Diagnostic Plots for Univariate and Bivariate Models (and Joint Models in Ecology)

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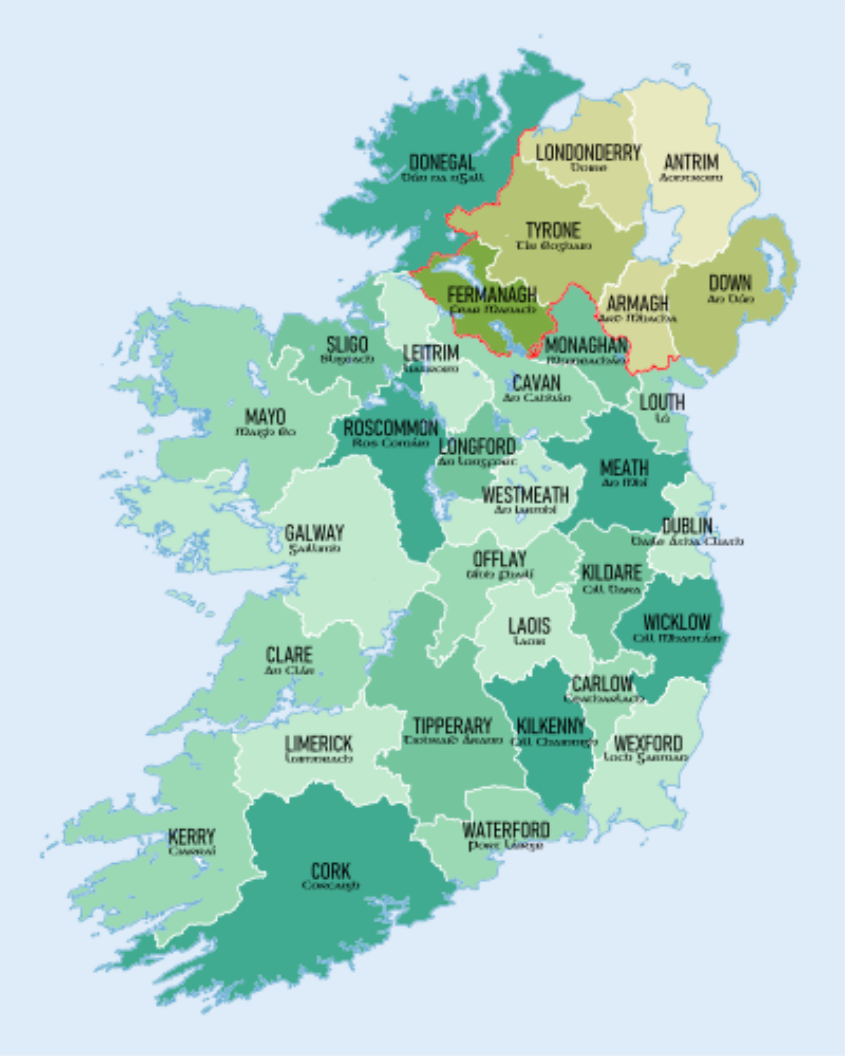
Warsaw University of Technology – December 2018

Ireland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KILDARE
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CARLOW
TIPPERARY
KILKENNY
WEXFORD
LIMERICK
WATERFORD
KERRY
CORK

Maynooth University

Maynooth University

Maynooth University

- Department of Mathematics and Statistics
- Currently in the Statistics group – 5 lecturers/1 professor, 5 PhD students, 1 post-doc
- Statistics applied to agriculture, biological control, ecology, climatology, methods for data visualisation, Bayesian methods, machine learning
- New programmes in Data Science and Analytics
- Involvement with the Hamilton Institute (Maynooth University), and Teagasc (Agriculture and Food Development Authority)

Outline

- Motivation Example – Bald Eagle Data
- Half-normal Plots with Simulation Envelopes
- Bivariate Plots with Simulation Polygons
- Example 1 – Bivariate Normal
- Example 2 – Bivariate Poisson
- Analysis of the Bald Eagle Data

Case study

- Bald eagles and mallards

Case study

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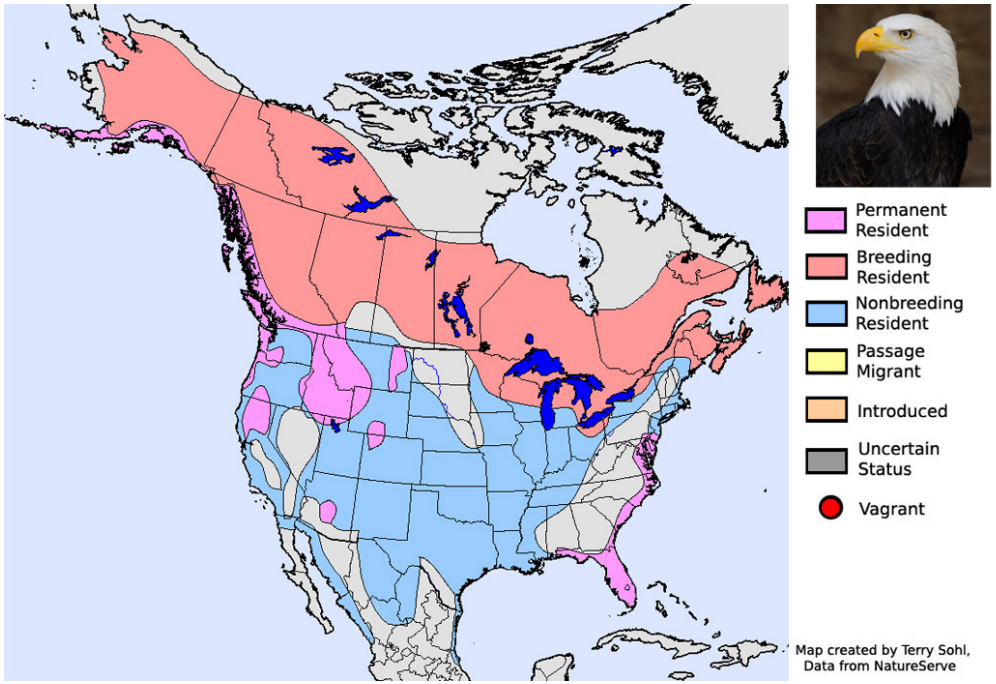
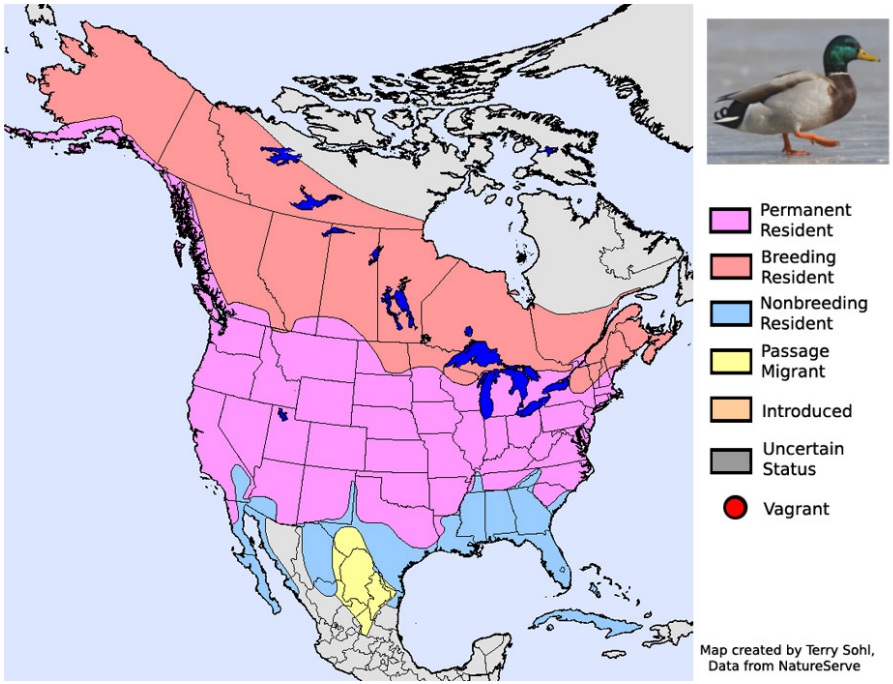
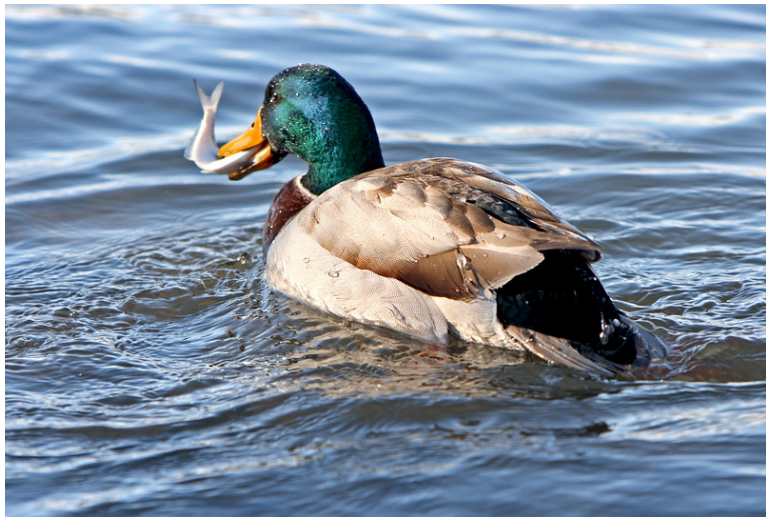
Case study





(Ridgely et al., 2003)

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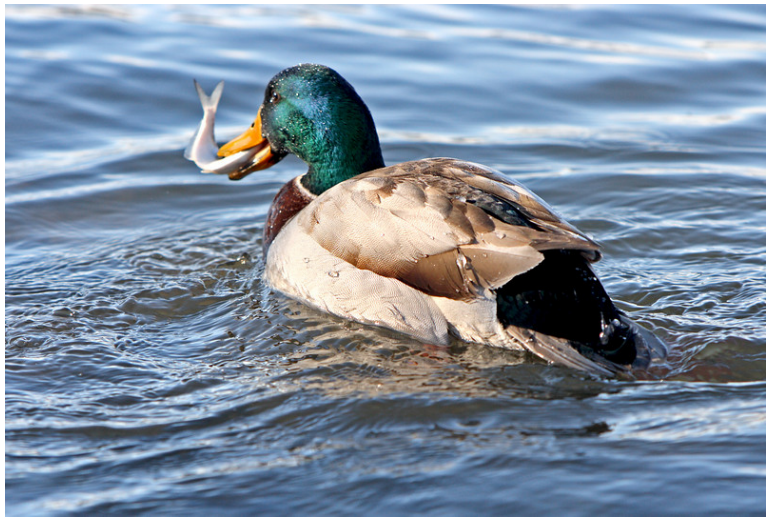

- How should we assess goodness-of-fit for joint models?

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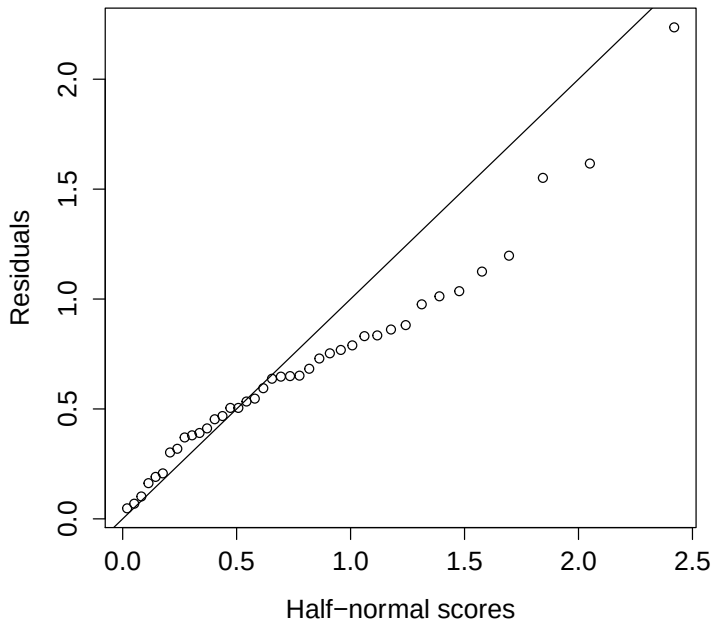
- Half-normal plots with simulation envelopes
- Ordered absolute values of a model diagnostic vs. expected order statistics of a half-normal distribution $\Phi^{-1}\left(\frac{i+n-\frac{1}{8}}{2n+\frac{1}{2}}\right)$



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

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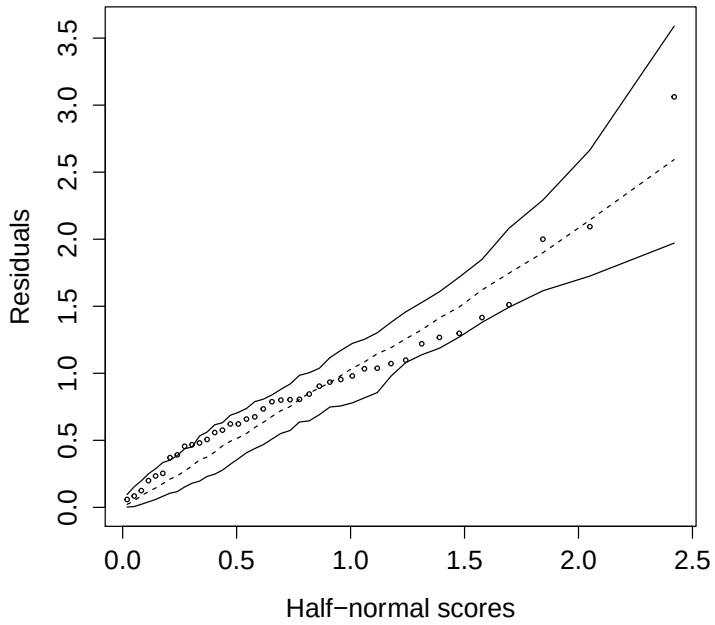

How can we do this for a bivariate model?

