

Start crop Point

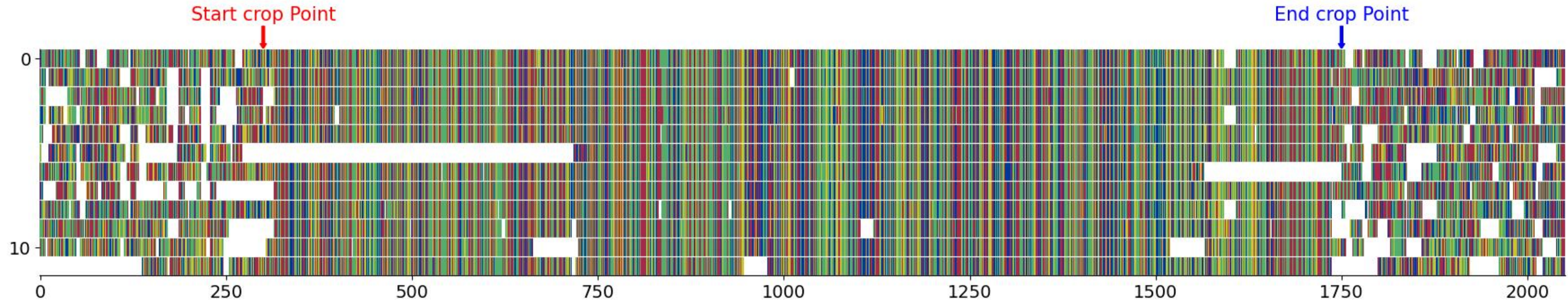
End crop Point

MSA length = 1451



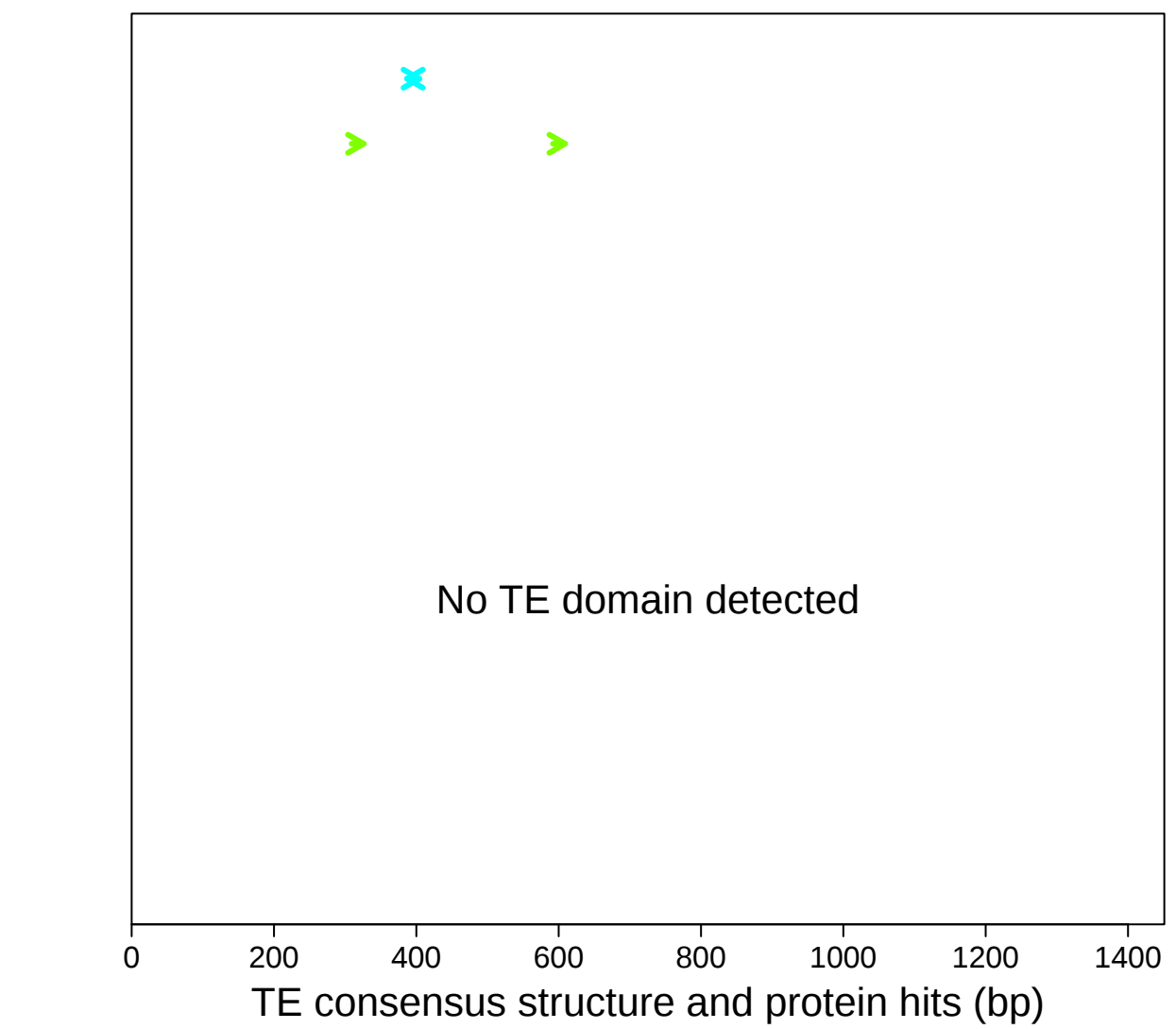
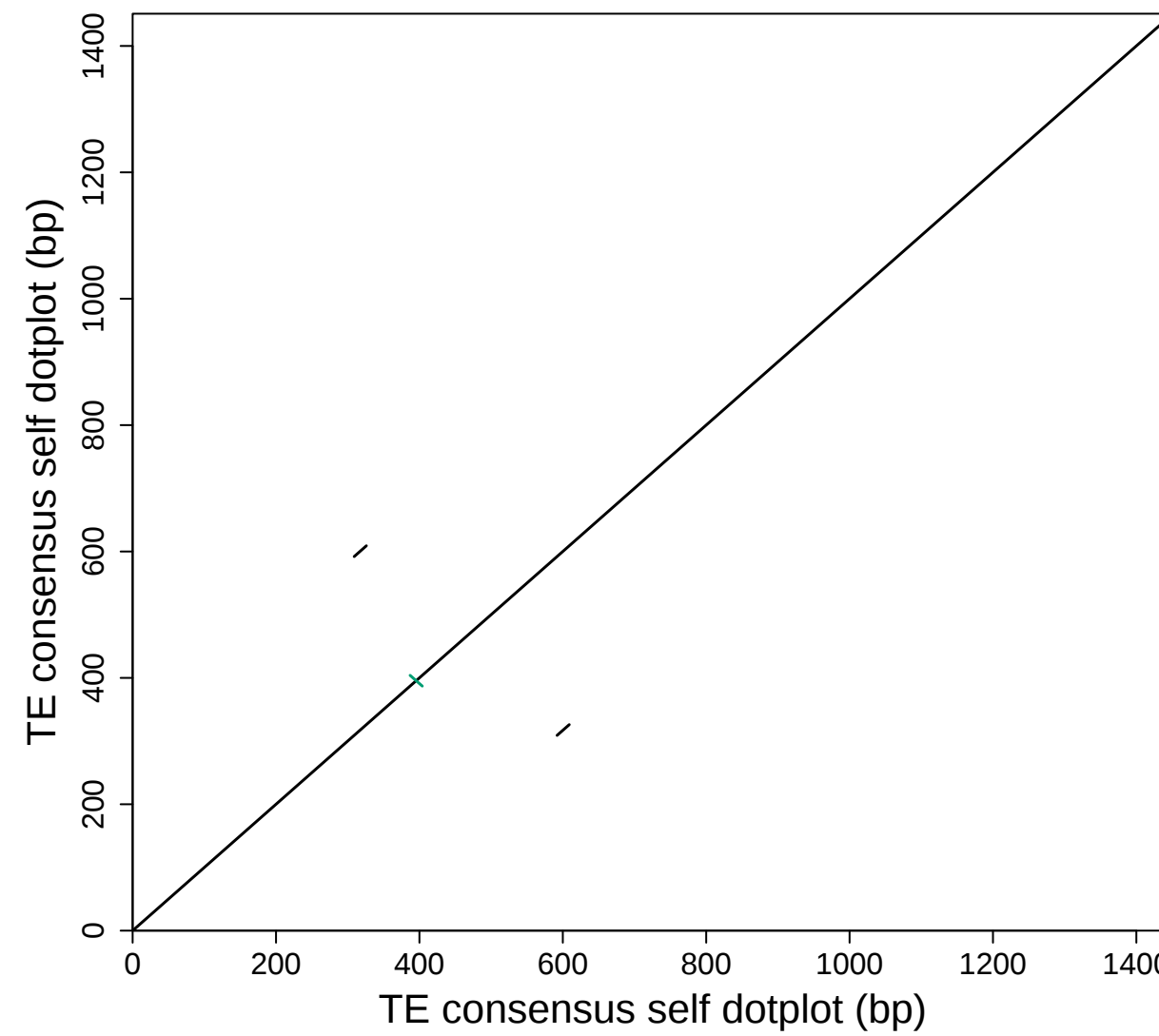
