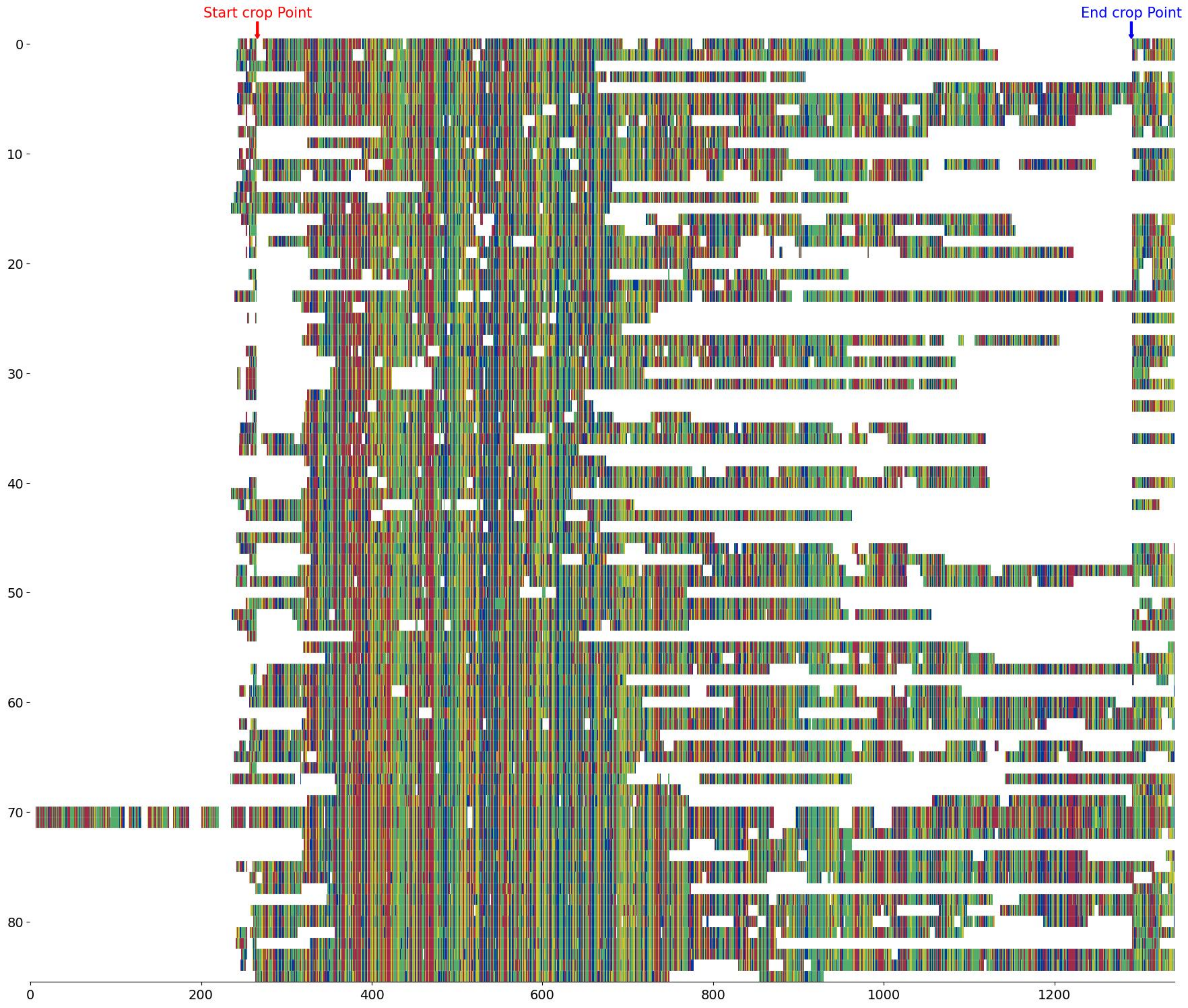
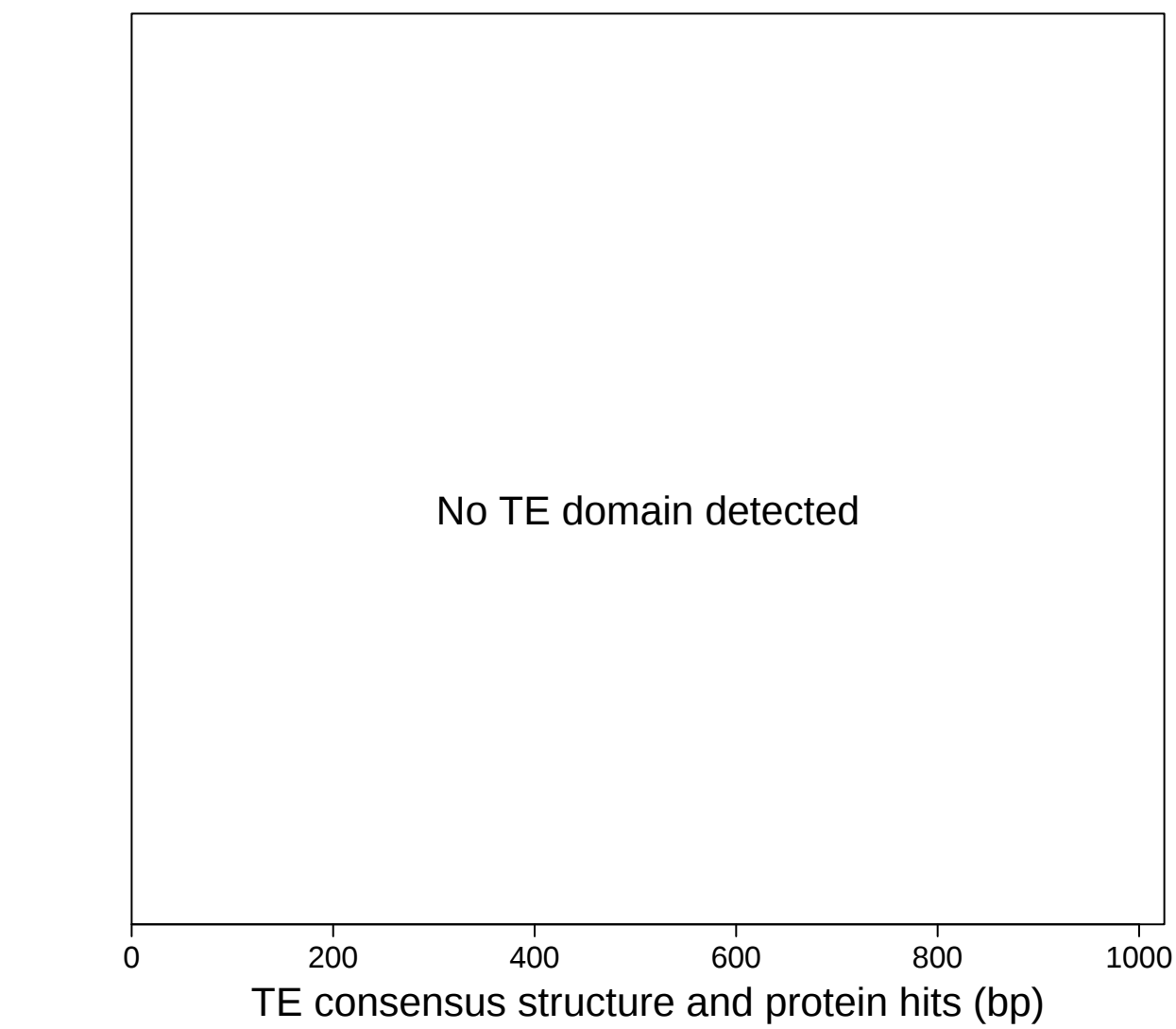
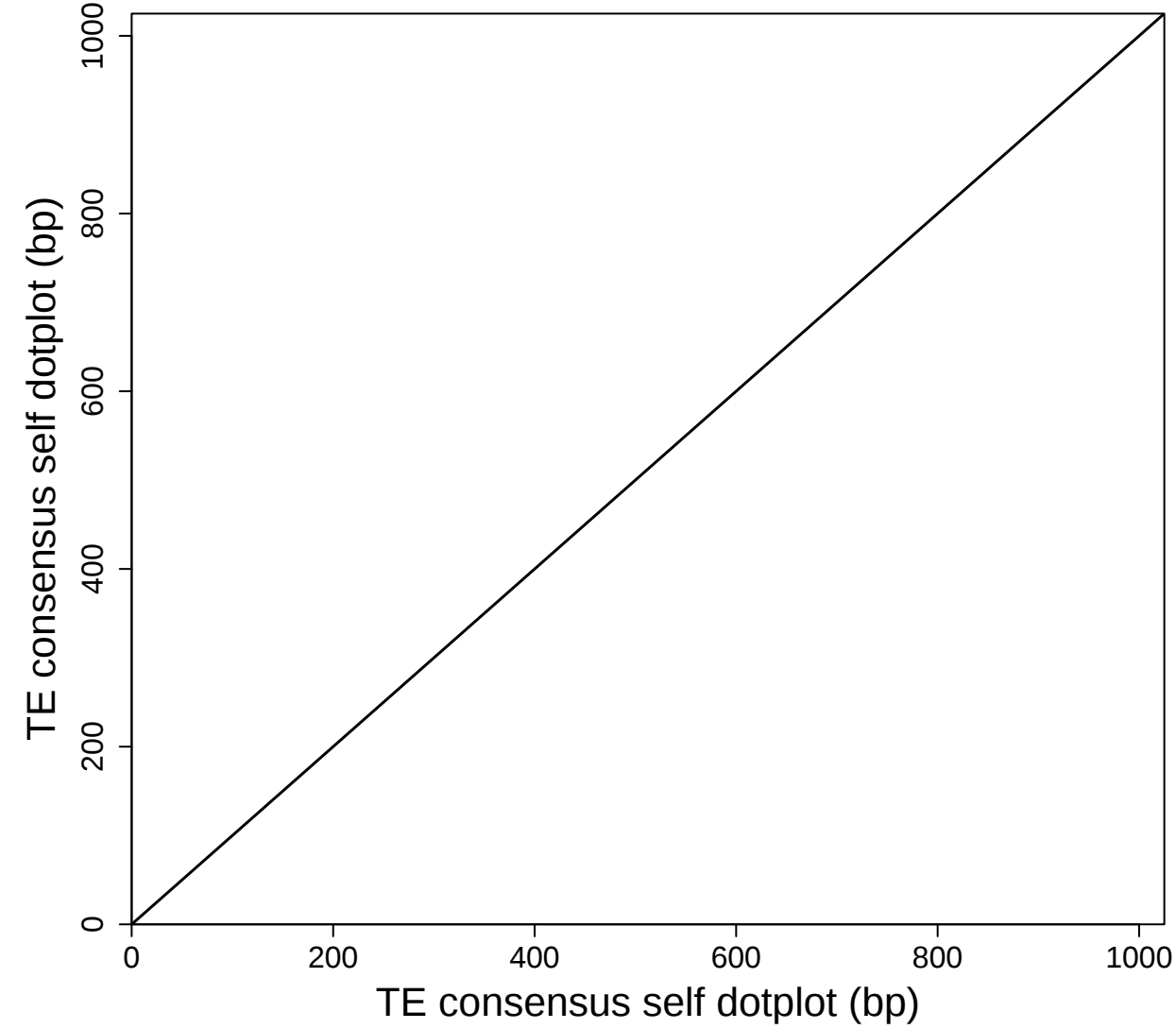
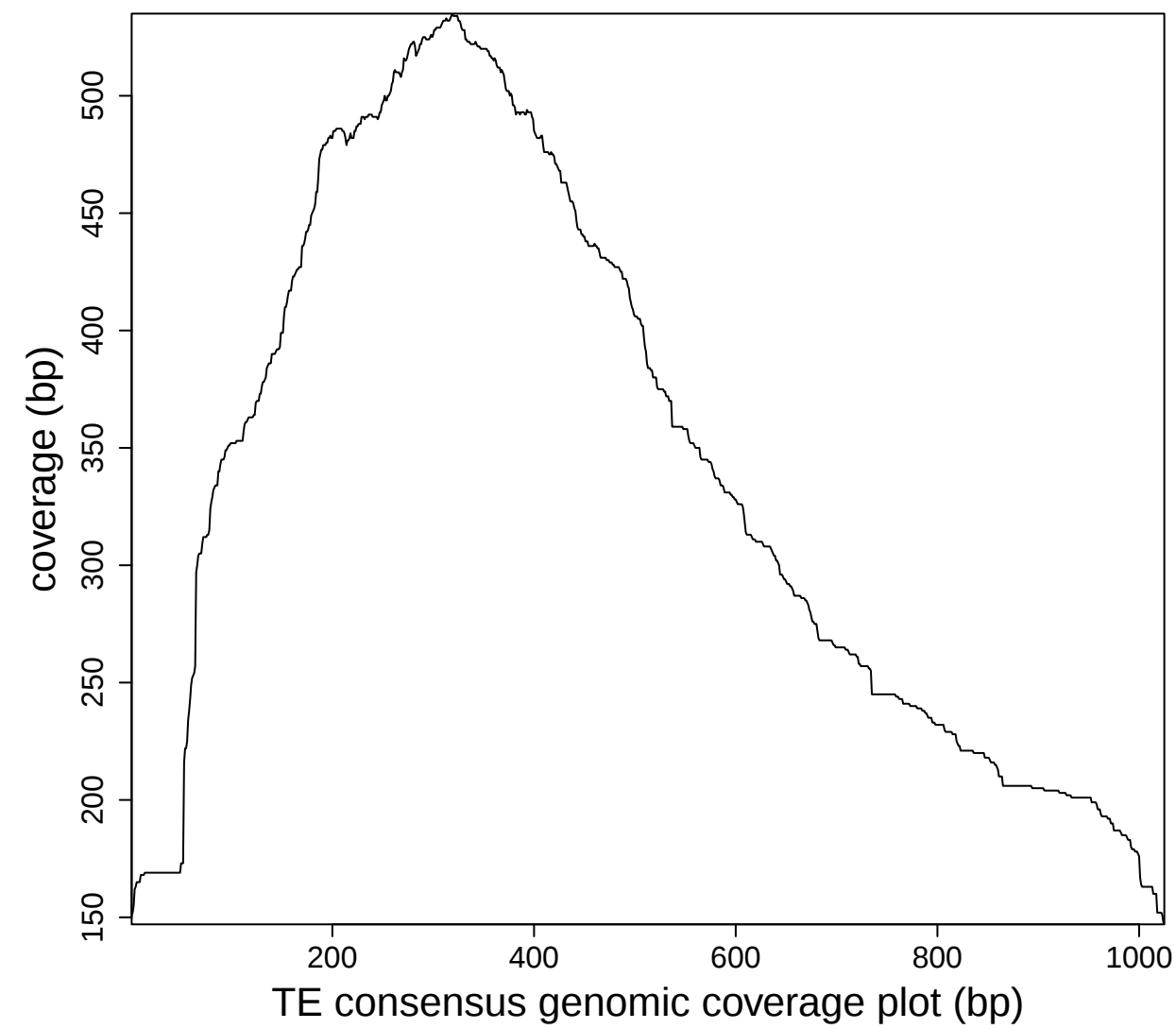
