Research Design, Fall 2021

04: a bit more probability review; sampling; statistical tests

Scott Spencer | Columbia University

general feedback for research reviews

Title: homework 1

This paper described A and …

Title: ~~homework 1~~

Why should I be reading this document? What is the main point?

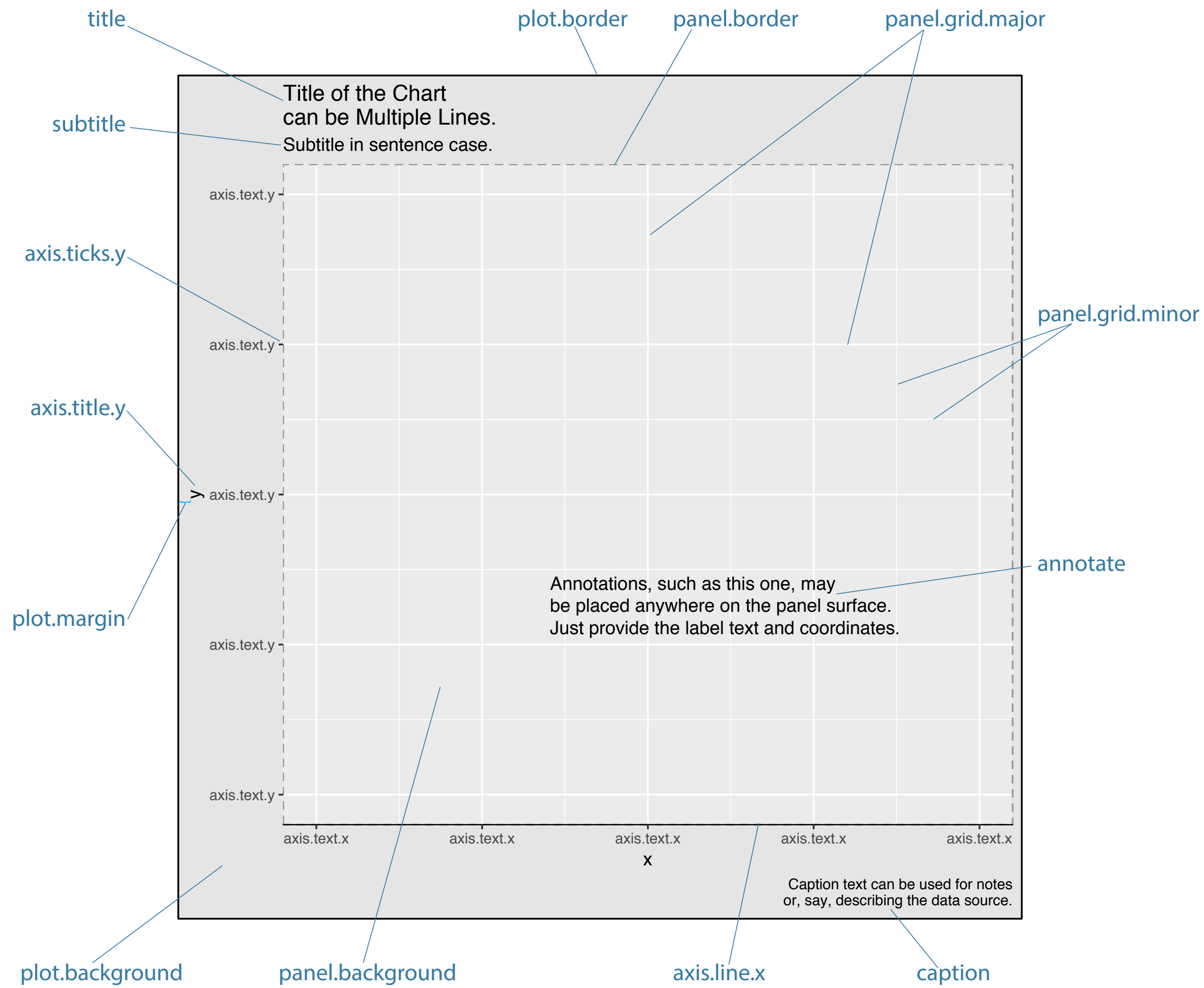The authors of ~~this paper~~ described A and …

To what paper are you referring? Always introduce new ideas before using language that refers back. And properly cite your sources.

Also review:

Harris, Joseph. Rewriting: *How to Do Things with Texts*.
Second edition. Logan: Utah State University Press, 2017.

tips on graphics as a tool for exploring and communicating research design

An aside on graphics to communicate research designs,

title

plot.border   panel.border   panel.grid.major

subtitle

Title of the Chart
can be Multiple Lines.

Subtitle in sentence case.

axis.text.y

axis.ticks.y

panel.grid.minor

axis.text.y

axis.title.y

> axis.text.y

Annotations, such as this one, may
be placed anywhere on the panel surface.
Just provide the label text and coordinates.

annotate

plot.margin

axis.text.y

axis.text.y

axis.text.x   axis.text.x   axis.text.x   axis.text.x   axis.text.x

x

Caption text can be used for notes
or, say, describing the data source.

plot.background   panel.background   axis.line.x   caption

Coding graphic elements, *example in R/GGplot2*

```r
# load grammar of graphics
library(ggplot2)


p <-

  # functions for data ink

  ggplot(data = <data>,
         mapping = aes(<aesthetic> = <variable>,
                       <aesthetic> = <variable>,
                       <...> = <...>) +
  geom_<type>(<...>) +
  scale_<mapping>_<type>(<...>) +
  coord_<type>(<...>) +
  facet_<type>(<...>) +
  <...> +


  # functions for non-data ink                element_blank()
                                              element_line(<...> = <...>)
  labs(<...>) +                               element_rect(<...> = <...>)
  theme(<...> = <...>) +                      element_text(<...> = <...>)
  annotate(<...>) +
  <...>
```

Drawing functions,

```r
stat_function(
  fun  = <function name>,
  args = list(parameter1 = <...>, parameter2 = <...>),
  geom = "<type>",
  ...)
```

a bit more statistics and probability

R's probability functions, **p**robability density (PDF), cumulative **d**istribution (CDF), **q**uantile, **r**andom generation

