

## subject6

```
data = read.csv(file = 'Subject6_smoking.csv')
head(data,5)
```

```
##   Smoker Status  Age
## 1   Yes  Alive 21.0
## 2   Yes  Alive 19.3
## 3    No   Dead 57.5
## 4    No  Alive 47.1
## 5   Yes  Alive 81.4
```

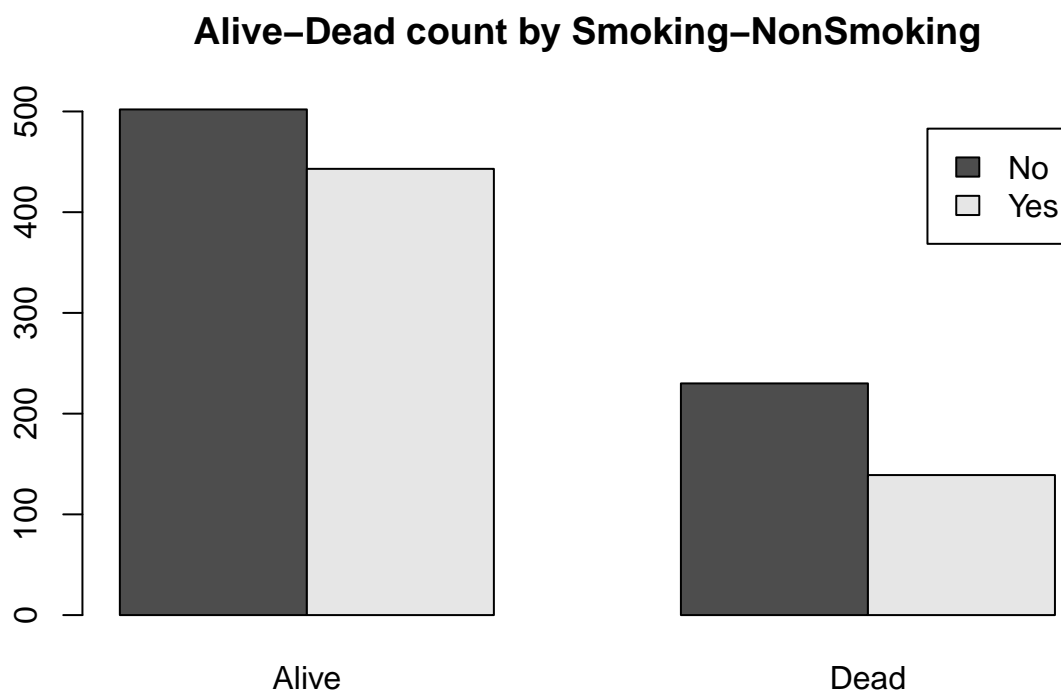
### 1

Two way table between if they are alive or dead and if yes or no they smoke

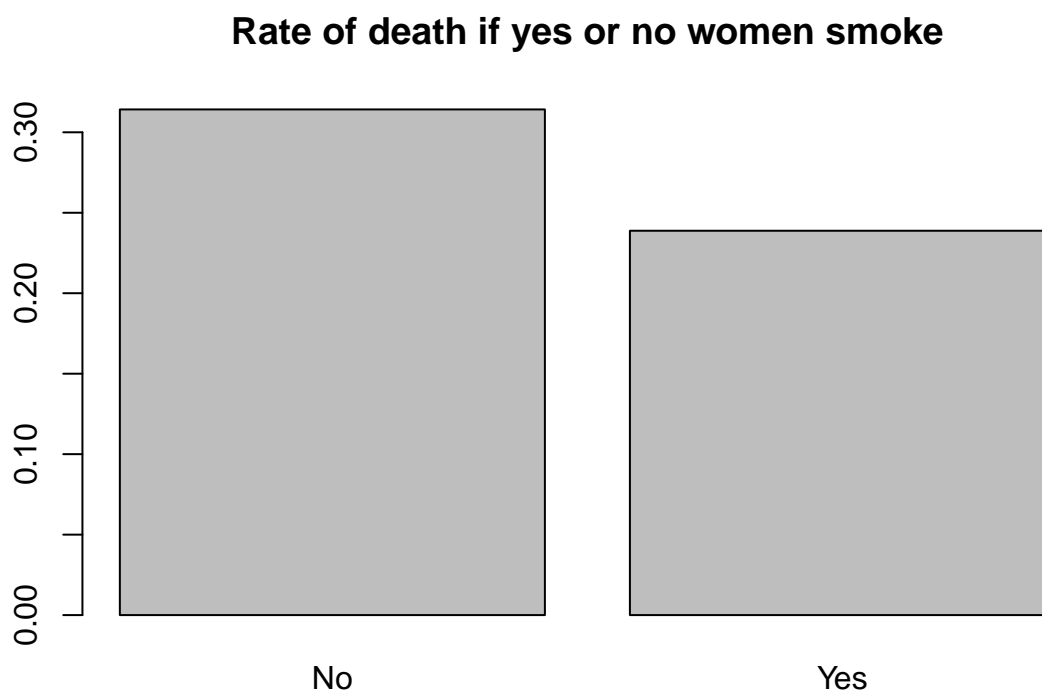
```
datatable = table(data$Smoker, data$Status)
addmargins(datatable)
```

```
##
##      Alive Dead  Sum
## No      502  230  732
## Yes     443  139  582
## Sum     945  369 1314
```

```
barplot(datatable, legend=TRUE, beside=TRUE, main='Alive-Dead count by Smoking-NonSmoking')
```



```
#rate of death in the smoking and non smoking categories  
