Class16

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Q. What does the star character accomplish here? Ask Barry, or your class neighbor, if you are not sure!

a powerful tool in shell commands to match multiple files or directories based on a pattern, in this case the .faa.gz

- Q. How many sequences are in this mouse. 1.protein.faa file? Hint: Try using grep to figure this out...
- Q. What happens if you run the above command without the > mm-first.fa part?

If you run the command without the > mm-first.fa part, the first 11 lines of the mouse.1.protein.faa file will be printed directly to your terminal.

Q. What happens if you were to use two '>' symbols (i.e. "> mm-first.fa)?

Adds the new output to the end of the existing file.

Reading Table

```
b <- read.table("myresults.tsv", sep="\t", header=TRUE)
head(data)

##

## 1 function (..., list = character(), package = NULL, lib.loc = NULL,
## 2 verbose = getOption("verbose"), envir = .GlobalEnv, overwrite = TRUE)

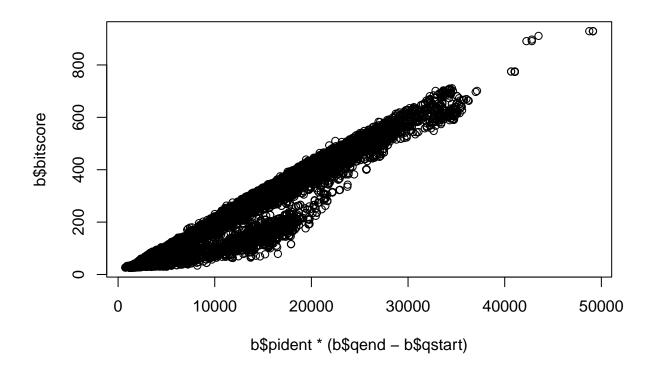
## 3 {
## 4 fileExt <- function(x) {
## 5 db <- grepl("\\\\.[^.]+\\\\.(gz|bz2|xz)$", x)
## 6 ans <- sub(".*\\\.", "", x)</pre>
```

Setting Colnames

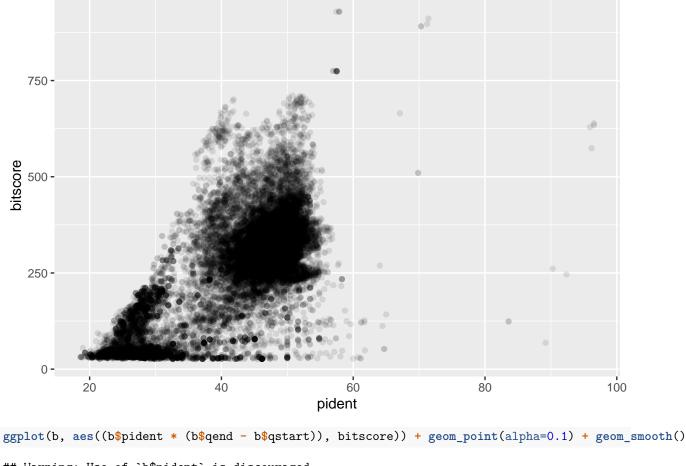
```
colnames(b) <- c("qseqid", "sseqid", "pident", "length", "mismatch", "gapopen", "qstart", "qend", "ssta</pre>
```

Plot() Function

```
plot(b$pident * (b$qend - b$qstart), b$bitscore)
```



```
# GGPlot
library(ggplot2)
ggplot(b, aes(pident, bitscore)) + geom_point(alpha=0.1)
```



```
## Warning: Use of `b$pident` is discouraged.
## i Use `pident` instead.
## Warning: Use of `b$qend` is discouraged.
## i Use `qend` instead.
## Warning: Use of `b$qstart` is discouraged.
## i Use `qend` instead.
## Warning: Use of `b$qstart` is discouraged.
## i Use `qitart` instead.
## Warning: Use of `b$pident` is discouraged.
## i Use `pident` instead.
## Warning: Use of `b$qend` is discouraged.
## i Use `qend` instead.
## Warning: Use of `b$qstart` is discouraged.
## i Use `qend` instead.
## Warning: Use of `b$qstart` is discouraged.
## i Use `qend` instead.
## warning: Use of `b$qstart` is discouraged.
## i Use `qstart` instead.
## `geom_smooth()` using method = 'gam' and formula = 'y ~ s(x, bs = "cs")'
```

