

Distance Sampling Simulations

Overview

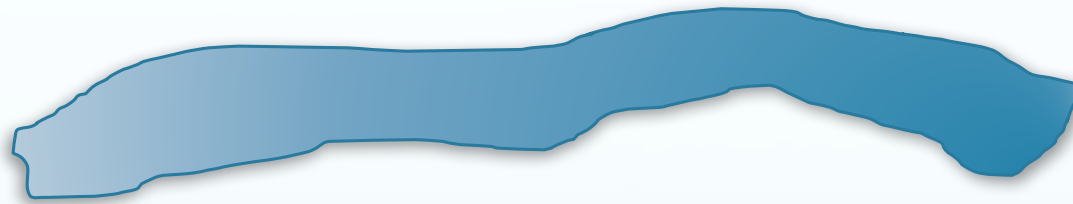
- Why simulate?
- How it works
- Automated survey design
 - Coverage probability
 - Which design?
 - Design trade-offs
- Defining the population
 - Population description
 - Detectability
- Example Simulations

Why Simulate?

- Surveys are expensive, we want to get them right!
(simulations cheap)
- Test different survey designs
- Test survey protocols
- Investigate violation of assumptions
- Investigate analysis properties

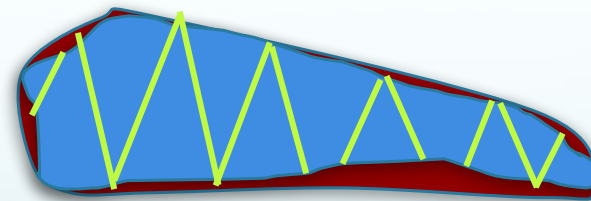
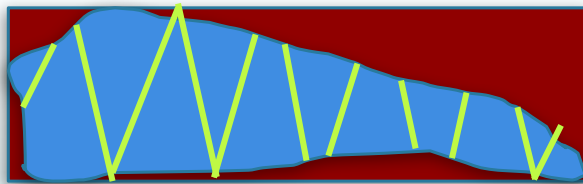
Why Simulate?

- I have a fairly long and narrow study region, are edge effects likely to be a problem?



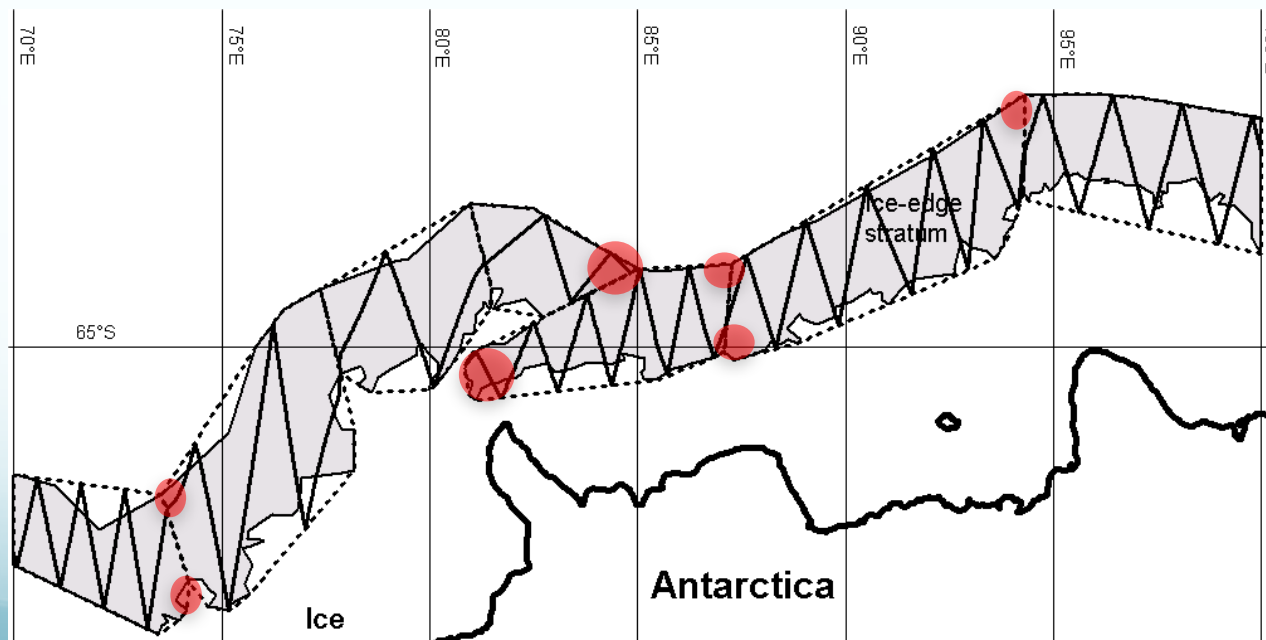
Why Simulate?

- Generating my equal spaced zig zag design in a convex hull gives better efficiency (less off effort transit time) but is this likely to introduce large amounts of bias due to non uniform coverage probability?



Why Simulate?

- What is the potential bias in this stratification technique?



Why Simulate?

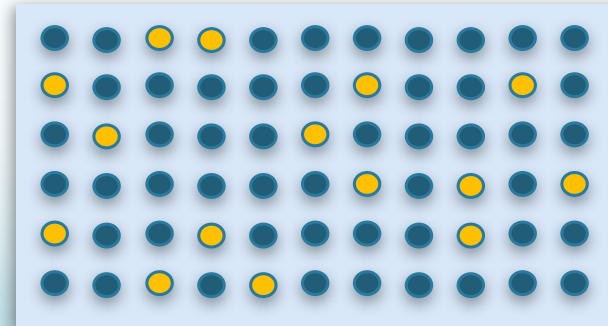
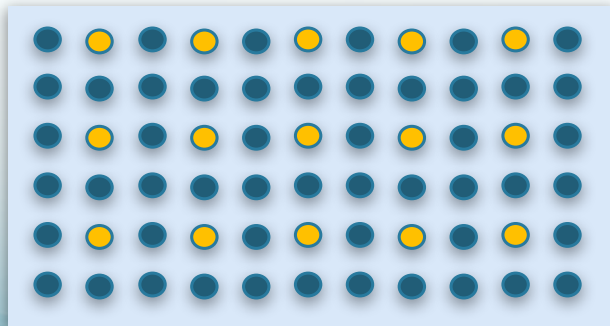
- From pilot study trials I know that there can be multiplicative error on recorded distances
- This error has a $\sim 15\%$ CV when collecting data in 3 bins or $\sim 30\%$ CV when attempting to collect exact distances... which is preferable (if we cannot improve accuracy or correct the measurements)?

Why Simulate?

- We suspect that the current survey design is less than ideal and may be introducing bias but people are reluctant to change...
- Simulate the current situation to get an idea of how bad things could be
- Simulate a new design to show how things could be improved

Why Simulate?