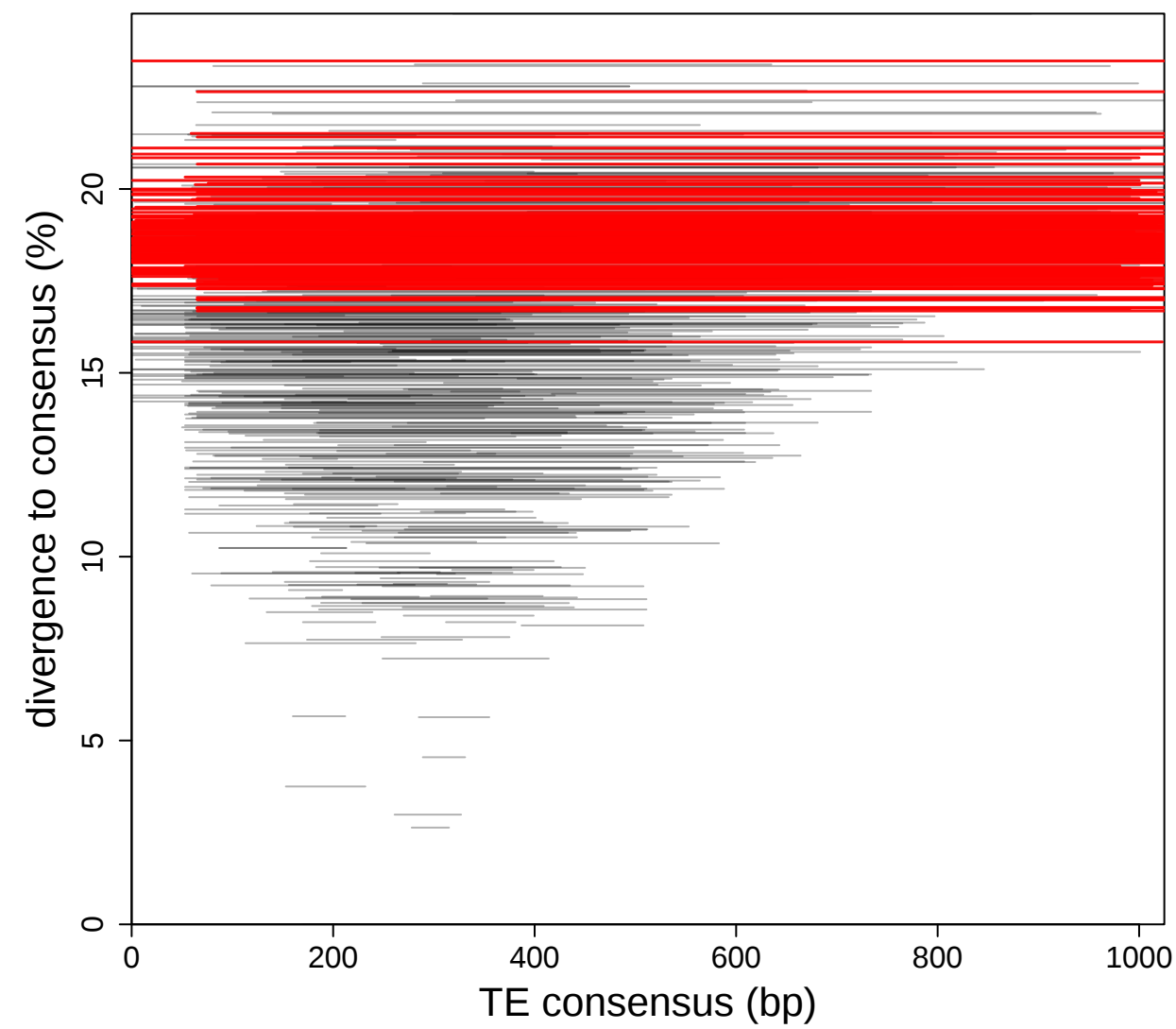






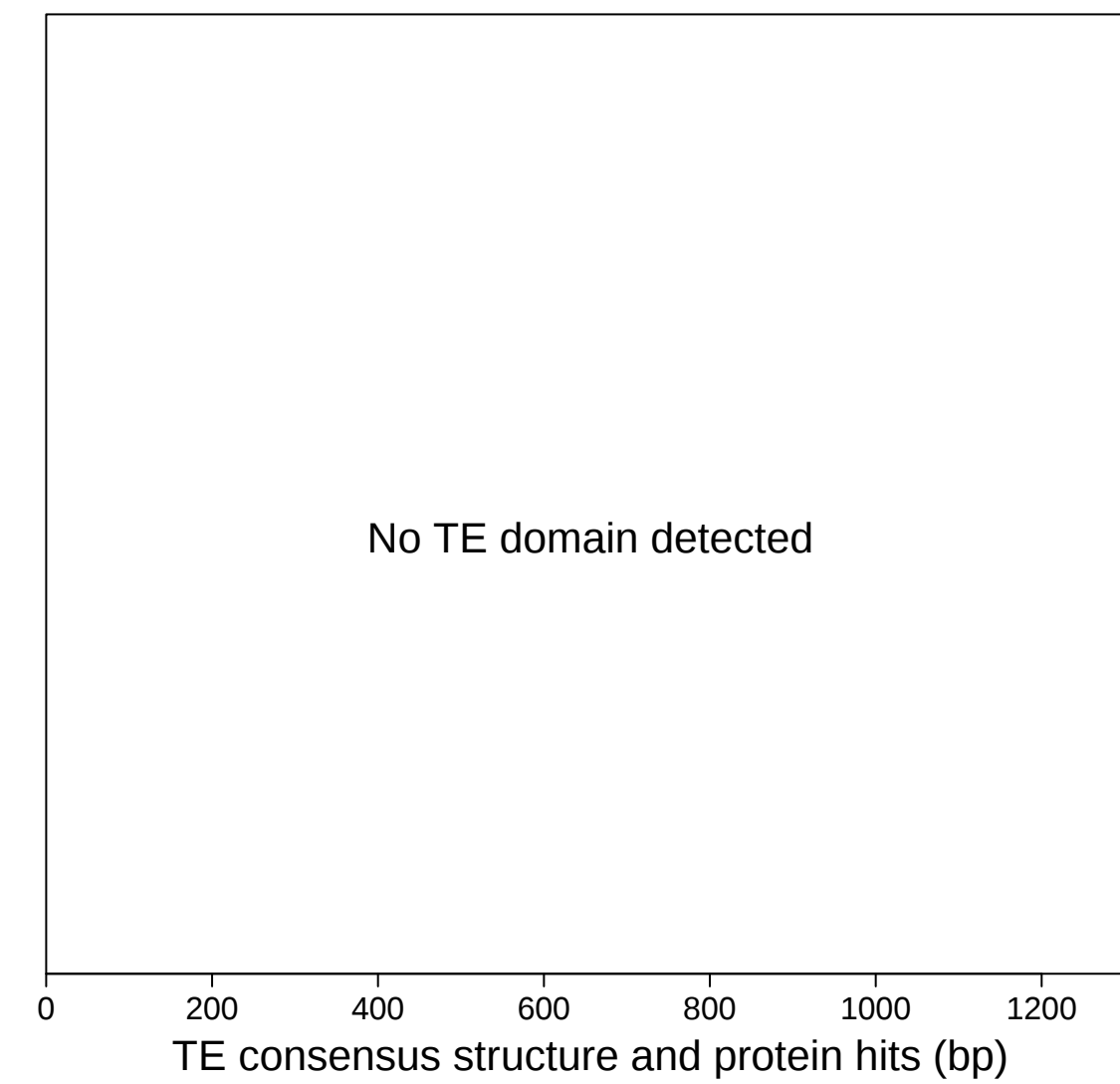
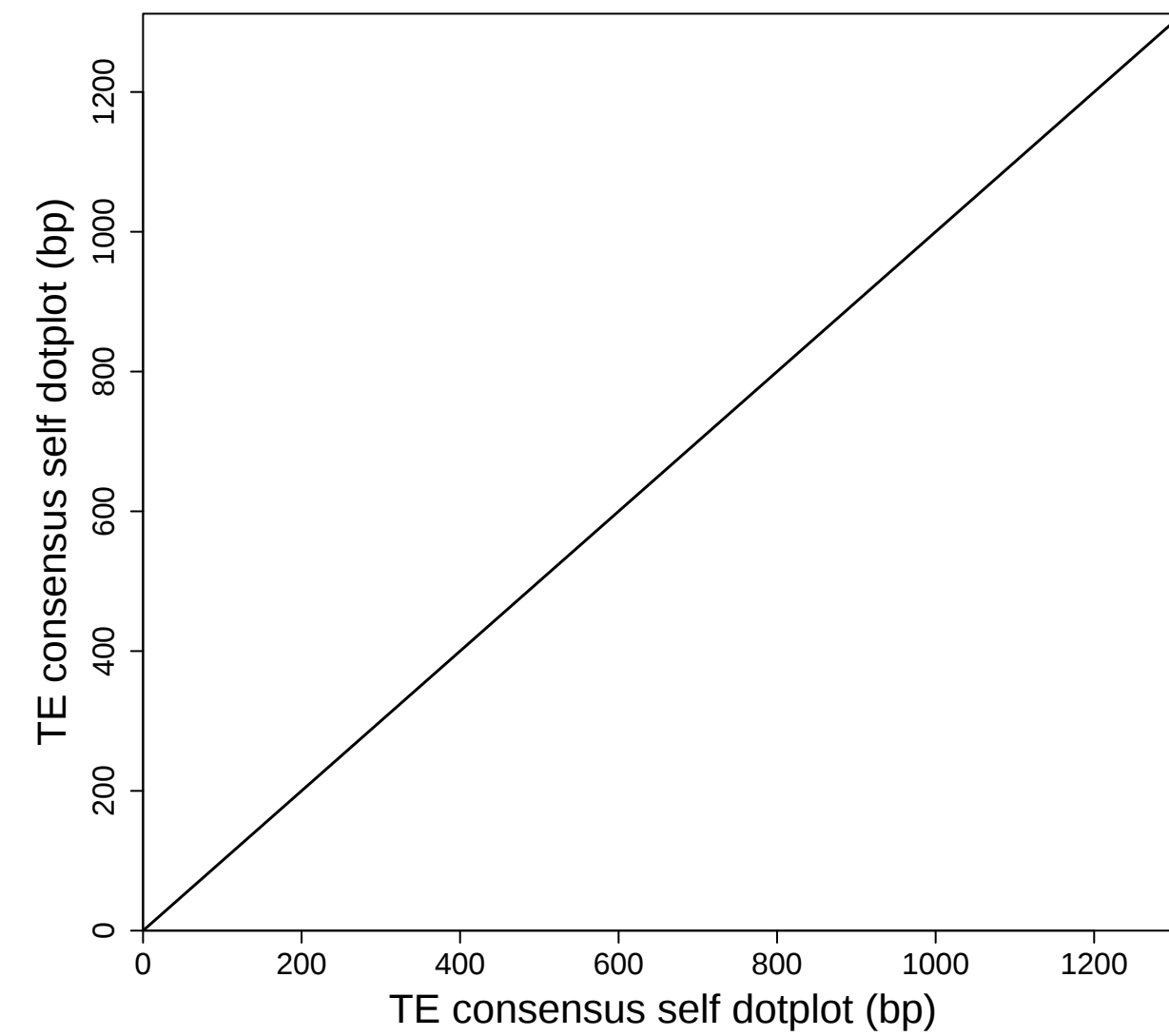
