

FIGURE S1 | ASAP delimitation results.

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ASAP Results

ASAP Web results

date:2023-05-22T10:11:52
input file:alinhamento_mamuruensis_asap.fasta
nb of sequences:17
length of seqs:621
subst. model:JC69 Jukes-Cantor
recurs split pval:1.000000e-02

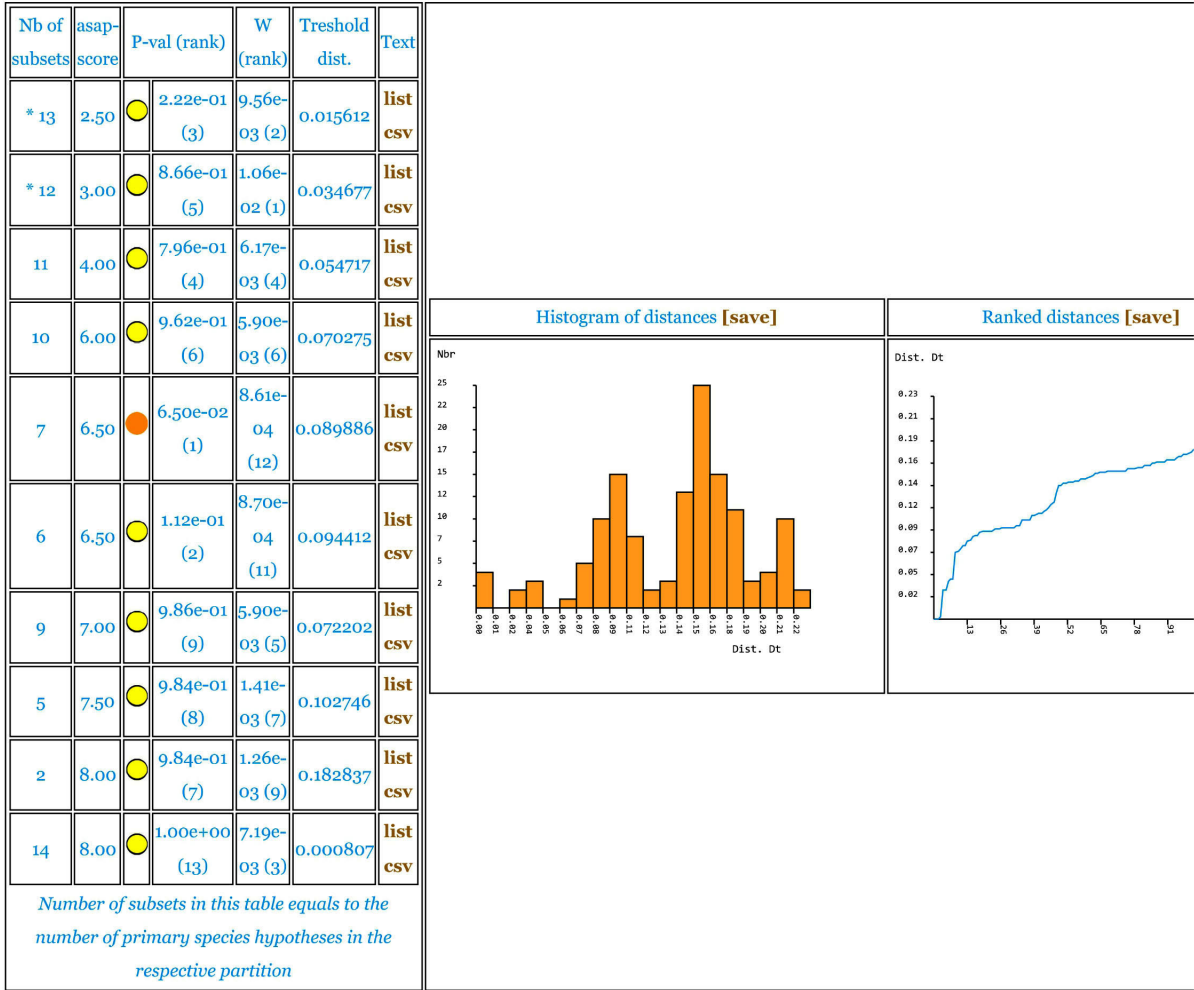
Split probability: 0.010000 sequence length: 621

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



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

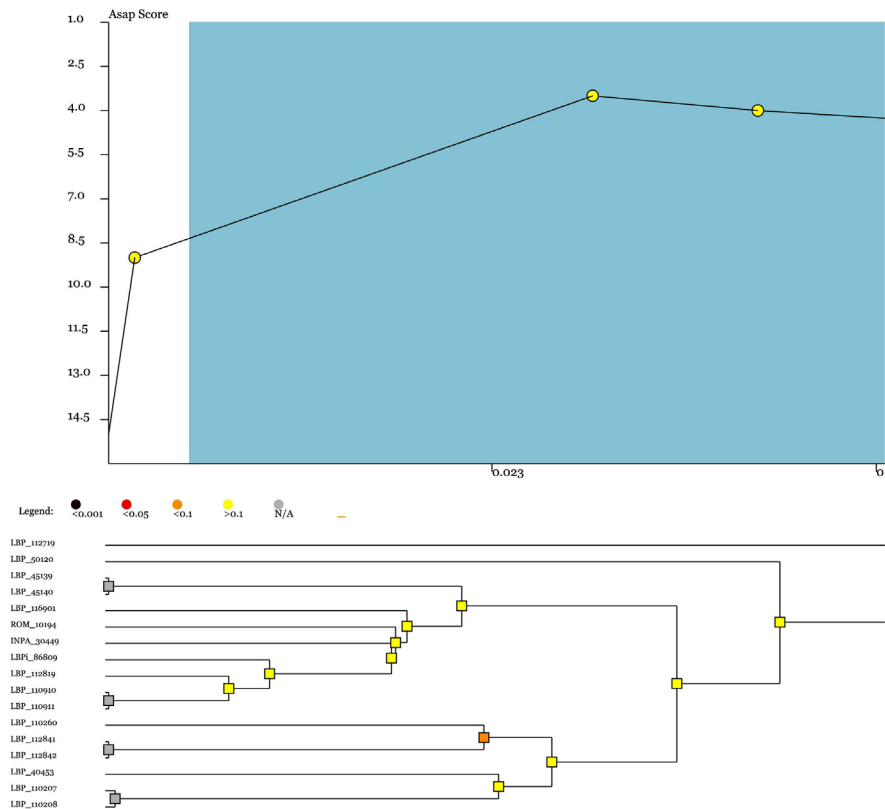
FIGURE S1 | (Continued)

<https://bioinfo.mnhn.fr/abi/public/asap/temp/11610.1804289383/results.html>

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ASAP Results



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

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

Running time: 0 min 0 seconds

Neotropical Ichthyology

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