

Cluster stability analysis

gap: 1.00, Tp: 2.00, single link

Cluster Number	J_median	D_median	A_median	J_avg	D_avg	A_avg	J_std	D_std	A_std
1 (1)	1.0	0.5	1.0	0.724	0.362	0.724	0.448	0.224	0.448
2 (1)	1.0	0.5	1.0	0.76	0.381	0.808	0.424	0.21	0.395
3 (1)	1.0	0.5	1.0	0.708	0.354	0.708	0.456	0.228	0.456
4 (1)	1.0	0.5	1.0	0.643	0.325	0.728	0.473	0.233	0.446
5 (1)	1.0	0.5	1.0	0.768	0.384	0.768	0.423	0.211	0.423
6 (1)	1.0	0.5	1.0	0.756	0.378	0.756	0.43	0.215	0.43
7 (1)	1.0	0.5	1.0	0.748	0.374	0.748	0.435	0.218	0.435
8 (1)	1.0	0.5	1.0	0.76	0.38	0.76	0.428	0.214	0.428
9 (1)	1.0	0.5	1.0	0.776	0.388	0.776	0.418	0.209	0.418
10 (1)	1.0	0.5	1.0	0.724	0.362	0.724	0.448	0.224	0.448
11 (1)	1.0	0.5	1.0	0.72	0.36	0.72	0.45	0.225	0.45
12 (2)	1.0	0.5	1.0	0.768	0.415	0.768	0.291	0.12	0.291
13 (6)	0.667	0.4	0.75	0.65	0.369	0.729	0.29	0.144	0.192
14 (15)	0.733	0.423	0.733	0.734	0.419	0.745	0.134	0.052	0.119
15 (390)	0.741	0.426	0.741	0.737	0.424	0.738	0.022	0.008	0.022