

# 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

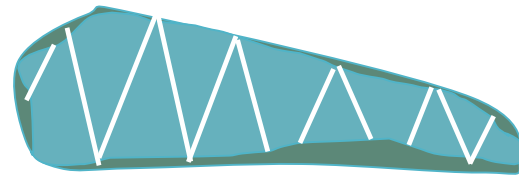
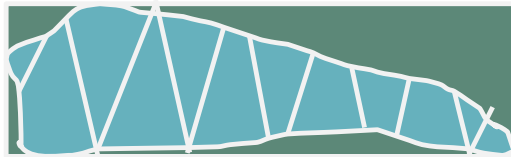
# Why Simulate?

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



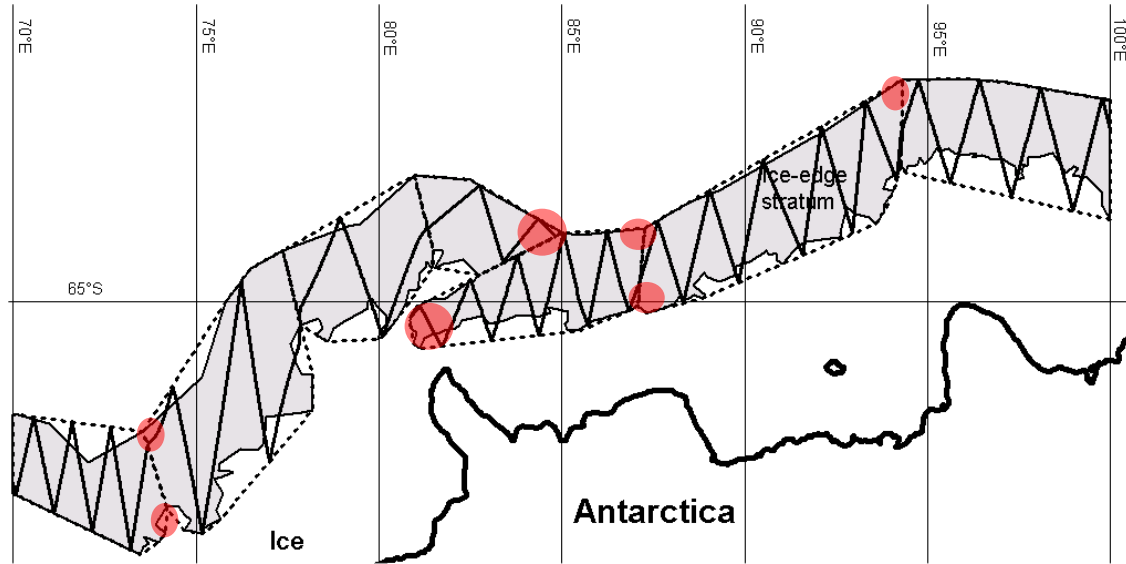
# Why Simulate?

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 ~15% CV when collecting data in 3 bins or ~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



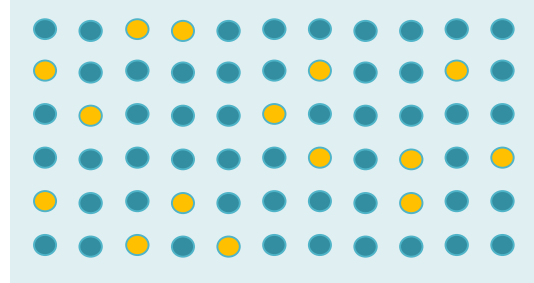
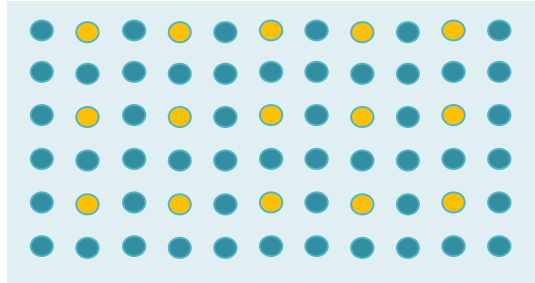
# Why Simulate?

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:

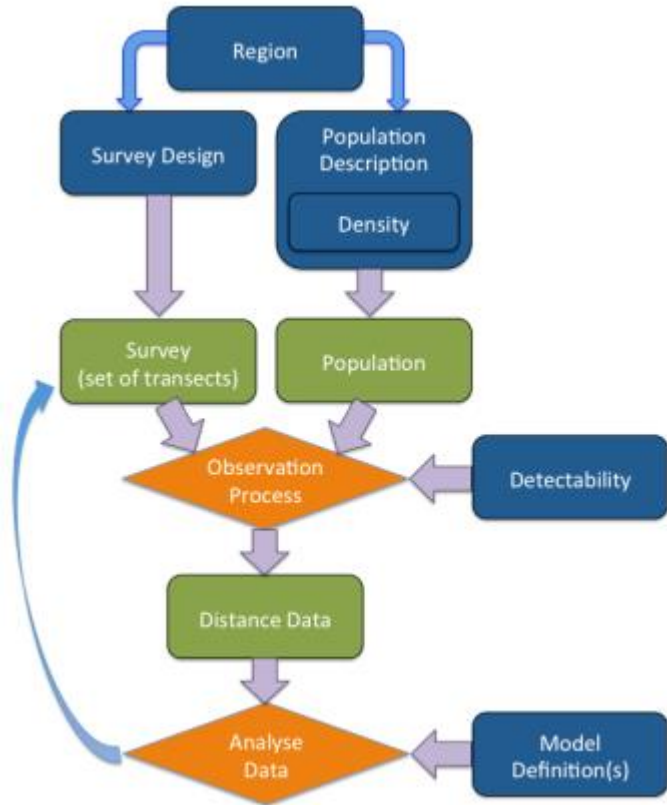
<https://github.com/DistanceDevelopment/DSsim/wiki>

