

Results of Ordered Likelihood Analysis

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```
library(here)

## here() starts at /home/devan/OneDriveUW0/0postdoc/sup
library(dplyr)

##
## Attaching package: 'dplyr'
##
## The following objects are masked from 'package:stats':
##
##   filter, lag
##
## The following objects are masked from 'package:base':
##
##   intersect, setdiff, setequal, union

library(tidyr)
csvs <- lapply(
  list.files(here("data/pangordlineages"),
    full.names = TRUE),
  read.csv
) %>%
bind_rows() %>%
separate(col = taxon,
  into = c("dummy", "acc", "lik", "subs"),
  sep = "_")

t1 <- csvs[csvs$acc == unique(csvs$acc)[1], ]
plot(y = t1$lik, x = seq_along(t1$lik))
```

