



DATA VISUALIZATION WITH GGPLOT2

# Graphics of Large Data

# Defining largeness

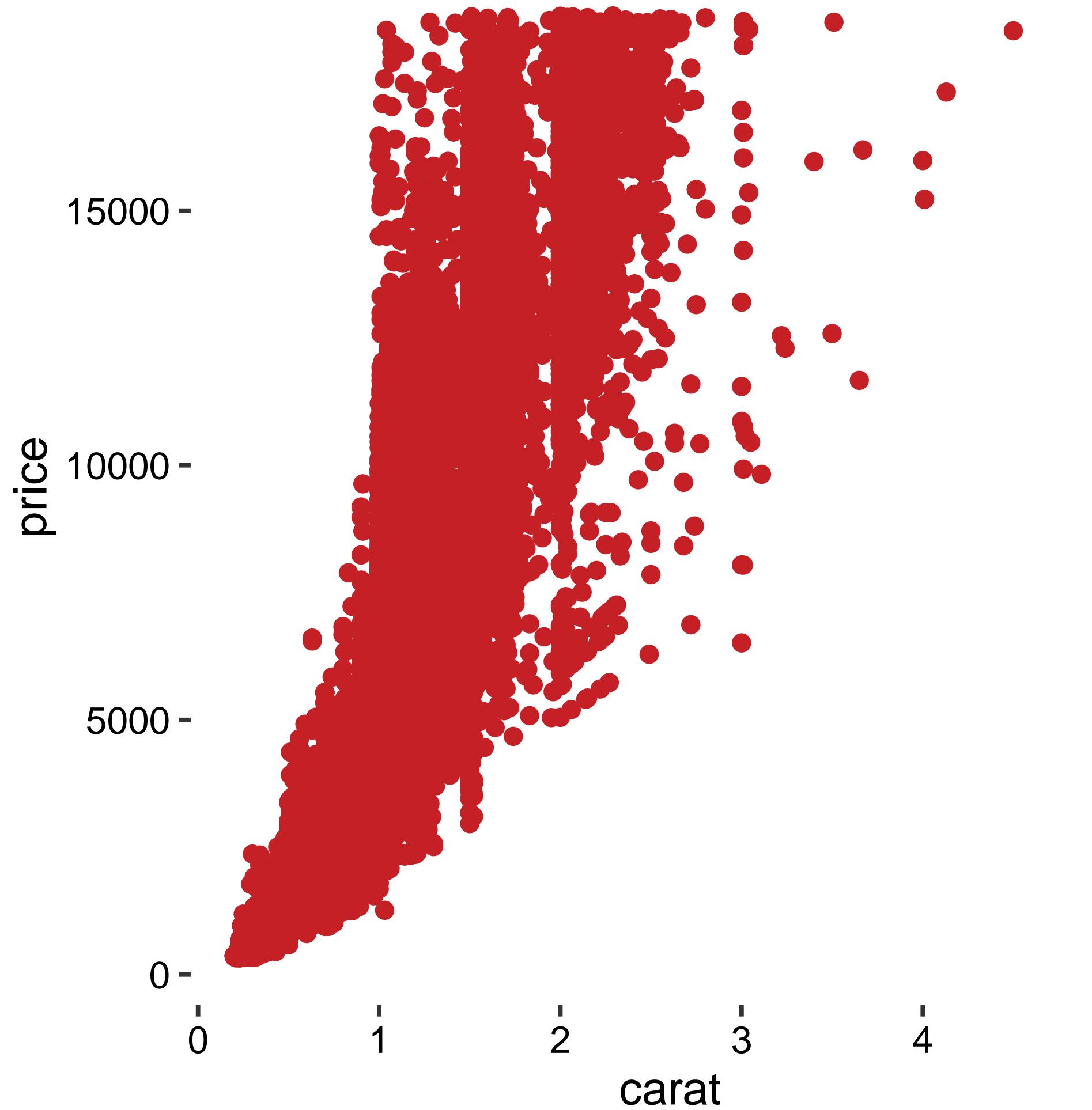
- Many observations
  - High-resolution time series
  - Large surveys
  - Website analytics
- Many variables
  - Multidimensional data
- Combination

# Many observations

```
> dim(diamonds)
[1] 53940    10

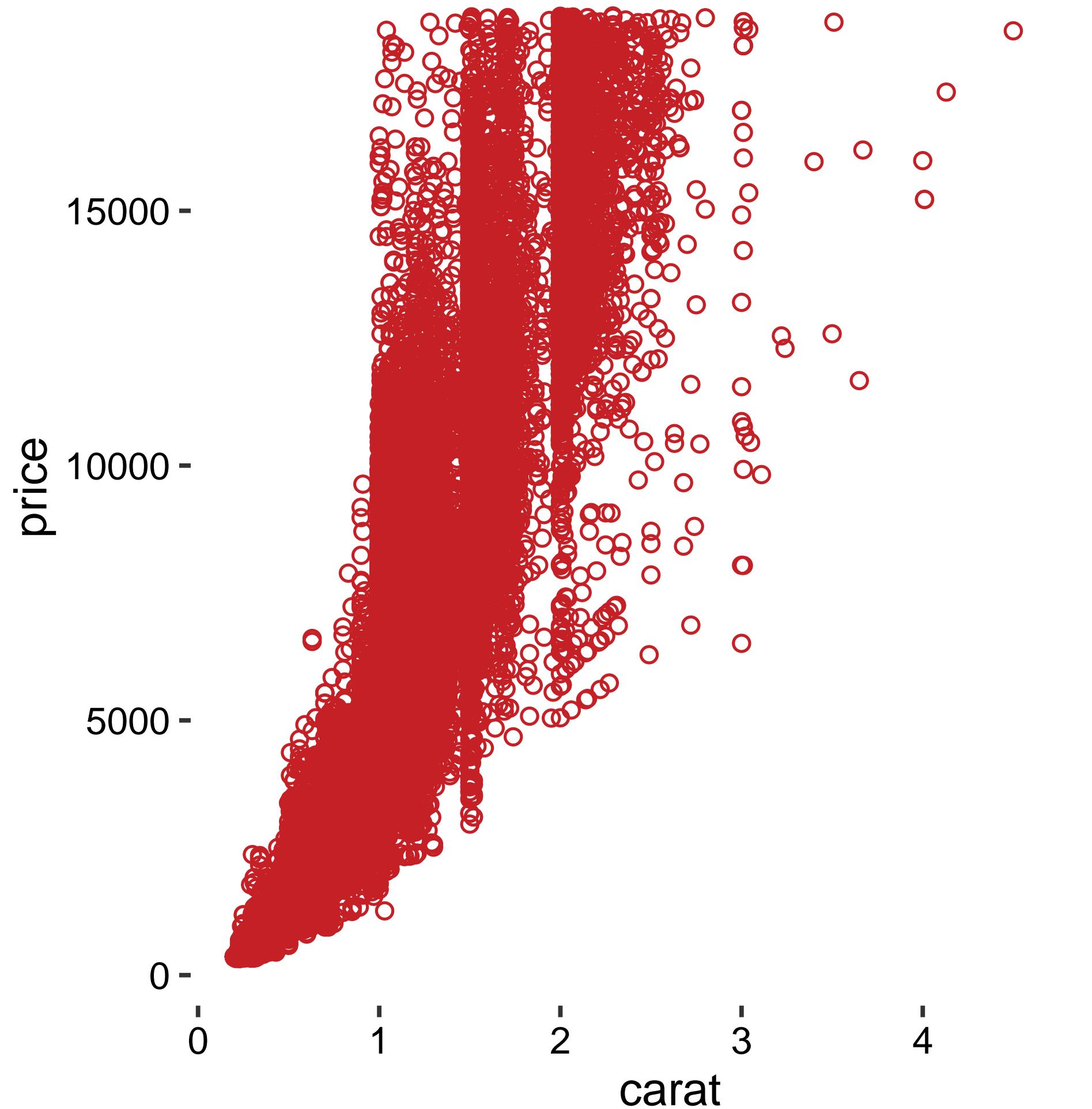
> head(diamonds)
  carat       cut color clarity depth table price     x     y     z
1  0.23      Ideal    E     SI2   61.5     55   326 326 3.95 3.98 2.43
2  0.21      Premium  E     SI1   59.8     61   326 326 3.89 3.84 2.31
3  0.23      Good    E     VS1   56.9     65   327 327 4.05 4.07 2.31
4  0.29      Premium  I     VS2   62.4     58   334 334 4.20 4.23 2.63
5  0.31      Good    J     SI2   63.3     58   335 335 4.34 4.35 2.75
6  0.24  Very Good  J     VVS2   62.8     57   336 336 3.94 3.96 2.48
```

# Scatter plot



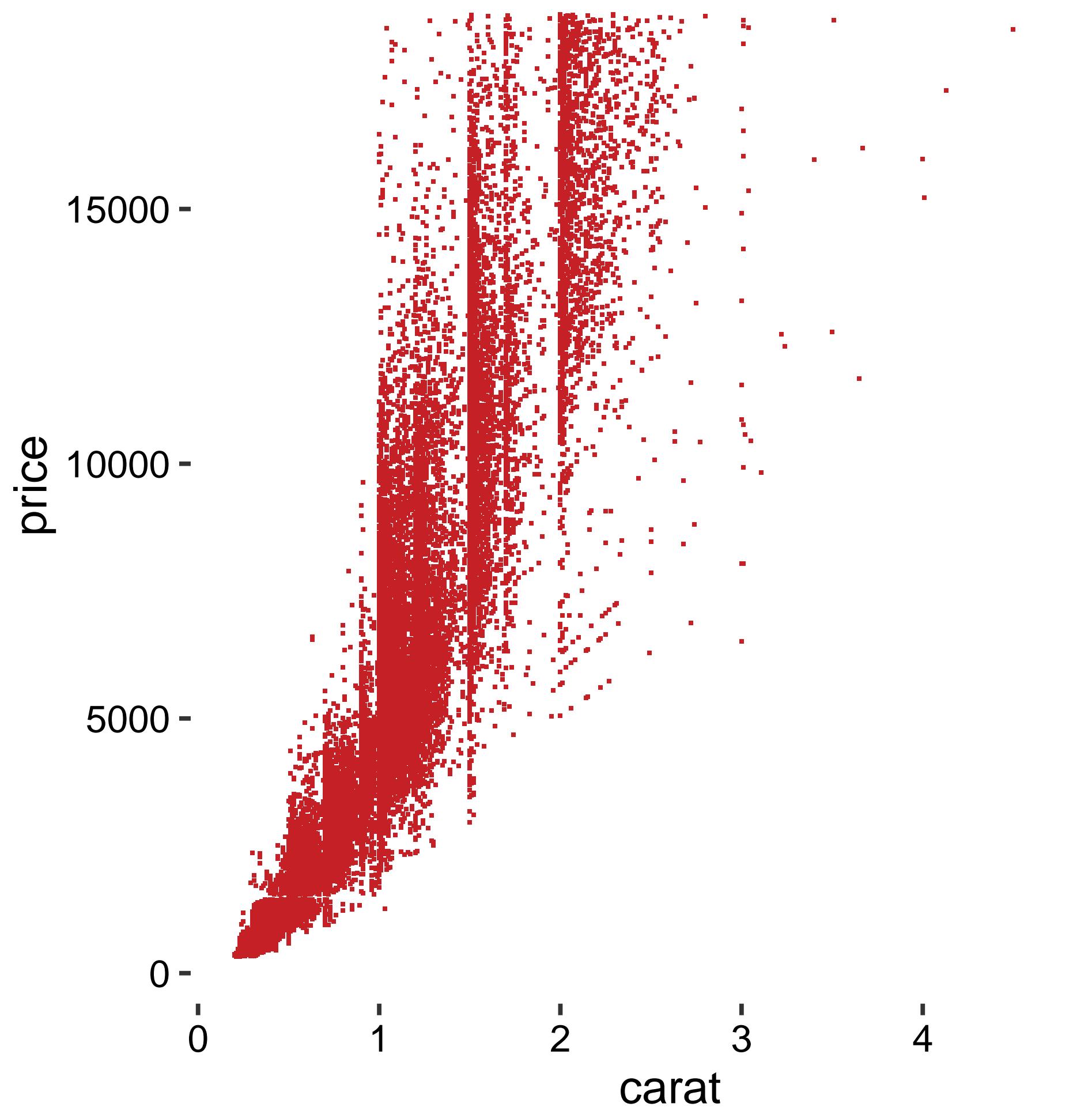
```
> ggplot(diamonds, aes(x = carat, y = price)) +  
  geom_point() # simplified
```

# Adjust symbol



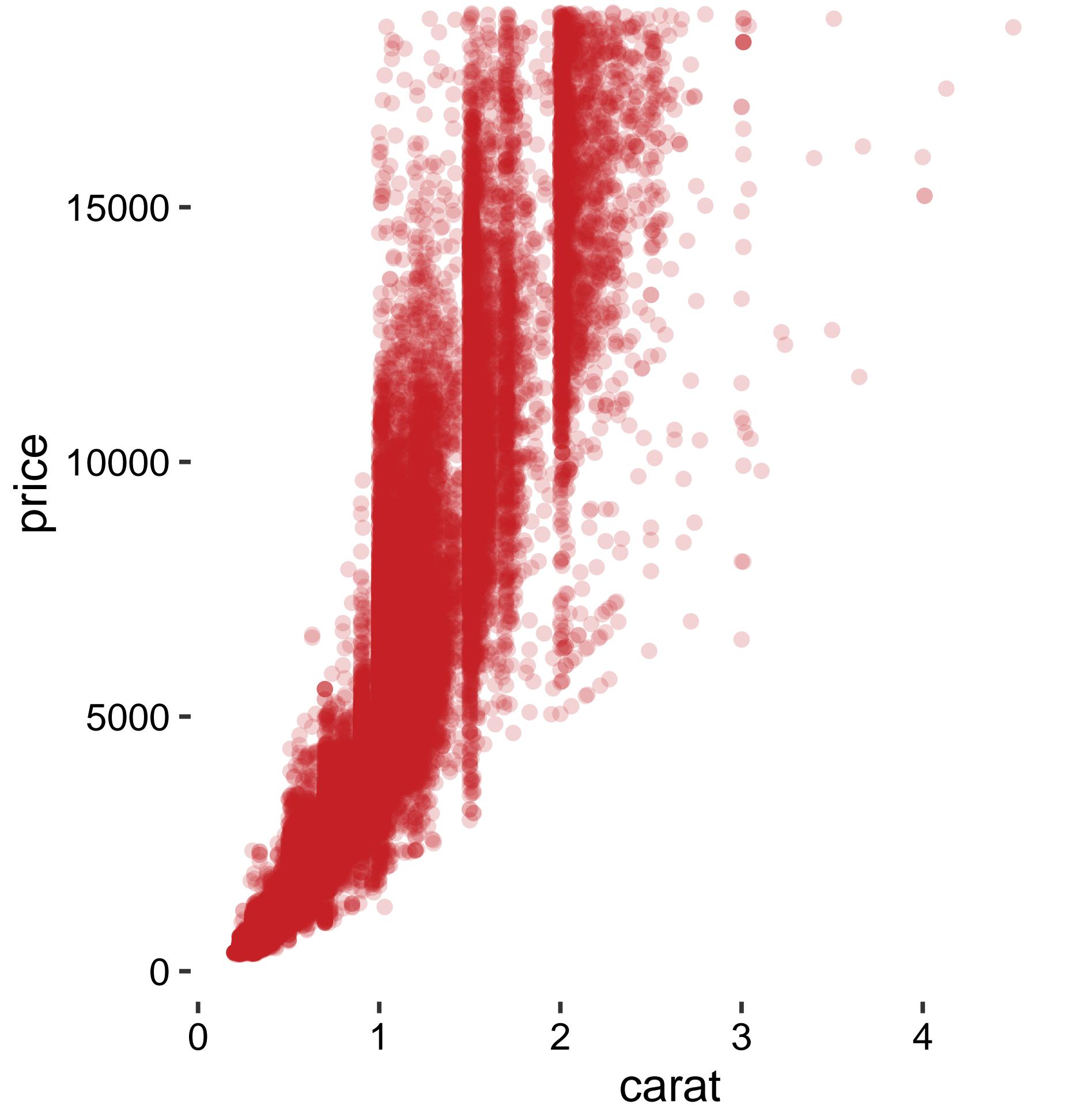
```
> ggplot(diamonds, aes(x = carat, y = price)) +  
  geom_point(shape = 1)
```

# Adjust size



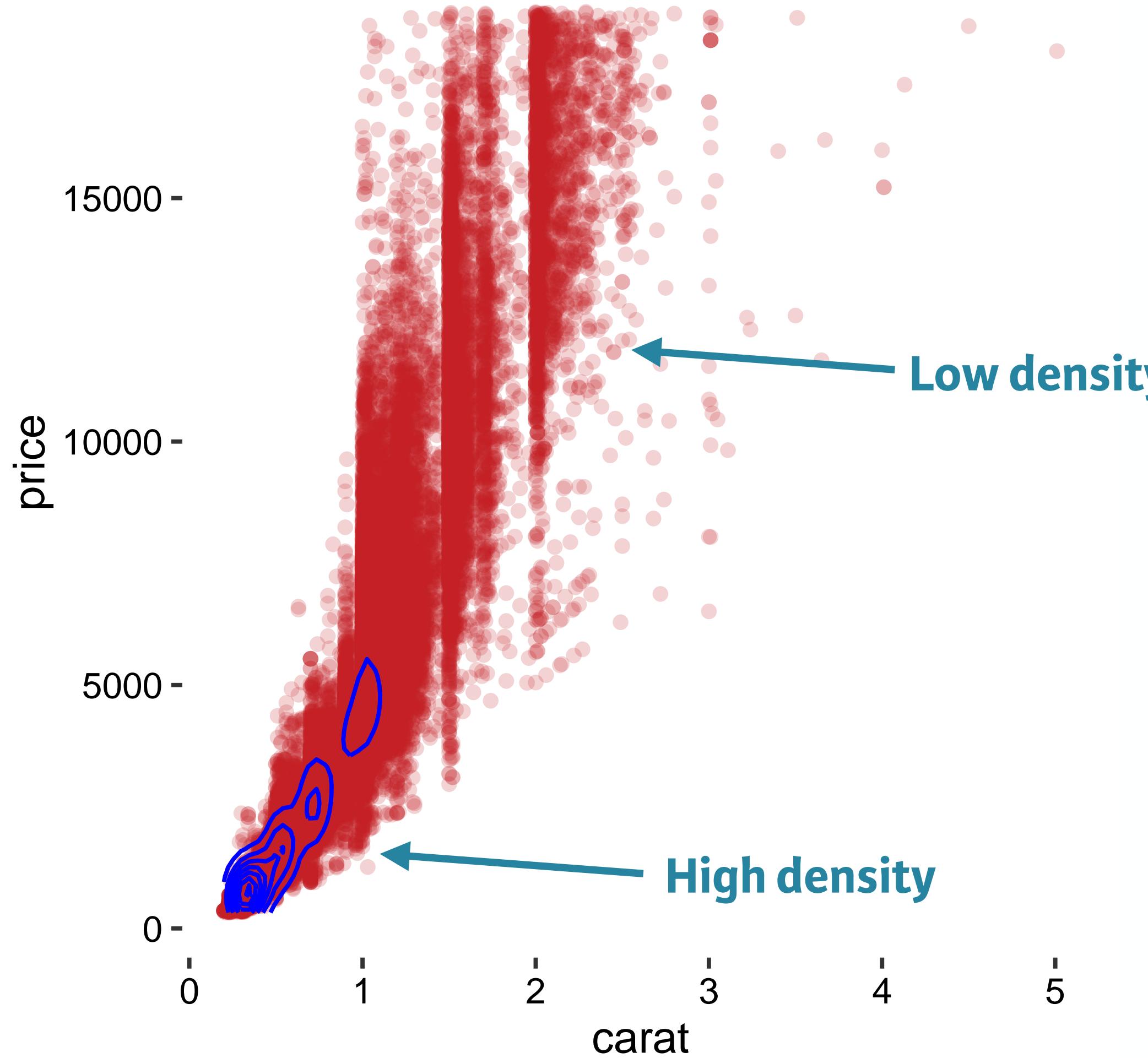
```
> ggplot(diamonds, aes(x = carat, y = price)) +  
  geom_point(shape = ".")
```

# Adjust opacity



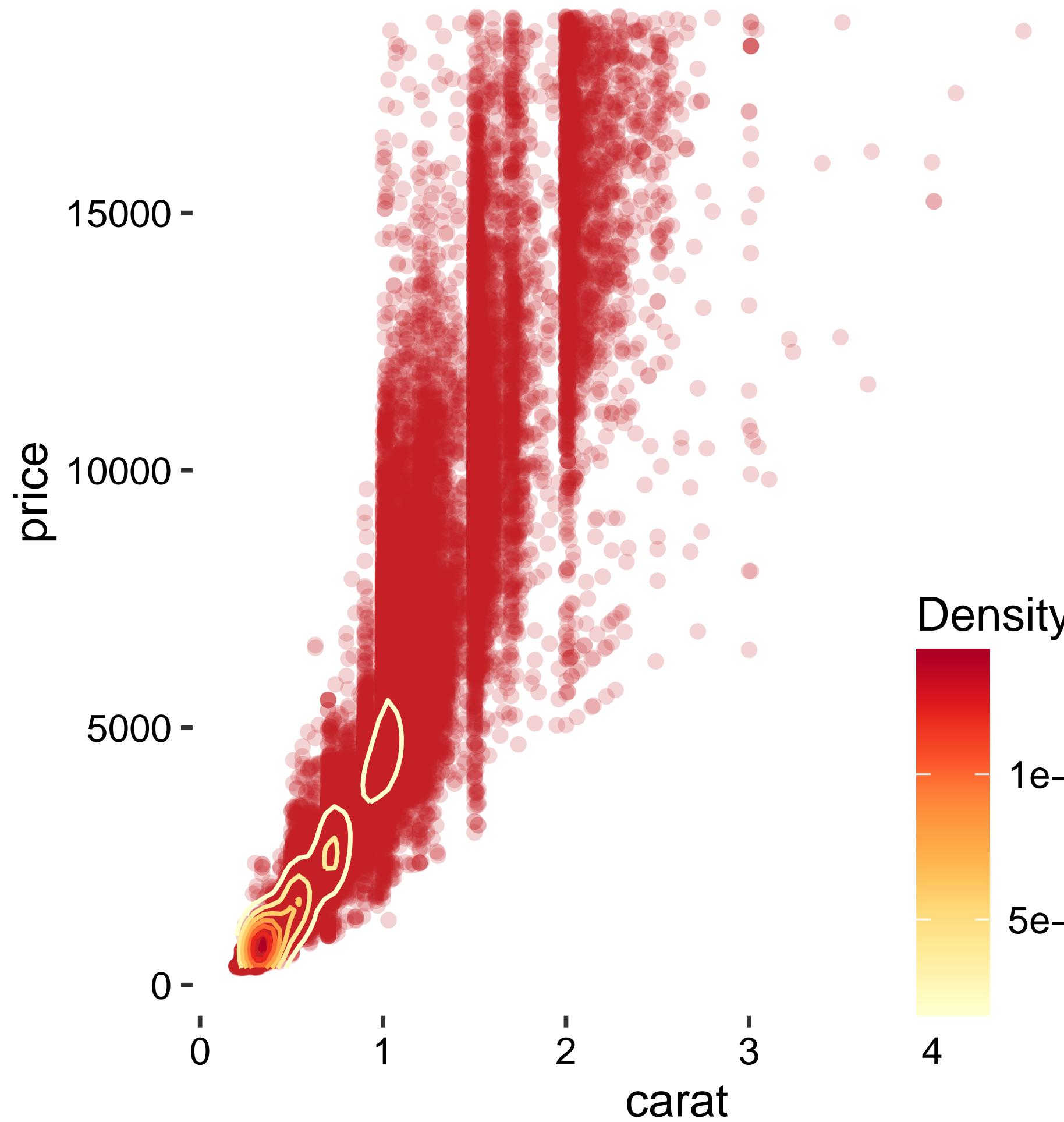
```
> ggplot(diamonds, aes(x = carat, y = price)) +  
  geom_point(shape = 16, alpha = 0.2)
```