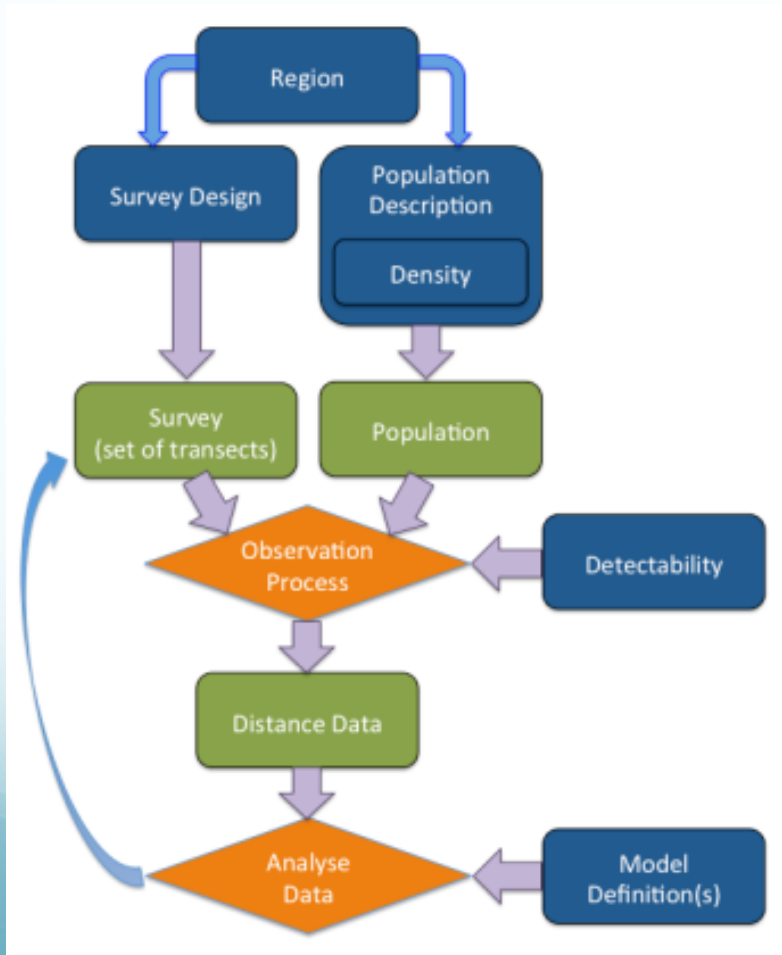
- I want to do an acoustic survey with two types of detectors.
 - The first records distances as per standard distance sampling requirements (standard detectors).
 - The second only records the presence of a sound (simple nodes).
- How many standard nodes do I need and how should I distribute them?



Why Simulate?

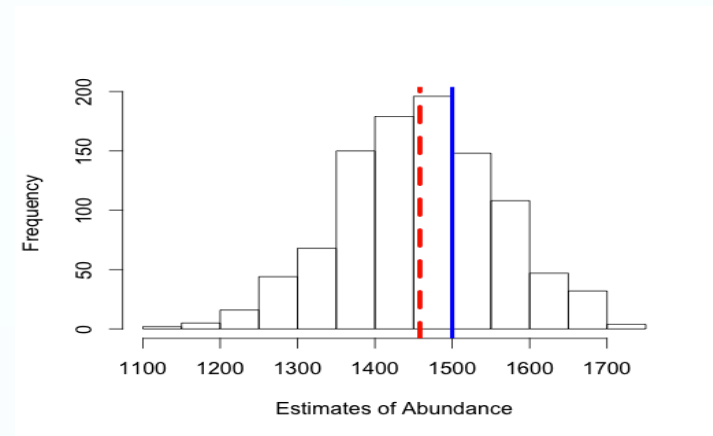
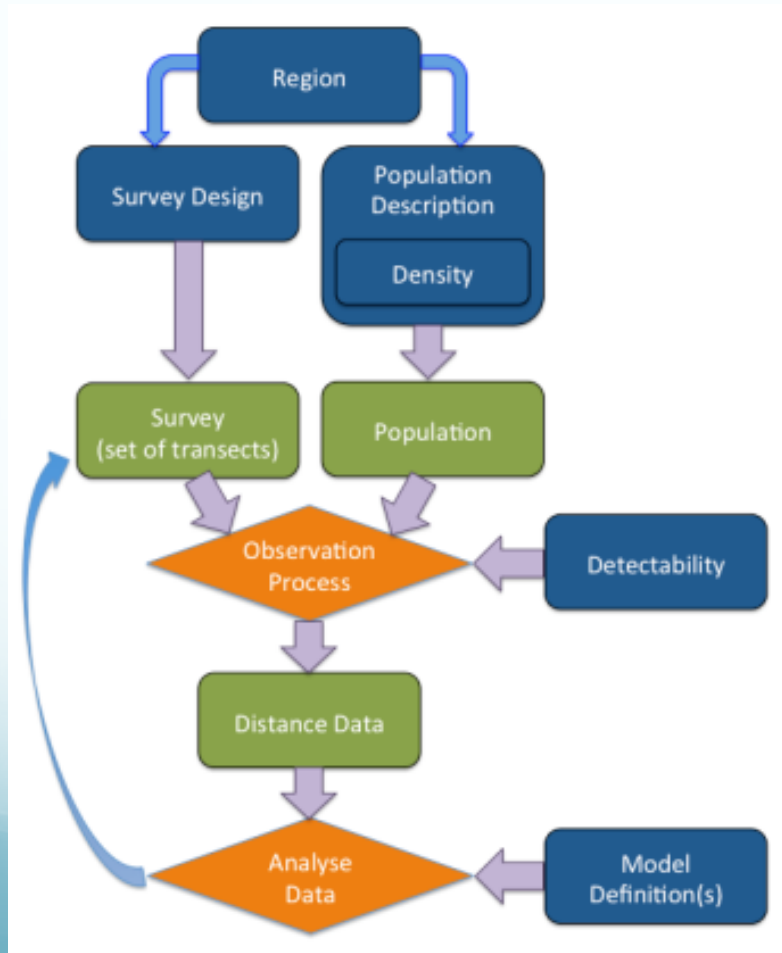
- I would like to use my data to generate both design (standard distance sampling) and model based (density surface model) estimates of density... which design will work best for my study?
- Hopefully coming soon to DSsim...
- Some example simulations can be found here:
<https://github.com/DistanceDevelopment/DSsim/wiki>

How it works



- Blue rectangles indicate information supplied by the user.
- Green rectangles are objects created by DSsim in the simulation process.
- Orange diamonds indicate the processes carried out by DSsim.

How it works



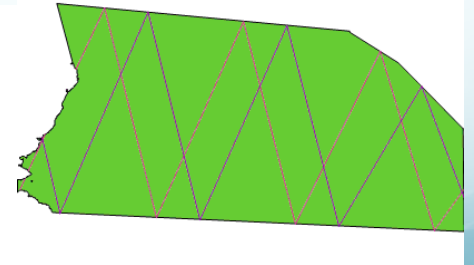
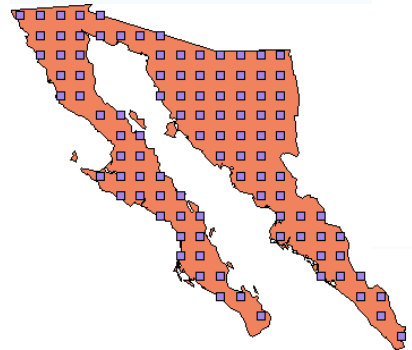
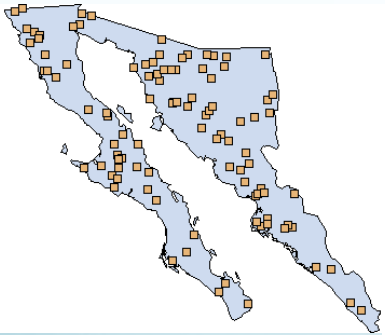
Assess:

- Bias
- Precision
- CI coverage

Across different designs/scenarios

Automated Survey Design

- Generate random sets of transects according to an algorithm
 - Assess design properties
 - Generate multiple transect sets for simulations

