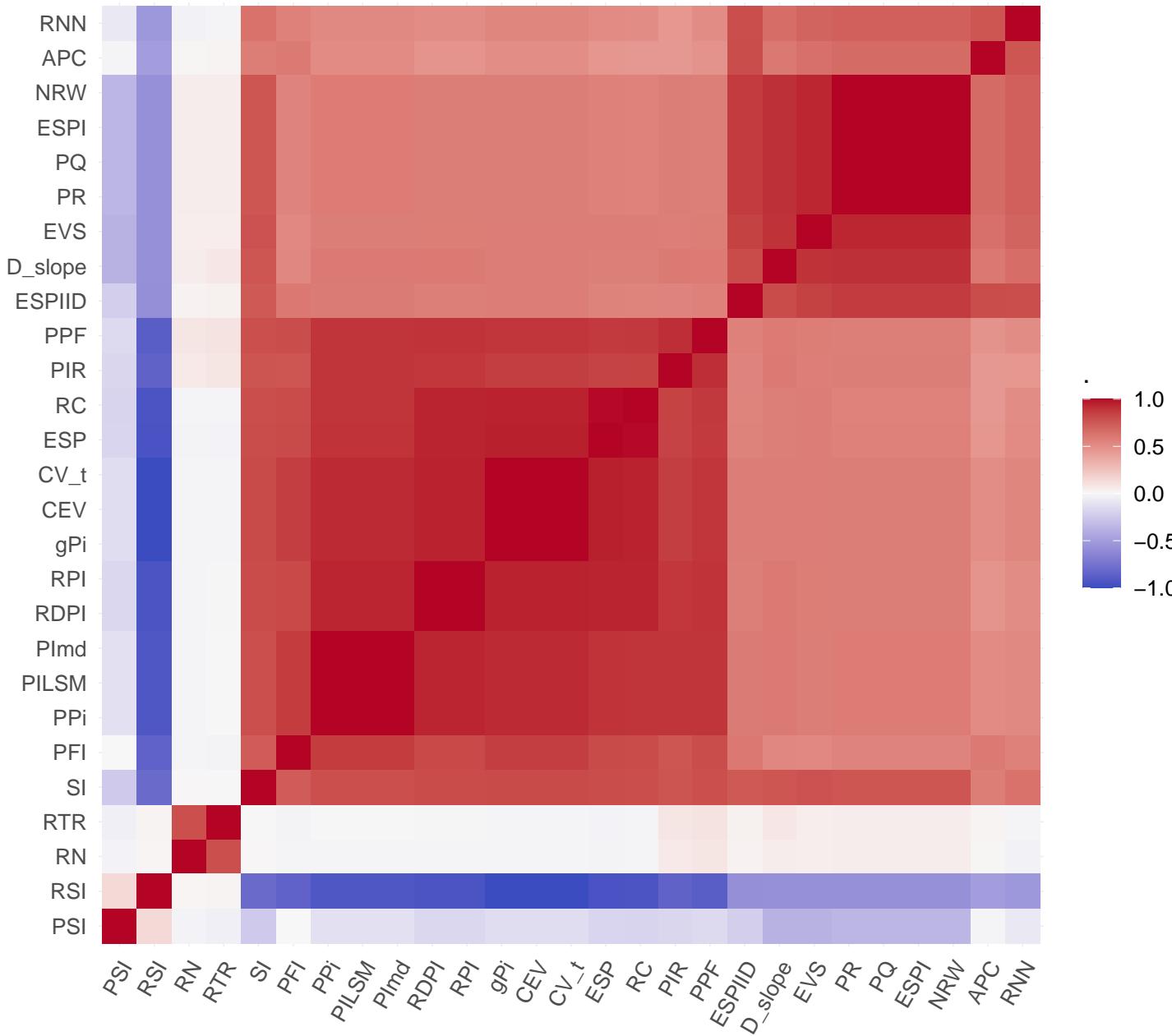
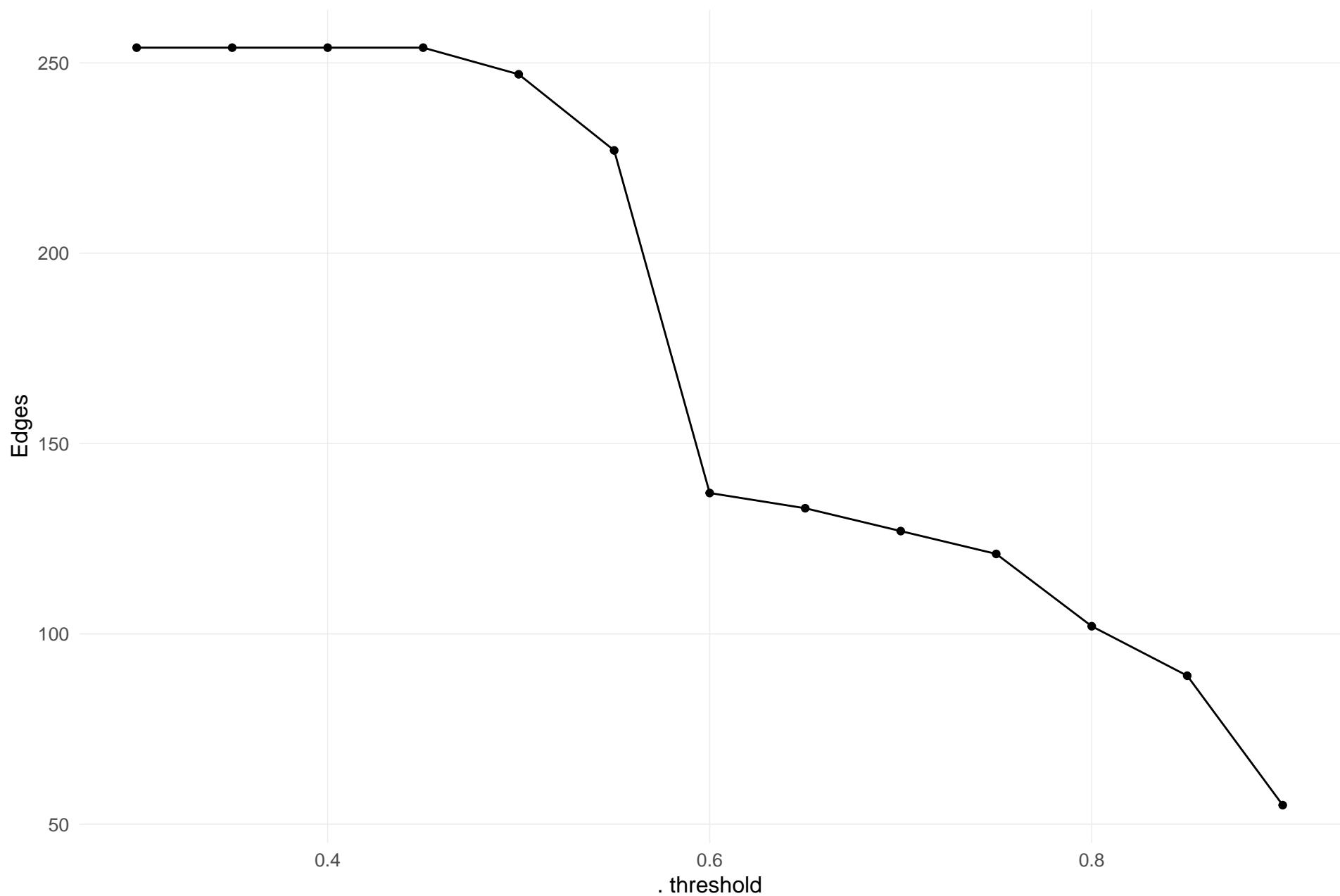


