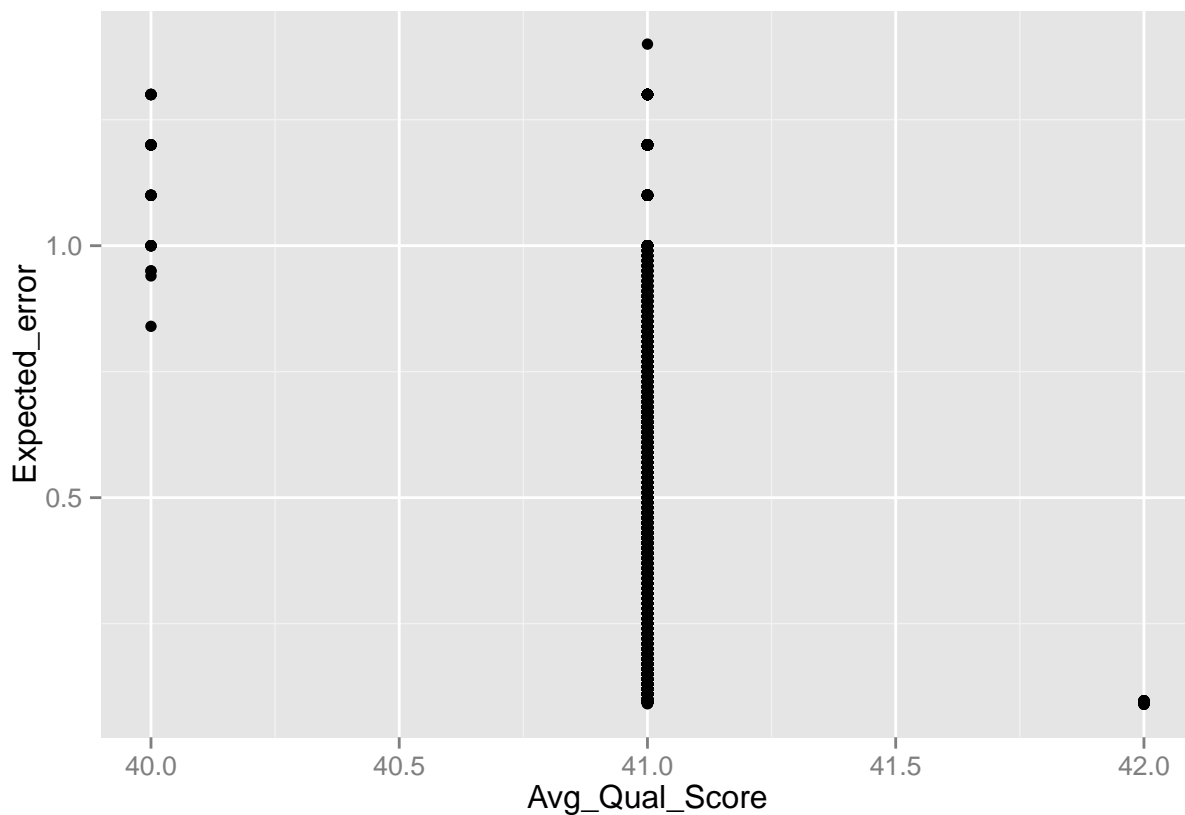# Mock Community from Pacific Biosciences (BEI)

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This is an R Markdown document. Markdown is a simple formatting syntax for authoring HTML, PDF, and MS Word documents. For more details on using R Markdown see http://rmarkdown.rstudio.com.

When you click the **Knit** button a document will be generated that includes both content as well as the output of any embedded R code chunks within the document. You can embed an R code chunk like this:

```r
library(ggplot2)
test <- read.delim("~/workspace/training/joshs_tasks/BEI_Mock_Community_Stats.txt")
qplot(data=test,y=Expected_error, x=Avg_Qual_Score, geom='point')
```


You can also embed plots, for example:

Note that the `echo = FALSE` parameter was added to the code chunk to prevent printing of the R code that generated the plot.