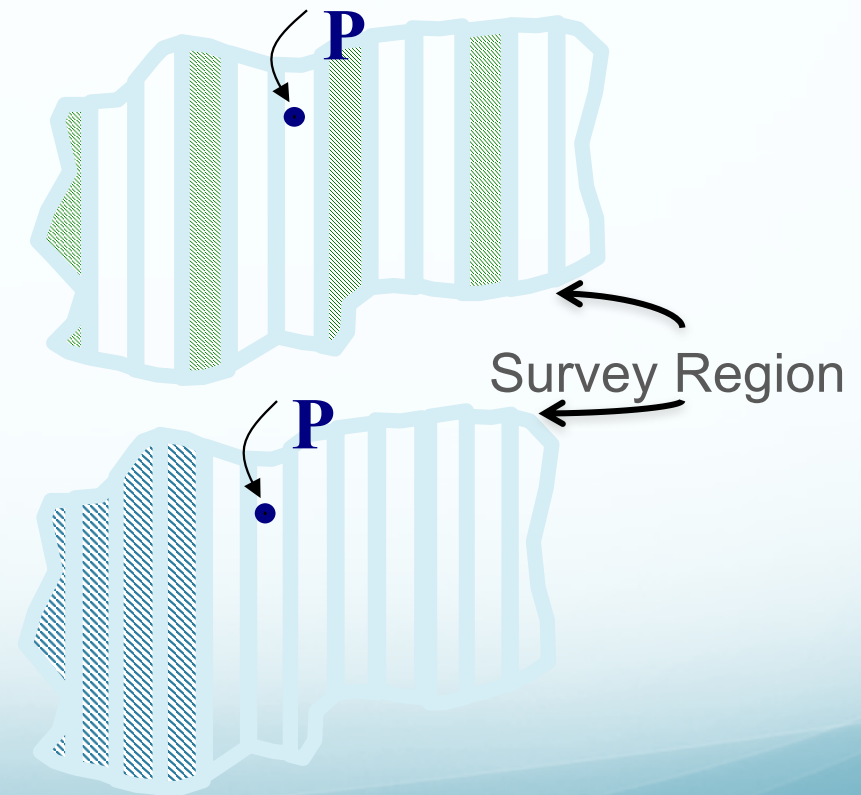


Automated Survey Design

- Coverage Probability

- Uniform coverage probability, $\pi = 1/3$
- Even coverage for any given realisation

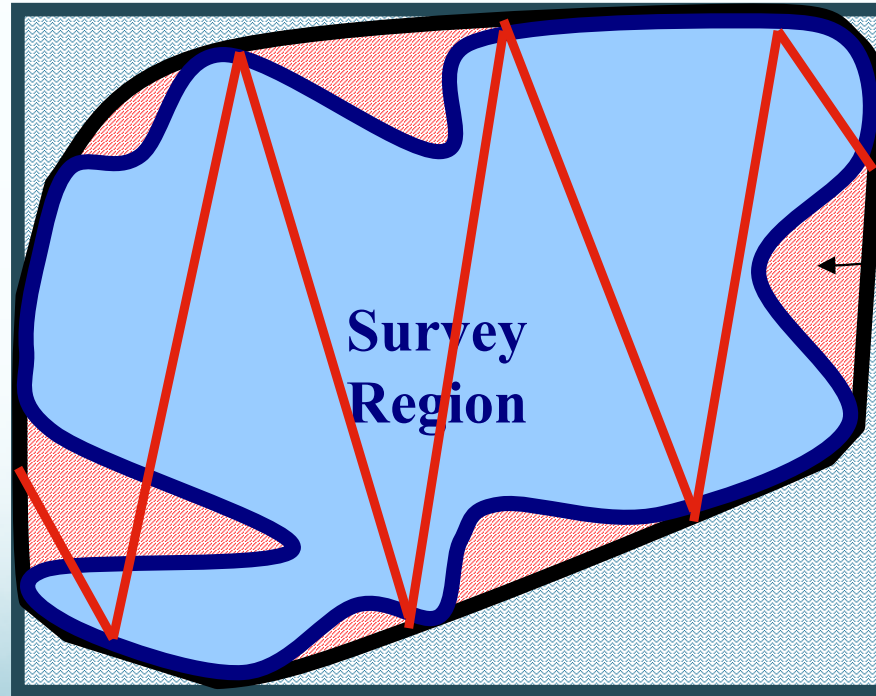
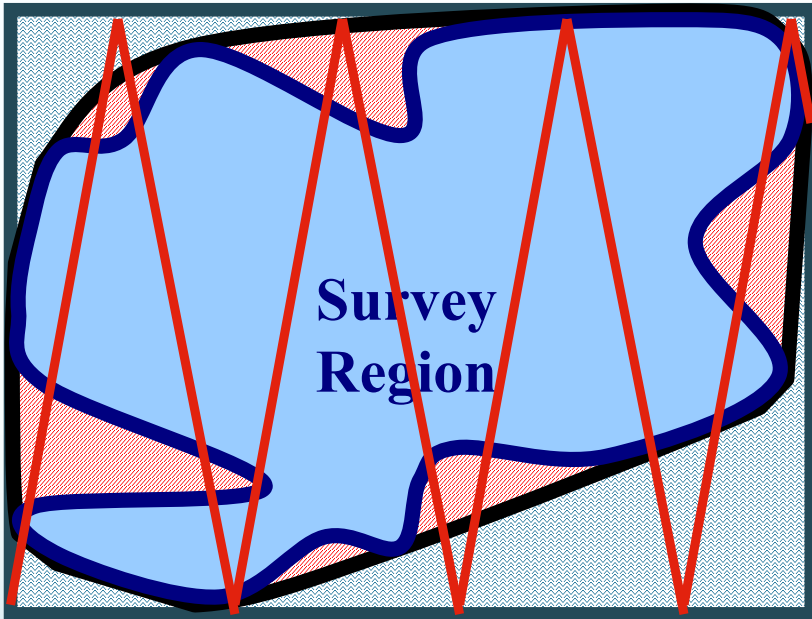
- Uniform coverage probability, $\pi = 1/3$
- Uneven coverage for any given realisation



Which Design?

- **Uniformity** of coverage probability
- **Even-ness** of coverage within any given realisation
- **Overlap** of samplers
- **Cost** of travel between samplers
- **Efficiency** when density varies within the region

Design Trade-Offs



Convex hull

Minimum bounding rectangle

Population Definition

- True population size?
- Occur as individuals or clusters?
- Covariates which will affect detectability?
- How is the population distributed within the study region?
 - Ideally have a previously fitted density surface
Otherwise test over a range of plausible distributions

Detectability

- Distance needs:
 - shape and scale parameters on the natural scale
 - covariate parameters on the log scale

Detectability

- Golftees project

$$\exp(0.268179) = 1.307581$$

Detection Fct/Global/Parameter Estimates (MCDS)

```

Effort      :    210.0000
# samples   :         1
Width      :    4.000000
# observations:    162

Model
Half-normal key, k(y) = Exp(-y**2/(2*s**2))

s = A(1) * Exp(fcn(A(2)) + fcn(A(3)))
Parameter A(1) is the intercept of the scale parameter s.
Parameter A(2) is the coefficient of covariate CLUSTER SIZE.
Parameter A(3) is the coefficient of level 0 of factor covariate SEX.
    
```