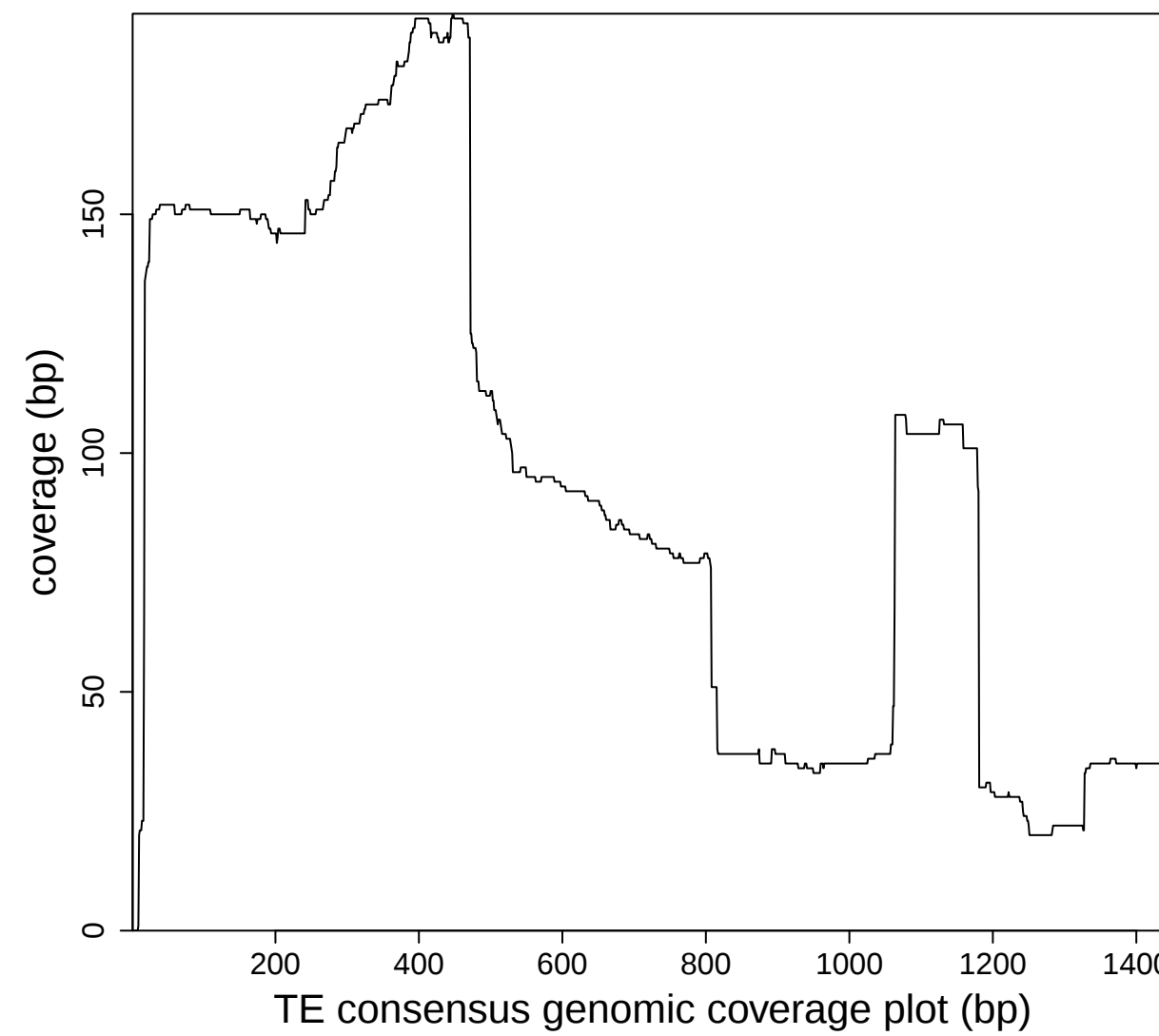
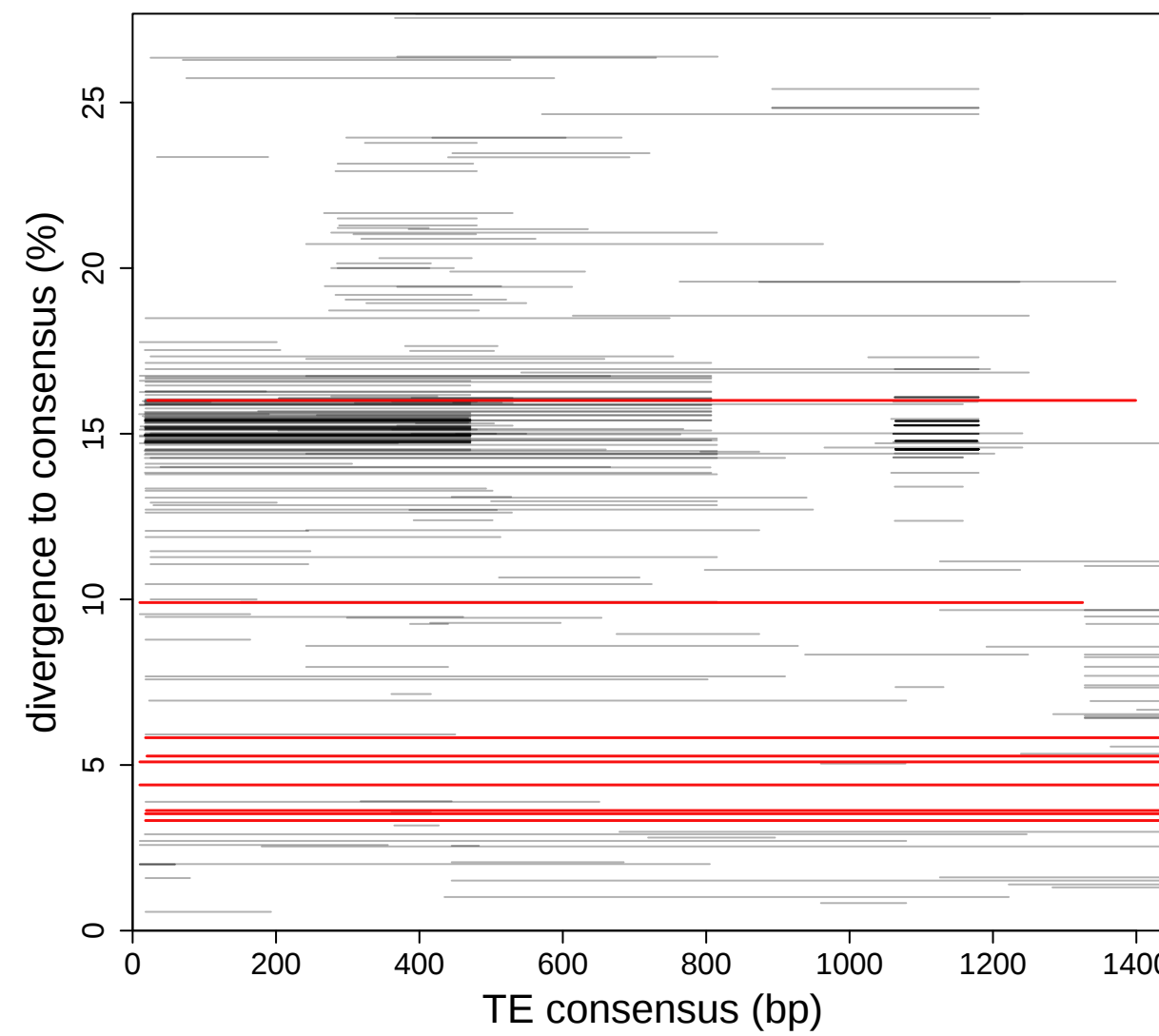
Start crop Point

End crop Point



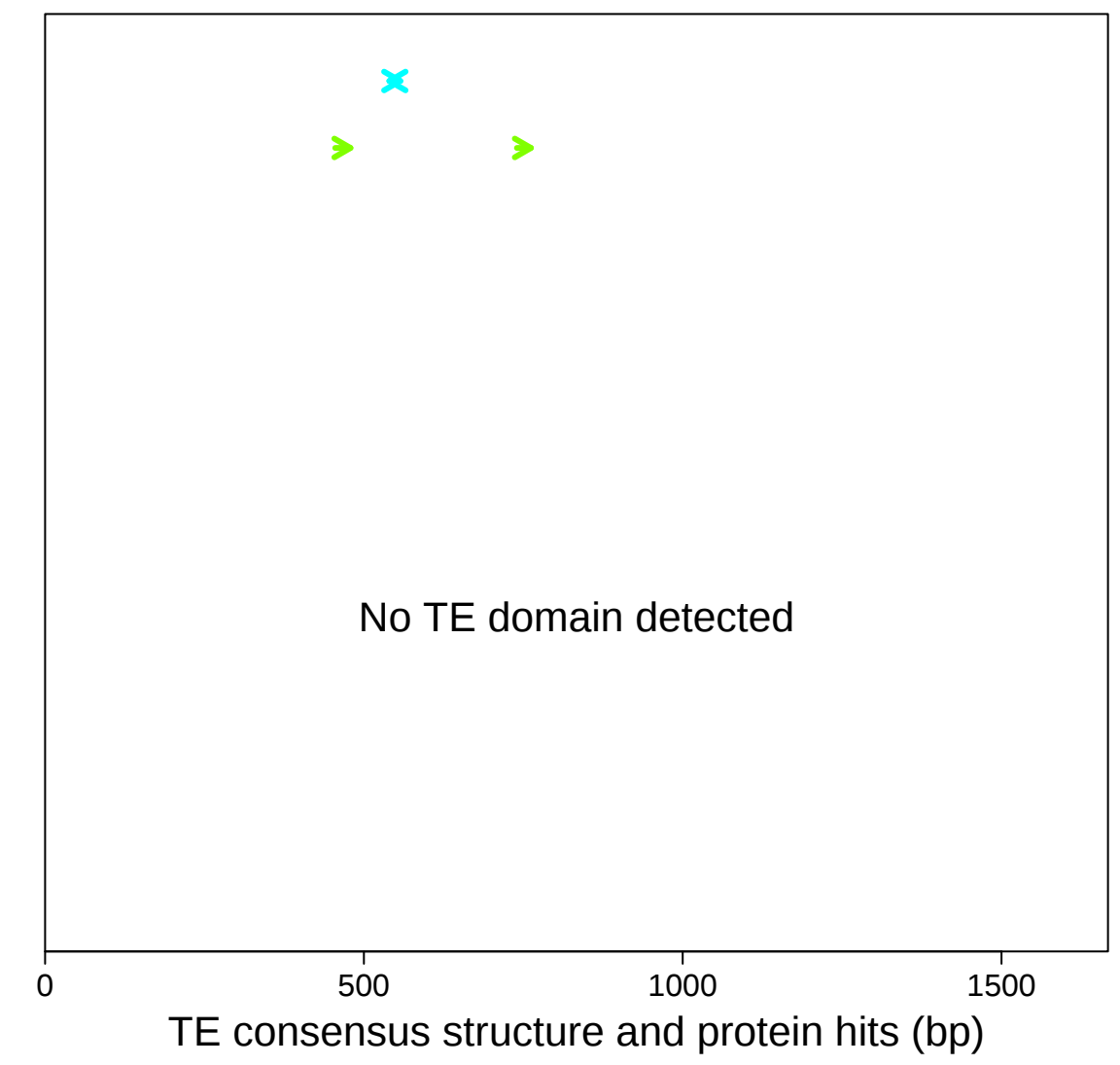
