Local and regional flood frequency analyses with BayesianMCMC

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With this document, we illustrate the use of the functions BayesianMCMC and BayesianMCMCreg. In the first section, flood frequency analysis is performed using the function BayesianMCMC with the data of the gauged station Saint Martin d'Ardèche (south-east of France), for which a lot of information on historical extreme floods is available (see Naulet, 2002). The methodology used for incorporation of historical information in BayesianMCMC is described in Payrastre et al. (2011). In the second section, a regional flood frequency analysis is performed using the function BayesianMCMCreg with data from several sites in the Ardèche region, which is exposed to frequent and severe flash floods, thus having a lot of records of extreme flood observations at different ungauged sites that were collected under the European Hydrate project (Borga et al., 2011; Gaume et al., 2009). Methodological aspects of this regional flood frequency approach are developped in Gaume et al. (2010); Nguyen et al. (2014).

1 Local flood frequency analysis with historical data

We use here systematic data and historical flood data as described respectively in Nguyen (2012) and Naulet (2002). Load the data available in the nsRFA package:

```
> library(nsRFA)
> data(Ardechedata)
> ls()
[1] "Ardeche_areas" "Ardeche_ungauged_extremes"
[3] "Beauvene_cont" "Chambonas_cont"
[5] "SaintLaurent_cont" "SaintMartin_cont"
[7] "SaintMartin_hist" "Vogue_cont"
```

The systematic data, including the maximum annual flood peaks (m^3/s) for the station Saint Martin d'Ardèche is (Table A.10, page 180 in Nguyen, 2012, catchment size 2240 km²):

> str(SaintMartin_cont)

'data.frame': 43 obs. of 2 variables: \$ year: int 1963 1964 1965 1966 1967 1968 1969 1970 1971 1972 ... \$ peak: int 3160 1930 1300 540 1050 1220 1330 3250 1500 1540 ...

At this location, a lot of information on extreme historical floods have been collected and analysed in Naulet (2002, Table 5.8, page 200; Fig. 5.30, page 208).

```
> str(SaintMartin_hist)
```

```
List of 2

$ peaks :'data.frame': 32 obs. of 2 variables:

...$ year: int [1:32] 1772 1827 1846 1853 1857 1859 1861 1878 1888 1890 ...

..$ peak: int [1:32] 6000 7400 6350 4000 4550 3950 3000 6350 3200 7550 ...

$ thresholds:'data.frame': 4 obs. of 4 variables:

...$ from.yr : num [1:4] 1645 1772 1827 1892

...$ to.yr : num [1:4] 1771 1826 1891 1962

...$ threshold: num [1:4] 7250 6000 5050 2400

...$ num.yr : num [1:4] 127 55 65 71
```

The perception thresholds in SaintMartin_hist\$thresholds indicate the dicharges that are assumed not to have been exceeded in the indicated time periods except for the historical floods in SaintMartin_hist\$peaks. The historical period has been stopped here in 1962, i.e. before the beginning of the systematic series.

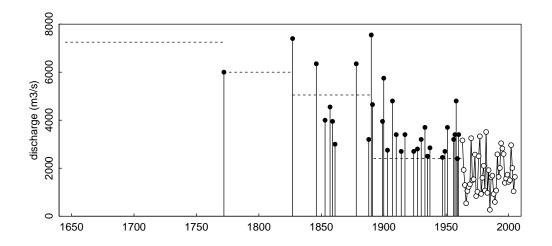
> plot(c(1640,2010), c(0, 8000), type="n", xlab="", ylab="discharge (m3/s)")

```
> points(SaintMartin_cont[,"year"], SaintMartin_cont[,"peak"], type="o", pch=21, bg="white")
```

- > points(SaintMartin_hist\$peaks[,"year"], SaintMartin_hist\$peaks[,"peak"], pch=16)
- > points(SaintMartin_hist\$peaks[,"year"], SaintMartin_hist\$peaks[,"peak"], type="h")
- > segments(x0=SaintMartin_hist\$thresholds[,"from.yr"],
- + x1=SaintMartin_hist\$thresholds[,"to.yr"],

+

```
y0=SaintMartin_hist$thresholds[,"threshold"], lty=2)
```



With the function BayesianMCMC we fit a flood frequency model to the flood data using a Bayesian Markov Chain Monte Carlo algorithm. The particularity of this function is that it takes into account historical information (see e.g. Payrastre et al., 2011), and that several perception thresholds can be specified. In order to do so, we have to organise the historical data into 3 vectors of equal length:

