

MCMC Diagnostics - IFLS data

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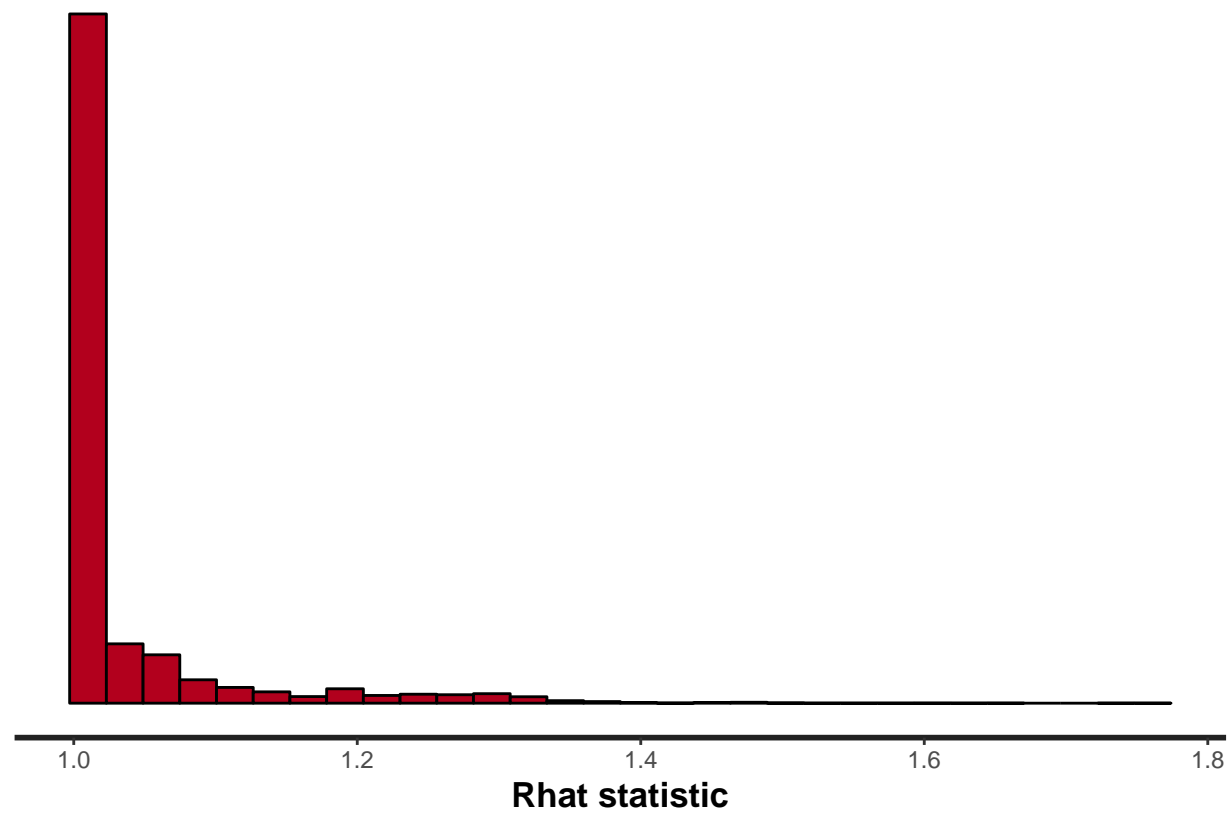
07/21/2020

```
K <- 7  
Ti <- 3  
N <- 1973
```

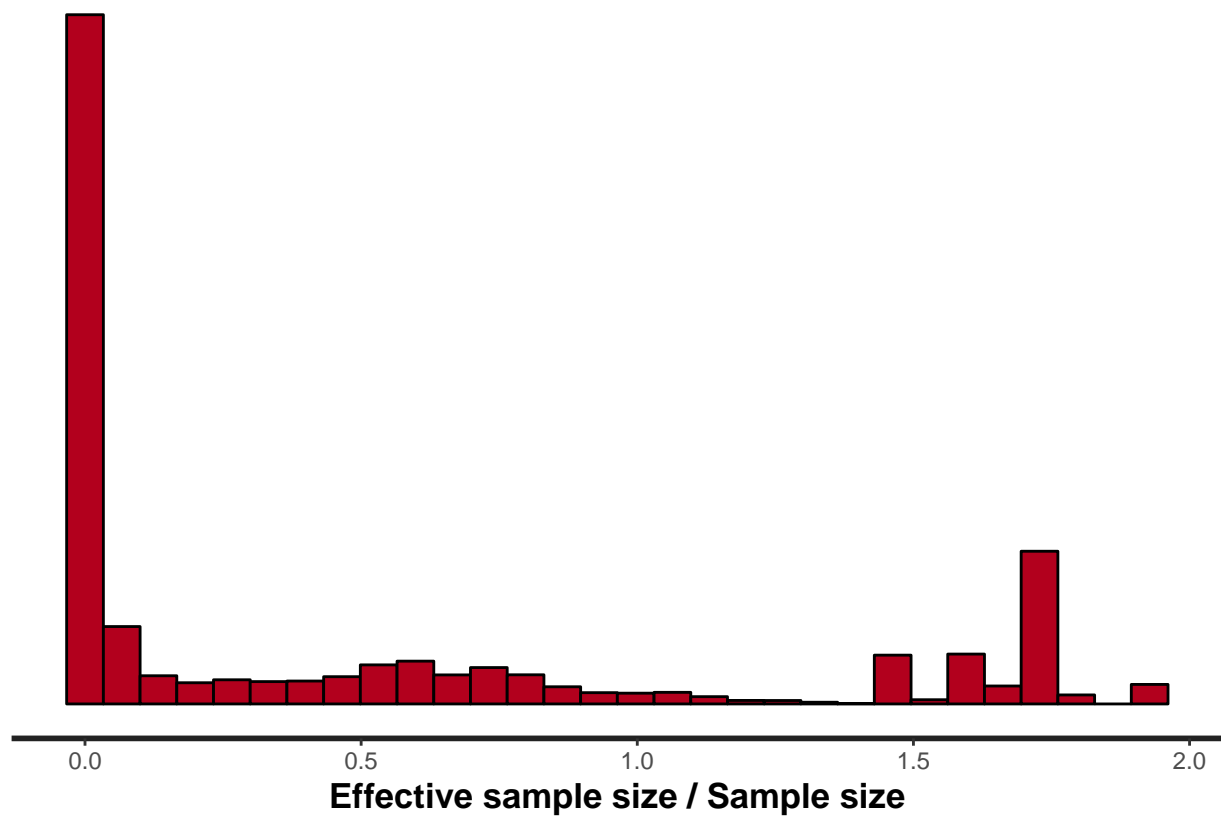
General MCMC diagnostic plots

Overall model diagnostics from rstan package.