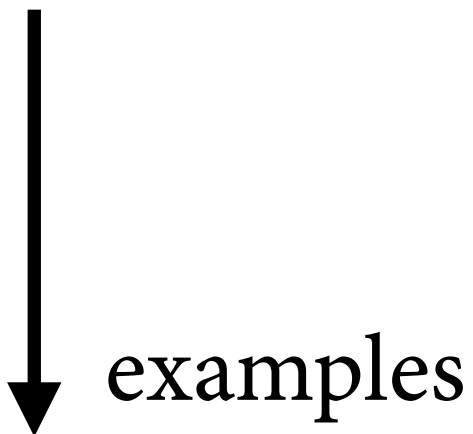**p**\<probability function name\>   **p**robability density function

d\<probability function name\>   cumulative **d**istribution function

q\<probability function name\>   **q**uantile function

r\<probability function name\>   **r**andom generation function

↓ examples

normal	*negative binomial*
student t	*gamma*
bernoulli	*cauchy*
binomial	*100s more*
poisson	*...*

law of large numbers, as $n \to \infty, \bar{x} \longrightarrow \mu$

```
set.seed(29914)

n     <- seq(1, 10000, by = 25)
mu    <- 0
sigma <- 1
x_bar <- sapply(n, FUN = function(n) {mean( rnorm(n, mu, sigma) )} )
```

```
library(ggplot2); library(ggthemes); library(latex2exp)
theme_set( theme_tufte(base_family = "sans") )

ggplot() +
  scale_x_continuous(labels = scales::comma) +
  geom_hline(yintercept = mu, color = "pink", lwd = 2) +
  geom_point(aes(n, x_bar), size = 0.5, color = "dodgerblue") +
  geom_line(aes(n, x_bar), alpha = 0.2, color = "dodgerblue") +
  annotate("text", 0, mu, hjust = 1, size = 16/.pt, label = TeX("$\\mu$") ) +
  labs(x = "sample size, n", y = TeX("sample mean, $\\bar{x}$"))
```
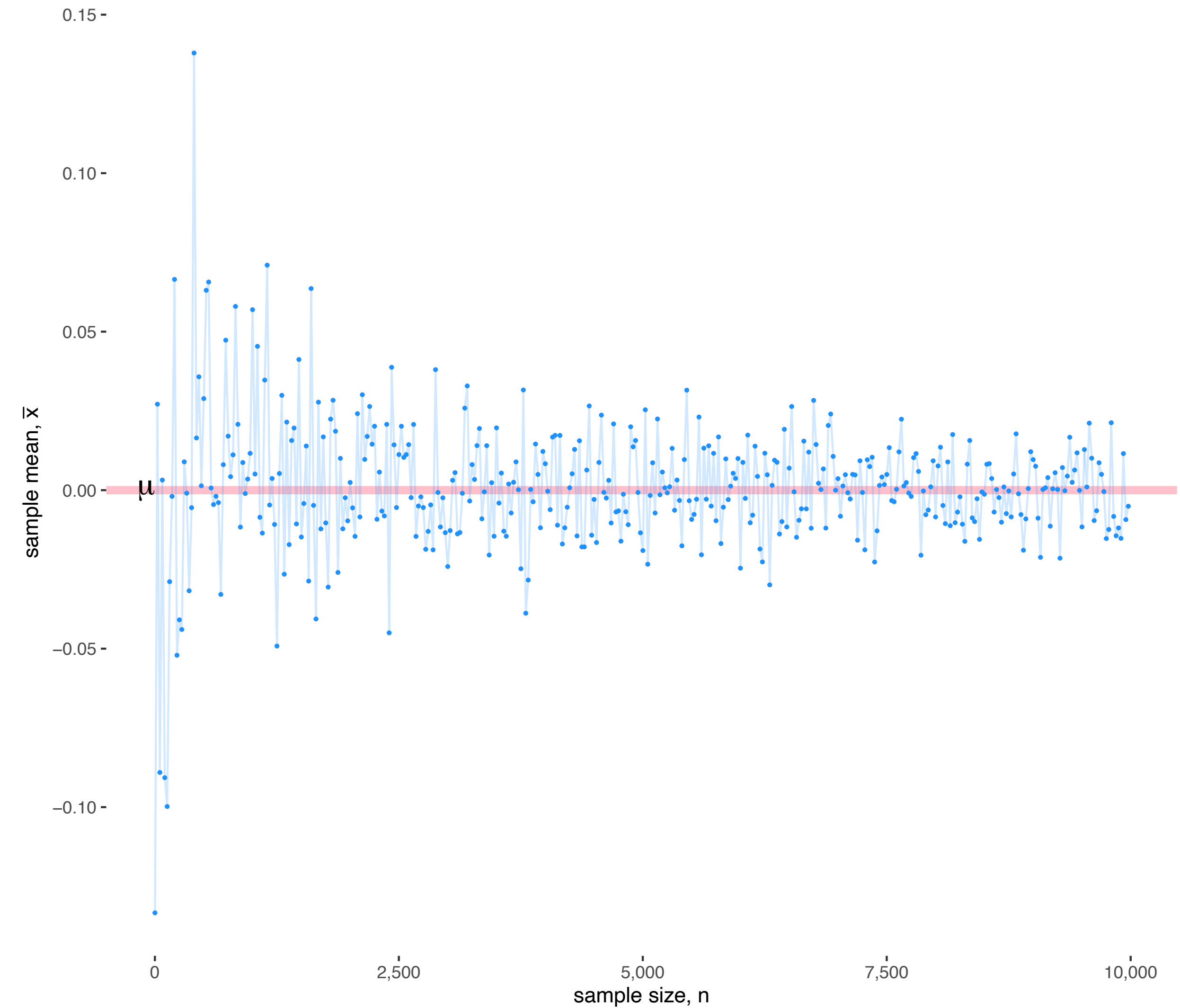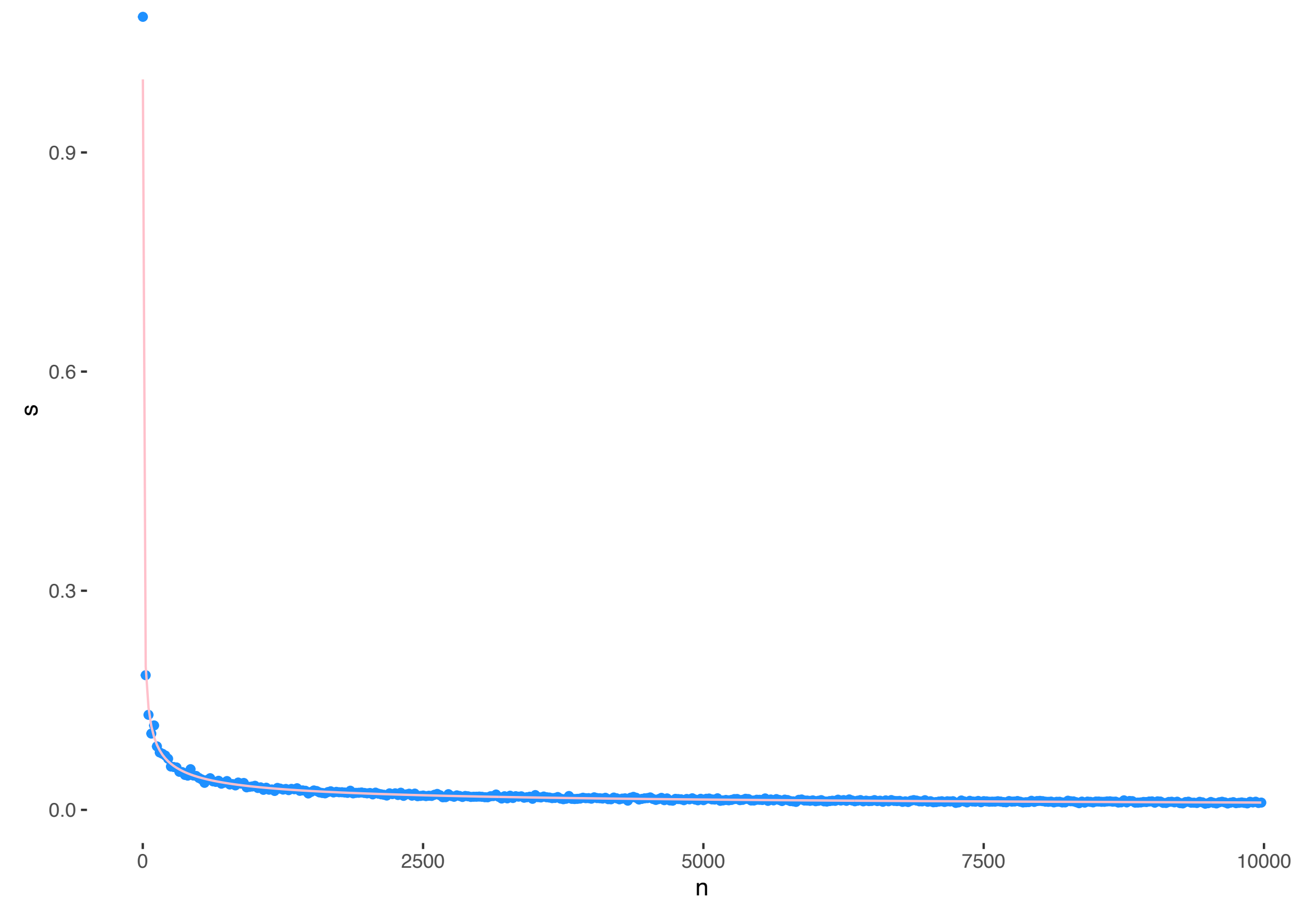
Experiment with different values of n, μ, and σ to see how $\bar{x}$ compares with μ.

