



Figure S1. Trypanosoma cruzi infectiveness rates (%) of each species at the tips of the phylogeny and values on the phylogenetic eigenvector 1 (P.E.V. 1) from Agnarsson et al. (2010). Infectiveness rates were determined as the total positive/total examined*100 in hemoculture or xenodiagnosis tests. Values are scales and normalized around 0 to have a comparable scale.

Reference:

Agnarsson I, Kuntner M, May-Collado LJ (2010) Dogs, cats, and kin: a molecular species-level phylogeny of Carnivora. *Mol Phylogenet Evol* 54: 726-745