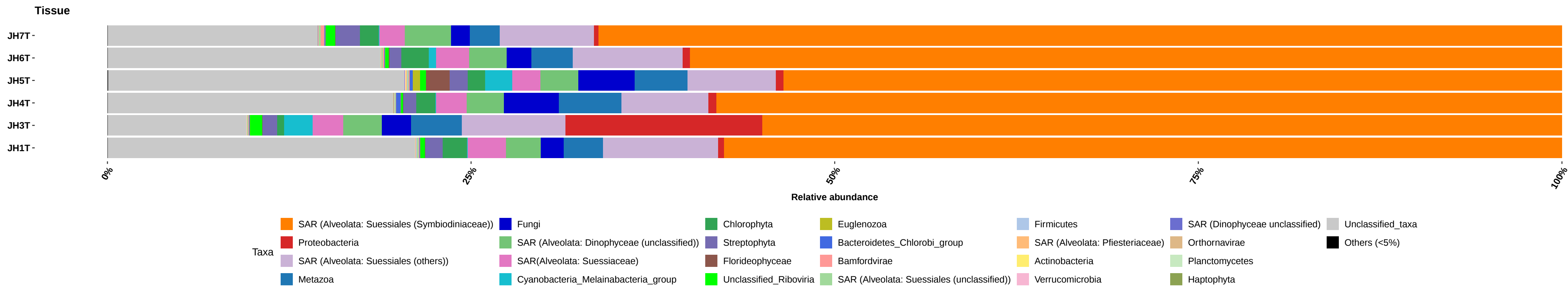
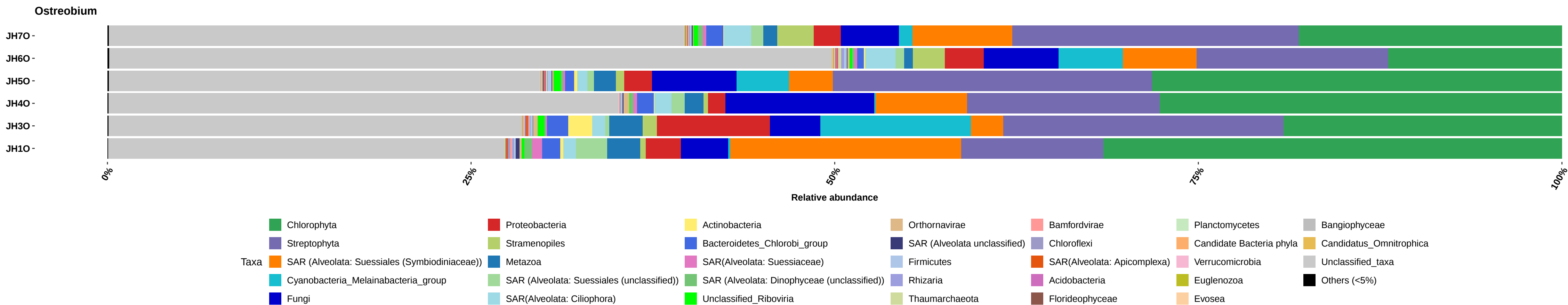


Meta-transcriptome based abundance profiles

A



B



C

