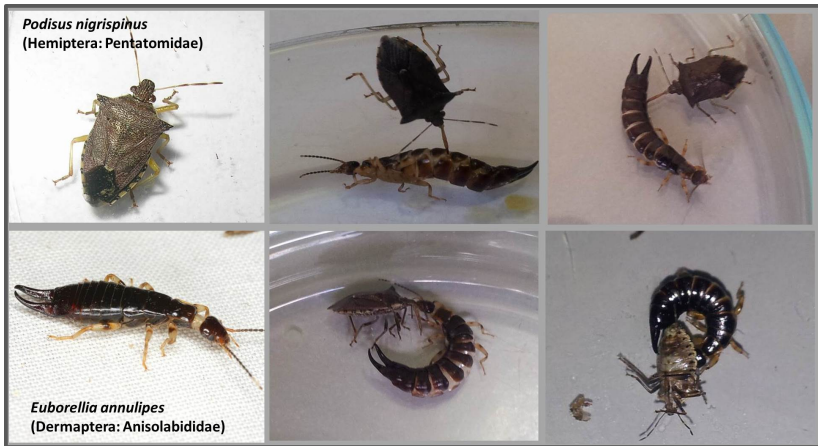
(a) No correlation



(b) Estimating correlation



# An example with real data



# Bivariate Poisson model

$$X_j \sim P(\lambda_j), \quad j = 0, 1, 2$$

$$Y_1 = X_0 + X_1$$

$$Y_2 = X_0 + X_2$$

$$(Y_1, Y_2) \sim \text{BP}(\lambda_0, \lambda_1, \lambda_2)$$

$$P(Y_1 = y_1, Y_2 = y_2) = e^{-(\lambda_0 + \lambda_1 + \lambda_2)} \frac{\lambda_1^{y_1} \lambda_2^{y_2}}{y_1! y_2!} \sum_{k=0}^{\min(y_1, y_2)} \binom{y_1}{k} \binom{y_2}{k} k! \left( \frac{\lambda_0}{\lambda_1 \lambda_2} \right)^k$$

$$Y_1 \sim P(\lambda_0 + \lambda_1)$$

$$Y_2 \sim P(\lambda_0 + \lambda_2)$$

# Model fitting

- Pseudo-likelihood maximization (Gourieroux et al., 1984)
- Newton-Raphson algorithm (Jung & Winkelmann, 1993; Kocherlakota & Kocherlakota, 2001)
- Generalized least squares (Ho & Singer, 2001)
- Bayesian methods (Tsionas, 2001)
- EM algorithm (Karlis & Ntzoufras, 2005)

# Model fitting

- Complete-data log-likelihood

$$\begin{aligned}l(\lambda_0, \lambda_1, \lambda_2) &= \sum_{i=1}^n \log\{P(X_{0i} = x_{0i}, X_{1i} = x_{1i}, X_{2i} = x_{2i})\} \\ &= \sum_{i=1}^n \log\{P(X_{0i} = x_{0i})P(X_{1i} = x_{1i})P(X_{2i} = x_{2i})\}\end{aligned}$$

- Maximisation is straightforward by fitting three independent Poisson GLMs
- One to variable  $x_{1i} = y_{1i} - x_{0i}$ , another to variable  $x_{2i} = y_{2i} - x_{0i}$ , and to variable  $x_{0i}$ , replaced by its conditional expectation  $z_i$

$$\begin{aligned}z_i &= E(X_{0i} | Y_{1i}, Y_{2i}) \\ &= \sum_{x_{0i}=0}^{\infty} x_{0i} P(X_{0i} = x_{0i} | Y_{1i} = y_{1i}, Y_{2i} = y_{2i}) \\ &= \lambda_0 \frac{P(Y_{1i} = y_{1i} - 1, Y_{2i} = y_{2i} - 1)}{P(Y_{1i} = y_{1i}, Y_{2i} = y_{2i})}\end{aligned}$$

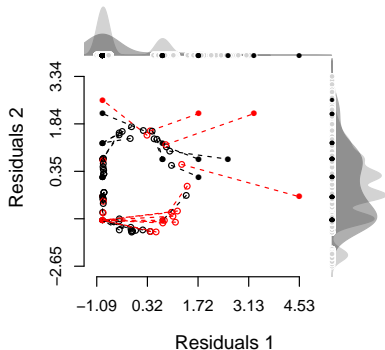
# Estimates

Table 2: Parameter estimates (standard errors) for both models fitted to the attack data

