

Figure S1

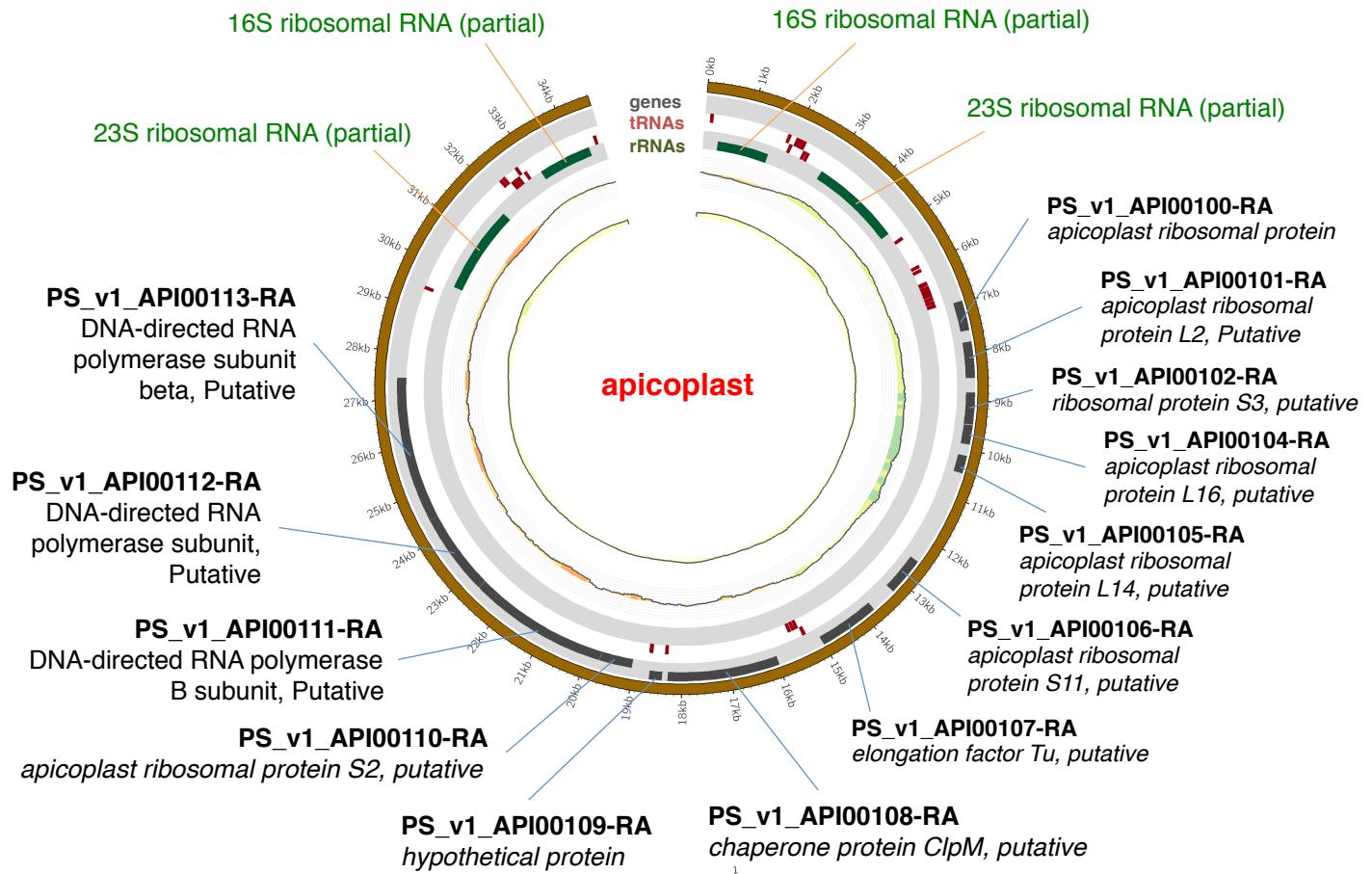


Figure S2

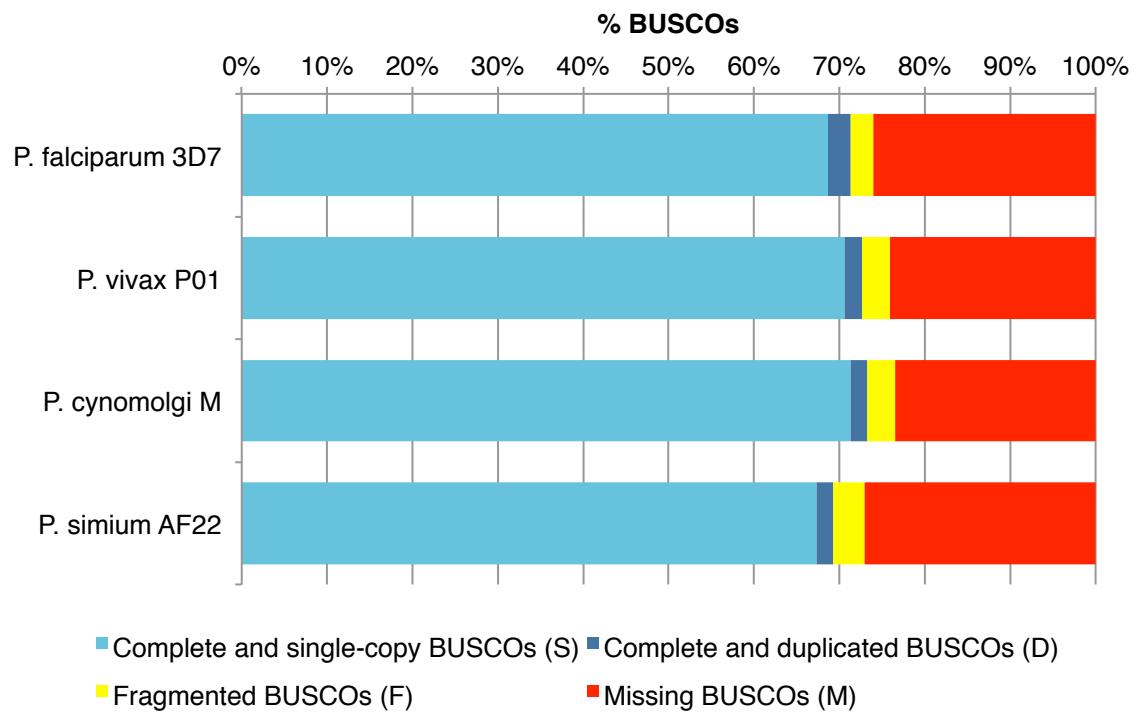


Figure S3

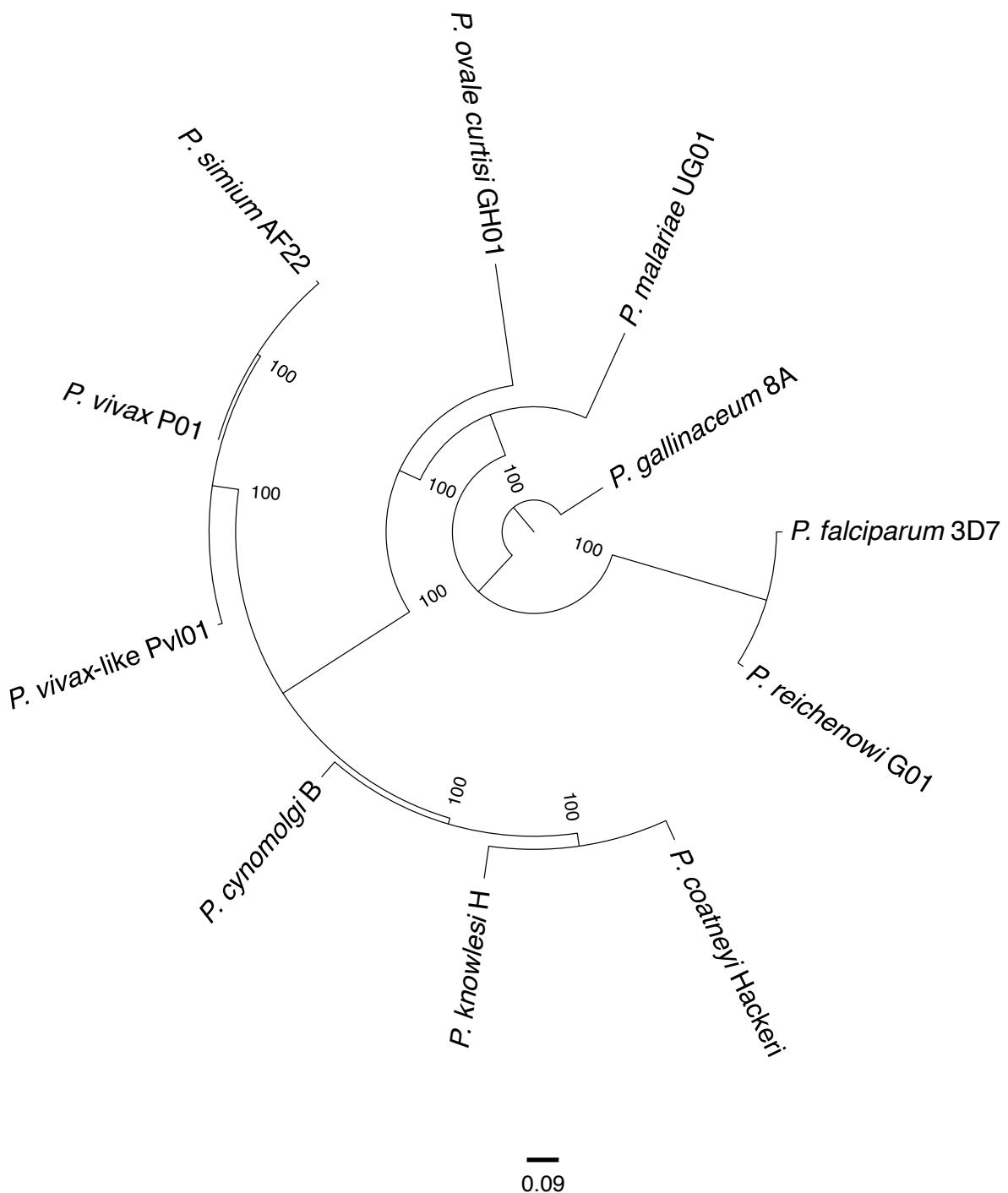
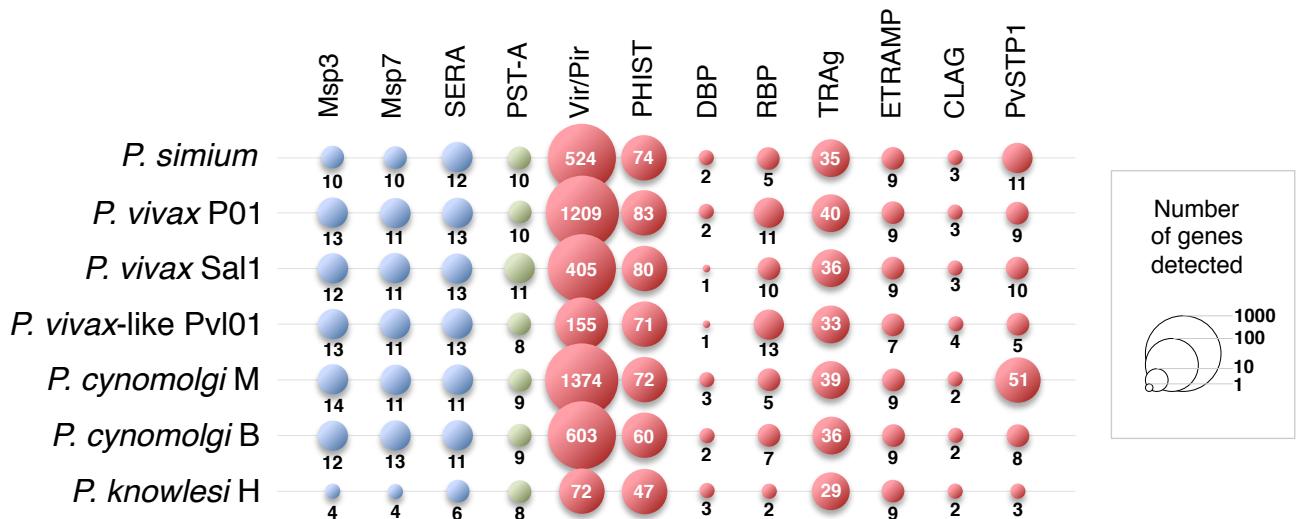


