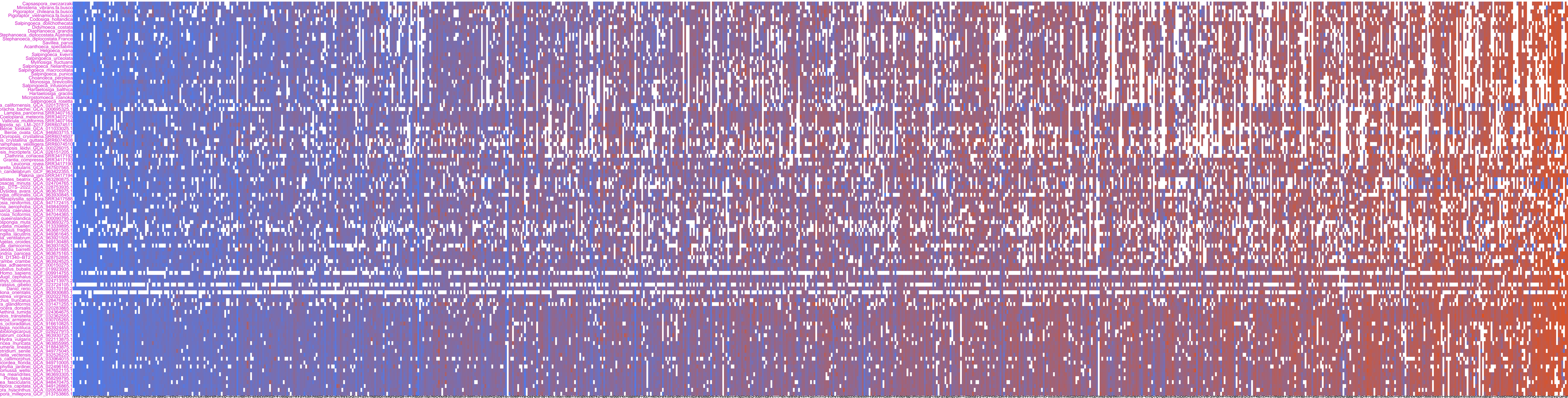


n taxa = 101

steenwyk2025_ClipKITkpic_concat.reorder.matrix.tab.gz



□ Absent (19%)
■ Partial (81%)
■ Complete (0%)

Gene partitions (n = 869)