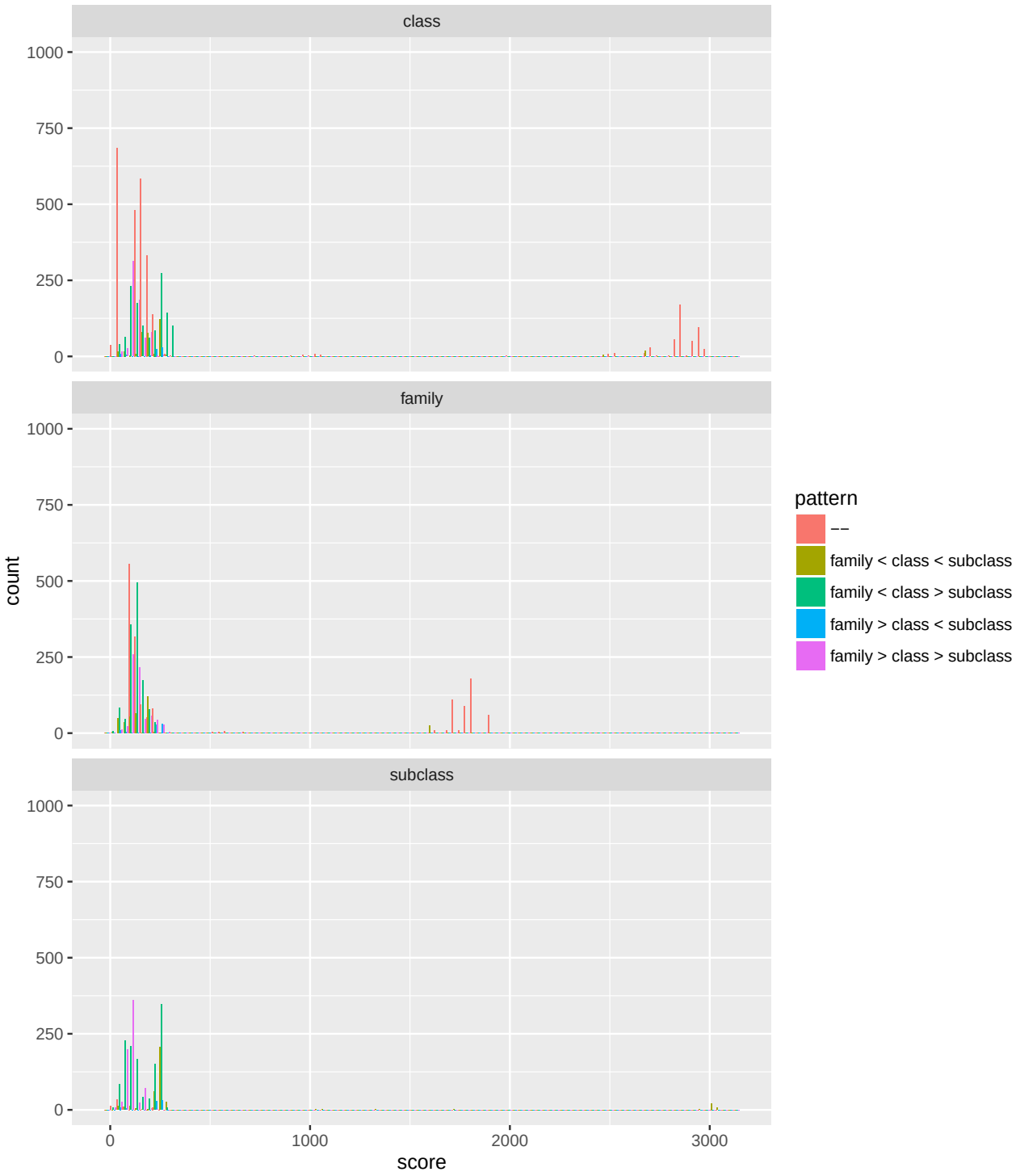
Pfitmap-eval: Distribution of scores per sequence and profile

