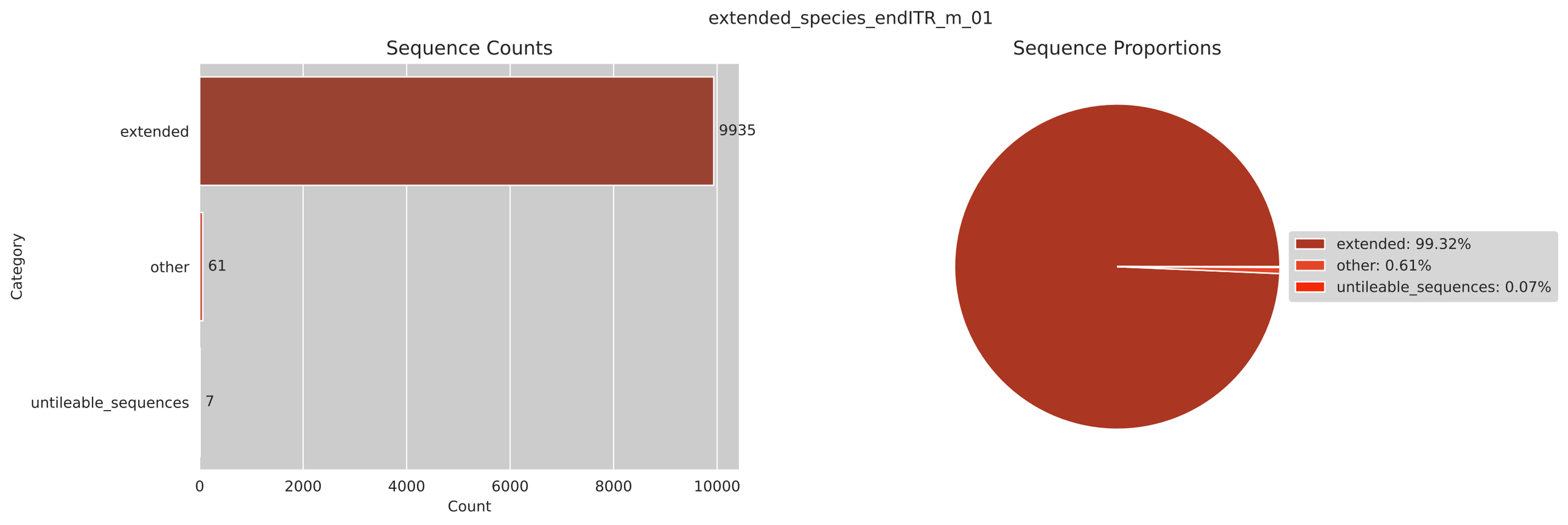
extended_species_endITR_m_01
Sequence Counts
extended
other
untileable_sequences
9935
61
7
Category
0
2000
4000
6000
8000
10000
Count
Sequence Proportions
extended: 99.32%
other: 0.61%
untileable_sequences: 0.07%