rate_vector = (addmargins(datatable)[, 'Dead'] / addmargins(datatable)[, 'Sum'])[c('No', 'Yes')]  
barplot(rate_vector, main="Rate of death if yes or no women smoke")
```



## Explanation

We can observe that the rate of mortality is higher in the women that didn't smoke and that is suprising because we would expect the contrary.

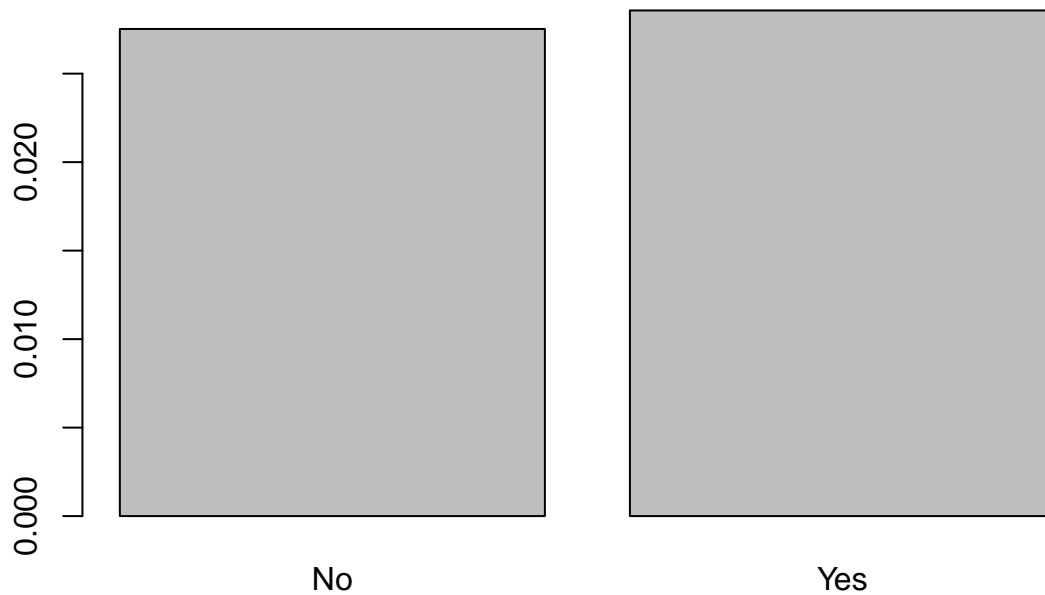
## 2

```
data_18_34 = data[data$Age > 18 & data$Age < 34, ]
datatable_18_34 = addmargins(table(data_18_34$Smoker, data_18_34$Status))
datatable_18_34
```

```
##
##      Alive Dead Sum
##   No    212    6 218
##   Yes   170    5 175
##   Sum   382   11 393
```

```
rate_vector_18_34 = (datatable_18_34[, 'Dead'] / datatable_18_34[, 'Sum'])[c('No', 'Yes')]
barplot(rate_vector_18_34, main="Rate of death for women between 18 and 34 if yes or no they smoke")
```

