

Estimating occupancy and fitting models

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Estimating
occupancy and
fitting models

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Occupancy and
detectability

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Edge solutions

Exact mean and
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Motivation

Why monitor populations?

- ▶ invasive species - foxes: threat to native wildlife e.g. lyrebirds



- ▶ endangered species - e.g. growling grass frog



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Objective and challenge

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Probability of presence

Probability of detection

ψ

p

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Estimate Occupancy in the presence of Imperfect detection

Objective

Challenge

- site = patch of land, fixed area of stream bank etc.
- site occasion = visit to a site.

Presence-absence data

A typical detection matrix

Site	Survey.1	Survey.2	Survey.3	Survey.4	
1	0	0	0	0	
2	0	0	0	0	
3	0	0	0	0	
4	0	0	0	0	
5	0	0	0	0	
6	0	0	0	0	
7	0	0	0	0	
8	1	← x_{ij}	1	1	
9	0	0	1	1	
10	1	1	1	1	
11	0	1	0	0	
12	0	1	0	0	$x_{i.} \neq 0$
13	1	1	1	1	
14	1	1	1	1	
15	1	1	1	1	
16	0	0	0	0	$x_{i.} = 0$
17	0	0	0	0	
18	1	1	1	1	
19	0	0	0	0	
20	1	1	1	1	
21	1	1	1	1	
22	0	0	0	0	
23	0	1	1	1	
24	0	1	1	1	
25	0	0	0	1	
26	1	1	1	1	
27	0	0	0	0	

Table: Capture histories for the growling grass frog. The 27 independent sites each were surveyed on 4 occasions at night within the 2002-2003 season (Heard et al., 2006).

The basic occupancy model (BOD) – ZIB

- The detection (p) and occupancy (ψ) probabilities remain constant over all N sites and T survey occasions.

The number of detections $X_{i\cdot}$, at site i is distributed as

$$X_{i\cdot} \stackrel{d}{=} \begin{cases} 0, & \text{with probability } (1 - \psi), \\ \text{Bi}(T, p), & \text{with probability } \psi. \end{cases}$$

$$\Pr(X_{i\cdot} = x_{i\cdot}) = \begin{cases} \psi(1 - p)^T + (1 - \psi), & x_{i\cdot} = 0; \\ \psi \binom{T}{x_{i\cdot}} p^{x_{i\cdot}} (1 - p)^{T - x_{i\cdot}}, & x_{i\cdot} = 1, 2, \dots, T. \end{cases}$$

Two states and three possible outcomes:

detection (x_{ij}) = 0 \rightarrow not present, OR present but not detected

detection (x_{ij}) = 1 \rightarrow present

Full likelihood (ZIB) under constant ψ & p

(MacKenzie et al., 2002)

$$\begin{aligned} L(\psi, p \mid \mathbf{X}) &= \prod_{i=1}^N L_i(\psi, p \mid x_i.) \\ &= \left(\underbrace{\psi(1-p)^T + (1-\psi)}_{\text{nondetections}} \right)^{N-k} \underbrace{\psi^k p^x (1-p)^{NT-x}}_{\text{detections}} \end{aligned}$$

- ▶ number of detected sites

$$k = \sum_{i=1}^N I(x_{i.} > 0)$$

- ▶ total number of detections

$$x = \sum_{i=1}^N x_{i.}$$

k and x sufficient for ψ, p

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Score equations – Full likelihood under constant ψ & p

$$\psi_s = \frac{k}{N\theta_s} \quad \text{and} \quad p_s = \frac{x\theta_s}{kT}$$

$$\theta_s = 1 - (1 - p_s)^T \quad (\text{prob. of at least one detected site.})$$

$$\text{Var}_{\text{Mac}}(\hat{\psi}) = \frac{\psi}{N} \left((1 - \psi) + \frac{1 - \theta}{\theta - Tp(1 - p)^{T-1}} \right).$$

BUT do not always give MLEs !!!

As $N, T \rightarrow \infty$ these are the MLE.

If N, T small then these will not apply.

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Limitations – Full likelihood (BOD)

1. Convergence – direct maximisation BOD does not always converge, no closed form solutions for ψ_s, p_s
2. Boundary issues – $\hat{\psi} > 1, \hat{p} > 1$
3. Standard errors – no closed form solutions for Var_{Mac} , hessian not always available

(Karavarsamis et al. (2013); Karavarsamis and Huggins (2019b), Karavarsamis and Huggins (2019a), Karavarsamis and Watson (b), Karavarsamis and Watson (a))

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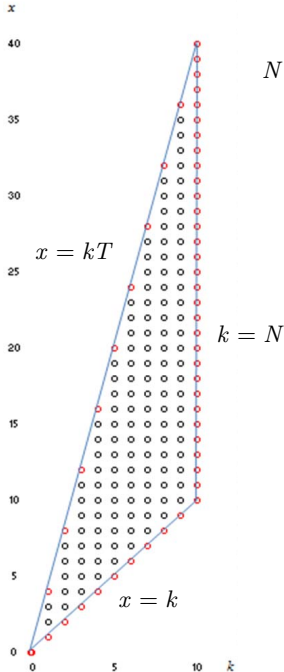
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2. Boundary problem

$$N = 10, T = 4$$



$$\psi_s = k/N\theta_s$$
$$p_s = x\theta_s/kT$$

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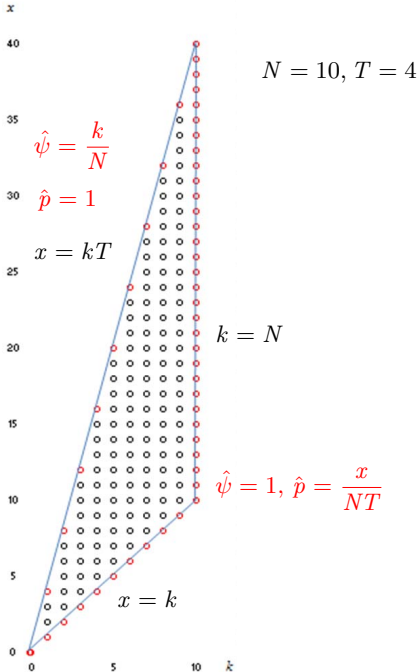
2. Boundary problem

‘Edge’ solutions

MLE is 1 but
soln to score eqn > 1

$$\psi_s = k/N\theta_s$$

$$p_s = x\theta_s/kT$$



This caused

1. non-convergence of the likelihood (too flat, multiple local maxima)
2. estimates that were greater than 1 i.e. $\hat{\psi} > 1$ or $\hat{p} > 1$
3. problems with interval estimators i.e. standard errors

