Choosing a Detection function





Overview of detection function modelling

Formal definition

Criteria for a good detection function model

Key functions and adjustment terms

Choosing the number of parameters

Comments about truncation





Formal definition

The detection function describes the relationship between distance and the probability of detection

Formally denoted by g(x) (usually referred to as 'g of x')

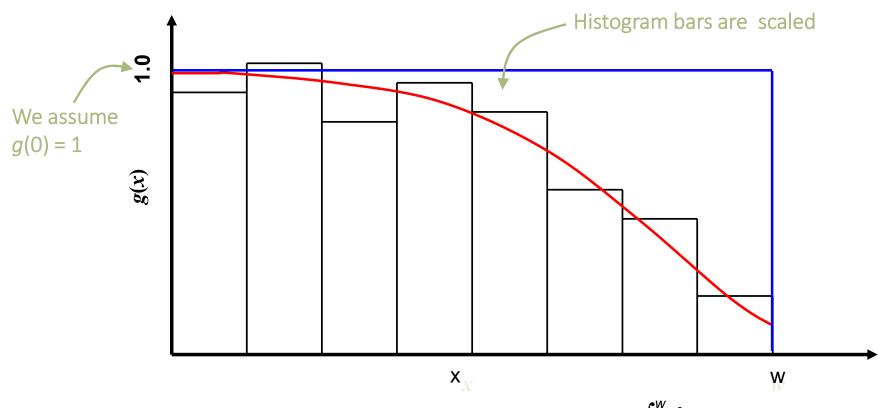
g(x) = the probability of detecting an animal, given that it is at distance x from the line

Key to the concept of distance sampling





The detection function, g(x)









Modelling g(x)

g(x) represents the underlying relationship between detection probability and distance

However, the true form of g(x) is unknown to us

We need to estimate g(x) by fitting a model to our data

i.e., we need to find a curve that will approximate the underlying relationship





Criteria for robust estimation

Four main criteria for a good model:

- 1. Model robustness use a model that will fit a wide variety of plausible shapes for g(x)
- 2. Shape criterion use a model with a 'shoulder' i.e. g'(0)=0
- 3. Pooling robustness use a model for the average detection function, even when many factors affect detectability
- 4. Estimator efficiency use a model that will lead to a precise estimator of density





Key functions





Key functions

The first step in constructing a model for g(x) is to choose a key function

This determines the basic model shape

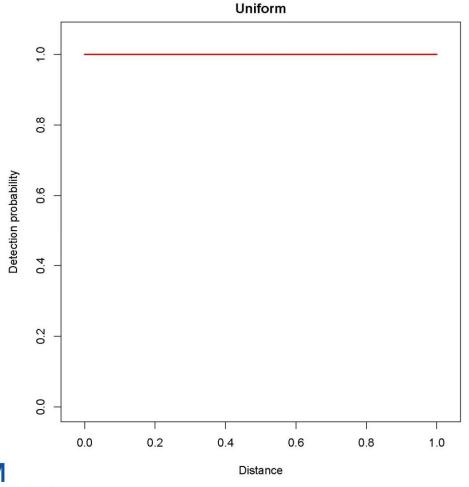
Three key functions available in distance sampling software:

- Uniform
- Half normal
- Hazard rate





Key functions (cont.)



• Model formula:

$$g(x) = 1, x \leq w$$

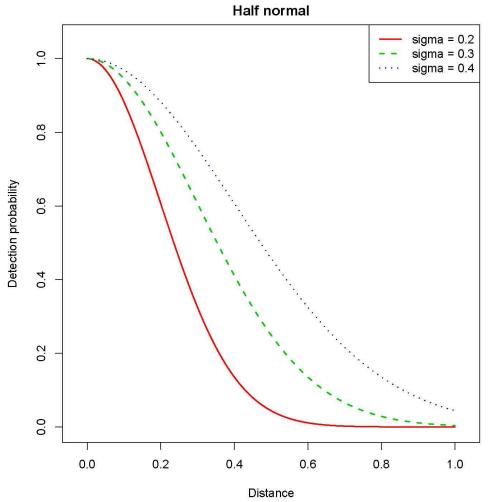
- Parameters = 0
- Shape criterion?Yes
- Model robust?

