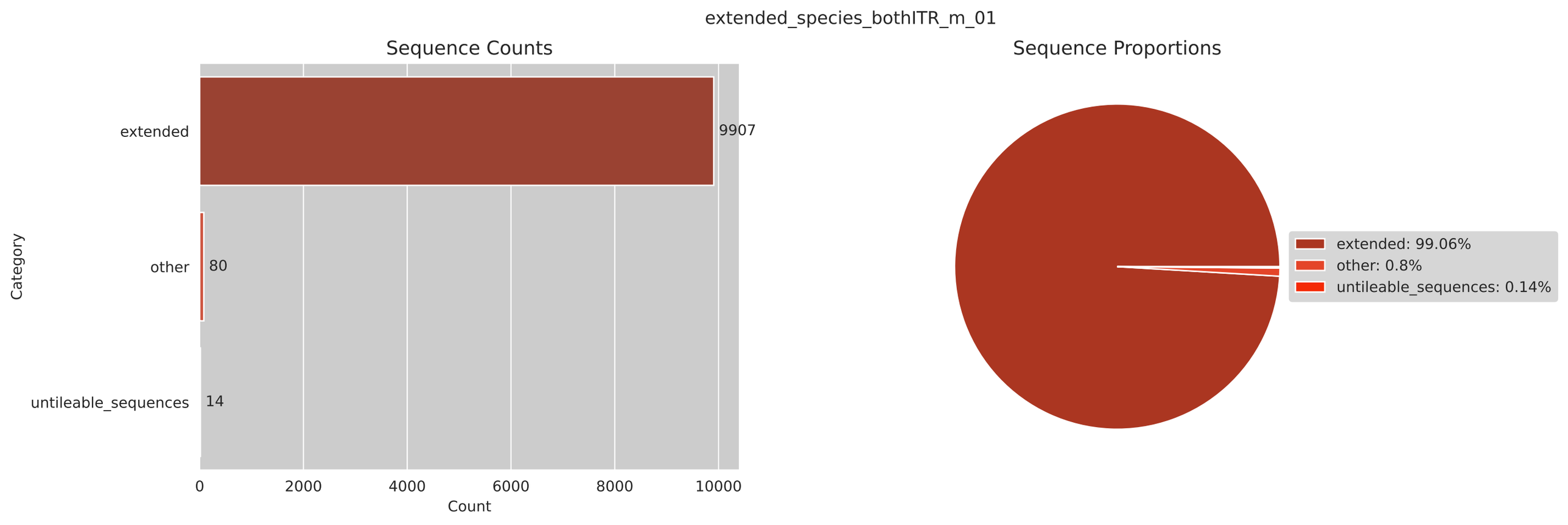
extended_species_bothITR_m_01
Sequence Counts
Category
extended
other
untileable_sequences
9907
80
14
0
2000
4000
6000
8000
10000
Count
Sequence Proportions
extended: 99.06%
other: 0.8%
untileable_sequences: 0.14%