

# Homework 02 - 3.2, 3.4, 3.18, 3.22, 3.38, 3.42

*Max Wagner*

*September 26, 2015*

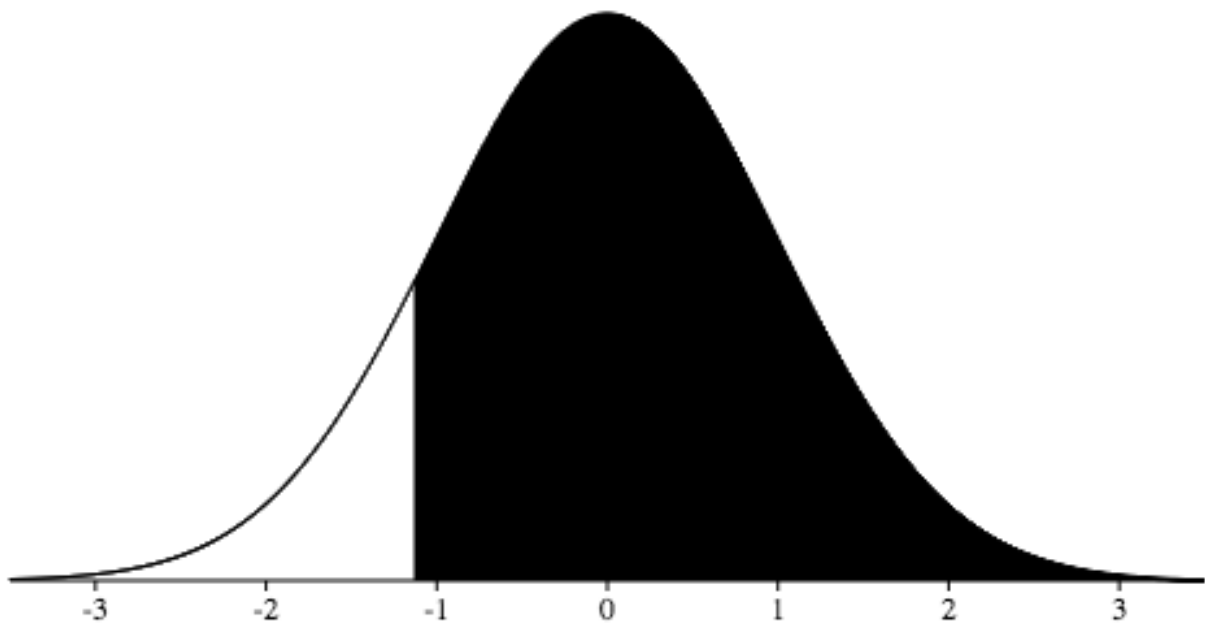
---

3.2

a.

```
1 - pnorm(-1.13)
```

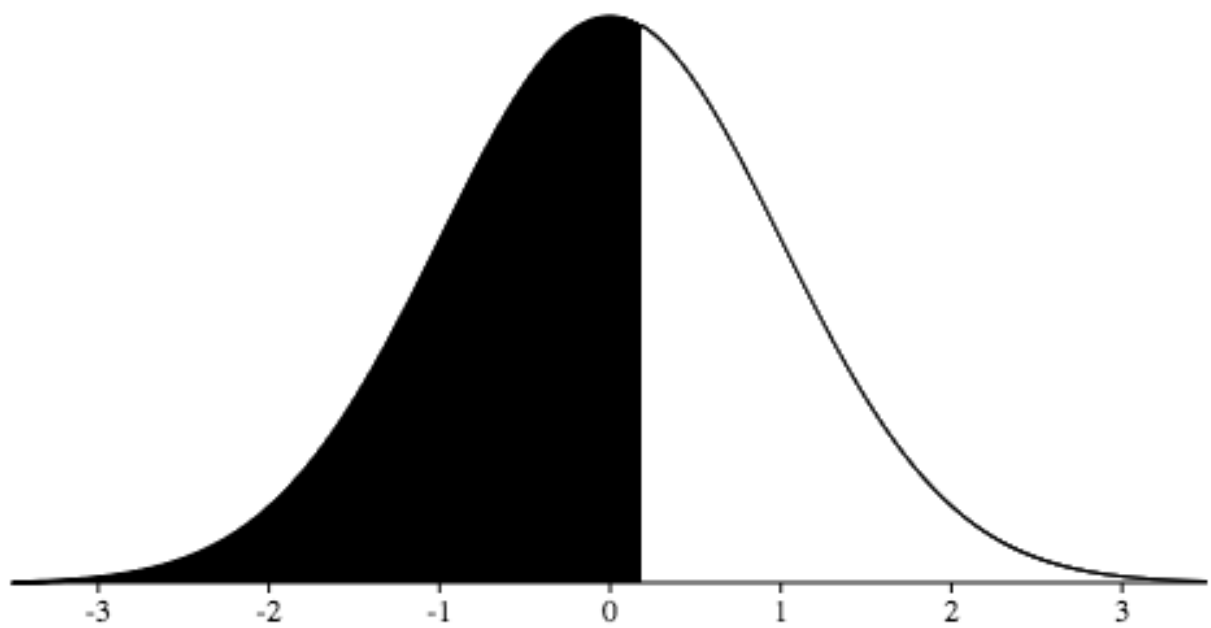
```
## [1] 0.8707619
```



b.

```
pnorm(.18)
```

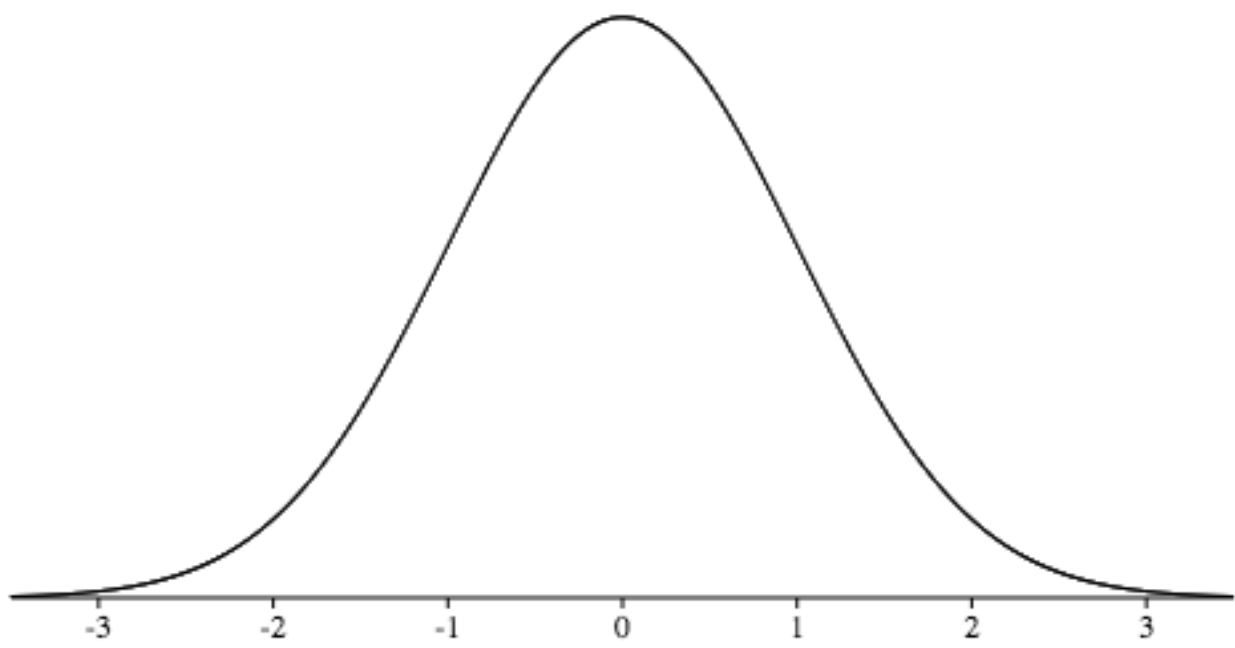
```
## [1] 0.5714237
```



c.

```
1 - pnorm(8)
```

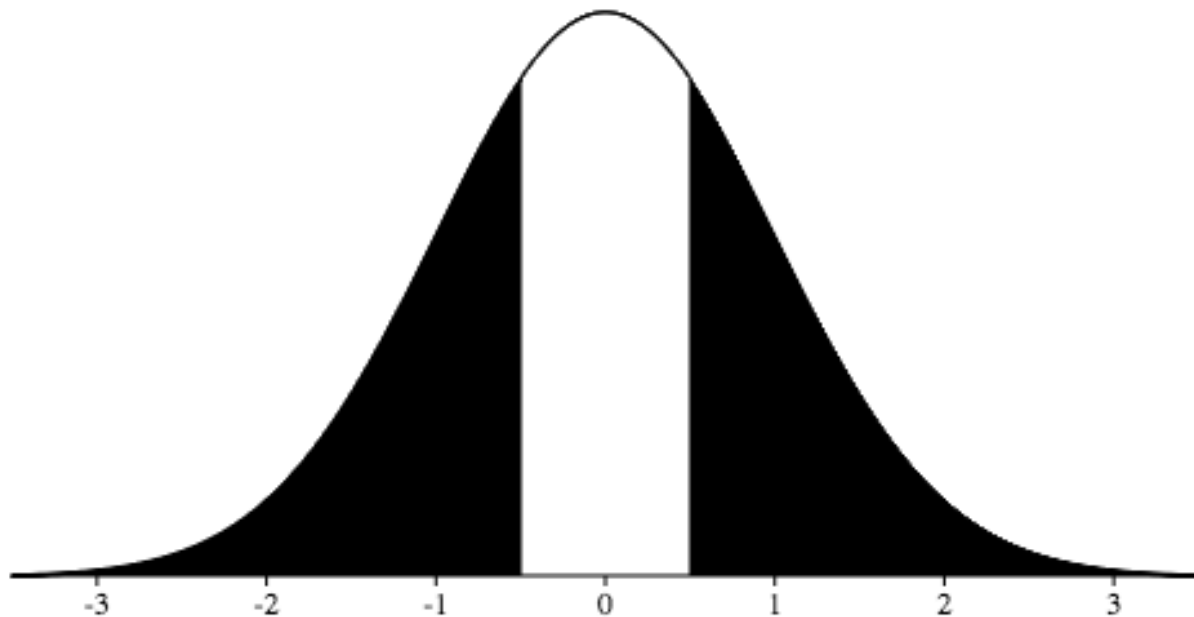
```
## [1] 6.661338e-16
```



d.

```
(1 - pnorm(.5)) + (pnorm(-.5))
```

```
## [1] 0.6170751
```



3.4

a.  $N(\mu = 4313, \sigma = 583)$ ,  $N(\mu = 5261, \sigma = 807)$

b. The z-scores tell me that Leo is 1.09 sd's from the mean, and Mary is .31 sd's from the mean.

```
leo_z <- (4948 - 4313) / 583; leo_z
```

```
## [1] 1.089194
```

```
mary_z <- (5513 - 5261) / 807; mary_z
```

```
## [1] 0.3122677
```

c. Mary ranked better than Leo. Both are positive, which means they are above the mean time, but Mary's z-score is lower than Leo's, meaning she is closer to the mean.

d. The percentage is the  $(1 - \text{pnorm})$ .

```
leo_per <- pnorm(leo_z); 1 - leo_per
```

```
## [1] 0.1380342
```

e. The percentage is the  $(1 - \text{pnorm})$ .

```
mary_per <- pnorm(mary_z); 1 - mary_per
```

```
## [1] 0.3774186
```

