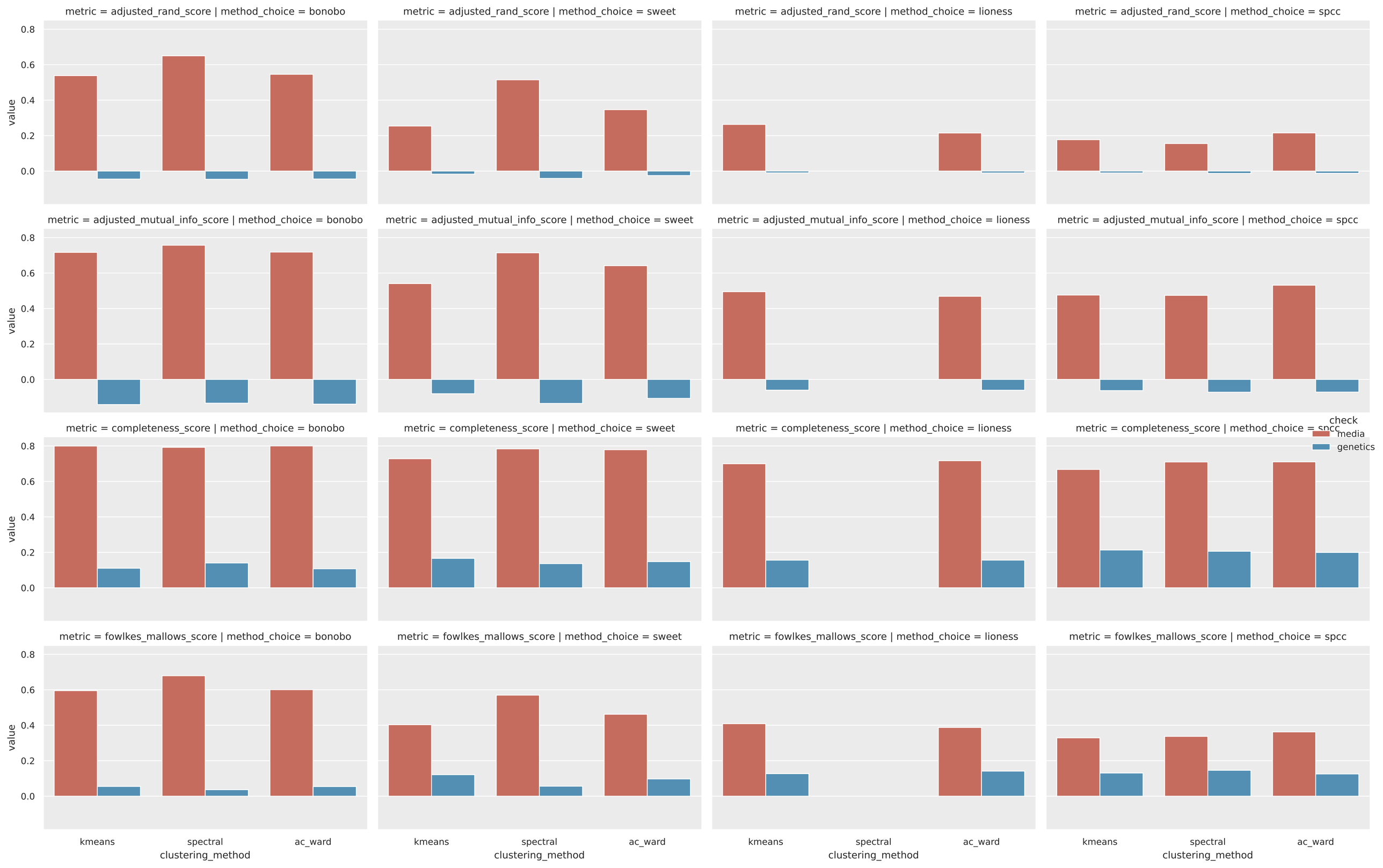
metric = adjusted_rand_score | method_choice = bonobo
metric = adjusted_rand_score | method_choice = sweet
metric = adjusted_rand_score | method_choice = lioness
metric = adjusted_rand_score | method_choice = spcc
metric = adjusted_mutual_info_score | method_choice = bonobo
metric = adjusted_mutual_info_score | 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
0.0
0.2
0.4
0.6
0.8
kmeans
spectral
ac_ward
clustering_method
check
media
genetics