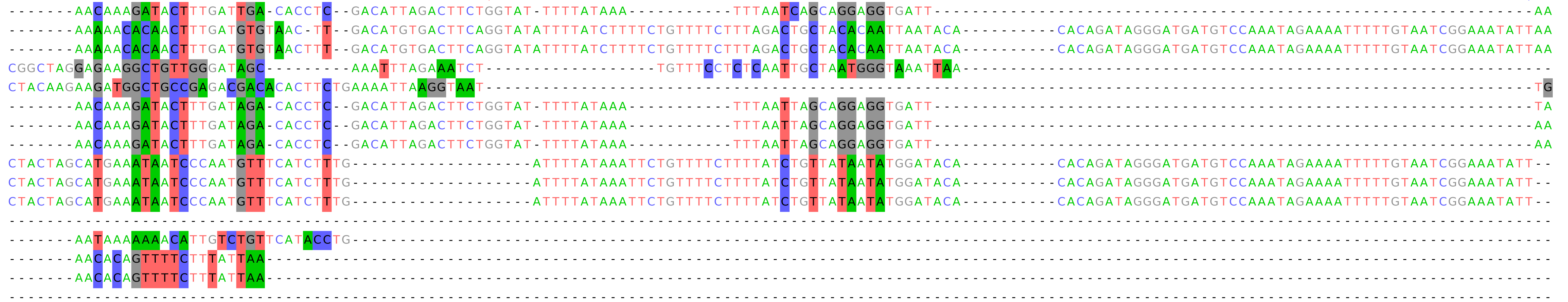


Start crop Point

End crop Point

MSA length = 5805



Start crop Point

End crop Point

0

10

0

