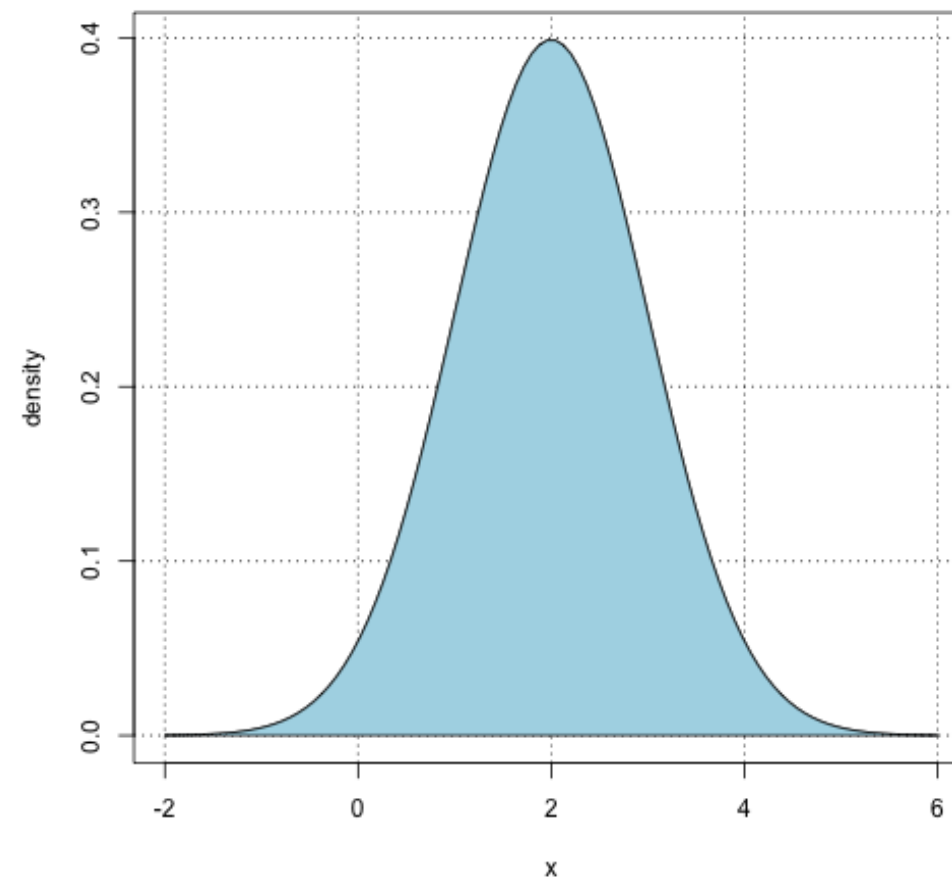




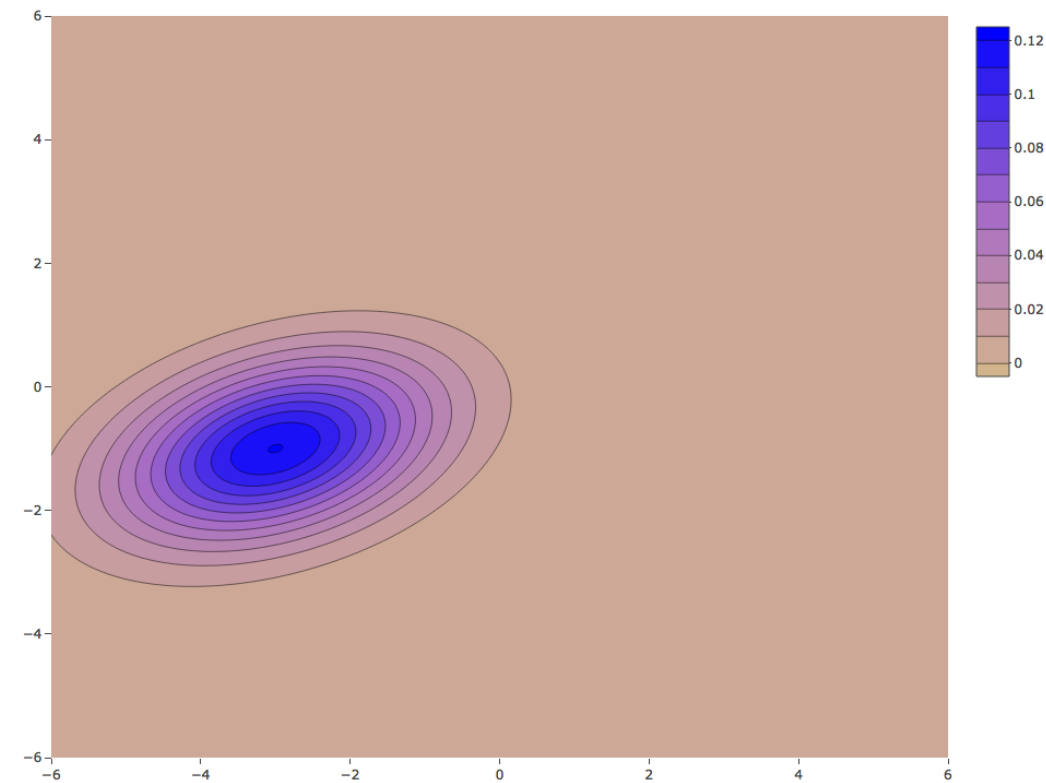
Univariate normal distribution



Univariate normal with mean 2 and variance 1

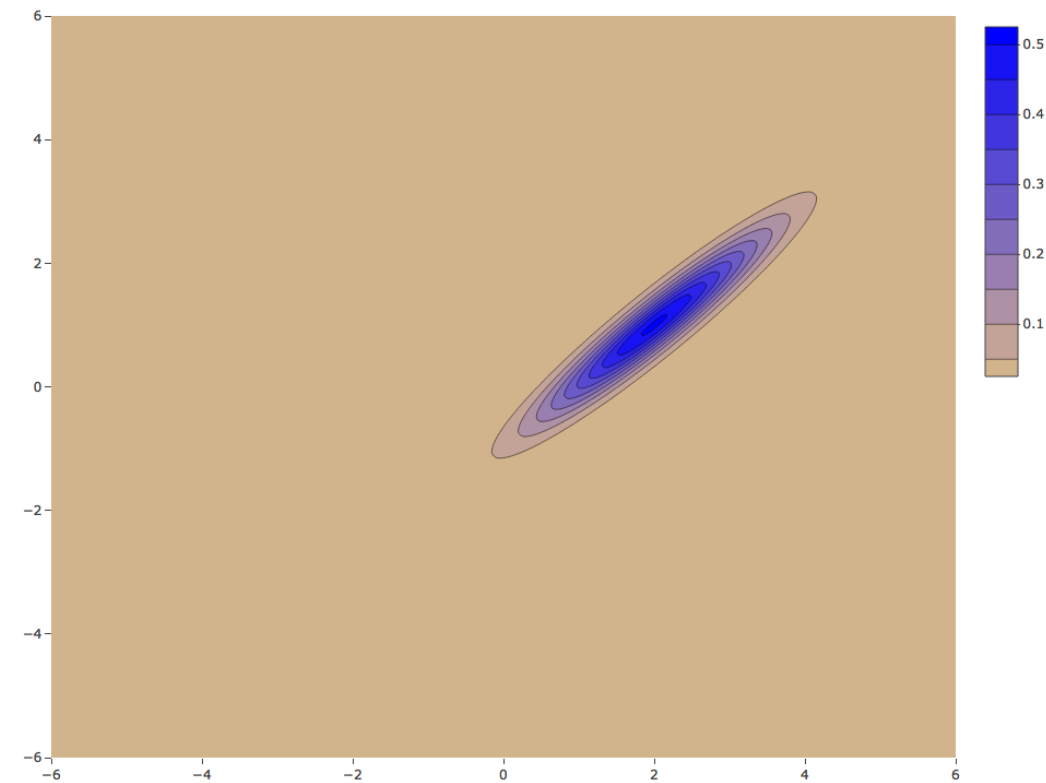
Bivariate normal density with a different mean

$$\mu = \begin{pmatrix} -1 \\ -3 \end{pmatrix}, \quad \Sigma = \begin{pmatrix} 1 & 0.5 \\ 0.5 & 2 \end{pmatrix}$$



Bivariate normal density with strong correlation

$$\mu = \begin{pmatrix} 1 \\ 2 \end{pmatrix}, \quad \Sigma = \begin{pmatrix} 1 & 0.95 \\ 0.95 & 1 \end{pmatrix}$$



Functions for statistical distributions in R

Normal

`rnorm`

`dnorm`

`pnorm`

`qnorm`

Multivariate Normal

`rmvnorm`

`dmvnorm`

`pmvnorm`

`qmvnorm`

t

`rt`

`dt`

`pt`

`qt`

Multivariate t

`rmvt`

`dmvt`

`pmvt`

`qmvt`

Functions for statistical distributions in R

Normal

`rnorm`
`dnorm`
`pnorm`
`qnorm`

Multivariate Normal

`rmvnorm`
`dmvnorm`
`pmvnorm`
`qmvnorm`

t

`rt`
`dt`
`pt`
`qt`

Multivariate t

`rmvt`
`dmvt`
`pmvt`
`qmvt`

- The first letter denotes

- `p` for "probability"
- `q` for "quantile"
- `d` for "density"
- `r` for "random"

- Followed by the distribution name

- `norm`
- `mvnorm`
- `t`
- `mvt`



The rmvnorm function

```
library(mvtnorm)
rmvnorm(n, mean, sigma)
```

Need to specify:

- `n` the number of samples
- `mean` the mean of the distribution
- `sigma` the variance-covariance matrix