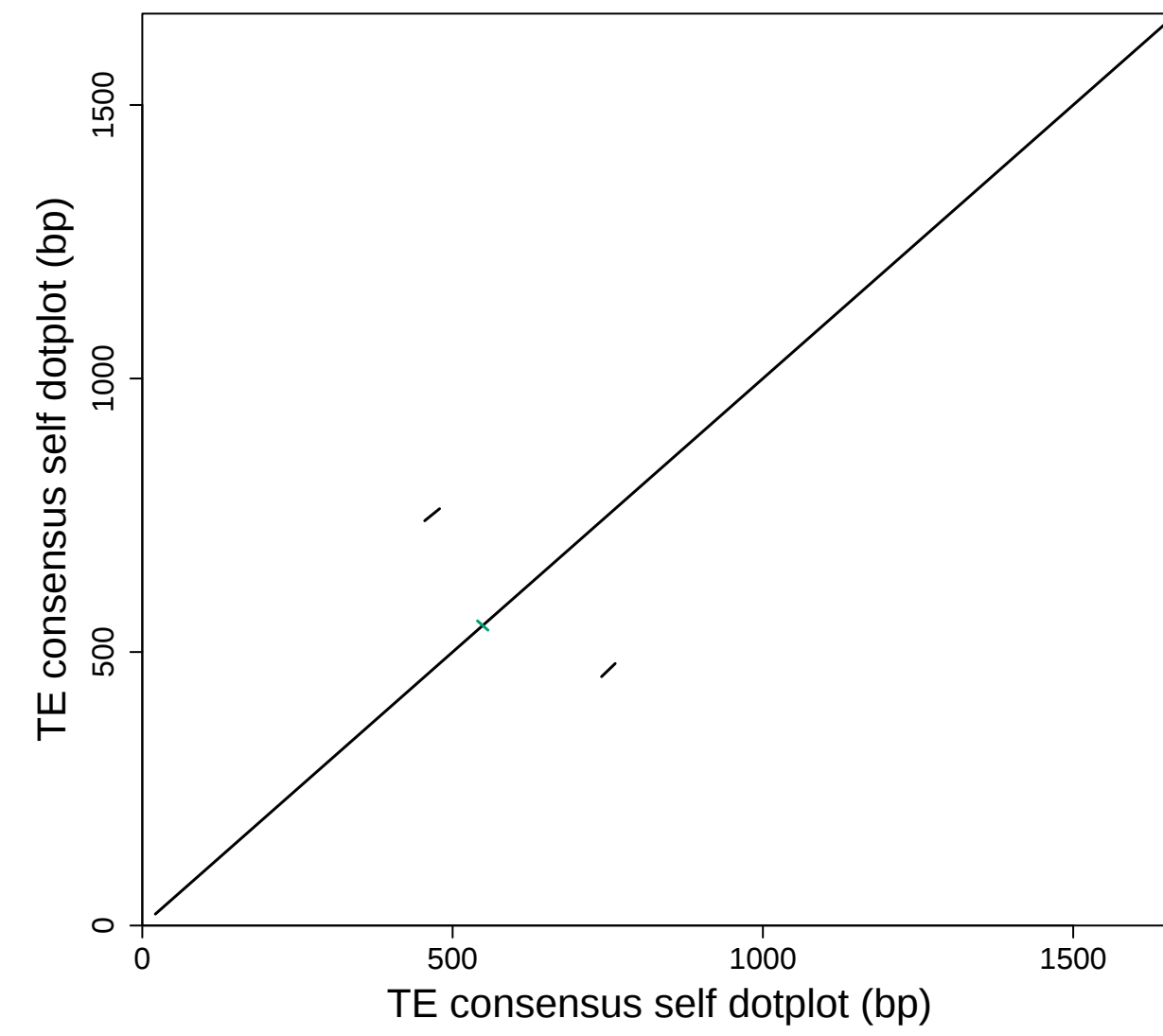
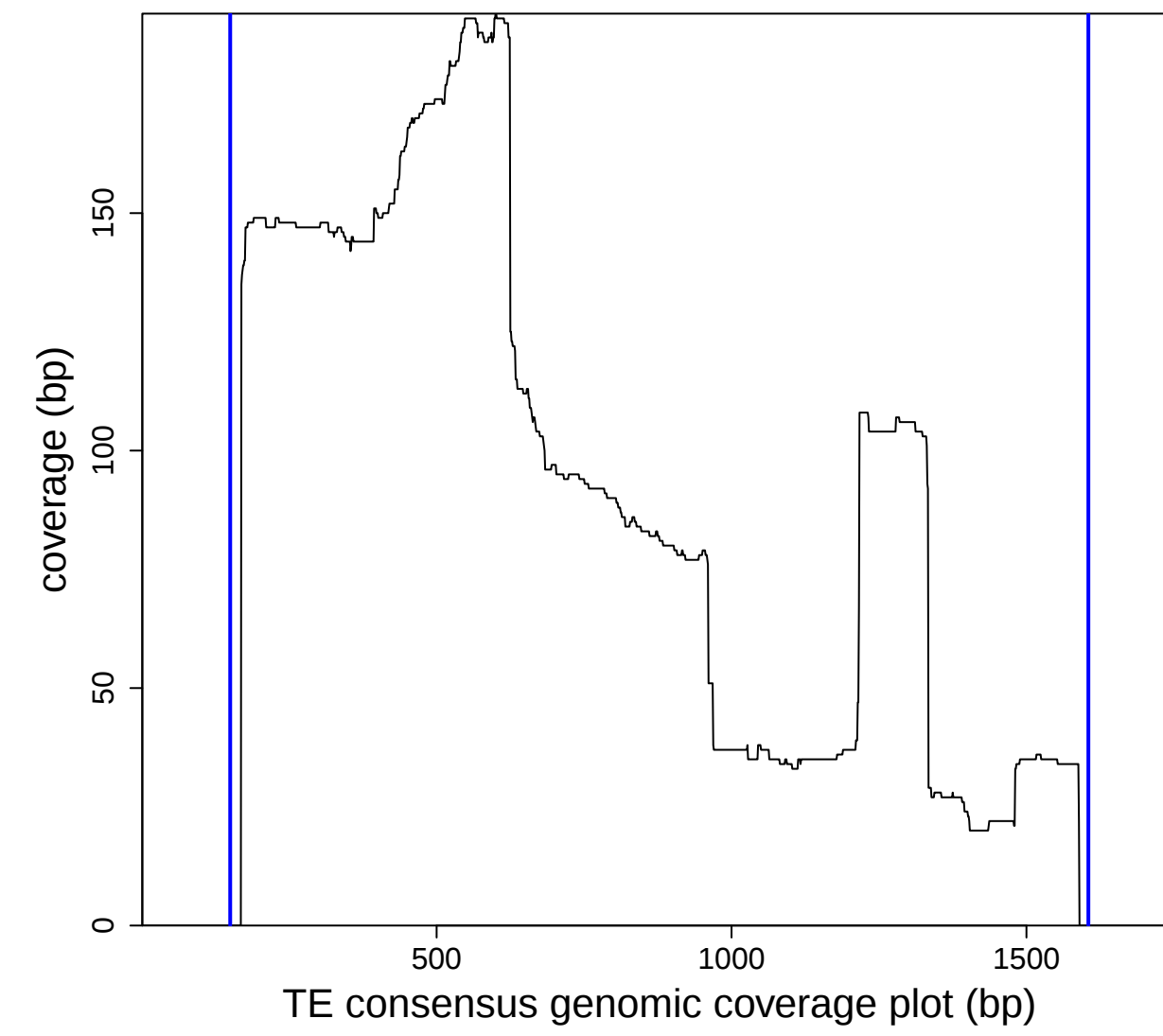
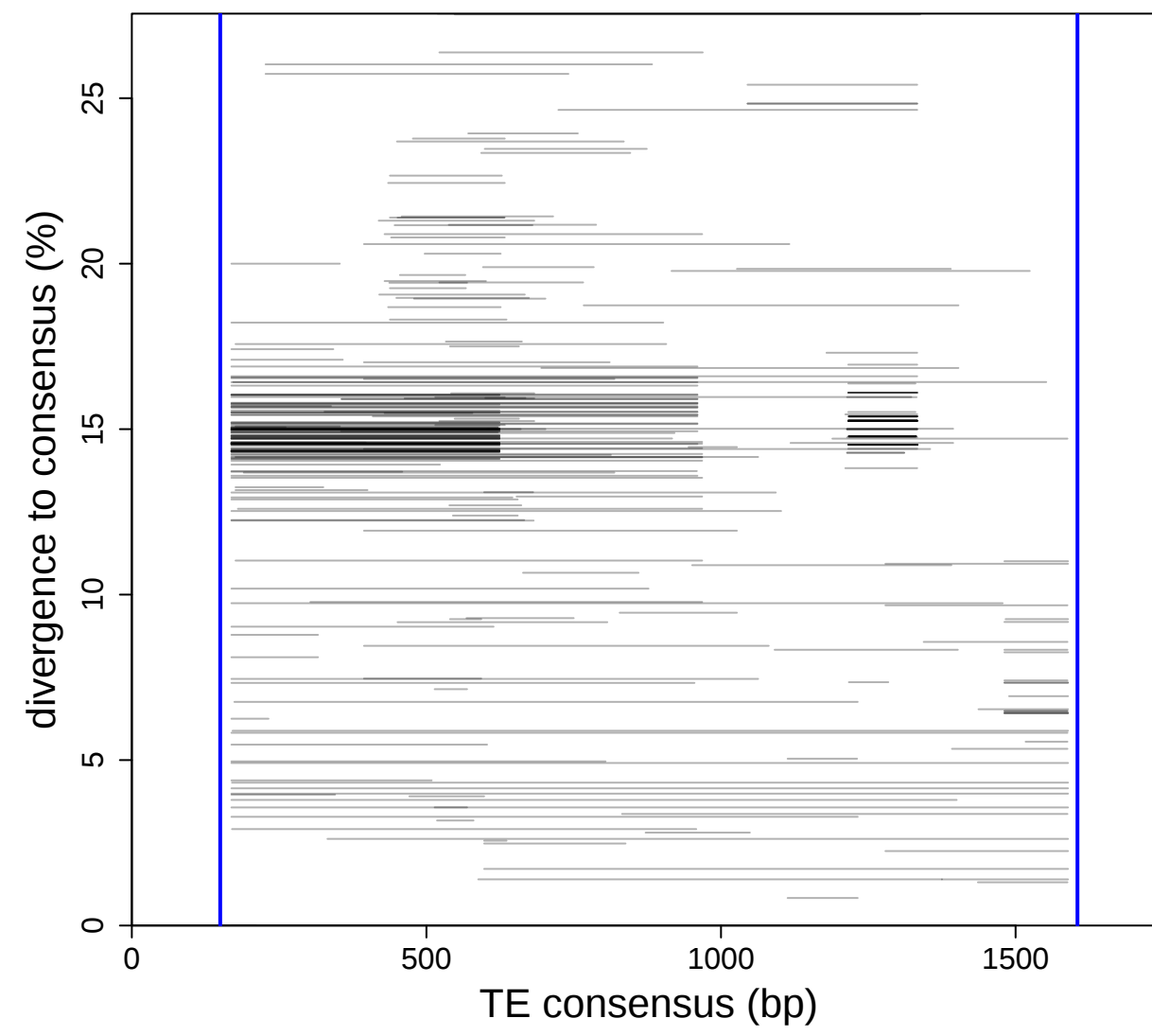
a.b