# 2D density



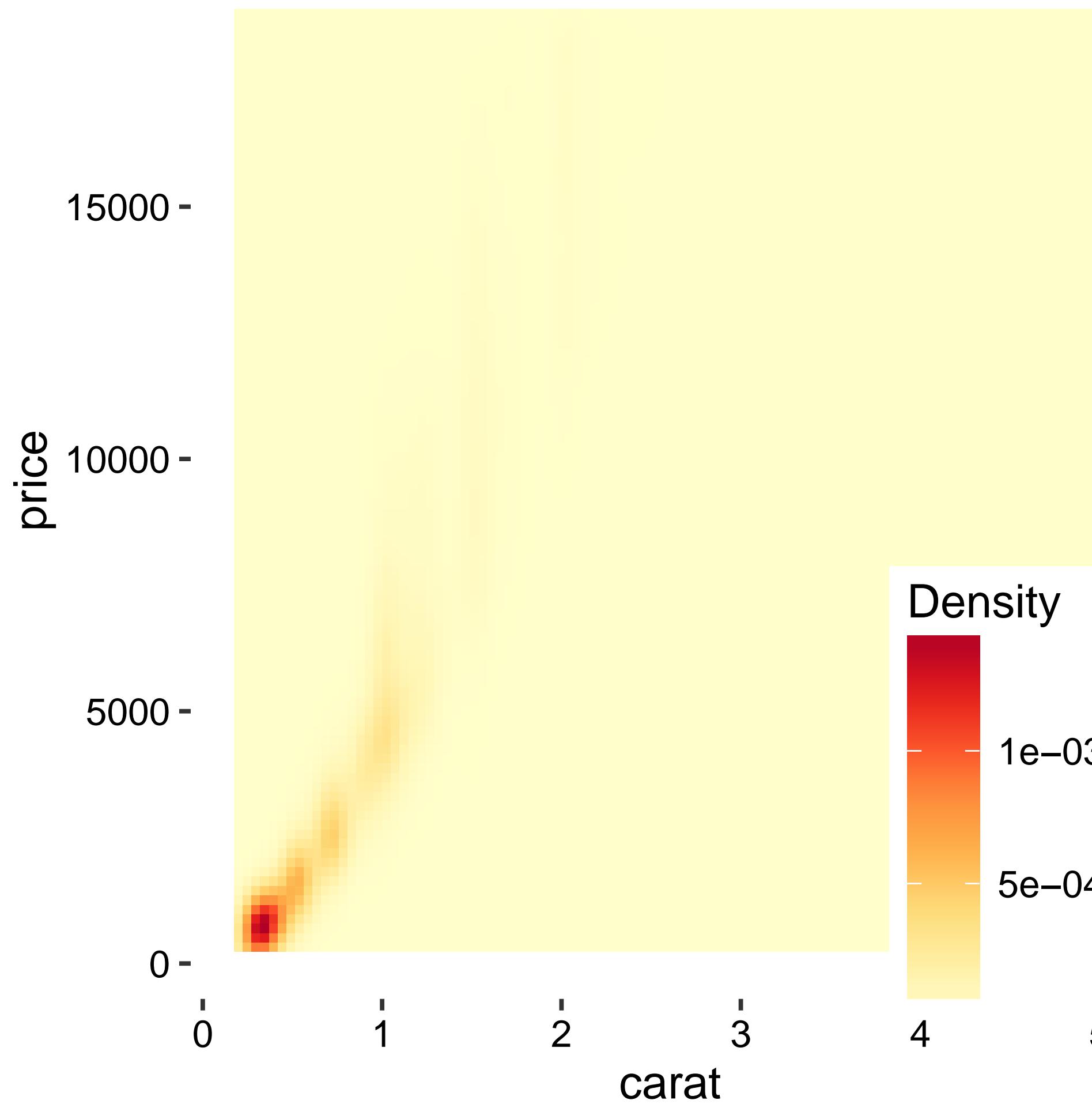
```
> ggplot(diamonds, aes(x = carat, y = price)) +  
  geom_point(shape = 16, alpha = 0.2) +  
  stat_density2d(col = "blue")
```

# 2D density (2)



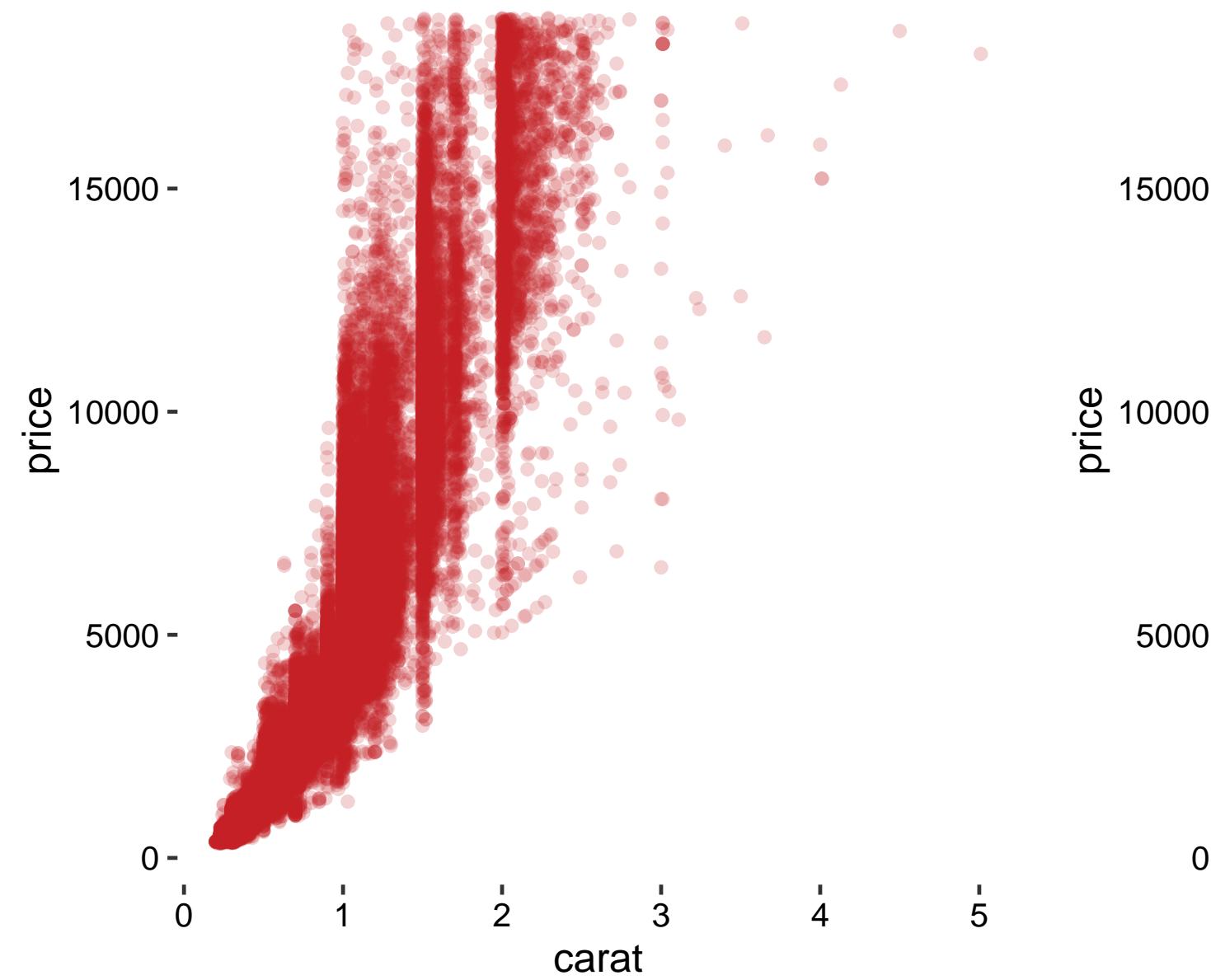
```
> ggplot(diamonds, aes(x = carat, y = price)) +  
  geom_point(shape = 16, alpha = 0.2) +  
  stat_density2d(aes(colour=..level..))
```

# 2D density (3)

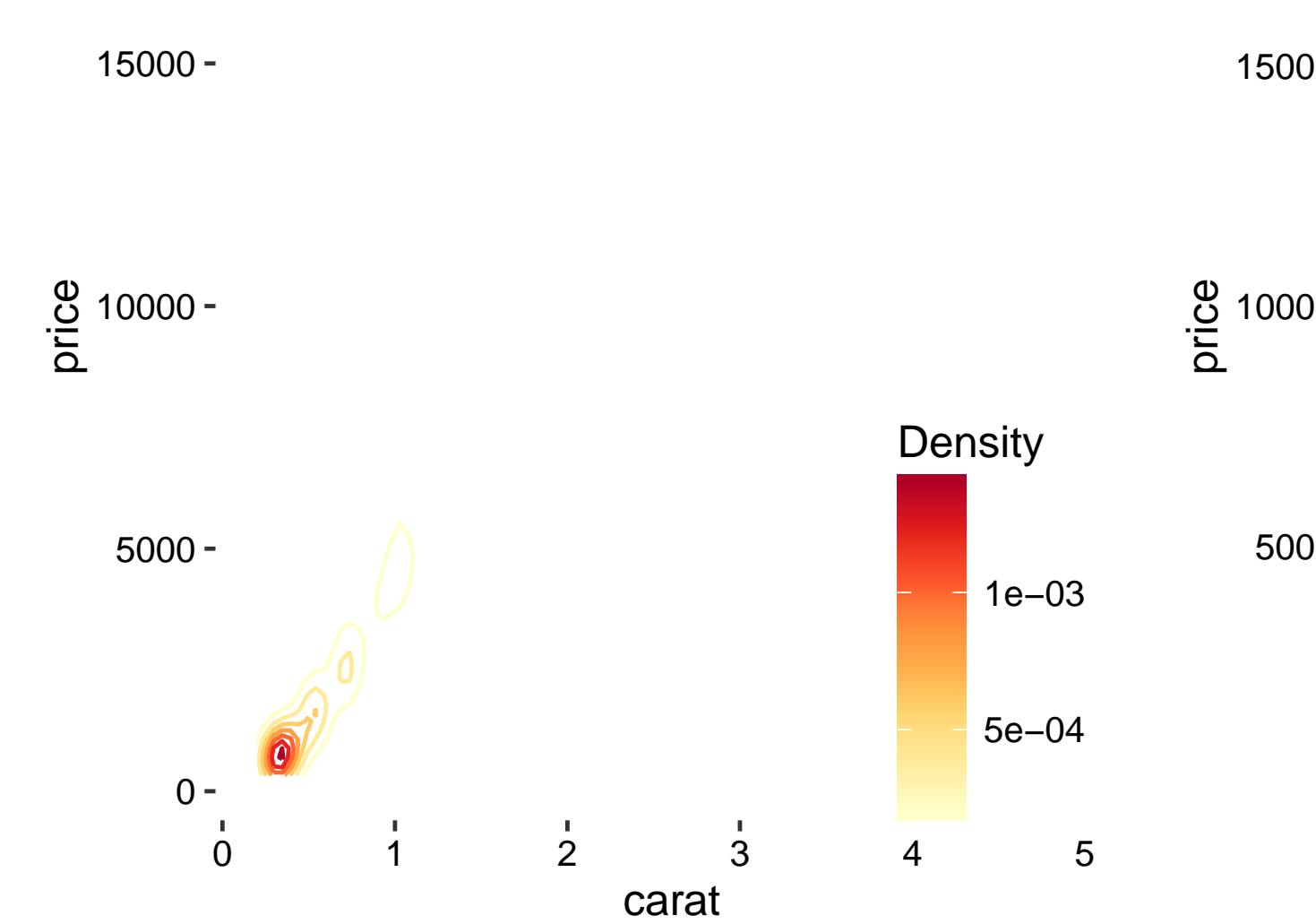


```
> ggplot(diamonds, aes(x = carat, y = price)) +  
  stat_density2d(geom = "tile",  
    aes(fill = ..density..),  
    contour = FALSE)
```

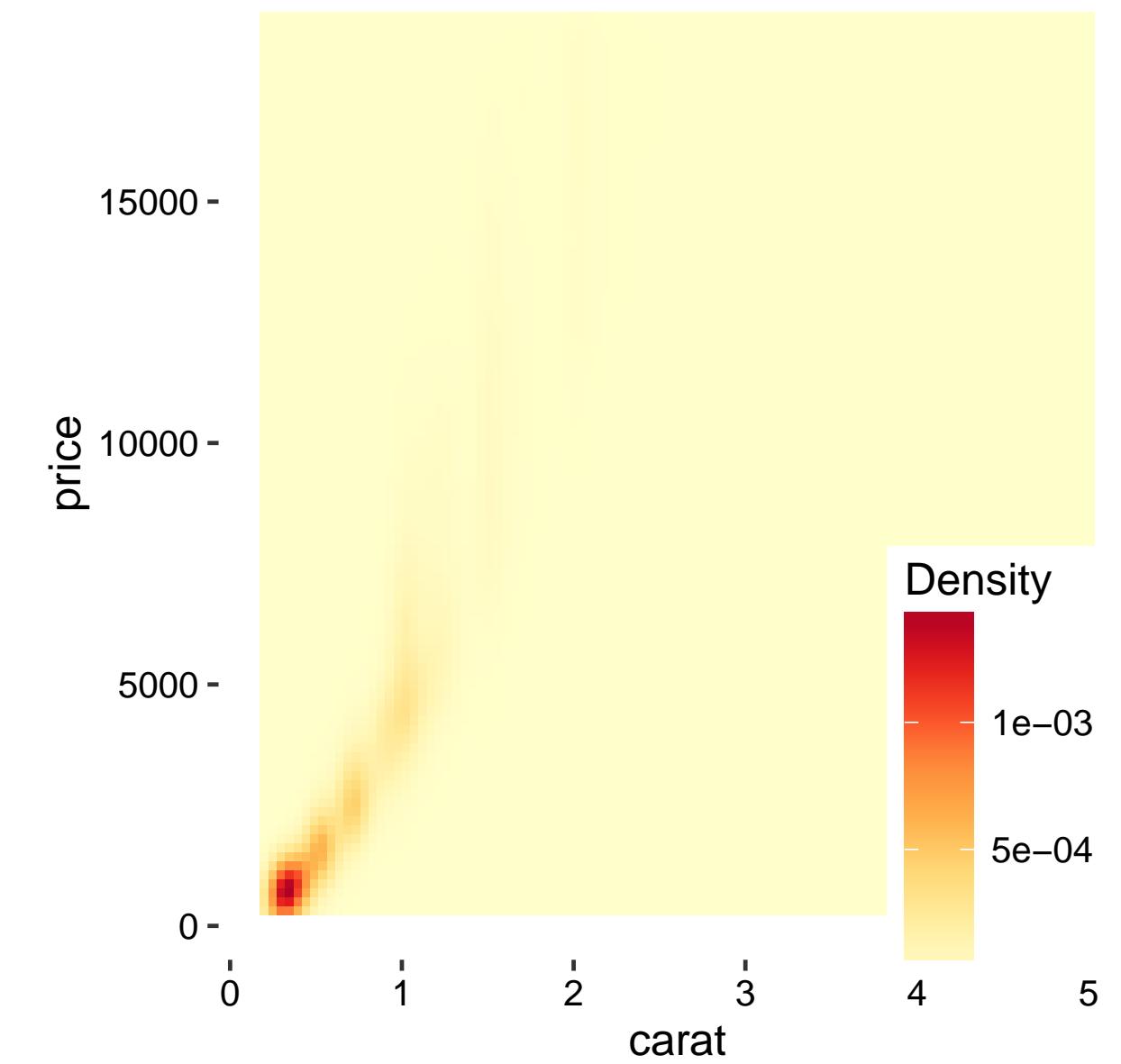
# Comparison



Opacity

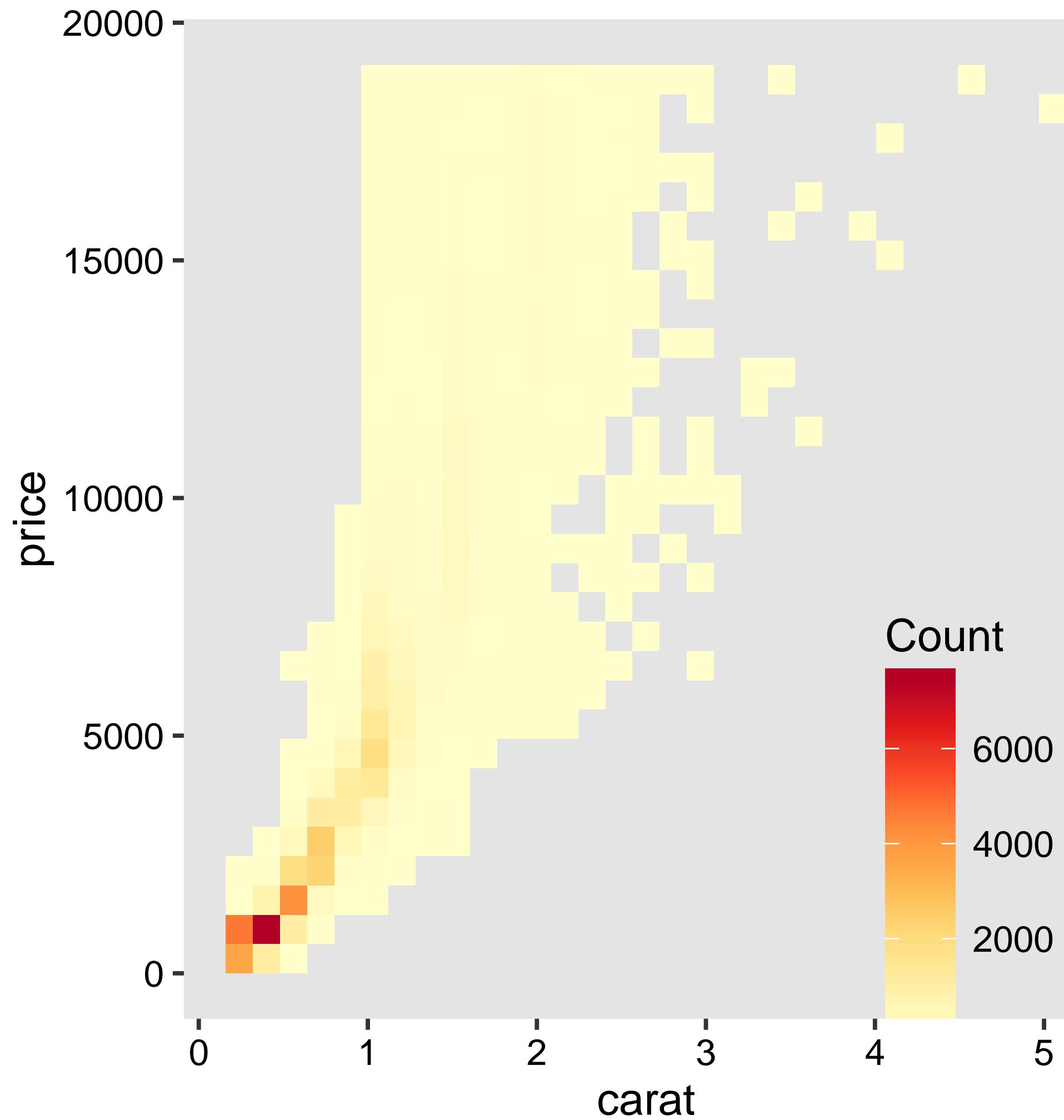


colored contours



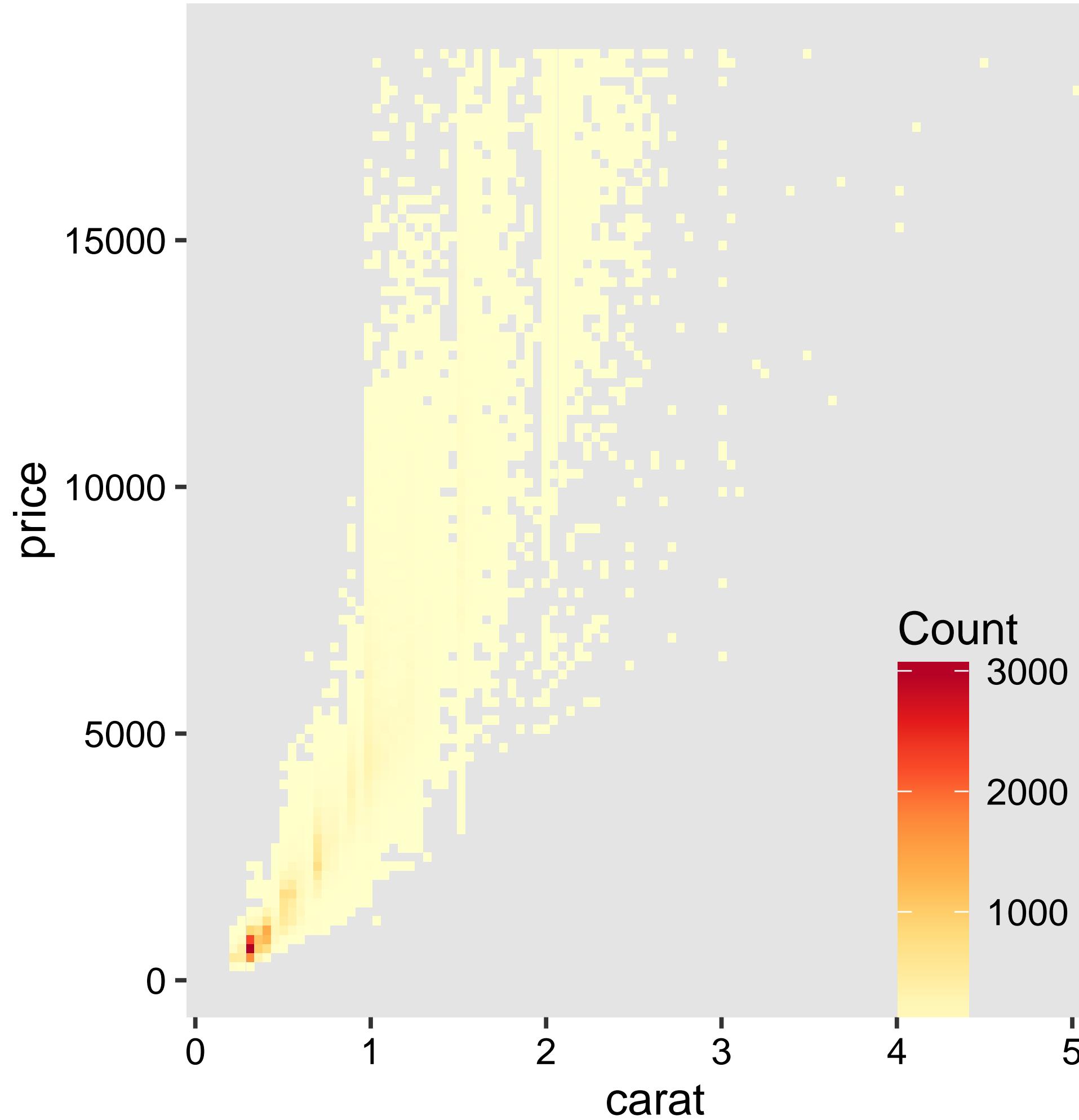
heatmap

# Binning (1)



```
> ggplot(diamonds, aes(x = carat, y = price)) +  
  geom_bin2d()
```

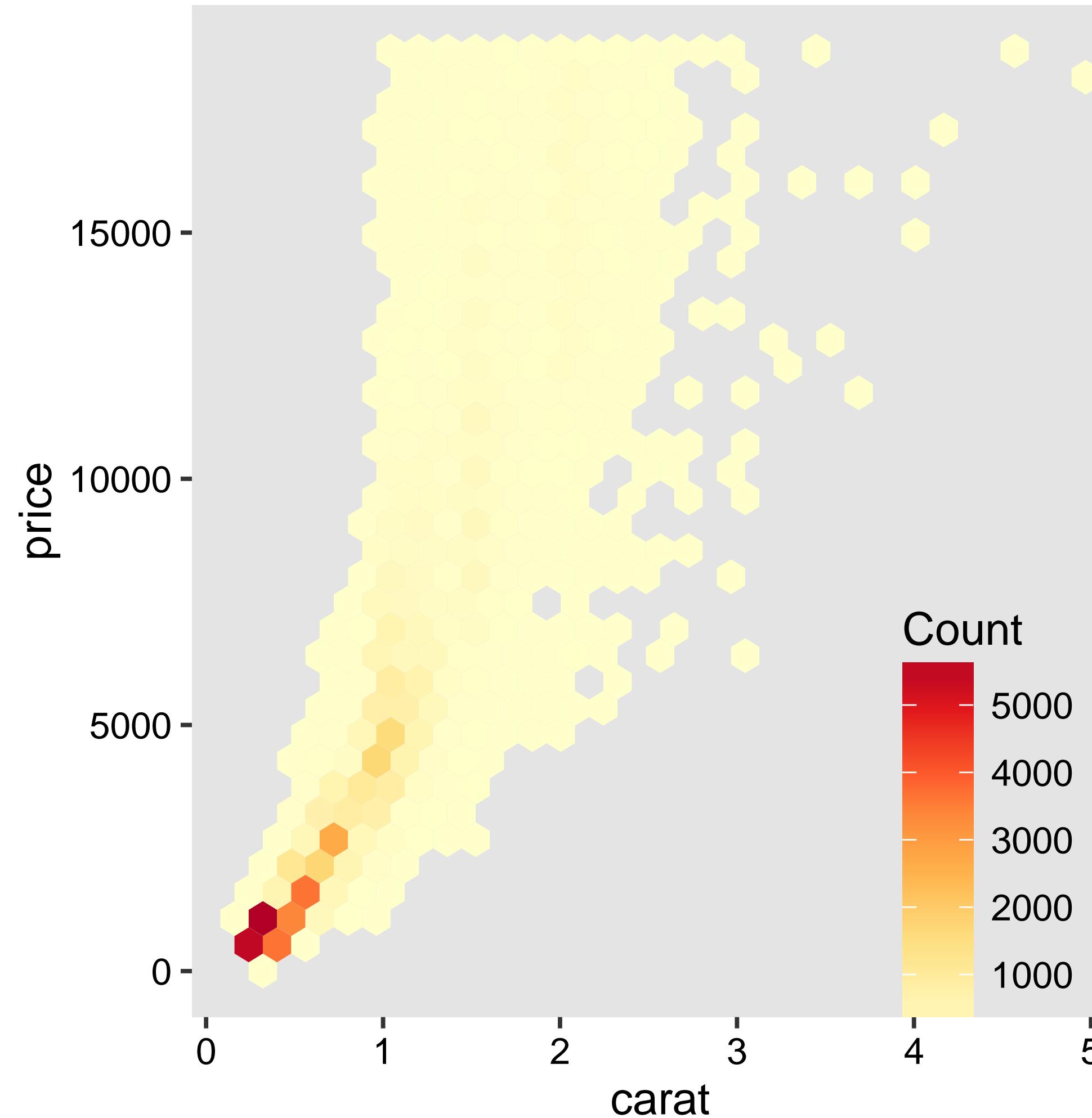
# Binning (2)