Analysis report . full_res10

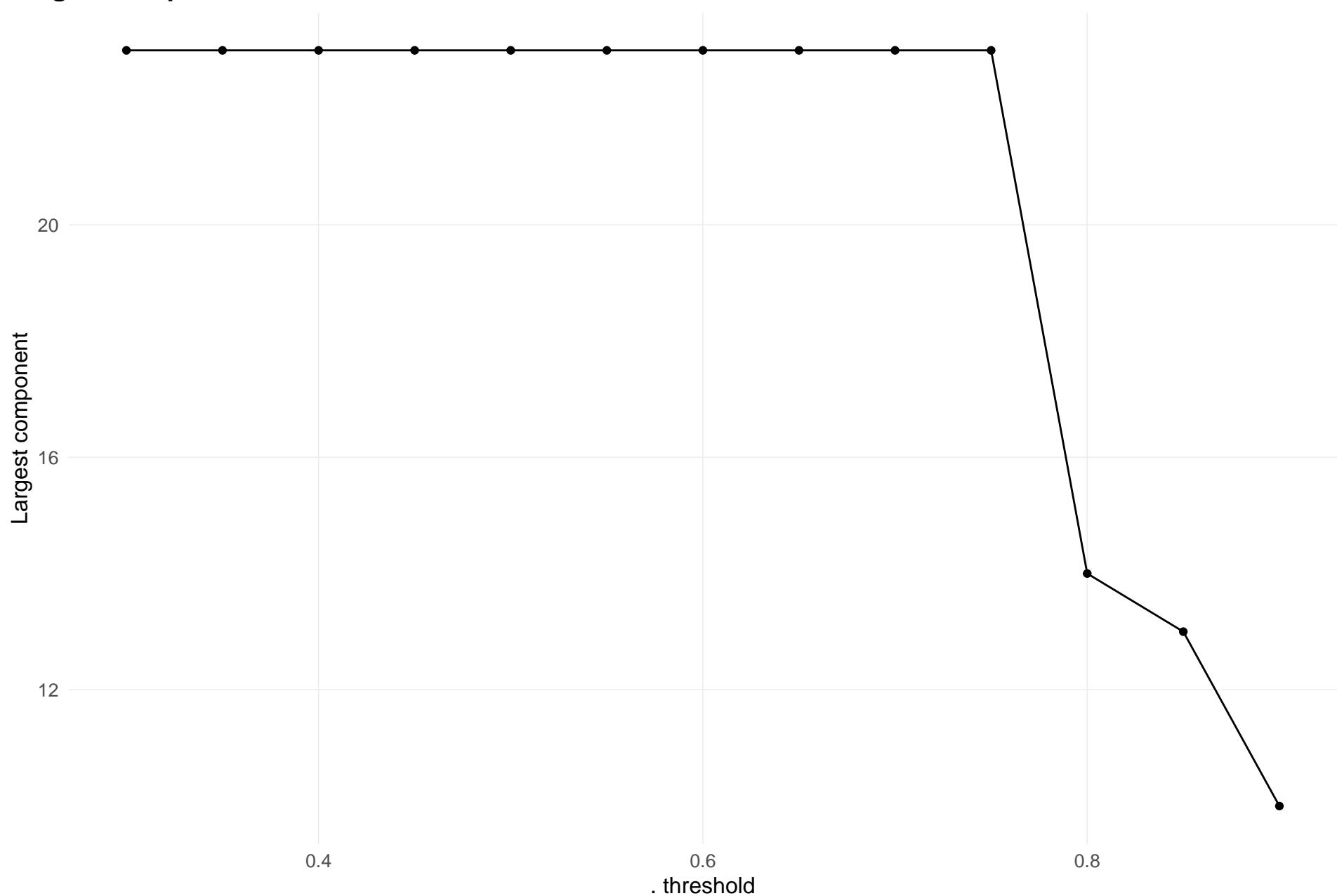
. threshold = 0.8 k = {3, 5, 10, 15, 20}

