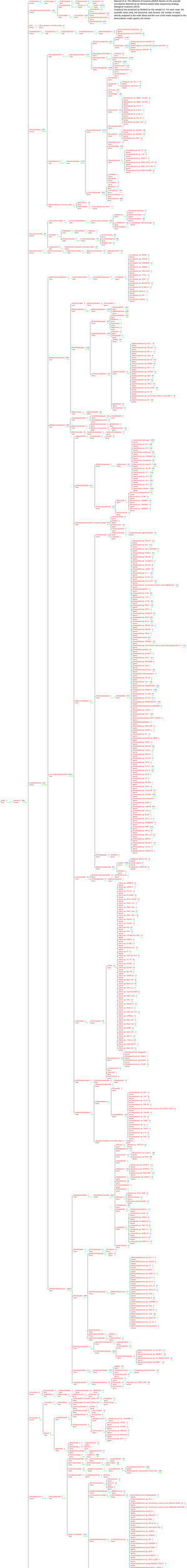
Mansour et al. The influence of invasive phyllish blooms on the aquatic microbiome: a multi-omics study using deep sequencing strategy. Environmental Research (2014)

Graphical tree produced by BioMed for the sample E1. For each node, the number of reads assigned to the node (clade) and the number of reads directly assigned to the node (blue) and the sum of the reads assigned to its descendants nodes (green) are shown.