Figure S9: Similarity trees of genome, chromosome II, and the *pot-2* locus

Isotypes are colored by their POT-2 F68I variant status. F=Blue, I=Red. Trees were generated from SNVs (A) genome-wide, (B) using SNVs located on the right arm of chromosome II (13Mb to the terminus), or (C) using SNVs located within the *pot-2* region (II:14,524,173-14,525,111).