- xhist containing the historical flood peak values;
- seuil containing the perception thresholds associated with each historical flood in xhist;
- nbans containing the number of years covered by the perception thresholds in seuil (number of years during with the thresholds have not been exceeded except for the floods listed in xhist).

```
> xhistSM <- c(-1,SaintMartin_hist$peaks[,"peak"])</pre>
```

```
> seuilSM <- c(7250, 6000, rep(5050, 10), rep(2400, 21))
```

> nbansSM <- c(127, 55, 65, rep(0,9), 71, rep(0, 20))

```
> data.frame(xhistSM, seuilSM, nbansSM)
```

	xhistSM	seuilSM	nbansSM
1	-1	7250	127
2	6000	6000	55
3	7400	5050	65
4	6350	5050	0
5	4000	5050	0
6	4550	5050	0
7	3950	5050	0
8	3000	5050	0
9	6350	5050	0
10	3200	5050	0
11	7550	5050	0
12	4650	5050	0

13	3950	2400	71
14	5750	2400	0
15	2750	2400	0
16	4800	2400	0
17	3400	2400	0
18	2700	2400	0
19	3400	2400	0
20	2700	2400	0
21	2800	2400	0
22	3200	2400	0
23	3700	2400	0
24	2500	2400	0
25	2850	2400	0
26	2450	2400	0
27	2700	2400	0
28	3700	2400	0
29	3200	2400	0
30	3400	2400	0
31	4800	2400	0
32	2400	2400	0
33	3400	2400	0

This presentation of the dataset gives directly the correpondance between each historical flood and the associated perception threshold. Since the same threshold value is repeated several times (once for each historical flood), the application period of each threshold has to be parted in nbansSM: the only requirement is that for each threshold, when you sum up the nbansSM values next to it, the sum is the total number of years covered by the threshold. For simplicity reasons, here we just put once in nbansSM the total number of years covered by each threshold, the other values being set to 0. Finally, in this step you just have to repeat in seuilSM the value of each perception threshold as many times as peak discharge information is available during the application period of the threshold, and then indicate once in nbansSM the duration of the application period for each threshold, the other values being set to zero.

You can also notice that the first value in xhistSM is equal to -1: this is because the perception threshold defined in seuilSM has never been exceeded and no information is available on floods that did not exceed the threshold (see help).

The Bayesian MCMC algorithm is applied as follows:

```
> fit <- BayesianMCMC(xcont=SaintMartin_cont[,"peak"],
+ xhist=xhistSM,
+ seuil=seuilSM,
+ nbans=nbansSM,
+ nbpas=1000, nbchaines=3, confint=c(0.05, 0.95), dist="GEV",
+ varparameters0=c(NA, NA, 0.5))
```

Here we choose a generalized extreme value (GEV) distribution as model, with 3 Markov chains, and a confidence interval of 90% (from 5% to 95%). Note that, to have better convergence, higher values of nbpas should be used (e.g., 10000).

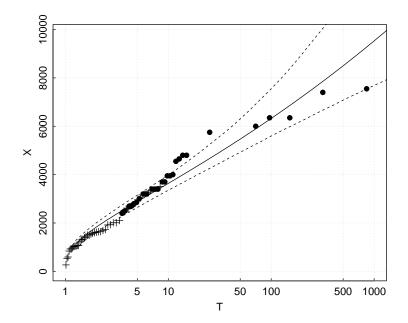
Usually you need to run it a second time to make sure the Bayesian MCMC algorithm has converged, that is what the BayesianMCMCcont function is for:

> fit <- BayesianMCMCcont(fit)</pre>

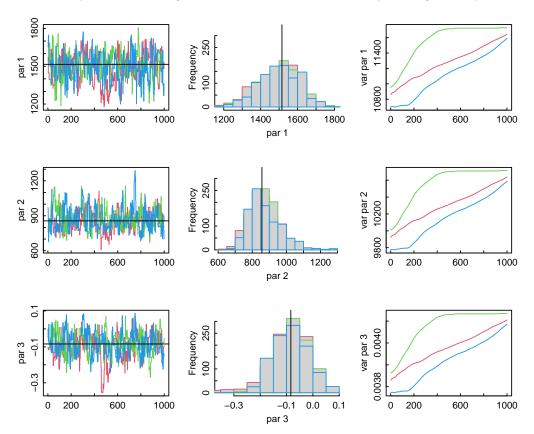
The results of the fitting and some diagnostic graphs can be plotted with the command:

> plot(fit, which=1:4, ask=TRUE)

The first plot which can be seen is the plotting position of the data, as well as the fitted distribution (with the maximum likelihood) and the confidence intervals:

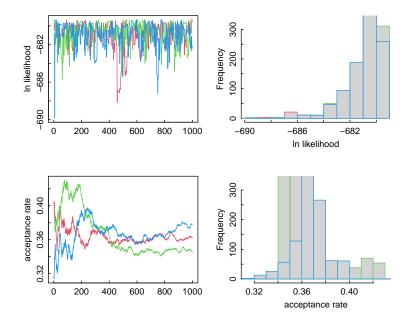


Note that the crosses are standing for the systematic data and the points for the historical data.

