

Bias

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```
dat <- as.data.table(readRDS(file.path(base_dir, dat_path, paste0(base_nm, '_tmp'), 'coverage.rds')))
```

Coverage

Percentile

```
tmp <- dat[, .(lower = boot::perc.ci(boot_reps)[4], upper = boot::perc.ci(boot_reps)[5]),  
  by = c('subset_num', 'replication', 'n'))[,  
  .(lower = mean(lower), upper = mean(upper)), by = c('replication', 'n')  
  ]  
tmp[, .(coverage = mean(lower <= te & upper >= te)), by = c('n')]
```

```
##          n coverage  
## 1: 20000    0.884  
## 2: 50000    0.864
```

Normal

```
tmp <- dat[, .(lower = mean(boot_reps) - 1.96*sd(boot_reps), upper = mean(boot_reps) + 1.96*sd(boot_reps)),  
  by = c('subset_num', 'replication', 'n'))[,  
  .(lower = mean(lower), upper = mean(upper)), by = c('replication', 'n')  
  ]  
tmp[, .(coverage = mean(lower <= te & upper >= te)), by = c('n')]
```

```
##          n coverage  
## 1: 20000    0.876  
## 2: 50000    0.850
```

Bias

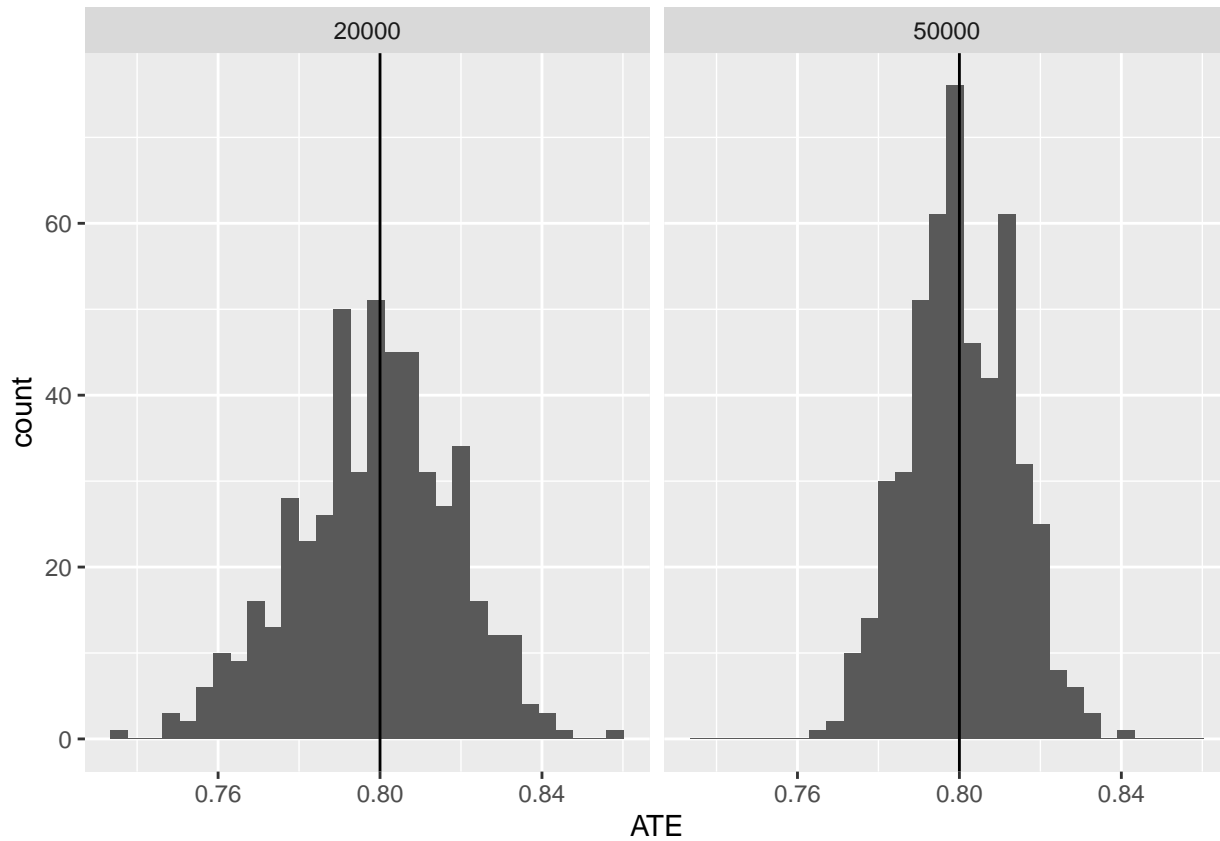
```
true_sd <- dat[, .(SD_full = mean(sd_full)), by = c('n')]
```

ATE

```
bias <- dat[, .(ATE = mean(boot_reps),  
  SE = sd(boot_reps)), by = c('subset_num', 'replication', 'n'))[,  
  .(ATE = mean(ATE), SE = mean(SE)), by = c('replication', 'n')  
  ]  
  
ggplot(bias, aes(x = ATE)) +
```

```
geom_histogram() +  
facet_wrap(~ n) +  
geom_vline(xintercept = te)
```

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



Standard Error

```
ggplot(bias, aes(x = SE)) +  
geom_histogram() +  
facet_wrap(~ n) +  
geom_vline(data = true_sd, aes(xintercept = sqrt(SD_full)))
```

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

