

# Distance Sampling Simulations

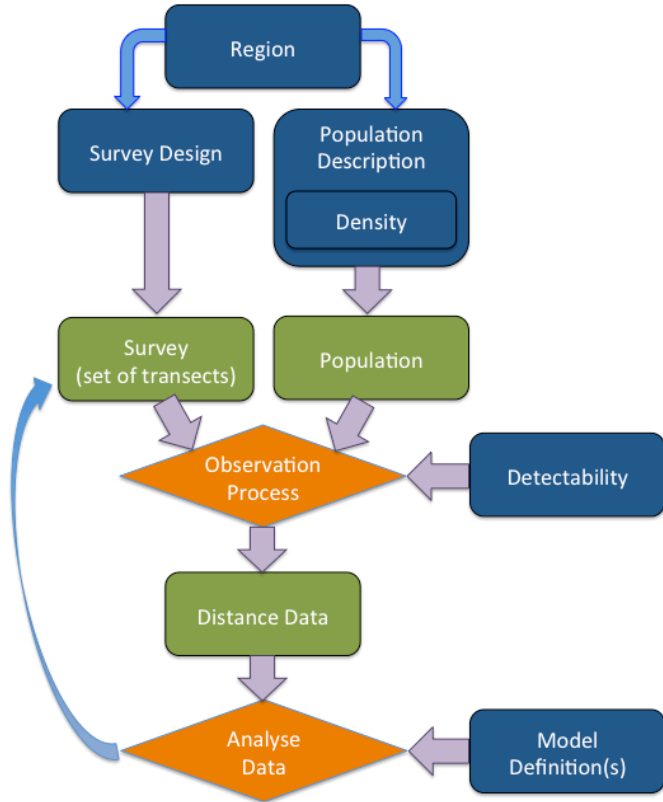
# Overview

- 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?

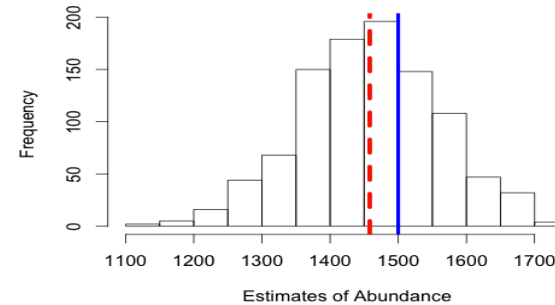
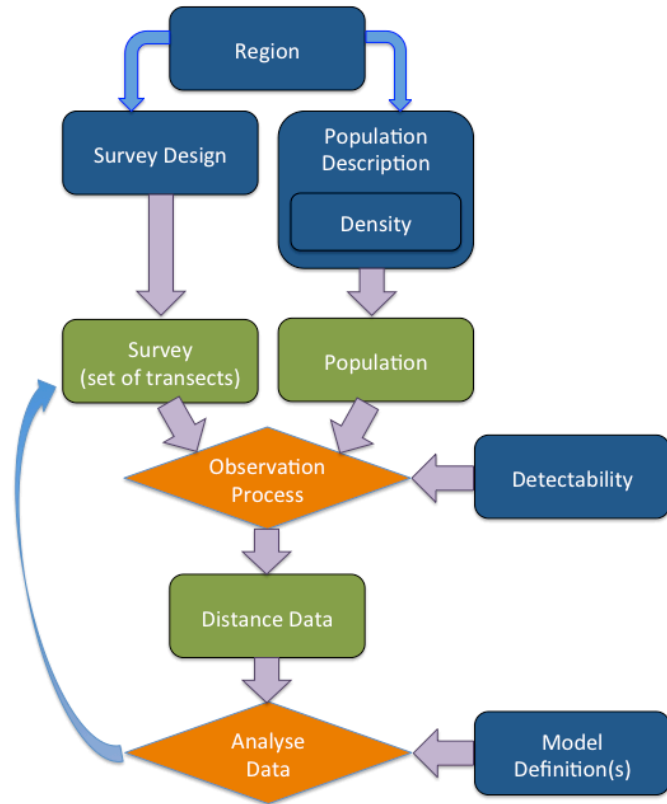
- Surveys expensive, simulations cheap!
- Test different survey designs
- Test survey protocols
- Investigate analysis properties
- Investigate violation of assumptions

