

Start crop Point



End crop Point

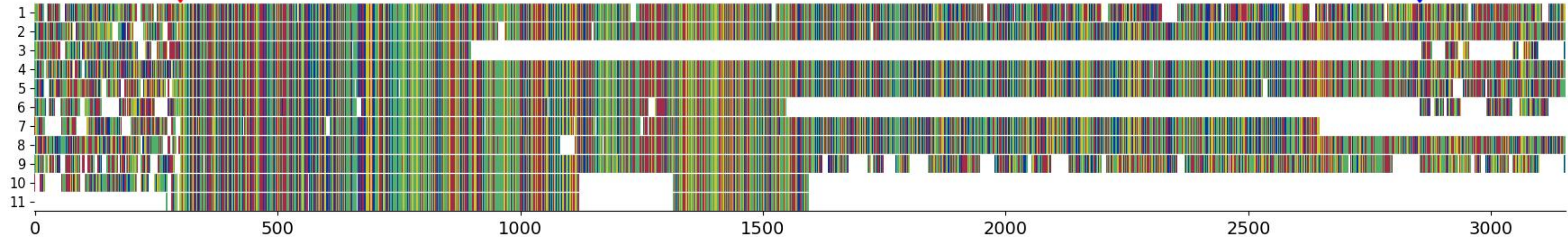


MSA length = 2554



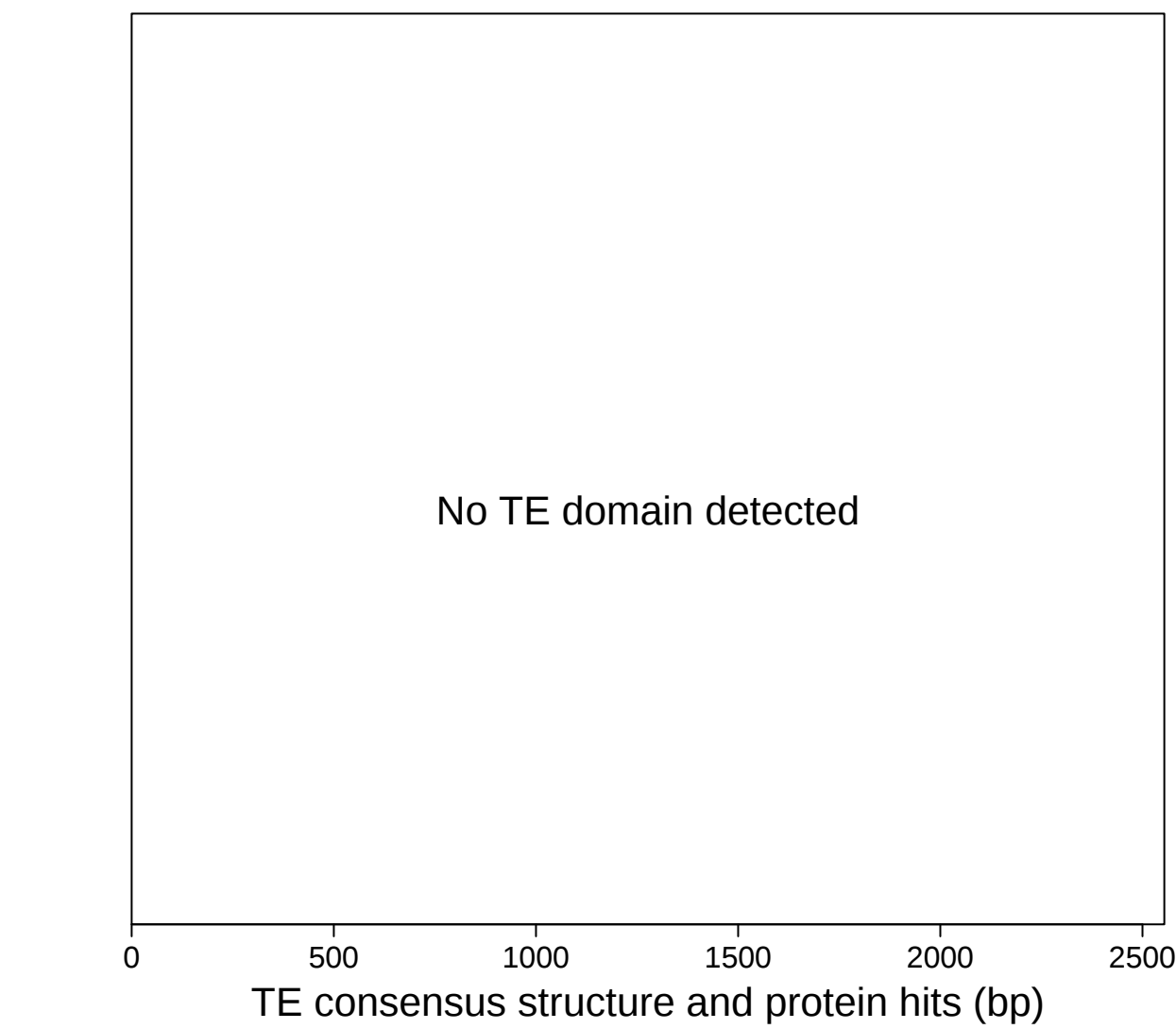