Experiment with different seed values and compare results. How would you describe the uncertainty of $\bar{x}$?

central limit theorem, **standard deviation** s of $\bar{x} = \dfrac{\sigma}{\sqrt{n}}$
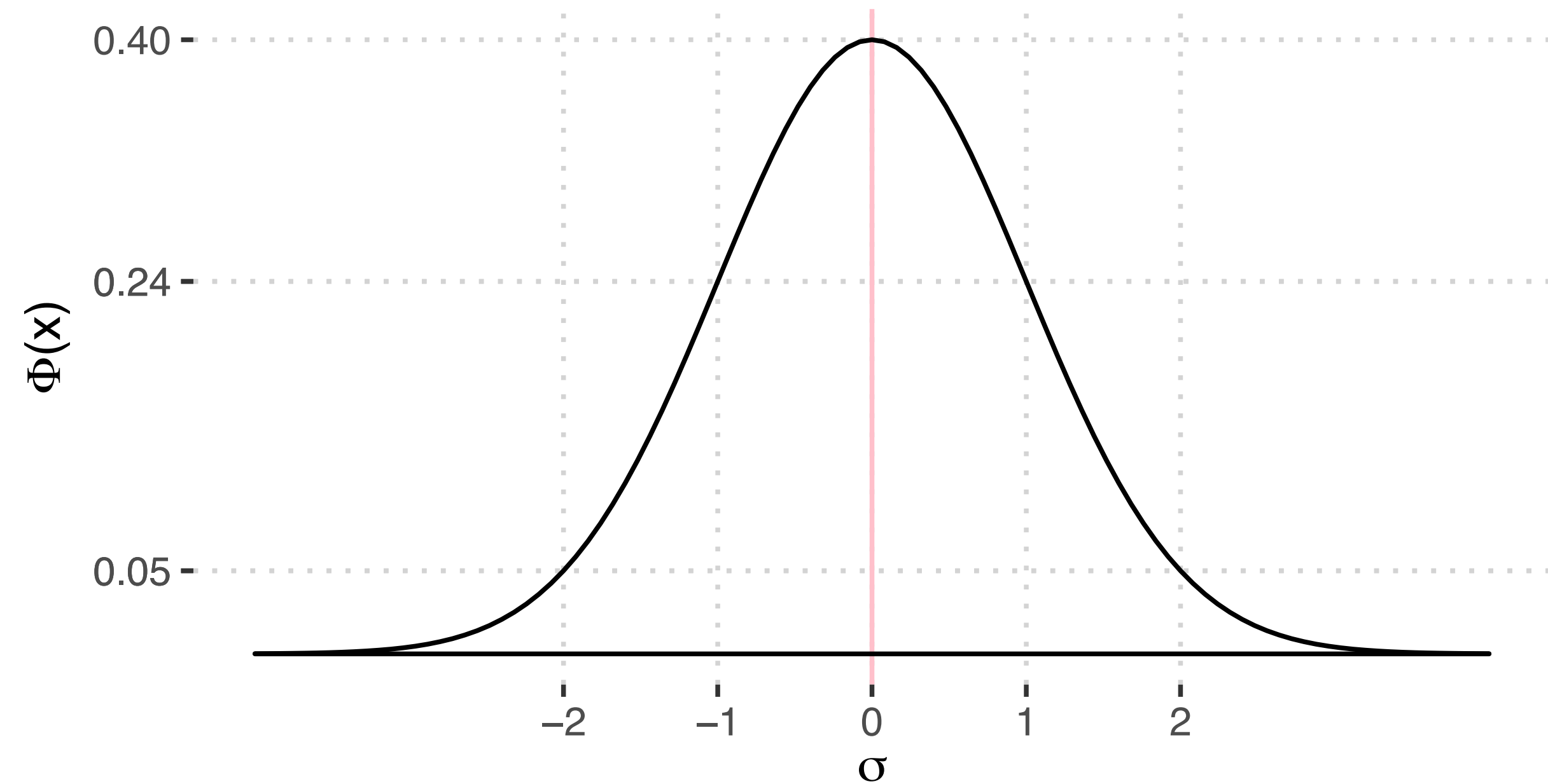
```r
sample_mean <- function() {
  sapply(n, FUN = function(n) {mean( rnorm(n, mu, sigma) )} )
  }

x_bar <- replicate(1000, sample_mean() )
s <- apply(x_bar, 1, sd)

ggplot() +
  geom_point(aes(n, s), color = "dodgerblue") +
  geom_line(aes(n, sigma / sqrt(n) ), color = "pink")
```


