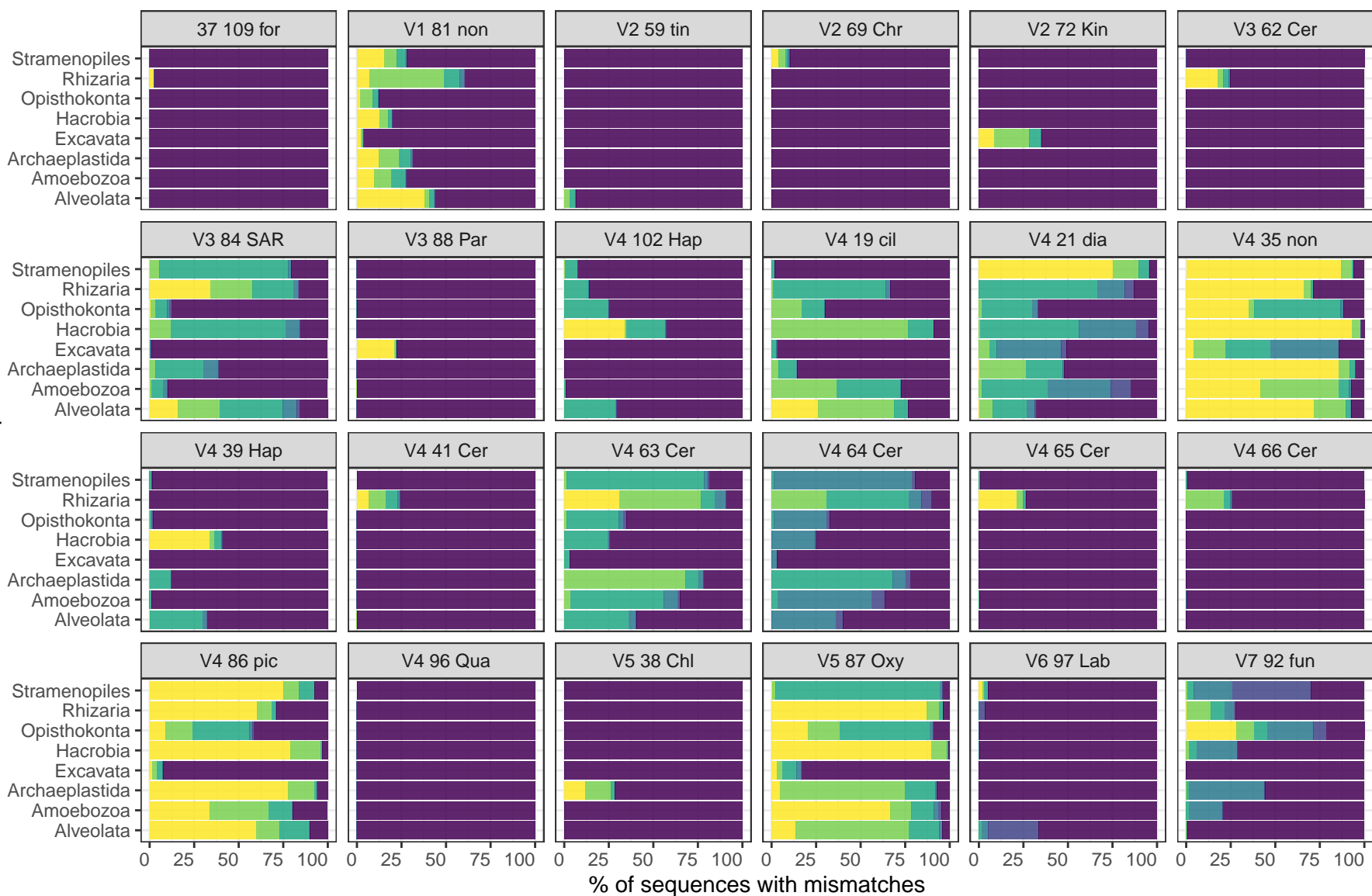


A

specific primers

Mismatches 5+ 4 3 2 1 0

Group

