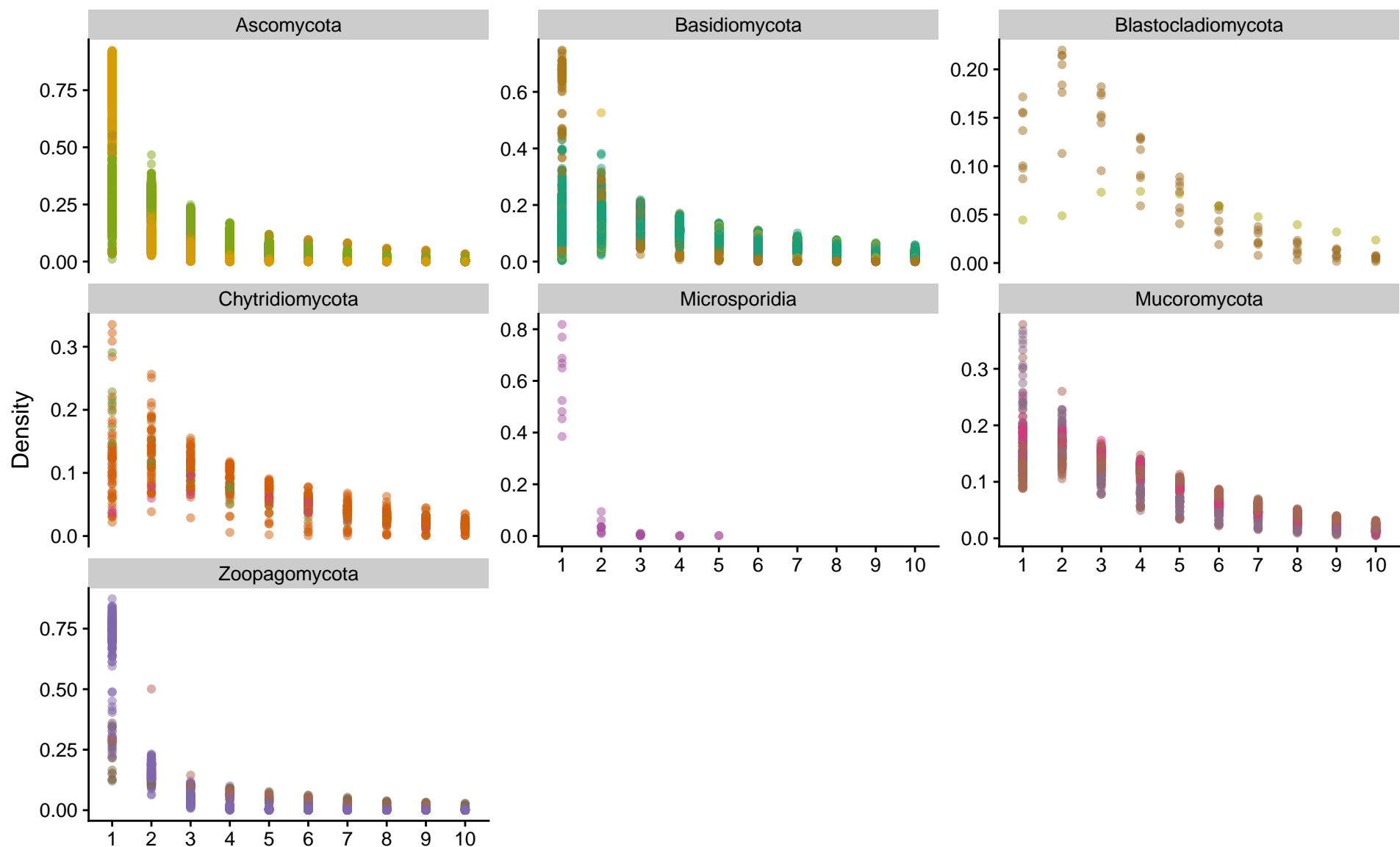


# Frequency of transcripts with exon counts with Orthologs



Transcript Exon Count

