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.