

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

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## Introduction

The Distance software (Thomas et al., 2010) has been downloaded >40,000 times in its 20-year history. Much of the underlying machinery is written in R. For some users, there may be benefits to performing the analysis with the underlying R code, rather than working with the graphical user interface (GUI).

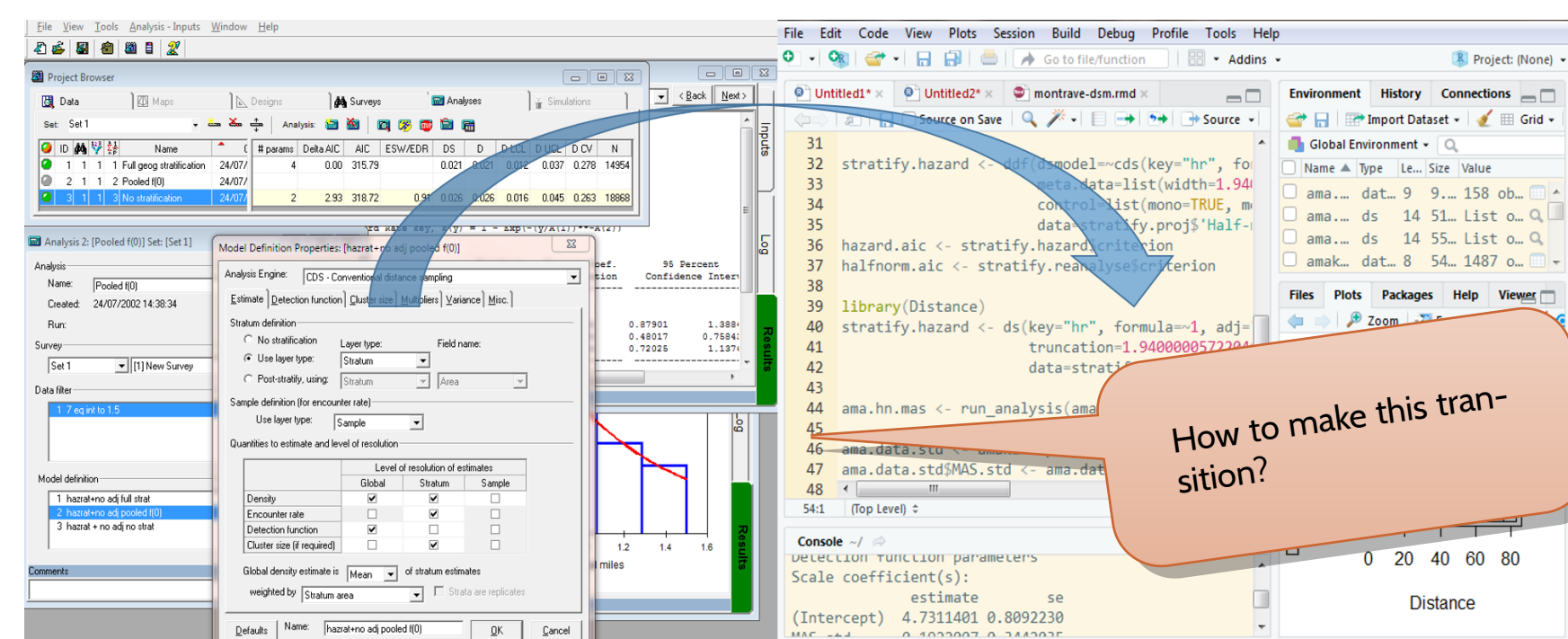


Fig. 1: Traditional Distance for Windows interface (left) and distance sampling analysis in R (right).

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.

## How to bridge between the two?

Distance GUI projects contain essential information necessary to conduct an analysis. The fundamental purpose of the `readdist` package is to access this information and place it into R objects for further scrutiny.

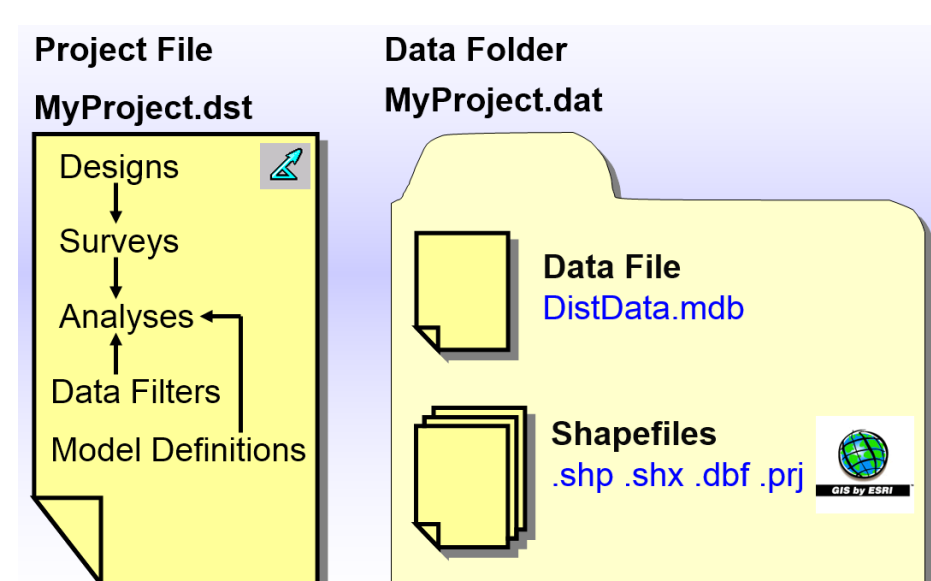


Figure 1: Database structure of a Distance project.

- Trick is to extract contents of (2) and translate into R code
- Target the results of previous step to contents of (1) the data
- Perhaps contrast results of R analysis with results stored in the Access database

The elements of a distance sampling analysis are contained inside Distance project files in the form of an Access database.

1. Data
2. model definitions
3. analysis results

Mining information from tables within the database, via either RODBC (Windows) or `mdb-tools` (Macs). Then it is a matter of performing the following three steps.

## Applications-Legacy projects

- We try to do this. what?
- we try to do that.

## Learning structure of R interface



Fig. 3: The organisation for which we work

## Comparative analysis of difficult data

- We try to do this. what?
- we try to do that.

## Caveats

`readdist` 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
- bootstraps for variance estimation.

## Additional information

### References

- Miller, D. L., 2017. *Package readdist*.
- 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 codes to package/website/bioRxiv

## Object produced by `convert_project()`

Figure 4 (left) shows the structure of an object of type `converted_distance_analysis`. There are as many of these produced by `convert_project()` as there were analyses in the Distance project.

The list contains all the salient information of a Distance analysis extracted from the Access database. We point out several of the list elements critical to subsequent processing of the now-extracted data.

**call** R code call to the function `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 hierarchical nature of the data base.

The `converted_distance_analysis` list can be passed as an argument to `run_analysis()`.

```
analysis_object
├── call
├── aic.select
├── status
├── env
│   ├── data
│   ├── obs.table
│   ├── region.table
│   └── sample.table
├── units
├── filter
├── group_size
├── detection_by
├── gof_intervals
├── estimation
├── ID
├── engine
├── project
└── project_file
```

Fig. 4:  
Structure of