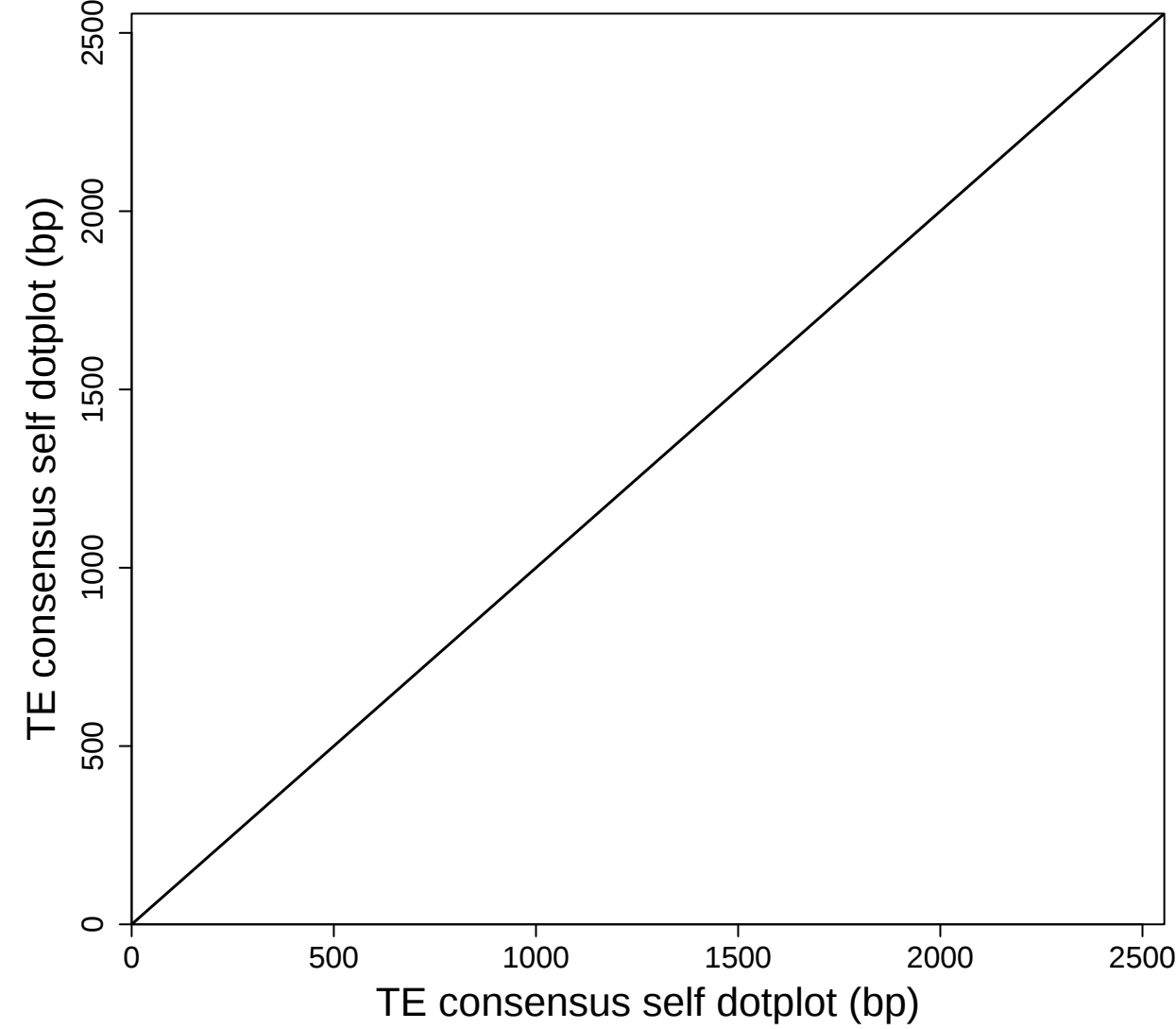
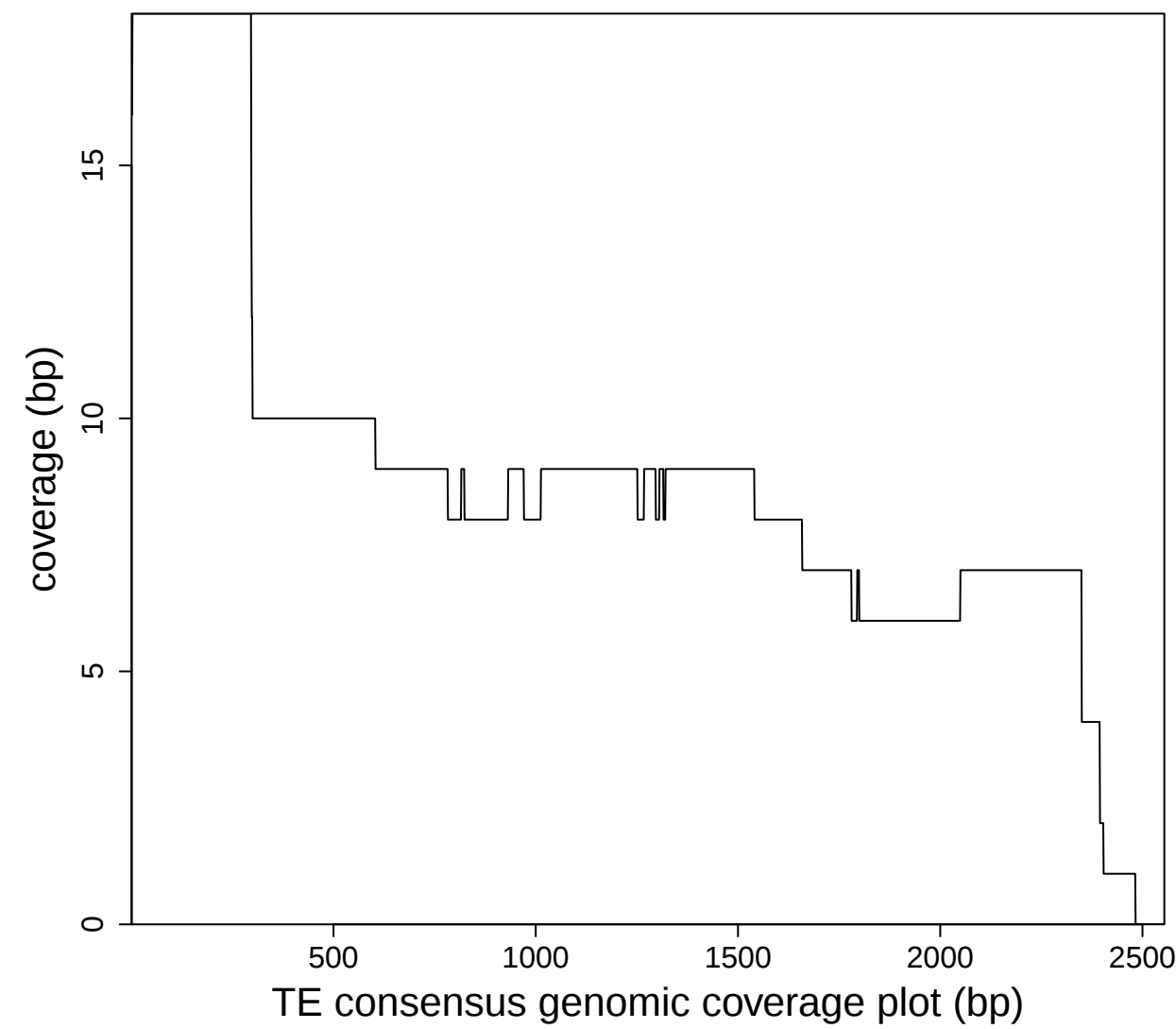
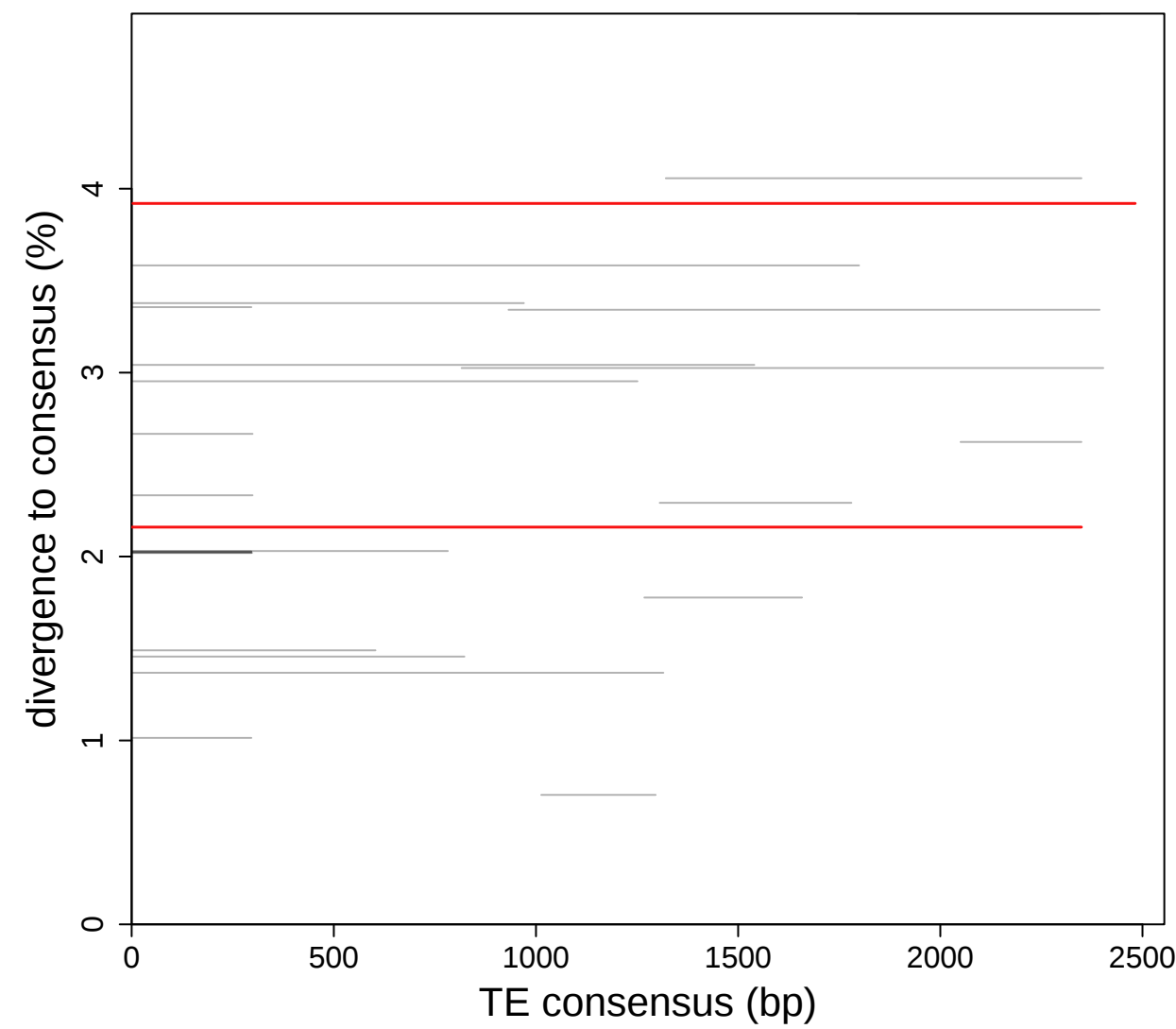
Start crop Point

