

# Sprint session

NetCDF functionality in the R statistical software

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# The afternoon programme

- Outline
- Installing R
- Introduction to R
- R language basics
- Exercises

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- Outline
  - Installing R
  - Introduction to R
  - R language basics
  - Exercises
- 
- Introduction to NetCDF
  - Using NetCDF format in R
  - Working with your own data

# Installing R Statistical Software

Downloading R from the R-project website: <http://www.r-project.org>

The screenshot shows a Windows Internet Explorer browser window displaying the R Project website. The address bar shows <http://www.r-project.org/>. The page title is "The R Project for Statistical Computing".

The main content area features the R logo on the left and a navigation menu with links: "About R", "What is R?", "Contributors", "Screenshots", "What's new?", "Download, Packages", "CRAN", "R Project", "Foundation", "Members & Donors", "Mailing Lists", "Bug Tracking", "Developer Page", "Conferences", "Search", "Documentation", "Manuals", "FAQs", "The R Journal", "Wiki", "Books", "Certification", "Other".

The central part of the page displays several statistical plots:

- PCA 5 vars**: A circular plot showing the distribution of variables: Family, Examination, Education, and Agriculture. A label "(1-3) 60%" is present.
- Clustering 4 groups**: A dendrogram showing hierarchical clustering of data points into four groups.
- Factor 1 [41%]** and **Factor 3 [19%]**: Two histograms showing the distribution of data points along these factors.

Below the plots is a "Getting Started:" section with the following text:

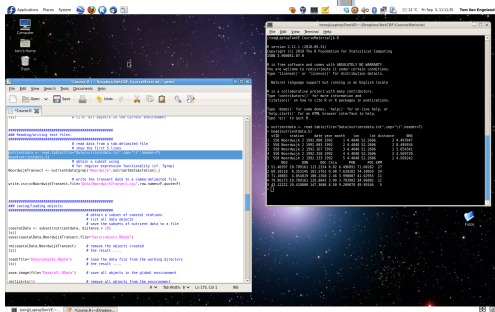
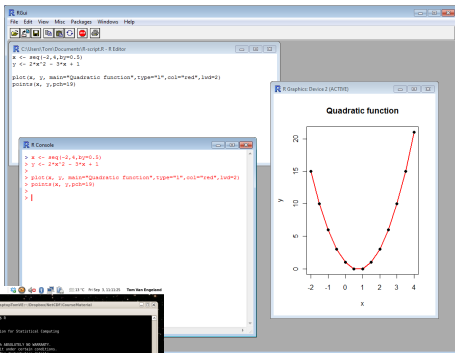
Getting Started:

- R is a free software environment for statistical computing and graphics. It compiles and runs on a wide variety of UNIX platforms, Windows and MacOS. To [download R](#), please choose your preferred [CRAN mirror](#).
- If you have questions about R like how to download and install the software, or what the license terms are, please read our [answers to frequently asked questions](#) before you send an email.

The browser's status bar at the bottom indicates "Internet | Protected Mode: On" and a zoom level of "100%".

# Installing R Statistical Software

- Rgui  
(Graphical User Interface)  
( !Windows only! )
- Rconsole  
(+ any lightweight text editor)  
( !All systems! )



# Installing R Statistical Software

- Dedicated GUIs and IDEs

- ▶ Windows

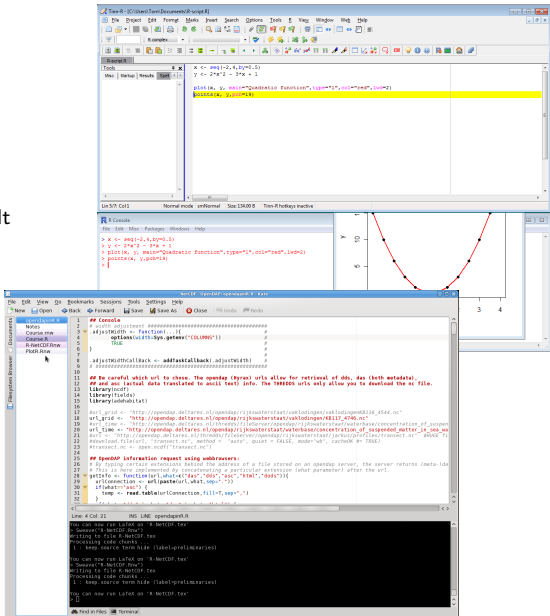
- ★ Tinn-R
- ★ WinEdt + RWinEdt
- ★ TextPad

- ▶ Linux

- ★ Kate
- ★ gEdit

- ▶ Multiplatform

- ★ Komodo Edit + SciViews-K
- ★ ESS for Emacs
- ★ Eclipse
- ★ jEdit
- ★ Alpha



# GNU's S

- Statistics and graphics
- Open source and distributable under GPL
- R and S code are largely exchangeable

# GNU's S

- Statistics and graphics
- Open source and distributable under GPL
- R and S code are largely exchangeable
- Plugin packages
- good connectivity with other software packages
- platform independent (Unix-alikes, MacOS, Windows)

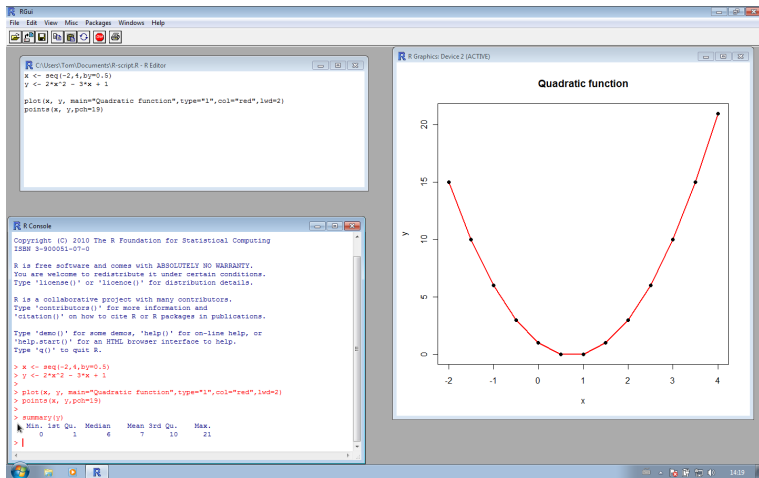


# Learning R

- Learning R consumes more time than for commercial packages
- Still many facilities for learning R
  - ▶ R help files
  - ▶ Tutorials
  - ▶ R Site Search (search engine)
  - ▶ Forums and mailinglists
  - ▶ Large community, producing and sharing code and small package-related tutorials
  - ▶ Best accessible with standard search engine (Google, ...)

