

CSC 587 Homework 1

Corey Osborne

We read Su_raw_matrix.txt into a data frame “su”

```
su <- read.delim("Su_raw_matrix.txt")
```

Gets mean and standard deviation of Liver_2.CEL column, as well as the column mean and sum, and then print the results

```
liver2_mean <- mean(su$Liver_2.CEL)
liver2_sd <- sd(su$Liver_2.CEL)

col_means <- colMeans(su)
col_sums <- colSums(su)

print(liver2_mean)
```

```
## [1] 241.8246
```

```
print(liver2_sd)
```

```
## [1] 1133.352
```

```
print(col_means)
```

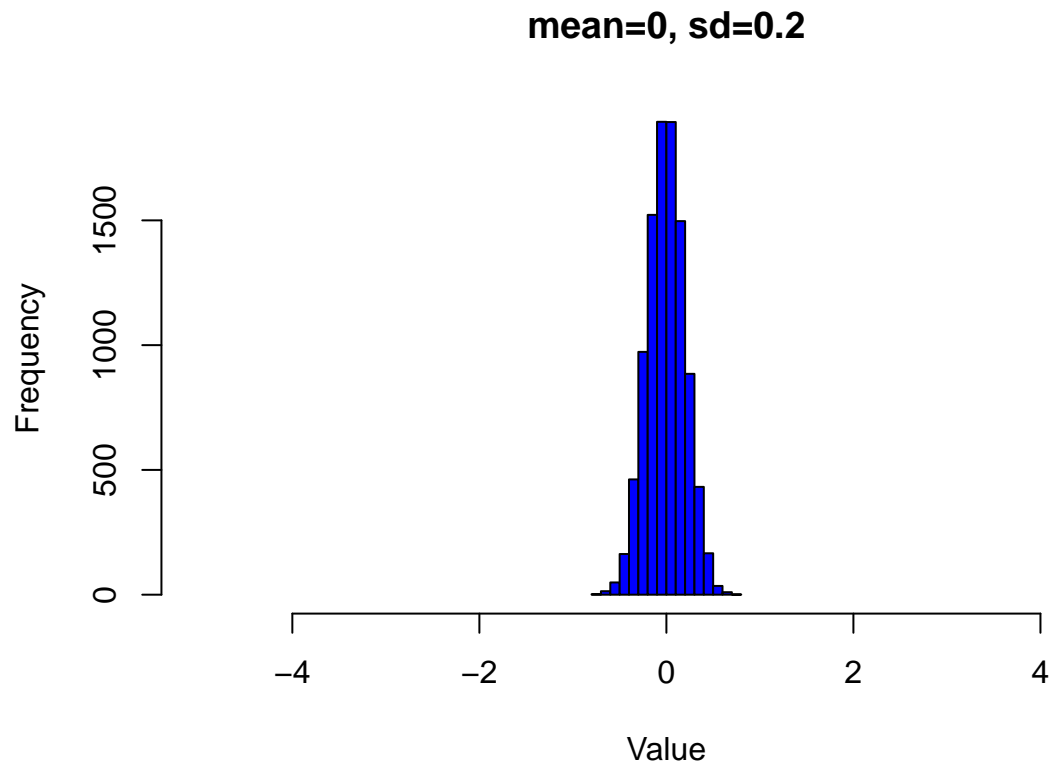
```
##      Brain_1.CEL      Brain_2.CEL Fetal_brain_1.CEL Fetal_brain_2.CEL
##      204.9763      315.0924      198.3439      267.6551
## Fetal_liver_1.CEL Fetal_liver_2.CEL      Liver_1.CEL      Liver_2.CEL
##      209.8722      399.1482      160.8558      241.8246
```

```
print(col_sums)
```

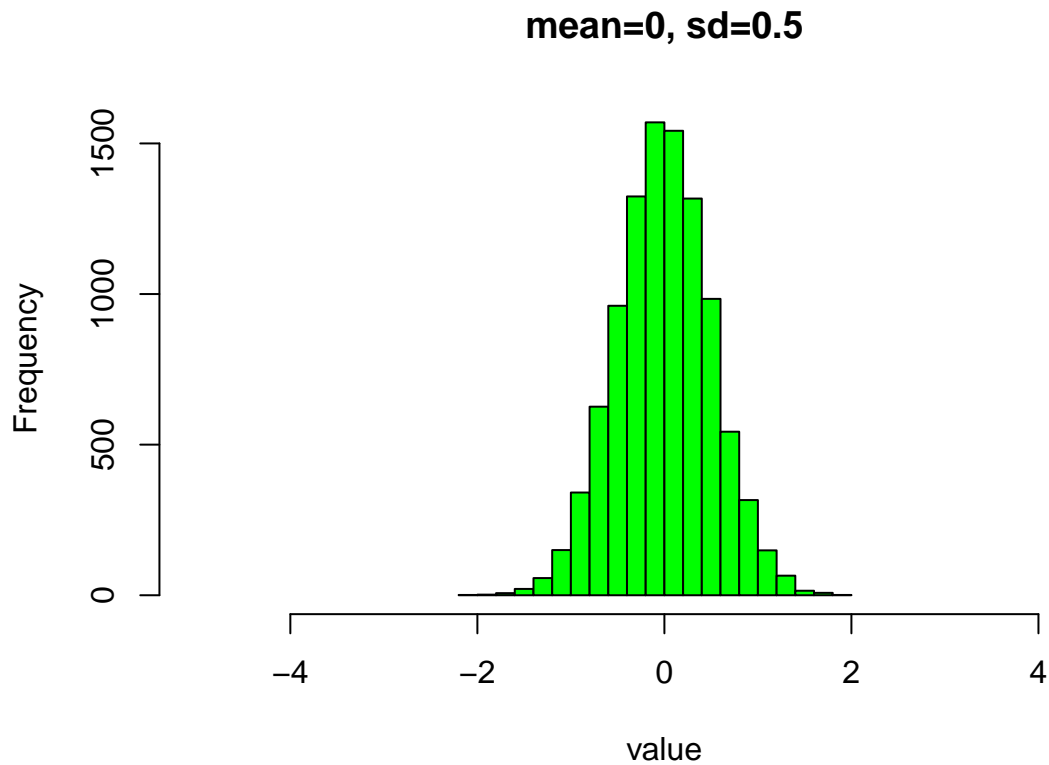
```
##      Brain_1.CEL      Brain_2.CEL Fetal_brain_1.CEL Fetal_brain_2.CEL
##      2588031      3978357      2504290      3379413
## Fetal_liver_1.CEL Fetal_liver_2.CEL      Liver_1.CEL      Liver_2.CEL
##      2649846      5039645      2030966      3053278
```

