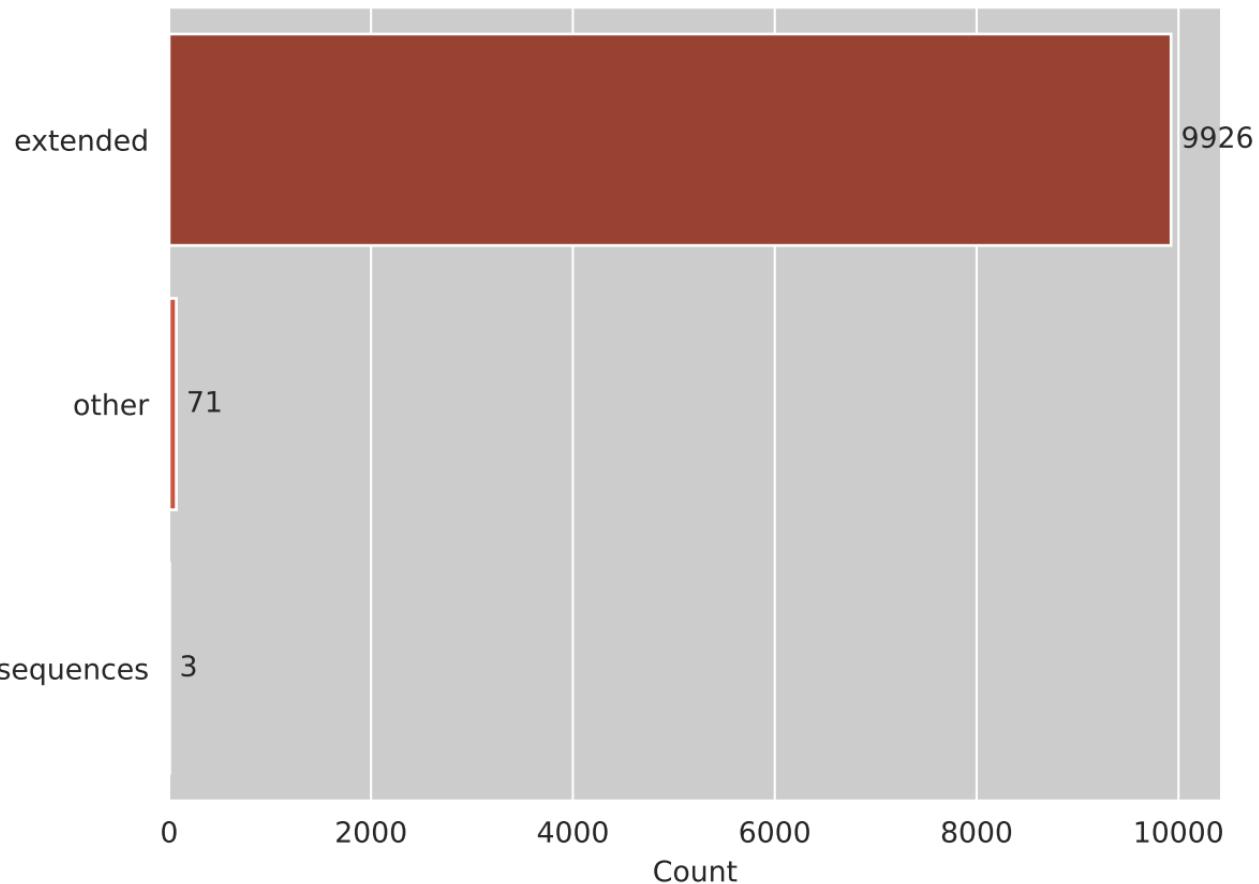


extended\_species\_startITR\_m\_0

Category

Sequence Counts



Sequence Proportions