# Talking to R

- A matter of question and answer



# Built-in R help facilities

- `?<command>`

Examples:

```
> ?help
> ?plot
> help(plot)
```

- `??<command>`, which searches for string matches in help files. The function names are returned:

```
> ??plot
> ??line
> help.search("line")
```

- R help starting page launched in a web browser:

```
> help.start()
```

# Built-in R help facilities

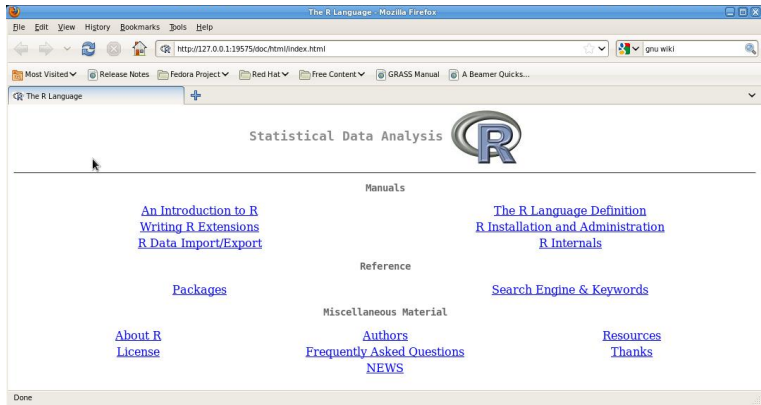


Figure: Start page for the R help files.

# The beginning

- Assignments, comments, and retrieving variables

```
> var1 <- "some value"
> var1                # type varname to retrieve content
[1] "some value"
> counter <<- 1
> counter
[1] 1
> temp = TRUE
> temp
[1] TRUE
```

- Exact meaning of different assignments: `help(assignOps)`

# Modes, types, and classes

- Data type determined upon assignment
- Internal storage mode
- Simple types combined in more complex data structures
- Single mode data structures (vectors, matrices, arrays)
- Multiple mode data structures (lists, data frames, externally defined classes)

Table: R data types and internal storage modes:

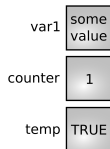
| <b>types</b> | <b>modes</b> |
|--------------|--------------|
| logical      | logical      |
| integer      | numeric      |
| double       | numeric      |
| complex      | complex      |
| character    | character    |
| symbol       | name         |
| raw          | raw          |
| list         | list         |
| NULL         | NULL         |
| ...          |              |

# Single values

```
> var1 <- "some value"  
> counter <- 1  
> temp = TRUE
```

This is actually a single valued vector:

```
> length(var1)  
[1] 1
```



# Vectors

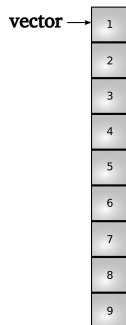
Extension in one dimension:

```
> b <- 1:9
> b
[1] 1 2 3 4 5 6 7 8 9

> c <- seq(0,3,by=0.5)
[1] 0.0 0.5 1.0 1.5 2.0 2.5 3.0

> temp = c(T,F,F,T)
> temp
[1] TRUE FALSE FALSE TRUE

> rep(c(1,3:1),times=2)
[1] 1 3 2 1 1 3 2 1
```





# Matrices

Extension to two dimensions:

```
> mat <- matrix(data=seq(1,9,by=1),nrow=3)
```

```
> mat
```

```
      [,1] [,2] [,3]  
[1,]    1    4    7  
[2,]    2    5    8  
[3,]    3    6    9
```

```
> dim(mat)
```

```
[1] 3 3
```

```
> mat[1,]
```

```
[1] 1 4 7
```

```
> mat[,1]
```

```
[1] 1 2 3
```

**matrix** →

|   |   |    |    |
|---|---|----|----|
| 1 | 5 | 9  | 13 |
| 2 | 6 | 10 | 14 |
| 3 | 7 | 11 | 15 |
| 4 | 8 | 12 | 16 |

# Arrays

Extension to n dimensions:

```
> mat4d <- array(data=seq(1,36,by=1),dim=c(3,3,2,2))
```

```
> mat4d
```

```
, , 1, 1
```

```
  [,1] [,2] [,3]
[1,]   1   4   7
[2,]   2   5   8
[3,]   3   6   9
```

```
, , 2, 1
```

```
  [,1] [,2] [,3]
[1,]  10  13  16
[2,]  11  14  17
[3,]  12  15  18
```

```
, , 1, 2
```

```
  [,1] [,2] [,3]
[1,]   1   4   7
[2,]   2   5   8
[3,]   3   6   9
```

```
, , 2, 2
```

```
  [,1] [,2] [,3]
[1,]  10  13  16
[2,]  11  14  17
[3,]  12  15  18
```

matrix →

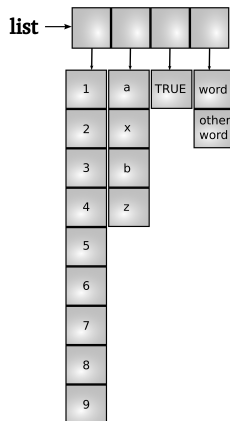
|   |   |    |    |
|---|---|----|----|
| 1 | 5 | 9  | 13 |
| 2 | 6 | 10 | 14 |
| 3 | 7 | 11 | 15 |
| 4 | 8 | 12 | 16 |

# Lists

- Combine different data types:

```
> datarecord <- list(name="Tom Van Engeland",  
+                   Department="Ecosystem Studies",  
+                   Functions=c(PhD=4,Postdoc=1),  
+                   Carpooling=T  
+                   )
```

