

WinBUGS : part 2

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Agenda

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- ▶ Hierarchical model: linear regression example
- ▶ R2WinBUGS



Linear Regression

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- ➊ Bayesian linear regression model :

- ➌ Likelihood

$$y_i = \alpha + \beta x_i + \varepsilon_i, \quad \varepsilon_i \sim N(0, \sigma^2)$$



$$y_i \sim N(\alpha + \beta x_i, \sigma^2) \quad for \quad i = 1, \dots, n$$

- ➌ Prior (non-informative):

$$\alpha \sim N(0, 10^4)$$

$$\beta \sim N(0, 10^4)$$

$$\tau \sim \text{Gamma}(0.0001, 0.0001) \quad \text{with } \tau = 1/\sigma^2$$



Linear Regression

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④ In WinBUGS :

```
model {  
  for (i in 1:n){  
    y[i]~dnorm(mu[i],tau)  
    mu[i] <- alpha + beta * x[i]  
  }  
}
```

- **Prior :**

```
alpha~dnorm(0, 0.0001)  
beta ~ dnorm (0, 0.0001)  
tau ~ dgamma (0.0001, 0.0001) }
```



Hierarchical model

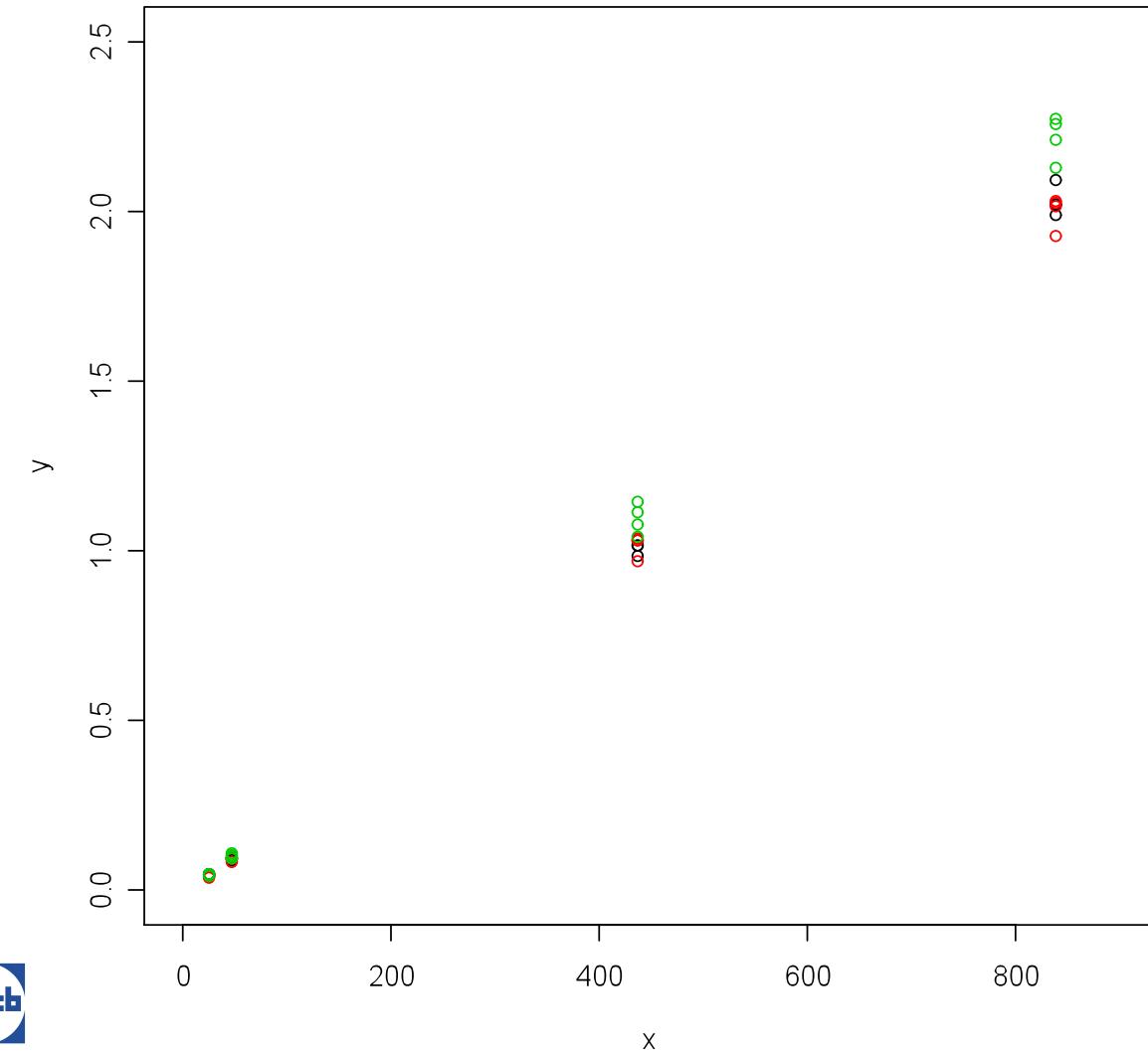
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- ◉ Calibration experiment:
- ◉ A new calibration curve is established every day
 - 3 DAYS of calibration
 - 4 levels of concentrations
 - 4 repetitions by level
- ◉ Linear calibration curve
- ◉ Calibration curve (intercept and slope) will slightly vary from day to day.