End crop Point



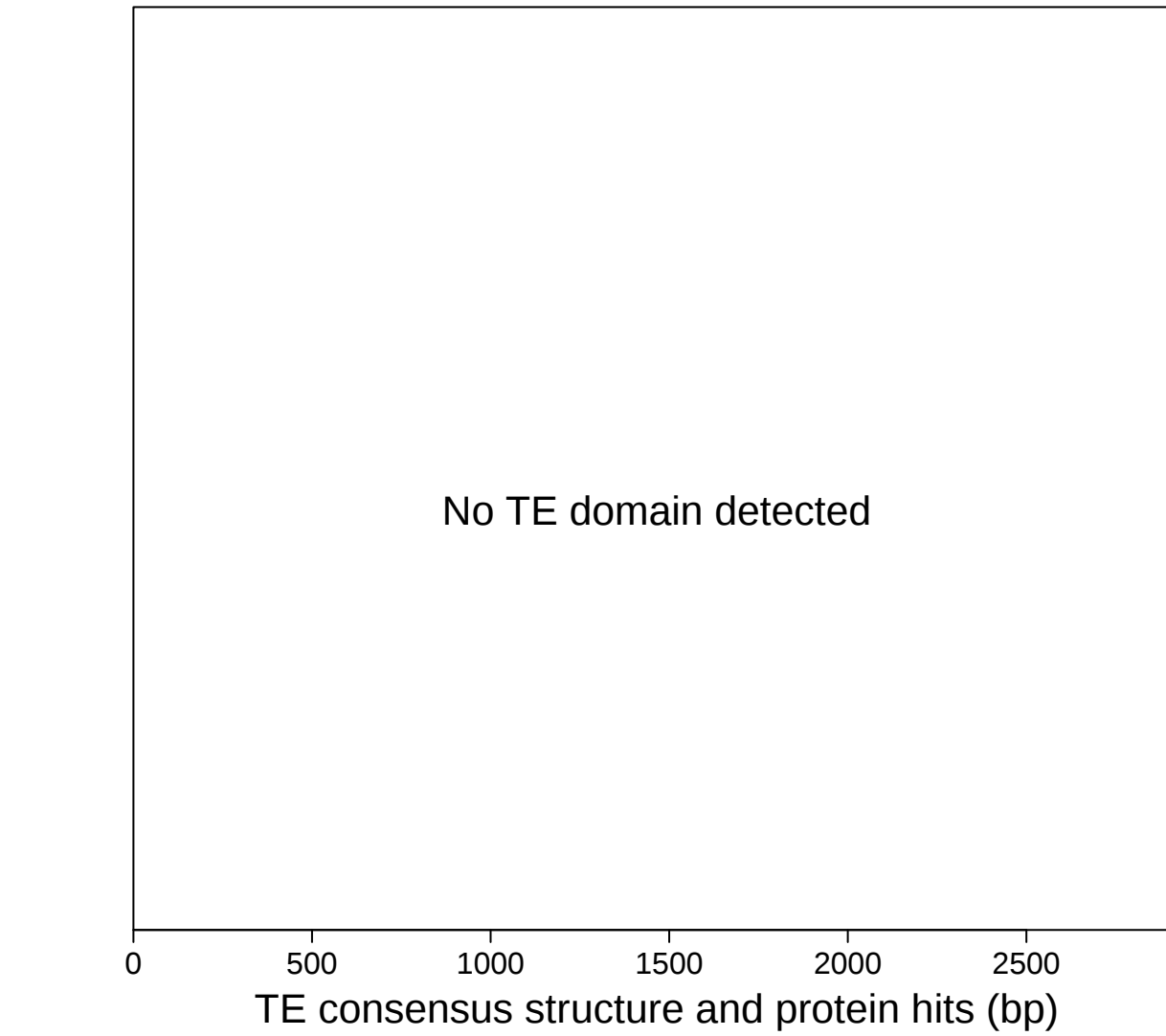
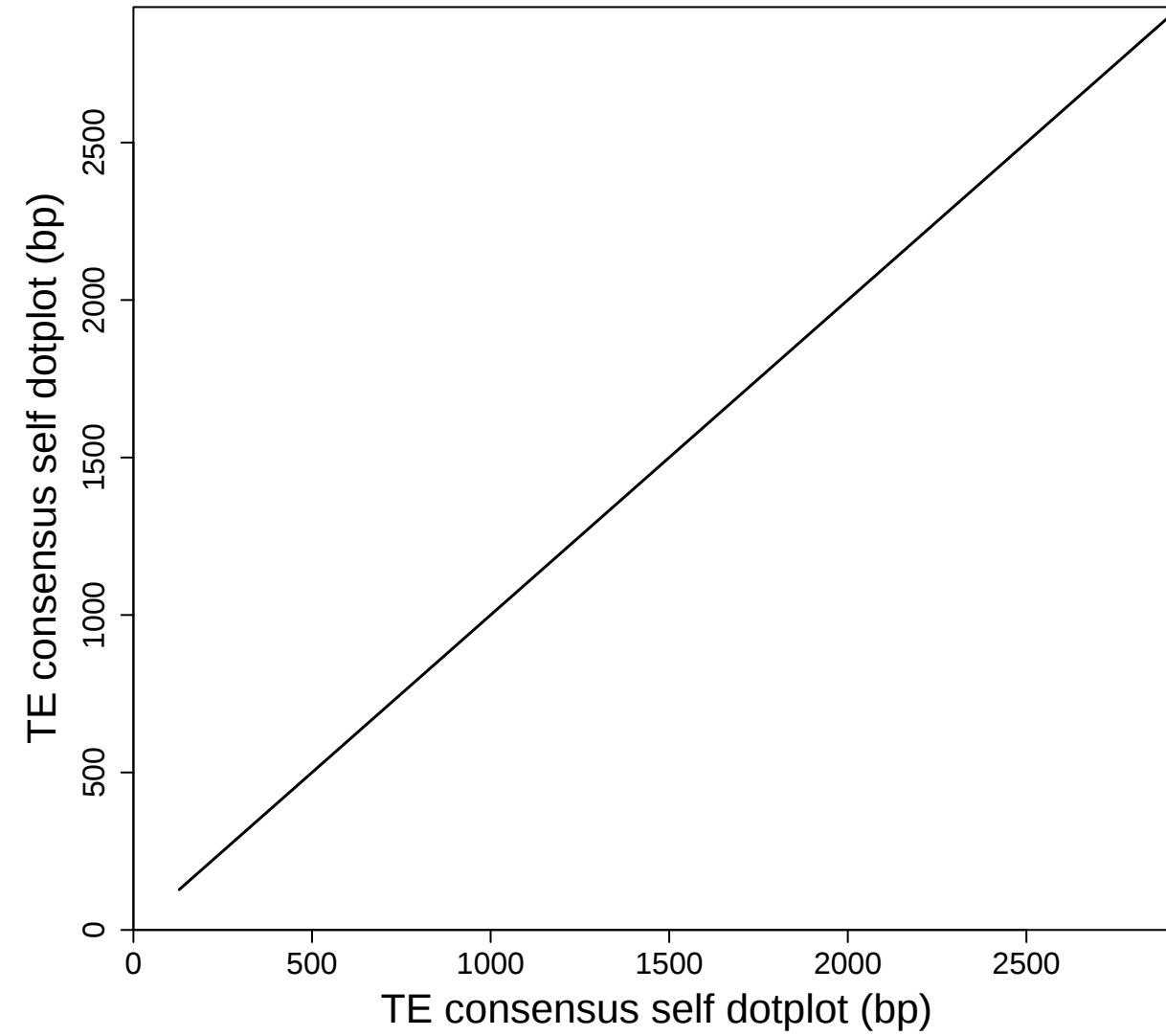
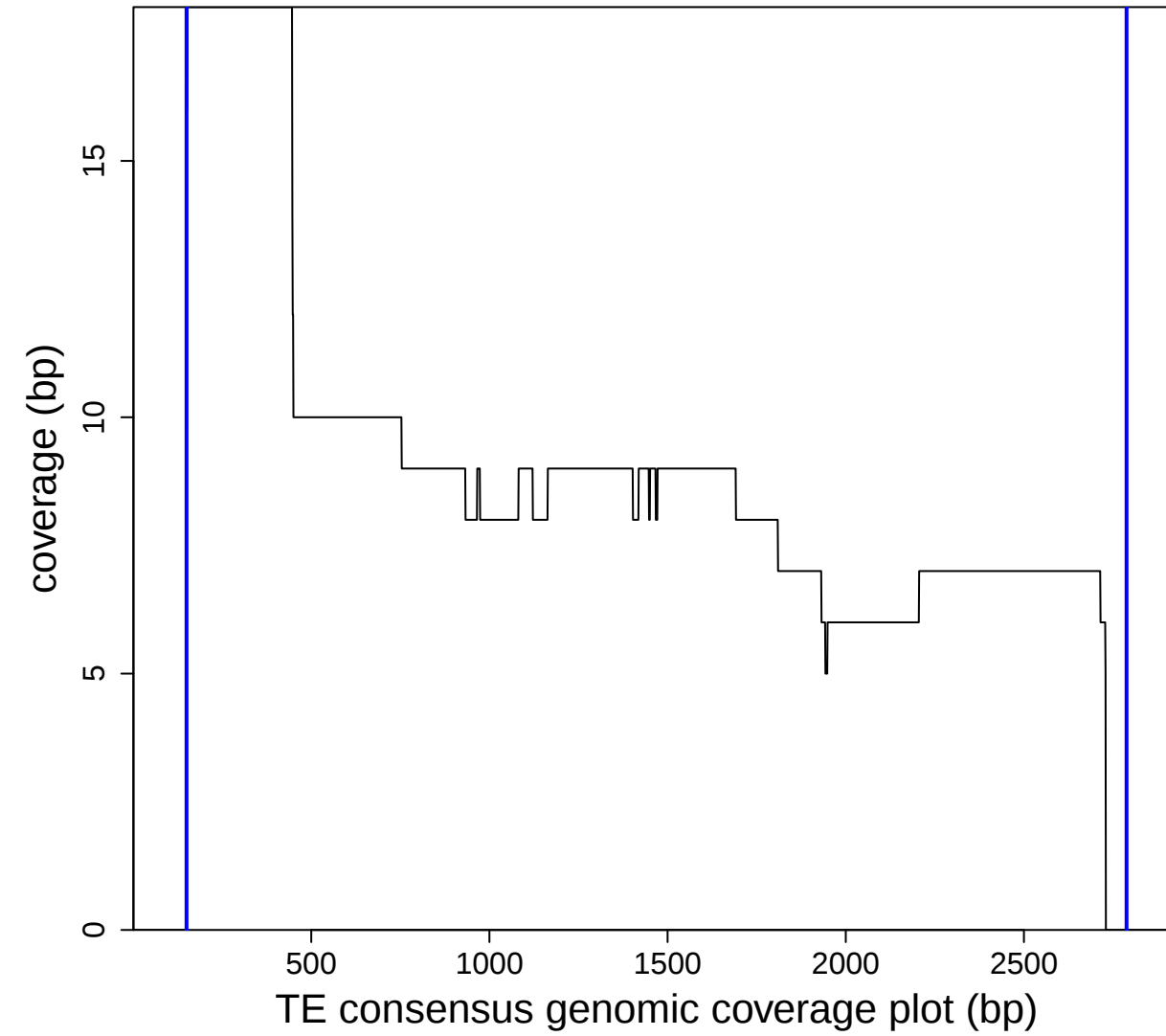
