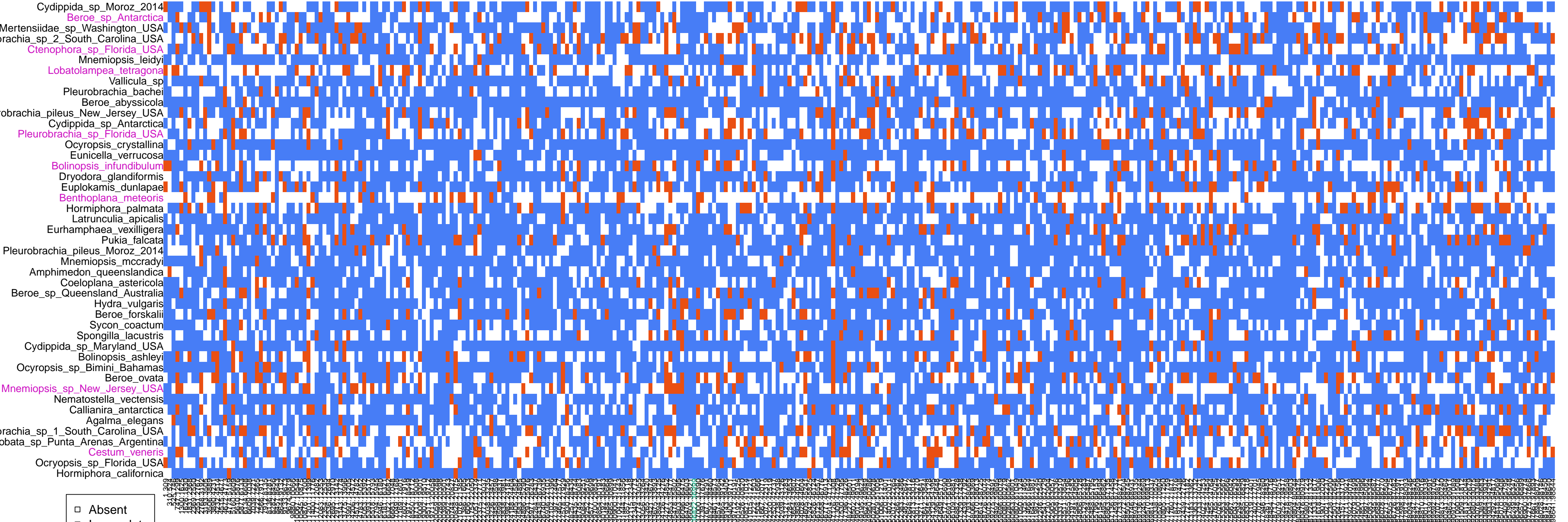


n taxa = 45

whelan2017_Ctenophore_full_matrix.tab



Gene partitions (n = 350)