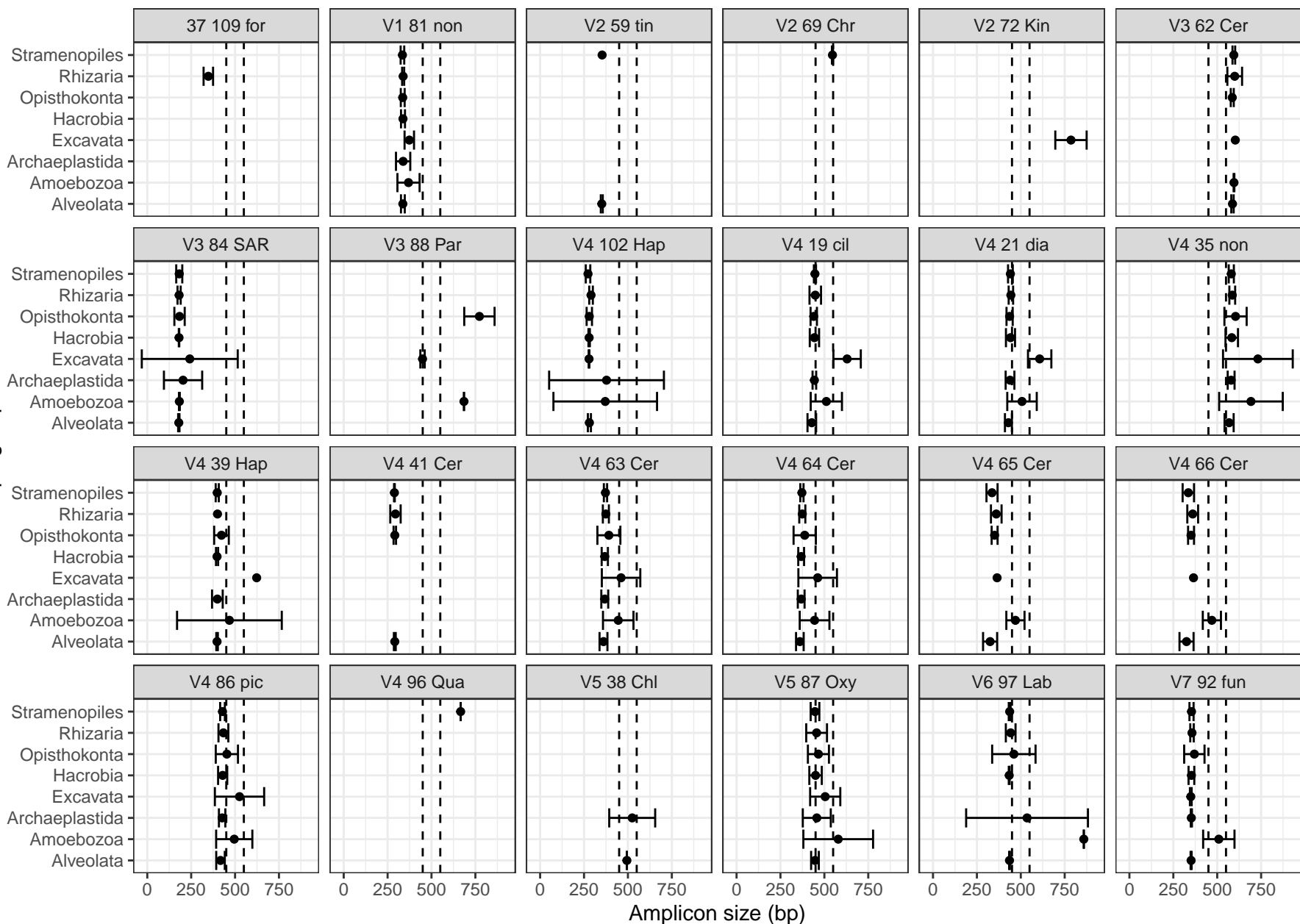


% of sequences with mismatches

B

specific primers

Supergroup



Amplicon size (bp)