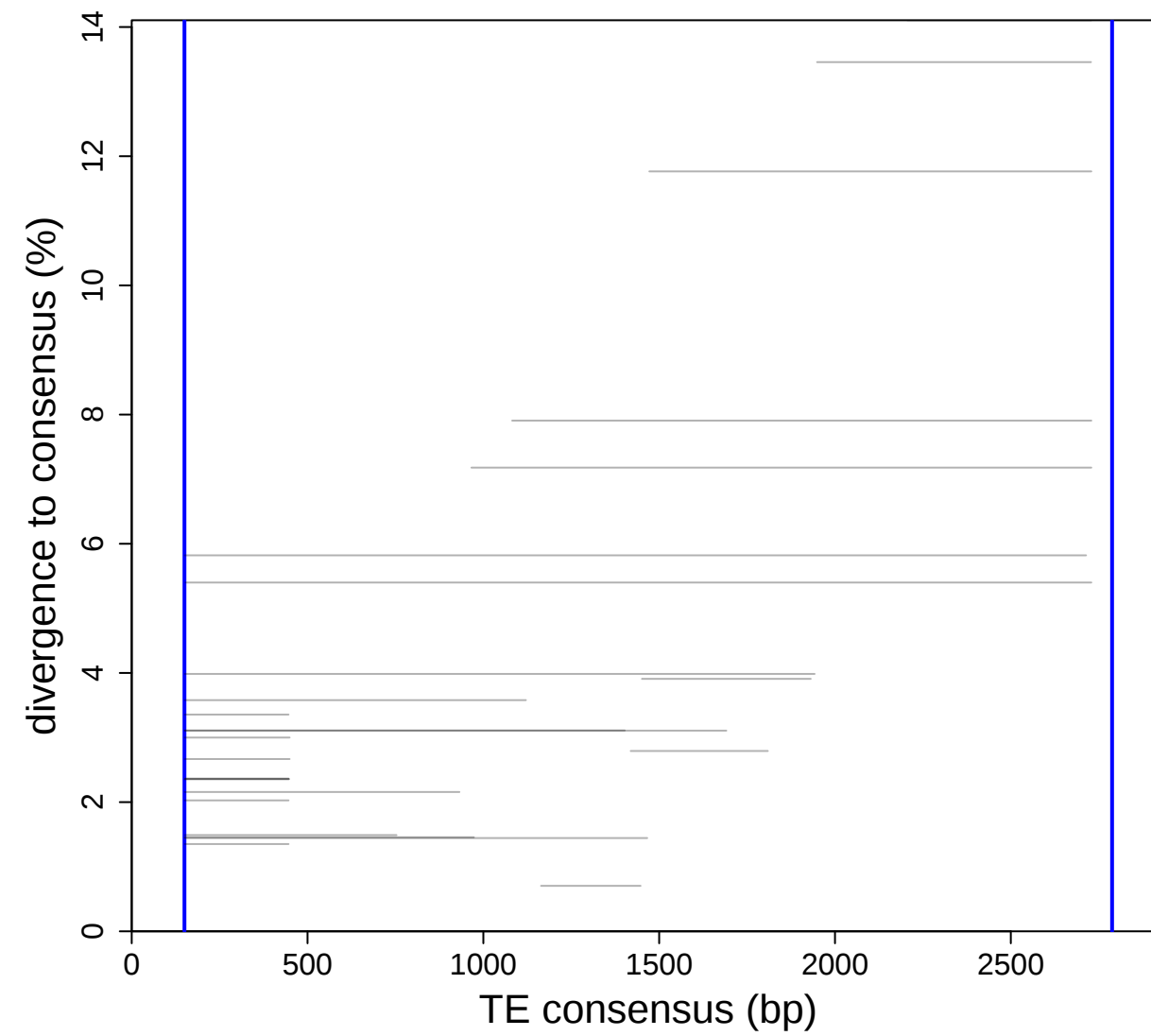
b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa
size: 2554bp; fragments: 26; full length: 2 (≥ 2298.6 bp)

