



Figure S3: Population structure analysis of *C. parapsilosis* strain genomes performed with ADMIXTURE (Alexander et al. 2011) reveals possible admixture in strain N3_182. Each color represents a different ancestral population (N=6) and the proportion of a color in each strain represents how much of the variation in its genome is attributed to that particular ancestral population. All strains except N3_182 are predicted to have a separate ancestral population.