Here we generate 10,000 random numbers to use for plotting histograms and normal distributions. We do one with a smaller standard deviation of 0.2, and another with a standard deviation of 0.5.

```
randomNum1 <- rnorm(10000, mean=0, sd=0.2)
hist(randomNum1, xlim=c(-5,5), main="mean=0, sd=0.2", xlab="Value", col="blue")
```



```
randomNum2 <- rnorm(10000, mean=0, sd=0.5)
hist(randomNum2, xlim=c(-5,5), main="mean=0, sd=0.5", xlab="value", col="green")
```

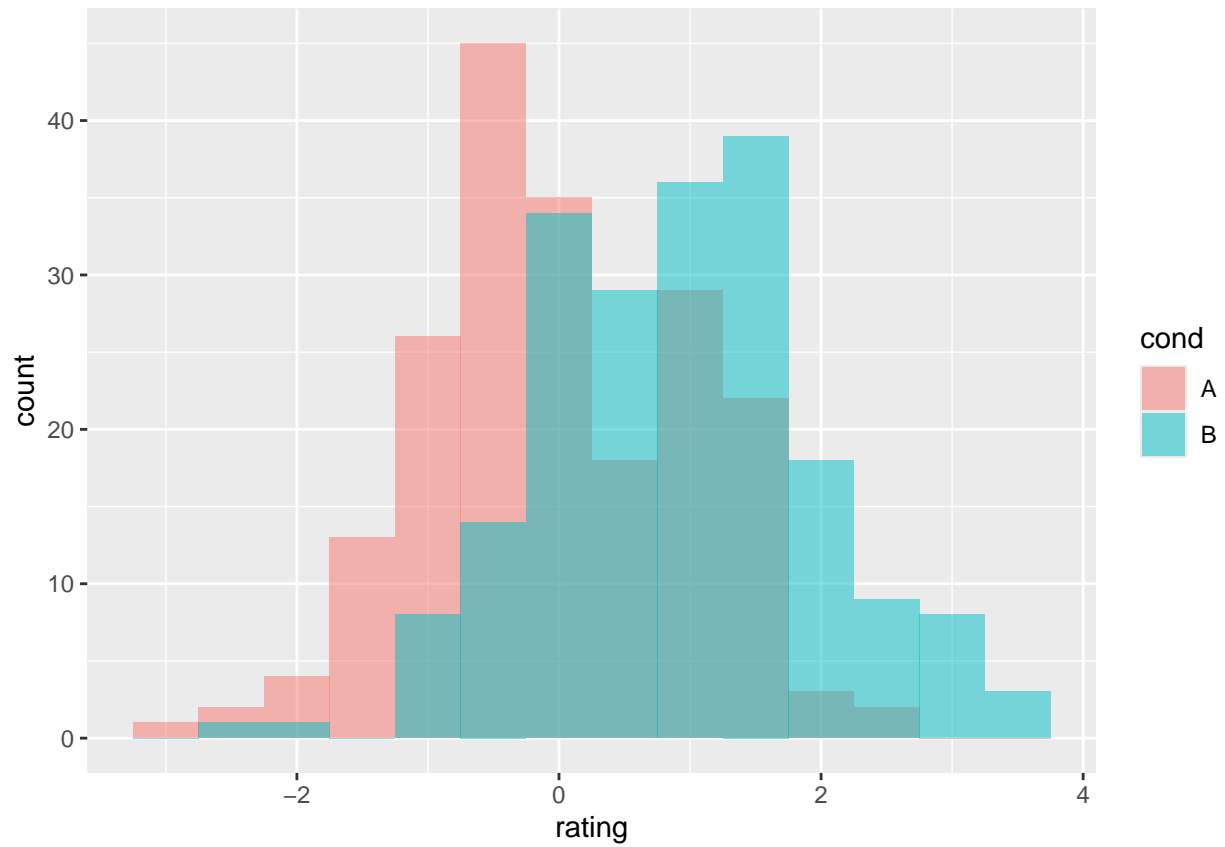


dat will be a sample dataset with two categories, A and B, which will contain 200 values each and then normally distributes, where A has a mean of 0 and B has a mean of 0.8.

```
library(ggplot2)
dat <- data.frame(cond = factor(rep(c("A", "B"), each=200)),
                  rating = c(rnorm(200), rnorm(200, mean=.8)))
```