Hierarchical model

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**3 days for
calibration**



Hierarchical model

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- ➊ Data :Practical exercices\Hierarchical reg\data_orig.xls



Hierarchical model

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- ➊ Bayesian hierarchical linear regression model :

$$y_{ij} = \alpha_j + \beta_j x_i + \varepsilon_{ij}$$

$$\alpha_j \sim N(\alpha_{mean}, \sigma_\alpha^2)$$

$$\beta_j \sim N(\beta_{mean}, \sigma_\beta^2)$$

$$\varepsilon_{ij} \sim N(0, \sigma^2)$$

for $i=1, \dots, n$ and $j=1, \dots, m$

n observations per day
m days



Hierarchical model

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► Bayesian hierarchical linear regression model :

- Prior (non-informative):

$$\alpha_{\text{mean}} \sim N(0, 10000)$$

$$\beta_{\text{mean}} \sim N(0, 10000)$$

$$\tau_\alpha \sim \text{Gamma}(1, 0.001) \quad \text{with } \tau_\alpha = 1/\sigma^2_\alpha$$

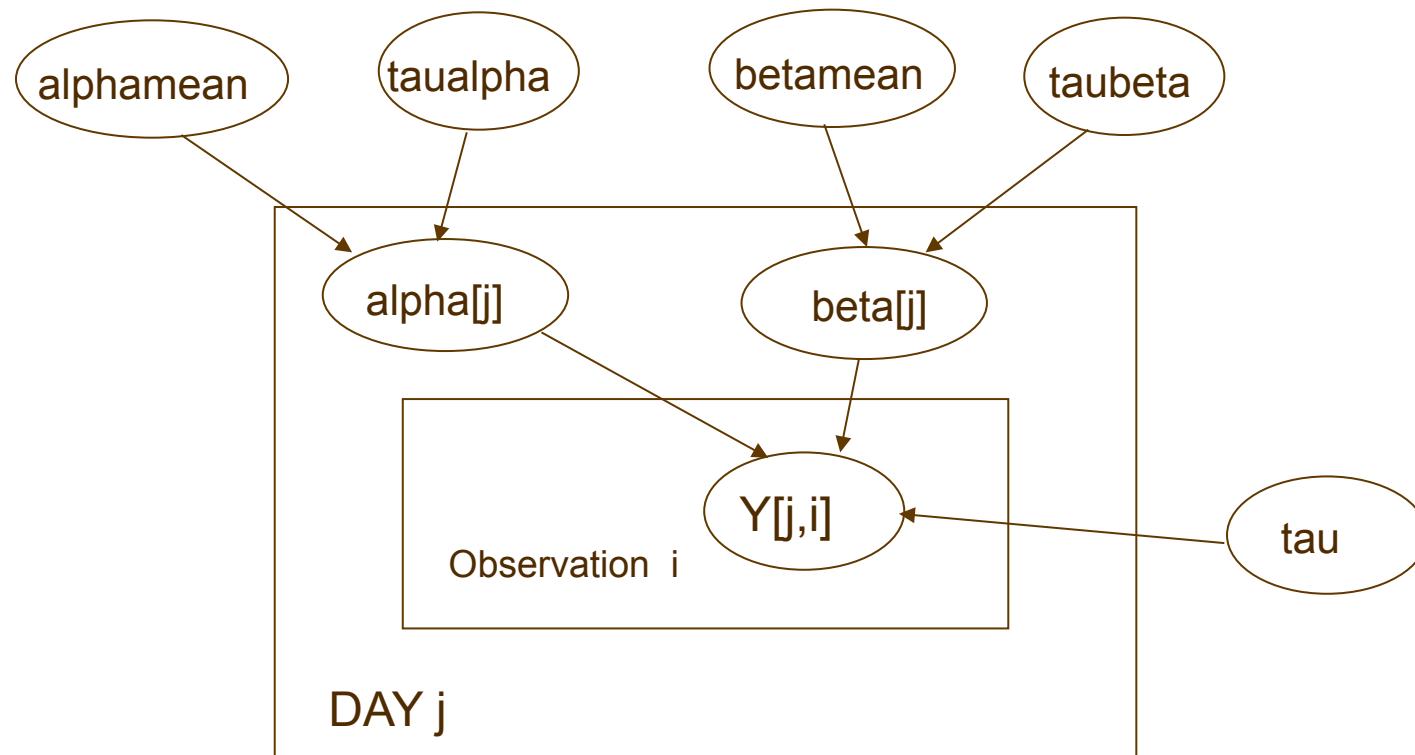
$$\tau_\beta \sim \text{Gamma}(1, 0.001) \quad \text{with } \tau_\beta = 1/\sigma^2_\beta$$

$$\tau \sim \text{Gamma}(1, 0.001) \quad \text{with } \tau = 1/\sigma^2$$



Graphical illustration

