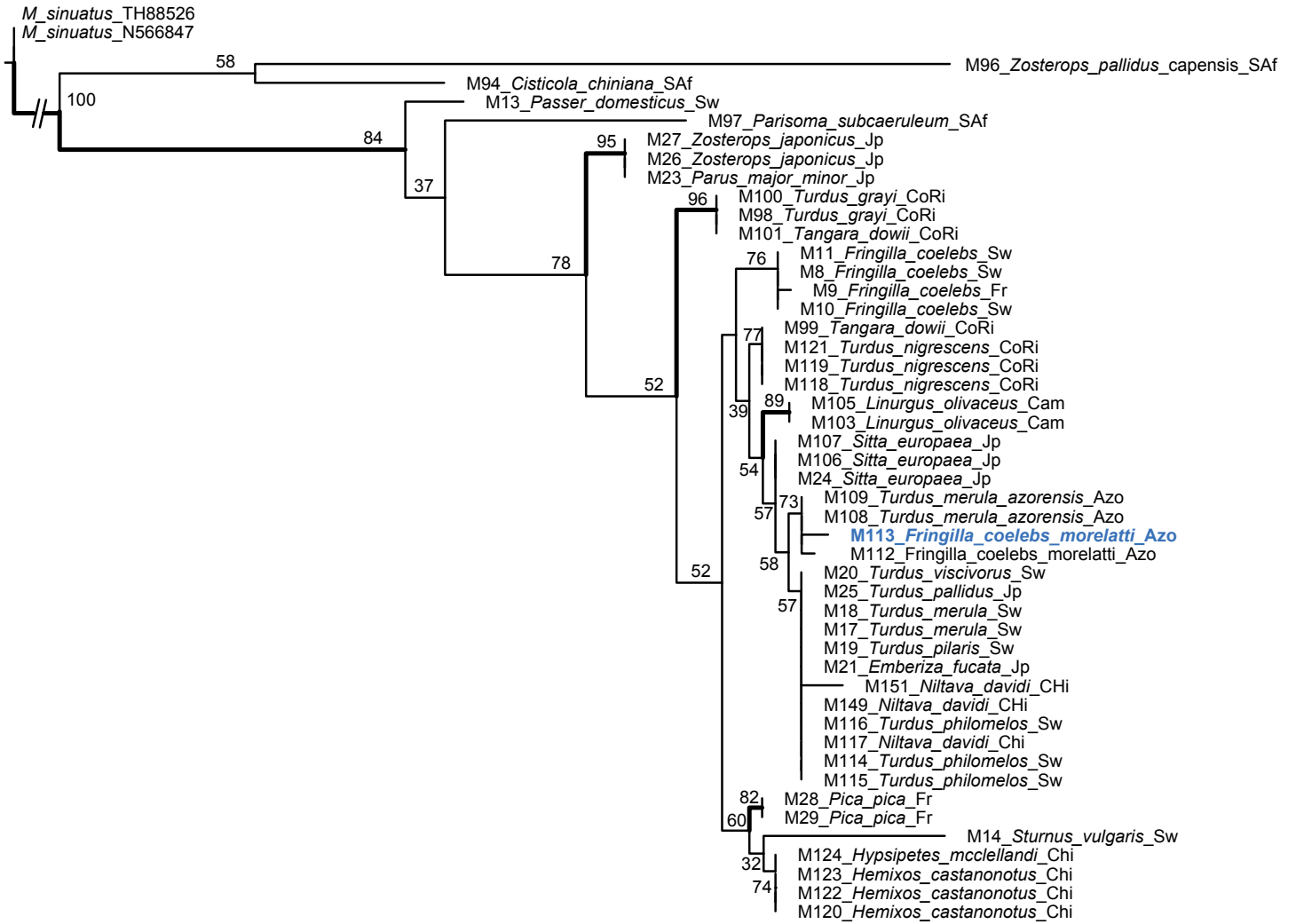


**Supplementary figure 1:** host phylogeny - relationships of *Menacathus eurysternus* samples. The sample used in this study for the metagenomic assembly printed in bold blue.



0.05