Design and Analysis of Distance Sampling Studies

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Part 4 - Mark-Recapture Distance Sampling

Part 4
Mark-recapture distance sampling

Mark-recapture Distance sampling (MRDS)

MRDS - Mark-Recapture Distance Sampling WARNING: Tough sledding ahead!

MRDS - Introduction

Mark-Recapture Distance Sampling

- Allows $g(0) \neq 1$ by using capture-recapture to estimate detection probability.
 - $E[\widehat{D}] = Dg(0)$ so bias can be considerable if g(0) < 1.
- Usually has two observers who can (independently) identity individual animals and match animals seen by both, seen by one and not the other. This gives an estimate of animals missed and hence g(0).
- 'Observer' is a generic term any two methods of observation along the transect will work.

Implemented in DISTANCE through a call to R, but not yet fully featured.

CHECK TO SEE THAT R is installed and working.

MRDS - Introduction

Types of observer data (which influences analysis)

- Independent configuration. Two observers record location and distance of animals without communication with each other.
 Treatment of observers is symmetric.
- Trial configuration. Observer 2 "seeds" animals which are then detected by Observer 1. E.g. Observer 2 could be radio detections. Observers are not treated symmetrically.
- Removal configuration. Observer 1 "removes" animals and Observer 2 only looks for new animals. E.g. Observer 1 could be map of known nests seen in previous surveys.

Removal configurations not yet implemented in DISTANCE.

MRDS data structure

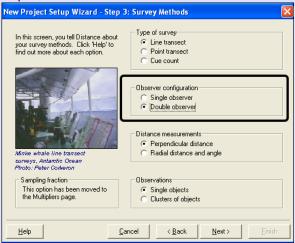
- Usual fields as before +
- NEW: Unique identifier for each object and 2 lines for each object (one for each observer)
- NEW: Field identifying the observer, the distance, and if each observer detected (0=no, 1=yes)

Refer to GolfTee example. **Independent configuration** of observers.

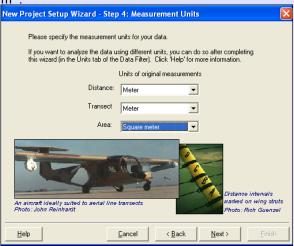
- There is a field for sex (I didn't know that golf tees had sexes)
 which actually is color (green=0 or yellow=1).
- There is a field for exposure if tees are above grass (exposed=1) or within grass (exposed=0).
- Two teams = two observers.

Notice that there is only 1 transect (not a good idea!).

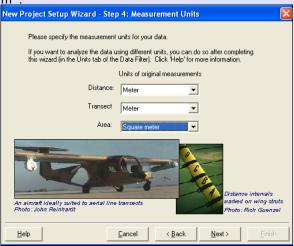
Import the data in the usual fashion.



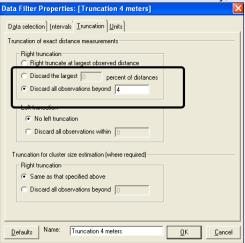
 \dots Import the data in the usual fashion - lengths in m, and area in $\underline{\mathsf{m}^2}$.



 \dots Import the data in the usual fashion - lengths in m, and area in $\underline{\mathsf{m}^2}$.



Create a Data Filter in the usual way, truncating at 4 m.



There are four detection functions!

- Observer 1 detection function $p_1(y, z)$ where z are covariates
- Observer 2 detection function $p_2(y, z)$
- Observer 1 detects Observer 2 detects $p_{1|2}(y,z)$
- Observer 2 detects Observer 1 detects $p_{2|1}(y,z)$

What is the shape of detection functions if Observer 1 is more experienced than Observer 2?

What is the shape of detection functions if Observer 1 and Observer 2 are both equally experienced.

The four detection functions!

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What is the shape of detection functions if Observer 1 is more experienced than Observer 2?

What is the shape of detection functions if Observer 1 and Observer 2 are both equally experienced.

