

pt.R

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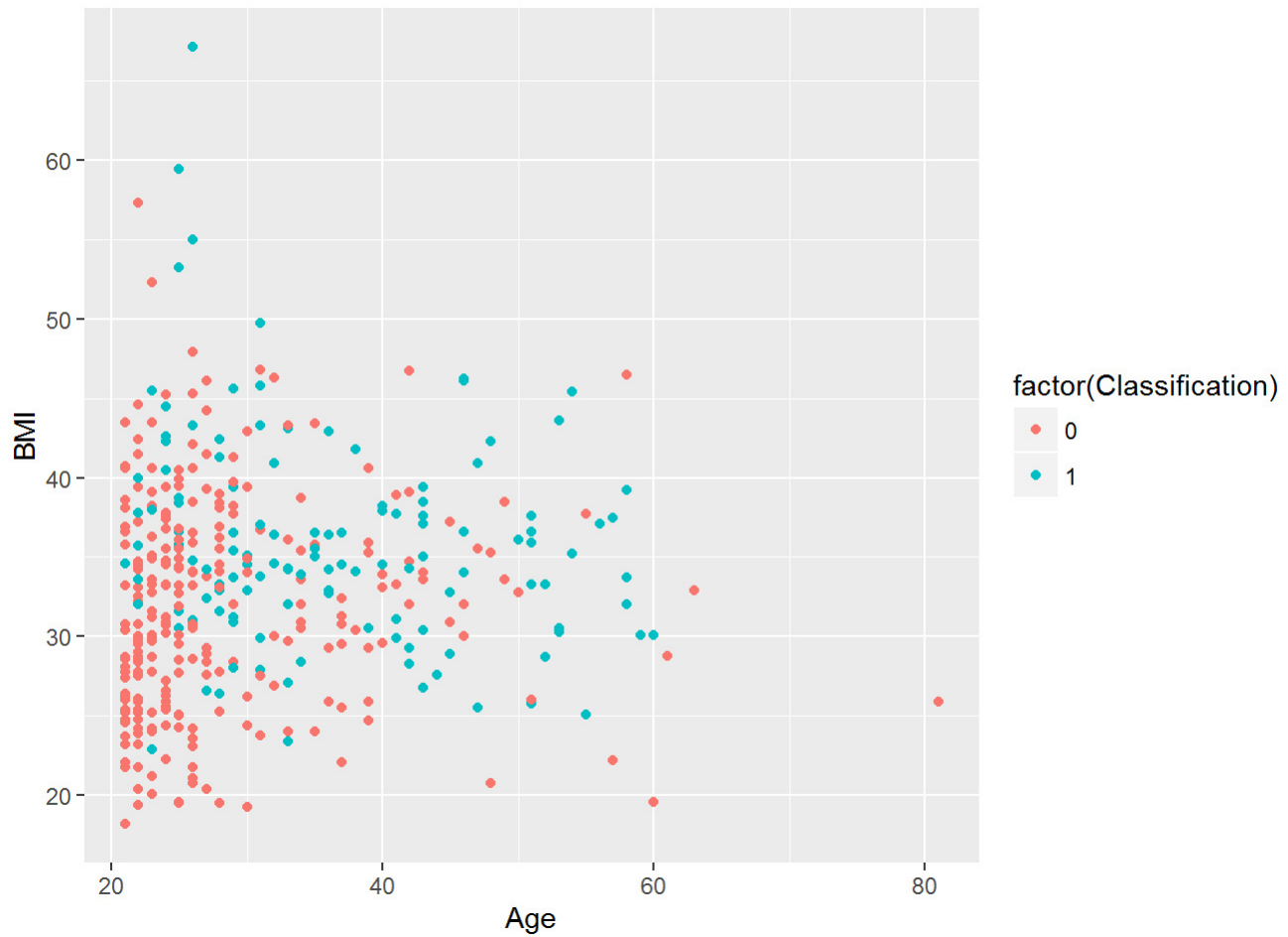
```
setwd("~/Research/Diabetes analysis2/NewDiabetes")
pima.indians.diabetes <- read.csv("C:/Users/IBM_ADMIN/Desktop/pima-indians-diabetes.d
ata", header=FALSE)
names(pima.indians.diabetes)<-c('TimesPregnant','GlucoseLV','DiastolicBP','TricepsThi
ckness','SerumInsulin','BMI','Heridarymarkup','Age','Classification')
View(pima.indians.diabetes)
library(ggplot2)
```

```
## Warning: package 'ggplot2' was built under R version 3.2.5
```

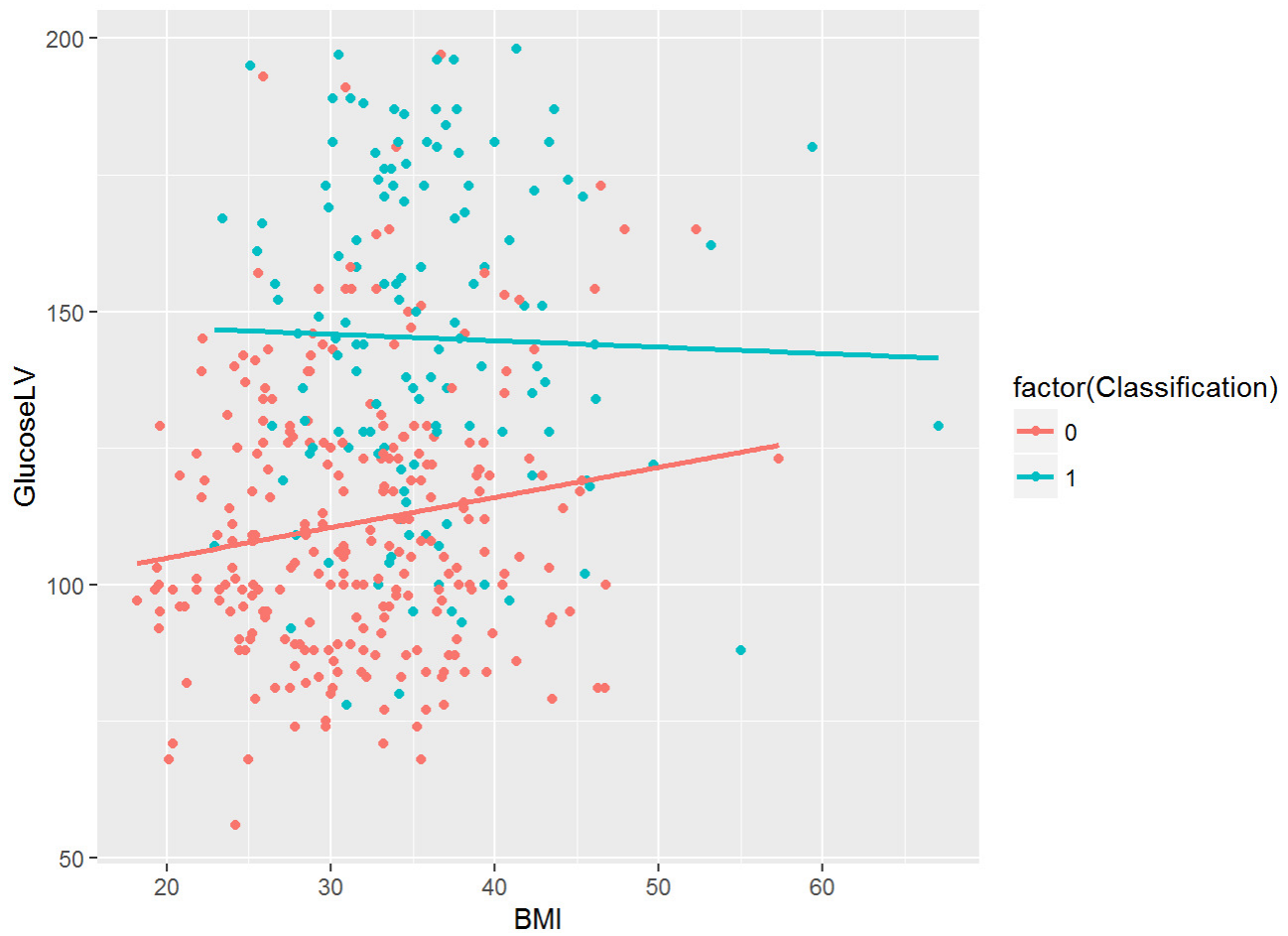
```
pima.indians.diabetes$DiastolicBP[pima.indians.diabetes$DiastolicBP==0]<-NA
pima.indians.diabetes$GlucoseLV[pima.indians.diabetes$GlucoseLV==0]<-NA