# 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



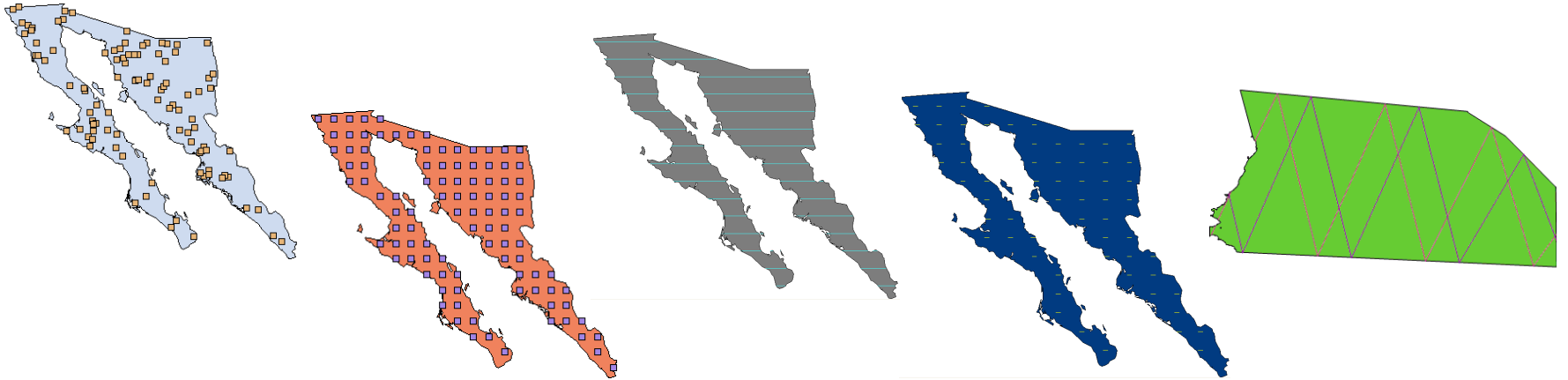
Assess:

