

#1 5.1



Making cereal for breakfast:

- ① get out a bowl & spoon
- ② pour cereal in the bowl
- ③ milk in the bowl
- ④ eat cereal
- ⑤ wash dishes

\* cereal comes first, don't fight me on that

$$F_{\text{system}} = 1 - \left[ (1 - F_1(+)) (1 - F_2(+)) (1 - F_3(+)) (1 - F_4(+)) (1 - F_5(+)) \right]$$

#2 5.8  $h(+): \left( \frac{f(t)}{R(t)} \right)^n$

$$f(t) \text{ w/ } \theta=0 \Rightarrow \lambda \beta(x)^{\beta-1} e^{-\lambda x^\beta} \quad \lambda, \beta > 0$$

$$R(t) = 1 - \left[ \int_0^t \lambda \beta x^{\beta-1} e^{-\lambda x^\beta} dx \right]$$

$$\lambda \beta \int_0^t x^{\beta-1} e^{-\lambda x^\beta} dx$$

$$(x^\beta)(x^{-1})(e^{-\lambda x^\beta})$$

$$u = x^\beta$$

$$du = \beta x^{\beta-1} dx$$

$$dx = \frac{1}{\beta x^{\beta-1}} du$$

$$x=0 \quad u=x=t^{\beta}$$

$$= \int (\bar{e}^{-\lambda u}) \cancel{(x^{\beta-1})} \left( \frac{1}{\beta x^{\beta-1}} \right) du = \int_0^{t^{\beta}} \frac{\bar{e}^{-\lambda u}}{\beta} du$$

$$\cancel{\beta \lambda} \frac{1}{\beta} \int_0^{t^{\beta}} \bar{e}^{-\lambda u} du \Rightarrow \lambda \left[ \frac{\bar{e}^{-\lambda u}}{-\lambda} \right] \Big|_0^{t^{\beta}}$$

$$= - \left[ \bar{e}^{\lambda u} \right] \Big|_0^{t^{\beta}} = - \left[ \bar{e}^{-\lambda t^{\beta}} - 1 \right]$$

$$h(t) = \left[ \frac{f(t)}{R(t)} \right]^n$$

$$\text{CDF} = -\bar{e}^{-\lambda t^{\beta}} + 1$$

Reliability

$$= \bar{e}^{\lambda t^{\beta}}$$

$$= \prod_{i=1}^n \lambda \beta x_i^{\beta-1} \bar{e}^{-\lambda x_i^{\beta}}$$

$$\overline{\prod_{i=1}^n \bar{e}^{-\lambda x_i^{\beta}}}$$

$$h(t) = \prod_{i=1}^n [\lambda \beta x_i^{\beta-1}] = \lambda^n \beta^n x^{\sum_{i=1}^n \beta_i - n}$$

# STAT 466 HW 12

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## Question 1

Draw a reliability block diagram describing how to successfully perform an everyday task. Include an equation which shows the system CDF comprised of the component CDFs.

## Question 2

Calculate the hazard function for a series system with n components when each component lifetime has a Weibull distribution. (use Parameterization 1 from Appendix B, with  $\theta = 0$ )

## Question 3

Fit a log-linear process to the data in Table 6.2. How does the fit compare with that of a PLP in terms of a Bayesian  $\chi^2$  goodness-of-fit test? Also compare the log-linear process and PLP fits using DIC discussed in Sect. 4.6.

```
super.comp <- read.csv('table62.txt', header=TRUE, sep = '', strip.white = TRUE)
head(super.comp)
```

```
## Month Cumulative.Day Failures Cumulative.Fail
## 1 Jul 31 5 5
## 2 Aug 62 4 9
## 3 Sep 92 6 15
## 4 Nov 123 1 16
## 5 Dec 153 2 18
## 6 Jan 184 1 19
```

```
cum.fail <- super.comp$Cumulative.Fail
cum.day <- super.comp$Cumulative.Day
failures <- super.comp$Failures
n <- length(failures)
z <- rep(0,n)

BMSmpLLM.HW <- "model {
  for(i in 1:(n-1)){
    cum.fail[i] ~ dpois(lam[i])
    lam[i] <- -(gam0+gam1*cum.day[i]) + C
  }
  cum.fail[n] ~ dpois(lam[n])
  lam[n] = exp(gam0)*(exp(gam1*cum.day[n])-1)/gam1 + C
  gam0 ~ dnorm(0,1/10)
  gam1 ~ dnorm(0,1/10)
  C = 1000
}"
"
```

```
jags.inits <- list(list(gam0=1,gam1=1),
                      list(gam0=0,gam1=-1),
                      list(gam0=-1,gam1=-1))
```

```
BMSmpLLM.HW.sim <- jags(
  data=c('cum.day','cum.fail','n'),
  parameters.to.save=c('gam0','gam1'),
  model.file=textConnection(BMSmpLLM.HW),
  inits=jags.inits,
  n.iter=32000,
  n.burnin=2000,
  n.chains=3,
  n.thin=1
)
```

```
## module glm loaded
```

```
## Compiling model graph
## Resolving undeclared variables
## Allocating nodes
## Graph information:
##   Observed stochastic nodes: 15
##   Unobserved stochastic nodes: 2
##   Total graph size: 101
##
## Initializing model
```

```
head(BMSmpLLM.HW.sim$BUGSoutput$sims.matrix)
```

```
##      deviance      gam0      gam1
## [1,] 26883.78 -3.293488  0.0002808562
## [2,] 26883.47 -3.513959 -0.0007114971
## [3,] 26883.82 -4.015364  0.0015455789
## [4,] 26886.85 -2.558481 -0.0040811540
## [5,] 26883.62 -3.057508 -0.0018198002
## [6,] 26883.20 -3.656092  0.0010440093
```

```
length(BMSmpLLM.HW.sim$BUGSoutput$sims.matrix[,1]) #90,000 samples obtained
```

```
## [1] 90000
```

```
gam0 <- BMSmpLLM.HW.sim$BUGSoutput$sims.matrix[,2]
gam1 <- BMSmpLLM.HW.sim$BUGSoutput$sims.matrix[,3]

gelman.diag(BMSmpLLM.HW.sim$BUGSoutput)
```

```
## Potential scale reduction factors:
##
##          Point est. Upper C.I.
## deviance      1      1
## gam0          1      1
## gam1          1      1
##
## Multivariate psrf
##
## 1
```