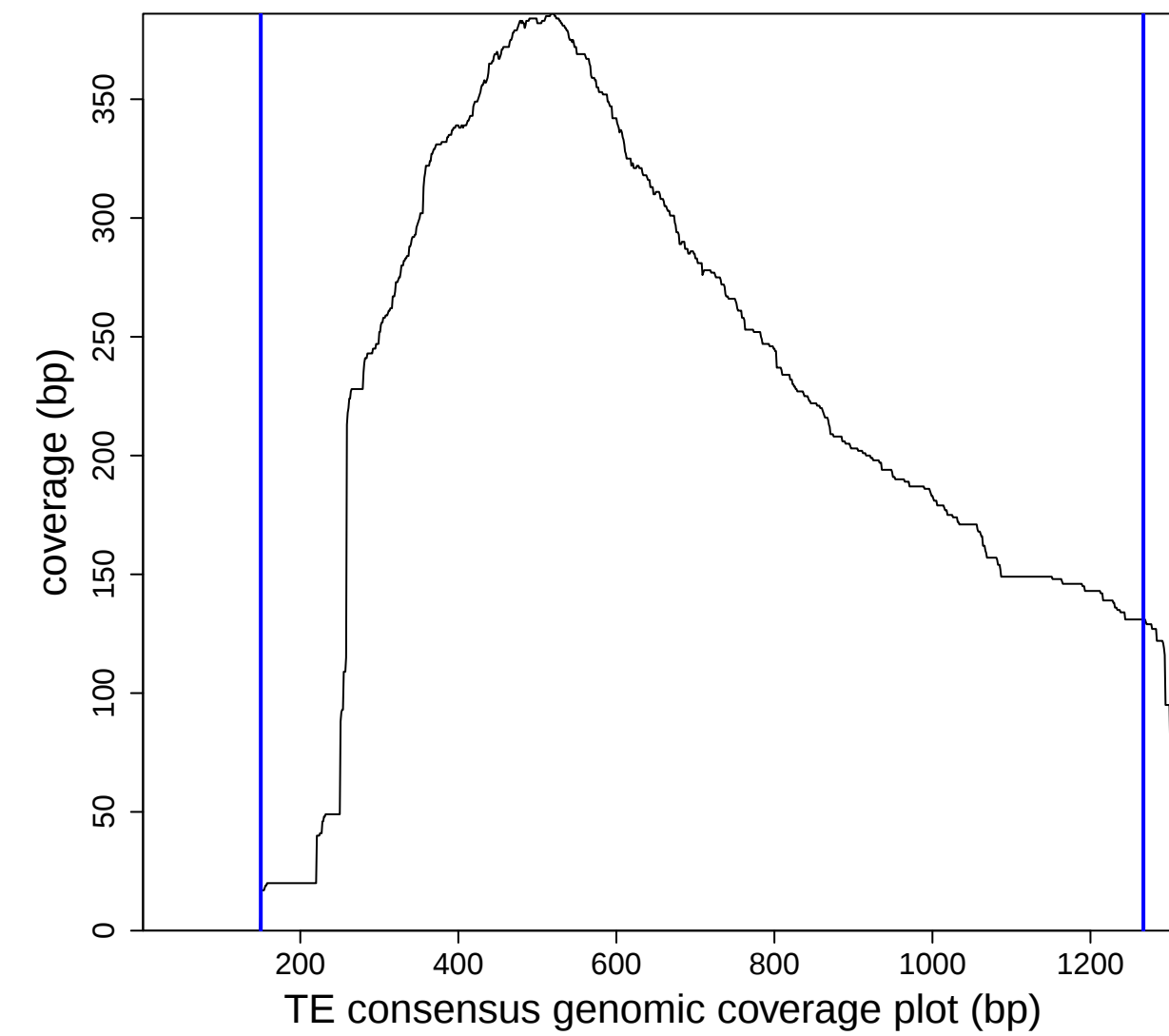
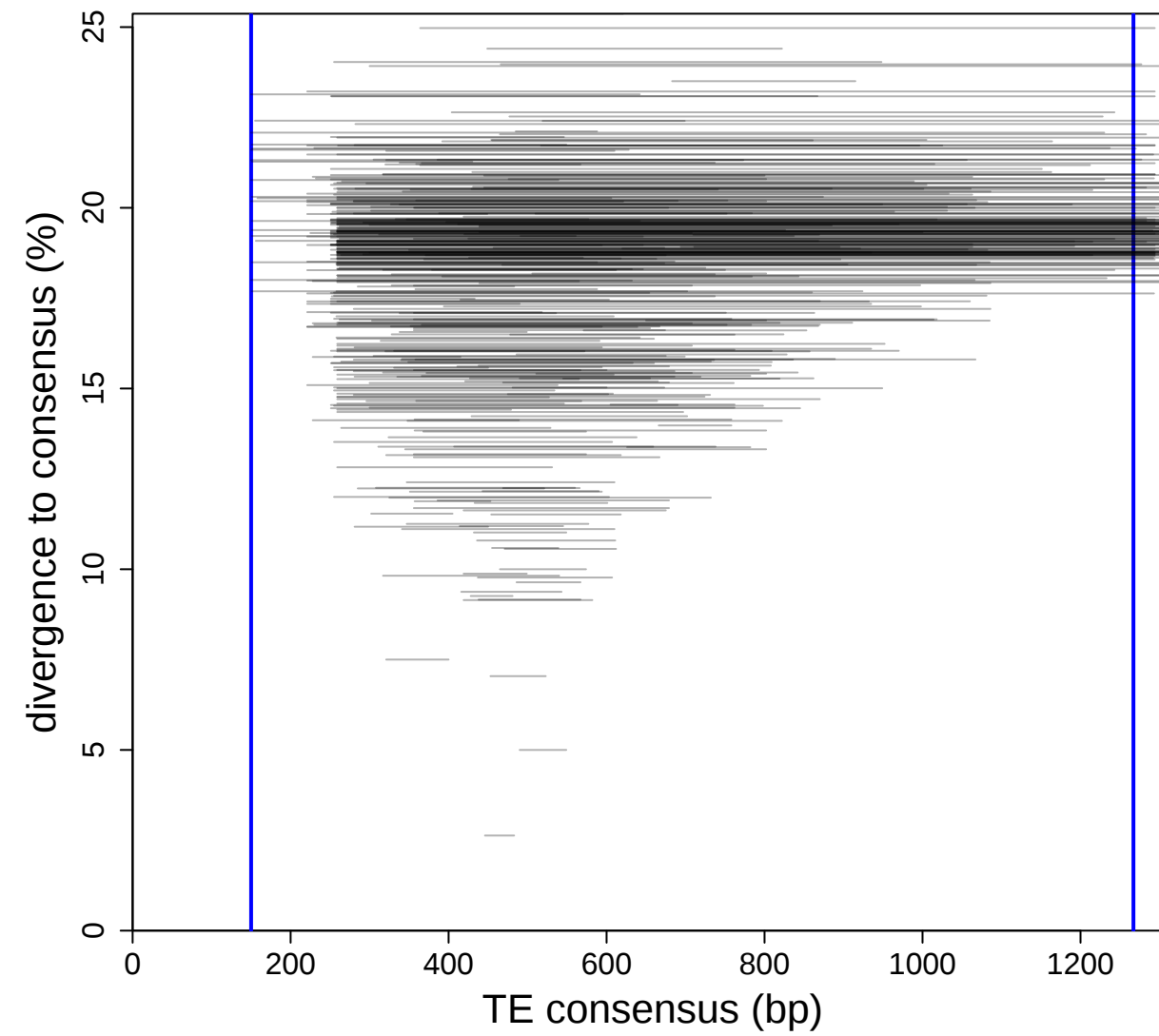
l.b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bc\nfa_aln.fa_cl.fa_gs.fa
