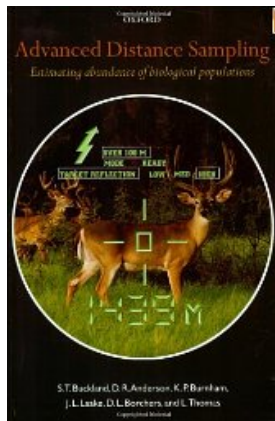
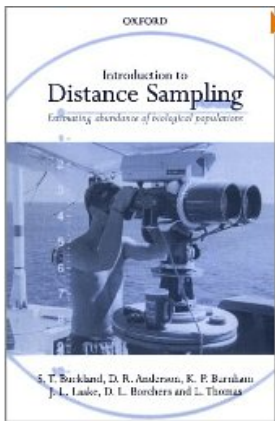


# Distance sampling

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Abundance estimation is often accomplished using this equation:

$$\hat{N} = \frac{n}{\hat{p}}$$

- $N$  is abundance (population size)
- $n$  is the number of individuals detected
- $\hat{p}$  is an estimate of detection probability: The probability of detecting an individual

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Most methods differ in how they estimate  $p$

## Challenges

- Detection probability is rarely constant

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- Detection probability is rarely constant
- It is a function of:
  - ▶ Age
  - ▶ Sex
  - ▶ Habitat
  - ▶ Distance

## Basic idea

- $p_i$  is the probability of detecting individual  $i$  at distance  $x_i$ .

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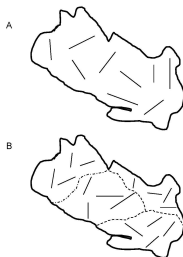
## Advantages of distance sampling

- Population size can be estimated from a single survey
- Explicit link between population size and density

- Randomly place line transects or points throughout the study area
- Record the distance to each animal detected

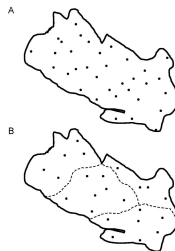
## Line transects

Figure 9.5



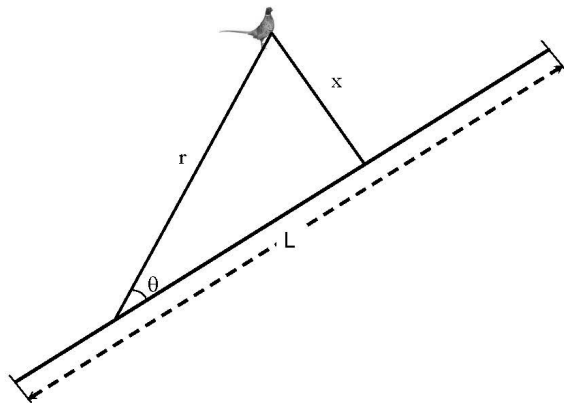
## Point transects

Figure 9.7



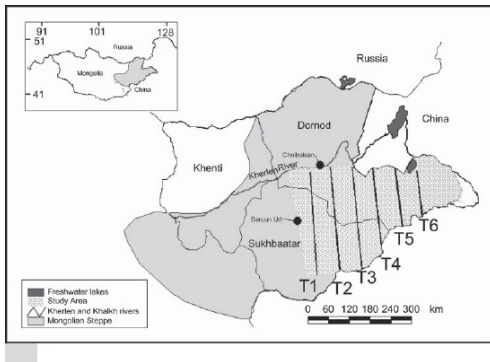
# LINE TRANSECTS

Figure 9.1



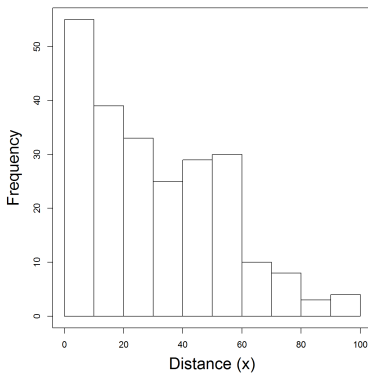
In line transect sampling, it is common to record the radial distance and bearing, rather than the perpendicular distance. However, the analysis must be conducted on the perpendicular distance data.