Figure S4

A



B

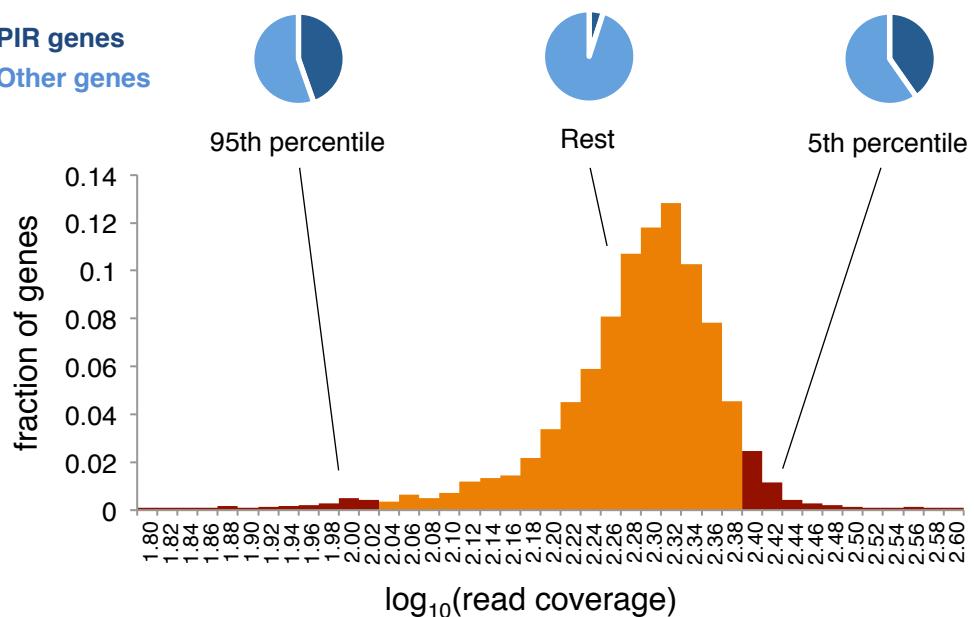


Figure S5

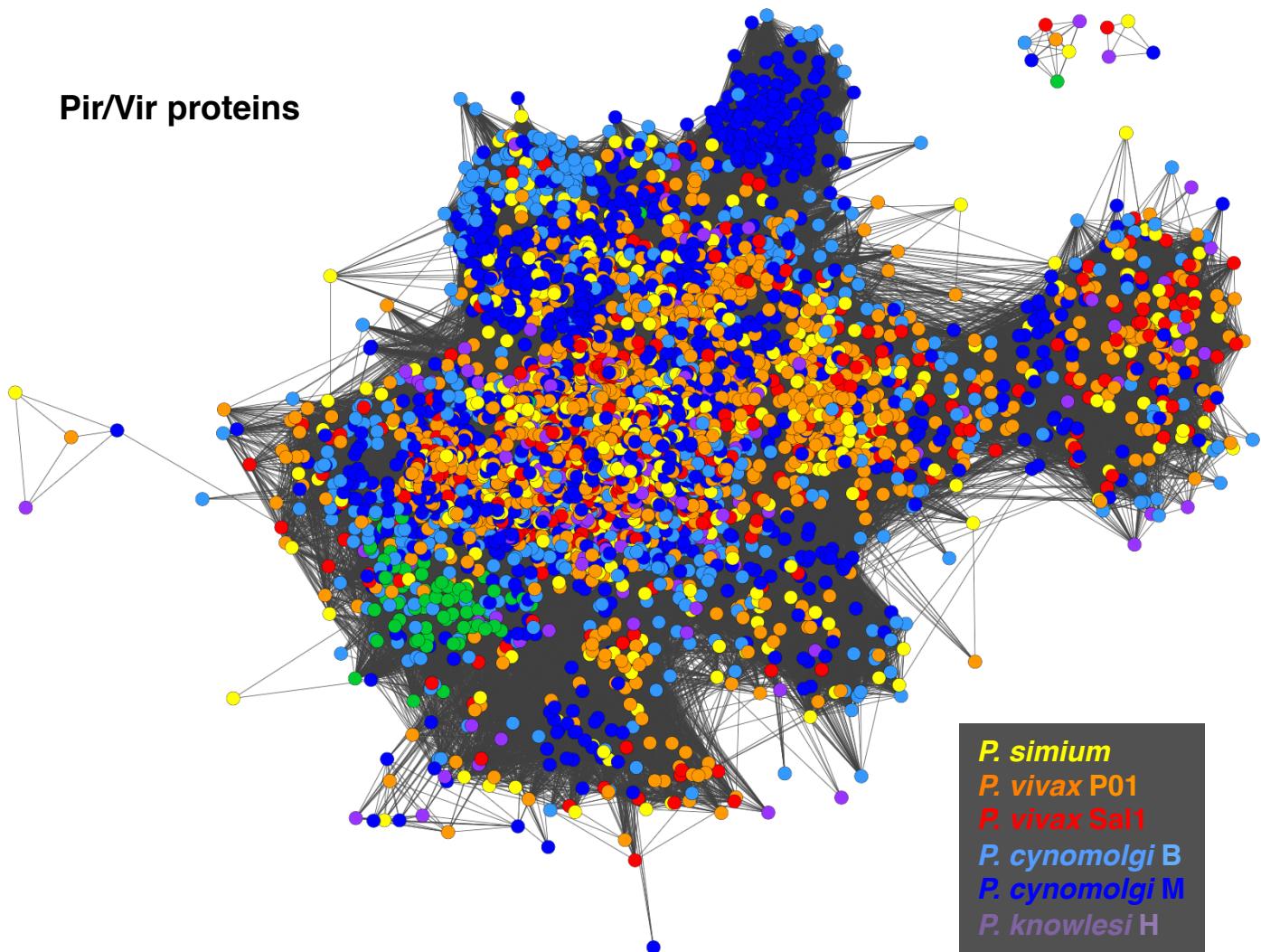


Figure S6

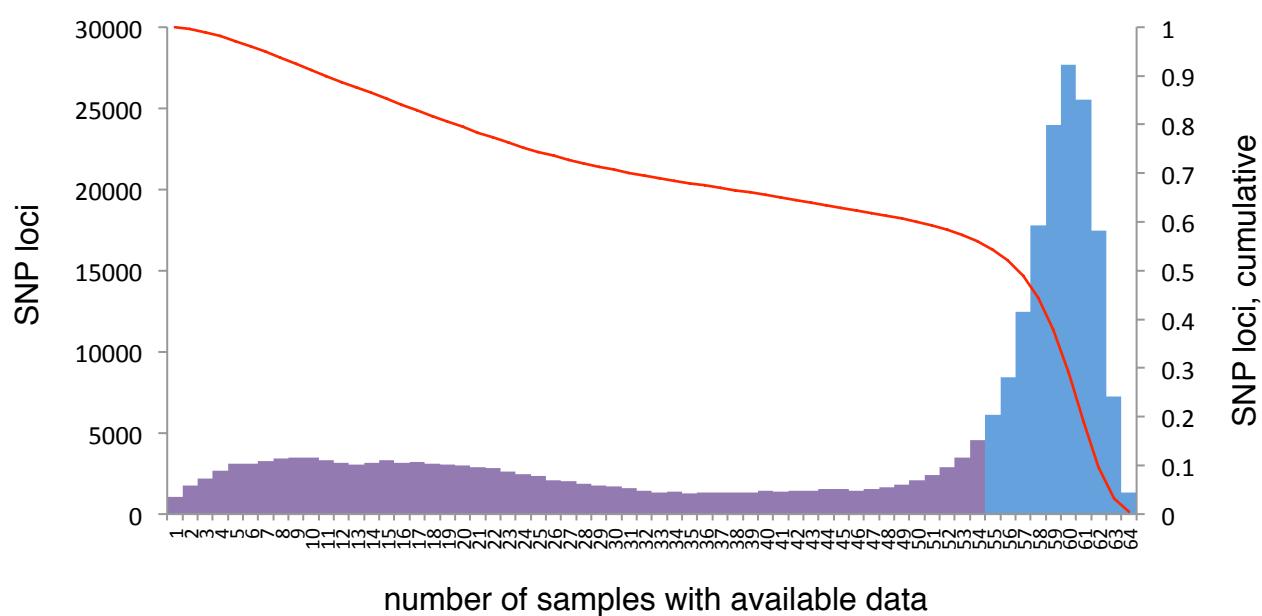
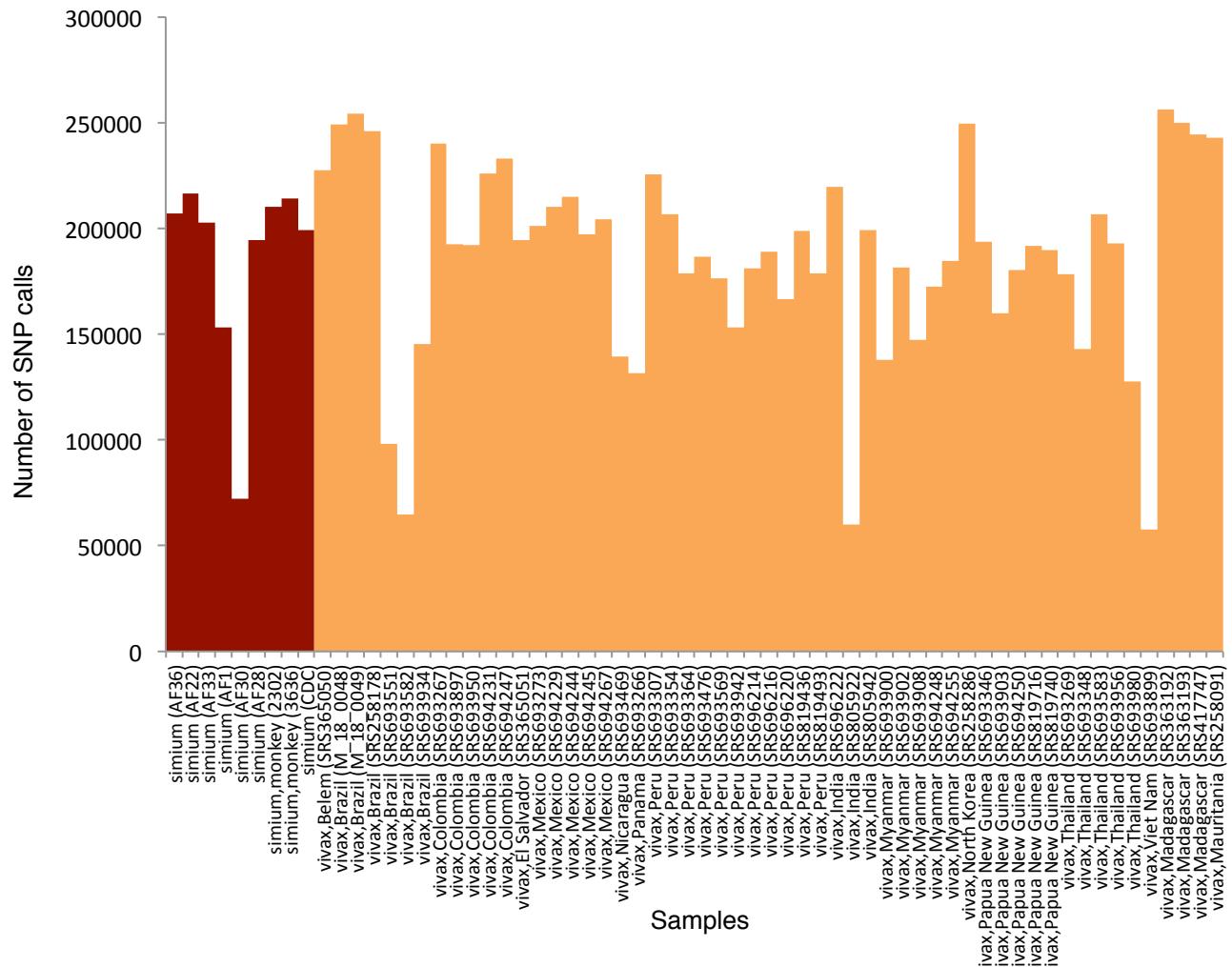
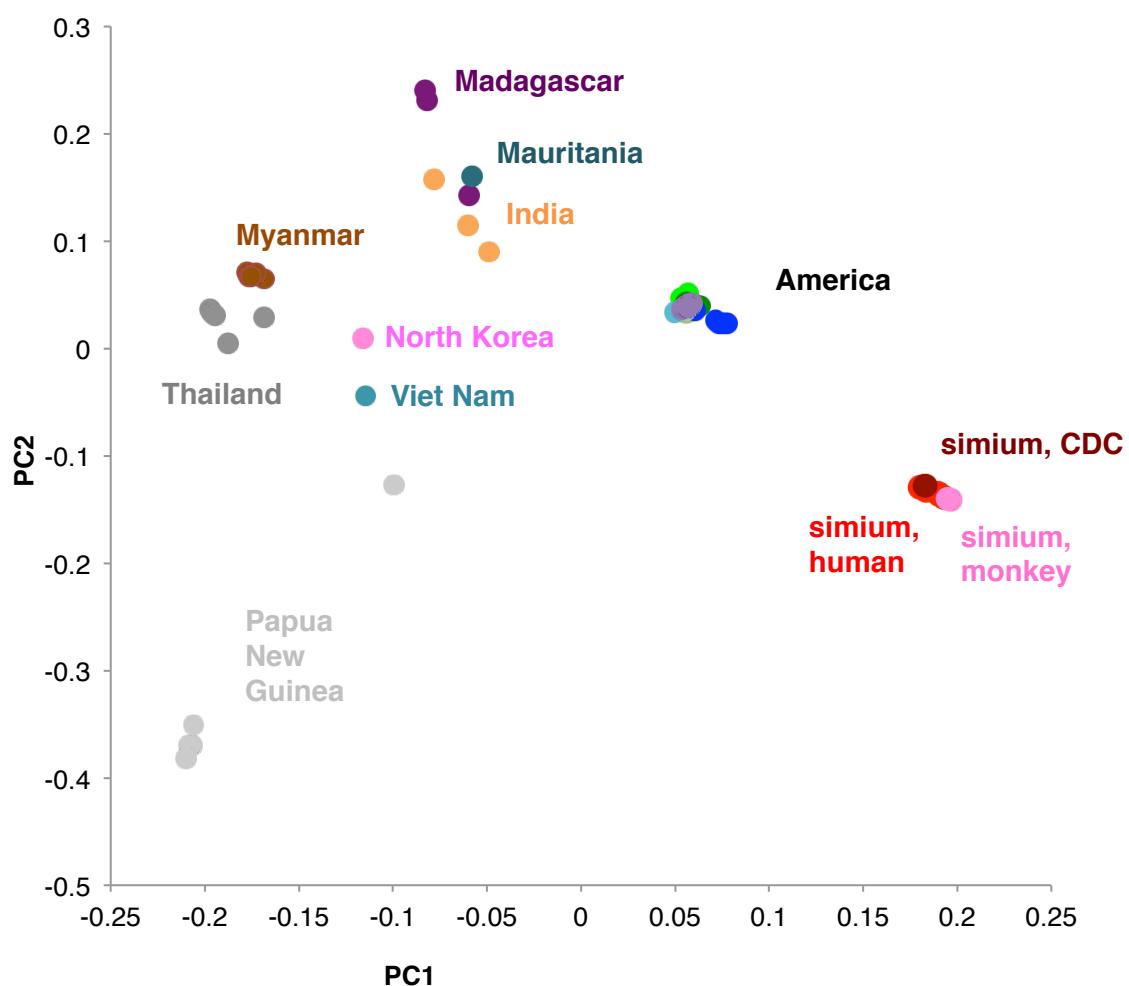
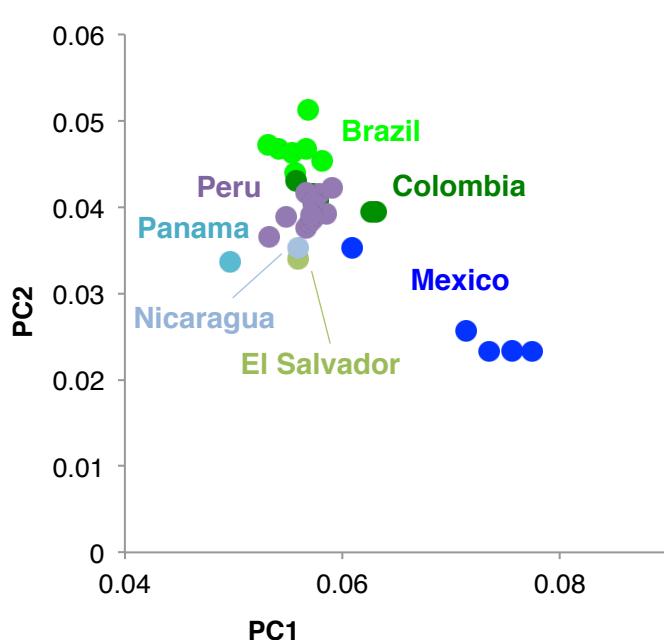