- Different types
- Different sizes



# Data frames

- List elements of equal size - a data table

```
> dataset <-  
+ data.frame(treatment=  
+           rep(c("control", "level 1", "level 2"), e:  
+           replicate=rep(1:3, times=3),  
+           value=c(2,6,4,8,9,8,3,2,1)  
+           )  
> dataset  
  treatment replicate value  
1 control           1     2  
2 control           2     6  
3 control           3     4  
4 level 1           1     8  
5 level 1           2     9  
6 level 1           3     8  
7 level 2           1     3  
8 level 2           2     2  
9 level 2           3     1
```

data.frame →

| a | 1 | TRUE  | 1 |
|---|---|-------|---|
| a | 2 | NA    | 2 |
| a | 3 | TRUE  | 3 |
| b | 1 | FALSE | 4 |
| b | 2 | TRUE  | 5 |
| b | 3 | NA    | 6 |
| c | 1 | FALSE | 7 |
| c | 2 | FALSE | 8 |
| c | 3 | NA    | 9 |

# Accessing objects

- to see the content:

```
varname
```

- to see the structure:

```
str(object)  
ls.str()
```

- to acquire a more specific part of the information:

```
summary(object)  
length(vector)  
ncol(matrix)  
nrow(matrix)  
colnames(matrix)  
rownames(matrix)  
names(list)  
dimnames(array)  
subset(dataframe, rowselection, select=columnnames)
```

- indexing

```
matrix[rownumber, columnnumbers]  
matrix[rownames, columnnames]  
list[indexnumber] (pointer to list element)  
list[[indexnumber]] (list element at place <indexnumber>)  
list$listelement
```

- See help files on package specific functions

# Operators

## Arithmetic operators

|                                |                               |
|--------------------------------|-------------------------------|
| <code>x + y</code>             | addition                      |
| <code>x - y</code>             | subtraction                   |
| <code>x * y</code>             | multiplication (element-wise) |
| <code>x / y</code>             | division                      |
| <code>x %/% y</code>           | integer division              |
| <code>x %% y</code>            | modulo                        |
| <code>x ^ y</code>             | power                         |
| <code>x %*% y</code>           | matrix multiplication         |
| Logic operators (element-wise) |                               |
| <code>! x</code>               | negation                      |
| <code>x &amp; y</code>         | AND                           |
| <code>x   y</code>             | OR                            |
| <code>xor(x,y)</code>          | XOR                           |

Type ?Arithmetic,?Logic,?Special,or ?Syntax for more information !

# Managing Your R Environment

- Get all objects from an environment:

```
> ls()
```

- Search for objects in an environment:

```
> find("search string")  
> apropos("search string")
```

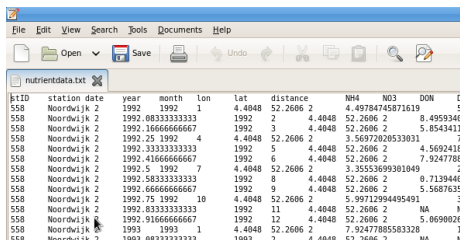
- Remove objects

```
> rm(object1, object2)  
> rm(list=ls())
```

# Text files

Provide a flexible means for exchange of small datasets

```
> nutrientdata <- read.table(file="Data/nutrientdata.txt", sep="\t", header=T)
> NoordwijkTransect <- nutrientdata[grep("Noordwijk", nutrientdata$station),]
> write.csv(x=NoordwijkTransect, file="Data/NoordwijkTransect.csv",
+          row.names=F, quote=F)
```



The screenshot shows a text editor window with a menu bar (File, Edit, View, Search, Tools, Documents, Help) and a toolbar with icons for Open, Save, Undo, Cut, Copy, Paste, Find, and Print. The main text area displays a tab-separated data file with the following content:

| stID | station     | date              | year | month | lon    | lat     | distance | NH4              | NO3 | DON       | 0 |
|------|-------------|-------------------|------|-------|--------|---------|----------|------------------|-----|-----------|---|
| 558  | Noordwijk 2 | 1992              | 1992 | 1     | 4.4048 | 52.2606 | 2        | 4.49784745871619 |     |           | 5 |
| 558  | Noordwijk 2 | 1992.083333333333 | 1992 | 2     | 4.4048 | 52.2606 | 2        | 52.2606          | 2   | 8.4959340 |   |
| 558  | Noordwijk 2 | 1992.166666666667 | 1992 | 3     | 4.4048 | 52.2606 | 2        | 52.2606          | 2   | 5.8543411 |   |
| 558  | Noordwijk 2 | 1992.25           | 1992 | 4     | 4.4048 | 52.2606 | 2        | 3.56972020533031 |     |           | 7 |
| 558  | Noordwijk 2 | 1992.333333333333 | 1992 | 5     | 4.4048 | 52.2606 | 2        | 52.2606          | 2   | 4.5692418 |   |
| 558  | Noordwijk 2 | 1992.416666666667 | 1992 | 6     | 4.4048 | 52.2606 | 2        | 52.2606          | 2   | 7.9247788 |   |
| 558  | Noordwijk 2 | 1992.5            | 1992 | 7     | 4.4048 | 52.2606 | 2        | 3.35553699301049 |     |           | 2 |
| 558  | Noordwijk 2 | 1992.583333333333 | 1992 | 8     | 4.4048 | 52.2606 | 2        | 52.2606          | 2   | 0.7139448 |   |
| 558  | Noordwijk 2 | 1992.666666666667 | 1992 | 9     | 4.4048 | 52.2606 | 2        | 52.2606          | 2   | 5.5687635 |   |
| 558  | Noordwijk 2 | 1992.75           | 1992 | 10    | 4.4048 | 52.2606 | 2        | 5.99712994495491 |     |           | 3 |
| 558  | Noordwijk 2 | 1992.833333333333 | 1992 | 11    | 4.4048 | 52.2606 | 2        | NA               |     |           | N |
| 558  | Noordwijk 2 | 1992.916666666667 | 1992 | 12    | 4.4048 | 52.2606 | 2        | 52.2606          | 2   | 5.0699026 |   |
| 558  | Noordwijk 2 | 1993              | 1993 | 1     | 4.4048 | 52.2606 | 2        | 7.92477885583328 |     |           | 1 |
| 558  | Noordwijk 2 | 1993.083333333333 | 1993 | 2     | 4.4048 | 52.2606 | 2        | NA               |     |           | N |