- Bias
- Precision
- CI coverage

Across different designs/  
scenarios

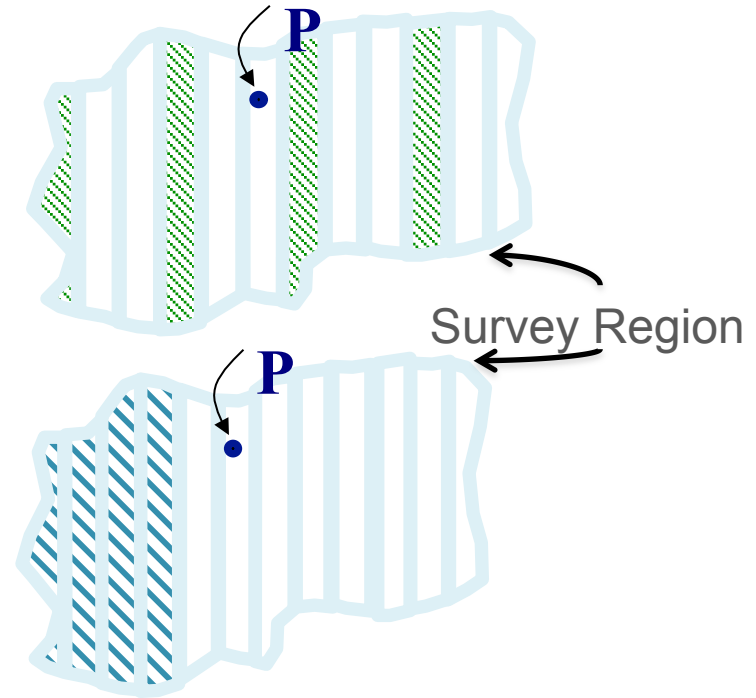
# Automated Survey Design

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

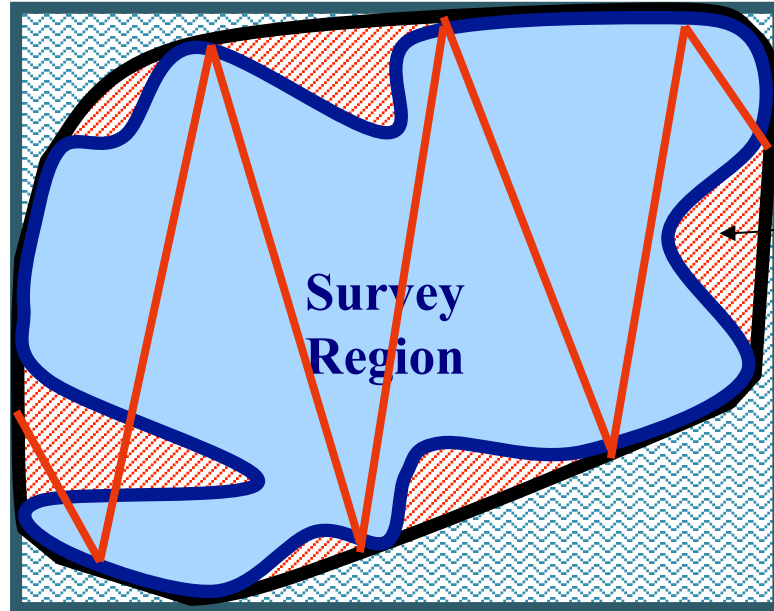
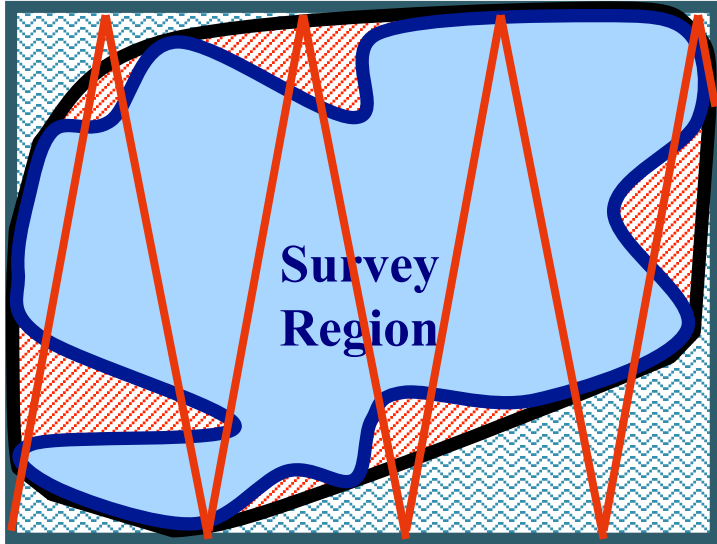


# Which Design?

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

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

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 Coef. 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')

