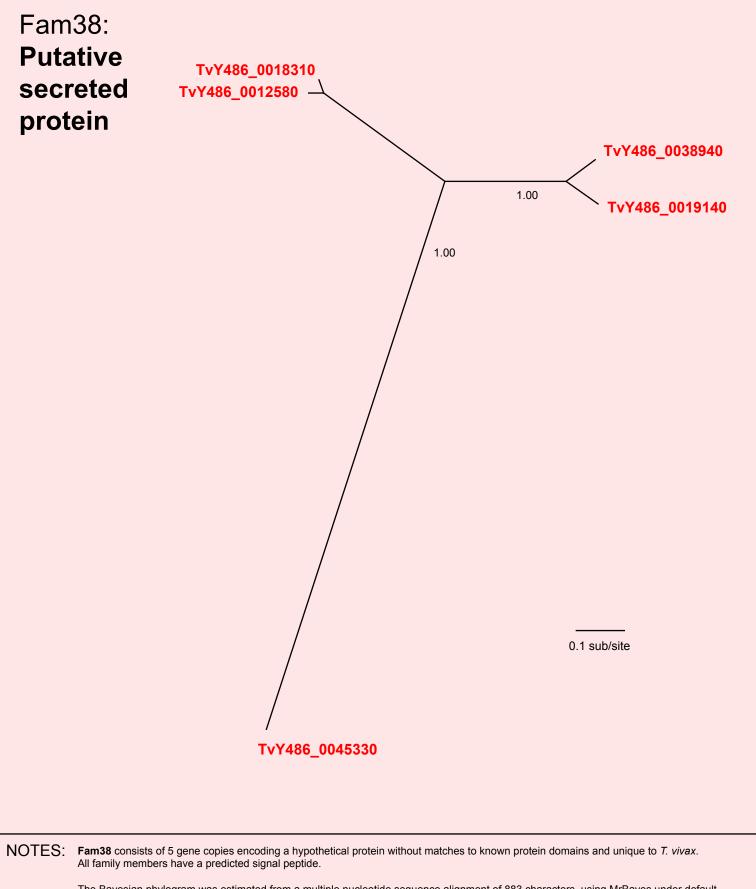
If you use this data, please cite: Jackson, AP et al. 2012. A cell surface phylome for African Trypanosomes. *Manuscript submitted.* 





The Bayesian phylogram was estimated from a multiple nucleotide sequence alignment of 883 characters, using MrBayes under default settings. The tree is unrooted. Selected nodes are supported by posterior probabilities.