# How it works

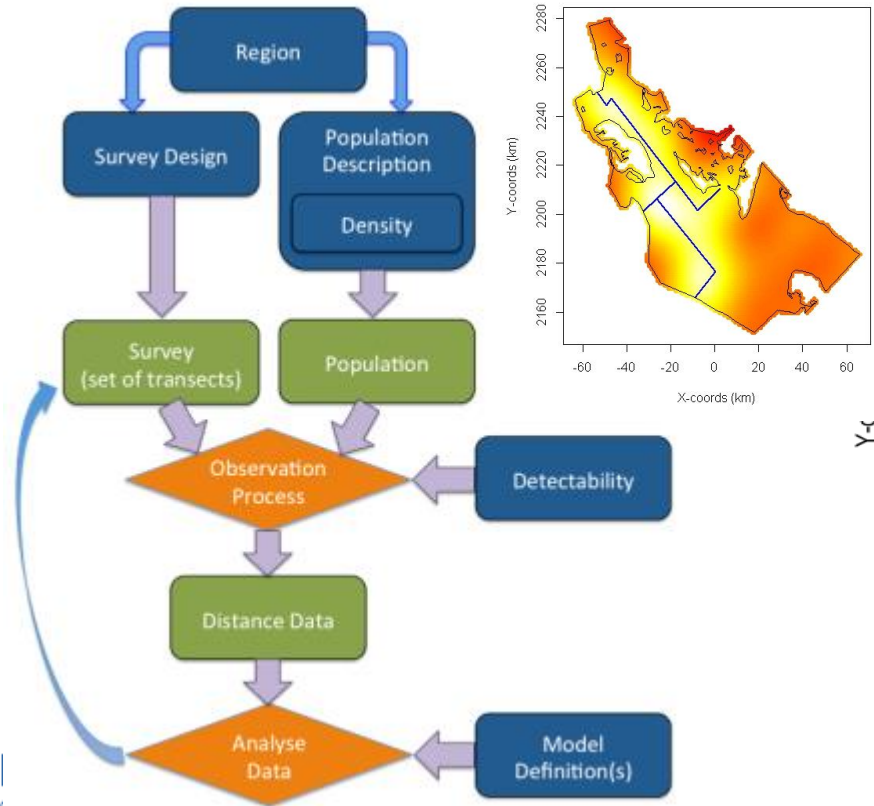
Blue rectangles indicate information supplied by the user.

Green rectangles are objects created by DSsim in the simulation process.

Orange diamonds indicate the processes carried out by DSsim.



# How it works



**Survey Region**

Population Description AIC = 2747

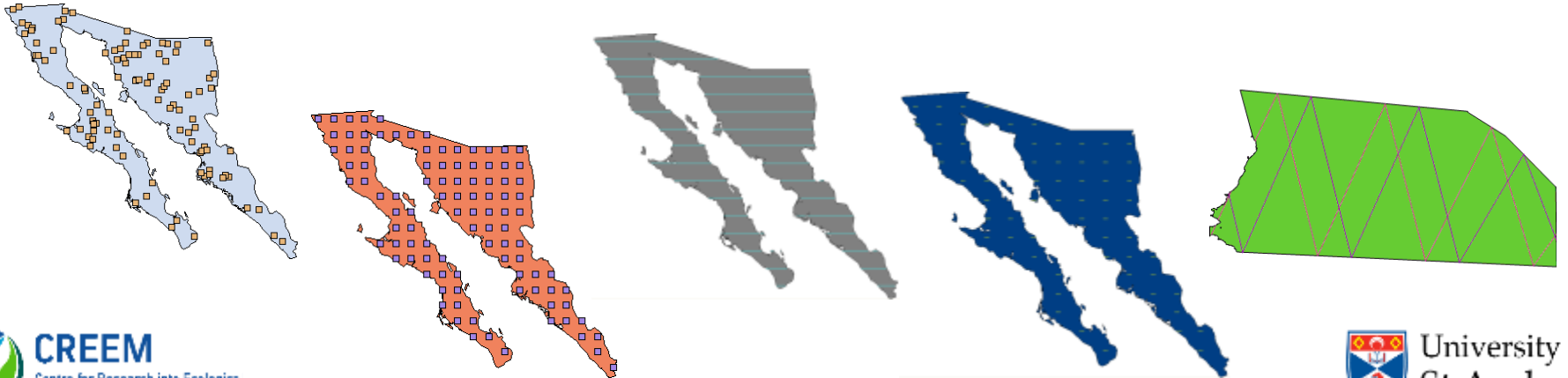
- Population size or density
- Zig zag design
- Density surface
- Equal Spaced
- Clusters?
- Spacing = 10km
- Covariates affecting?
- Minus sampling