Using rmvnorm to generate random samples

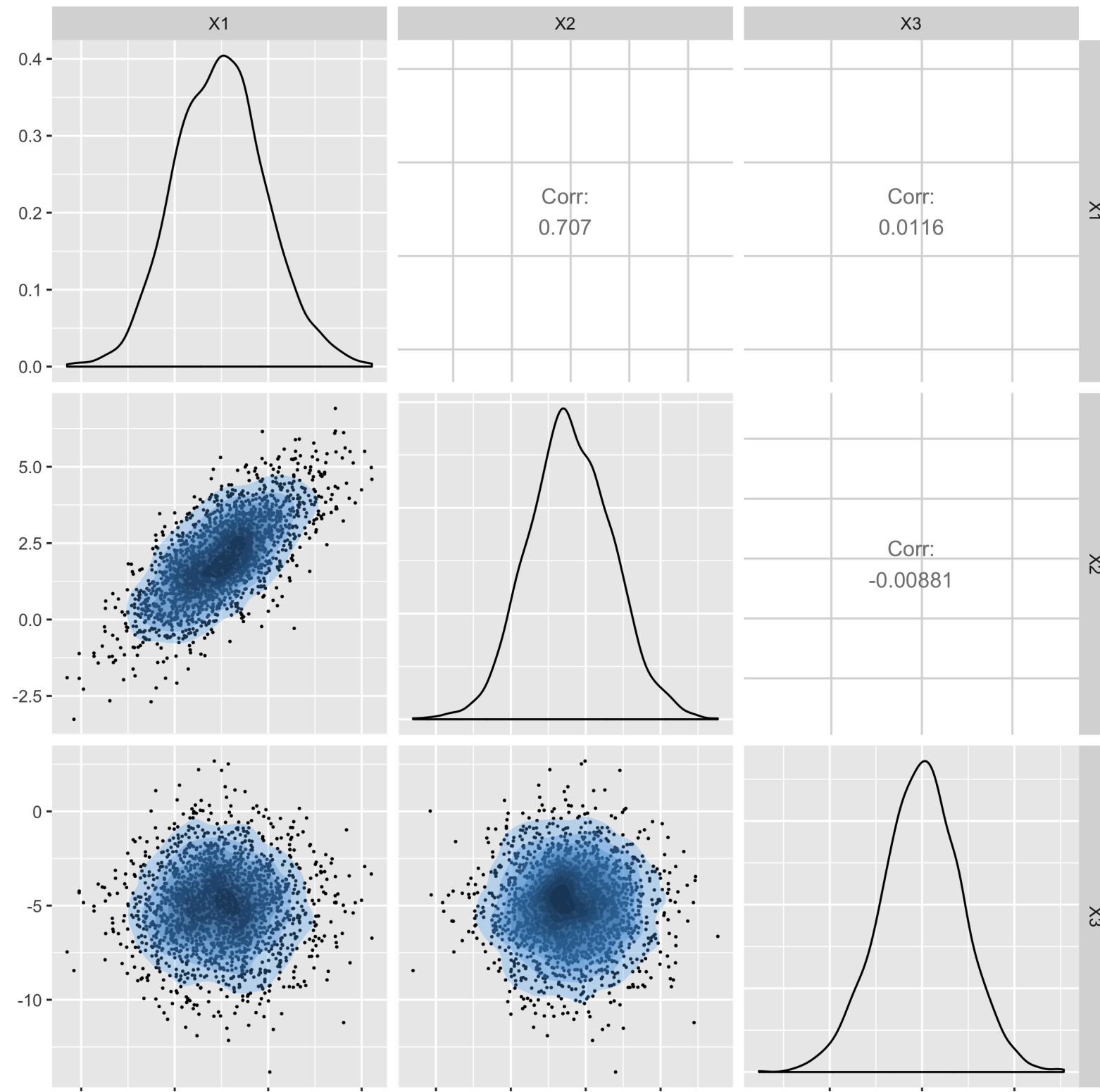
Generate 1000 samples from a 3 dimensional normal with

$$\mu = \begin{pmatrix} 1 \\ 2 \\ -5 \end{pmatrix} \quad \Sigma = \begin{pmatrix} 1 & 1 & 0 \\ 1 & 2 & 0 \\ 0 & 0 & 5 \end{pmatrix}$$

```
mu1 <- c(1, 2, -5)
sigma1 <- matrix(c(1,1,0,
                  1,2,0,
                  0,0,5), 3, 3)

set.seed(34)
rmvnorm(n = 1000, mean = mu1, sigma = sigma1)
```

Plot of generated samples





MULTIVARIATE PROBABILITY DISTRIBUTIONS IN R

**Let's practice simulating
from a multivariate normal
distribution!**



MULTIVARIATE PROBABILITY DISTRIBUTIONS IN R

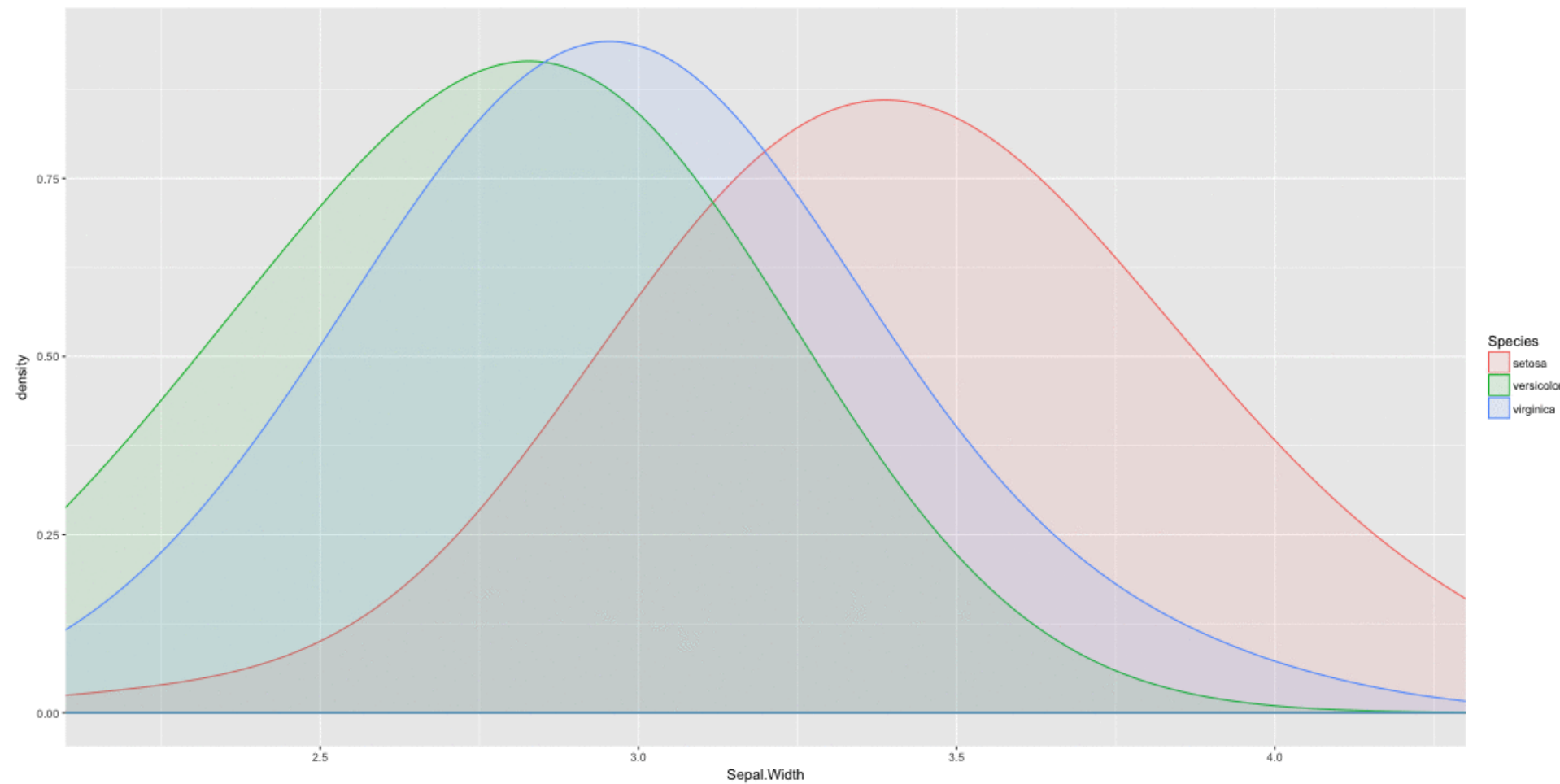
Density of a multivariate normal distribution

Surajit Ray

Senior Lecturer, University of Glasgow

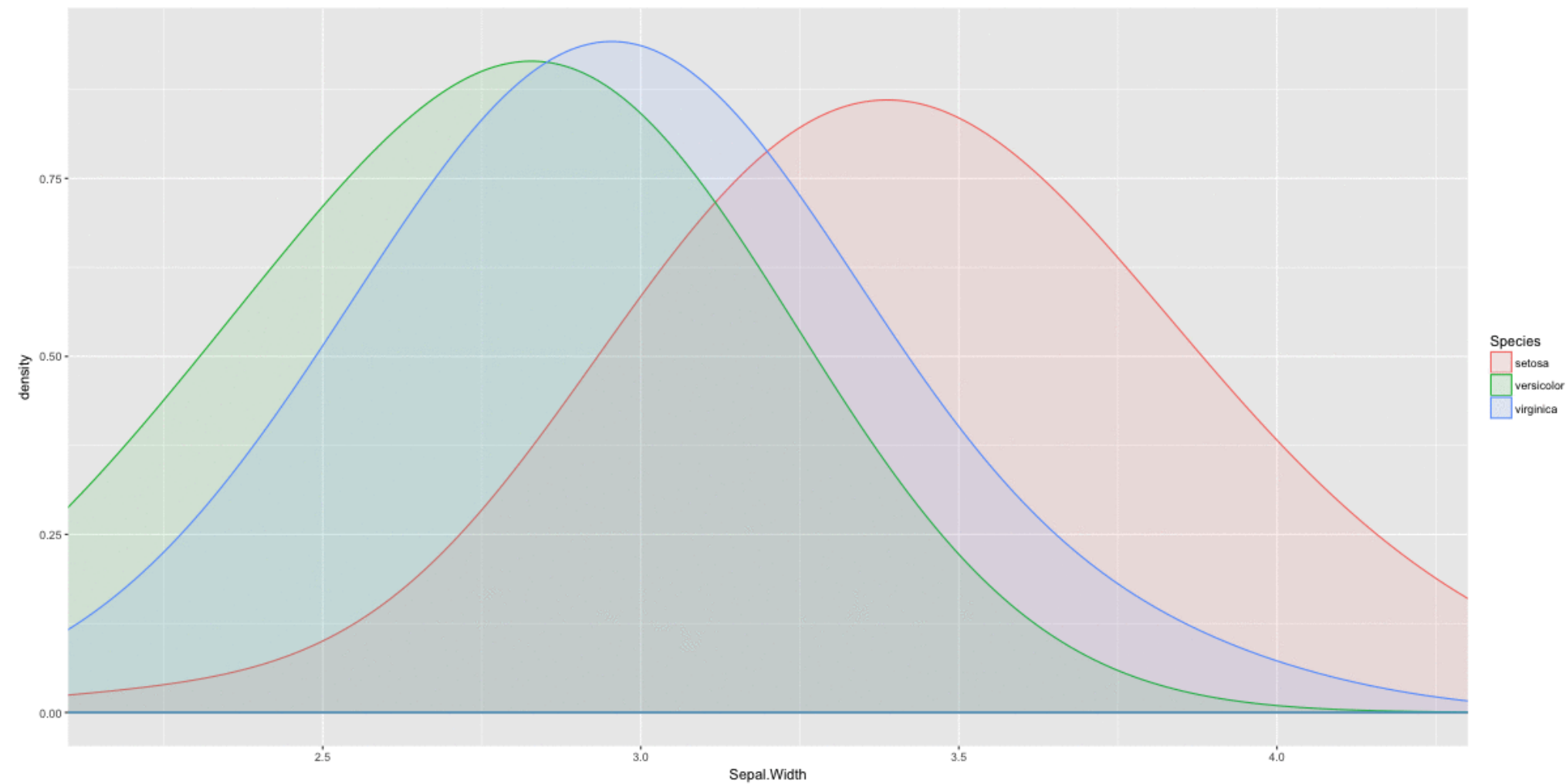


Why calculate the density of a distribution?