After TETrimmer 2554 bp

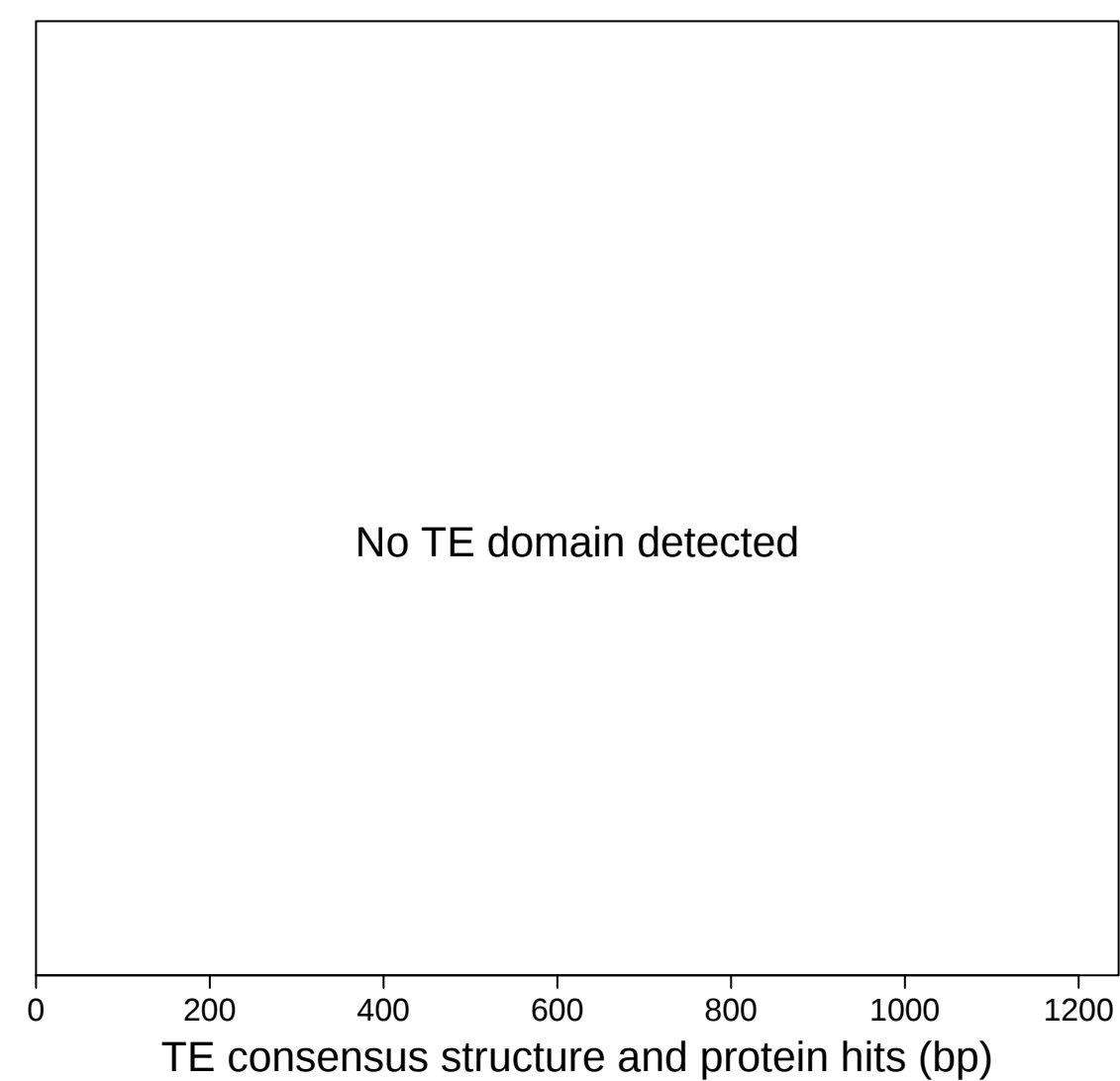
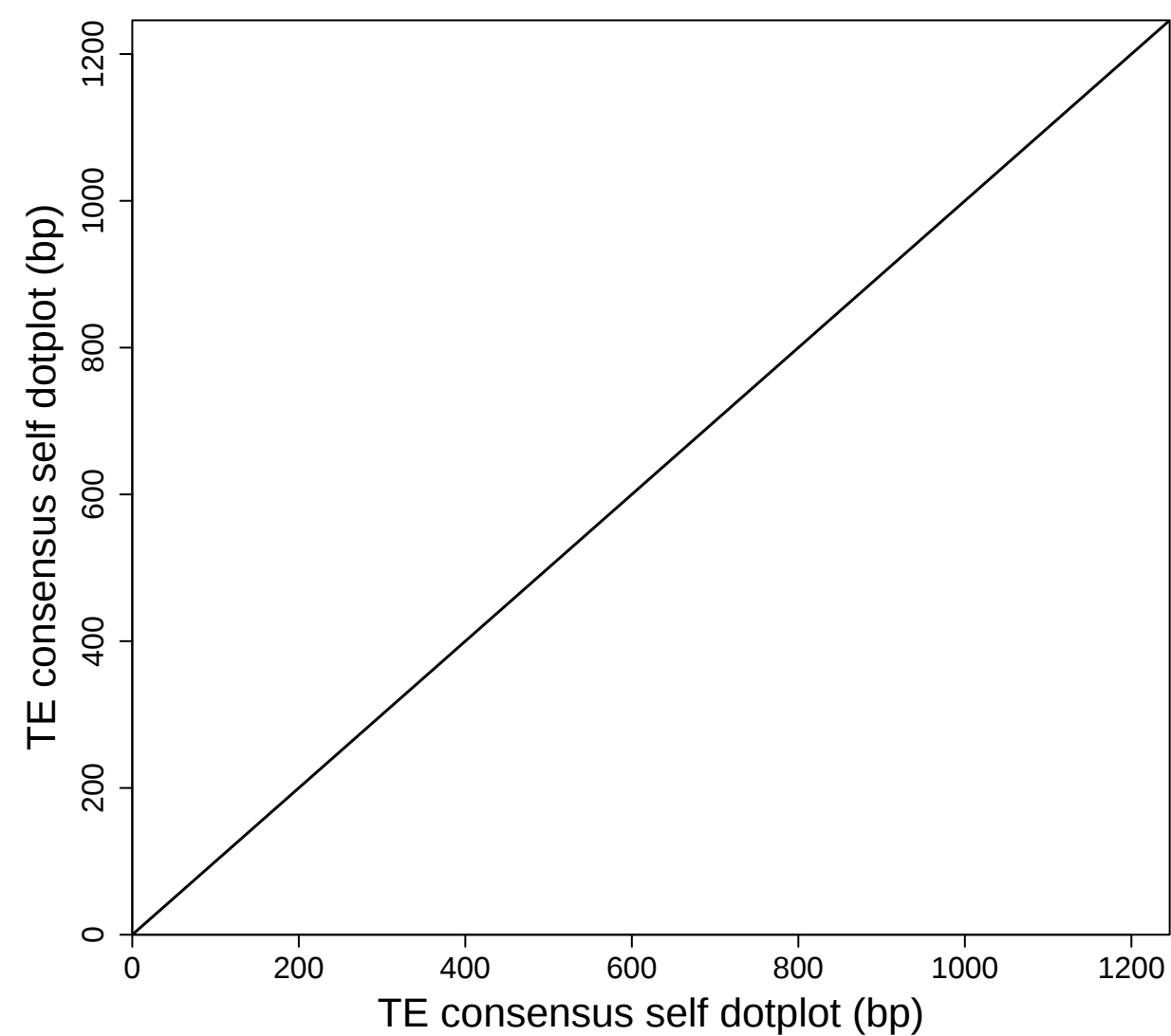
