Untitled

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Loading packages

```
library("ggplot2")
library("ggpubr")

#set.seed(159)
```

Loading data

```
samp25_mean <- replicate(50, mean(rnorm(25, 0, 1)))
samp200_mean <- replicate(50, mean(rnorm(200, 0, 1)))
samp500_mean <- replicate(50, mean(rnorm(500, 0, 1)))
samp1000_mean <- replicate(50, mean(rnorm(1000, 0, 1)))
samp10000_mean <- replicate(50, mean(rnorm(10000, 0, 1)))</pre>
```

Loading plots

```
p1 <- ggplot(NULL, aes(x = samp25_mean, ..density..)) + geom_histogram(binwidth = 0.05) + xlim(c(-.5, ... p2 <- ggplot(NULL, aes(x = samp200_mean, ..density..)) + geom_histogram(binwidth = 0.035) + xlim(c(-.35 p3 <- ggplot(NULL, aes(x = samp500_mean, ..density..)) + geom_histogram(binwidth = 0.025) + xlim(c(-0.2 p4 <- ggplot(NULL, aes(x = samp1000_mean, ..density..)) + geom_histogram(binwidth = 0.015) + xlim(c(-.1 p5 <- ggplot(NULL, aes(x = samp10000_mean, ..density..)) + geom_histogram(binwidth = 0.005) + xlim(c(-... ps <- ggplot(NULL, aes(x = samp10000_mean, ..density..)) + geom_histogram(binwidth = 0.005) + xlim(c(-... ps <- ggplot(NULL, aes(x = samp10000_mean, ..density..)) + geom_histogram(binwidth = 0.005) + xlim(c(-... ps <- ggplot(NULL, aes(x = samp10000_mean, ..density..)) + geom_histogram(binwidth = 0.005) + xlim(c(-... ps <- ggplot(NULL, aes(x = samp10000_mean, ..density..)) + geom_histogram(binwidth = 0.005) + xlim(c(-... ps <- ggplot(NULL, aes(x = samp10000_mean, ..density..)) + geom_histogram(binwidth = 0.005) + xlim(c(-... ps <- ggplot(NULL, aes(x = samp10000_mean, ..density..)) + geom_histogram(binwidth = 0.005) + xlim(c(-... ps <- ggplot(NULL, aes(x = samp10000_mean, ..density..)) + geom_histogram(binwidth = 0.005) + xlim(c(-... ps <- ggplot(NULL, aes(x = samp10000_mean, ..density..)) + geom_histogram(binwidth = 0.005) + xlim(c(-... ps <- ggplot(NULL, aes(x = samp10000_mean, ..density..)) + geom_histogram(binwidth = 0.005) + xlim(c(-... ps <- ggplot(NULL, aes(x = samp1000_mean, ..density..)) + geom_histogram(binwidth = 0.005) + xlim(c(-... ps <- ggplot(NULL, aes(x = samp1000_mean, ..density..)) + geom_histogram(binwidth = 0.005) + xlim(c(-... ps <- ggplot(NULL, aes(x = samp1000_mean, ..density..)) + geom_histogram(binwidth = 0.005) + xlim(c(-... ps <- ggplot(NULL, aes(x = samp1000_mean, ..density..)) + geom_histogram(binwidth = 0.005) + xlim(c(-... ps <- ggplot(NULL, aes(x = samp1000_mean, ..density..)) + geom_histogram(binwidth = 0.005) + xlim(c(-... ps <- ggplot(NULL, aes(x = samp1000_mean, .
```

Combining plots

```
ggarrange(p1, p2, p3, p4, p5)

## Warning: Removed 2 rows containing missing values (geom_bar).
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## Warning: Removed 1 rows containing non-finite values (stat_bin).
## Warning: Removed 2 rows containing missing values (geom_bar).
## Removed 2 rows containing missing values (geom_bar).
```