size: 1025bp; fragments: 605; full length: 150 (>=922.5bp)

After TEtrimmer 1025 bp



2.fasta.b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa
size: 1317bp; fragments: 415; full length: 0 (>=1185.3bp)

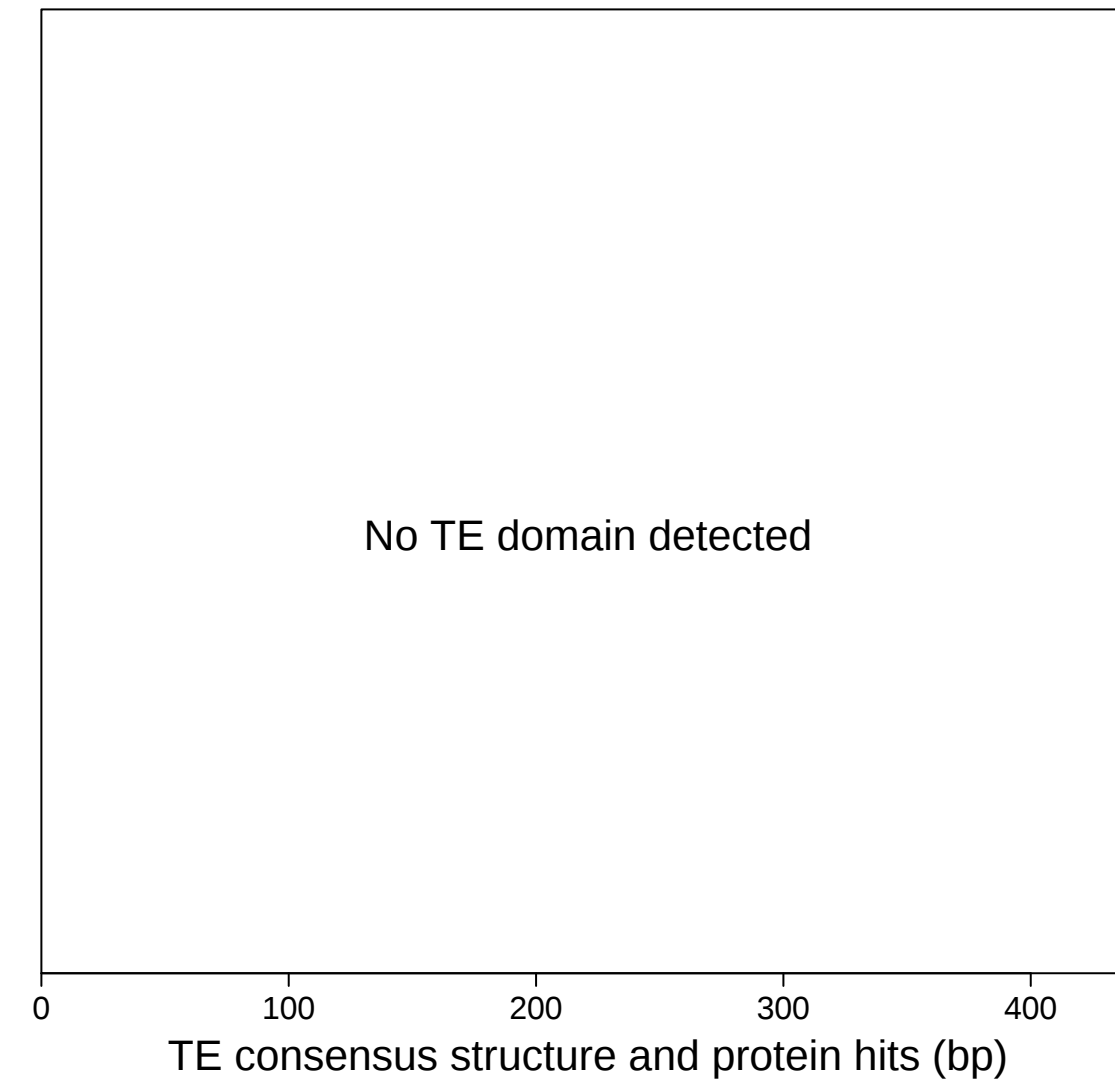
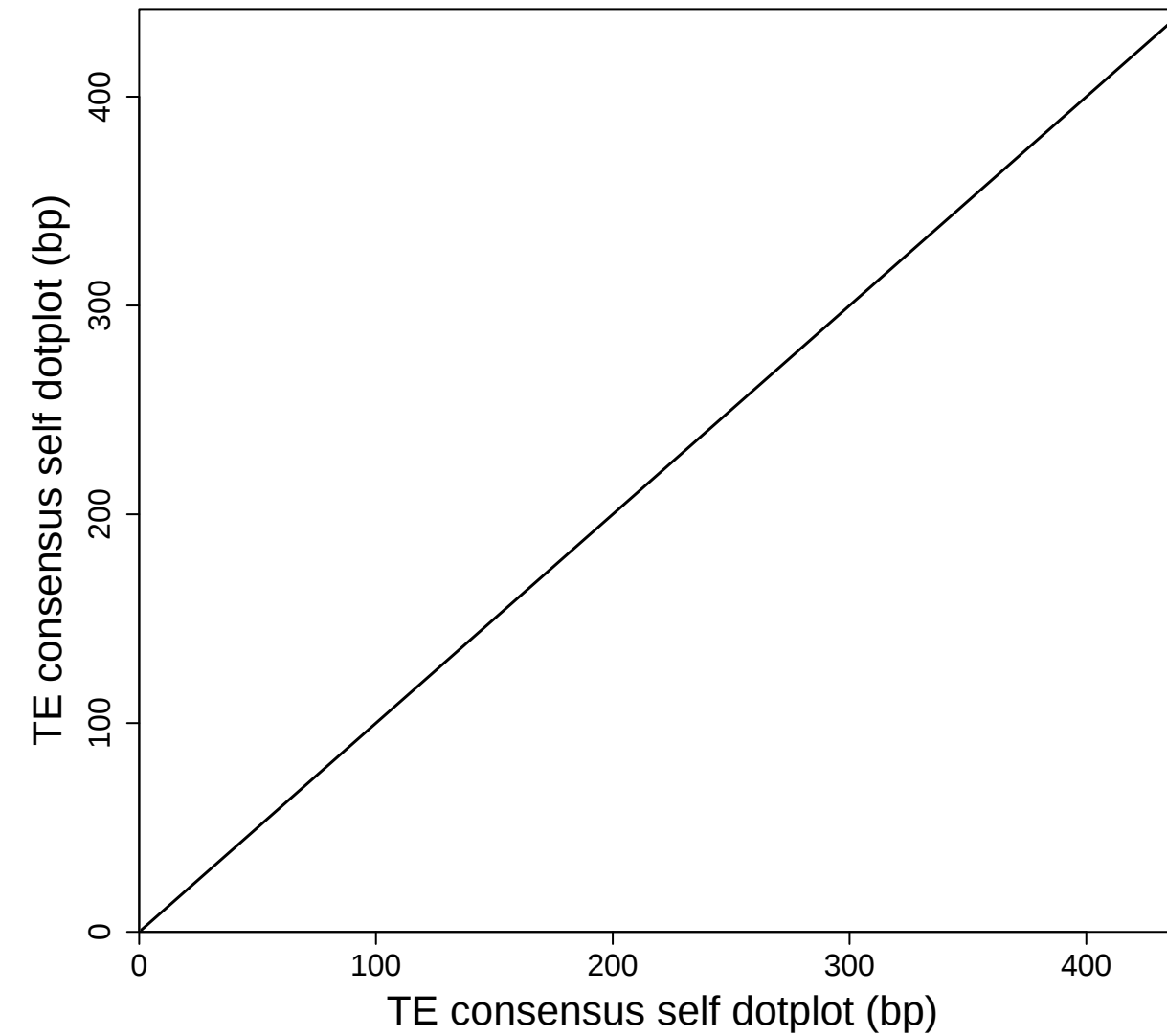