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

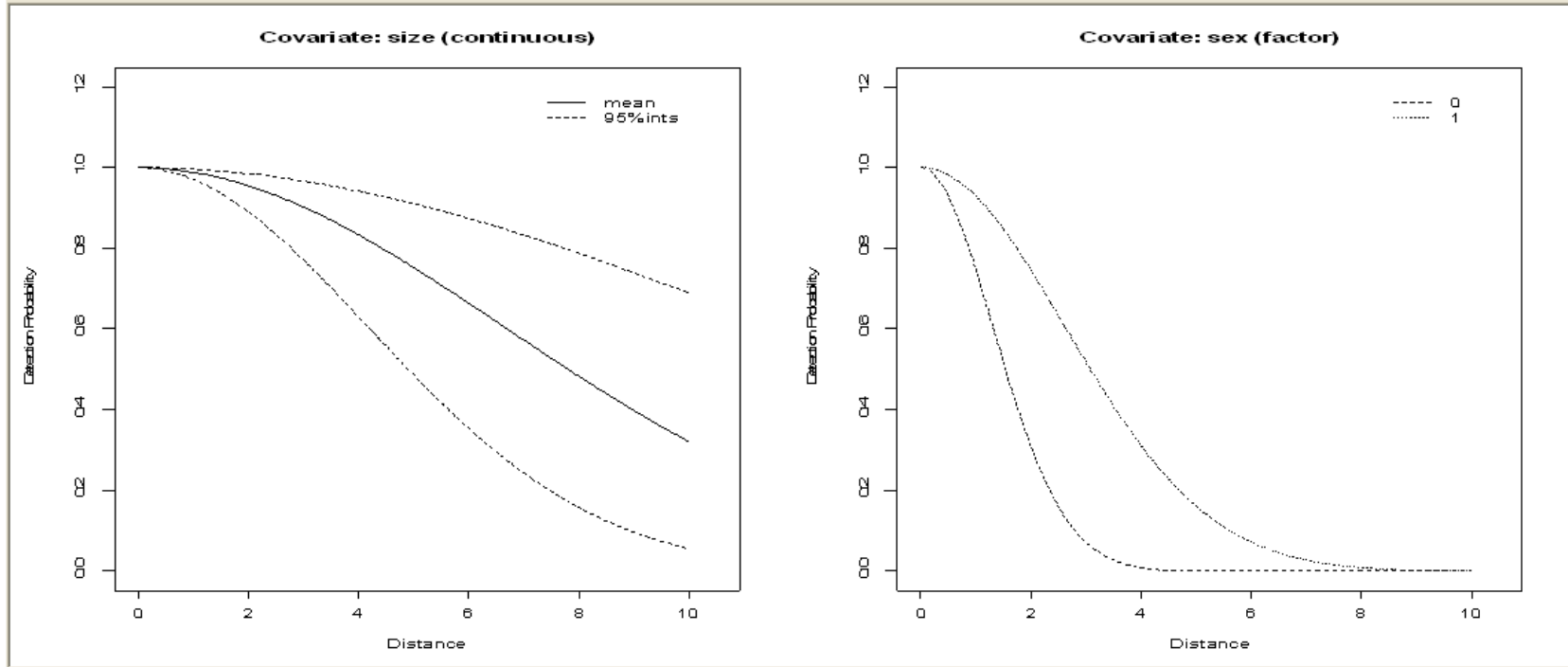
$$\exp(\log(2.622)-0.696) = 1.307265$$

# 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...



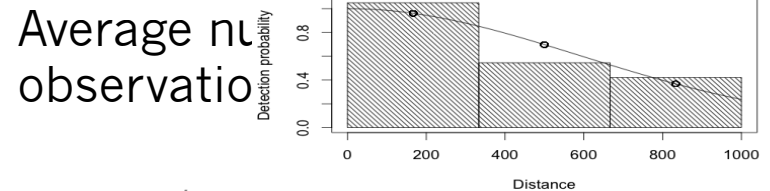
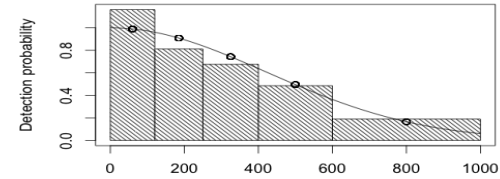
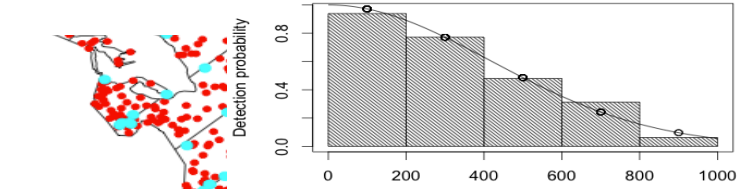
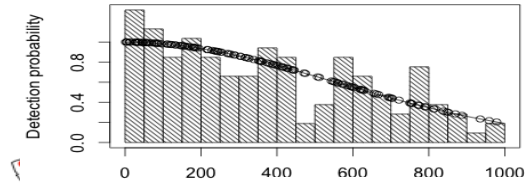
# Example Simulations

- To bin or not to bin?
- Testing pooling robustness in relation to truncation distance.
- Comparison of subjective and random 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