Simplest (unreasonable) Case

Suppose that all animals were equally detectable at all distances, and observers are independent, but a different detection probability for each observer?

What do the detection functions look like?

Then we get for each animal:

•
$$P(1,0) = p_1(1-p_2)$$

•
$$P(0,1) = (1-p_1)p_2$$

•
$$P(1,1) = p_1p_2$$

Now a standard mark-recapture experiment and $\widehat{N} = \frac{n_1 n_2}{n_{12}}$, i.e. Petersen estimator

More complex (but still unreasonable) Case Suppose that detectability of animals varies by distance, and observers are independent, but a different detection probability for each observer (that varies by distance)?

Then we get for each animal:

•
$$P(1,0)(y) = p_1(y)(1-p_2(y))$$

What do the detection functions look like?

•
$$P(0,1)(y) = (1-p_1(y)) p_2(y)$$

•
$$P(1,1)(y) = p_1(y)p_2(y)$$

Break distances into bands, do a separate Petersen in each band, and add them up.

Reality Case

Suppose that detectability of animals varies by distance, and observers are NOT independent, but a different detection probability for each observer (that varies by distance)? What do the detection functions look like?

Then we get for each animal:

- $P(1,0)(y) = p_1(y) (1 p_{2|1}(y))$
- $P(0,1)(y) = (1 p_{1|2}(y)) p_2(y)$
- $P(1,1)(y) = p_1(y)p_{2|1}(y) = p_{1|2}(y)p_2(y)$

Now we have to estimate all 4 detection functions!

Observers acting independently does NOT imply that detection functions are independent.

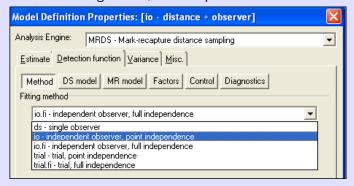
Suppose detection depends on distance y and cluster size s with perfect detection at the origin. If observer 1 detects an object at distance y it is more likely to be a larger cluster which implies that observer 2 will detect it.

We call this *Point Independence* (independence only at y=0) to distinguish it from *Complete Independence*

Specifying models in DISTANCE in MRDS.

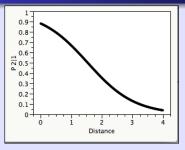
Four types of models depending on configuration and type of independence assumed:

- independent configuration, point independence
- independent configuration, full independence
- trial configuration, point independence
- trail configuration, full independence



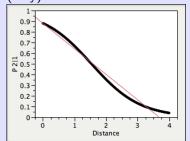
Two different models for detection must be specified:

- Distance Sampling (DS) model $(p_1(y, z))$ and $p_2(y, z)$.
- Conditional probability of detection models (MR) $(p_{1|2}(y,z))$ and $p_{2|1}(y,z)$



Develop a model for this curve?

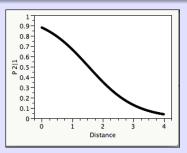
Try a straight line, $P_{2|1} = \beta_0 + \beta_1(distance)$, but not satisfactory (why)

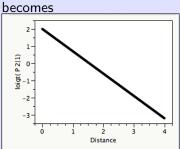


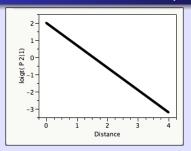
To model these 'S'-shaped curves and prevent the line from going $>1\ {\rm or}<0$, we often model

$$logit(p) = ln\left(\frac{p}{1-p}\right)$$

Probability	Odds	Logit
.01	1:99	-4.59
.1	1:9	-2.20
.5	1:1	0
.6	6:4 or 3:2 or 1.5	.41
.9	9:1	2.20
.99	99:1	4.59







Now modeling on logit-scale

$$logit(p_{2|1}) = \beta_0^* + \beta_1^*(distance)$$

keeps 0 .

The reverse transformation can be written in two ways:

$$p = \frac{e^{log\text{-}odds}}{1 + e^{log\text{-}odds}} = \frac{1}{1 + e^{-log\text{-}odds}}$$

Specifying a model for the MR component in DISTANCE

Need to use a generalized linear model syntax for the right hand part of the model.