f. Z-scores would still be able to be calculated, but d and e would not be possible without a normal plot.

3.18

a. 68% of score fall within  $1\sigma$ , 96% within  $2\sigma$ , and 100% within  $3\sigma$ .

```
range_68 <- c((61.52 - 4.58), (61.52 + 4.58)); range_68
```

```
## [1] 56.94 66.10
```

```
per_68 <- (21 - 4) / 25; per_68
```

```
## [1] 0.68
```

```
range_95 <- c((61.52 - 4.58 * 2), (61.52 + 4.58 * 2)); range_95
```

```
## [1] 52.36 70.68
```

```
per_95 <- 24 / 25; per_95
```

```
## [1] 0.96
```

```
range_997 <- c((61.52 - 4.58 * 3), (61.52 + 4.58 * 3)); range_997
```

```
## [1] 47.78 75.26
```

b. The distribution is unimodal and symmetrical, with the curve looking normal. The normal probability plot follows a relatively straight line, with 2 outliers, neither of which are extreme.

3.22

a.

```
x <- ((1 - .02) ^ 9) * .02; x
```

```
## [1] 0.01667496
```

b.

```
x <- (1 - .02) ^ 100; x
```

```
## [1] 0.1326196
```

c.

```
expected <- (1 / .02); expected
```

```
## [1] 50
```

```
sd <- sqrt((1 - .02) / (.02 ^ 2)); sd
```

```
## [1] 49.49747
```

d.

```
expected <- (1 / .05); expected
```

```
## [1] 20
```

```
sd <- sqrt((1 - .05) / (.05 ^ 2)); sd
```

```
## [1] 19.49359
```

e. The mean and sd of a success are lower when the probability is higher.

3.38

a.

```