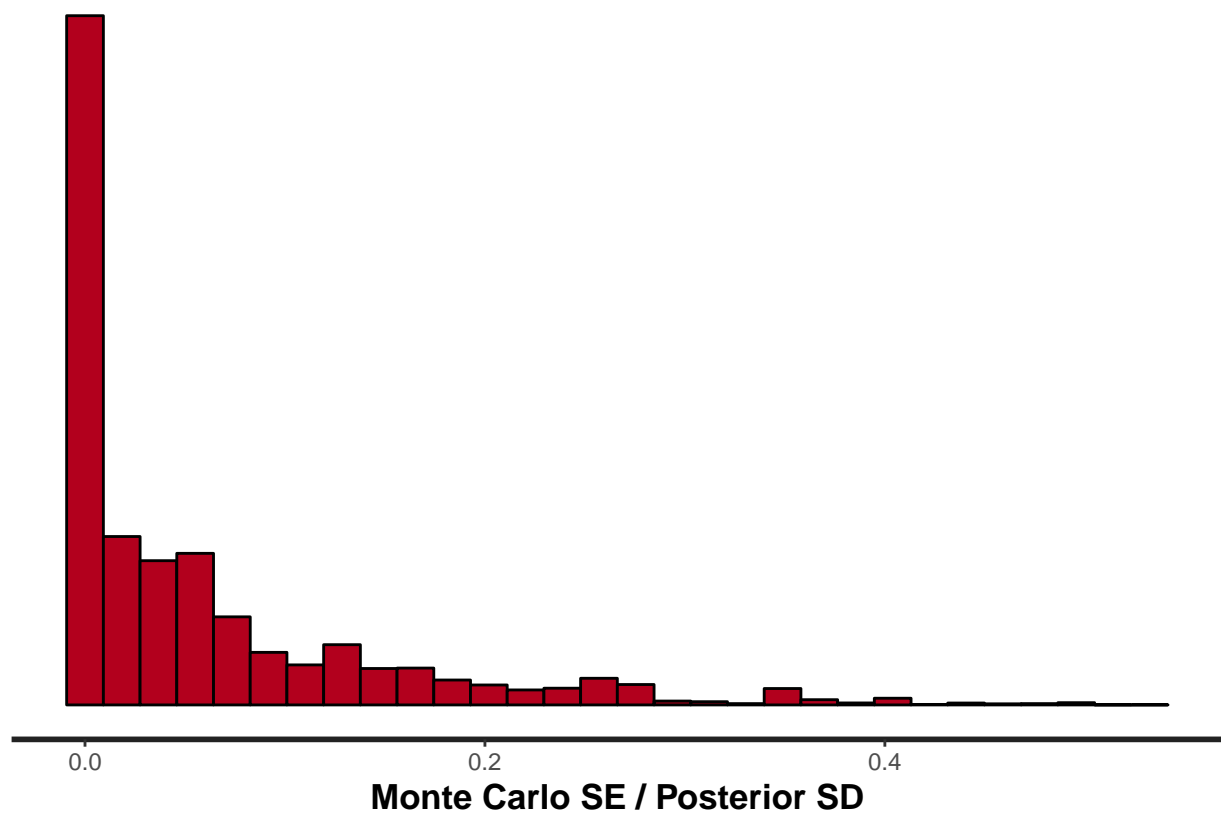
```
## `stat_bin()` using `bins = 30`. Pick better value with `binwidth`.
```



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```



Individual Parameter Diagnostics

Individual parameter plots. Autocorrelation and trace plots for individual parameters, and histograms of posterior medians for group parameters.

```
get_single_plots <- function(fit, param) {
  print(fit_summ[param,c(1,2,3,5,6,7,9,10)])
  print(stan_ac(fit, pars = param))
  print(rstan::traceplot(fit, pars = param))
}

get_aggreg_plots <- function(fit, param, trim = F, trim_amount) {
  ind <- grep(paste0("^",param), rownames(as.data.frame(summary(fit)$summary)))
  medians <- data.frame(avg = as.data.frame(summary(fit)$summary)$`50%`[ind])
  title <- paste0("Posterior Medians of ",param)
  print(ggplot(medians, aes(x = avg)) + geom_histogram(bins = 30) + ggtitle(title) +
    xlab("Medians") + ylab("Count"))
  print(" ")
  if (trim == T) {
    lim <- quantile(abs(medians$avg), probs = trim_amount)
    meds_trim <- medians %>% filter(abs(medians$avg) < lim)
    print(ggplot(meds_trim, aes(x = avg)) + geom_histogram(bins = 60) +
      ggtitle(paste0(title, " Without Extreme ",100*(1-trim_amount),"%")))
  }
  means <- data.frame(avg = as.data.frame(summary(fit)$summary)$`mean`[ind])
  title <- paste0("Posterior Means of ",param)
  print(ggplot(means, aes(x = avg)) + geom_histogram(bins = 30) + ggtitle(title) +
    xlab("Means") + ylab("Count"))
  print(" ")
  sds <- data.frame(avg = as.data.frame(summary(fit)$summary)$`sd`[ind])
  title <- paste0("Posterior Standard Deviations of ",param)
  print(ggplot(sds, aes(x = avg)) + geom_histogram(bins = 30) + ggtitle(title) +
    xlab("Standard Deviations") + ylab("Count"))
}

plot_fit <- function(fit) {
  get_single_plots(fit, tau_params)
  get_single_plots(fit, beta)
  get_aggreg_plots(fit, "w")
  get_aggreg_plots(fit, "z")
  get_aggreg_plots(fit, "p")
}

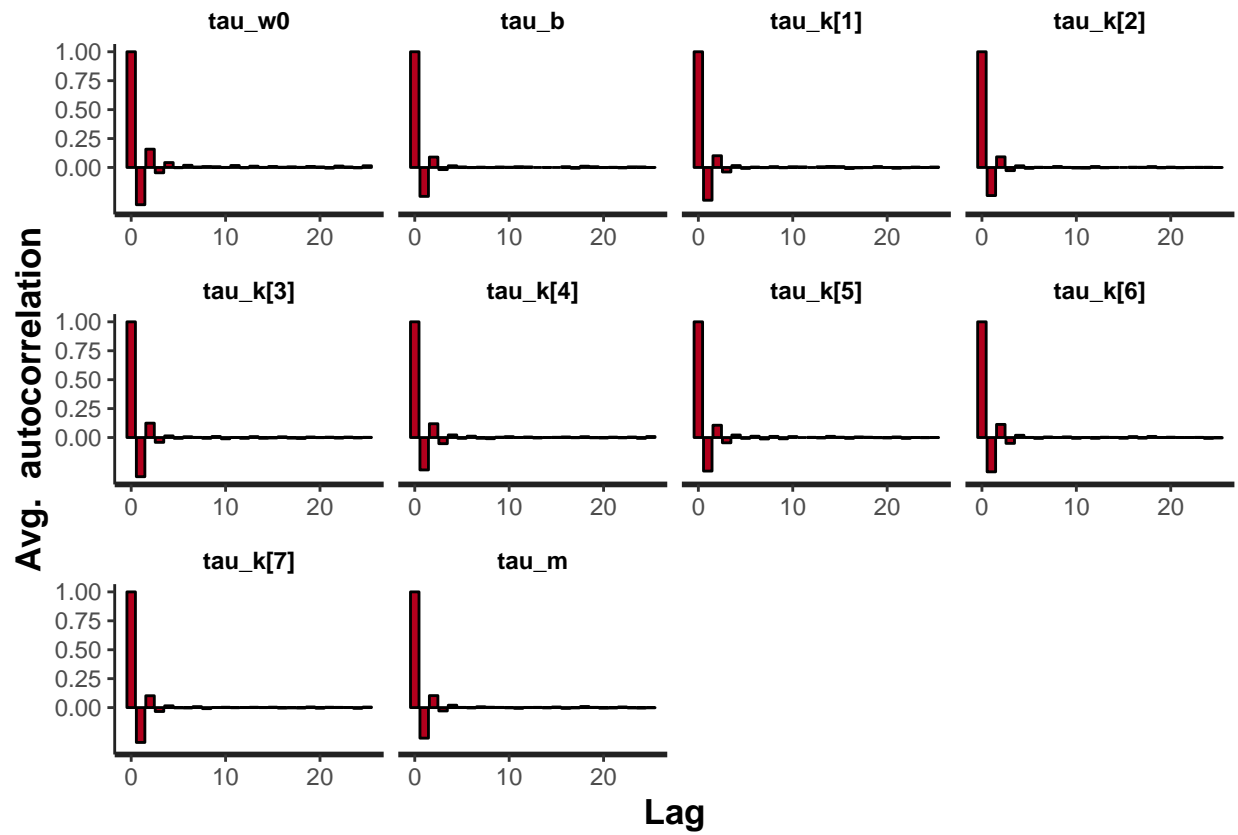
plot_fit(fit)
```