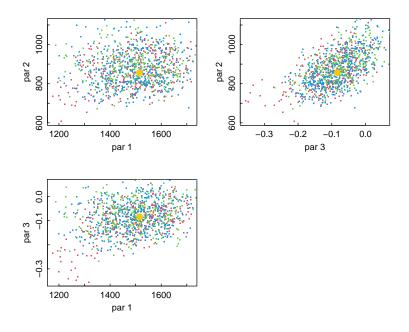


The second plot allows a diagnostic of the MCMC simulation by looking at the parameters:

The third plot is also for a diagnostic of the MCMC simulation but looking at the likelihood and MCMC acceptance rate:



Finally, the fourth plot is showing as cloud plots the posterior distribution of parameters obtained with the MCMC simulation:



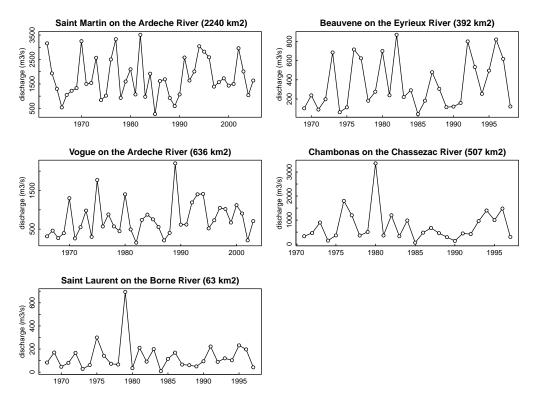
2 Regional flood frequency analysis with extreme floods in ungauged catchments

We use here data from 5 streamgauges in the Ardèche region as described in Nguyen (2012, pages 180-184).

```
> SaintMartin_cont
```

- > Vogue_cont
- > SaintLaurent_cont
- > Beauvene_cont
- > Chambonas_cont

```
plot(SaintMartin_cont[,"year"], SaintMartin_cont[,"peak"], type="o", pch=21, bg="white",
>
      xlab="", ylab="discharge (m3/s)", main=" Saint Martin on the Ardeche River (2240 km2)")
 plot(Vogue_cont[,"year"], Vogue_cont[,"peak"], type="o", pch=21, bg="white",
>
      xlab="", ylab="discharge (m3/s)", main="Vogue on the Ardeche River (636 km2)")
+
 plot(SaintLaurent_cont[,"year"], SaintLaurent_cont[,"peak"], type="o", pch=21, bg="white",
>
      xlab="", ylab="discharge (m3/s)", main="Saint Laurent on the Borne River (63 km2)")
+
 plot(Beauvene_cont[,"year"], Beauvene_cont[,"peak"], type="o", pch=21, bg="white",
>
      xlab="", ylab="discharge (m3/s)", main="Beauvene on the Eyrieux River (392 km2)")
 plot(Chambonas_cont[,"year"], Chambonas_cont[,"peak"], type="o", pch=21, bg="white",
>
       xlab="", ylab="discharge (m3/s)", main="Chambonas on the Chassezac River (507 km2)")
```



On top of that, extreme flood events have been reconstructed for several ungauged catchments, or for periods preceeding the systematic records in gauged catchments (see Table 4.2, page 127, in Nguyen, 2012).

> Ardeche_areas # km2

	station	aroa	
1	Rosieres		
-			
2	Joyeuse		
3	Vans		
4	PontRolandy		
5	Dorne		
6	Meyras	130	
7	Meysee	56	
8	Rieutord	62	
9	Burzet	47	
10	Lamastre	242	
11	Pouzin	140	
12	Chambon	11	
13	Saliouse	61	
14	PontCezar	635	
15	Chambonas	507	
16	Aubenas	480	
17	BarrageCollanges	343	
18	SaintMartin	2240	
19	Vogue	636	
20	SaintLaurent	63	
21	Beauvene	392	
> A	rdeche_ungauged_e	extre	nes # m3/s
	station	peak	associated.period
1	Rosieres	1820	50
2	Joyeuse	1000	50
3	Vans	130	50