1000

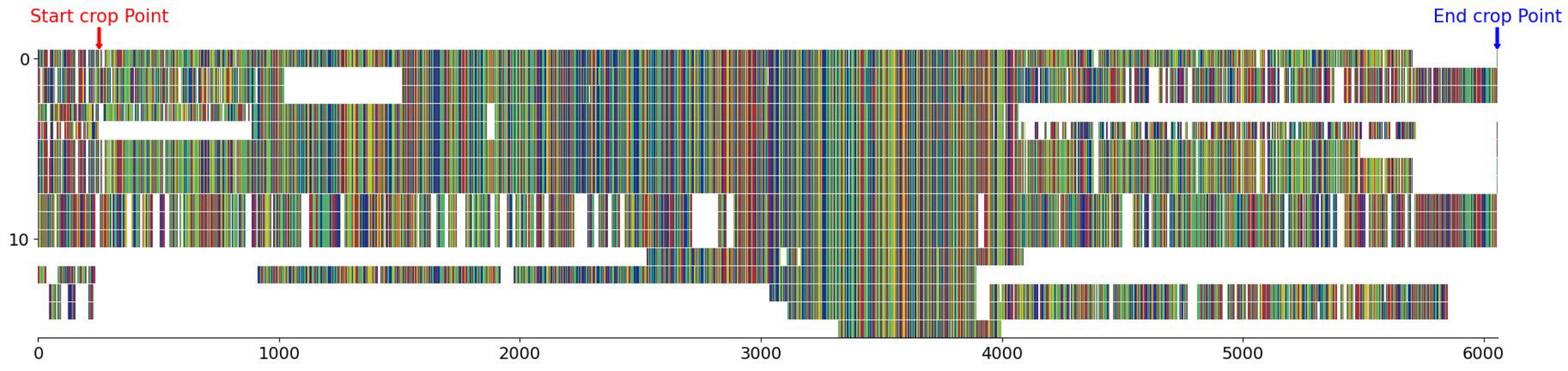
2000

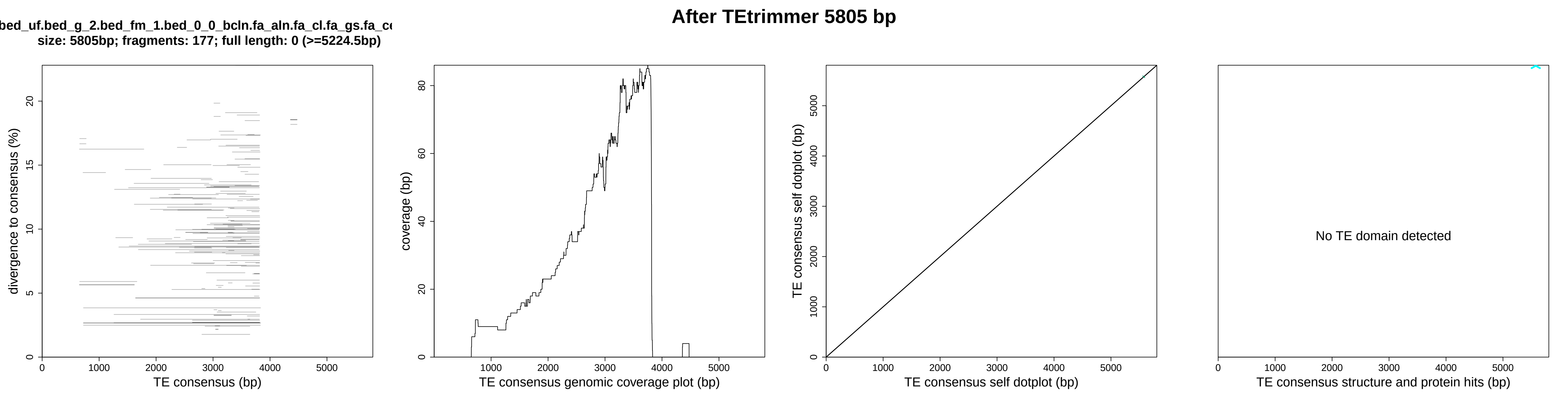
3000

4000

5000

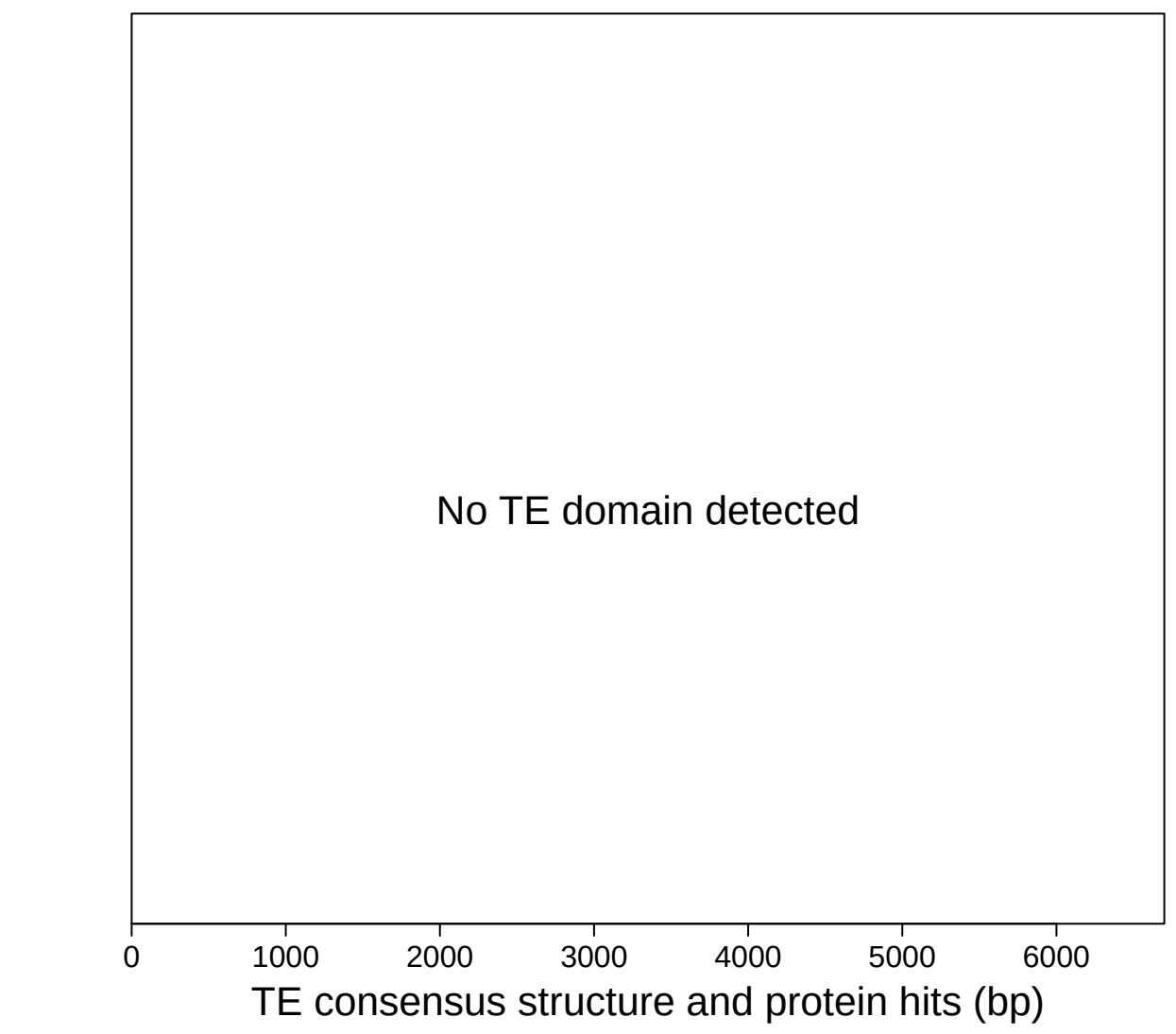
