Assignment 2 MATH 208 (Question 2)

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Question 2

a)

```
data(Diabetes)
df_diabetes = as_tibble(Diabetes)
df_grouped = df_diabetes %>%
  group_by(group) %>%
  summarise_all(list(
    Avg=mean,
    Med=median
    )
  ) %>%
  pivot_longer(
    col=contains('_'),
    names_to = "Measure") %>%
pivot_wider(
  id_cols = Measure,
  names_from = group
  ) %>%
  arrange(
    desc(Measure)
    )
df_grouped
```

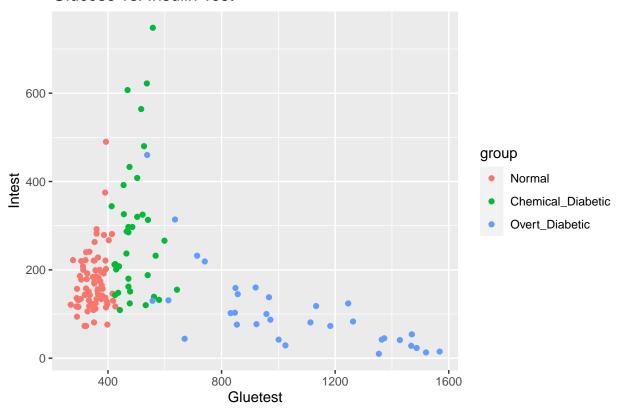
```
## # A tibble: 10 x 4
                   Normal Chemical_Diabetic Overt_Diabetic
##
     Measure
##
                    <dbl>
                                      <dbl>
                                                      <dbl>
      <chr>
## 1 sspg_Med
                  105
                                     223
                                                    320
                  114
                                     209.
                                                   319.
## 2 sspg_Avg
## 3 relwt_Med
                    0.95
                                       1.06
                                                      0.98
## 4 relwt_Avg
                    0.937
                                       1.06
                                                      0.984
## 5 instest_Med 157
                                     252.
                                                    83
## 6 instest_Avg 173.
                                     288
                                                    106
## 7 glutest_Med 353
                                                   972
                                     476.
## 8 glutest_Avg 350.
                                     494.
                                                   1044.
## 9 glufast_Med 90
                                      99.5
                                                   203
## 10 glufast_Avg 91.2
                                      99.3
                                                   218.
```

The variable that seem to differentiate amongst the different types of diabetes is glutest, intest, glufast and sspg

b)

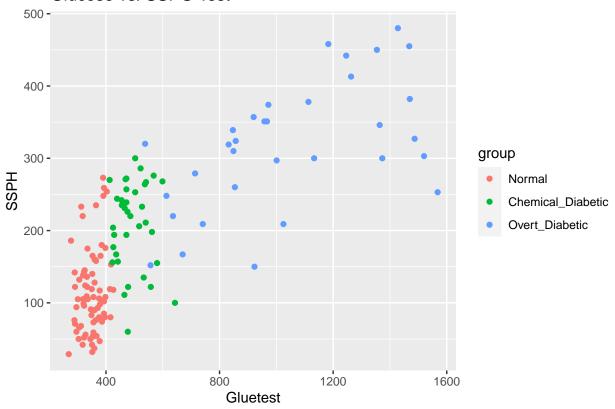
```
df_test = df_diabetes %>% group_by(group)
ggplot(df_diabetes, aes(x=glutest, y=instest, col = group)) + geom_point() +
labs(title="Glucose vs. Insulin Test", x="Gluetest", y="Intest")
```