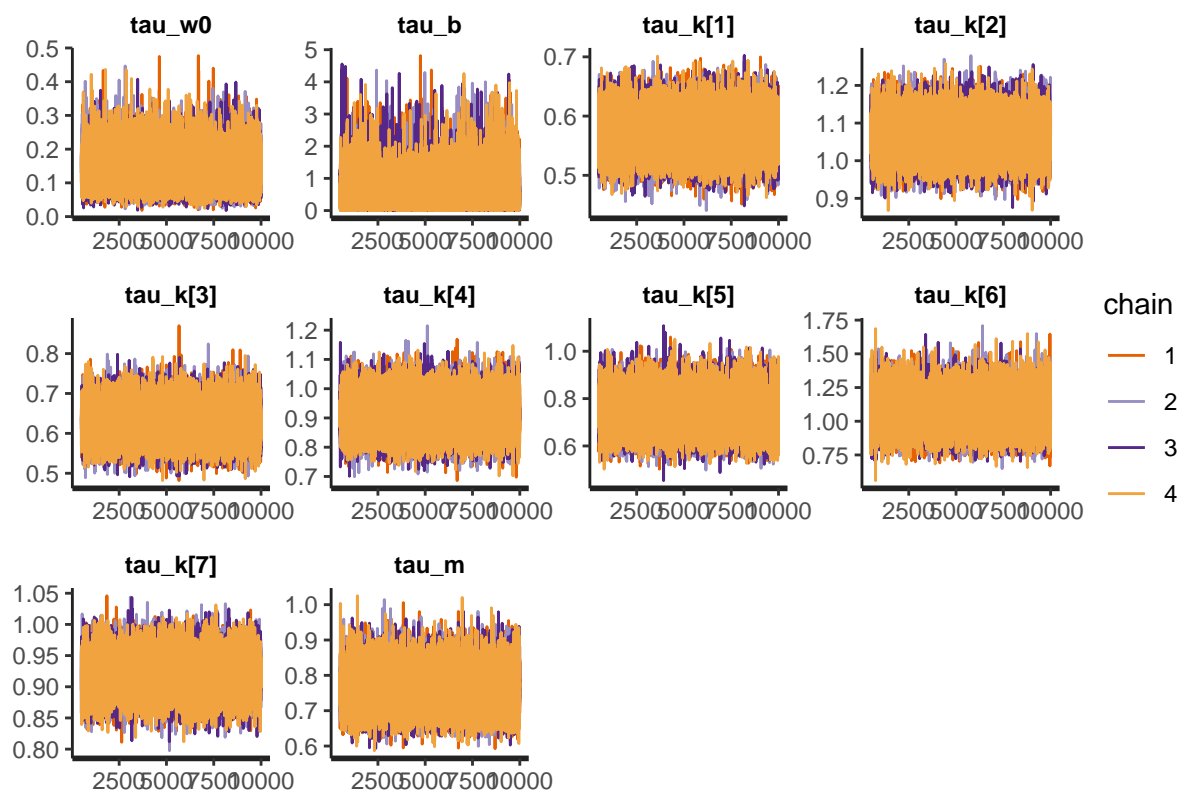
##		mean	se_mean	sd	25%	50%	75%
##	tau_w0	0.1393623	0.0004433818	0.05233179	0.1014644	0.1322487	0.1702399
##	tau_b	0.5003613	0.0023003079	0.50118804	0.1454357	0.3494985	0.6880240
##	tau_k[1]	0.5711584	0.0001315253	0.03348011	0.5483916	0.5703450	0.5933404
##	tau_k[2]	1.0653977	0.0002090100	0.04977134	1.0309337	1.0647099	1.0987120
##	tau_k[3]	0.6293228	0.0001575772	0.04265028	0.5998784	0.6280303	0.6574728
##	tau_k[4]	0.9124367	0.0002497892	0.06203567	0.8696811	0.9110412	0.9536499
##	tau_k[5]	0.7431449	0.0002882225	0.07443530	0.6918243	0.7406955	0.7915888
##	tau_k[6]	1.0656795	0.0004929857	0.12668075	0.9781905	1.0602587	1.1469348
##	tau_k[7]	0.9205138	0.0001134266	0.02926739	0.9005347	0.9201386	0.9402220
##	tau_m	0.7625909	0.0002250807	0.05309478	0.7257083	0.7601524	0.7965980
##		n_eff	Rhat				

```

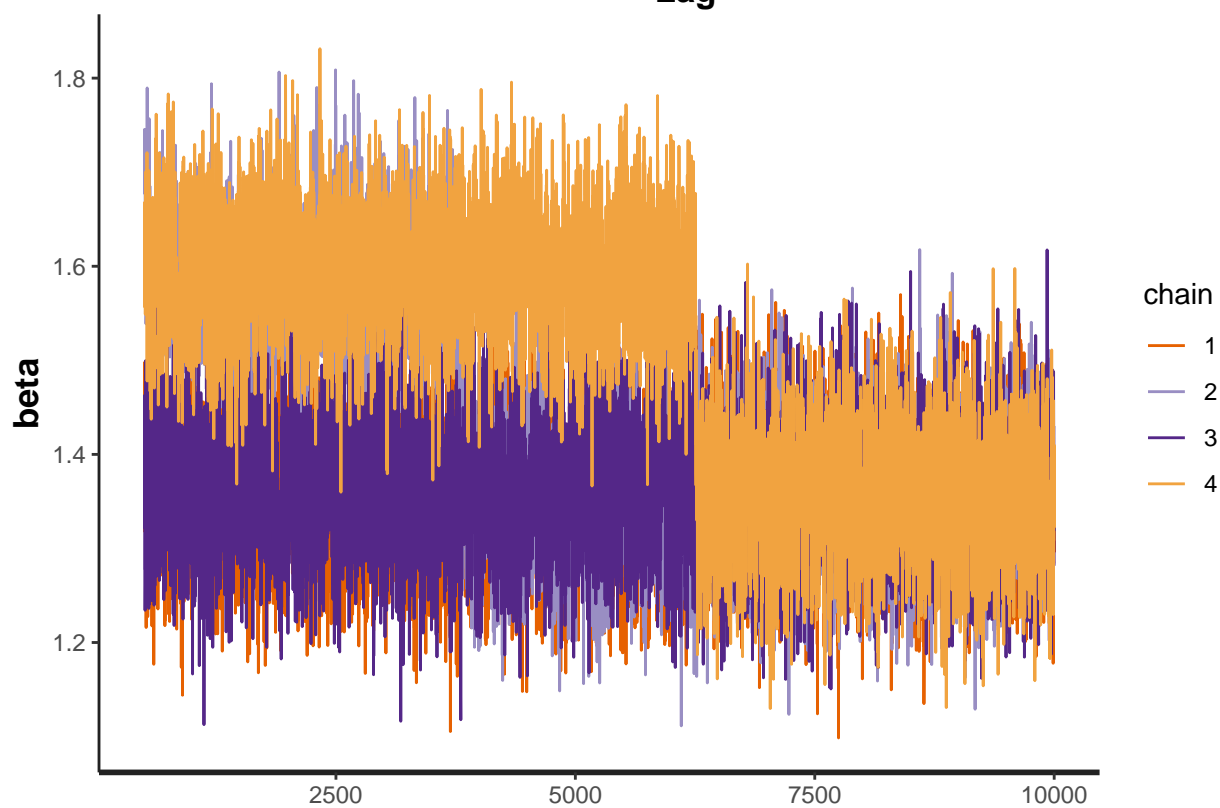
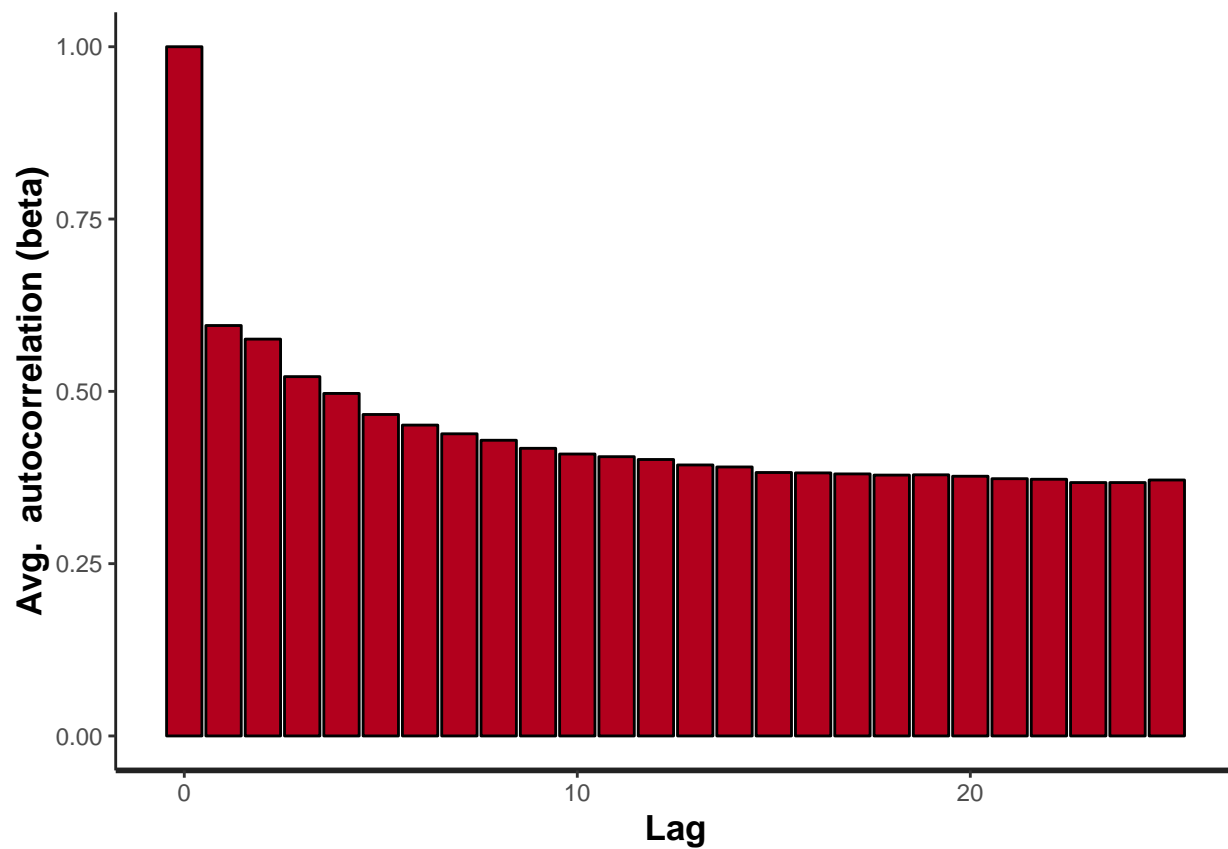
## tau_w0    13930.78 1.0030183
## tau_b     47471.12 1.0017243
## tau_k[1]   64797.08 0.9998993
## tau_k[2]   56705.43 1.0000874
## tau_k[3]   73258.29 0.9999264
## tau_k[4]   61678.78 0.9999273
## tau_k[5]   66696.35 0.9999715
## tau_k[6]   66031.73 0.9999950
## tau_k[7]   66579.13 0.9999999
## tau_m      55645.14 0.9999861

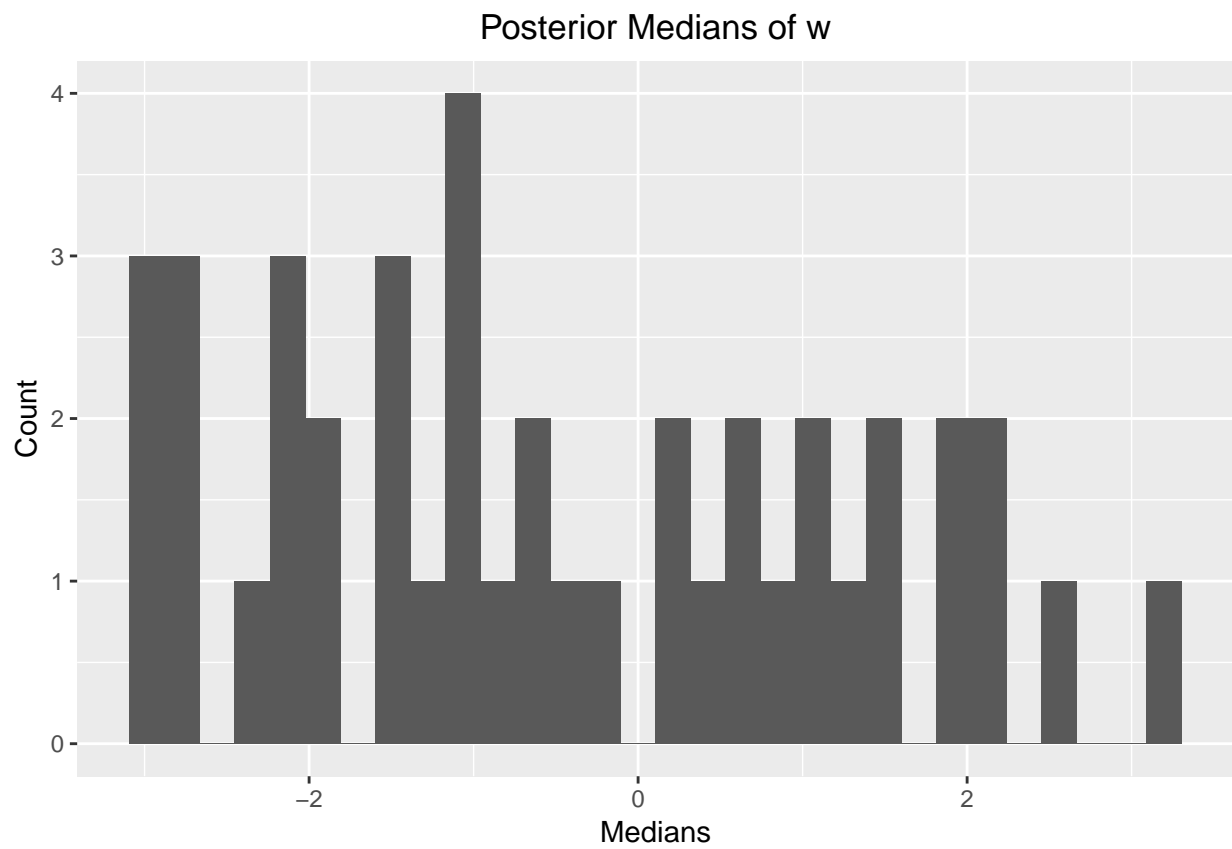
```