Parameter	Point Estimate	Standard Error	Percent of Variation	95 Percent Confidence Interval	
A(1)	2.622	0.8370			
A(2)	0.9294E-01	0.8172E-01			
A(3)	-0.6951	0.2937			
f(0)	0.36330	0.17850E-01	4.91	0.32972	0.40030
p	0.68814	0.33810E-01	4.91	0.62454	0.75821
ESW	2.7525	0.13524	4.91	2.4981	3.0329

Natural scale

Log scale

Detection Fct/Summary (MRDS)

```

Summary for ds object
Number of observations :    162
Distance range       :    0 - 4
AIC                  :    428.572

Detection function:
Half-normal key function

Detection function parameters
Scale coefficient(s):
              estimate      se
(Intercept) 0.26817900 0.27140001
size         0.09314751 0.08176431
sex1         0.69600047 0.29401571

              Estimate      SE      CV
Average p     0.6882835 0.05258548 0.07640090
N in covered region 235.3681131 21.00939868 0.08926187
    
```

Detectability

- In simulation:

Detectability

Detection function model: Half-Normal

Define parameters for each stratum

Region	Study Area
Scale	1.31
Shape	
cluster size	0.093
sex.0	0
sex.1	0.696

(The units for the detection function are 'Meter')



Detectability

Detection function model: Half-Normal

Define parameters for each stratum

Region	Study Area
Scale	2.62
Shape	
cluster size	0.093
sex.0	-0.696
sex.1	0

(The units for the detection function are 'Meter')

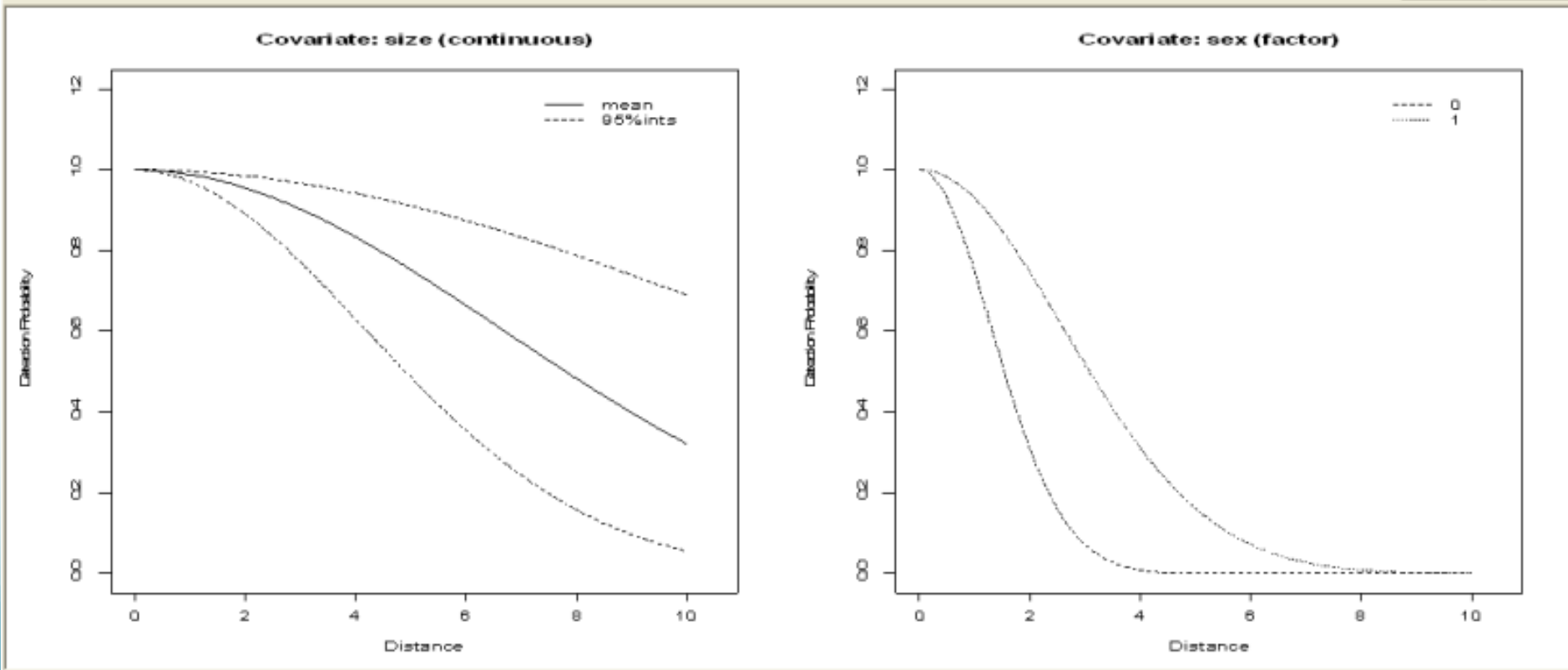
$$\exp(\log(1.307581)+0.696) = 2.622633$$

$$\exp(\log(2.622)-0.696) = 1.307265$$

Detectability

Plot: Probability of Detection

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Analysis

- **Data Filter** must specify a right truncation distance
- **Model Definition** must be either MRDS or MA
 - MRDS – for fitting a specific model
 - MA – for model selection (Note: MA model definitions require the creation of analyses)

Any questions so far...

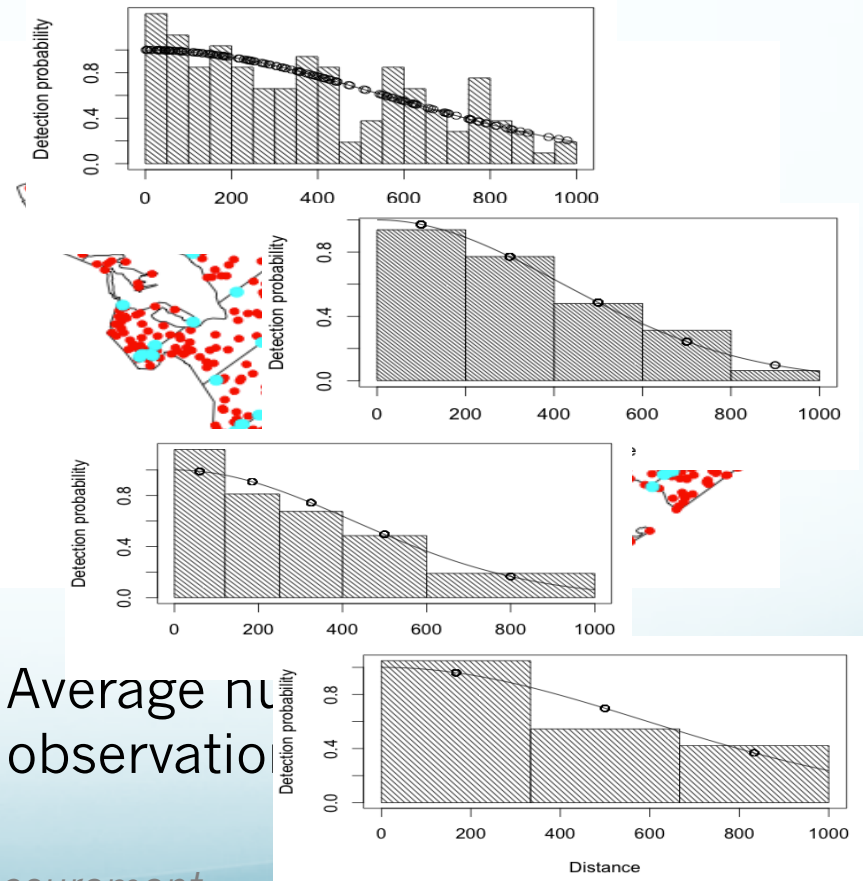
Example Simulations

- To bin or not to bin?
 - It is better to collect binned data accurately than attempt to collect exact distances and introduce measurement error!
- Testing pooling robustness in relation to truncation distance.
 - Demonstrating why you shouldn't be scared to truncate distance sampling data
- Comparison of subjective and random designs.
 - How wrong can you go with a subjective design?
 - Comparing zig zag and parallel designs.