Kendall's . . full_res10

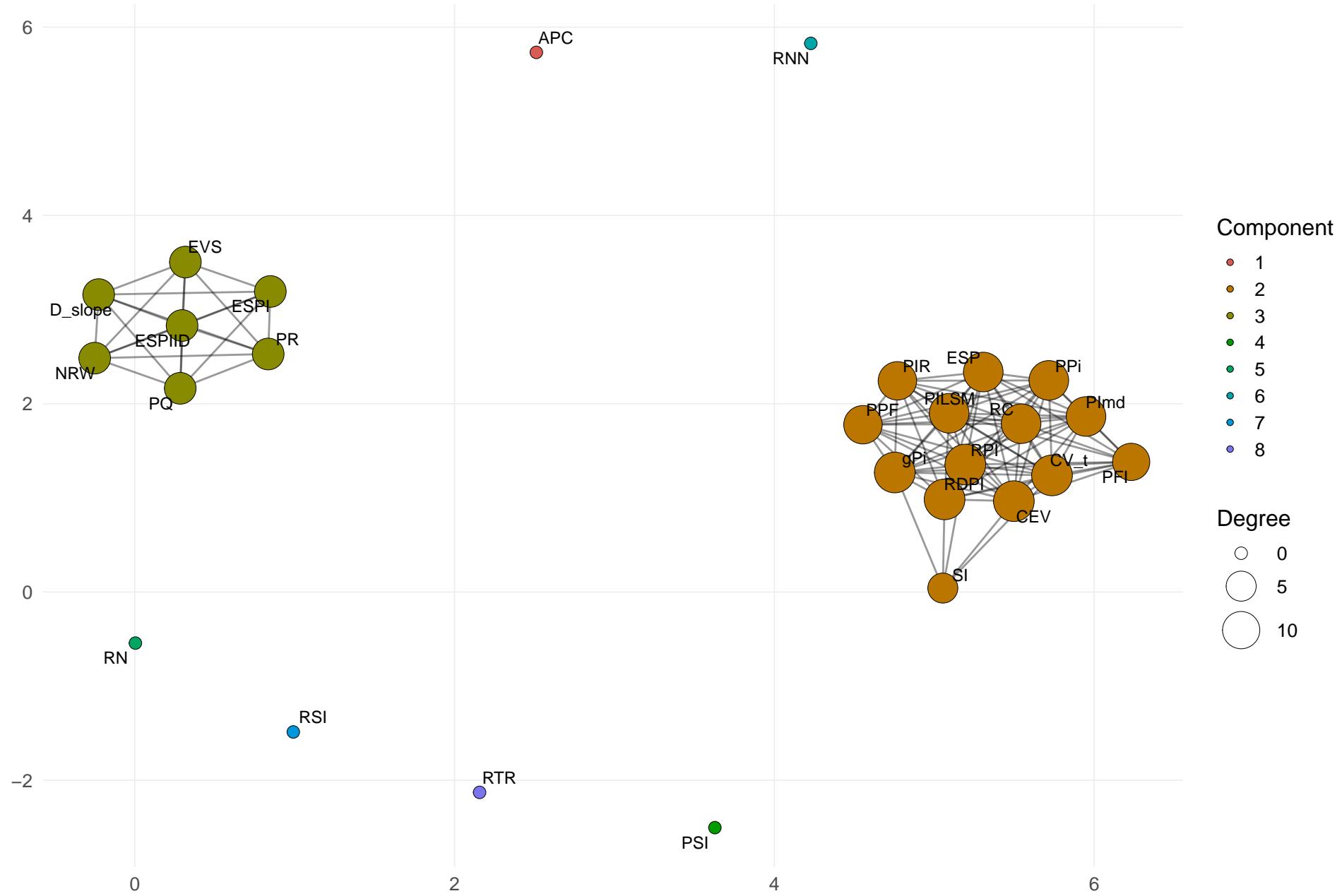
Edge count vs . . full_res10



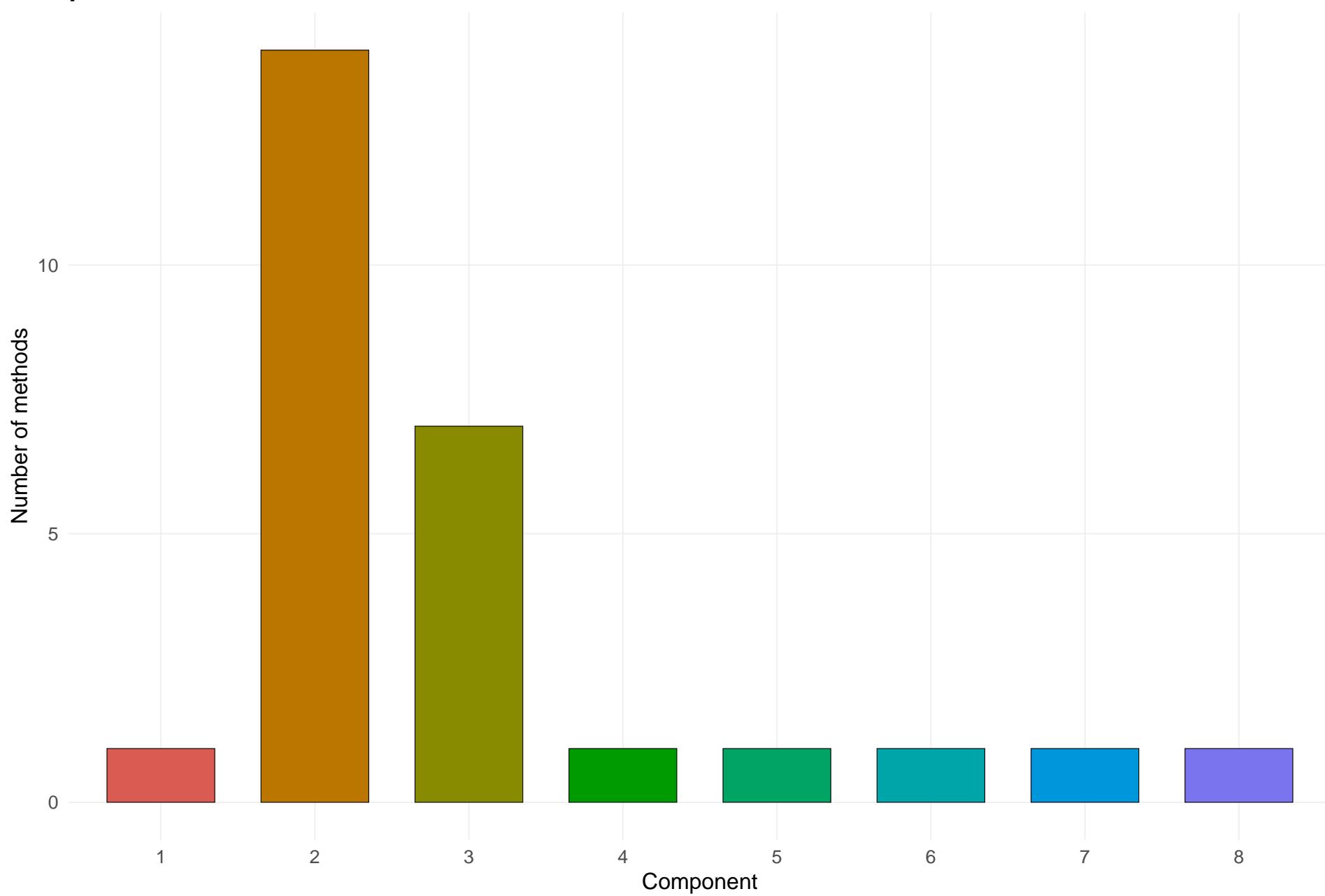
Largest component size vs . . full_res10