daniel.lundin@dbb.su.se

2016-04-16

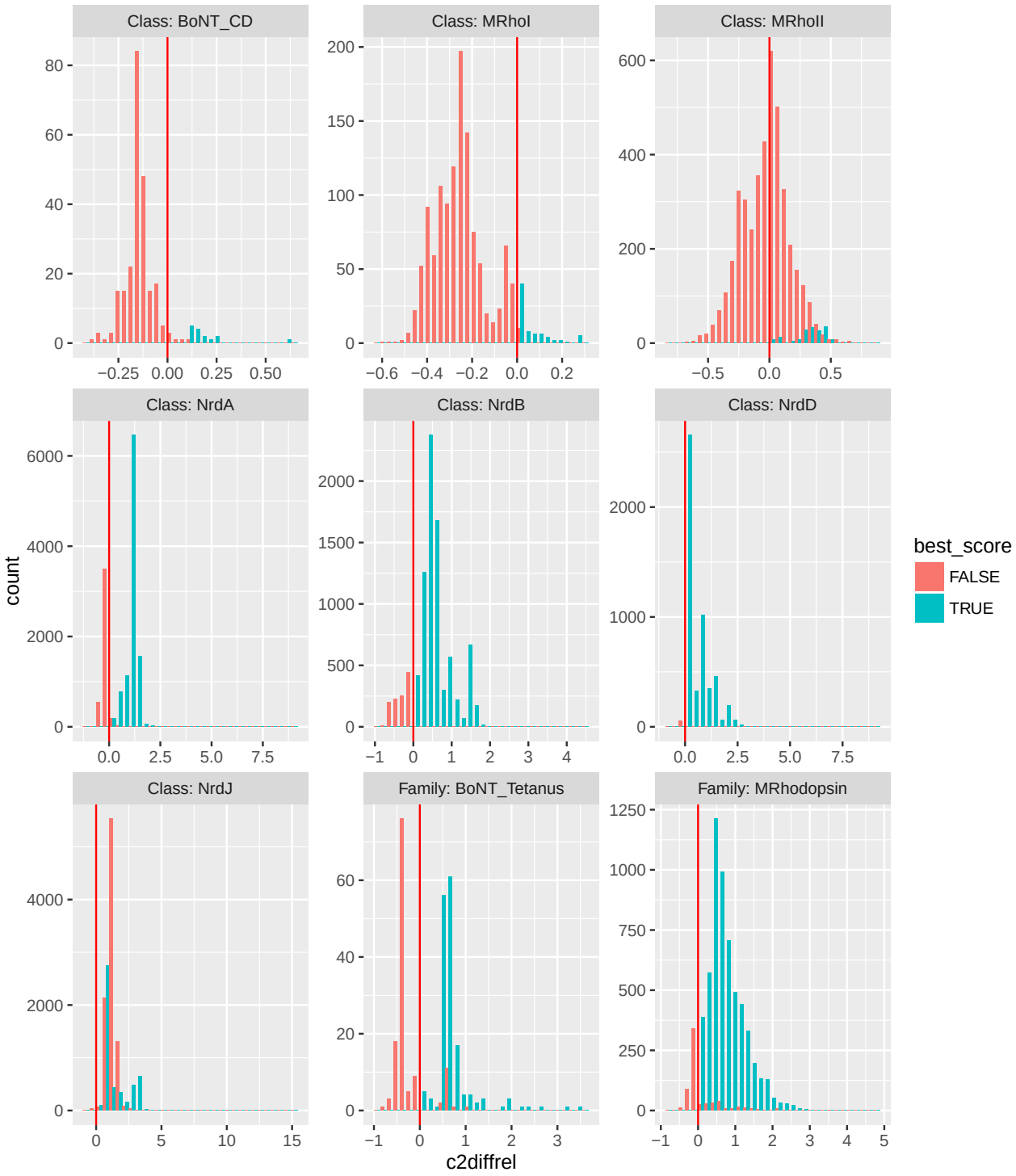
Contents

```
## Warning: Removed 10137 rows containing non-finite values (stat_bin).
```




```
## `stat_bin()` using `bins = 30`. Pick better value with `binwidth`.
```

```
## Warning: Removed 2 rows containing non-finite values (stat_bin).
```


Class: BoNT_CD
Class: MRhoI
Class: MRhoII
Class: NrdA
Class: NrdB
Class: NrdD
Class: NrdJ
Family: BoNT_Tetanus
Family: MRhodopsin
count
c2diffrel
best_score
FALSE
TRUE