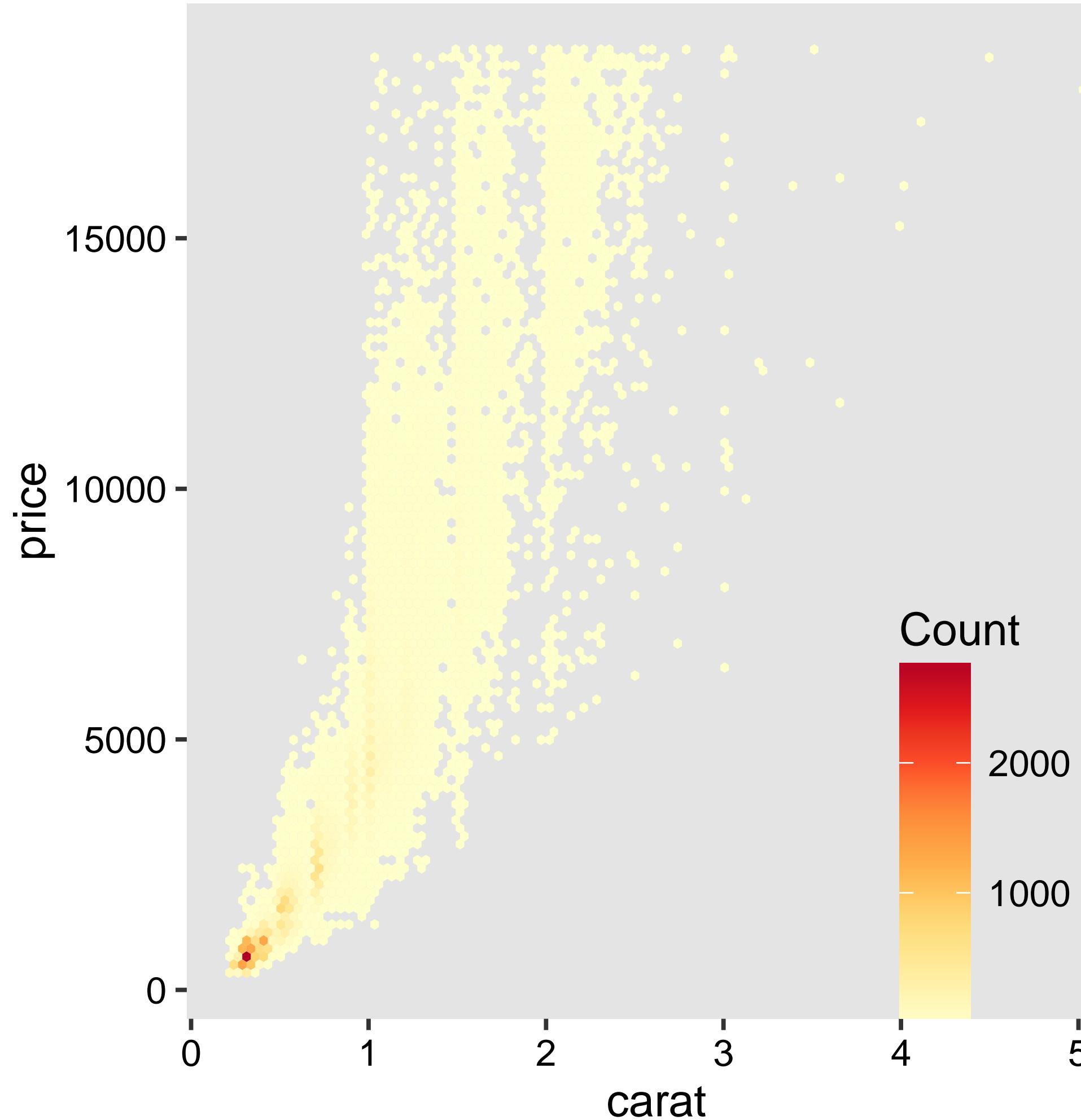
```
> ggplot(diamonds, aes(x = carat, y = price)) +  
  geom_bin2d(bins = 100)
```

# Hex-binning (1)

```
> ggplot(diamonds, aes(x = carat, y = price)) +  
  geom_hex()
```



# Hex-binning (2)



```
> ggplot(diamonds, aes(x = carat, y = price)) +  
  geom_hex(bins = 100)
```

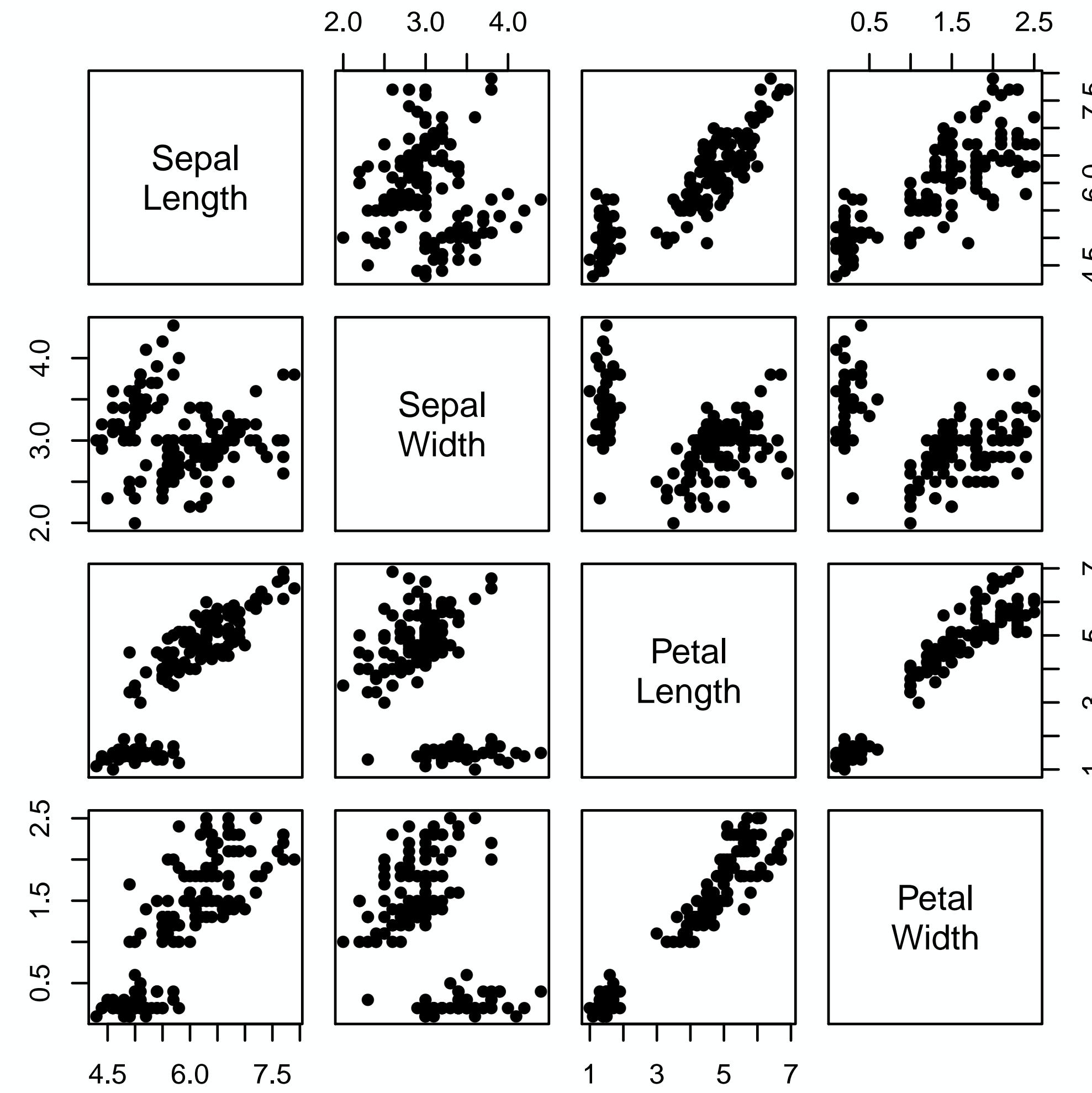
# Many observations

- Reducing over-plotting
- Reducing amount of information that is plotted
- Aggregates

# Many variables

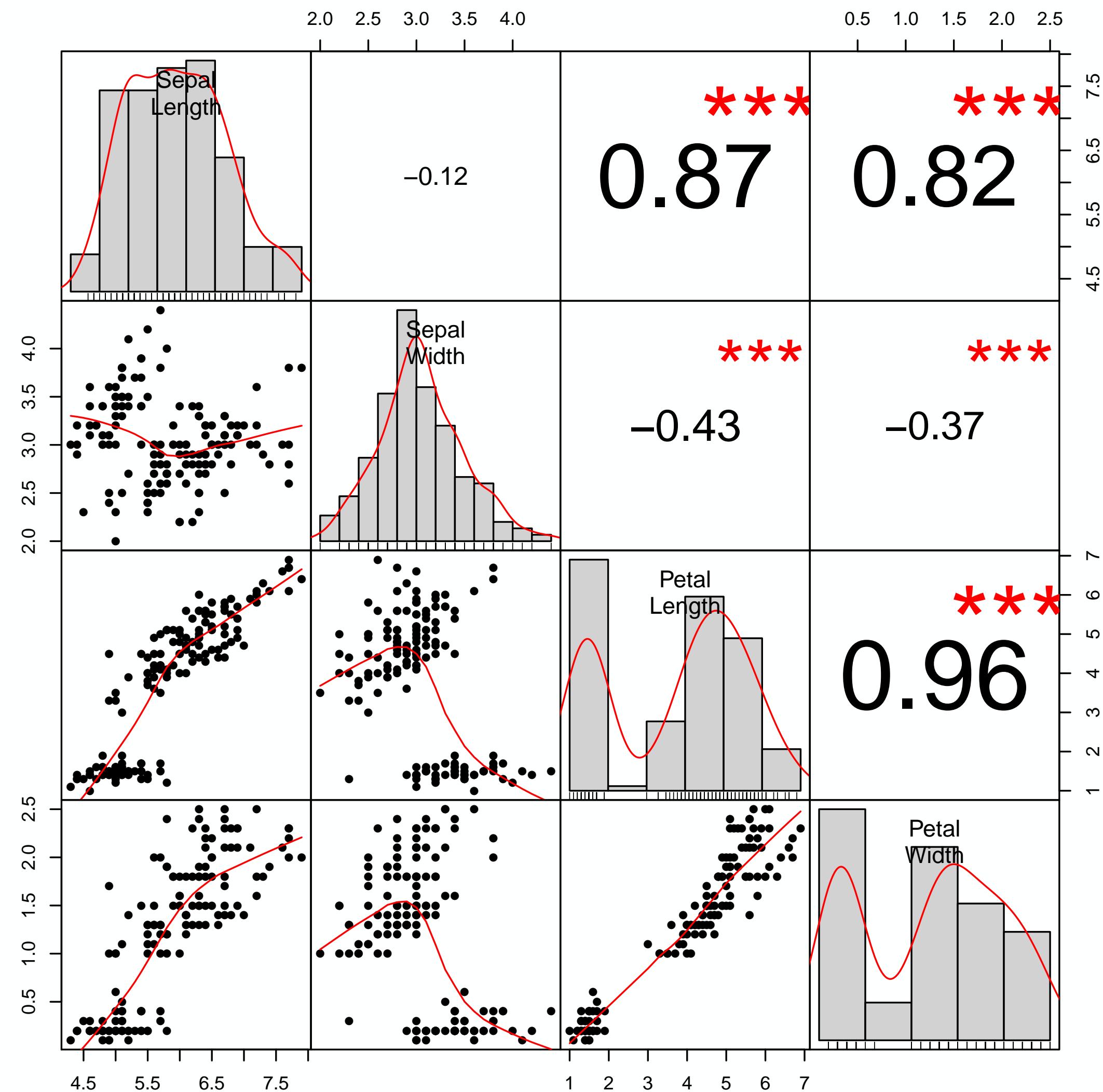
- Multi-variate or high-dimensional data
- Data reduction methods
- Previously: facets
- Two plot types

# SPLOM - Scatter PLOT Matrix



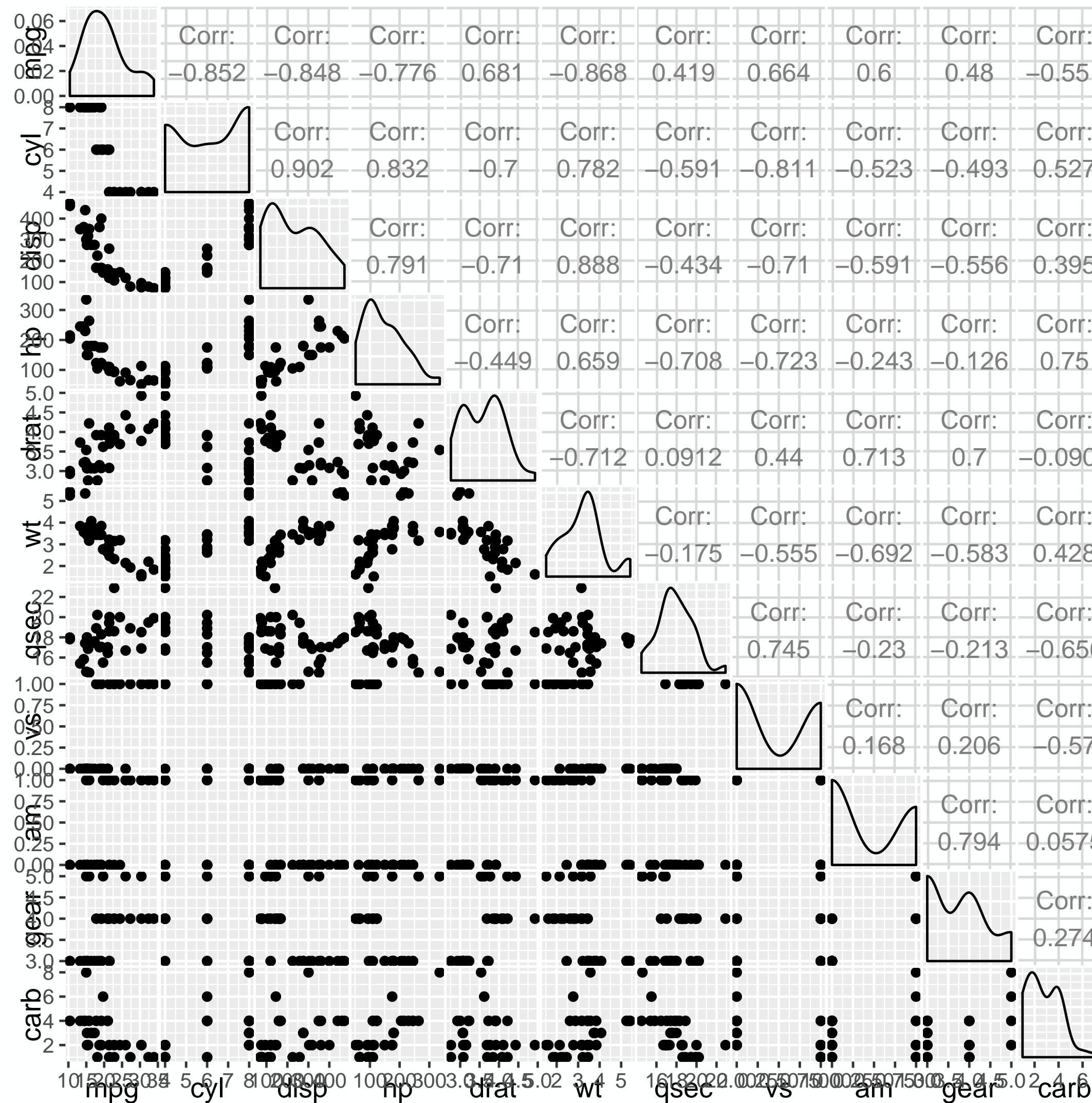
```
> pairs(iris[-5])
```

# Correlation matrix



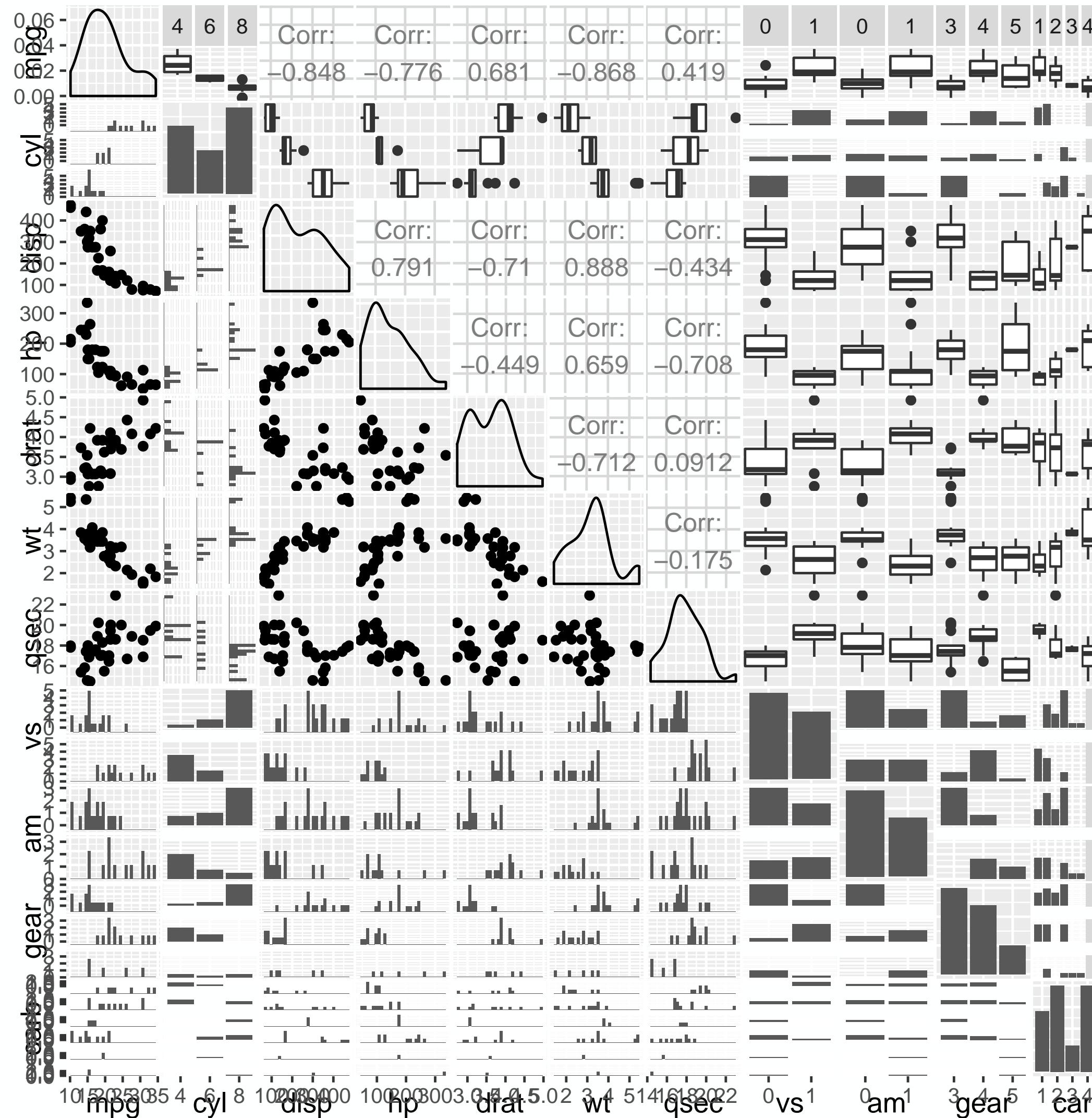
```
> library(PerformanceAnalytics)  
> chart.Correlation(iris[-5])
```

# ggAlly::ggpairs()



```
> library(GGally)  
> ggpairs(mtcars)
```

# ggAlly::ggpairs()



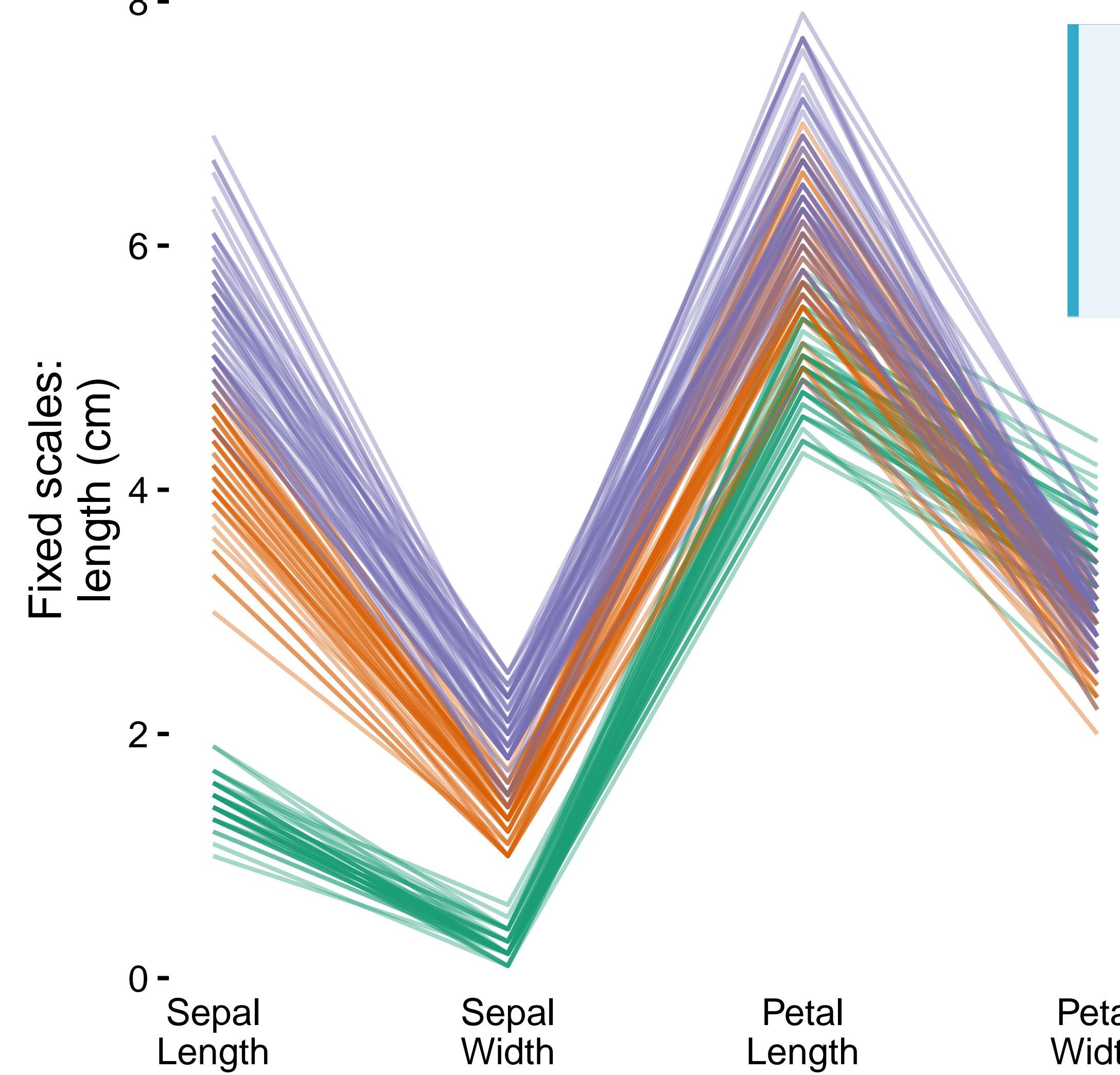
```
> library(GGally)
```

```
> # make columns categorical
```

```
> ... # omitted
```

```
> ggpairs(mtcars)
```

