

ASAP Web results

date:2023-05-22T10:11:52
 input file:Richtersius_COI.fas
 nb of sequences:42
 length of seqs:641
 subst. model:JC69 Jukes-Cantor
 recurs split pval:1.000000e-02

Split probability: 0.010000 sequence length: 641

Save Spart text File [here](#)

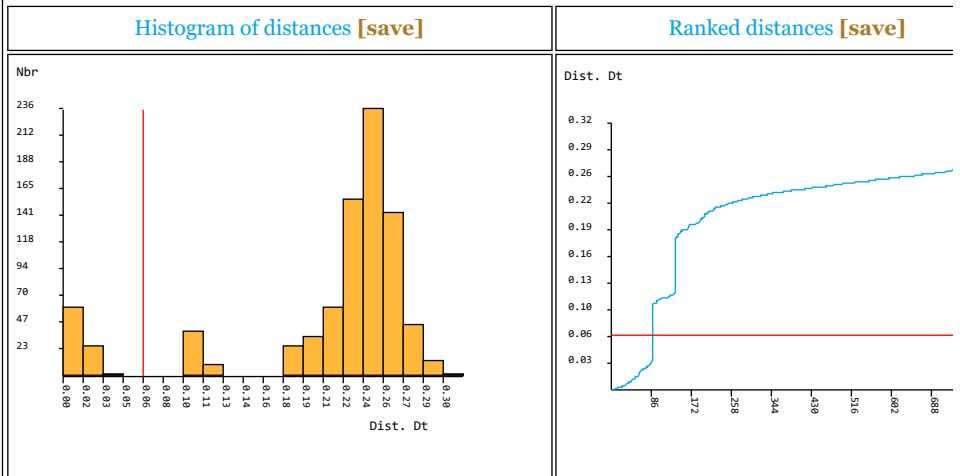
Save Spart XML File [here](#)

10 best partitions found by ASAP (see [FAQ](#) for more details)

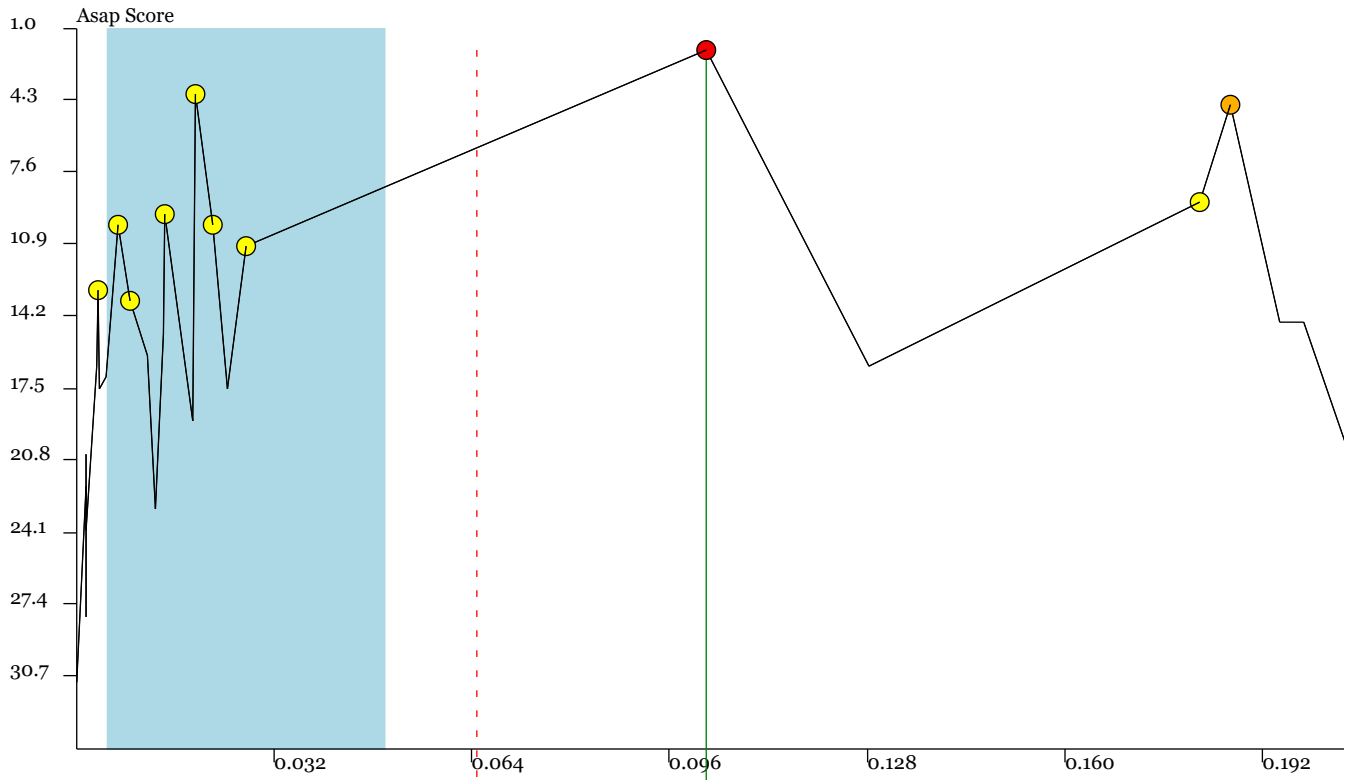
The lower the score, the better the partition