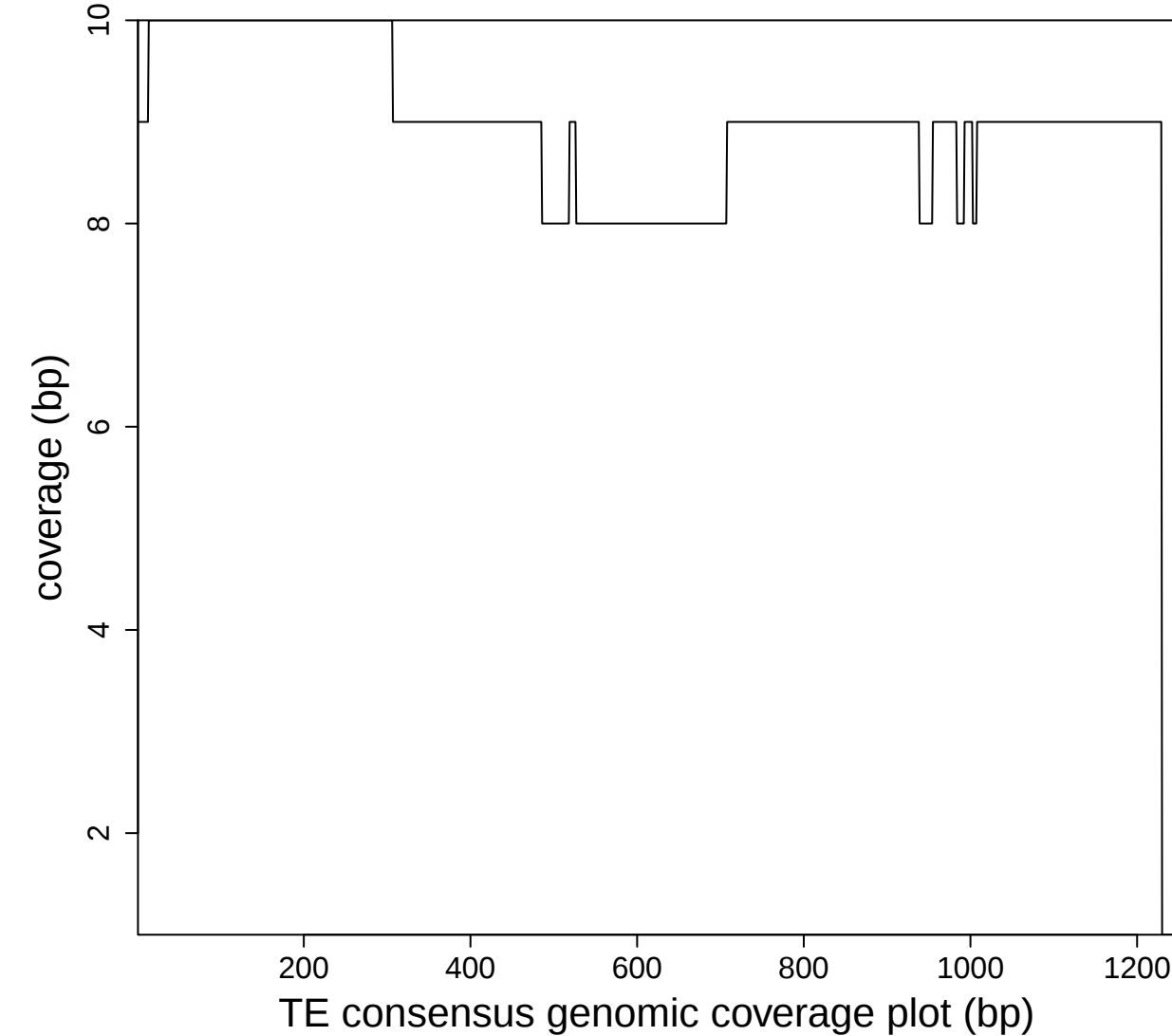
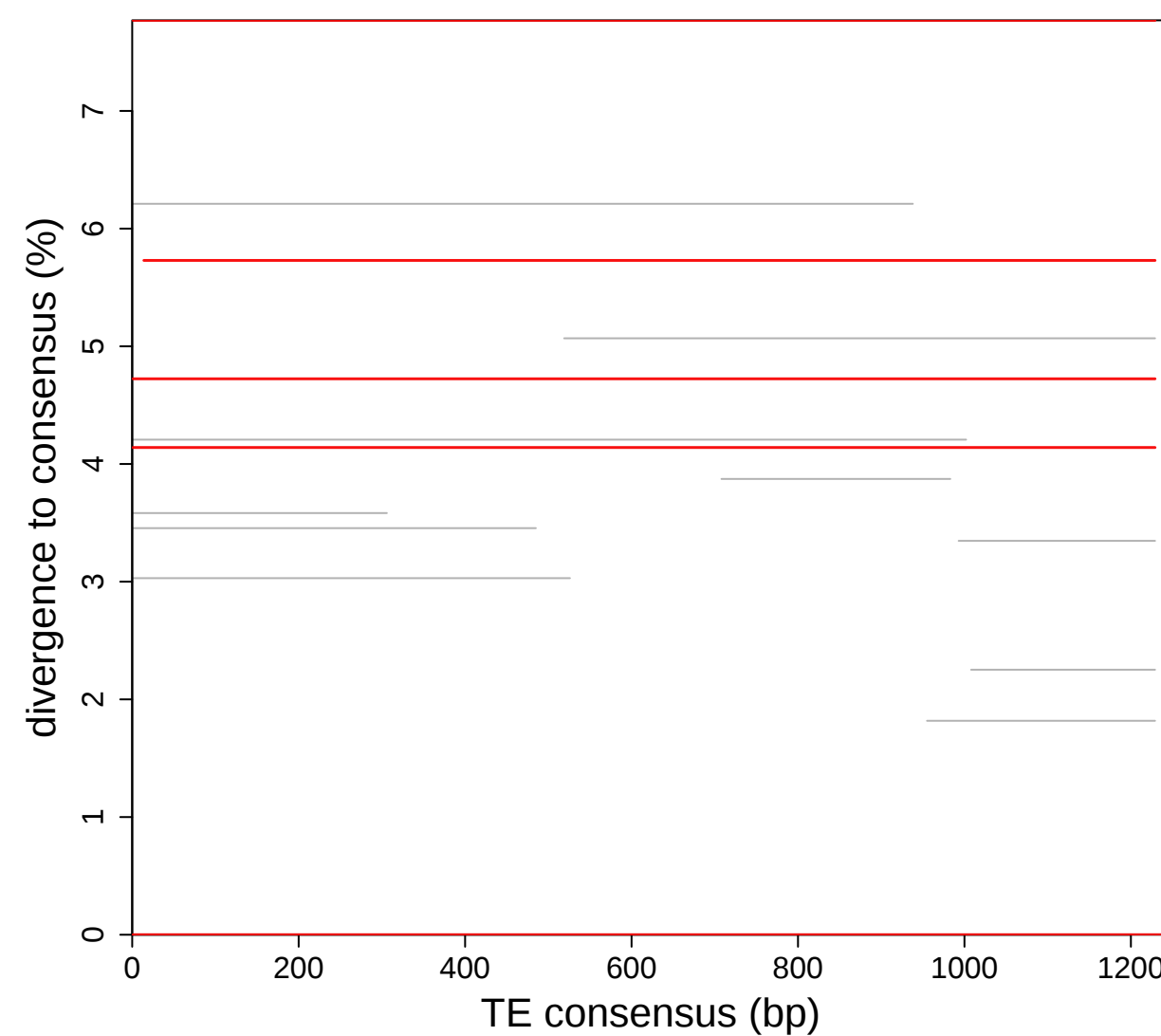