## Rate of death for women between 18 and 34 if yes or no they smoke

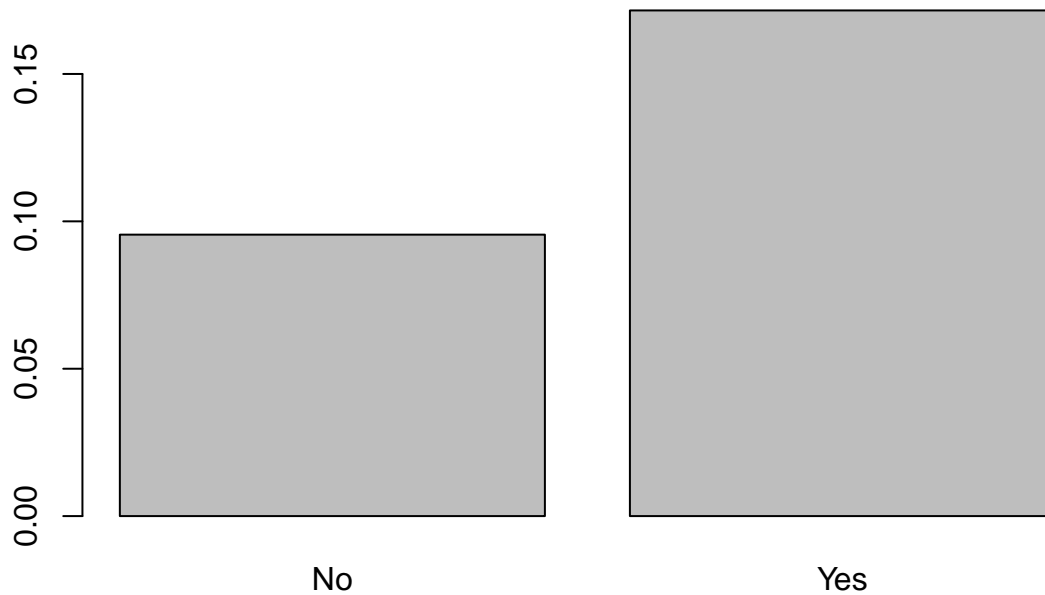


```
data_34_54 = data[data$Age >= 34 & data$Age < 54, ]
datatable_34_54 = addmargins(table(data_34_54$Smoker, data_34_54$Status))
datatable_34_54
```

```
##
##      Alive Dead Sum
##   No    180   19 199
##   Yes   198   41 239
##   Sum   378   60 438
```

```
rate_vector_34_54 = (datatable_34_54[, 'Dead'] / datatable_34_54[, 'Sum'])[c('No', 'Yes')]
barplot(rate_vector_34_54, main="Rate of death for women between 34 and 54 if yes or no they smoke")
```