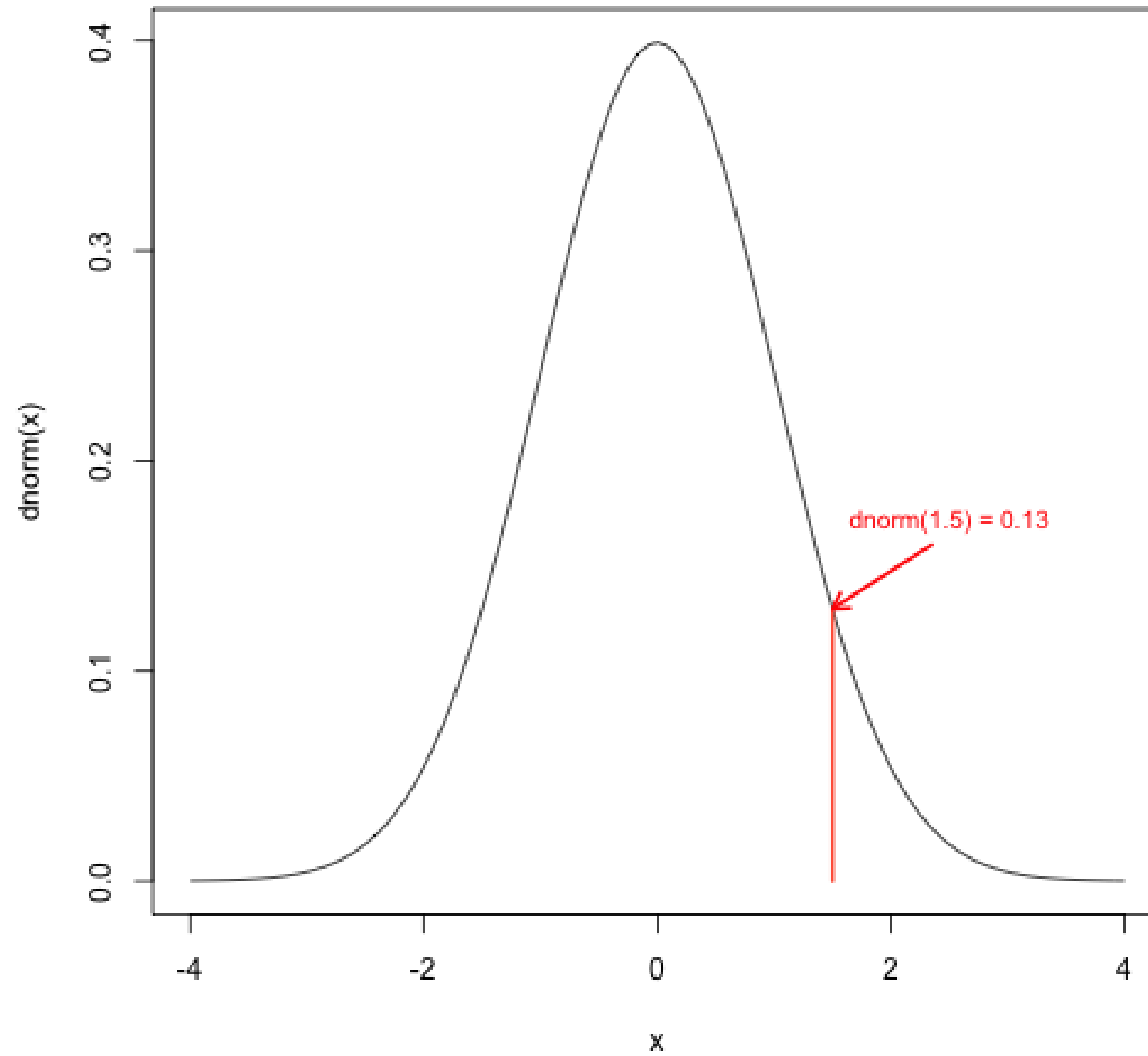


Why calculate the density of a distribution?





Univariate normal functions `dnorm()`



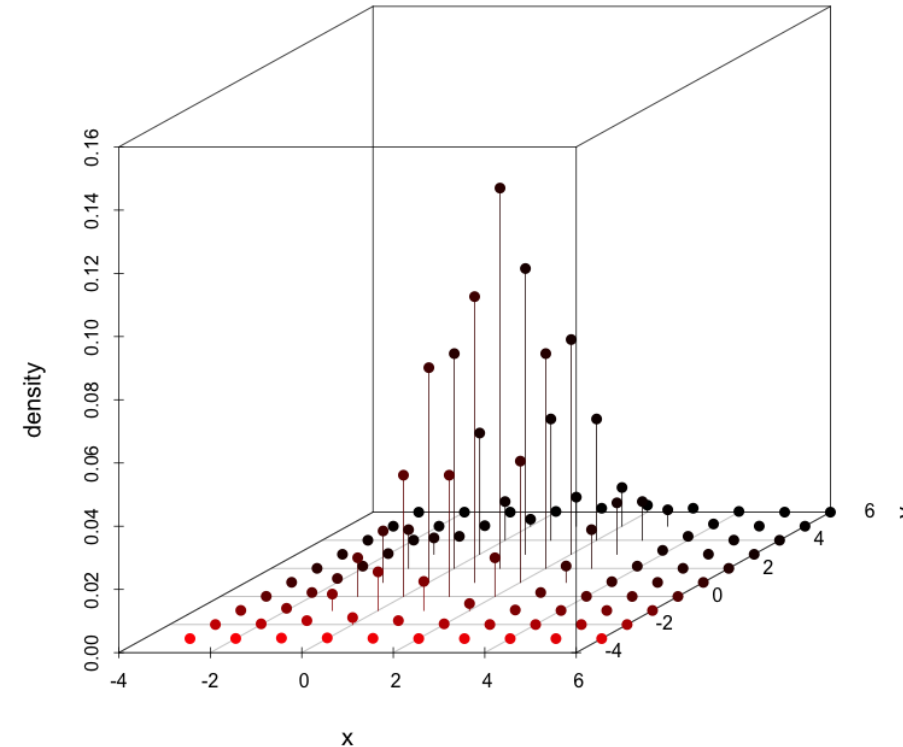
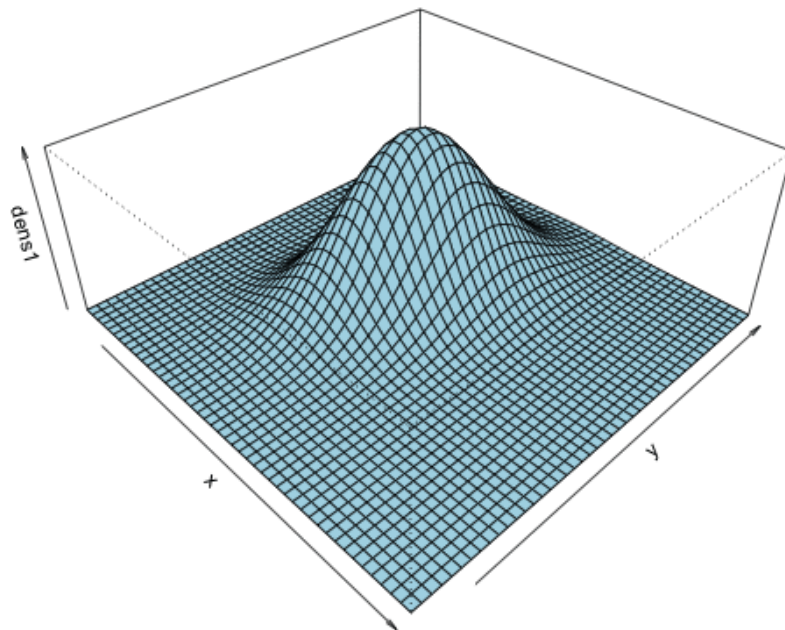


Probability density of a bivariate normal

Standard bivariate normal with

$$\mu = \begin{pmatrix} 0 \\ 0 \end{pmatrix}, \Sigma = \begin{pmatrix} 1 & 0 \\ 0 & 1 \end{pmatrix}$$

Density heights calculated at several locations (xy coordinates)





Density using dmvnorm

```
library(mvtnorm)
dmvnorm(x, mean, sigma)
```

- x can be a row vector or a matrix

```
mu1 <- c(1, 2)
sigma1 <- matrix(c(1, .5, .5, 2), 2)
dmvnorm(x = c(0, 0), mean = mu1, sigma = sigma1)
```

```
0.0384
```



Density at multiple points using dmvnorm

```
x <- rbind(c(0, 0), c(1, 1), c(0, 1)); x
```

```
[1,] 0 0  
[2,] 1 1  
[3,] 0 1
```

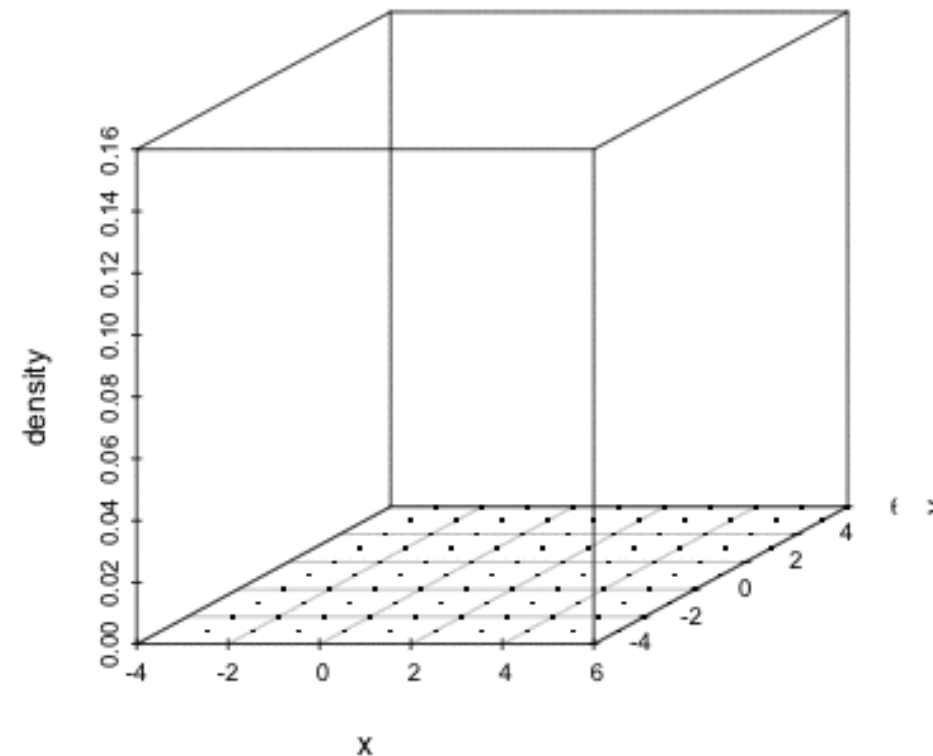
```
dmvnorm(x = x, mean = mu, sigma = sigma)
```

```
[1] 0.0384 0.0904 0.0679
```

