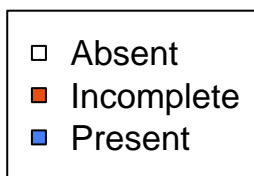
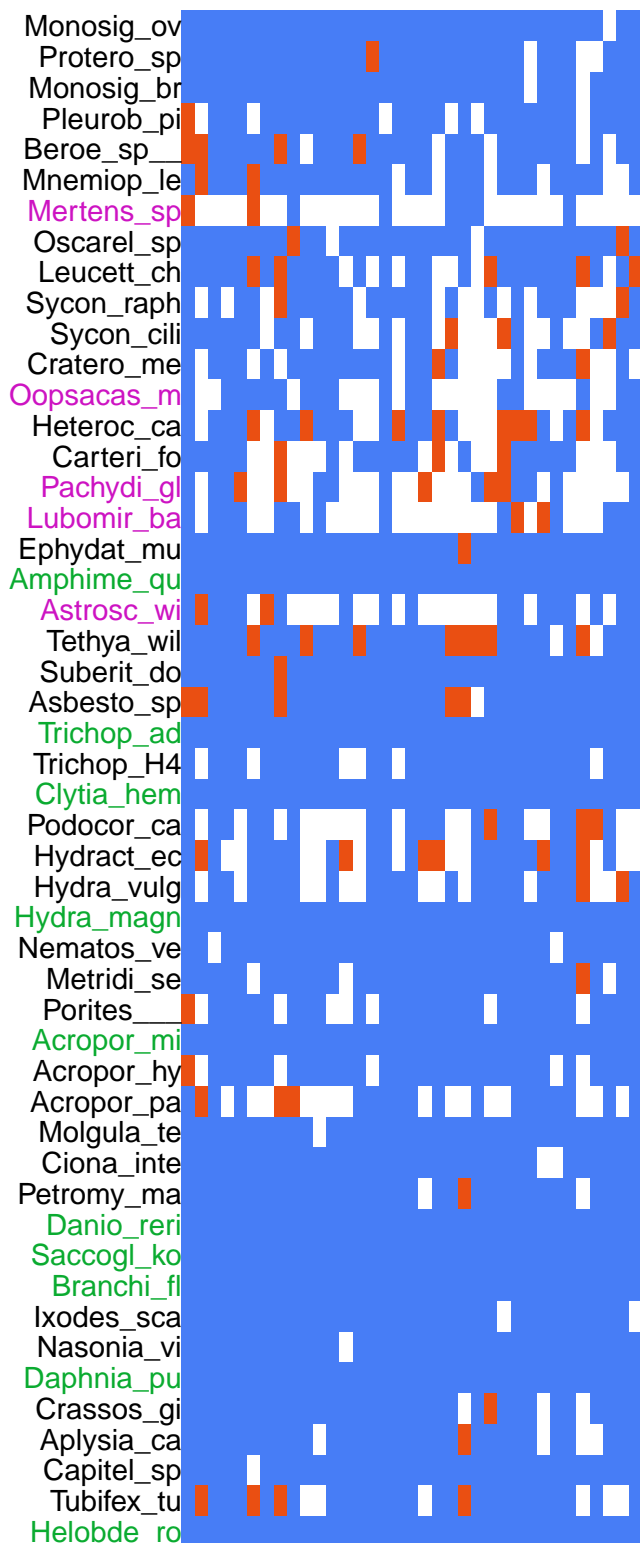


n taxa = 50 nosenko2013_nonribosomal_matrix



Gene partitions (n = 35)