# Parallel coordinate plot



```
> ggparcoord(iris, columns = 1:4,  
  groupColumn = 5,  
  scale = "globalminmax",  
  order = "anyClass", alphaLines = 0.4)
```



DATA VISUALIZATION WITH GGPLOT2

**Let's practice!**



DATA VISUALIZATION WITH GGPLOT2

# Ternary Plots

# Ternary plot

- Triangle plot
- Compositional trivariate data
- Add up to 100%
- Typical example: soil composition
  - Sand
  - Silt
  - Clay

# africa

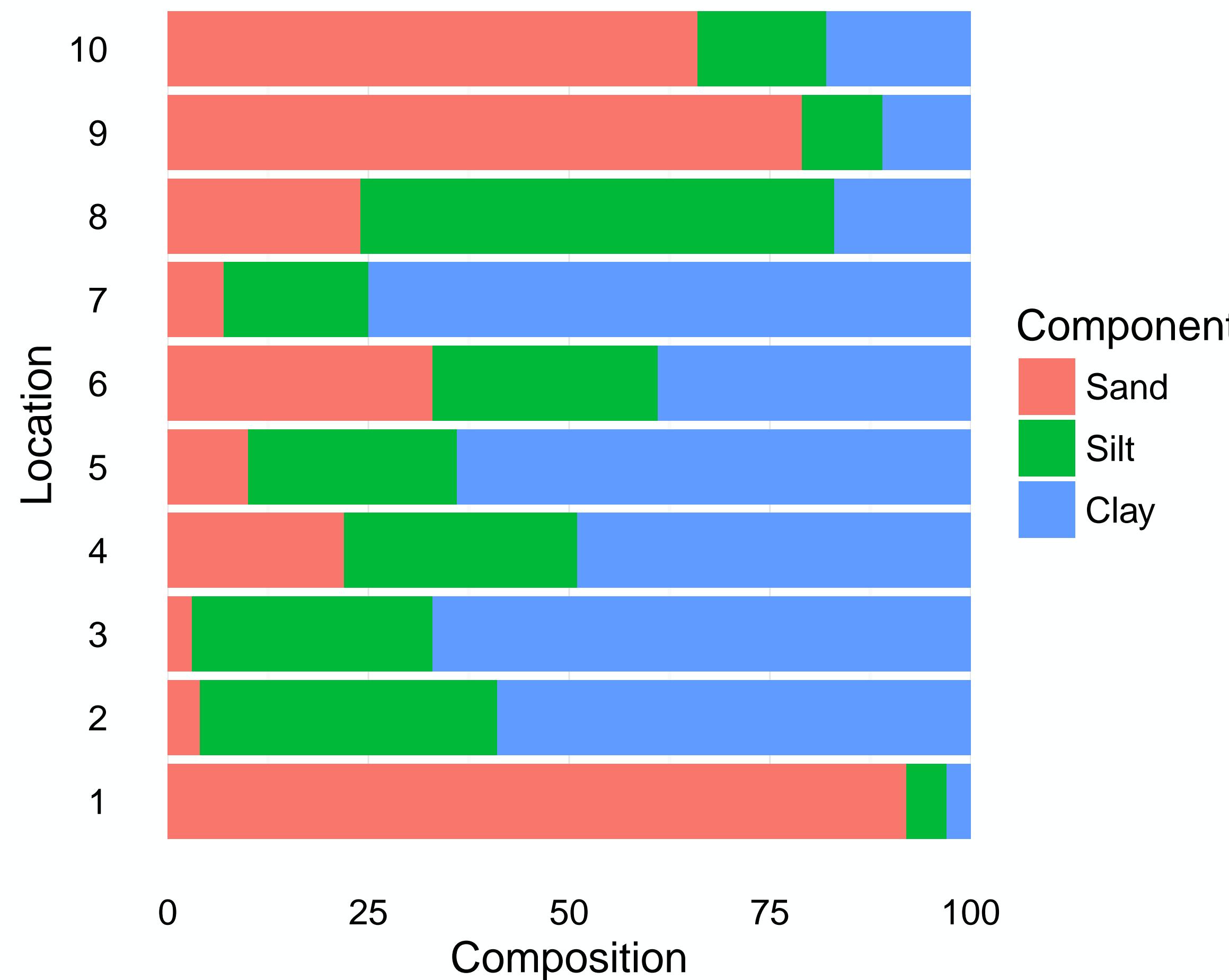
```
> dim(africa)
[1] 40093      3

> head(africa)
  Sand Silt Clay
18100   24   12   64
18103   36   14   50
18113   56   18   26
18115   52   21   27
18142   65    3   32
18158   43   14   43
```

# africa.sample

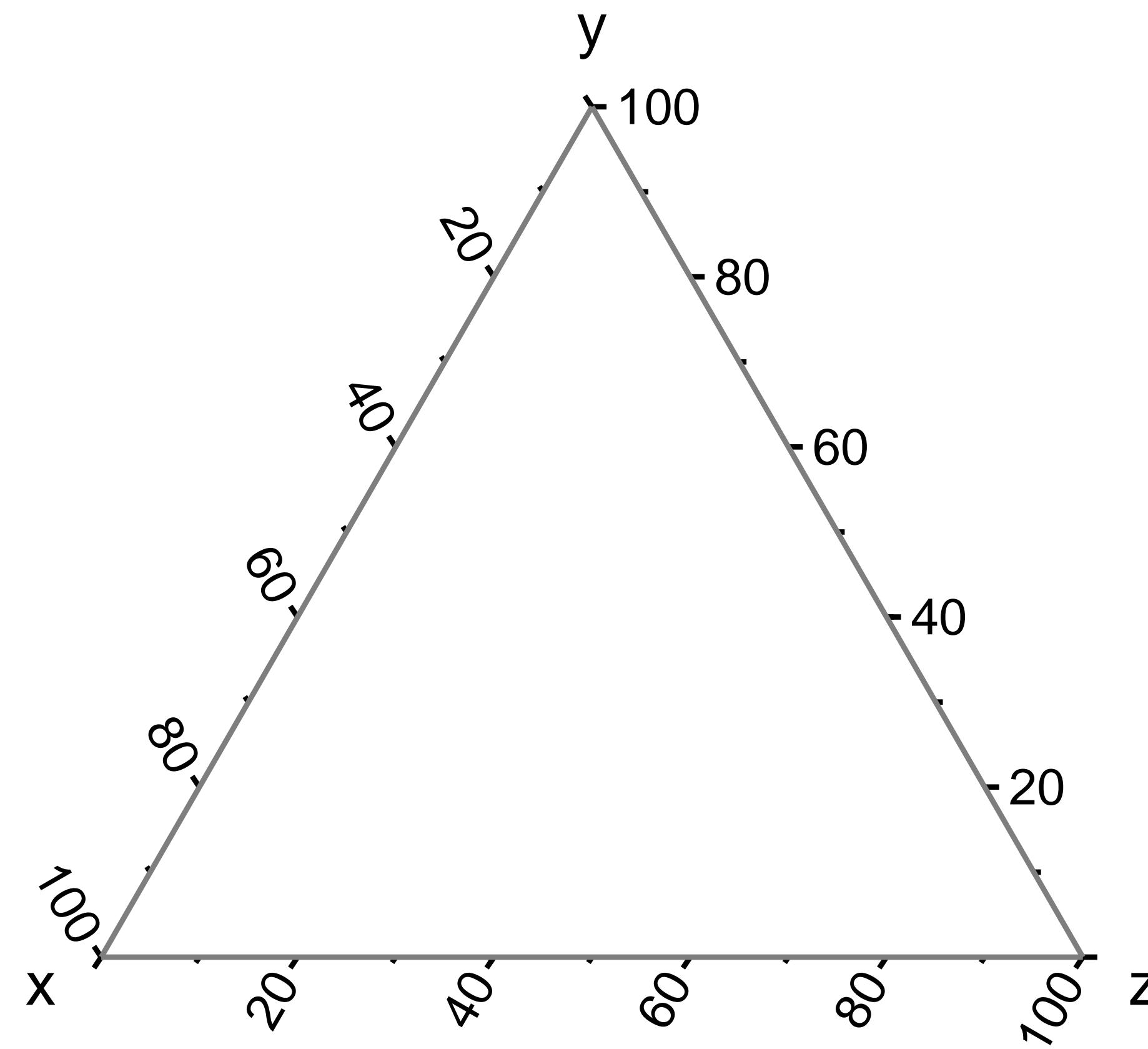
```
> africa.sample
   Sand Silt Clay ID
1 23678    92     5    3  1
2 27995     4    37   59  2
3 35263     3    30   67  3
4 41380    22    29   49  4
5 47814    10    26   64  5
6 62640    33    28   39  6
7 67707     7    18   75  7
8 80337    24    59   17  8
9 84911    79    10   11  9
10 85014   66    16   18 10
```

# Stacked bar plot

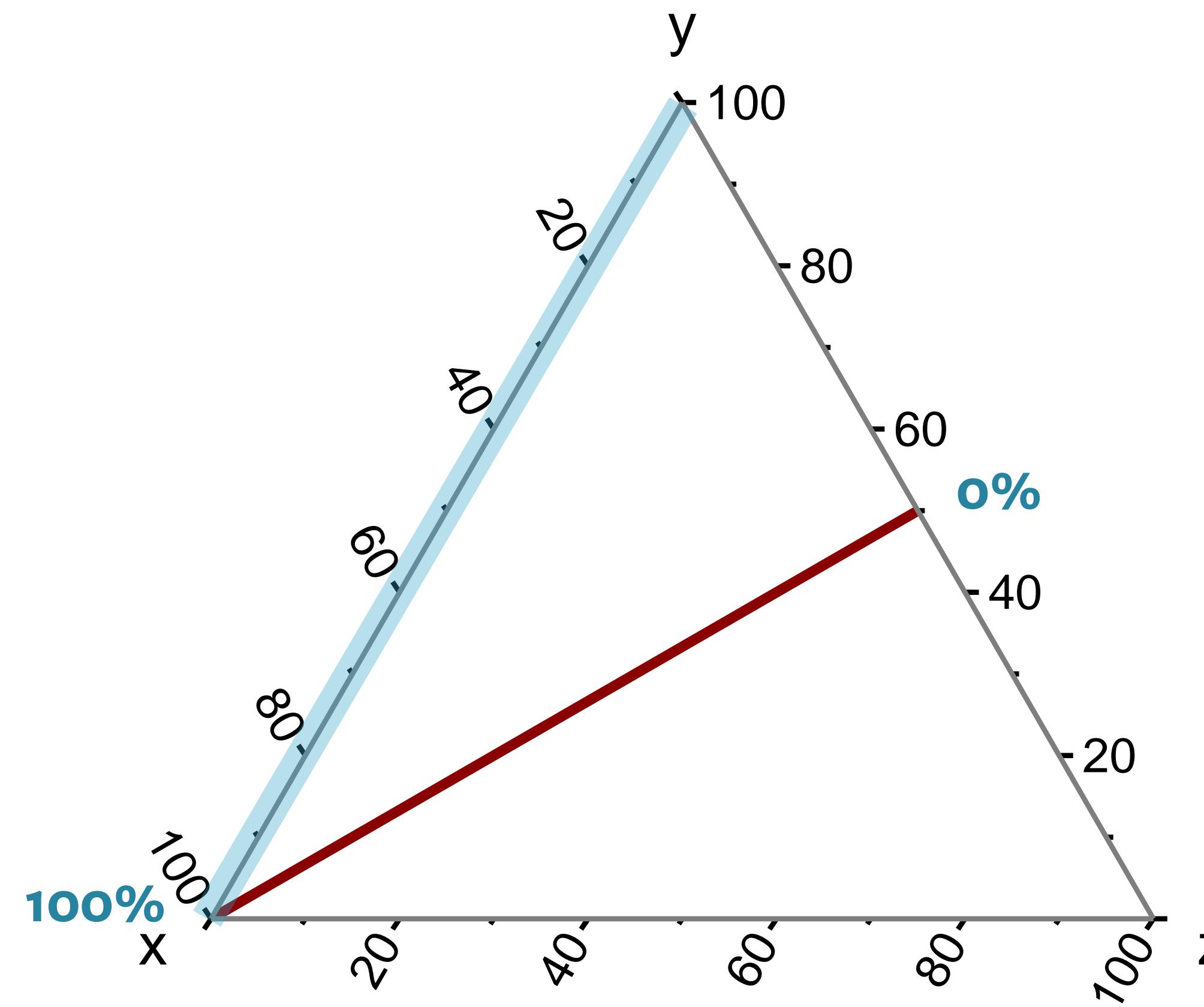


> africa.sample			
	Sand	Silt	Clay
ID			
23678	92	5	3
27995	4	37	59
35263	3	30	67
41380	22	29	49
...			

