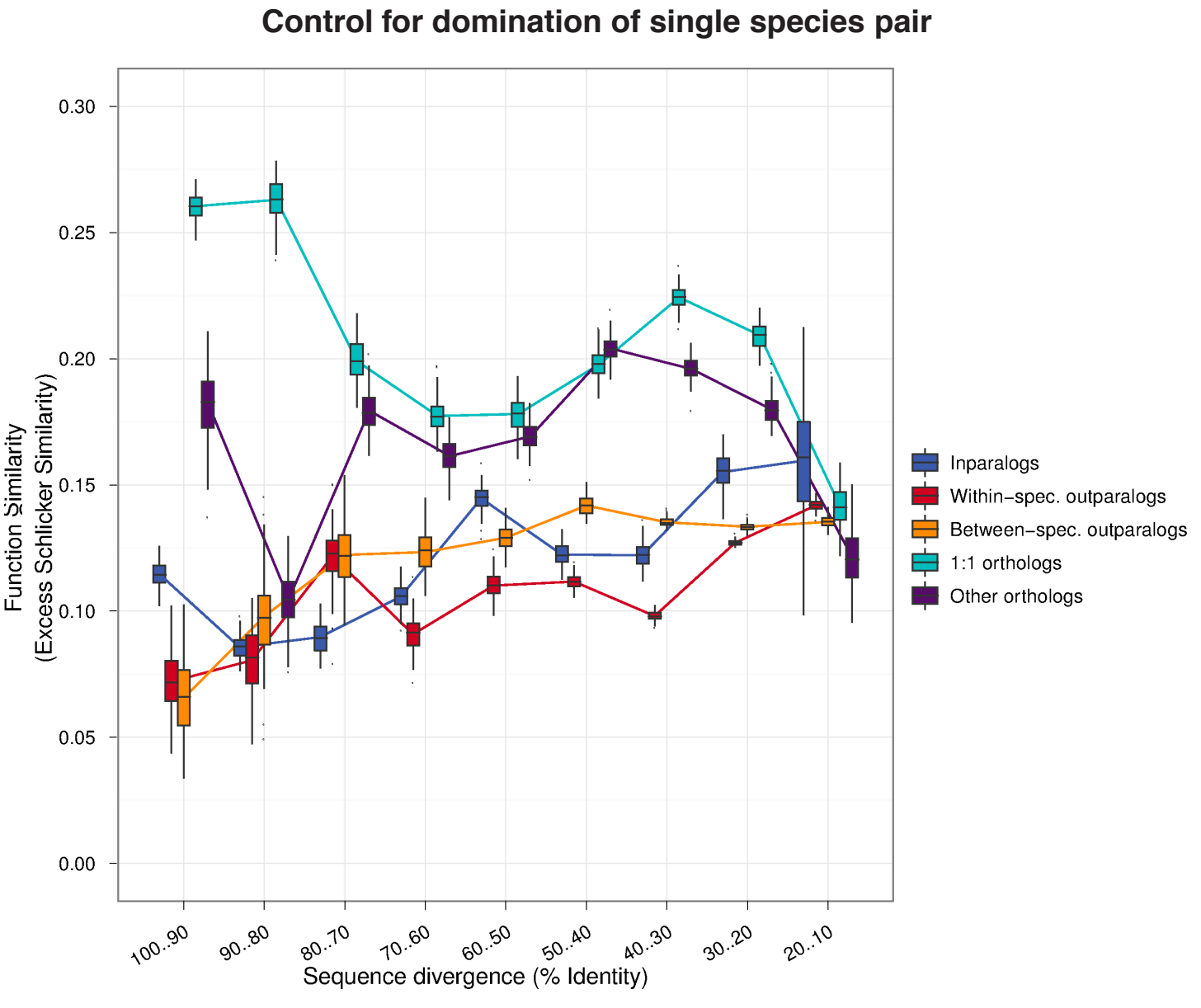
Control for domination of single species pair
Function Similarity
(Excess Schlicker Similarity)
0.30
0.25
0.20
0.15
0.10
0.05
0.00
100..90
90..80
80..70
70..60
60..50
50..40
40..30
30..20
20..10
Sequence divergence (% Identity)
Inparalogs
Within−spec. outparalogs
Between−spec. outparalogs
1:1 orthologs
Other orthologs