

ASAP Web results

date:2023-05-22T10:11:52
input file:extra-mohave.afas
nb of sequences:100
length of seqs:1140
subst. model:K80 Kimura
recurs split pval:1.000000e-02











Split probability: 0.010000 sequence length: 1140

Save Spart text File [here](#)

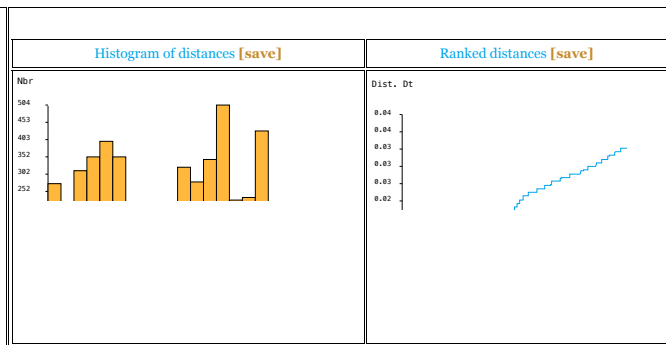
Save Spart XML File [here](#)

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

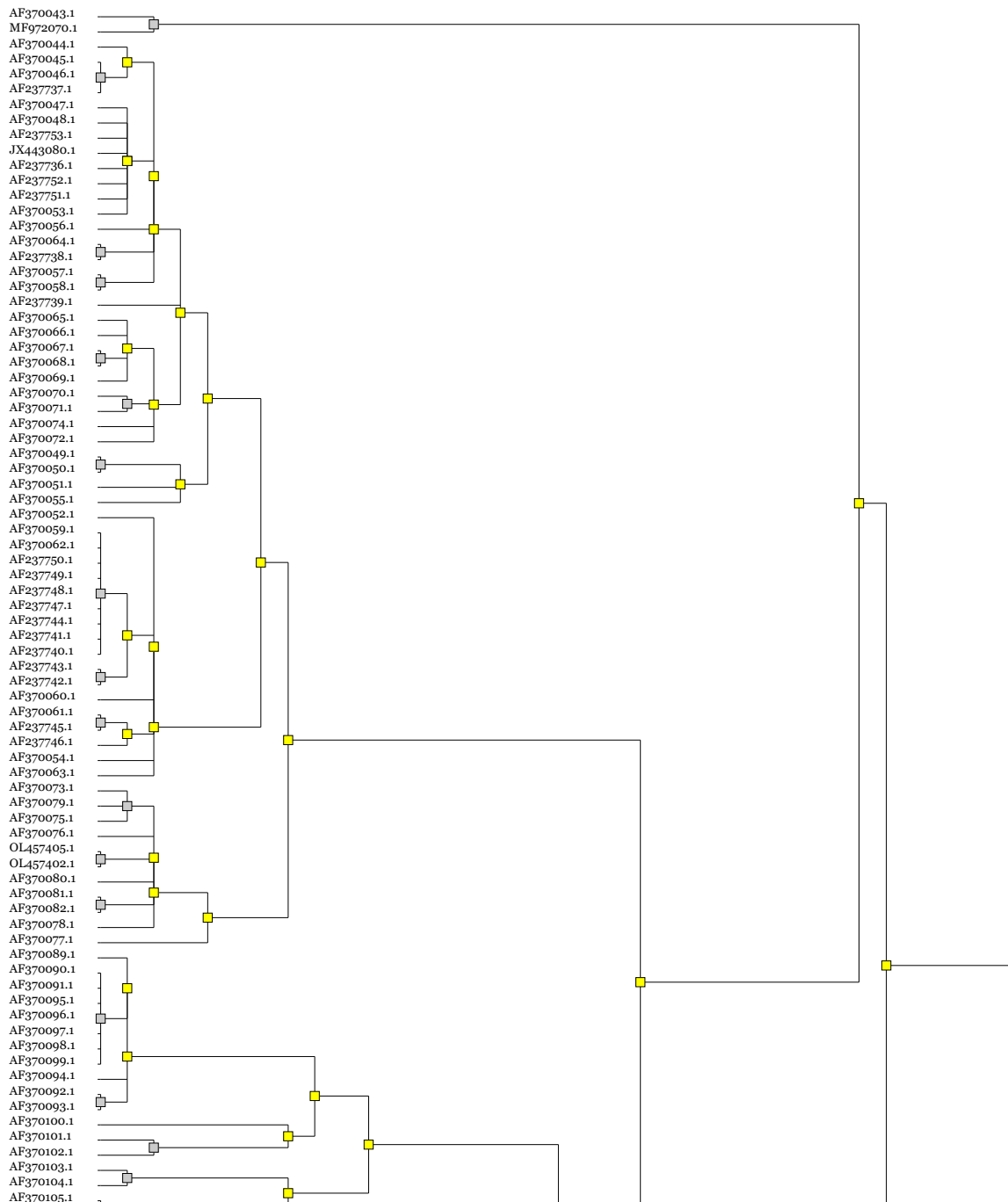
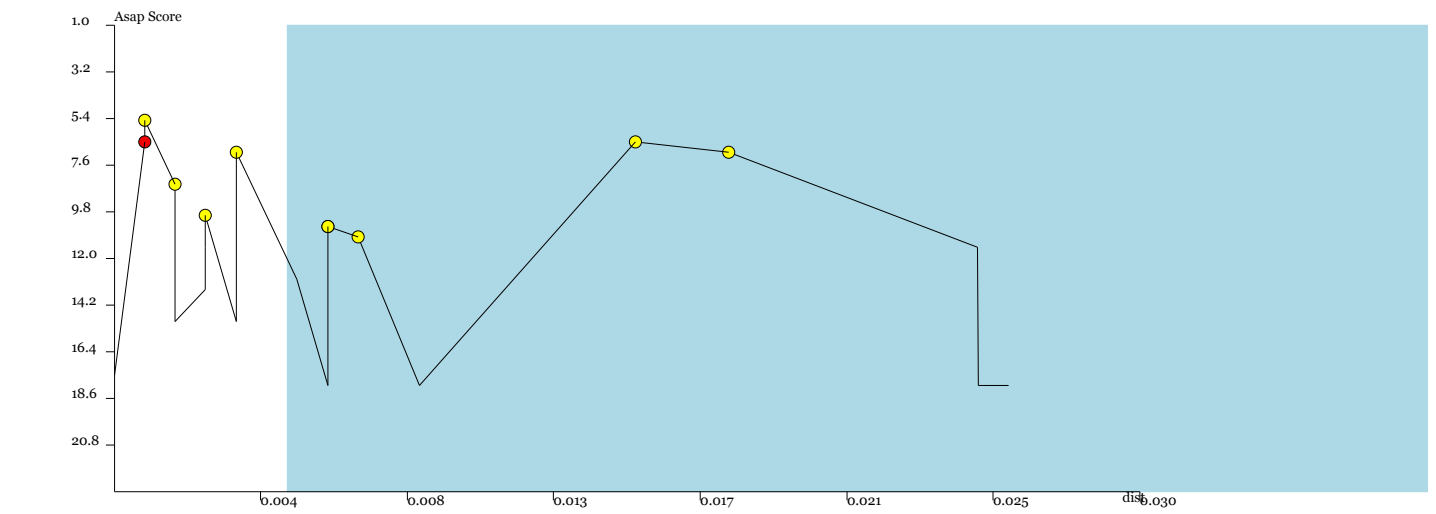
The lower the score, the better the partition