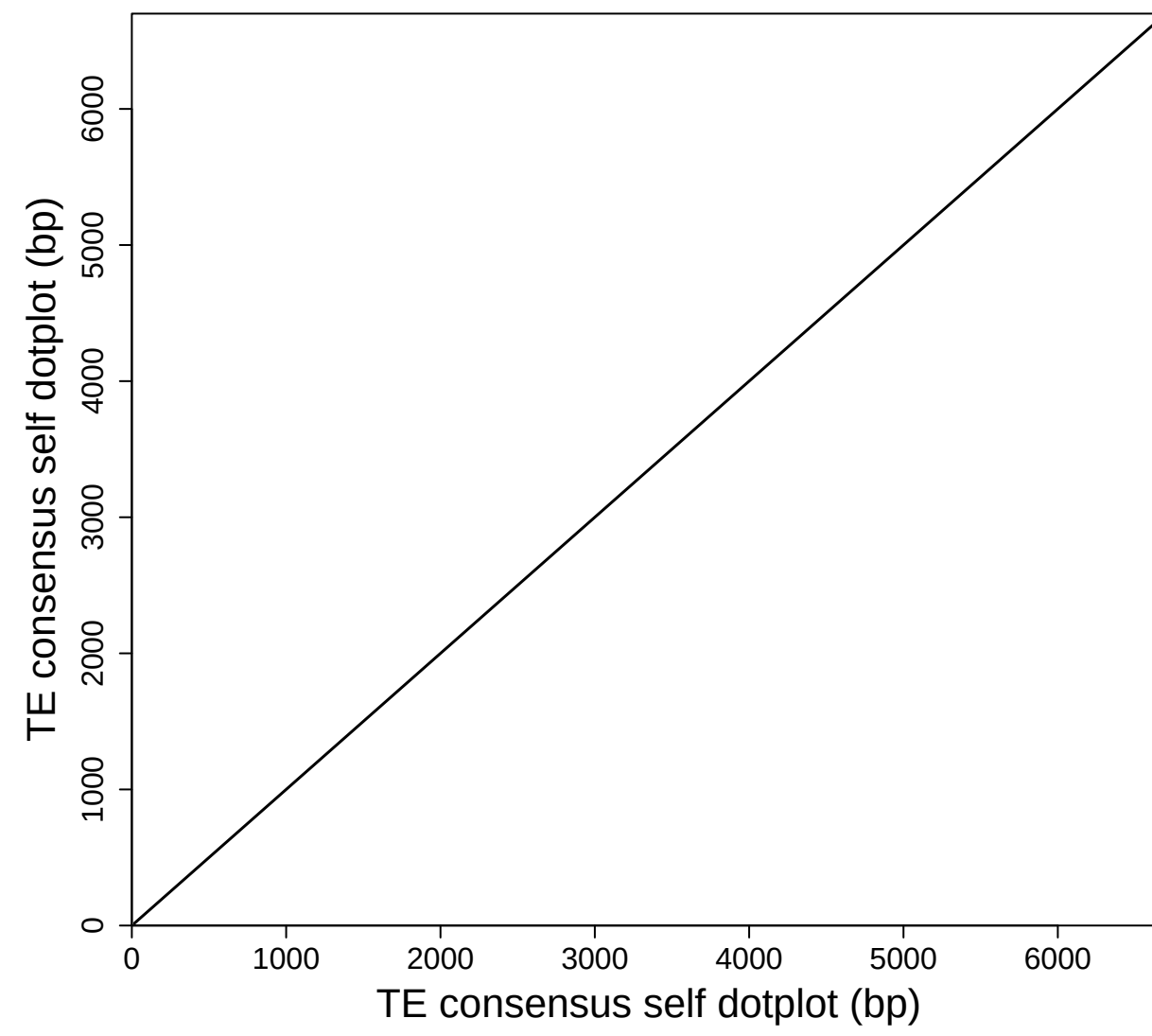
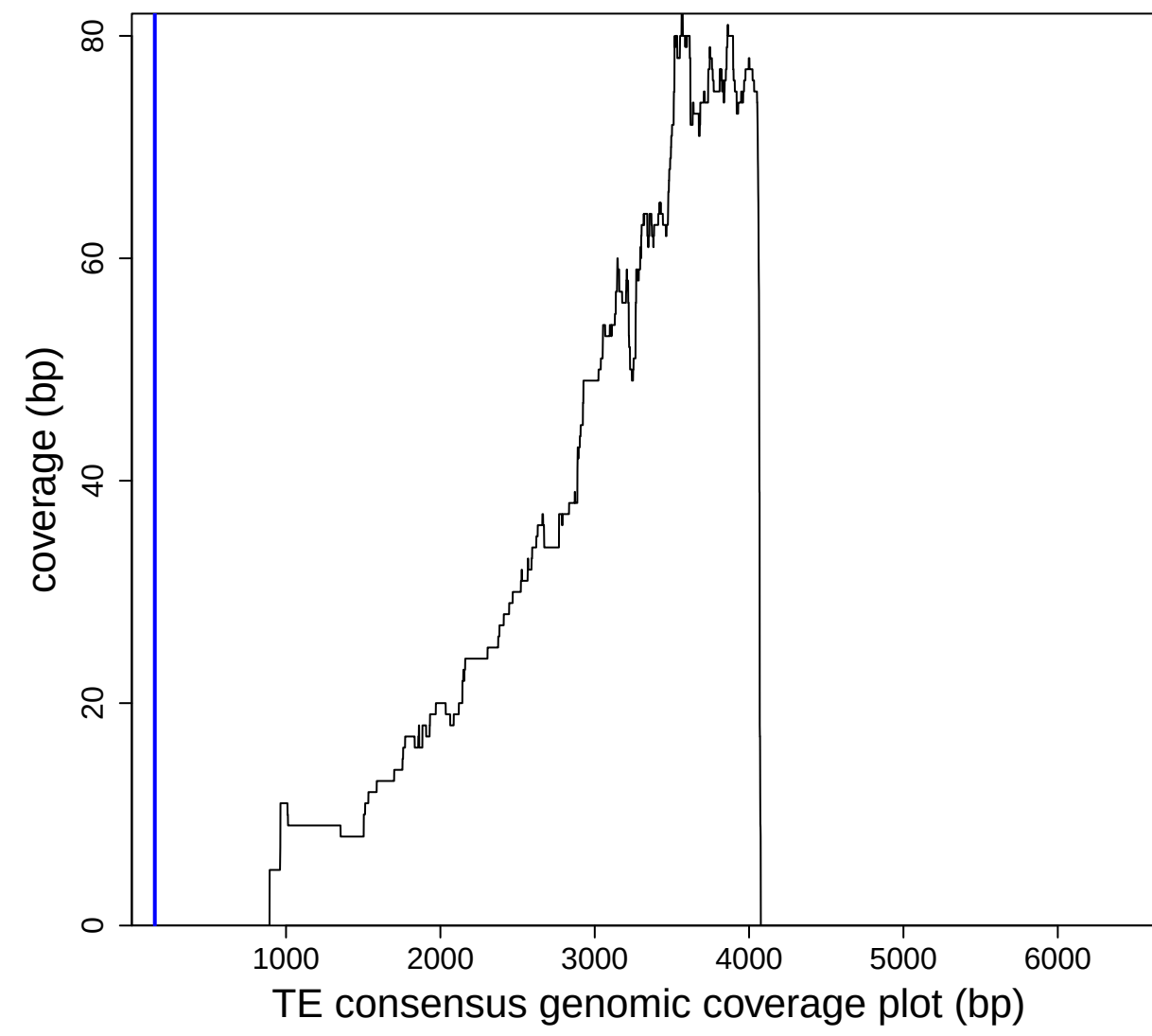
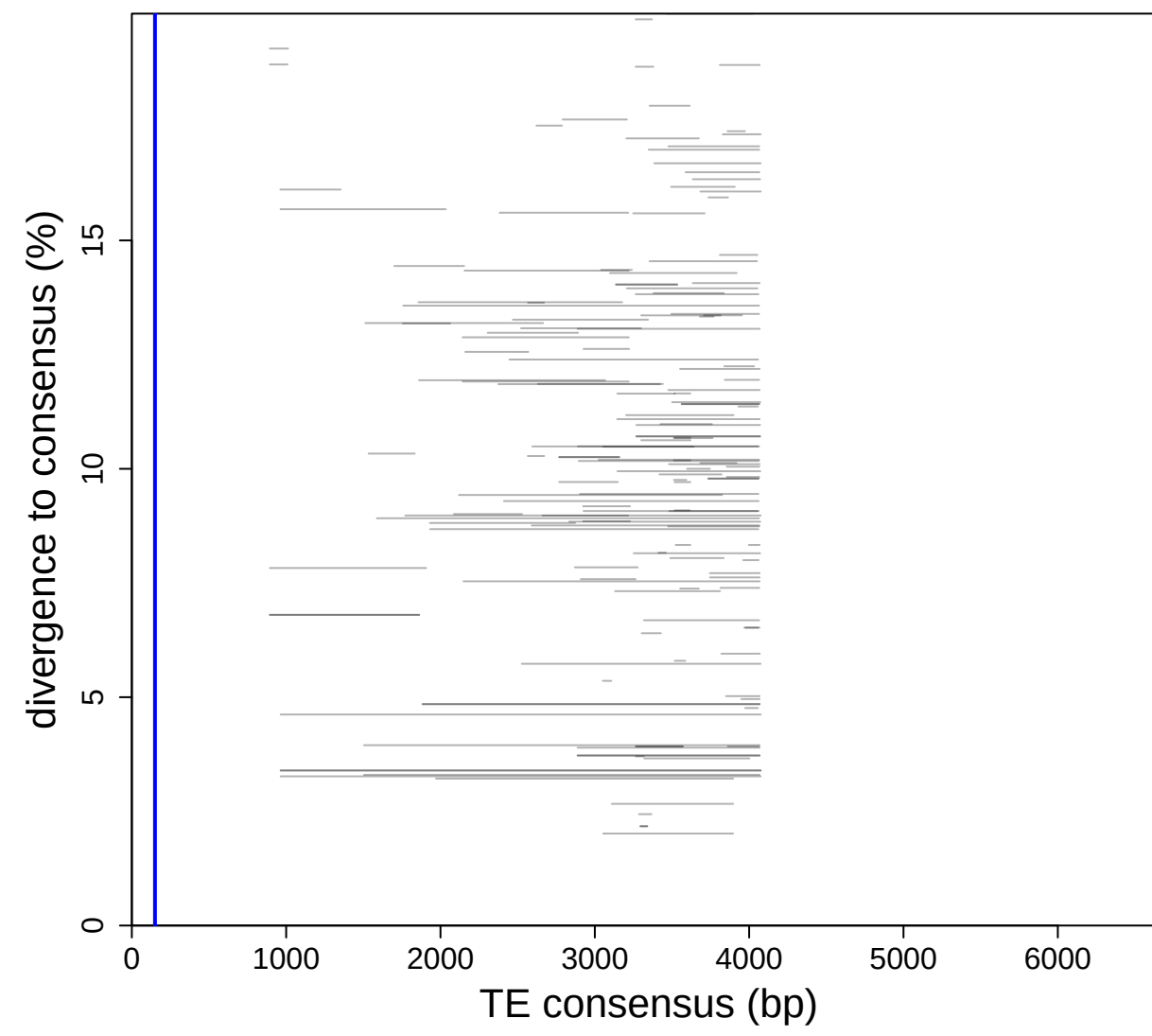
6000



