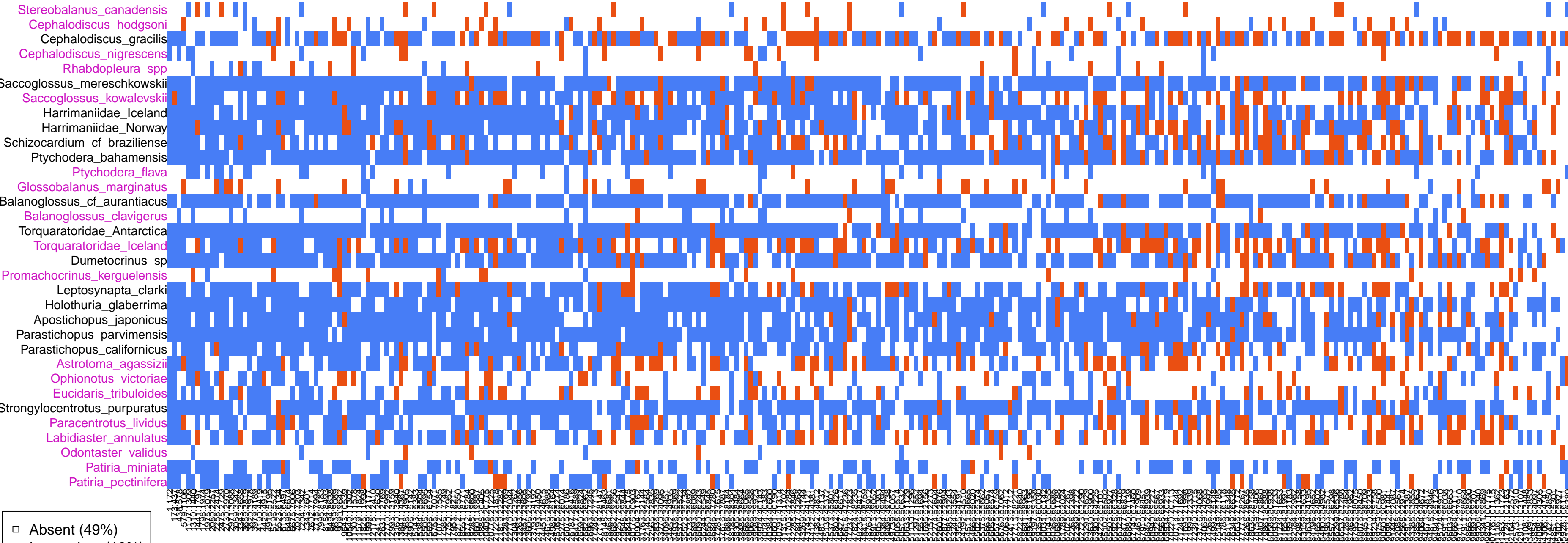


n taxa = 33

cannon2014\_299-33\_matrix.tab



Absent (49%)

Incomplete (10%)

Present (40%)

Gene partitions ( n = 299 )