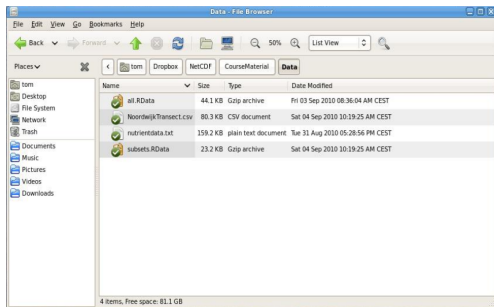
# .RData files

- Write individual data objects to a R-specific binary data format (.RData):

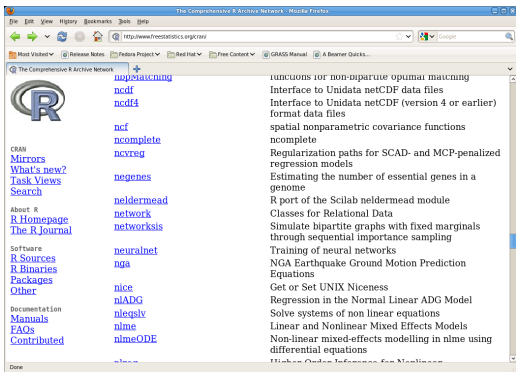
```
> coastalData <- subset(nutrientdata, distance < 10)
> save(coastalData, NoordwijkTransect, file="Data/subsets.RData")
> load(file="Data/subsets.RData")
```

- Save all objects from an environment to an .RData file:

```
> save.image(file="Data/all.RData")
> load(file="Data/all.RData")
```



# Packages



The screenshot shows a web browser window displaying the CRAN website. The browser's address bar shows the URL <http://www.freeStatistics.org/cran/>. The page content is organized into several sections:

- CRAN**: Includes links for [Mirrors](#), [What's new?](#), [Task Views](#), and [Search](#).
- About R**: Includes links for [R Homepage](#) and [The R Journal](#).
- Software**: Includes links for [R Sources](#), [R Binaries](#), [Packages](#), and [Other](#).
- Documentation**: Includes links for [Manuals](#), [FAQs](#), and [Contributed](#).

The main content area displays a list of R packages with their names and brief descriptions:

- [nnpMatchng](#): Functions for non-parametric optimal matching
- [ncdf](#): Interface to Unidata netCDF data files
- [ncdf4](#): Interface to Unidata netCDF (version 4 or earlier) format data files
- [ncf](#): spatial nonparametric covariance functions
- [ncomplete](#): ncomplete
- [ncvreg](#): Regularization paths for SCAD- and MCP-penalized regression models
- [negenes](#): Estimating the number of essential genes in a genome
- [neldermead](#): R port of the Scilab neldermead module
- [network](#): Classes for Relational Data
- [networksis](#): Simulate bipartite graphs with fixed marginals through sequential importance sampling
- [neuralnet](#): Training of neural networks
- [nga](#): NGA Earthquake Ground Motion Prediction Equations
- [nice](#): Get or Set UNIX Niceness
- [niADG](#): Regression in the Normal Linear ADG Model
- [nleqslv](#): Solve systems of non linear equations
- [nlme](#): Linear and Nonlinear Mixed Effects Models
- [nlmeODE](#): Non-linear mixed-effects modelling in nlme using differential equations

- Only loaded when needed
- About 25 default packages
- Other packages managed on the CRAN website
- Install binary or from source after compilation

# Packages

- Installation

```
> install.packages("nlme",dependencies=T)
```

- Loading

```
> library(nlme)
```

```
> require(nlme)
```

Require is mainly intended to be called inside other functions, to allow for exception handling and a decent termination of the relevant function.

- Other functionalities:

```
> update.packages()
```

```
> available.packages()
```

```
> installed.packages()
```

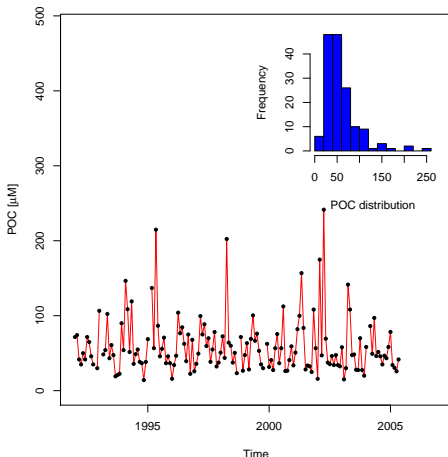
```
> remove.packages()
```

# Statistics and Graphs with R

- main purpose = data analysis
- production of publication quality graphs

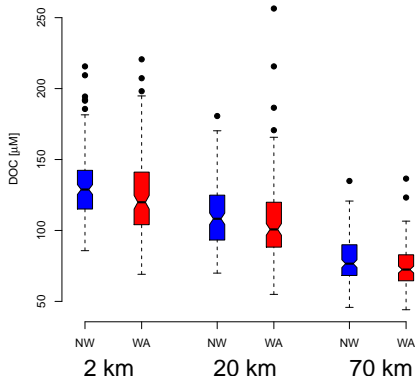
# Example 1:

```
> Noordwijk <- subset(coastalData,
+                       station=="Noordwijk 2")
> par(mar=c(4,4,0.5,0.5))
> plot(Noordwijk$date, Noordwijk$POC,
+       xlab="Time",
+       ylab=expression("POC ["*mu*"M]"),
+       type="l", col="red",
+       ylim=c(0, max(Noordwijk$POC, na.rm=T)*2))
> points(Noordwijk$date, Noordwijk$POC,
+        pch=19, cex=0.6)
> par(new=T, fig=c(0.55, 0.95, 0.55, 0.95),
+       mar=c(4, 4, 0, 0))
> hist(Noordwijk$POC, main="", col="blue",
+       xlab="POC distribution")
```



