**ARB DATABASE FOR SPONGE-DERIVED 16S rRNA SEQUENCES**

From:  Taylor, M.W., Radax, R., Steger, D., and Wagner, M. (2007) Sponge-associated microorganisms: evolution, ecology and biotechnological potential. *Microbiology and Molecular Biology Reviews* 71: 295-347.

A few minor things to know when using this database:

(1) All of the maximum-likelihood trees from the MMBR article are included.

(2) Under "Species → Configurations" you can find the various (sponge-derived) microbial taxa. By marking a particular configuration you can select all sponge-derived sequences from that taxon.

(3) These fields contain useful sponge-related information (in addition to the standard ARB fields such as Name, Fullname etc.):

Keywords: species of host sponge (if known)
Habitat: sample collection location (if known)


If you need any additional information please contact Mike Taylor at mw.taylor@auckland.ac.nz