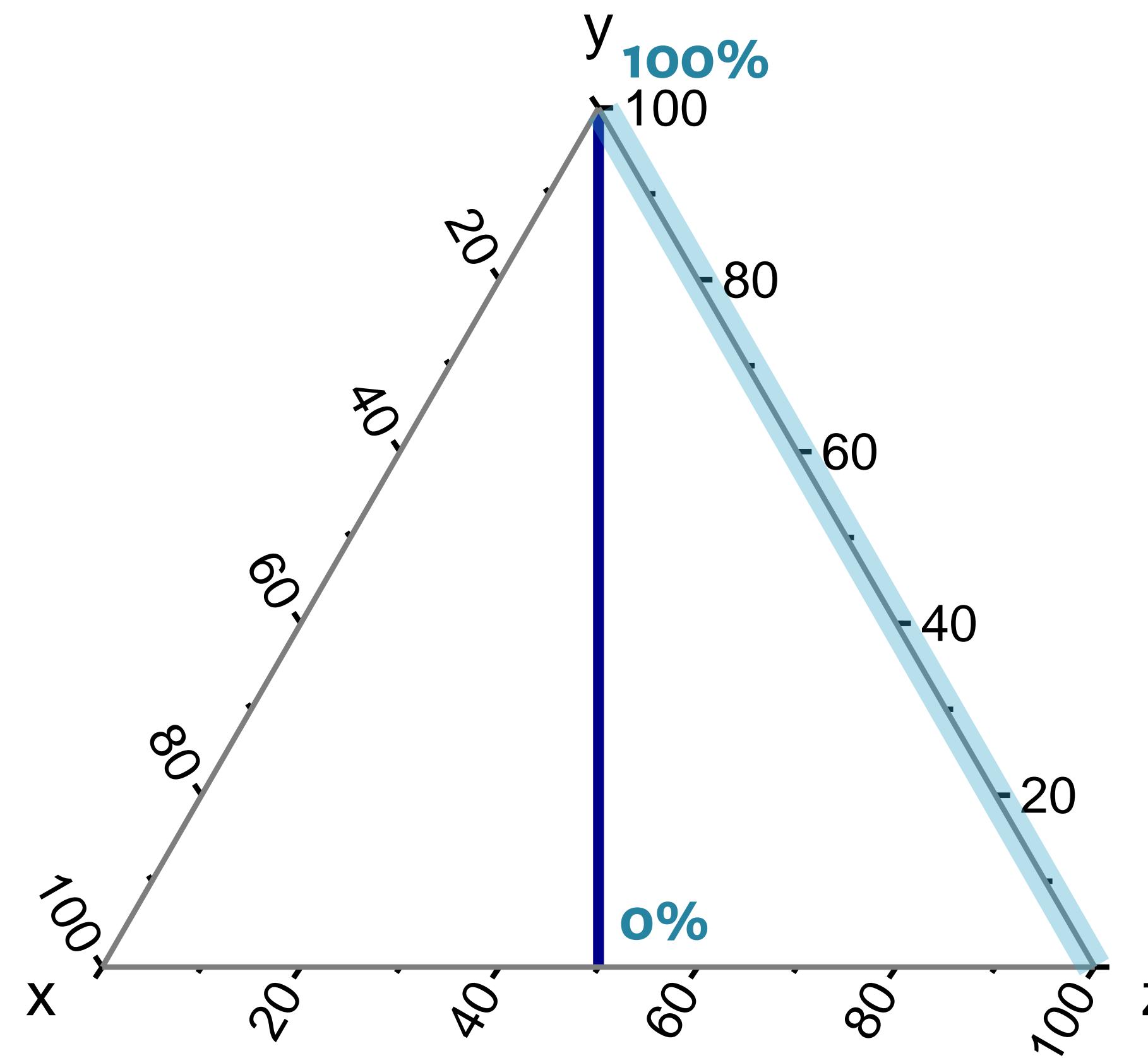
# Ternary plot



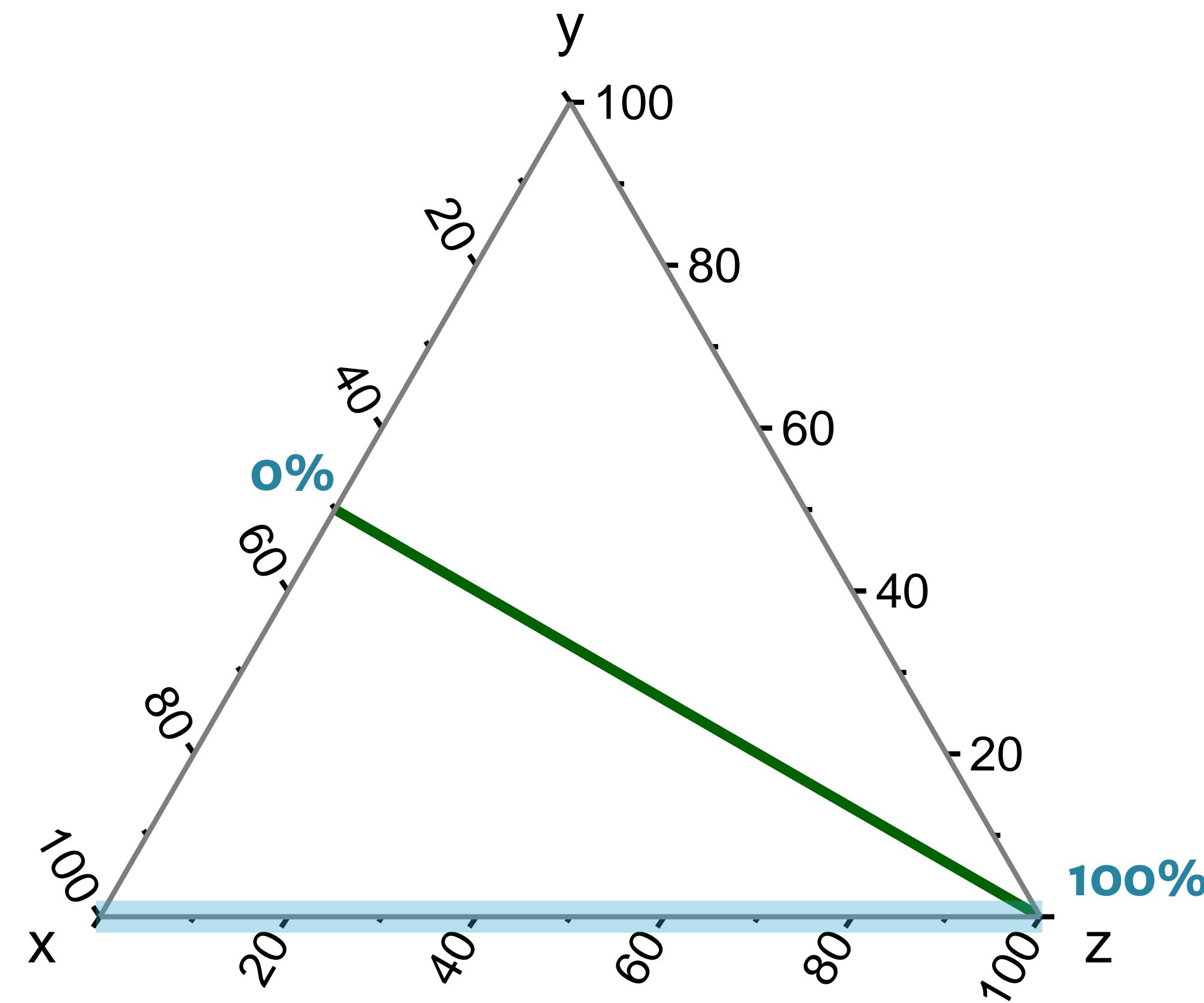
# Ternary plot - X



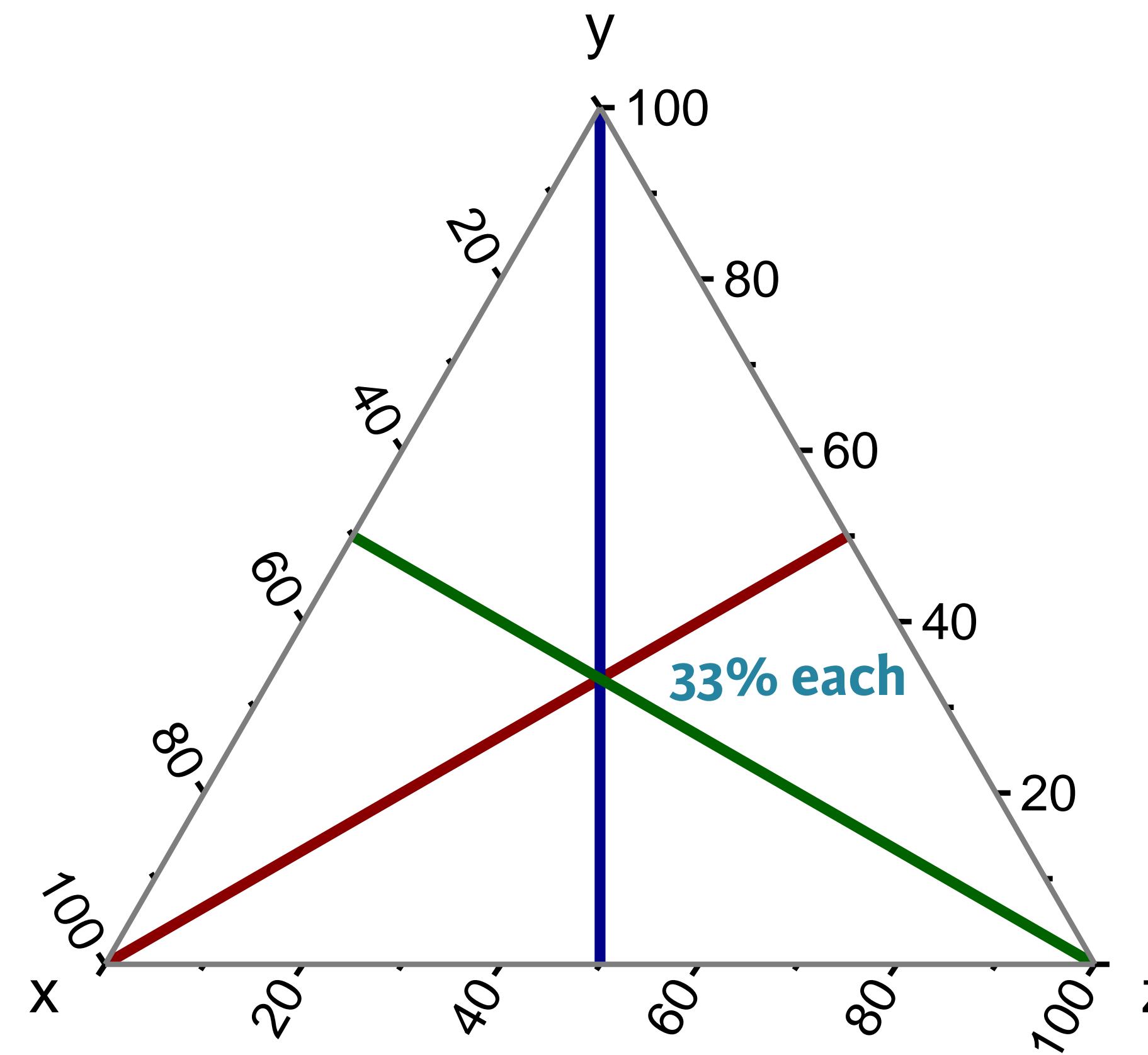
# Ternary plot - Y



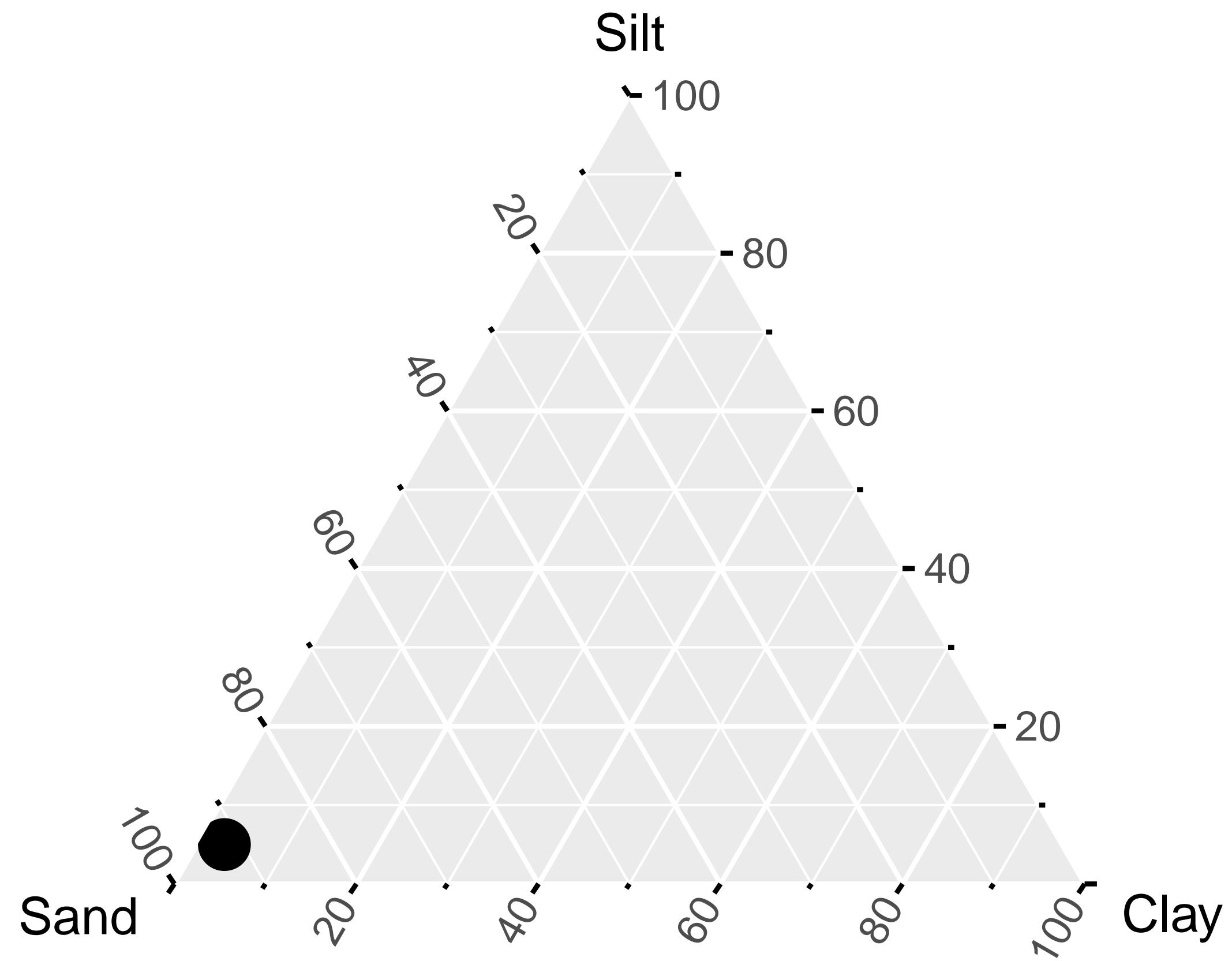
# Ternary plot - Z



# Ternary plot - 3 axes

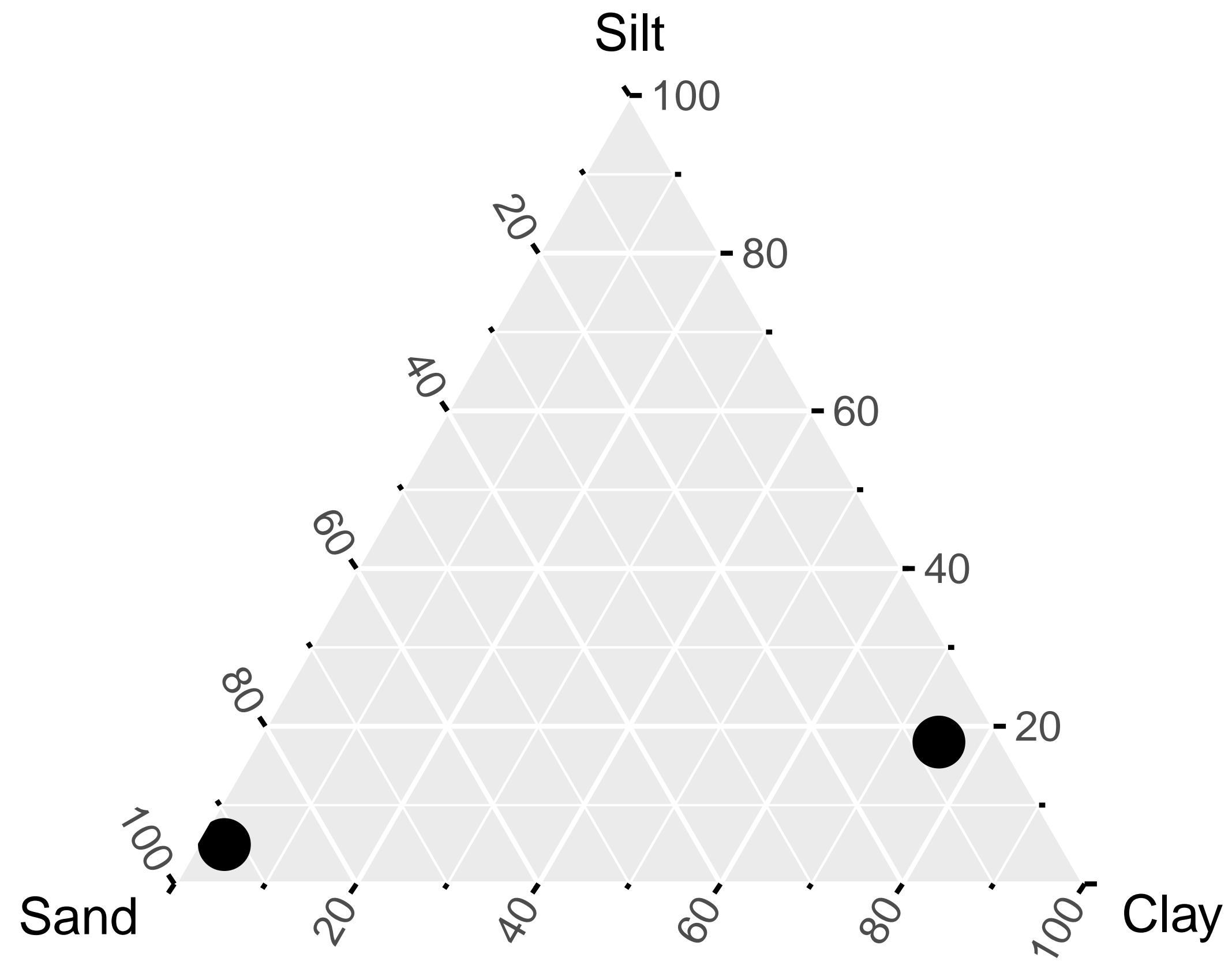


# Location 1



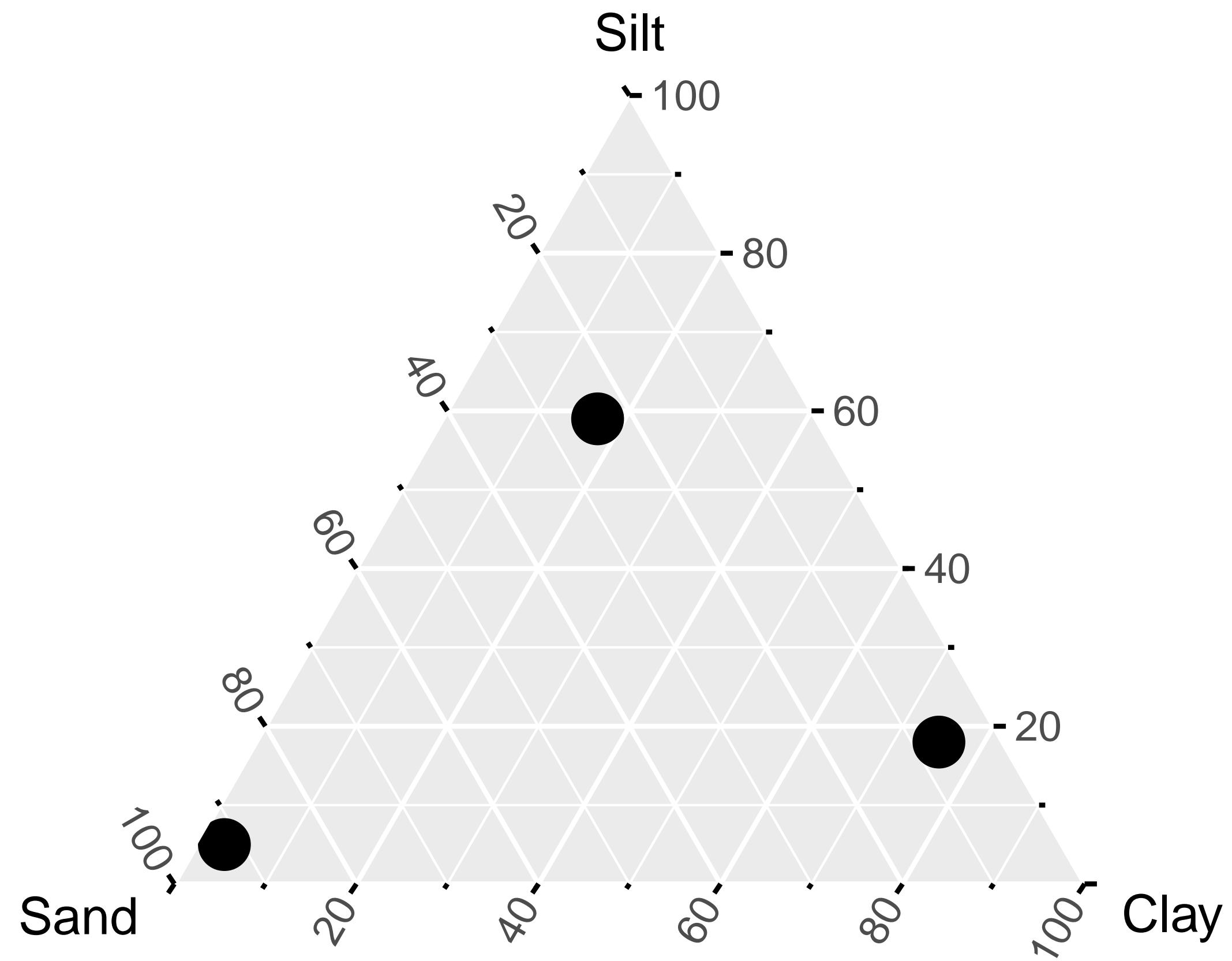
> africa.sample				
	Sand	Silt	Clay	ID
23678	92	5	3	1
27995	4	37	59	2
35263	3	30	67	3
41380	22	29	49	4
...				

# Location 7



```
> africa.sample  
  Sand Silt Clay ID  
...  
62640   33   28   39   6  
67707     7   18   75   7  
80337   24   59   17   8  
...
```

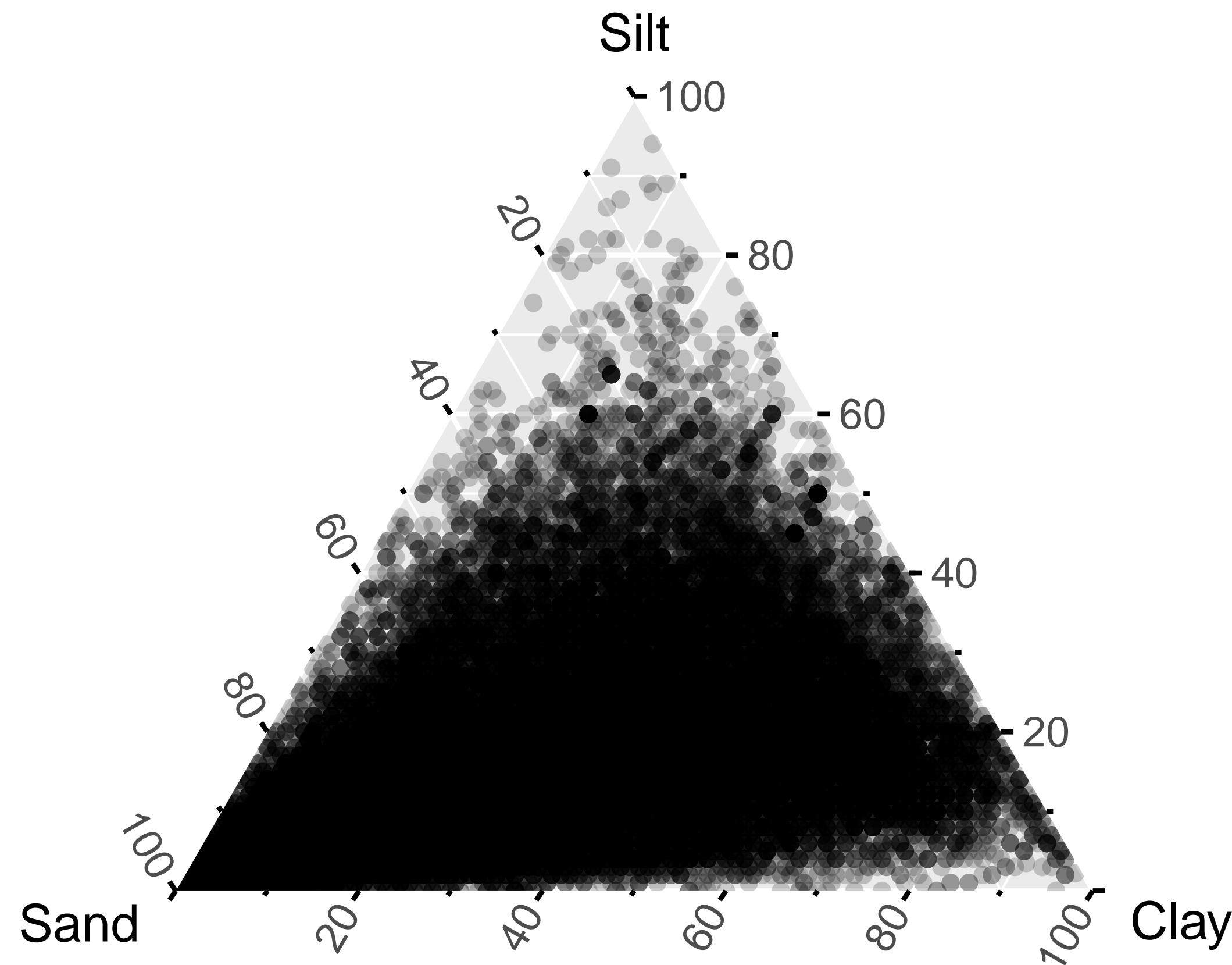
# Location 8



```
> africa.sample  
  Sand Silt Clay ID  
...  
62640   33   28   39   6  
67707    7   18   75   7  
80337   24   59   17   8  
...
```

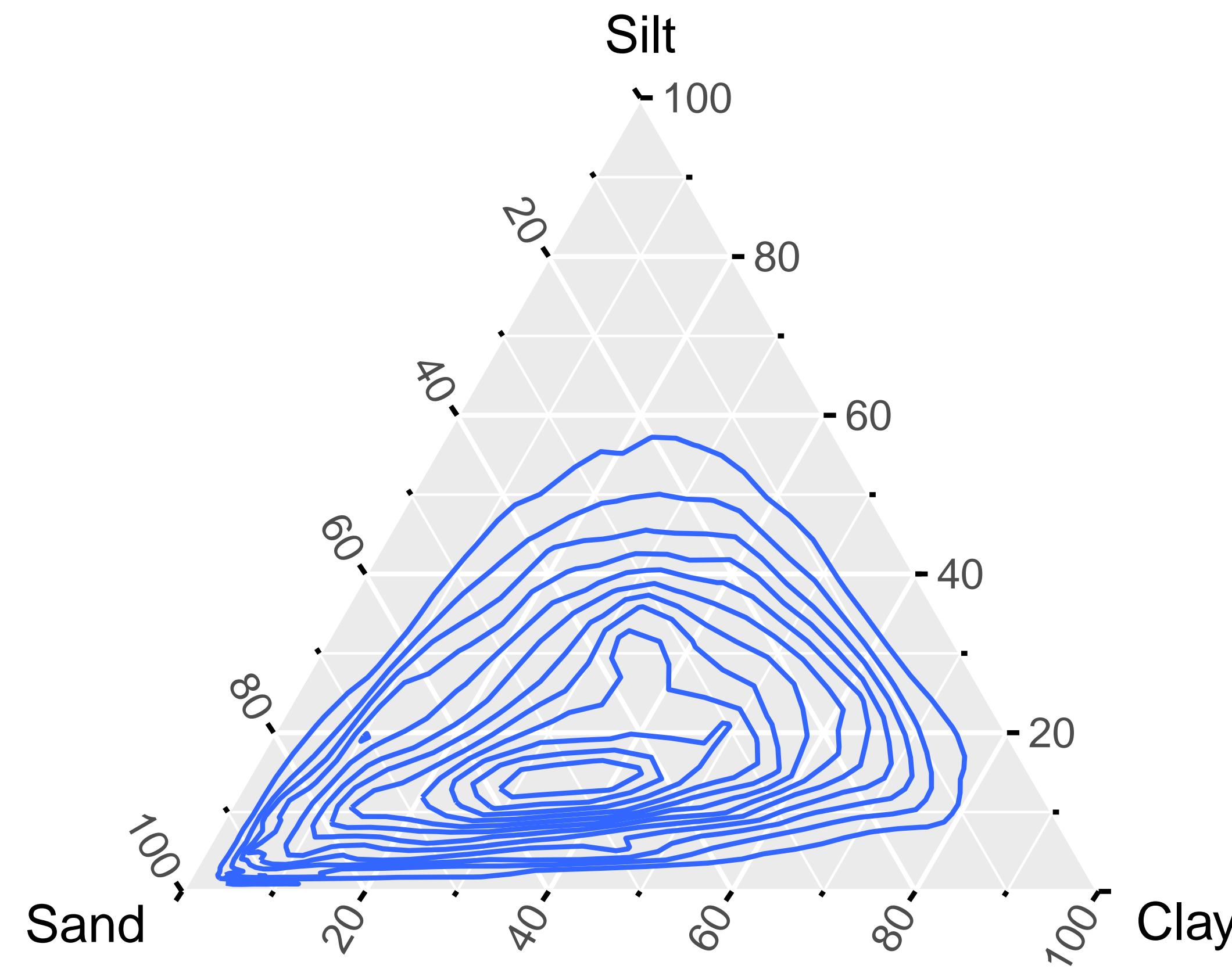
# All data (1)

```
> library(ggtern)
> ggtern(africa, aes(Sand, Silt, Clay)) +
  geom_point(alpha = 0.2, shape = 16)
```

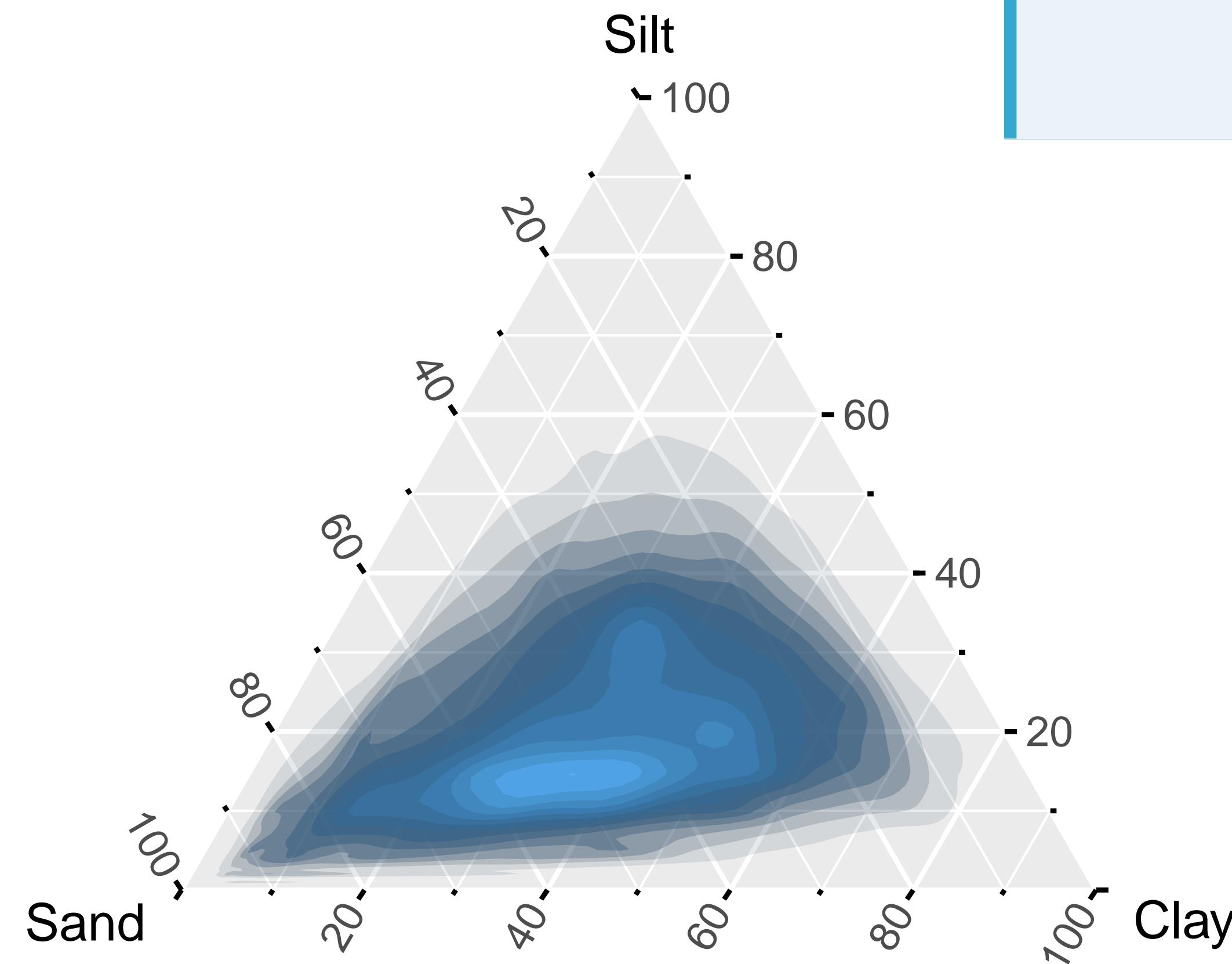


# All data (2)

```
> library(ggtern)
> ggtern(africa, aes(Sand, Silt, Clay)) +
  geom_density_tern()
```



# All data (3)



```
> library(ggtern)
> ggtern(africa, aes(Sand, Silt, Clay)) +
  stat_density_tern(geom = "polygon",
    n = 200,
    aes(fill = ..level..,
        alpha = ..level..))
```



DATA VISUALIZATION WITH GGPLOT2

**Let's practice!**



DATA VISUALIZATION WITH GGPLOT2

# Network Plots

# Blood types

- Four main types
  - A, B, AB, O
- Rho negative or positive
- 8 different types
- Not all compatible
- Which blood type can be used as a donor?