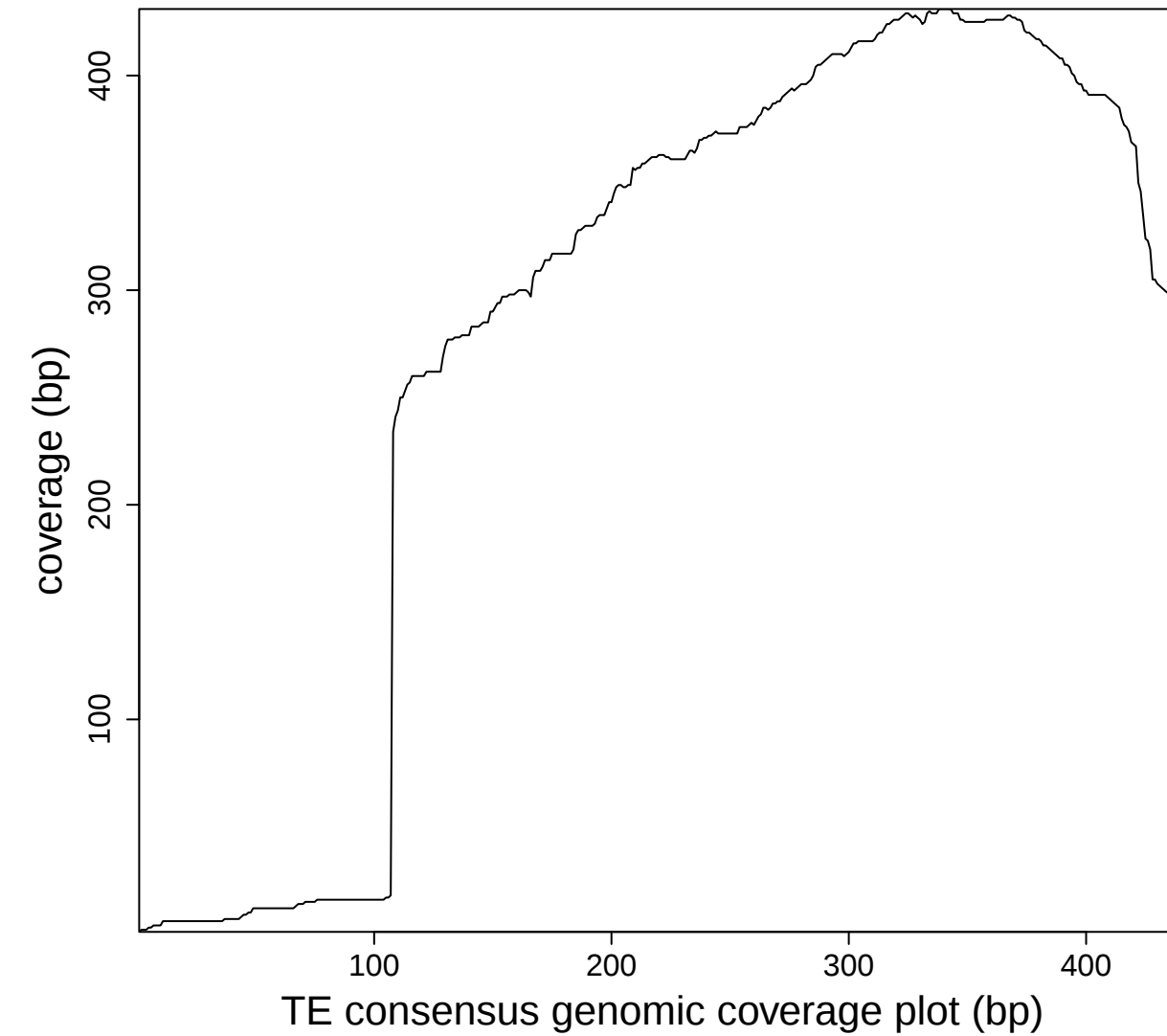
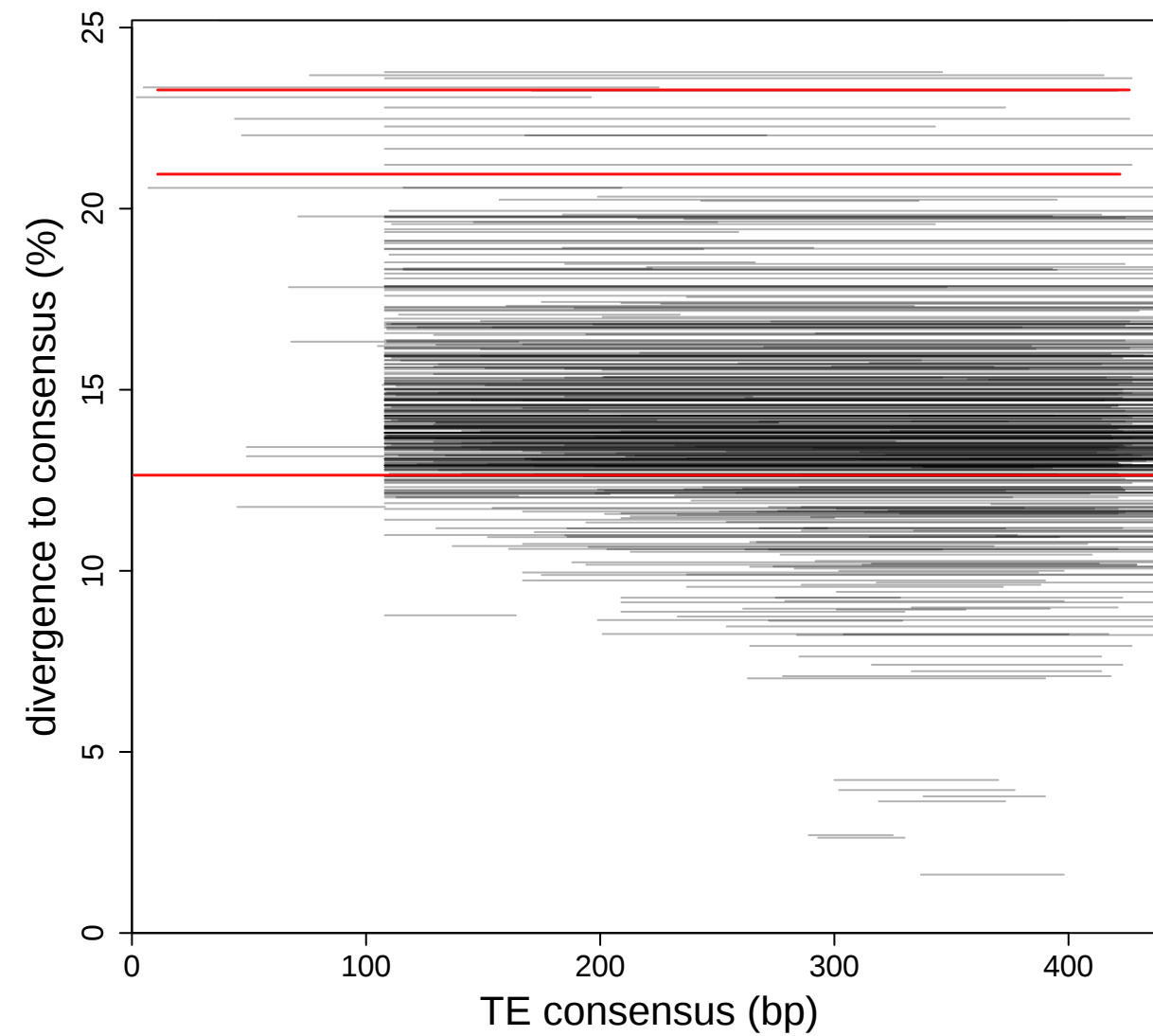
After TEtrimmer Extended plot Blue lines are boundaries



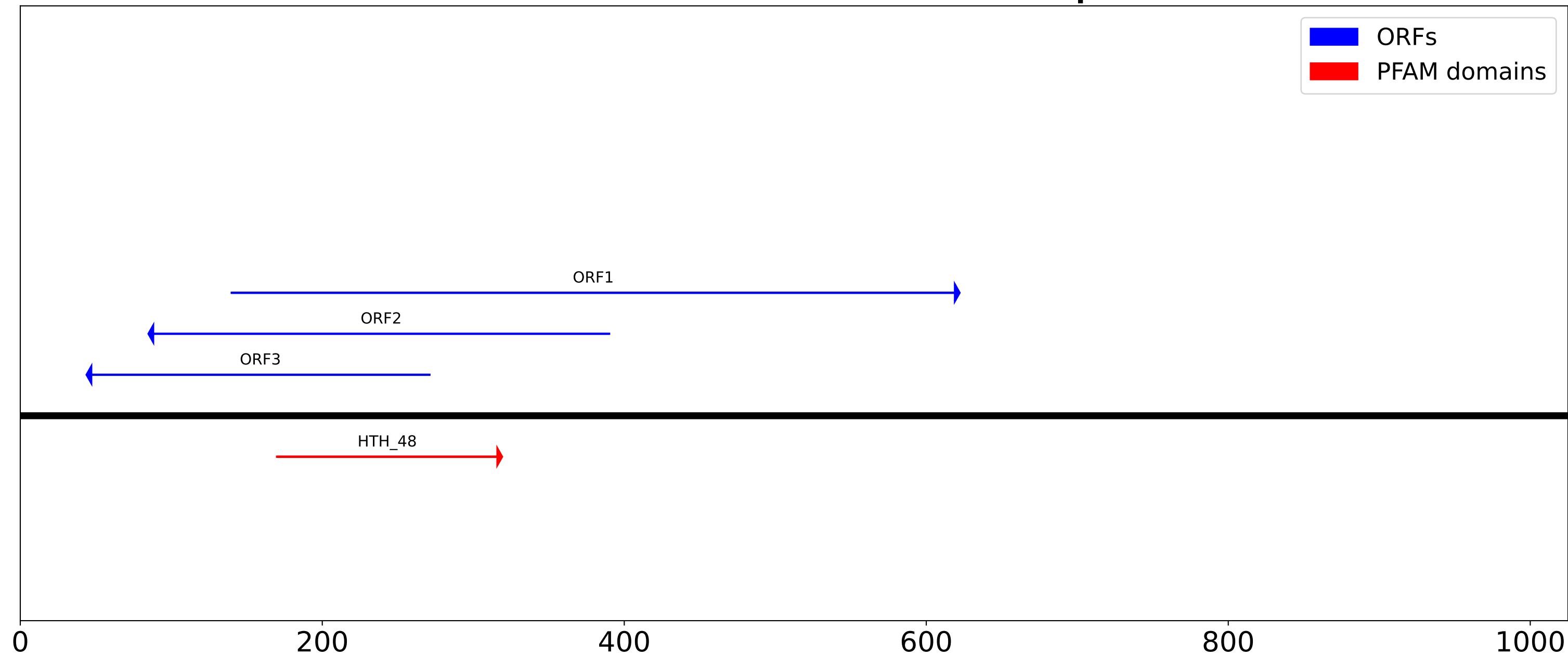
