

# RCaN

## Supplementary Material

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### 1 Introduction

1. Goals: an example of a RCaN study.
2. The case study: the Barents sea (main text).
3. The RCaN file has been previously built. It is joined.
4. All following commands in joined R script.
5. A first run with all main steps.
6. A second run after removing some constraints
7. Comparisons between both runs and interpretation

### 2 Preliminary: R Environment

A few libraries are to be loaded.

```
> library(RCaN) #the main package
> library(ggplot2) #to draw results
> library(coda) #to explore mcmc
> library(dplyr) #to manipulate data frame
> library(xtable) #to create latex tables
> library(xlsx) # to import excel files
```

### 3 The RCaN file

Parameters, observations and constraints have been gathered in an Excel file with a specific structure.

```
> NAMEFILE <- '/Users/christianmullon/Desktop/Ocean/BarentsSeaReconstructions_01_02_21.xlsx'
> # NAMEFILE <- '/Users/christianmullon/Desktop/Ocean/CaN_template_mini.xlsx'
```

### 3.1 Components

	Component	Inside	AssimilationE	Digestibility	OtherLosses
1	PP	0.00	0.00	0.65	0.00
2	H zoo	1.00	1.00	0.90	8.40
3	O zoo	1.00	1.00	0.90	5.50
4	Benthos	1.00	0.94	0.60	1.50
5	PelF	1.00	0.90	0.90	2.85
6	DemF	1.00	0.93	0.85	1.65
7	MM	1.00	1.00	0.00	5.50
8	Birds	1.00	0.84	0.00	60.00
9	Fisheries	0.00	0.00	0.00	0.00
10	NorSeaZoo	0.00	0.00	0.00	0.00

Table 1: Components

### 3.2 Fluxes

### 3.3 Observations

### 3.4 Constraints

	Flux	From	To	Trophic
1	PP_Hzoo	PP	Hzoo	1.00
2	PP_Ozoo	PP	Ozoo	1.00
3	PP_Benthos	PP	Benthos	1.00
4	Hzoo_Ozoo	Hzoo	Ozoo	1.00
5	Hzoo_PelF	Hzoo	PelF	1.00
6	Ozoo_Ozoo	Ozoo	Ozoo	1.00
7	Ozoo_PelF	Ozoo	PelF	1.00
8	Ozoo_DemF	Ozoo	DemF	1.00
9	Ozoo_MM	Ozoo	MM	1.00
10	Ozoo_Birds	Ozoo	Birds	1.00
11	Benthos_Benthos	Benthos	Benthos	1.00
12	Benthos_DemF	Benthos	DemF	1.00
13	PelF_PelF	PelF	PelF	1.00
14	PelF_DemF	PelF	DemF	1.00
15	PelF_MM	PelF	MM	1.00
16	PelF_Birds	PelF	Birds	1.00
17	DemF_DemF	DemF	DemF	1.00
18	DemF_MM	DemF	MM	1.00
19	NorSeaZoo_Hzoo	NorSeaZoo	Hzoo	0.00
20	NorSeaZoo_Ozoo	NorSeaZoo	Ozoo	0.00
21	PelF_Fisheries	PelF	Fisheries	0.00
22	DemF_Fisheries	DemF	Fisheries	0.00
23	MM_Fisheries	MM	Fisheries	0.00
24	Ozoo_Fisheries	Ozoo	Fisheries	0.00

Table 2: Fluxes

	Year	Prod_Sat	Hzoo_Biomass	Ozoo_Biomass	Pelagics
1	1988.00		25432.12	24275.61	428.28
2	1989.00		31987.20	16130.85	864.52
3	1990.00		23027.73	7481.54	5831.66
4	1991.00		21188.34	16833.36	7288.56
5	1992.00		27314.02	7940.31	5152.50
6	1993.00		37612.31	11880.41	799.64
7	1994.00		72438.19	22699.62	203.94
8	1995.00		57941.78	23526.60	195.66
9	1996.00		38465.04	24633.25	504.21
10	1997.00		43364.75	19153.71	912.15