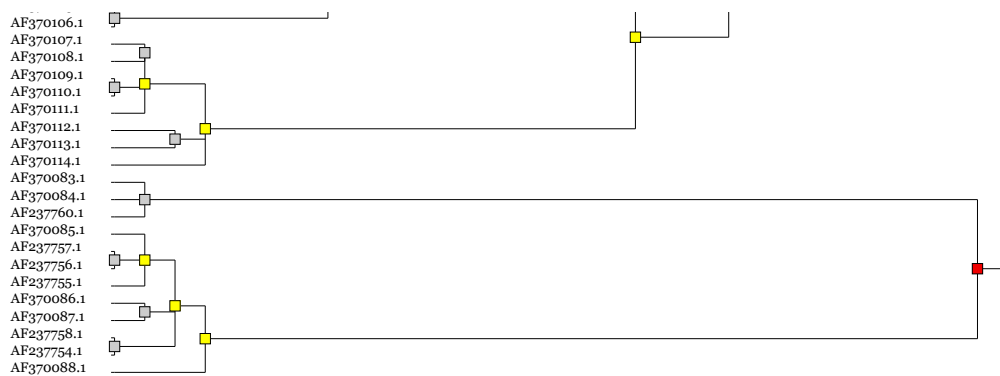
Nb of subsets	asap-score	P-val (rank)	W (rank)	Threshold dist.	Text
53	4.50	 8.00e-01 (7)	2.62e-05 (2)	0.000878	list csv
71	5.50	 1.28e-02 (1)	1.26e-05 (10)	0.000439	list csv
* 6	5.50	 8.52e-01 (10)	2.74e-05 (1)	0.011930	list csv
* 5	6.00	 8.28e-01 (8)	2.44e-05 (4)	0.016382	list csv
13	6.00	 8.48e-01 (9)	2.47e-05 (3)	0.003516	list csv
43	7.50	 7.94e-01 (6)	1.28e-05 (9)	0.001317	list csv
15	9.00	 8.82e-01 (11)	1.53e-05 (7)	0.002638	list csv
* 10	9.50	 3.73e-01 (4)	1.03e-05 (15)	0.006163	list csv
* 9	9.50	 7.27e-01 (5)	1.03e-05 (14)	0.006168	list csv
* 8	10.00	 1.34e-01 (3)	8.59e-06 (17)	0.006610	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](#)





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

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

Running time: 0 min 16 seconds