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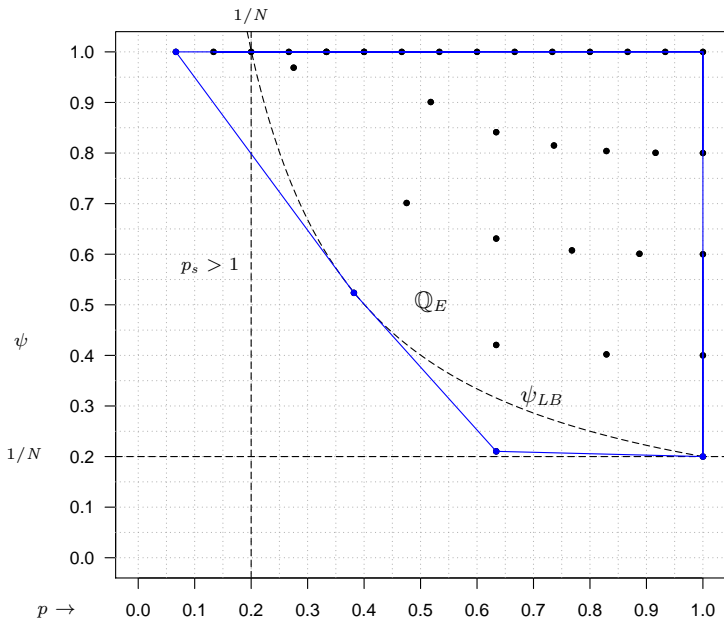
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Plausible region – MLEs always exist ($N = 5, T = 3$)



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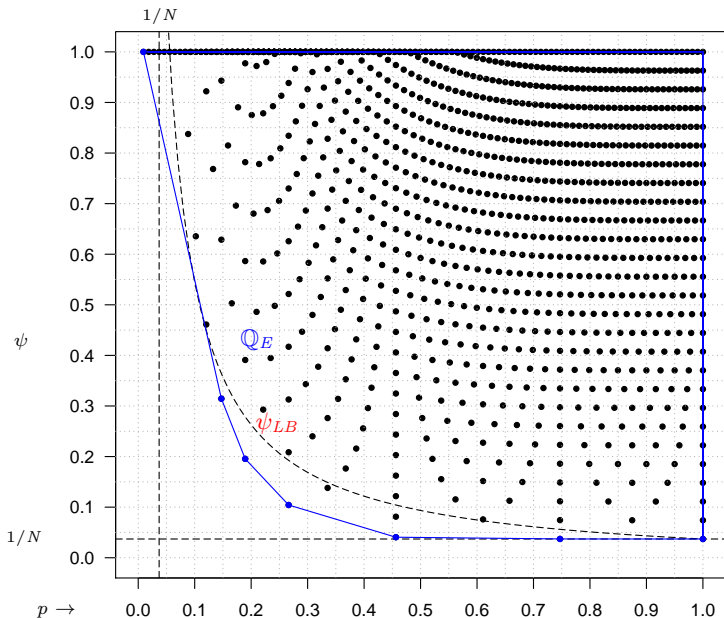
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Plausible region – MLEs always exist ($N = 27, T = 4$)



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Problem 3. Standard errors – exact mean and variance

We derived an expression for the joint pmf of (X, K) .

This allowed us to evaluate the

- ▶ bias of $\hat{\psi} = \hat{\psi}(x, k)$
- ▶ exact variance of $\hat{\psi}$

Results

- ▶ bias-corrections for $\hat{\psi}$ not effective when N, T , or p small because not enough information in (x, k)
- ▶ asymptotic variance underestimates actual variance

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Bias - $N = 5, T = 3$

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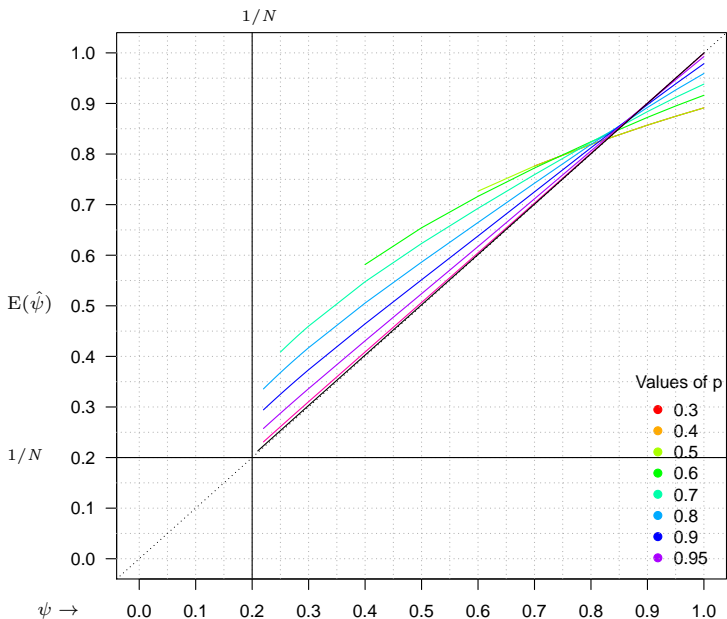
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Bias - $N = 27, T = 4$

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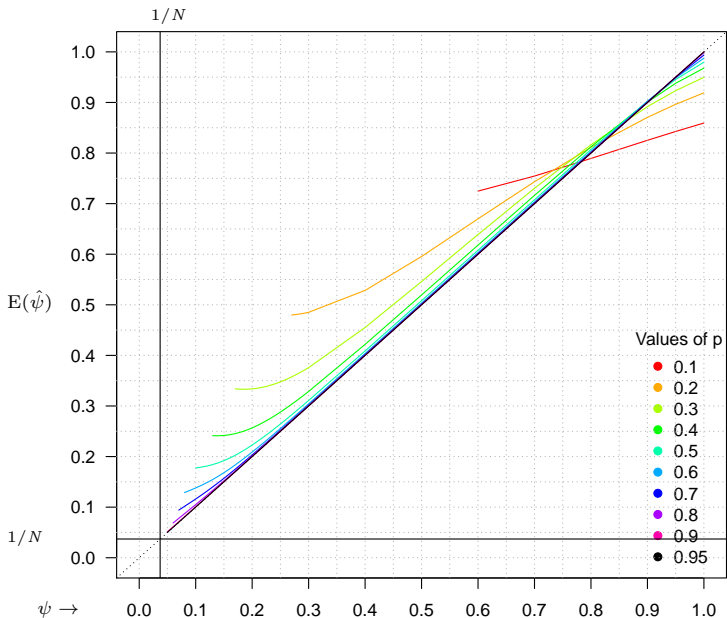
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Asymptotic and exact variance for $\hat{\psi}$: $N = 5, T = 3$