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In WinBUGS

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WinBUGS model

```
model{  
  for(i in 1:n){  
    y[i]~dnorm(mu[i], tau)  
    mu[i]<-alpha[serie[i]] + (beta[serie[i]]*(x[i]-mean(x[])))  
  }  
}  
  
for(j in 1:J){  
  alpha[j]~dnorm(alphamean, taualpha)  
  beta[j]~dnorm(betamean, taubeta) }
```

Loop on the observations

Residual variance

Individual parameters normally distributed around the population mean with precision taualpha/taubeta

```
alphamean~dnorm(0, 0.0001)  
betamean~dnorm(0, 0.0001)
```

Prior for the mean parameters

```
tau~dgamma(1,0.001)
```

Prior for the residual variability

```
taualpha~dgamma(1,0.001)  
taubeta~dgamma(1,0.001)
```

Prior for the “inter-day variability”

```
sigma<-1/sqrt(tau)  
sigmaalpha<-1/sqrt(taualpha)  
sigmabeta<-1/sqrt(taubeta)}
```

Derive parameters

Model

Prior



Exercice

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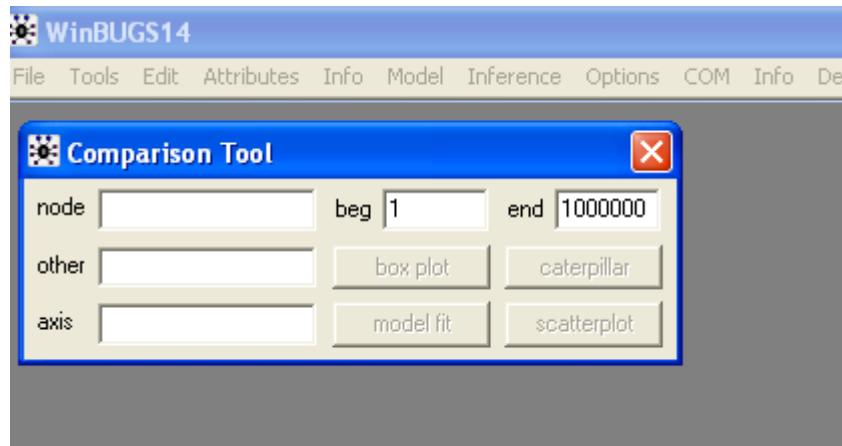
- ▶ Open data.txt, inits1.txt, inits2.txt (and model.txt)
- ▶ Run the model in WinBUGS with 1000 iterations for burnin, 5000 iterations for inference
- ▶ Monitor the parameters:
 - alpha, beta,
 - alphamean,betamean,
 - taualpha,taubeta,
 - mu,
 - tau