AIC = 2748

Across different designs/scenarios

# Automated Survey Design

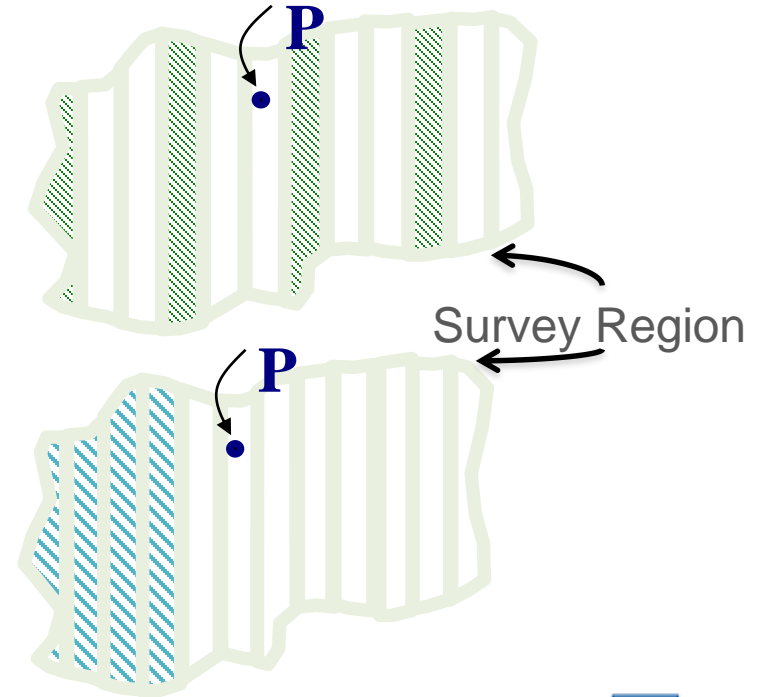
- Generate random sets of transects according to an algorithm
- Assess design properties
- Generate multiple transect sets for simulations



# Automated Survey Design

## Coverage Probability

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



# Which Design?

**Uniformity** of coverage probability

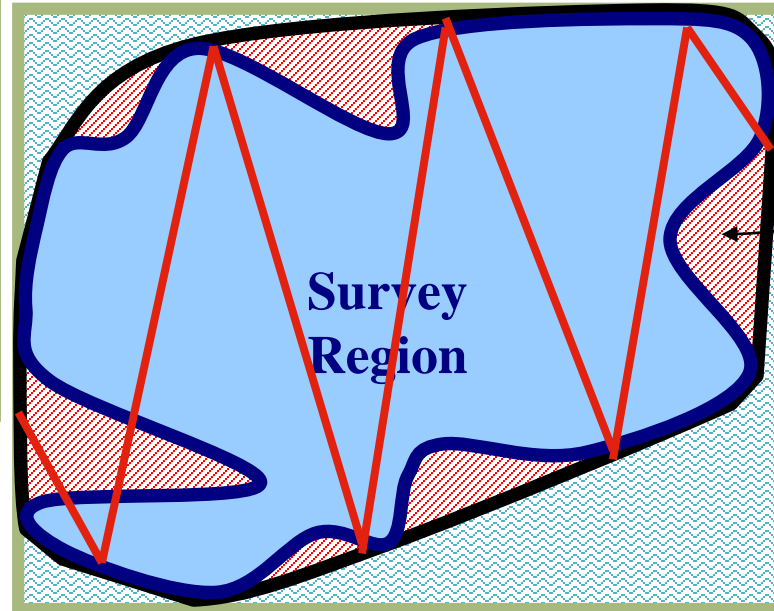
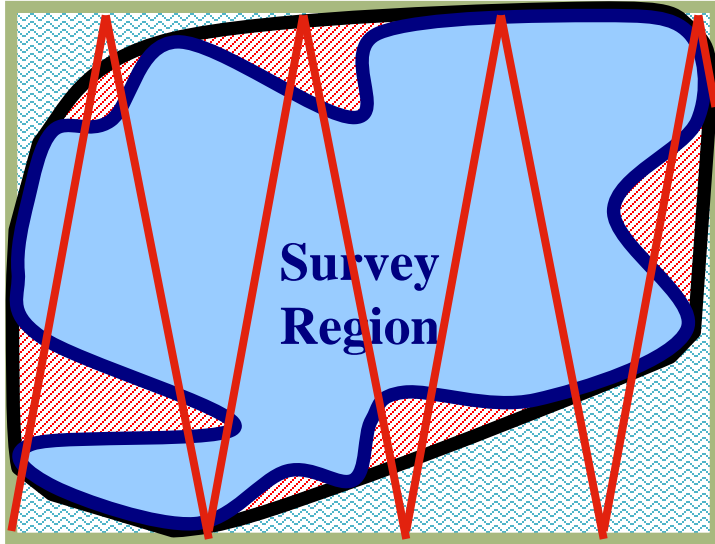
**Even-ness** of coverage within any given realisation

**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

DSsim needs:

shape and scale parameters on the natural scale  
and covariate parameters on the log scale

# Detectability

Golftees project

Detection Fct/Global/Parameter Estimates (MCDS)

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

Natural  
scale

Model

Half-normal key,  $k(y) = \text{Exp}(-y^2/(2*s^2))$

$s = A(1) * \text{Exp}(fcn(A(2)) + fcn(A(3)))$

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.

