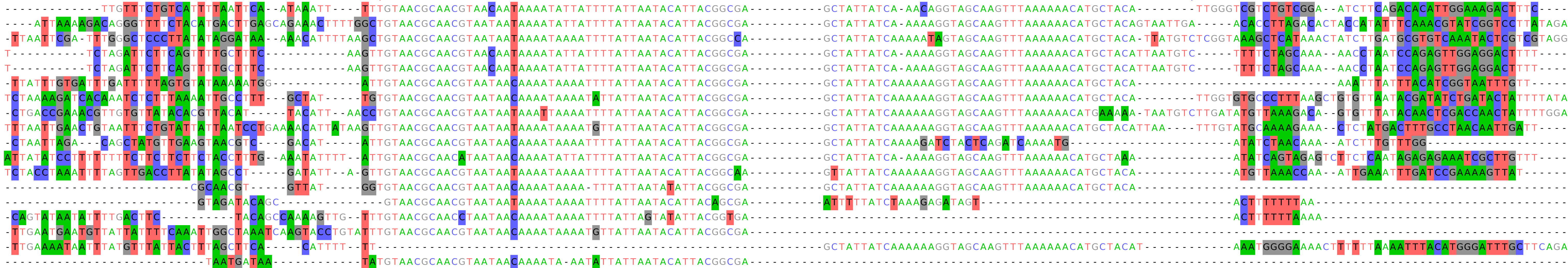


MSA length = 1450