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.f
size: 1451bp; fragments: 339; full length: 9 (>=1305.9bp)

After TEtrimmer 1451 bp



6.fasta.b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcIn.fa_aln.fa_cl.f
size: 1755bp; fragments: 336; full length: 0 (>=1579.5bp)

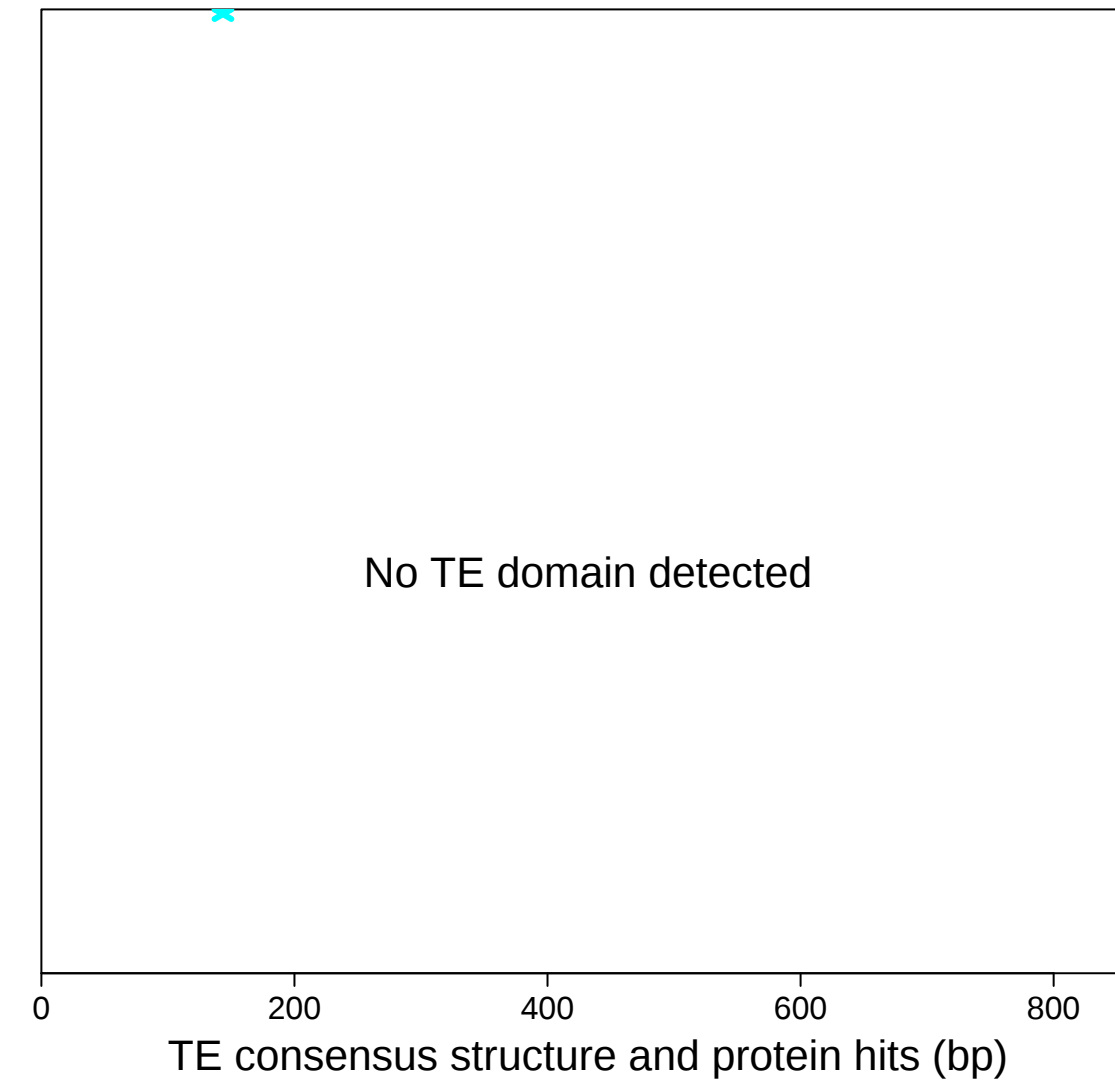
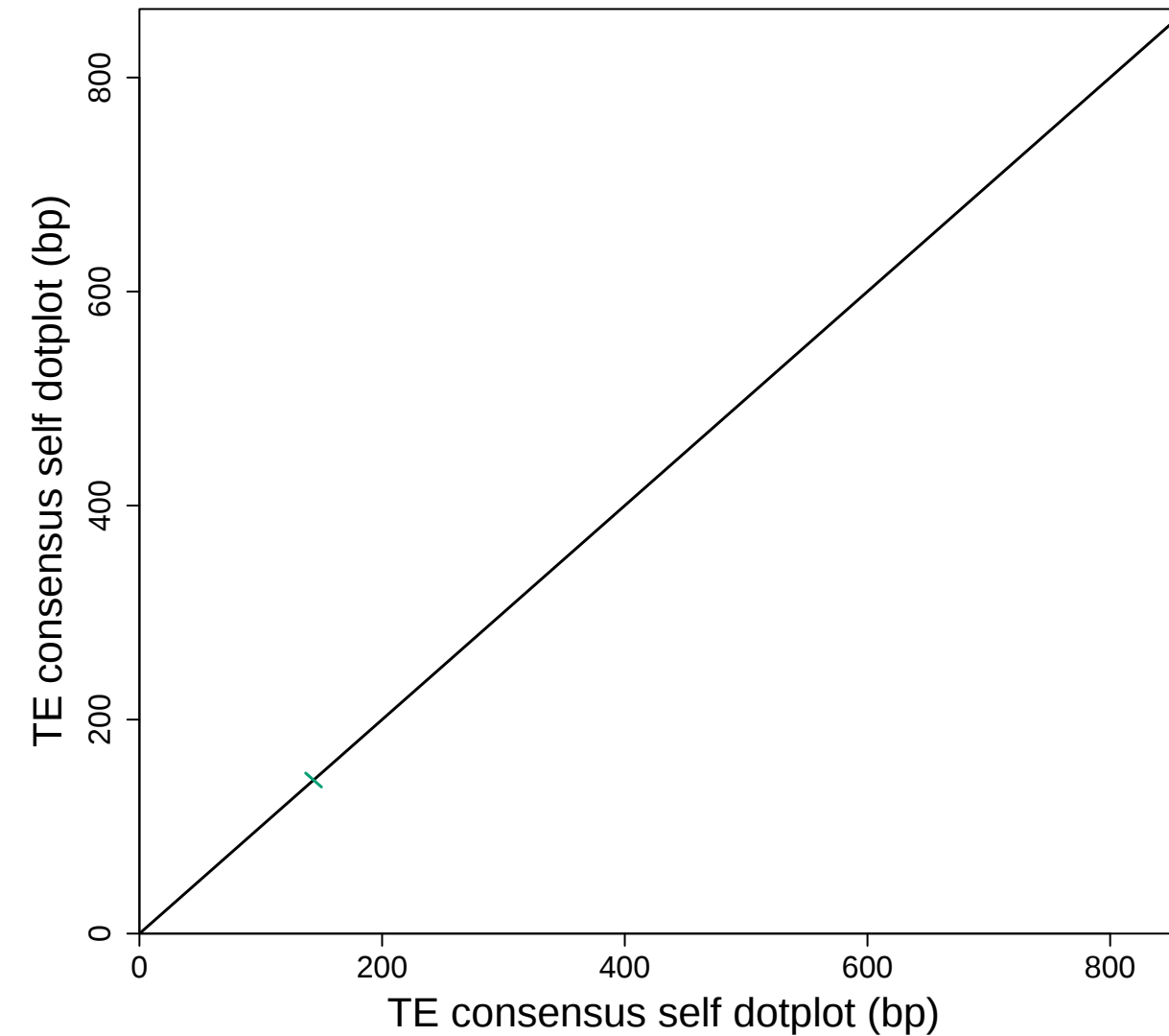