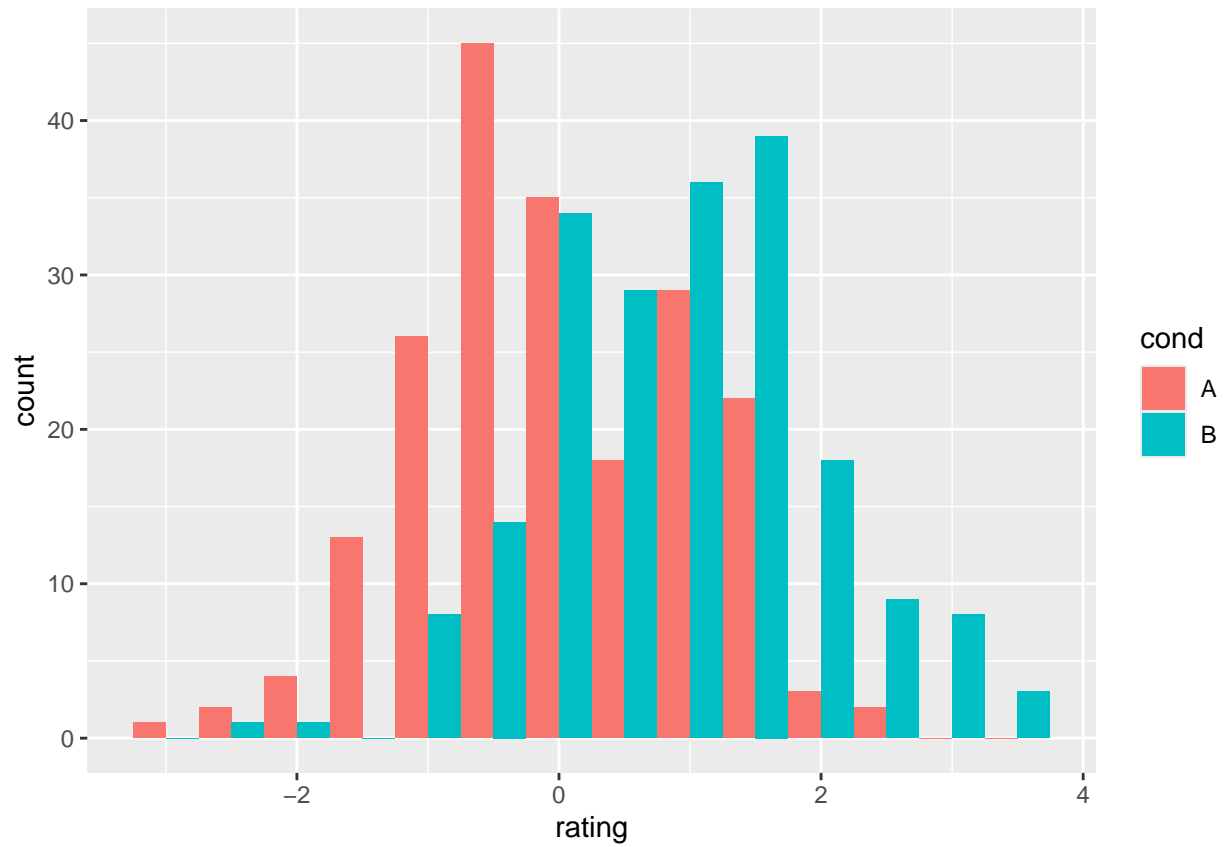
This will produce an overlaid histogram, plotting 'rating' attribute, with bins of 0.5 from the dat sample dataset

```
ggplot(dat, aes(x=rating, fill=cond)) +
  geom_histogram(binwidth=.5, alpha=.5, position="identity")
```



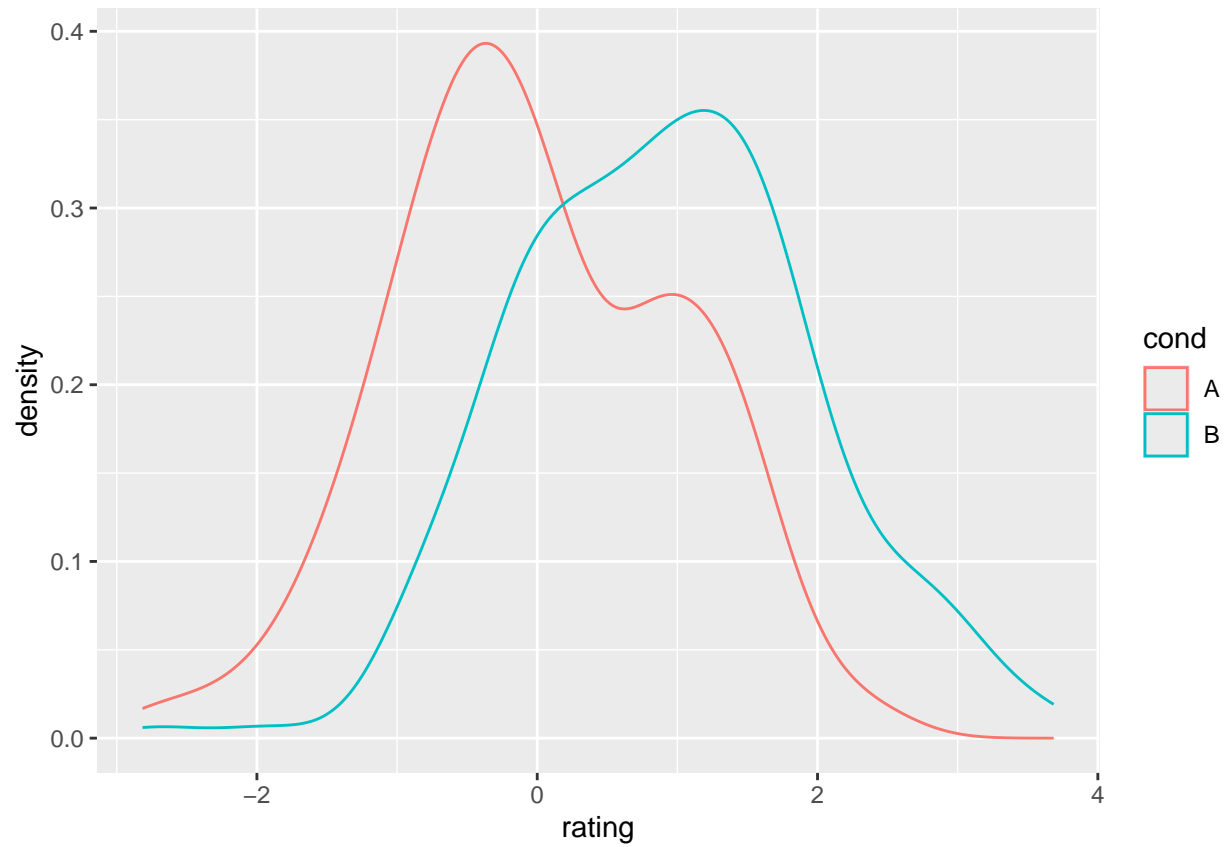
An interleaved histogram of the dat dataset, plotting 'rating' attribute (two side-by-side histograms)

```
ggplot(dat, aes(x=rating, fill=cond)) +  
  geom_histogram(binwidth=.5, position="dodge")
```