Start crop Point

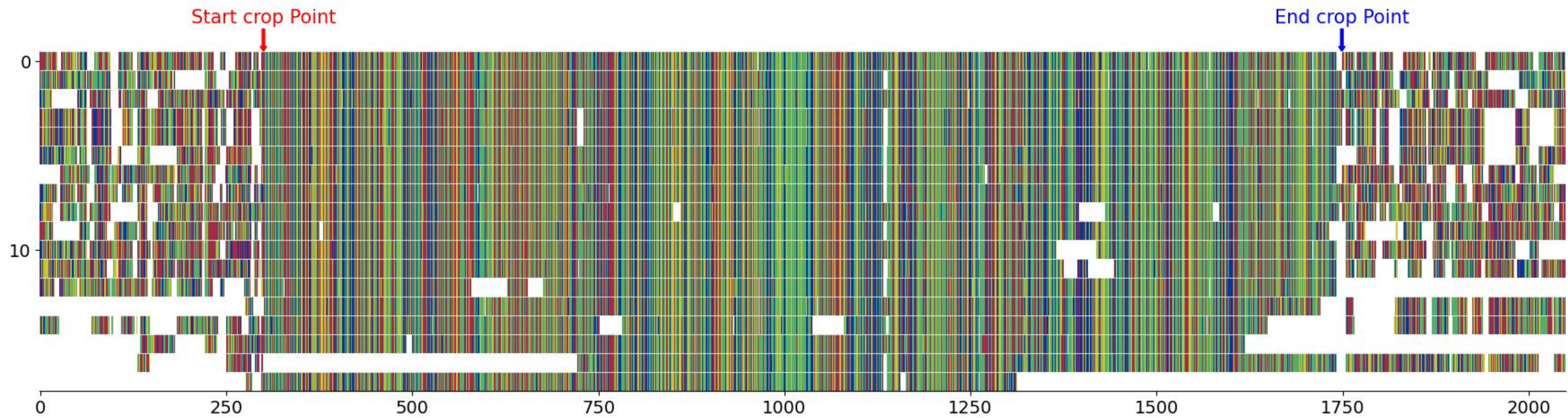


