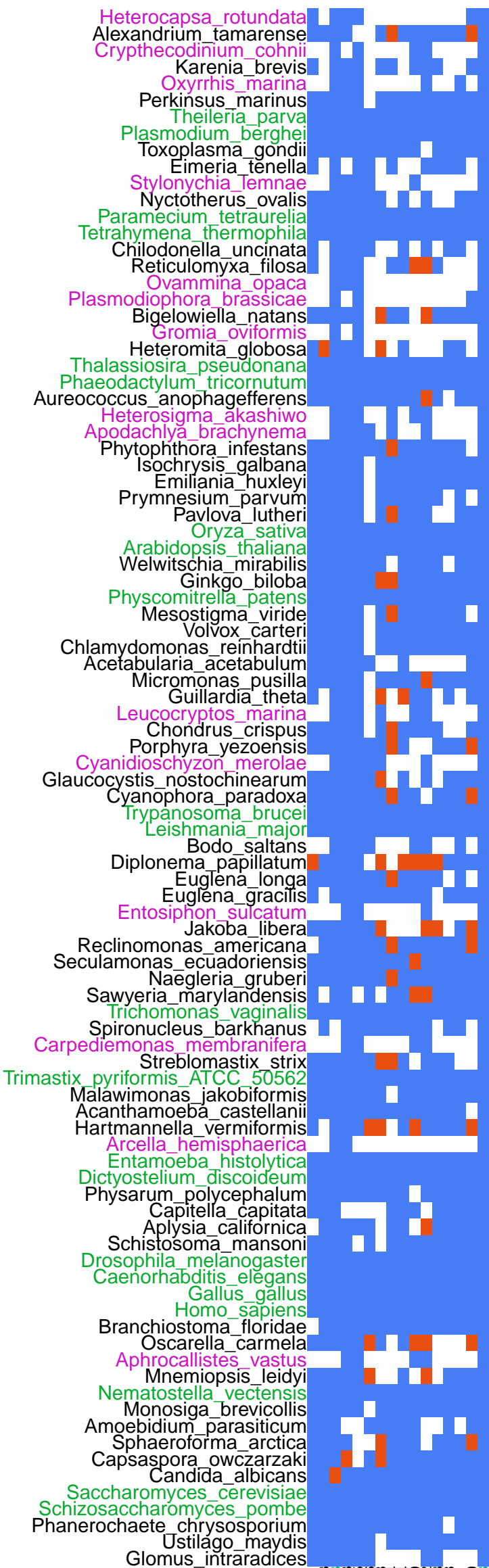


n taxa = 94

parfrey2011_fig2_matrix



□ Absent (21%)
■ Incomplete (4%)
■ Present (76%)

Gene partitions (n = 57)