Figure S7

A



B



C

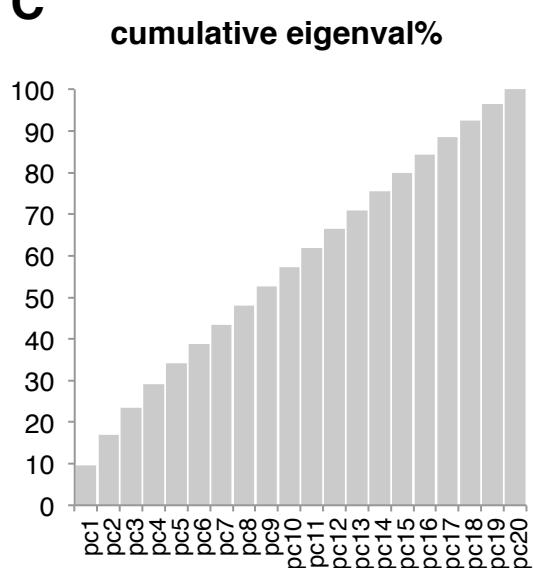


Figure S8

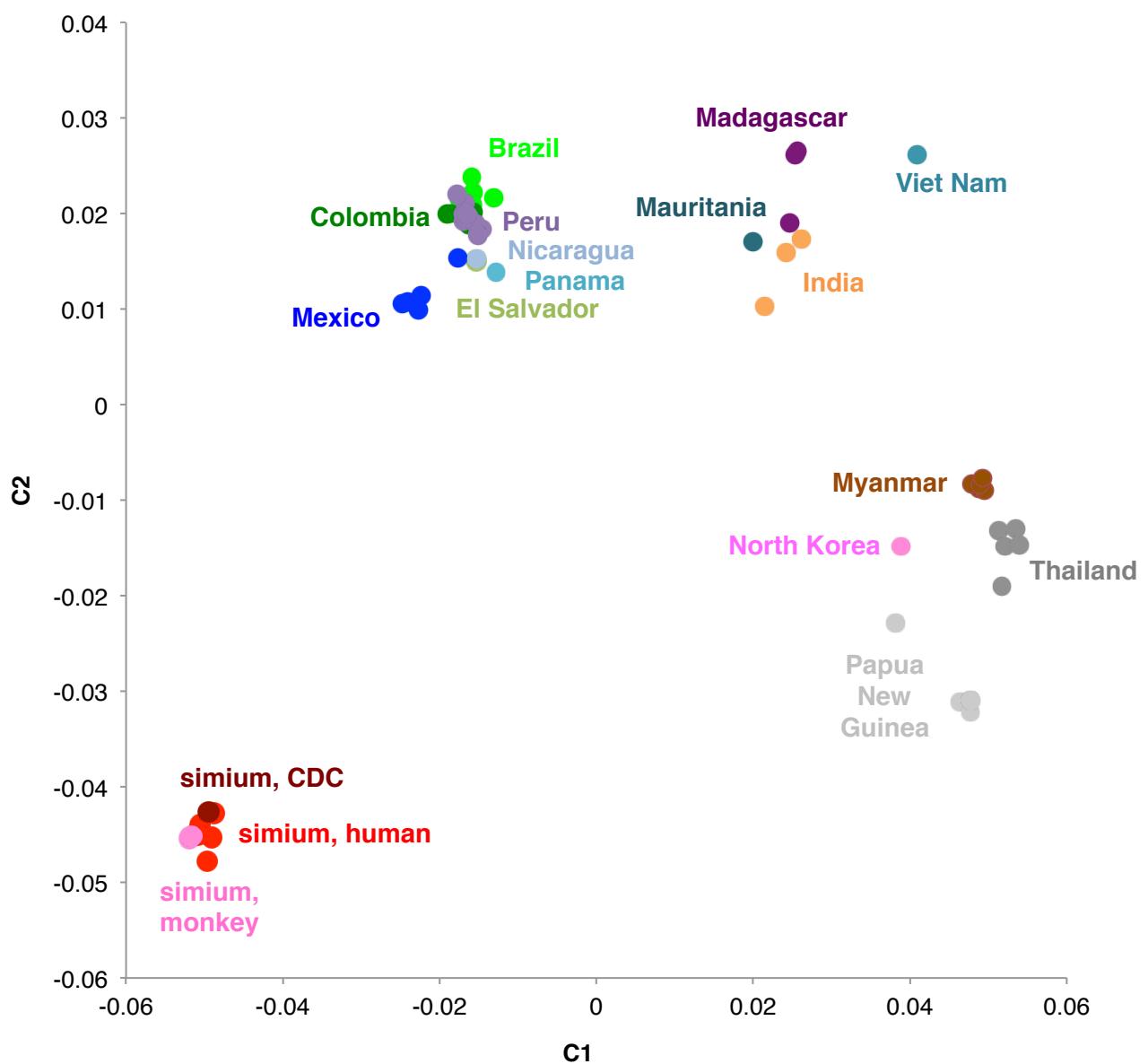


Figure S9

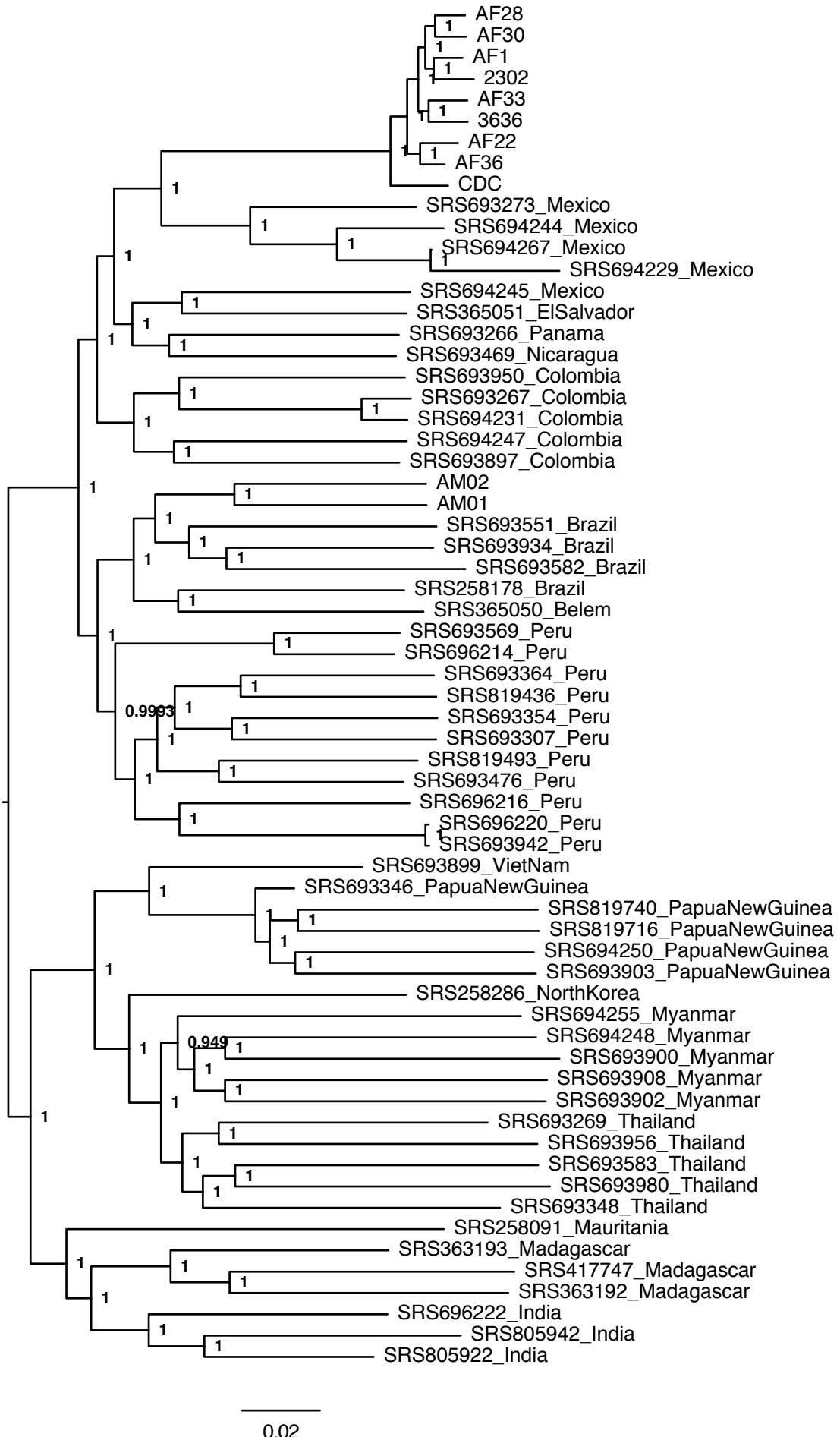


Figure S10

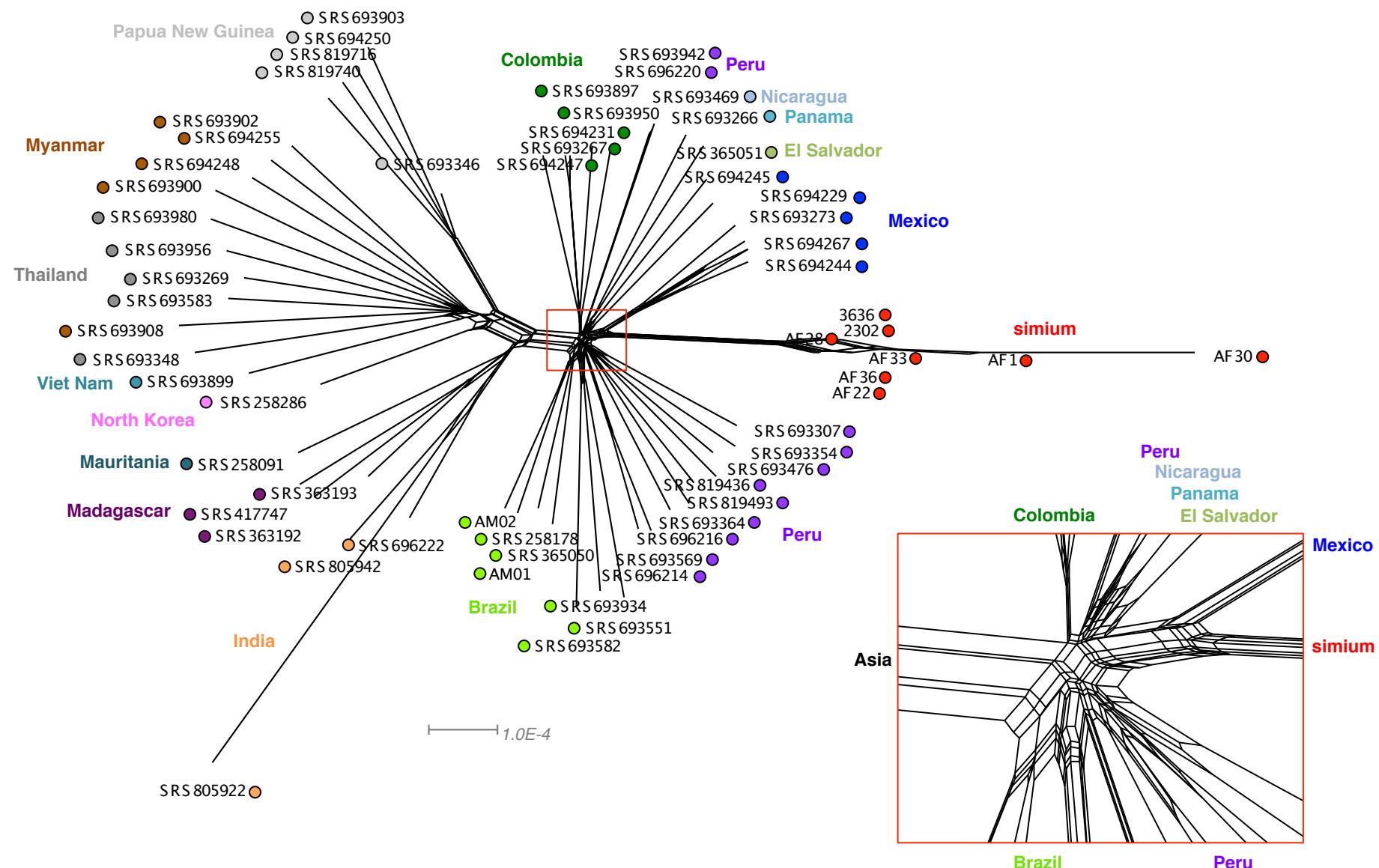