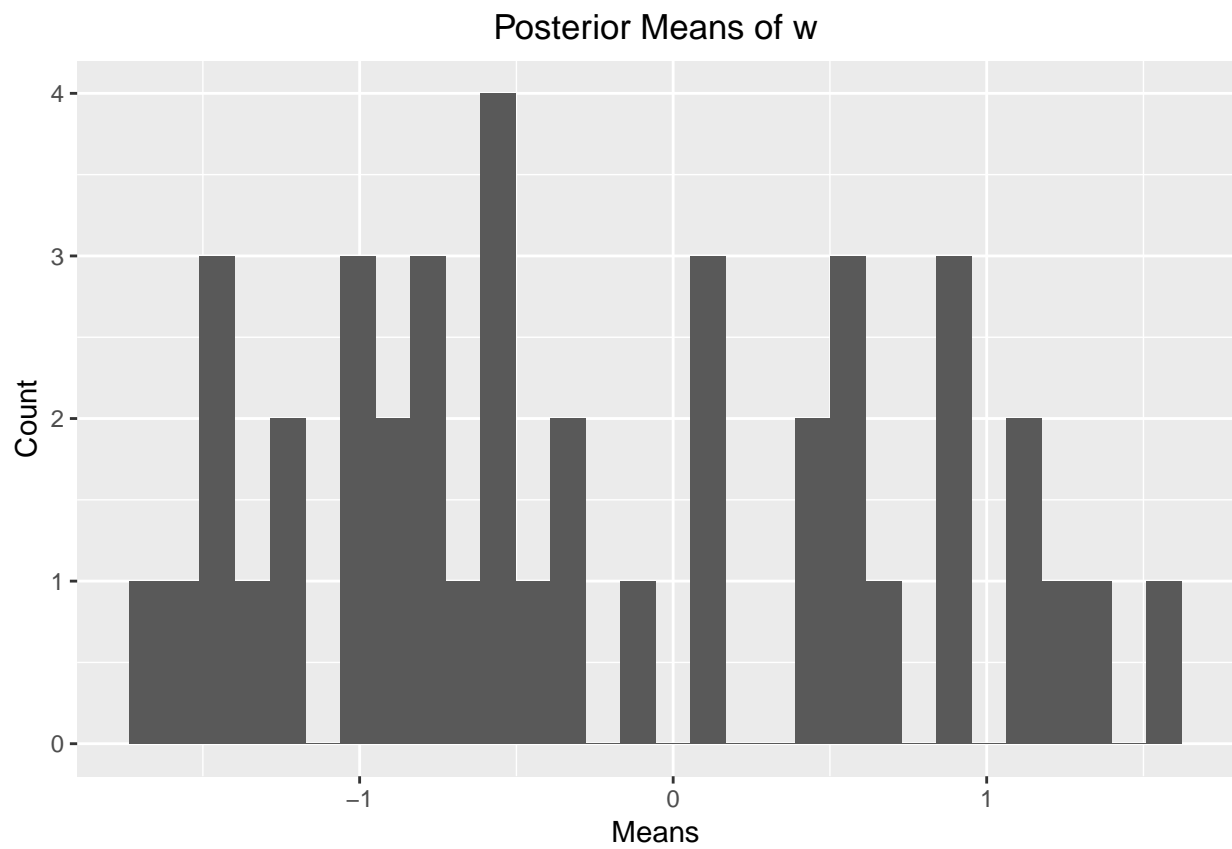


```
##          mean    se_mean      sd    25%    50%    75%   n_eff
## beta 1.41419 0.03730978 0.1193964 1.328311 1.384897 1.484532 10.24087
##          Rhat
## beta 1.482525
```