Log scale

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

$\text{exp}(0.268179) = 1.307581$

(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:

```
cov.param <- list()
cov.param$size <- 0.093
cov.param$sex <- data.frame(level = c(0,1),
                             param = c(-0.696, 0))
```

```
detect <- make.detectability(key.function = "hn",
                             scale.param = 2.62,
                             cov.param = cov.param,
                             truncation = 4)
```

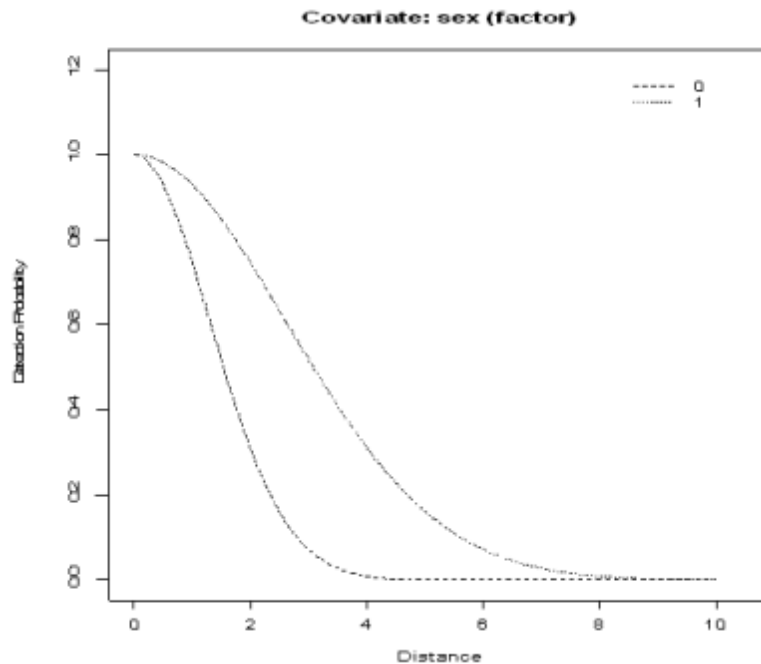
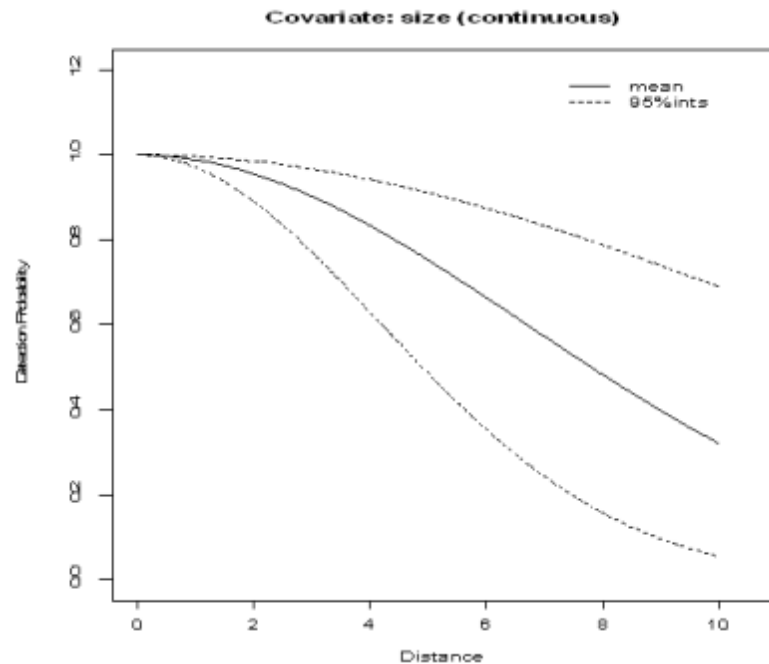
$\exp(\log(1.307581)+0.696) = 2.622633$

```
cov.param <- list()
cov.param$size <- 0.093
cov.param$sex <- data.frame(level = c(0,1),
                             param = c(0,0.696))
```

```
detect <- make.detectability(key.function = "hn",
                             scale.param = 1.31,
                             cov.param = cov.param,
                             truncation = 4)
```