Plotting bivariate densities with perspective plot

Steps:

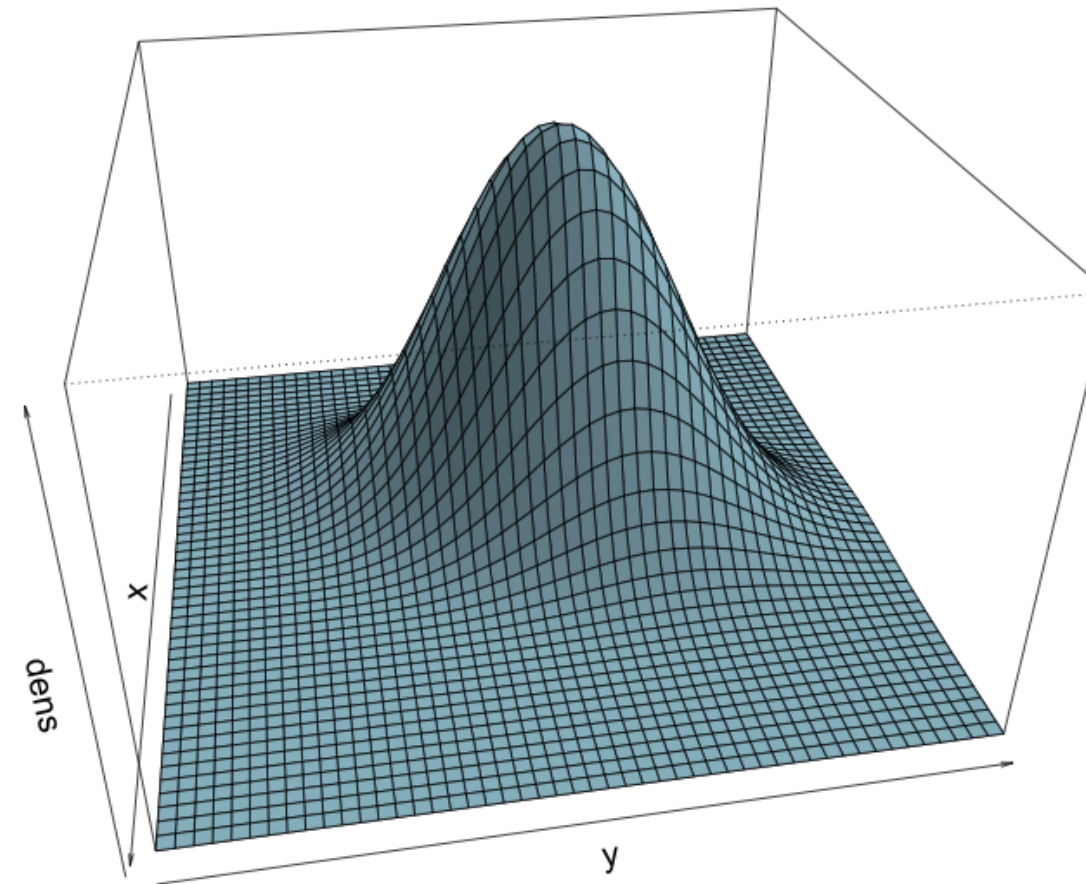
- Create grid of x and y coordinates
- Calculate density on grid



Plotting bivariate densities with perspective plot

Steps:

- Create grid of x and y coordinates
- Calculate density on grid
- Convert densities into a matrix
- Create perspective plot using `persp()` function



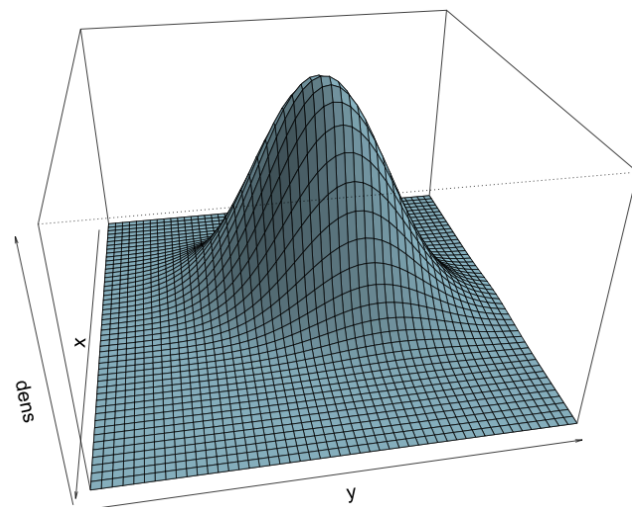
Code for plotting bivariate densities

```
# Create grid
d <- expand.grid(seq(-3, 6, length.out = 50 ), seq(-3, 6, length.out = 50))

# Calculate density on grid
dens1 <- dmvnorm(as.matrix(d), mean=c(1,2), sigma=matrix(c(1, .5, .5, 2), 2))

# Convert to matrix
dens1 <- matrix(dens1, nrow = 50 )

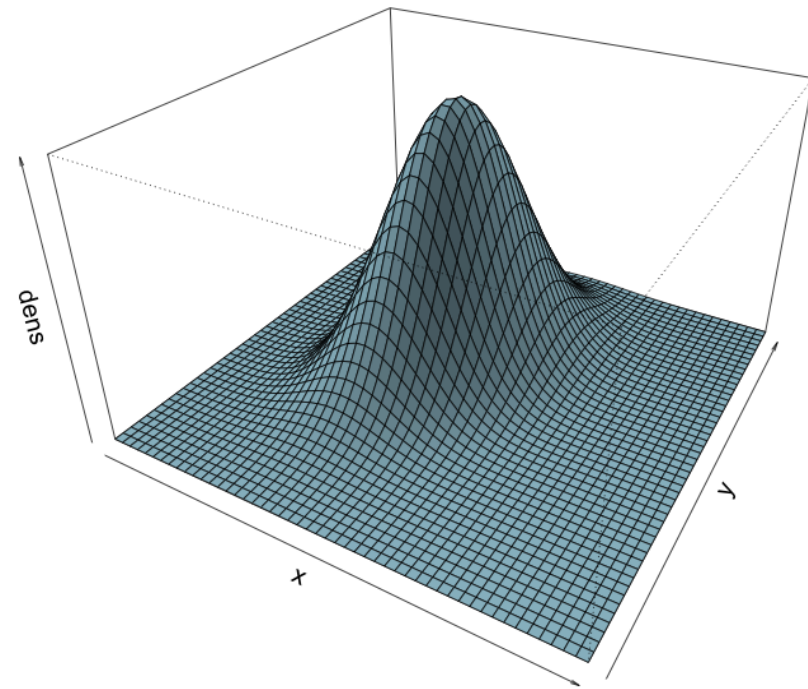
# Use perspective plot
persp(dens1, theta = 80, phi = 30, expand = 0.6, shade = 0.2,
      col = "lightblue", xlab = "x", ylab = "y", zlab = "dens")
```



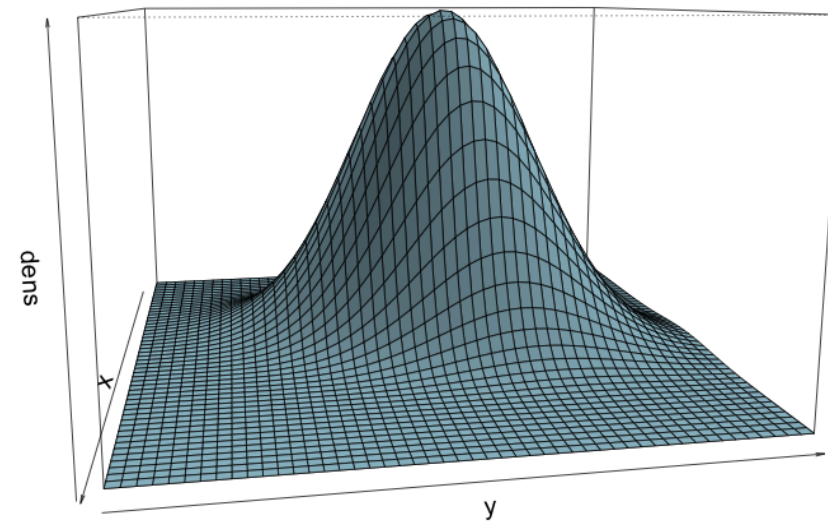


Changing viewing angle in perspective plot

`persp()` **with** `theta = 30`, `phi = 30`



`persp()` **with** `theta = 80`, `phi = 10`





MULTIVARIATE PROBABILITY DISTRIBUTIONS IN R

Let's practice!



