

Final Report

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Load Packages

```
library(tidyverse)
```

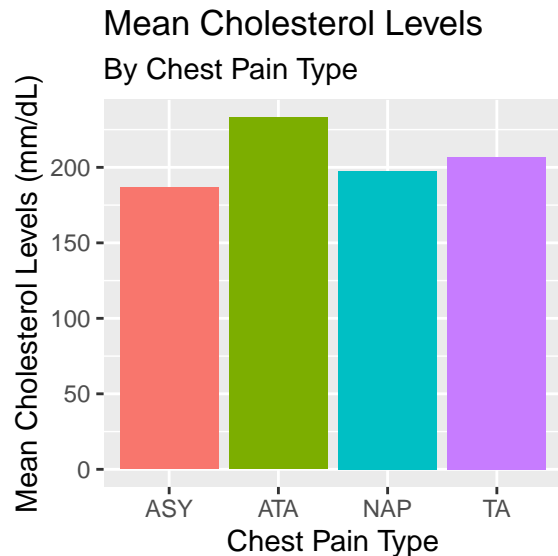
Load Data

```
heart <- readr::read_csv("heart.csv")
```

```
mean_cholesterol <- heart %>%  
  group_by(ChestPainType) %>%  
  summarize(mean_cholesterol = mean(Cholesterol))%>%  
  print()
```

```
## # A tibble: 4 x 2  
##   ChestPainType mean_cholesterol  
##   <chr>          <dbl>  
## 1 ASY           187.  
## 2 ATA           233.  
## 3 NAP           197.  
## 4 TA            207.
```

```
mean_cholesterol %>%  
  ggplot()+  
  geom_col(mapping = aes(x = ChestPainType, y = mean_cholesterol, fill = ChestPainType), position = "dodge")+  
  theme(legend.position = "none")+  
  labs(title = "Mean Cholesterol Levels",  
        subtitle = "By Chest Pain Type",  
        x = "Chest Pain Type",  
        y = "Mean Cholesterol Levels (mm/dL)")
```



```
heart_grouped <- heart %>%
  mutate(chol_level = cut(Cholesterol,
                        breaks = c(-Inf, 120, 200, 239, Inf),
                        labels = c("Low", "Normal", "Intermediate", "High"),
                        right=FALSE))
```

```
heart_grouped <- heart_grouped %>%
  mutate(press_level = cut(RestingBP,
                        breaks = c(-Inf, 120, 130, 140, 180, Inf),
                        labels = c("Normal", "Elevated", "Hypertension 1", "Hypertension 2", "Hyperten"),
                        right=FALSE))
```

```
heart_grouped <- heart_grouped %>%
  mutate(Sex, sex_factor=ifelse(Sex=="M", 0,1)) %>%
  mutate(ExerciseAngina, exer_factor=ifelse(ExerciseAngina=="N", 0,1))
```

```
chol_grouped <- heart_grouped %>%
  group_by(ChestPainType, chol_level)%>%
  summarize(count = n())
```

`summarise()` has grouped output by 'ChestPainType'. You can override using the `.groups` argument.

```
RBP_grouped <- heart_grouped %>%
  group_by(ChestPainType, press_level) %>%
  summarize(count = n())
```

`summarise()` has grouped output by 'ChestPainType'. You can override using the `.groups` argument.

```
exer_grouped <- heart_grouped %>%
  group_by(ChestPainType, exer_factor) %>%
  summarize(count = n())
```

`summarise()` has grouped output by 'ChestPainType'. You can override using the `.groups` argument.

```
sex_grouped <- heart_grouped %>%
  group_by(ChestPainType, sex_factor) %>%
  summarize(count = n())
```

`summarise()` has grouped output by 'ChestPainType'. You can override using the `.groups` argument.