Can use continuous and categorical (factors) in the model.

Examples:

- $logit(p_{2|1}) = 1$ corresponds to $logit = \beta_0$.
- $logit(p_{2|1}) = distance$ corresponds to $logit = \beta_0 + \beta_1(distance)$
- $logit(p_{2|1}) = sex$ corresponds to $logit = \beta_0 + \beta_1(I(sex = f))$ where I() is an indicator function
- $logit(p_{2|1}) = distance + sex$ corresponds to $logit = \beta_0 + \beta_1(distance) + \beta_2(I(sex = f))$
- $logit(p_{2|1}) = distance + sex + distance : sex$ corresponds to $logit = \beta_0 + \beta_1(distance) + \beta_2(I(sex = f)) + \beta_3(I(sex = f))(distance)$

Draw the curves (in logit space) for each of these.

Creating MR component in MRDS

Define the component and independence structure. We will start with independent configuration and full independence among observers.

```
Model Definition Properties: [io - distance + observer]
Analysis Engine:
                   MRDS - Mark-recapture distance sampling
  Estimate Detection function | Variance | Misc.
   Method DS model MR model Factors
                                             Control
                                                      Diagnostics
  Fitting method
     io.fi - independent observer, full independence
```

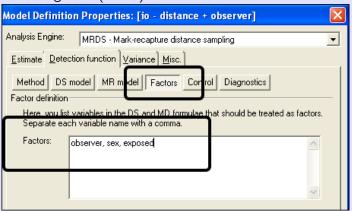
Notice poor choice of words for "io" option for configuration.

Creating MR component in MRDS



CAUTION: Variable names are a bit trick - see later slides.

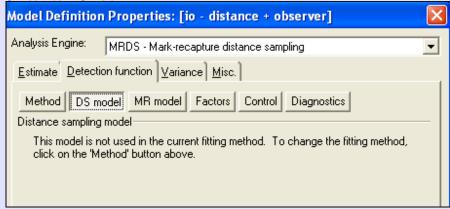
Creating MR component in MRDS
List categorical (factor) variables in FACTORS tab.



CAUTION: Variable names are a bit trick – see later slides. No harm in listing all variables in dataset here even if not used in model.

Creating DS component in MRDS

Switch to DS Tab:

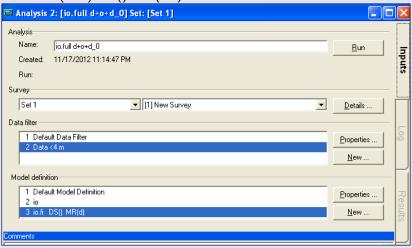


Why? If fully independent, then MR functions provides ALL information needed.

Combination of DS and MR components needed:

Fitting method	Configuration	Independence	DS model	MR model	
ds	single ¹	-	yes	no	
io	independent	point	yes	yes	
io.fi	independent	full	no	yes	
trial	trial	point	yes	yes	
trial.fi	trial	full	no	yes	
removal ²	removal	point	yes	yes	
removal.fi ²	removal	full	no	yes	

io.fi Data(< 4) DS() MR(dis): Fit Model ...



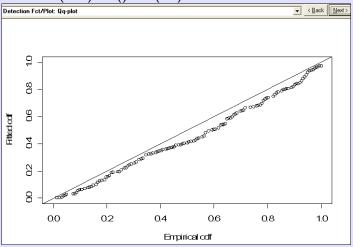
```
io.fi Data(< 4) DS() MR(dis): ... Results ...
Here is a list of the variable (field) names you can use in the
model. See reference manual for rules.
The following fields will be written to the data file, and
formulae. Note that you should use the new names, not the o
formulae, and that formulae names are case sensitive.
  Format: [layer name].[field name] AS new name
  [Observation].[Perp distance] AS distance
  [Observation].[object] AS object
  [Observation].[observer] AS observer
  [Observation].[detected] AS detected
  [Observation].[Cluster Size] AS cluster.size
  [Observation].[Sex] AS sex
  [Observation].[Exposure] AS exposure
  [Line transect].[Label] AS label
  [Line transect].[Line length] AS line.length
  [Region].[Label] AS stratum.label
  [Region].[Area] AS area
  [Study area].[Label] AS qlobal.label
```