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

# Detectability



# 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  $U \sim \text{Beta}(\theta, \theta)$ <sup>1</sup>

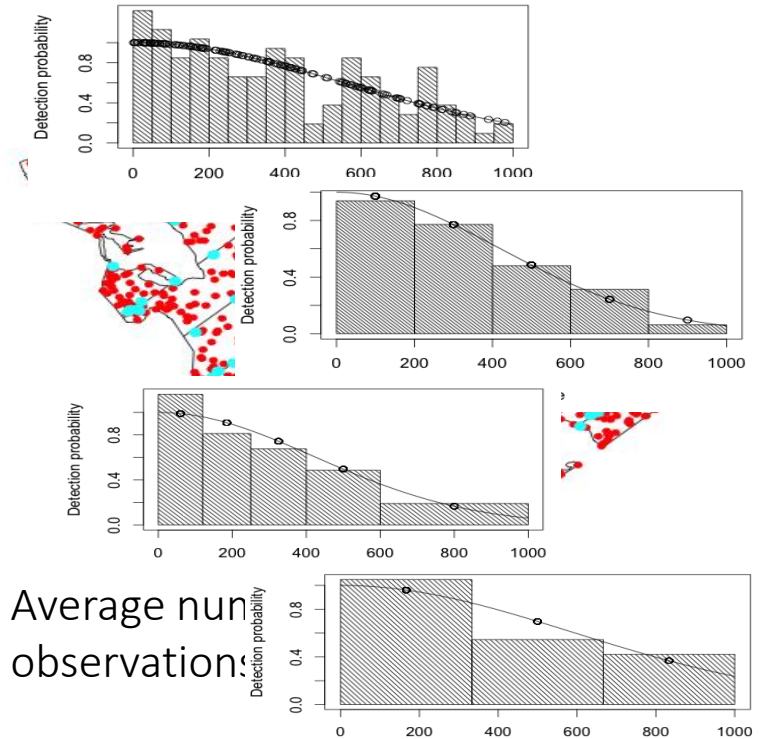
No error, ~15% CV ( $\theta = 5$ ), ~30% CV ( $\theta = 1$ )

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



Average number of observations:

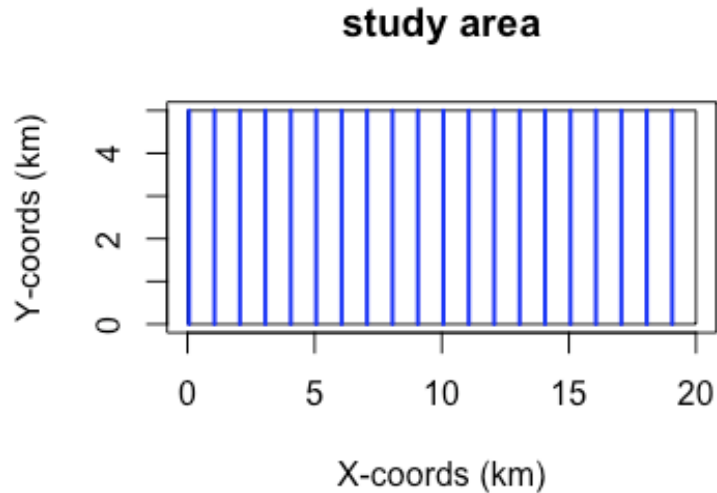
# To Bin or Not to Bin Results

	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

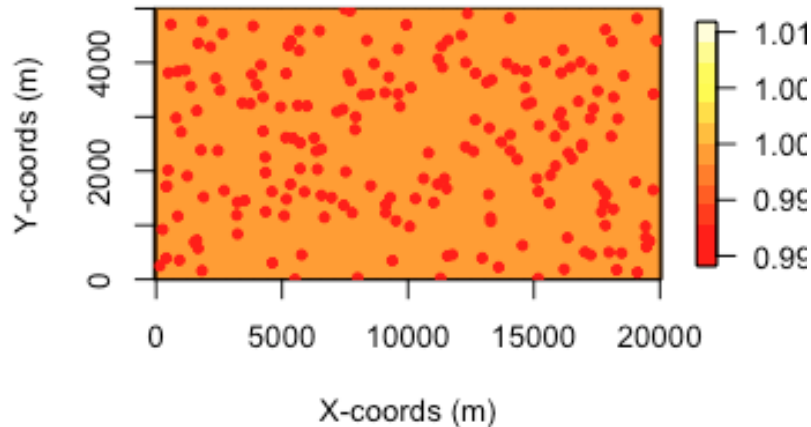


- Rectangular study region
- Systematic parallel transects with a spacing of 1000m

# Pooling Robustness and Truncation

DSsim vignette

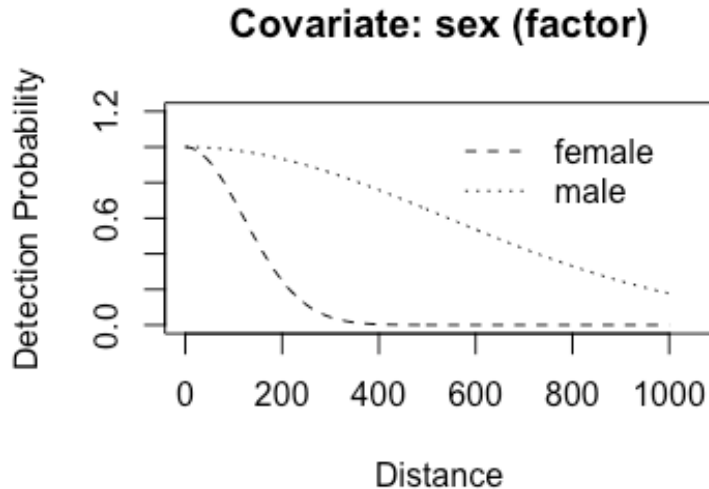
**Density Surface with Example Population**