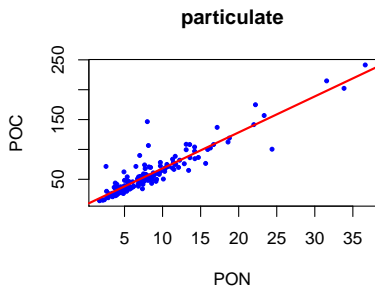
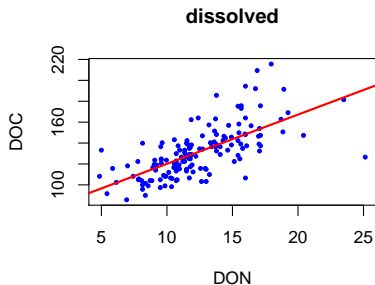
## Example 2:

```
> pdf(file="Illustrations/boxplots.pdf",
+     width=6.5,height=6.5)
> boxplot(DOC~station*distance,
+         data=nutrientdata,notch=T,
+         col=c("blue","blue","blue",
+              "red","red","red"),
+         axes=F,
+         ylab=expression("DOC ["*mu*"M]"),
+         pch=19)
> axis(2)
> axis(1,at=c(1,4,8,11,15,18),
+      labels=c("NW","WA","NW",
+              "WA","NW","WA"))
> mtext(1,outer=T,line=-2,
+       at=c(0.23,0.53,0.83),
+       text=c("2 km","20 km","70 km"),
+       cex=2)
> dev.off()
```



## Example 3:

```
> Dmodel <- lm(DOC ~ DON, data=Noordwijk)
> Pmodel <- lm(POC ~ PON, data=Noordwijk)
> par(mfrow=c(2,1))
> plot(Noordwijk$DON, Noordwijk$DOC,
+      main="dissolved", xlab="DON", ylab="DOC",
+      type="p", pch=19, cex=0.5, col="blue")
> abline(coef(Dmodel), col="red", lwd=2)
> plot(Noordwijk$PON, Noordwijk$POC,
+      main="particulate", xlab="PON", ylab="POC",
+      type="p", pch=19, cex=0.5, col="blue")
> abline(coef(Pmodel), col="red", lwd=2)
> mtext(side=1, outer=T, line=1, at=0.5,
+       text="organic matter C/N ratios")
```



# Writing your own functions

- to repeat certain tasks

```
someFunctionName <- function (set, of, paramaters) {  
  code block  
}
```

- Example:

```
CNratioExt <- function(carbon, nitrogen, inverse=F) {  
  if(!(is.vector(carbon) & is.vector(nitrogen))) {  
    print("This function only accepts vectors")  
    return()  
  }  
  # Test vector types  
  if(length(carbon) != length(nitrogen)) {  
    print("carbon and nitrogen dataset not of same size")  
    return()  
  }  
  # Checking equality of their lengths  
  if(inverse) {  
    return(nitrogen/carbon)  
  }  
  else {  
    return(carbon/nitrogen)  
  }  
}
```



# Flow control

- for-loop

```
for (i in seq(2,10,by=0.01)) {code block}
```

- while-loop

```
while (test == TRUE) {code block}
```

- if-then-else construct

```
if (test == TRUE) {code block}  
else {other code block}
```

- switch construct

```
switch(expression,  
choice1 = {code block},  
choice2 = {code block},  
choice3 = {code block})
```

- ?Control for more information

# R Exercise

# NetCDF and OpenDAP

# NetCDF

NetCDF = Network Common Data Form

Unidata (UCAR; University Corporation for Atmospheric Research) <http://www.unidata.ucar.edu>

Set of

- data formats
- programming interfaces
- software libraries (abstract data type)

- binary file
- regular N-dimensional arrays
- contains metadata (self-descriptiveness)  
= data on the data (crucial side-information)
- platform-independent (portability)

# The data model

- dimensions

- ▶ describe the extent of the variables' arrays
- ▶ axes of variability
- ▶ declare lengths of variables
- ▶ name + length
- ▶ fixed or unlimited
  - a dynamically adjustable dimension that can change throughout the creation of the file with no performance penalty

- variables

- ▶ the actual n-dimensional arrays of data
- ▶ fixed data type (char, byte, short, int, float, double)
- ▶ coordinate variables: data counterpart of dimensions (same name)
- ▶ and just "variables"

- attributes

- ▶ metadata
- ▶ scalar or 1D-array (vector)
- ▶ associated with variable or entire dataset (global)
- ▶ some reserved

# NetCDF versions

- Classical netCDF data model
  - ▶ data model as presented here
  - ▶ severe size restrictions  
(mainly for physico-chemical data; less for biological data)
  - ▶ only one unlimited dimension
  - ▶ only predefined set of base types
- Version 3.6.0
  - ▶ allowed for 64-bit based addresses  
=> max size doubled (4 GB)
- Common Data Model and HDF5 (NetCDF 4.1.1)
  - ▶ more data structuring (grouping; hierarchical structuring)
  - ▶ ?addition information on the data?
  - ▶ more unlimited dimensions
  - ▶ new base types
  - ▶ user-defined data types

# NetCDF versions

- Classical netCDF data model
  - ▶ data model as presented here
  - ▶ severe size restrictions  
(mainly for physico-chemical data; less for biological data)
  - ▶ only one unlimited dimension
  - ▶ only predefined set of base types
- Version 3.6.0
  - ▶ allowed for 64-bit based addresses  
=> max size doubled (4 GB)
- Common Data Model and HDF5 (NetCDF 4.1.1)
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**Backward** compatible



# NetCDF versions

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Versatility

Portability

**Backward** compatible

# What about relational databases?

- no relationships between data points
- no efficient storage and access to gridded data
- extra costs involved in DBMS do not pay-off  
as useful functionality (concurrency control, access rights, views, ...)

NetCDF is intended for gridded data which are more efficiently presented and managed using (ir)regular grids than by relationship (relational databases)

**BUT** Relational databases to link dataset through metadata?

# Metadata

- Rules of application partially delineated but not enforced  
tension between self-descriptiveness, dedication, and flexibility
- Provision left to user
- Standards and conventions needed
- Examples: <http://www.unidata.ucar.edu/netcdf/conventions.html>
- Which metadata needed for ecological variables?