To Bin or Not to Bin?

Simulation:

- Generated 999 datasets
- Added multiplicative measurement error
 - Distance = True Distance * R
 - $R = (U + 0.5)$, where $U \sim \text{Beta}(\theta, \theta)^1$
 - No error, ~15% CV ($\theta = 5$), ~30% CV ($\theta = 1$)
- Analysed them in difference ways
 - Exact distances, 5 Equal bins, 5 Unequal bins, 3 Equal bins
- Model selection on minimum AIC
 - Half-normal v Hazard rate



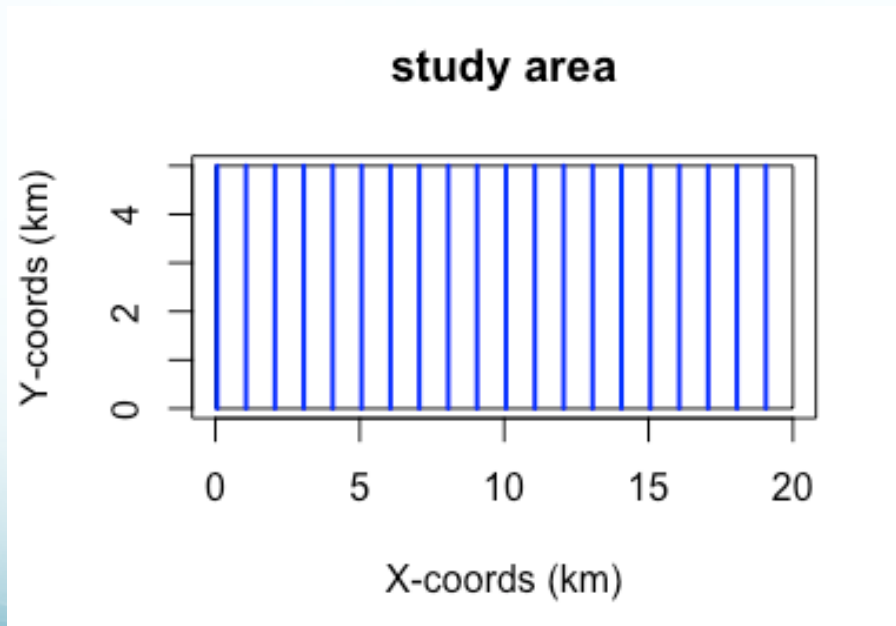
¹Marques T. (2004) Predicting and correcting bias caused by measurement error in line transect sampling using multiplicative error models *Biometrics* **60**:757--763

To Bin or Not to Bin Results

	Exact Distances	5 Equal Bins	5 Unequal Bins	3 Equal Bins
No Error	-1.16% bias 210 SE	-1.11% bias 217 SE	-0.16% bias 221 SE	-0.19% bias 255 SE
15% CV	0.48% bias 214 SE	0.5% bias 221 SE	1.36% bias 221 SE	1.72% bias 264 SE
30% CV	6.66% bias 237 SE	6.61% bias 250 SE	7.43% bias 262 SE	8.20% bias 338 SE

Pooling Robustness and Truncation

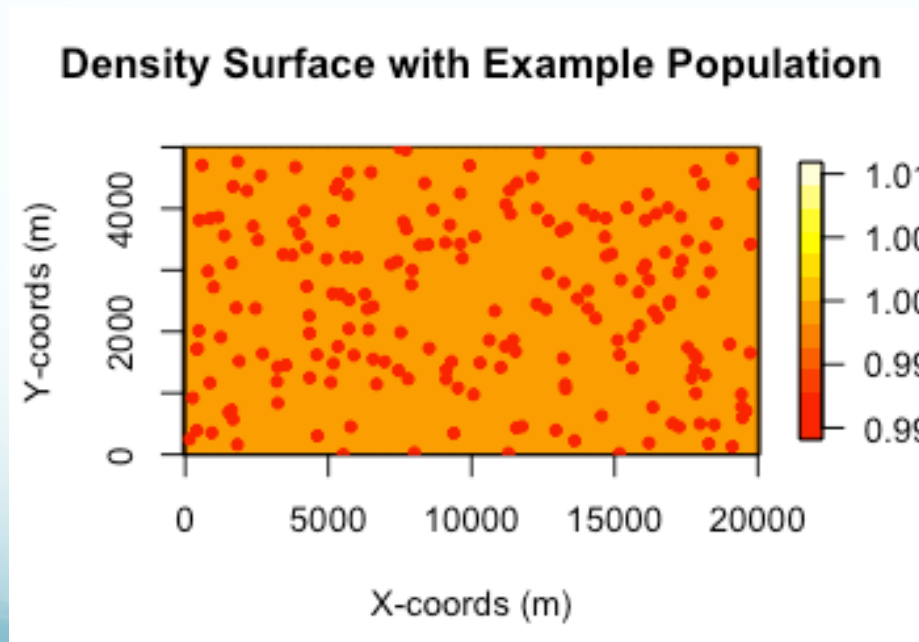
- DSsim vignette



- Rectangular study region
- Systematic parallel transects with a spacing of 1000m

Pooling Robustness and Truncation

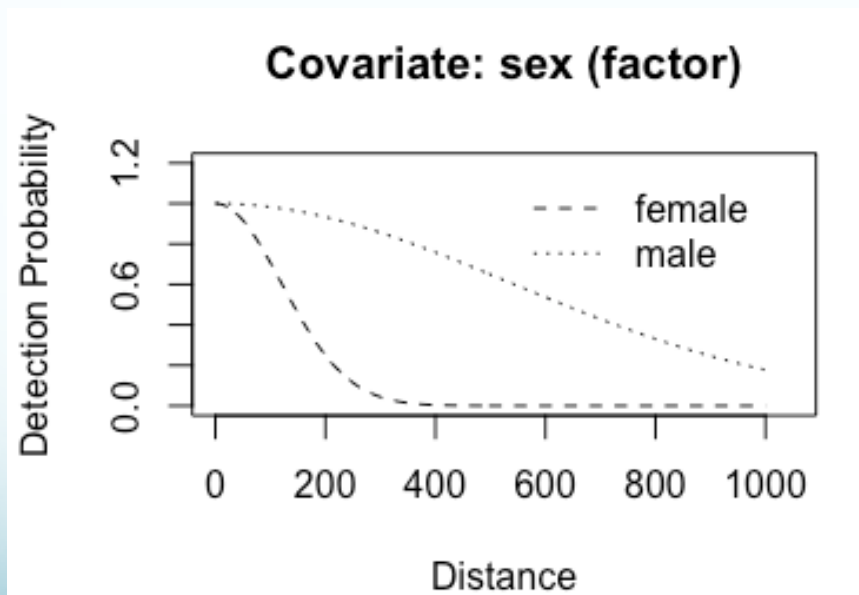
- DSsim vignette



- Uniform density surface
- Population size of 200
- 50% male, 50% female

Pooling Robustness and Truncation

- DSsim vignette



- Half-normal shape for detectability
- Scale parameter of 120 for the females
- Scale parameter of ~540 for the males