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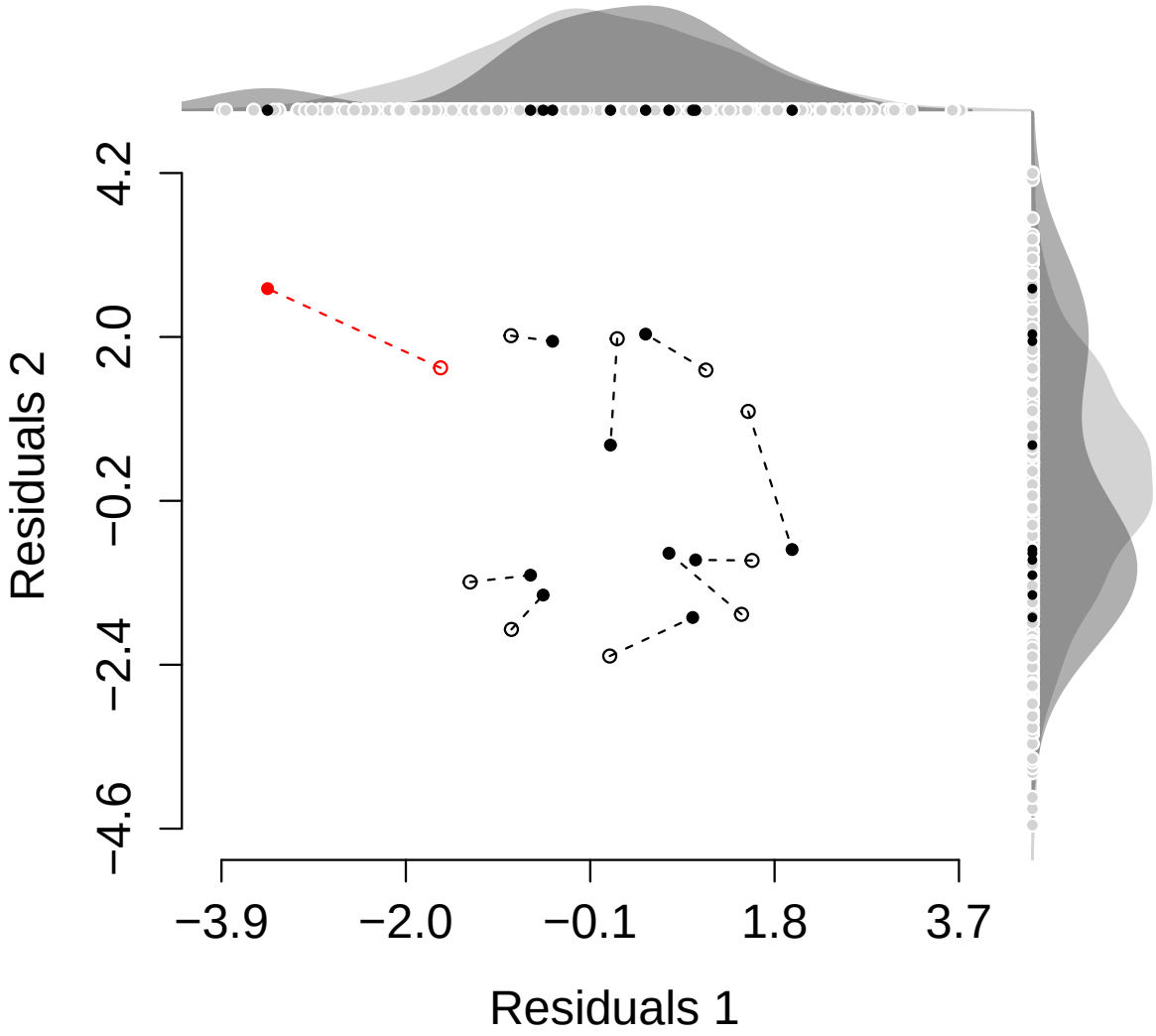
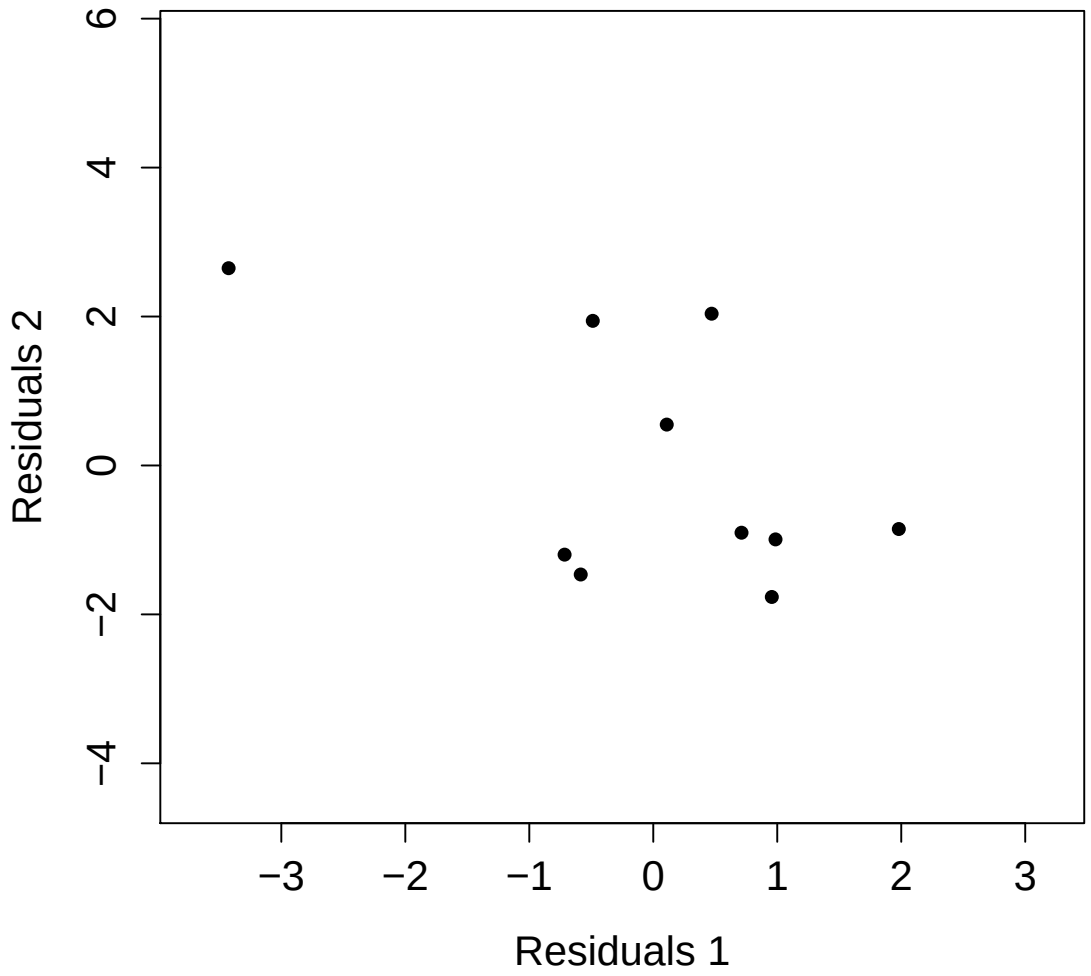
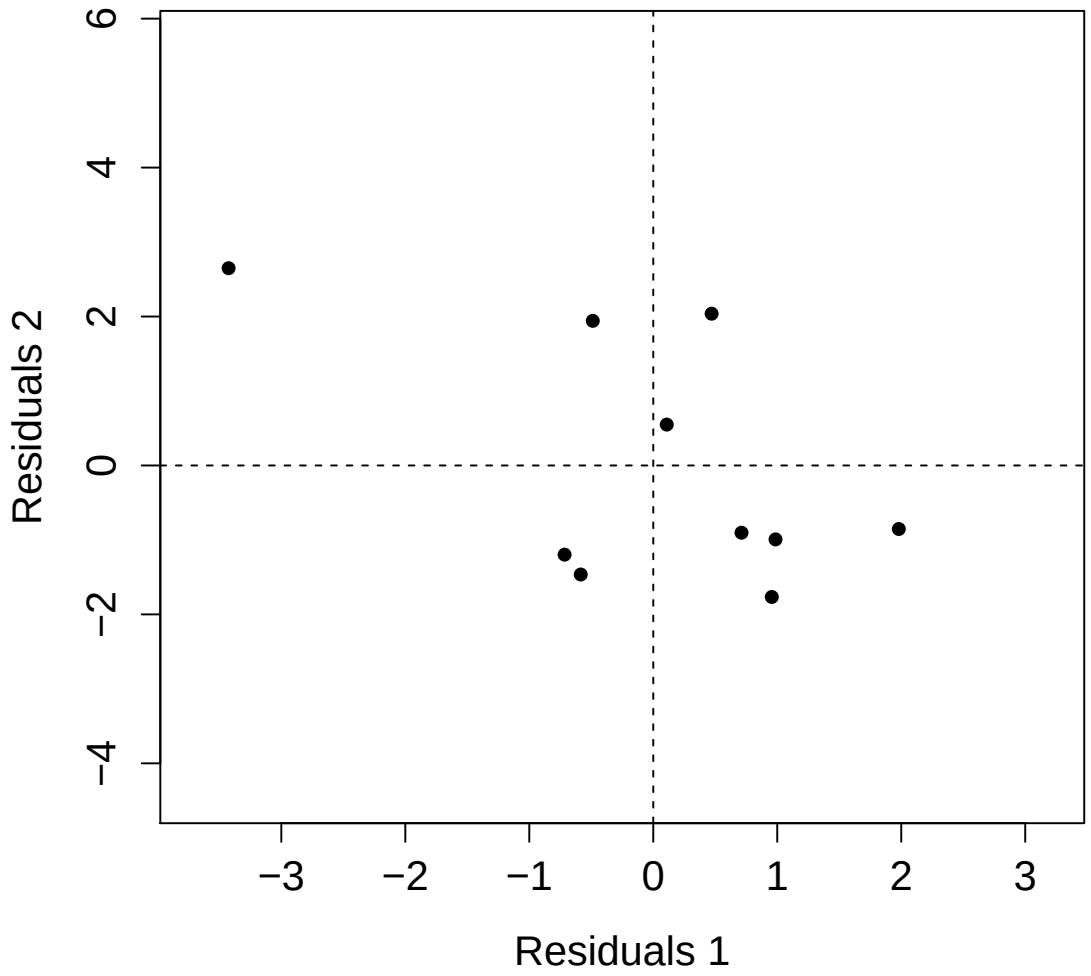


Bivariate residuals



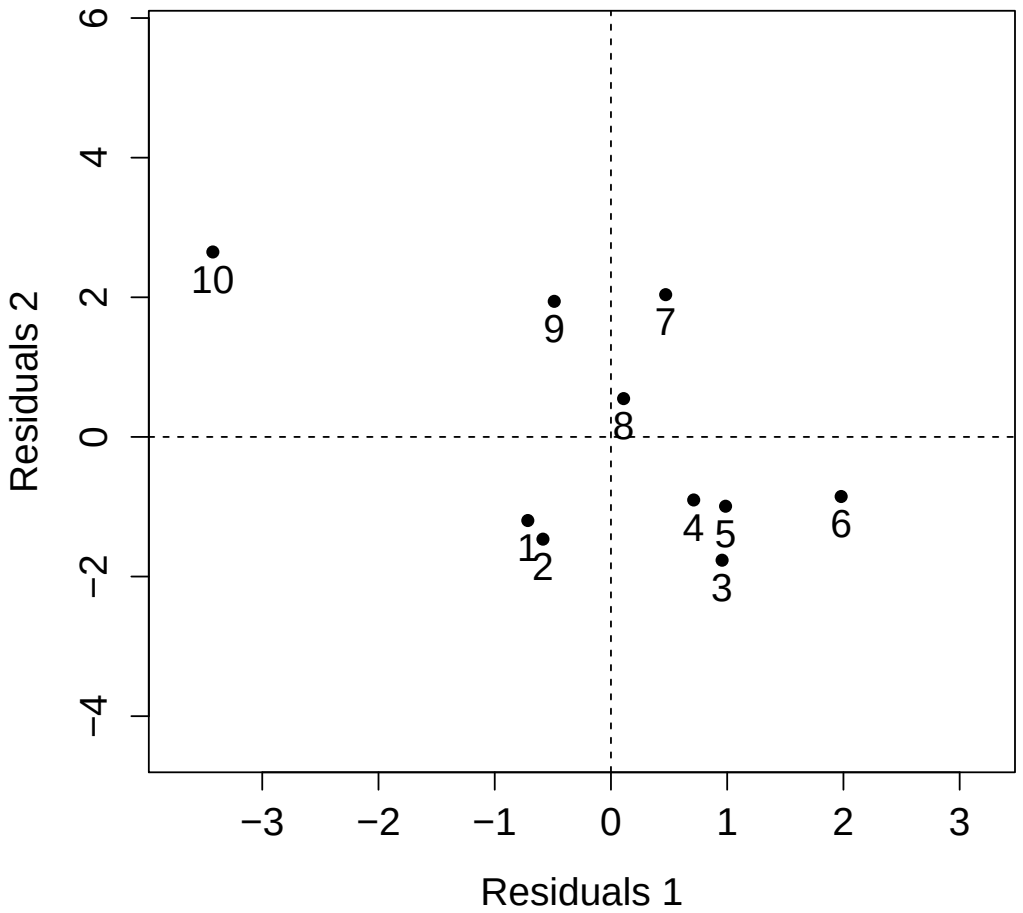
We need to order them



Ordering by angles

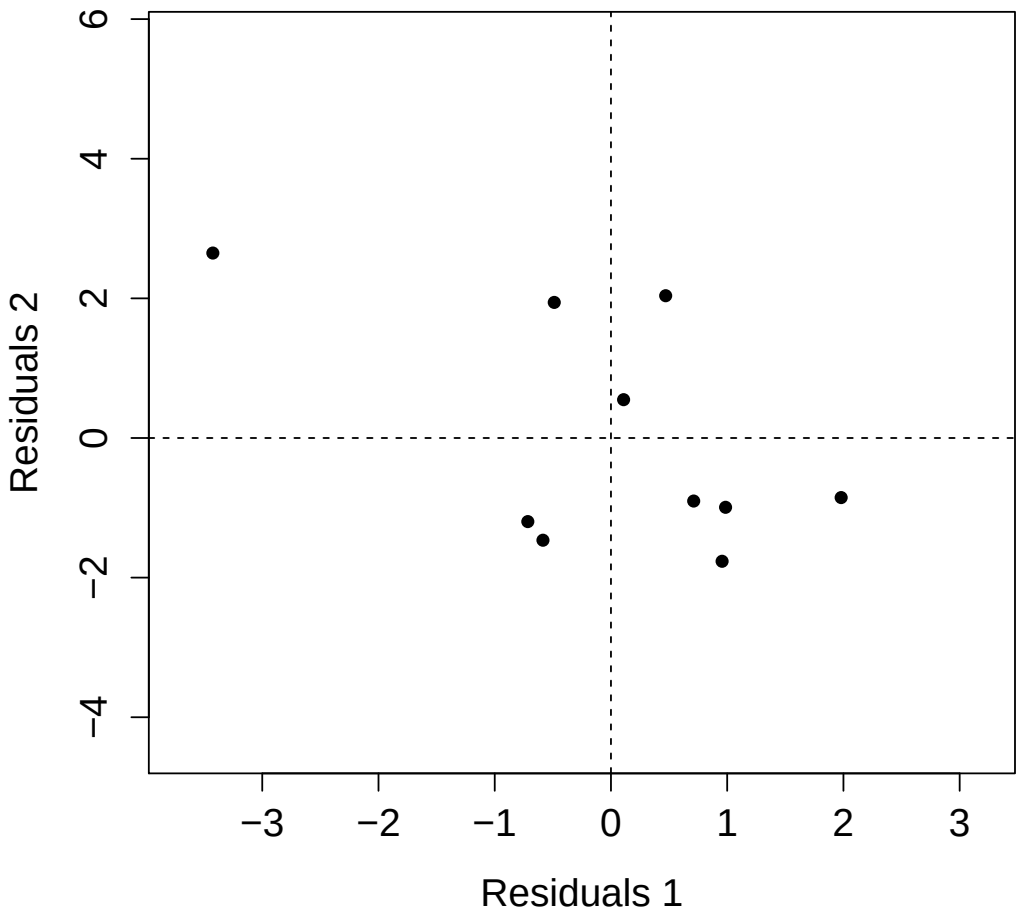
$$\alpha_i = \begin{cases} \tan^{-1}\left(\frac{y_i}{x_i}\right), & x > 0 \\ \tan^{-1}\left(\frac{y_i}{x_i}\right) + \pi, & x < 0 \text{ and } y \geq 0 \\ \tan^{-1}\left(\frac{y_i}{x_i}\right) + \pi, & x < 0 \text{ and } y < 0 \\ \pm\frac{\pi}{2}, & x = 0 \text{ and } y \gtrless 0 \\ \text{undefined},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