MULTIVARIATE PROBABILITY DISTRIBUTIONS IN R

Cumulative Distribution and Inverse CDF

Surajit Ray

Senior Lecturer, University of Glasgow

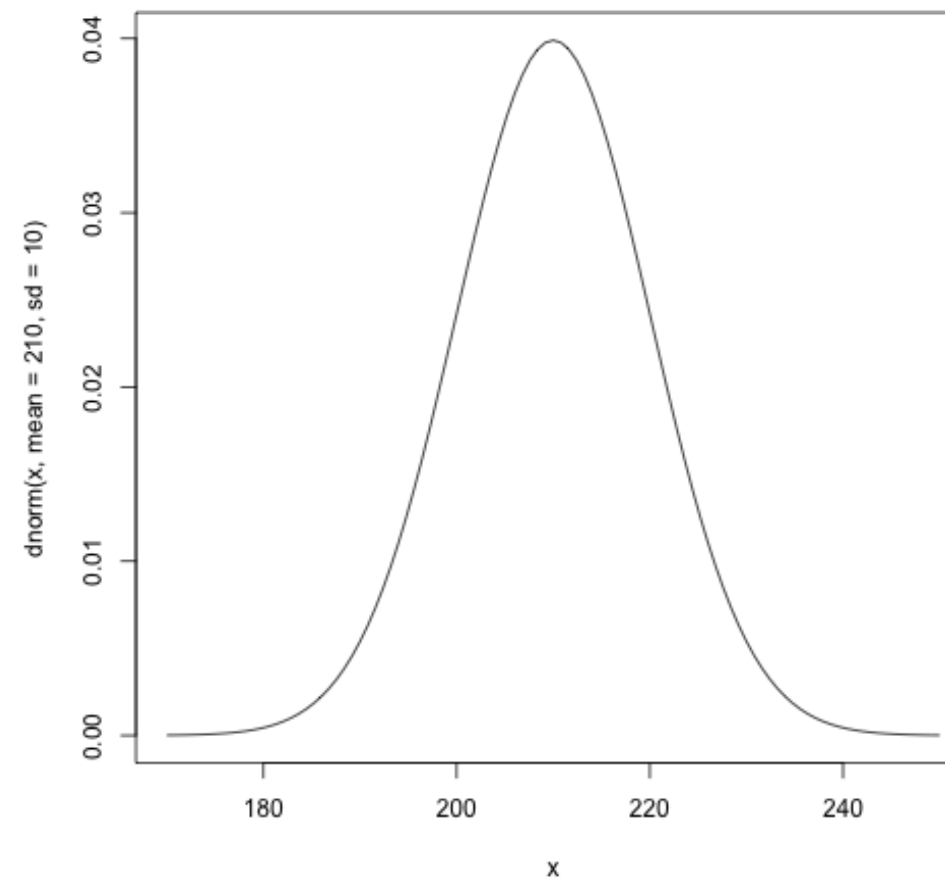


When do we need to calculate CDF and inverse CDF?



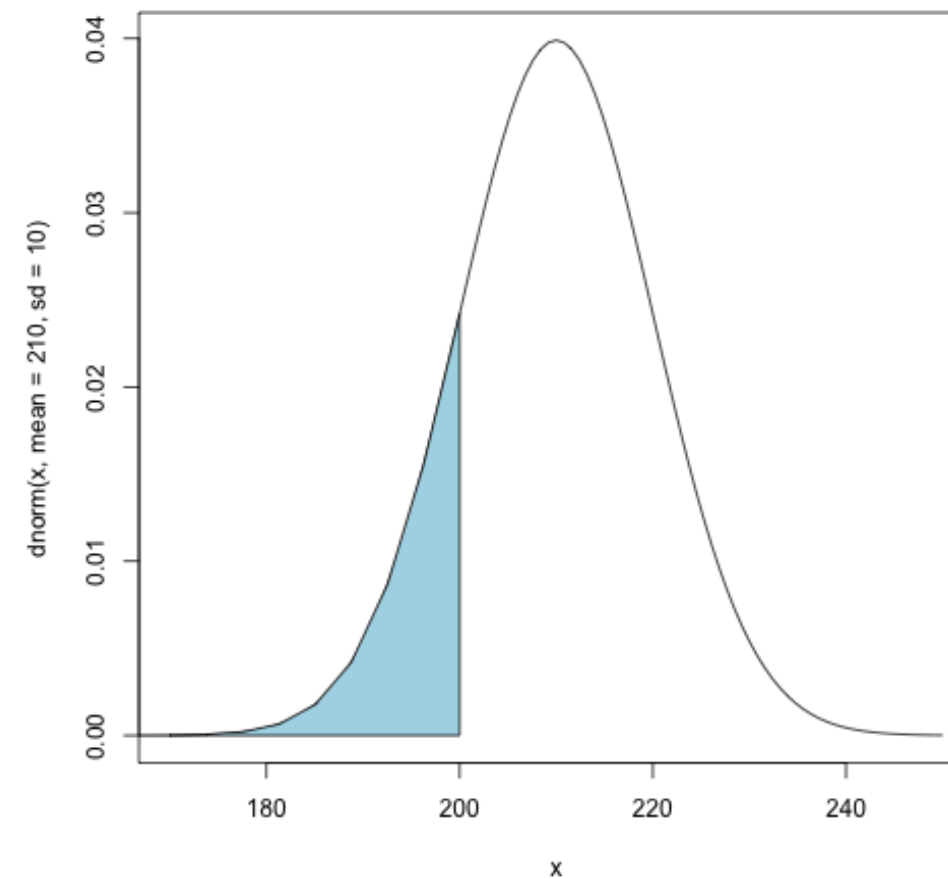


When do we need to calculate CDF and inverse CDF?



Normal density with $\mu = 210$ and $\sigma = 10$

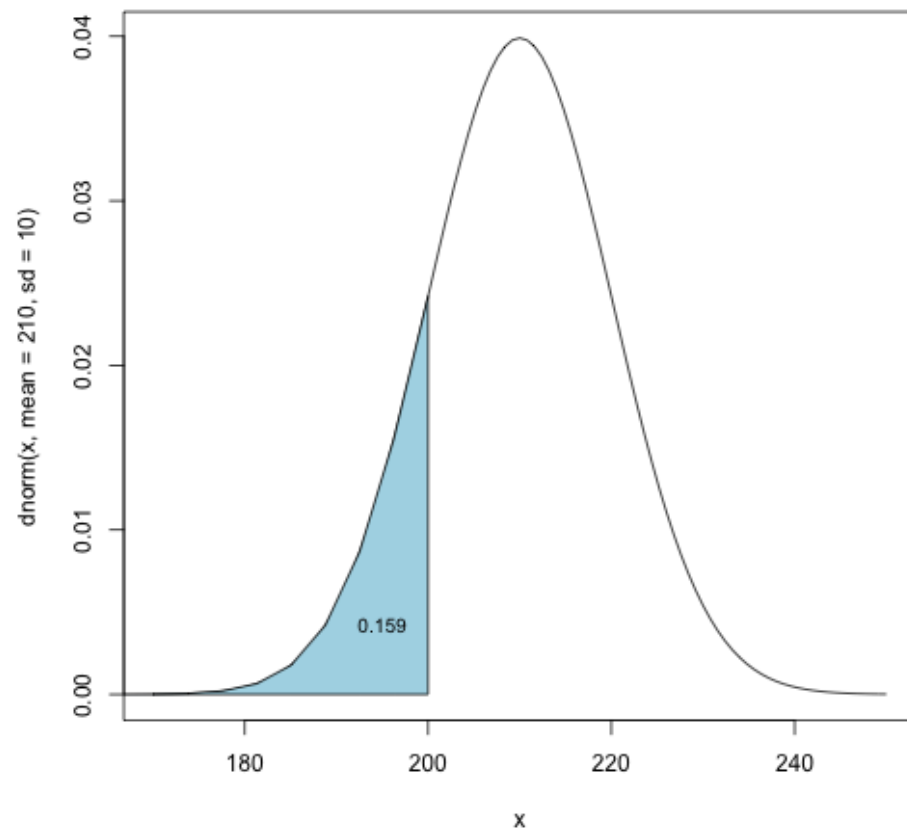
When do we need to calculate CDF and inverse CDF?



Area under the curve for $x < 200$

When do we need to calculate CDF and inverse CDF?

```
pnorm(200, mean = 210, sd = 10)  
[1] 0.159
```

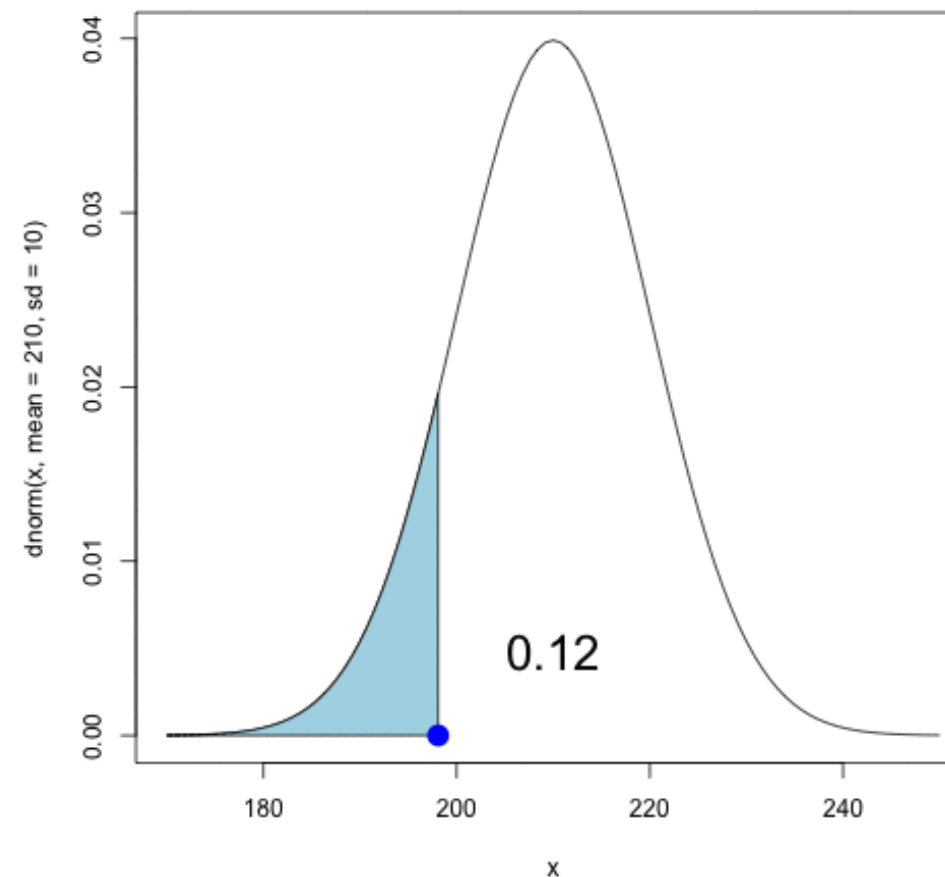


When do we need to calculate CDF and inverse CDF?