.fasta.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa
size: 2938bp; fragments: 26; full length: 0 (>=2644.2bp)

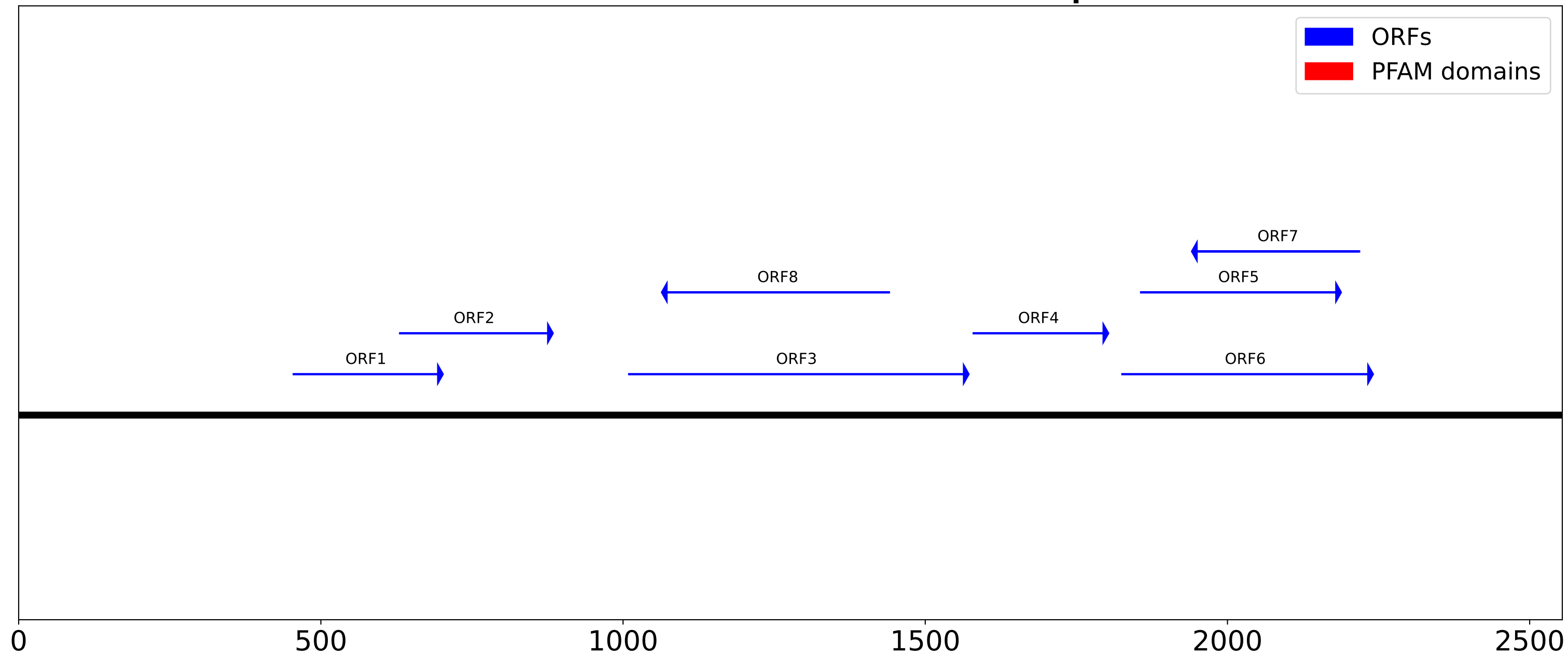
After TEtrimmer Extended plot Blue lines are boundaries