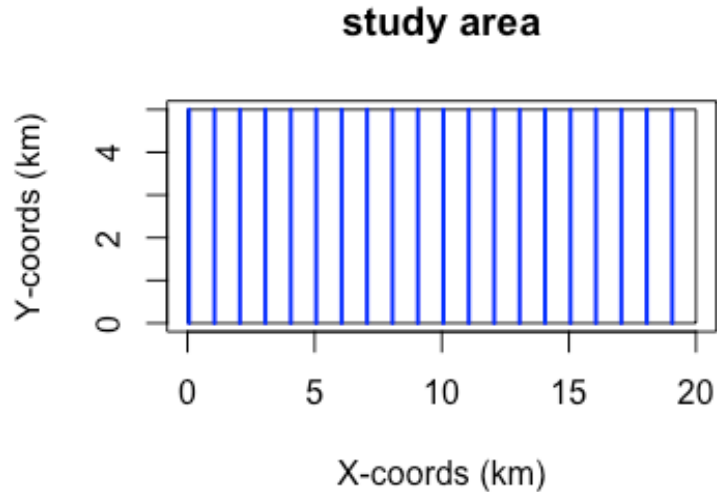
<sup>1</sup>Marques T. (2004) Predicting and correcting bias caused by measurement error in line transect sampling using multiplicative error models *Biometrics* 60:757--763

# 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

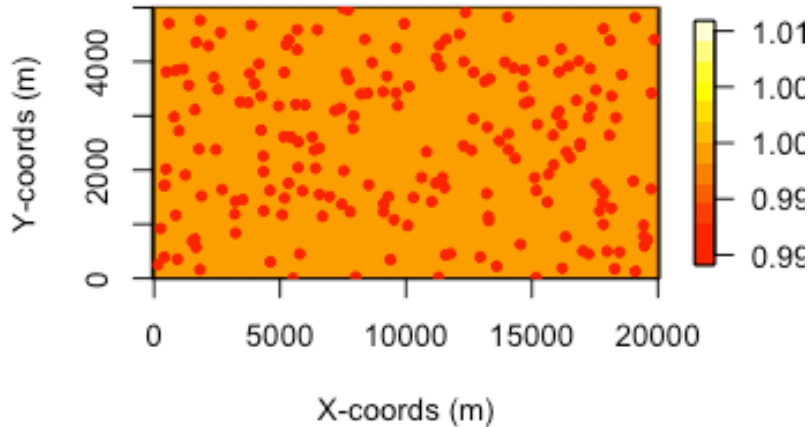


- Rectangular study region
- Systematic parallel transects with a spacing of 1000m

# Pooling Robustness and Truncation

- DSsim vignette

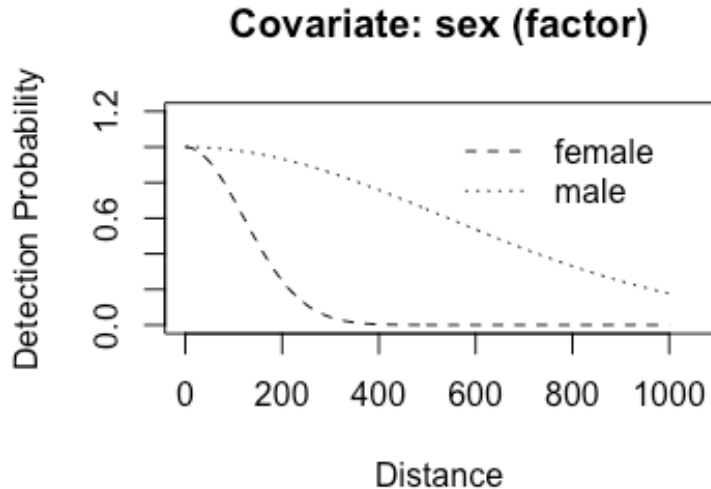
**Density Surface with Example Population**



- 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 ~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))