Glucose vs. Insulin Test

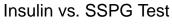


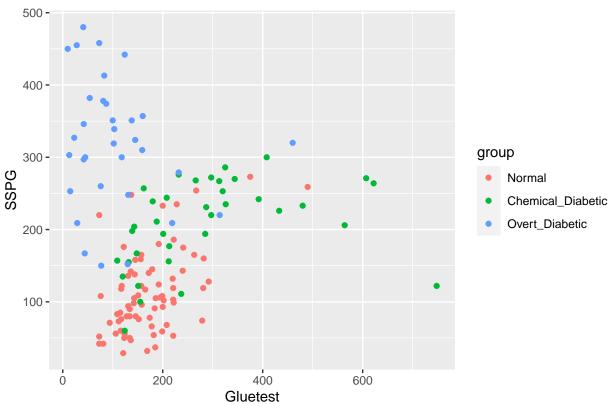
```
ggplot(df_diabetes, aes(x=glutest, y = sspg, col = group)) + geom_point() +
labs(title="Glucose vs. SSPG Test", x="Gluetest", y="SSPH")
```

Glucose vs. SSPG Test



```
ggplot(df_diabetes, aes(x=instest, y = sspg, col = group)) + geom_point() +
labs(title="Insulin vs. SSPG Test", x="Gluetest", y="SSPG")
```



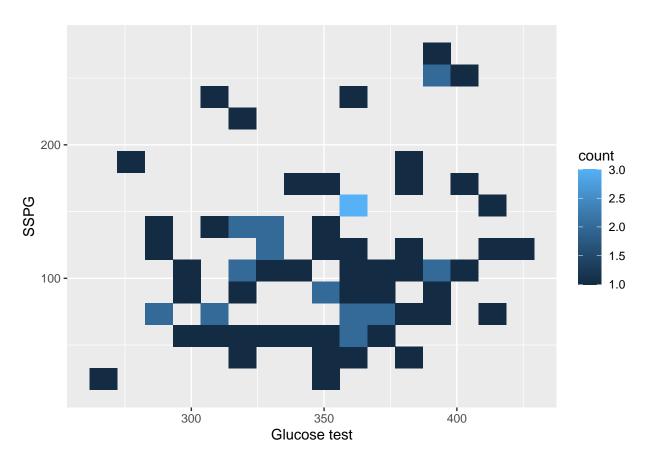


Glucose vs. SSPG test seem to return the greatest distinction looking at the plots above

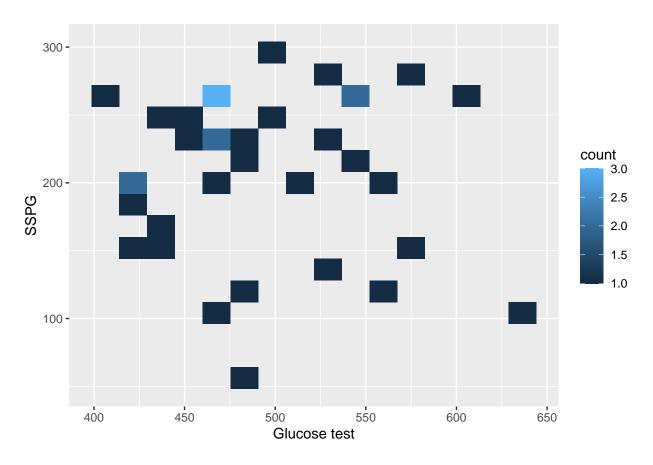
c)

```
# Group data
group_normal = df_diabetes[df_diabetes$group == "Normal",]
group_chemical = df_diabetes[df_diabetes$group == "Chemical_Diabetic",]
group_overt = df_diabetes[df_diabetes$group == "Overt_Diabetic",]

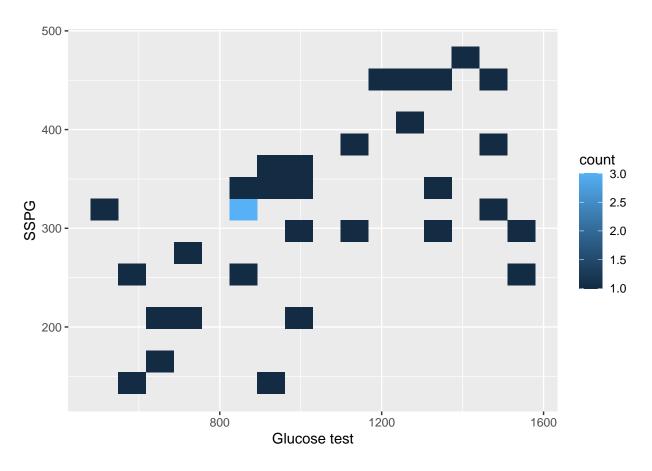
# Histograms
ggplot(group_normal, aes(x=glutest, y=sspg)) + geom_bin2d(bins=15) +
labs(x="Glucose test", y="SSPG")
```



```
ggplot(group_chemical, aes(x=glutest, y=sspg)) + geom_bin2d(bins=15) +
labs(x="Glucose test", y="SSPG")
```



