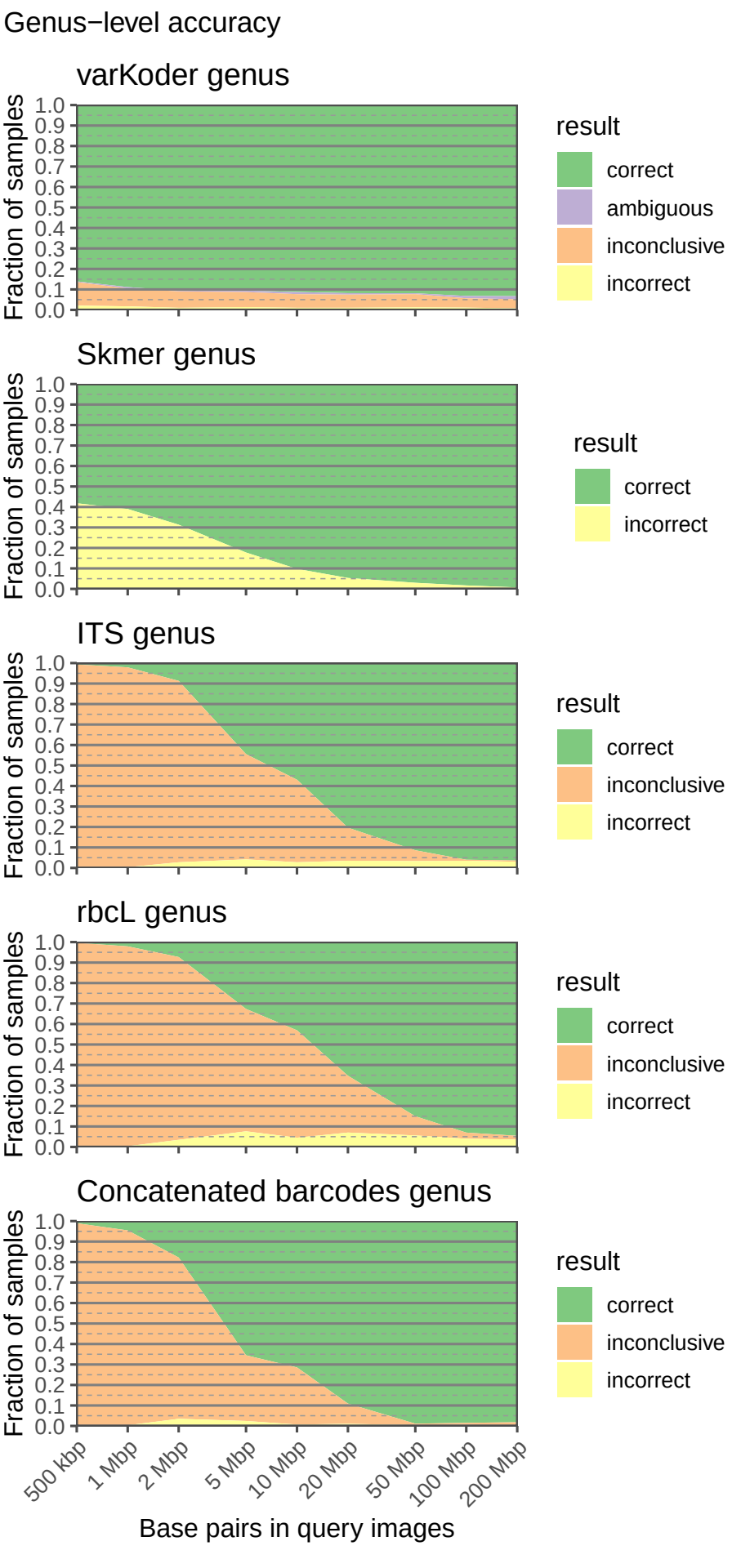
Genus−level accuracy
varKoder genus
Skmer genus
ITS genus
rbcL genus
Concatenated barcodes genus
Fraction of samples
0.0
0.1
0.2
0.3
0.4
0.5
0.6
0.7
0.8
0.9
1.0
result
correct
ambiguous
inconclusive
incorrect
500 kbp
1 Mbp
2 Mbp
5 Mbp
10 Mbp
20 Mbp
50 Mbp
100 Mbp
200 Mbp
Base pairs in query images