Before TEtrimmer 442 bp

TE: rnd_3_family_432

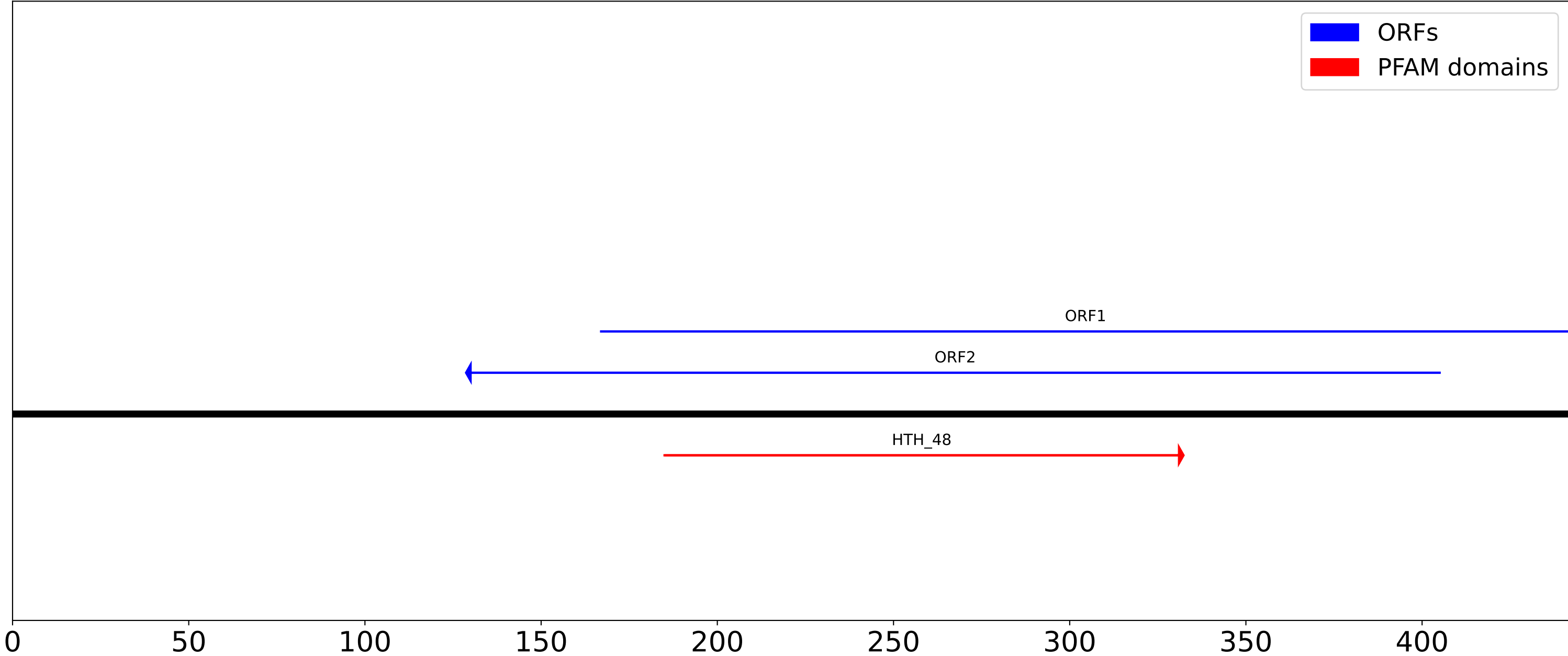
size: 442bp; fragments: 468; full length: 3 (≥ 397.8 bp)



