## Computer Lab 4 732A54 - Bayesian Learning

Jooyoung Lee - joole336 Zuxiang Li - zuxli371

May 24, 2020

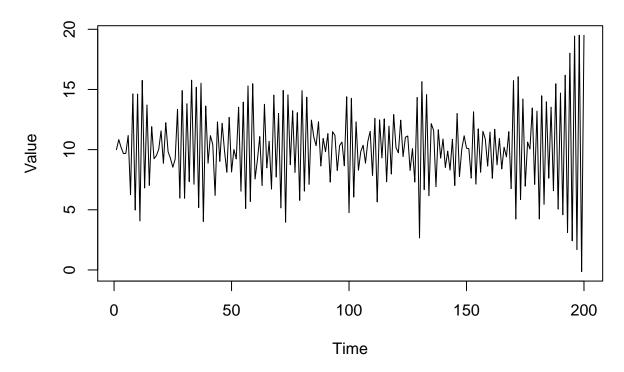
#### 1. Time series models in Stan

(a) Write a function in R that simulates data from the AR(1)-process

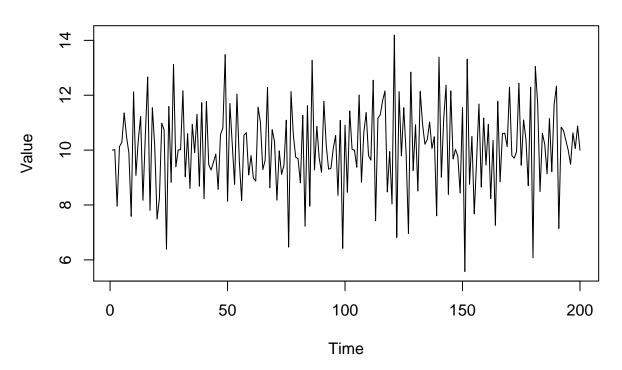
$$x_t = \mu + \phi(x_{t-1} - \mu) + \epsilon_t, \epsilon_t \sim N(0, \sigma^2)$$

for given values of  $\mu, \phi$  and  $\sigma^2$ . Start the process at  $x_1 = \mu$  and then simulate values for  $x_t$  for  $t = 2, 3, \dots, T$  and return the vector  $x_{1:T}$  containing all time points. Use  $\mu = 10$ ,  $\sigma^2 = 2$  and T = 200 and look at some different realizations (simulations) of  $x_{1:T}$  for values of  $\phi$  between -1 and 1 (this is the interval of  $\phi$  where the AR(1)-process is stable). Include a plot of at least one realization in the report. What effect does the value of  $\phi$  have on  $x_{1:T}$ 