# Advantages & disadvantages

- freely available
- data + variable description + creation history + other attributes
- reasonably compact portable binary format
- allows extraction of parts of large data file
- possibility to write your own interface (e.g. R-packages)

## **BUT**

- no 'ragged' objects
- extensive use of special 'missing values' may be needed

# NetCDF with R

# R packages

- Available packages

- ▶ ncdf
- ▶ RNetCDF
  - Not for OSX
- ▶ ncdf4
  - ★ Has netCDF 4 capability
  - ★ Has OpenDAP capability if compiled properly
  - ★ Does currently not work on Windows

- Installation

- ▶ Windows: binaries at CRAN
- ▶ Linux and OSX: need to install NetCDF libraries first

```
R CMD INSTALL --configure-args="  
-with-netcdf_incldir=/path/to/netcdf/include  
-with-netcdf_libdir=/path/to/netcdf/lib"  
ncdf_xx.yy.tar.gz
```

# The ncdf package

- reading `open.ncdf(...)` `get.var.ncdf(...)`
- writing `open.ncdf(...,write=T)` `put.var.ncdf(...)`
- creation `create.ncdf(...)` `dim.def.ncdf(...)` `var.def.ncdf(...)`

# Downloading and opening a NetCDF file

```
> url_grid <- "http://opendap.deltares.nl/opendap/  
+           rijkswaterstaat/vaklodingen/KB117_4746.nc"  
> download.file(url=url_grid,  
+             destfile="vaklodingenKB117_4746.nc",  
+             method = "auto",  
+             quiet = FALSE,  
+             mode="wb",  
+             cacheOK = TRUE)  
grid.nc <- open.ncdf("vaklodingenKB117_4746.nc")
```



# Querying a local NetCDF file

```
names(grid.nc)
```

| Name       | Description  |
|------------|--|
| id         | a file id  |
| ndims      | number of dimensions                                 |
| natts      | number of global attributes                          |
| unlimdimid | id of unlimited dimension or -1 if not present       |
| filename   | filename on disk                                     |
| writable   | file opened for writing of not                       |
| dim        | dimensions (named list of objects of class dim.ncdf) |
| nvars      | number of non-coordinate variables                   |
| var        | variables (named list of objects of class var.ncdf)  |

# Querying a local NetCDF file

Finding the variables and dimensions:

```
> names(grid.nc$var)
[1] "crs" "z" "lon" "lat"
> names(grid.nc$dim)
[1] "time" "y" "x"
> grid.nc$dim$time
$name
[1] "time"
$len
[1] 60
$unlim
[1] TRUE
$id
[1] 1
$dimvarid
[1] 2
$units
[1] "days since 0001-01-01 00:00:00 +1:00"
$svals
 [1] 713684 714415 714780 715145 715510 715876 716241 716606 716971 717337 717702 718067
[13] 718432 718798 719163 719528 719893 720259 720624 720989 721354 721720 722085 722450
[25] 722815 723546 724276 724642 725007 725372 725737 726103 726468 726833 727198 727564
[37] 727929 728294 728659 729025 729030 729176 729390 729395 729541 729755 729760 729906
[49] 730120 730125 730272 730486 730851 730856 731002 731216 731221 731367 731581 731947
$create_dimvar
[1] TRUE

attr(,"class")
[1] "dim_ncdf"
```

# Querying a local NetCDF file

Extraction of the variables:

Try `grid.nc$var$z`

... There is no data?

```
> x <- get.var.ncdf(nc=grid.nc,varid="x")
> y <- get.var.ncdf(grid.nc,varid="y")
> time <- get.var.ncdf(grid.nc,"time")
> z <- get.var.ncdf(grid.nc,"z")
> list(x=dim(x),y=dim(y),z=dim(z))
```

```
$x
[1] 500
```

```
$y
[1] 625
```

```
$z
[1] 500 625 60
```

Variables are read from the local file...

# Querying a local NetCDF file

- What about the attributes?
- No function to get a list of attributes?
- Attributes are scatter all over the object...

```
library(RNetCDF)
alternative <- open.nc("/home/tom/Dropbox/NetCDF/vaklodingenKB117_4746.nc")
print.nc(alternative)
```

# Querying a local NetCDF file

- What about the attributes?
- No function to get a list of attributes?
- Attributes are scatter all over the object...

```
library(RNetCDF)
alternative <- open.nc("/home/tom/Dropbox/NetCDF/vaklodingenKB117_4746.nc")
print.nc(alternative)
```

- Or ask them from the OpenDAP server...

# Querying a local NetCDF file

```
> getInfo(url_grid,what="dds") # cf. course
```

```
Dataset {  
  Int32 crs;  
  Grid {  
    ARRAY:  
      Float64 lat[y = 625][x = 500];  
    MAPS:  
      Float64 y[y = 625];  
      Float64 x[x = 500];  
  } lat;  
  Grid {  
    ARRAY:  
      Float64 z[time = 10][y = 625][x = 500];  
    MAPS:  
      Float64 time[time = 10];  
      Float64 y[y = 625];  
      Float64 x[x = 500];  
  } z;  
  Grid {  
    ARRAY:  
      Float64 lon[y = 625][x = 500];  
    MAPS:  
      Float64 y[y = 625];  
      Float64 x[x = 500];  
  } lon;  
  Float64 y[y = 625];  
  Float64 x[x = 500];  
  Float64 time[time = 10];  
} opendap/rijkswaterstaat/vaklodingen/vaklodingenKB117_4746.nc;
```

Try `getInfo(url_grid,what="das") ...`

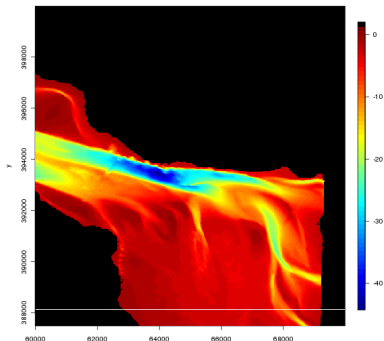
# Querying a local NetCDF file

Other extensions for OpenDAP querying:

- .help : generates a web based help document
- .html : generates a web page containing information and action buttons over the netcdf file
- .info : returns the netcdf file metadata
- .dds : returns information about the variables
- .das : returns information about the variable attributes
- .ver : returns the OpenDAP server version
- .asc : returns the netcdf file in ASCII format

