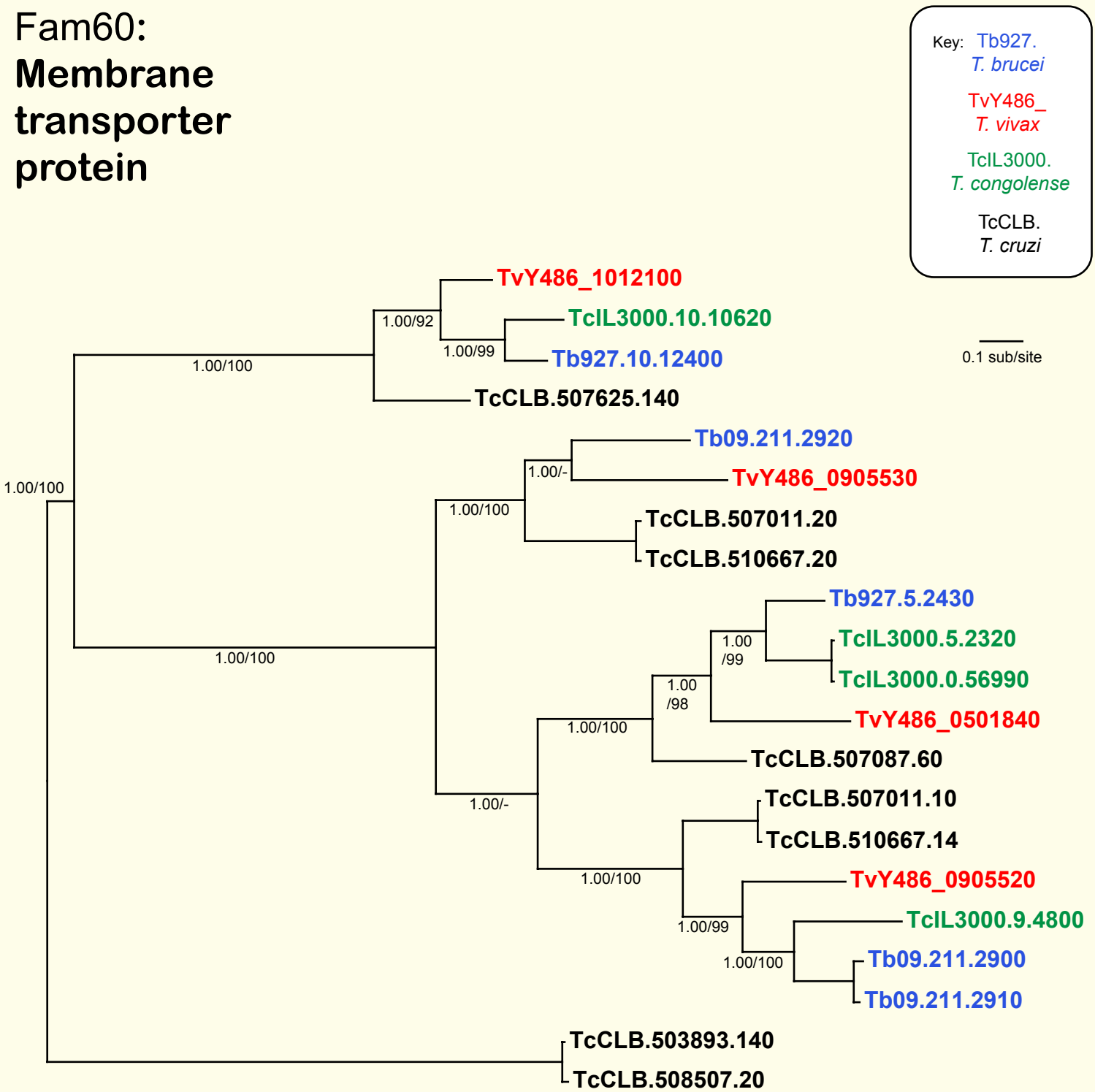
If you use these data, please cite:

Jackson, AP et al. 2012. A cell surface phylome for African Trypanosomes. *Manuscript submitted.*

# Fam60: **Membrane transporter protein**




NOTES: **Fam60** includes various putative membrane transporter genes conserved across *Trypanosoma*. All genes have multiple transmembrane helices.

The Bayesian phylogram was estimated from a multiple nucleotide sequence alignment of 1335 characters, using MrBayes under default settings. The tree is rooted using divergent *T. cruzi* sequences as an outgroup. Selected nodes are supported by posterior probabilities and non-parametric bootstraps generated from a maximum likelihood analysis under a GTR+Γ model.