End crop Point



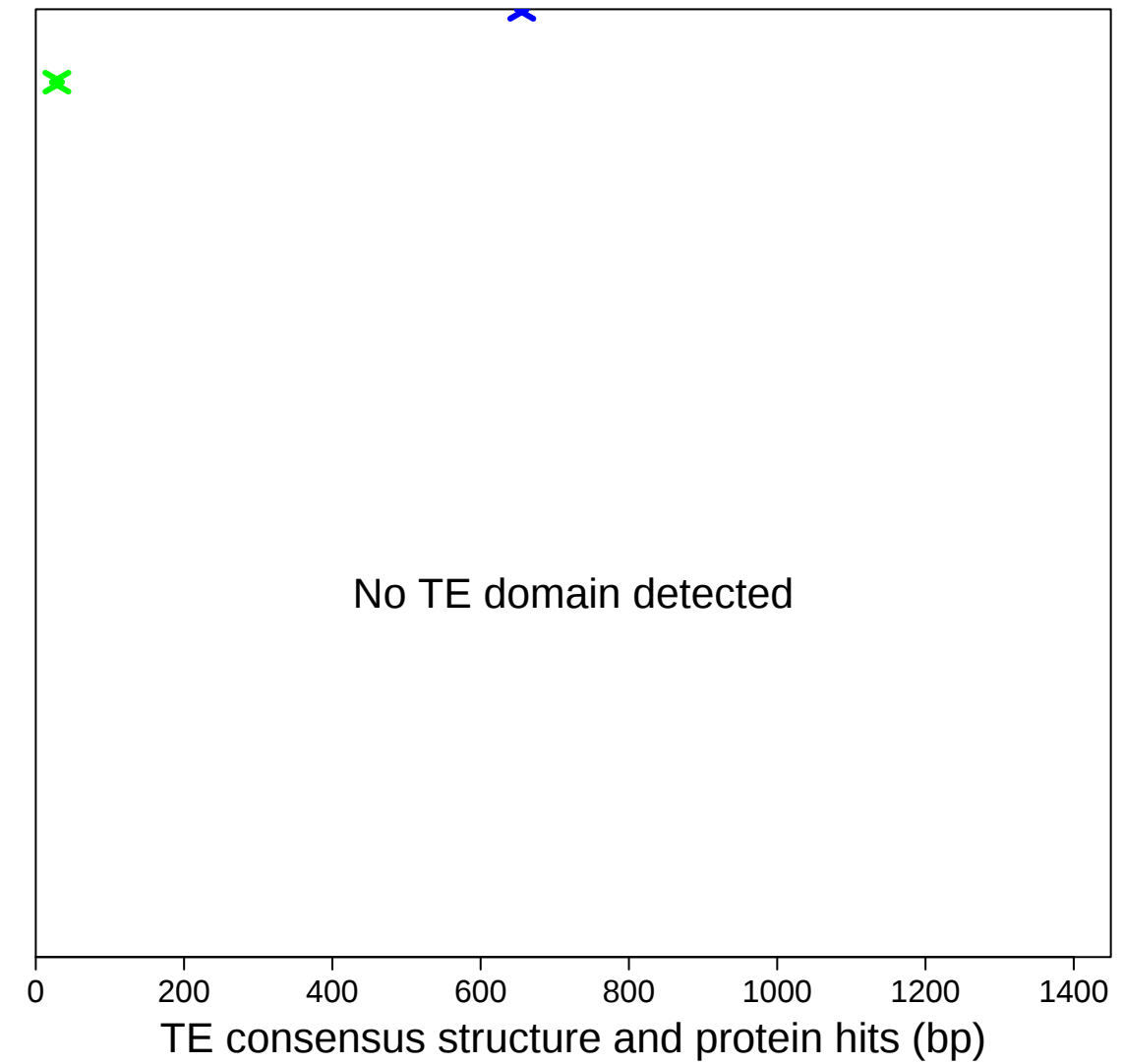