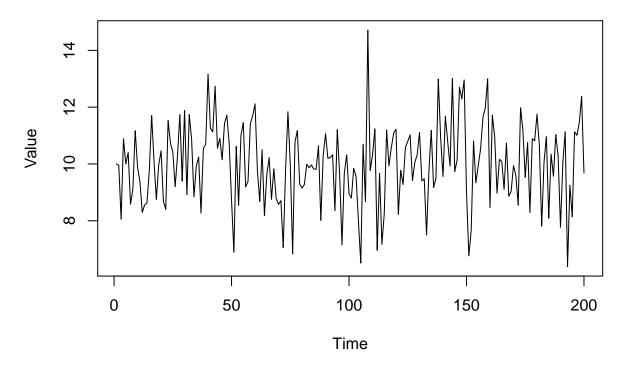
### phi=-1



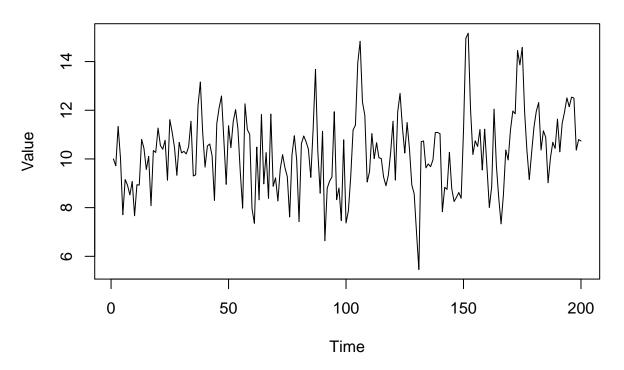




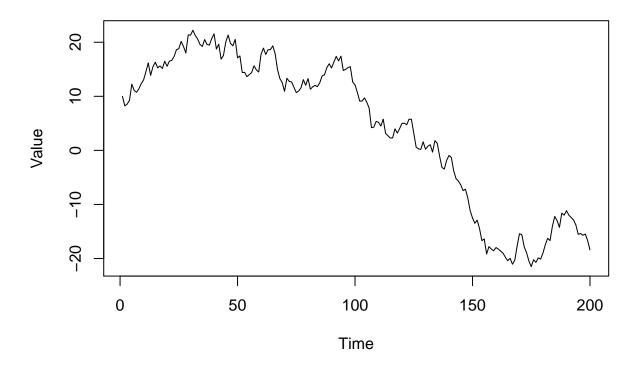








### phi=1



As the absolute value of  $\phi$  decreases, the simulated result shows more random result.

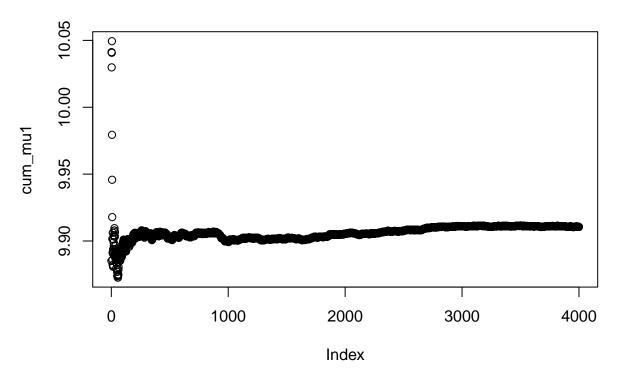
- (b) Use your function from a) to simulate two AR(1)-processes,  $x_{1:T}$  with  $\phi = 0.3$  and  $y_{1:T}$  with  $\phi = 0.95$ . Now, treat your simulated vectors as synthetic data, and treat the values of  $\mu$ ,  $\phi$  and  $\sigma^2$  as unknown and estimate them using MCMC. Implement Stan-code that samples from the posterior of the three parameters, using suitable non-informative priors of your choice. [Hint: Look at the time-series models examples in the Stan user's guide/reference manual, and note the different parameterization used here.]
  - i. Report the posterior mean, 95% credible intervals and the number of effective posterior samples for the three inferred parameters for each of the simulated AR(1)-process. Are you able to estimate the true values?
- ii. For each of the two data sets, evaluate the convergence of the samplers and plot the joint posterior of  $\mu$  and  $\phi$ . Comments?
- ## Warning: There were 193 divergent transitions after warmup. Increasing adapt\_delta above 0.8 may help ## http://mc-stan.org/misc/warnings.html#divergent-transitions-after-warmup
- ## Warning: Examine the pairs() plot to diagnose sampling problems
- ## Warning: Bulk Effective Samples Size (ESS) is too low, indicating posterior means and medians may be
- ## Running the chains for more iterations may help. See
- ## http://mc-stan.org/misc/warnings.html#bulk-ess

## Warning: Tail Effective Samples Size (ESS) is too low, indicating posterior variances and tail quant
## Running the chains for more iterations may help. See
## http://mc-stan.org/misc/warnings.html#tail-ess

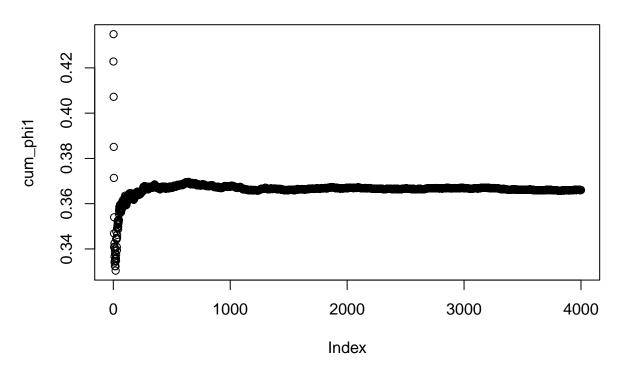
```
##
                             se_mean
                                                         2.5%
