Exercise 46 on page 411.

The 99% confidence interval for the standard deviation is:

Lower limit:

$$\sqrt{\frac{(n-1)s^2}{\chi^2_{\alpha/2,n-1}}} = \sqrt{\frac{(19-1)7.234}{38.58}} = 1.887$$

Upper limit:

$$\sqrt{\frac{(n-1)s^2}{\chi^2_{1-\alpha/2,n-1}}} = \sqrt{\frac{(19-1)7.234}{6.844}} = \textbf{4.481}$$

This confidence interval is **not valid** whatever the nature of the distribution. It is only valid for the normal distribution. The formula used to compute the interval relies on the fact that the distribution of a squred normal distribution is a chi-squared distribution (from page 315 of textbook).

```
alpha <- 0.01
vals <- c(
    19.75, 21.25, 21.5, 22.50, 23.25, 23.5, 24.00, 24, 24.25,
    24.5, 25.00, 26.0, 26.25, 26.25, 27.0, 27.75, 28, 28.00,
    28.25, 30
)
n <- length(vals)
s2 <- var(vals)

chi_high <- qchisq(alpha/2, df=n-1)
chi_low <- qchisq(1 - alpha/2, df=n-1)
lower <- (n-1)*s2/chi_low
upper <- (n-1)*s2/chi_high

lower_s <- sqrt(lower) # lower limit
upper_s <- sqrt(upper) # upper limit</pre>
```

Exercise 8 on page 435.

The null hypothesis H_0 is that the average warpage of the special laminate μ_s will be equal to the warpage of the regular laminage μ_r .

$$H_0: \quad \mu_s = \mu_r$$

The alternative hypothesis H_a is that the warpage of the special laminate μ_s is less than the warpage of the regular laminate μ_r .

$$H_a: \mu_s < \mu_r$$

A **type I error** in this context would be determining that the average warpage of the special laminate is less than the average warpage of the regular laminate when it really is not.

A type II error in this context would be determining that the average warpage of the special laminate is not less than the warpage of the regular laminate when it really is.

Exercise 12a, b, c on page 436.

 \mathbf{a}

The parameter of iterest μ is the real average breaking distance at 40mph using the new design.

$$H_0: \mu = 120 \text{ feet}$$

$$H_a: \mu < 120 \text{ feet}$$

b)

The appropriate regection region is R_2 . We want to reject the only if there is a reduction in braking distance so the regection region should be lower tailed.

c)

The signifigance level is:

$$\alpha = \Phi\left(\frac{\bar{x} - \mu}{\sigma/\sqrt{n}}\right) = \Phi\left(\frac{115.20 - 120}{10/\sqrt{36}}\right) = 1.9884 \times 10^{-3}$$

To acheive $\alpha = 0.001$:

$$\Phi(z) = 0.001 \implies z = -3.090$$

$$z = \frac{\bar{x} - \mu}{\sigma / \sqrt{n}} \implies \bar{x} = \frac{z\sigma}{\sqrt{n}} + \mu$$

$$\bar{x} = \frac{-3.090 \cdot 10}{\sqrt{36}} = 114.85$$

```
sigma <- 10;
n <- 36;
mu <- 120;
x <- 115.20;
z <- (115.20 - 120) / (sigma/sqrt(n))
alpha <-pnorm(z)
z2 <- qnorm(0.001)
x2 <- z2 * sigma / sqrt(n) + mu</pre>
```

Exercise 28 on page 449.

The data suggests the true average lateral recumebcy time is less than 20 mins.

Hypotheses

$$H_0: \quad \mu = \mu_0 = 20 \text{ mins}$$

$$H_a: \quad \mu < \mu_0$$

Rejection region

$$z \leqslant -z_{\alpha}$$
$$-z_{\alpha} = -z_{0.1} = -1.28$$

Test statistic

$$z = \frac{\bar{x} - \mu_0}{s/\sqrt{n}} = \frac{18.6 - 20}{8.6/\sqrt{73}} = -1.39$$

The null hypothesis should be rejected

$$-1.39 < -1.28 \implies z < -z_{\alpha}$$

```
n <- 73;
x_bar <- 18.6;
S <- 8.6;
mu_0 <- 20;
alpha <- 0.1;
z_alpha <- qnorm(0.1)
z <- (x_bar - mu_0)/(S/sqrt(n))
reject_h0 <- z < z_alpha # TRUE</pre>
```

Exercise 32a on page 449.

a)

The data does suggest the population mean differs from 100 using $\alpha = 0.05$

Hypotheses

$$H_0: \quad \mu = \mu_0$$

$$H_a: \quad \mu \neq \mu_0$$

Rejection region

$$t \geqslant t_{\alpha/2, n-1}$$
 or $t \leqslant -t_{\alpha/2, n-1}$

$$t_{\alpha/2,n-1} = t_{0.05/2,12-1} = 2.201$$

Test statistic

$$t = \frac{\bar{x} - \mu_0}{s/\sqrt{n}} = \frac{98.46 - 100}{6.142/\sqrt{12}} = -0.870$$

The null hypothesis should **not** be rejected

$$-2.201 < -0.870 < 2.201 \implies -t_{\alpha/2,n-1} < t < t_{\alpha/2,n-1}$$

```
data <- c(
    105.6, 90.9, 91.2, 96.9, 96.5, 91.3,
    101.1, 105.0, 99.6, 107.7, 103.3, 92.4
);

# calcuate test statistic value
n <- length(data);
x_bar <- mean(data);
s <- sqrt(var(data));
mu_0 <- 100;
t <- (x_bar - mu_0)/(s/sqrt(n))

# calculate rejection region
alpha <- 0.05;
t_alpha <- qt(alpha/2, n-1, lower.tail=FALSE)

# check whether should reject
reject_h0 <- t > t_alpha || t < -1*t_alpha # <- FALSE</pre>
```