Experiment with different values of σ and μ and
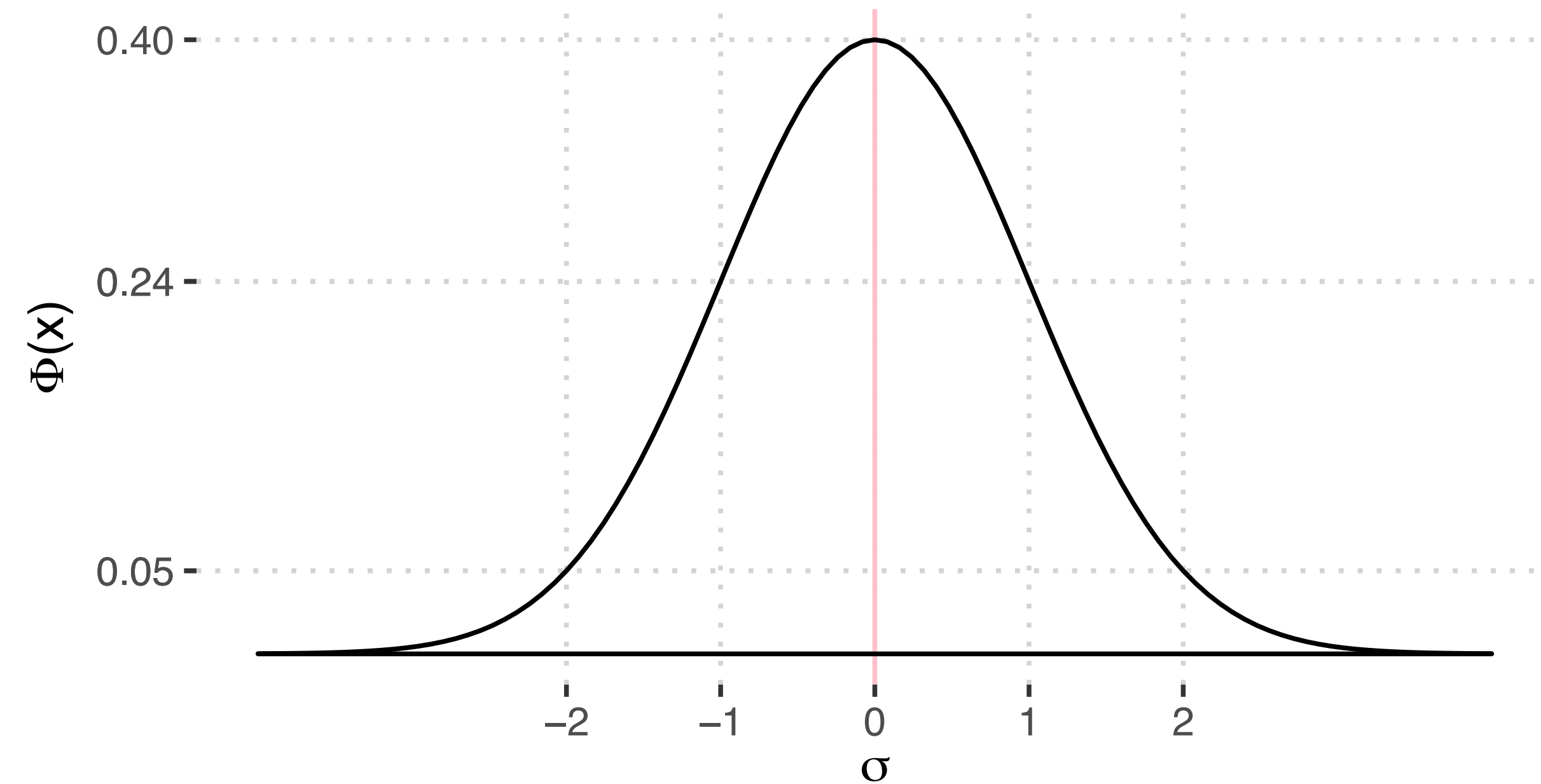replications to check the relationships empirically.

```r
pdf <- ggplot() +
  theme(panel.grid.major = element_line(color = "lightgray", linetype = "dotted")) +
  scale_x_continuous(breaks = seq(-2, 2)) +
  scale_y_continuous(breaks = dnorm(seq(-2, 2)), labels = scales::comma) +
  geom_vline(xintercept = 0, color = "pink") +
stat_function(fun  = dnorm,
              args = list(mean = 0, sd = 1),
              geom = "density",
              xlim = c(-4,4)) +
labs(x = TeX("$\\sigma$"), y = TeX("$\\Phi(x)$"))
```

```
pdf <- ggplot() +
  theme(panel.grid.major = element_line(color = "lightgray", linetype = "dotted")) +
  scale_x_continuous(breaks = seq(-2, 2)) +
  scale_y_continuous(breaks = dnorm(seq(-2, 2)), labels = scales::comma) +
  geom_vline(xintercept = 0, color = "pink") +
  stat_function(fun  = dnorm,
                args = list(mean = 0, sd = 1),
                geom = "density",
                xlim = c(-4,4)) +
  labs(x = TeX("$\\sigma$"), y = TeX("$\\Phi(x)$"))
```

```
cdf <- ggplot() +
  theme(panel.grid.major = element_line(color = "lightgray", linetype = "dotted")) +
  scale_x_continuous(breaks = seq(-2, 2)) +
  scale_y_continuous(breaks = pnorm(seq(-2, 2)),
                     labels = scales::label_percent(accuracy = 1)) +
  geom_vline(xintercept = 0, color = "pink") +
  stat_function(fun  = pnorm,
                args = list(mean = 0, sd = 1),
                geom = "density",
                xlim = c(-4,4)) +
  labs(x = TeX("$\\sigma$"), y = TeX("$F_\\Phi(x)$"))
```
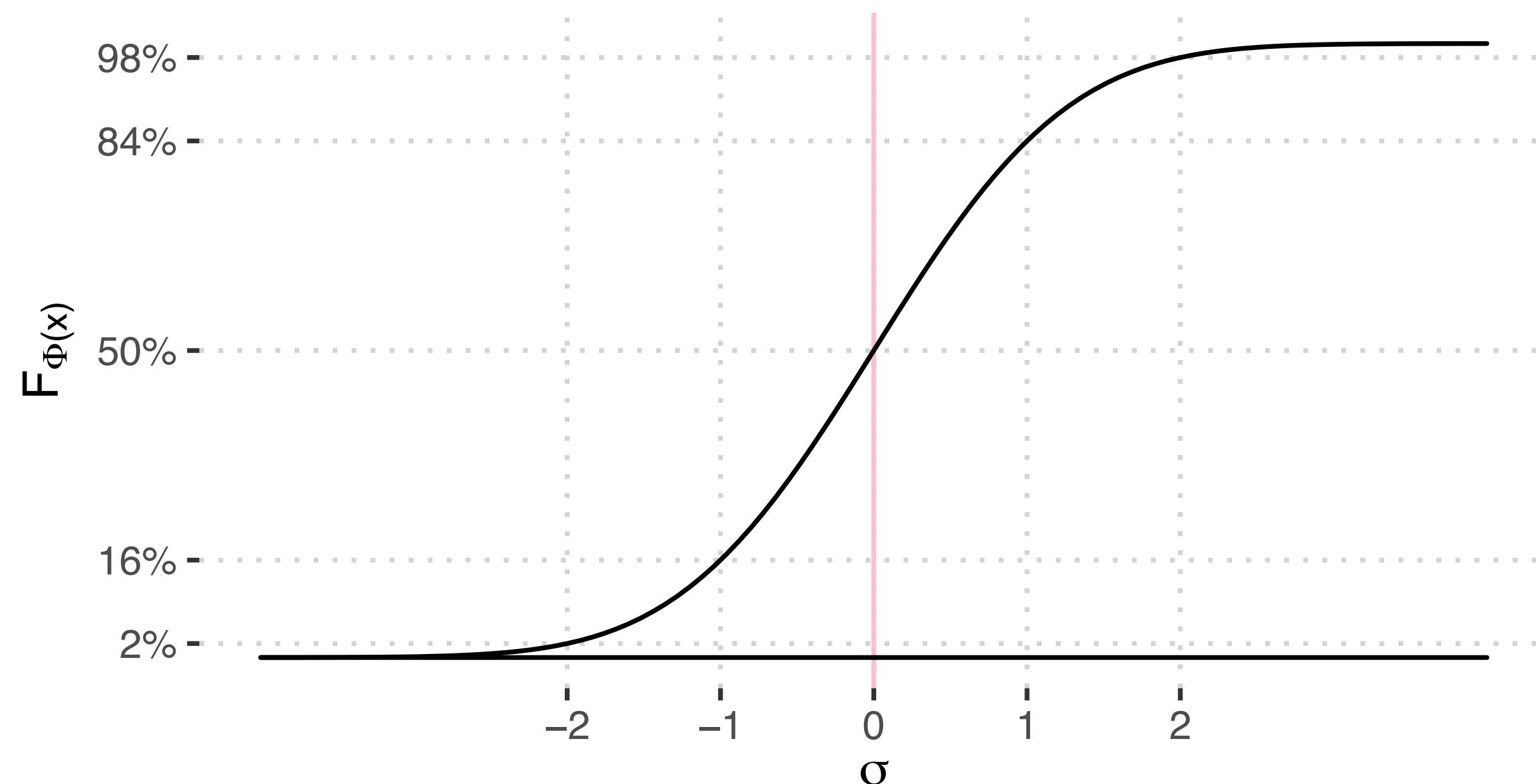
more on sampling