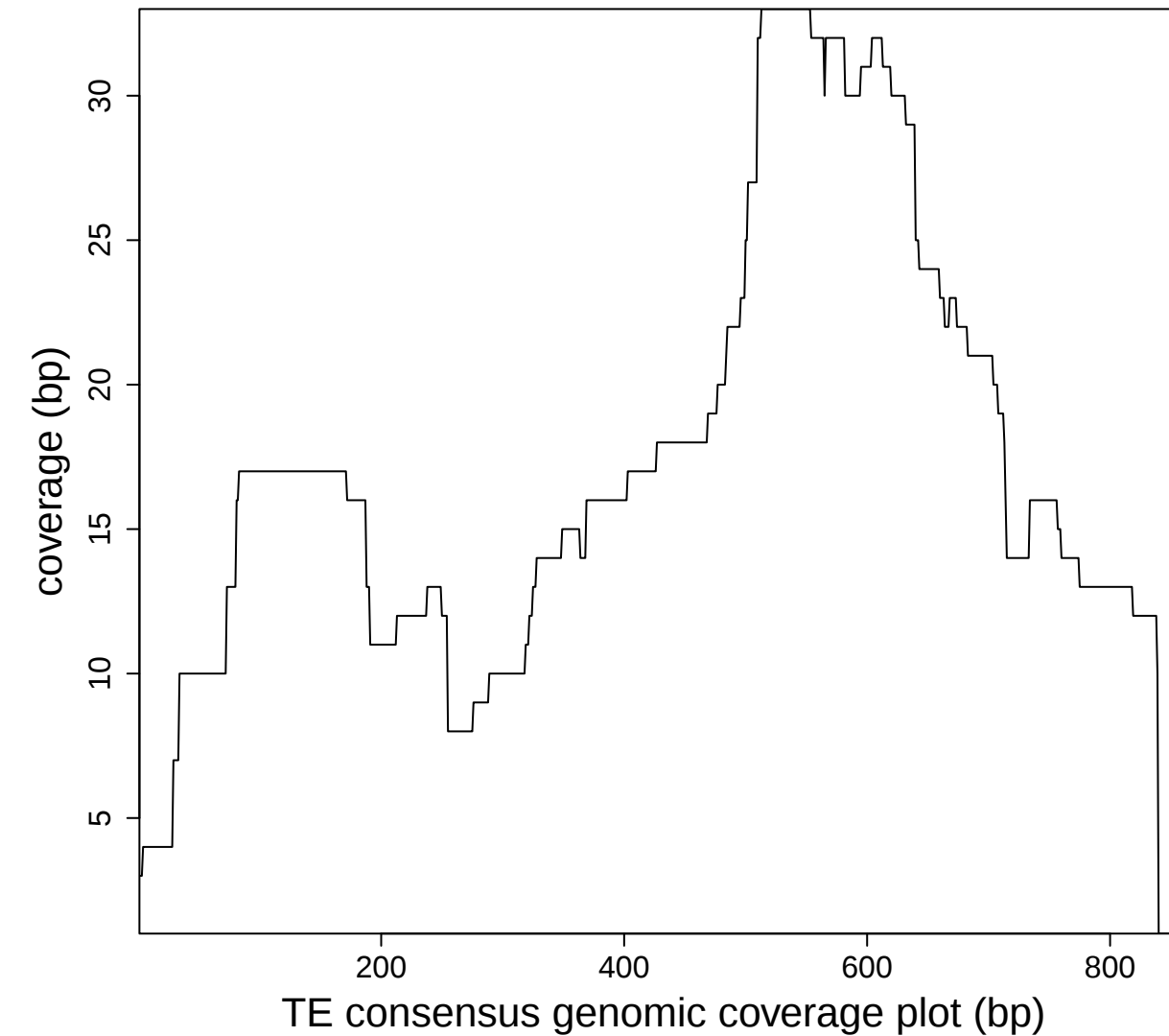
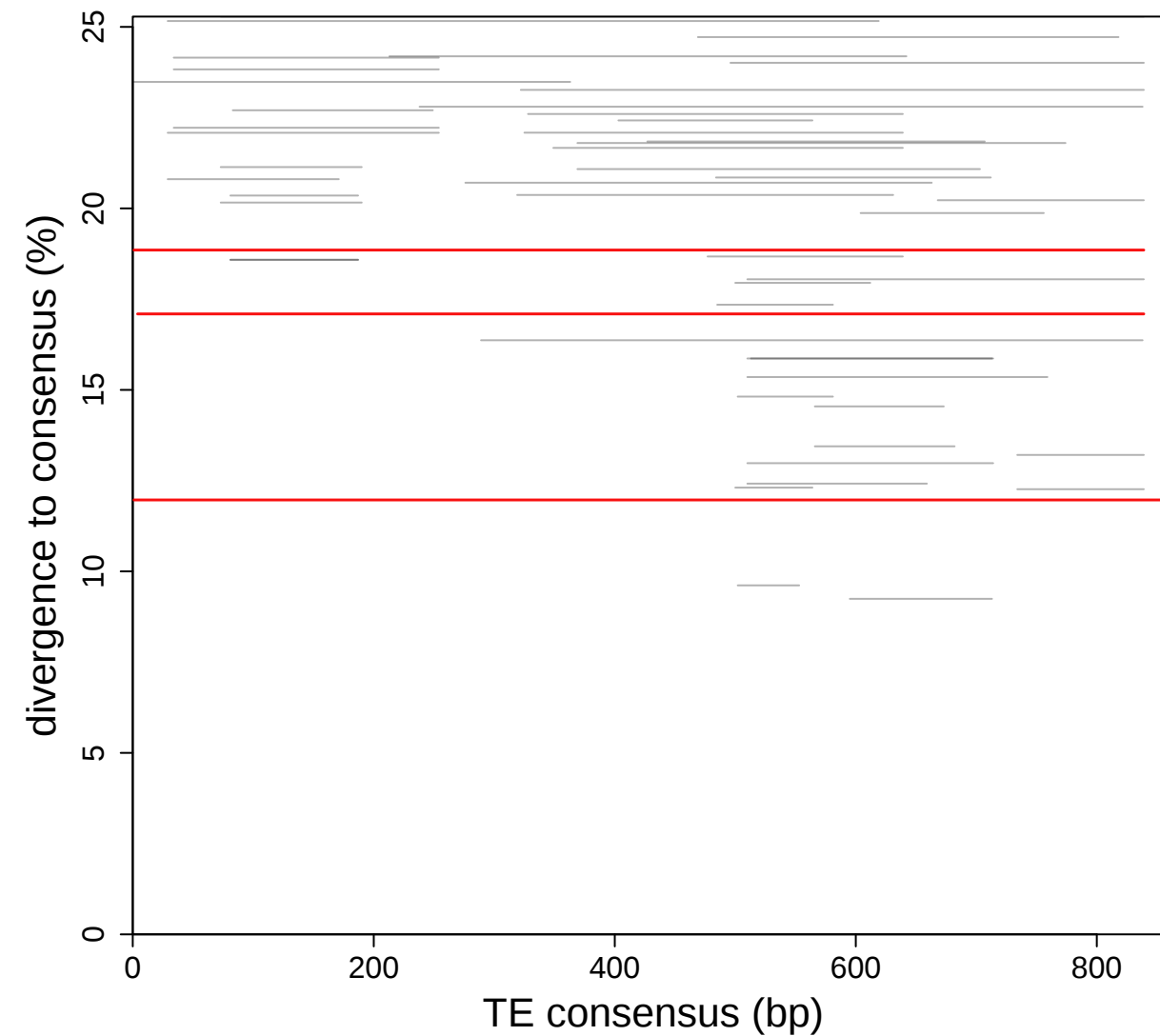
After TEtrimmer Extended plot Blue lines are boundaries