What is the x_0 such that the cumulative probability at x_0 is 0.95?

```
qnorm( p = 0.95, mean = 210, sd = 10)  
[1] 226.45
```

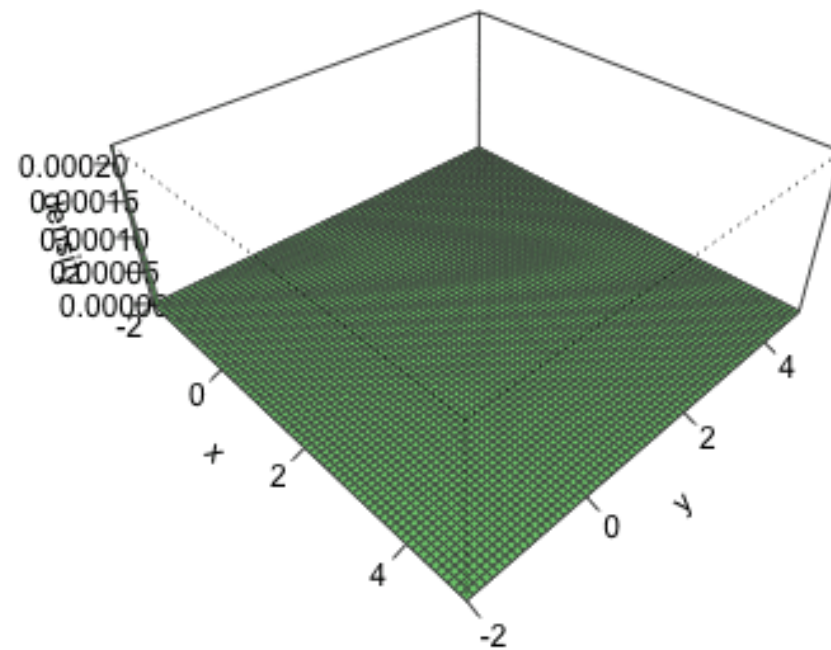
⇒ 95% of the coffee jars will have less than **226.45** grams of coffee





Cumulative distribution for a bivariate normal

Bivariate CDF at $x = 2$ and $y = 4$ for a normal with $\mu = \begin{pmatrix} 1 \\ 2 \end{pmatrix}$, $\Sigma = \begin{pmatrix} 1 & .5 \\ .5 & 2 \end{pmatrix}$



Cumulative distribution using pmvnorm

Bivariate CDF at $x = 2$ and $y = 4$ for a normal with $\mu = \begin{pmatrix} 1 \\ 2 \end{pmatrix}$, $\Sigma = \begin{pmatrix} 1 & 0.5 \\ 0.5 & 2 \end{pmatrix}$

```
mu1 <- c(1, 2)
sigma1 <- matrix(c(1, 0.5, 0.5, 2), 2)

pmvnorm(upper = c(2, 4), mean = mu1, sigma = sigma1)
```

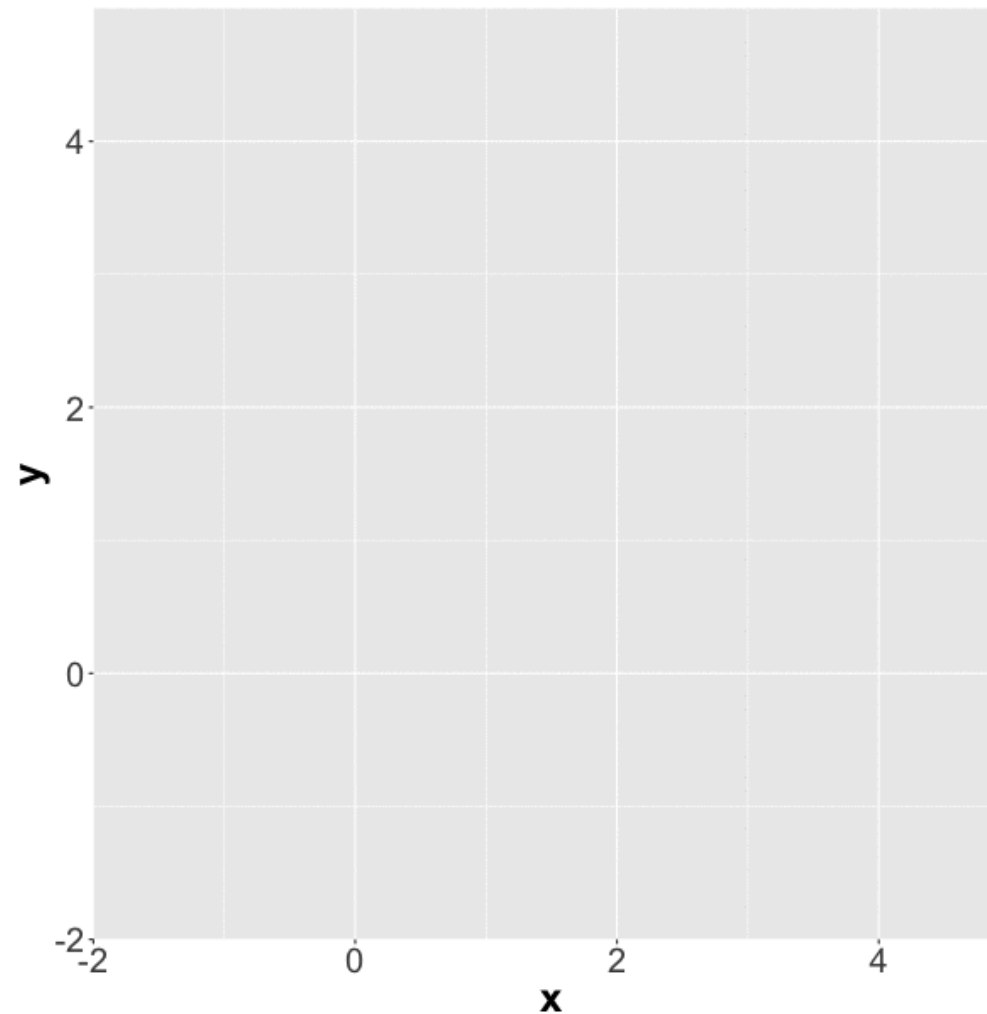
```
[1] 0.79
attr(,"error")
[1] 1e-15
attr(,"msg")
[1] "Normal Completion"
```



Probability between two values using pmvnorm

Probability of $1 < x < 2$ and $2 < y < 4$

```
pmvnorm(lower = c(1, 2),  
        upper = c(2, 4),  
        mean = mu1,  
        sigma = sigma1)
```



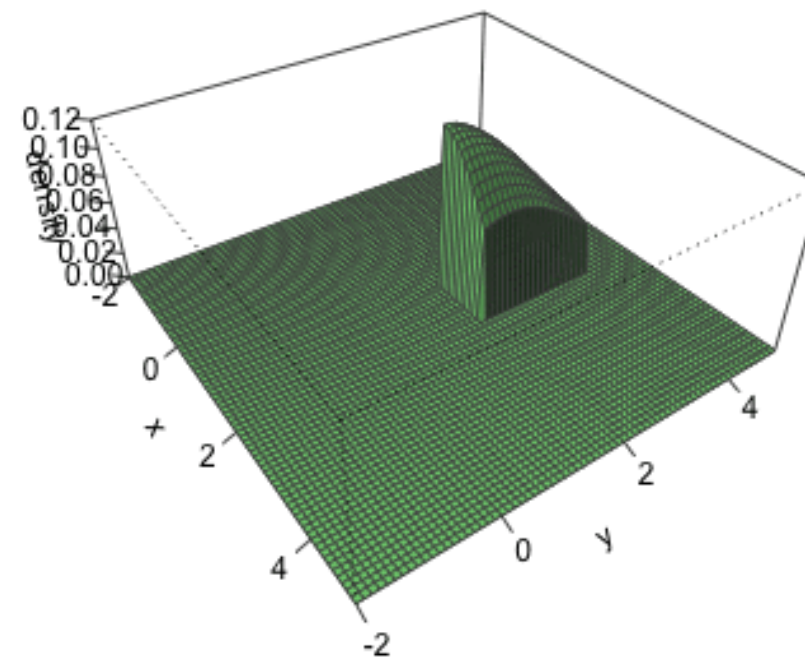


Probability between two values using pmvnorm

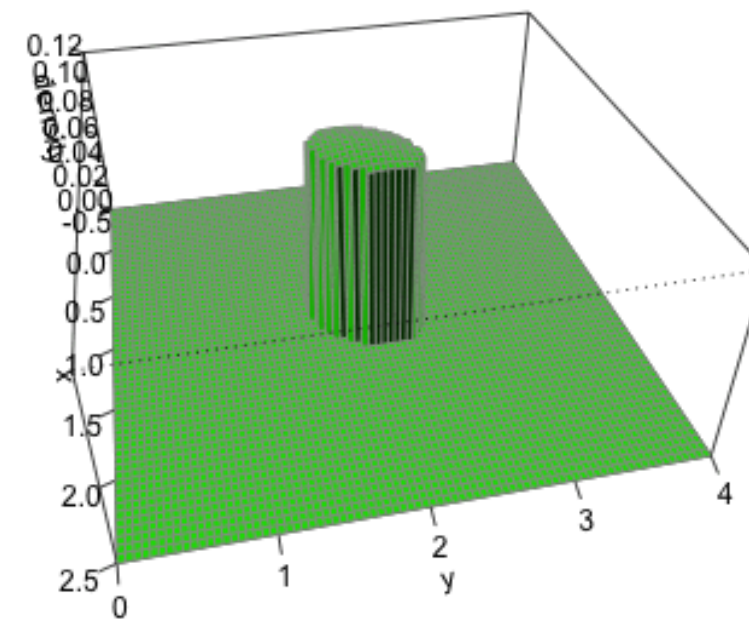
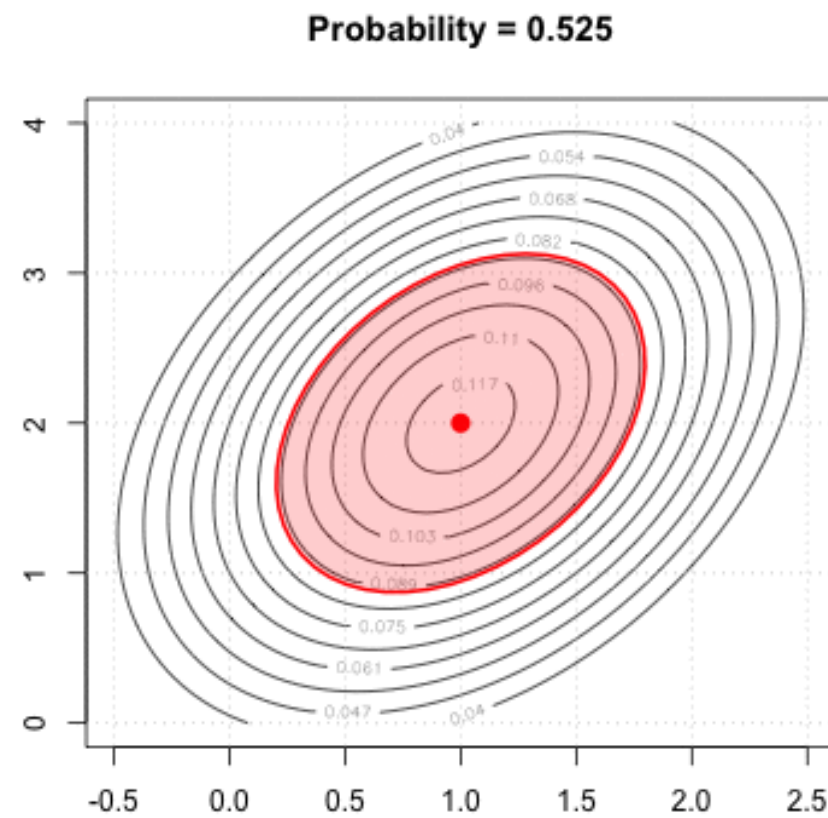
Probability of $1 < x < 2$ and $2 < y < 4$

```
pmvnorm(lower = c(1, 2),  
        upper = c(2, 4),  
        mean = mu1,  
        sigma = sigma1)
```

```
[1] 0.163
```