Exercice

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- To check the fit for serie 1:
 - Go into: *Inference -> compare*

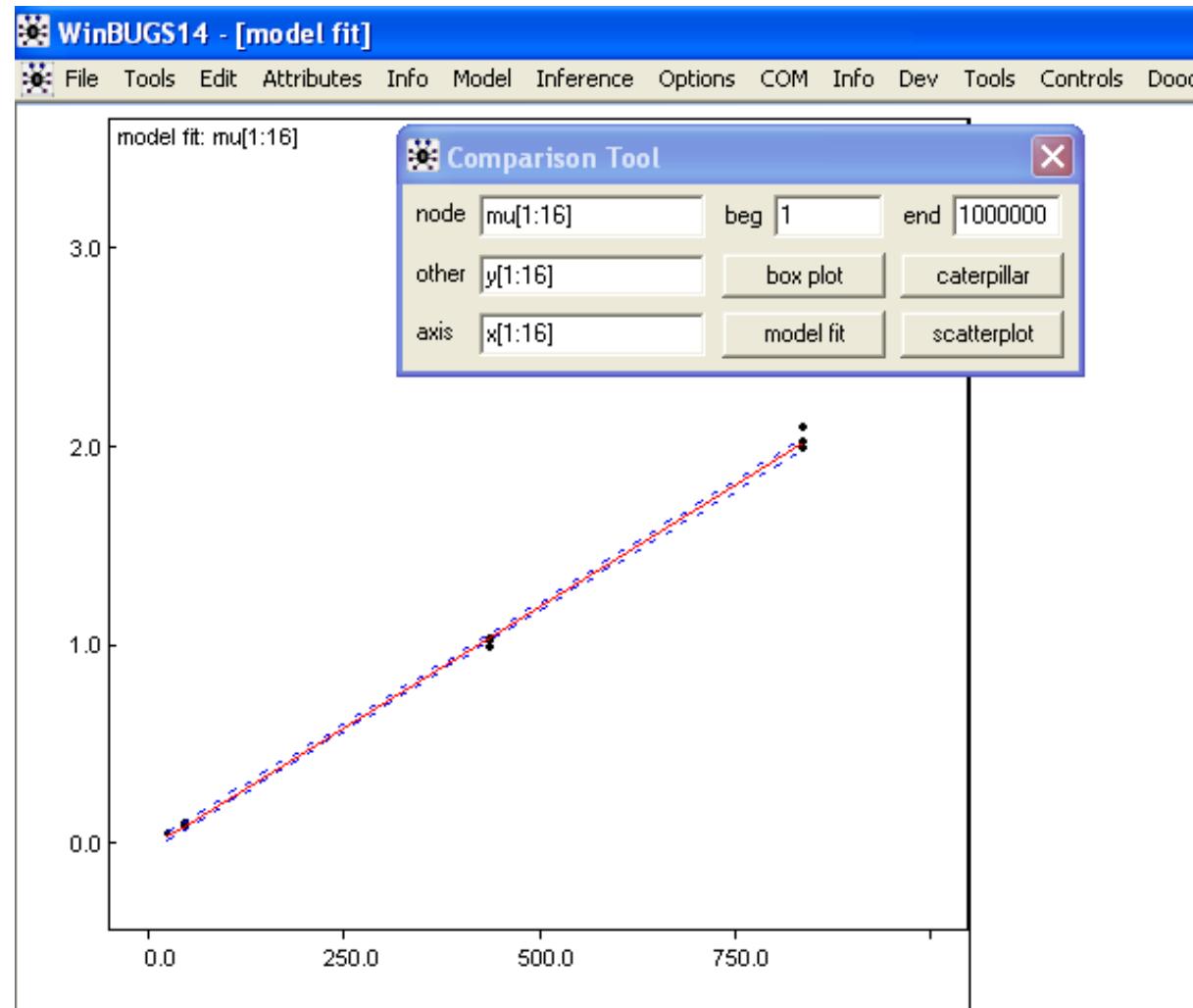


- In node: $\mu[1:16]$
- In other: $y[1:16]$
- In axis: $x[1:16]$
- Click on "model fit"



Exercice

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R2WinBUGS



Gabriele,
living with rheumatoid arthritis



R2WinBUGS: presentation

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► Drawbacks with WinBUGS:

- You have to write the data and initial values.
- You have to specify the parameters to be monitored in each run.
- The outputs are standards.

► Interesting to save the output and read it into R for further analyses :

- R2WinBUGS allows WinBUGS to be run from R
- Possibility to have the results of the MCMC and work from them (plot, convergence diagnostics...)



R2WinBUGS: steps

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① 1- Create a .txt file

- The model is saved in a .txt file :

```
model{
  for(i in 1:n){
    y[i]~dnorm(mu[i], tau)
    mu[i]<-alpha[serie[i]] + (beta[serie[i]]*(x[i]-mean(x[])))}

    for(j in 1:J){
      alpha[j]~dnorm(alphamean, taualpha)
      beta[j]~dnorm(betamean, taubeta) }

  alphamean~dnorm(0, 0.0001)
  betamean~dnorm(0, 0.0001)

  tau~dgamma(1,0.001)
  taualpha~dgamma(1,0.001)
  taubeta~dgamma(1,0.001)

  sigma<-1/sqrt(tau)
  sigmaalpha<-1/sqrt(taualpha)
  sigmabeta<-1/sqrt(taubeta)}
```



R2WinBUGS: steps

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- ④ 2- In R, the working directory is the one where the model is saved :

```
setwd("C:\\BAYES2010\\Exercices\\WinBUGS_Part2\\\nExercice")
```



R2WinBUGS: steps

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- ➊ 3- Load the R2WinBUGS packages :

