

SUPPLEMENTARY FIGURE 5

Phylogeny for scytonemin-dedicated AAAB genes of AroB amino acid sequences derived from BEAST analyses. Entries in blue type correspond to homologs found within full scytonemin operons. Bayesian posterior probabilities (BPP) at the nodes are color coded as follows: red for $BPP \geq 0.8$, pink for $0.8 > BPP \geq 0.5$ and white for $BPP \leq 0.5$.