io.fi Data(< 4) DS() MR(dis): ... Results ...

```
Summary for io.fi object
Number of observations : 162
Number seen by primary : 124
Number seen by secondary: 142
Number seen by both : 104
                        : 701.3888
AIC
Conditional detection function parameters:
            estimate
                            SA
(Intercept) 2.8806278 0.3439334
distance -0.9179038 0.1478512
                        Estimate
                                         SE
                     0.8705244 0.024878308 0.028578531
Average p
Average primary p(0) 0.9468804 0.015767751 0.016652315
Average secondary p(0) 0.9468804 0.015767751 0.016652315
Average combined p(0) 0.9971783 0.002726864 0.002734580
N in covered region
                     186.0947321 7.480834001 0.040199064
```

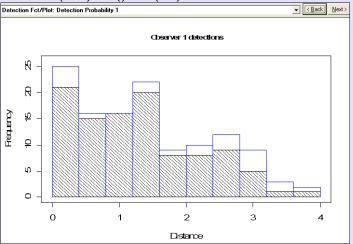
What does this model say?

io.fi Data(< 4) DS() MR(dis): ... Results ...



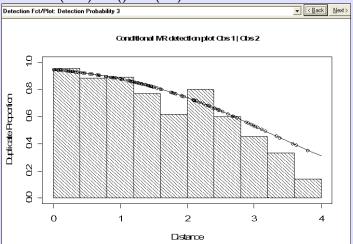
What does this model say?

io.fi Data(< 4) DS() MR(dis): ... Results ...



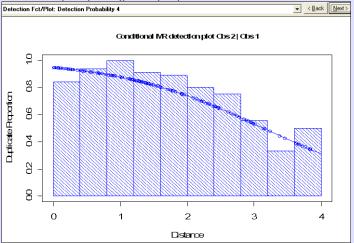
What does QQ-plot indicate?

io.fi Data(< 4) DS() MR(dis): ... Results ...



What does this and next plot say?

io.fi Data(< 4) DS() MR(dis): ... Results ...



What does this and previous plot say?

io.fi Data(< 4) DS() MR(dis): ... Results ...

```
< Back
Density Estimates and associated quantities
Summary statistics:
  Region Area CoveredArea Effort n k ER se.ER cv.ER
                            210 162 1 0.7714286 NaN
       1 1680
                  1680
Abundance:
 Label Estimate
                                          1.01
1 Total 186.0947 14.64169 0.07867871 105.2985 328.8866 1.326888
Density:
 Label Estimate
                                                1.01
                                                          1101
Total 0.1107707 0.008715294 0.07867871 0.06267765 0.1957658 1.326888
```

Notice that because there was only 1 transect, variance is understated.

Fit the io.fi Data(< 4) DS() MR(dis+obs) model.

Don't forget to specify observer as a factor (category). What do you conclude?

io.fi Data(< 4) DS() MR(dis+obs): ... Results ...

What does this model look like?

```
io.fi Data(< 4) DS() MR(dis+obs): ... Results ... What do you conclude?
```

Fit the io.fi Data(< 4) DS() MR(dis+obs+dist:obs) model. Don't forget to specify observer as a factor (category). Interaction denoted by colon in model formulae.

What do you conclude?

```
Fit the io.fi Data(< 4) DS() MR(dis+obs+dist:obs+color+exposure)
```

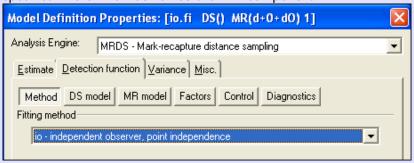
Don't forget to specify observer, color, exposure as a factor (category).