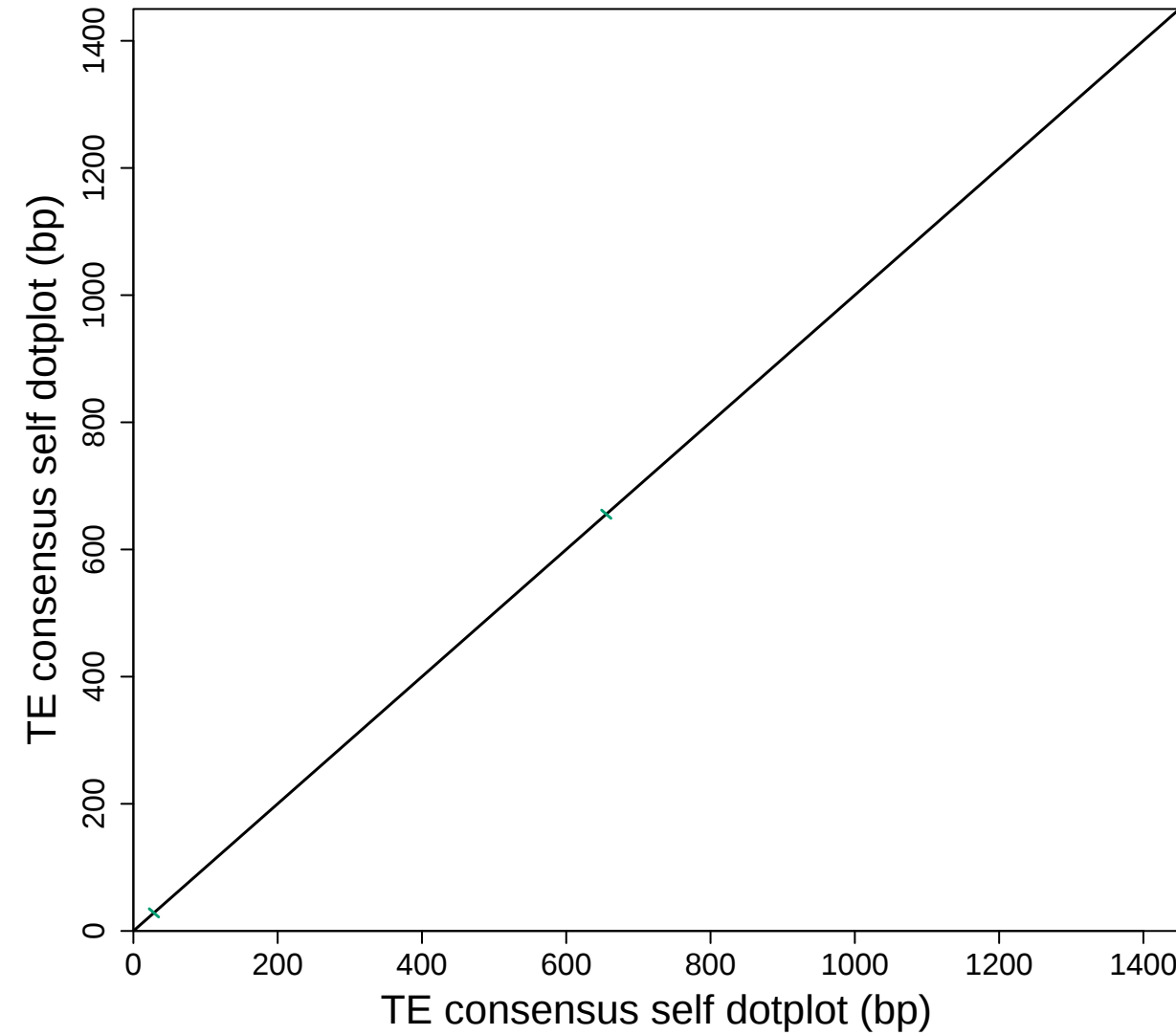