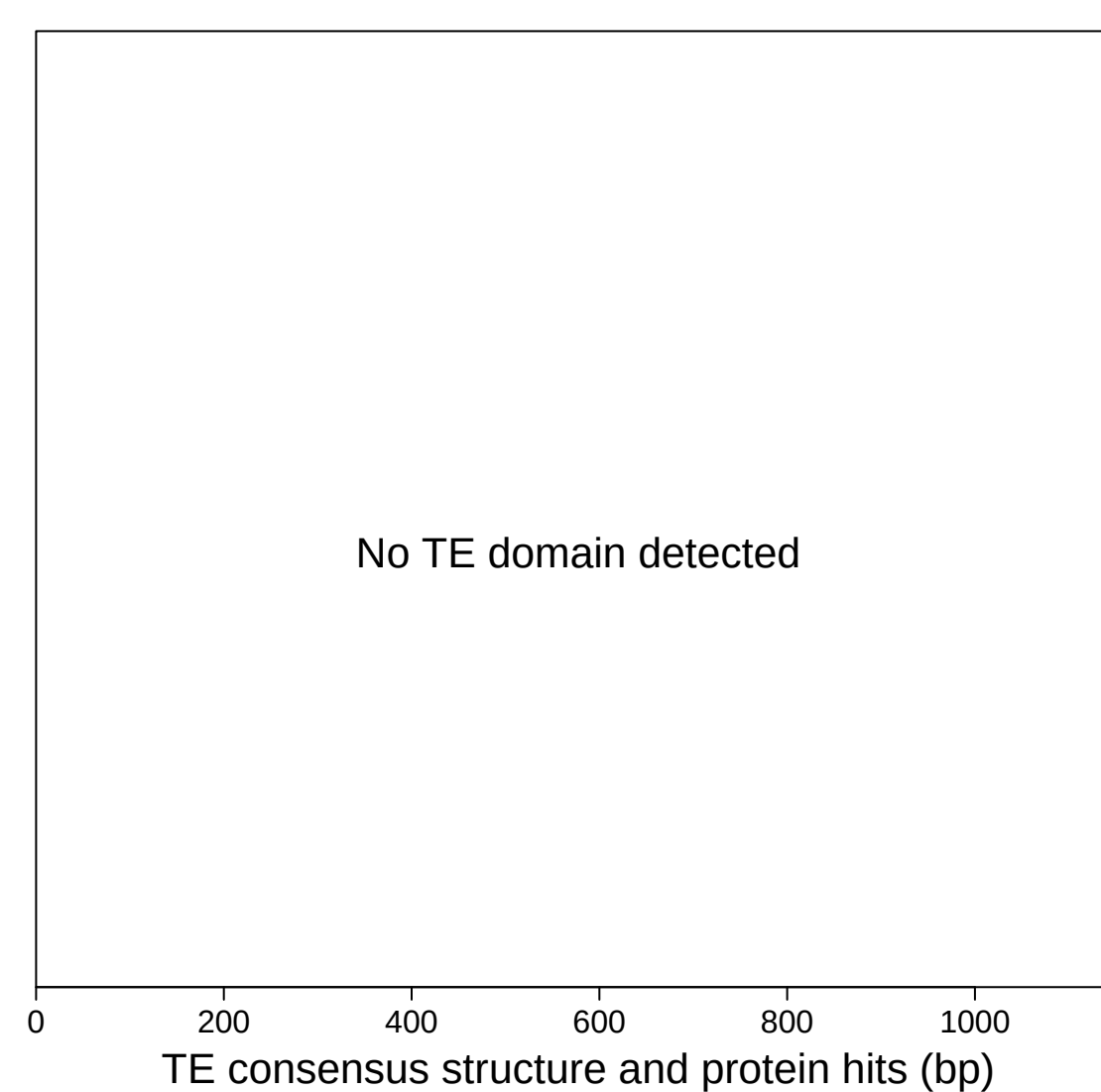
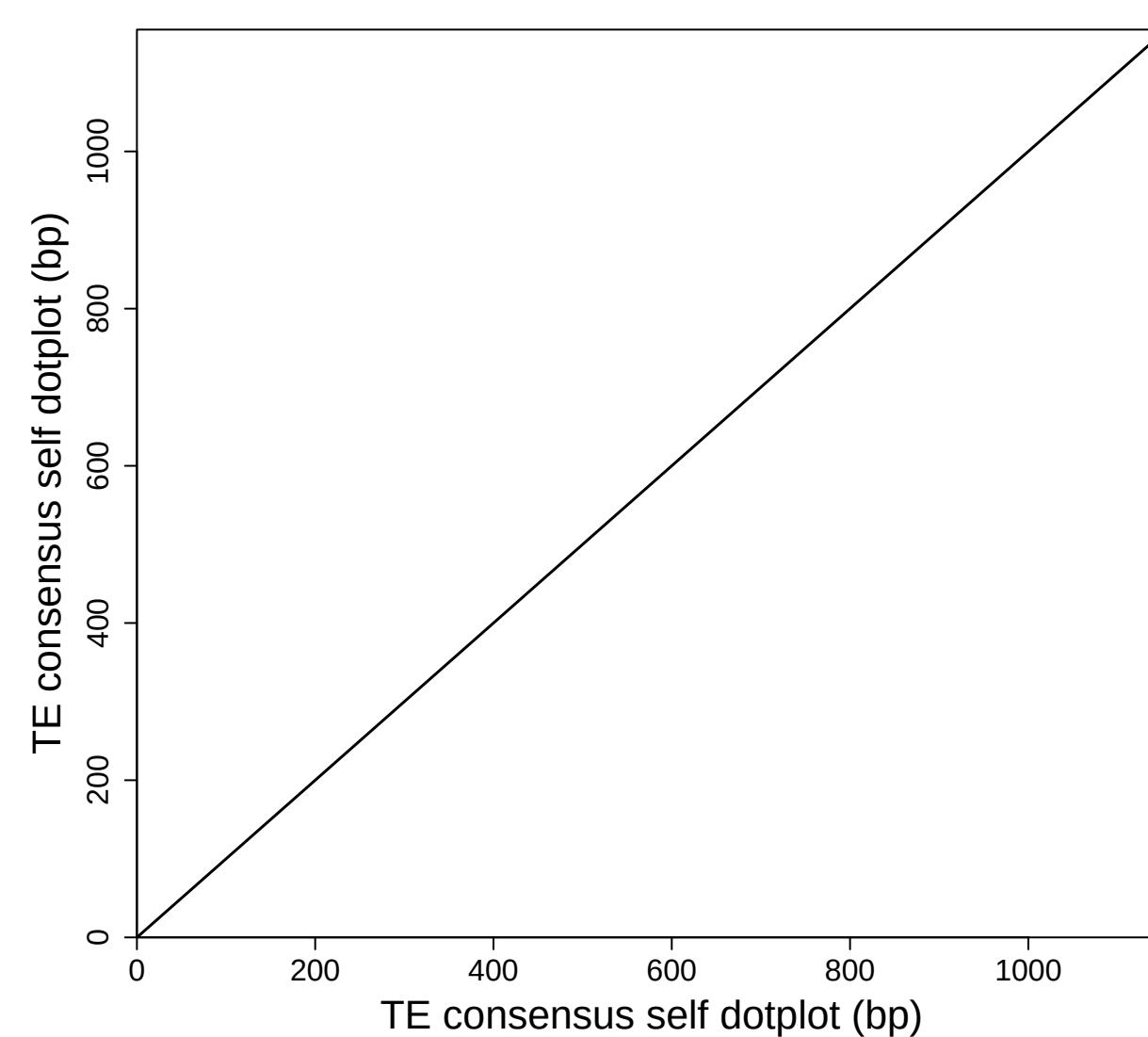