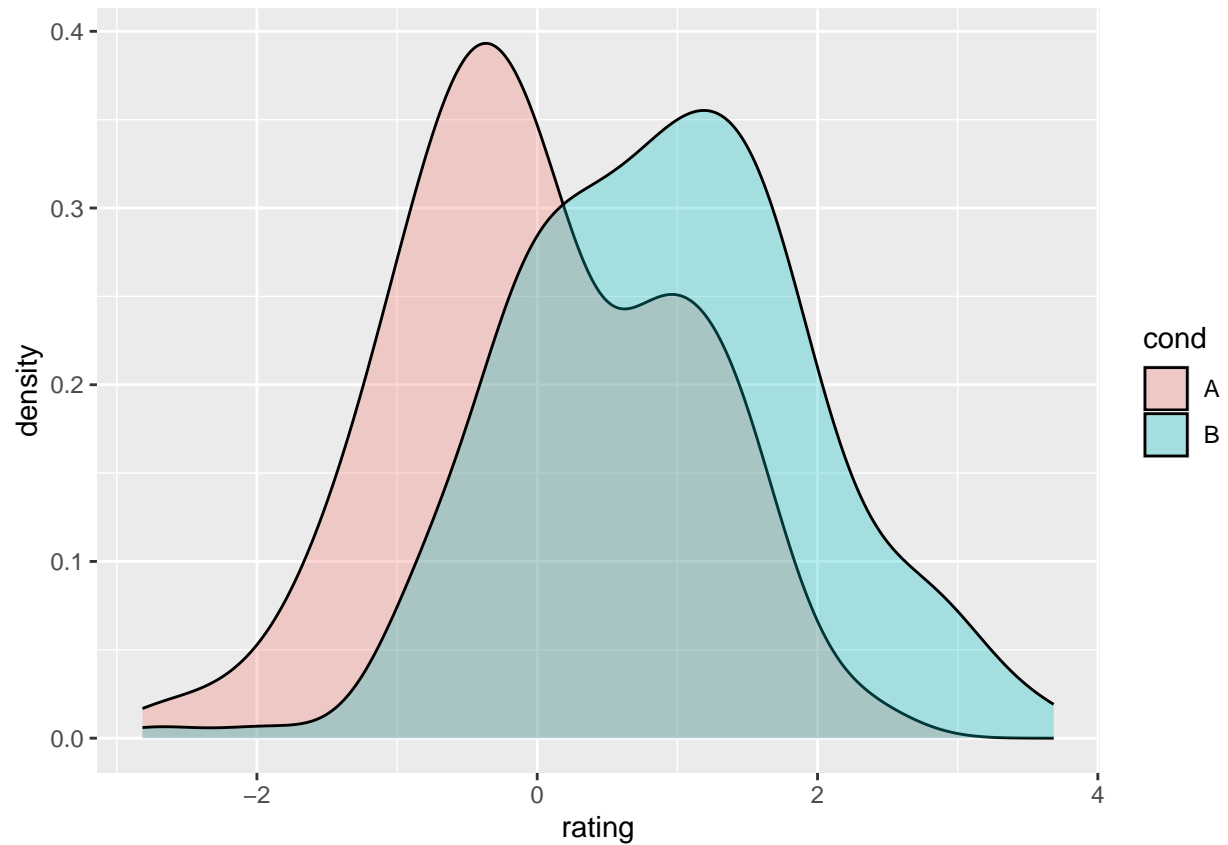
Density plots of dat

```
ggplot(dat, aes(x=rating, colour=cond)) + geom_density()
```



Density plots of dat with fill

```
ggplot(dat, aes(x=rating, fill=cond)) + geom_density(alpha=.3)
```