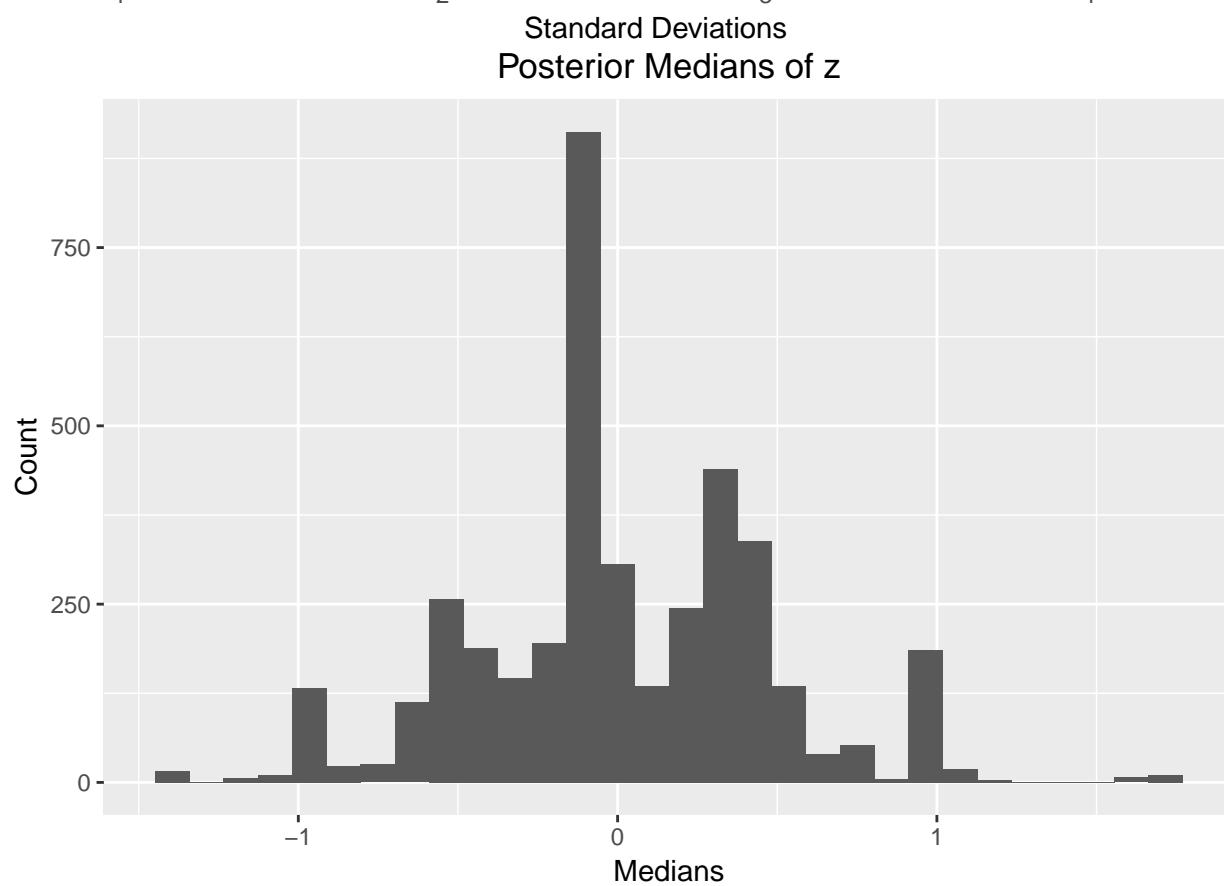
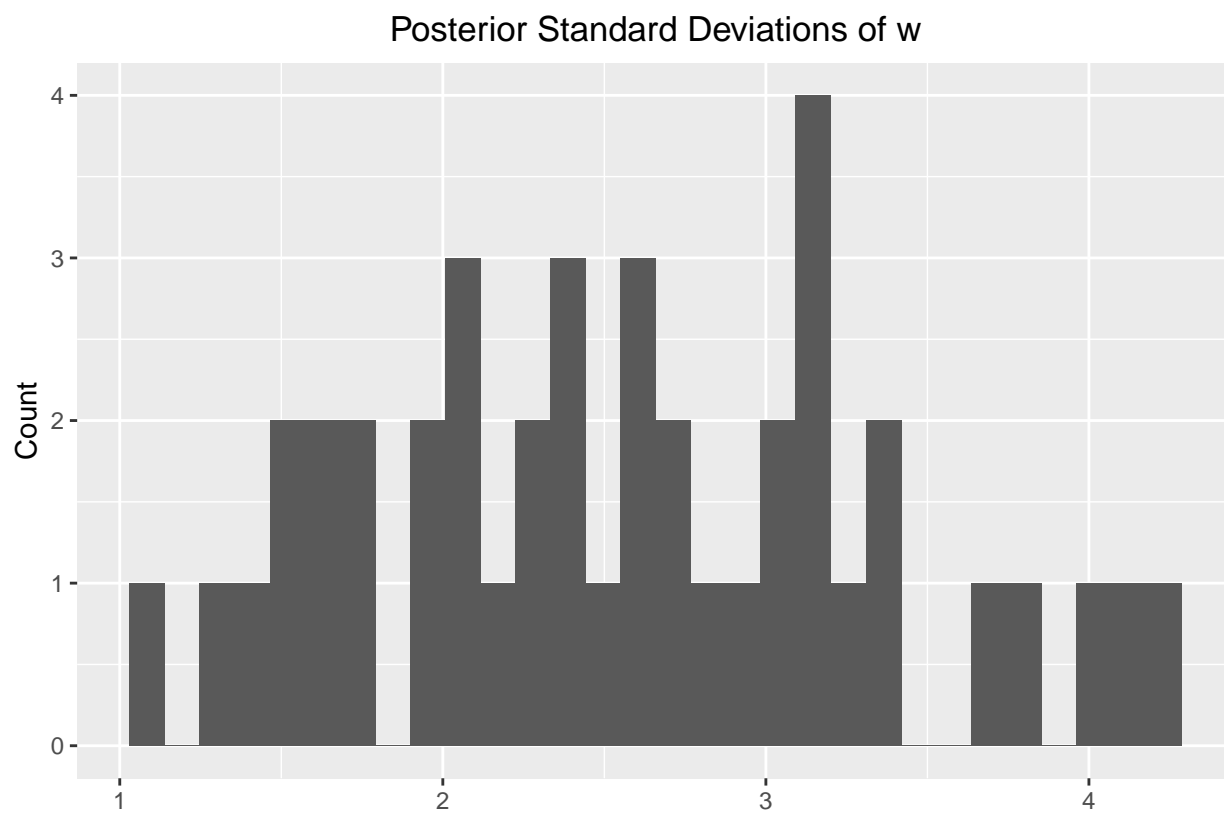