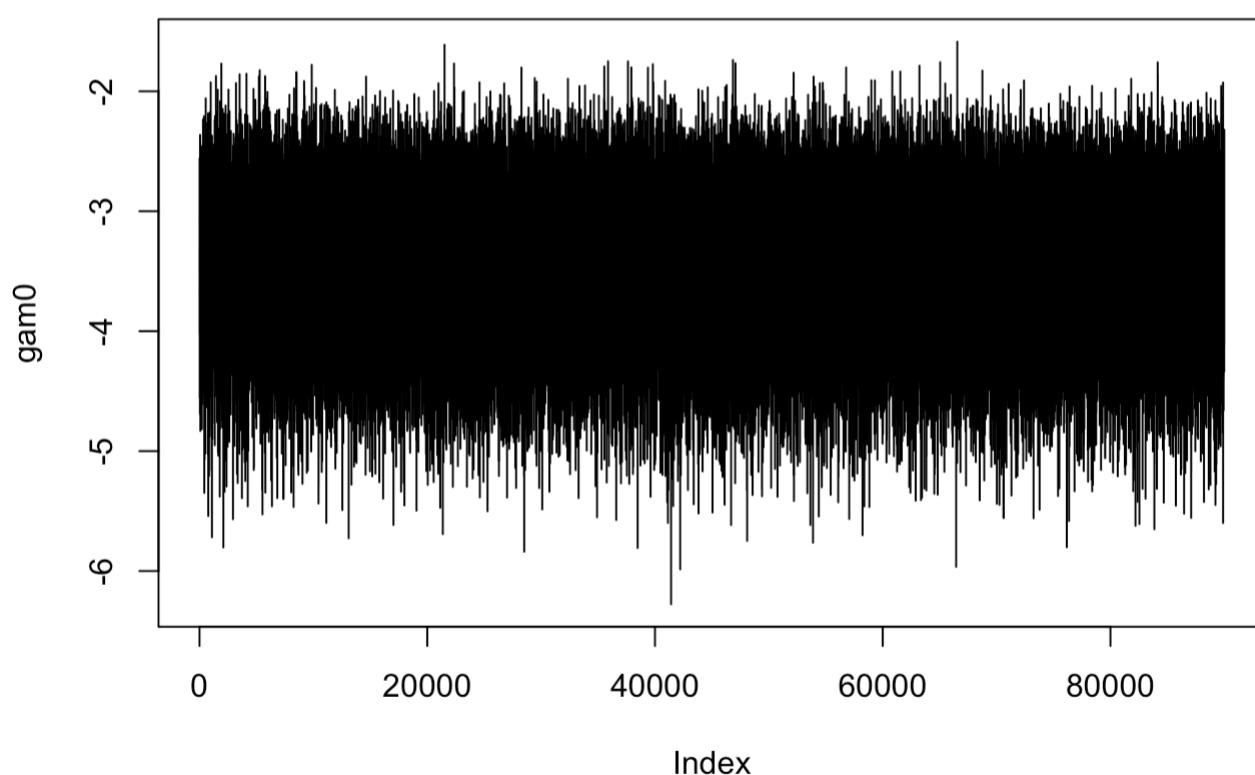
```
BMSmpLLM.HW.sim$BUGSoutput$DIC
```

```
## [1] 26886.85
```