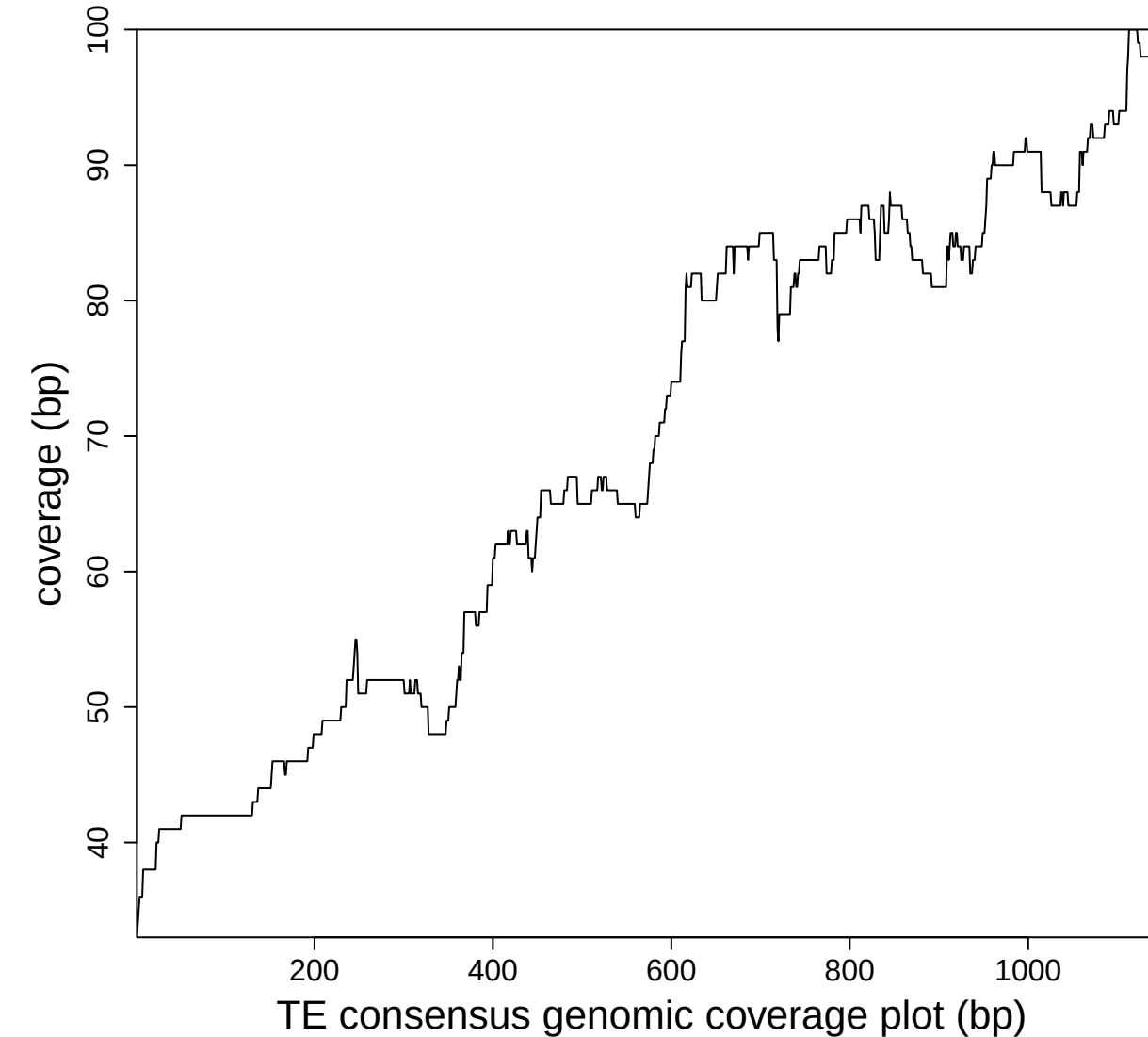
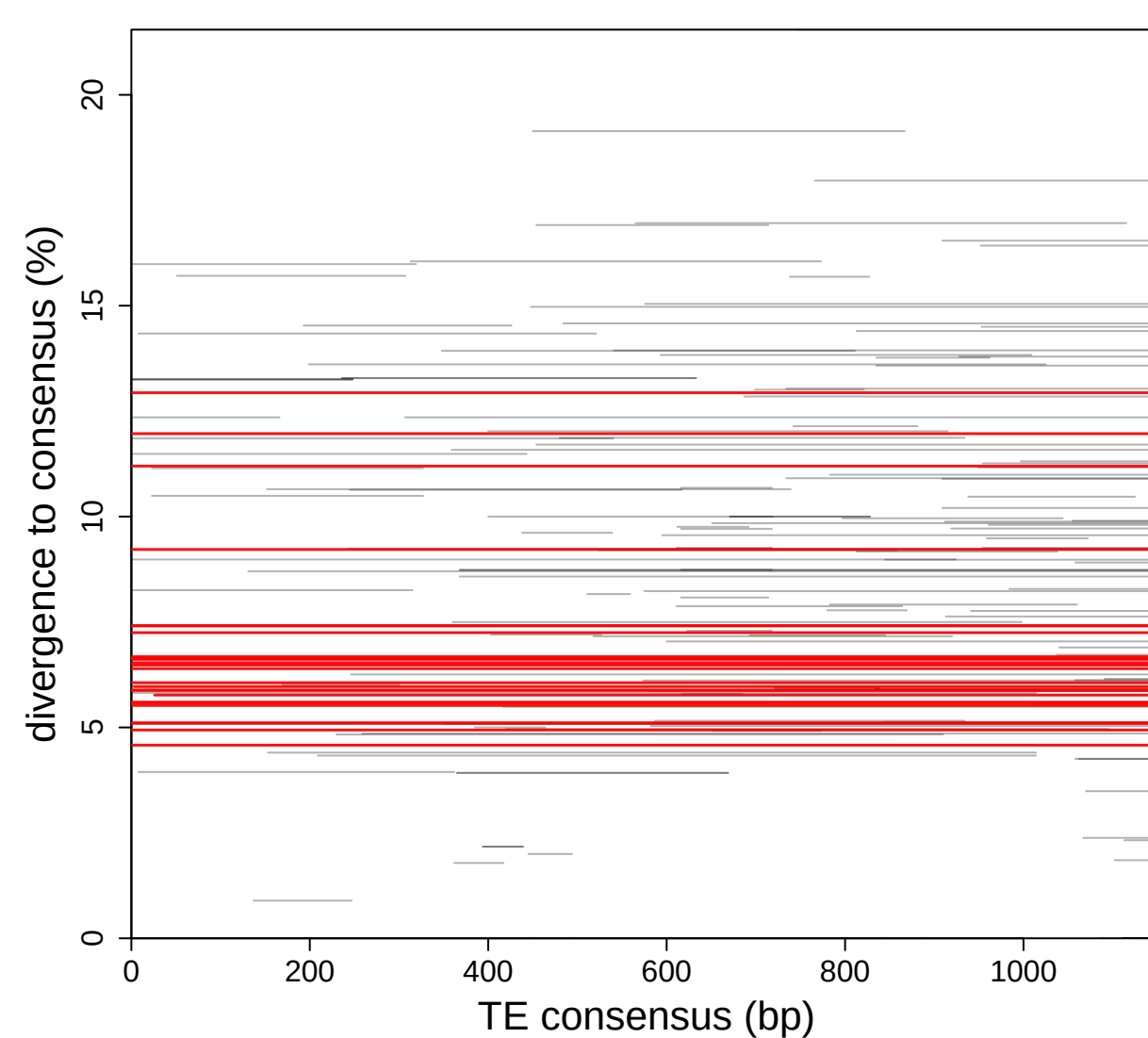