sampling, a few of many approaches

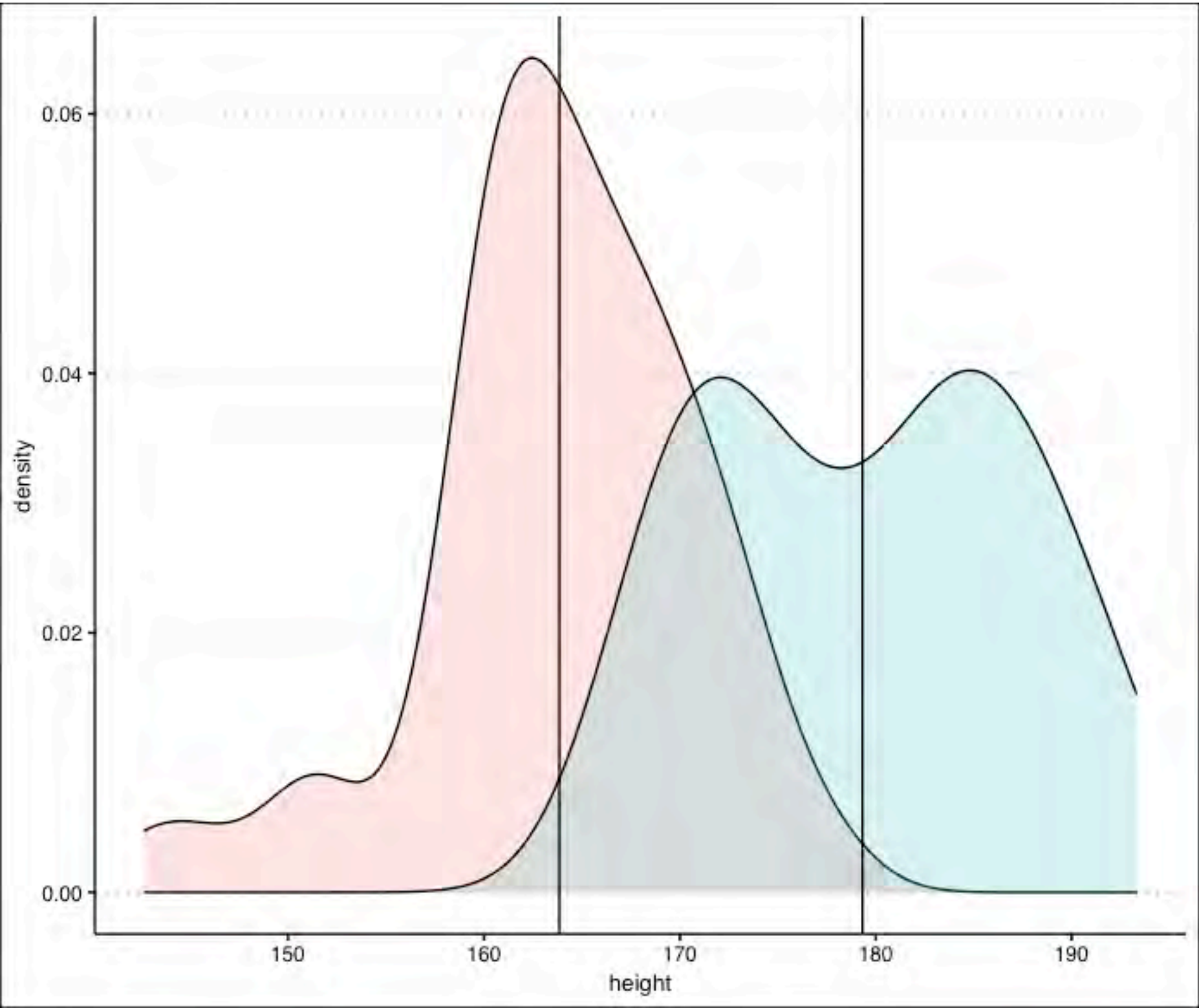simple random

convenience

interval

cluster

quota

stratified

quasi-experimental

statistical tests

Example, recall our random sample of heights in NYC

Sample mean for both groups,

```
|male  |    x_bar|
|:-----|--------:|
|FALSE | 163.8453|
|TRUE  | 179.3198|
```


How do we decide whether to reject H_0 in favor of H_A?

$$H_0 : \overline{height}_{men} = \overline{height}_{women}$$

$$H_A : \overline{height}_{men} \neq \overline{height}_{women}$$

We need some kind of test!

A general procedure for a statistical test

Assume an appropriate *probability model* to describe the behavior of the random variable under investigation.

Define a *null* hypothesis and an *alternative* hypothesis that permits meaningful conclusions.

Specify a *test statistic*.

Choose a *level of significance α* for the test.

Determine a *distribution* and *critical region* of the test statistic.

Calculate a value of the test statistic from a random sample of data.

Accept of reject H_0 by comparing the calculated value of the test statistic with the values defining the critical region.