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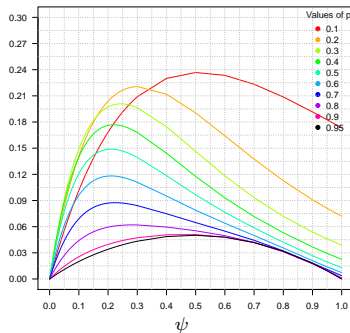
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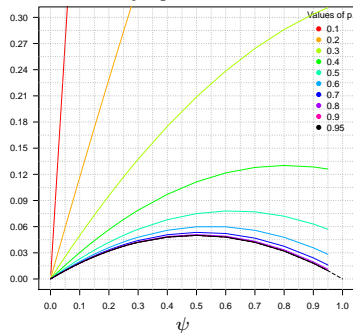
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Exact variance



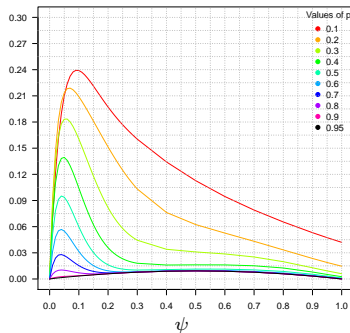
Asymptotic variance



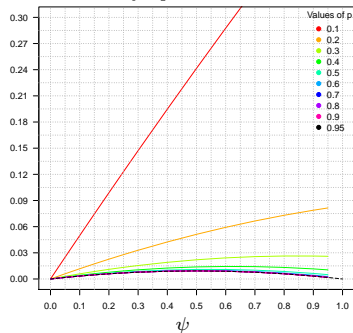
Asymptotic and exact variance for $\hat{\psi}$: $N = 27, T = 4$

Frogs $\hat{\psi} = 0.557, \hat{p} = 0.782, ase = 0.096, se = 0.095$

Exact variance



Asymptotic variance



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Full likelihood limitations and solutions — summary

1. Non-convergence of the likelihood (identifiability)

1.1 next...

2. Boundary issues ($\psi_s, p_s > 1$)

2.1 Edge solutions and Plausible region (Karavarsamis & Watson, 2019 – in prep.)

3. Standard errors

3.1 Exact variance showed asymptotic not good (Karavarsamis *et al.*, 2013)

4. Too hard to include covariates

4.1 next...

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Characteristics

- ▶ Repeated visits to a site (introduces heterogeneity — may be solved with covariates)
- ▶ Observe presence–absence of a species
- ▶ Covariate information
 - ▶ site characteristics — geographic...
 - ▶ species characteristics
 - ψ : habitat type, patch size, age, gender...
 - p : weather, site accessibility, detection methods...

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Full and partial likelihoods

Full likelihood

- ▶ estimates highly variable
- ▶ too hard to include covariates e.g. Welsh et al. (2013), and not a GLM

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Existing methods

- ▶ **unmarked**, bootstrap, Bayesian methods

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Partial likelihood (two-stage approach)

- ▶ easy to include non-linear covariates i.e. GAMs !
- ▶ resolves limitations e.g. efficient closed form variance approximations
- ▶ reduces the dimension of models

Partial likelihoods – benefits

What we want:

- ▶ to include non-linear covariates with GAMs and have full use of GLM methodology

How to achieve goals:

- ▶ partials suit this well and allow to consider ψ and p separately
- ▶ repeated observations at each site give more info on detections
- ▶ more info on detections encourages us to consider (i.e. to estimate) detections separately from occupancy which ignores info on 1st detections
- ▶ achieve this with partial likelihoods (they simplify complex likelihoods and deal with nuisance params) need to ignore info on first detections but no great loss of efficiency
- ▶ two-stage estimation for ψ and p gives full use of GLMs etc
- ▶ now can get variance approximations too
- ▶ standard errors are readily obtainable, unlike those obtained from inverting the hessian of the full likelihood that may fail especially near the boundaries of the parameter space in about 5% – 20% of cases

Homogenous case – ψ and p constant

Partial likelihood

$$\begin{aligned}L(\psi, p) &\propto (1 - \psi + \psi(1 - p)^T)^{N-k} \prod_{i=1}^k \psi p^{x_i} \cdot (1 - p)^{T-x_i} \\&= (1 - \psi\theta)^{N-k} \psi^k \prod_{i=1}^k (1 - p)^{a_i-1} p^{x-k} (1 - p)^{b-(x-k)} \\&= L_1(\psi, p) L_2(p)\end{aligned}$$

- omit first detections, a_i
- total number of occasions after a_i is b

Now estimate ψ and p SEPARATELY !

Homogenous case – ψ and p constant

Partial likelihood

$$\begin{aligned}L(\psi, p) &\propto (1 - \psi + \psi(1 - p)^T)^{N-k} \prod_{i=1}^k \psi p^{x_i} \cdot (1 - p)^{T-x_i} \\&= (1 - \psi\theta)^{N-k} \psi^k \times p^{x-k} (1 - p)^{b-(x-k)} \\&= L_1(\psi, p) \times L_2(p)\end{aligned}$$

- omit first detections, a_i
- total number of occasions after a_i is b

Now estimate ψ and p SEPARATELY !

Score equations — homogeneous partial likelihood

Stage 1: $L_2(p)$ gives

$$\hat{p} = \frac{x - k}{b}, \quad \text{Var}(\hat{p}) = \hat{p}(1 - \hat{p})/b$$

Stage 2: $L_1(\psi, \hat{p})$ gives

$$\hat{\psi} = \frac{k}{N\hat{\theta}}$$

and

$$\begin{aligned} \text{Var}(\hat{\psi}) &= \text{Var} \left\{ \text{E}(\hat{\psi} \mid b, k) \right\} + \text{E} \left\{ \text{Var}(\hat{\psi} \mid b, k) \right\} \\ &\approx \frac{\psi(1 - \psi\theta)}{N\theta} + \left(\frac{\psi(1 - \psi\theta)}{N\theta} + \psi^2 \right) \frac{T^2(1 - p)^{2(T-1)}}{\theta^2} \frac{p(1 - p)}{b} \end{aligned}$$

Now we have closed form solutions, yipee!!!