## Rate of death for women between 34 and 54 if yes or no they smoke



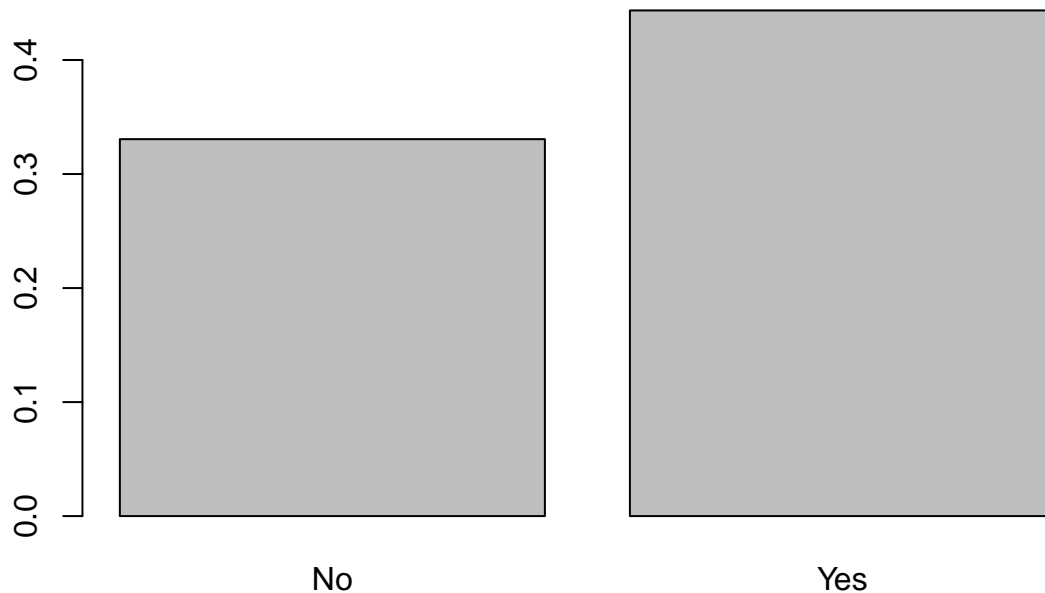
Women who smoke die more in this category.

```
data_54_65 = data[data$Age >= 54 & data$Age < 65, ]
datatable_54_65 = addmargins(table(data_54_65$Smoker, data_54_65$Status))
datatable_54_65
```

```
##
##      Alive Dead Sum
##   No      81   40 121
##   Yes      64   51 115
##   Sum     145   91 236
```

```
rate_vector_54_65 = (datatable_54_65[, 'Dead'] / datatable_54_65[, 'Sum'])[c('No', 'Yes')]
barplot(rate_vector_54_65, main="Rate of death for women between 54 and 65 if yes or no they smoke")
```

## Rate of death for women between 54 and 65 if yes or no they smoke



Women who smoke die more in this category.

```
data_65 = data[data$Age >= 65, ]
datatable_65 = addmargins(table(data_65$Smoker, data_65$Status))
datatable_65
```

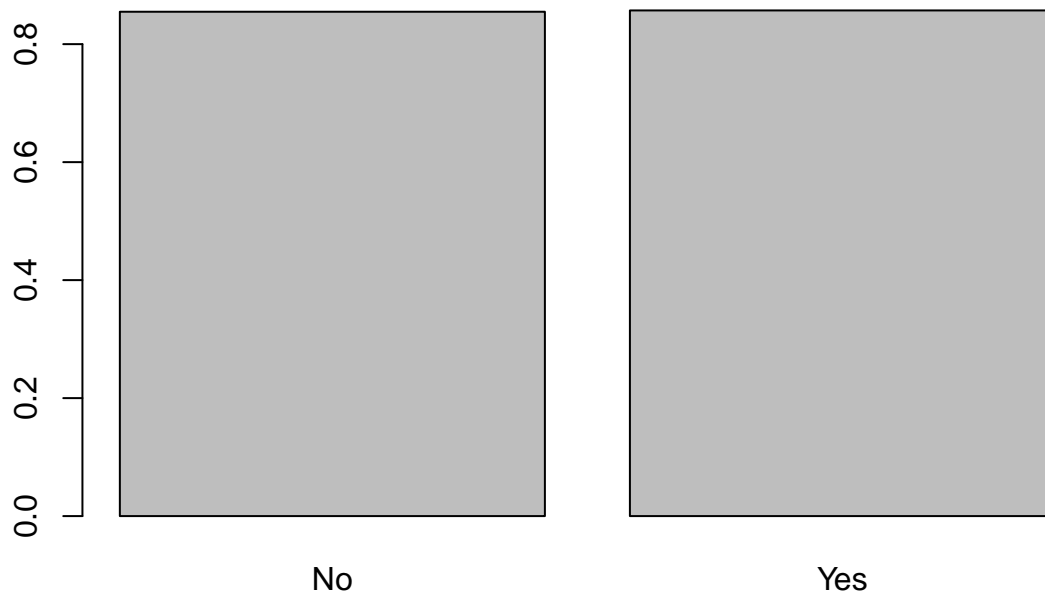
```
##
##      Alive Dead Sum
##   No      28  165 193
##   Yes       7   42  49
##   Sum      35  207 242
```

