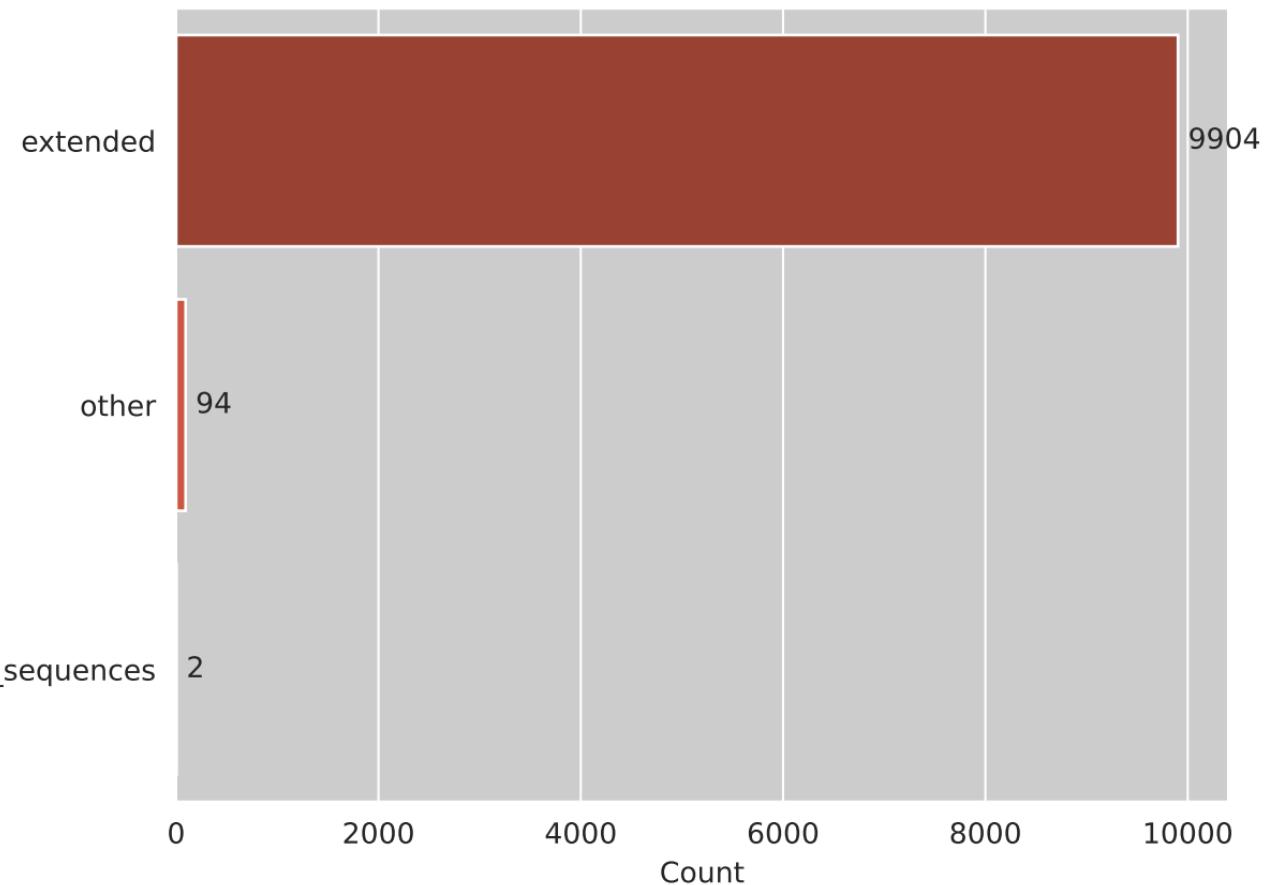


extended\_species\_bothITR\_m\_0

Category

Sequence Counts



Sequence Proportions