- 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  $\sim 540$  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))

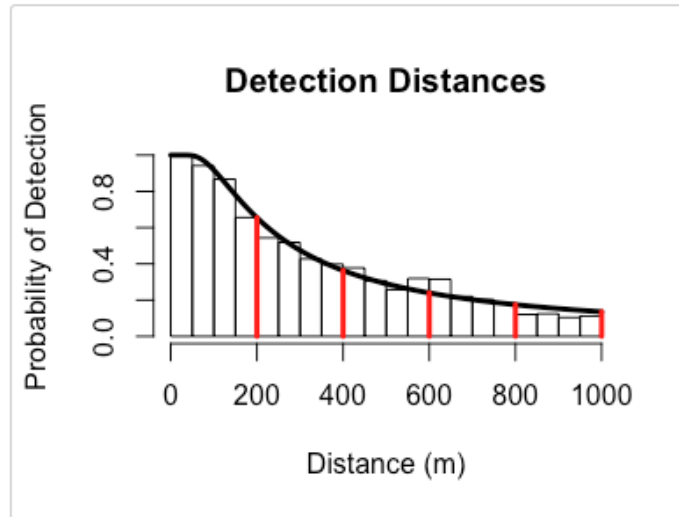
detect.cov <- make.detectability(key.function = "hn" ,
                                 scale.param = 120,
                                 cov.param = cov.params,
                                 truncation = 1000)
```

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

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

# Pooling Robustness and Truncation

DSsim vignette



