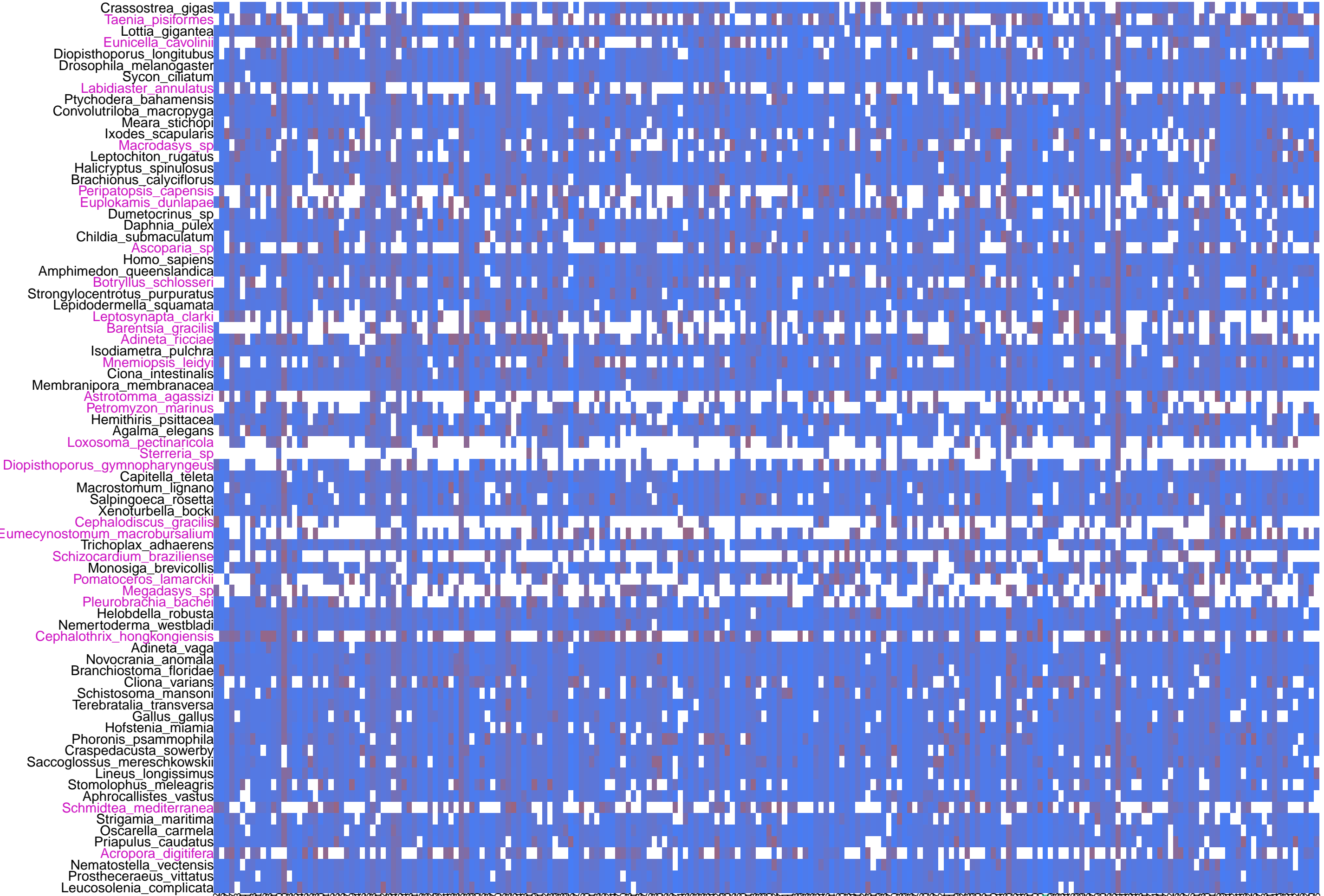


n taxa = 78

cannon2016_percent_matrix.tab



□ Absent (20%)
■ Partial (80%)
■ Complete (1%)

Gene partitions (n = 212)