3.fasta.b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.f
size: 6700bp; fragments: 165; full length: 0 (>=6030bp)

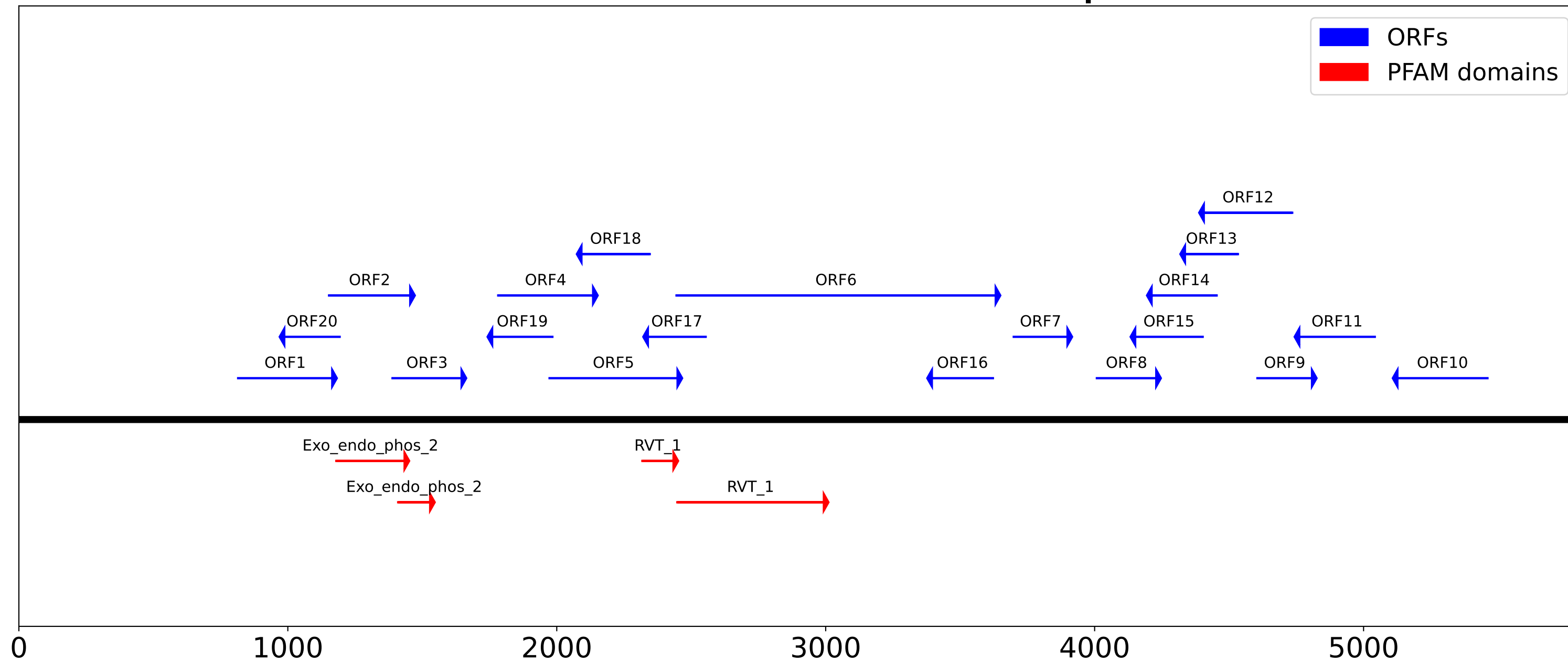
After TEtrimmer Extended plot Blue lines are boundaries