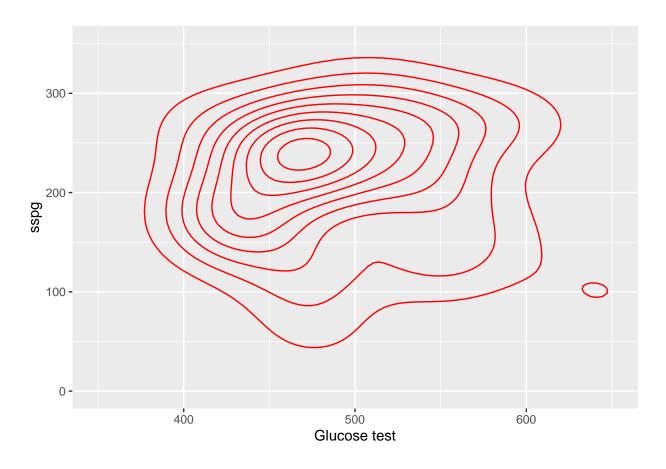
```
ggplot(group_overt, aes(x=glutest, y=sspg)) + geom_bin2d(bins=15) +
labs(x="Glucose test", y="SSPG")
```



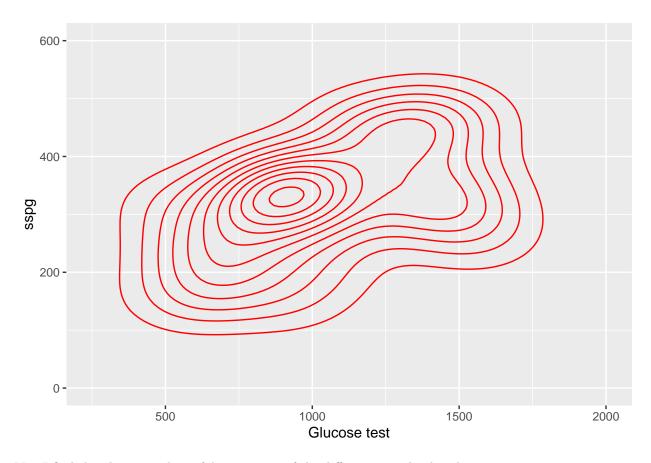
```
# contour plot
ggplot(group_normal, aes(x=glutest,y=sspg)) + geom_density_2d(col="red") +
labs(x="Glucose test", y="sspg") + xlim(c(200,500)) + ylim(c(-50,350))
```



```
ggplot(group_chemical, aes(x=glutest,y=sspg)) + geom_density_2d(col="red") +
labs(x="Glucose test", y="sspg") + xlim(c(350,650)) + ylim(c(0,350))
```



```
ggplot(group_overt, aes(x=glutest,y=sspg)) + geom_density_2d(col="red") +
labs(x="Glucose test", y="sspg") + xlim(c(250,2000)) + ylim(c(0,600))
```



Yes, I find the plots provide useful summaries of the differences in the distributions.