Generic normal distribution, distance to zero, at a 95 percent *confidence interval* and 80 percent *power*
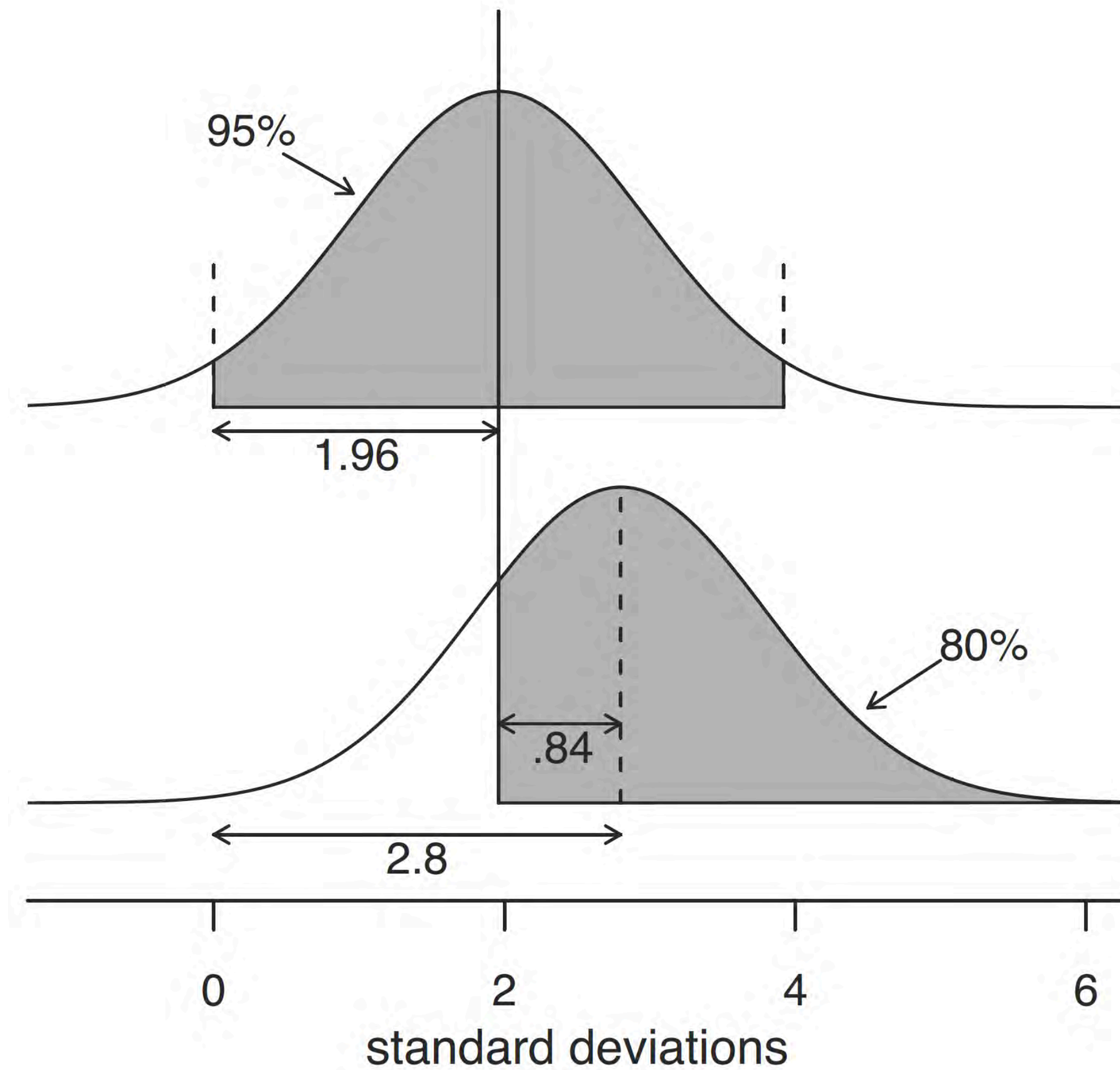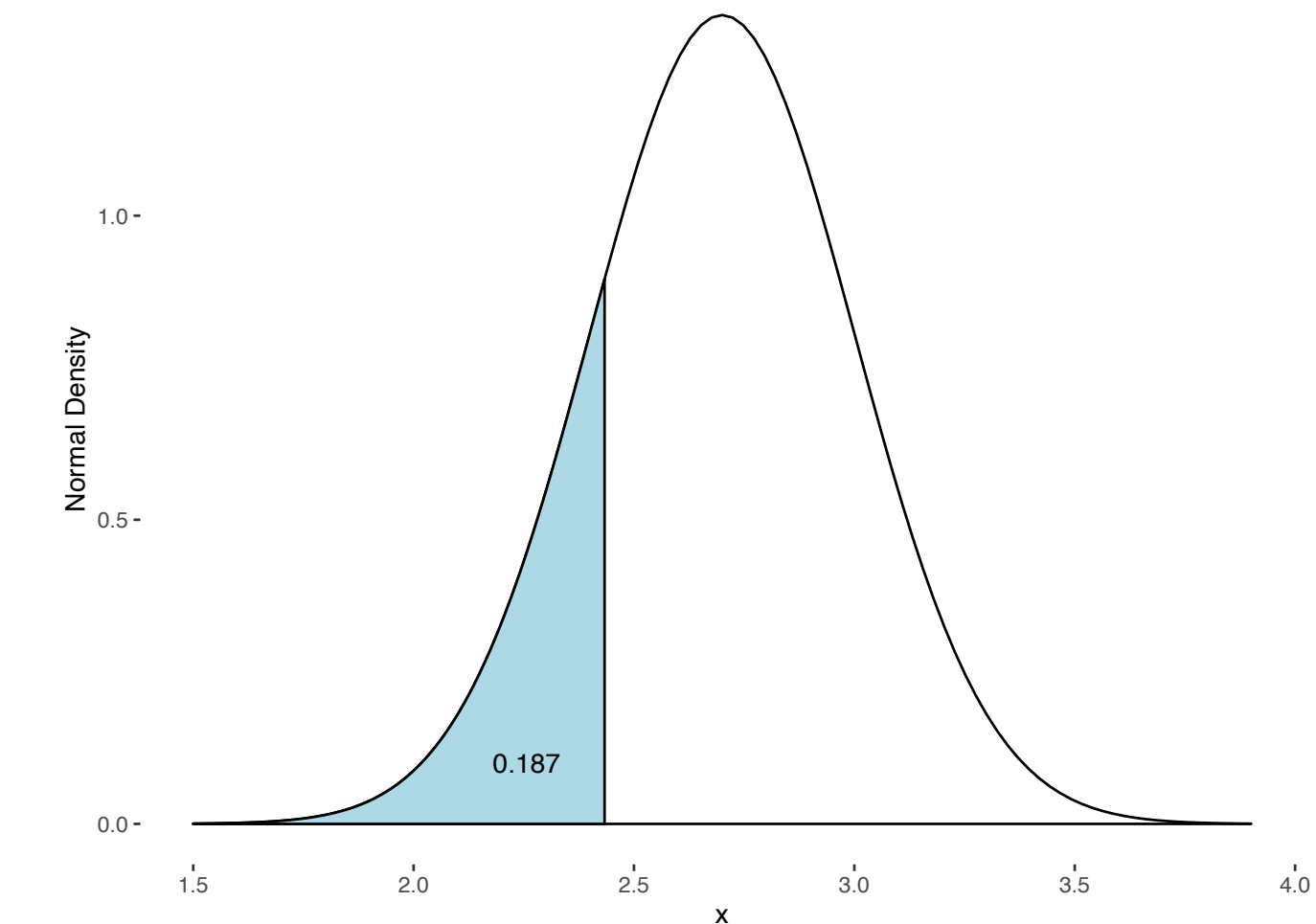
$$H_0 : \mu = 2.7, H_A : \mu < 2.7$$

```
set.seed(92)

n <- 30
mu <- lambda <- 2.7
sigma <- sqrt(lambda)
x <- rpois(n, lambda)
```

$$z = \frac{\bar{x} - \mu}{\sigma/\sqrt{n}}, \ p = F_\Phi(z)$$

```
x_bar <- mean(x)

z <- ( x_bar - mu ) / ( sigma / sqrt(n) )
p <- pnorm(z)
```