Pooling Robustness and Truncation

- DSsim vignette

```
# Create the covariate parameter list
cov.params <- list()
# Note the covariate parameters are supplied on the log scale
cov.params$sex = data.frame(level = c("female", "male"),
                             param = c(0, 1.5))

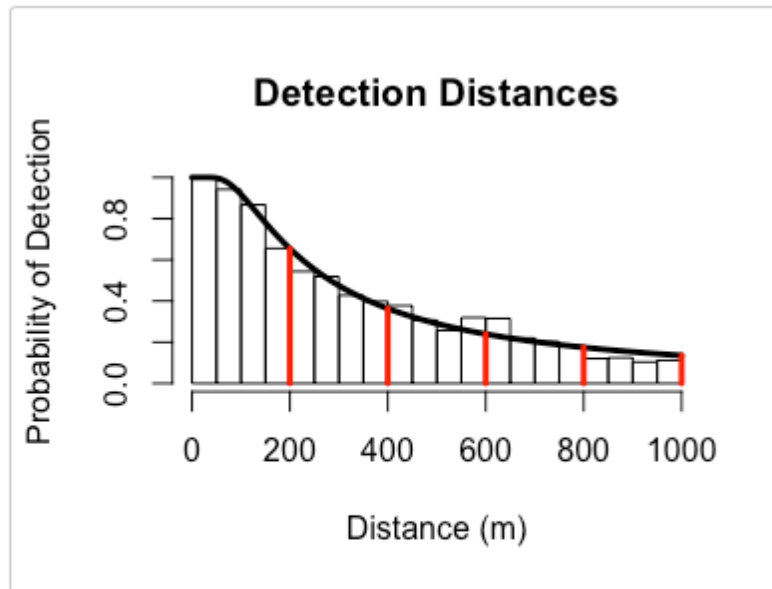
detect.cov <- make.detectability(key.function = "hn" ,
                                scale.param = 120,
                                cov.param = cov.params,
                                truncation = 1000)
```

$$\exp(\log(120)+1.5) = 537.8$$

- Half-normal shape for detectability
- Scale parameter of 120 for the females
- Scale parameter of ~540 for the males

Pooling Robustness and Truncation

- DSsim vignette



- Two types of analyses:
 - **hn v hr**
 - $hn \sim \text{sex}$
- Selection criteria: AIC

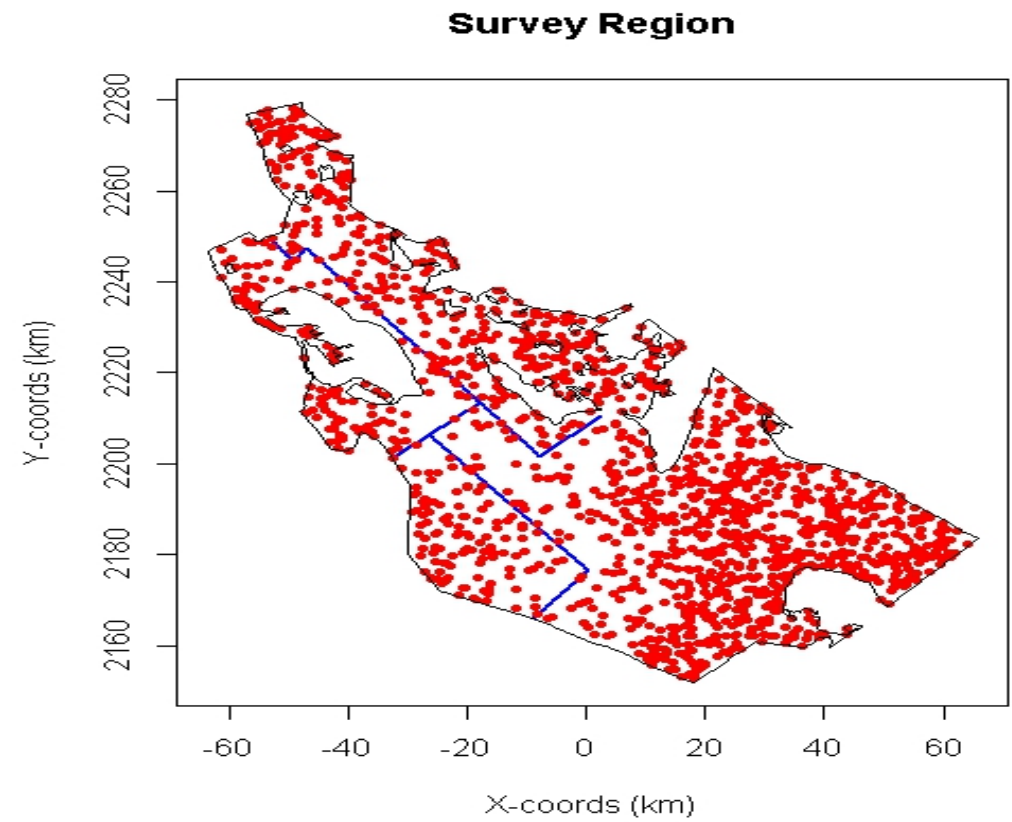
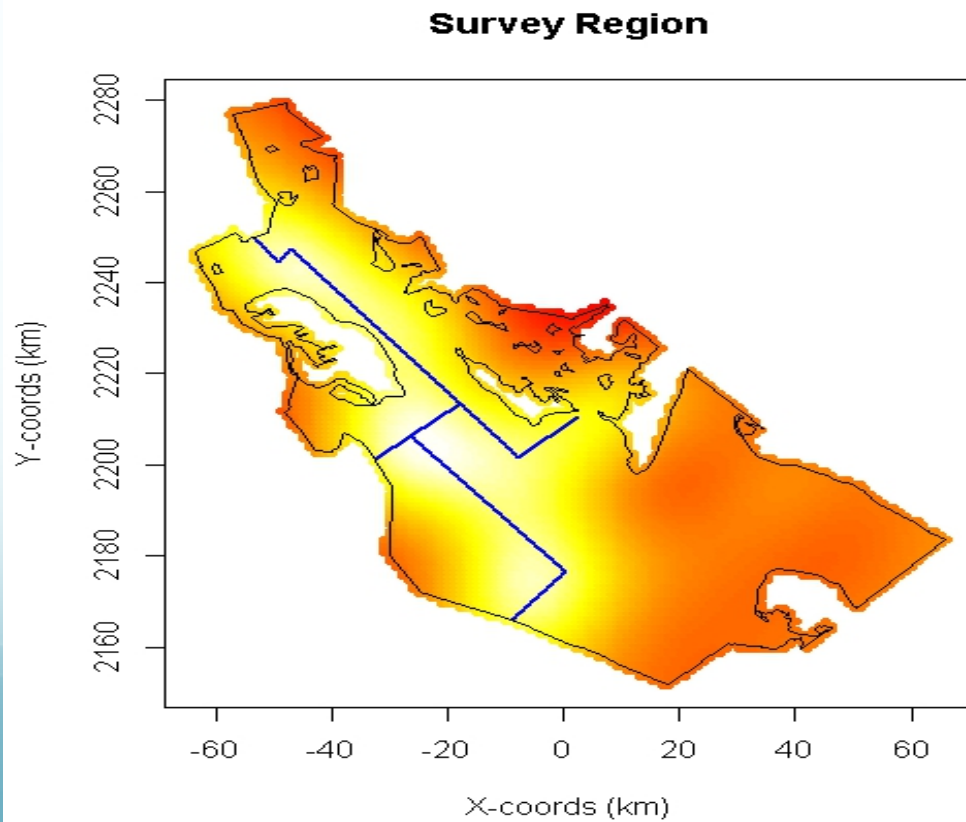
Histogram of data from covariate simulation with manually selected candidate truncation distances.

