

```
PFs <- read.delim("~/Documents/transcriptomic-exercise/output/annotation/PFsofAnnotation.txt",  
quote="\"", comment.char="", header = F, as.is = T)  
sort(PFs[, 1])  
hist(table(PFs[, 1]), main = "Frecuencia de Pfams")
```

Frecuencia de Pfams

