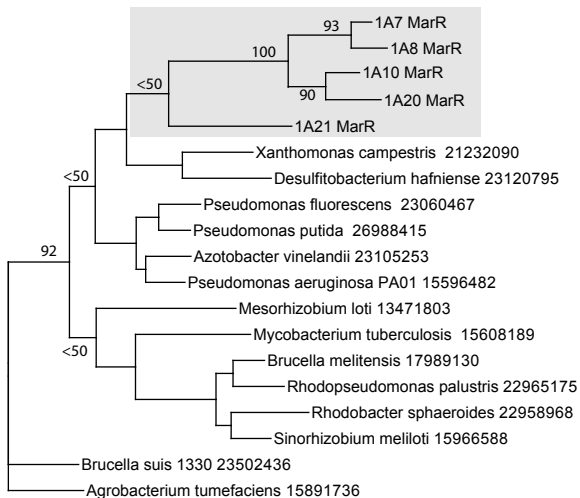
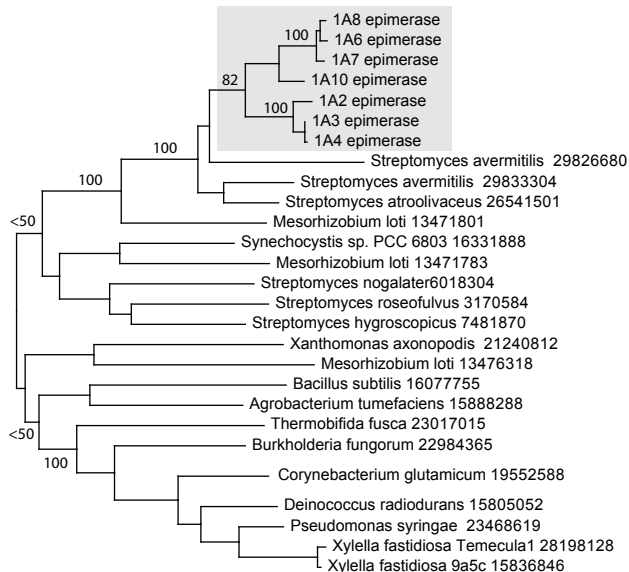


PF01047: MarR transcriptional regulator
 COG1846: MarR transcriptional regulator



Ln Likelihood = -3711.40

PF01370: NAD dependent epimerase/dehydratase
 COG0702: Predicted nucleoside-diphosphate-sugar epimerases



Ln Likelihood = -11080.81284