After TEtrimmer ORF and PFAM domain plot



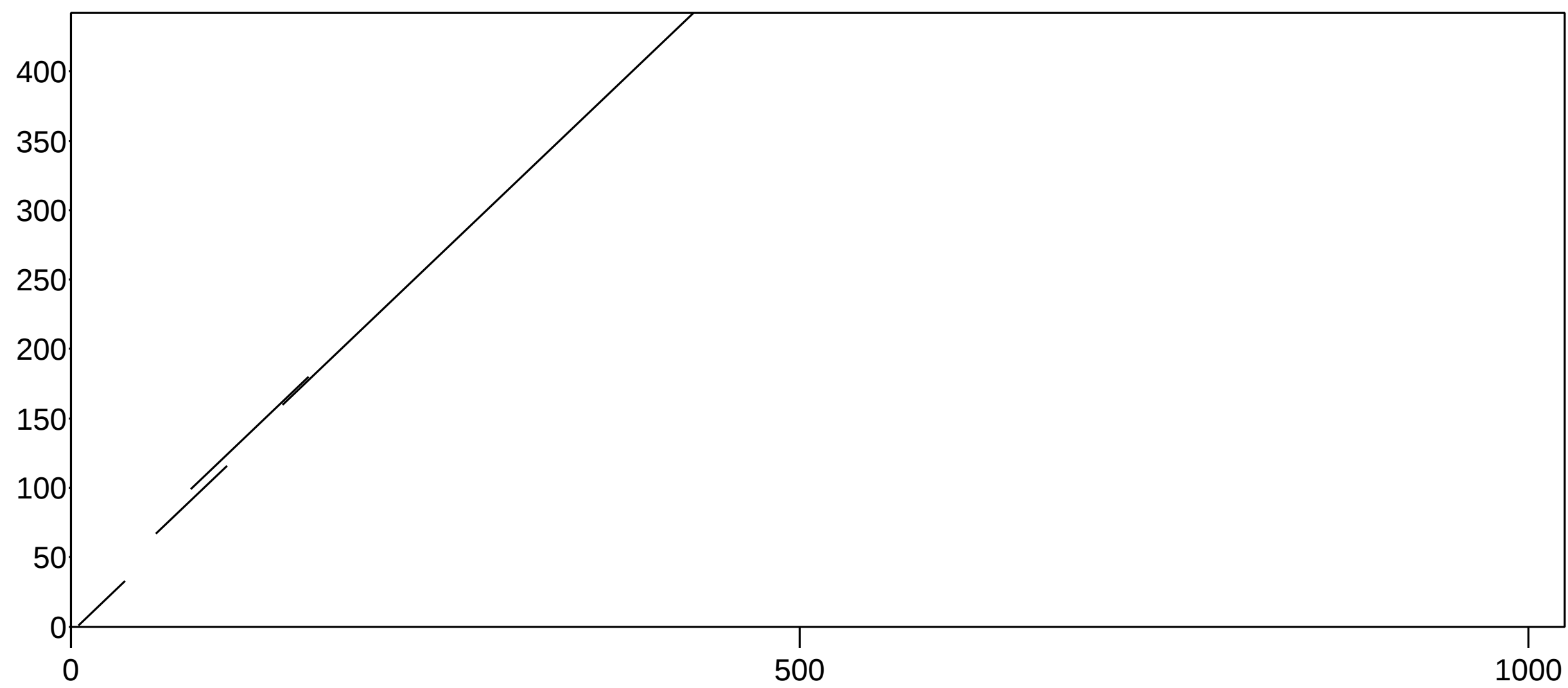
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)