Figure S12

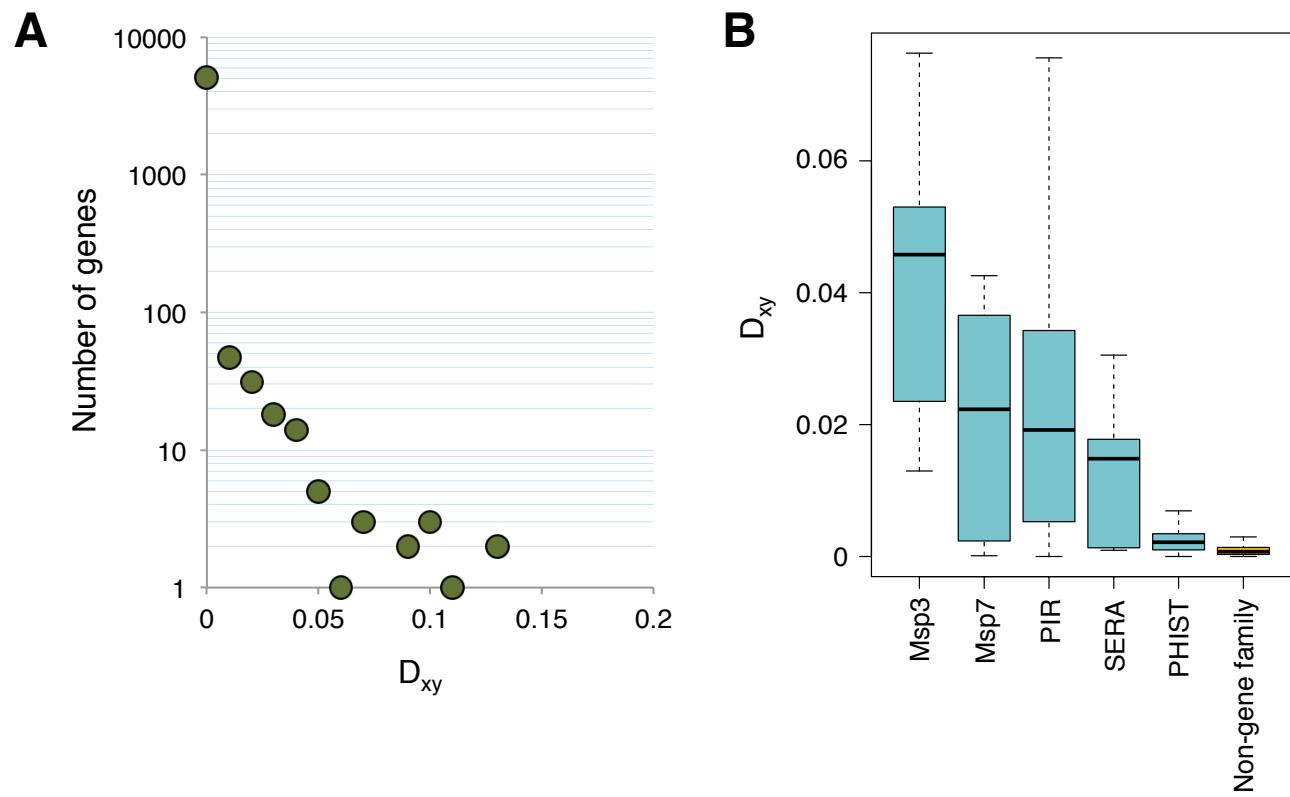


Figure S13

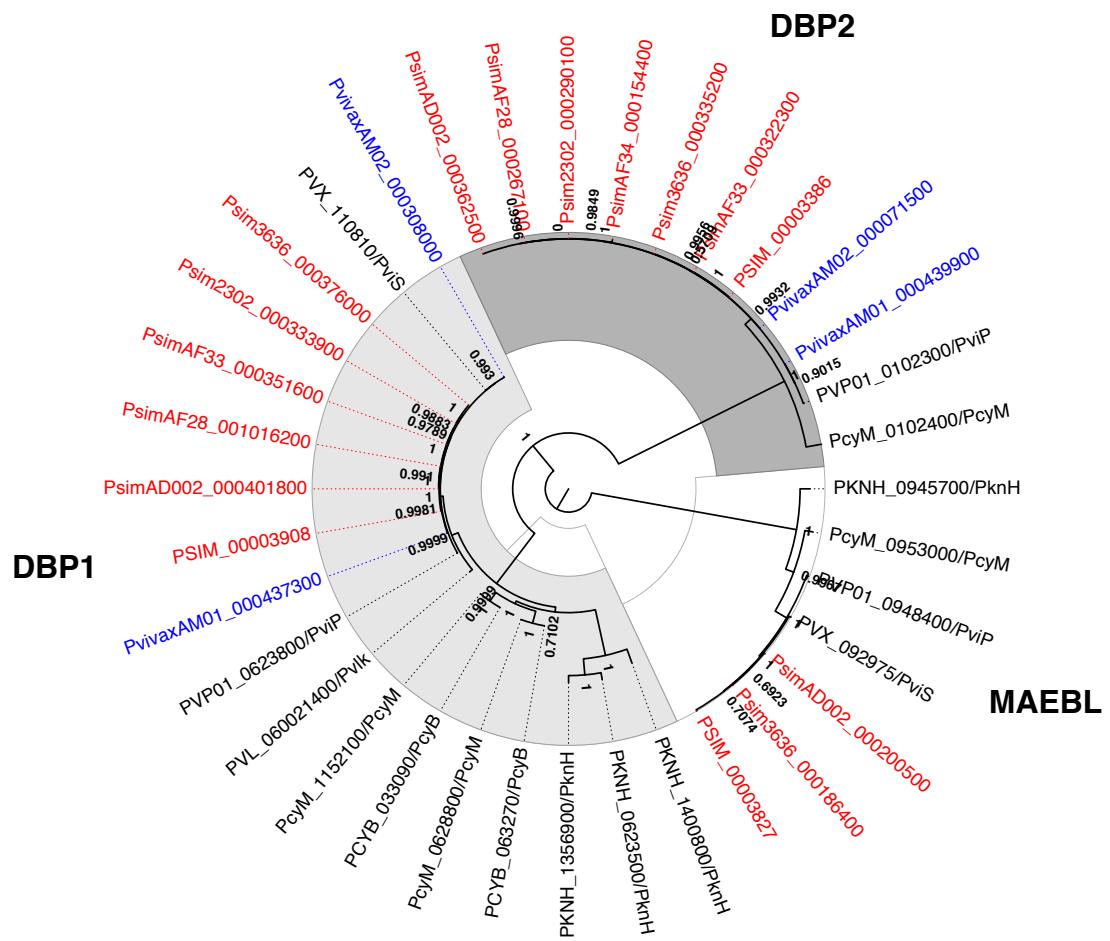


Figure S14

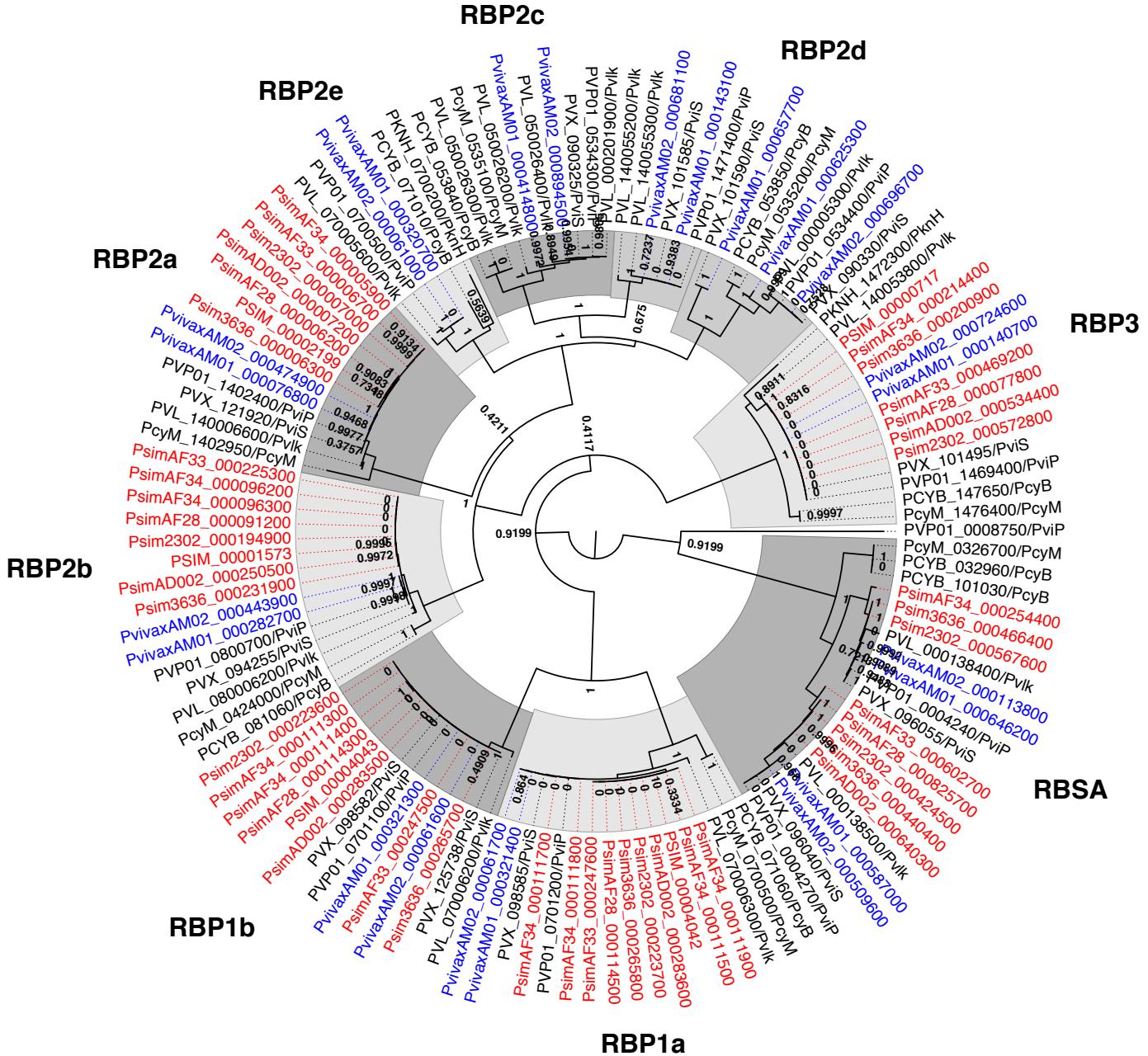


Figure S15

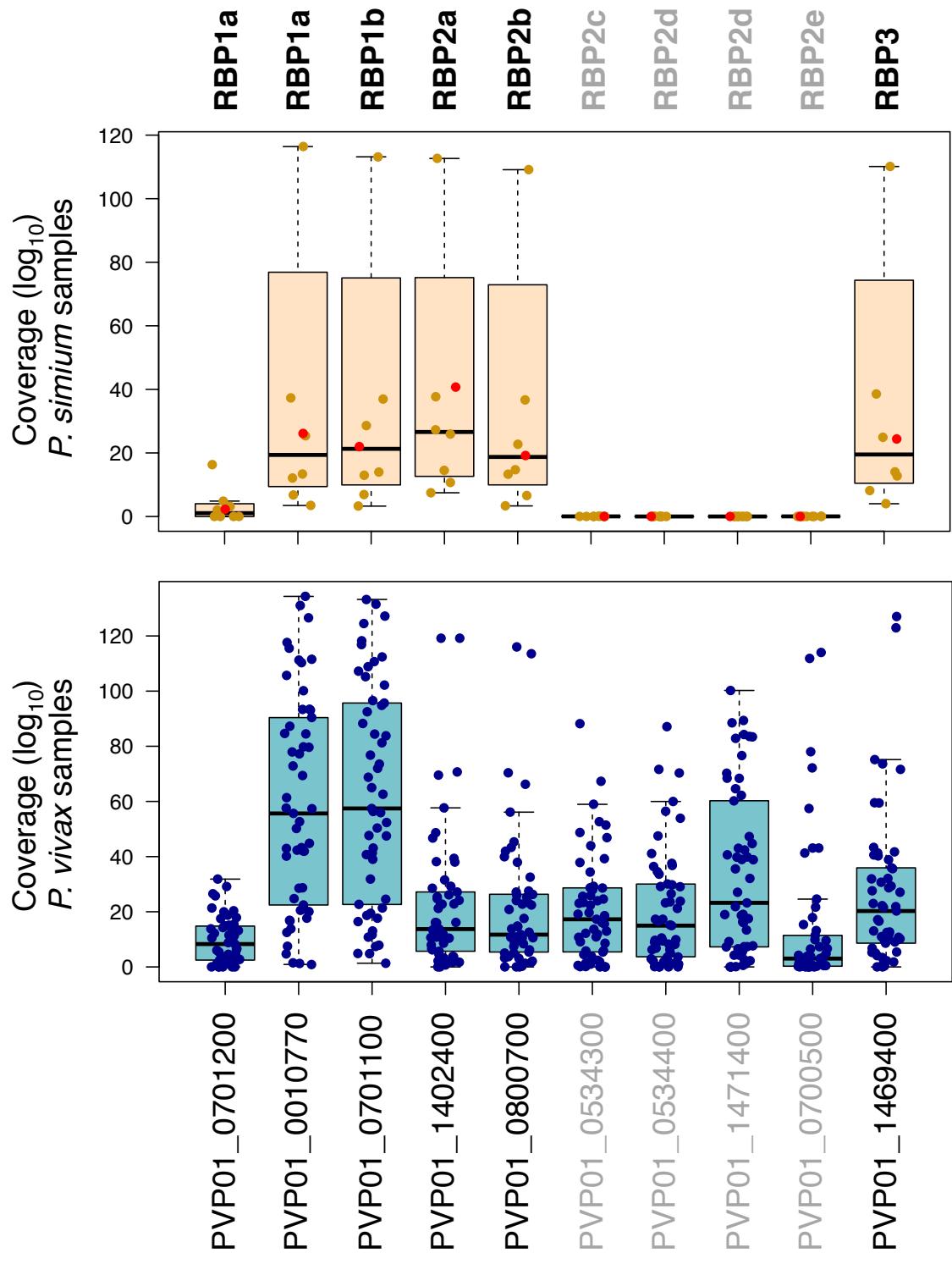