Nb of subsets	asap-score	P-val (rank)	W (rank)	Threshold dist.	Text
12	1.00	6.69e-03 (1)	1.68e-03 (1)	0.064821	list csv
* 16	3.00	4.01e-01 (4)	1.53e-03 (2)	0.019095	list csv
9	3.50	9.44e-02 (2)	9.52e-04 (5)	0.184329	list csv
11	8.00	7.50e-01 (12)	1.23e-03 (4)	0.155090	list csv
* 18	8.50	5.53e-01 (7)	5.97e-04 (10)	0.014312	list csv
* 24	9.00	4.95e-01 (6)	5.48e-04 (12)	0.005852	list csv
* 15	9.00	6.63e-01 (9)	6.22e-04 (9)	0.020699	list csv
* 13	10.00	3.51e-01 (3)	3.96e-04 (17)	0.025974	list csv
27	12.00	4.71e-01 (5)	3.52e-04 (19)	0.003417	list csv
* 23	12.50	7.72e-01 (14)	5.90e-04 (11)	0.007760	list csv

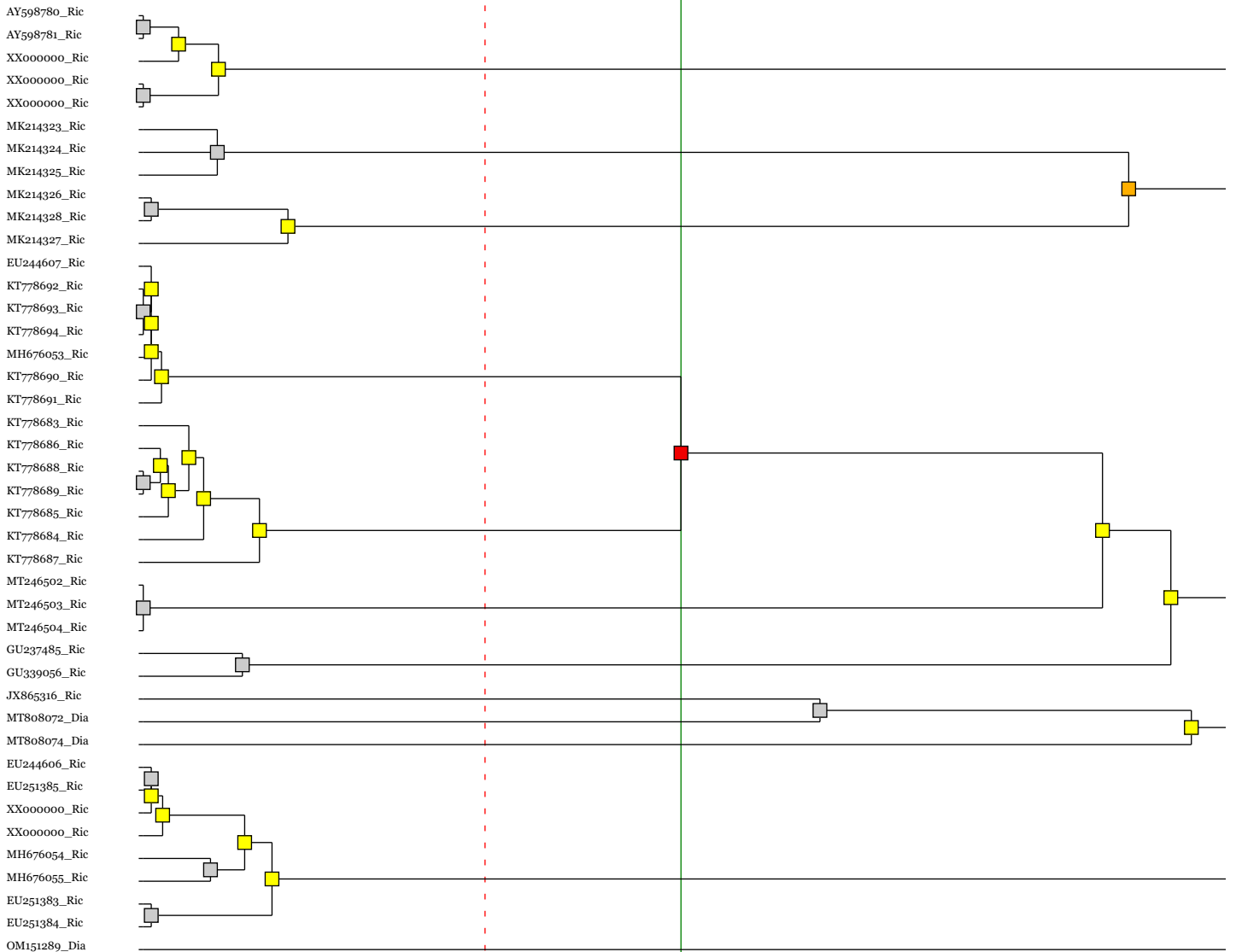
Number of subsets in this table equals to the number of primary species hypotheses in the respective partition



[View/Save Boxed subsets graph here](#)



Legend: ● <0.001 ● <0.05 ● <0.1 ● >0.1 ● N/A



[View/save curves and dendrogram here](#)

Responsive Crossing lines Legend: Green Line= grouping distance (Dc)- Red line = treshold distance(Dt)

Running time: 0 min 4 seconds