Ordering by angles



Now we simulate

- Simulate 99 bivariate responses and refit model



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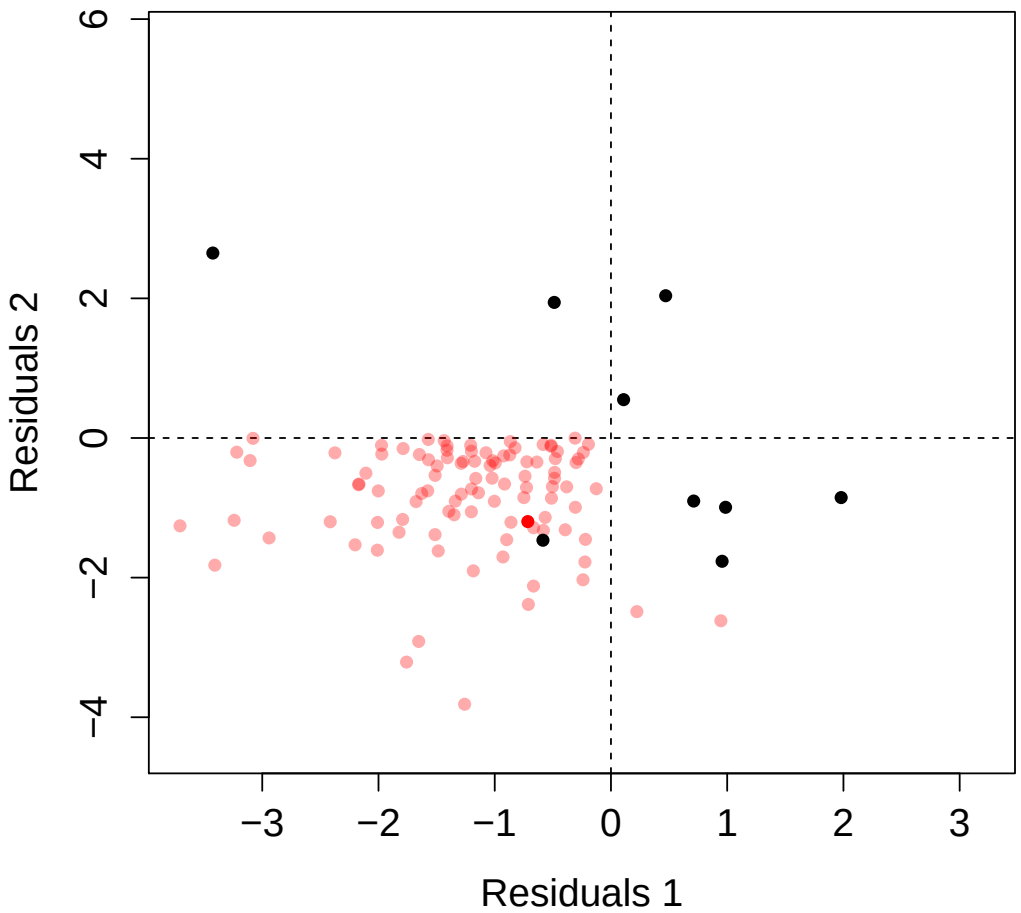

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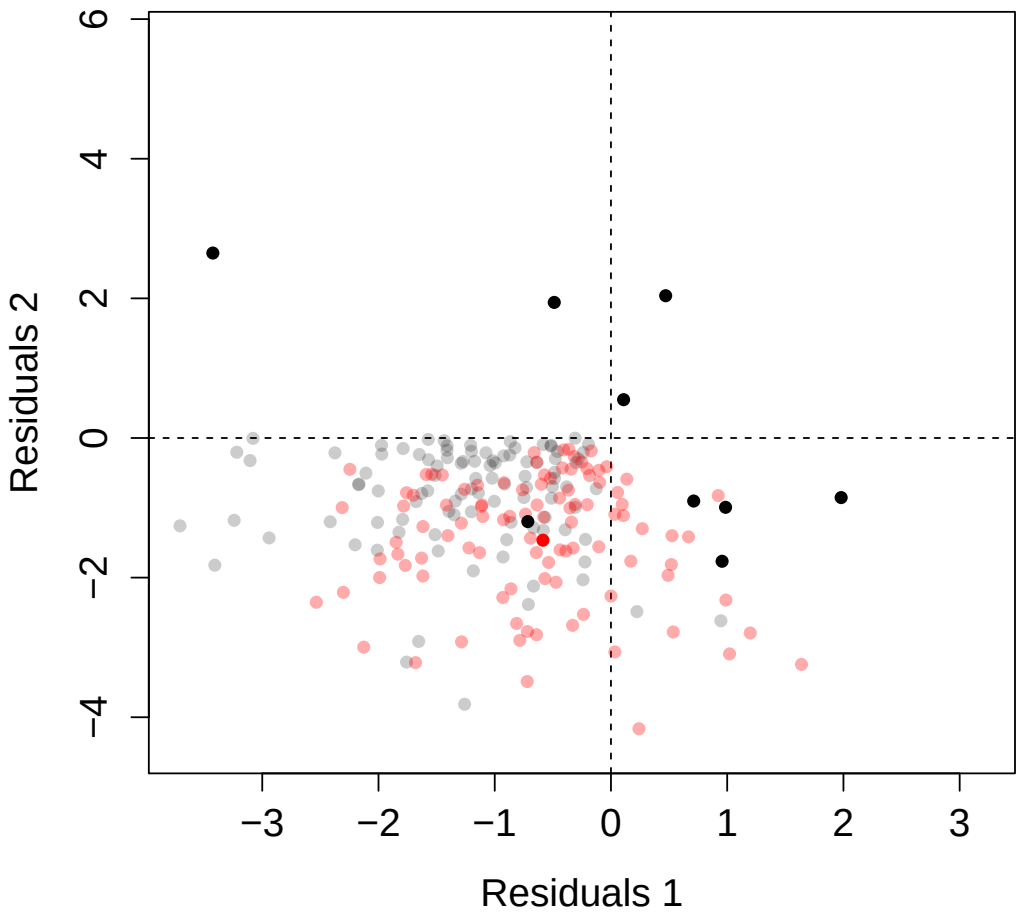

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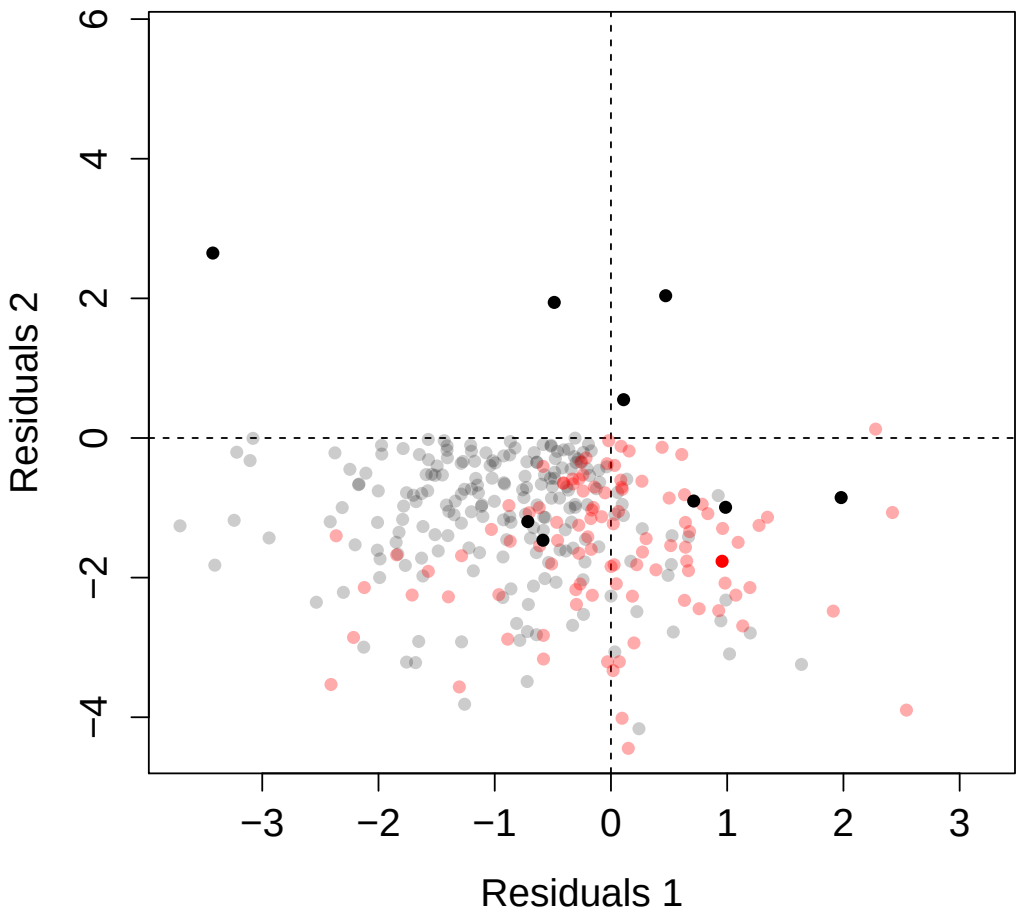