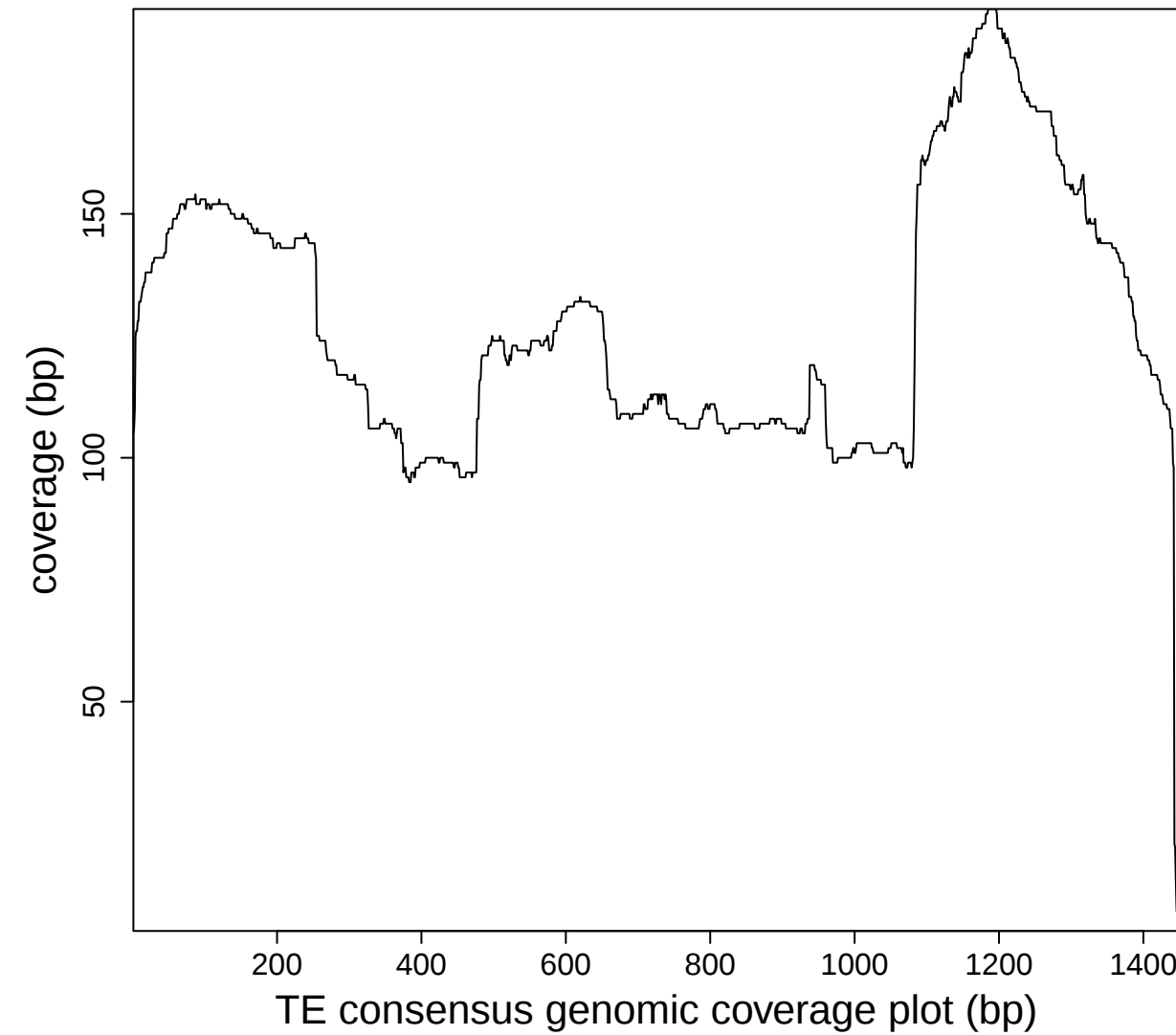
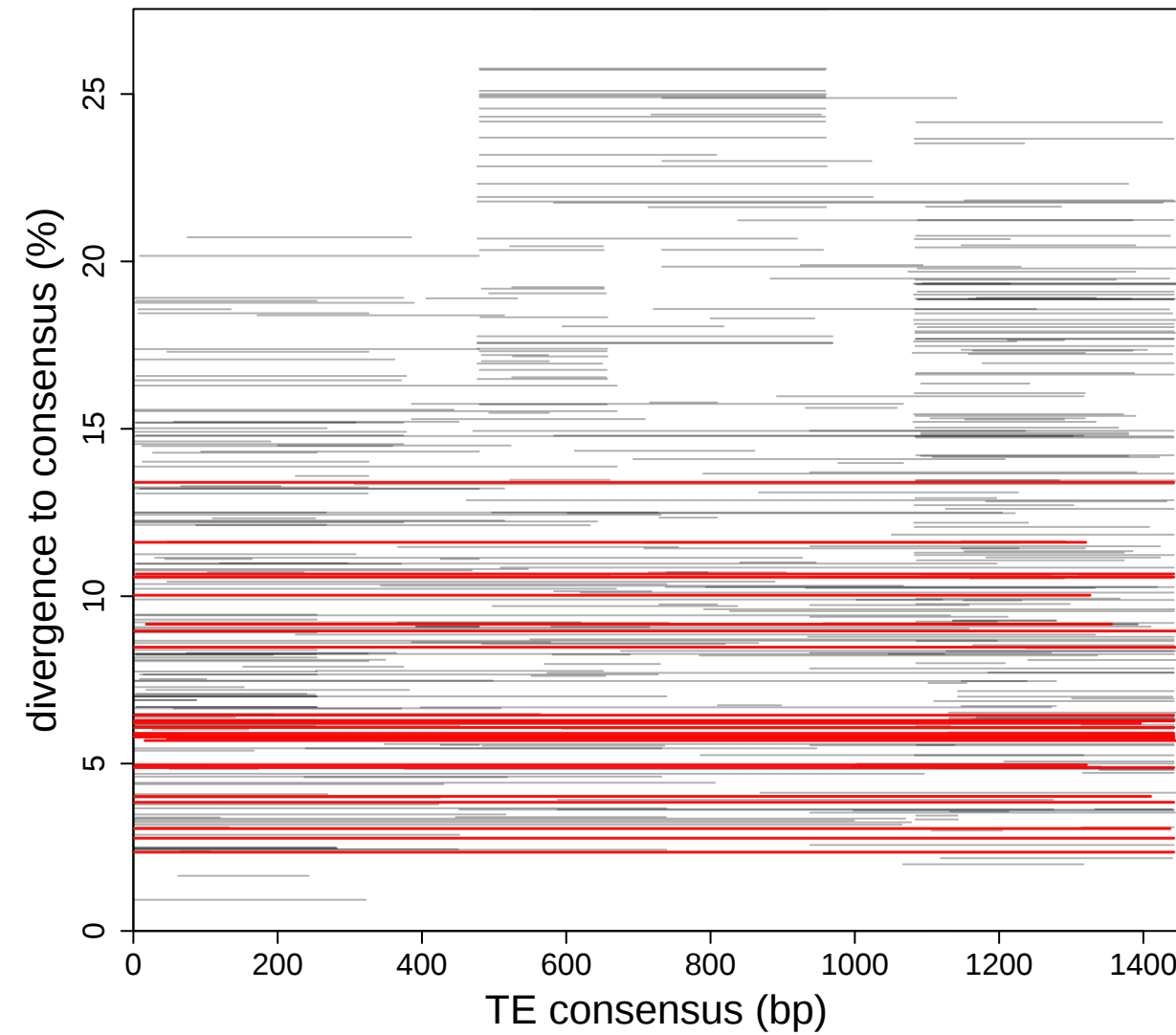