```
library(R2WinBUGS)
```

- ➋ 4- Create the data for WinBUGS :

```
donnee=read.table("data_orig.txt",header=TRUE)
```

```
x=donnee$concentration
```

```
y=donnee$resp
```

```
serie=donnee$serie
```

```
n=length(y)
```

```
data <- list(n=n,J=3, x=x, y=y,serie=serie)
```



R2WinBUGS: steps

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④ 5- Load the initials values in a list :

- One list for each set of initial values
- “One list of the lists”

```
inits1=list(alphamean=0,betamean=1,tau=1,alpha=rep(0,3),beta=
  rep(1,3),taualpha=1,taubeta=1)
inits2=list(alphamean=0,betamean=1,tau=0.1,alpha=rep
  (0,3),beta=rep(0.5,3),taualpha=10,taubeta=10)
inits=list(inits1,inits2)
```

⑤ 6- Specify the parameters to monitor in a list.

```
parameters=list("alpha","beta","tau","alphamean","betamean",
  "taualpha","taubeta")
```



R2WinBUGS: steps

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- ④ 7- Create a bugs object called “sims1”

```
sims1<-bugs(data=data,inits=inits,parameters=parameters,  
model.file="model.txt",n.chains=2,n.iter=5000,n.burnin=  
1000)
```



Outputs



Gabriele,
living with rheumatoid arthritis



R2WinBUGS: outputs

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► `names(sims1):`

- "n.chains" "n.iter" "n.burnin"
- "sims.array" "sims.list" "sims.matrix" : all the iterations
- "summary" "mean" "sd" "median"
- "pD" "DIC"



R2WinBUGS: get the chains

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- ④ 3 ways to get the chains :

```
dim(sims1$sims.array)
```

```
[1] 4000 2 12
```

```
dim(sims1$sims.matrix)
```

```
[1] 8000 12
```

```
names(sims1$sims.list)
```

```
"alpha" "beta" "tau" "alphamean" "betamean"  
"taualpha" "taubeta" "deviance"
```