pima.indians.diabetes$TricepsThickness[pima.indians.diabetes$TricepsThickness==0]<-NA
pima.indians.diabetes$SerumInsulin[pima.indians.diabetes$SerumInsulin==0]<-NA
pima.indians.diabetes$BMI[pima.indians.diabetes$BMI==0]<-NA
pima.indian.diabetes2<-na.omit(pima.indians.diabetes)
summary(pima.indian.diabetes2)
```

```
## TimesPregnant      GlucoseLV      DiastolicBP      TricepsThickness
## Min.      : 0.000   Min.      : 56.0   Min.      : 24.00   Min.      : 7.00
## 1st Qu.: 1.000   1st Qu.: 99.0   1st Qu.: 62.00   1st Qu.:21.00
## Median : 2.000   Median :119.0   Median : 70.00   Median :29.00
## Mean    : 3.301   Mean    :122.6   Mean    : 70.66   Mean    :29.15
## 3rd Qu.: 5.000   3rd Qu.:143.0   3rd Qu.: 78.00   3rd Qu.:37.00
## Max.    :17.000   Max.    :198.0   Max.    :110.00   Max.    :63.00
## SerumInsulin      BMI      Heridarymarkup      Age
## Min.      : 14.00   Min.    :18.20   Min.    :0.0850   Min.    :21.00
## 1st Qu.: 76.75   1st Qu.:28.40   1st Qu.:0.2697   1st Qu.:23.00
## Median :125.50   Median :33.20   Median :0.4495   Median :27.00
## Mean    :156.06   Mean    :33.09   Mean    :0.5230   Mean    :30.86
## 3rd Qu.:190.00   3rd Qu.:37.10   3rd Qu.:0.6870   3rd Qu.:36.00
## Max.    :846.00   Max.    :67.10   Max.    :2.4200   Max.    :81.00
## Classification
## Min.      :0.0000
## 1st Qu.:0.0000
## Median :0.0000
## Mean    :0.3316
## 3rd Qu.:1.0000
## Max.    :1.0000
```

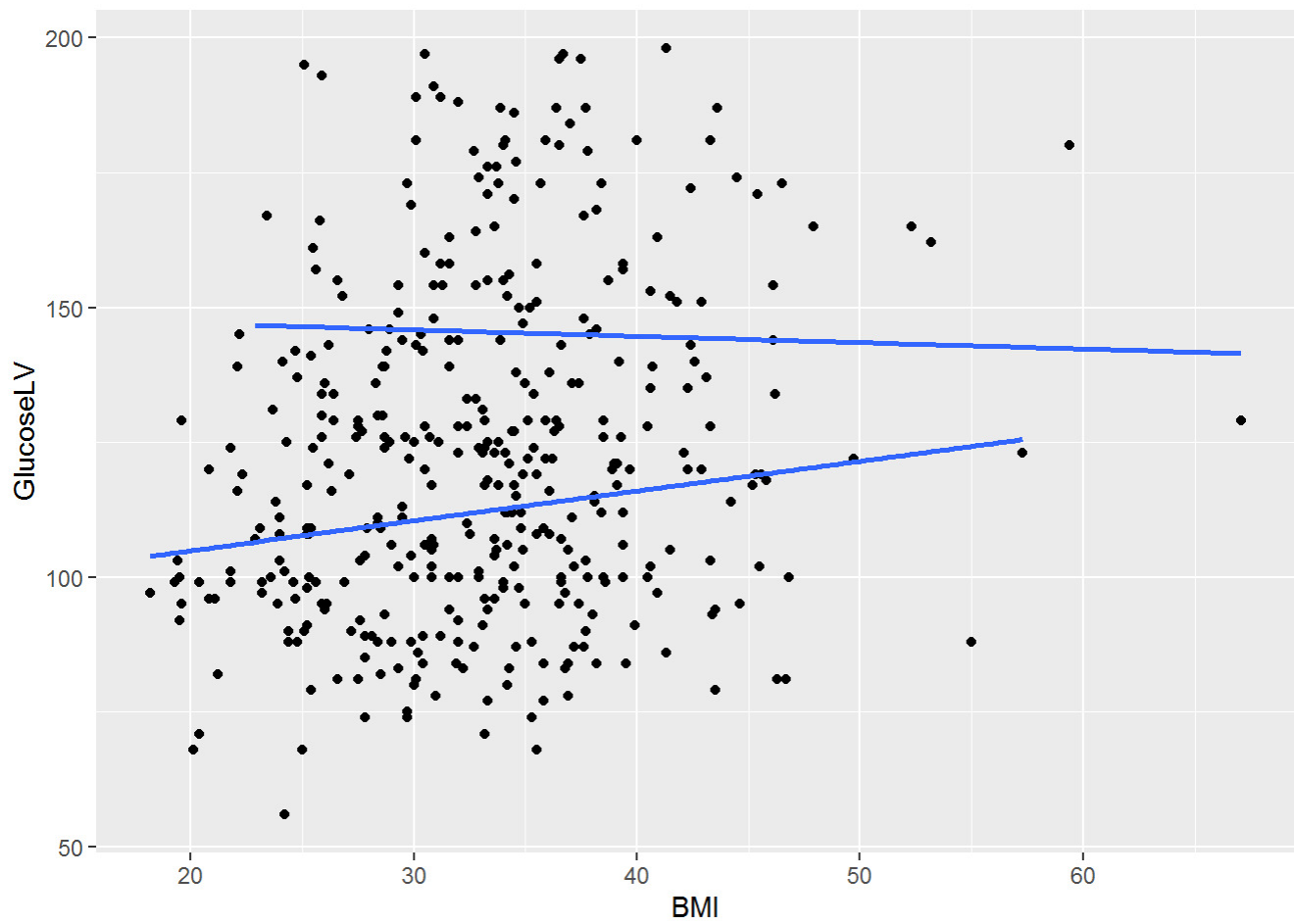
```
View(pima.indian.diabetes2)
ggplot(pima.indian.diabetes2, aes(x=Age, y=BMI, col=factor(Classification)))+geom_point()
```



