

Cluster stability analysis
gap: 1.00, Tp: 0.25, 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	0.696	0.348	0.696	0.461	0.23	0.461
2 (1)	1.0	0.5	1.0	0.716	0.358	0.716	0.452	0.226	0.452
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	0.692	0.348	0.768	0.458	0.226	0.423
5 (1)	1.0	0.5	1.0	0.724	0.362	0.724	0.448	0.224	0.448
6 (1)	1.0	0.5	1.0	0.768	0.384	0.768	0.423	0.211	0.423
7 (1)	1.0	0.5	1.0	0.7	0.35	0.7	0.459	0.23	0.459
8 (1)	1.0	0.5	1.0	0.768	0.384	0.768	0.423	0.211	0.423
9 (2)	1.0	0.5	1.0	0.712	0.389	0.738	0.329	0.146	0.311
10 (2)	0.5	0.333	1.0	0.532	0.289	0.778	0.425	0.213	0.297
11 (4)	0.75	0.429	0.75	0.755	0.42	0.755	0.222	0.081	0.222
12 (13)	0.692	0.409	0.769	0.655	0.376	0.745	0.254	0.13	0.125
13 (18)	0.714	0.417	0.778	0.616	0.358	0.748	0.267	0.14	0.103
14 (31)	0.71	0.415	0.71	0.692	0.404	0.713	0.131	0.063	0.079
15 (346)	0.726	0.421	0.734	0.713	0.416	0.722	0.052	0.02	0.052