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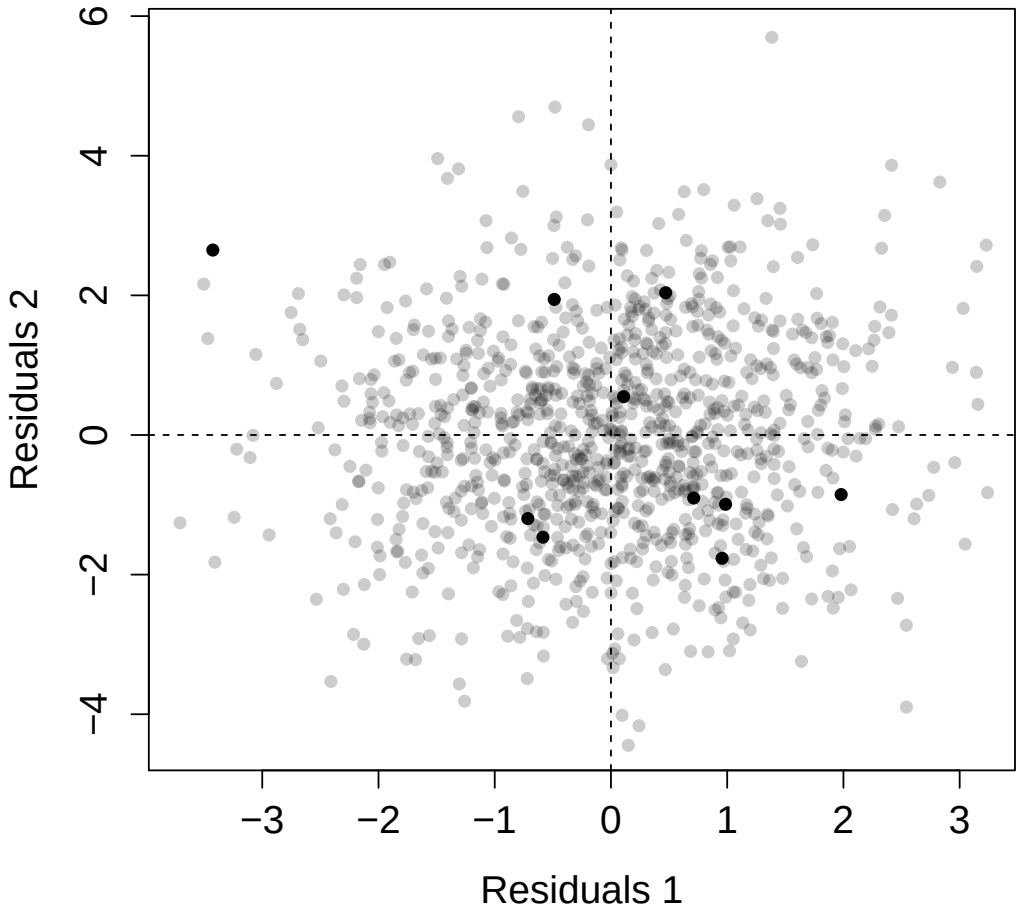
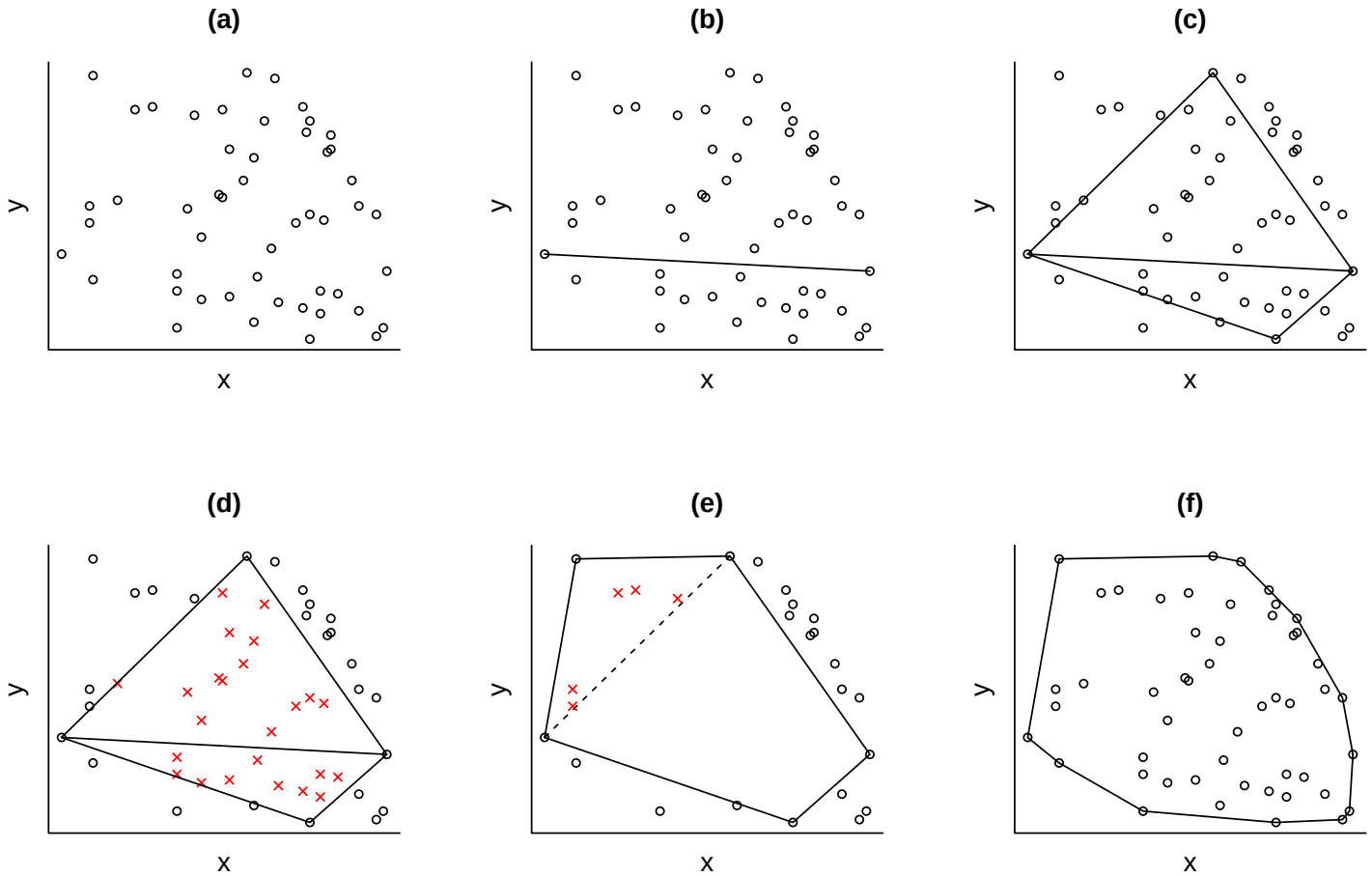
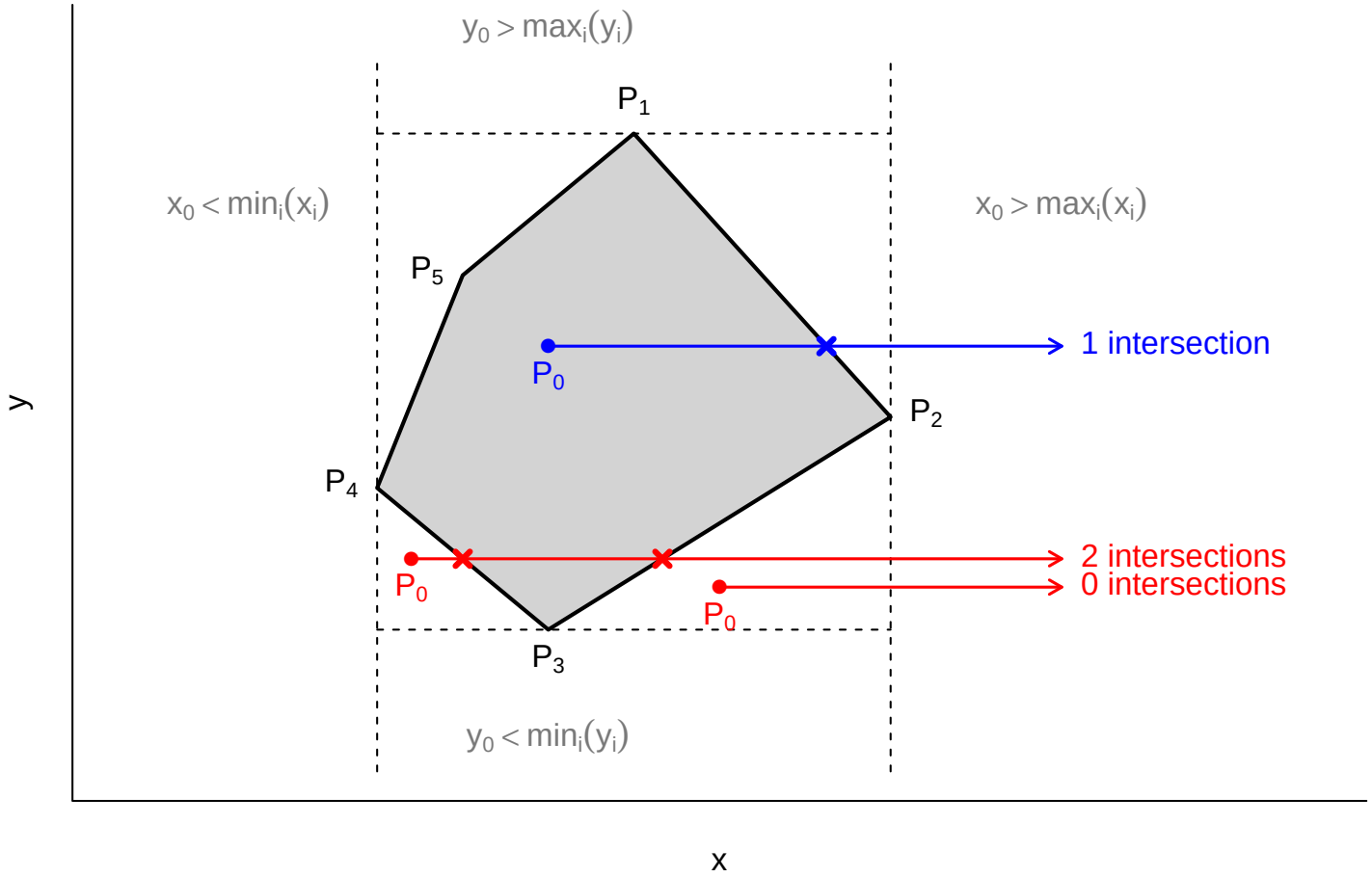


Now we must obtain our “envelope”

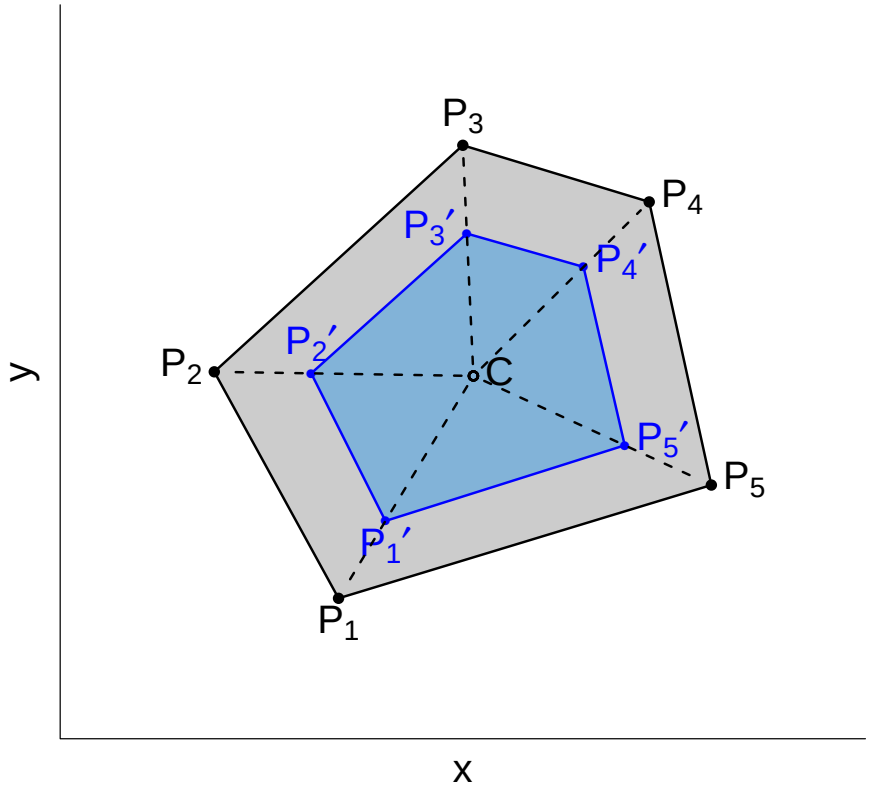
- Convex Hull



Is the point inside?

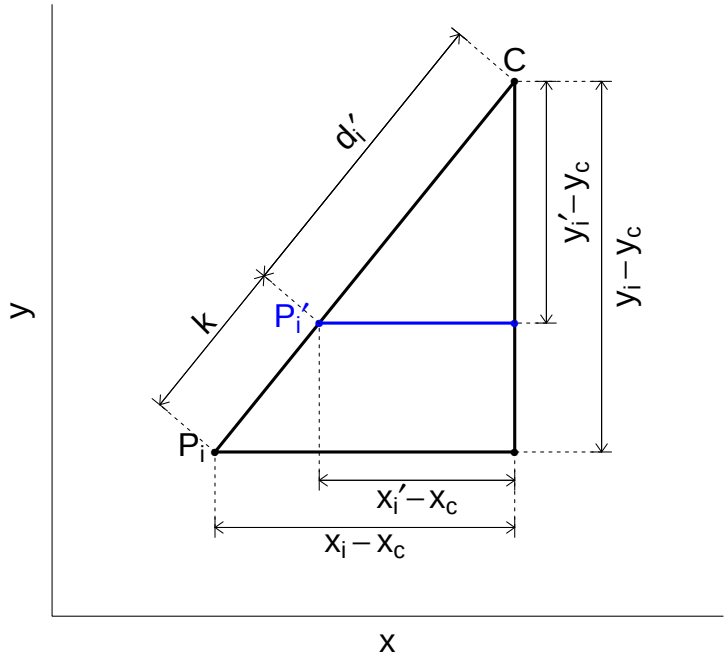


95% of a polygon



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

95% of a polygon



$$\frac{x_i' - x_C}{x_i - x_C} = \frac{y_i' - y_C}{y_i - y_C} = \frac{d_i'}{d_i}$$

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

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

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

95% of a polygon

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

$$\begin{aligned}
a &= \sum_{i=1}^{n-1} \frac{\tilde{x}_i \tilde{y}_{i+1} - \tilde{x}_{i+1}\tilde{y}_i}{d_i d_{i+1}} + \frac{\tilde{x}_n \tilde{y}_1 - \tilde{x}_1 \tilde{y}_n}{d_n d_1} \\
b &= \sum_{i=1}^{n-1} \left\{ \frac{d_i(\tilde{x}_{i+1} y_i - x_i \tilde{y}_{i+1}) + d_{i+1}(x_{i+1}\tilde{y}_i - \tilde{x}_i y_{i+1})}{d_i d_{i+1}} \right. \\
&\qquad \left. + \frac{d_n(\tilde{x}_1 y_n - x_n \tilde{y}_1) + d_1(x_1 \tilde{y}_n - \tilde{x}_n y_1)}{d_n d_1} \right\} \\
c &= 2(A \pm \gamma A_{\mathbf{P}})
\end{aligned}$$

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

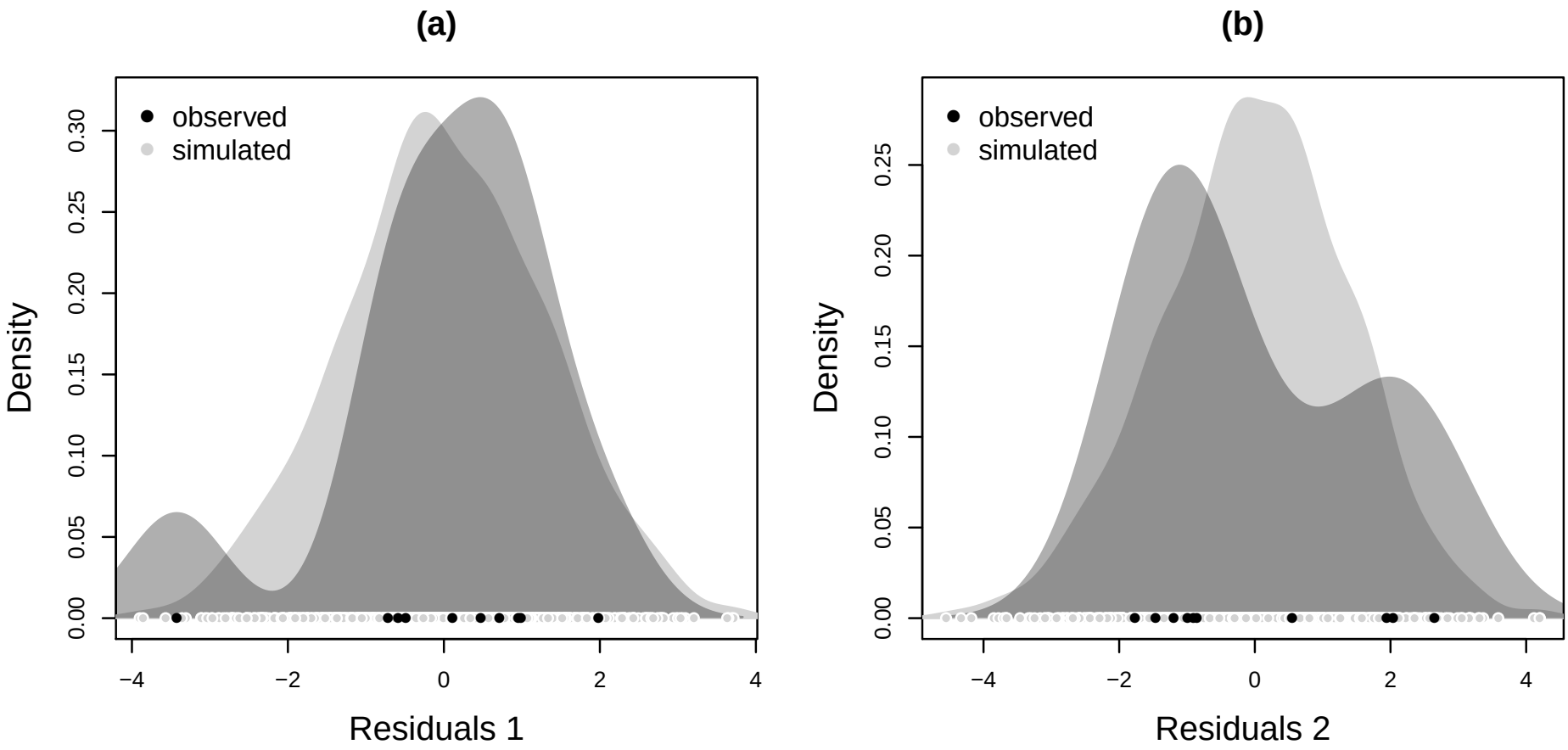
95% of a polygon

- 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:

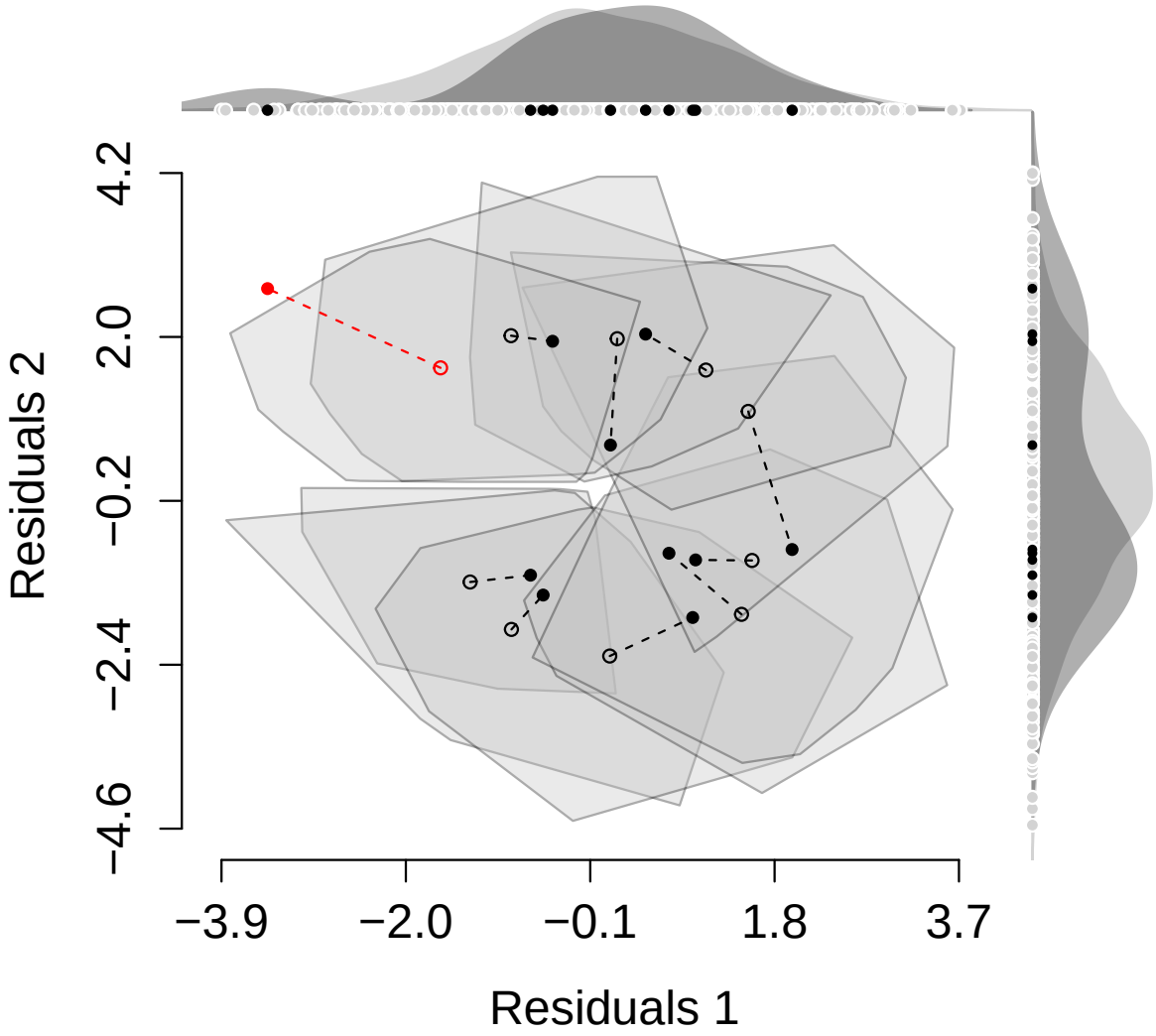
$$\begin{aligned} x_i^* &= \sqrt{\alpha}(x_i - x_C) + x_C \\ y_i^* &= \sqrt{\alpha}(y_i - y_C) + y_C \end{aligned}$$

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

Adding density plots



Final display



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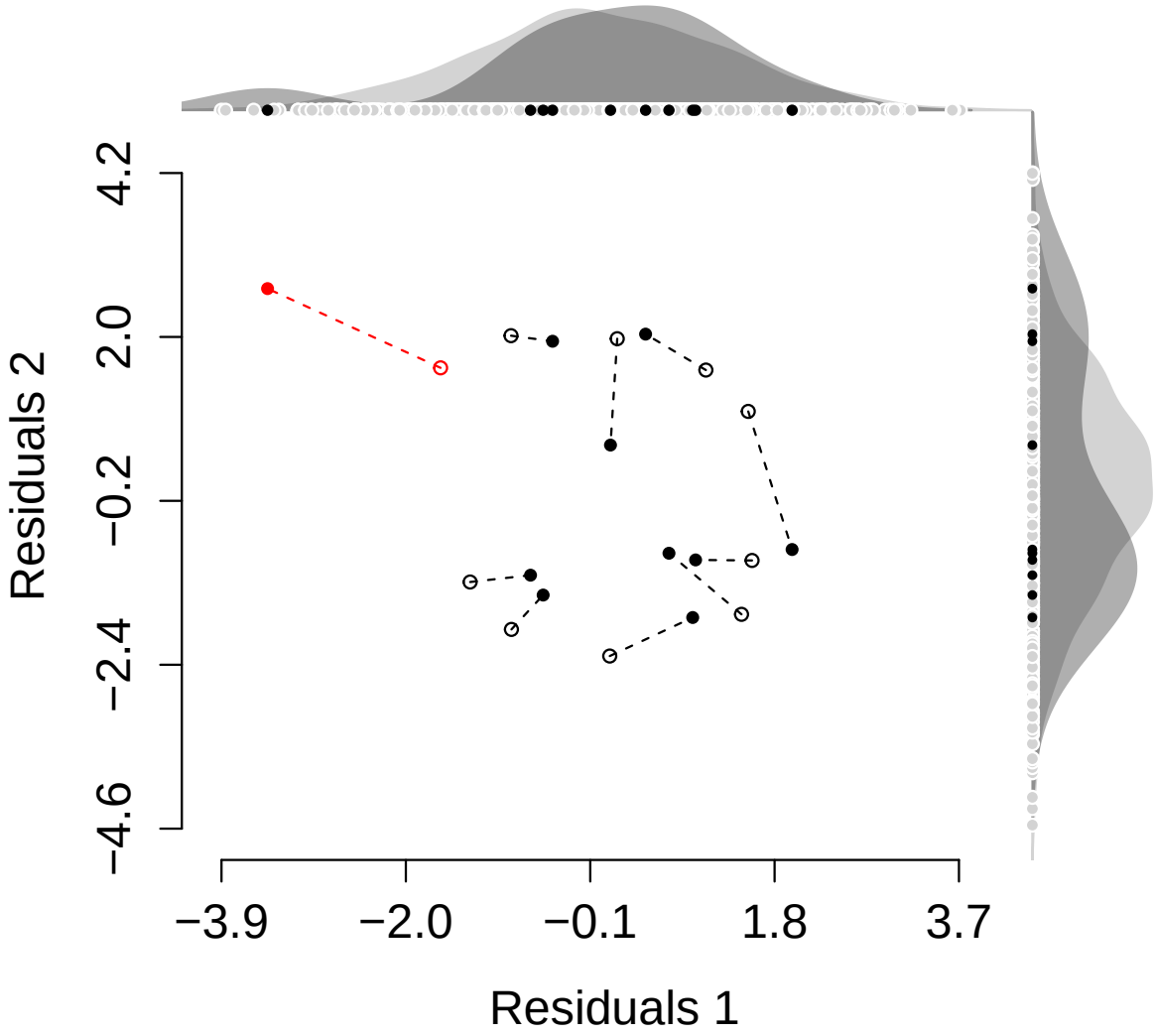
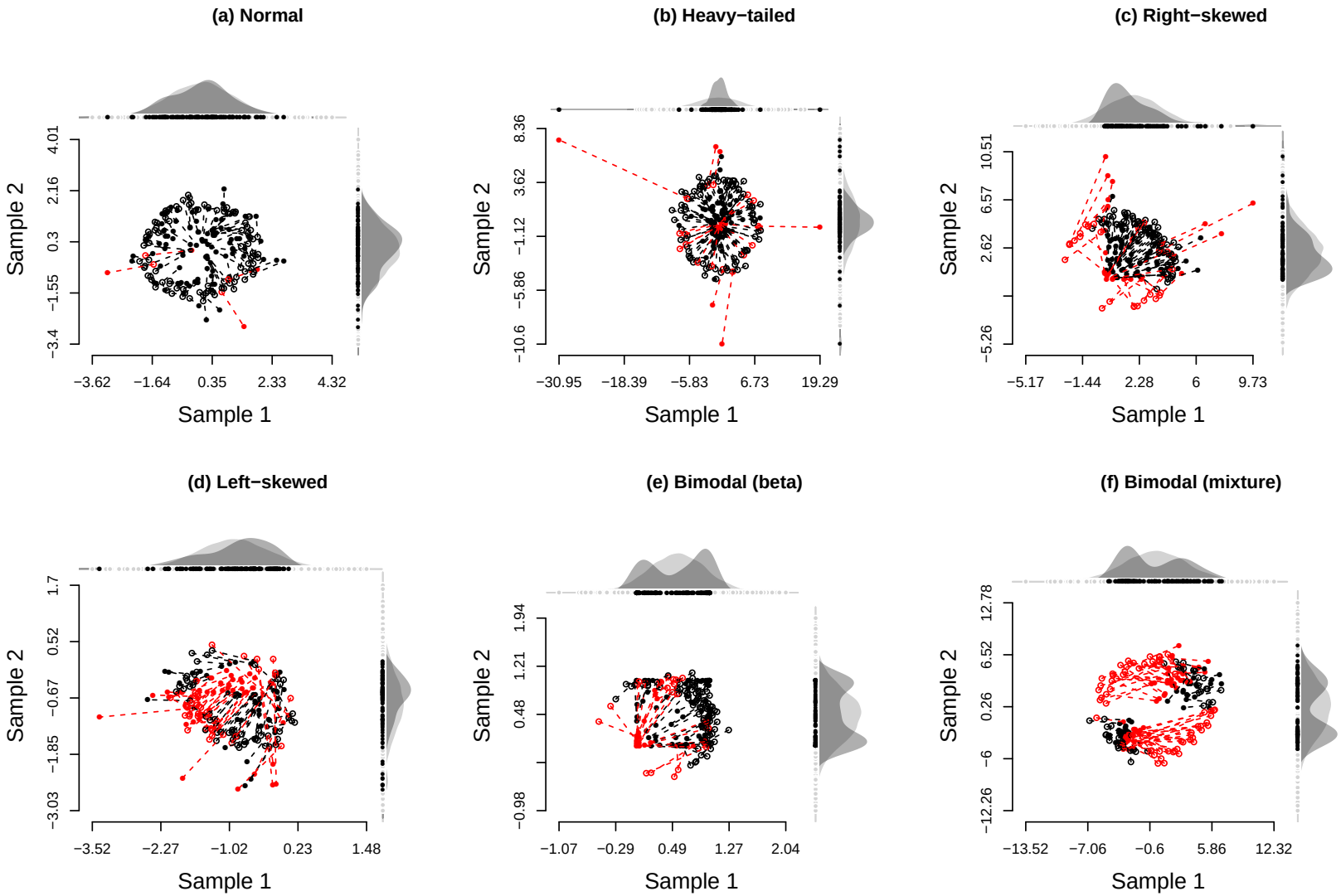


Expected shapes



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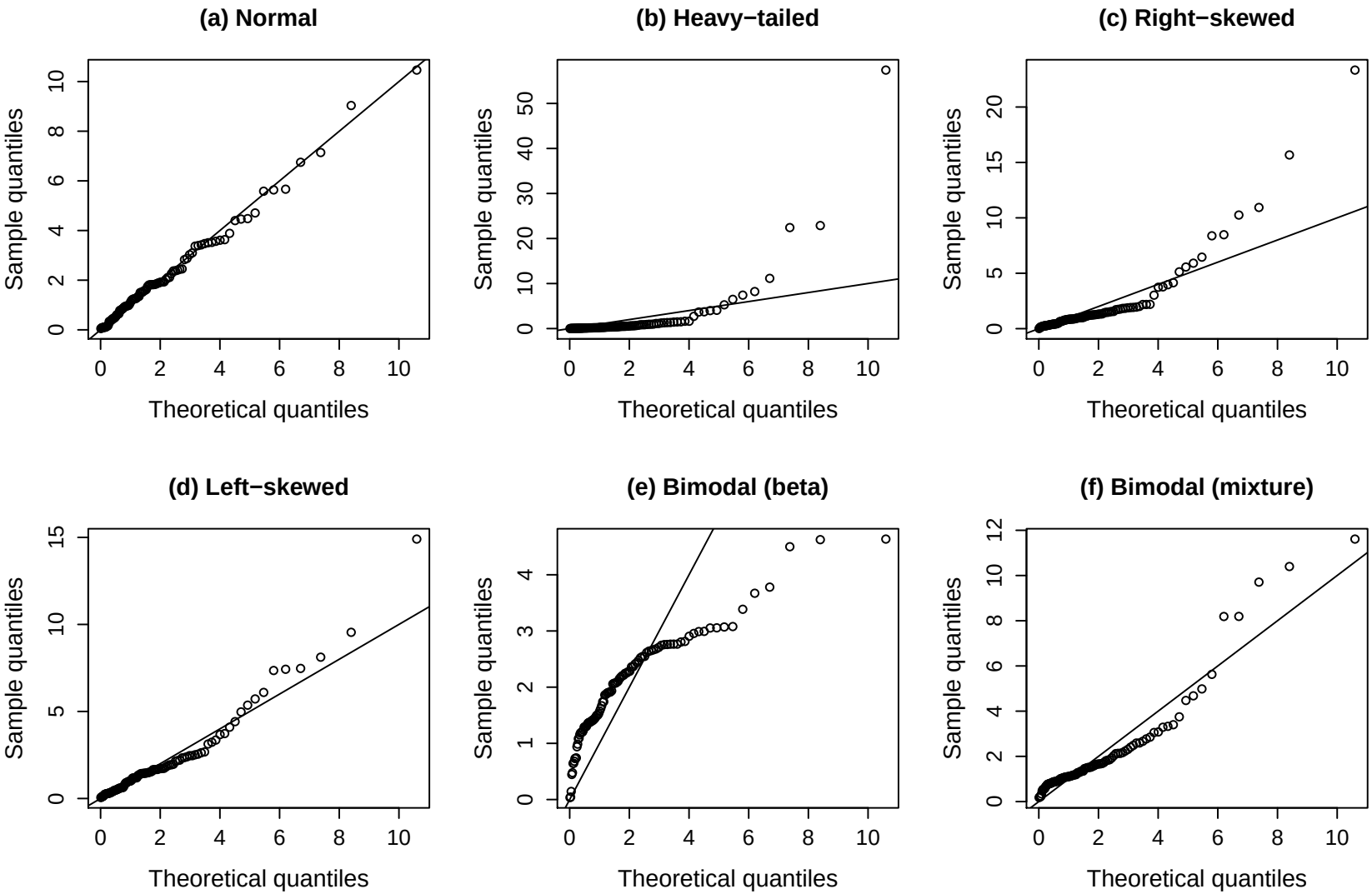