No





Key functions (cont.)



• Model formula:

$$g(x) = \exp\left(\frac{-x^2}{2\sigma^2}\right), x \le w$$

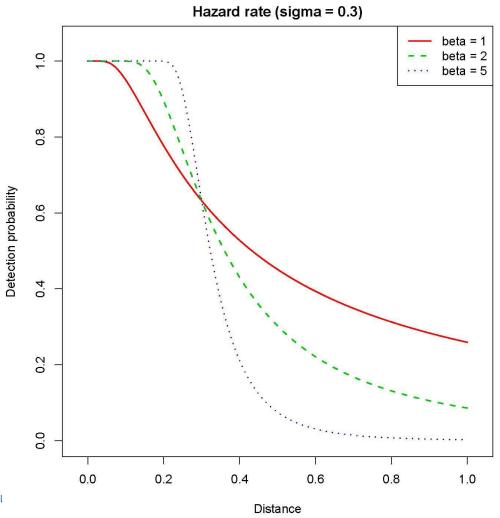
- Parameters = 1
- Shape criterion?Yes
- Model robust?

Somewhat





Key functions (cont.)



Model formula:

$$g(x) = 1 - \exp \left[-\left(\frac{x}{\sigma}\right)^{-\beta} \right], x \le w$$

- Parameters = 2
- Shape criterion?Yes
- Model robust?

Yes





Adjustment terms





Adjustment terms

Models can be made more robust by adding a series of adjustment terms (also called series expansion or series adjustment) to the key function

Key function \times (1 + Series)

Series = $\alpha_1 \times \text{term}_1 + \alpha_2 \times \text{term}_2 + \dots$ etc.

The α_i parameters must be estimated

Resulting curve model is scaled so that g(0)=1

The number of adjustment terms needs to be chosen





Adjustment terms

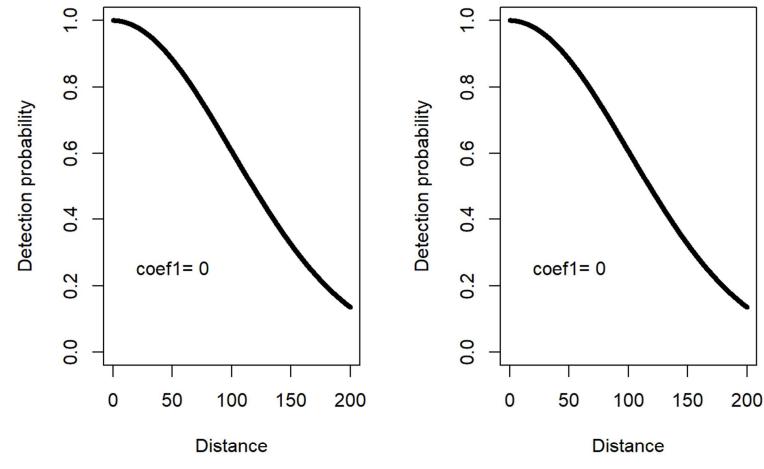
Distance allows the selection of three types of series (one type per model)

Key function	Series adjustment	
Uniform*	Cosine*	
Half normal [†]	Hermite polynomial [†]	
Hazard rate	Simple polynomial	





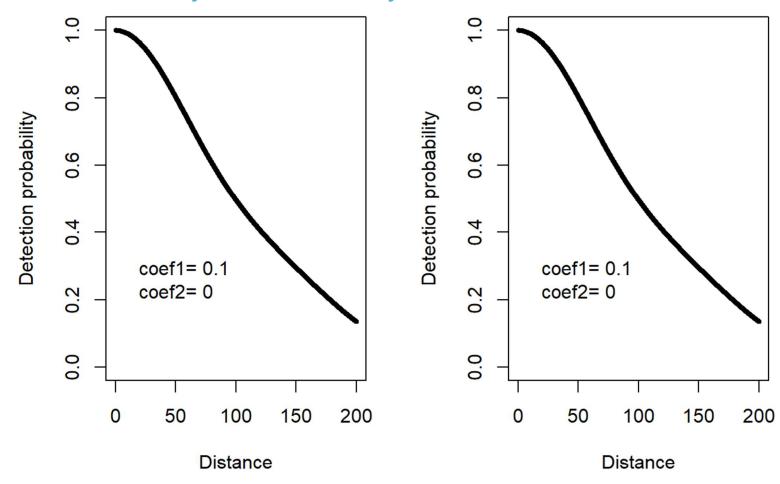
Half normal key, single cosine adjustment term