Figure S16

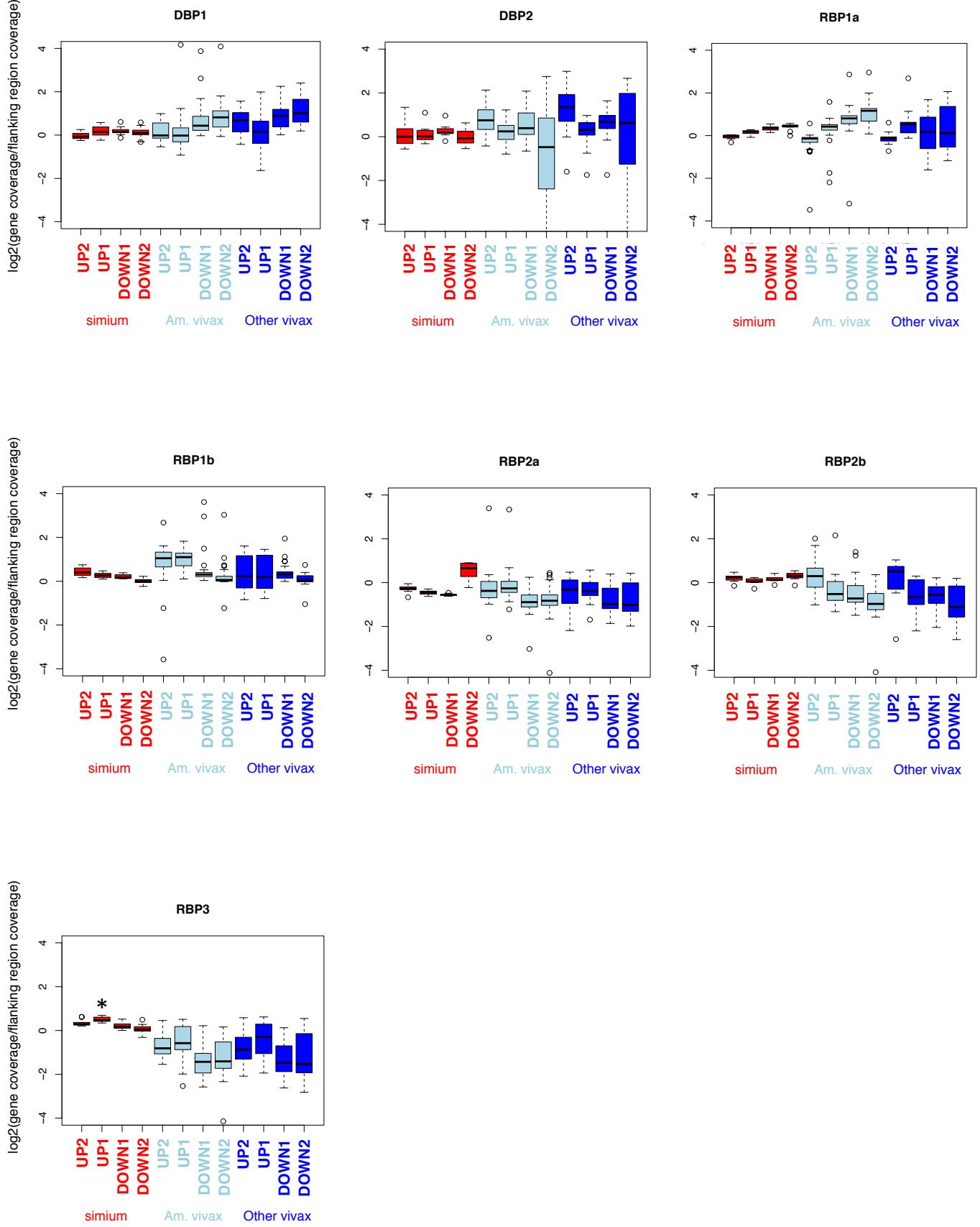


Figure S18

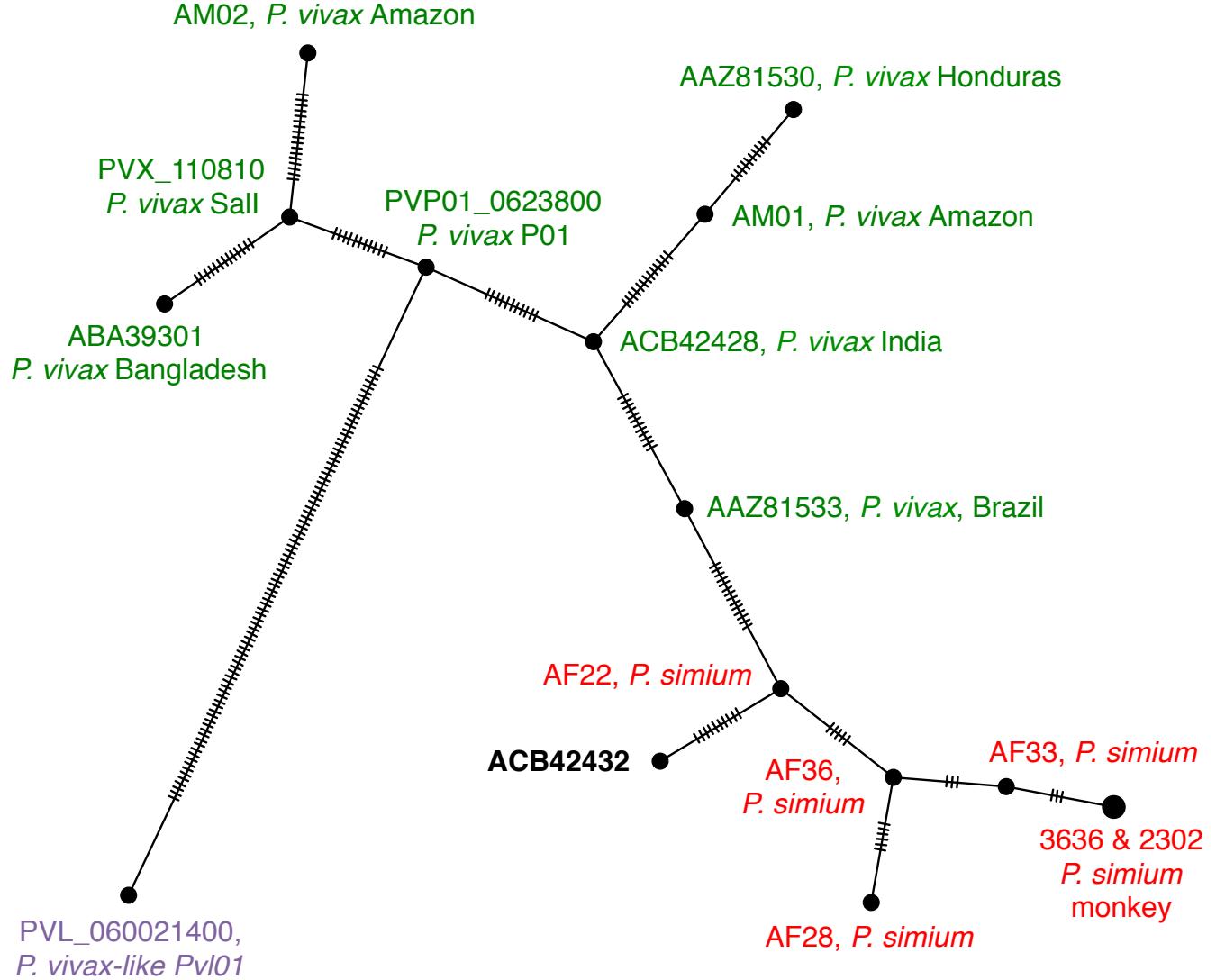


Figure S19

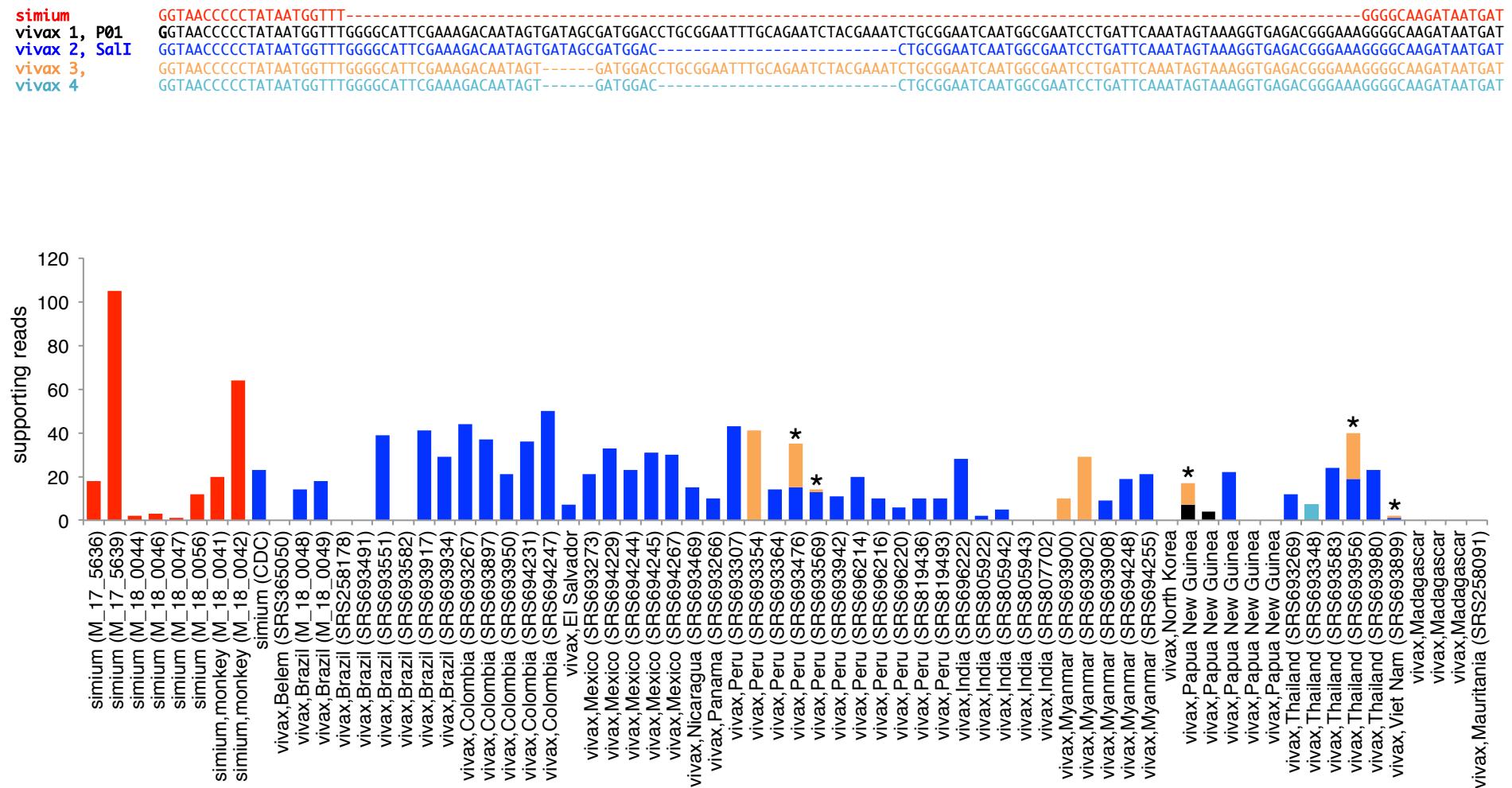
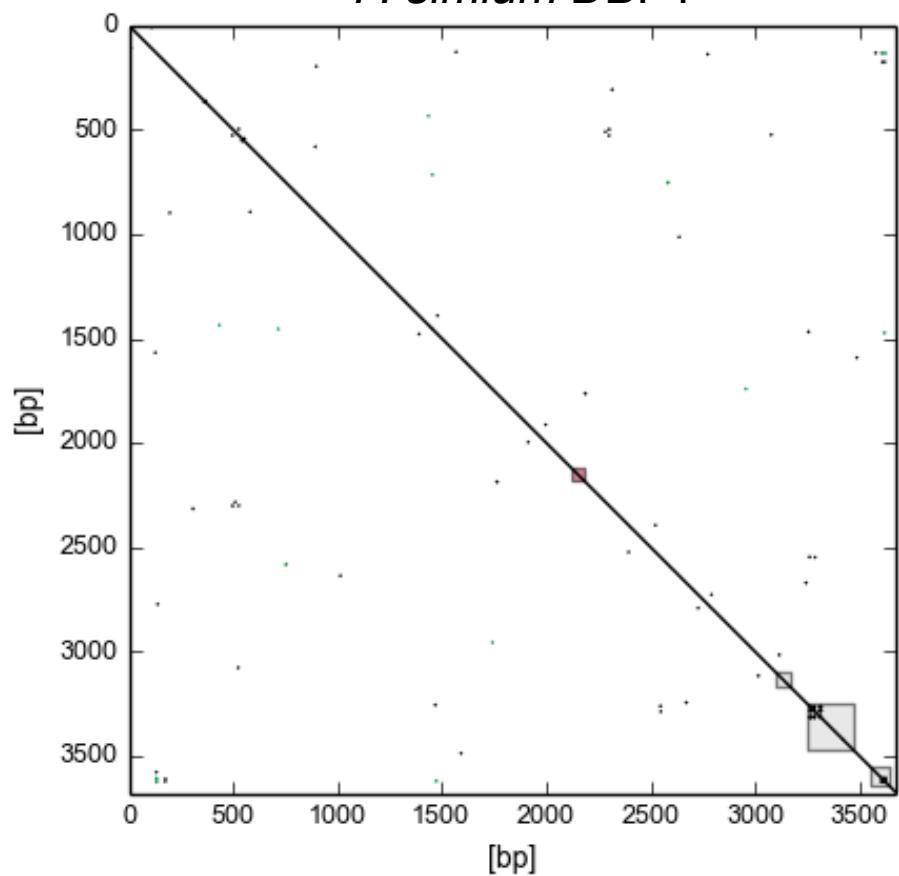