Given the null hypothesis, about
19 percent of experiments have $\bar{x} \leq 2.4\bar{3}$


```
ggplot() +
  stat_function(fun  = dnorm,
                args = list(mean = mu, sd = sigma  / sqrt(n)),
                geom = "density",
                fill = "white",
                xlim = c(mu - 4 * sigma / sqrt(n),
                         mu + 4 * sigma / sqrt(n)) ) +
  stat_function(fun  = dnorm,
                args = list(mean = mu, sd = sigma / sqrt(n)),
                geom = "density",
                fill = "lightblue",
                xlim = c(mu - 4 * sigma / sqrt(n), x_bar)) +
  annotate("segment", x = x_bar, xend = x_bar,
           y = 0, yend = dnorm(x_bar, mu, sigma  / sqrt(n))) +
  annotate("text", x = x_bar - 0.1, y = 0.1, hjust = 1,
           label = format(p, digits = 3)) +
  scale_x_continuous(breaks = seq(-1.5, 4, by = 0.5)) +
  labs(y = "Normal Density")
```
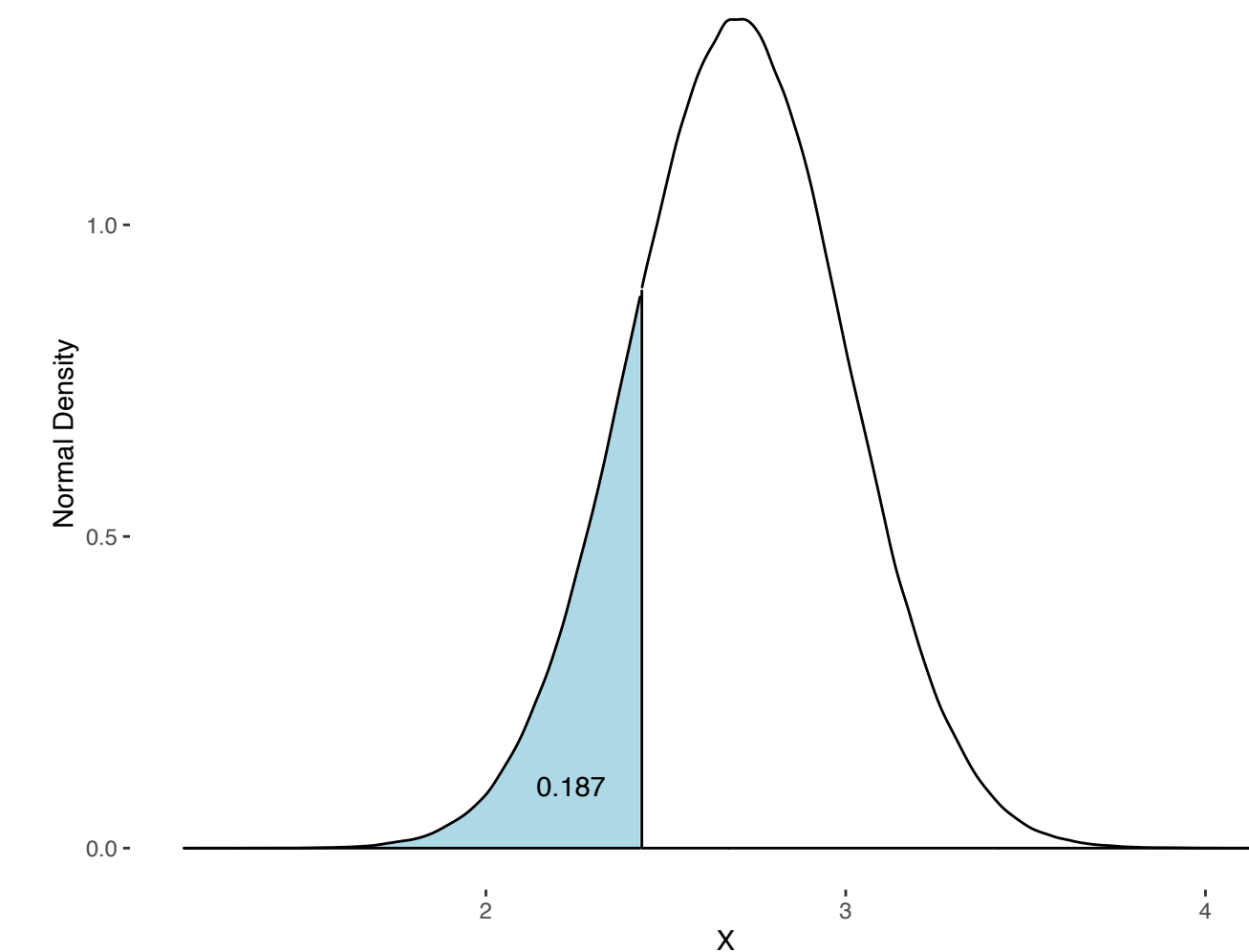
$$H_0 : \mu = 2.7, H_A : \mu < 2.7$$

```
X <- rnorm(1e6, mu, sigma / sqrt(n) )

d <- data.frame( density(X)[1:2] )
```


$$z = \frac{\bar{x} - \mu}{\sigma/\sqrt{n}}, \; p = F_\Phi(z)$$

```
ggplot(d) +
  theme(legend.position = "") +
  geom_ribbon(aes(x = x,
                  ymin = 0,
                  ymax = y,
                  fill = x < x_bar),
              color = "black") +
  scale_x_continuous(breaks = seq(-1.5, 4, by = 0.5)) +
  scale_fill_manual(values = c("white", "lightblue")) +
  annotate("segment", x = x_bar, xend = x_bar,
           y = 0, yend = dnorm(x_bar, mu, sigma  / sqrt(n))) +
  annotate("text",
           x = x_bar - 0.1, y = 0.1, hjust = 1,
           label = format( mean( X < x_bar ), digits = 3 ) ) +
  labs(x = "X", y = "Normal Density")
```

Simulation gives us the same answer:
given the null hypothesis, about 19 percent of
experiments have $\bar{x} \leq 2.4\bar{3}$

statistical tests, if **unknown** σ, can use sample standard deviation *s* and student's t distribution — *t-statistic*