R2WinBUGS: Get the chains of the different parameters

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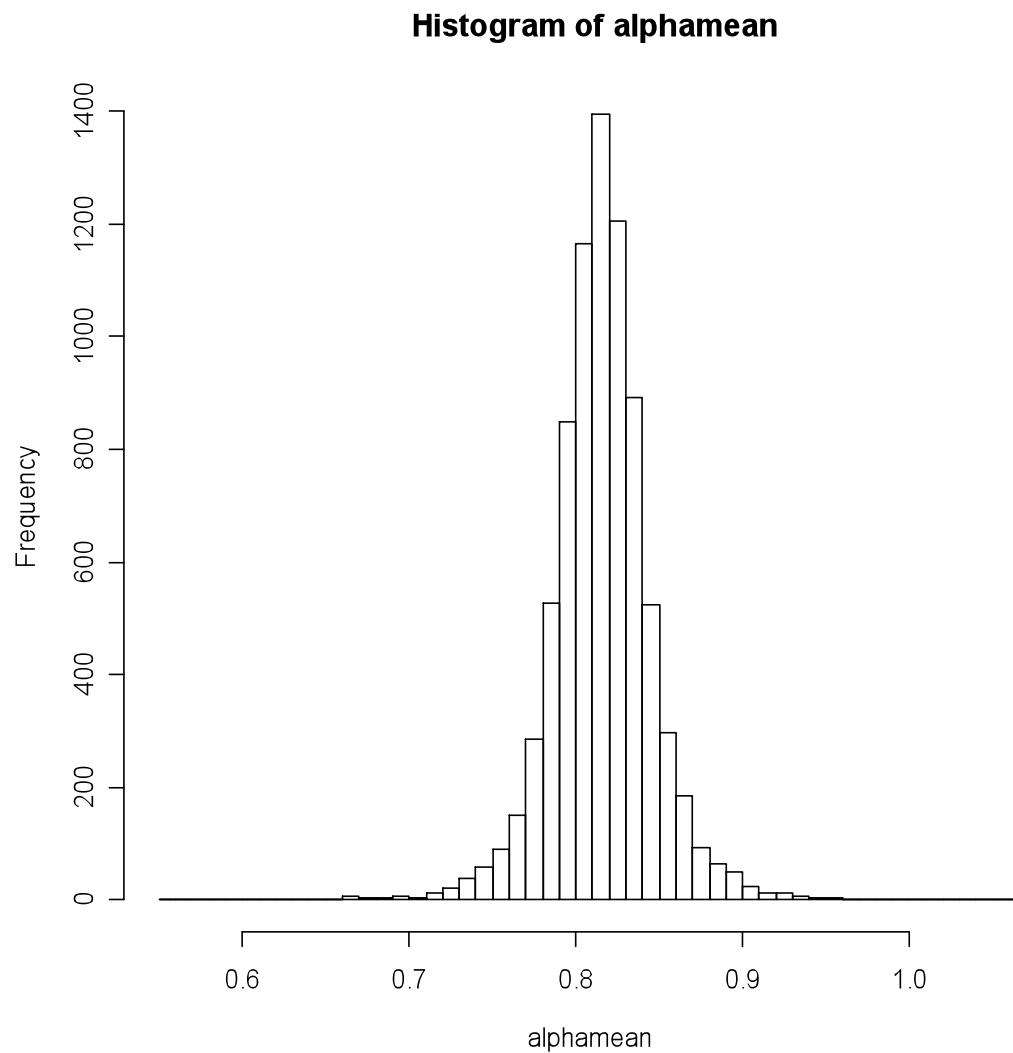
```
alphamean<-sims1$sims.list$alphamean  
betamean<-sims1$sims.list$betamean  
alpha<-sims1$sims.list$alpha  
beta<-sims1$sims.list$beta  
tau<-sims1$sims.list$tau
```