Figure S20

P. simium DBP1



P. vivax DBP1

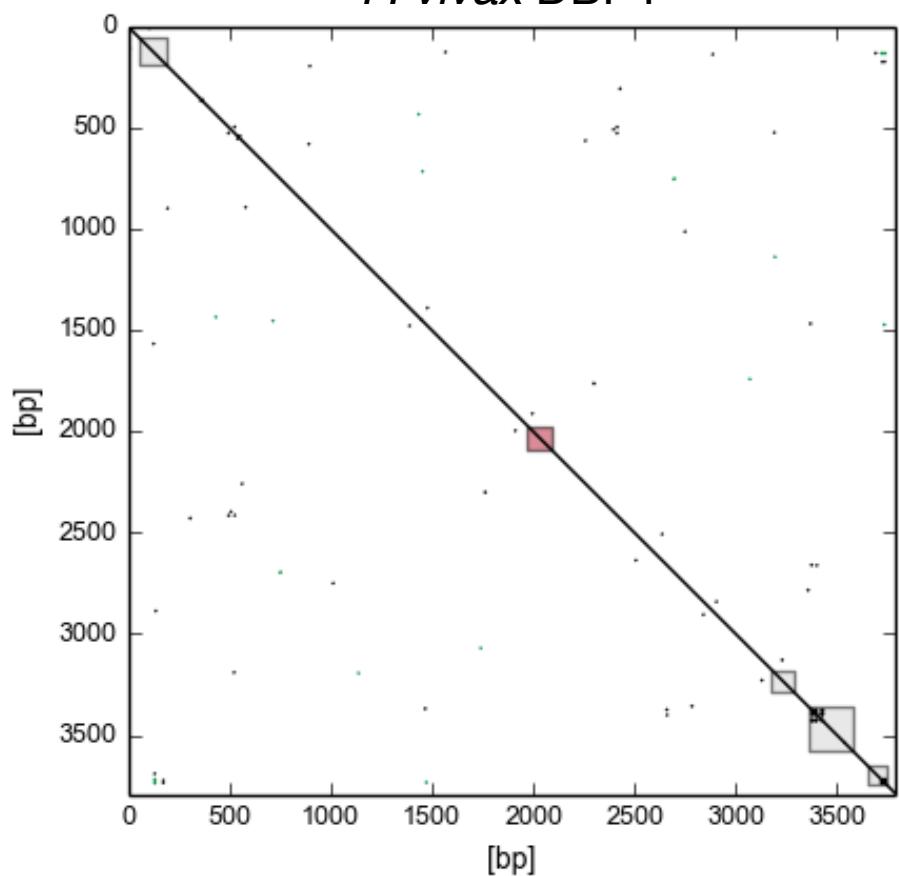


Figure S21

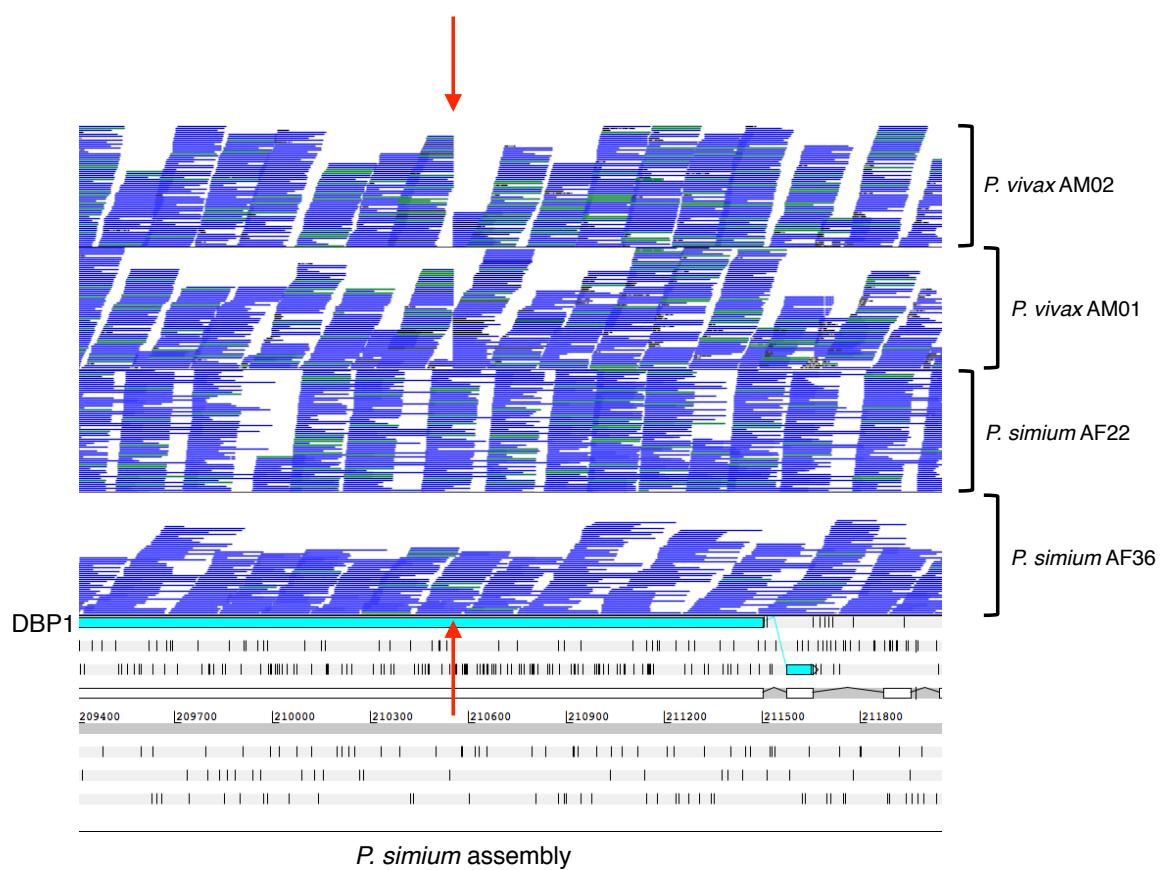


Figure S22

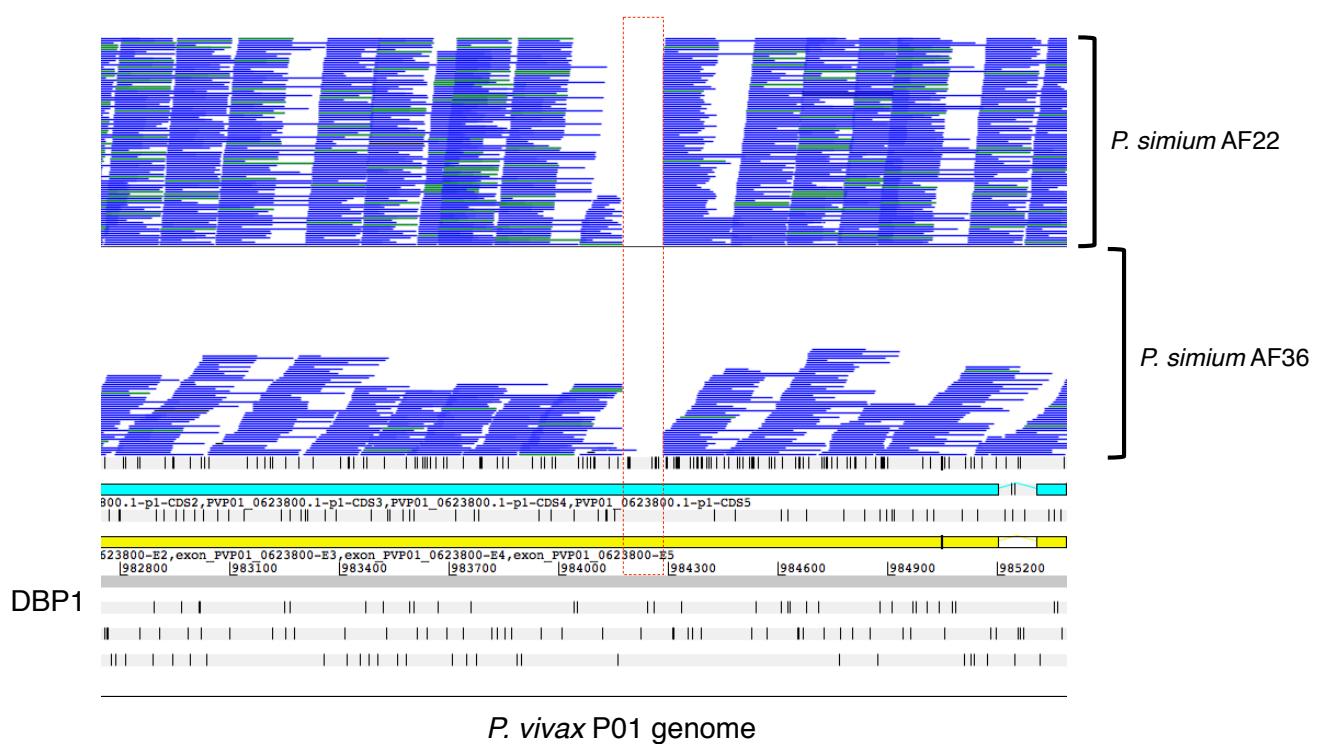


Figure S23

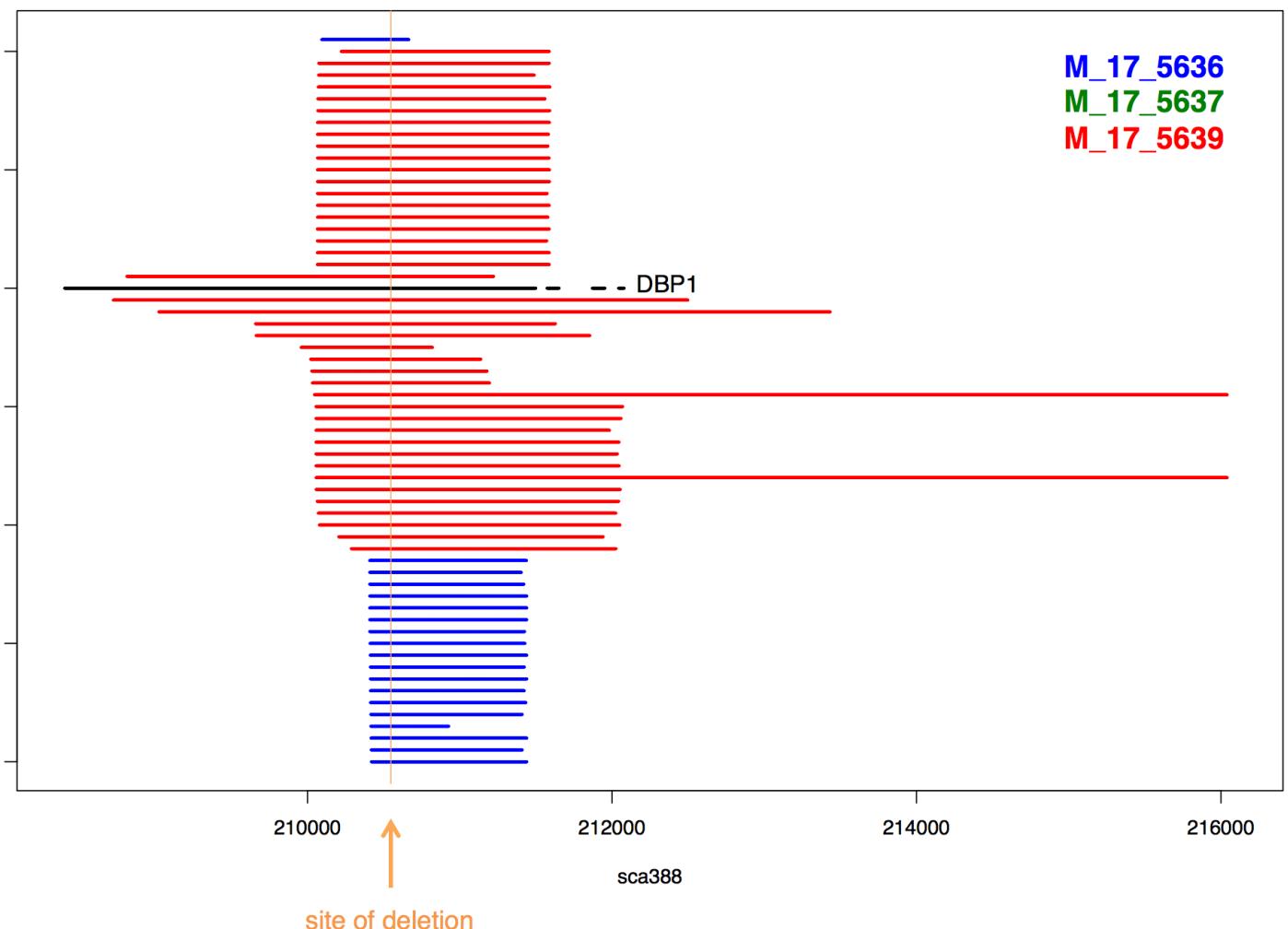
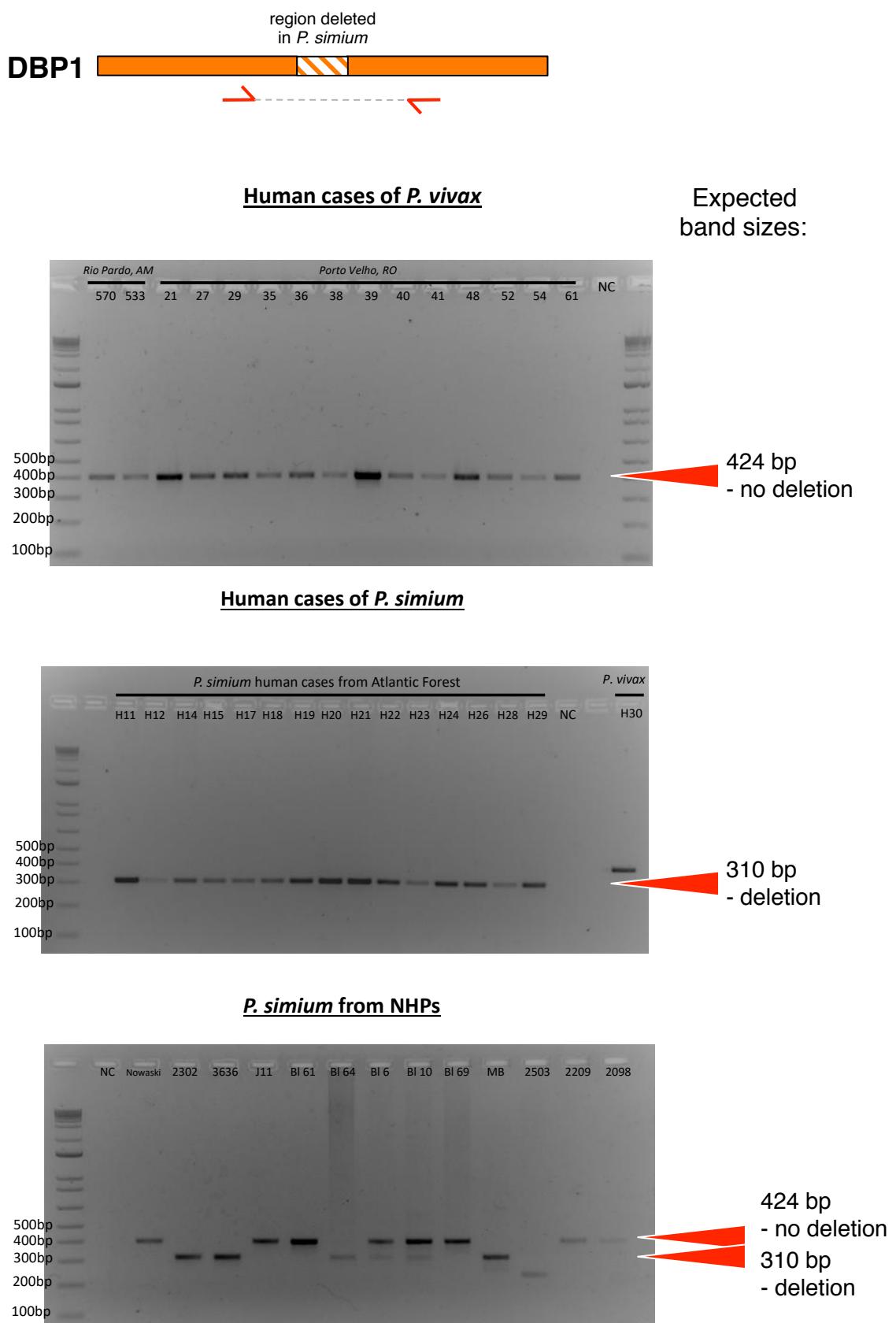