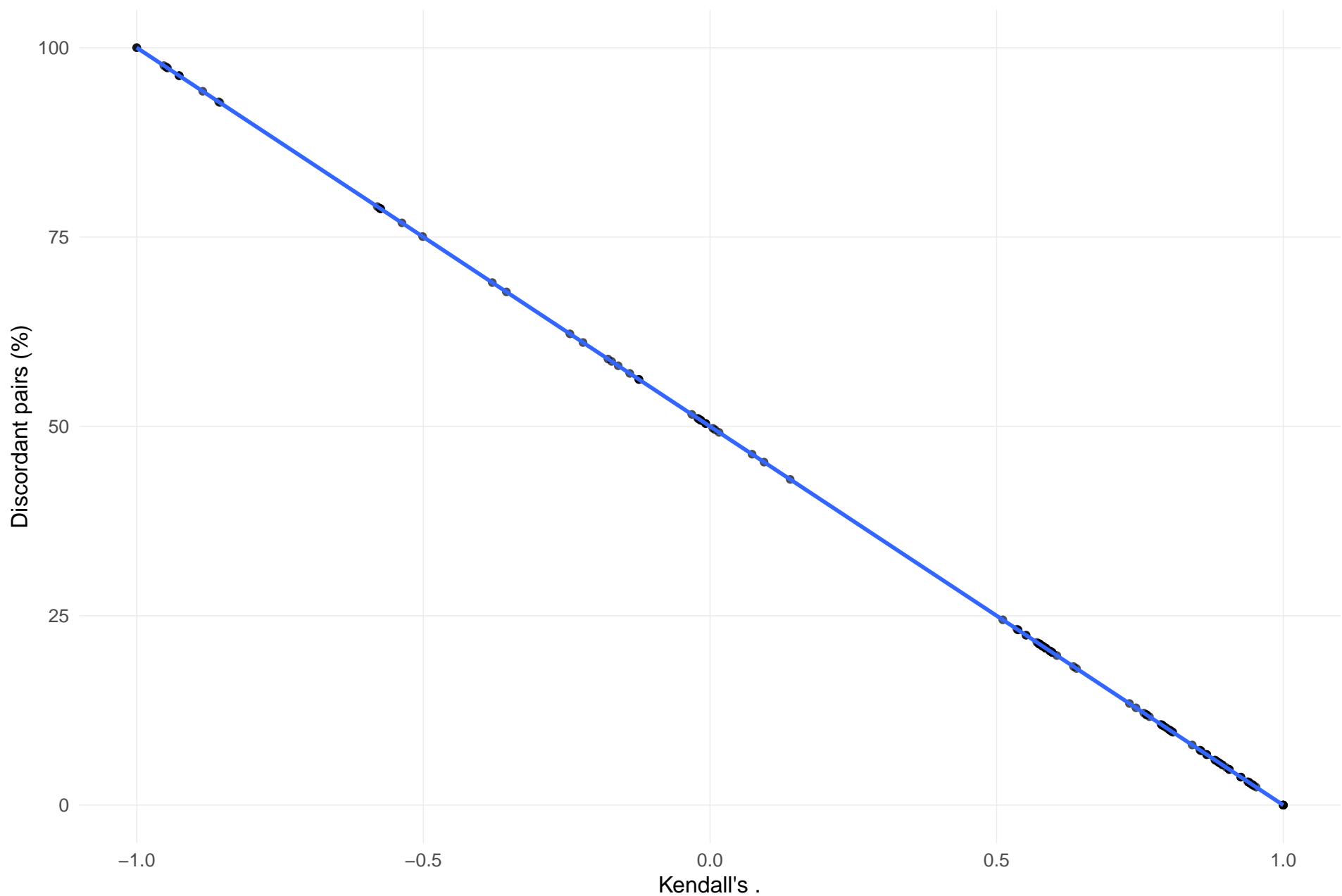
Agreement network (. >= 0.8) . full_res10