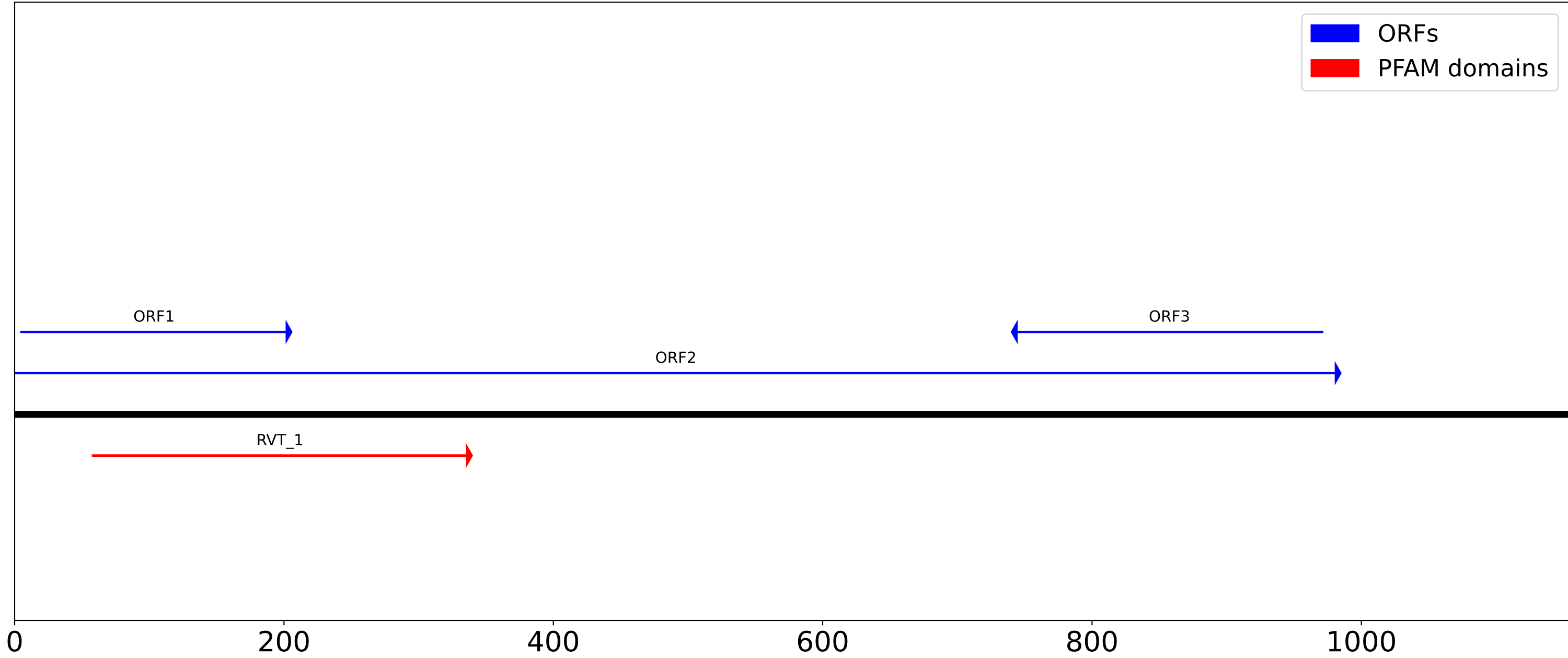
TE: rnd_5_family_2733
size: 1155bp; fragments: 177; full length: 26 (≥ 1039.5 bp)