Pooling Robustness and Truncation

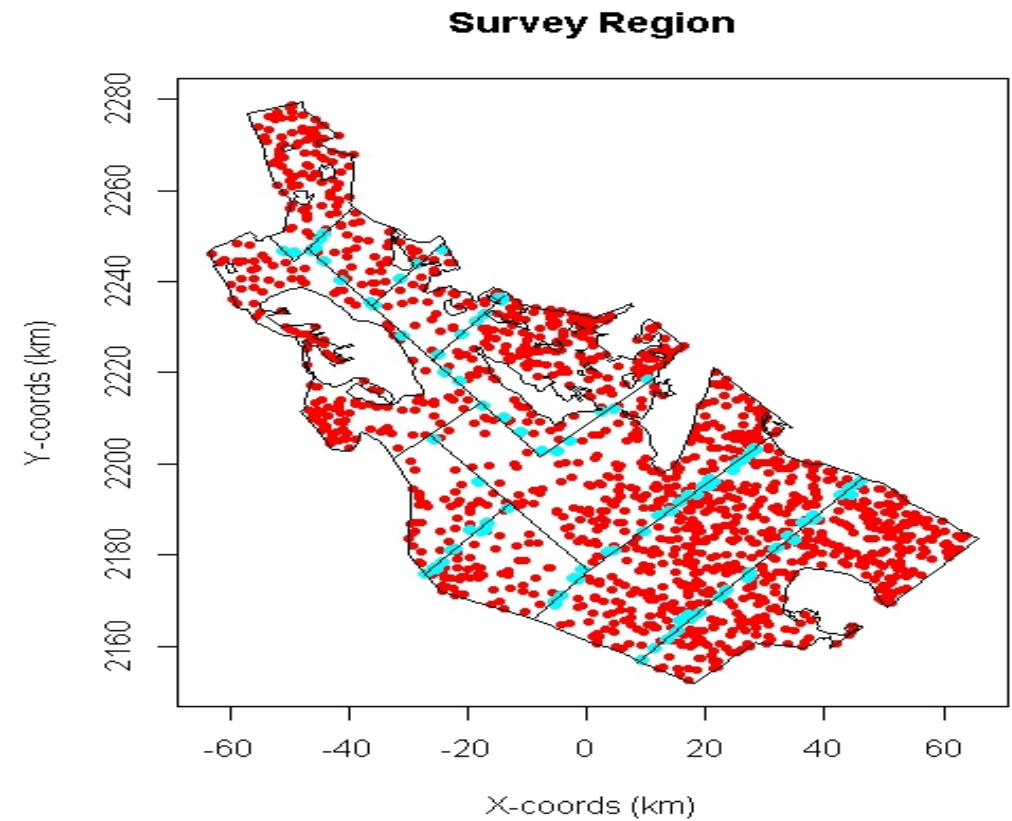
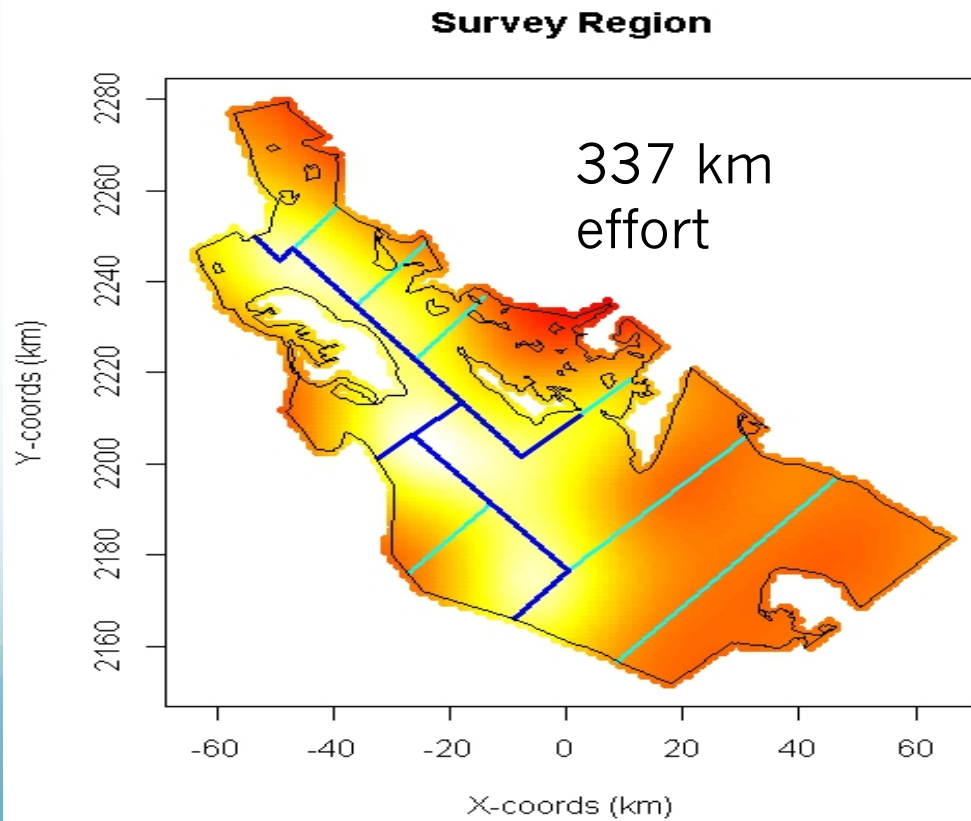
- Results HN v HR:

<i>Truncation</i>	<i>mean n</i>	<i>mean \hat{N}</i>	<i>mean se</i>	<i>SD(\hat{N})</i>	<i>%Bias</i>	<i>RMSE</i>	<i>% CI Coverage</i>
200	66	197	34.27	34.05	-1.32	34.13	97.5
400	102	190	31.06	34.79	-5.13	36.25	87.9
600	128	190	34.04	35.27	-5.24	36.77	81.9
800	144	190	34.31	36.61	-5.10	37.99	77.1
1000	154	184	30.93	39.49	-7.76	42.42	68.1