## Mongolian gazelle *Procapra gutturosa*



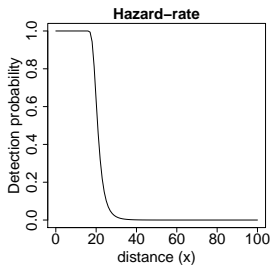
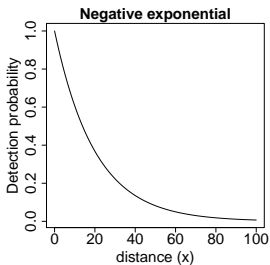
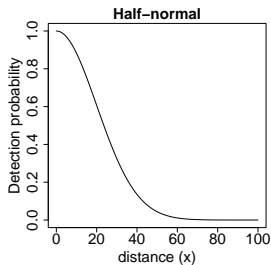
# LINE TRANSECTS

Data	
Animal	Distance ( $x$ )
1	4.4
2	25.3
3	41.8
4	3.1
5	78.5
$\vdots$	
$n$	4.4



# ESTIMATING $\bar{p}$ , AVERAGE DETECTION PROBABILITY

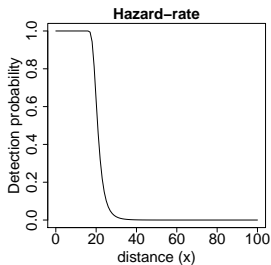
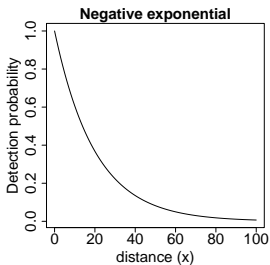
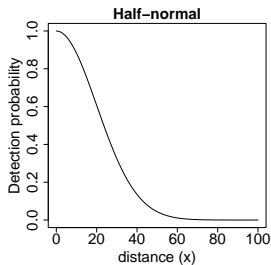
- Fit a detection function,  $g(x)$ , to the data
  - ▶ Assume  $g(0) = 1$





# ESTIMATING $\bar{p}$ , AVERAGE DETECTION PROBABILITY

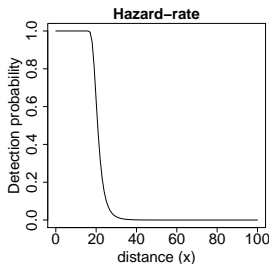
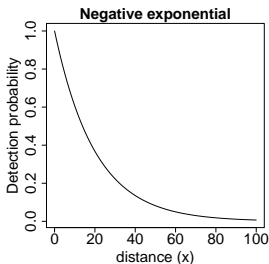
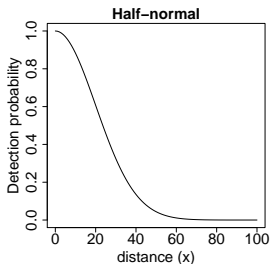
- Fit a detection function,  $g(x)$ , to the data
  - ▶ Assume  $g(0) = 1$
- Assume individuals are “uniformly” distributed with respect to the transect (valid under random sampling):  $p(x) = 1/B$  where  $B$  is the transect half-width.



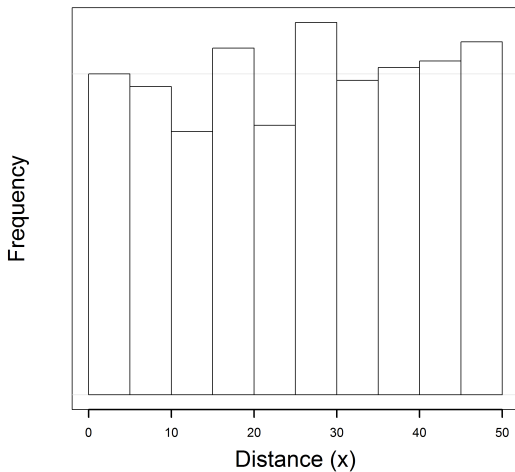
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- Assume individuals are “uniformly” distributed with respect to the transect (valid under random sampling):  $p(x) = 1/B$  where  $B$  is the transect half-width.
- $\bar{p}$  is then the average detection function, weighted by  $p(x)$ .

