

How to use the R code that accompanies “Quantifying sources of uncertainty in projections of future climate”

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1 Required software

R is a free software environment for statistical computing and graphics. It can be installed from [www.http://cran.r-project.org/](http://cran.r-project.org/).

Our code uses the R packages `lme4` (Bates and Maechler, 2009) <http://cran.r-project.org/web/packages/lme4/index.html> and `arm` (Gelman et al., 2010) <http://cran.r-project.org/web/packages/arm/index.html>. The latter uses the package `R2WinBUGS` (Sturtz et al., 2005) <http://cran.r-project.org/web/packages/R2WinBUGS/index.html> to run WinBUGS from R. The WinBUGS software performs Bayesian analyses using Markov chain Monte Carlo (MCMC) methods. It can be installed from <http://www.mrc-bsu.cam.ac.uk/bugs/winbugs/contents.shtml>.

The R commands used below are given in the file `examplecode.R`.

2 Getting started

The R package `arm` and WinBUGS must be installed first. Then we

- load the `arm` package (which automatically loads the `lm4` package) and `R2WinBUGS` packages;
- set the working directory to one containing the files `examplecode.fns` and `superandfinite.bug` and the data files `tempdata.Rdata` and `raindata.Rdata`;
- read in the functions contained in `examplecode.fns`.

```
> library(arm)
> library(R2WinBUGS)
> setwd("Z:/CLIMATE/IMSC/FOR_REC")
> # input functions MCMC.half.Cauchy.finite(), two.way.sim() and REML.results()
> source("examplecode.fns")
```

3 Analyses of temperature indices

We load and inspect the temperature data. The value of the index is named `y`.

```
> load(file="tempdata.Rdata") # load the temperature data
> summary(all.temp.data) # see which variables are present
```

	y	scenario	GCM	run	region	period
Min.	:0.1133	A1B:2438	ncar_ccsm3_0 : 920	run1 :2806	Min. : 0	Min. :1.0
1st Qu.	:1.1145	A2 :1656	cccma_cgcm3_1 : 690	run2 :1104	1st Qu.: 5	1st Qu.:1.0

Median :1.6957	B1 :2024	mri_cgcm2_3_2a : 690	run3 :1012	Median :11	Median :1.5
Mean :1.9993		mpi_echam5 : 460	run4 : 460	Mean :11	Mean :1.5
3rd Qu.:2.5819		miroc3_2_medres: 414	run5 : 460	3rd Qu.:17	3rd Qu.:2.0
Max. :8.5460		miub_echo_g : 414	run6 : 92	Max. :22	Max. :2.0
		(Other) :2530	(Other): 184		

```
> all.temp.data[1:10,] # look at first 10 rows of data
```

	y	scenario	GCM	run	region	period
1	0.8937861	A1B	bccr_bcm2_0	run1	0	1
2	1.1928243	A1B	cccma_cgcm3_1	run1	0	1
3	1.2127593	A1B	cccma_cgcm3_1	run2	0	1
4	1.2114399	A1B	cccma_cgcm3_1	run3	0	1
5	1.1877431	A1B	cccma_cgcm3_1	run4	0	1
6	1.2191221	A1B	cccma_cgcm3_1	run5	0	1
7	1.4497154	A1B	cccma_cgcm3_1_t63	run1	0	1
8	1.2101686	A1B	cnrm_cm3	run1	0	1
9	0.6423304	A1B	csiro_mk3_0	run1	0	1
10	1.2490990	A1B	csiro_mk3_5	run1	0	1

We select the period (1 for 2020–2049, 2 for 2069–2098) and region (0 for global, 1 for region 1 etc.) and extract the data for this combination. Here we analyse global temperature indices for period 1 (2020–2049).

```
> which.period <- 1 # 1 for 2020-2049, 2 for 2069-2098
> which.reg <- 0 # 0 for global, 1 for region 1 etc.
> cat(paste("region",which.reg,", period",which.period),"\n")
```

```
region 0 , period 1
```

```
> cond <- which(all.temp.data$region==which.reg & all.temp.data$period==which.period)
> temp <- all.temp.data[cond,] # extract relevant data
```

3.1 Analysis using REML

We use the function `lmer` in the package `lme4`. The function `REML.results` extracts the REML estimates of the fixed effect μ and the random effects $\sigma_G, \sigma_S, \sigma_{GS}$ and σ_R . Setting the argument `CI` to `TRUE` means that simulation is used to estimate confidence intervals for the parameters. The argument `conf` (default 95) gives the desired confidence level and `n.sim` (default 1000) the number of simulations on the which the intervals are based. After every hundredth simulation the simulation number is printed to the screen. There is an option (`my.seed`) to specify a user-defined random number seed.

