

SUPPLEMENTARY MATERIAL

ASAP method

Split probability: 0.010000 sequence length: 677

Save Spart File here

9 best partitions found by ASAP (see FAQ for more details)

The lower the score, the better the partition

Nb of species	asap-score	P-val (rank)	W (rank)	Threshold dist.	Text
* 7	2.50	8.03e-04 (1)	4.91e-04 (4)	0.012491	list csv
* 7	3.50	1.82e-01 (2)	4.75e-04 (5)	0.020786	list csv
* 8	3.50	3.07e-01 (4)	5.26e-04 (3)	0.005892	list csv
3	3.50	5.39e-01 (6)	9.35e-04 (1)	0.057153	list csv
* 4	5.00	7.37e-01 (8)	5.73e-04 (2)	0.030217	list csv
* 6	5.50	5.33e-01 (5)	4.34e-04 (6)	0.025861	list csv
14	6.00	2.30e-01 (3)	1.33e-04 (9)	0.001403	list csv
9	7.00	6.89e-01 (7)	3.96e-04 (7)	0.004215	list csv
2	8.50	9.60e-01 (9)	2.79e-04 (8)	0.087565	list csv

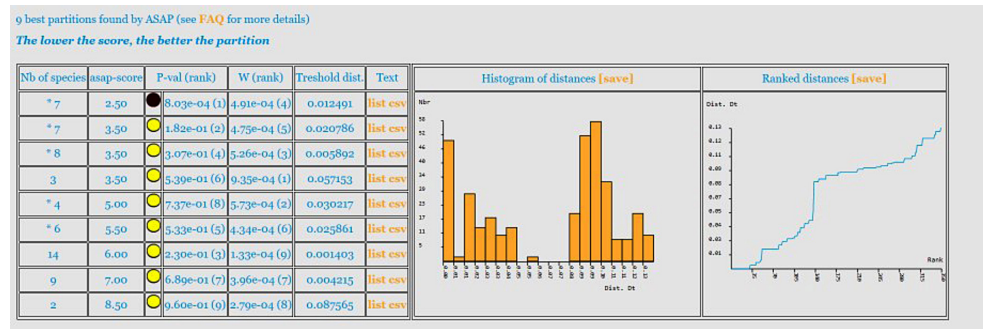


FIGURE S3 | Number of groups obtained with ASAP method.



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