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Full versus partial likelihood - homogeneous case

Simulations – small p , large N

Examine efficiency of partial likelihood

	Partial		Full	
	\hat{p}	$\hat{\psi}$	\hat{p}	$\hat{\psi}$
$N = 1000, T = 5$	0.050	0.400	0.050	0.400
Med. est.	0.049	0.407	0.049	0.407
Median s.e.	0.016	0.124	0.015	0.120
Mad	0.016	0.127	0.015	0.123
Efficiency	1.021	0.988		
$N = 100, T = 5$	0.200	0.600	0.200	0.600
Med. est.	0.198	0.609	0.197	0.609
Median s.e.	0.040	0.106	0.037	0.101
Mad	0.051	0.103	0.036	0.101
Efficiency	0.843	0.909		

Full versus partial likelihood - homogeneous case

Simulations for $N = 27$, $T = 4$.

	Partial		Full	
	\hat{p}	$\hat{\psi}$	\hat{p}	$\hat{\psi}$
Value	0.600	0.600	0.600	0.600
Median estimate	0.600	0.604	0.600	0.604
Mad	0.078	0.104	0.065	0.105
Median s.e.	0.078	0.097	0.066	0.097
Efficiency	0.709	0.991		

Application – Frogs

▶ $\hat{\psi}_{Part} = 0.556$, s.e. = 0.098

$\hat{\psi}_{Full} = 0.557$, s.e. = 0.096

▶ $\hat{p}_{Part} = 0.889$, s.e. = 0.052

$\hat{p}_{Full} = 0.780$, s.e. = 0.054

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- ▶ Equations are straight forward and can be used directly for estimation (R code). Full likelihood uses optim.
- ▶ Eventhough we ignored 1st detections, the partial works well, no significant loss of efficiency
- ▶ Analytic forms for the estimators means more stable than full likelihood
- ▶ These results are encouraging to pursue with the two-stage approach for heterogeneous case and for including covariates

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Site inhomogeneity

- ψ, p vary between sites but constant within sites.

Contribution of a single site is

$$\begin{aligned} L_i(\psi_i, p_i) &= \left\{ 1 - \psi_i + \psi_i(1 - p_i)^\tau \right\}^{1-w_i} \left\{ \binom{\tau}{x_{i\cdot}} \psi_i p_i^{x_{i\cdot}} (1 - p_i)^{\tau - x_{i\cdot}} \right\}^{w_i} \\ &\propto (1 - \psi_i \theta_i)^{1-w_i} \psi_i^{w_i} \left\{ p_i (1 - p_i)^{(a_i - 1)} \right\}^{w_i} \left\{ p_i^{(x_{i\cdot} - 1)} (1 - p_i)^{b_i - x_{i\cdot} + 1} \right\}^{w_i} \\ &= L_{1i}(\psi_i, p_i) L_{2i}(p_i) \end{aligned}$$

Site inhomogeneity

- ψ, p vary between sites but constant within sites.

Contribution of a single site is

$$\begin{aligned} L_i(\psi_i, p_i) &= \left\{ 1 - \psi_i + \psi_i(1 - p_i)^\tau \right\}^{1-w_i} \left\{ \binom{\tau}{x_{i\cdot}} \psi_i p_i^{x_{i\cdot}} (1 - p_i)^{\tau - x_{i\cdot}} \right\}^{w_i} \\ &\propto (1 - \psi_i \theta_i)^{1-w_i} \psi_i^{w_i} \times \left\{ p_i^{(x_{i\cdot} - 1)} (1 - p_i)^{b_i - x_{i\cdot} + 1} \right\}^{w_i} \\ &= L_{1i}(\psi_i, p_i) \times L_{2i}(p_i) \end{aligned}$$

... ignore first detections

$$L_i(\eta_i, p_i) = L_{1i}(\eta_i) \times L_{2i}(p_i), \quad \eta_i = \psi_i \theta_i$$

Now we can easily include covariates...

Dimension of models to check reduces significantly!
($w_i = \text{presence}$)

Benefits of two-stage approach

- ▶ not impacted by boundary conditions
- ▶ full likelihood numerically unstable eg without constraints non-convergence, local maxima, or ests outside parameter space or extreme ests
- ▶ Bayes method may underestimate variance of posterior distribution
- ▶ penalized likelihood methods to help with instability eg occuPEN, occuPEN_CV, two-stage
- ▶ faster than full likelihood (occu in unmarked via optim) as it reduces dimension of parameter space
- ▶ covariates may be related to detection or occupancy to be associated with each site separately
- ▶ covariates may vary with time
- ▶ full access to R glm machinery, vglm and vgam etc

(Karavarsamis & Huggins (2019) (CSDA))

Conditional likelihood for detection

We model detections with a conditional likelihood

$$\begin{aligned}L(\eta_s, \mathbf{p}_s) &= (1 - \eta_s)^{z_s} \eta_s^{1-z_s} \times \left\{ \frac{\prod_{j=1}^{\tau} p_{sj}^{y_{sj}} (1 - p_{sj})^{1-y_{sj}}}{\theta_s} \right\}^{1-z_s} \\ &= L_1(\eta_s) L_2(\mathbf{p}_s).\end{aligned}$$

The contribution of site s to the log-likelihood is then

$$\ell(\eta_s, \mathbf{p}_s) = z_s \log(1 - \eta_s) + (1 - z_s) \log(\eta_s) \quad (1)$$

$$+ (1 - z_s) \left\{ \sum_{j=1}^{\tau} y_{sj} \log(p_{sj}) + \sum_{j=1}^{\tau} (1 - y_{sj}) \log(1 - p_{sj}) - \log(\theta_s) \right\}. \quad (2)$$

Use (2) to get $\hat{\beta}$ (\hat{p}) then use these ests to obtain $\hat{\alpha}$ ($\hat{\psi}$) from (1).

Replace η_s by $\tilde{\eta}_s = \psi_s \hat{\theta}_s$ in the log-partial likelihood (1) and maximise this to estimate $\boldsymbol{\alpha}$.