- Two types of analyses:
  - $h_n \nu h_r$
  - $h_n \sim \text{sex}$
- Selection criteria: AIC

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

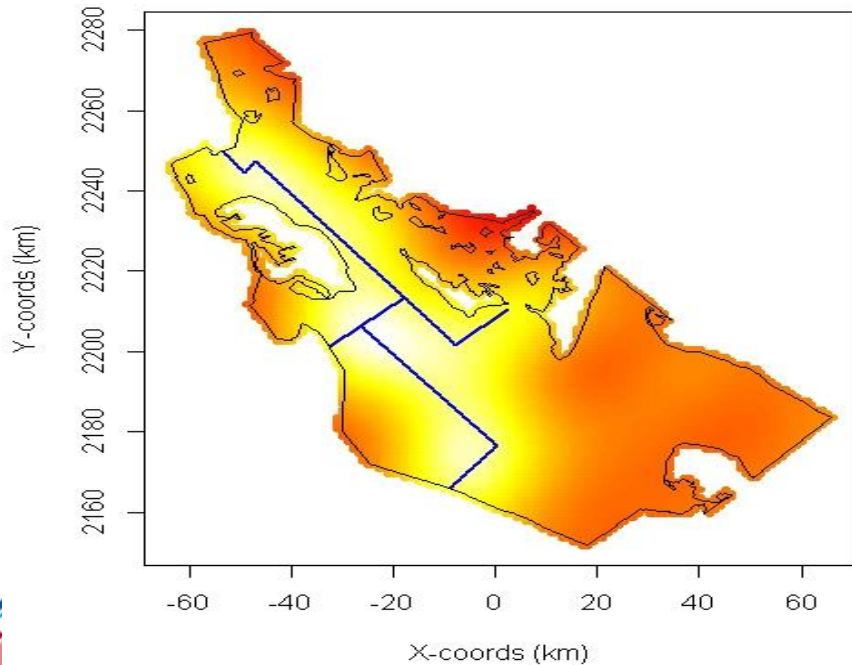
# Pooling Robustness and Truncation

Results HN v HR:

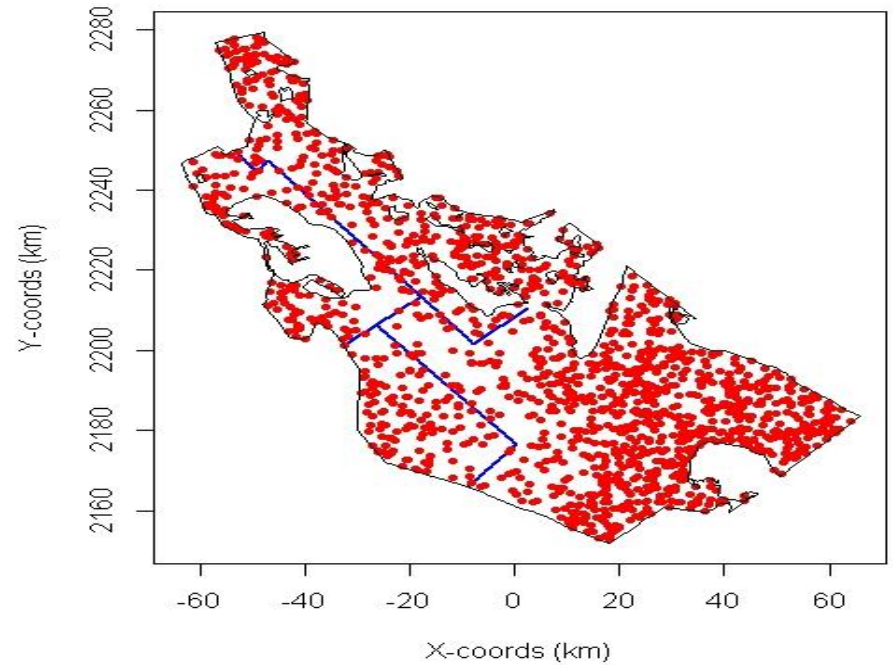
<i>Truncation</i>	<i>mean n</i>	<i>mean <math>\hat{N}</math></i>	<i>mean se</i>	<i>SD(<math>\hat{N}</math>)</i>	<i>%Bias</i>	<i>RMSE</i>	<i>% CI Coverage</i>
200	66	197	34.27	34.05	-1.32	34.13	97.5
400	102	190	31.06	34.79	-5.13	36.25	87.9
600	128	190	34.04	35.27	-5.24	36.77	81.9