Table 3: Observations

	Id	Constraint
1	C01	$PP\_H zoo + PP\_O zoo + PP\_B enthos \leq Prod\_Sat * 1.5$
2	C02	$PP\_H zoo + PP\_O zoo + PP\_B enthos \geq Prod\_Sat / 1.5$
3	C03	$PP\_H zoo + PP\_O zoo + PP\_B enthos \leq 2000000$
4	C04	$PP\_H zoo + PP\_O zoo + PP\_B enthos \geq 500000$
5	C05	$NorSeaZoo\_H zoo = 8 * 1600$
6	C06	$NorSeaZoo\_O zoo = 2 * 1600$
7	C07	$PelF\_Fisheries \geq Pel\_landings$

Table 4: Constraints

## 4 Building polytope

```
> begin <- Sys.time()
> POLYTOPE <- buildCaN(NAMEFILE)
> end <- Sys.time()
> end-begin
```

Time difference of 3.074418 mins

```
> summary(POLYTOPE)
```

	Length	Class	Mode
components_param	10	data.frame	list
species	7	-none-	character
fluxes_def	4	data.frame	list
flow	24	-none-	character
series	22	data.frame	list
ntstep	1	-none-	numeric
data_series_name	21	-none-	character
constraints	5	data.frame	list
H	49	-none-	numeric
N	168	-none-	numeric
A	2009575	dgCMatrix	S4
AAll	2009575	dgCMatrix	S4
C	49600	dgCMatrix	S4
CAll	49600	dgCMatrix	S4
v	64	-none-	numeric
vAll	64	-none-	numeric
L	173600	dgCMatrix	S4
b	2593	-none-	numeric
bAll	2593	-none-	numeric
symbolic_enviro	903	-none-	environment

## 5 Structure of polytope

The polytope is defined by two pairs of a matrix and a vector.  $F$  being the vector of all flows at all timesteps, first one  $(A, b)$  is an equality  $A.F = b$ , second one  $(C, v)$  is an equality  $C.F \leq v$ . For the Barents sea, we have: :

```
> dim(POLYTOPE$A)
```

```
[1] 2593 775
```

```
> length(POLYTOPE$b)
```

```
[1] 2593
```

```
> dim(POLYTOPE$C)
```

```
[1] 64 775
```

```
> length(POLYTOPE$v)
```

```
[1] 64
```

## 6 Checking polytope

As it is defined in the RCaN file for the Barents' sea, the polytope is non-empty and bounded:

```
> checkPolytopeStatus(POLYTOPE)
```

```
[1] "polytope ok"
```

Limits of the Barents' sea polytope in all dimensions are obtained with `getAllBoundsParam`:

```
> BOUNDS <- getAllBoundsParam(POLYTOPE)
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```

```
> summary(BOUNDS)
```

param	lowerbound	upperbound
Length:775	Min. : 0.0	Min. : 0
Class :character	1st Qu.: 0.0	1st Qu.: 2667
Mode :character	Median : 0.0	Median : 8629
	Mean : 12566.2	Mean : 311440
	3rd Qu.: 906.5	3rd Qu.: 86908
	Max. :448256.5	Max. :7983360

Function plotPolytope2DCaNmod allows seeing the polytope in the plane defined by two parameters. In its first two dimensions, for the second 1990, the Barents sea polytope dimensions appears as.

