

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	0.732	0.366	0.732	0.444	0.222	0.444
2 (1)	1.0	0.5	1.0	0.752	0.376	0.752	0.433	0.216	0.433
3 (1)	1.0	0.5	1.0	0.708	0.354	0.712	0.455	0.227	0.454
4 (1)	1.0	0.5	1.0	0.668	0.336	0.744	0.468	0.232	0.437
5 (1)	1.0	0.5	1.0	0.772	0.386	0.772	0.42	0.21	0.42
6 (1)	1.0	0.5	1.0	0.724	0.362	0.724	0.448	0.224	0.448
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	0.76	0.38	0.76	0.428	0.214	0.428
9 (2)	1.0	0.5	1.0	0.712	0.388	0.742	0.335	0.15	0.314
10 (2)	0.5	0.333	1.0	0.51	0.282	0.76	0.413	0.208	0.301
11 (4)	0.75	0.429	0.75	0.785	0.433	0.785	0.195	0.067	0.195
12 (13)	0.692	0.409	0.769	0.638	0.367	0.74	0.266	0.138	0.125
13 (18)	0.722	0.419	0.778	0.632	0.367	0.755	0.256	0.133	0.101
14 (31)	0.71	0.415	0.71	0.694	0.405	0.717	0.136	0.064	0.086
15 (346)	0.726	0.421	0.734	0.713	0.416	0.722	0.05	0.019	0.05