p1 <- factorial(3) / (factorial(2) * factorial(3 - 2))
```

```
p2 <- (.51 ^ 2) * (.49 ^ 1)
```

```
answer <- p1 * p2; answer
```

```
## [1] 0.382347
```

b. In any of these cases, the probability will remain the same, as they all include 2 successes and 1 failure. If the parents stopped having children after the 2nd boy, it would be different.

```
matrix(c("G", "B", "B", "B", "G", "B", "B", "B", "G"), byrow = T, nrow = 3)
```

```
##      [,1] [,2] [,3]
```

```
## [1,] "G"  "B"  "B"
```

```
## [2,] "B"  "G"  "B"
```

```
## [3,] "B"  "B"  "G"
```

Using the addition rule for disjoint outcomes works out the same answer.

```
.49 * .51 * .51 * 3
```

```
## [1] 0.382347
```

c. Part a. only requires one formula, and one probability, whereas part b. require entering multiple probabilities multiple times.

3.42

a.

```
p1 <- (factorial(10 - 1)) / (factorial(3 - 1) * factorial(10 - 3))
p2 <- (.15 ^ 3) * (.85 ^ 7)
answer <- p1 * p2; answer
```

```
## [1] 0.03895012
```

b.

```
p1 <- factorial(10) / (factorial(3) * factorial(10 - 3))
p2 <- (.15 ^ 3) * (.85 ^ 7)
answer <- p1 * p2; answer
```

```
## [1] 0.1298337
```

- c. The difference between a. and b., is that a. is calculated with a good serve on the 10th trial, whereas b. is calculated without a preset 10th trial.