($z_s = 1 - w_s$ indicator of no detections)

Stage 1: Estimate detection p and coefficients

$$L_2(\boldsymbol{\beta}) = \prod_{s=1}^O \frac{p_s^{y_s} (1 - p_s)^{\tau - y_s}}{\theta_s},$$

a function of the number of detections at each site where there was at least one detection, i.e. y_s , $s = 1, \dots, O$.

- model redetections with logistic regression (positive binomial family in vgam)
- simple binomial function e.g. GLMs, GAMs, VGAMs etc.
- $p(u_s, \boldsymbol{\beta}) = h(u_s^T \boldsymbol{\beta})$, $j = 1, \dots, \tau$, vector of covariates u_s and coefficient vector $\boldsymbol{\beta} \in \mathbb{R}^q$. $h(x) = (1 + \exp(-x))^{-1}$, logistic function.

Now can use GLM family and covariates to get

- $\hat{\boldsymbol{\beta}}$ unlike the simple homogeneous model the conditional likelihood estimators will not be the mle's
- estimated covariance $\widehat{V}_{\boldsymbol{\beta}}$ for $\hat{\boldsymbol{\beta}}$
 - \hat{p}
 - $\text{Var}(\hat{p}) = \hat{\boldsymbol{\beta}}^T \widehat{V}_{\boldsymbol{\beta}} \hat{\boldsymbol{\beta}}$

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Stage 1: Estimate detection p and coefficients

- ▶ Fitting the detection model for time homogeneous covariates.
- ▶ With the univariate response Y , the implementation is very similar to `glm`.
- ▶ Term `omit.constant=TRUE` does not affect the fitting but removes the constant terms from the computation of the AIC.
- ▶ Data frame 'data' is a reduced data frame that contains data from the sites where occupancy was detected.
- ▶ Variable Y is the number of times the species was detected at each occupied site,
- ▶ $\tau(= 3)$ is the number of visits to each site s , ($S = 1, \dots, 656$ sites)
- ▶ Site covariates are `vegcov1, vegcov2, \dots, vegcov6`.

```
> V.out=vglm(cbind(Y,3-Y) vegcov1+vegcov2+vegcov3+vegcov4  
+vegcov5+vegcov6, family=posbinomial(omit.constant=TRUE),data=data)
```

```
> coef(V.out) (Intercept) vegcov1 vegcov2 vegcov3 vegcov4  
1.5590909 0.5493825 -0.2512287 -0.1048756 0.1656597  
vegcov5 vegcov6  
0.1186192 -0.1277806
```

Hutchinson et al. (2015) avian data

Stage 1: Time Dependent Covariates for Detection

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$$L_2(\mathbf{p}_s) = \frac{\prod_{j=1}^{\tau} p_{sj}^{y_{sj}} (1 - p_{sj})^{1-y_{sj}}}{\theta_s}.$$

- ▶ distinct probabilities p_{sj} , $j = 1, \dots, \tau$ for different visits to site s .
- ▶ detections form a sequence of independent Bernoulli trials, we observe the outcome if there is at least one detection ie redetections
- ▶ covariate vector u_{sj} contains an indicator of the visit time
- ▶ modelled by allowing the intercept to vary with the visit j , and easily implemented in VGAM package.
- ▶ four time dependent covariates measured for each visit to each site: `time`, `temp`, `cloud`, `julian`
- ▶ `time1, ..., time2, ..., julian2, julian3`

Stage 1: Time Dependent Covariates for Detection

- ▶ fitting these models is more complex as many more models are available and the response consists of the detections on each visit to the site and is hence multivariate.
- ▶ time dependent intercepts and the relationship with the site covariates remains independent of time
- ▶ specified through the `parallel.t` argument to the `posbernoulli.t` family.
- ▶ `parallel.t=FALSE~1` is the default for the `posbernoulli.t` family

```
> V.out=vglm(cbind(survey1,survey2,survey3) ~  
vegcov1+vegcov2+vegcov3+vegcov4+vegcov5+vegcov6,  
family=posbernoulli.t(parallel.t=FALSE ~ 1), data = data)  
> coef(V.out)  
(Intercept):1 (Intercept):2 (Intercept):3 vegcov1 vegcov2  
1.8583766 1.5130892 1.3527893 0.5515551 -0.2522520  
vegcov3 vegcov4 vegcov5 vegcov6  
-0.1052631 0.1663444 0.1190595 -0.1282785
```

Stage 1: Time Dependent Covariates for Detection

- time dependent covariates are time, temp, cloud and julian measured for each visit to each site
- included in the data data frame as time1, time2, ... julian2, julian3
- fit time varying covariates but constant intercept
- requires use of the xij and form2 arguments in VGAM

```
> V.out=vglm(cbind(survey1,survey2,survey3)
~vegcov1+vegcov2+vegcov3+vegcov4
+vegcov5+vegcov6+time.tij+temp.tij+cloud.tij+julian.tij,
data=Data.all,
xij=list(time.tij~time1+time2+time3-1,temp.tij~temp1+temp2+temp3-
1,
cloud.tij~cloud1+cloud2+cloud3-1,julian.tij~julian1+julian2
+julian3-1),
family=posbernoulli.t(parallel.t=FALSE~0),
form2=~vegcov1+vegcov2+vegcov3+vegcov4+vegcov5+vegcov6+time.tij
+temp.tij+cloud.tij+julian.tij+time1+time2+time3+temp1+temp2
+temp3+cloud1+cloud2+cloud3+julian1+julian2+julian3)
> coef(V.out)
(Intercept)  vegcov1    vegcov2    vegcov3    vegcov4
 1.60651791  0.54525171 -0.24061702 -0.08727207  0.16955603
vegcov5  vegcov6  time.tij  temp.tij  cloud.tij  julian.tij
0.108527 -0.112085 -0.069456 -0.238605 -0.161028 -0.264658
```

Figure: Fitting a model using with time varying covariates but constant intercept for the two-stage approach in vglm.