4	PontRolandy	1150	50
5	Dorne	630	50
6	Meyras	900	50
7	Meysee	500	50
8	Rieutord	444	50
9	Burzet	350	50
10	Lamastre	970	50
11	Pouzin	700	50
12	Chambon	100	50
13	Saliouse	300	50
14	PontCezar	1500	50
15	Chambonas	3360	50
16	Aubenas	2200	50
17	BarrageCollanges	1685	50
18	SaintMartin	4500	50

> Ardeche_ungauged_extremes_new <- merge(Ardeche_ungauged_extremes, Ardeche_areas)</p>

In the following, the usage of the function BayesianMCMCreg is illustrated to provide a regional flood frequency analysis including both systematic and extreme flood data (see Gaume et al., 2010). This function uses the catchment areas to draw an index flood relationship on the form $\mu = S^{\beta}$. Therefore, the datasets have first to be organised in the following way:

- one vector **xcont** including all the systematic series;
- one vector **scont** including the catchment surfaces for each record in **xcont**;
- one vector **xhist** including the peak discharges of extreme floods;
- one vector shist including the catchment surfaces for each extreme flood in xhist;
- one vector seuil including the perception thresholds associated with each extreme flood in xhist;
- one vector **nbans** including the period of application of each perception threshold, i.e. the period during which the threshold has been exceeded only by the extreme flood in **xhist**

```
> xcont <- c(SaintMartin_cont[,2],</pre>
+
              Vogue_cont[,2],
              SaintLaurent_cont[,2],
+
+
              Beauvene_cont[,2], Chambonas_cont[,2])
  scont <- c(rep(2240, length(SaintMartin_cont[,2])),</pre>
>
              rep(636, length(Vogue_cont[,2])),
+
+
              rep(63, length(SaintLaurent_cont[,2])),
              rep(392, length(Beauvene_cont[,2])),
+
              rep(507, length(Chambonas_cont[,2])))
> xhist <- Ardeche_ungauged_extremes_new[,"peak"]</pre>
> xhist[Ardeche_ungauged_extremes_new[,"station"]=="Chambonas"] <- -1</pre>
> shist <- Ardeche_ungauged_extremes_new[,"area"]</pre>
> nbans <- Ardeche_ungauged_extremes_new[,"associated.period"]</pre>
> seuil <- Ardeche_ungauged_extremes_new[,"peak"]</pre>
```

BayesianMCMCreg can then be applied in the following way:

>	fitreg <- BayesianMCMCreg	r(xcont=xcont,	
+		<pre>scont=scont,</pre>	
+		xhist=xhist,	
+		shist=shist,	
+		nbans=nbans,	
+		seuil=seuil,	
+		<pre>nbpas=1000, nbchaines=3, confint=c(0.05)</pre>	,0.95), dist="GEV",

+ varparameters0=c(NA, NA, NA, 0.5))
> fitreg <- BayesianMCMCregcont(fitreg)
> fitreg <- BayesianMCMCregcont(fitreg)</pre>

Note that, to have better convergence, higher values of nbpas should be used (e.g., 10000).

The method assumes that, once the data are scaled by the catchment area (to the power of β), the peaks can be pooled into a single sample and the regional growth curve can be estimated using it. Several assumptions are therefore made: that the scaled samples are homogeneous, identically distributed and independent.

One good thing is that the method uses the MCMC algorithm not only for the regional growth curve parameters, but also to estimate β .

Also, the method allows to include historical data (associated with gauged series) as well as extreme discharge events available at ungauged sites, as in this example, which are treated as the historical ones. In the example, it is assumed that extreme floods observed have not been exceeded in the last 50 years (ungauged sites), or during the 50 years before the beginning of the gauged series (historical floods). Therefore, here seuil=xhist).

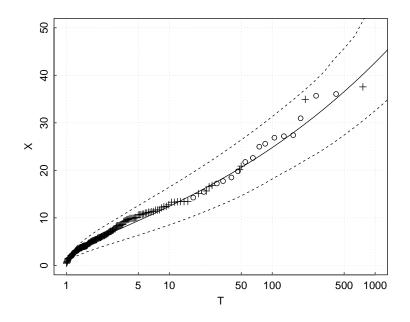
Since one of the extreme floods recorded is already included in the systematic gauged series (1980 flood at Chambonas), its value has been replaced here by -1 in xhist, and the corresponding lines in nbans and seuil are kept unchanged to indicate that this extreme flood has not been exceeded during a 50-year period preceeding the systematic series (see above).