# Plotting NetCDF data

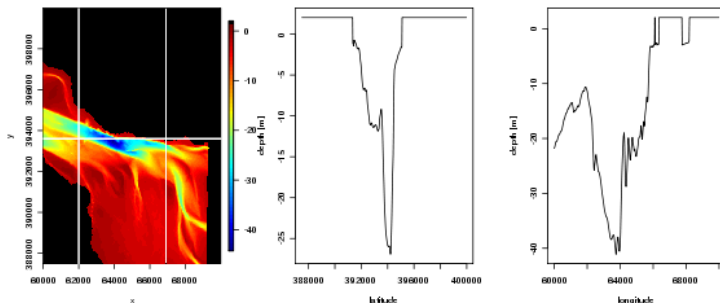
```
> # The plot function that is used needs variable values in ascending order
> y <- rev(y)
> z <- z[, (ncol(z):1), ]
> # select only the slices, containing data
> selection <- which(apply(z, MARGIN=3,
+                         FUN=function(a) diff(range(a, na.rm=T))) != 0,
+                         arr.ind=T)
> z <- z[, , selection]
> # Replace -9999 with some value closer to the relevant range
> z[z == -9999] <- max(z, na.rm=T)+1
> # the dimensions as headers in the depth array is more informative
> # Standard calendar has 365.242 days in a year...
> dimnames(z) <- list(x, y, round(
+                       time[selection]/365.242))
> # Plot the first non-empty time slice
> image.plot(x, y, z[, , 1],
+           col = c(tim.colors(), "black"))
```





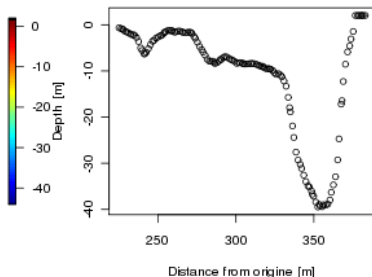
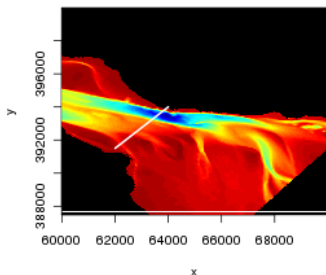
# Working with a data subset

```
> par(mfrow=c(1,3))  
> image.plot(x,y,z[, ,1],col = c(tim.colors(),"black"))  
> abline(v=62000,h=393600,col="white",lwd=2)  
> plot(y,z["62000", ,1],type="l",xlab="latitude",ylab="depth [m]")  
> plot(x,z[, "393600", 1],type="l",xlab="longitude",ylab="depth [m]")
```



# Working with a data subset

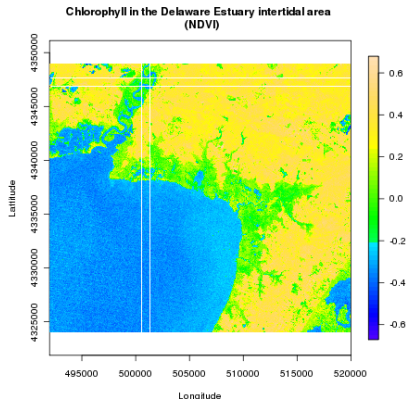
```
> depthProfile <- transect(datamatrix=z[,,"1988"],
+                           startpoint = c(62000,391500),
+                           endpoint = c(64000,394000),
+                           method="dimension")
> # cf. course for function...
> par(mfrow=c(1,2))
> image.plot(x,y,z[,,"1988"],col = c(tim.colors(),"black"))
> segments(x0=depthProfile$start[1],y0=depthProfile$start[2],
+          x1=depthProfile$end[1],
+          y1=depthProfile$end[2],lwd=2,col="white")
> plot(depthProfile$distance,depthProfile$values,
+       xlab="Distance from origine [m]",ylab="Depth [m]")
```



# Writing NetCDF files

Suppose: Landsat 7 spectral bands as ArcInfo ASCII raster files

```
> library(adehabitat)
> ETM3 <- import.asc(file="/home/tom/Desktop/ETM3.asc")
> ETM4 <- import.asc(file="/home/tom/Desktop/ETM4.asc")
> NDVI <- (ETM4-ETM3)/(ETM4+ETM3) # NDVI = (NIR - red) / (NIR + red)
> image.plot(NDVI,col=topo.colors(64),
+           main="Chlorophyll in the Delaware Estuary
+               intertidal area \n (NDVI)",
+           xlab="Longitude",ylab="Latitude")
```



# Writing NetCDF files

- We want to store this NDVI map in a NetCDF file.
- Steps to follow:
  - ▶ Create dimensions (`dim.ncdf` class)
  - ▶ Create non-coordinate variables (`var.ncdf` class)
  - ▶ Create ncdf file in memory (`create.ncdf()`)
  - ▶ Add the variable content to the NetCDF file (`put.var.ncdf()`)
  - ▶ Close the file, causing the entire buffer to be written to disk
  - ▶ Add attributes (metadata) to the file after re-opening in writeable mode

# Writing NetCDF files

- Create dimensions

```
# Define R variables with the dimension values, derived from the ESRI file  
x <- seq(attr(NDVI,"xll"),length.out=dim(NDVI)[1],by=attr(NDVI,"cellsize"))  
y <- seq(attr(NDVI,"yll"),length.out=dim(NDVI)[2],by=attr(NDVI,"cellsize"))
```

```
# Define the dimension objects (dim.ncdf), based on the R variables above  
x_dim <- dim.def.ncdf(name="x",units="m", vals=x, unlim=F, create_dimvar=T)  
y_dim <- dim.def.ncdf(name="y",units="m", vals=y, unlim=F, create_dimvar=T)
```

# Writing NetCDF files

- Create non-coordinate variables

```
# Define a 0 dimensional variable (var.ncdf)  
# Only needed to attach metadata, cf. later)  
crs_var <- var.def.ncdf.no_dim(name="crs",units="",dim=numeric(0),  
                               longname="UTM 18N",prec="integer",  
                               missval=NA)  
  
  # This is a derived function from the ncdf package  
  # see remark 2 for function  
  
# Define a NetCDF variable  
ndvi_var <- var.def.ncdf(name="ndvi",units="",dim=list(x_dim,y_dim),  
                        longname="Normalized Difference Vegetation Index",  
                        prec="double",missval=-9999)
```