Inverse CDF for bivariate normal



Dark red ellipse is the 0.95 quantile

Implementing qmvnorm to calculate quantiles

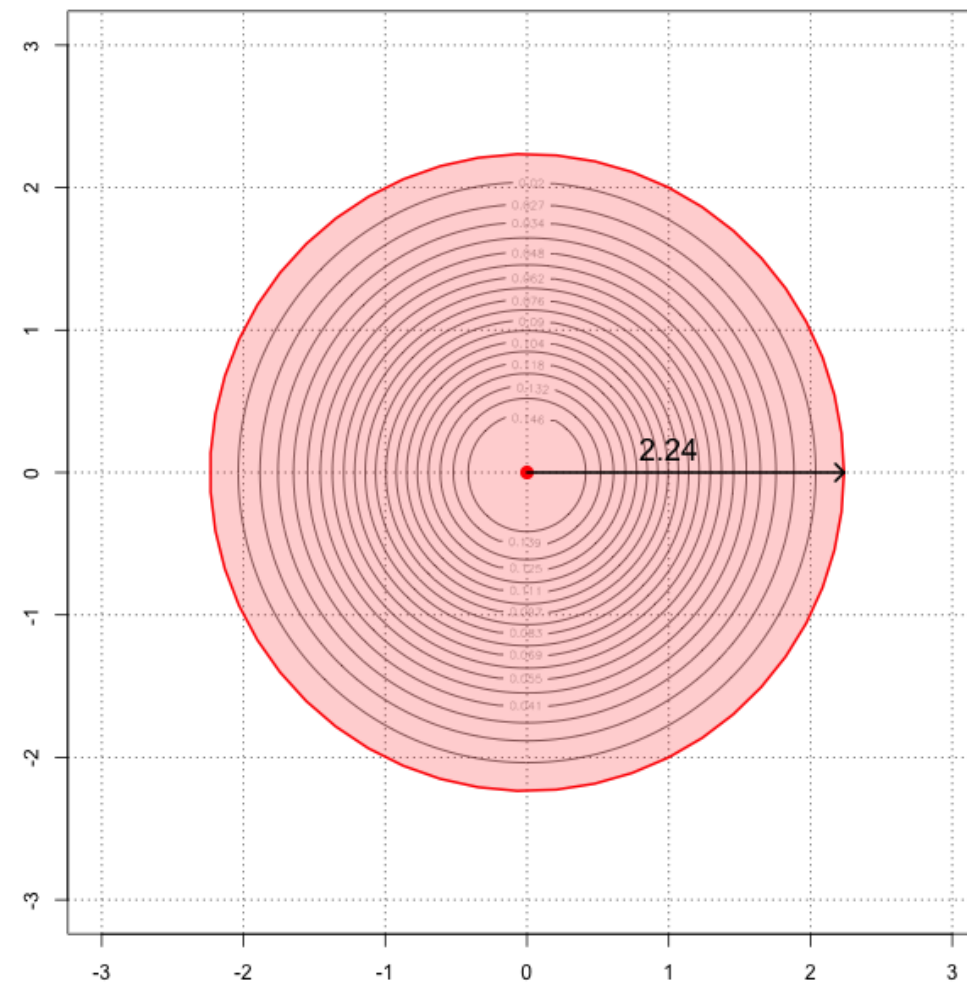
```
sigma1 <- diag(2)
sigma1
      [,1] [,2]
[1,]    1    0
[2,]    0    1
```

```
qmvnorm(p = 0.95, sigma = sigma1,
        tail = "both")
```

```
$quantile
[1] 2.24

$f.quantile
[1] -1.31e-06

attr(,"message")
[1] "Normal Completion"
```



The red circle with radius 2.24 contains 0.95 of the probability



MULTIVARIATE PROBABILITY DISTRIBUTIONS IN R

Let's practice!



MULTIVARIATE PROBABILITY DISTRIBUTIONS IN R

Checking normality of multivariate data

Surajit Ray

Senior Lecturer, University of Glasgow



Why check normality?

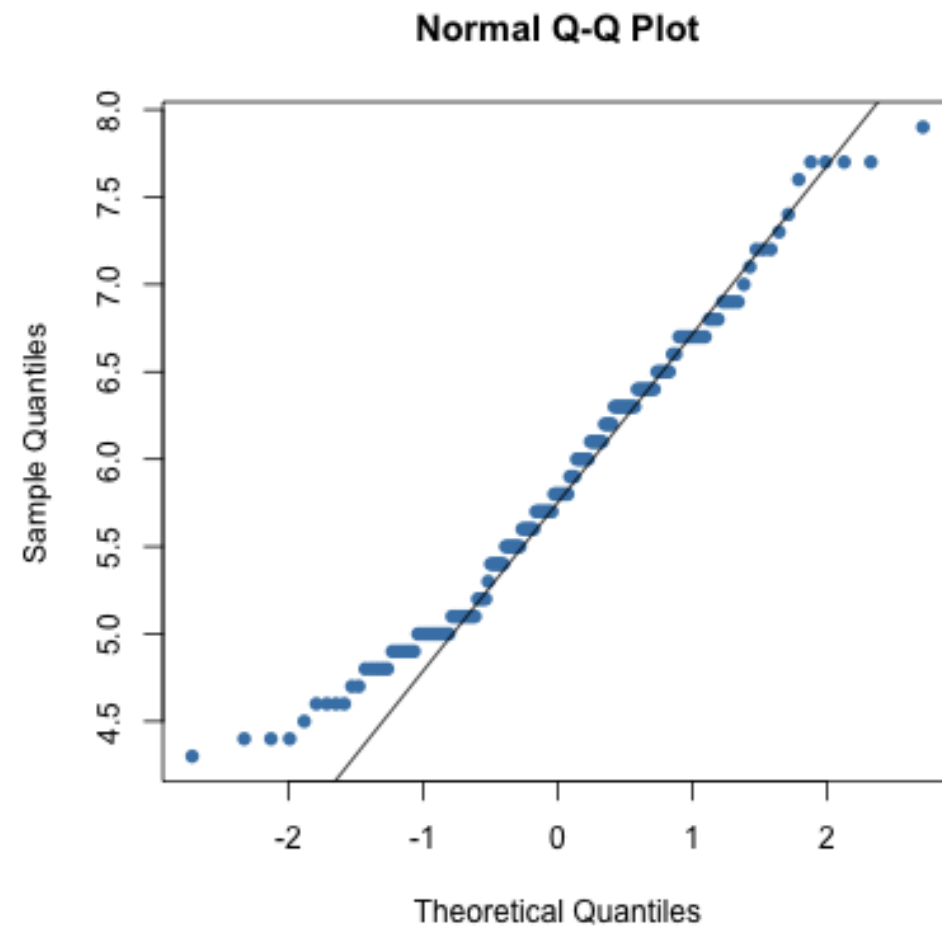
- Classical statistical techniques that assume univariate/multivariate normality:
 - Multivariate regression
 - Discriminant analysis
 - Model-based clustering
 - Principal component analysis (PCA)
 - Multivariate analysis of variance (MANOVA)



Review: univariate normality tests

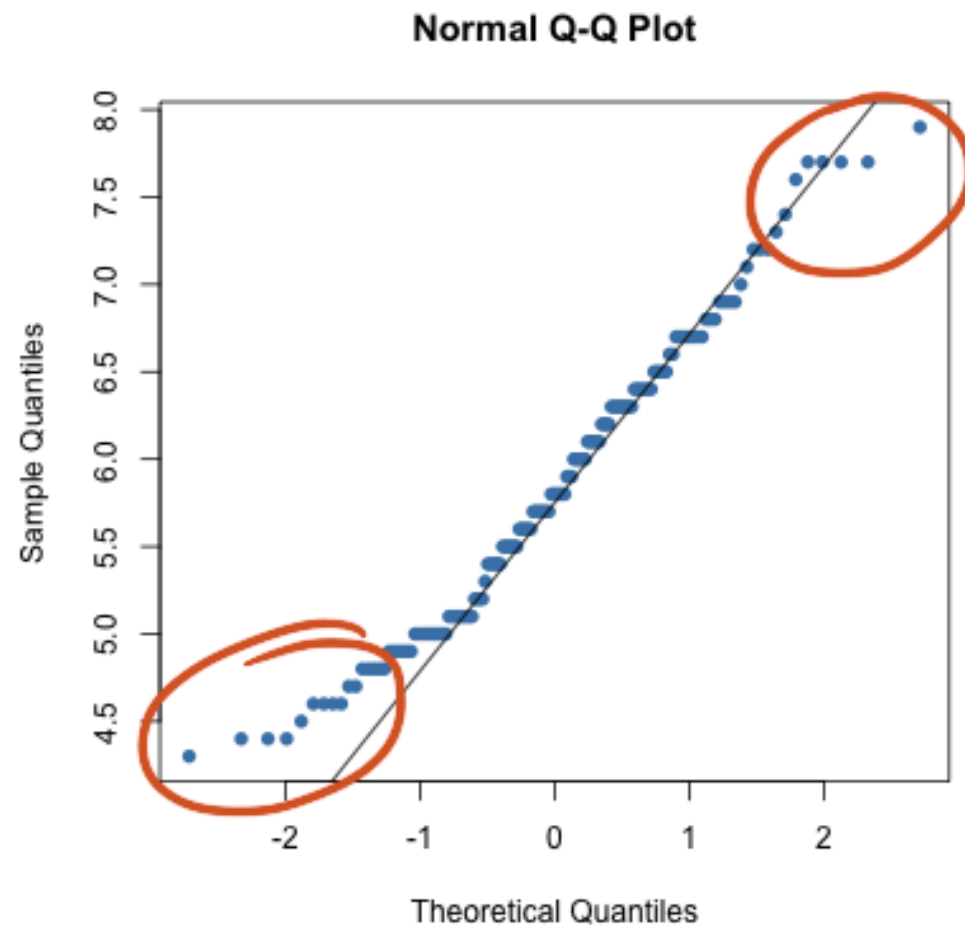
```
qqnorm(iris_raw[, 1])  
qqline(iris_raw[, 1])
```

- If the values lie along the reference line the distribution is close to normal



Review: univariate normality tests

```
qqnorm(iris_raw[, 1])
qqline(iris_raw[, 1])
```