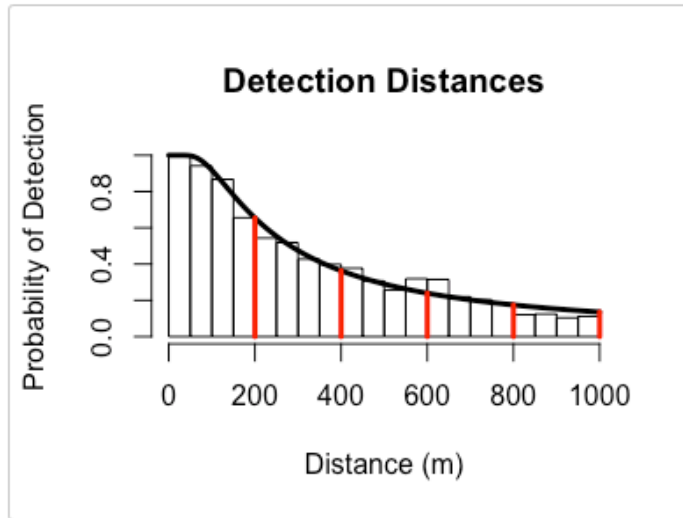
detect.cov <- make.detectability(key.function = "hn" ,
                                scale.param = 120,
                                cov.param = cov.params,
                                truncation = 1000)
```

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

- 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 ~ sex**
- Selection criteria: AIC

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