R2WinBUGS: Histograms

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`hist(alphamean)`



R2WinBUGS: Draw some traces

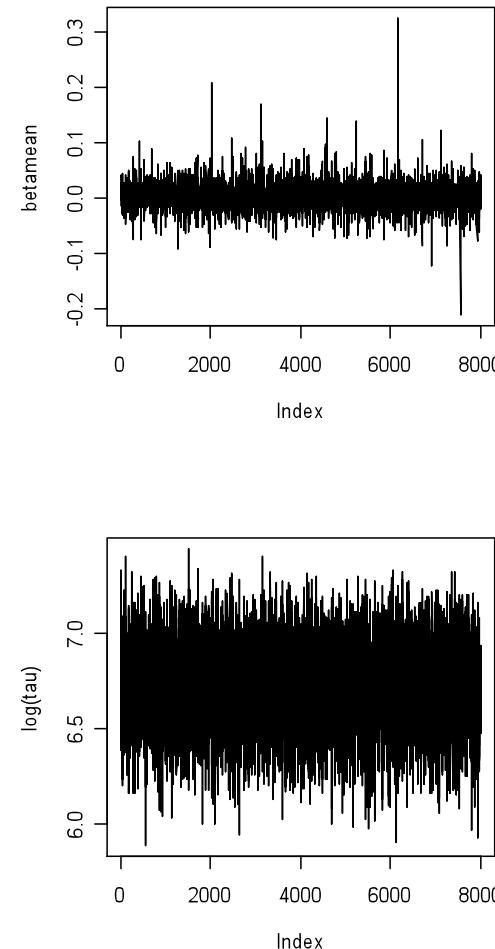
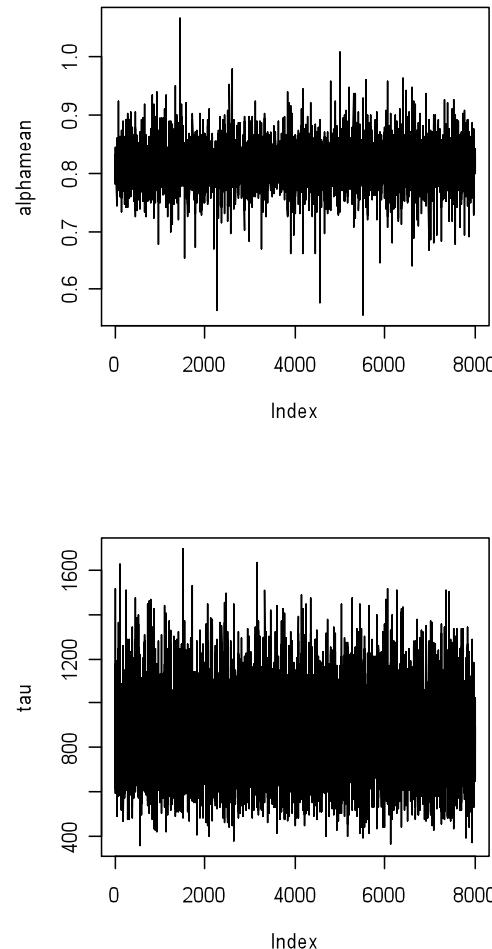
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```
par(mfrow=c(2,2))
plot(alphamean,type='l')
plot(betamean,type='l')
plot(tau,type='l')
plot(log(tau),type='l')
```



R2WinBUGS: Draw the traces

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R2WinBUGS: fitted curves

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```
#Compute the median a posteriori  
alphaest<-apply(alpha,2,function(x){quantile(x,0.5)})  
betaest<-apply(beta,2,function(x){quantile(x,0.5)})  
alphameanest<-quantile(alphamean,0.5)  
betameanest<-quantile(betamean,0.5)  
  
#Graphical illustration  
plot(data$x[serie==1],data$y[serie==1],type="n",xlab="x",ylab="y",xlim=c(0,900),ylim=c(0,2.5))  
for (j in 1:3)  
{  
  lines(data$x[serie==j],alphaest[j]+betaest[j]*(data$x[serie==j]),col=j)  
  points(data$x[serie==j],data$y[serie==j],col=j)  
}
```



R2WinBUGS: fitted curves

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