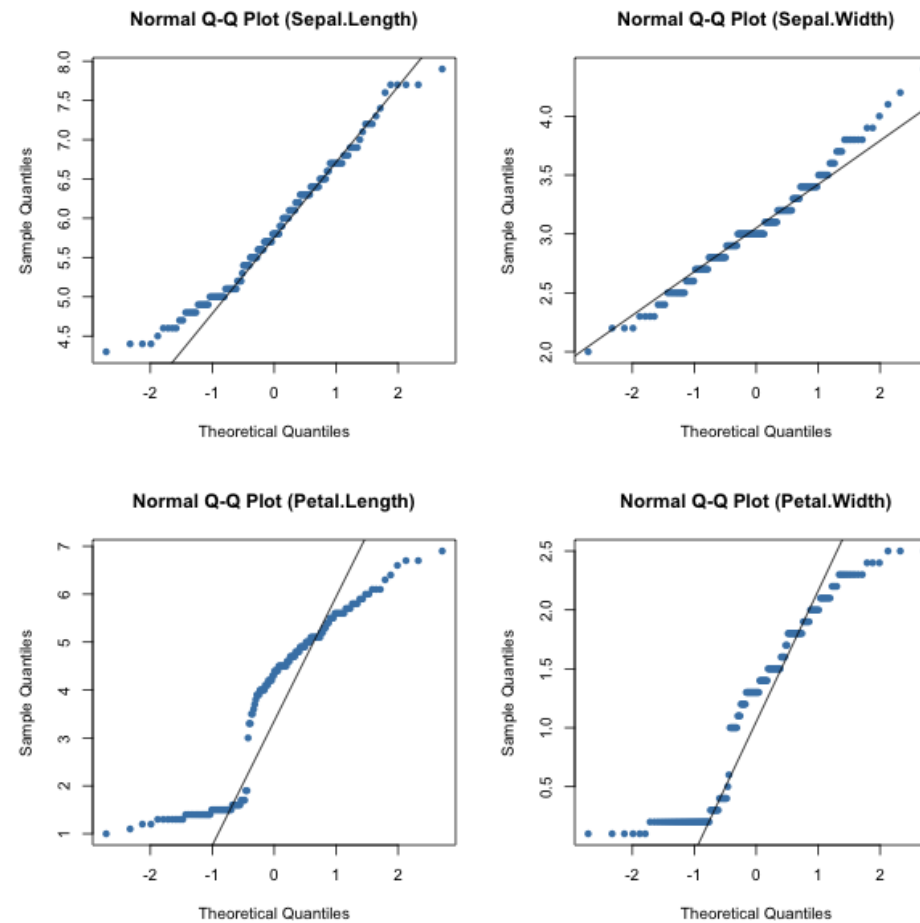


- If the values lie along the reference line the distribution is close to normal
- Deviation from the line might indicate
 - heavier tails
 - skewness
 - outliers
 - clustered data



qqnorm of all variables

```
uniPlot(iris_raw[, 1:4])
```





MVN library multivariate normality test functions

- Multivariate normality tests by
 - Mardia
 - Henze-Zirkler
 - Royston
- Graphical approaches
 - chi-square Q-Q
 - perspective
 - contour plots



MVN library multivariate normality test functions

- Multivariate normality tests by
 - Mardia ✓
 - Henze-Zirkler ✓
 - Royston
- Graphical approaches
 - chi-square Q-Q ✓
 - perspective
 - contour-plots

Using mardiaTest to check multivariate normality

```
mardiaTest(iris_raw[, 1:4])
```

```
Mardia Multivariate Normality Test
```

```
-----  
data : iris_raw[, 1:4]
```

```
g1p      : 2.697  
chi.skew : 67.43  
p.value.skew : 4.758e-07
```

```
g2p      : 23.74  
z.kurtosis : -0.2301  
p.value.kurt : 0.818
```

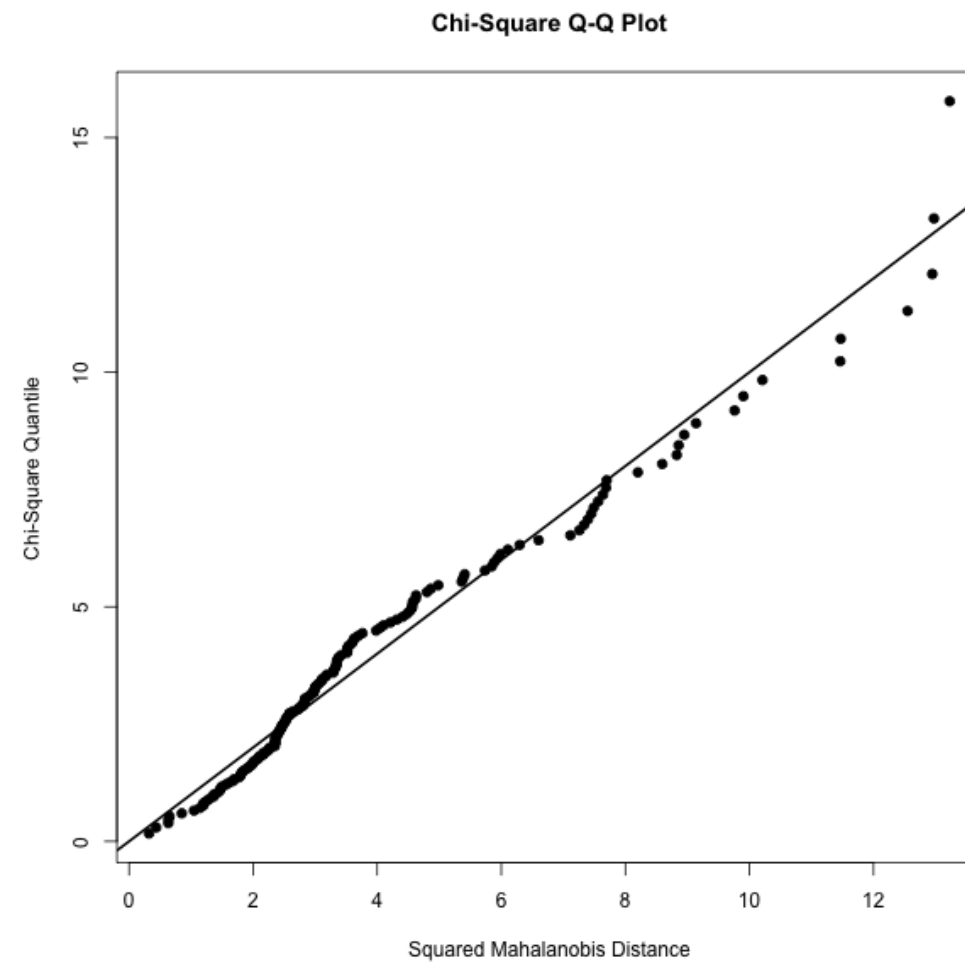
```
chi.small.skew : 69.33  
p.value.small : 2.342e-07
```

```
Result      : Data are not multivariate normal.  
-----
```



Using qqplot from mardiaTest to check multivariate normality

```
mardiaTest(iris_raw[, 1:4], qqplot = TRUE)
```





Using hzTest to check multivariate normality

```
hzTest(iris_raw[,1:4])
```

```
Henze-Zirkler's Multivariate Normality Test
```

```
-----  
data : iris_raw[, 1:4]
```

```
HZ      : 2.333269
```

```
p-value : 0
```

```
Result  : Data are not multivariate normal.  
-----
```

Testing multivariate normality by species

```
mardiaTest(iris[iris_raw$Species  
              == "setosa", 1:4])
```

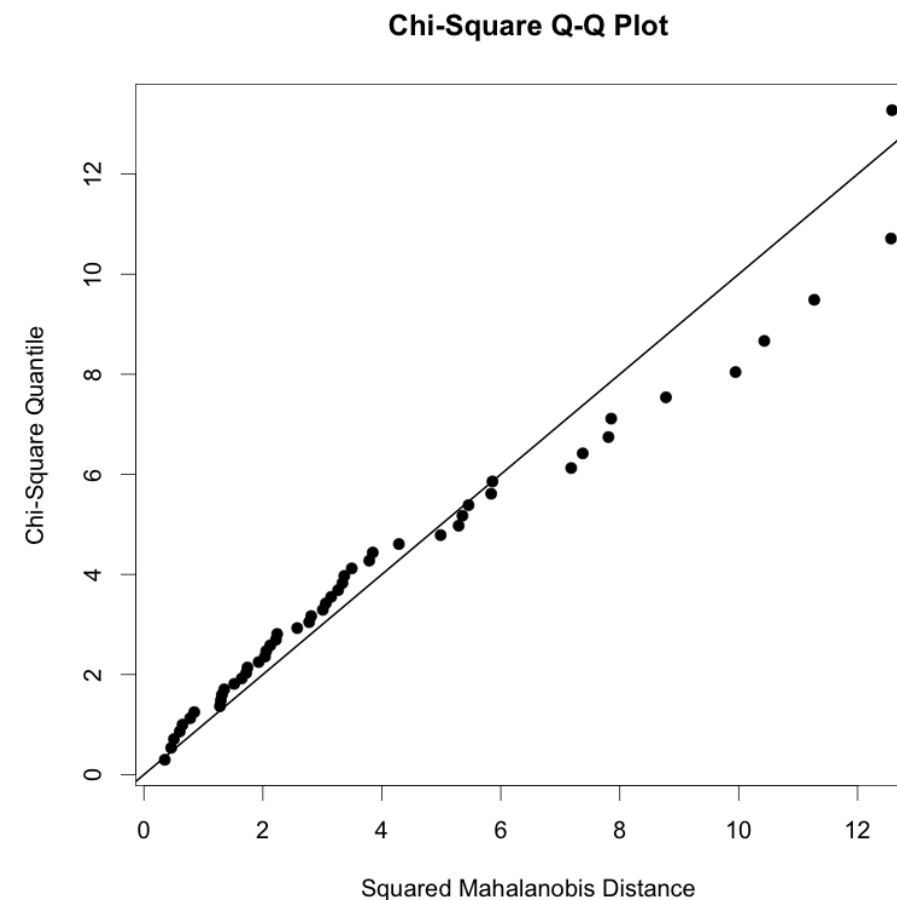
Mardia's Multivariate Normality Test

```
g1p      : 3.08  
chi.skew : 25.7  
p.value.skew : 0.177
```

```
g2p      : 26.5  
z.kurtosis : 1.29  
p.value.kurt : 0.195
```

```
chi.small.skew : 27.85973  
p.value.small  : 0.1127617
```

```
Result      : Data are  
              multivariate normal.
```





MULTIVARIATE PROBABILITY DISTRIBUTIONS IN R

**Let's make use of the tests
for multivariate normality!**