metric = adjusted_rand_score | method_choice = bonobo

metric = adjusted_rand_score | method_choice = sweet

metric = adjusted_rand_score | method_choice = lioness

metric = adjusted_rand_score | method_choice = spcc

metric = adjusted_mutual_info_score | method_choice = bonobo

metric = adjusted_mutual_info_score | method_choice = sweet

metric = adjusted_mutual_info_score | method_choice = lioness

metric = adjusted_mutual_info_score | method_choice = spcc

metric = completeness_score | method_choice = bonobo

metric = completeness_score | method_choice = sweet

metric = completeness_score | method_choice = lioness

metric = completeness_score | method_choice = spcc

metric = fowlkes_mallows_score | method_choice = bonobo

metric = fowlkes_mallows_score | method_choice = sweet

metric = fowlkes_mallows_score | method_choice = lioness

metric = fowlkes_mallows_score | method_choice = spcc

value

0.8
0.6
0.4
0.2
0.0
−0.2

kmeans
spectral
ac_ward

clustering_method

check
media
genetics