                                                                     97.5%
                                                                              n_eff
                   mean
                                             sd
## mu
              9.9105327 0.002607351 0.16681281
                                                   9.5803863
                                                                10.2502626 4093.169
              0.3660179 0.001052926 0.06570282
                                                   0.2393487
                                                                 0.4970067 3893.791
## phi
              2.0685545 0.003485378 0.20848421
                                                   1.6954845
                                                                 2.5057457 3578.052
## sigmasq
           -172.8497650 0.028467281 1.27242066 -176.2412856 -171.4430240 1997.880
## lp__
##
                Rhat
## mu
           1.0010368
## phi
           1.0002323
## sigmasq 0.9992364
## lp__
           0.9998927
##
                             se_mean
                                                          2.5%
                                                                      97.5%
                                                                               n_eff
                   mean
## mu
              5.9961859 0.858325888 17.11822032
                                                                 27.0287142 397.7524
                                                  -26.1287080
## phi
              0.9514007 0.002430702 0.02985416
                                                    0.8959984
                                                                  0.9988678 150.8504
              2.0383176 0.008875215 0.21412869
                                                    1.6723550
                                                                  2.5318175 582.0924
## sigmasq
## lp__
           -174.1090357 0.238838693 2.42935403 -180.3007247 -171.5303495 103.4599
```