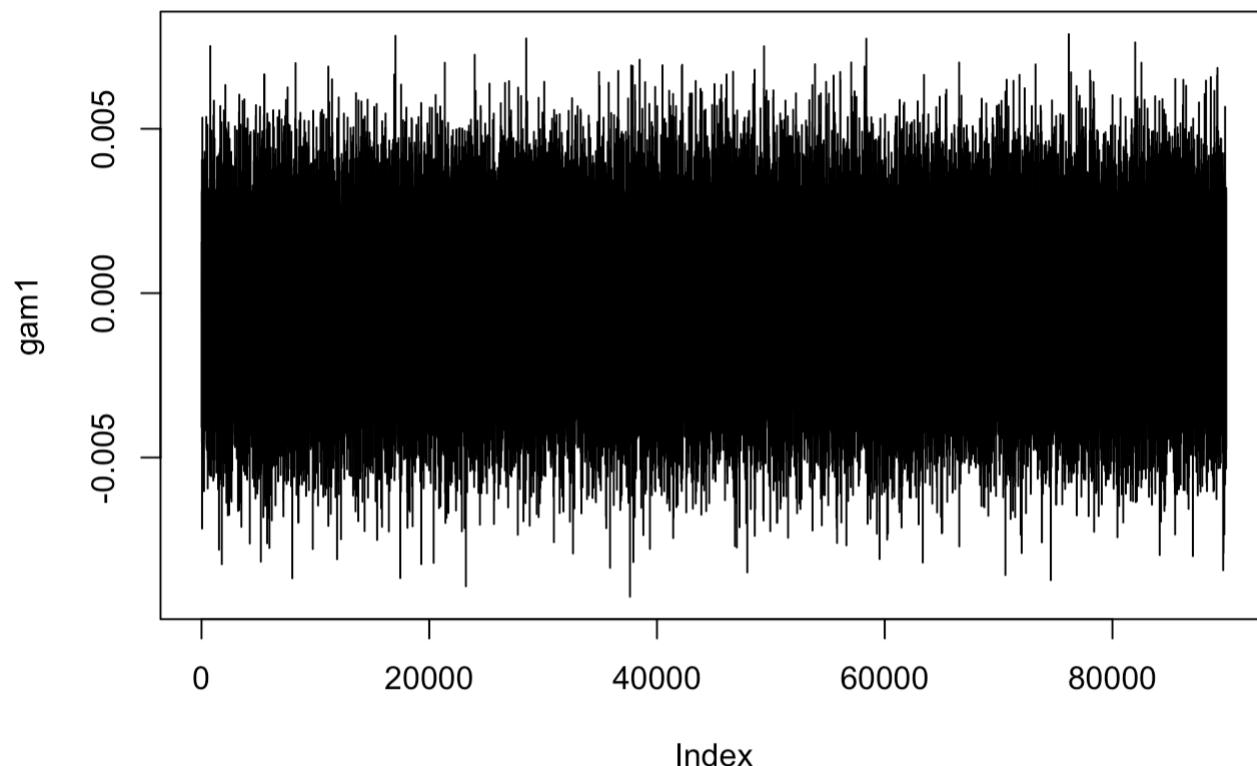
```
effectiveSize(BMSmpLLM.HW.sim)
```

```
## deviance      gam0      gam1
## 19649.474  8221.810  8298.632
```