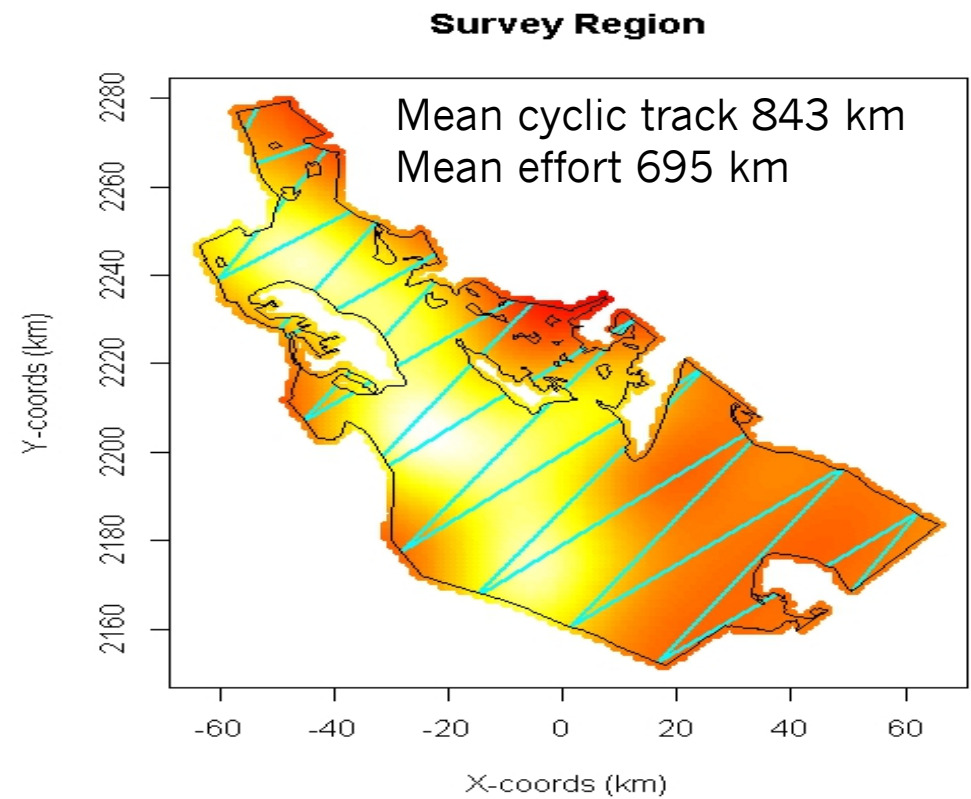
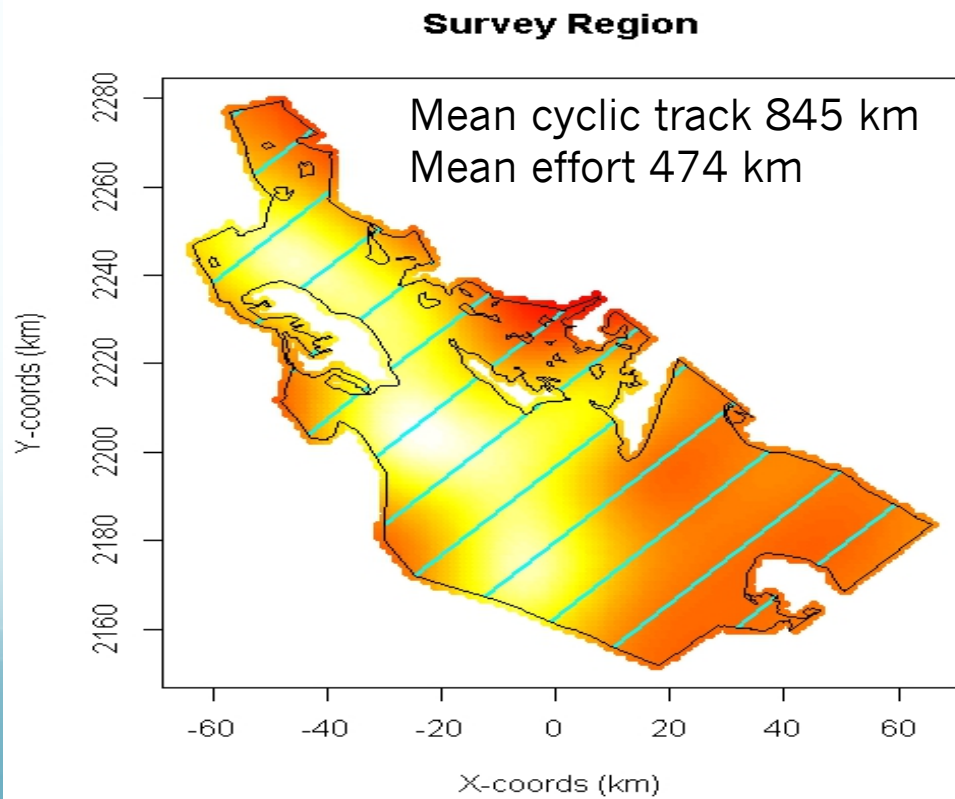
Example Simulation



Subjective survey design



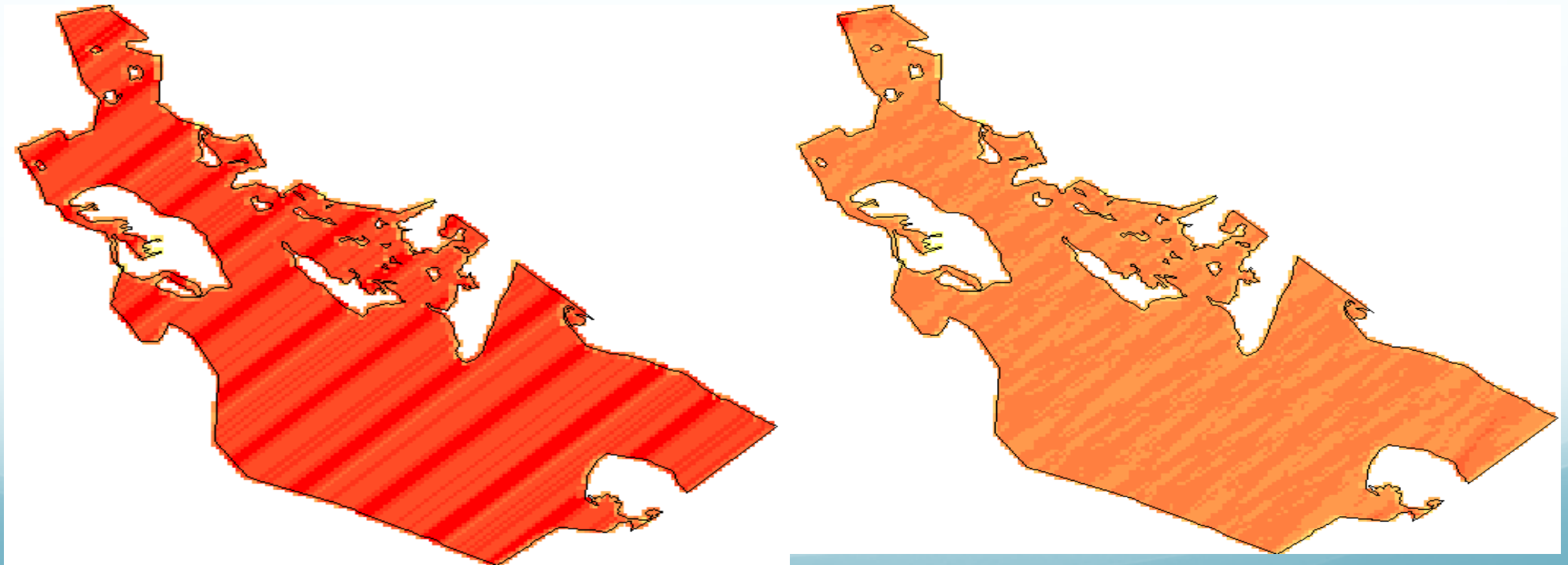
Random Designs



Coverage probability

Systematic Parallel Design

Equal Spaced Zigzag Design



Simulation

- Generates a realisation of the population based on a fixed N of 1500
- Generates a realisation of the design
 - Different each time for the random designs
 - The same each time for the subjective design
- Simulates the detection process
- Analyses the results
 - Half-normal
 - Hazard-rate
- Repeats a number of times

Practical

- Now attempt the DSsim practical:
 - *R version – subjective design and parallel v zig zag*
 - *Distance version – parallel v zig zag only*
- You will need the library *shapefiles*.