Figure S24



forward primer: ATCGCGACAAAGCTACTGCC
reverse primer: TCCCACACAATCCCTGCTG

Figure S26

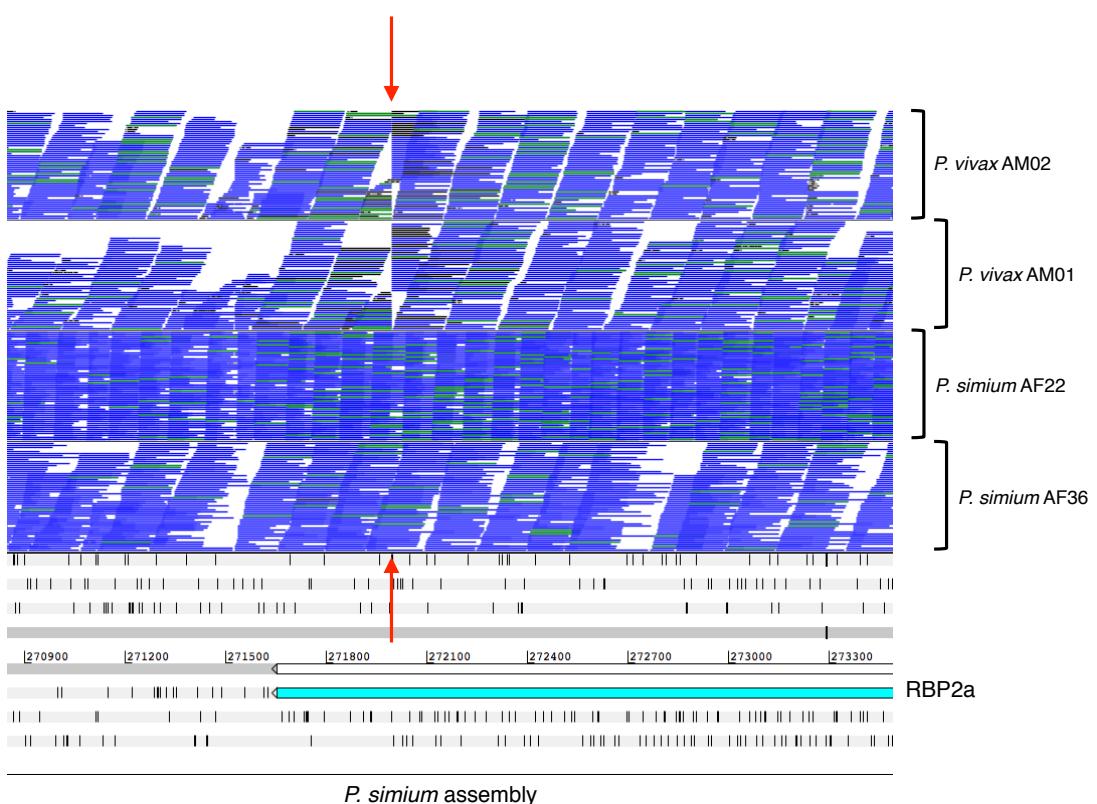


Figure S27

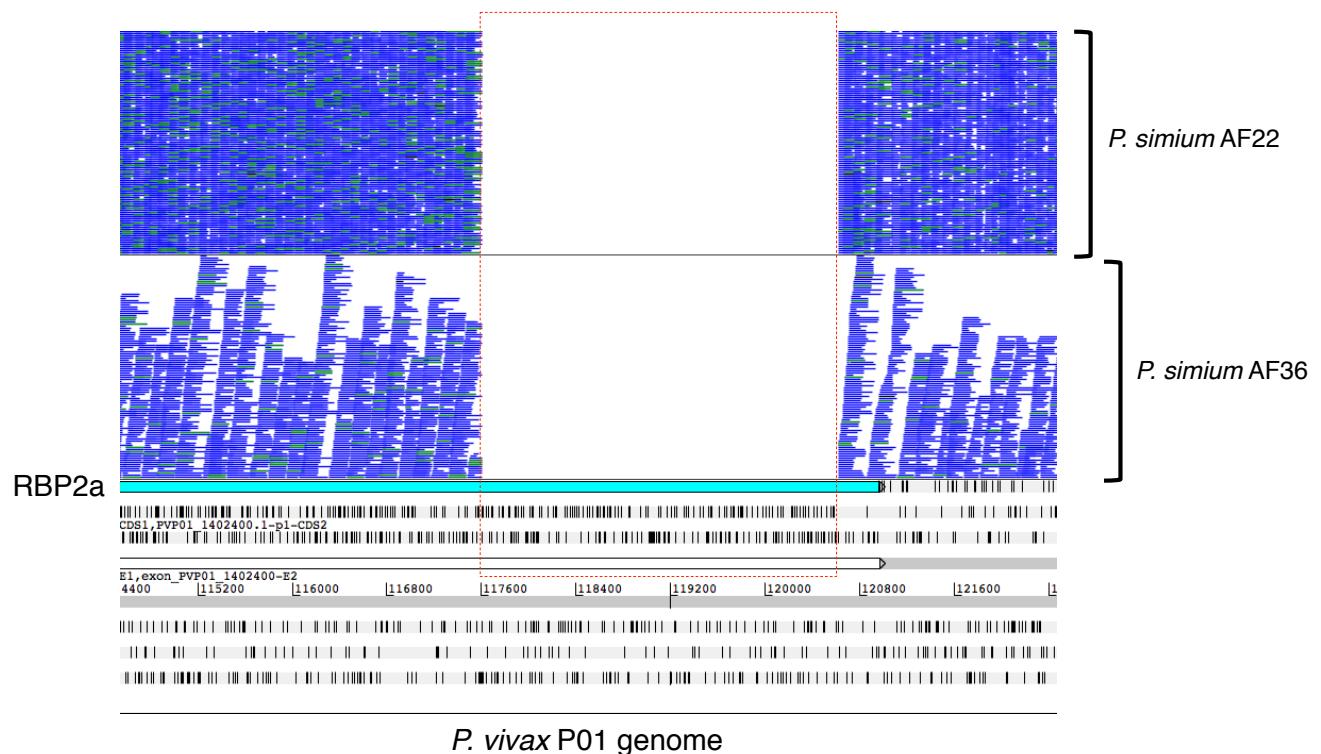


Figure S28

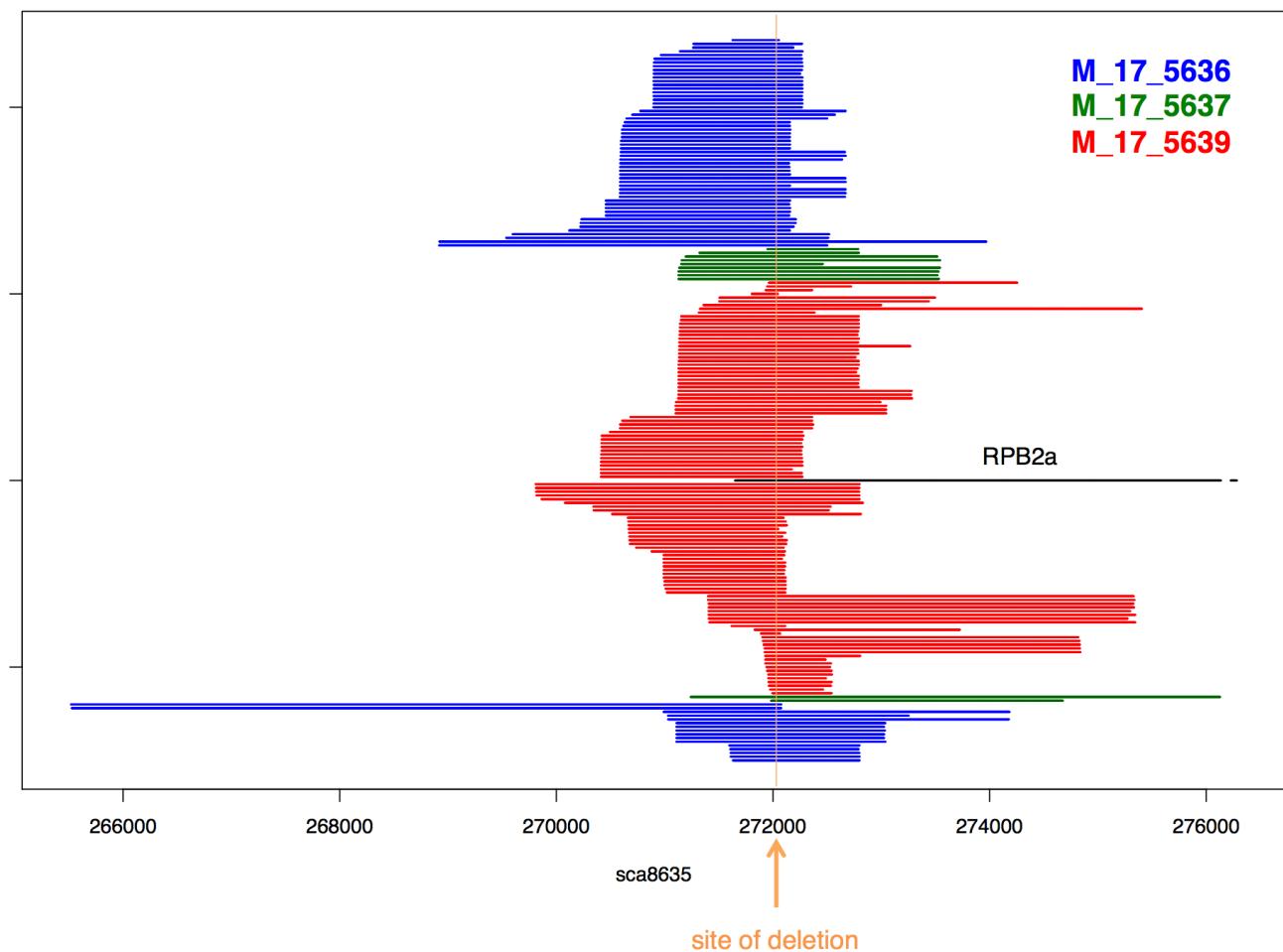


Figure S29

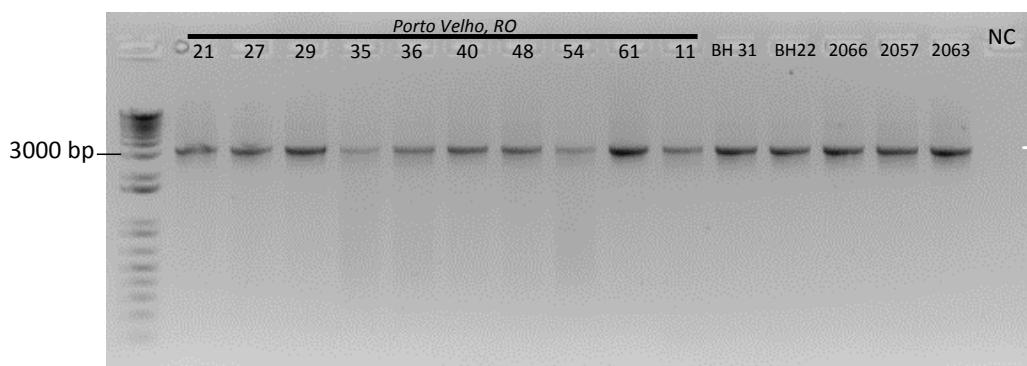
RBP2a

region deleted in *P. simium*

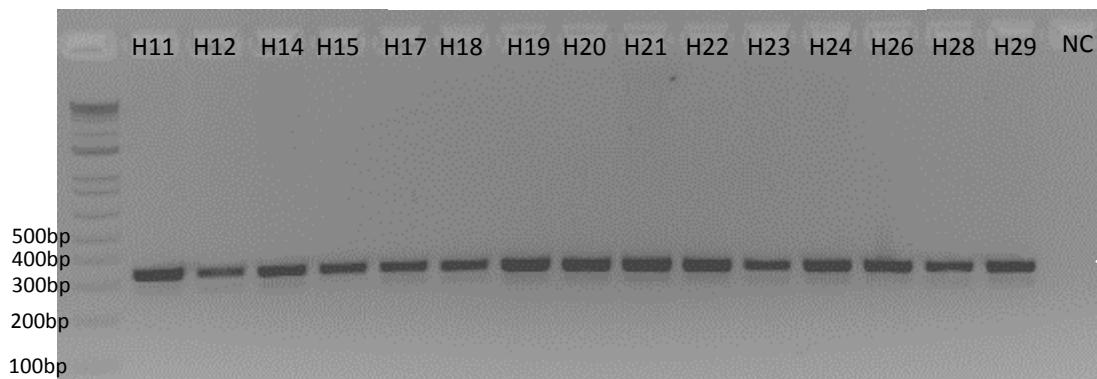


Human *P. vivax*

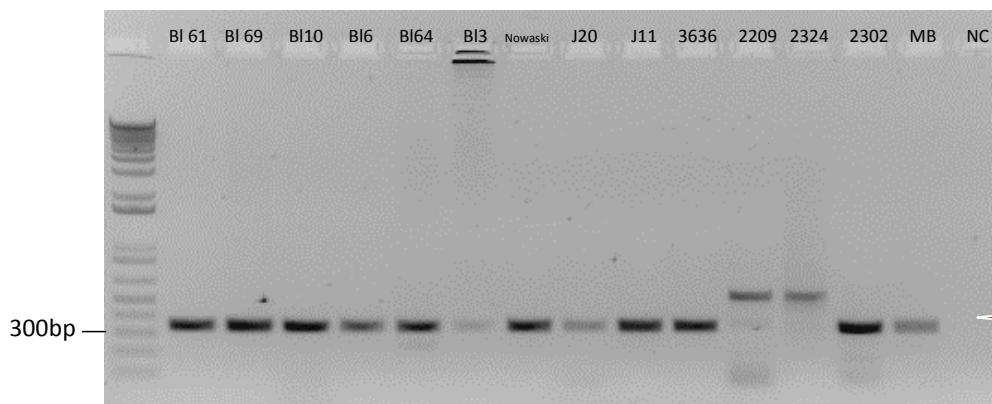
Expected
band sizes:



Human *P. simium*



NHP *P. simium*

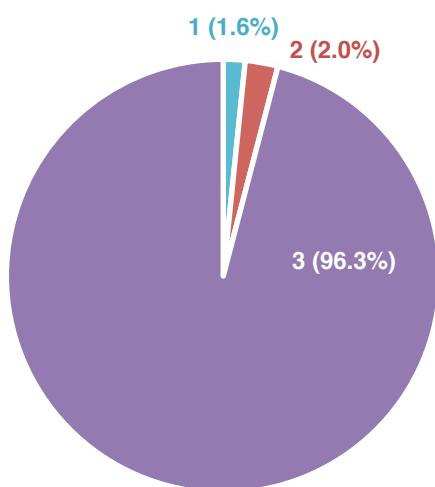


forward primer: AGTGCAAAAGCAGAGGAGG
reverse primer: CCTGCTGAAAACGGACATT

Figure S30

A

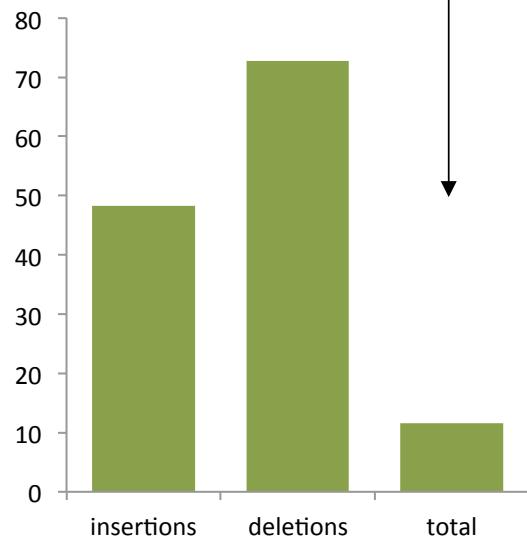
Percentages of indels in genes with sizes being integers of 1, 2 and 3.



