# STAT 8010 R Lab 6: Normal Distributions

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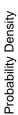
## Contents

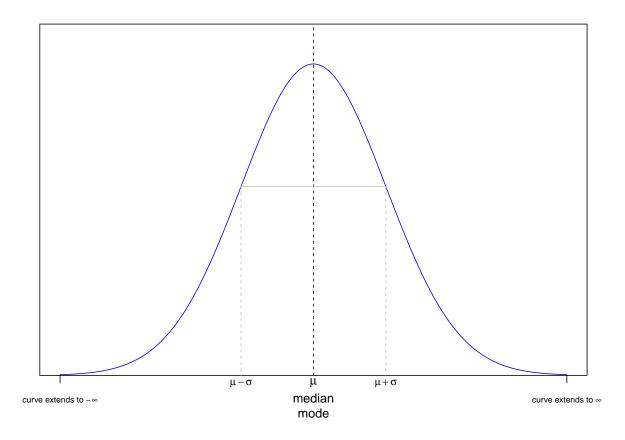
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#### Normal distribution

## Normal density curve

```
xg \leftarrow seq(-3.5, 3.5, 0.01)
yg <- dnorm(xg)
par(las = 1)
plot(xg, yg, type = "l", xlab = "",
     ylab = "Probability Density",
     col = "blue", xaxt = "n",
    yaxt = "n", yaxs = "i",
    ylim = c(0, 0.45))
abline(v = 0, lty = 2)
axis(1, at = 0, label = expression(mu),
    tick = F, line = -1)
axis(1, at = 0, label = "median", line = 0,
    tick = F)
axis(1, at = 0, label = "mode", line = 1,
axis(1, at = -3.5, labels = expression(paste("curve extends to ", -infinity)), cex.axis = 0.7)
axis(1, at = 3.5, labels = expression(paste("curve extends to ", infinity)), cex.axis = 0.7)
segments(0, dnorm(1), -1, col = "gray")
segments(0, dnorm(1), 1, col = "gray")
segments(-1, dnorm(1), -1, 0, lty = 2,
         col = "gray")
segments(1, dnorm(1), 1, 0, lty = 2,
        col = "gray")
axis(1, at = -1, label = expression(mu - sigma), tick = F, cex.axis = 0.8, line = -1)
axis(1, at = 1, label = expression(mu + sigma), tick = F, cex.axis = 0.8, line = -1)
```





## Standard Normal: Z N(0,1)

We use  $\Phi(\cdot)$  to denote the cdf of the standard normal distribution

- 1.  $\Phi(0) = .50 \Rightarrow$  Mean and Median (50<sub>th</sub> percentile) for standard normal are both 0
- 2.  $\Phi(-z) = 1 \Phi(z)$
- 3.  $\mathbb{P}(Z > z) = 1 \Phi(z) = \Phi(-z)$

## pnorm(0)

## [1] 0.5

pnorm(-1)

## [1] 0.1586553

1 - pnorm(1)

## [1] 0.1586553

pnorm(1, lower.tail = F)

## [1] 0.1586553

pnorm(-1.75)

## [1] 0.04005916

pnorm(2) - pnorm(-2)

## [1] 0.9544997

```
pnorm(0.5)
## [1] 0.6914625
STAT 8020 exam score example
mu = 78; sigma2 = 36
1 - pnorm(84, mu, sqrt(sigma2))
## [1] 0.1586553
(pnorm(84, mu, sqrt(sigma2)) - pnorm(75, mu, sqrt(sigma2))) / (1 - pnorm(75, mu, sqrt(sigma2)))
## [1] 0.7705512
Standard normal percentiles
qnorm(c(.1, .55, .9))
## [1] -1.2815516 0.1256613 1.2815516
General normal percentiles
qnorm(.8, 20, 7)
## [1] 25.89135
20 + 7 * qnorm(.8)
## [1] 25.89135
Normal approximation of Binomial Distribution
n = 400; p = 0.93
sum(dbinom(370:373, n, p))
## [1] 0.3009909
pbinom(373, n, p) - pbinom(369, n, p)
## [1] 0.3009909
mu = n * p; sigma = sqrt(n * p * (1 - p))
pnorm(373.5, mu, sigma) - pnorm(369.5, mu, sigma)
## [1] 0.3035037
plot(0:400, dbinom(0:400, n, p), type = "h", las = 1,
    xlim = c(344, 400), xlab = "x", ylab = "", col = "blue")
xg <- seq(344, 400, 0.1); yg <- dnorm(xg, mu, sigma)
lines(xg, yg, col = "blue", lty = 2)
abline(v = c(369.5, 373.5), lty = 2, col = "red")
lines(370:373, dbinom(370:373, n, p), type = "h", col = "red")
```