Start crop Point

End crop Point



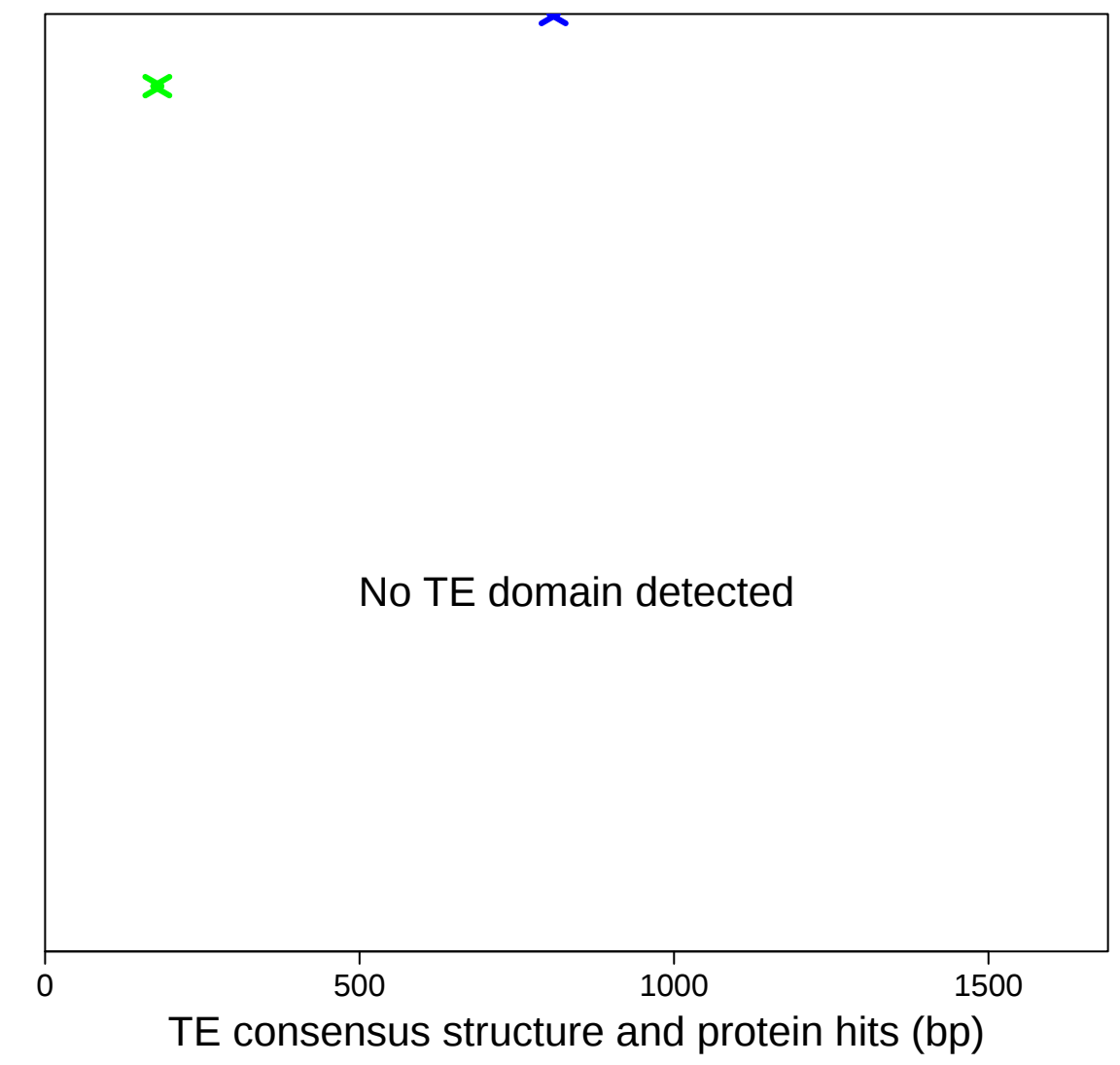