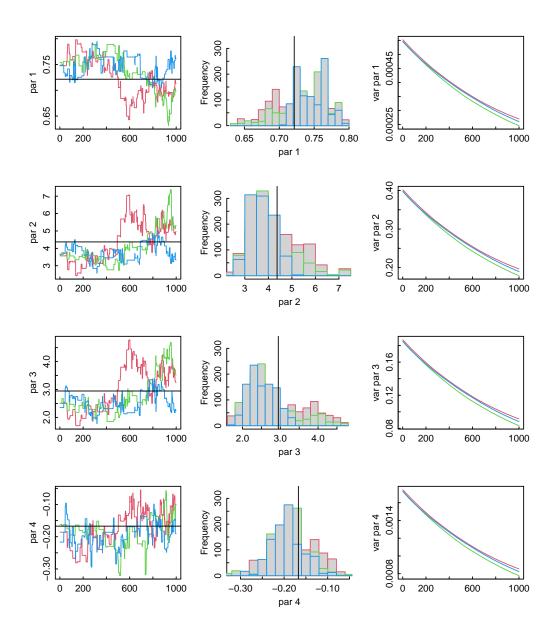
Note that once again you may need to run the Bayesian MCMC algorithm a second time to make sure it converged, this time with the function BayesianMCMCregcont.

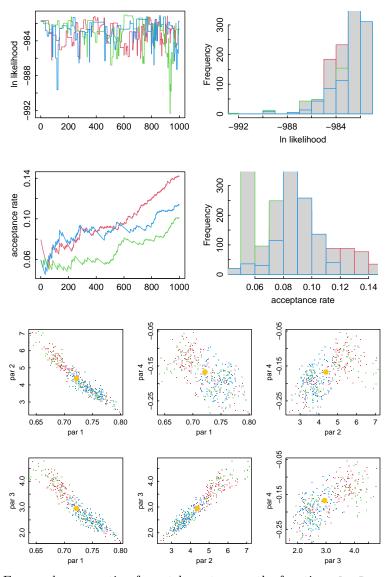
The same graphics, as the ones plotted for the local analysis, can be plotted here.

> plot(fitreg, which=1:4, ask=TRUE)

The first one refers to the regional growth curve, valid for an ungauged site with area 1 km^2 :

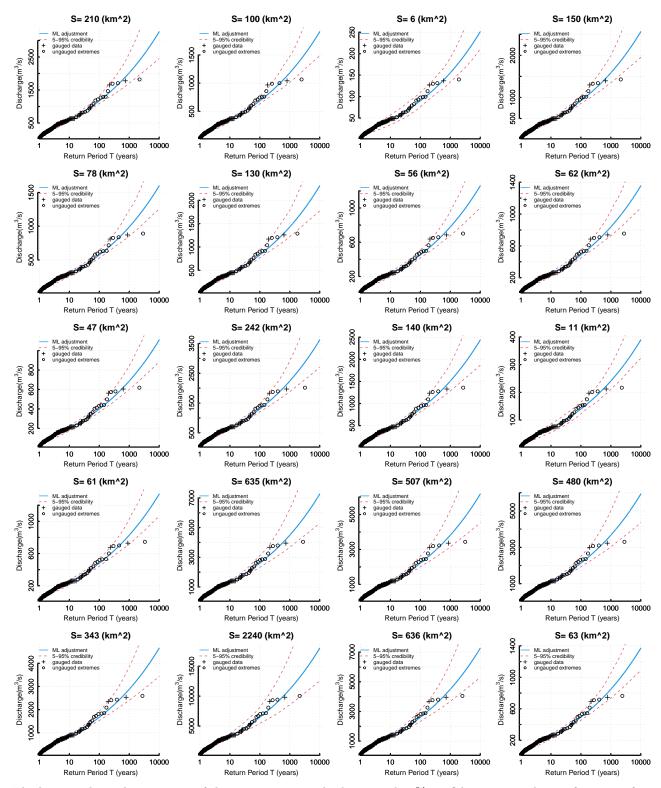






For graphs accounting for catchment areas, the function plotBayesianMCMCreg_surf can be used. Here we will do it for every catchment surface of the gauged stations:

> plotBayesianMCMCreg_surf(fitreg, surf=unique(Ardeche_areas\$area))



The last graph produces means of the previous mean discharge and 90% confidence intervals as a function of the surface. It efficiently shows how the surfaces of the chosen catchments have an influence on the algorithm. For more information about how the regional version works, you can see the PhD thesis Nguyen (2012), and Nguyen et al. (2014).

References

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