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	0.8	0.4	0.8	0.401	0.2	0.401
2 (1)	1.0	0.5	1.0	0.744	0.372	0.744	0.437	0.219	0.437
3 (1)	1.0	0.5	1.0	0.764	0.382	0.768	0.425	0.212	0.423
4 (1)	1.0	0.5	1.0	0.613	0.309	0.724	0.483	0.239	0.448
5 (1)	1.0	0.5	1.0	0.748	0.374	0.748	0.435	0.218	0.435
6 (1)	1.0	0.5	1.0	0.704	0.352	0.704	0.457	0.229	0.457
7 (1)	1.0	0.5	1.0	0.78	0.39	0.78	0.415	0.208	0.415
8 (1)	1.0	0.5	1.0	0.804	0.402	0.804	0.398	0.199	0.398
9 (2)	1.0	0.5	1.0	0.708	0.386	0.744	0.335	0.15	0.311
10 (2)	0.5	0.333	1.0	0.542	0.3	0.742	0.404	0.202	0.301
11 (4)	0.75	0.429	0.75	0.768	0.426	0.768	0.206	0.074	0.206
12 (13)	0.714	0.417	0.769	0.663	0.379	0.751	0.252	0.13	0.118
13 (18)	0.667	0.4	0.75	0.598	0.35	0.742	0.276	0.145	0.11
14 (31)	0.71	0.415	0.71	0.697	0.406	0.723	0.134	0.065	0.077
15 (346)	0.728	0.421	0.737	0.716	0.416	0.725	0.054	0.022	0.053