



# MIGRATING DISTANCE SAMPLING PROJECTS FROM DISTANCE FOR WINDOWS TO THE DISTANCE R PACKAGE

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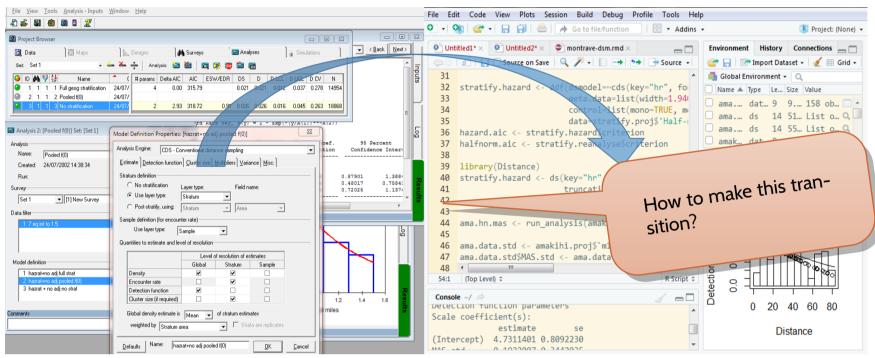
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### Introduction

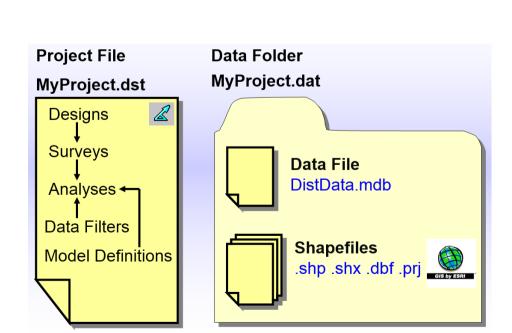
The Distance software (Thomas et al., 2010) has been downloaded >40,000 times in its 20-year history. It consists of a Windows-based Graphical User Interface (GUI) that gives users ready access to survey design tools (written in Visual Basic) and analysis engines (written in FORTRAN and, more recently, R). For some users, there my be benifits to performing their analyses directly with the underlying R code, rather than working with the GUI.



Challenges hindering the transition between analysis with the GUI and analyses in R are two-fold:

- Legacy data reside in Distance (GUI) projects, unavailable for importing into R, and
- Analyses that are easily described using the GUI may be difficult to specify, particularly if analyst is not proficient in R.

## Getting data out of Distance



Distance sampling analysis are defined in Distance project files (an Access database).

- Data
- Model definitions
- Analysis results

readdst uses RODBC (Windows, the 32-bit version of R must be used for RODBC to function properly) or mdb-tools (Mac) to mine the information

from the database. With the database components available, a sequence of translation steps are undertaken.

- Translate model definitions into R code
- Save data into an R environment
- Extract results from the database for completed analyses

## **Converting Distance projects to R**

Workhorse function in readdst package is <code>convert\_projects()</code>. Its single argument is simply the path to an existing Distance GUI project (created by Distance 6 or 7). <code>convert\_projects()</code> interrogates the Access database and translates the contents of database tables into an R object of class <code>converted\_distance\_analyses</code>, a list of lists of class <code>converted\_distance\_analyses</code>.

## Object produced by convert\_project()

The list returned by <code>convert\_project()</code>, for a given entry (analysis), contains all the salient information of a Distance analysis extracted from the Access database. We point out two elements of the list critical to subsequent processing of the now-extracted data.

**call** R code call to the function ddf() in package mrds to duplicate the analyses done in the Distance GUI.

**env** An *environment* consisting of a set of data frames. These data frames include the original data extracted from the Distance project, along with the observation, sample and region tables describing the hierarchial nature of the data base.

The converted\_distance\_analysis list can be passed as an argument to run\_analysis(). Given object has been created by a call to convert\_project, run\_analysis() will perform an mrds analysis by processing data in object\$env using ddf() syntax in object\$call to conduct a distance sampling analysis.

# Learning distance sampling in R

Distance GUI users can use converted\_distance\_analysis objects to learn how to perform corresponding analyses using the mrds R package:

```
> library(readdst)
> stratify.proj <- convert_project("C:/Distance Projects/myproject")</pre>
> stratify.proj
  ID
                                                            Status
1 13 Half-normal cosine no stratification exact
                                                            Ran OK
                Half-normal cosine pooled detfn
                                                            Ran OK
2 16
        Half-normal cosine strat-specific detfn Ran with warnings
> stratify.proj[[1]]
Model name : Half-normal cosine no stratification exact
Data filter :
           : mrds::ddf(dsmodel=~cds(key="hn", formula=~1,
mrds call
```

adj.series="cos", adj.order=NULL),

meta.data=list(width=NA, left=0),

method="ds", data=data)

Note too that components from objects of type converted\_distance\_analysis can be analysed using the ds() function from the Distance package.

control=list(mono=TRUE, mono.strict=TRUE),

# Comparative analysis of data sets

The test\_stats() function in readdst performs the analysis residing in the call element of the converted\_distance\_analysis, producing estimates of parameters of distance sampling analyses (e.g. AIC scores,  $\hat{P}_a$ ,  $\hat{D}$ ,  $\hat{D}$  and associated measures of precision). These results are contrasted with estimates of the same parameters stored in the Access data base, as computed by the Distance GUI.

The resulting comparison is produced in tabular form with the difference between estimates presented as proportions. Those comparative estimates within a specified tolerance are designated with **X**.

Statisti	c Distance_valu	ie mrds_valu	e Differenc	e Pass
n	90	90	0	X
parameters	1	1	0	X
AIC	63.880100250244	63.879819152981	0.000004396471	X
Chi^2 p	0.1707298	0.9906514	4.802452	
P_a	0.451959013939	0.451956754149	0.000004969147	X
CV(P_a)	0.07679188997	0.07679084338	0.00001362931	X
log-likelihood	-30.94005	-30.93991	0	X
C-vM p	1	0.9	0.1	
density	0.05059615	0.02768243	0.4528748	
CV(density)	0.2693232	0.3070729	0.1401652	
individuals	34658	18962.4623604	0.4528691	
V(individuals)	0.2693232	0.3070729	0.1401652	

This testing capability is part of our strategy to migrate software development from the existing Distance GUI to R packages. We can check consistency of results between the Distance GUI and developing R code.

#### **Caveats**

readdst is not able to translate all GUI analyses into R code. Current limitations are inability to translate

- analyses using the dsm, mads and Dssim engines,
- analyses using post-stratification and
- bootstrapping for variance estimation.

#### **Additional information**

#### References

Miller, D. L., E. Rexstad, L. Thomas, L. Marshall, and J. Laake. 2016. Distance Sampling in R. bioRxiv.

Thomas, L., S. T. Buckland, E. A. Rexstad, J. L. Laake, S. Strindberg, S. L. Hedley, J. R. Bishop, T. A. Marques, and K. P. Burnham. 2010. Distance software: design and analysis of distance sampling surveys for estimating population size. *Journal of Applied Ecology*, 47(1):5-14.



QR code readdst Github



QR code Miller et al. (bioRxiv)



QR code
Thomas et al.
(2010)