Firstly, we calcute estimates only (no confidence intervals).

```
> fit.REML <- lmer(y~1+ (1|scenario)+(1|GCM)+(1|factor(scenario):factor(GCM)), data=temp, REML=T)
> # no confidence intervals (CI=F)
> REML.res <- REML.results(fit.REML,x1=temp$GCM,x2=temp$scenario,CI=F)
> REML.res
```

mu	sigma_G	sigma_S	sigma_GS	sigma_R
1.08454952	0.23077883	0.09655014	0.03948617	0.04533721

Now, we add estimated 95% confidence intervals.

```
> # conf% confidence intervals (CI=T, n.sim simulations, my.seed=(optional) random number seed)
> # The simulation number is printed to the screen (every 100 iterations).
> # With n.sim=1000, for example, this will take a few minutes
> REML.res <- REML.results(fit.REML,x1=temp$GCM,x2=temp$scenario,CI=TRUE,conf=95,n.sim=10,my.seed=1)
> REML.res
```

	estimate	SE	lower limit	upper limit
mu	1.08454952	0.051410567	0.97722435	1.11217234
sigma_G	0.23077883	0.039885309	0.17331124	0.30127709
sigma_S	0.09655014	0.048361379	0.01591441	0.16051255
sigma_GS	0.03948617	0.008523832	0.02486251	0.05082362
sigma_R	0.04533721	0.002537944	0.04336055	0.05123868

3.2 Bayesian analyses with half-Cauchy weakly-informative priors

Firstly, set the scale parameter of the half-Cauchy priors for super-population standard deviations.

```
> if (which.period==1) my.A <- 0.5 # set scale parameter of half-Cauchy prior for SDs
> if (which.period==2) my.A <- 1
```

We use the function `bugs()` in the package `R2WinBUGS` to draw an approximate sample from the posterior distribution of the parameters using MCMC. This `bugs` function is called within the function `MCMC.half.Cauchy.centre`. Use

```
> help(bugs)
```

to see what are the arguments to, and outputs from, `bugs`.

Note in particular that `R2WinBUGS` needs to know where `WinBUGS` is installed on your computer. By default it will look in somewhere like `"C:/Program Files/WinBUGS14/"` but if `WinBUGS` does not exist in this location then the code will fail with error messages that include `"WinBUGS executable does not exist in c:/Program Files/WinBUGS14/"`.

You can tell `R2WinBUGS` where to look for `WinBUGS` using the `bugs.directory` argument to `bugs()`. The code is set up so that you can pass this argument to `bugs()` via `MCMC.half.Cauchy.finite()`. First we define an object containing the location of `WinBUGS` on the computer.

```
> my.bugs <- "C:/Users/paul/Documents/WinBUGS14"
```

Then in the calls to `MCMC.half.Cauchy.finite()` below we add the argument `bugs.directory=my.bugs`. Removing this argument will mean that `bugs()` will look for `WinBUGS` in the default location.

Here we simulate 5 independent Markov chains of length 2500 observations each and discard the first 500 observations from each chain. We use all the $5 \times 2000 = 10000$ observations to estimate the posterior distribution of the parameters (the fixed effect μ and the super-population standard deviations and finite-population standard deviations. When `MCMC.half.Cauchy.centre` is called a `WinBUGS` window will open and close when the calculations finish.

```
> #----- Super-population and finite-population standard deviations -----#
> set.seed(200)
> my.sim.fin <- MCMC.half.Cauchy.finite(temp,A=my.A,n.iter=2500,n.chains=5,n.burnin=500,
+   n.thin=1,bugs.directory=my.bugs)
> Bayes.res.fin <- my.sim.fin$summary[-10,]
> row.names(Bayes.res.fin) <- c("mu","sigma_G","sigma_S","sigma_GS","sigma_R",
+   "s_G","s_S","s_GS","s_R")
> print(round(Bayes.res.fin,3)) # summary of the results
```

	mean	sd	2.5%	25%	50%	75%	97.5%	Rhat	n.eff
mu	1.091	0.176	0.754	1.022	1.090	1.160	1.422	1.005	3200
sigma_G	0.239	0.038	0.177	0.212	0.234	0.261	0.327	1.012	270
sigma_S	0.217	0.204	0.058	0.105	0.156	0.254	0.723	1.010	340
sigma_GS	0.041	0.009	0.025	0.035	0.040	0.047	0.060	1.011	310
sigma_R	0.046	0.004	0.040	0.044	0.046	0.049	0.055	1.005	670
s_G	0.231	0.009	0.214	0.225	0.231	0.236	0.248	1.002	2200

```
s_S      0.097 0.009 0.079 0.091 0.097 0.103 0.114 1.004 990
s_GS     0.040 0.008 0.026 0.035 0.040 0.045 0.056 1.009 350
s_R      0.046 0.002 0.042 0.044 0.046 0.047 0.051 1.004 950
```

```
> #summary(my.sim.fin)           # my.sim contains lots of other stuff!
```

This is a summary of the posterior sample mean, standard deviation and various quantiles for each of the parameters. `Rhat` is a standard diagnostic for MCMC convergence. As a rough guide values of less than 1.1 suggest that approximate convergence has been attained. `n.eff` is the effective sample size on which the estimated posterior distribution of the parameter is based.