## Rhat
## mu 1.011453
## phi 1.013877
## sigmasq 1.000070
## lp\_\_ 1.017550

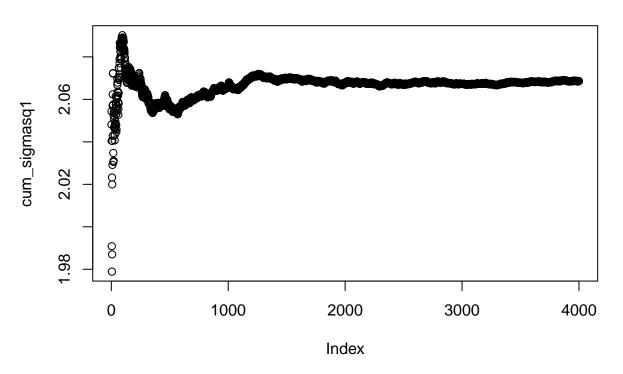
#### mu1



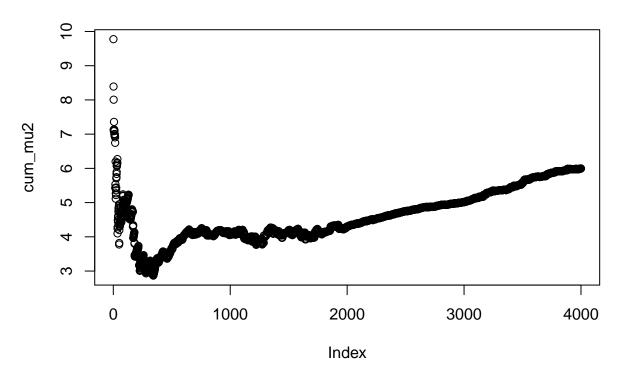


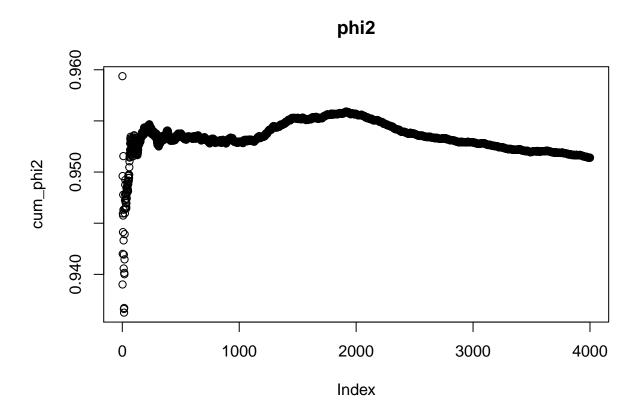


# sigmasq1

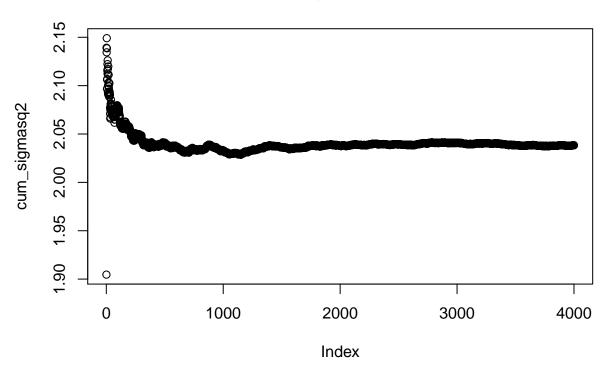


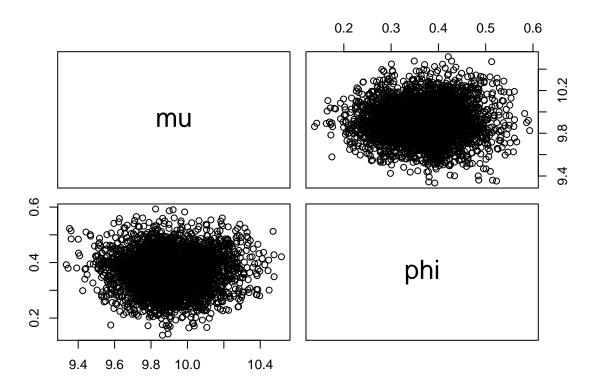


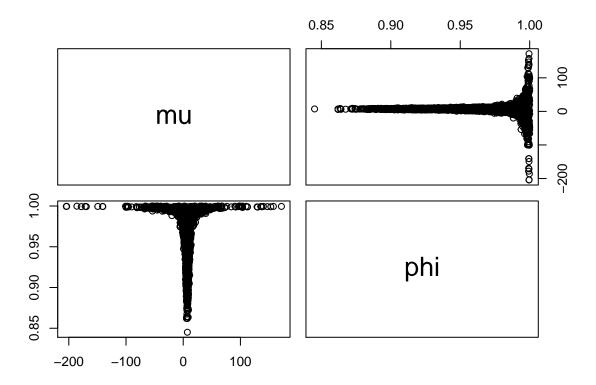




# sigmasq2







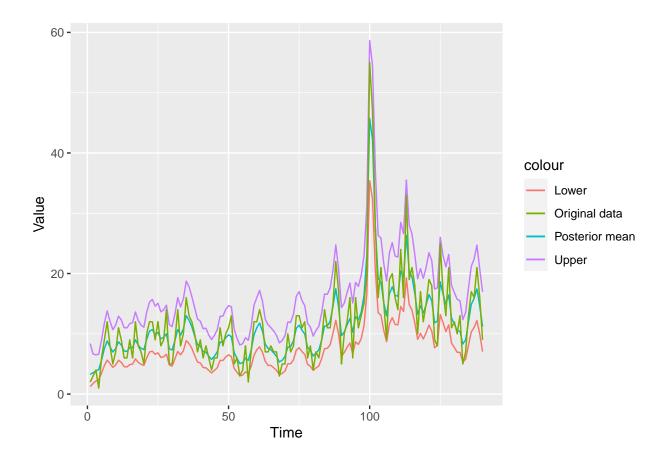
The posterior using the data which  $\phi = 0.3$  shows converging simulated parameters. The joint posterior of  $\mu 1$  and  $\phi 1$  are concentrated around parameters' mean. The joint posterior distribution seems normally distributed. Since we know the data is simulated using normal distribution, it is possible to conclude that the model constructed for sampling described the data well.

However, the posterior using the data which  $\phi = 0.95$  showed different result to the one using data with  $\phi = 0.3$ . Sampled  $\mu$ 2 and  $\phi$ 2 tend to show trends, rather than converging to certain values. This is because the randomness involved in the data for deriving posterior is significantly less than the one with  $\phi = 0.3$ .

(c) The data campy.dat contain the number of cases of campylobacter infections in the north of the province Quebec (Canada) in four week intervals from January 1990 to the end of October 2000. It has 13 observations per year and 140 observations in total. Assume that the number of infections  $c_t$  at each time point follows an independent Poisson distribution when conditioned on a latent AR(1)-process  $x_t$ , that is