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Stage 2: Estimate occupancy and coefficients

Three methods:

1) Direct maximisation of the 1st partial likelihood, $L_{1s}(\psi_s, p_s)$ as a function of ψ_s , $(1 - \psi_s \theta_s)^{1-w_s} \psi_s^{w_s}$

2) Iterative Weighted Least Squares

3) Iterative method

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Stage 2: Numeric maximisation

To estimate α maximise the partial likelihood

$$\prod_{s=1}^S L_{1s}(\tilde{\eta}_s)$$

where \mathbf{p}_s and hence θ_s has been replaced by its estimator from Stage 1: $\hat{\mathbf{p}}_s = \mathbf{p}_s(\hat{\beta})$.

$$L_1(\alpha) = \prod_{s=1}^S L_{1s}(\tilde{\eta}_s) \propto \prod_{s=1}^S (1 - \psi_s \hat{\theta}_s)^{z_s} \psi_s^{1-z_s}$$

Let $w_s = 1 - z_s$, then the log-partial likelihood is

$$\ell(\alpha) = \sum_{s=1}^S \left\{ (1 - w_s) \log(1 - \psi_s \hat{\theta}_s) + w_s \log(\psi_s) \right\}.$$

This may be maximised numerically using the `optim` function in R. However, there are two other possible approaches.

$$\psi_s = h(x_s^T \alpha)$$

where x_s is a vector of covariates associated with site s and $\alpha \in \mathbb{R}^p$ is a vector of coefficients.

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Stage 2: IWLS

For a logistic model, let matrix X have s th column x_s

- ▶ $\mathbf{w} = (w_1, \dots, w_S)^T$, $E(w_s) = \eta_s = \theta_s \psi_s$, $\boldsymbol{\eta} = (\eta_1, \dots, \eta_S)^T$
- ▶ Then, as θ_s is not a function of $\boldsymbol{\alpha}$, maximising the partial log-likelihood is equivalent to maximising $\ell(\boldsymbol{\eta}) = \sum_{s=1}^S \{(1 - w_s) \log(1 - \eta_s) + w_s \log(\eta_s)\}$.
- ▶ $\boldsymbol{\eta}(\boldsymbol{\alpha})$ be $\boldsymbol{\eta}$ evaluated at $\boldsymbol{\alpha}$.
- ▶ Set $V = \text{diag}\{(1 - \boldsymbol{\eta})\boldsymbol{\eta}\}$ and $U = \text{diag}\{\theta_s \psi_s (1 - \psi_s)\}$.
- ▶ $\boldsymbol{\alpha}^{(k)}$ is estimate at the k th step and let $\mathbf{Z} = UX\boldsymbol{\alpha}^{(k)} + \mathbf{w} - \boldsymbol{\eta}(\boldsymbol{\alpha}^{(k)})$.
- ▶ $\mathbf{u}(\boldsymbol{\alpha})$ are the partial score equations

Then the estimate at the $(k + 1)$ th is

$$\begin{aligned}\boldsymbol{\alpha}^{(k+1)} &= \boldsymbol{\alpha}^{(k)} + J(\boldsymbol{\alpha})^{-1} \mathbf{u}(\boldsymbol{\alpha}^{(k)}) \\ &= \left(XUV^{-1}UX^T \right)^{-1} XUV^{-1}U\mathbf{Z}\end{aligned}$$

The IWLS estimate is obtained by repeating this step until convergence.

An estimate of the expected Fisher information corresponding to the partial likelihood, $E\{I(\boldsymbol{\alpha}, \boldsymbol{\beta})\}$, is given by $\tilde{I}(\boldsymbol{\alpha}, \boldsymbol{\beta}) = XUV^{-1}UX^T$.

Derivations in Karavarsamis & Huggins (2015), CSDA

Method 3: Iterative method

Under the logistic model

$$\psi_i(x) = \frac{\exp(\alpha^T x_i)}{1 + \exp(\alpha^T x_i)}$$

$$\psi_i(x)\theta_i = \frac{\exp(\alpha^T x_i + \log(\theta_i))}{1 + \exp(\alpha^T x_i)}.$$

If

$$a_i = \log(\theta_i) - \log\{1 + \exp(\alpha^T x_i)(1 - \theta_i)\}$$

Method 3: Iterative method

Under the logistic model

$$\psi_i(x) = \frac{\exp(\alpha^T x_i)}{1 + \exp(\alpha^T x_i)}$$

$$\psi_i(x)\theta_i = \frac{\exp(\alpha^T x_i + \log(\theta_i))}{1 + \exp(\alpha^T x_i)}.$$

If

$$a_i = \log(\theta_i) - \log\{1 + \exp(\alpha^T x_i)(1 - \theta_i)\}$$

then

$$\psi_i\theta_i = \frac{\exp(\alpha^T x_i + a_i)}{1 + \exp(\alpha^T x_i + a_i)}$$

- a_i is function of linear predictor $\alpha^T x_i$.

∴ Iterative approach

Method 3: Iterative method

- $\hat{\alpha}_0$: initial estimate for α from GLM **without** offset

1. $\hat{\alpha}_{s-1}$ is estimate of α from previous step, $s - 1$, then

$$a_i^{(s)} = \log(\theta_i) - \log\{1 + \exp(\alpha_{s-1}^T x_i)(1 - \theta_i)\}.$$

2. Fit GLM to the w_i with offset $a_i^{(s)}$ to produce a new $\hat{\alpha}_i$.

Repeat steps 1 and 2 until convergence

Variance for $\hat{\alpha}$ and $\hat{\psi}$

$$\widehat{\text{Var}}\{\hat{\alpha}(\hat{\beta})\} \approx$$

$$I\{\hat{\alpha}(\hat{\beta}), \hat{\beta}\}^{-1} + I\{\hat{\alpha}(\hat{\beta}), \hat{\beta}\}^{-1} \tilde{B}\{\hat{\alpha}(\hat{\beta}), \hat{\beta}\} \hat{V}_{\beta} \tilde{B}\{\hat{\alpha}(\hat{\beta}), \hat{\beta}\}^T I\{\hat{\alpha}(\hat{\beta}), \hat{\beta}\}^{-1},$$

gives

$$\widehat{\text{Var}}(\hat{\psi}_i) = \{\hat{\psi}_i(1 - \hat{\psi}_i)\}^2 x_i^T \widehat{\text{Var}}\{\hat{\alpha}(\hat{\beta})\} x_i$$

- \hat{V}_{β} covariance for $\hat{\beta}$
- $Q(\alpha, \beta) = \partial l(\alpha, \beta) / \partial \alpha$ - partial score function
- $I(\alpha, \beta) = - \partial Q(\alpha, \beta) / \partial \alpha$
- $\tilde{B}\{\alpha(\beta), \beta\} = \partial Q(\alpha, \beta) / \partial \beta$

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Method 3: Iterative method - Results

1000 simulations for large N and small T with 2 covariates

- Relatively unbiased
- s.e. reasonable

	$\hat{\alpha}_0$	$\hat{\alpha}_1$	$\hat{\alpha}_2$
$N = 1000, T = 4$	1.000	0.500	0.500
Median	1.007	0.509	0.508
mad	0.207	0.095	0.094
Med s.e.	0.199	0.093	0.097