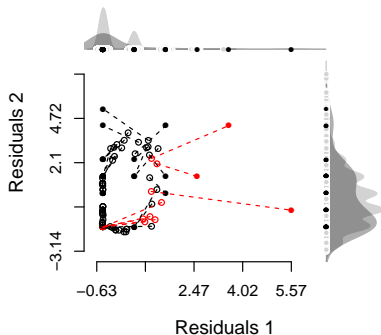
Parameter	Assuming independence	Estimating covariance
$\log \lambda_1$ (stinkbug)	-0.85 (0.20)	-1.43 (0.04)
$\log \lambda_2$ (earwig)	1.00 (0.08)	0.93 (0.00)
$\log \lambda_0$	0.00 (-)	-1.66 (0.05)
$-2 \times \log \text{lik}$	332.32	327.83

# Bivariate residual plots with simulation polygons

(a) No correlation



(b) Estimating correlation





# Using PIT diagnostics

- Randomized PIT diagnostics for discrete models

$$r_{ji}^{rand.pit} = F(y_{ji} - 1; \hat{\theta}_{ji}) + u_i \{F(y_{ji}; \hat{\theta}_{ji}) - F(y_{ji} - 1; \hat{\theta}_{ji})\},$$

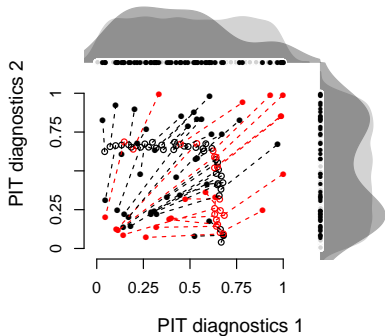
with  $F(-1) = 0$ , and  $u_i$  is a realization of  $U_i \sim \text{Uniform}(0, 1)$ .

- For the bivariate Poisson model:

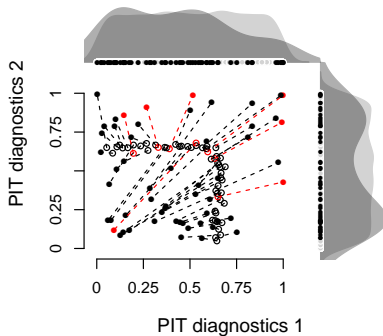
$$F(y_{ji}; \hat{\theta}_{ji}) = e^{-\hat{\mu}_j} \sum_{k=0}^{y_{ji}} \frac{\hat{\mu}_j^k}{k!}$$

# Bivariate residual plots with simulation polygons

(c) No correlation

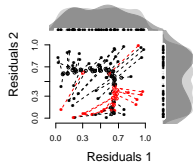


(d) Estimating correlation

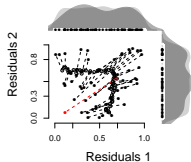


# Line-up test

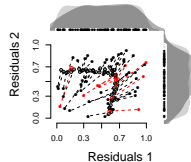
Suspect: 1



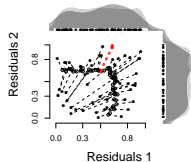
Suspect: 2



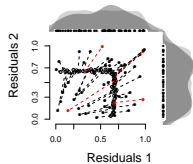
Suspect: 3



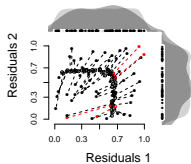
Suspect: 4



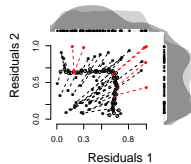
Suspect: 5



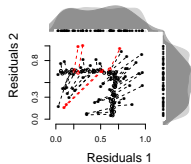
Suspect: 6



Suspect: 7



Suspect: 8



# Code availability

- bivrp package on CRAN

[bivrp-package](#)

[add.dplots.plot](#)

[add.dplots.prep](#)

[bivrp](#)

[chp.perpoint](#)

[get.k](#)

[get.newpolygon](#)

[is.point.inside](#)

[plot.bivrp](#)

[polygon.area](#)

[sorttheta](#)