What do you conclude?

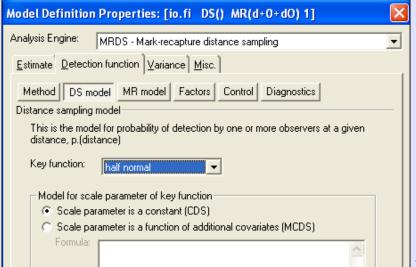
Point independence.

Here you assume observers have independent detections only a ONE point (typically the transect line).

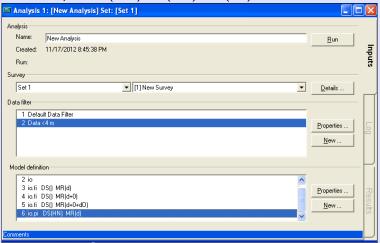
This now requires a separate model for the detection function specified in a similar fashion as the MR component.



Fit the
io.pi Data(< 4) DS(HN)
MR(dis+obs+dist:obs+color+exposure)



Fit the io.pi Data(< 4) DS(HN) MR(dis)



Model: io.pi Data(< 4) DS(HN) MR(dis)

What do you conclude?

More complex models(!)

	PI	PI		
$p_{j 3-j}(y,\underline{z})$	$g_{\cdot}(y,\underline{z})$	$\Delta { m AIC}$	\widehat{N}	$\Delta { m AIC}$
D D	1	64.2	232.0	67.6
D D	C	59.6	236.9	
D D	S	66.2	232.0	
■ D	E	66.2	232.0	
D	C + E	60.7	237.0	
b D	C + S	60.5	236.0	
D+P	C	55.9	236.8	
D + C	C	54.4	237.0	55.4
■ D+S	C	59.2	236.9	
D + E	C	25.4	237.5	
D+C+P	C	50.7	236.9	
D+S+P	C	55.5	236.9	63.7
D+E+P	C	21.8	237.4	
$\mathbf{M} D + C + P + E$	C	6.7		14.5
D+S+P+E	C	19.9	237.5	
D+S+P+C	C	49.8	237.0	50.5
D+P+C+S+E	C	4.3	239.0	11.5
D + P + C + S + E + C : E	C	2.8	250.8	
D + P + C + S + E + S : E	C	6.2	239.1	13.4
D + P + C + S + E + S : C	C	1.7	239.0	14.9
D+P+C+S+E+S:C+C:E	C	0.0	252.0	16.1
D+P+C+S+E+S:C+S:E	C	2.7	240.7	17.8
D + P + C + S + E + E : C + S : E	C	4.7	251.2	14.8
D + P + C + S + E + E : C + S : E + C : S	C	0.7	271.8	18.2

P=Obs; C=Color; S=Size; E=Exposure

MRDS - Exercise

Harbour Porpoise sample data from 1994 SCANS survey

- Independent configuration of observers.
- 98 transect lines, 1 km half-width of varying lengths (km)
- Recorded distance to group, group size, sex (groups are composed of a single sex), and exposure (a detectability factor related to weather and other attributes)
- Total survey region is 889600 km2

Data set is large, so model take a few seconds to fit!

MRDS - Summary

Deals with $g(0) \neq 1$. Survey Protocol

- Independent configuration.
- Trial configuration.
- Removal configuration (not yet implemented in MRDS).

MRDS - Summary

Data structure:

- The usual + object identifier +
- Field for Observer (limit currently is 2) + Detected (1=yes, 0=no)

Group the data together in the usual way. Import in the usual way.

MRDS - Summary

Model Building.

- Start with small models and build up to more complex models
 - Full independence vs. Point independence
 - Few covariates vs. many covariates
 - Complex models lead to cases where categories of animals seen by one observer are never seen by the other observer leading to an infinite population size!
- DS (HN or HR, but no series); MR (logistic regression)
- Use Linear Model syntax, e.g. distance + sex + distance:sex
- AIC for model comparison over all options within same data filter

R package has more features, but less easy to use.