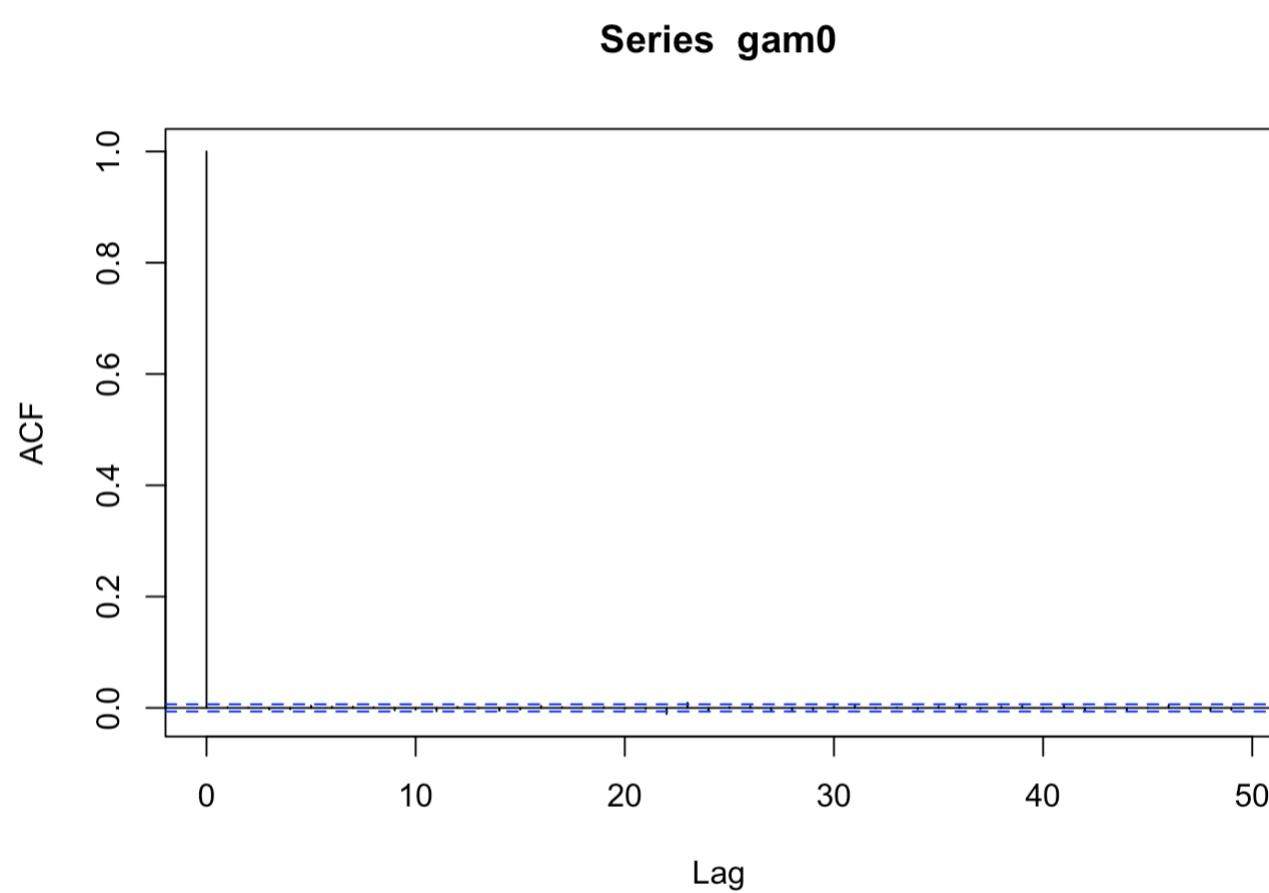
```
plot(gam0, type='l')
```



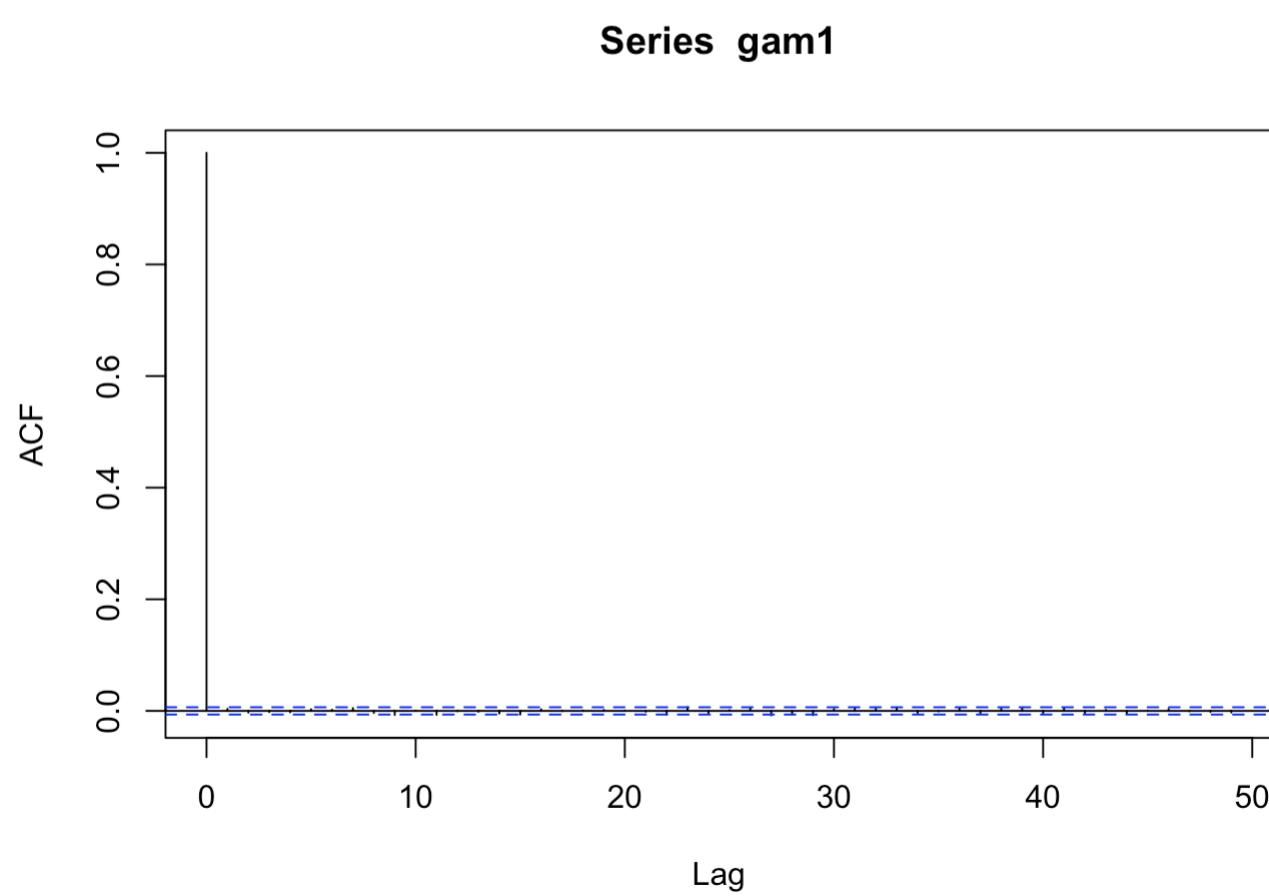
```
plot(gam1, type='l')
```



```
acf(gam0)
```



```
acf(gam1)
```



```

mcmcChain <- BMSmpLLM.HW.sim$BUGSoutput$sims.matrix[,3]
C <- 1000

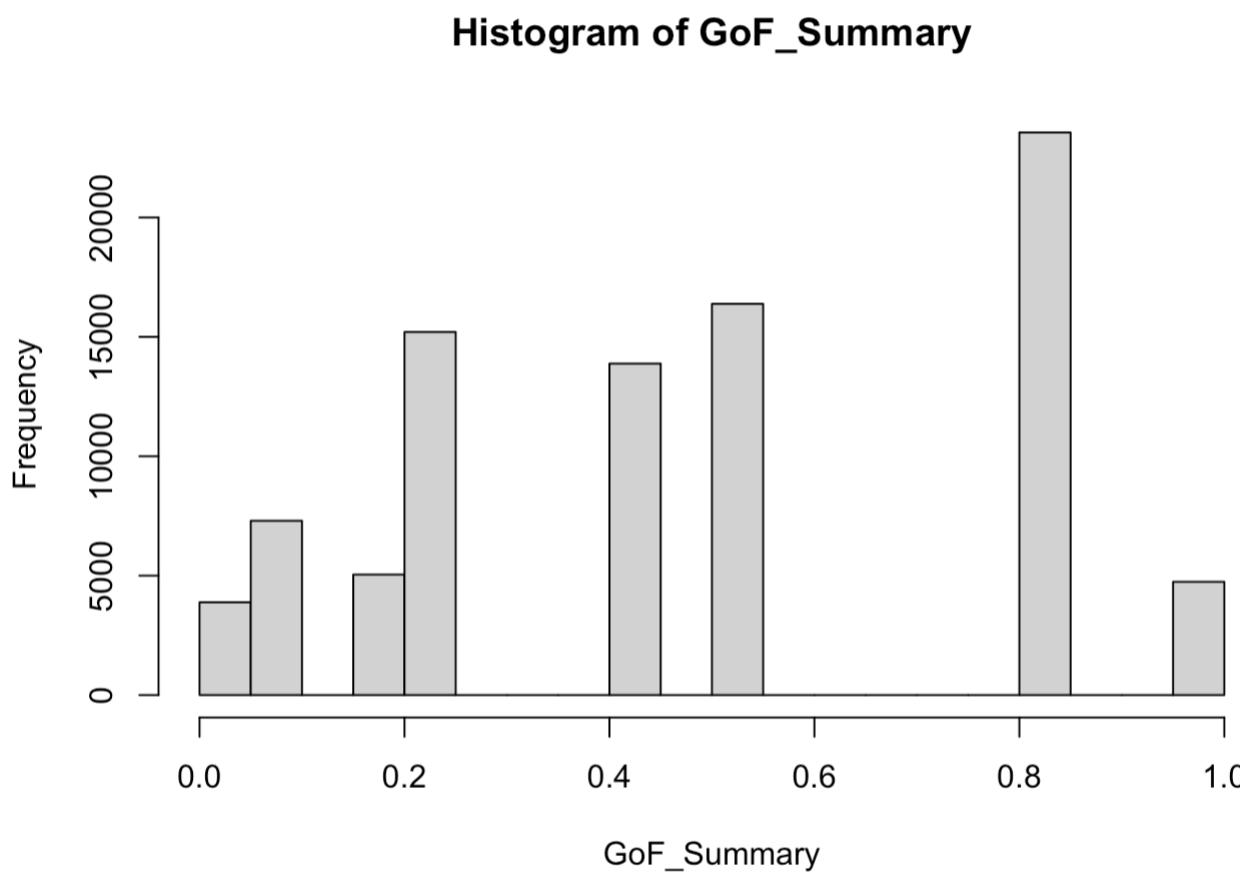
GoF <- matrix(NA, ncol=length(cum.fail), nrow=length(mcmcChain))
for (i in 1:length(mcmcChain)) {
  sim.fail <- numeric(length(cum.fail))
  for (j in 1:length(cum.fail)) { #exp(gam0)*(exp(gam1*cum.day[n])-1)/gam1 + C
    sim.fail[j] <- exp(gam0[i])*(exp(gam1[i]*cum.day[j])-1)/gam1[i] + C
  }
  GoF[i,] <- ppois(cum.fail,sim.fail)
  temp <- runif(length(cum.fail),GoF[i,],1)
  GoF[i,] <- pmax(GoF[i,],temp)
}

# Function requires fitted quantiles and returns a p-value
GoF_Test <- function(fitted_quantiles) {
  n <- length(fitted_quantiles)
  K <- round((n)^(0.4))
  mK <- table(cut(fitted_quantiles,(0:K)/K))
  np <- n/K
  RB <- sum(((mK-np)^2)/np)
  return(1-pchisq(RB,K-1))
}

GoF_Summary <- apply(GoF,1,GoF_Test)

hist(GoF_Summary,xlim=c(0,1))