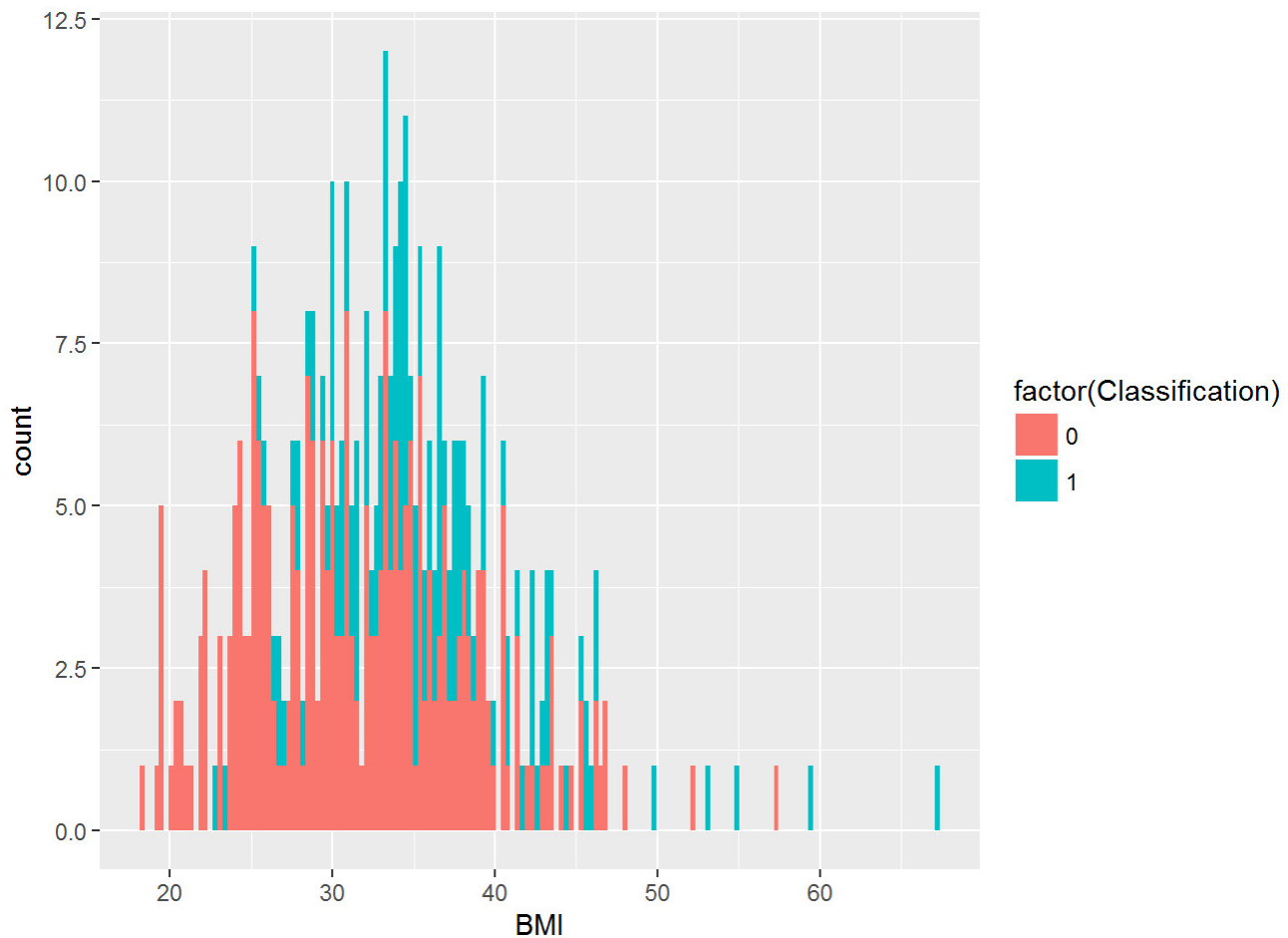
```
ggplot(pima.indian.diabetes2, aes(x=BMI, y=GlucoseLV, col= factor(Classification)))+geom_point()+geom_smooth(method="lm", se=FALSE)
```



```
ggplot(pima.indian.diabetes2, aes(x=BMI, y=GlucoseLV, Col=factor(Classification))) + geom_point() + geom_smooth(method="lm", se=FALSE)
```



```
ggplot(pima.indian.diabetes2, aes(x=BMI, fill=factor(Classification))) + geom_histogram(b  
inwidth = 0.3)
```



```
summary(pima.indian.diabetes2)
```

```
## TimesPregnant      GlucoseLV      DiastolicBP      TricepsThickness
## Min.   : 0.000      Min.   : 56.0      Min.   : 24.00      Min.   : 7.00
## 1st Qu.: 1.000      1st Qu.: 99.0      1st Qu.: 62.00      1st Qu.:21.00
## Median : 2.000      Median :119.0      Median : 70.00      Median :29.00
## Mean   : 3.301      Mean   :122.6      Mean   : 70.66      Mean   :29.15
## 3rd Qu.: 5.000      3rd Qu.:143.0      3rd Qu.: 78.00      3rd Qu.:37.00
## Max.   :17.000      Max.   :198.0      Max.   :110.00      Max.   :63.00
## SerumInsulin      BMI      Heridarymarkup      Age
## Min.   : 14.00      Min.   :18.20      Min.   :0.0850      Min.   :21.00
## 1st Qu.: 76.75      1st Qu.:28.40      1st Qu.:0.2697      1st Qu.:23.00
## Median :125.50      Median : 33.20      Median :0.4495      Median :27.00
## Mean   :156.06      Mean   : 33.09      Mean   :0.5230      Mean   :30.86
## 3rd Qu.:190.00      3rd Qu.:37.10      3rd Qu.:0.6870      3rd Qu.:36.00
## Max.   :846.00      Max.   : 67.10      Max.   :2.4200      Max.   :81.00
## Classification
## Min.   :0.0000
## 1st Qu.:0.0000
## Median :0.0000
## Mean   :0.3316
## 3rd Qu.:1.0000
## Max.   :1.0000
```

```
library(dplyr)
```

```
## Warning: package 'dplyr' was built under R version 3.2.5
```

```
##  
## Attaching package: 'dplyr'
```

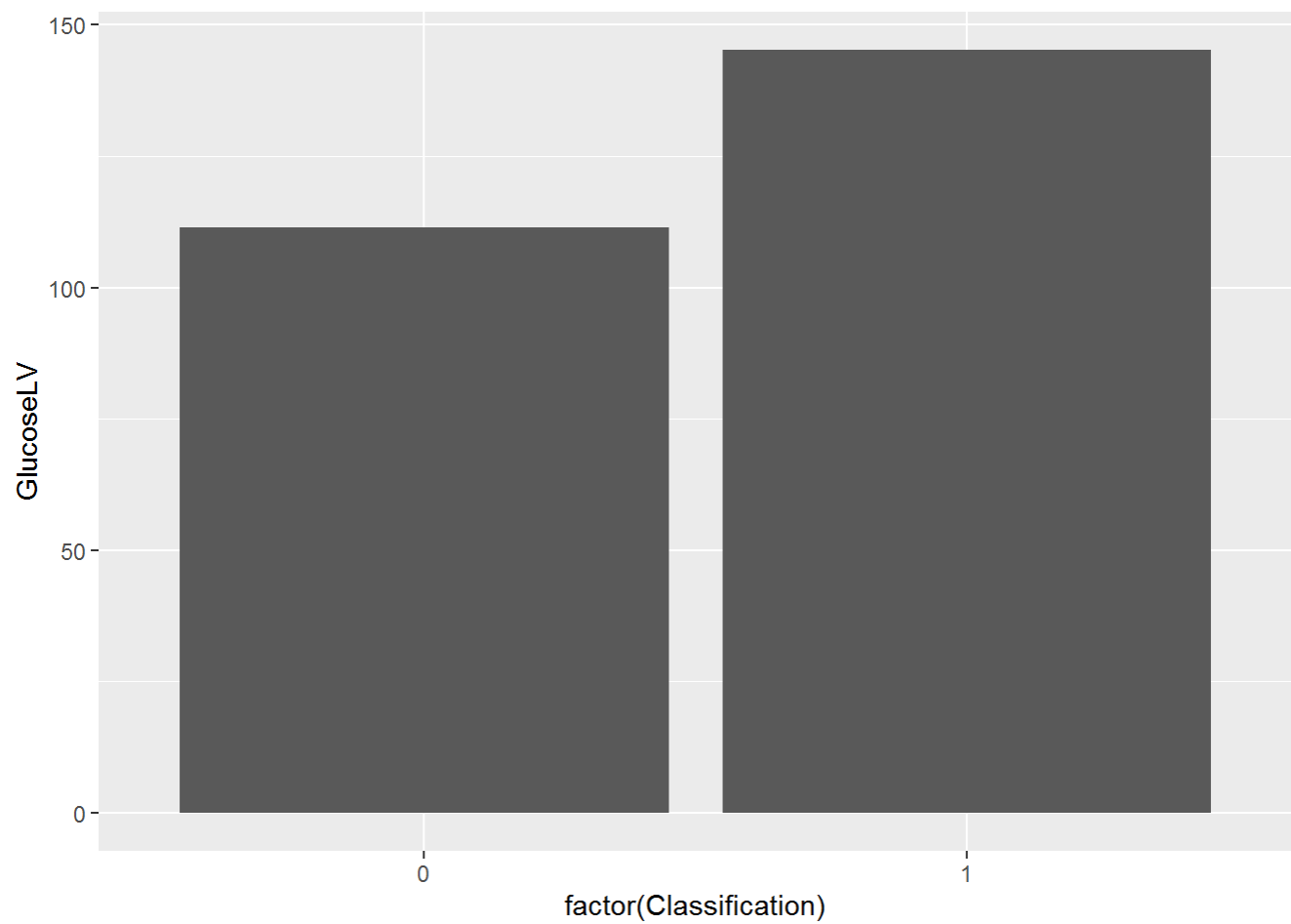
```
## The following objects are masked from 'package:stats':  
##  
##   filter, lag
```

```
## The following objects are masked from 'package:base':  
##  
##   intersect, setdiff, setequal, union
```

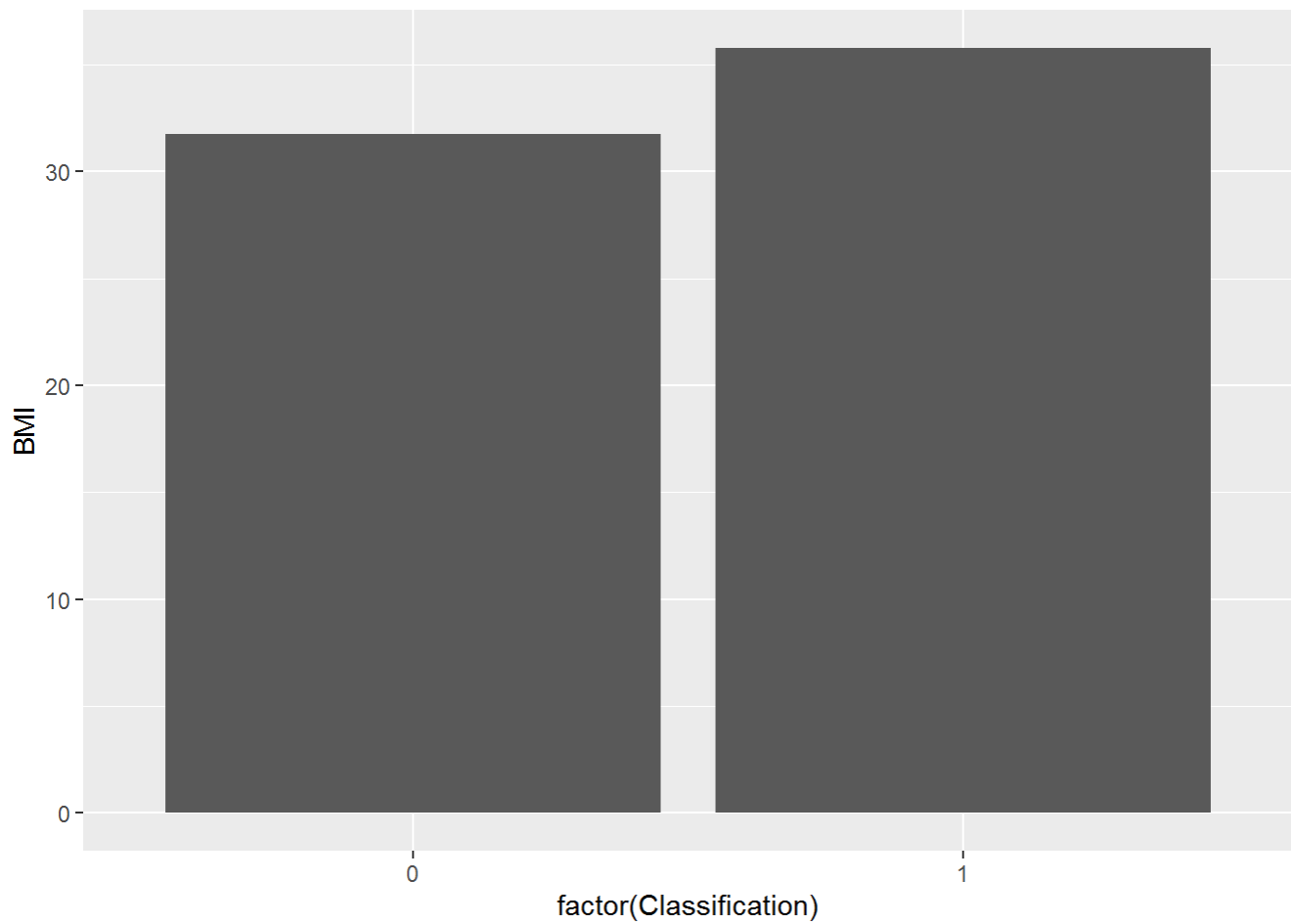
```
pima.indian.diabetes3<-pima.indian.diabetes2 %>% group_by(Classification)%>% summaris  
e_each(funs(mean),GlucoseLV,DiastolicBP,SerumInsulin,BMI,TricepsThickness,Age)  
View(pima.indian.diabetes3)  
str(pima.indian.diabetes3)
```

```
## Classes 'tbl_df', 'tbl' and 'data.frame':   2 obs. of  7 variables:  
## $ Classification : int  0 1  
## $ GlucoseLV : num  111 145  
## $ DiastolicBP : num  69 74.1  
## $ SerumInsulin : num  131 207  
## $ BMI : num  31.8 35.8  
## $ TricepsThickness: num  27.3 33  
## $ Age : num  28.3 35.9  
## - attr(*, "na.action")=Class 'omit' Named int [1:376] 1 2 3 6 8 10 11 12 13 16 .  
..  
## .. ..- attr(*, "names")= chr [1:376] "1" "2" "3" "6" ...
```

```
ggplot(pima.indian.diabetes3,aes(x=factor(Classification),y=GlucoseLV))+geom_bar(stat  
="identity")
```



```
ggplot(pima.indian.diabetes3, aes(x=factor(Classification), y=BMI)) + geom_bar(stat="identity")
```



```
ggplot(pima.indian.diabetes2, aes(x=GlucoseLV, fill=Classification)) + geom_bar(position="fill")
```