CWVS: Critical Window Variable Selection

CWVS_Example

- [1] Simulate data for analysis:
 - Setting the reproducibility seed and initializing packages for data simulation:

```
set.seed(4679)
library(CWVS)
library(boot) #Inverse logit transformation
```

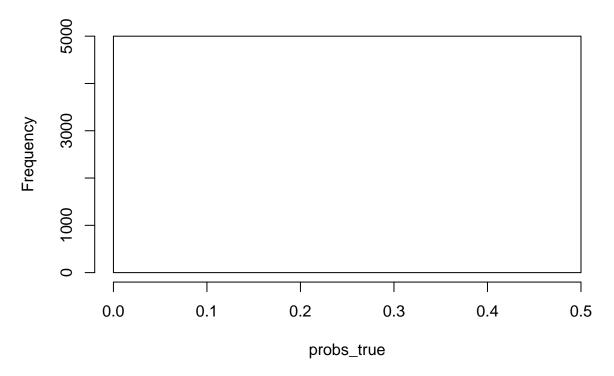
• Setting the global data values:

```
n<-5000 #Sample size
m<-27 #Number of exposure time periods
x<-matrix(1,
nrow=n,
ncol=1) #Covariate design matrix
z<-matrix(rnorm(n=(n*m)),
nrow=n,
ncol=m) #Exposure design matrix
for(j in 1:m){
z[,j]<-(z[,j] - median(z[,j]))/IQR(z[,j]) #Data standardization (interquartile range)
}</pre>
```

• Setting the values for the statistical model parameters:

```
beta_true<- -0.30
theta_true<-rep(0.00, times=m)
gamma_true<-c(rep(0, times=12),
rep(1, times=4),
rep(0, times=11))
alpha_true<-gamma_true*theta_true
logit_p_true<-x%*%beta_true +
z%*%alpha_true
probs_true<-inv.logit(logit_p_true)
hist(probs_true)</pre>
```

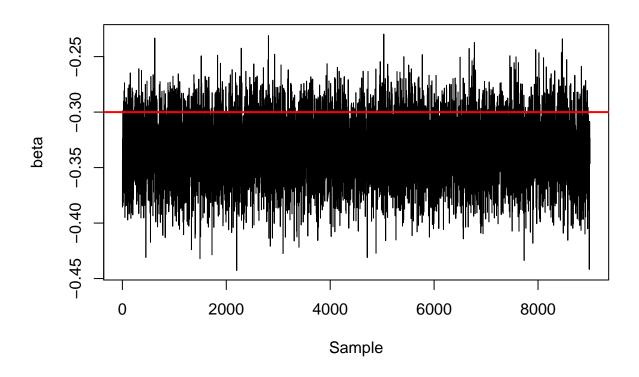
Histogram of probs_true

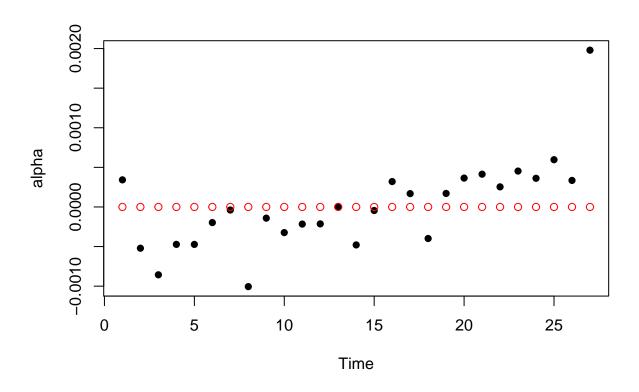


• Simulating the analysis dataset:

[2] Fit CWVS to identify/estimate critical windows of susceptibility:

```
## phi1 Acceptance: 40%
## phi2 Acceptance: 33%
## A11 Acceptance: 92%
## A22 Acceptance: 9%
## *********
## Progress: 40%
## phi1 Acceptance: 39%
## phi2 Acceptance: 34%
## A11 Acceptance: 92%
## A22 Acceptance: 11%
## *********
## Progress: 50%
## phi1 Acceptance: 39%
## phi2 Acceptance: 36%
## A11 Acceptance: 91%
## A22 Acceptance: 16%
## *********
## Progress: 60%
## phi1 Acceptance: 39%
## phi2 Acceptance: 35%
## A11 Acceptance: 91%
## A22 Acceptance: 15%
## **********
## Progress: 70%
## phi1 Acceptance: 39%
## phi2 Acceptance: 35%
## A11 Acceptance: 91%
## A22 Acceptance: 14%
## *********
## Progress: 80%
## phi1 Acceptance: 40%
## phi2 Acceptance: 35%
## A11 Acceptance: 91%
## A22 Acceptance: 12%
## **********
## Progress: 90%
## phi1 Acceptance: 40%
## phi2 Acceptance: 35%
## A11 Acceptance: 91%
## A22 Acceptance: 11%
## *********
## Progress: 100%
## phi1 Acceptance: 40%
## phi2 Acceptance: 35%
## A11 Acceptance: 91%
## A22 Acceptance: 11%
## **********
plot(results$beta[1, 1001:10000],
    type="1",
    ylab="beta",
    xlab="Sample")
abline(h=beta_true,
      col="red",
```





```
plot(rowMeans(results$gamma[,1001:10000]),
          pch=16,
          ylab="gamma",
          xlab="Time")
points(gamma_true,
          col="red")
```

