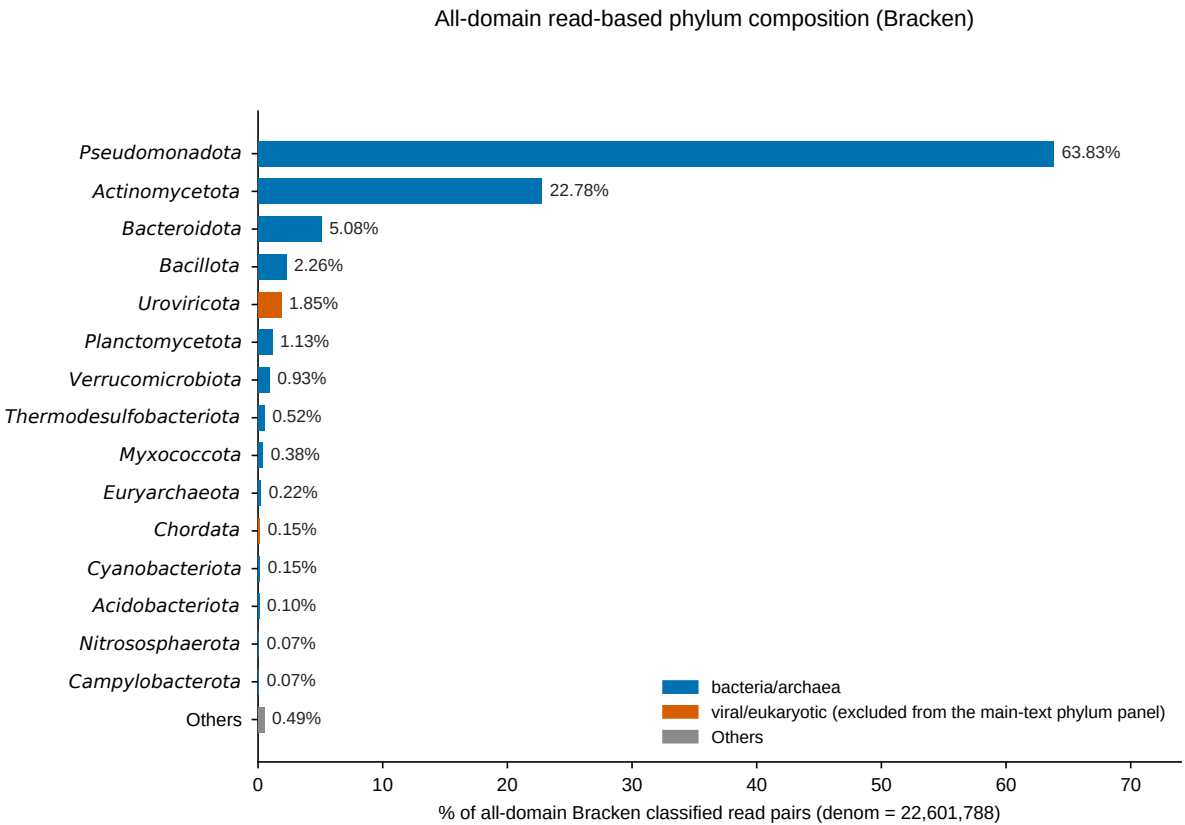
All-domain read-based phylum composition (Bracken)
Pseudomonadota
Actinomycetota
Bacteroidota
Bacillota
Uroviricota
Planctomycetota
Verrucomicrobiota
Thermodesulfobacteriota
Myxococcota
Euryarchaeota
Chordata
Cyanobacteriota
Acidobacteriota
Nitrososphaerota
Campylobacterota
Others
63.83%
22.78%
5.08%
2.26%
1.85%
1.13%
0.93%
0.52%
0.38%
0.22%
0.15%
0.15%
0.10%
0.07%
0.07%
0.49%
bacteria/archaea
viral/eukaryotic (excluded from the main-text phylum panel)
Others
0
10
20
30
40
50
60
70
% of all-domain Bracken classified read pairs (denom = 22,601,788)