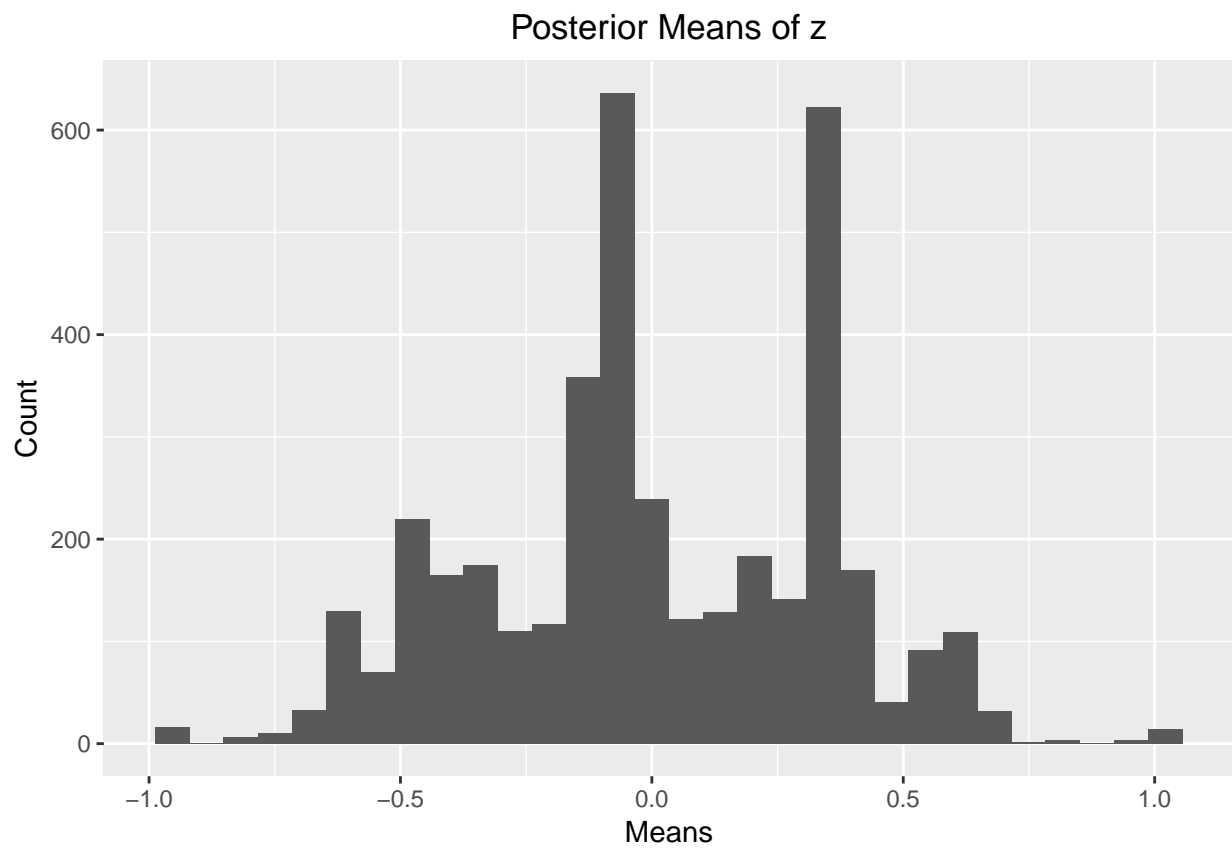
```
## [1] " "
```



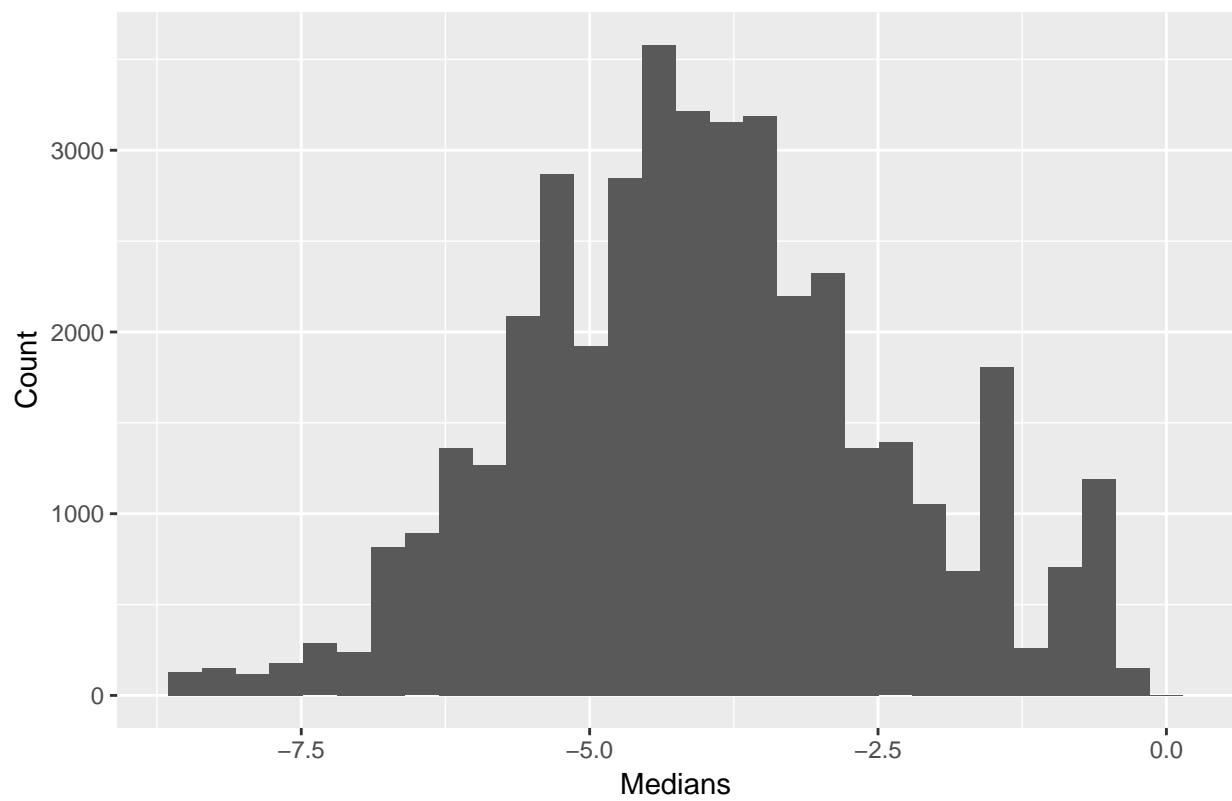
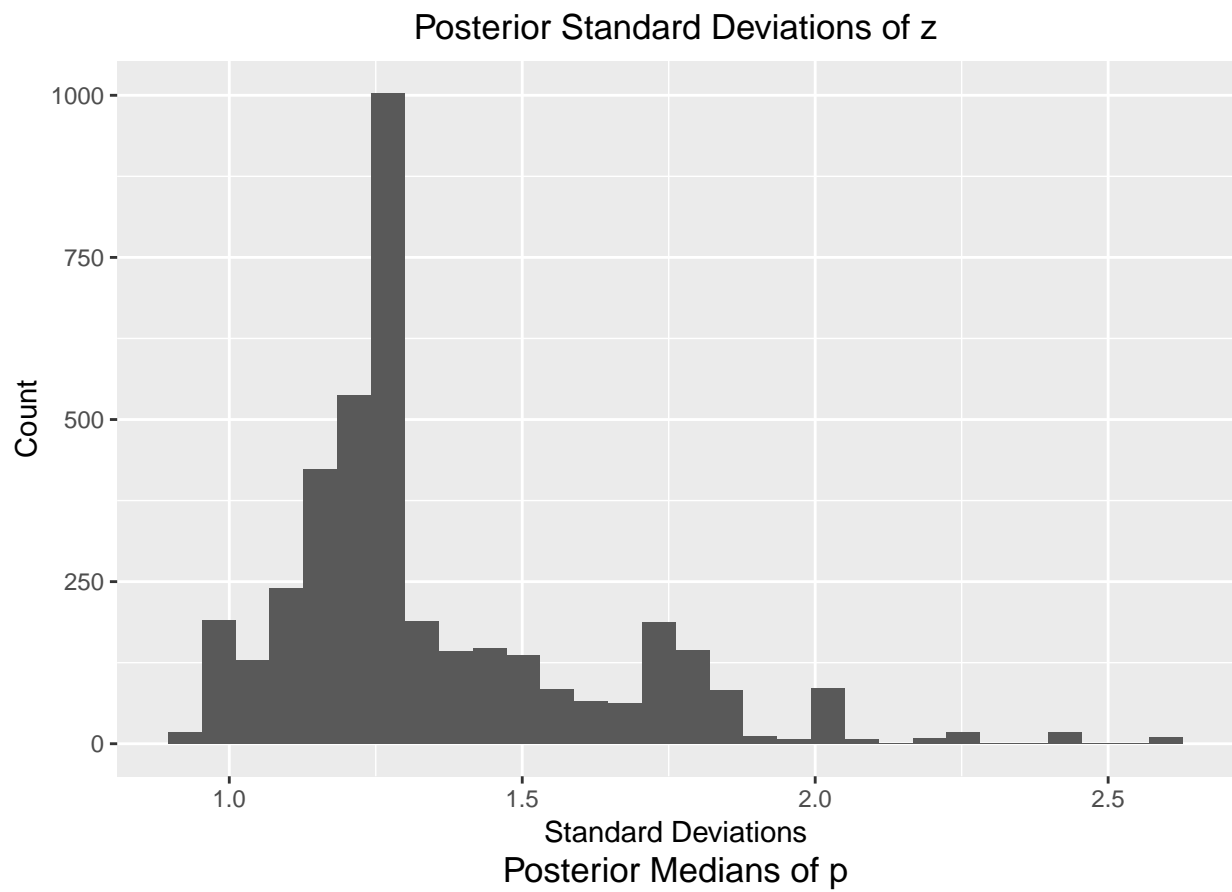
```
## [1] "    "
```