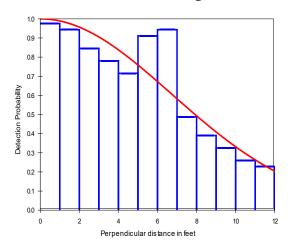
Half normal key, two cosine adjustment terms

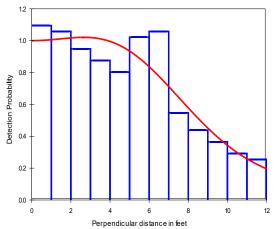


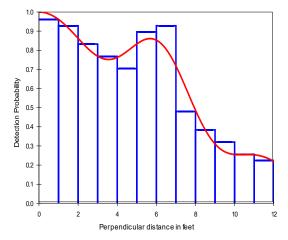




Adjustment terms – how many?







Half normal	Half normal	Half normal
0 adjustment terms	1 adjustment term	5 adjustment terms
1 parameter	2 parameters	6 parameters
$\hat{P}_a = 0.65$	$\hat{P}_a = 0.72$	$\hat{P}_a = 0.63$
$CV(\hat{P}_a) = 5.8\%$	$CV(\hat{P}_a) = 11.6\%$	$CV(\hat{P}_a) = 19.9\%$



Note: There is a monotonicity constraint in Distance that is switched on by default to prevent detection functions from increasing. The constraint had to be turned off to produce the third plot. The third plot is for demonstration only – it would not be a good detection function to choose (unless there was a biological reason why detection probability would increase at those distances).



Bias vs variance tradeoff





How many parameters?

- Models with too few parameters will not be flexible enough to describe the underlying relationship
- Adding parameters will improve the fit
- But models with too many parameters will be too flexible and will also describe the random noise in the data
- We generally seek models with an intermediate number of parameters



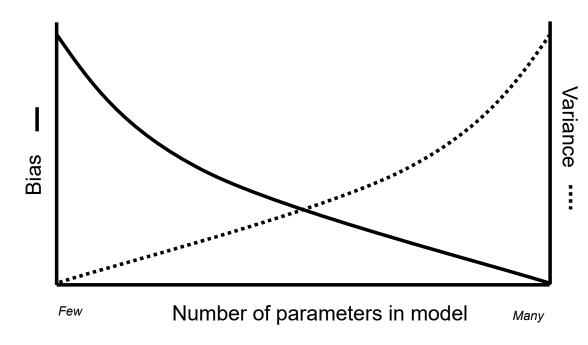


How many parameters?

This problem can also be expressed as a trade-off between bias and variance

Models with too few parameters tend to produce estimates with low variance and high bias

Models with too many parameters tend to produce estimates with low bias and high variance (note the increasing CV for the estimate of P_a on the earlier slide)







Data truncation prior to model fitting





Truncation

$$\widehat{N} = \frac{nA}{2wL\widehat{P}_a}$$

Need to choose the value of w (right truncation)

Detections at large distances contribute little to estimating the shape of g(x) at small distances (i.e. the shoulder) and may lead to poor fit and high variance

Typically, we might truncate around 5% of observation for line transects (perhaps nearer 10% for point transects)

Can also use estimated values of g(x) from fitted model as truncation criterion; truncate at w when g(w)=0.15





