

SuppFig 1: Supertree of 28S rRNA, Consensus of 498000 trees.

GenBank accession numbers in sequence name.

Numbers on branches give percent of topologies supporting a branch.

SSD : Sargasso Sea Database scaffold.

*SSD : SSD scaffold for which topology enables to assign taxonomic affiliation (branch supported by 80% of topologies)

S6 : paired reads from sample 6 (see Method)

A : Anchor dataset sequence.

BBH : GenBank Best Blast Hit sequence.