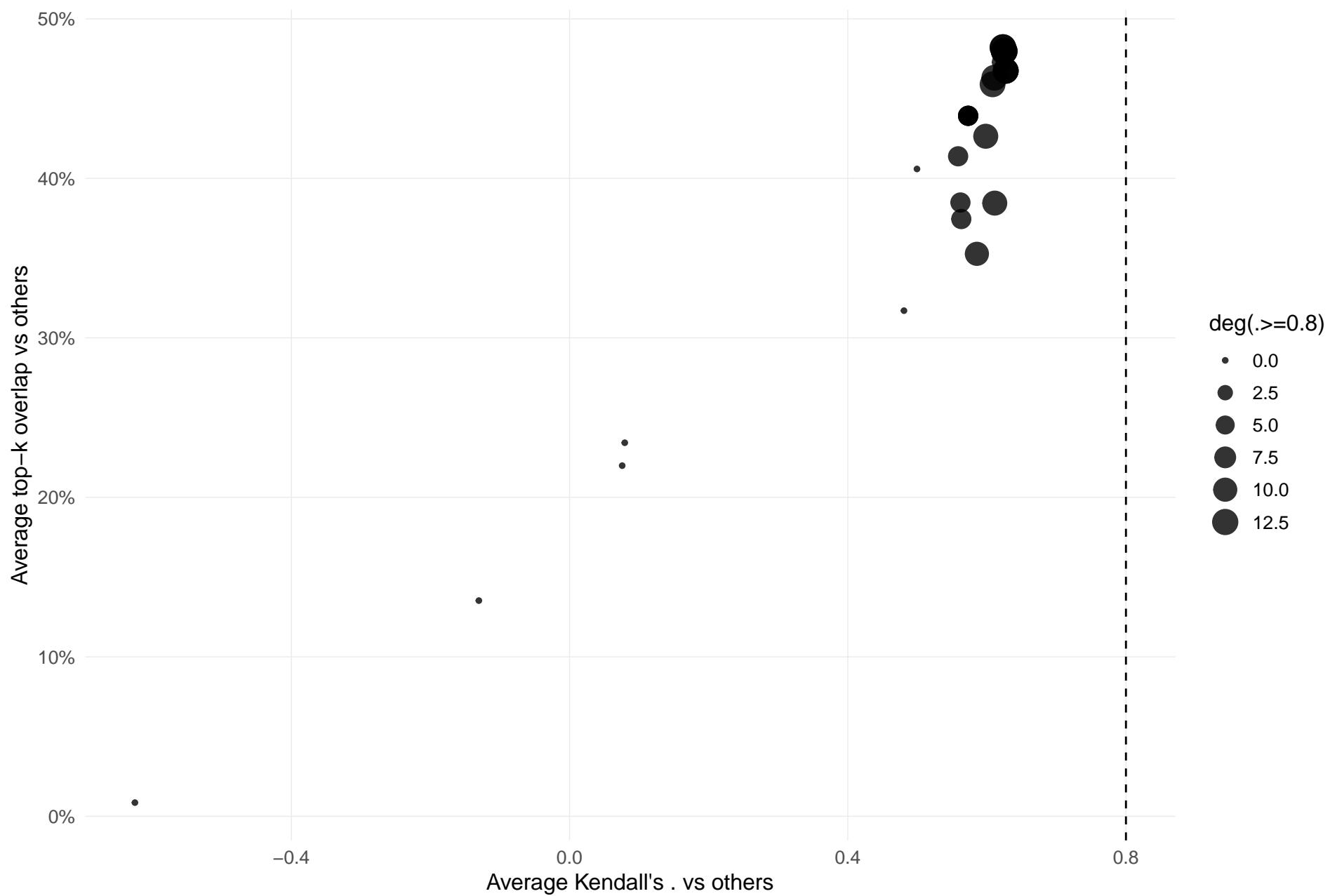
Component sizes . full_res10



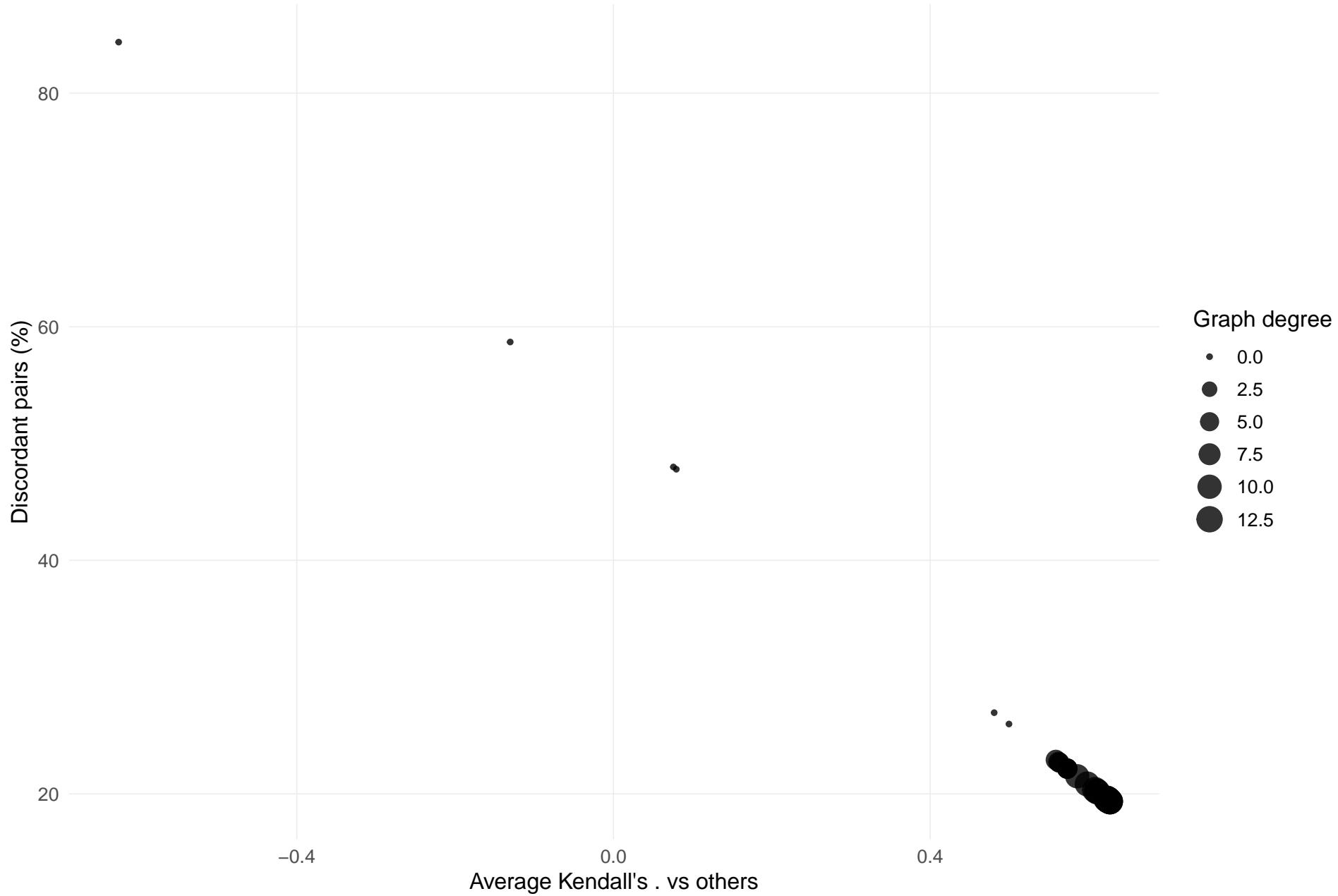
Discordant pairs vs Kendall's . . full_res10