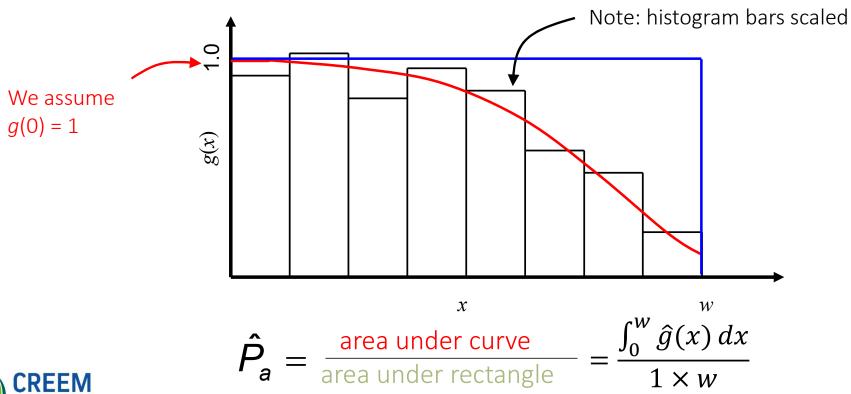
Alternative derivations for understanding detectability





1. The detection function, g(x)

g(x) = probability of detecting an animal, given that it is at distance x from the line



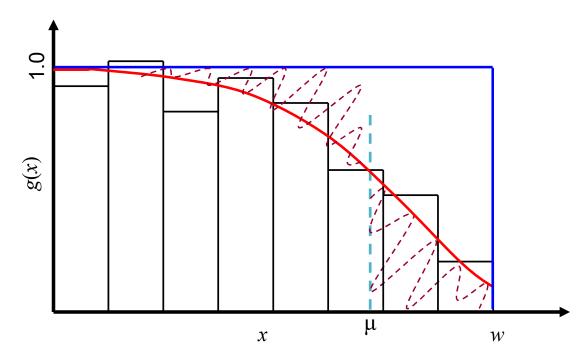




2. Effective strip (half) width, μ

 Instead of a <u>line transect</u> out to w, where proportion P_a objects are seen, think of a <u>strip transect</u> out to some distance μ.

area under curve area under rectangle



The ESW, μ , is the distance at which as many objects are detected beyond μ as are missed within μ

Line transect out to w

$$\hat{N} = \frac{nA}{2wL\hat{P}_a}$$

Area covered

$$=\frac{\int_0^w \hat{g}(x)dx}{w} = \frac{\hat{\mu}}{w}$$

Strip transect out to μ

$$\hat{N} = \frac{nA}{2\hat{\mu}L}$$

Area <u>effectively</u> covered



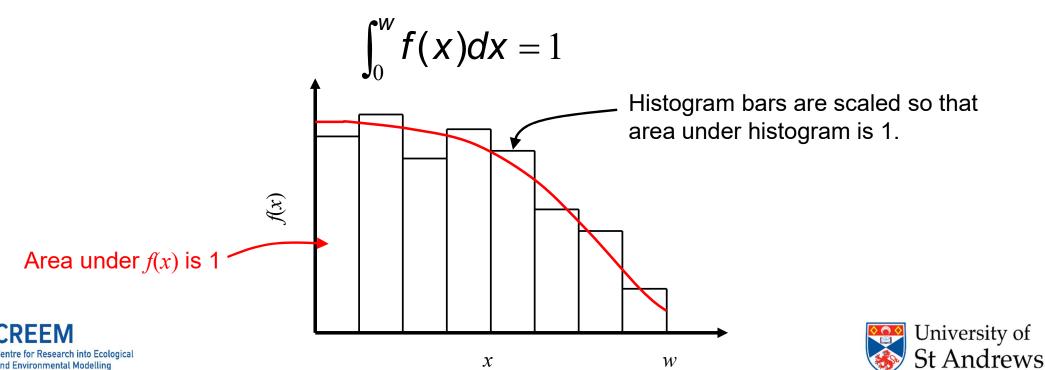


3. The probability density function, f(x)

f(x)dx = probability of observing an animal between distance x and x+dx, given it was observed somewhere in (0,w)

f(x) is called the probability density function (pdf) of the observed distances

Because observations are between 0 and w, the area under f(x) is 1.0



Why is f(x) useful?