Bivariate Residual Plots with Simulation Polygons

Internal functions to prepare 'bivrp' objects

Internal functions to prepare 'bivrp' objects

Bivariate Residual Plots with Simulation Polygons

Internal functions to prepare 'bivrp' objects

Polygon operations

Polygon operations

Determine if point is inside or outside a simple polygon area

Plot Method for bivrp Objects

Polygon operations

Internal functions to prepare 'bivrp' objects

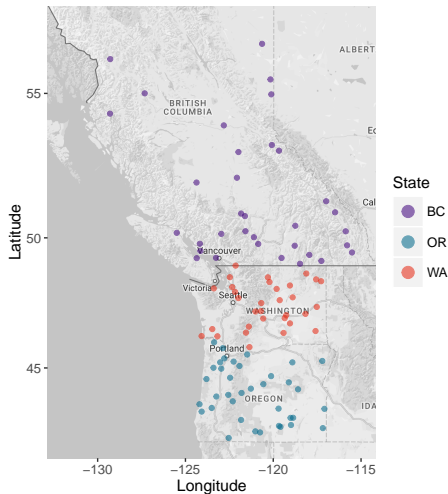
# Final considerations

- This is not a formal test!
- Simple tool for assessing goodness-of-fit of bivariate models
- Use of different diagnostics is recommended (e.g. PIT diagnostics)
- Drawbacks include computational burden for complex models and the way outliers may influence convex hulls
- Problematic extension to big data
- Extension to the  $n$ -variate setting
- Complementary and (hopefully) helpful approach

But what about the eagles?



# Back to our first motivation example



# Imperfect detection

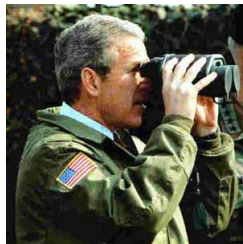




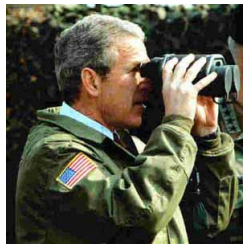
# Imperfect detection



# Imperfect detection



# Imperfect detection



Can you spot the cat?



# Can you spot the cat?



# Case study



Route	Stops 1-10	Stops 11-20	Stops 21-30	Stops 31-40	Stops 41-50
4	0	0	0	1	4
6	0	0	0	1	0
16	0	0	0	1	0
17	0	0	0	0	0
...					
407	0	0	0	1	5
409	0	0	0	0	3
...					



Route	Stops 1-10	Stops 11-20	Stops 21-30	Stops 31-40	Stops 41-50
4	0	0	1	0	4
6	0	0	0	0	0
16	0	0	0	0	0
17	0	2	4	0	0
...					
407	0	0	0	1	0
409	1	0	0	0	7
...					

# Joint model formulation (Moral et al., 2018)

- Bivariate N-mixture model

$$\begin{aligned} Y_{1it} | N_{1i} &\sim \text{Binomial}(N_{1i}, p_{1it}) \\ N_{1i} &\sim \text{Poisson}(\lambda_{1i}) \text{ or } \text{NB}(\lambda_{1i}, \phi_1) \end{aligned}$$

$$\begin{aligned} Y_{2it} | N_{1i}, N_{2i} &\sim \text{Binomial}(N_{2i}, p_{2it}) \\ N_{2i} | N_{1i} &\sim \text{Poisson}(\psi_i + \lambda_{2i} N_{1i}) \text{ or } \text{NB}(\psi_i + \lambda_{2i} N_{1i}, \phi_2) \end{aligned}$$

## Assumptions

- independence among sites
- closed population; no migration
- $\lambda_{2i} = 0 \Rightarrow$  no correlation between species

# Joint N-mixture model

- Likelihood:

$$L(\mathbf{p}_1, \mathbf{p}_2, \boldsymbol{\theta}_1, \boldsymbol{\theta}_2, \boldsymbol{\psi} | \{y_{1it}\}, \{y_{2it}\}) =$$
$$\prod_{i=1}^R \left\{ \sum_{n_{1i}=\max_t\{y_{1it}\}}^{\infty} \left[ \prod_{t=1}^T \text{Bin}(y_{1it}; n_{1i}, p_{1it}) \right] f_{N_{1i}}(n_{1i}; \boldsymbol{\theta}_{1i}) \times \right.$$
$$\left. \sum_{n_{1i}=\max_t\{y_{1it}\}}^{\infty} f_{N_{1i}}(n_{1i}; \boldsymbol{\theta}_{1i}) \sum_{n_{2i}=\max_t\{y_{2it}\}}^{\infty} \left[ \prod_{t=1}^T \text{Bin}(y_{2it}; n_{2i}, p_{2it}) \right] f_{N_{2i}}(n_{2i}; \boldsymbol{\psi}_i, \boldsymbol{\theta}_{2i}) \right\}$$



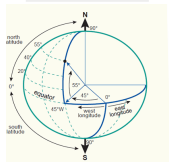
**Total variation in *observed* eagle abundance**