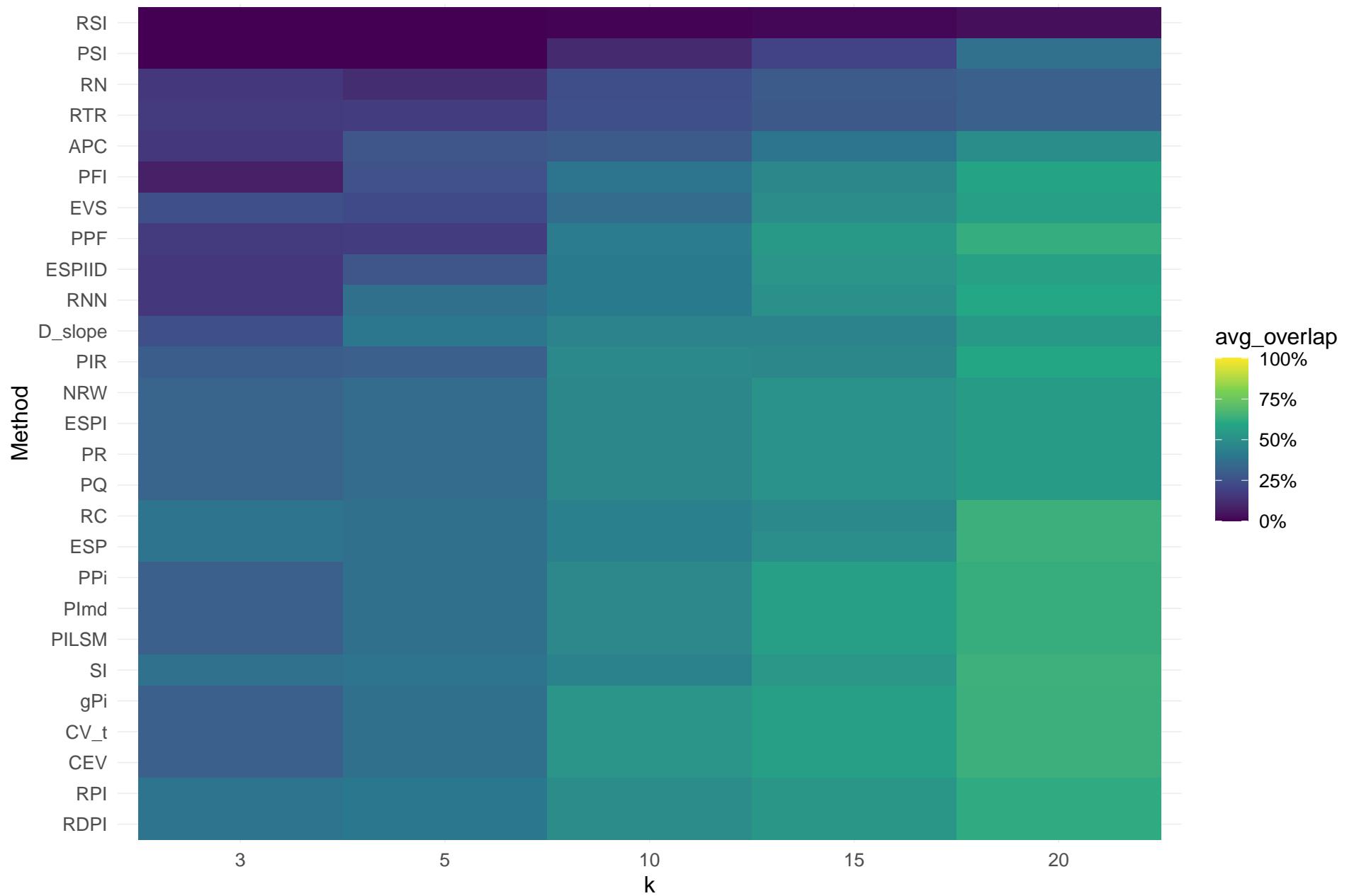
Method agreement summary . full_res10



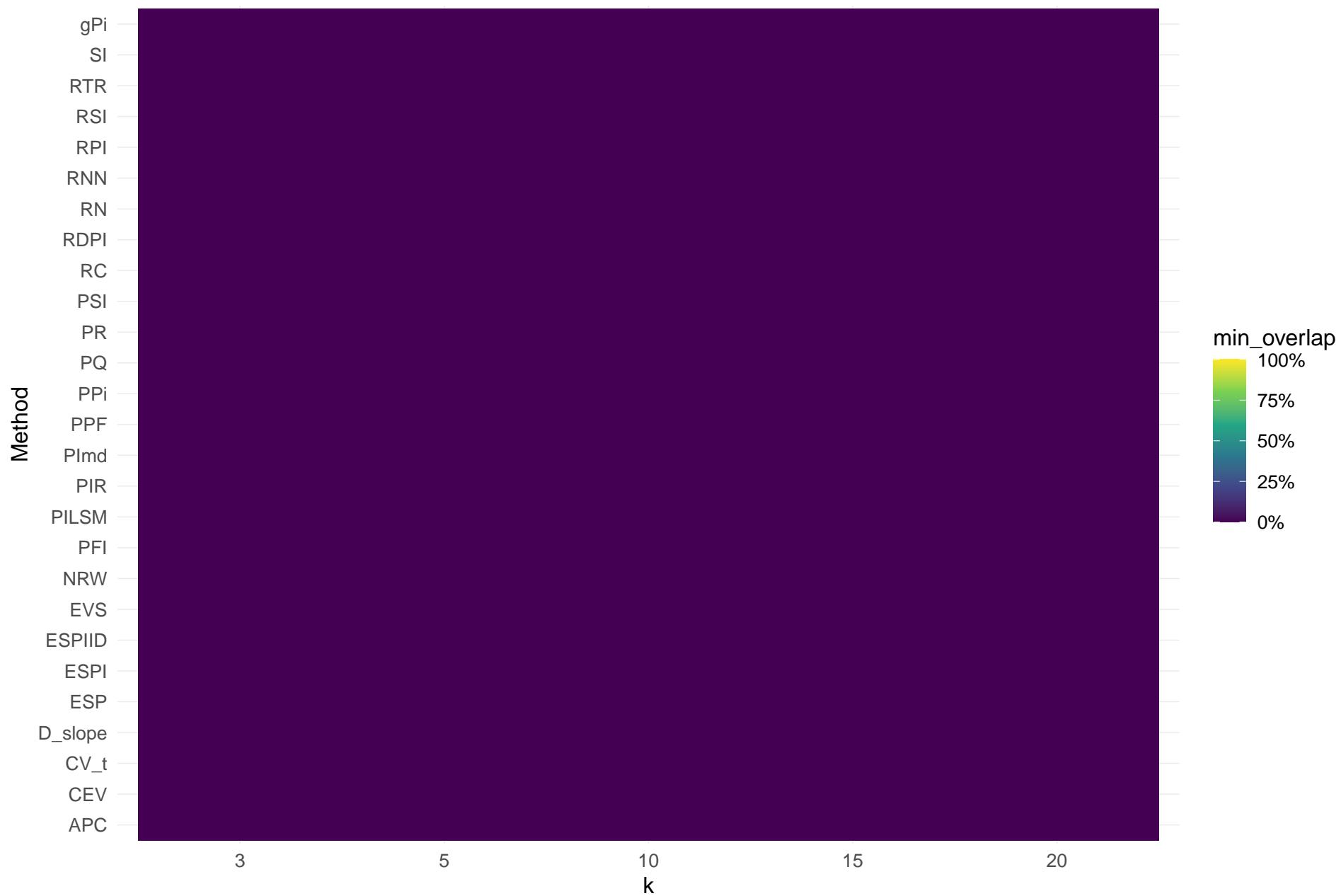
. vs discordance . full_res10



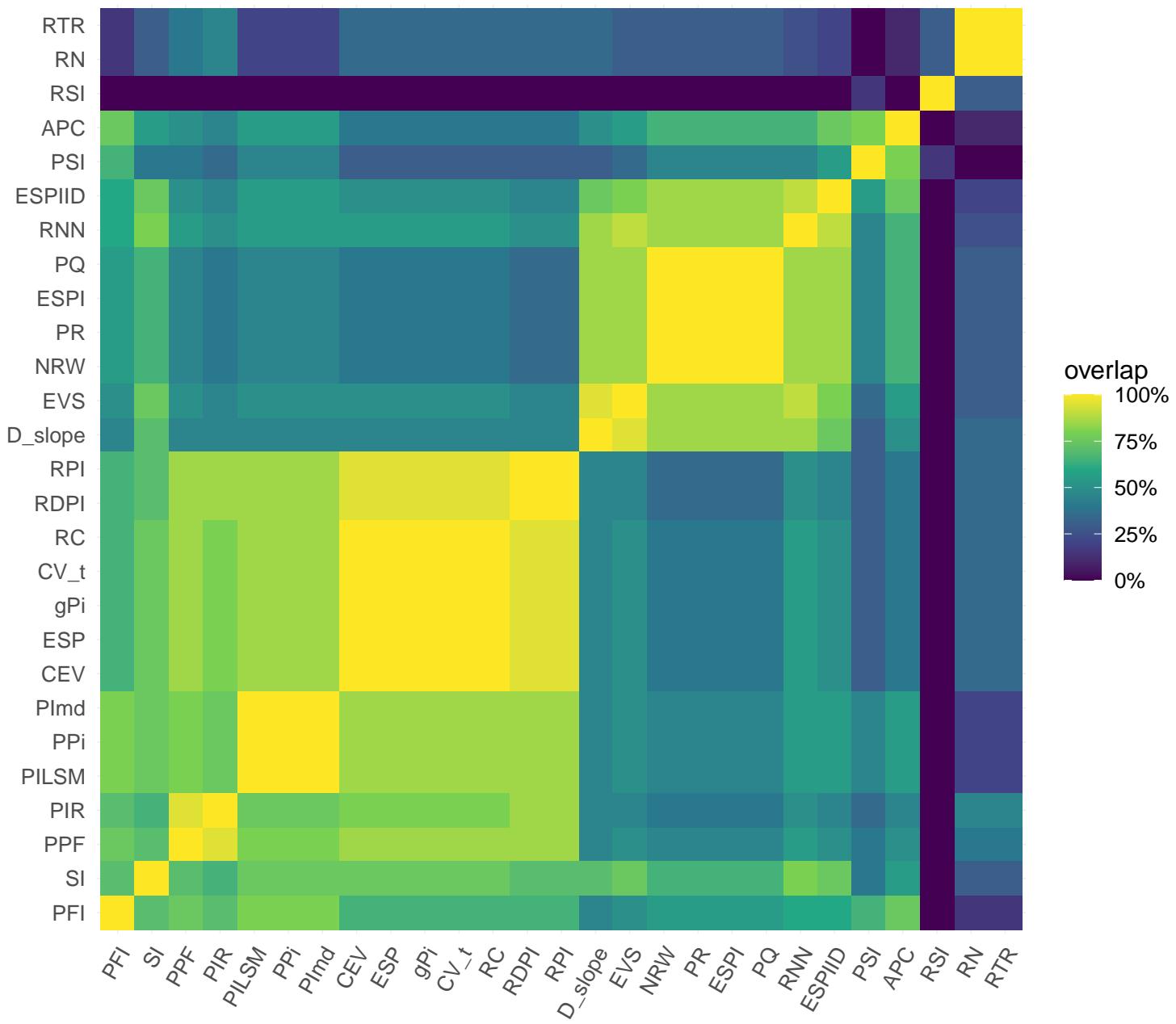
Average top-k overlap vs others (global) . full_res10



