

Cluster stability analysis

gap: 1.00, Tp: 1.00, 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	0.708	0.354	0.708	0.456	0.228	0.456
2 (1)	1.0	0.5	1.0	0.78	0.39	0.78	0.415	0.208	0.415
3 (1)	1.0	0.5	1.0	0.764	0.382	0.764	0.425	0.213	0.425
4 (1)	1.0	0.5	1.0	0.632	0.321	0.684	0.47	0.232	0.466
5 (1)	1.0	0.5	1.0	0.605	0.305	0.72	0.485	0.24	0.45
6 (1)	1.0	0.5	1.0	0.744	0.372	0.744	0.437	0.219	0.437
7 (1)	1.0	0.5	1.0	0.752	0.376	0.752	0.433	0.216	0.433
8 (1)	1.0	0.5	1.0	0.708	0.354	0.708	0.456	0.228	0.456
9 (1)	1.0	0.5	1.0	0.712	0.356	0.712	0.454	0.227	0.454
10 (1)	1.0	0.5	1.0	0.813	0.407	0.816	0.389	0.194	0.388
11 (1)	1.0	0.5	1.0	0.752	0.376	0.752	0.433	0.216	0.433
12 (1)	1.0	0.5	1.0	0.527	0.269	0.784	0.49	0.24	0.412
13 (1)	1.0	0.5	1.0	0.744	0.372	0.752	0.437	0.219	0.433
14 (1)	1.0	0.5	1.0	0.764	0.382	0.764	0.425	0.213	0.425
15 (4)	0.75	0.429	0.75	0.74	0.415	0.751	0.22	0.082	0.22
16 (15)	0.733	0.423	0.733	0.711	0.411	0.716	0.133	0.052	0.121
17 (41)	0.732	0.423	0.732	0.719	0.416	0.721	0.092	0.036	0.084
18 (175)	0.743	0.426	0.743	0.728	0.42	0.739	0.066	0.026	0.034
19 (175)	0.651	0.394	0.651	0.662	0.397	0.673	0.081	0.032	0.06