

The Tree of Life

This tree was generated by David M. Hillis, Derrick Zwickl, and Robin Gutell (University of Texas) based on an analysis of small subunit rRNA sequences obtained from about 3,000 species from throughout the Tree of Life. When possible, sequences were chosen so that the number of species included in a major group is roughly proportional to the number of known species in that group. The total number of species included is approximately the square-root of the number of species thought to exist on Earth.