# Pooling Robustness and Truncation

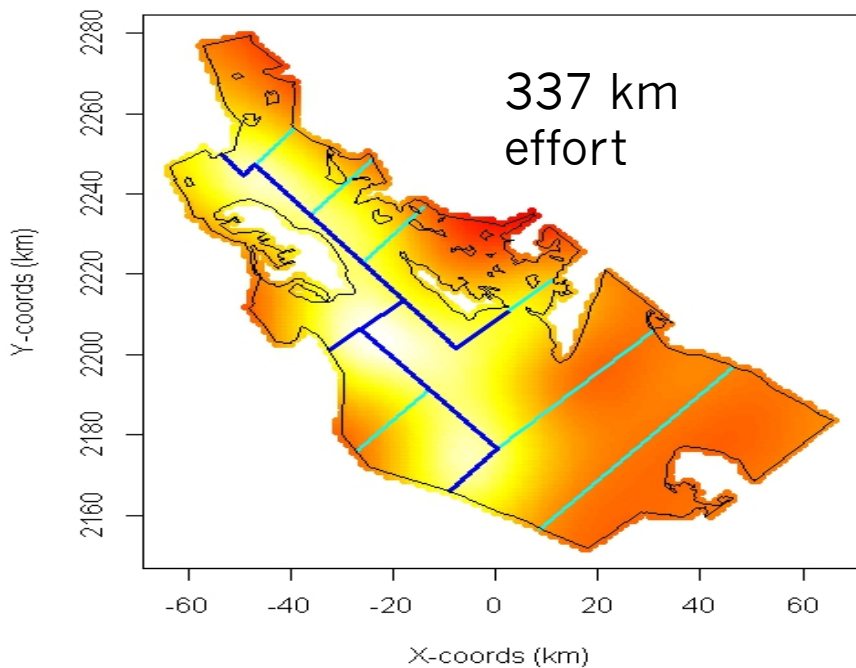
- Results HN v HR:

<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

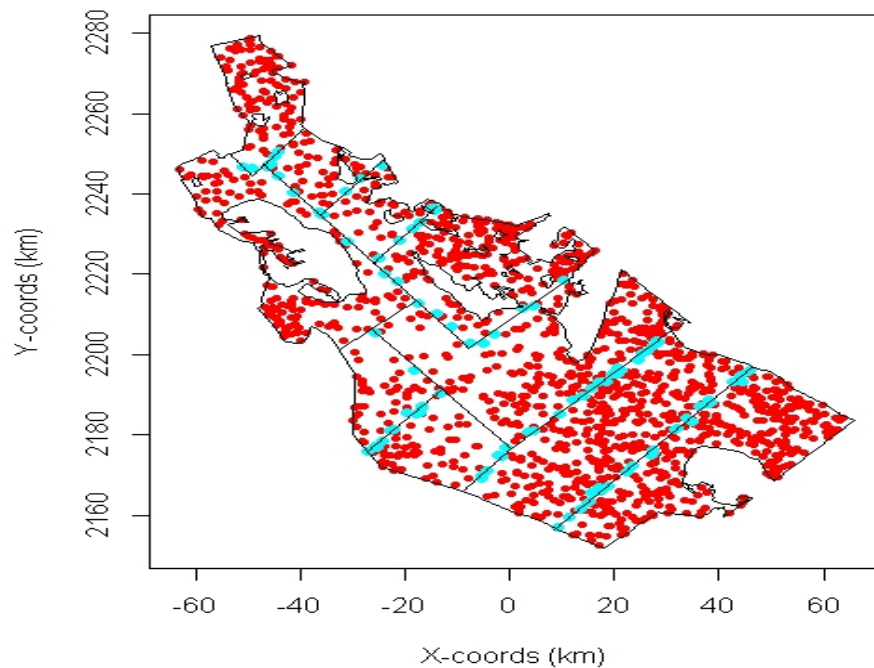


# Subjective survey design

**Survey Region**

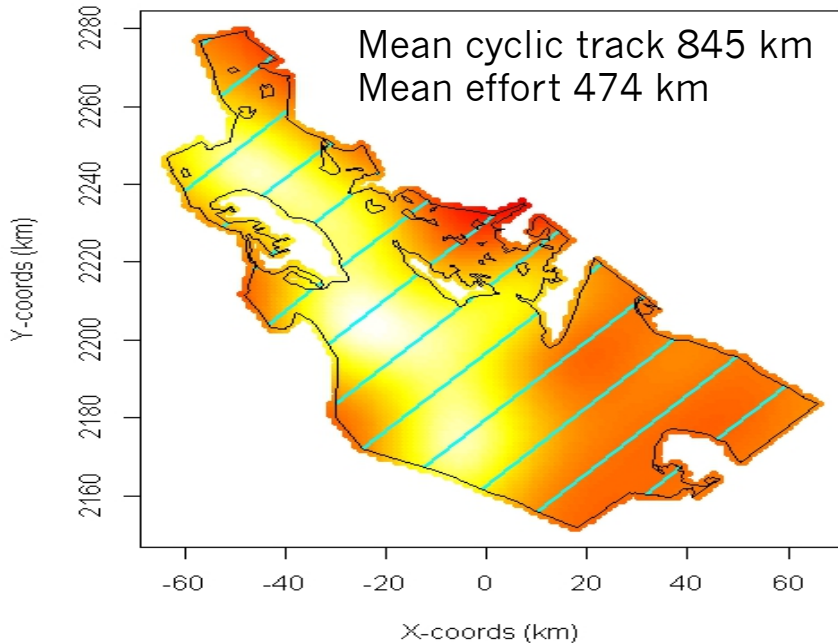


**Survey Region**

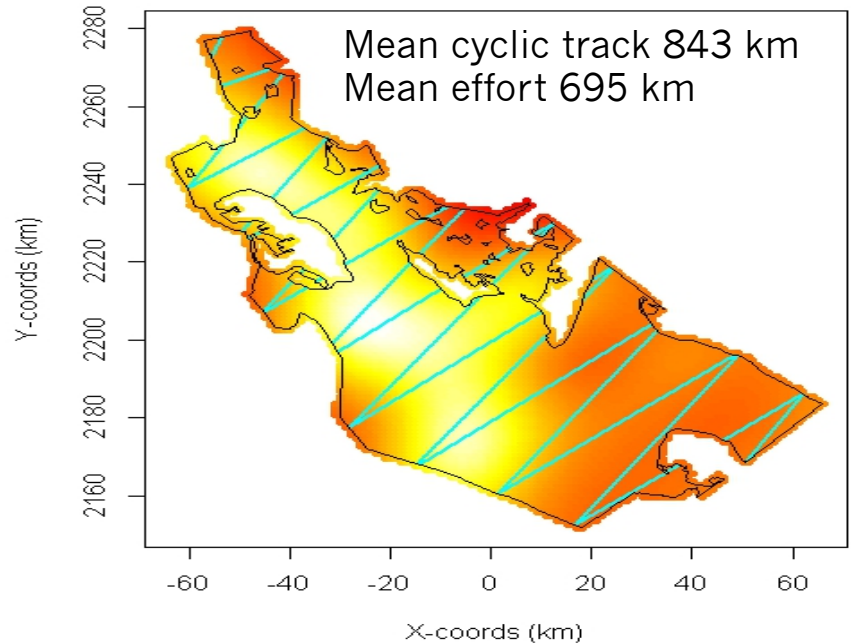


# Random Designs

**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*.