```



```
mean(GoF_Summary < 0.05)
```

```
## [1] 0.04315556
```

```

# Power Law Process
cum.fail <- super.comp$Cumulative.Fail
cum.day <- super.comp$Cumulative.Day
failures <- super.comp$Failures
n <- length(cum.day)
z <- rep(0,n)

BMSmpPLP <- "model {
  for (i in 1:(n-1)) {
    z[i] ~ dpois(lam[i])
    lam[i] = -log(L[i]) + C
    L[i] = phi/eta * (cum.fail[i]/eta)^(phi-1)
  }
  z[n] ~ dpois(lam[n])
  lam[n] = L[n]
  L[n] = (cum.fail[n]/eta)^(phi)
  eta ~ dexp(1)
  phi ~ dexp(1)
  C = 100
}
"

```

```

BMSmpPLP.sim <- jags(
  data = c('cum.fail', 'z', 'n'),
  parameters.to.save = c('eta', 'phi'),
  model.file = textConnection(BMSmpPLP),
  n.iter = 32000,
  n.burnin = 2000,
  n.chains = 3,
  n.thin = 1
)

```

```

## Compiling model graph
## Resolving undeclared variables
## Allocating nodes
## Graph information:
##   Observed stochastic nodes: 15
##   Unobserved stochastic nodes: 2
##   Total graph size: 123
##
## Initializing model

```

```
head(BMSmpPLP.sim$BUGSoutput$sims.matrix)
```

```

##      deviance     eta      phi
## [1,] 2853.809 2.876219 1.208531
## [2,] 2856.138 1.406667 0.843632
## [3,] 2853.466 3.548596 1.062315
## [4,] 2851.409 4.741009 1.474377
## [5,] 2851.229 4.001995 1.258556
## [6,] 2852.588 2.927208 1.127695

```

```
length(BMSmpPLP.sim$BUGSoutput$sims.matrix[,1]) #90,000 samples obtained
```

```
## [1] 90000
```

```

eta <- BMSmpPLP.sim$BUGSoutput$sims.matrix[,2]
phi <- BMSmpPLP.sim$BUGSoutput$sims.matrix[,3]

gelman.diag(BMSmpPLP.sim$BUGSoutput)

```

```

## Potential scale reduction factors:
## 
##      Point est. Upper C.I.
## deviance      1      1
## eta          1      1
## phi          1      1
## 
## Multivariate psrf
## 
## 1

```

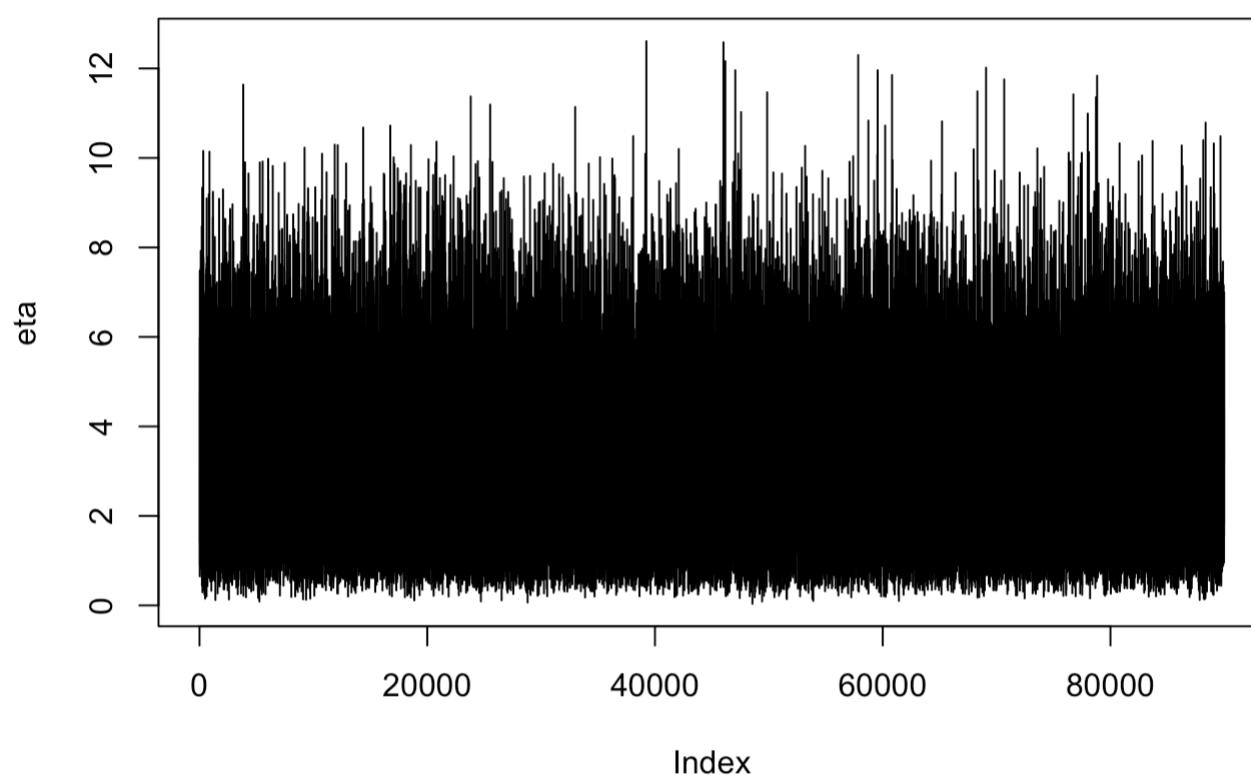
```
BMSmpPLP.sim$BUGSoutput$DIC
```

```
## [1] 2858.168
```