800	144	190	34.31	36.61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

# Example Simulation

**Survey Region**

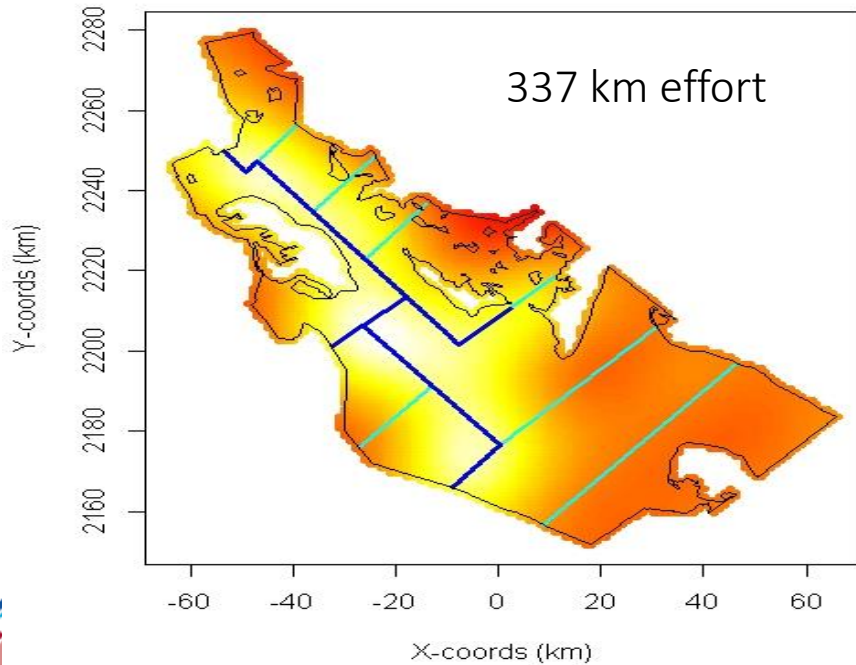


**Survey Region**

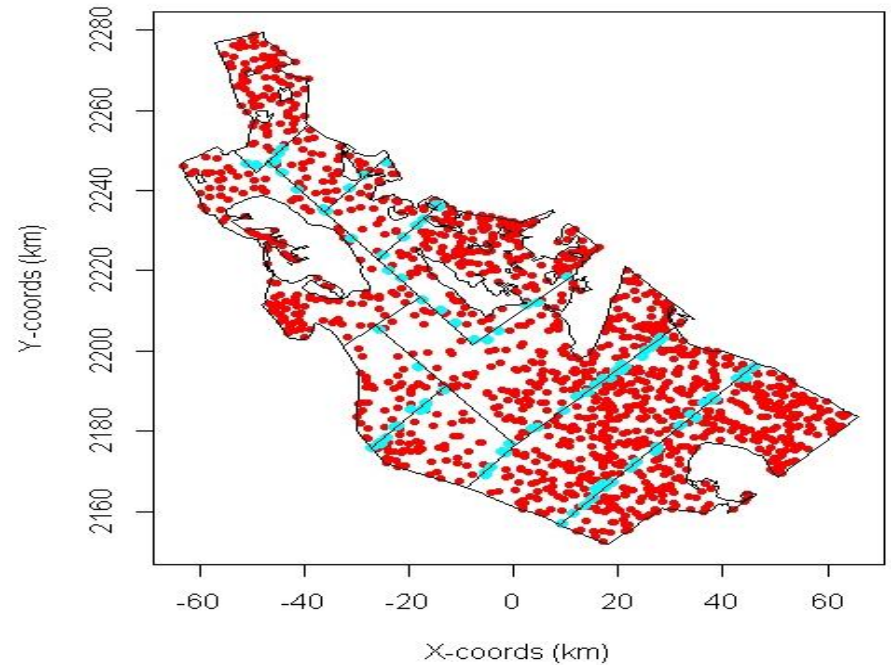


# Subjective survey design

**Survey Region**



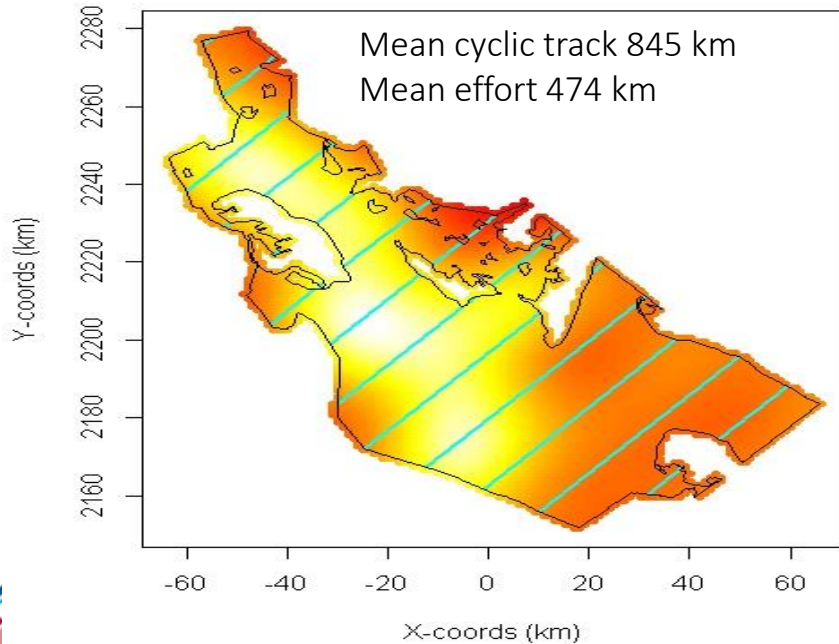
**Survey Region**



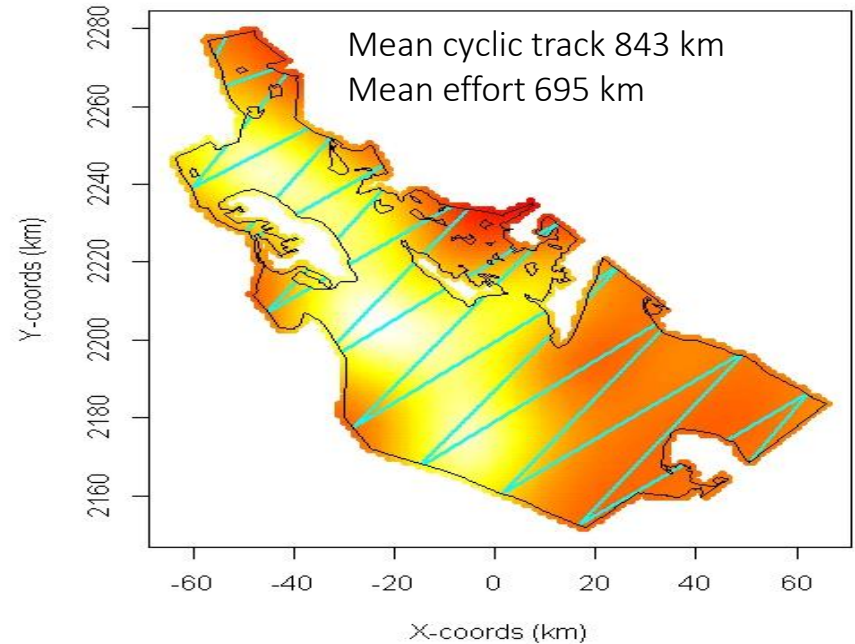


# Random Designs

**Survey Region**



**Survey Region**

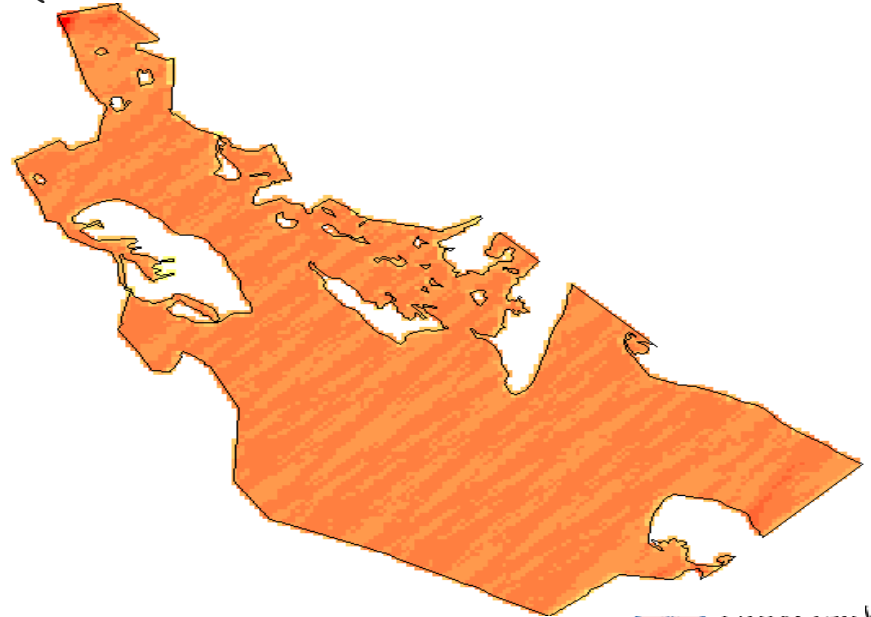


# Coverage probability

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*.