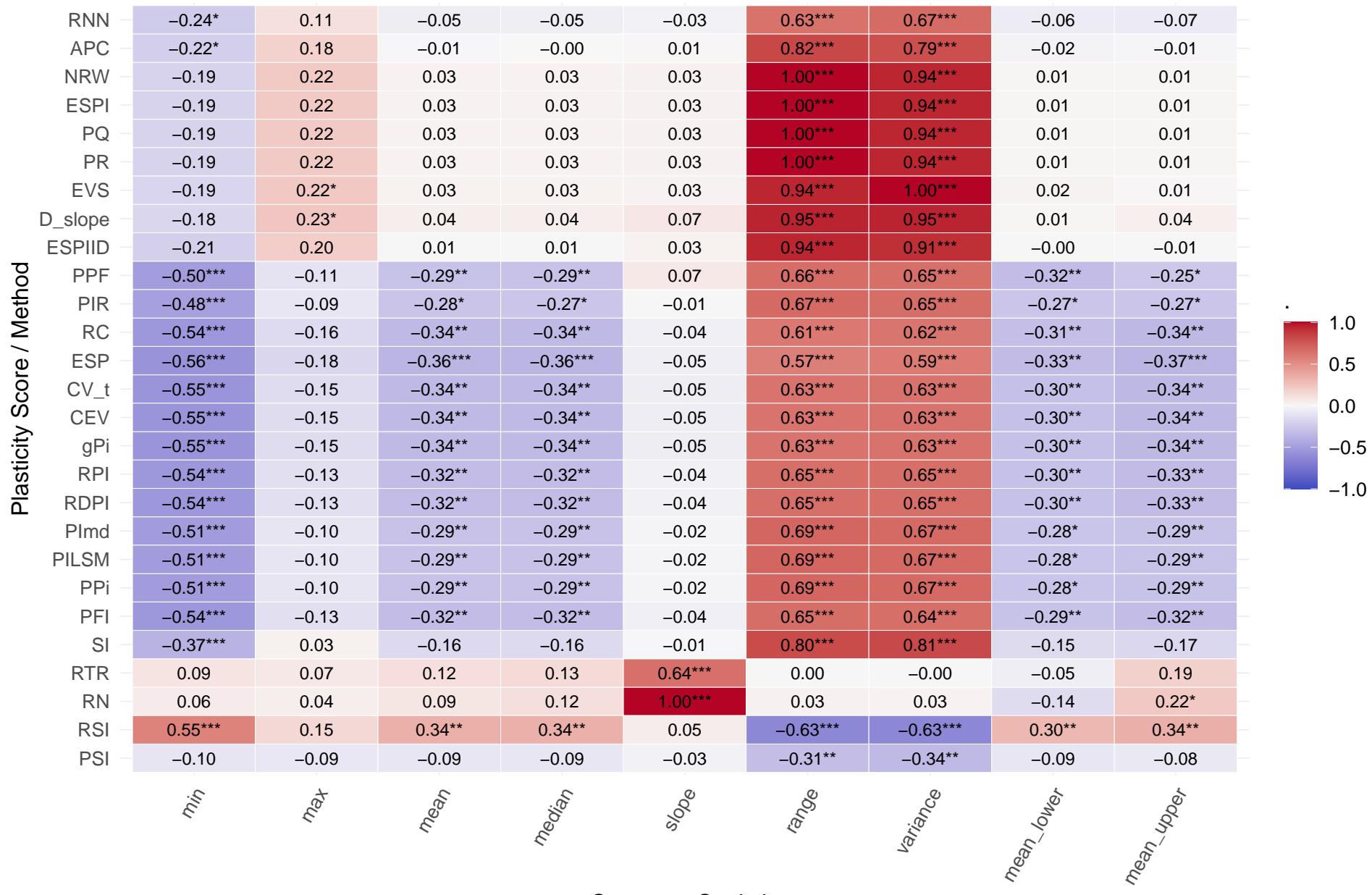
Minimum top-k overlap vs others (global) . full_res10