## Explanations

We can observe that for women older than 65, there are much more deaths in the category that don't smoke. This makes the data unbalanced. This high number of deaths brings the total rate as observed previously to a high number.

```
rate_vector_65 = (datatable_65[, 'Dead'] / datatable_65[, 'Sum'])[c('No', 'Yes')]
barplot(rate_vector_65, main="Rate of death for women of more than 65 if yes or no they smoke")
```

## Rate of death for women of more than 65 if yes or no they smoke



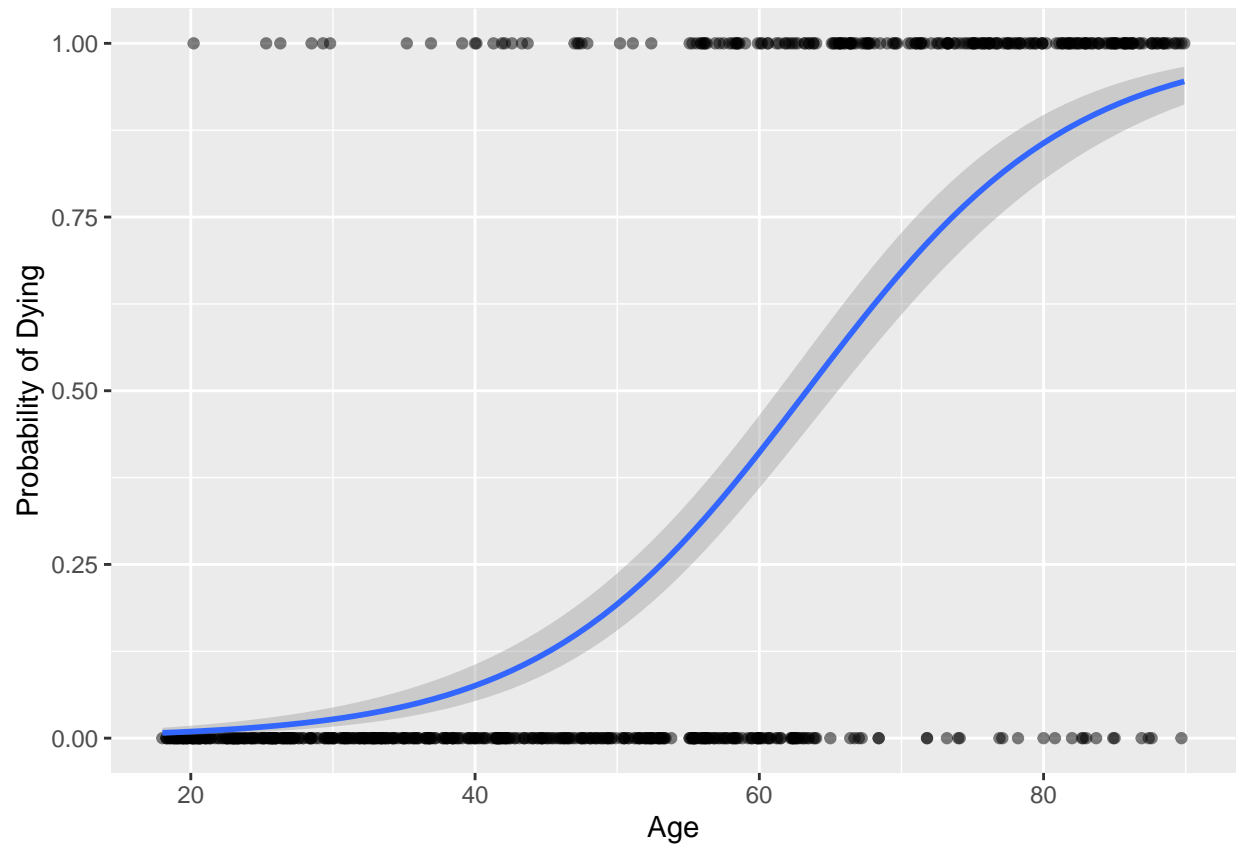
The relative rate in this category is similar thought.

