

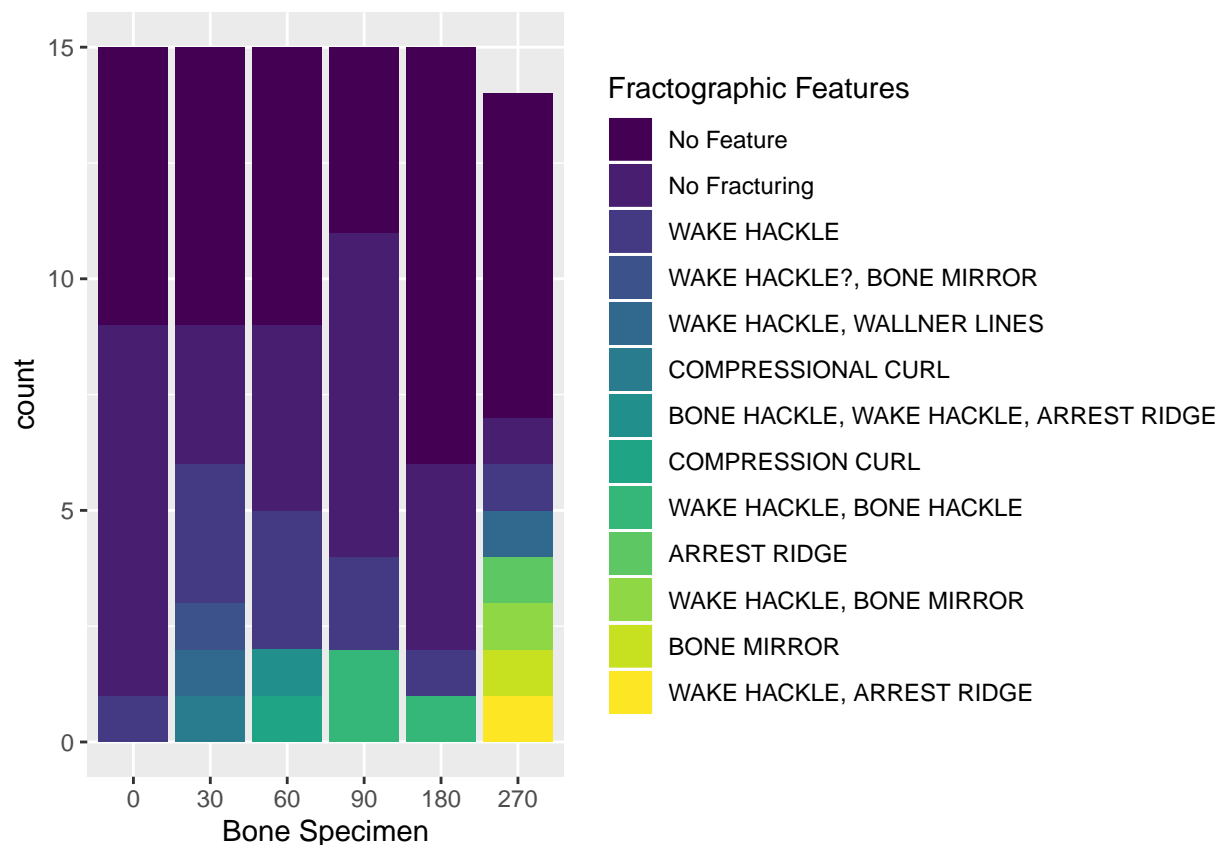
inital EDA

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EDA

```
ggplot(data = bone) + geom_bar(mapping = aes(x = `Bone Specimen`, fill = `Fractographic Features`))
```



Models

```
fit1<-glm(bone$`Bisecting Fracture`~bone$days,family = binomial())  
fit2<-glm(bone$`Fracture-like`~bone$days,family = binomial())  
summary(fit1)
```

```
##  
## Call:  
## glm(formula = bone$`Bisecting Fracture` ~ bone$days, family = binomial())  
##  
## Deviance Residuals:  
##      Min       1Q   Median       3Q      Max   
## -1.3753  -1.2920   0.9917   1.0670   1.1187
```

```
##
## Coefficients:
##           Estimate Std. Error z value Pr(>|z|)
## (Intercept)  0.07685    0.48625   0.158   0.874
## bone$days   0.06285    0.12664   0.496   0.620
##
## (Dispersion parameter for binomial family taken to be 1)
##
##      Null deviance: 121.47  on 88  degrees of freedom
## Residual deviance: 121.23  on 87  degrees of freedom
## AIC: 125.23
##
## Number of Fisher Scoring iterations: 4
summary(fit2)

##
## Call:
## glm(formula = bone$`Fracture-like` ~ bone$days, family = binomial())
##
## Deviance Residuals:
##      Min       1Q   Median       3Q      Max
## -1.8587  -1.2733   0.7036   0.8810   1.0844
##
## Coefficients:
##           Estimate Std. Error z value Pr(>|z|)
## (Intercept) -0.03916    0.51010  -0.077   0.9388
## bone$days   0.26179    0.14229   1.840   0.0658 .
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
##      Null deviance: 109.24  on 88  degrees of freedom
## Residual deviance: 105.69  on 87  degrees of freedom
## AIC: 109.69
##
## Number of Fisher Scoring iterations: 4
```