# Writing NetCDF files

- Create ncdf file in memory

```
> ndvi.nc <- create.ncdf("/home/tom/Dropbox/NetCDF/CourseMaterial/Data/NDVI.nc",  
+                        vars=list(crs_var,ndvi_var),verbose=T)
```

- Add the variable content to the NetCDF file

```
> put.var.ncdf(nc=ndvi.nc, varid="ndvi", vals=NDVI,start=c(1,1),count=c(-1,-1))
```

- Close the file, allowing for the entire buffer to be written to disk

```
> close.ncdf(ndvi.nc)
```

# Writing NetCDF files

- Re-open the file to add extra attributes

```
> ndvi.nc <- open.ncdf("/home/tom/Dropbox/NetCDF/CourseMaterial/Data/NDVI.nc",  
+                       write=T)
```

- Which attributes to add?
- Different conventions and standards  
(CF convention is used frequently)
- Here we will work in three step:
  - ▶ Global attributes  
(refer to the dataset as a whole)
  - ▶ Non-coordinate variable attributes  
(refer to individual variables)
  - ▶ Information on the coordinate system  
(implemented by adding an extra variable)



# Writing NetCDF files

- Global attributes according to the CF convention  
(title, institution, source, history, references, comment)

```
att.put.ncdf(nc=ndvi.nc, varid=0, attname="title",
             attval="NDVI of the Delaware Estuary Intertidal Area")
att.put.ncdf(nc=ndvi.nc, varid=0, attname="institution", attval="NIOO-CEME")
att.put.ncdf(nc=ndvi.nc, varid=0, attname="source",
             attval="Landsat 7 Enhanced Thematic Mapper")
att.put.ncdf(nc=ndvi.nc, varid=0, attname="history",
             attval="The image cube was downloaded from the NASA website
                    and processed using GRASS 6.3")
att.put.ncdf(nc=ndvi.nc, varid=0, attname="references", attval="http://www.nasa.gov/")
att.put.ncdf(nc=ndvi.nc, varid=0, attname="comment",
             attval='The original Landsat image cube, downloaded as Erdas
                    Imagine format (.img), was imported into GRASS 6.3 using
                    the GDAL libraries for conversion. The bands 3 (red) and
                    4 (NIR; near infra-red) were selected for export as
                    ESRI ArcInfo ASCII files. These were read into the R
                    Statistical Software via the import.asc function of
                    the adehabitat package. The NDVI was calculated from
                    these bands, and exported as this NetCDF file.')
```

# Writing NetCDF files

- Information on the coordinate system
- In CF convention implemented as attributes to a non-sense variable
- Problem: ncdcf requires non-zero dimensions (bug or programming decision?)
- Flexibility  $\Leftrightarrow$  stringency in metadata provision
- Should a general-purpose package enforce extra requirements without taking into account generally applied conventions and standards?
- On the other hand: Should we leave metadata provision up to the discretion of the individual scientist?

"Adding all this extra information is time consuming and not productive" on the short term!

"I know my data ..."

# Writing NetCDF files

- Information on the coordinate system
- After some change to the some relevant ncdf functions (cf. course remarks)

```
att.put.ncdf(nc=ndvi.nc, varid="crs", attname="spatial_ref", attval='
"PROJCS["WGS 84 / UTM zone 18N",
  GEOGCS["WGS 84",
    DATUM["WGS_1984",
      SPHEROID["WGS 84",6378137,298.257223563,
        AUTHORITY["EPSG","7030"]],
      AUTHORITY["EPSG","6326"]],
    PRIMEM["Greenwich",0,
      AUTHORITY["EPSG","8901"]],
    UNIT["degree",0.01745329251994328,
      AUTHORITY["EPSG","9122"]],
    AUTHORITY["EPSG","4326"]],
  UNIT["metre",1,
    AUTHORITY["EPSG","9001"]],
  PROJECTION["Transverse_Mercator"],
  PARAMETER["latitude_of_origin",0],
  PARAMETER["central_meridian",-75],
  PARAMETER["scale_factor",0.9996],
  PARAMETER["false_easting",500000],
  PARAMETER["false_northing",0],
  AUTHORITY["EPSG","32618"],
  AXIS["Easting",EAST],
  AXIS["Northing",NORTH]]"
')
```

# Writing NetCDF files

- Information on the coordinate system
- After some change to the some relevant ncdf functions (cf. course remarks)

```
....  
    UNIT["degree",0.01745329251994328,  
        AUTHORITY["EPSG","9122"]],  
    AUTHORITY["EPSG","4326"]],  
UNIT["metre",1,  
    AUTHORITY["EPSG","9001"]],  
PROJECTION["Transverse_Mercator"],  
PARAMETER["latitude_of_origin",0],  
PARAMETER["central_meridian",-75],  
PARAMETER["scale_factor",0.9996],  
PARAMETER["false_easting",500000],  
PARAMETER["false_northing",0],  
AUTHORITY["EPSG","32618"],  
AXIS["Easting",EAST],  
AXIS["Northing",NORTH]]"  
) # nesting of ' and "  
att.put.ncdf(nc=ndvi.nc,varid="crs",  
             attname="grid_mapping_name",  
             attval="transverse_mercator")  
att.put.ncdf(nc=ndvi.nc,varid="crs",  
             attname="scale_factor_at_central_meridian",  
             attval="0.9996")  
....
```

# Writing NetCDF files

- Information on the coordinate system
- Additions to the coordinate values

```
# x coordinate
att.put.ncdf(ndvi.nc, varid="x", attname="standard_name",
             attval="projection_x_coordinate")
att.put.ncdf(ndvi.nc, varid="x", attname="_FillValue",
             attval="-9999")

# y coordinate
att.put.ncdf(ndvi.nc, varid="y", attname="standard_name",
             attval="projection_y_coordinate")
att.put.ncdf(ndvi.nc, varid="y", attname="_FillValue",
             attval="-9999")

close.ncdf(ndvi.nc)    # clear buffer and write to disk...
```

- Under certain conditions the standard\_name attribute should have a predetermined value
- \_FillValue is obligatory in the CF conventions, whereas missing\_value is abandoned as obligatory

**Thank you for your attention !**

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