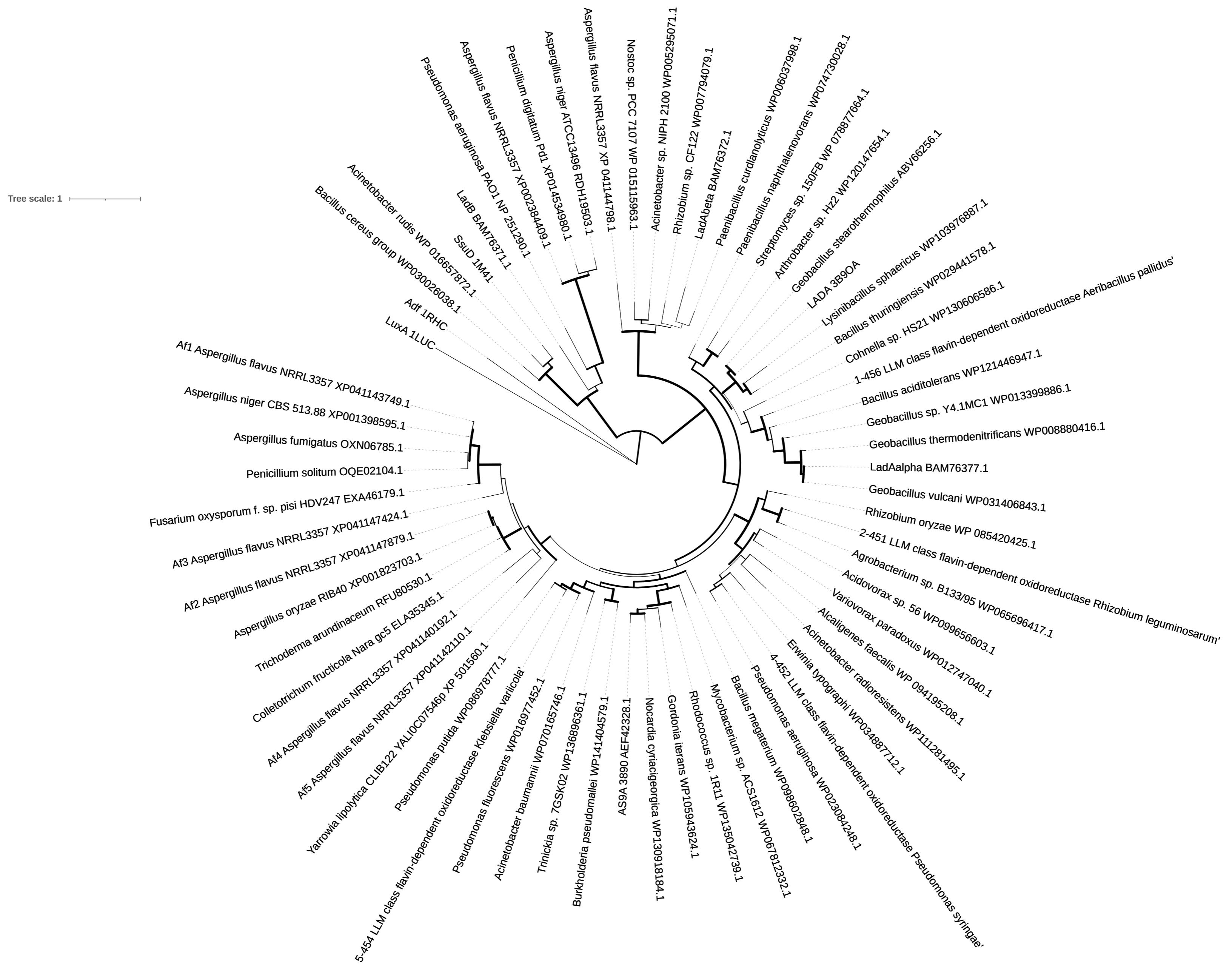


Supplementary Figure 2. Maximum Likelihood phylogenetic tree with GenBank accession numbers correspond to all leaf labels in Figure 2



**Supplementary Figure 2B. Similarity matrix of the A representative sample (n=63) of the 260 derived protein sequences of the identified LadA-like FMN-dependent monooxygenases in bacteria and fungi**