# blood\_donors

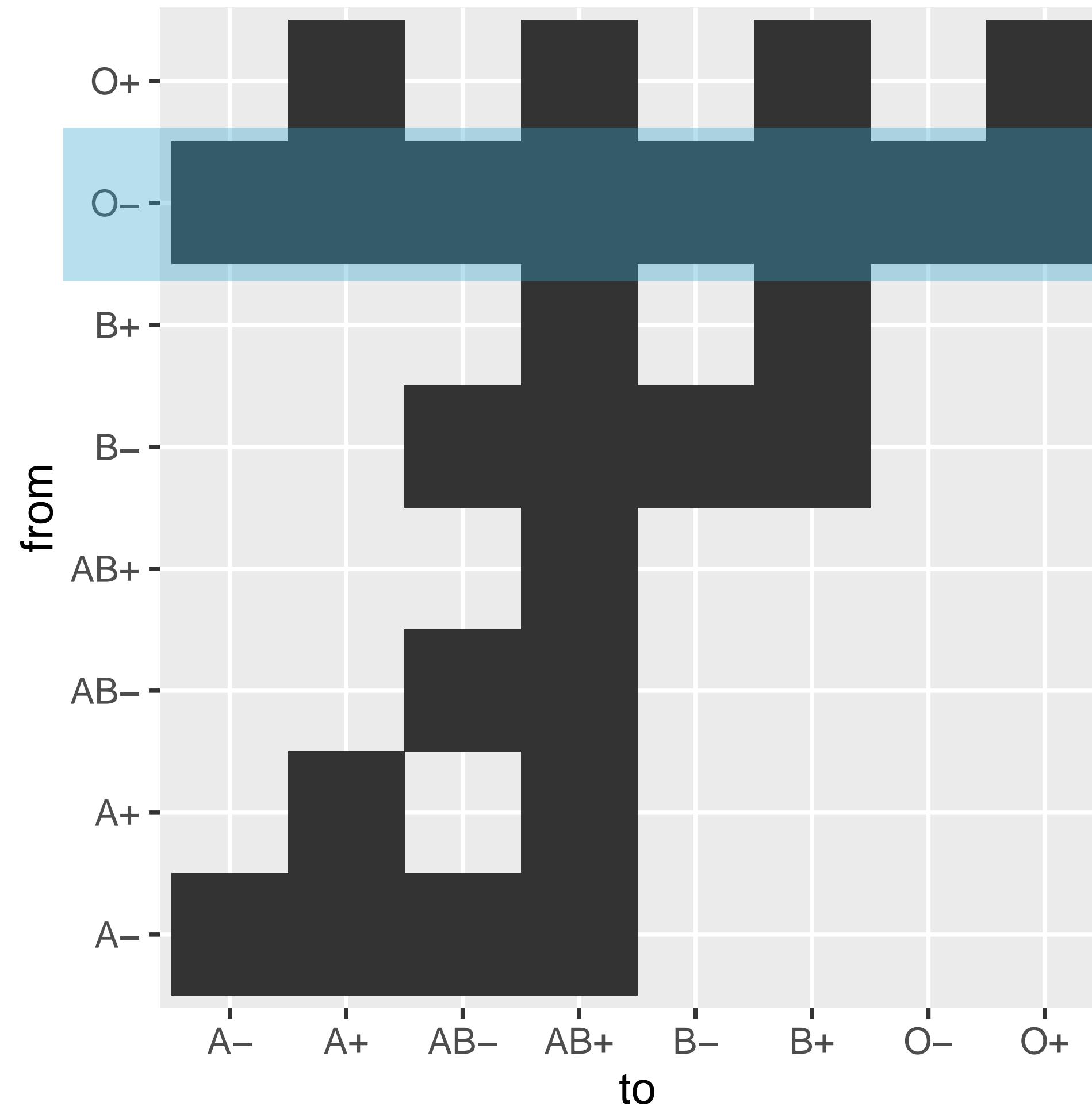
```
> dim(blood_donors)
[1] 27  2

> head(blood_donors)
  from   to
1  AB-  AB+
2  AB-  AB-
3  AB+  AB+
4  A-   AB+
5  A-   AB-
6  A-   A+
```

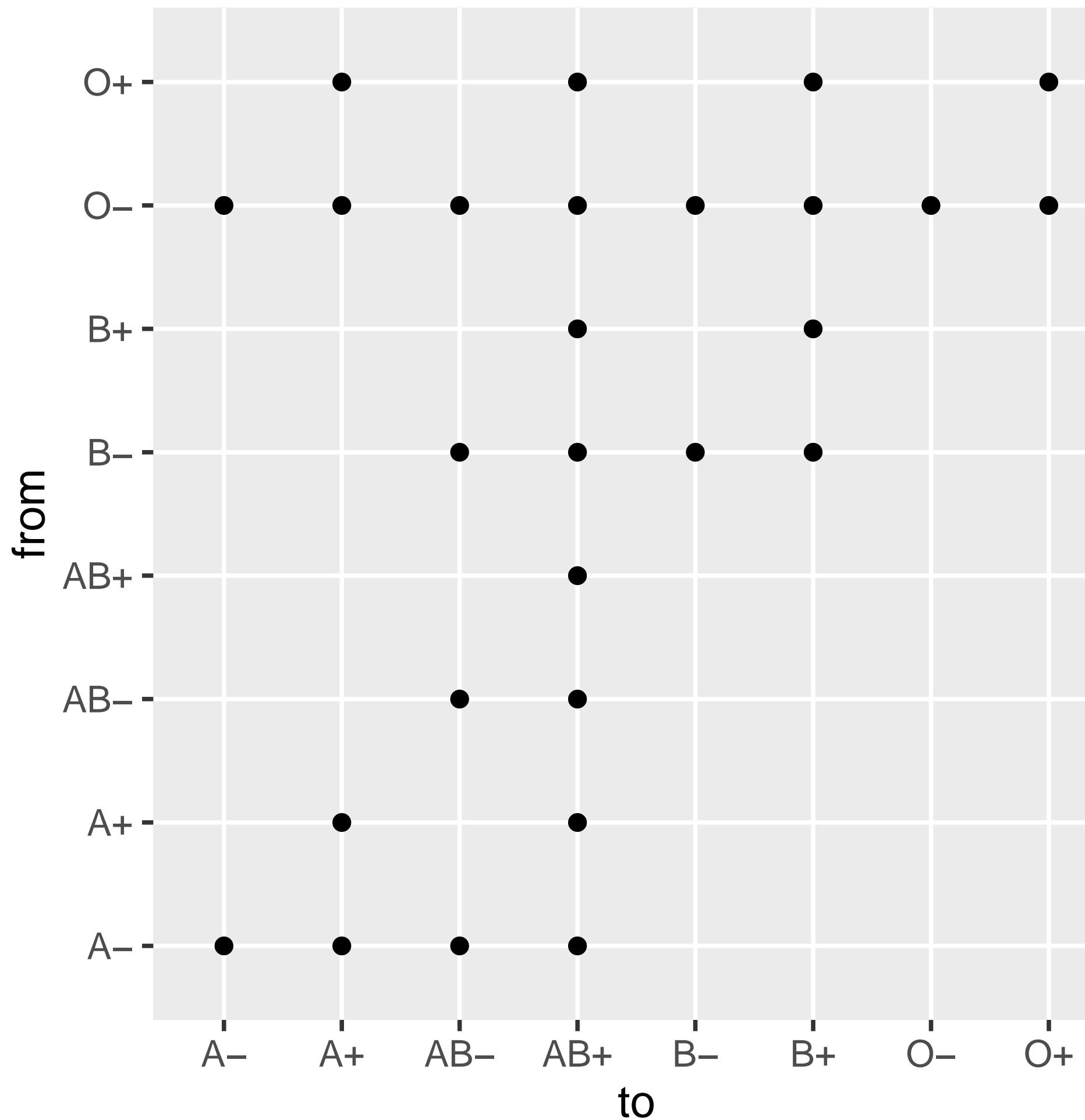
```
> levels(blood_donors$from)
[1] "A-"  "A+"  "AB-" "AB+" "B-"  "B+"  "O-"  "O+"
```

# Heatmap



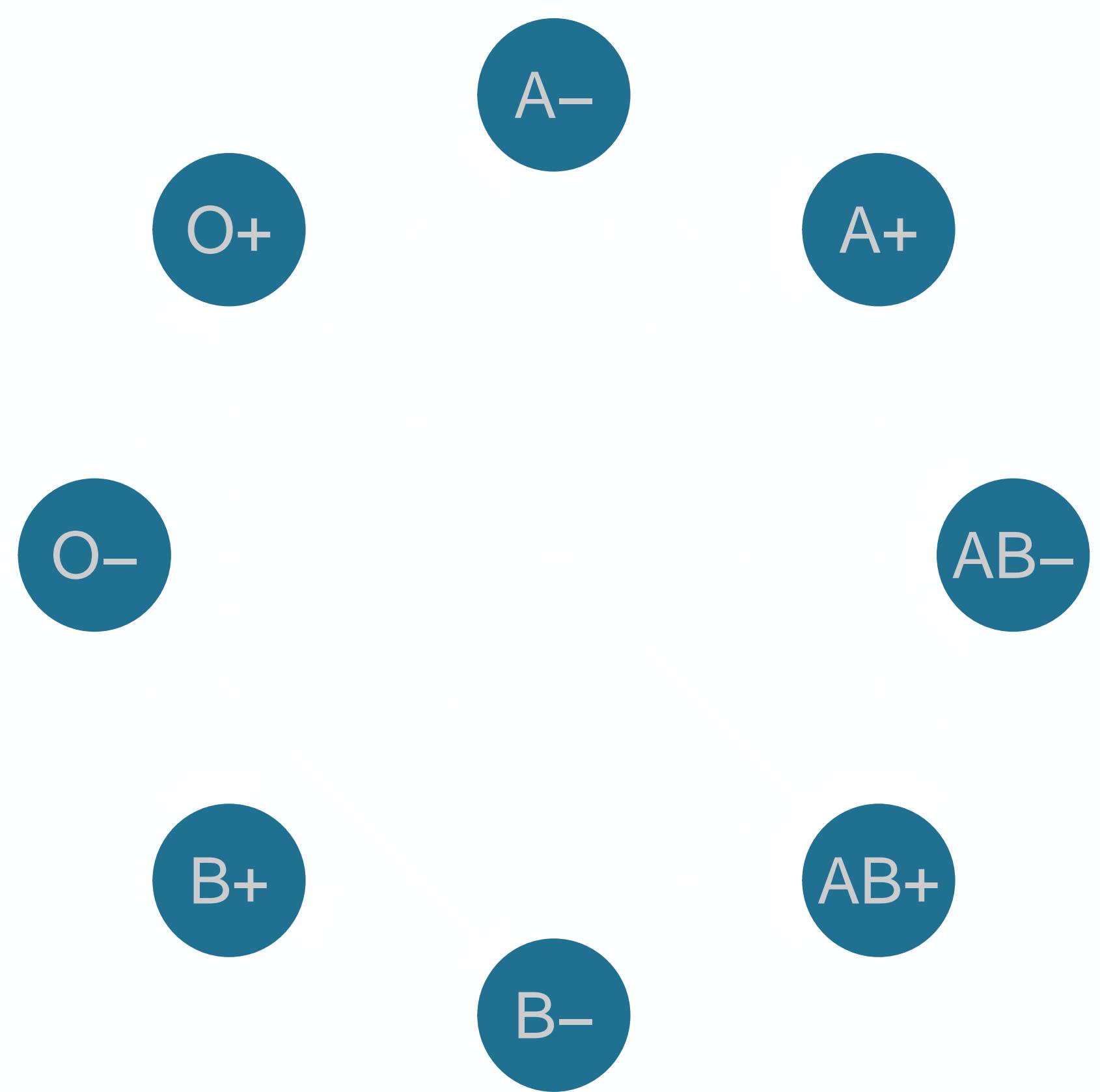
```
> ggplot(Blood_donors, aes(x = to, y = from)) +  
  geom_tile()
```

# Scatter plot

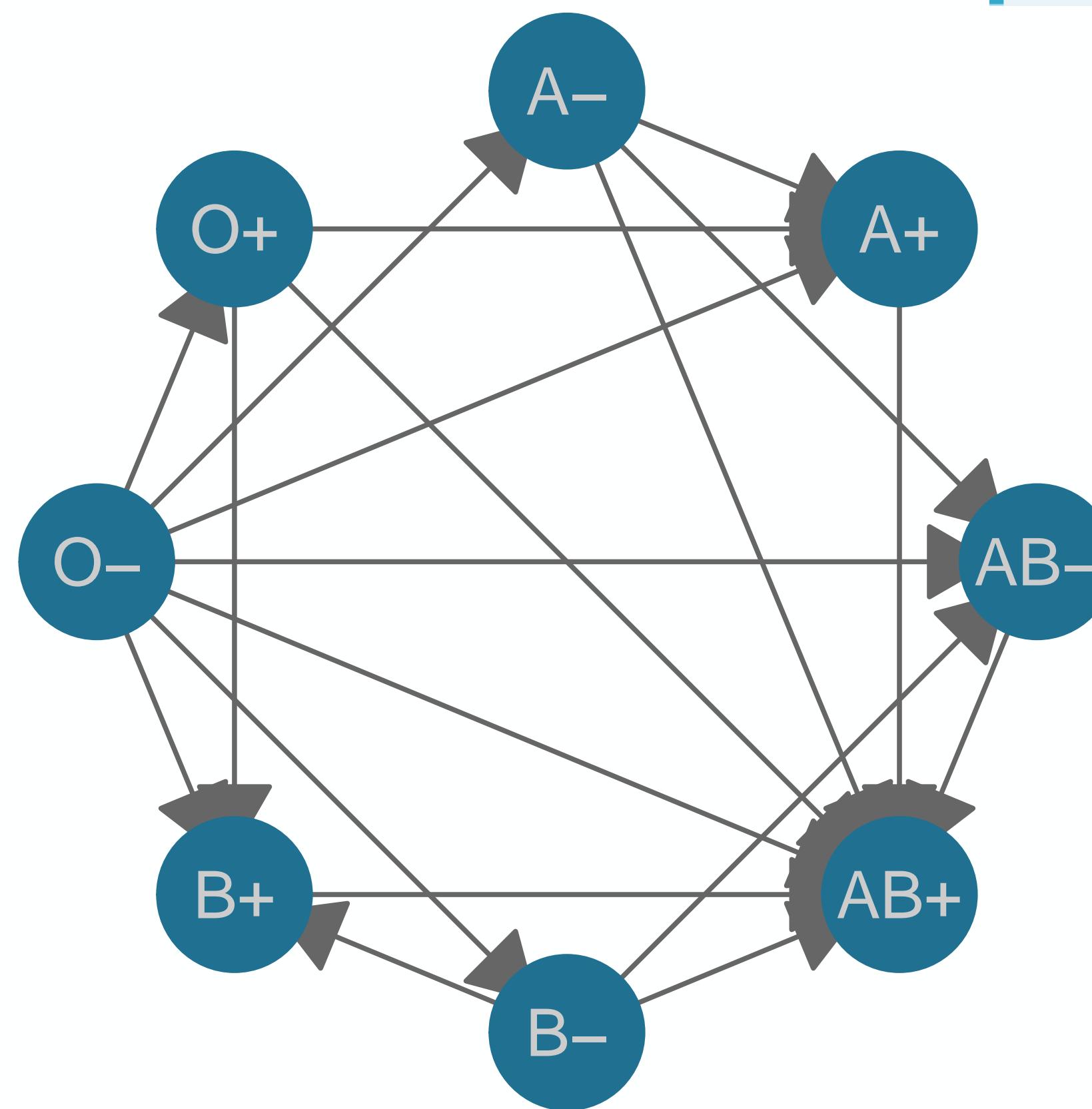


```
> ggplot(Blood_donors, aes(x = to, y = from)) +  
  geom_point()
```

# Network

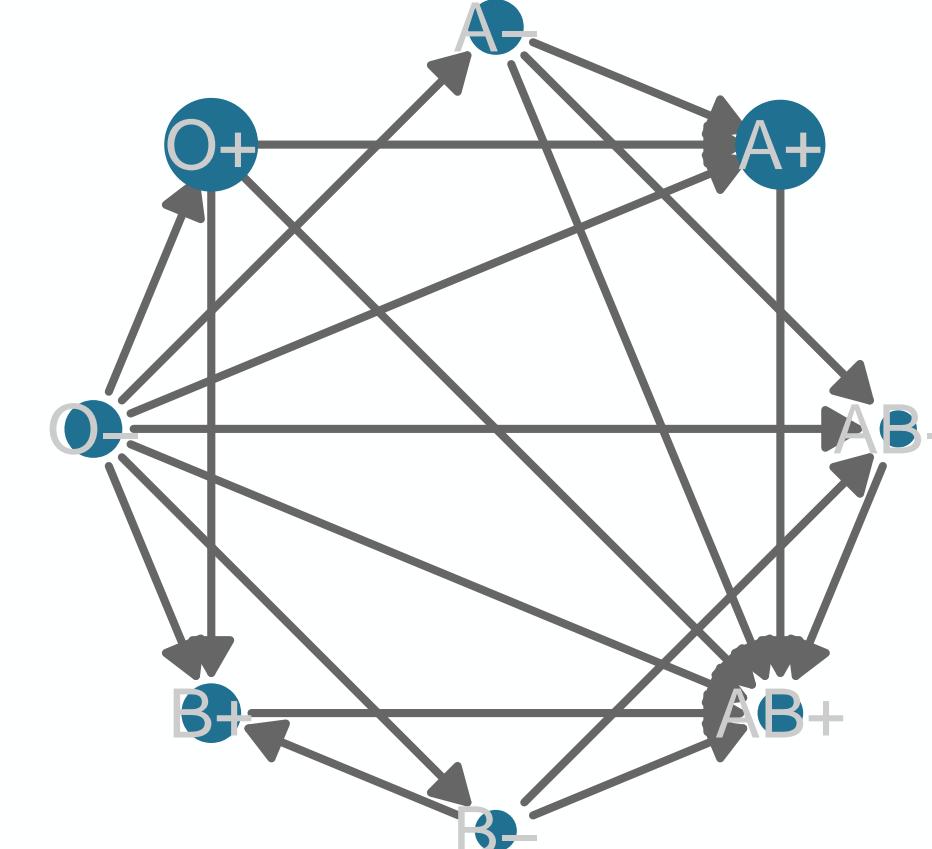


# Network

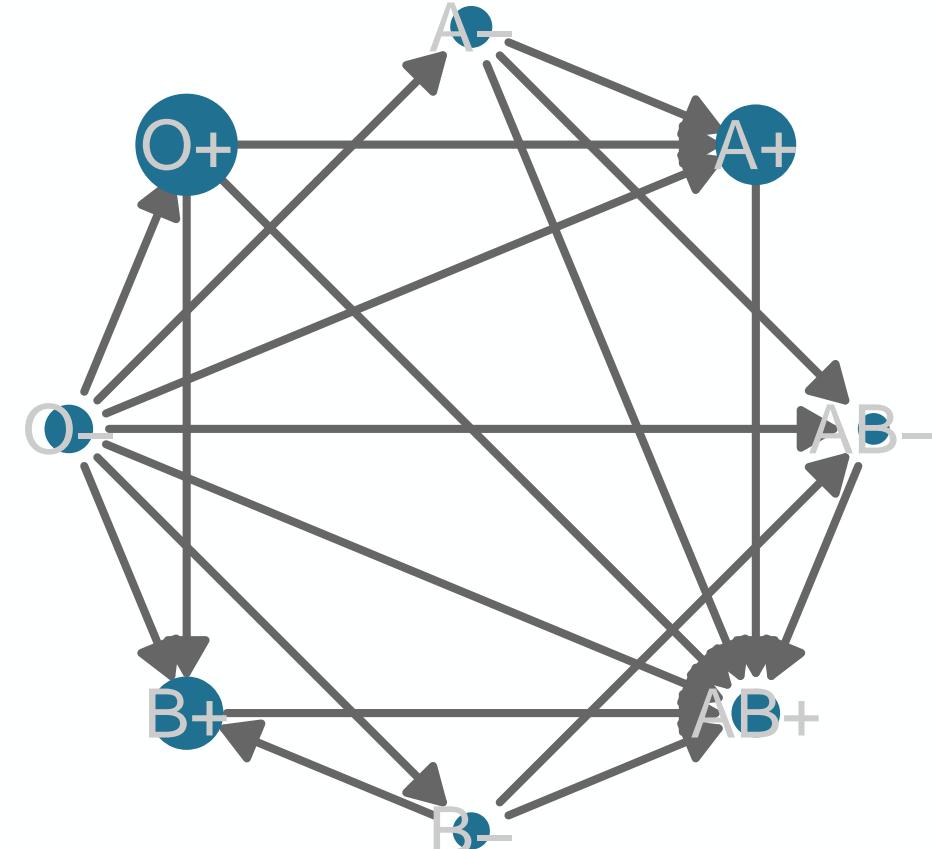


```
> library(geomnet)
> ggplot(blood_donors, aes(from_id = from, to_id = to)) +
  geom_net(layout = "circle",
    label = TRUE, directed = TRUE)
```