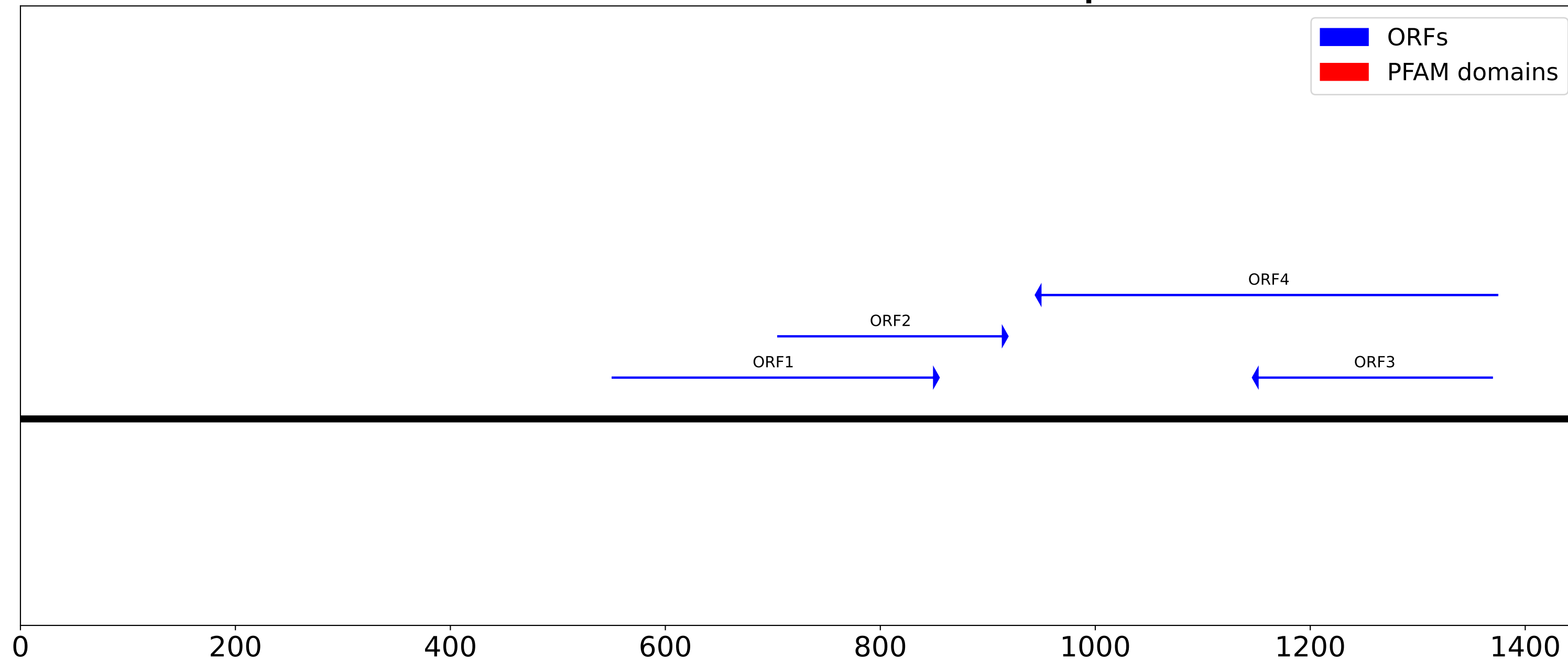
Before TEtrimmer 864 bp

TE: rnd_4_family_2806

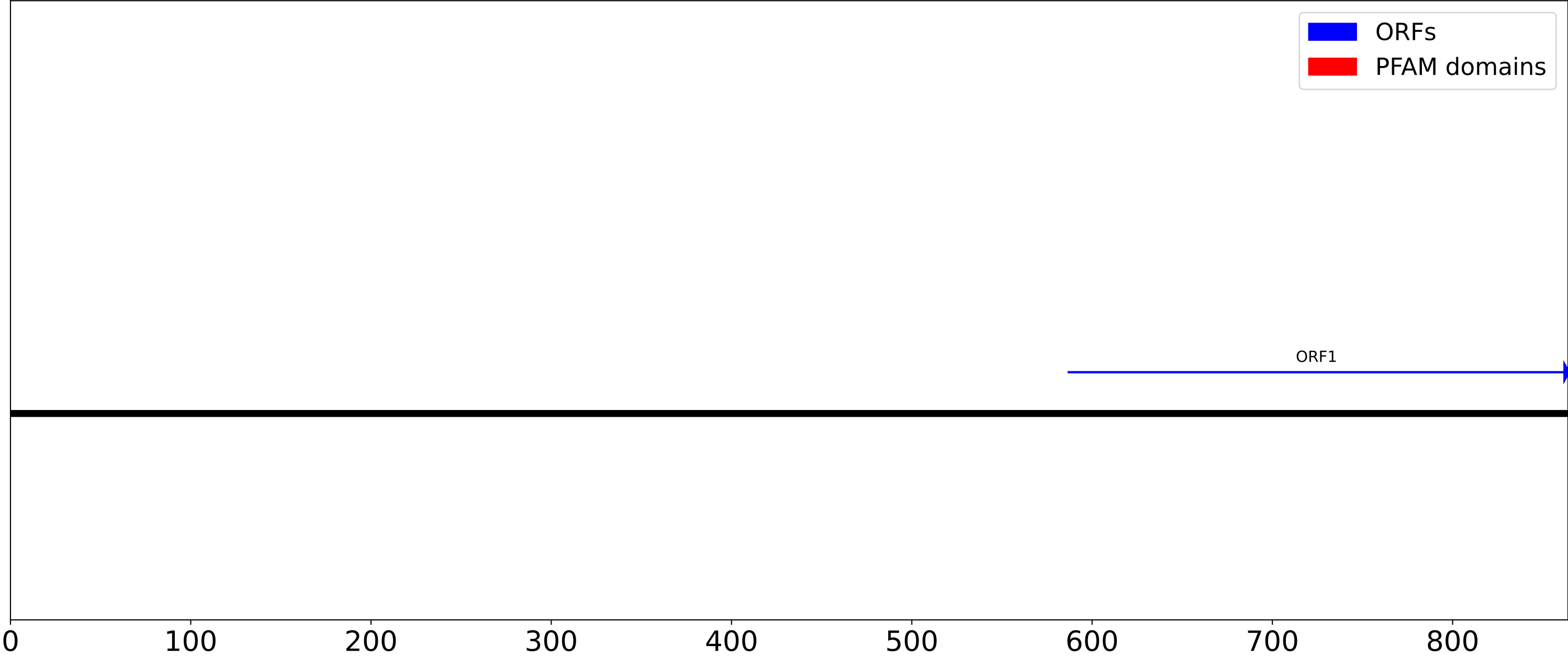
size: 864bp; fragments: 52; full length: 3 (≥ 777.6 bp)