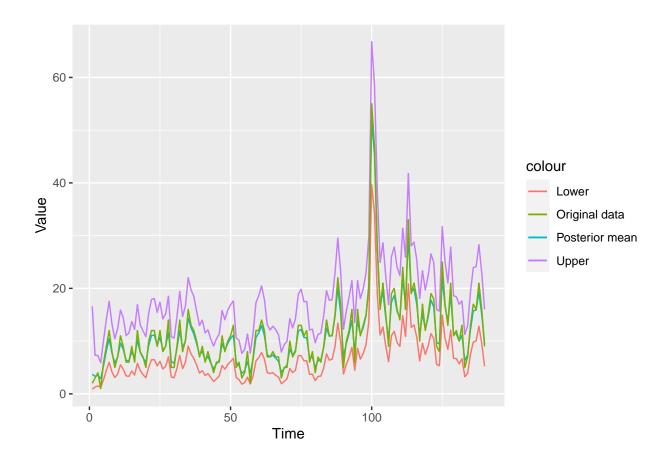
$$c_t | x_t \sim Poisson(exp(x_t))$$

where  $x_t$  is an AR(1)-process as in a). Implement and estimate the model in Stan, using suitable priors of your choice. Produce a plot that contains both the data and the posterior mean and 95% credible intervals for the latent intensity  $\theta_t = exp(x_t)$  over time. [Hint: Should  $x_t$  be seen as data or parameters?]



In this case,  $x_t$  should be seen as parameter; the resulting value that we want to simulate is  $c_t$ , which is a variable depending on value of  $x_t$ .

(d) Now, assume that we have a prior belief that the true underlying intensity  $\theta_t$  varies more smoothly than the data suggests. Change the prior for  $\sigma^2$  so that it becomes informative about that the AR(1)-process increments  $\epsilon_t$  should be small. Re-estimate the model using Stan with the new prior and produce the same plot as in c). Has the posterior for  $\theta_t$  changed?



Increased degrees of freedom in simulating sigmasq represents more informative prior for it. In this question we suppose nu=20 on prior. Means of posterior thetas are more similar to actual data. 95% credible interval seems wider than before; but such credible interval covers almost all actual data points.