Caucasians



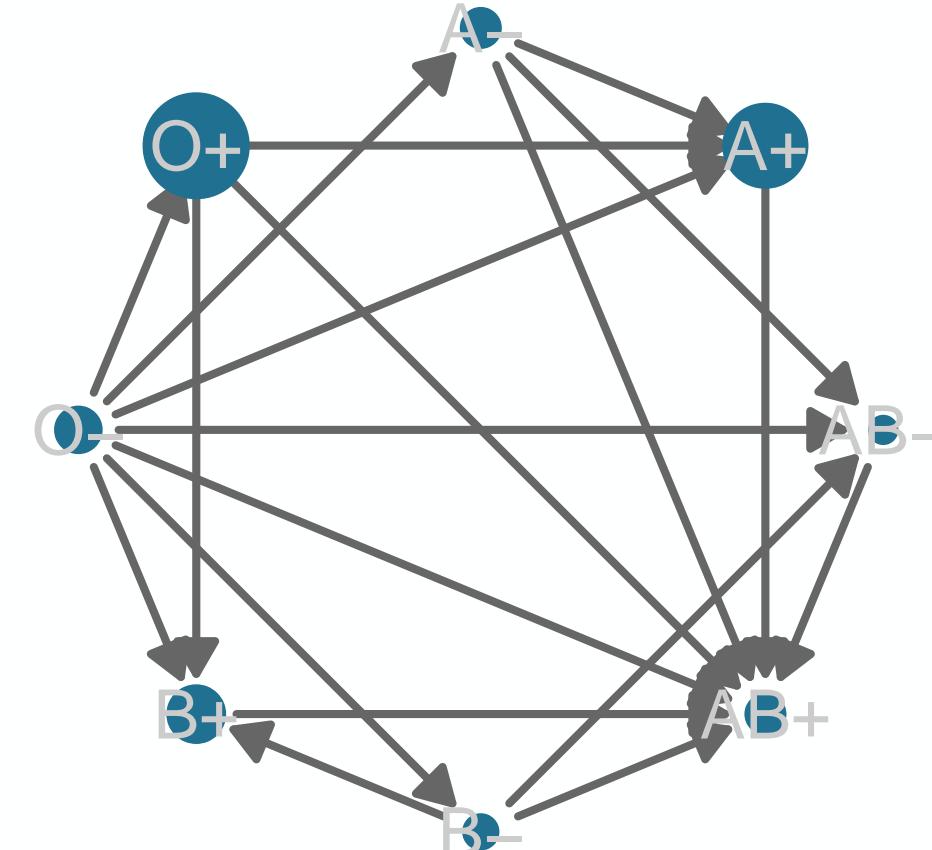
African.American



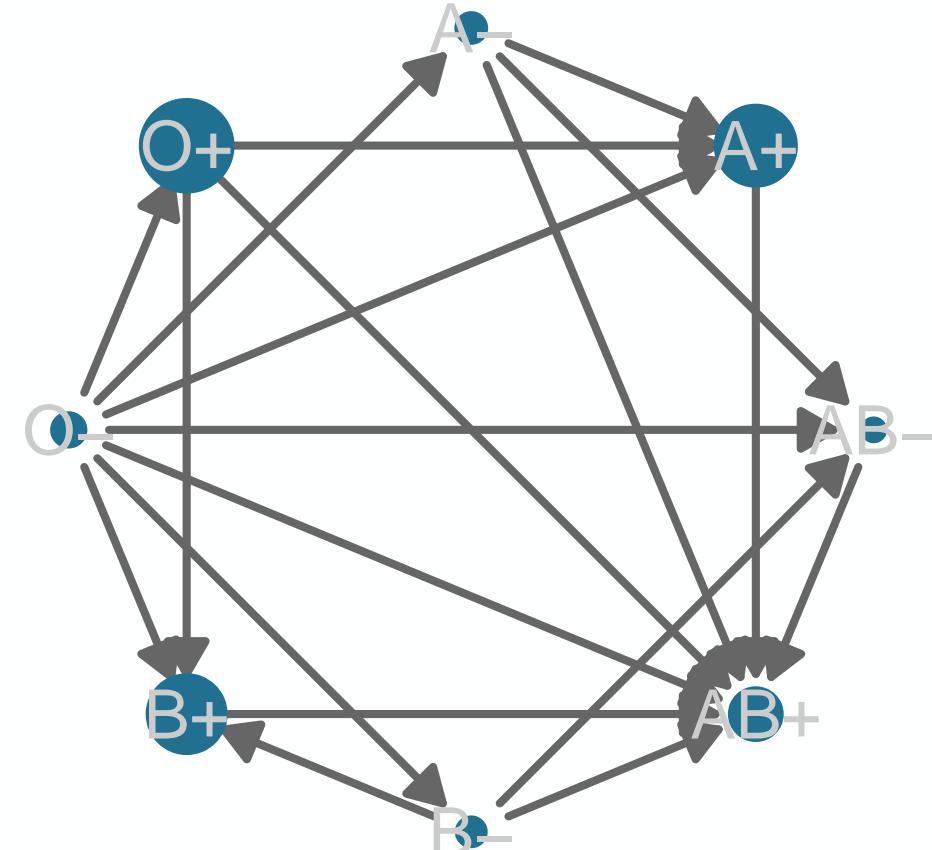
## Predominance

- 10
- 20
- 30
- 40
- 50

Hispanic



Asian





DATA VISUALIZATION WITH GGPLOT2

**Let's practice!**



DATA VISUALIZATION WITH GGPLOT2

# Diagnostic Plots

# Diagnostic Plots

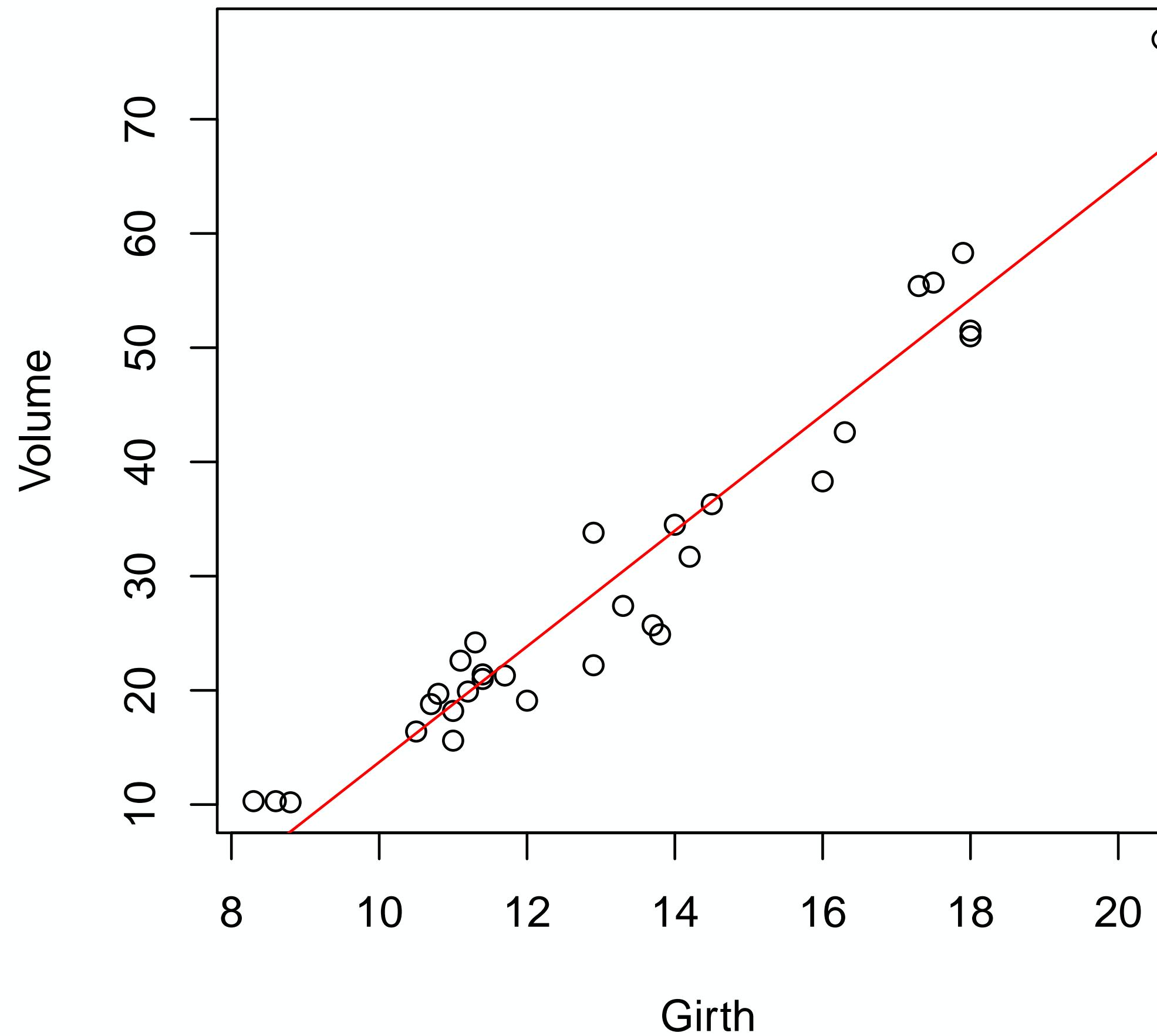
- Quantitative data
- Ordinary least squares model
- Assess how the model fits the data

# trees

```
> dim(trees)
[1] 31  3

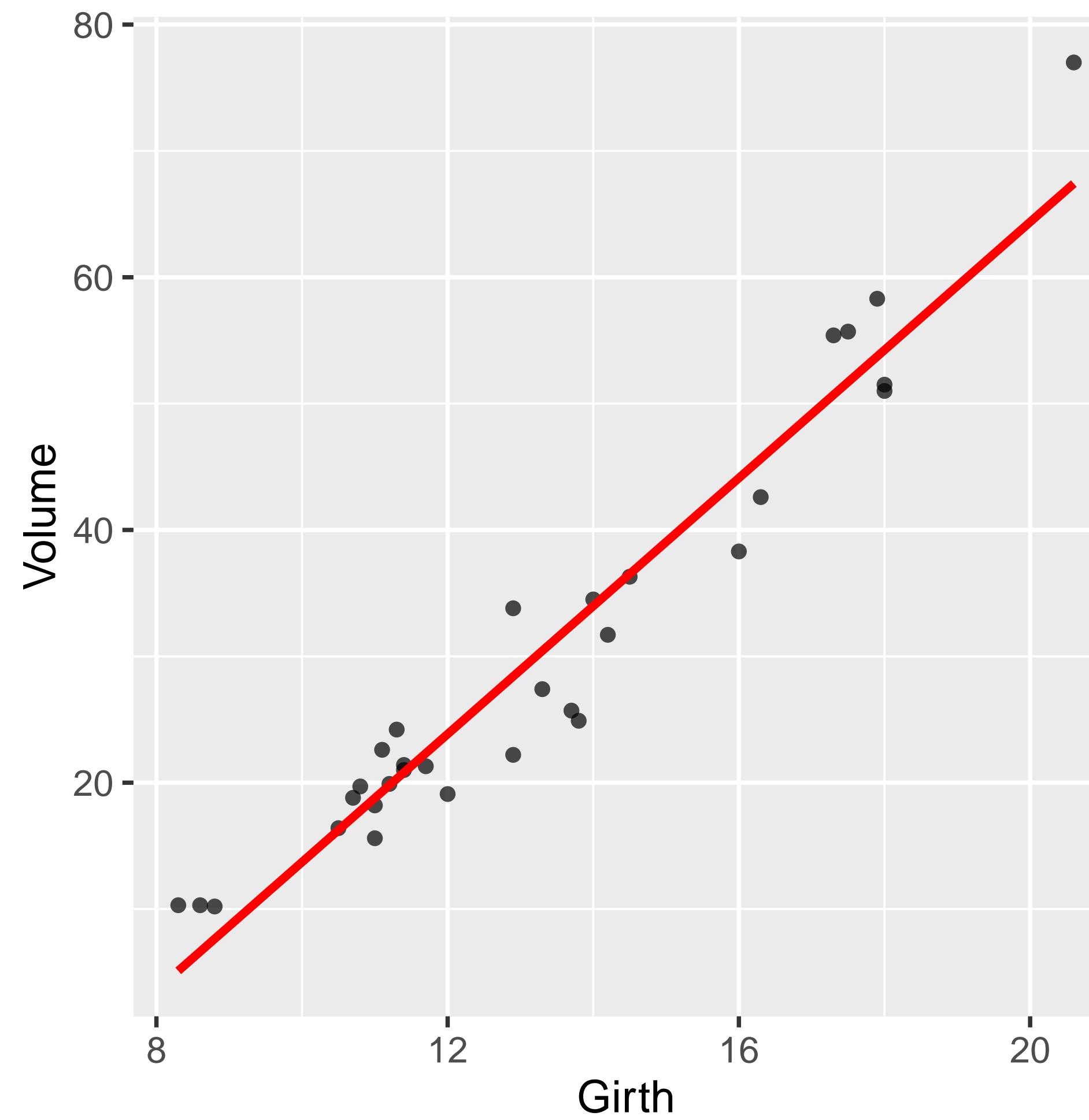
> head(trees)
  Girth Height Volume
1   8.3     70   10.3
2   8.6     65   10.3
3   8.8     63   10.2
4  10.5     72   16.4
5  10.7     81   18.8
6  10.8     83   19.7
```

# Linear model



```
> res <- lm(Volume ~ Girth, data = trees)
> plot(Volume ~ Girth, data = trees)
> abline(res, col = "red")
```

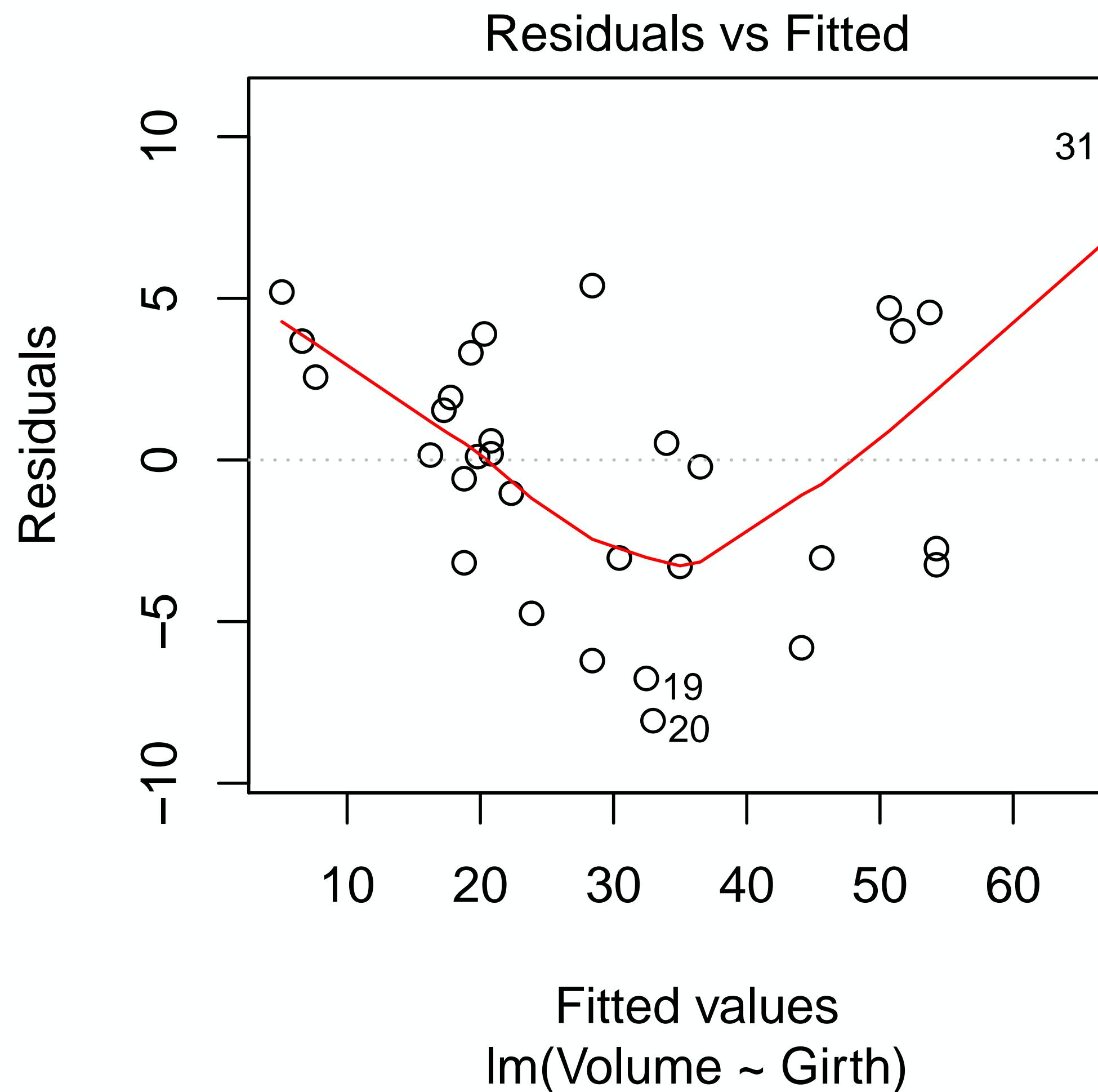
# Linear model



```
> ggplot(trees, aes(Girth, Volume)) +  
  geom_point(shape = 16, alpha = 0.7) +  
  stat_smooth(method = "lm", se = F, col = "red")
```

# plot model (1)

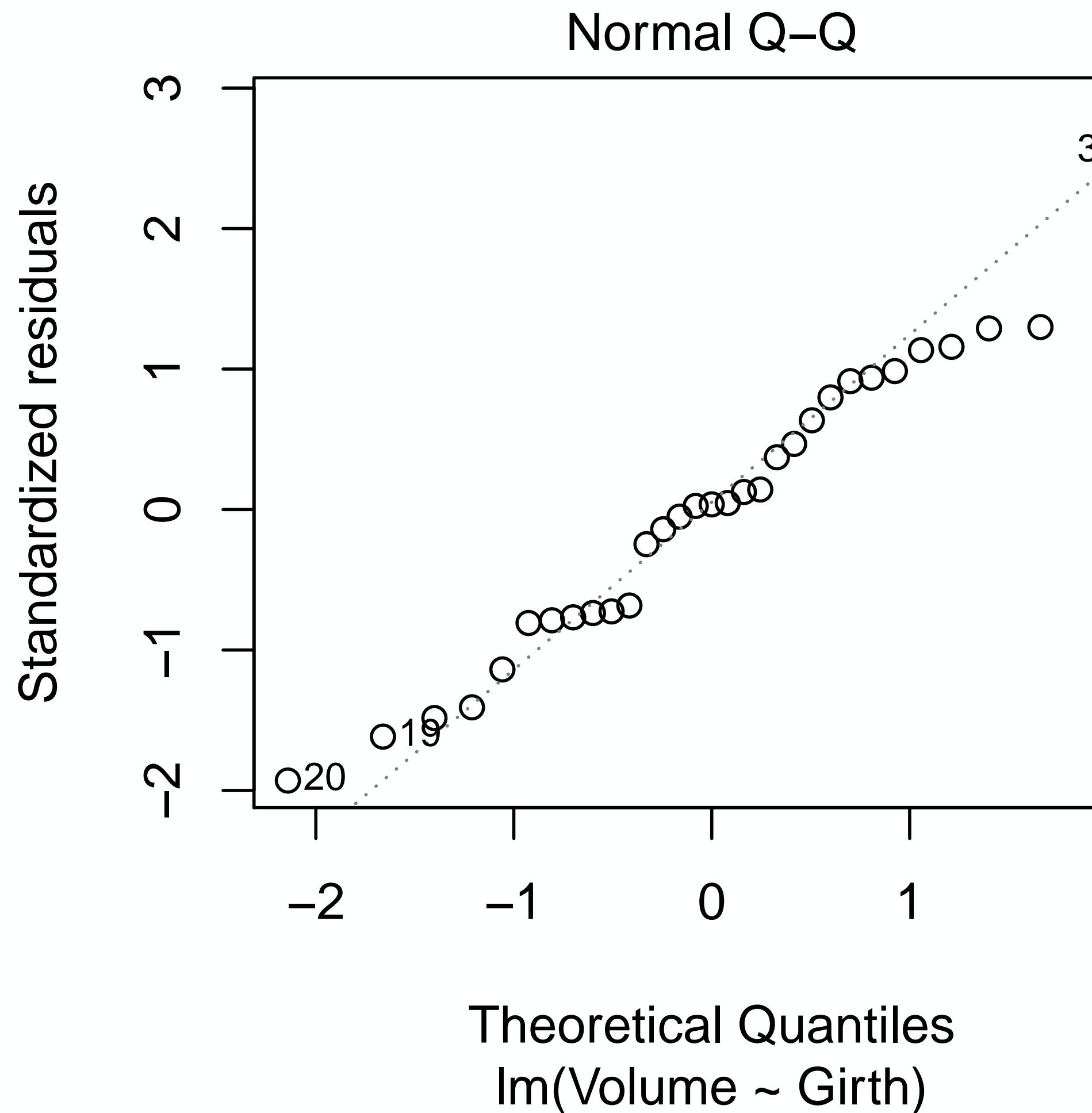
```
> res <- lm(Volume ~ Girth, data = trees)  
> plot(res)
```



Ideally no clear trend

# plot model (2)

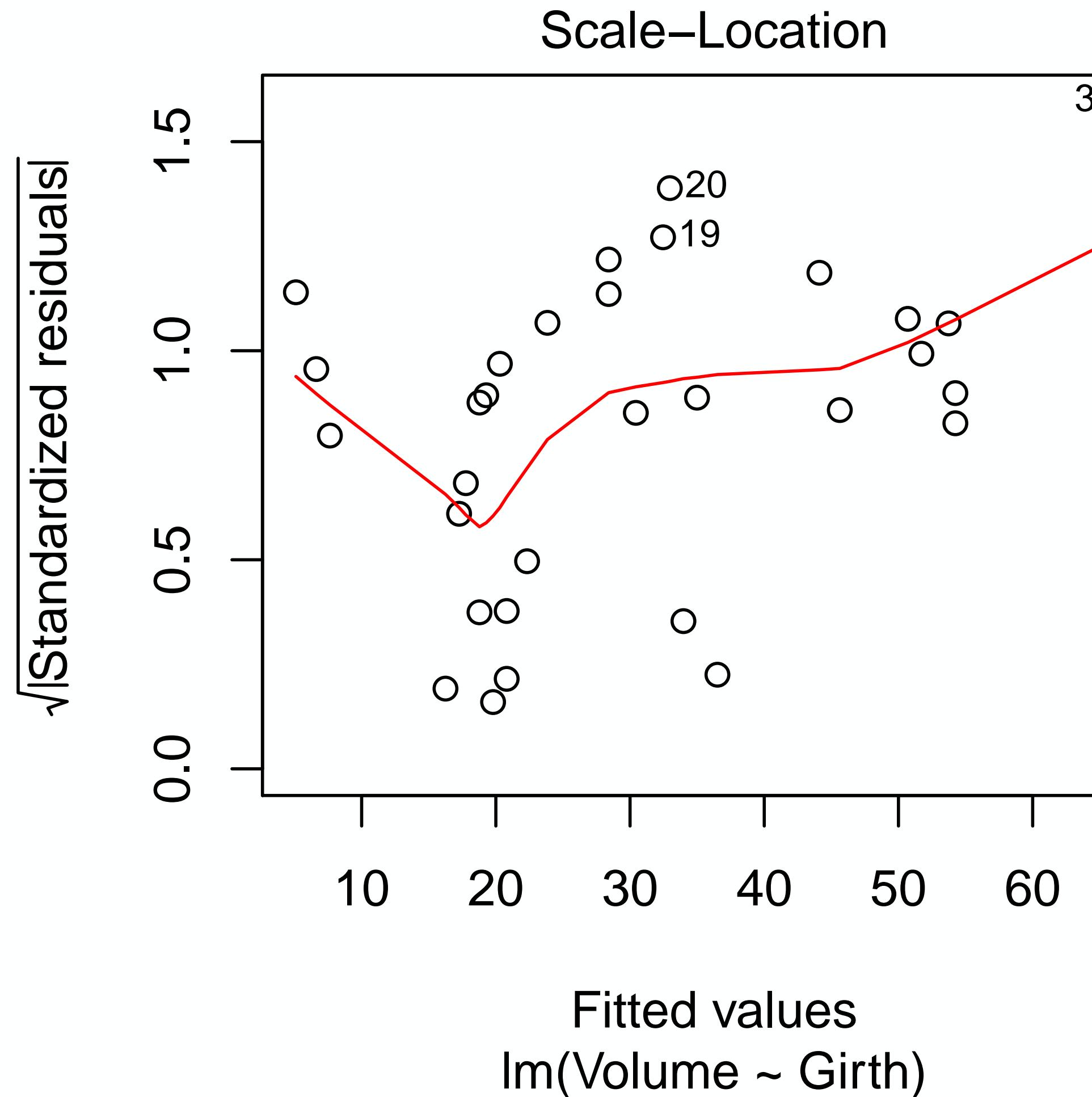
```
> res <- lm(Volume ~ Girth, data = trees)  
> plot(res)
```



Are residuals normally distributed?

