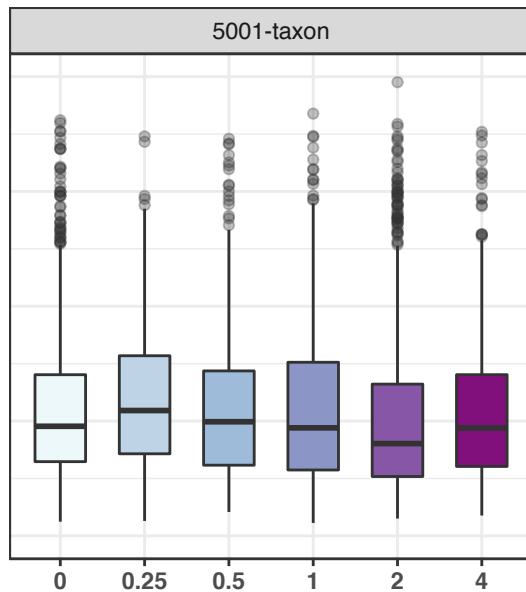
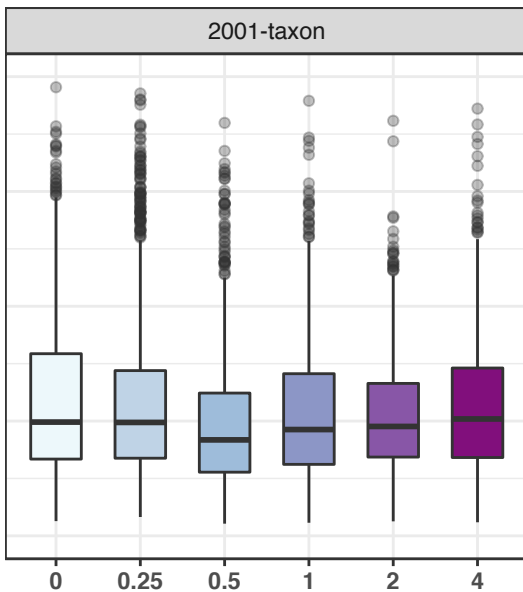
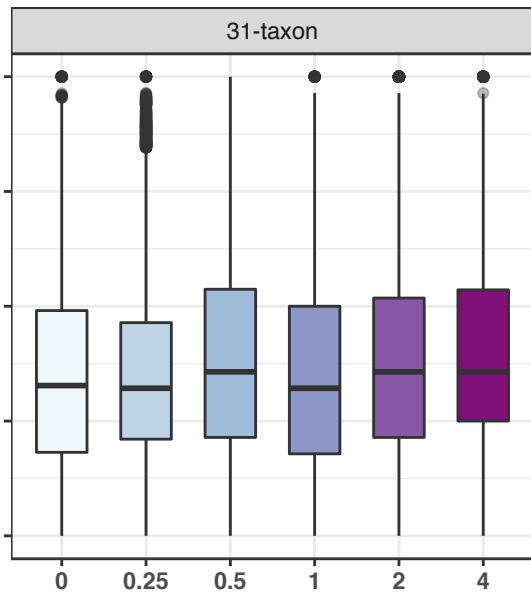


RF distance (true vs estimated gene trees)



Root to crown ratio

0	0.5	2
0.25	1	4