	$\hat{\alpha}_0$	$\hat{\alpha}_1$	$\hat{\alpha}_2$
$N = 1000, T = 6$	1.000	0.500	0.500
Median	1.016	0.503	0.504
mad	0.120	0.075	0.075
Med s.e.	0.132	0.076	0.076

Fitting the Full Likelihood with occu

- matrix of factors, `Visit` corresponding to the three visits
- list `Obs` that contains data frames of the time varying covariates

```
> Visit=matrix(as.factor(c(rep("a",656),rep("b",656),rep("c",656))),
  ncol=3)
> Obs=list(time=as.data.frame(Model.out@T.ij[,c(1,5,9)]),
  temp=as.data.frame(Model.out@T.ij[,c(2,6,10)]),
  cloud=as.data.frame(Model.out@T.ij[,c(3,7,11)]),
  julian=as.data.frame(Model.out@T.ij[,c(4,8,12)]),
  Visit=as.data.frame(Visit))
> D=unmarkedFrameOccu(y=Model.out@Detect,
  siteCovs=as.data.frame(Model.out@X[,-1]),obsCovs=Obs)
> 0.5.out=occu(~Visit+vegcvov1+vegcvov2+vegcvov3+vegcvov4+vegcvov5+vegcvov6
  +time+temp+cloud+julian-1~vegcvov1+vegcvov2+vegcvov3+vegcvov4+vegcvov5
  +vegcvov6,data=D,engine=c("C"))
> 0.5.out@estimates
```

Figure: Fitting a model with `occu` for time varying covariates on the full model.

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Single simulated data for $N = 150$ and $T = 4$

- Estimates roughly the same

		β	s.e. β	t_β	α	s.e. α	t_α
Iterative	Inter.	-0.273	0.177	-1.540	0.206	0.229	0.900
	x	0.442	0.169	2.620	-0.075	0.224	-0.335
	x.1	0.283	0.158	1.798	0.623	0.208	2.995
Unmarked	Inter.	-0.228	0.153	-1.489	0.198	0.224	0.882
	x	0.459	0.142	3.221	-0.081	0.221	-0.368
	x.1	0.085	0.126	0.673	0.710	0.210	3.380

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IWLS and occu

- Default settings in `occu` the estimates did not converge but two-stage was fine
- “Nelder-Mead” method set to a maximum of 2000 iterations
- For the standardised data, `occu` with the default options did converge

Table: Occupancy and detection estimates for full likelihood and two-stage approaches for the (a) unstandardised and (b) standardised brook trout data. For each covariate, we report its: estimate (Estimate), standard error (se), Student’s t -statistic (t), and p -value (p). Occupancy for the two-stage approach estimated with IWLS method.

Parameter	Estimate	Full Likelihood			Two-stage			
		se	t	p	Estimate	se	t	
(a) Unstandardised								
Occupancy ψ								
Intercept	-3.9716	0.6858	-5.7914	0.0000	-4.0452	1.1218	-3.6060	0.0003
Ele	0.0013	0.0003	4.5338	0.0000	0.0013	0.0004	3.6441	0.0003
Detection p								
Intercept	0.0580	0.7352	0.0788	0.9372	-0.1609	1.2397	-0.1298	0.8938
Ele	0.0004	0.0002	1.9697	0.0489	0.0004	0.0003	1.2516	0.2167
CSA	-0.8325	0.2822	-2.9503	0.0032	-0.7438	0.2873	-2.5888	0.0096
(b) Standardised								
Occupancy ψ								
Intercept	-0.19	0.36	-0.52	0.60	-0.34	0.32	-1.04	0.30
Ele	1.53	0.45	3.42	0.00	1.48	0.40	3.71	0.00
Detection p								
Intercept	-0.14	0.35	-0.38	0.70	-0.16	0.36	-0.44	0.66
Ele	0.36	0.35	1.04	0.30	0.43	0.37	1.18	0.24
CSA	-0.82	0.28	-2.97	0.00	-0.80	0.28	-2.81	0.00

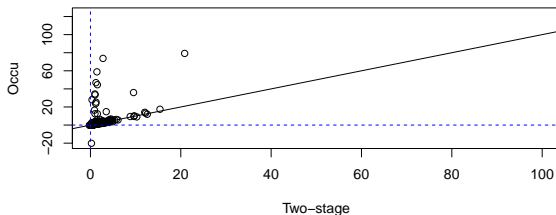
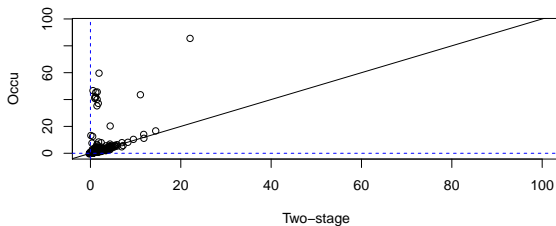


Figure: Comparison of estimated occupancy parameters ($\hat{\alpha}$) between occu and two-stage (IWLS) for 1000 simulations with $\alpha = (1, 1)$, $\beta = (-1.5, -0.5, -0.5)$. Top figure shows intercept estimates and bottom figure estimates for the slope parameter.

Agreement for intercept estimates greater, or less, than three when the actual value to be estimated is $\alpha_1 = 1$.

- number of estimates that are either both or neither greater than three ($\hat{\alpha}_1 > 3$), less than or equal to three ($\hat{\alpha}_1 \leq 3$), or when these disagree
- `occu` gives estimates $\hat{\alpha}_1 > 3$ that are large four times more often than IWLS (36 vs 12)
- no universal best method for finding estimates for occupancy
- if IWLS then try `optim` (or `occu`)

Two-stage IWLS	occu method	
	$\hat{\alpha}_1 \leq 3$	$\hat{\alpha}_1 > 3$
$\hat{\alpha}_1 \leq 3$	832	36
$\hat{\alpha}_1 > 3$	12	37

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$$p_i : \quad \kappa_i = u_i^T \alpha + f_1(v_{i1}) + \cdots + f_K(v_{iK}) \quad \begin{cases} u_i & = \text{parametric,} \\ v_i & = \text{nonparametric} \end{cases}$$

$$\psi_i : \quad \nu_i = x_i^T \alpha + g_1(r_{i1}) + \cdots + g_J(r_{iJ}) \quad \begin{cases} x_i & = \text{parametric,} \\ g_i & = \text{nonparametric} \end{cases}$$