Total variation in *observed* eagle abundance

Habitat suitability



Geographical coordinates



Total variation in *observed* eagle abundance

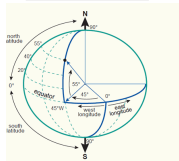
Habitat suitability

Environmental  
covariates

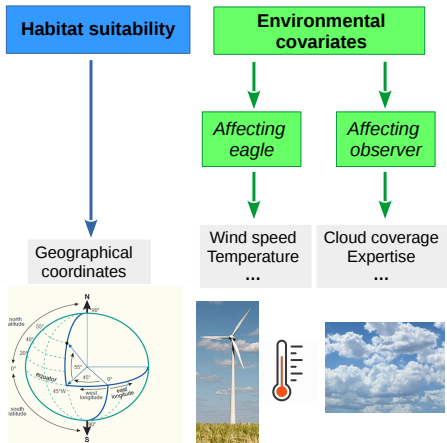
Geographical  
coordinates

Wind speed  
Temperature  
...

*Affecting  
eagle*



Total variation in *observed* eagle abundance

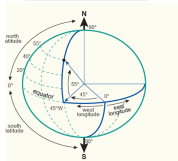


Total variation in *observed* eagle abundance

Habitat suitability



Geographical coordinates



Environmental covariates



Affecting eagle

Affecting observer



Wind speed  
Temperature  
...

Cloud coverage  
Expertise  
...



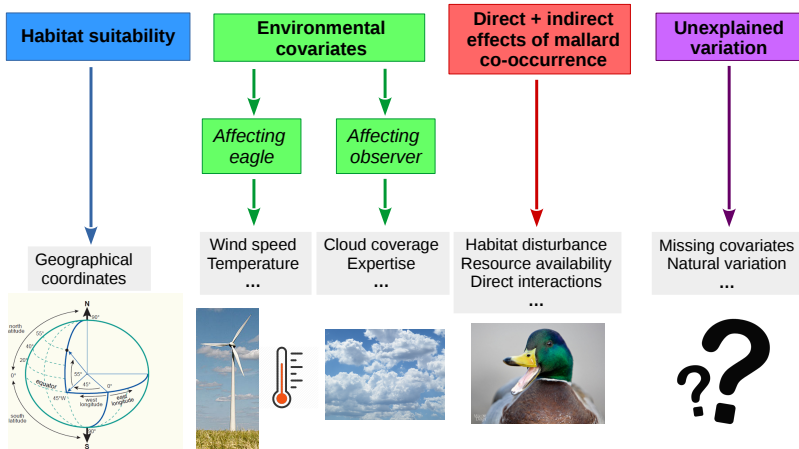
Unexplained variation



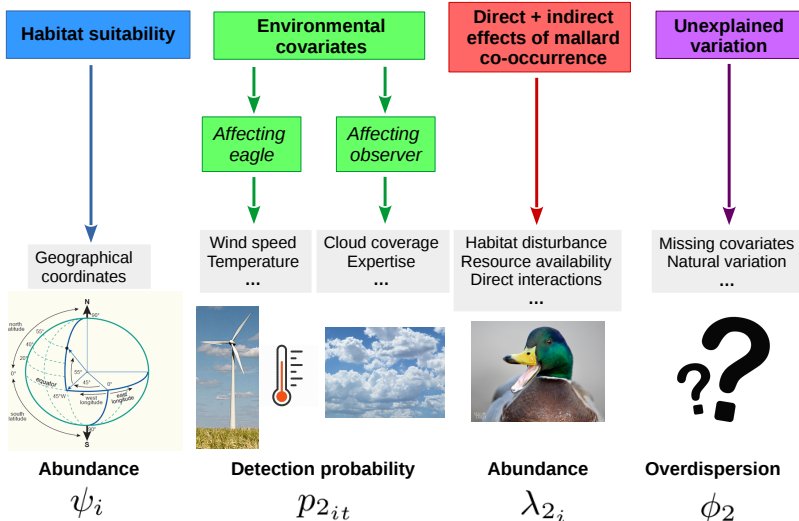
Missing covariates  
Natural variation  
...