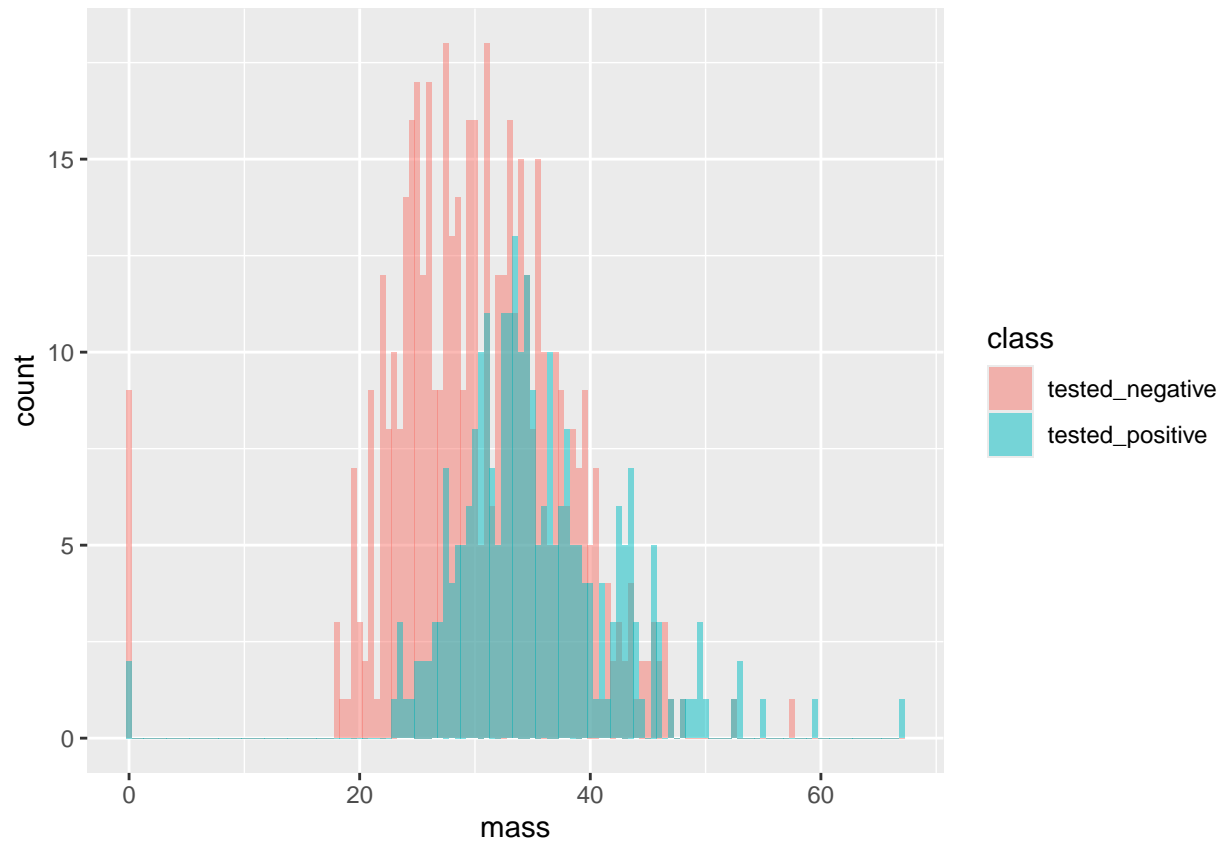


Instead of using random sample set (dat), we'll use a diabetes dataset produced from diabetes_train.csv

```
diabetes <- read.csv("diabetes_train.csv")
```

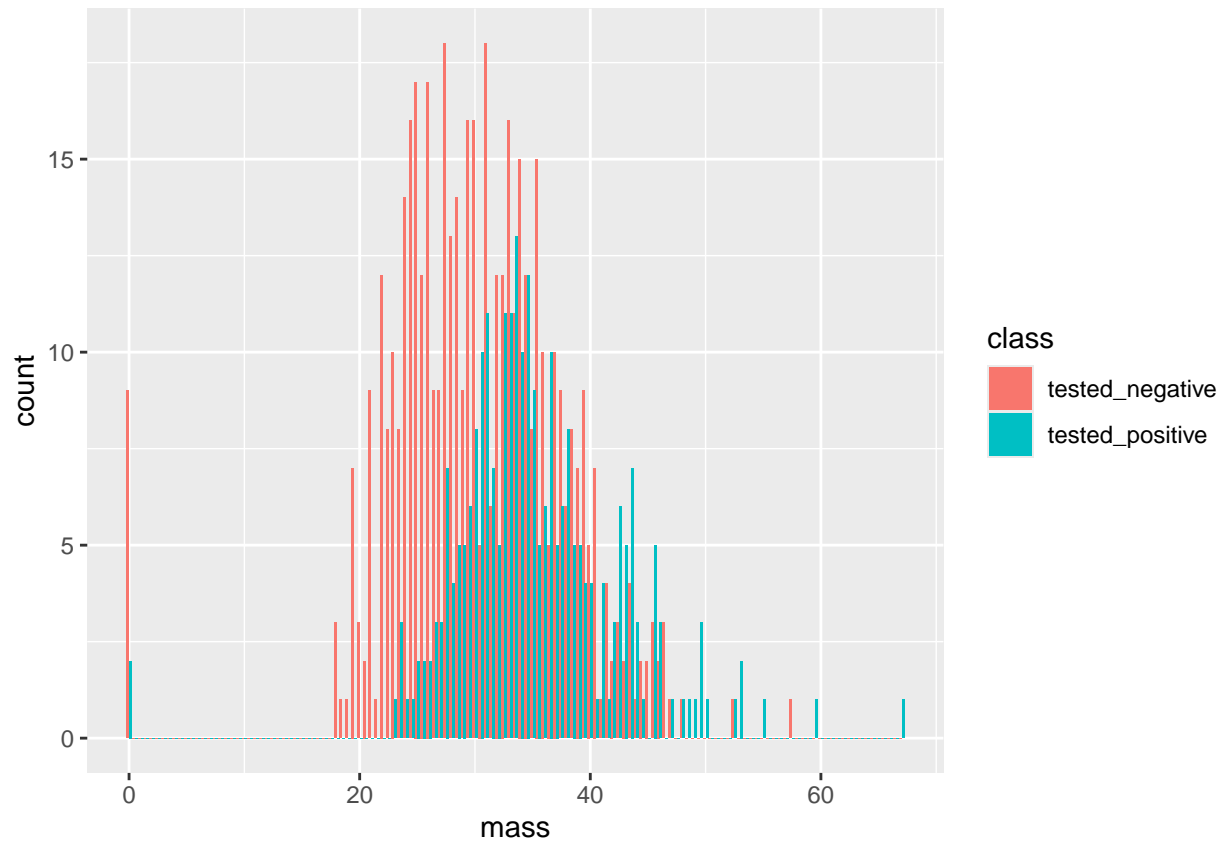
Overlaid histogram of diabetes dataset, plotting mass attribute

```
ggplot(diabetes, aes(x=mass, fill=class)) +  
  geom_histogram(binwidth=.5, alpha=.5, position="identity")
```



