

Semi-quantification

```
## # A tibble: 38 x 3
##   Compound      Homologue      pattern_CF2CF2
##   <chr>         <chr>         <chr>
## 1 11C1-PF30UdS  9C1-PF30NS    smaller
## 2 4:2 FTS      6:2 FTS      bigger
## 3 6:2 diPAP    6:2/8:2 diPAP bigger
## 4 6:2 FTS      4:2 FTS      smaller
## 5 6:2 FTS      8:2 FTS      bigger
## 6 6:2/8:2 diPAP 6:2 diPAP    smaller
## 7 6:2/8:2 diPAP 8:2 diPAP    bigger
## 8 8:2 diPAP    6:2/8:2 diPAP smaller
## 9 8:2 FTS      6:2 FTS      smaller
## 10 9C1-PF30NS   11C1-PF30UdS bigger
## # ... with 28 more rows
```