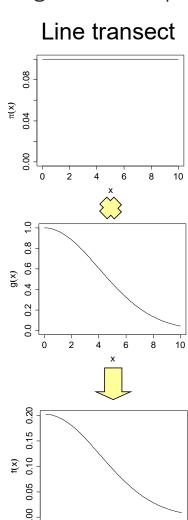
1. Useful for point transects, as it gives the expected distribution of detection distances

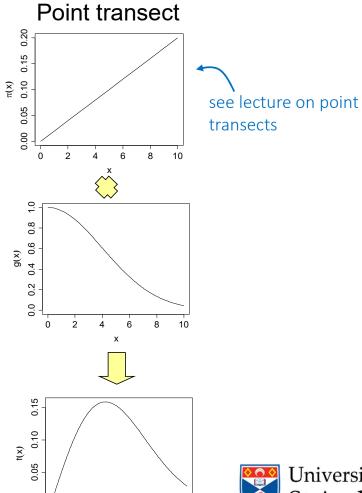
True distribution of animals

Detection function, g(x)

Observed distribution, f(x)



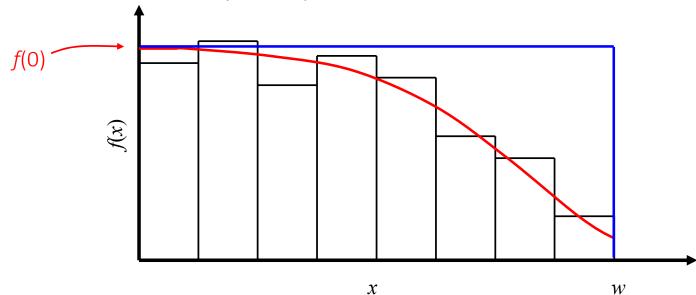






Why is f(x) useful?

2. Gives another way to estimate P_a Lots of statistical machinery to fit pdfs

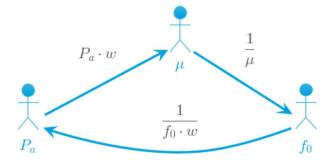


$$\hat{P}_a = \frac{\text{area under curve}}{\text{area under rectangle}} = \frac{1}{\hat{f}(0)W}$$
 $\hat{N} = \frac{nA}{2wL\hat{P}_a} = \frac{nA}{2wL\left(\frac{1}{\hat{f}(0)W}\right)} = \frac{nA\hat{f}(0)}{2L}$





Formulae – line transects



Three ways to think about line transects

1. Proportion seen or average probability of detection in covered region, P_{α}

$$\hat{N} = \frac{nA}{2wL\hat{P}_a} \qquad \qquad \hat{D} = \frac{n}{2wL\hat{P}_a}$$

$$\hat{D} = \frac{n}{2wL\hat{P}_a}$$

2. Effective strip (half-)width, ESW, μ.

$$P_{a} = \frac{\mu}{W}$$

$$\hat{N} = \frac{nA}{2\hat{\mu}L}$$

$$\hat{D} = \frac{n}{2\hat{\mu}L}$$

3. Pdf of observed distances, f(x), evaluated at 0 distance

$$f(0) = \frac{1}{\mu}$$

$$\hat{N} = \frac{n\hat{f}(0)A}{2L}$$

$$\hat{D} = \frac{nf(0)}{2L}$$





Notation (knowns and unknowns)





Notation – line transects

Known constants and data:

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k = \text{number of lines}
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$$I_j$$
 = length of j^{th} line, $j=1,...,k$

$$L = \Sigma I_i = \text{total line length}$$

n = number of animals or clusters detected

 x_i = distance of i^{th} detected animal or cluster from the line, i=1,...,n

w = truncation distance for x

A =size of region of interest

a = area of "covered" region = 2wL

 s_i = size of ith detected cluster, i=1,...,n





Notation – line transects

Parameters and functions:

N = population size / abundance of animals

 N_s = abundance of clusters

D = density = animals per unit area = N/A

 D_s = density of clusters

g(x) = detection function

f(x) = probability density function (pdf) of observed distances

f(0) = f(x) evaluated at 0 distance

 μ = effective strip (half-)width

 P_a = probability of detecting an animal or cluster given it is in the covered area a

E(s) = mean size of clusters in the population



