

Woods Hole

## Getting data in and getting results out

Anders Nielsen

# Getting data in

- We need to get data into R first — OK, or look into?
- We further need to pass data to the cpp program
- TMB uses 'computer memory'
- In other words no need to write stuff to files on disk
- A simple example:

42

```
library(TMB)
compile("scalar.cpp")
dyn.load(dynlib("scalar"))

Y<-scan("scalar.dat")

data <- list()
data$Y <- Y

param <- list()
param$mu <- 0

obj <- MakeADFun(data, param, DLL="scalar")
opt <- nlminb(obj$par, obj$fn, obj$gr)
opt$par
```

```
#include <TMB.hpp>
template<class Type>
Type objective_function<Type>::operator()()
{
  DATA_SCALAR(Y);
  PARAMETER(mu);
  Type nll = pow(Y-mu,2);
  return nll;
}
```

# Transfers basic objects

What	R side (example)	c++ side
Number	1	DATA_SCALAR
Vector	c(1,2,3)	DATA_VECTOR
Matrix	matrix(c(1,2,3,4), nrow=2, ncol=2)	DATA_MATRIX
Array	matrix(c(1,2,3,4), nrow=2, ncol=2)	DATA_ARRAY
Integer	1	DATA_INTEGER
Integer Vector	c(1,2,3)	DATA_IVECTOR
Integer Matrix	matrix(c(1,2,3,4), nrow=2, ncol=2)	DATA_IMATRIX
Integer Array	matrix(c(1,2,3,4), nrow=2, ncol=2)	DATA_IARRAY
Factor	factor(c("a","b"))	DATA_FACTOR
String	"a"	DATA_STRING

# Dimensions set by data

```
library(TMB)
compile("size.cpp")
dyn.load(dynlib("size"))
data <- list()
data$V <- 1:3
data$M <- matrix(1:6, nrow=2, ncol=3)
data$A <- array(1:6, dim=c(1,2,3))
```

```
param <- list()
param$mu <- 0
obj <- MakeADFun(data, param)
obj$report()
```

```
#Rout> $Ad
#Rout> [1] 1 2 3
#Rout>
#Rout> $Vs
#Rout> [1] 3
#Rout>
#Rout> $Ms
#Rout> [1] 6
#Rout>
#Rout> $Mnrow
#Rout> [1] 2
#Rout>
#Rout> $As
#Rout> [1] 6
#Rout>
#Rout> $Mncol
#Rout> [1] 3
#Rout>
```

```
#include <TMB.hpp>
template<class Type>
Type objective_function<Type>::operator()()
{
  DATA_VECTOR(V);
  DATA_MATRIX(M);
  DATA_ARRAY(A);

  int Vs=V.size();
  REPORT(Vs);

  int Ms=M.size();
  REPORT(Ms);
  int Mnrow=M.rows();
  REPORT(Mnrow);
  int Mncol=M.cols();
  REPORT(Mncol);

  int As=A.size();
  REPORT(As);
  vector<int> Ad(3);
  //Ad(0)=A.dim[0]; Ad(1)=A.dim[1]; Ad(2)=A.dim[2];
  Ad=A.dim;
  REPORT(Ad);

  PARAMETER(mu);
  Type nll = pow(mu,2);
  return nll;
}
```

# Checking what is read in

- Useful to verify when developing models
- Can use `std::cout<<`, but `REPORT` is often easier:

```
library(TMB)
compile("verify.cpp")
dyn.load(dynlib("verify"))
data <- list()
data$V <- 1:3
data$M <- matrix(1:6, nrow=2, ncol=3)
data$A <- array(1:6, dim=c(1,2,3))

param <- list()
param$mu <- 0
obj <- MakeADFun(data, param)
out <- obj$report()
out$M==data$M

#Rout>      [,1] [,2] [,3]
#Rout> [1,] TRUE TRUE TRUE
#Rout> [2,] TRUE TRUE TRUE
```

```
#include <TMB.hpp>
template<class Type>
Type objective_function<Type>::operator()()
{
  DATA_VECTOR(V);
  DATA_MATRIX(M);
  DATA_ARRAY(A);
  REPORT(V);
  REPORT(M);
  REPORT(A);

  PARAMETER(mu);
  Type nll = pow(mu,2);
  return nll;
}
```

# Indexing from 0

- In  $C^{++}$  the first element is **number 0**
- Different from R, so difficult to remember in the beginning

## In R

```
data <- list()  
data$y <- c(1.1, 2.2)  
data$z <- mymatrix
```

```
y[1] ... y[n]  
z[1,1] ... z[m,n]
```

## In $C^{++}$ -part

```
DATA_VECTOR(y)  
DATA_ARRAY(z)
```

```
y(0) ... y(n-1)  
z(0,0) ... z(m-1,n-1)
```

# Exercise: No ragged arrays

- TMB has no ragged arrays.
- Consider how we would represent this data set in TMB?

```
> set.seed(123)
> ex <- lapply(c(7,9,13), function(n)round(rnorm(n),3))
> ex

[[1]]
[1] -0.560 -0.230  1.559  0.071  0.129  1.715  0.461

[[2]]
[1] -1.265 -0.687 -0.446  1.224  0.360  0.401  0.111 -0.556  1.787

[[3]]
[1]  0.498 -1.967  0.701 -0.473 -1.068 -0.218 -1.026 -0.729 -0.625 -1.687
[11]  0.838  0.153 -1.138
```

# Getting results out

- If estimated standard errors are not needed, then the
  - `REPORT(X)` in the  $C^{++}$  file
  - `obj$report()`\$X in the  $R$  file
- If estimated standard errors are needed, then use
  - `ADREPORT(X)` in the  $C^{++}$  file (for derived quantities)
  - `summary(sdreport(obj))` in the  $R$  file
- To get estimates and standard deviations in the same format as they entered the parameter list, try:
  - Parameter list: `p1 <- as.list(sdreport(obj),"Est")`
  - Parameter Sd list: `plsd <- as.list(sdreport(obj),"Std")`



# Exercise: Beverton-Holt stock recruitment model

- The Beverton-Holt model can be written (slightly re-parametrized) as:

$$\log R_i = \log(a) + \log(ssb_i) - \log(1 + \exp(\log(b))ssb_i) + \varepsilon_i \text{ where } \varepsilon_i \sim \mathcal{N}(0, \sigma^2) \text{ independent}$$

- Estimate the model parameters  $\log(a)$  and  $\log(b)$  and  $\log(\sigma)$ .
- A data set of SSB and  $\log(R)$  is found in the shared folder