table 1

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```r
cancer <- read.csv("desktop/cancer csv.csv")
#view(cancer)
#summary(cancer)
Mean_area <- cancer$Mean_Area
Class_grade <- cancer$ClassGrade
ER <- cancer$ER
Mean_mean_R <- cancer$Mean_mean_R
Mean_mean_G <- cancer$Mean_mean_G
Mean_mean_B <- cancer$Mean_mean_B
Mean_mean_HSV <- cancer$Mean_mean_HSV
Mean_mean_Lab <- cancer$Mean_mean_Lab
Mean_mean_HE <- cancer$Mean_mean_HE
Mean_mean_BR <- cancer$Mean_mean_BR
mean(Mean_area)
```

```
## [1] 659.8554
```

```r
sd(Mean_area)
```

```
## [1] 152.3151
```

```r
range(Mean_area)
```

```
## [1]   375.5158 1419.0733
```

```r
mean(Mean_mean_R)
```

```
## [1] 101.5842
```

```r
sd(Mean_mean_R)
```

```
## [1] 11.32527
```

```r
range(Mean_mean_R)
```

```
## [1]   74.96138 135.61285
```

```r
mean(Mean_mean_G)
```

```
## [1] 73.90454
```

```r
sd(Mean_mean_G)
```

```
## [1] 12.42109
```

```r
range(Mean_mean_G)
```

```
## [1]  47.49451 110.67578
```

```r
mean(Mean_mean_B)
```

```
## [1] 94.72136
```

```r
sd(Mean_mean_B)
```

```
## [1] 10.84549
```

```r
range(Mean_mean_B)
```

```
## [1]  72.08346 128.92483
```

```r
mean(Mean_mean_HSV)
```

```
## [1] 90.31242
```

```r
sd(Mean_mean_HSV)
```

```
## [1] 13.5191
```

```r
range(Mean_mean_HSV)
```

```
## [1]  64.13022 131.27174
```

```r
mean(Mean_mean_Lab)
```

```
## [1] 86.01685
```

```r
sd(Mean_mean_Lab)
```

```
## [1] 12.15921
```

```r
range(Mean_mean_Lab)
```

```
## [1]   62.90402 121.67888
```

```r
mean(Mean_mean_HE)
```

```
## [1] 135.9635
```

```r
sd(Mean_mean_HE)
```

```
## [1] 25.31569
```

```r
range(Mean_mean_HE)
```

```
## [1]   65.96322 192.18241
```

```r
mean(Mean_mean_BR)
```

```
## [1] 7.98594
```

```r
sd(Mean_mean_BR)
```

```
## [1] 11.35399
```

```r
range(Mean_mean_BR)
```

```
## [1]   0.00000 67.65289
```