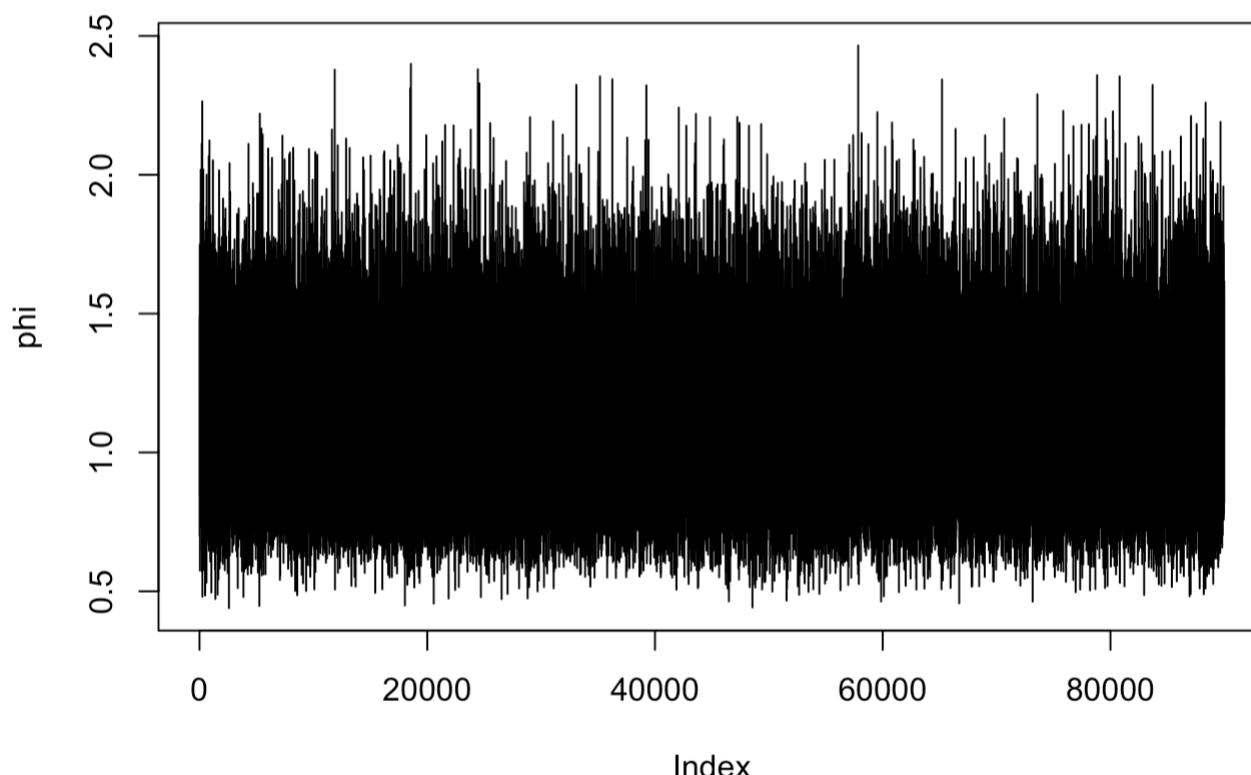
```
effectiveSize(BMSmpPLP.sim)
```

```
## deviance      eta      phi
## 6078.055 4429.091 4514.328
```

```
plot(eta, type='l')
```