Stage 1: \hat{p}_i

- ▶ GAMS to redetections r_i

Stage 2: $\hat{\psi}_i$

- ▶ GAMS to presence-absences w_i via iterative method with offset

$$a_i^{(s)} = \log(\hat{\theta}_i) - \log\{1 + \exp(\nu_i^{(s-1)})(1 - \hat{\theta}_i)\}$$

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$$\widehat{\text{Var}}(\widehat{\alpha}_\lambda(\widehat{\beta})) \approx I_\lambda\{\widehat{\alpha}(\widehat{\beta}), \widehat{\beta}\}^{-1} I\{\widehat{\alpha}(\widehat{\beta}), \widehat{\beta}\} I_\lambda\{\widehat{\alpha}(\widehat{\beta}), \widehat{\beta}\}^{-1} \\ + I_\lambda\{\widehat{\alpha}(\widehat{\beta}), \widehat{\beta}\}^{-1} \widetilde{B}\{\widehat{\alpha}(\widehat{\beta}), \widehat{\beta}\} \widehat{V}_\beta \widetilde{B}\{\widehat{\alpha}(\widehat{\beta}), \widehat{\beta}\}^T I_\lambda\{\widehat{\alpha}(\widehat{\beta}), \widehat{\beta}\}^{-1},$$

gives

$$\widehat{\text{Var}}(\widehat{\psi}_s^{(\lambda)}) = \{\widehat{\psi}_s(1 - \widehat{\psi}_s)\}^2 x_s^T \widehat{\text{Var}}\{\widehat{\alpha}_\lambda(\widehat{\beta})\} x_s$$

where $I_\lambda(\alpha, \widehat{\beta}) = I(\alpha, \widehat{\beta}) + \lambda_S \mathcal{P}^*$.

- V_β^* covariance for $\widehat{\beta}$
- $Q(\alpha, \beta) = \partial l(\alpha, \beta) / \partial \alpha$
- $I(\alpha, \widehat{\beta}) = - \partial Q(\alpha, \beta) / \partial \alpha$
- $\widetilde{B}\{\widehat{\alpha}(\widehat{\beta}), \widehat{\beta}\} = \partial Q(\alpha, \beta) / \partial \beta$
- λ_S smooth
- \mathcal{P}^* penalty matrix

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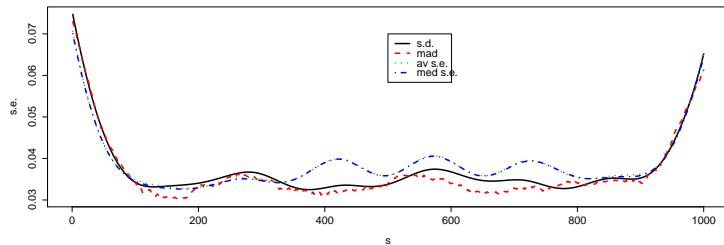
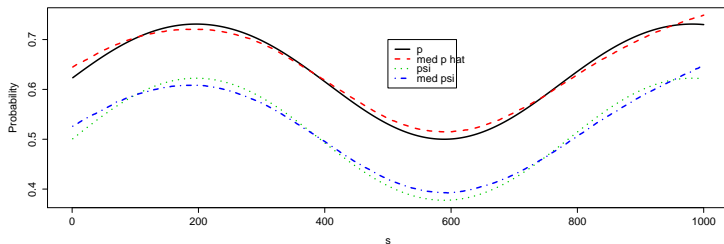
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		beta	se.beta	t.beta	alpha	se.alpha	t.alpha
Iter.	(Intercept)	-2.700	11.350	-0.238	147.663	1242.713	0.119
	ele	1.896	31.177	0.061	319.120	—	—
	forest	0.673	0.650	1.035	-0.719	19.815	-0.036
	s(ele).1	-18.100	30.030	-0.603	-102.503	2116.133	-0.048
	s(ele).2	2.749	58.341	0.047	-472.991	1207.377	-0.392
	s(ele).3	21.185	64.824	0.327	-563.946	2289.131	-0.246
	s(ele).4	12.077	47.065	0.257	-603.291	3555.318	-0.170
	s(ele).5	15.019	27.374	0.549	-725.366	2411.729	-0.301
	s(ele).6	6.219	29.648	0.210	-689.764	4113.145	-0.168
	s(ele).7	29.652	35.357	0.839	-775.511	1446.579	-0.536
s(ele).8	-14.317	26.192	-0.547	-768.387	2917.795	-0.263	
s(ele).9	-49.528	73.198	-0.677	2314.131	10896.921	0.212	
occu	int	-1.742	0.238	-7.311	3.950	1.931	2.045
	ele	0.829	0.392	2.112	2.141	0.862	2.484
	ele.sq	0.745	0.445	1.672	-3.549	1.533	-2.316
	forest	0.131	0.187	0.702	0.609	0.737	0.826

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- ▶ Occupancy models appear simple but are harder than expected
 - ▶ We resolved problems with construction of estimators and interval estimators
- ▶ Full likelihood possible but does not allow easy access to GLM machinery
- ▶ Welsh et al. (2013) show that problems with the full likelihood feed through to the covariate model
 - ▶ Partial likelihood allows full access to GLM machinery at both stages
 - ▶ Estimators from both stages are probabilities (so naturally constrained to between 0 and 1)

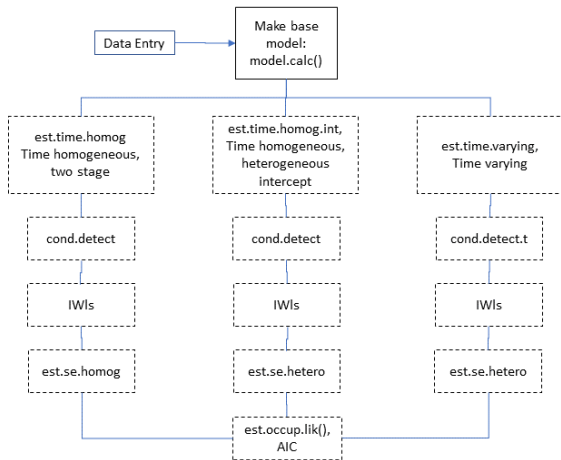


Figure: Flowchart of twostage algorithm

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Thank you

About me and contact info at

<https://natalie-karavarsamis.github.io>

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