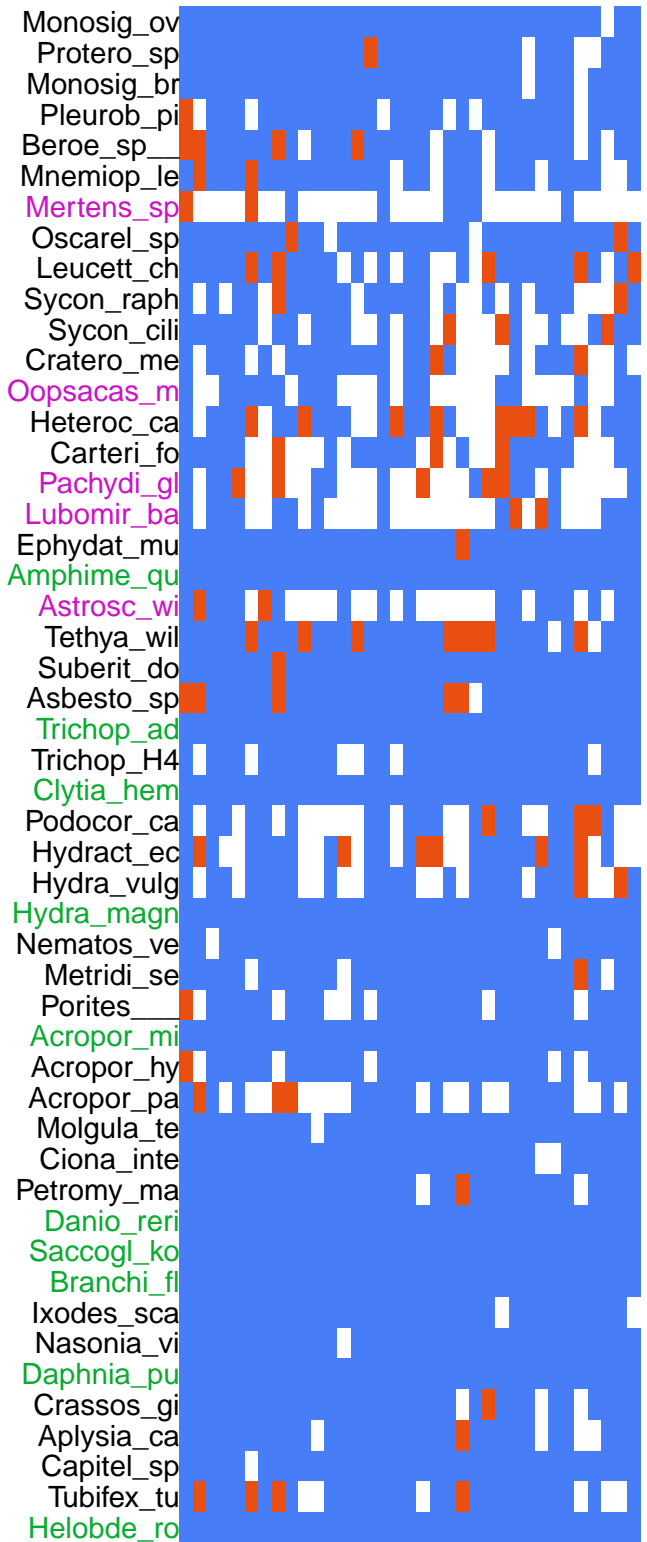


n taxa = 50 nosenko2013_nonribosomal_matrix



- Absent (17%)
- Incomplete (5%)
- Present (79%)

Gene partitions (n = 35)