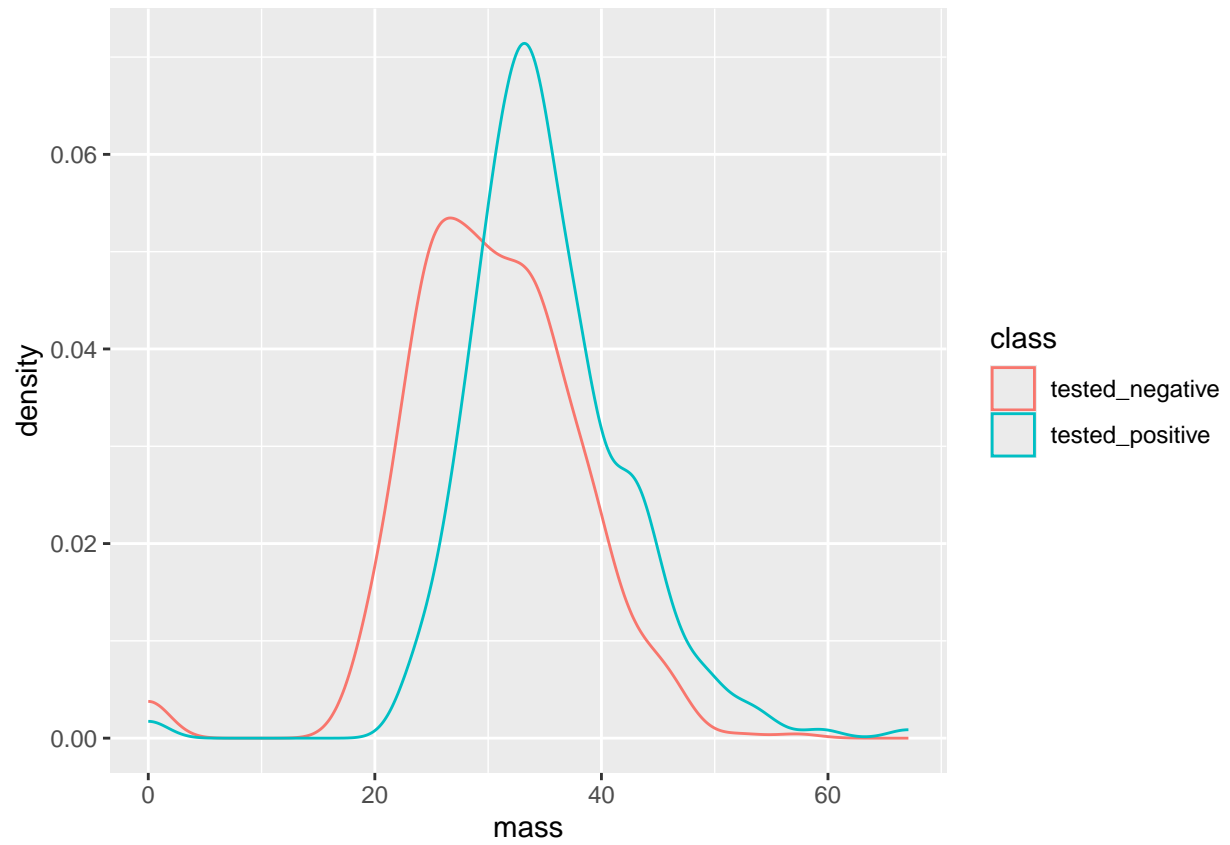
Interleaved histogram of diabetes dataset, plotting mass attribute

```
ggplot(diabetes, aes(x=mass, fill=class)) +  
  geom_histogram(binwidth=.5, position="dodge")
```