After TETrimmer ORF and PFAM domain plot



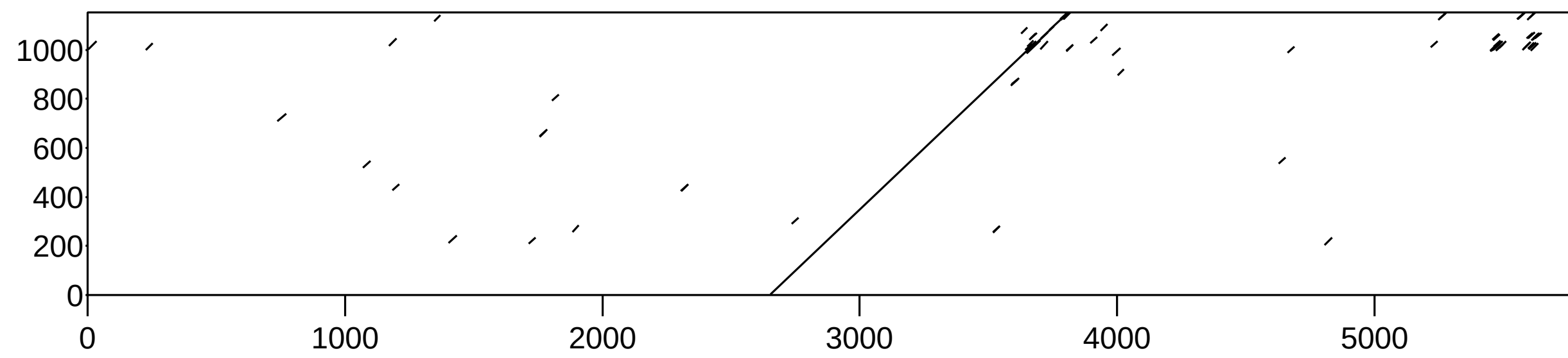
Before TEtrimmer ORF and PFAM domain plot



Dotplot

(windowsize = 25, threshold = 50.00 10/10/25)

nsensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)