# plot model (3)

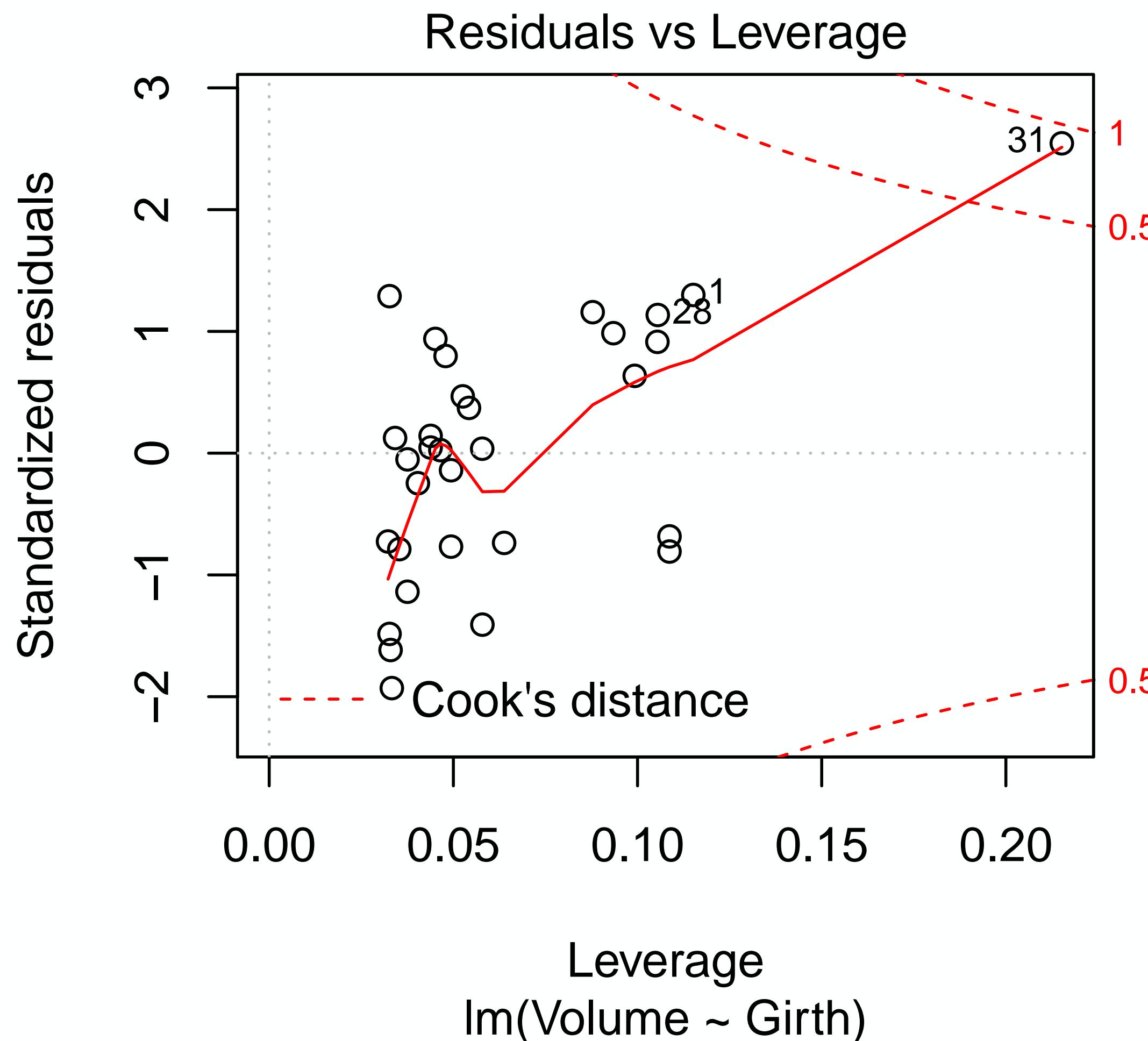
```
> res <- lm(Volume ~ Girth, data = trees)  
> plot(res)
```

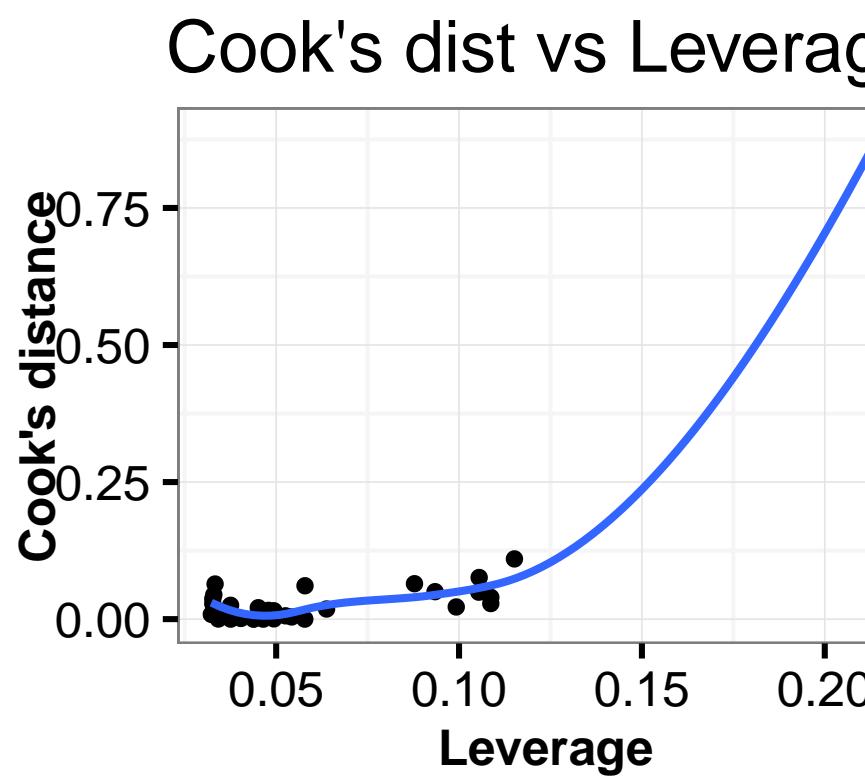
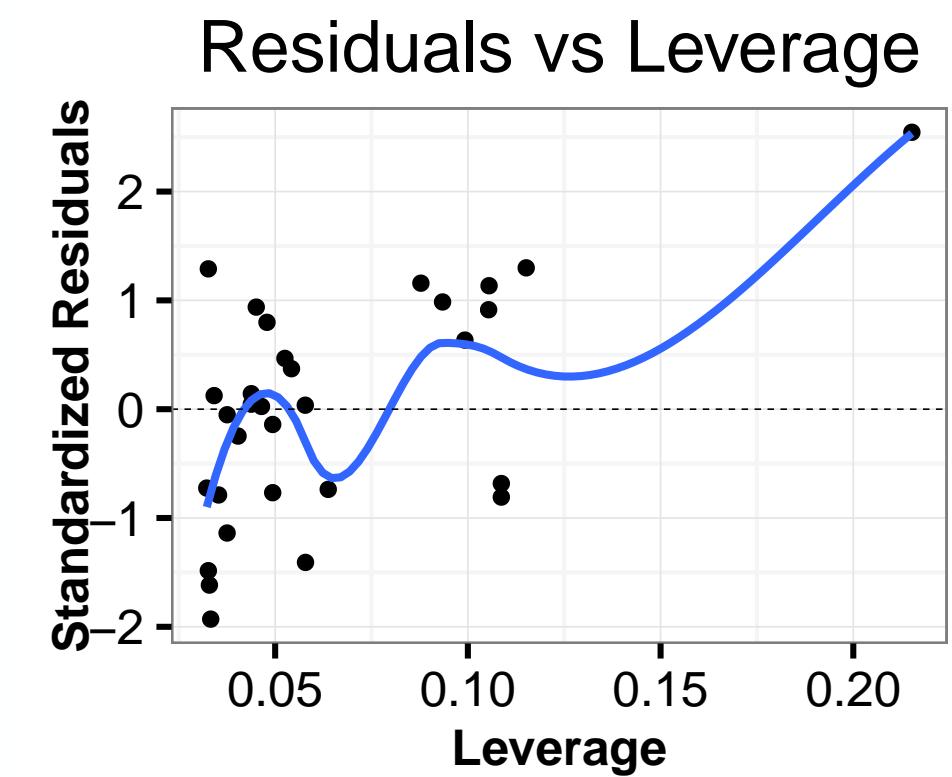
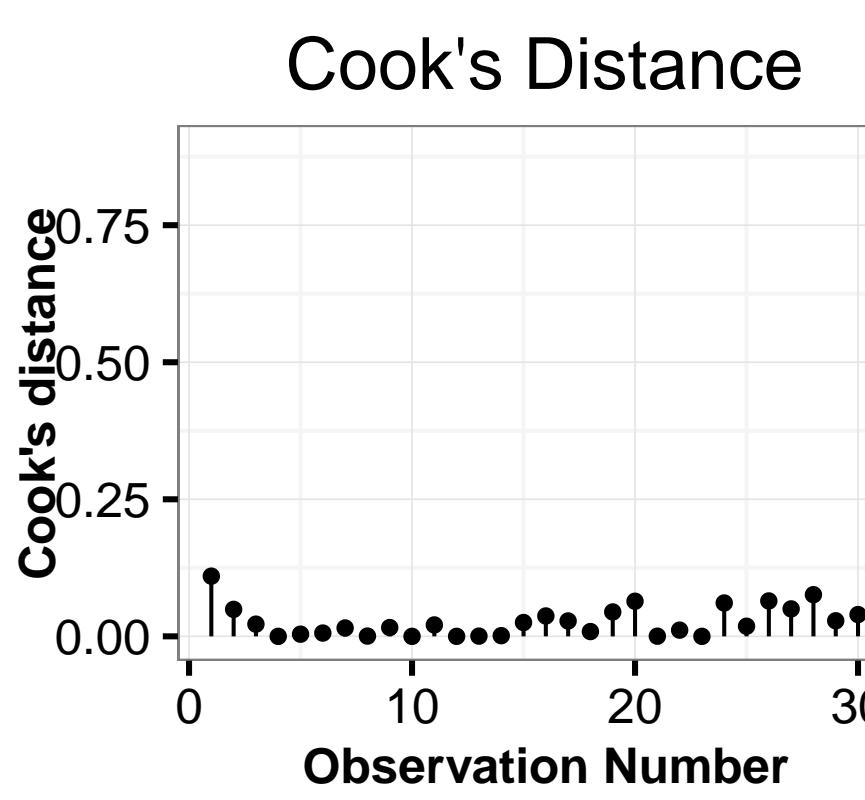
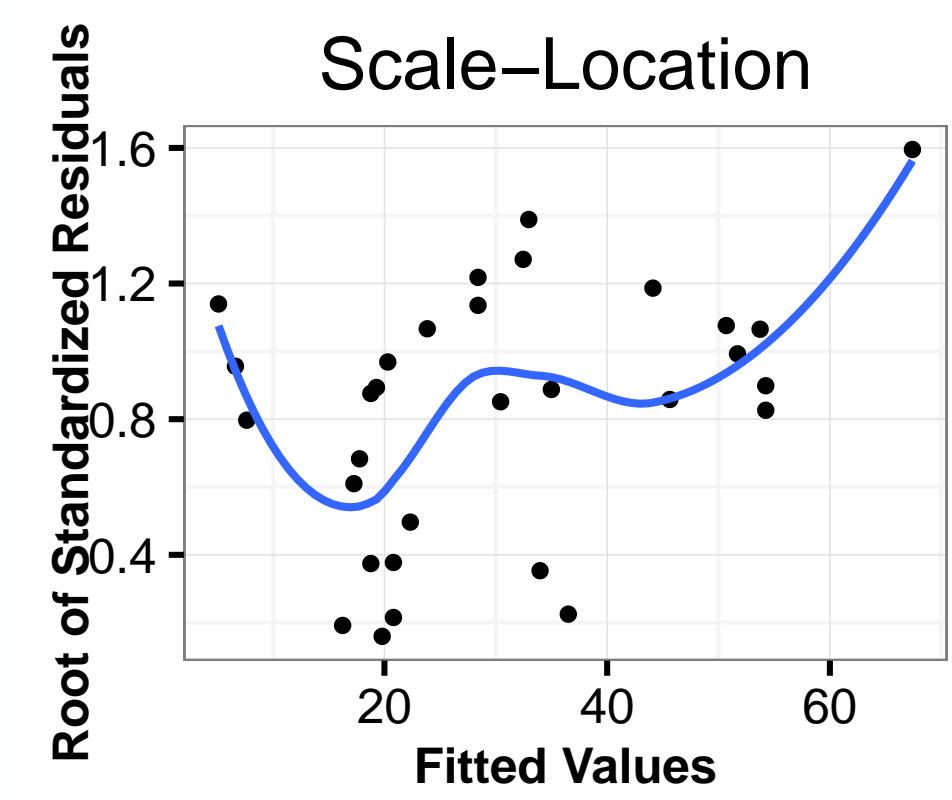
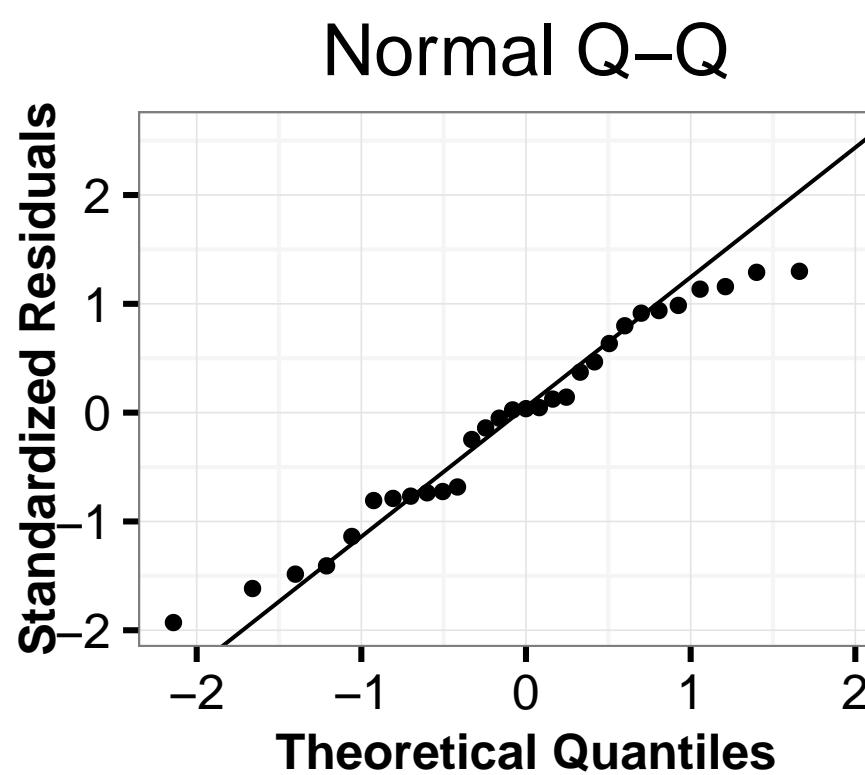
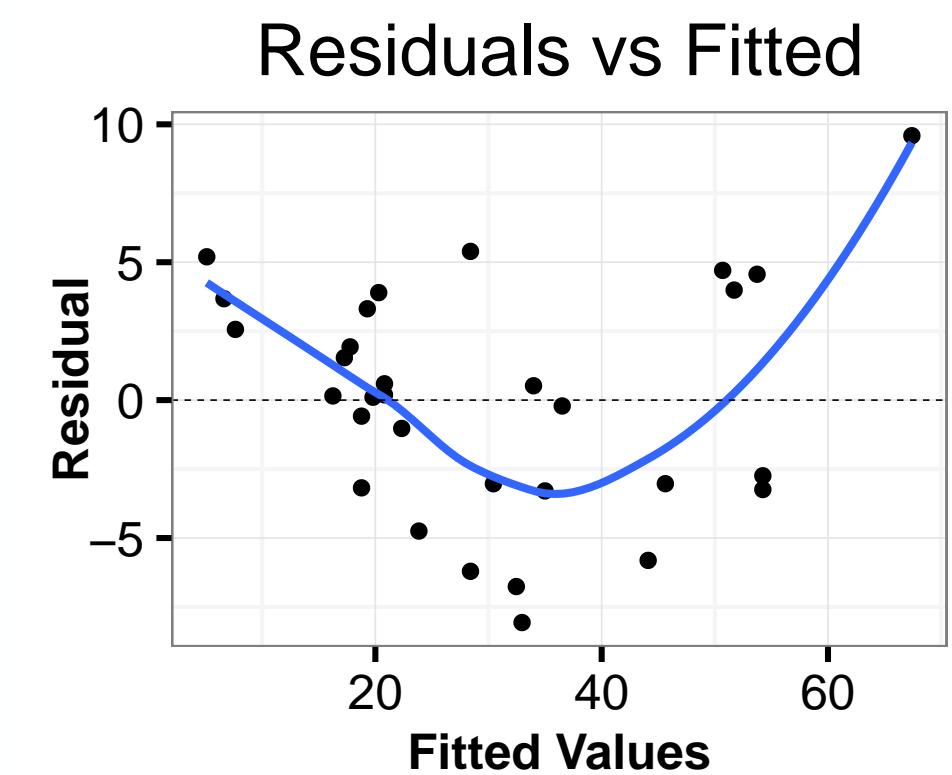


Assess homoscedasticity

# plot model (4)

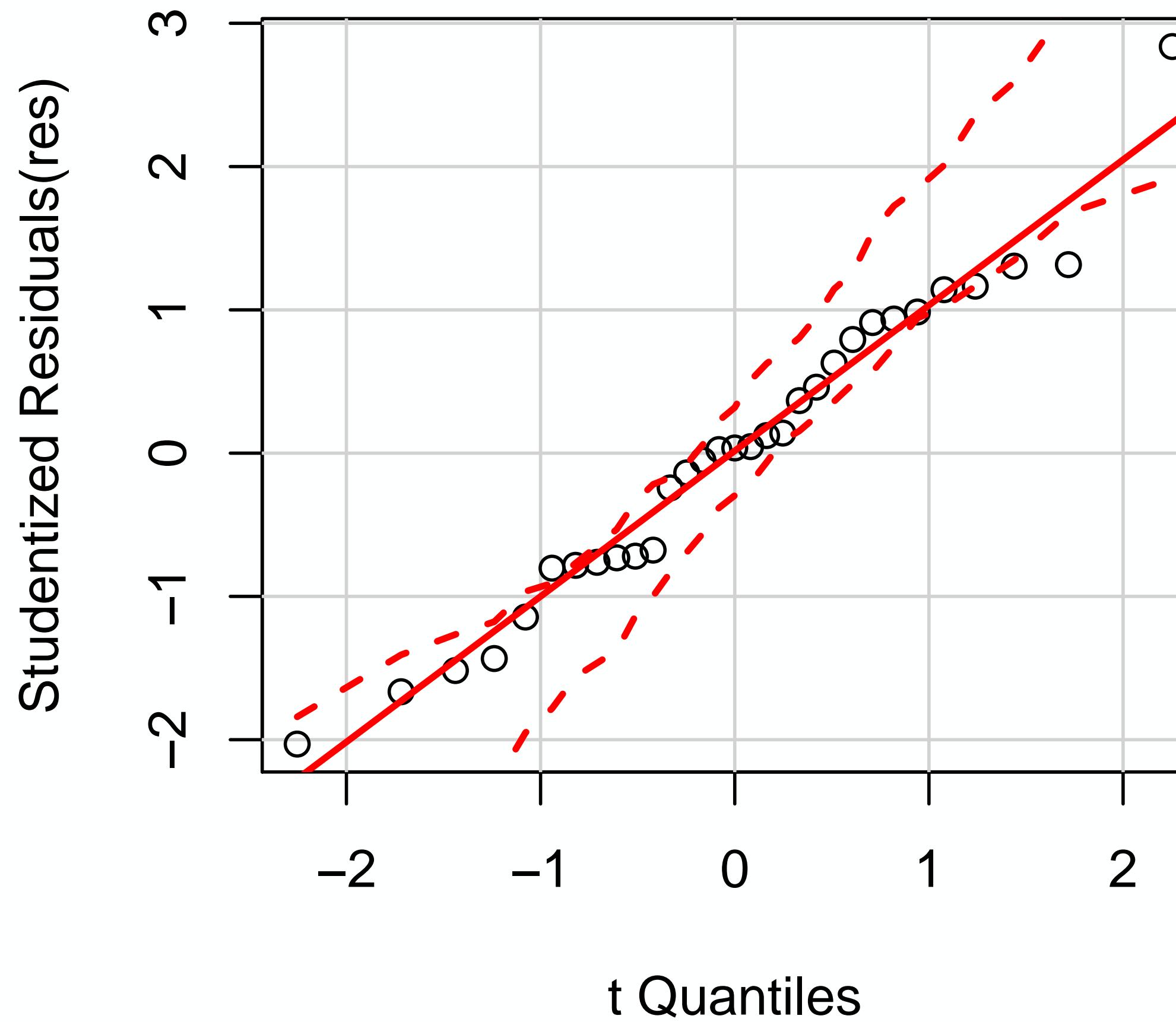
```
> res <- lm(Volume ~ Girth, data = trees)  
> plot(res)
```





# car package

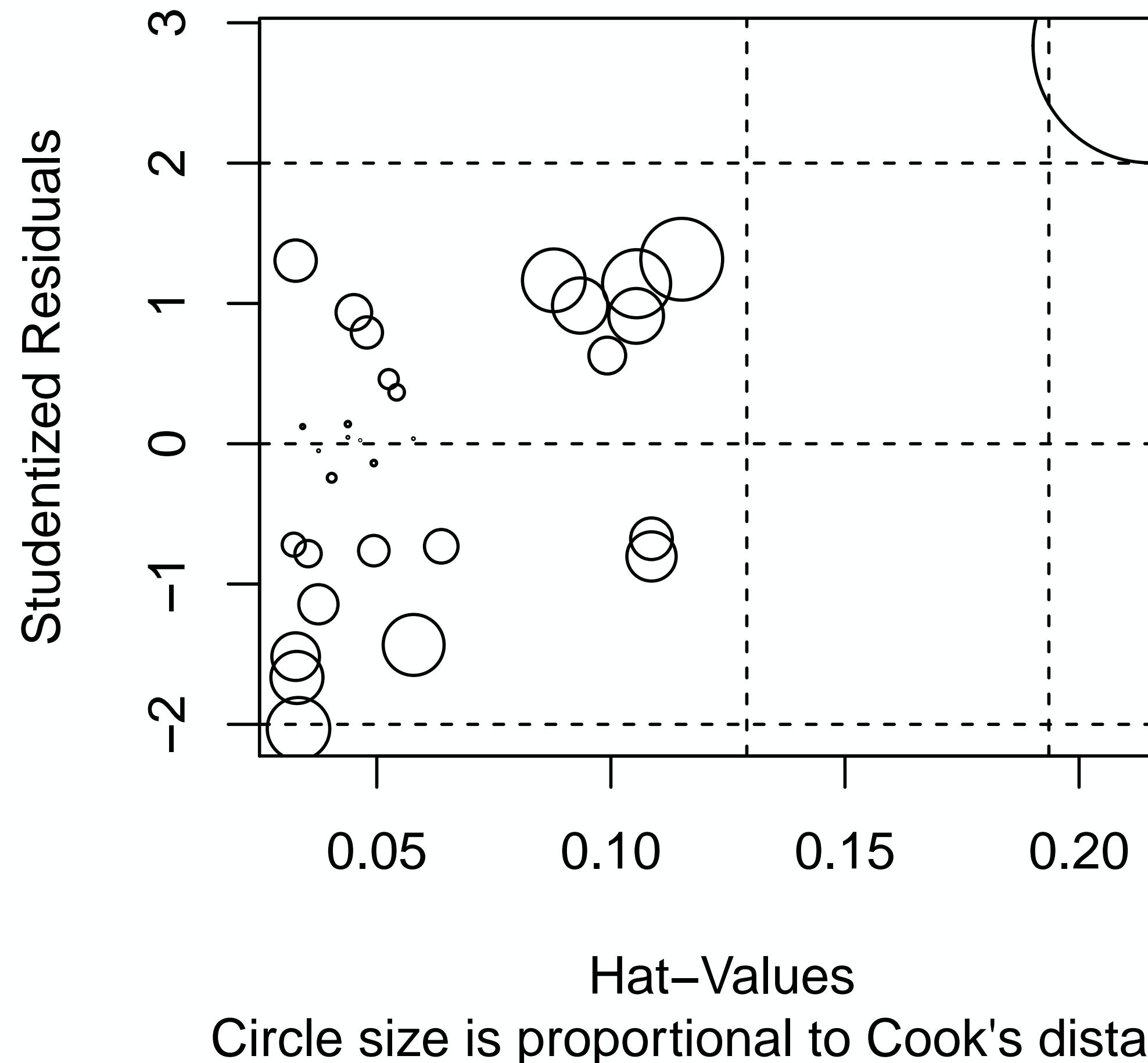
## Q-Q Plot



```
> library(car)
> qqPlot(res, id.method = "identify",
  simulate = TRUE, main = "Q-Q Plot")
```

# car package

## Influence Plot



```
> library(car)
> influencePlot(res, id.method = "identify",
  main = "Influence Plot")
```

# Recent development

- John Fox
- Michael Friendly
- **matplotlib**



DATA VISUALIZATION WITH GGPLOT2

**Let's practice!**