B

Percentage of all proteins being low-complexity regions

Percentage of indel sites overlapping low-complexity regions



C

Protein size distributions for genes with and without indels

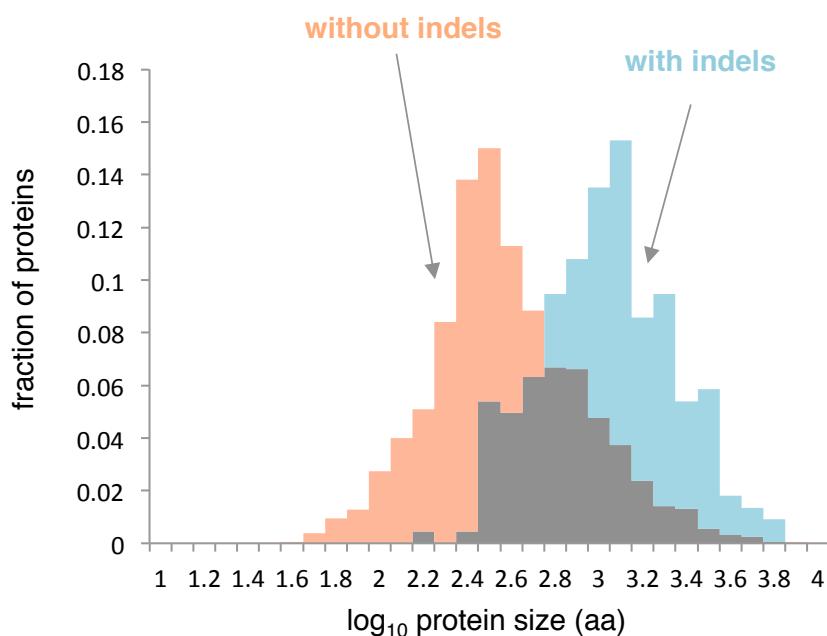


Figure S31

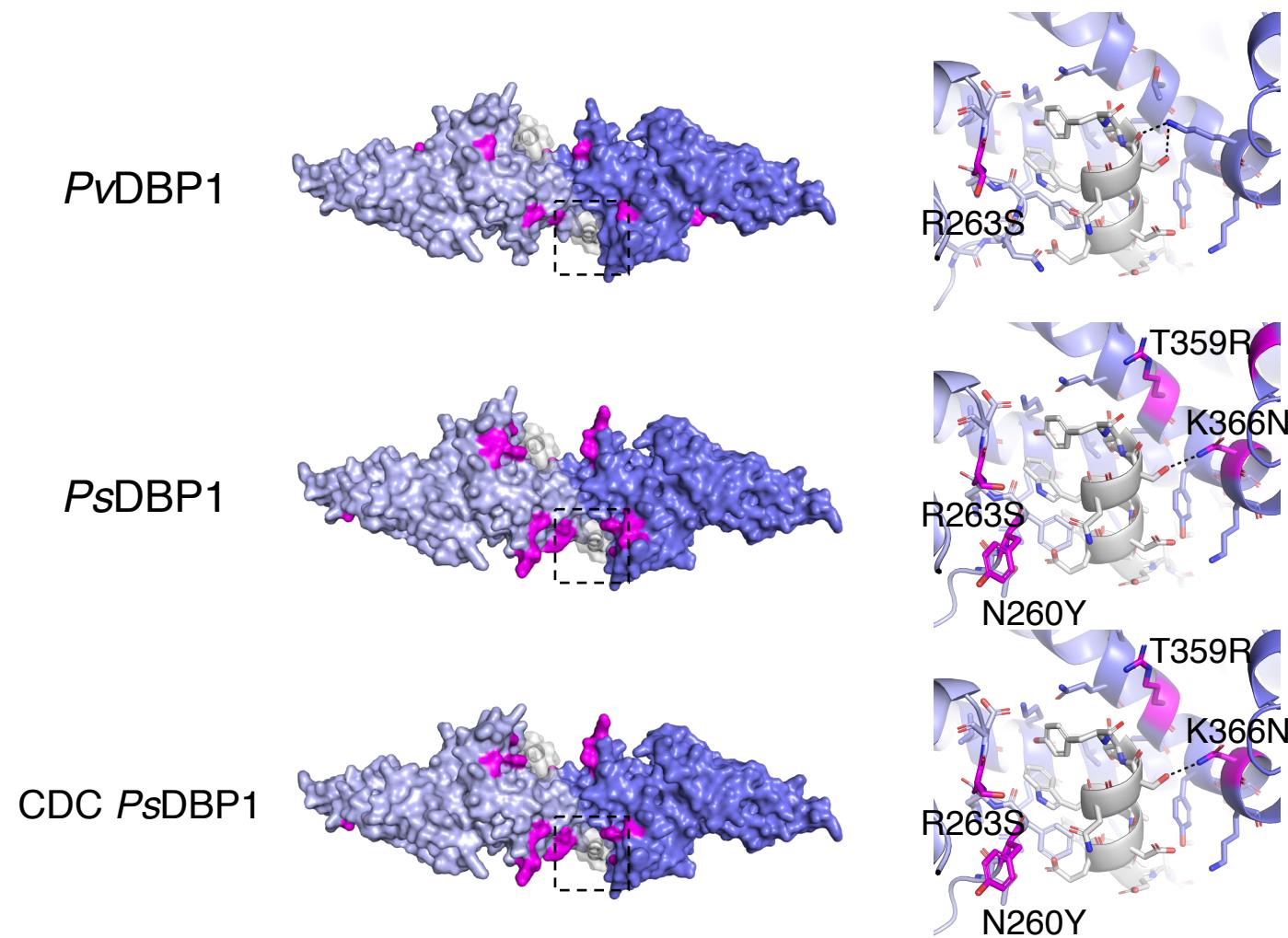


Figure S32

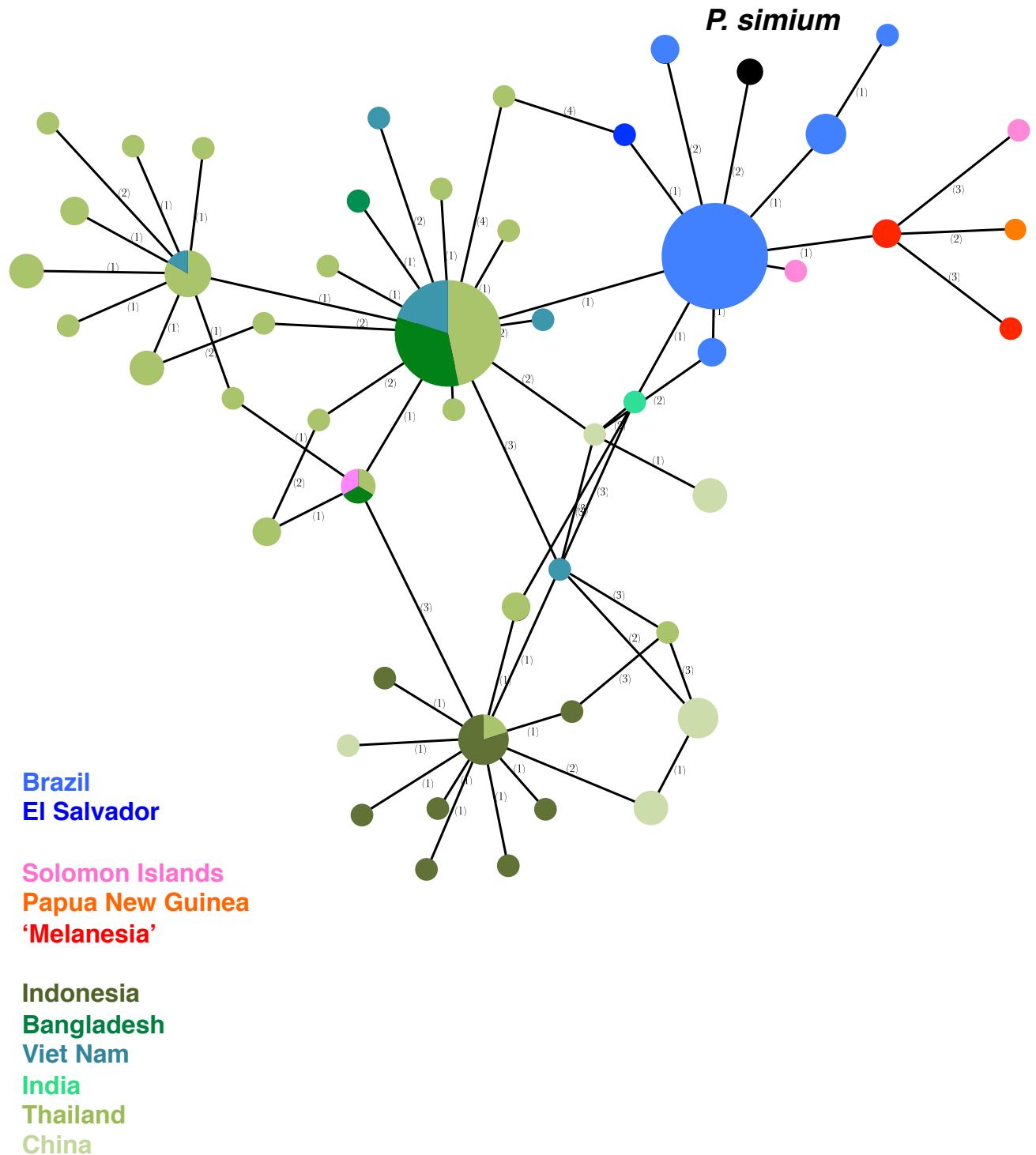


Figure S33

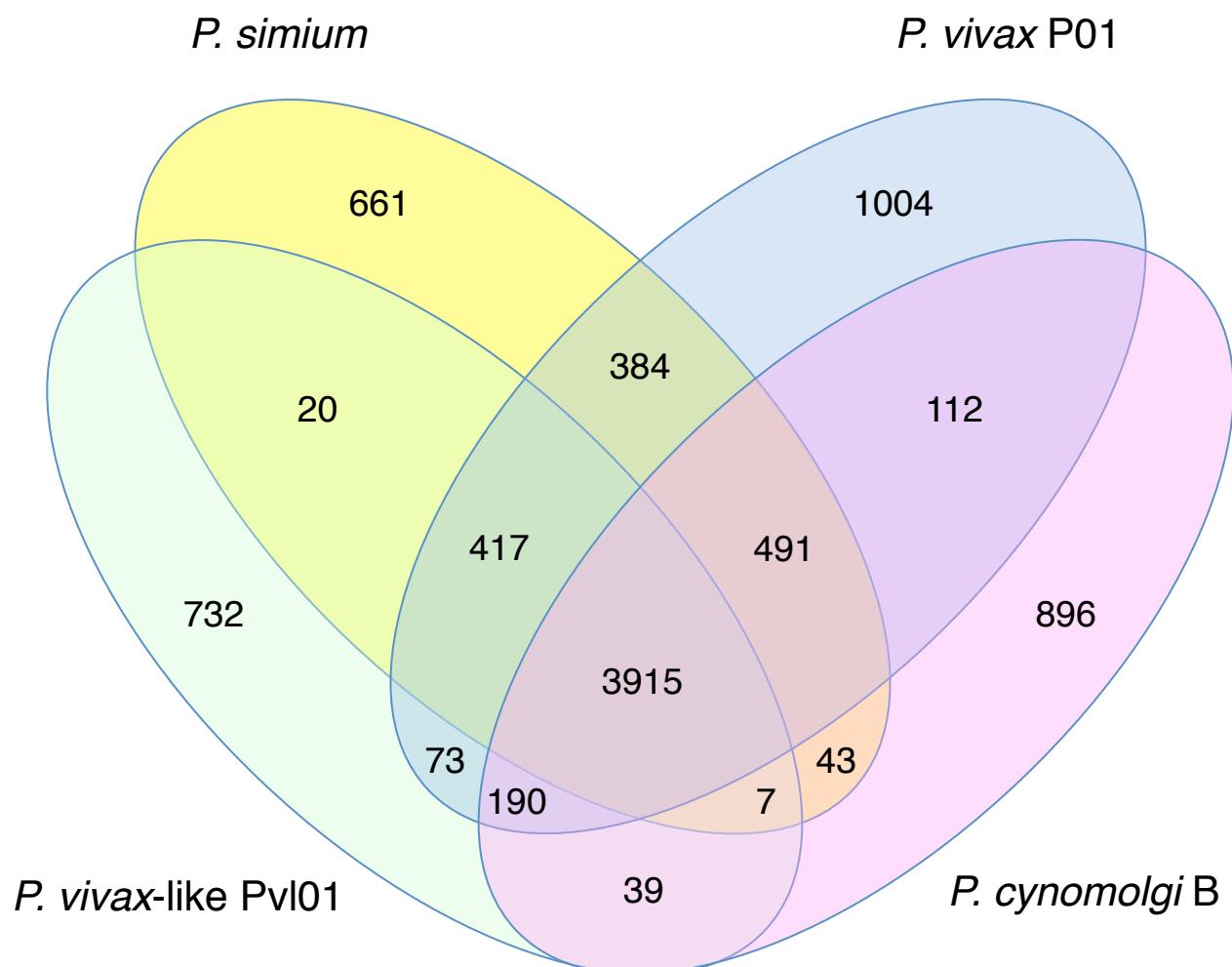


Figure S34