After TEtrimmer ORF and PFAM domain plot



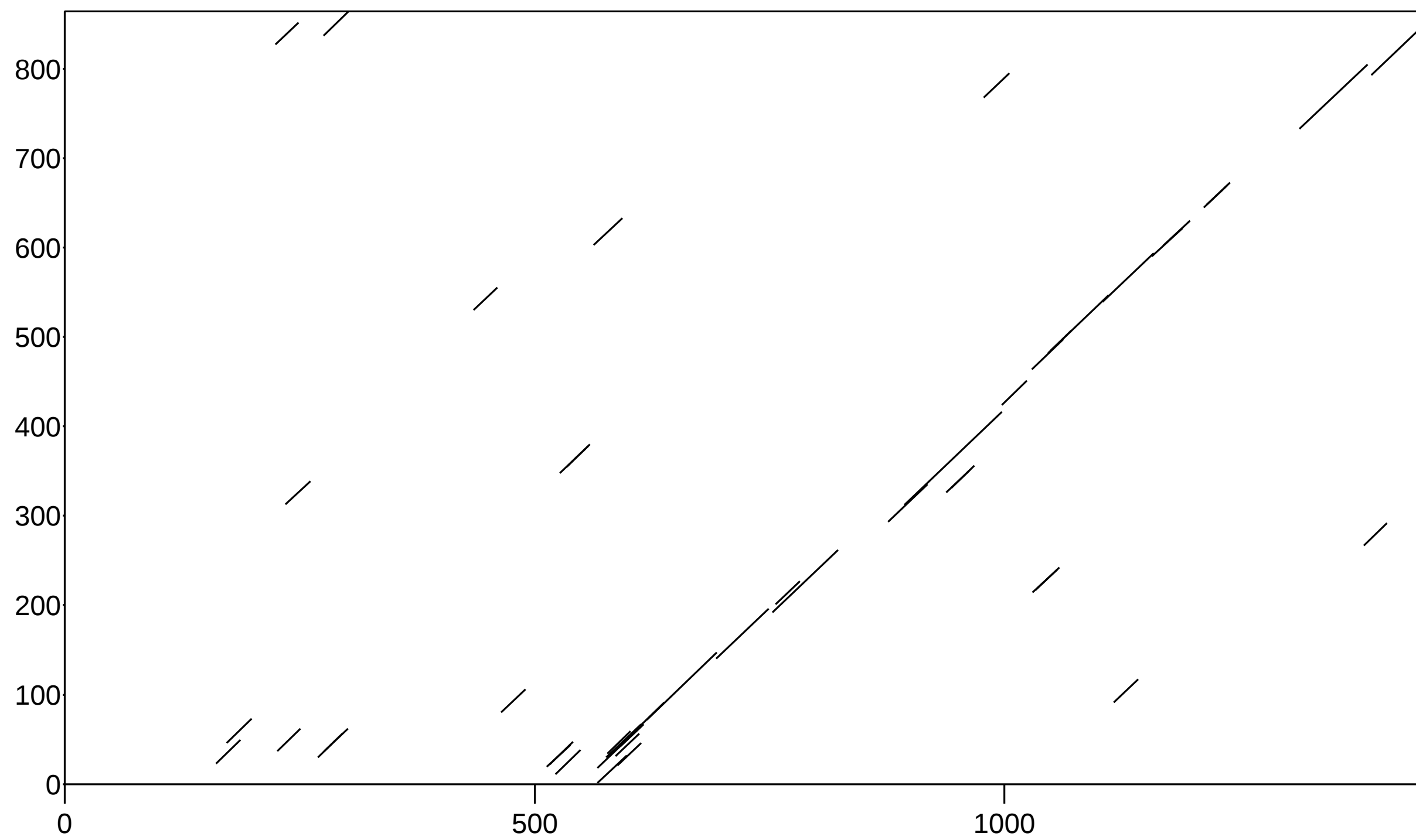
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)