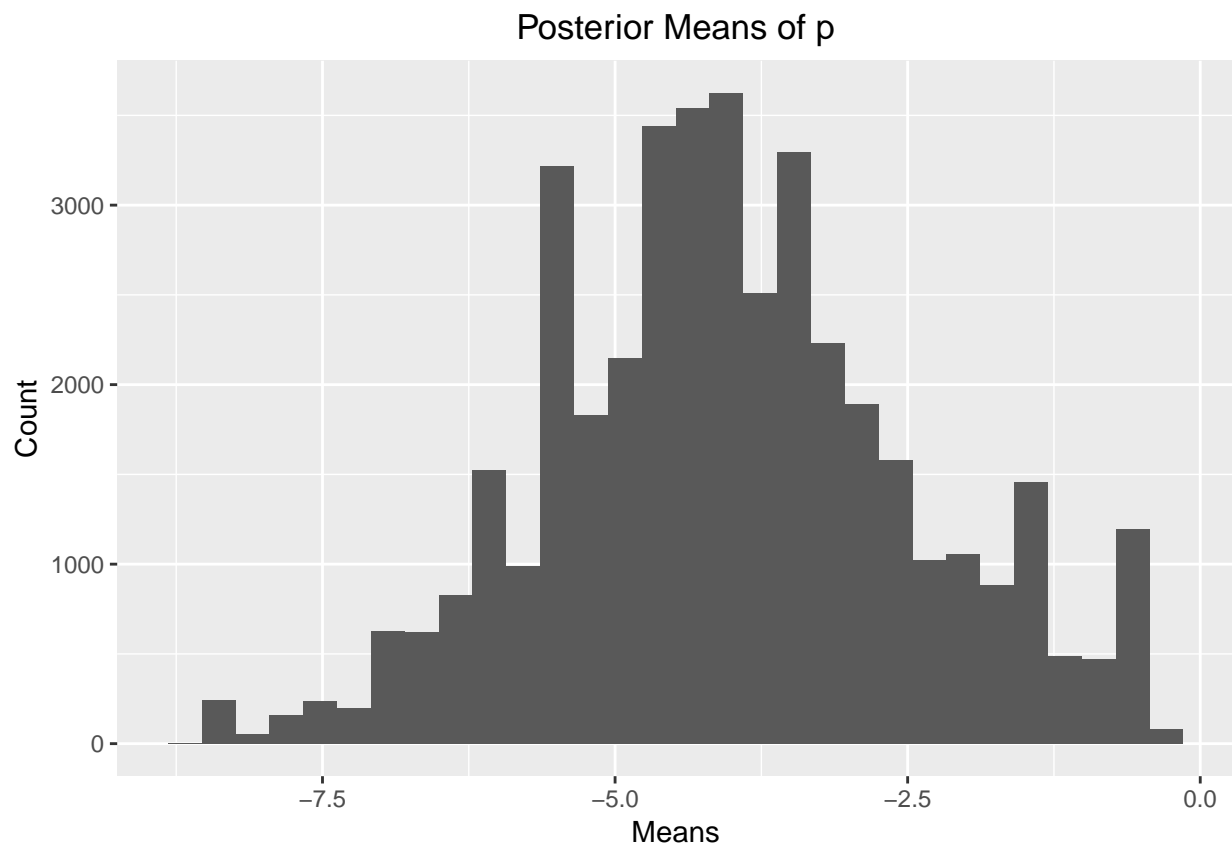
```
## [1] " "
```



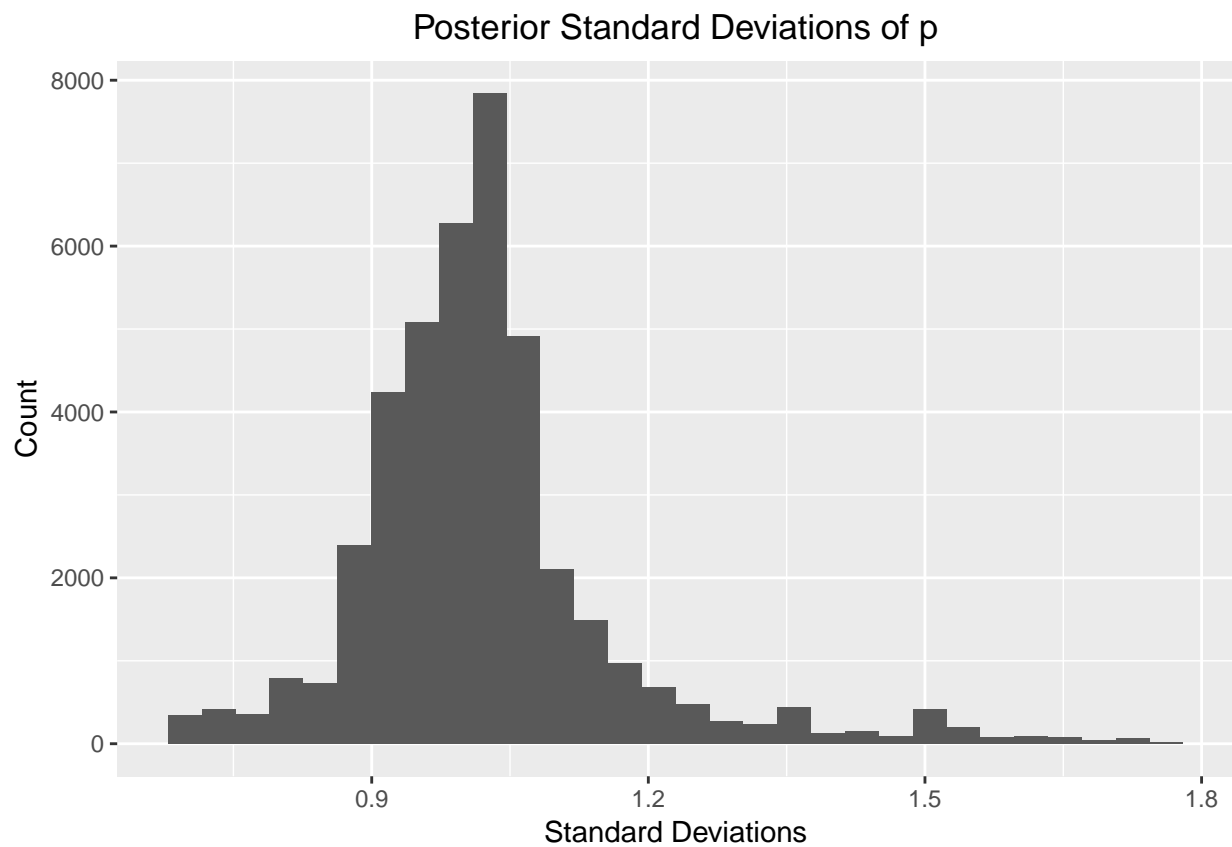
```
## [1] " "
```



```
## [1] " "
```