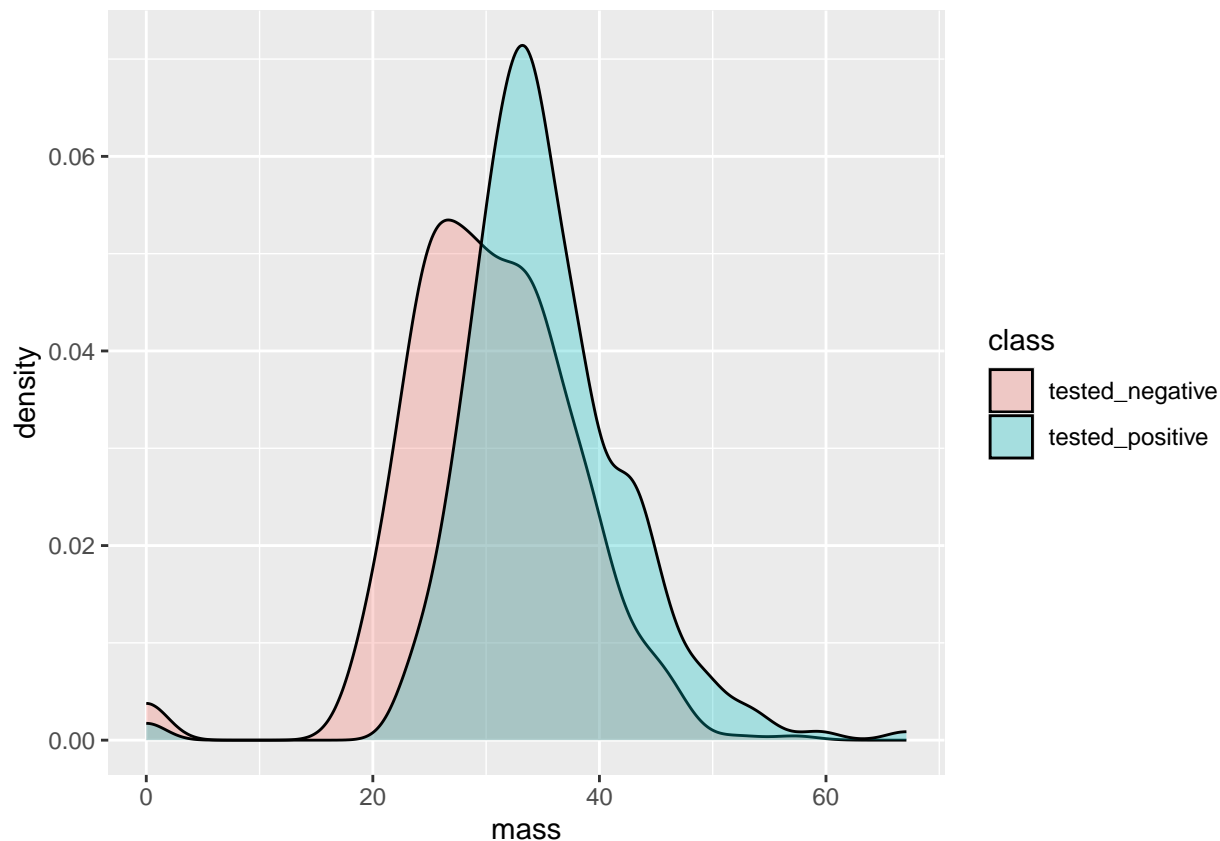
Diabetes dataset density plot

```
ggplot(diabetes, aes(x=mass, colour=class)) + geom_density()
```



Diabetes dataset density plot with fill

```
ggplot(diabetes, aes(x=mass, fill=class)) + geom_density(alpha=.3)
```



```
library(tidyverse)
```

```
## -- Attaching core tidyverse packages ----- tidyverse 2.0.0 --
## v dplyr      1.1.4      v readr      2.1.5
## v forcats    1.0.0      v stringr    1.5.1
## v lubridate  1.9.4      v tibble     3.2.1
## v purrr      1.0.2      v tidyr      1.3.1
## -- Conflicts ----- tidyverse_conflicts() --
## x dplyr::filter() masks stats::filter()
## x dplyr::lag()     masks stats::lag()
## i Use the conflicted package (<http://conflicted.r-lib.org/>) to force all conflicts to become errors
```

```
passengers <- read.csv("titanic.csv")
```

Here we remove any rows with missing values and provide a summary of statistical information

```
passengers %>% drop_na() %>% summary()
```

```
##           X           PassengerId         Survived         Pclass
##  Min.      : 0.0   Min.      : 1.0   Min.      :0.0000   Length:714
##  1st Qu.:221.2   1st Qu.:222.2   1st Qu.:0.0000   Class :character
##  Median :444.0   Median :445.0   Median :0.0000   Mode  :character
##  Mean    :447.6   Mean    :448.6   Mean    :0.4062
##  3rd Qu.:676.8   3rd Qu.:677.8   3rd Qu.:1.0000
```

```
## Max. :890.0 Max. :891.0 Max. :1.0000
## Name Sex Age SibSp
## Length:714 Length:714 Min. : 0.42 Min. :0.0000
## Class :character Class :character 1st Qu.:20.12 1st Qu.:0.0000
## Mode :character Mode :character Median :28.00 Median :0.0000
## Mean :29.70 Mean :0.5126
## 3rd Qu.:38.00 3rd Qu.:1.0000
## Max. :80.00 Max. :5.0000
## Parch Ticket Fare Cabin
## Min. :0.0000 Length:714 Min. : 0.00 Length:714
## 1st Qu.:0.0000 Class :character 1st Qu.: 8.05 Class :character
## Median :0.0000 Mode :character Median : 15.74 Mode :character
## Mean :0.4314 Mean : 34.69
## 3rd Qu.:1.0000 3rd Qu.: 33.38
## Max. :6.0000 Max. :512.33
## Embarked
## Length:714
## Class :character
## Mode :character
##
##
##
```