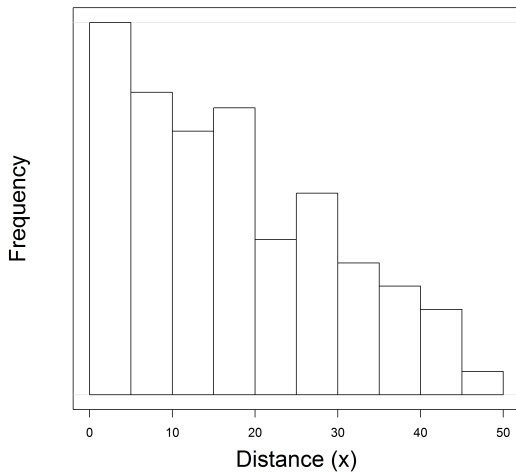
$$\bar{p} = \int_0^B g(x)p(x)dx$$



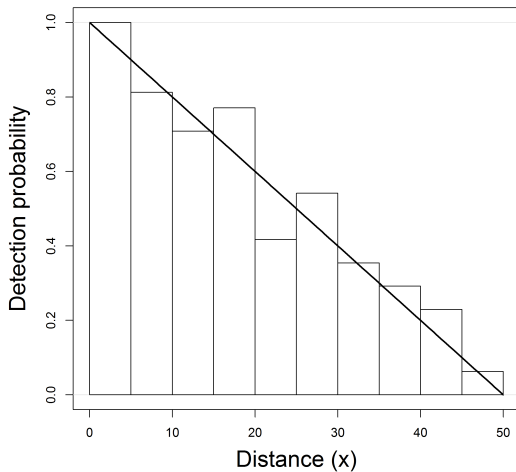
# COMPUTING $\bar{p}$ , AVERAGE DETECTION PROBABILITY



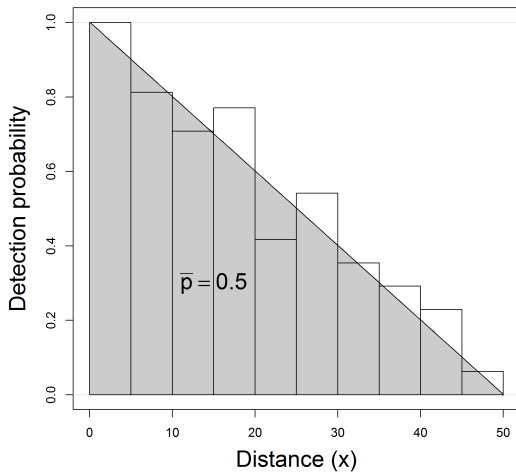
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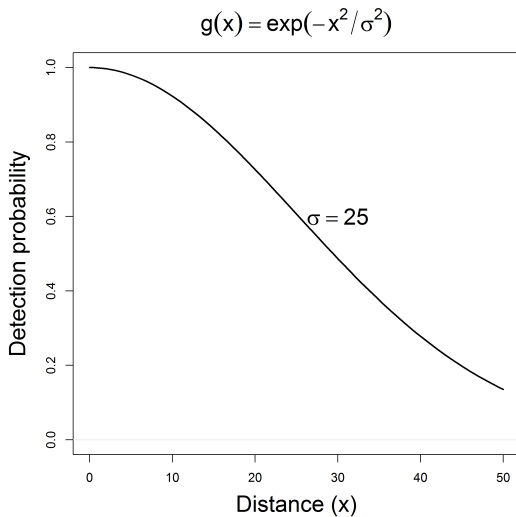
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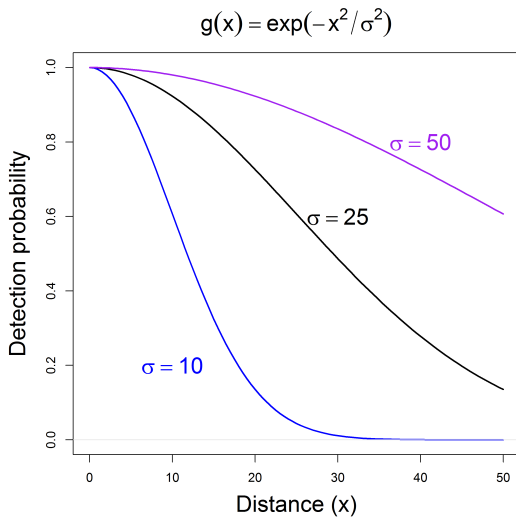
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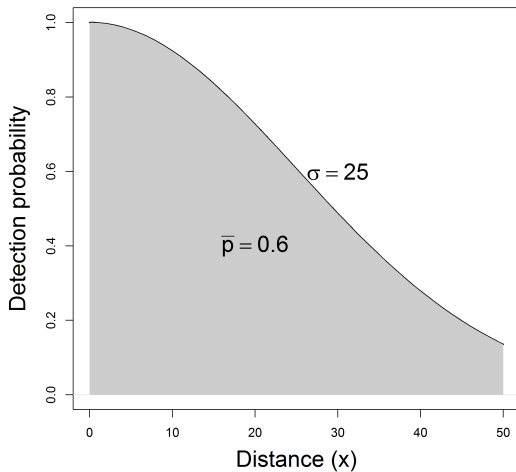