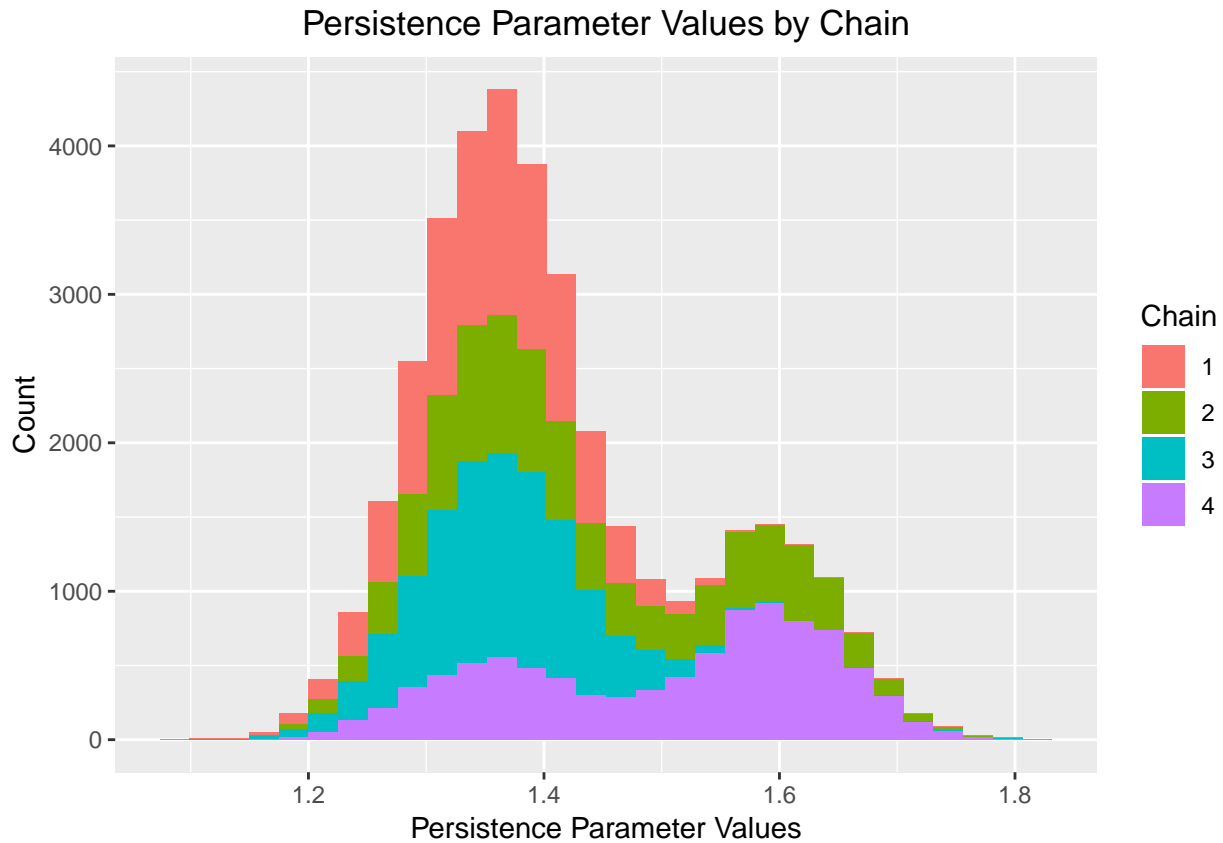
```
## [1] " "
```



Histograms for β values and w , and z posterior means across chains.

```
draws <- as.matrix(fit)
beta_col <- draws[,which(colnames(draws) == "beta")]
l <- length(beta_col)/4
plot_dat <- data.frame(val = beta_col, Chain = as.factor(c(rep(1,l),rep(2,l),rep(3,l),rep(4,l))))
ggplot(plot_dat, aes(x = val, fill = Chain)) + geom_histogram() +
  xlab("Persistence Parameter Values") + ylab("Count") + ggtitle("Persistence Parameter Values by Chain")

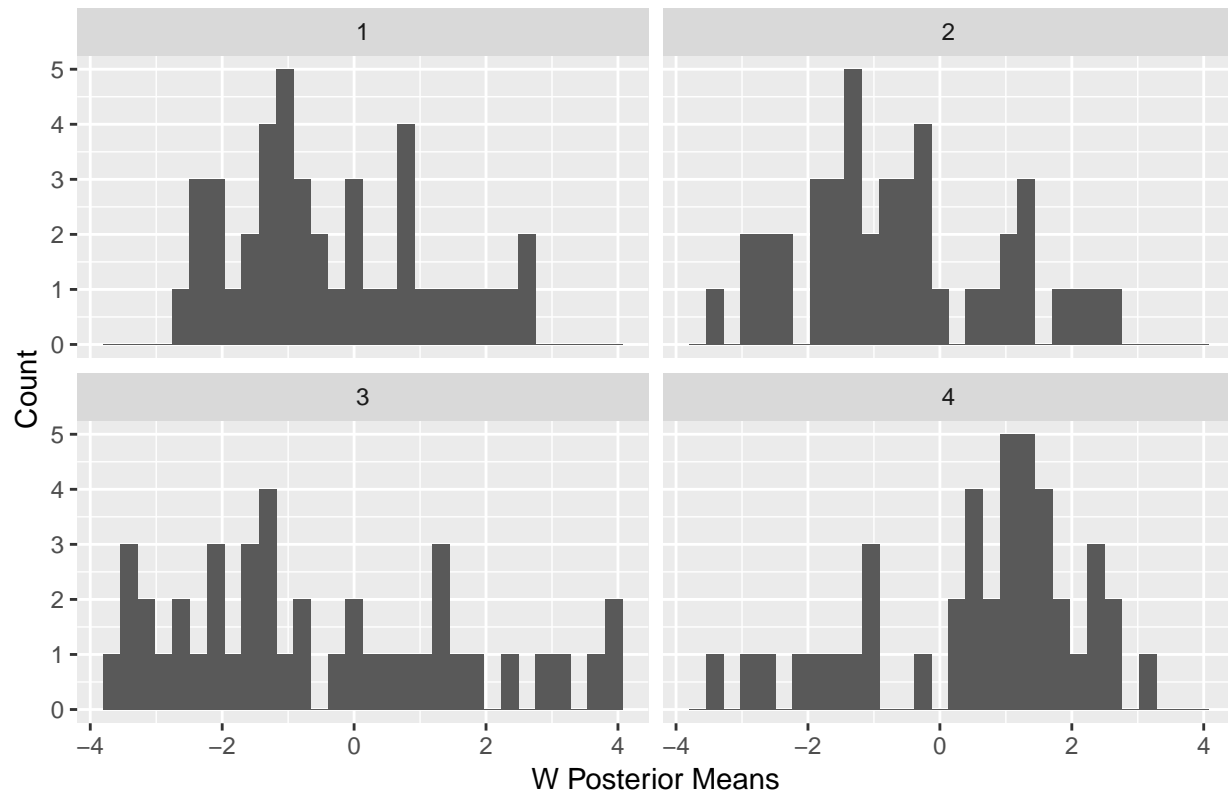
## `stat_bin()` using `bins = 30`. Pick better value with `binwidth`.
```



```
w_ind <- grep("^w", colnames(draws))
z_ind <- grep("^z", colnames(draws))
w_dat <- data.frame(chain1 = colMeans(draws[1:l,w_ind]),
                   chain2 = colMeans(draws[(l+1):(2*l),w_ind]),
                   chain3 = colMeans(draws[(2*l+1):(3*l),w_ind]),
                   chain4 = colMeans(draws[(3*l+1):(4*l),w_ind])) %>%
  pivot_longer(cols = everything(), names_to = "Chain", names_prefix = "chain") %>%
  mutate(Chain = as.factor(Chain))
z_dat <- data.frame(chain1 = colMeans(draws[1:l,z_ind]),
                   chain2 = colMeans(draws[(l+1):(2*l),z_ind]),
                   chain3 = colMeans(draws[(2*l+1):(3*l),z_ind]),
                   chain4 = colMeans(draws[(3*l+1):(4*l),z_ind])) %>%
  pivot_longer(cols = everything(), names_to = "Chain", names_prefix = "chain") %>%
  mutate(Chain = as.factor(Chain))
ggplot(w_dat, aes(x = value)) +
  geom_histogram() + facet_wrap(~Chain) + xlab("W Posterior Means") + ylab("Count") +
  ggtitle("W Posterior Means by Chain")
```

```
## `stat_bin()` using `bins = 30`. Pick better value with `binwidth`.
```

W Posterior Means by Chain



```
ggplot(z_dat, aes(x = value)) +
  geom_histogram() + facet_wrap(~Chain) +
  xlab("Z Posterior Means") + ylab("Count") +
  ggtitle("Z Posterior Means by Chain")
```

`stat_bin()` using `bins = 30`. Pick better value with `binwidth`.

Z Posterior Means by Chain