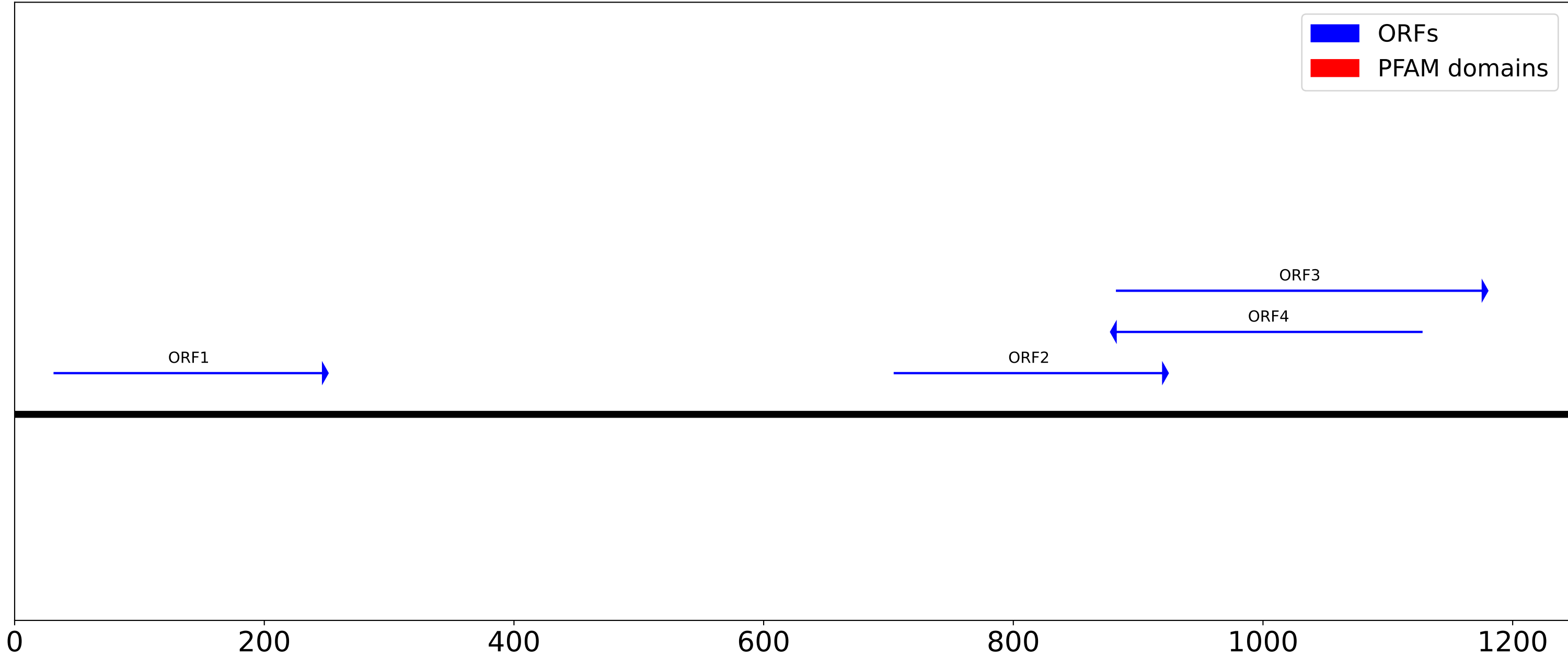
TE: ltr_1_family_886
size: 1246bp; fragments: 15; full length: 5 (≥ 1121.4 bp)



After TEtrimmer ORF and PFAM domain plot



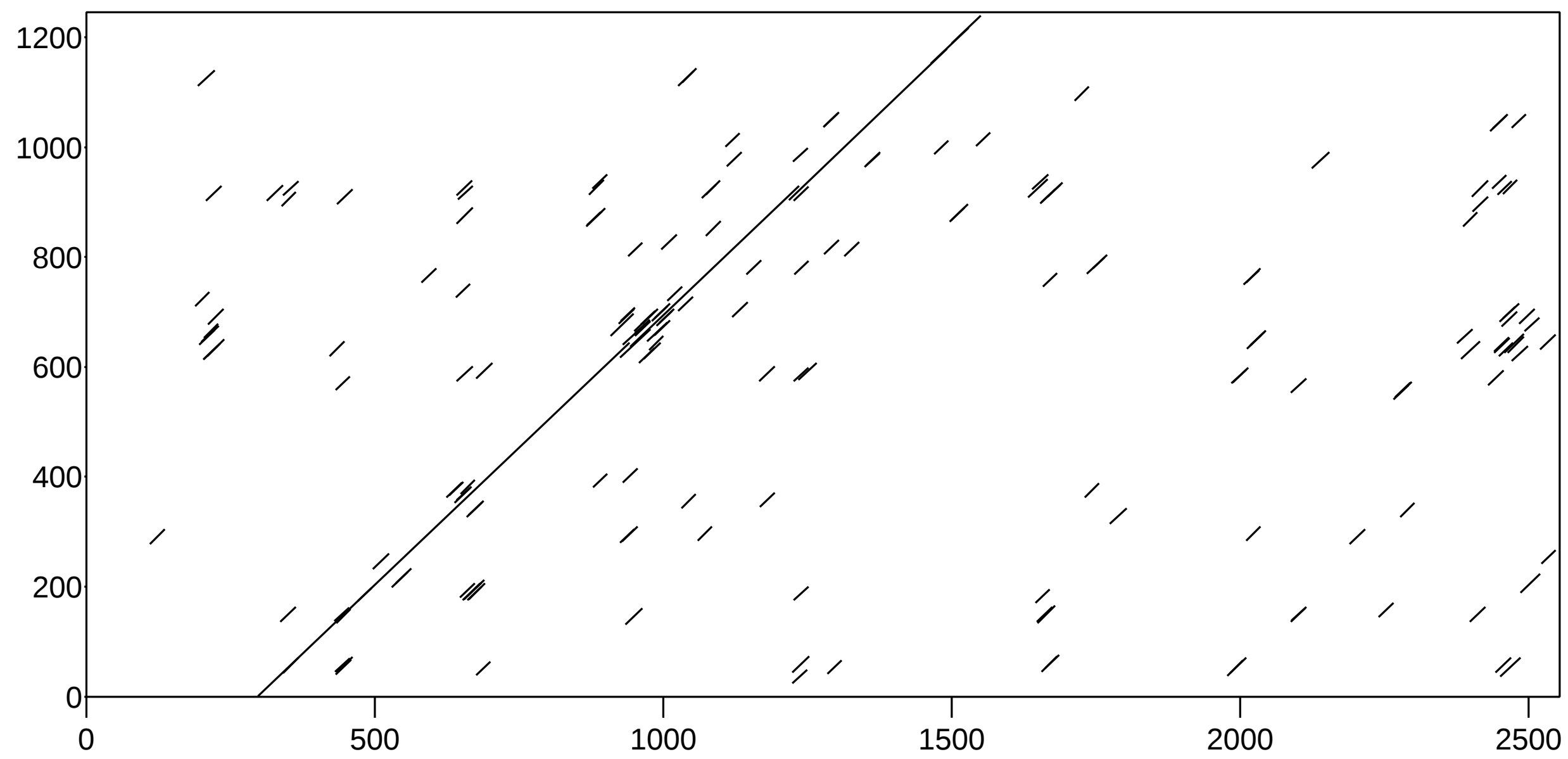
Before TEtrimmer ORF and PFAM domain plot



Dotplot

(windowsize = 25, threshold = 50.00 09/10/25)

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)