Total variation in *observed* eagle abundance



Total variation in *observed* eagle abundance

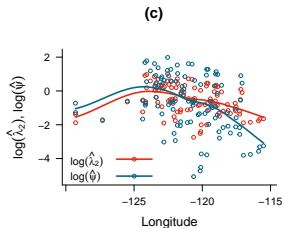
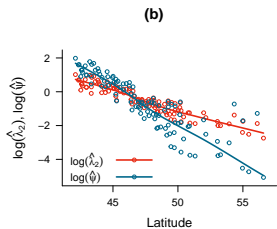
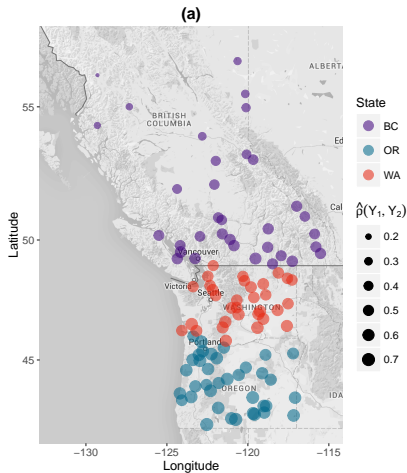


# Case study: Model selection

Parameter	Univariate models				Joint models			
	P-P	NB-P	P-NB	NB-NB	P-P	NB-P	P-NB	NB-NB
<b>Detection</b>								
(Mallard - $p_{1it}$ )								
Intercept	-4.49	-3.17	-4.49	-3.17	-2.69	-3.14	-4.49	-3.15
Temperature	0.25	0.36	0.25	0.36	0.28	0.36	0.25	0.36
Wind speed	-0.15	-0.28	-0.15	-0.28	-0.11	-0.29	-0.15	-0.28
<b>(Bald eagle - <math>p_{2it}</math>)</b>								
Intercept	-4.09	-4.09	-2.88	-2.88	-3.07	-2.87	-2.89	-2.88
Temperature	0.00	0.00	-0.15	-0.15	-0.04	-0.13	-0.19	-0.11
Wind speed	0.14	0.14	0.34	0.34	0.11	0.33	0.34	0.37
<b>Abundance</b>								
<b>(Mallard - <math>\lambda_{1i}</math>)</b>								
Intercept	3.92	2.72	3.92	2.72	2.10	2.63	3.92	2.67
Latitude	0.22	0.06	0.22	0.06	0.37	0.05	0.22	0.05
Longitude	0.32	0.17	0.32	0.17	0.30	0.15	0.32	0.15
Lat $\times$ Long	0.04	0.09	0.04	0.09	0.14	0.08	0.04	0.09
<b>(Bald eagle - <math>\psi_i</math>)</b>								
Intercept	3.22	3.22	2.14	2.14	-22.50	-0.76	2.04	-0.48
Latitude	-0.86	-0.86	-0.86	-0.86	-1.86	-1.59	-0.99	-1.65
Longitude	-0.13	-0.13	-0.23	-0.23	12.37	-0.67	-0.42	-0.94
Lat $\times$ Long	-0.12	-0.12	0.13	0.13	0.91	-0.16	0.05	-0.01
<b>(Bald eagle - <math>\lambda_{2i}</math>)</b>								
Intercept	—	—	—	—	0.12	-0.53	-28.96	-0.69
Latitude	—	—	—	—	-1.37	-0.83	-6.58	-0.78
Longitude	—	—	—	—	-0.59	-0.34	13.05	-0.27
Lat $\times$ Long	—	—	—	—	-0.31	0.04	3.49	0.07
<b>(Dispersion)</b>								
$\phi_1$	—	0.51	—	0.51	—	0.39	—	0.46
$\phi_2$	—	—	0.46	0.46	—	—	0.48	1.71
$-2 \times \log \text{lik}$	3350.28	2986.90	2994.67	2631.29	3196.88	2620.96	2990.69	2617.46

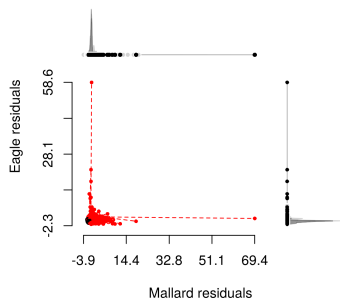


# Results



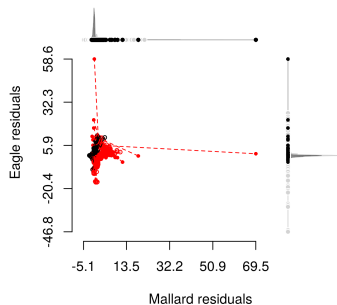
# Case study: bivariate residual plots

No correlation



368/515 (71%)

Joint model



197/515 (38%)

Thank you!



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