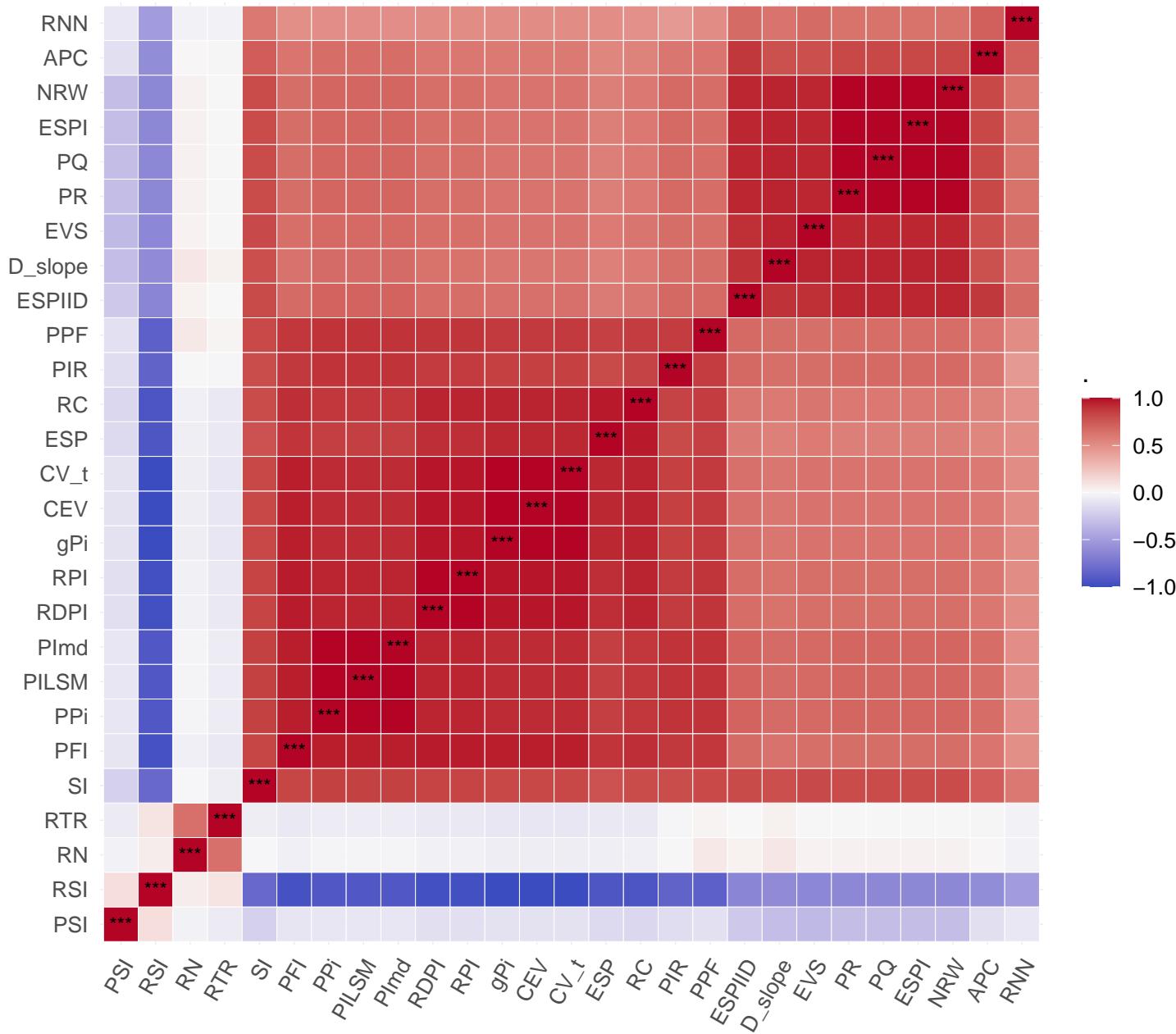
Pairwise top-20 overlap (methods x methods) . full_res10



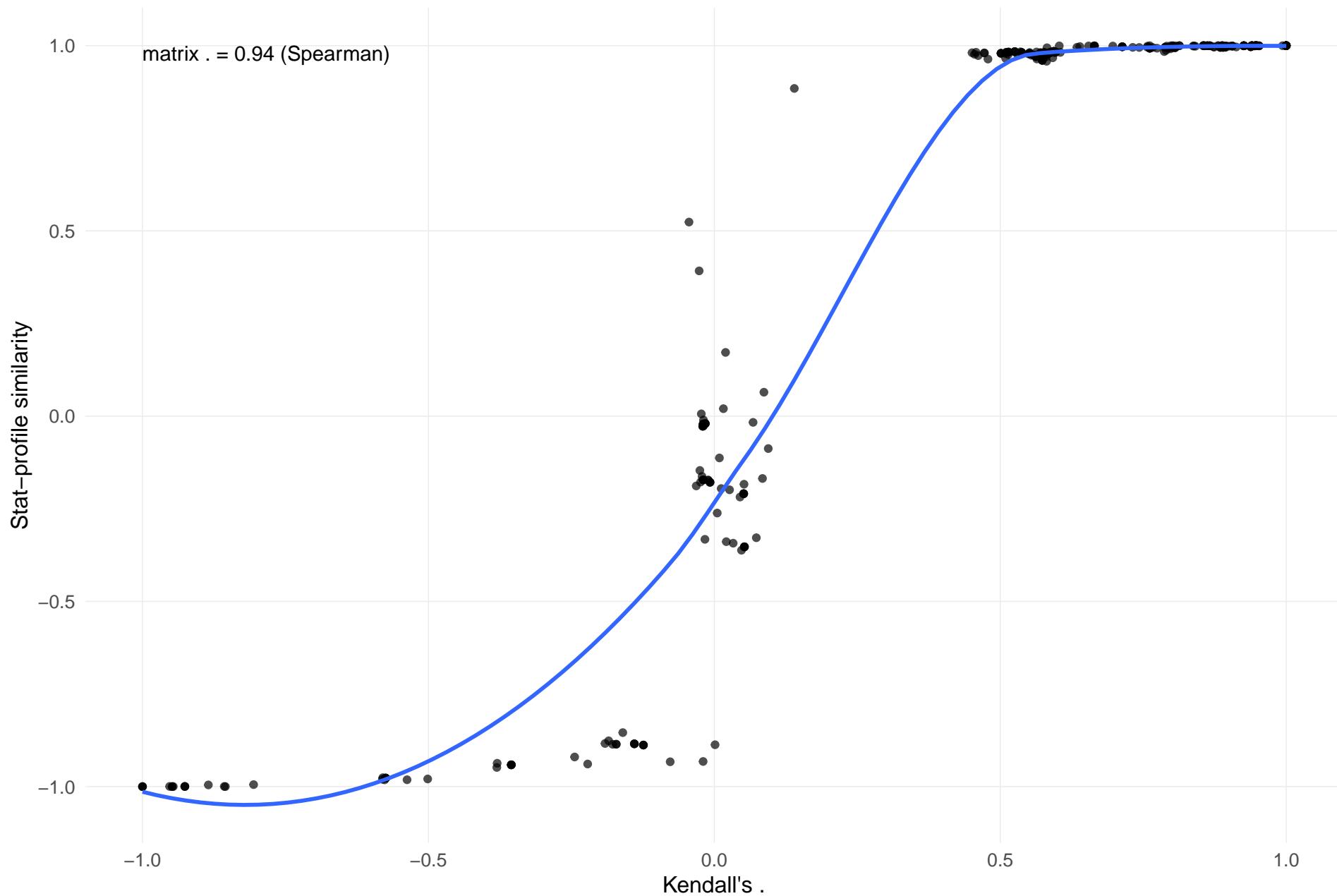
Method x Summary-stat Kendall correlation (with p-stars) . full_res10



Kendall correlation between plasticity score realisations . full_res10



Linking . to stat-profile similarity . full_res10



PCA biplot of method–stat Kendall correlations . full_res10

