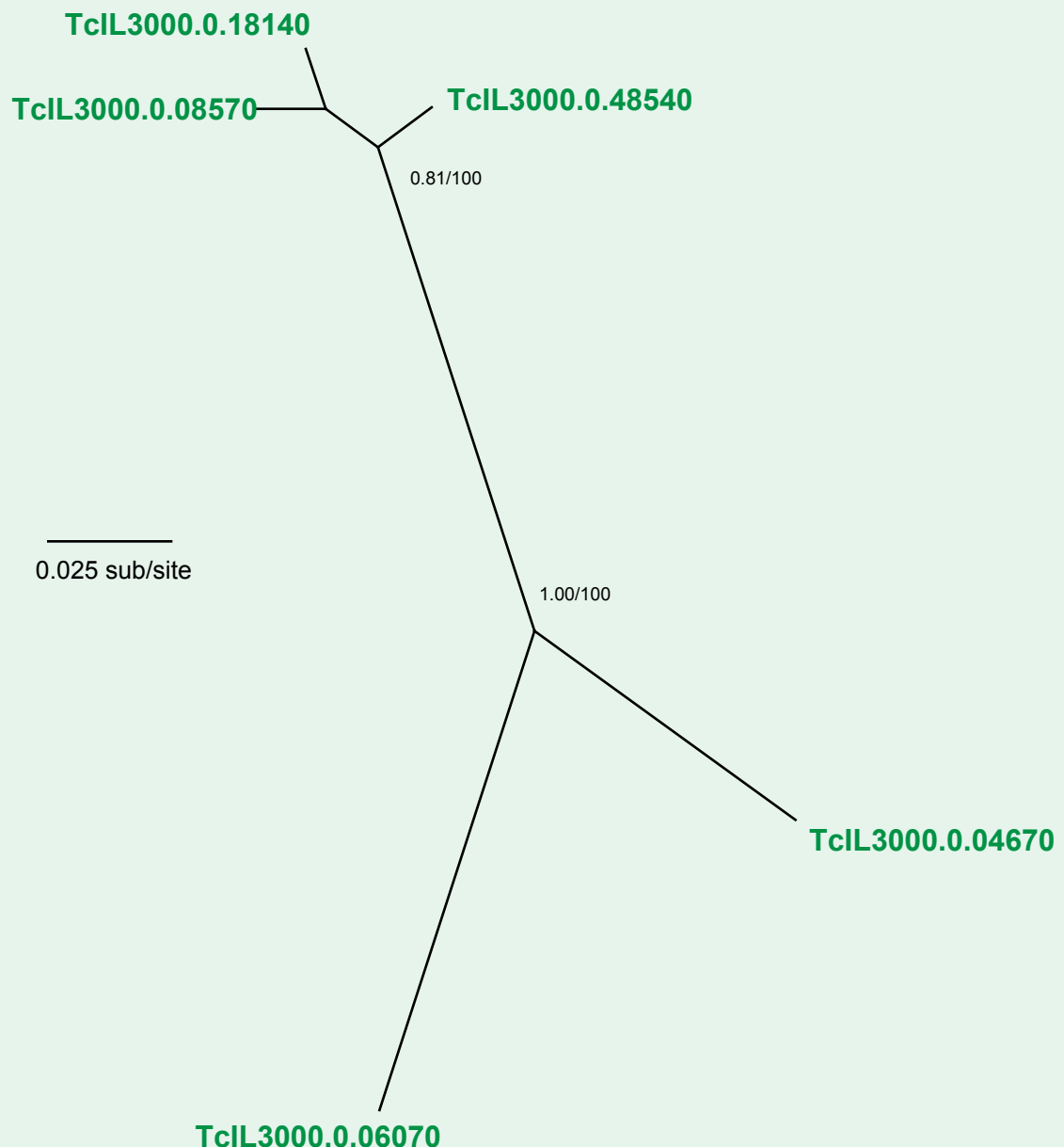


If you use this data, please cite:

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

Fam20: Hypothetical protein



NOTES: **Fam20** comprises five genes encoding hypothetical proteins of 107 amino acids with predicted signal peptides. All gene copies are currently unplaced.

The Bayesian phylogram was estimated from a multiple nucleotide sequence alignment of 321 characters, using MrBayes and a GTR+ Γ model. Selected nodes are supported by posterior probabilities and non-parametric bootstraps generated from a maximum likelihood analysis under a GTR+ Γ model. The tree is unrooted.