```

> fluxX <- paste(FLUXES[1,1], '[1990] ', sep="")
> fluxY <- paste(FLUXES[2,1], '[1990] ', sep="")
> plotPolytope2D(POLYTOPE, c(fluxX, fluxY))

```

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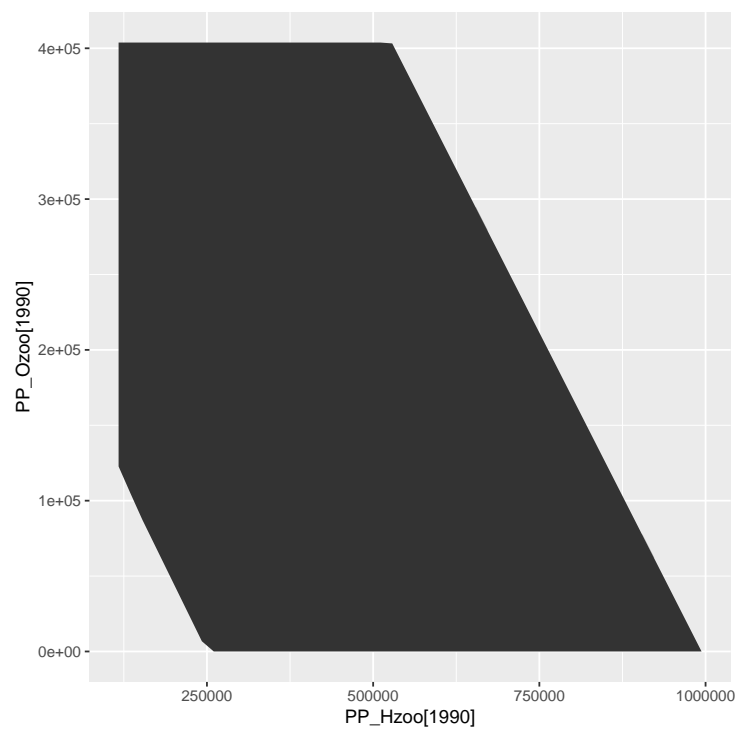
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## 7 Sampling polytope

### 7.1 Sampling

```
> begin = Sys.time()
> SAMPLE <- sampleCaN(POLYTOPE,
+                      N=100,thin=100,
+                      nchain=2,
+                      ncore=2)
> end=Sys.time()
> end-begin
```

Time difference of 16.11554 mins

## 7.2 Convergence

```
> nchain(SAMPLE$mcmc)

[1] 2

> # summary(SAMPLE$mcmc)

Gelman diagnostics

> fluxY <- paste(FLUXES[2,1], '[1990] ', sep="")
> gelman.diag(SAMPLE$mcmc[,fluxY])

Potential scale reduction factors:

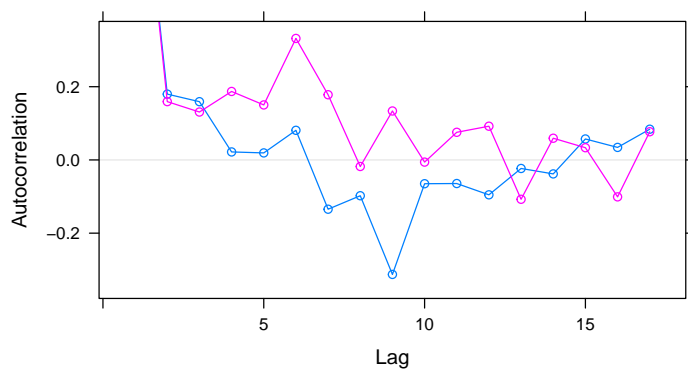
      Point est. Upper C.I.
[1,]      1.03      1.16

Autocorrelation function

> fluxZ <- paste(FLUXES[3,1], '[1990] ', sep="")
> thinned_SAMPLE <- window(SAMPLE$mcmc, thin=2)
> thin(thinned_SAMPLE)