An example using simulated data

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

Marginally

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

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

Model fitting

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

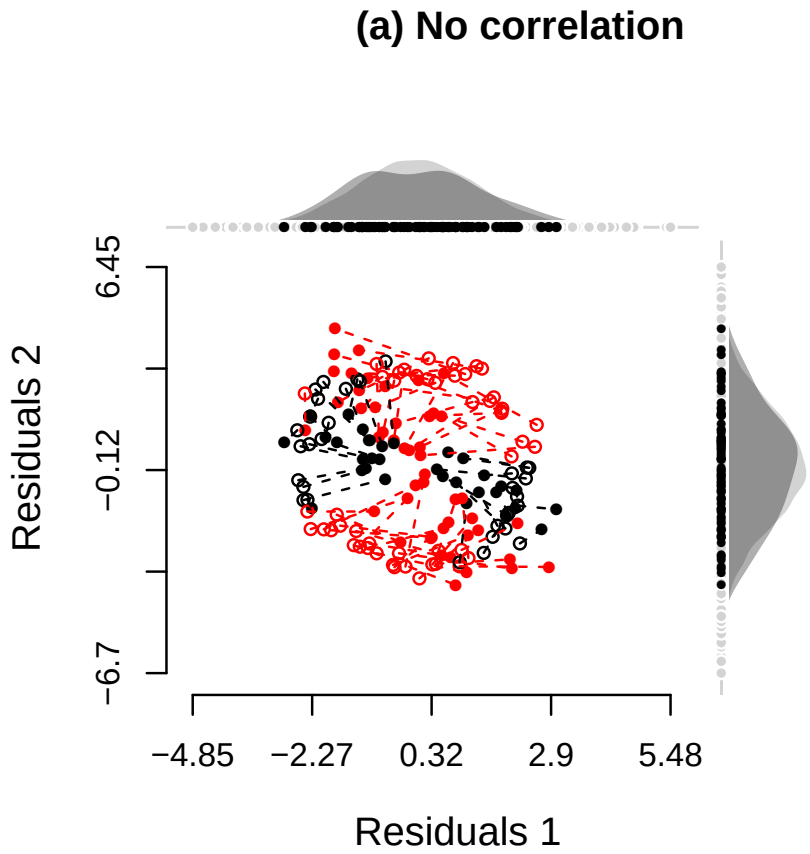
$$L(\boldsymbol{\beta}, \boldsymbol{\Sigma}) = \prod_{i=1}^{n} (2\pi)^{-1} |\boldsymbol{\Sigma}|^{-\frac{1}{2}} \exp\left\{ -\frac{1}{2} (\mathbf{y}_i - \boldsymbol{\mu}_i)^\top \boldsymbol{\Sigma}^{-1} (\mathbf{y}_i - \boldsymbol{\mu}_i) \right\}$$

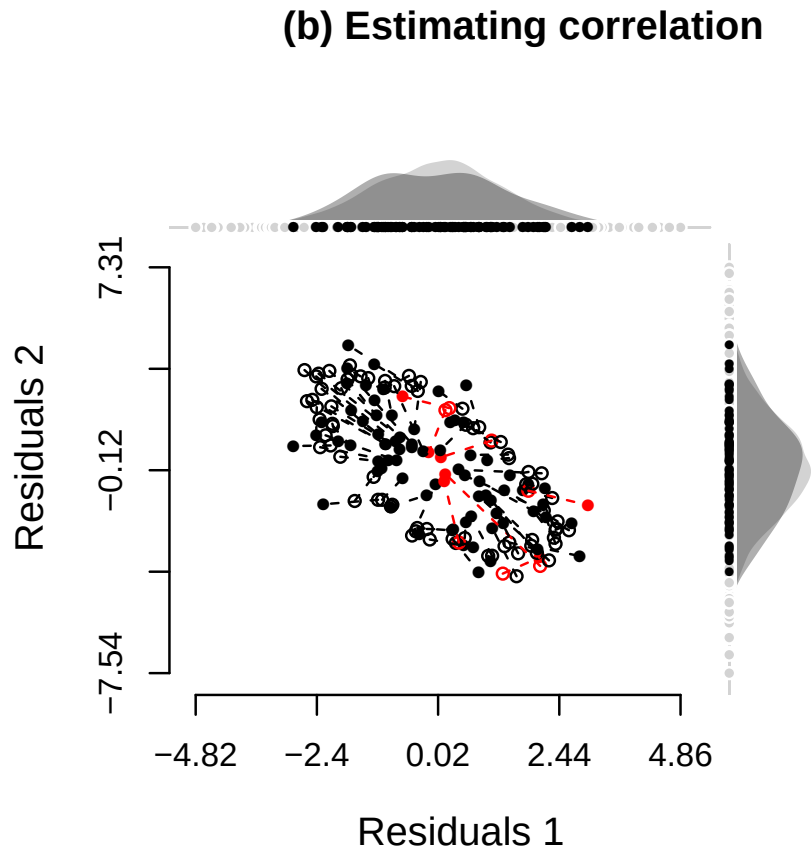
Estimates

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

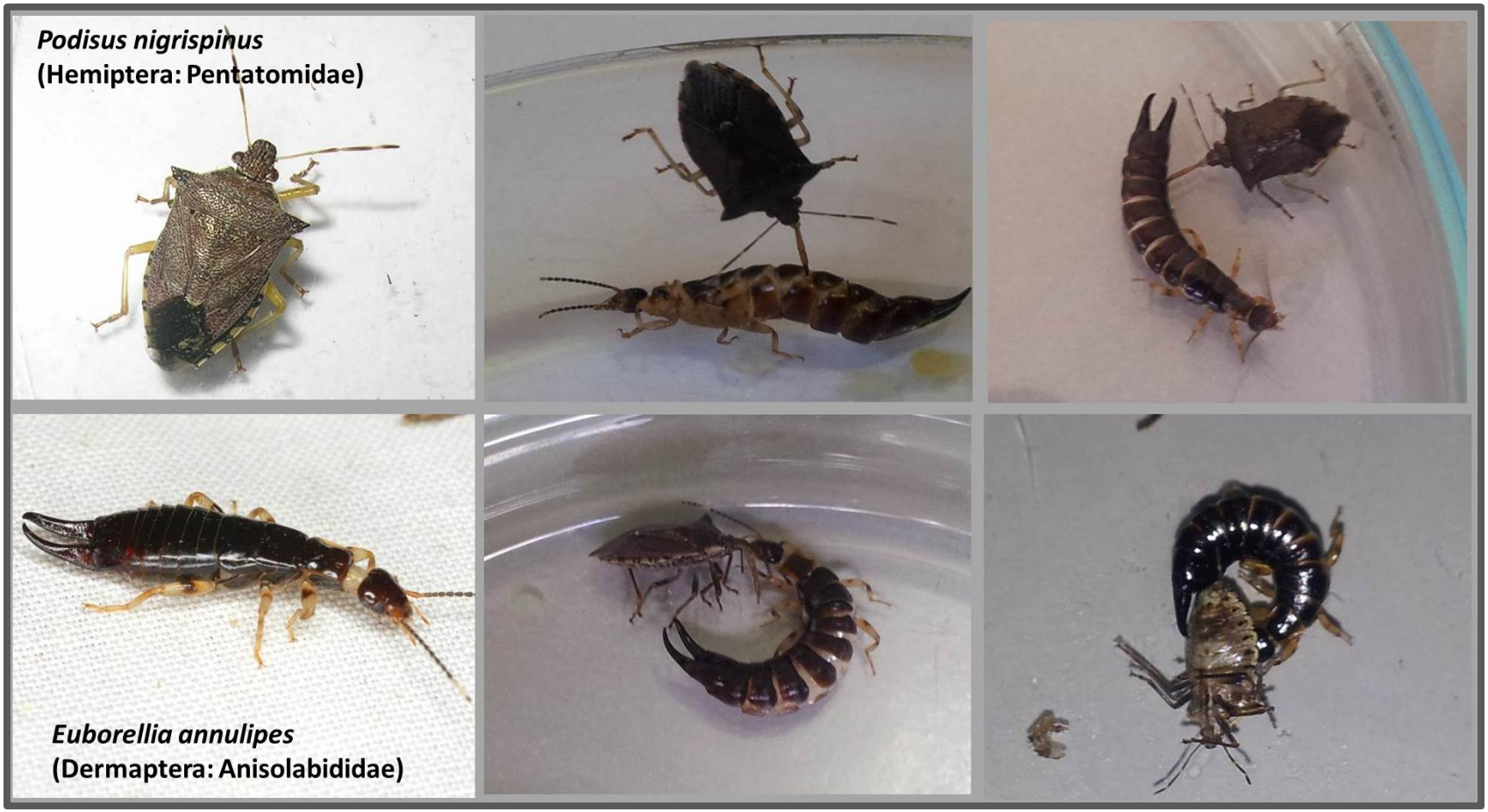
Parameter	Assuming independence	Estimating covariance	True value
β_{10}	1.50 (0.35)	1.50 (0.35)	2.00
β_{11}	0.43 (0.06)	0.43 (0.06)	0.40
β_{20}	0.78 (0.48)	0.78 (0.48)	0.20
β_{21}	0.18 (0.08)	0.18 (0.08)	0.20
σ_1^2	1.79 (0.28)	1.79 (0.28)	2.00
σ_2^2	3.48 (0.55)	3.49 (0.55)	3.00
σ_{12}	0.00 (−)	−1.61 (0.33)	−1.70
$-2\times$loglik	600.42	557.10	−

Bivariate residual plots with simulation polygons





An example with real data



Bivariate Poisson model

$$\begin{aligned} X_j &\sim \mathsf{P}(\lambda_j), \quad j = 0, 1, 2 \\ Y_1 &= X_0 + X_1 \\ Y_2 &= X_0 + X_2 \end{aligned}$$

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

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

$$\begin{aligned} Y_1 &\sim \mathsf{P}(\lambda_0 + \lambda_1) \\ Y_2 &\sim \mathsf{P}(\lambda_0 + \lambda_2) \end{aligned}$$

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

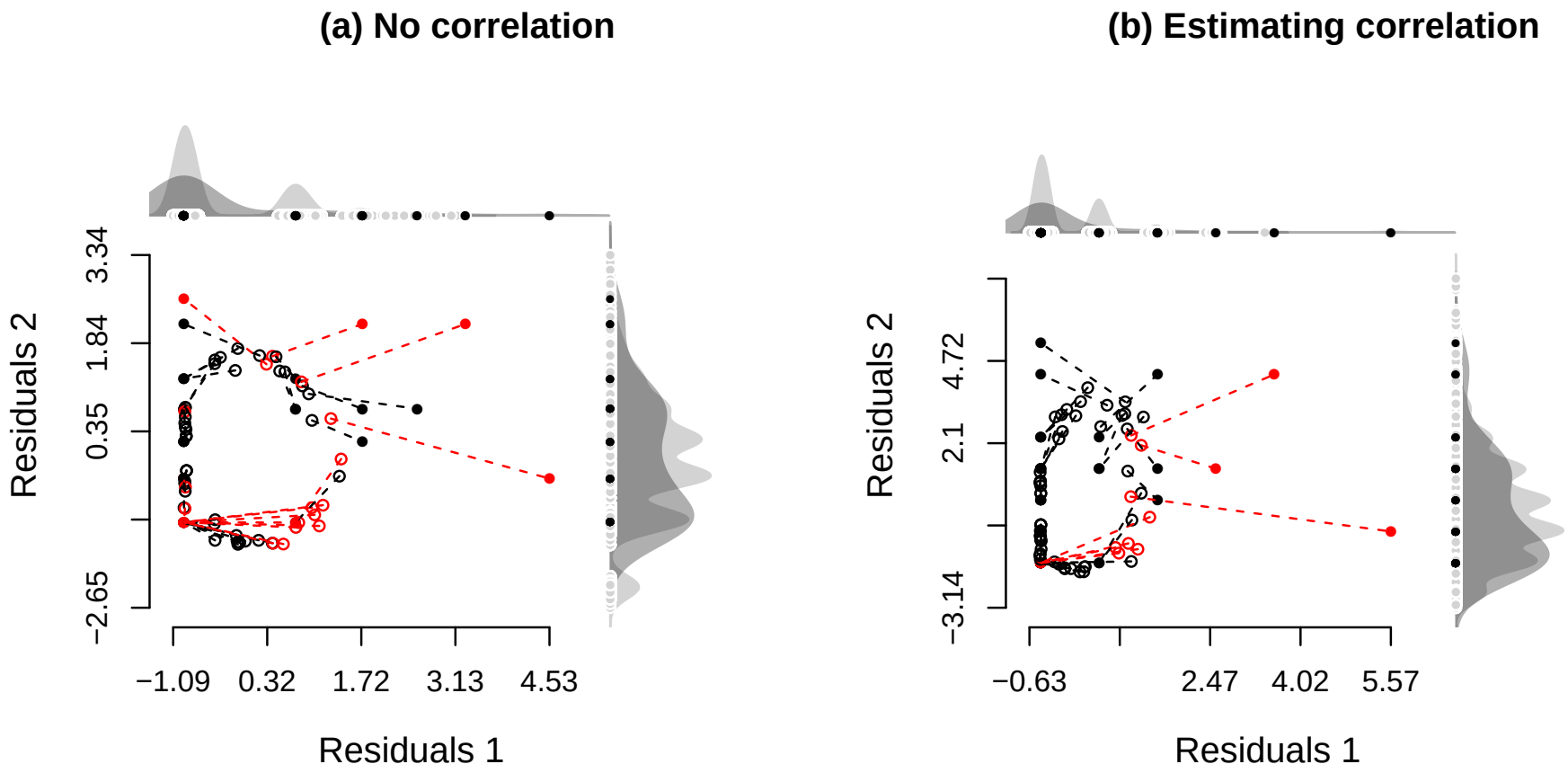
$$\begin{aligned} z_i &= \mathsf{E}(X_{0i}|Y_{1i}, Y_{2i}) \\ &= \sum_{x_{0i}=0}^{\infty} x_{0i} P(X_{0i} = x_{0i}|Y_{1i} = y_{1i}, Y_{2i} = y_{2i}) \\ &= \lambda_0 \frac{P(Y_{1i} = y_{1i} - 1, Y_{2i} = y_{2i} - 1)}{P(Y_{1i} = y_{1i}, Y_{2i} = y_{2i})} \end{aligned}$$