This filters and then creates a new dataset containing only male passengers (only 10 displayed)

```
passengers %>% filter(Sex == "male") %>% head(10)
```

```
## X PassengerId Survived Pclass Name Sex Age SibSp
## 1 0 1 0 3 Braund, Mr. Owen Harris male 22 1
## 2 4 5 0 3 Allen, Mr. William Henry male 35 0
## 3 5 6 0 3 Moran, Mr. James male NA 0
## 4 6 7 0 1 McCarthy, Mr. Timothy J male 54 0
## 5 7 8 0 3 Palsson, Master. Gosta Leonard male 2 3
## 6 12 13 0 3 Saundercock, Mr. William Henry male 20 0
## 7 13 14 0 3 Andersson, Mr. Anders Johan male 39 1
## 8 16 17 0 3 Rice, Master. Eugene male 2 4
## 9 17 18 1 2 Williams, Mr. Charles Eugene male NA 0
## 10 20 21 0 2 Fynney, Mr. Joseph J male 35 0
## Parch Ticket Fare Cabin Embarked
## 1 0 A/5 21171 7.2500 S
## 2 0 373450 8.0500 S
## 3 0 330877 8.4583 Q
## 4 0 17463 51.8625 E46 S
## 5 1 349909 21.0750 S
## 6 0 A/5. 2151 8.0500 S
## 7 5 347082 31.2750 S
## 8 1 382652 29.1250 Q
## 9 0 244373 13.0000 S
## 10 0 239865 26.0000 S
```

We then sort all passengers by Fare, in descending order (the highest-paying passengers will appear first))
(only 10 displayed)

```
passengers %>% arrange(desc(Fare)) %>% head(10)
```

##	X	PassengerId	Survived	Pclass	Name	Sex
## 1	258	259	1	1	Ward, Miss. Anna	female
## 2	679	680	1	1	Cardeza, Mr. Thomas Drake Martinez	male
## 3	737	738	1	1	Lesurer, Mr. Gustave J	male
## 4	27	28	0	1	Fortune, Mr. Charles Alexander	male
## 5	88	89	1	1	Fortune, Miss. Mabel Helen	female
## 6	341	342	1	1	Fortune, Miss. Alice Elizabeth	female
## 7	438	439	0	1	Fortune, Mr. Mark	male
## 8	311	312	1	1	Ryerson, Miss. Emily Borie	female
## 9	742	743	1	1	Ryerson, Miss. Susan Parker "Suzette"	female
## 10	118	119	0	1	Baxter, Mr. Quigg Edmond	male

##	Age	SibSp	Parch	Ticket	Fare	Cabin	Embarked
## 1	35	0	0	PC 17755	512.3292		C
## 2	36	0	1	PC 17755	512.3292	B51 B53 B55	C
## 3	35	0	0	PC 17755	512.3292	B101	C
## 4	19	3	2	19950	263.0000	C23 C25 C27	S
## 5	23	3	2	19950	263.0000	C23 C25 C27	S
## 6	24	3	2	19950	263.0000	C23 C25 C27	S
## 7	64	1	4	19950	263.0000	C23 C25 C27	S
## 8	18	2	2	PC 17608	262.3750	B57 B59 B63 B66	C
## 9	21	2	2	PC 17608	262.3750	B57 B59 B63 B66	C
## 10	24	0	1	PC 17558	247.5208	B58 B60	C

This calculates the total number of family members onboard, where Parch is parents/children, and SibSp is siblings/spouse (only 10 displayed)

```
passengers %>% mutate(Famsize = Parch + SibSp) %>% head(10)
```

##	X	PassengerId	Survived	Pclass	Name	Sex	Age	SibSp	Parch
## 1	0	1	0	3					
## 2	1	2	1	1					
## 3	2	3	1	3					
## 4	3	4	1	1					
## 5	4	5	0	3					
## 6	5	6	0	3					
## 7	6	7	0	1					
## 8	7	8	0	3					
## 9	8	9	1	3					
## 10	9	10	1	2					

##	Name	Sex	Age	SibSp	Parch
## 1	Braund, Mr. Owen Harris	male	22	1	0
## 2	Cumings, Mrs. John Bradley (Florence Briggs Thayer)	female	38	1	0
## 3	Heikkinen, Miss. Laina	female	26	0	0
## 4	Futrelle, Mrs. Jacques Heath (Lily May Peel)	female	35	1	0
## 5	Allen, Mr. William Henry	male	35	0	0
## 6	Moran, Mr. James	male	NA	0	0
## 7	McCarthy, Mr. Timothy J	male	54	0	0
## 8	Palsson, Master. Gosta Leonard	male	2	3	1
## 9	Johnson, Mrs. Oscar W (Elisabeth Vilhelmina Berg)	female	27	0	2
## 10	Nasser, Mrs. Nicholas (Adele Achem)	female	14	1	0

```
##           Ticket      Fare Cabin Embarked Famsize
## 1      A/5 21171   7.2500      S         1
## 2      PC 17599  71.2833    C85         1
## 3 STON/O2. 3101282   7.9250      S         0
## 4      113803  53.1000   C123         1
## 5      373450   8.0500      S         0
## 6      330877   8.4583      Q         0
## 7      17463  51.8625    E46         0
## 8      349909  21.0750      S         4
## 9      347742  11.1333      S         2
## 10     237736  30.0708      C         1
```

Here we check the average ticket fare by gender, as well as find the total number of survivors per gender

```
passengers %>% group_by(Sex) %>%
  summarise(meanFare = mean(Fare), numSurv = sum(Survived))
```

```
## # A tibble: 2 x 3
##   Sex    meanFare numSurv
##   <chr>    <dbl>   <int>
## 1 female    44.5     233
## 2 male     25.5     109
```

Here we use quantile to calculate the 10th, 30th, 50th, and 60th percentiles of the skin attribute from the diabetes dataset

```
percentiles <- quantile(diabetes$skin, probs = c(0.1, 0.3, 0.5, 0.6))
print(percentiles)
```

```
## 10% 30% 50% 60%
##   0  10  23  27
```