[1] 2

> acfplot(thinned_SAMPLE[,fluxZ])
```



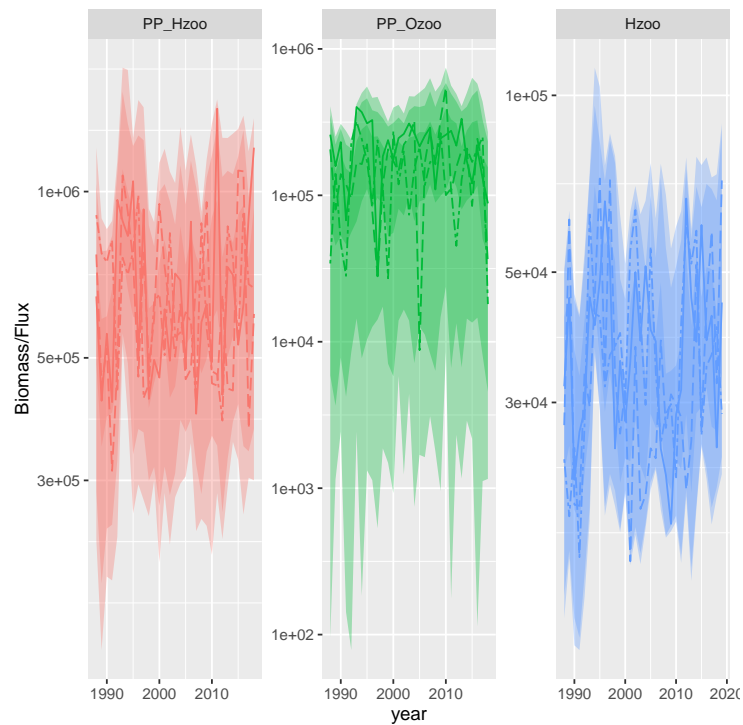
### 7.3 Dynamics

For several variables or flux, plots of sampled dynamics.

```
> fluxX <- FLUXES[1,1]
> fluxY <- FLUXES[2,1]
> compA <- COMPONENTS[2,1]
> c(fluxX,fluxY,compA)

[1] "PP_Hzoo" "PP_Ozoo" "Hzoo"

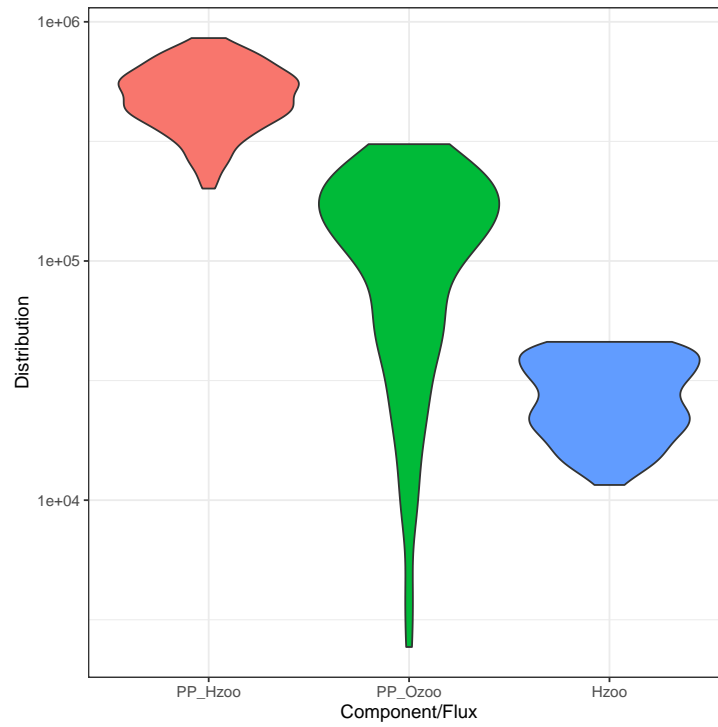
> g <- ggSeries(SAMPLE, c(fluxX,fluxY,compA), TRUE)
> g + scale_y_log10() + guides(color = FALSE, fill = FALSE)
```



## 7.4 Distribution

For a component or a flux, for a year, the distribution of sampled values.

```
> ggViolin(SAMPLE,c(fluxX,fluxY,compA),year=1990,TRUE)
```



## 7.5 Diet relationships

```
> ggDiet(SAMPLE, compA)
```