$$z = \frac{\bar{x} - \mu}{\sigma/\sqrt{n}}, \ p = F_{\Phi}(z)$$

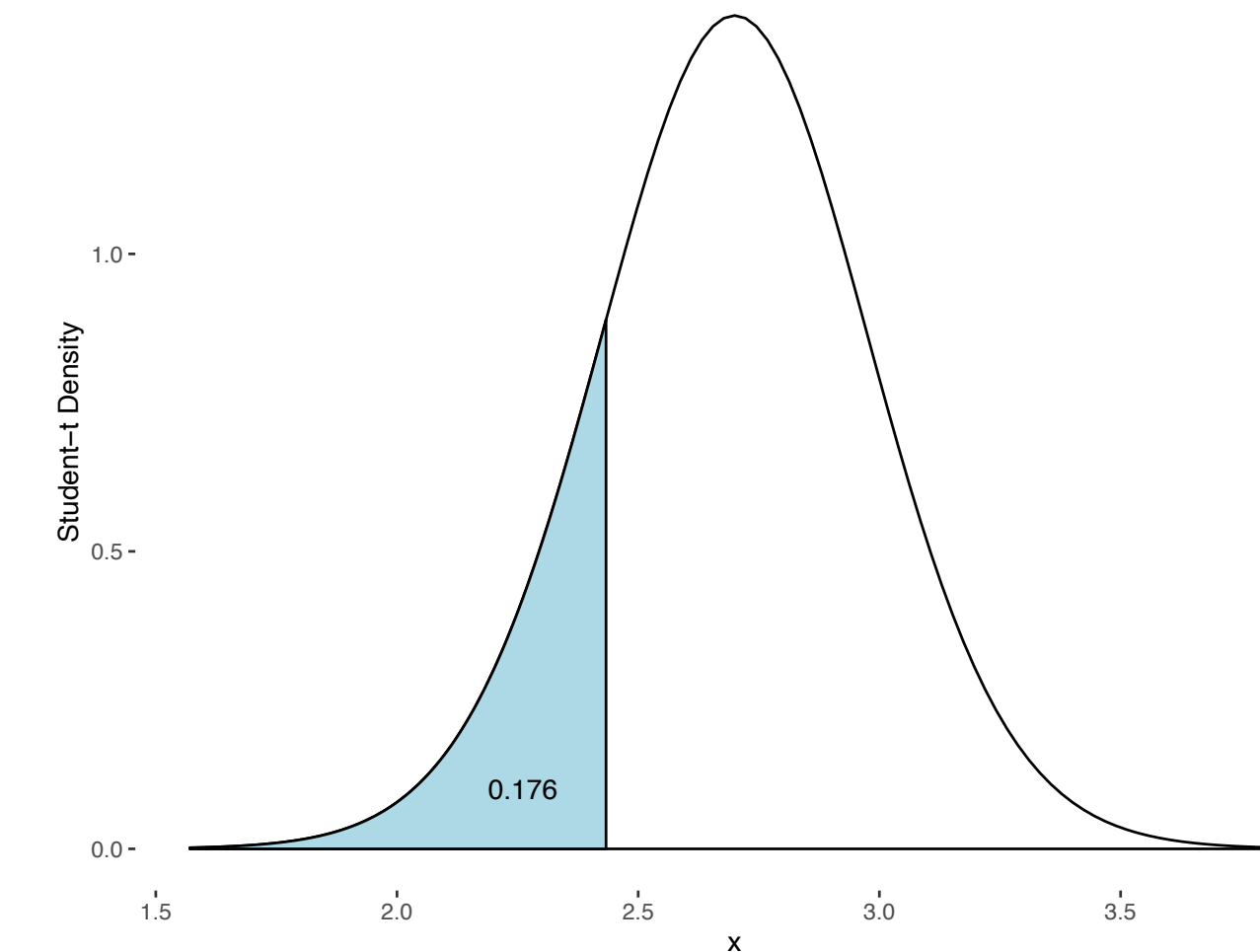$$t = \frac{\bar{x} - \mu}{s/\sqrt{n}}, \ p = F_{T}(t)$$

For ease of specifying student-t distribution parameters, here are functions for its four functions, just wrapping the base R versions:

```r
dstudent_t <-
  function(x, df, mu = 0, sigma = 1, log = FALSE) {
  if (log) {
    dt( (x - mu) / sigma, df = df, log = TRUE ) - log(sigma)
  } else {
    dt( (x - mu) / sigma, df = df ) / sigma
  }
}

pstudent_t <-
  function(q, df, mu = 0, sigma = 1, lower.tail = TRUE, log.p = FALSE) {
  pt( (q - mu) / sigma, df = df, lower.tail = lower.tail, log.p = log.p )
}

qstudent_t <-
  function(p, df, mu = 0, sigma = 1, lower.tail = TRUE, log.p = FALSE) {
  qt( p, df = df, lower.tail = lower.tail, log.p = log.p ) * sigma + mu
}

rstudent_t <-
  function(n, df, mu = 0, sigma = 1) {
  rt( n, df = df ) * sigma + mu
}
```

Of note: as $s \to \sigma \mid n \to \infty$, $t \to z$ and student's t distribution converges towards the normal distribution.

statistical tests, if **unknown** σ, can use sample standard deviation s and student's t distribution — *t-statistic*

$$H_0 : \mu = 2.7, H_A : \mu < 2.7$$

$$t = \frac{\bar{x} - \mu}{s/\sqrt{n}}, \ p = F_T(t)$$


```
s <- sd(x)

t <- ( x_bar - mu ) / ( s / sqrt(n) )
p <- pstudent_t(t, df = n - 1)
```

Given the null hypothesis, about
17 percent of experiments have $\bar{x} \leq 2.4\bar{3}$

```
ggplot() +
  stat_function(fun  = dstudent_t,
                args = list(df = n - 1, mu = mu, sigma = s  / sqrt(n)),
                geom = "density",
                fill = "white",
                xlim = c(mu - 4 * s / sqrt(n),
                         mu + 4 * s / sqrt(n)) ) +
  stat_function(fun  = dstudent_t,
                args = list(df = n - 1, mu = mu, sigma = s  / sqrt(n)),
                geom = "density",
                fill = "lightblue",
                xlim = c(mu - 4 * s / sqrt(n), x_bar)) +
  annotate("segment", x = x_bar, xend = x_bar,
           y = 0, yend = dstudent_t(x_bar, n - 1, mu, s  / sqrt(n))) +
  annotate("text", x = x_bar - 0.1, y = 0.1, hjust = 1,
           label = format(p, digits = 3)) +
  labs(y = "Student-t Density")
```

Firstly,

$$P(D\,|\,H) \neq P(H\,|\,D)$$

$$P(H\,|\,D) = \frac{P(D\,|\,H)P(H)}{P(D\,|\,H)P(H) + P(D\,|\,\neg H)P(\neg H)}$$

Secondly,

Dichotomous tests are common in the literature. We typically read comparisons that ask whether the probability of obtaining a value more "extreme" than the sample mean from the null hypothesis, and compare this to an *arbitrary* p value of 0.05 (conventionally denoted α), rejecting the null hypothesis if this z or t value of smaller, not rejecting otherwise.

But — again — this threshold test is *arbitrary*. Consider that a difference between probabilities 0.049 and 0.051, for example, is not itself typically significant.

At least report the probability of obtaining the sample value given the selected null probability distribution, *not* just the result of some dichotomous test, and consider the probability of obtaining the sample value if the hypothesis is *not* true. That's a future topic.

breakout session for group project kickoff

References

Blitzstein, Joseph K., and Jessica Hwang. *Introduction to Probability*. Second edition. Boca Raton: Taylor & Francis, 2019.

Gelman, Andrew, Jennifer Hill, and Aki Ventari. *Regression and Other Stories*. S.l.: Cambridge University Press, 2020.

Gelman, Andrew. "The Problems With P-Values Are Not Just With P-Values." The American Statistician, April 2016, 1–2.

Harris, Joseph. *Rewriting: How to Do Things with Texts*. Second edition. Logan: Utah State University Press, 2017.

McShane, Blakeley B., David Gal, Andrew Gelman, Christian Robert, and Jennifer L. Tackett. "Abandon Statistical Significance." The American Statistician 73, no. sup1 (March 29, 2019): 235–45.