4 Analysis of precipitation data

We repeat the Bayesian analysis the global precipitation indices for period 1 (2020–2049).

Firstly, we set up the data and select the region and time period.

```
> #===== Precipitation analyses (global or regional) =====#
> load(file="raindata.Rdata") # load the rainfall data
> summary(all.rain.data)
```

y	scenario	GCM	run	region	period
Min. : -40.8427	A1B:2254	ncar_ccsm3_0 : 920	run1 : 2714	Min. : 0	Min. : 1.0
1st Qu.: -0.2442	A2 :1656	cccma_cgcm3_1 : 690	run2 : 1104	1st Qu.: 5	1st Qu.: 1.0
Median : 3.0333	B1 :2024	mri_cgcm2_3_2a : 690	run3 : 966	Median : 11	Median : 1.5
Mean : 3.6342		mpi_echam5 : 460	run4 : 460	Mean : 11	Mean : 1.5
3rd Qu.: 7.0937		miroc3_2_medres : 414	run5 : 414	3rd Qu.: 17	3rd Qu.: 2.0
Max. : 37.8825		miub_echo_g : 414	run6 : 92	Max. : 22	Max. : 2.0
		(Other) : 2346	(Other): 184		

```
> which.period <- 1           # 1 for 2020-2049, 2 for 2069-2098
> which.reg <- 0              # 0 for global, 1 for region 1 etc.
> cat(paste("region",which.reg,", period",which.period),"\n")
```

region 0 , period 1

```
> cond <- which(all.rain.data$region==which.reg & all.rain.data$period==which.period)
> rain <- all.rain.data[cond,] # extract relevant data
> #----- Bayesian analysis -----#
>
> if (which.period==1) my.A <- 2.5 # set scale parameter of half-Cauchy prior for SDs
> if (which.period==2) my.A <- 5
>
> #----- Super-population standard deviations only -----#
```

Now, use MCMC as before.

```
> set.seed(200)
> my.sim.fin.prec <- MCMC.half.Cauchy.finite(rain,A=my.A,n.iter=2500,n.chains=5,n.burnin=500,
+ n.thin=1,bugs.directory=my.bugs)
> Bayes.prec.res.fin <- my.sim.fin.prec$summary[-10,]
> row.names(Bayes.prec.res.fin) <- c("mu","sigma_G","sigma_S","sigma_GS","sigma_R",
+ "s_G","s_S","s_GS","s_R")
> print(round(Bayes.prec.res.fin,3)) # summary of the results
```

	mean	sd	2.5%	25%	50%	75%	97.5%	Rhat	n.eff
mu	1.543	0.394	0.840	1.387	1.534	1.700	2.289	1.029	480
sigma_G	0.647	0.109	0.473	0.571	0.634	0.709	0.897	1.007	510
sigma_S	0.402	0.529	0.060	0.138	0.237	0.467	1.679	1.023	140
sigma_GS	0.199	0.032	0.144	0.176	0.197	0.218	0.269	1.006	600
sigma_R	0.111	0.010	0.094	0.105	0.111	0.117	0.132	1.004	1100
s_G	0.615	0.031	0.552	0.595	0.616	0.637	0.676	1.003	1600
s_S	0.124	0.035	0.055	0.101	0.124	0.147	0.191	1.016	240
s_GS	0.194	0.025	0.149	0.177	0.192	0.210	0.248	1.005	1000
s_R	0.110	0.007	0.099	0.105	0.109	0.114	0.125	1.002	2200

References

- Bates, D. and M. Maechler (2009). *lme4: Linear mixed-effects models using Eigen and Eigen++*. R package version 0.999375-32.
- Gelman, A., Y.-S. Su, M. Yajima, J. Hill, M. G. Pittau, J. Kerman, and T. Zheng (2010). *arm: Data Analysis Using Regression and Multilevel/Hierarchical Models*. R package version 1.3-02.
- Sturtz, S., U. Ligges, and A. Gelman (2005). R2WinBUGS: A package for running WinBUGS from R. *Journal of Statistical Software* 12(3), 1–16.