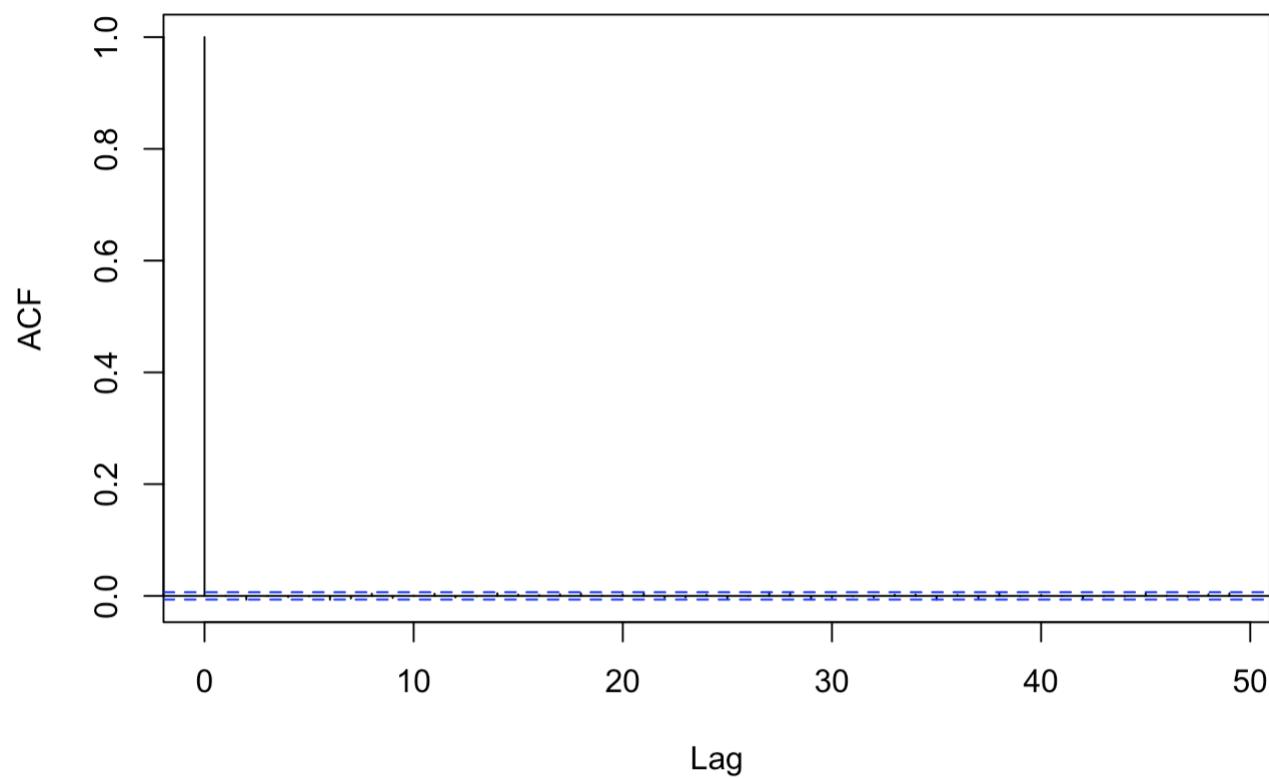


```
plot(phi, type='l')
```



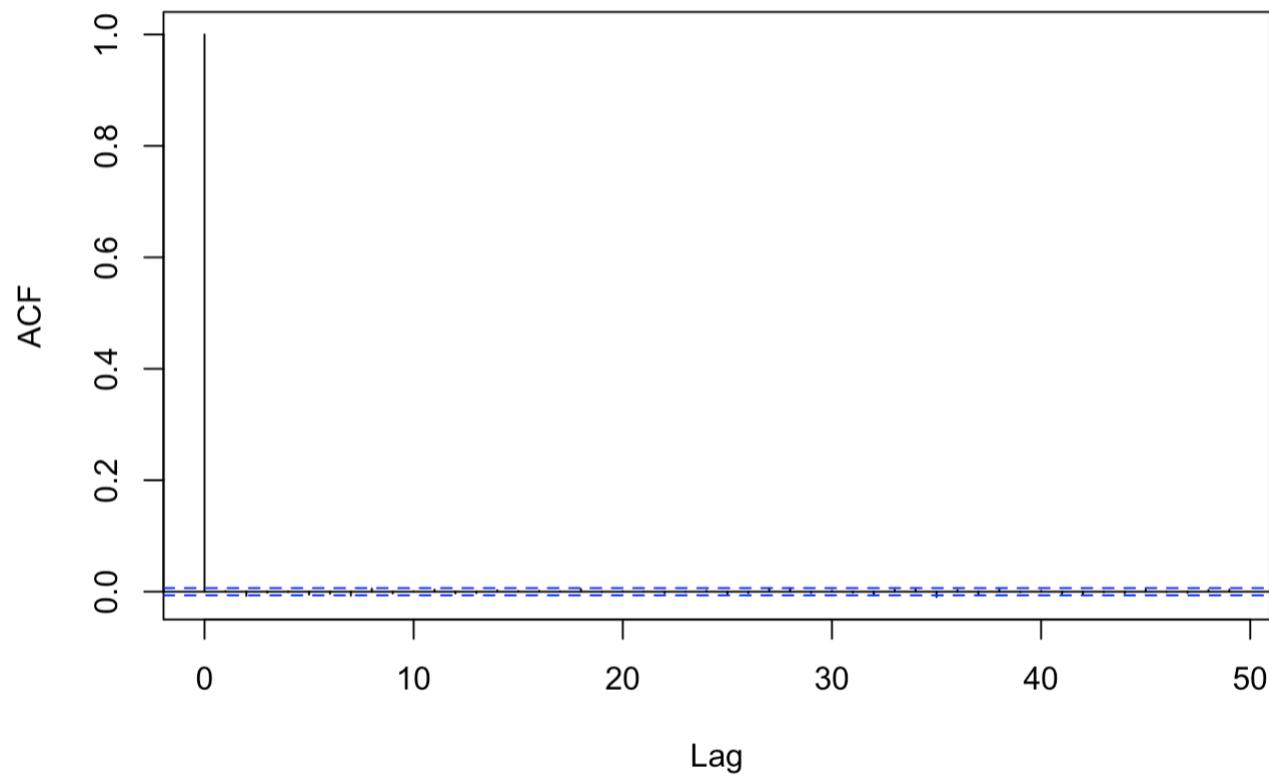
```
acf(eta)
```

**Series eta**



```
acf(phi)
```

**Series phi**



```

mcmcChainPLP <- BMSmpPLP.sim$BUGSoutput$sims.matrix[,3]

GoF <- matrix(NA, ncol=length(cum.fail), nrow=length(mcmcChainPLP))
for (i in 1:length(mcmcChainPLP)) {
  sim.fail <- numeric(length(cum.fail))
  for (j in 1:length(cum.fail)) { #phi/eta * (cum.fail[i]/eta)^{phi-1}
    sim.fail[j] <- (phi[i]/eta[i])*(cum.fail[j]/eta[i])^{(phi[i]-1)}
  }
  GoF[i,] <- ppois(cum.fail,sim.fail)
  temp <- runif(length(cum.fail),GoF[i,],1)
  GoF[i,] <- pmax(GoF[i,],temp)
}

# Function requires fitted quantiles and returns a p-value
GoF_Test <- function(fitted_quantiles) {
  n <- length(fitted_quantiles)
  K <- round((n)^{0.4})
  mK <- table(cut(fitted_quantiles,(0:K)/K))
  np <- n/K
  RB <- sum(((mK-np)^2)/np)
  return(1-pchisq(RB,K-1))
}

GoF_Summary <- apply(GoF,1,GoF_Test)

#hist(GoF_Summary,xlim=c(0,1))

#mean(GoF_Summary < 0.05)
# couldn't get this to run properly

```

DIC for the PLP is lower, but the log-linear process GoF is really good. The model fits the data well.