```
#Check for assumptions
plot(fit_c, which =1)
```
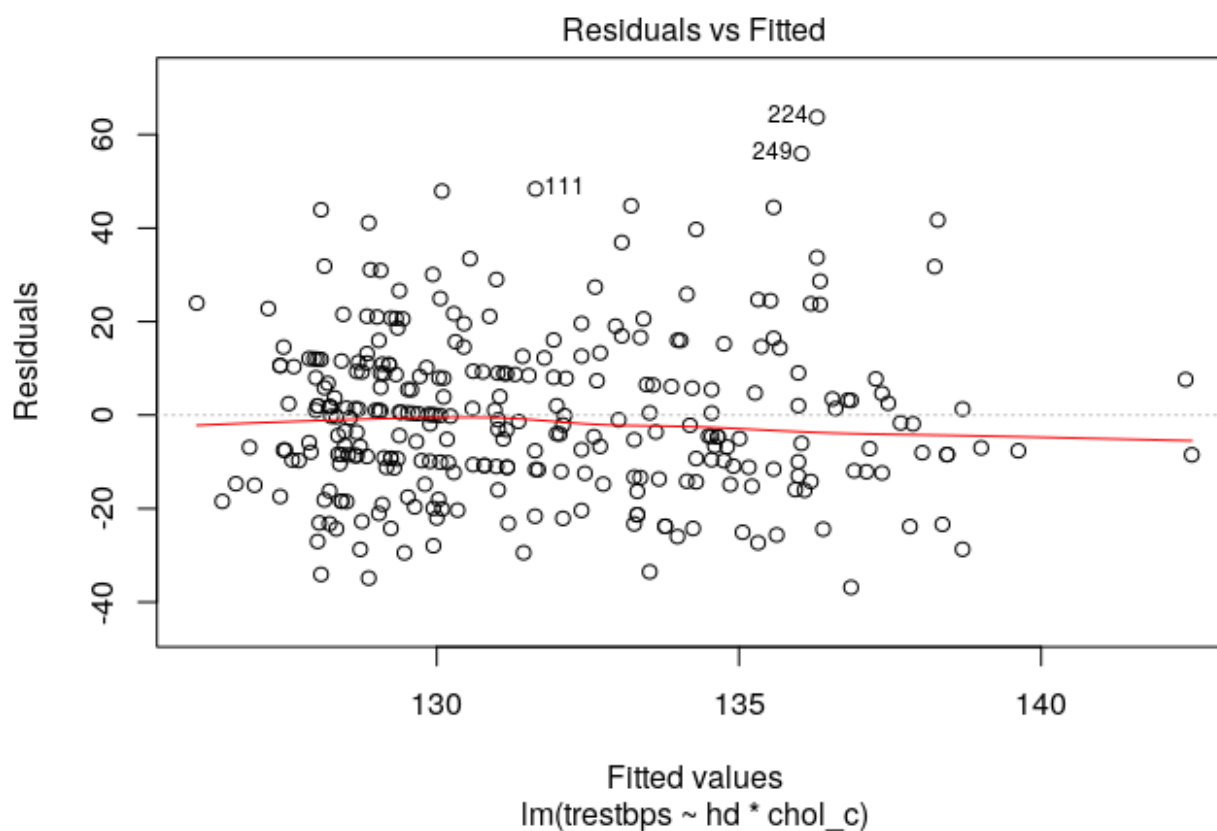


```
#Histogram of residuals '
hist(fit_c$residuals)
```