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	0.74	0.375	0.772	0.421	0.208	0.42
2 (1)	1.0	0.5	1.0	0.736	0.368	0.736	0.442	0.221	0.442
3 (1)	1.0	0.5	1.0	0.652	0.326	0.756	0.477	0.238	0.43
4 (1)	1.0	0.5	1.0	0.724	0.362	0.728	0.448	0.224	0.446
5 (1)	1.0	0.5	1.0	0.704	0.357	0.736	0.439	0.218	0.442
6 (1)	1.0	0.5	1.0	0.736	0.368	0.736	0.442	0.221	0.442
7 (1)	1.0	0.5	1.0	0.788	0.394	0.788	0.41	0.205	0.41
8 (1)	1.0	0.5	1.0	0.7	0.35	0.708	0.459	0.23	0.456
9 (416)	0.743	0.426	0.743	0.736	0.424	0.736	0.012	0.004	0.013