# COMPUTING $\bar{p}$ , AVERAGE DETECTION PROBABILITY

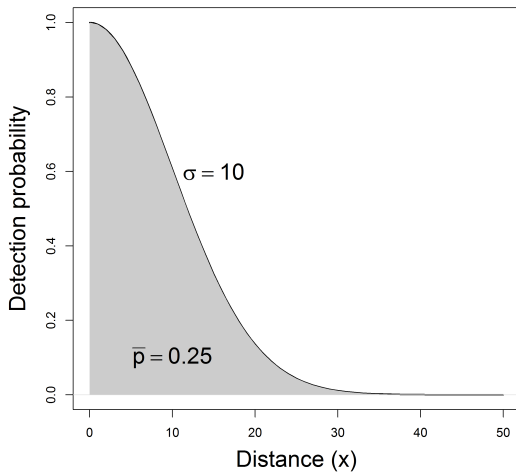




# COMPUTING $\bar{p}$ , AVERAGE DETECTION PROBABILITY



# COMPUTING $\bar{p}$ , AVERAGE DETECTION PROBABILITY



<https://richard-chandler.shinyapps.io/distance-sampling/>

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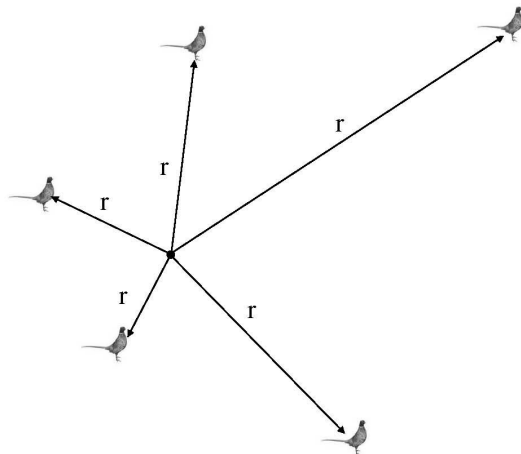
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- (3) Distance is measured accurately
- (4) Transect lines are placed randomly with respect to the animals
  - ▶ This ensures that individuals will be uniformly distributed with respect to the transect
- (5) Detections (of individuals or groups) are independent of one another

# POINT TRANSECTS

Figure 9.6



We still use the same type of detection function (half-normal, hazard, etc. . . ).

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We don't expect a “flat” histogram of distances when  $\bar{p} = 1$ .  
Why?

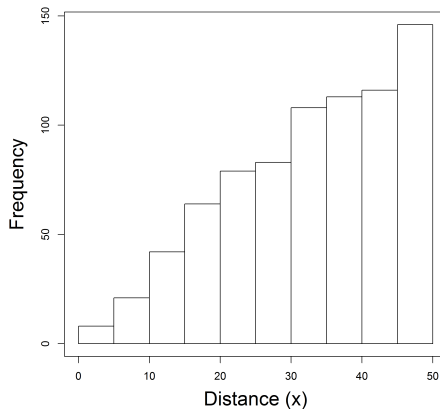
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More individuals occur far from the observer than close to the observer because area increases with distance.

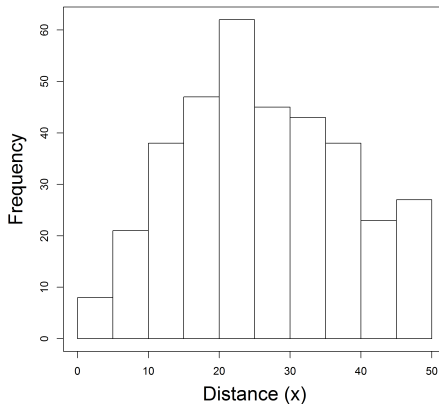
# POINT TRANSECTS. $\bar{p} = 1$

When average detection probability is 1, the histogram of distances will look like this because there is more area (and hence more individuals) far from the point.

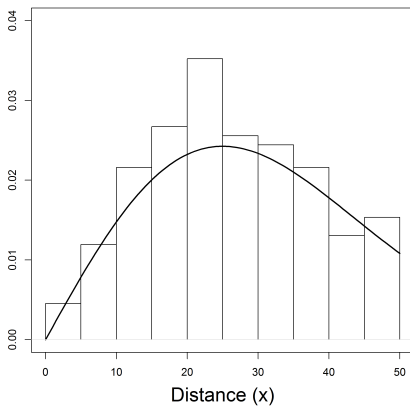


# POINT TRANSECTS. $\bar{p} < 1$

If average detection probability is less than 1, the histogram will increase and then decline.



The fitted curve will look like this





Read Chapter 10 – Capture-Mark-Recapture