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)
$-2\times$loglik	332.32	327.83

Bivariate residual plots with simulation polygons



Using PIT diagnostics

- Randomized PIT diagnostics for discrete models

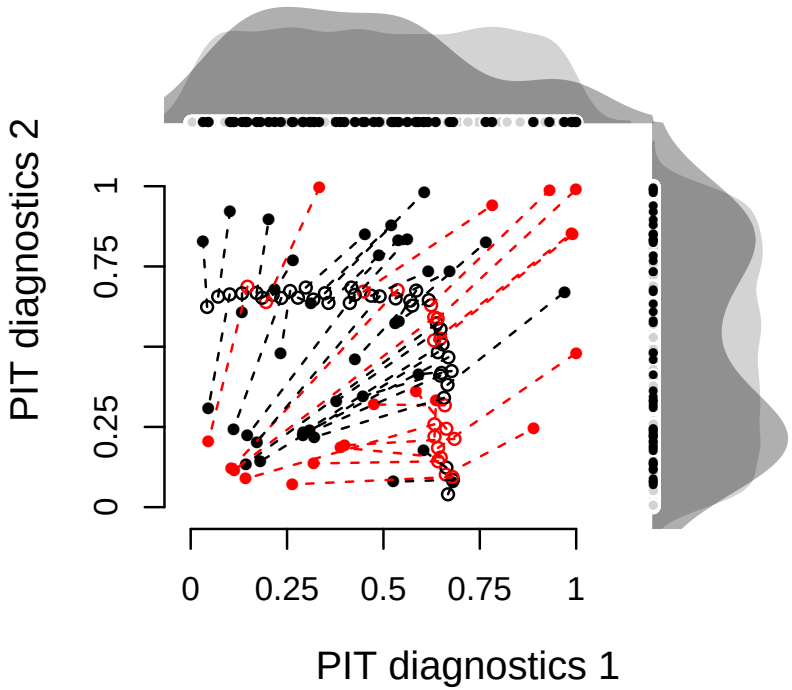
$$r_{ji}^{rand.pit} = F(y_{ji} - 1; \hat{\boldsymbol{\theta}}_{ji}) + u_i\{F(y_{ji}; \hat{\boldsymbol{\theta}}_{ji}) - F(y_{ji} - 1; \hat{\boldsymbol{\theta}}_{ji})\},$$

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

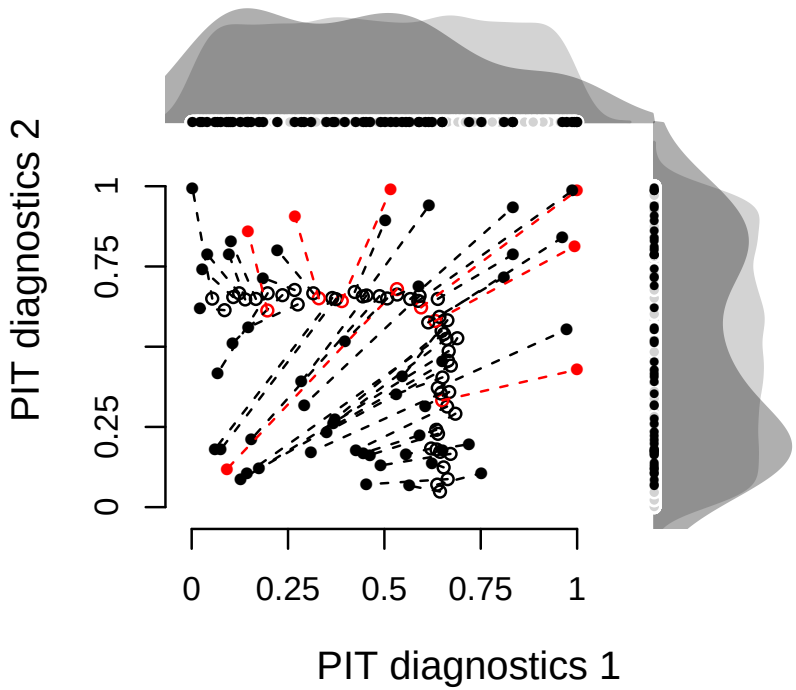
$$F(y_{ji}; \hat{\boldsymbol{\theta}}_{ji}) = e^{-\hat{\mu}_j} \sum_{k=0}^{y_{ji}} \frac{\hat{\mu}_j^k}{k!}$$

Bivariate residual plots with simulation polygons

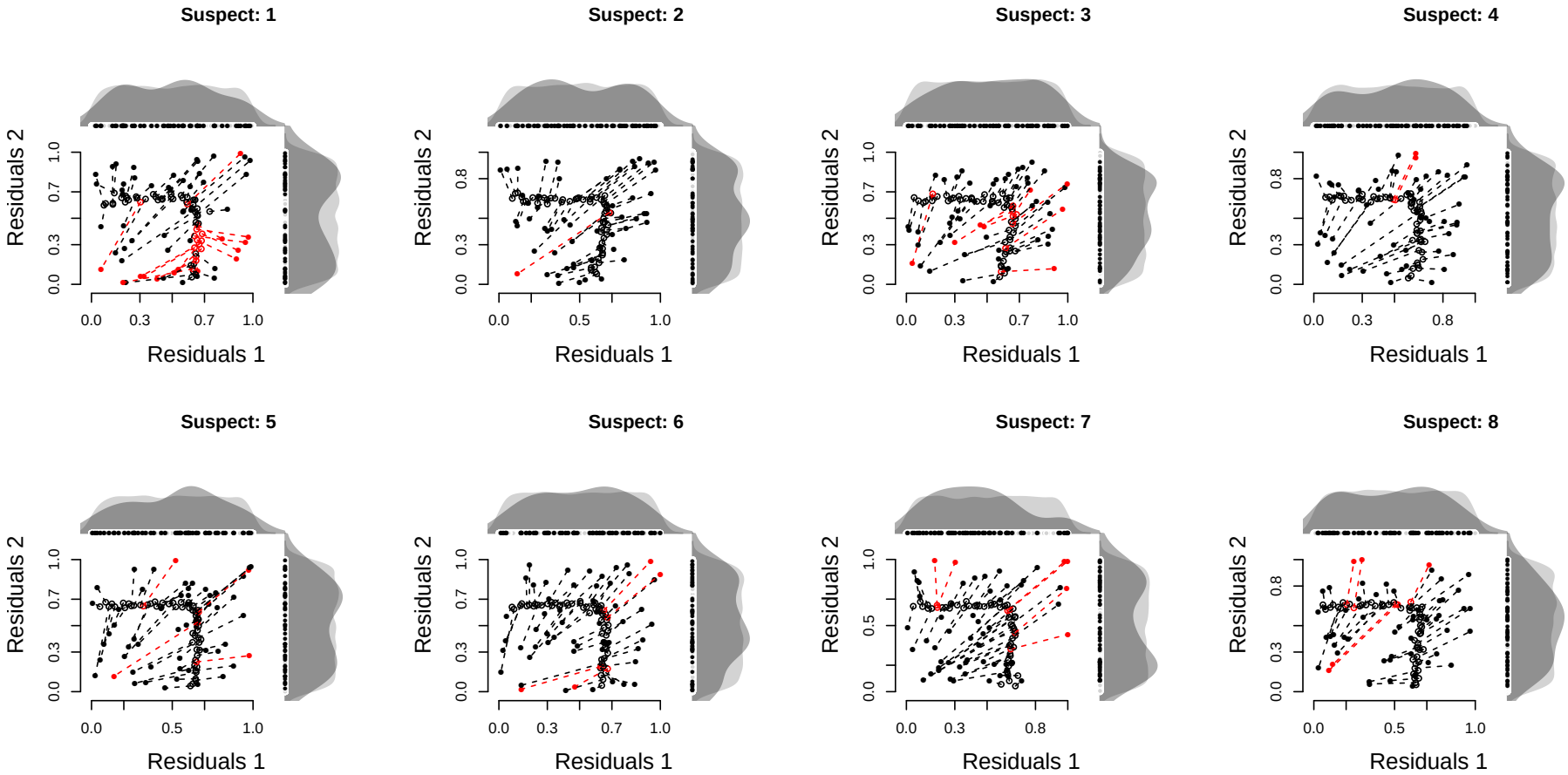
(c) No correlation



(d) Estimating correlation



Line-up test


Suspect: 1
Suspect: 2
Suspect: 3
Suspect: 4
Suspect: 5
Suspect: 6
Suspect: 7
Suspect: 8
Residuals 1
Residuals 2

Code availability

- `bivrp` package on CRAN

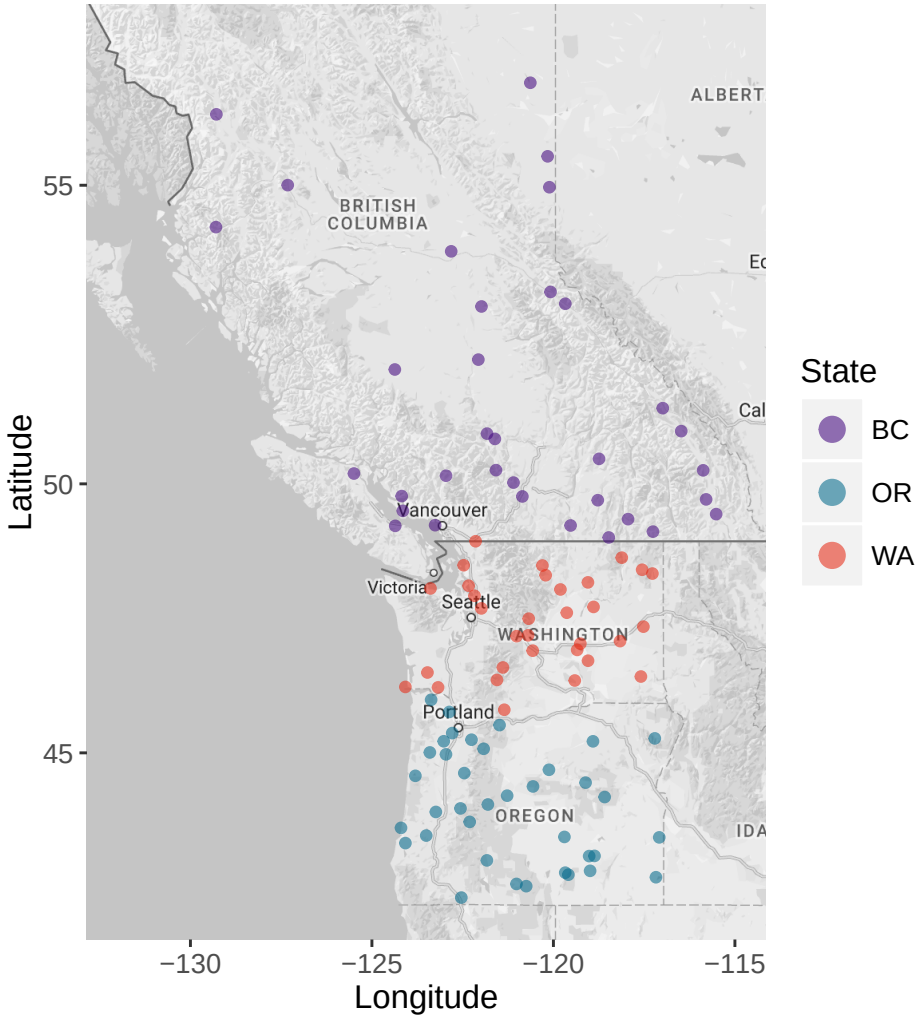
bivrp-package	Bivariate Residual Plots with Simulation Polygons
add.dplots.plot	Internal functions to prepare 'bivrp' objects
add.dplots.prep	Internal functions to prepare 'bivrp' objects
bivrp	Bivariate Residual Plots with Simulation Polygons
chp.perpoint	Internal functions to prepare 'bivrp' objects
get.k	Polygon operations
get.newpolygon	Polygon operations
is.point.inside	Determine if point is inside or outside a simple polygon area
plot.bivrp	Plot Method for bivrp Objects
polygon.area	Polygon operations
sorttheta	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?

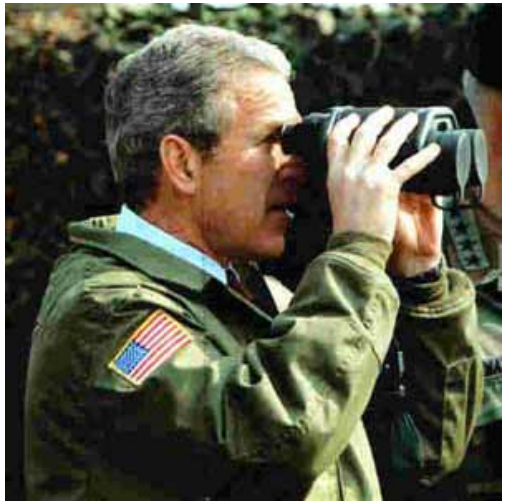
Back to our first motivation example



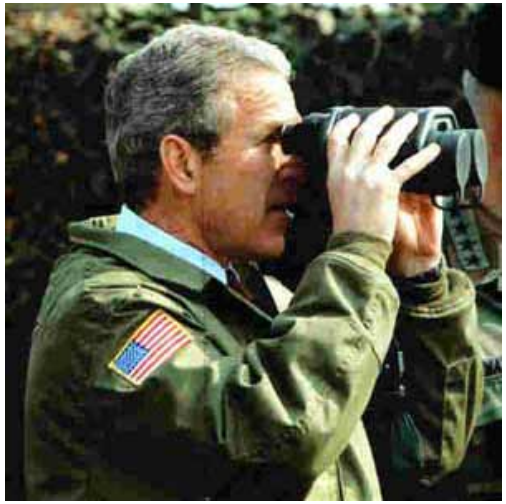
Imperfect detection

Imperfect detection

Imperfect detection



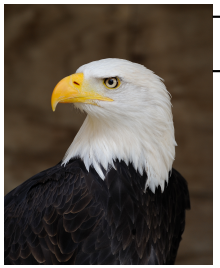
Imperfect detection



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

$$
\begin{aligned}
Y_{1_{it}}|N_{1_i} &\sim \text{Binomial}(N_{1_i}, p_{1_{it}}) \\
N_{1_i} &\sim \text{Poisson}(\lambda_{1_i}) \text{ or NB}(\lambda_{1_i}, \phi_1) \\
\\
Y_{2_{it}}|N_{1_i}, N_{2_i} &\sim \text{Binomial}(N_{2_i}, p_{2_{it}}) \\
N_{2_i}|N_{1_i} &\sim \text{Poisson}(\psi_i + \lambda_{2_i} N_{1_i}) \text{ or NB}(\psi_i + \lambda_{2_i} N_{1_i}, \phi_2)
\end{aligned}
$$