### Appendix A: Code Question 1

```
ar1_process=function(t,mu,phi,sigmasq){
  res=vector(length=t)
  res[1]=mu
  for(i in 2:t){
    res[i]=mu+phi*(res[i-1]-mu)+rnorm(1,0,sqrt(sigmasq))
  return(res)
}
mu=10
sigmasq=2
t=200
tmp2=ar1_process(t,mu,-1,sigmasq)
plot(x=seq(1,t),y=tmp2,type="l",xlab = "Time",ylab="Value",main = "phi=-1")
tmp3=ar1 process(t,mu,-0.5,sigmasq)
plot(x=seq(1,t),y=tmp3,type="1",xlab = "Time",ylab="Value",main = "phi=-0.5")
tmp4=ar1_process(t,mu,0,sigmasq)
plot(x=seq(1,t),y=tmp4,type="1",xlab = "Time",ylab="Value",main = "phi=0")
tmp1=ar1_process(t,mu,0.5,sigmasq)
plot(x=seq(1,t),y=tmp1,type="l",xlab = "Time",ylab="Value",main = "phi=0.5")
tmp5=ar1_process(t,mu,1,sigmasq)
plot(x=seq(1,t),y=tmp5,type="1",xlab = "Time",ylab="Value",main = "phi=1")
x1t=ar1_process(200,10,0.3,2)
y1t=ar1_process(200,10,0.95,2)
StanModel = '
data {
  int<lower=0> N;
  vector[N] y;
parameters {
 real mu;
 real <lower=-1,upper=1>phi;
 real<lower=0> sigmasq;
model {
 mu~ normal(1,100);
  phi~ normal(1,100);
  sigmasq ~ scaled_inv_chi_square(1,2);
 for (n in 2:N)
    y[n] ~ normal(mu + phi * (y[n-1]-mu), sqrt(sigmasq));
}
data1 = list(N=length(x1t), y=x1t)
data2 = list(N=length(y1t), y=y1t)
burnin = 1000
niter = 2000
fit1 = stan(model_code=StanModel,data=data1, warmup=burnin,iter=niter,chains=4,refresh=0)
fit2 = stan(model_code=StanModel,data=data2, warmup=burnin,iter=niter,chains=4,refresh=0)
```

```
df1=summary(fit1,probs = c(0.025, 0.975))$summary
print(df1)
df2=summary(fit2,probs = c(0.025, 0.975))$summary
print(df2)
# phi=0.3
mu1<- extract(fit1, 'mu')$mu</pre>
cum mu1 = cumsum(mu1)
for (i in 1:length(cum_mu1)){
  cum_mu1[i] = cum_mu1[i]/i
plot(cum_mu1,main="mu1")
phi1 <- extract(fit1, 'phi')$phi</pre>
cum_phi1 = cumsum(phi1)
for (i in 1:length(cum_phi1)){
  cum_phi1[i] = cum_phi1[i]/i
plot(cum_phi1,main="phi1")
sigmasq1 <- extract(fit1, 'sigmasq')$sigmasq</pre>
cum_sigmasq1 = cumsum(sigmasq1)
for (i in 1:length(cum_sigmasq1)){
  cum_sigmasq1[i] = cum_sigmasq1[i]/i
plot(cum_sigmasq1,main="sigmasq1")
# phi=0.95
mu2<- extract(fit2, 'mu')$mu</pre>
cum_mu2 = cumsum(mu2)
for (i in 1:length(cum_mu2)){
  cum_mu2[i] = cum_mu2[i]/i
plot(cum_mu2,main="mu2")
phi2 <- extract(fit2, 'phi')$phi</pre>
cum_phi2 = cumsum(phi2)
for (i in 1:length(cum phi2)){
  cum_phi2[i] = cum_phi2[i]/i
plot(cum_phi2,main="phi2")
sigmasq2 <- extract(fit2, 'sigmasq')$sigmasq</pre>
cum_sigmasq2= cumsum(sigmasq2)
for (i in 1:length(cum_sigmasq2)){
  cum_sigmasq2[i] = cum_sigmasq2[i]/i
plot(cum_sigmasq2,main="sigmasq2")
pairs(extract(fit1, c('mu','phi')))
pairs(extract(fit2, c('mu', 'phi')))
data=read.table("campy.dat",header = TRUE)
data = list(N=length(data$c), ct=data$c)
```

```
StanModel ='
data {
     int<lower=0> N;
     int ct[N];
parameters {
    real mu;
    real <lower=-1,upper=1>phi;
    real<lower=0> sigma;
     real xt[N];
}
model {
     mu~ normal(1,100);
     phi~ normal(1,100);
     sigma ~ scaled_inv_chi_square(1,2);
     xt[1]~normal(mu,1);
     ct[1]~poisson(exp(mu));
     for (n in 2:N){
           xt[n]~normal(mu + phi * (xt[n-1]-mu), sigma);
           ct[n] ~ poisson(exp(xt[n]));
     }
}
burnin = 1000
niter = 2000
fit1 = stan(model_code=StanModel,data=data, warmup=burnin,iter=niter,chains=4,refresh=0)
df=as.data.frame(summary(fit1,probs = c(0.025, 0.975))$summary)
df=df[c(-1,-2,-3,-144),]
df$mean=exp(df$mean)
df$ 2.5% = exp(df$ 2.5%)
df$ 97.5% = exp(df$ 97.5%)
ggplot()+geom_line(aes(x=c(1:length(df$mean)),y=df$mean,colour="Posterior mean"))+geom_line(aes(x=c(1:length(df$mean)),y=df$mean,colour="Posterior m
data=read.table("campy.dat",header = TRUE)
data = list(N=length(data$c), ct=data$c)
StanModel ='
data {
     int<lower=0> N;
     int ct[N];
parameters {
    real mu;
     real <lower=-1,upper=1>phi;
    real<lower=0> sigma;
    real xt[N];
}
model {
     mu~ normal(1,100);
     phi~ normal(1,100);
     sigma ~ scaled_inv_chi_square(20,2);
     xt[1]~normal(mu,1);
      ct[1]~poisson(exp(mu));
```

```
for (n in 2:N){
    xt[n]-normal(mu + phi * (xt[n-1]-mu), sigma);
    ct[n] ~ poisson(exp(xt[n]));
}

burnin = 1000
niter = 2000
fit1 = stan(model_code=StanModel,data=data, warmup=burnin,iter=niter,chains=4,refresh=0)

df=as.data.frame(summary(fit1,probs = c(0.025, 0.975))$summary)
df=df[c(-1,-2,-3,-144),]
df$mean=exp(df$mean)
df$^2.5%^=exp(df$^2.5%^)
df$^97.5%^=exp(df$^97.5%^)
ggplot()+geom_line(aes(x=c(1:length(df$mean)),y=df$mean,colour="Posterior mean"))+geom_line(aes(x=c(1:length(df$mean)),y=df$mean,colour="Posterior mean"))+geom_line(aes(x=c(1:length(df$mean)),y=df$mea
```