### 3

```
#change Alive and Dead in 0 and 1 in the Status column
data1 <- data %>%
  mutate(Status = ifelse(Status == "Alive",0,1))

data_no = data1[data1$Smoker == 'No',]
#plot logistic regression curve
ggplot(data_no, aes(x=Age, y=Status)) +
  geom_point(alpha=.5) +
  stat_smooth(method="glm", method.args = list(family=binomial))+
  labs(
    y = "Probability of Dying"
  )
```

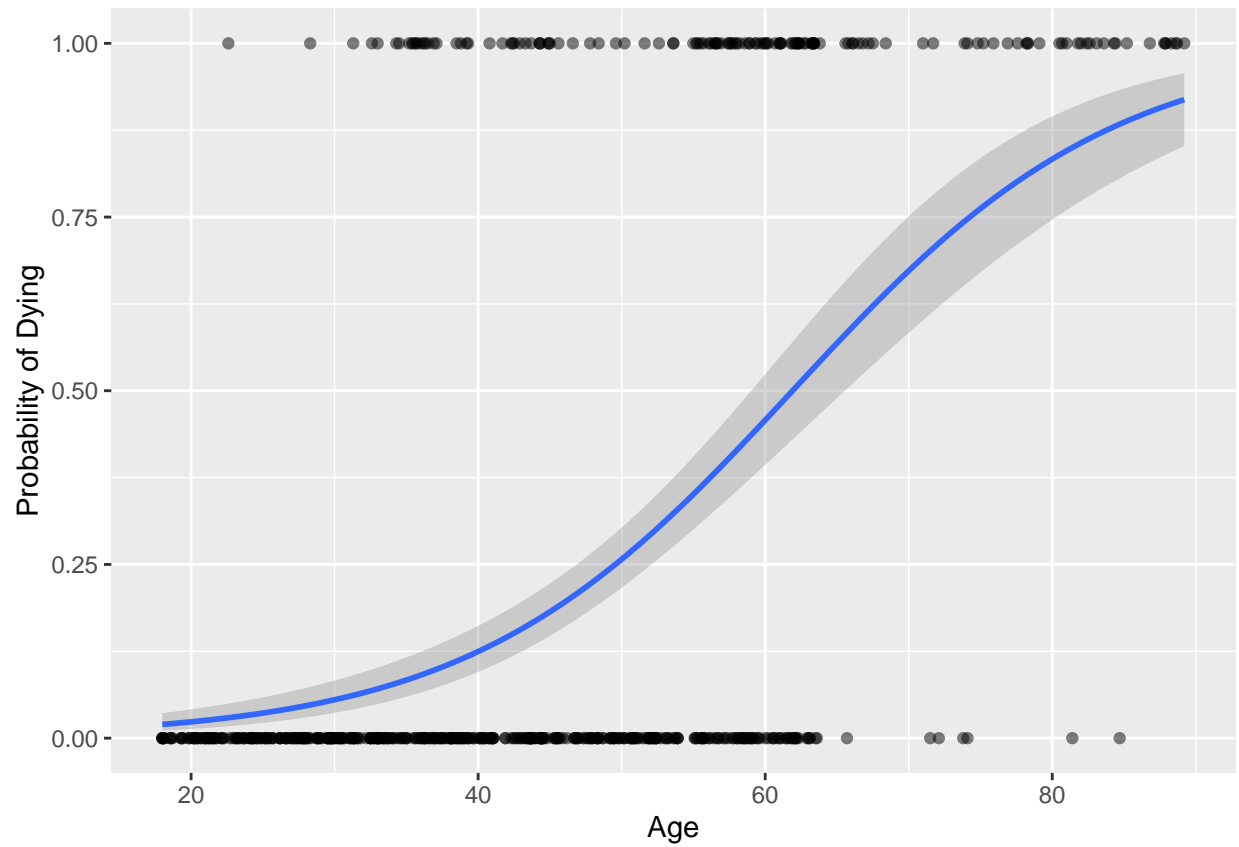
```
## 'geom_smooth()' using formula 'y ~ x'
```



```
data_yes = data1[data1$Smoker == 'Yes',]  
#plot logistic regression curve  
ggplot(data_yes, aes(x=Age, y=Status)) +  
  geom_point(alpha=.5) +  
  stat_smooth(method="glm", method.args = list(family=binomial)) +  
  labs(  
    y = "Probability of Dying"  
  )
```

```
## 'geom_smooth()' using formula 'y ~ x'
```





## Explanation

We can observe that the probability of death increase more quickly for the womens that smoke