Assumptions

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

Joint N-mixture model

- Likelihood:

$$L(\mathbf{p_1}, \mathbf{p_2}, \boldsymbol{\theta_1}, \boldsymbol{\theta_2}, \boldsymbol{\psi} | \{y_{1_{it}}\}, \{y_{2_{it}}\}) =$$

$$\prod_{i=1}^{R} \left\{ \sum_{n_{1_i}=\max_t\{y_{1_{it}}\}}^{\infty} \left[\prod_{t=1}^{T} \text{Bin}(y_{1_{it}}; n_{1_i}, p_{1_{it}}) \right] f_{N_{1_i}}(n_{1_i}; \boldsymbol{\theta_1}_i) \times \right.$$

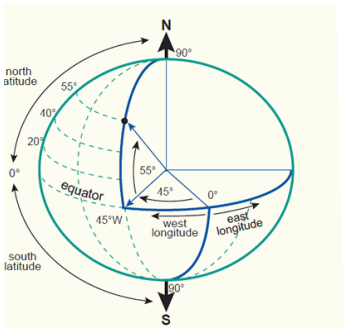
$$\left. \sum_{n_{1_i}=\max_t\{y_{1_{it}}\}}^{\infty} f_{N_{1_i}}(n_{1_i}; \boldsymbol{\theta_1}_i) \sum_{n_{2_i}=\max_t\{y_{2_{it}}\}}^{\infty} \left[\prod_{t=1}^{T} \text{Bin}(y_{2_{it}}; n_{2_i}, p_{2_{it}}) \right] f_{N_{2_i}}(n_{2_i}; \psi_i, \boldsymbol{\theta_2}_i) \right\}$$

Total variation in *observed* eagle abundance

Total variation in *observed* eagle abundance

Habitat suitability

Geographical coordinates



Total variation in *observed* eagle abundance

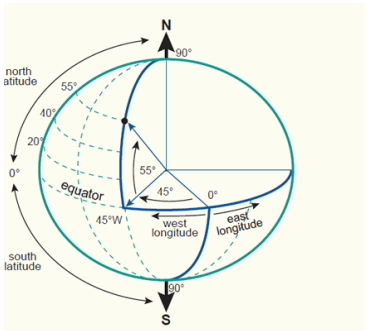
Habitat suitability

Environmental covariates

Affecting eagle

Geographical coordinates

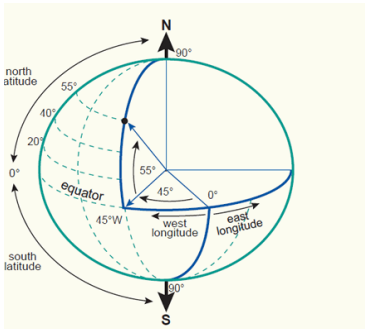
Wind speed
Temperature
...



Total variation in *observed* eagle abundance

Habitat suitability

Geographical coordinates



Environmental covariates

Affecting eagle

Wind speed
Temperature
...

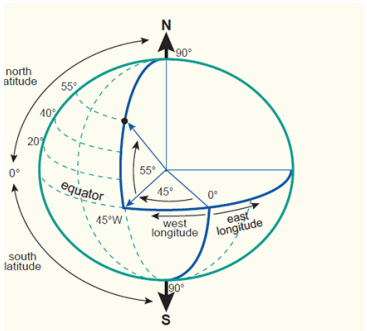
Affecting observer

Cloud coverage
Expertise
...

Total variation in *observed* eagle abundance

Habitat suitability

Geographical coordinates



Environmental covariates

Affecting eagle

Wind speed
Temperature
...

Affecting observer

Cloud coverage
Expertise
...

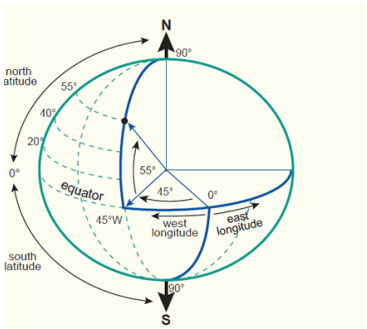
Unexplained variation

Missing covariates
Natural variation
...

Total variation in *observed* eagle abundance

Habitat suitability

Geographical coordinates



Environmental covariates

Affecting eagle

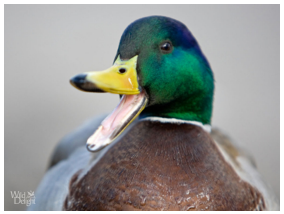
Wind speed
Temperature
...

Affecting observer

Cloud coverage
Expertise
...

Direct + indirect effects of mallard co-occurrence

Habitat disturbance
Resource availability
Direct interactions
...



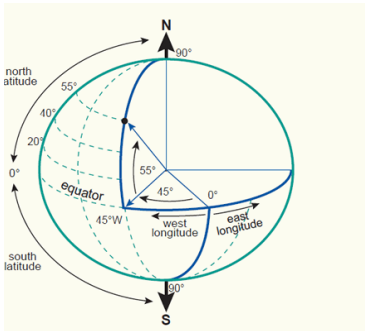
Unexplained variation

Missing covariates
Natural variation
...

Total variation in *observed* eagle abundance

Habitat suitability

Geographical coordinates



Abundance

ψ_i

Environmental covariates

Affecting eagle

Wind speed
Temperature
...

Affecting observer

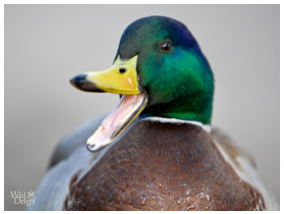
Cloud coverage
Expertise
...

Detection probability

$p_{2_{it}}$

Direct + indirect effects of mallard co-occurrence

Habitat disturbance
Resource availability
Direct interactions
...



Abundance

λ_{2_i}

Unexplained variation

Missing covariates
Natural variation
...

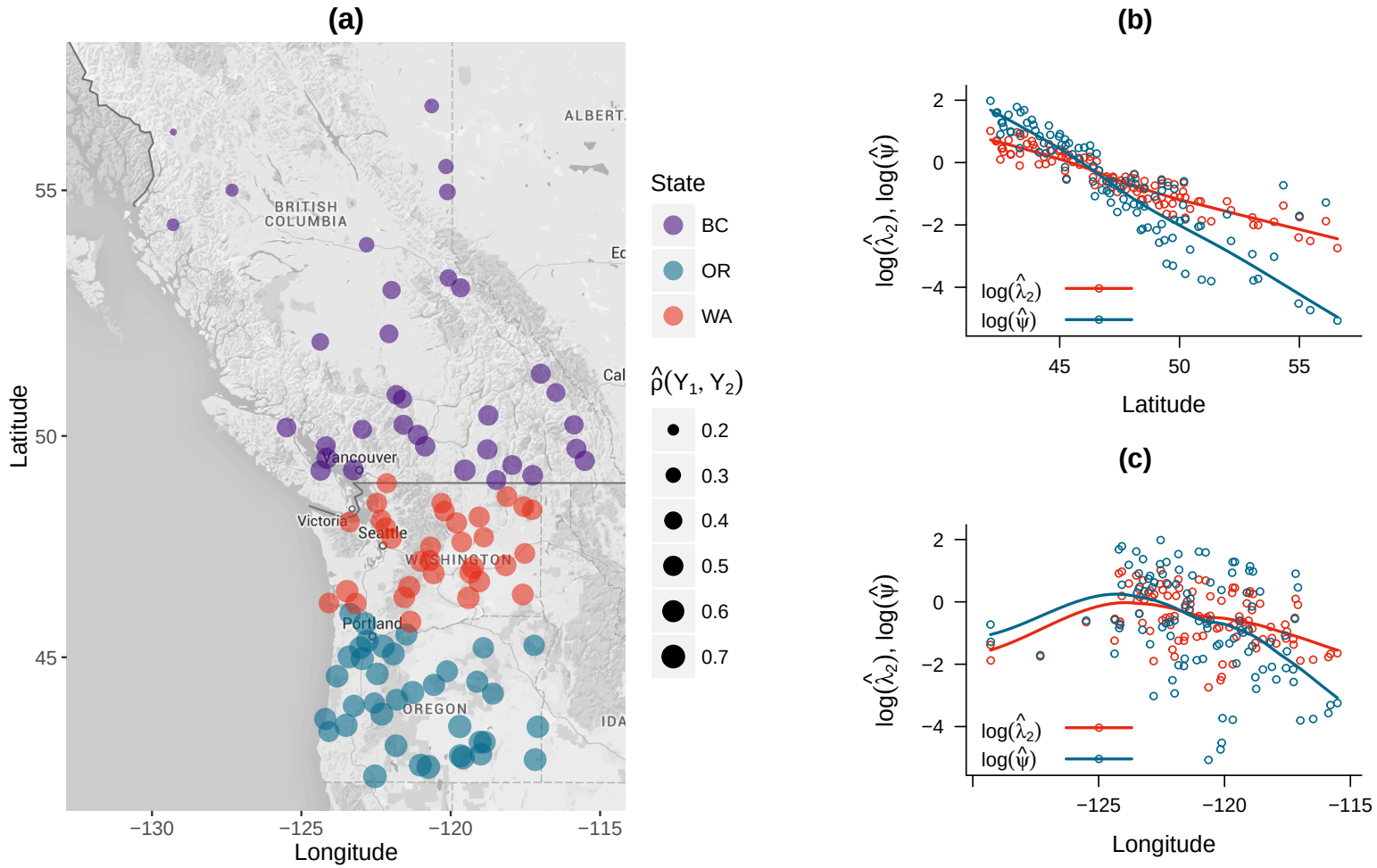
Overdispersion

ϕ_2

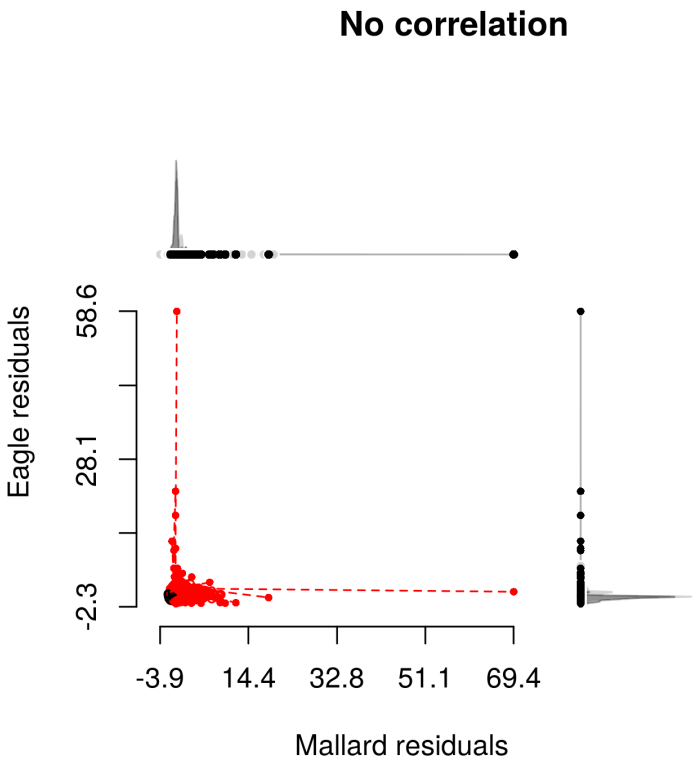
Case study: Model selection

Parameter	Univariate models				Joint models			
	P–P	NB–P	P–NB	NB–NB	P–P	NB–P	P–NB	NB–NB
Detection								
(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
(Bald eagle – p_{2it})								
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
Abundance								
(Mallard – λ_{1i})								
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×Long	0.04	0.09	0.04	0.09	0.14	0.08	0.04	0.09
(Bald eagle – ψ_i)								
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×Long	−0.12	−0.12	0.13	0.13	0.91	−0.16	0.05	−0.01
(Bald eagle – λ_{2i})								
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×Long	—	—	—	—	−0.31	0.04	3.49	0.07
(Dispersion)								
ϕ_1	—	0.51	—	0.51	—	0.39	—	0.46
ϕ_2	—	—	0.46	0.46	—	—	0.48	1.71
$-2 \times$ loglik	3350.28	2986.90	2994.67	2631.29	3196.88	2620.96	2990.69	2617.46

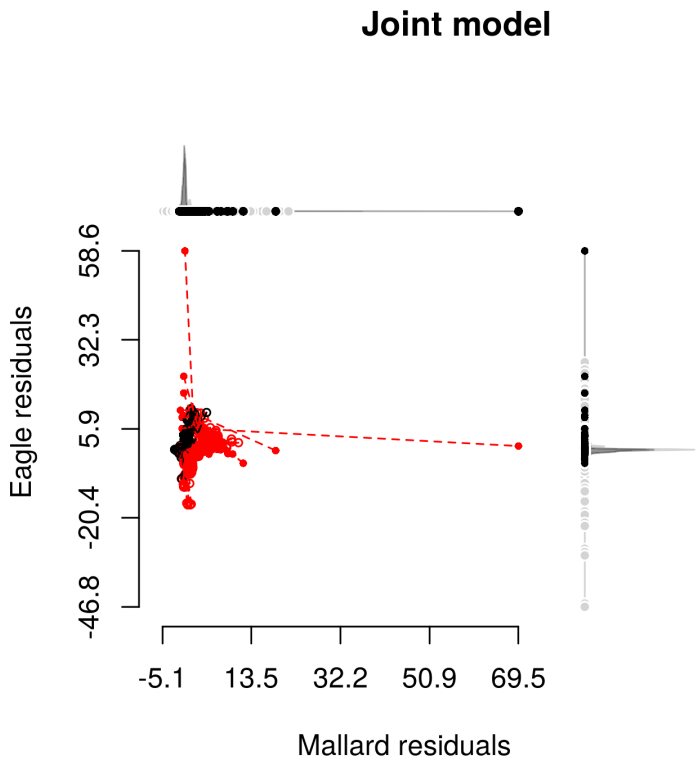
Results


(a)
ALBERT
BRITISH COLUMBIA
Vancouver
Victoria
Seattle
WASHINGTON
Portland
OREGON
IDA
55
50
45
Latitude
−130
−125
−120
−115
Longitude
State
BC
OR
WA
$\hat{\rho}(Y_1, Y_2)$
0.2
0.3
0.4
0.5
0.6
0.7
(b)
$\log(\hat{\lambda}_2), \log(\hat{\psi})$
2
0
−2
−4
$\log(\hat{\lambda}_2)$
$\log(\hat{\psi})$
45
50
55
Latitude
(c)
$\log(\hat{\lambda}_2), \log(\hat{\psi})$
2
0
−2
−4
$\log(\hat{\lambda}_2)$
$\log(\hat{\psi})$
−125
−120
−115
Longitude

Case study: bivariate residual plots



368/515 (71%)



197/515 (38%)

Thank you!

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