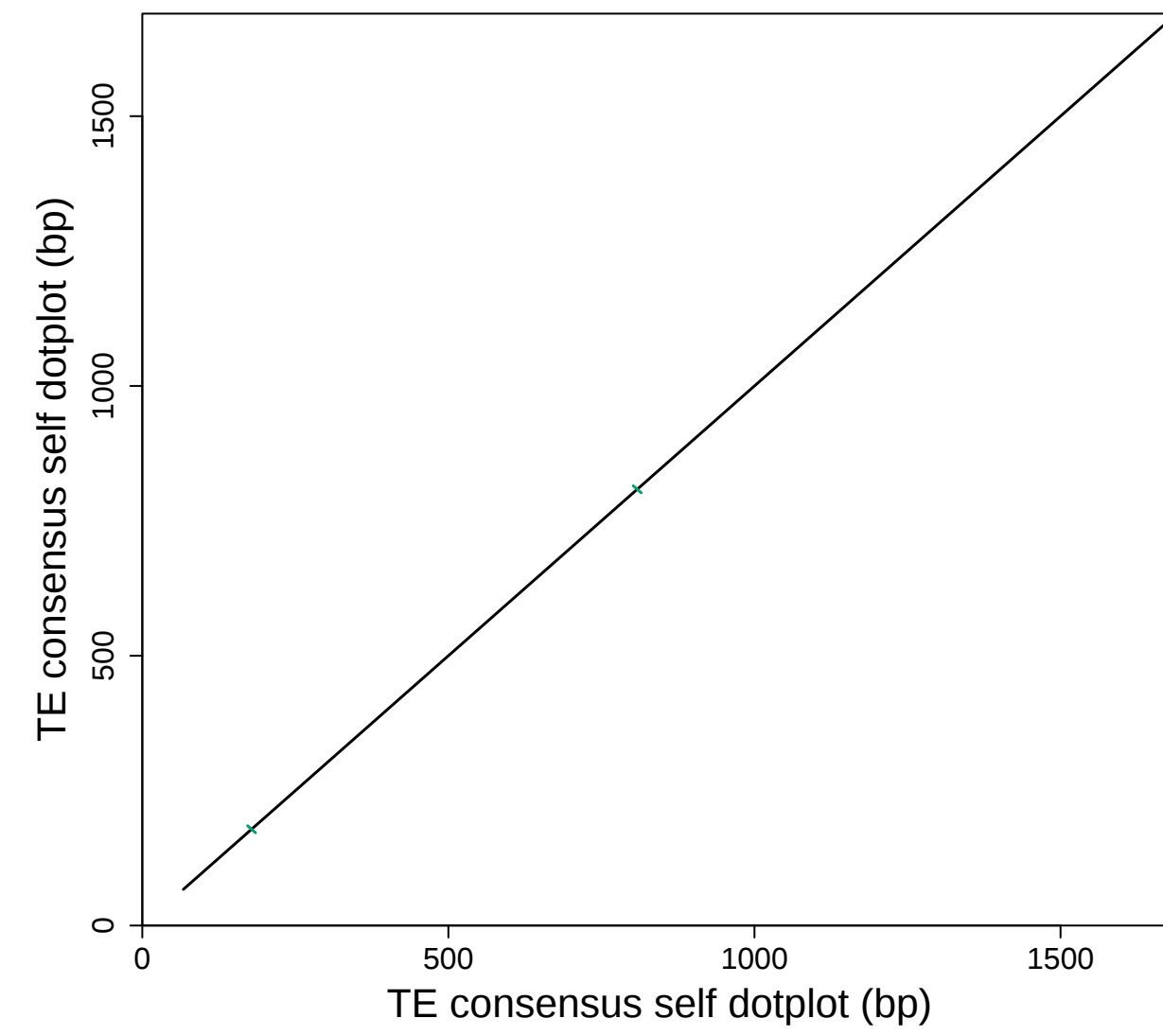
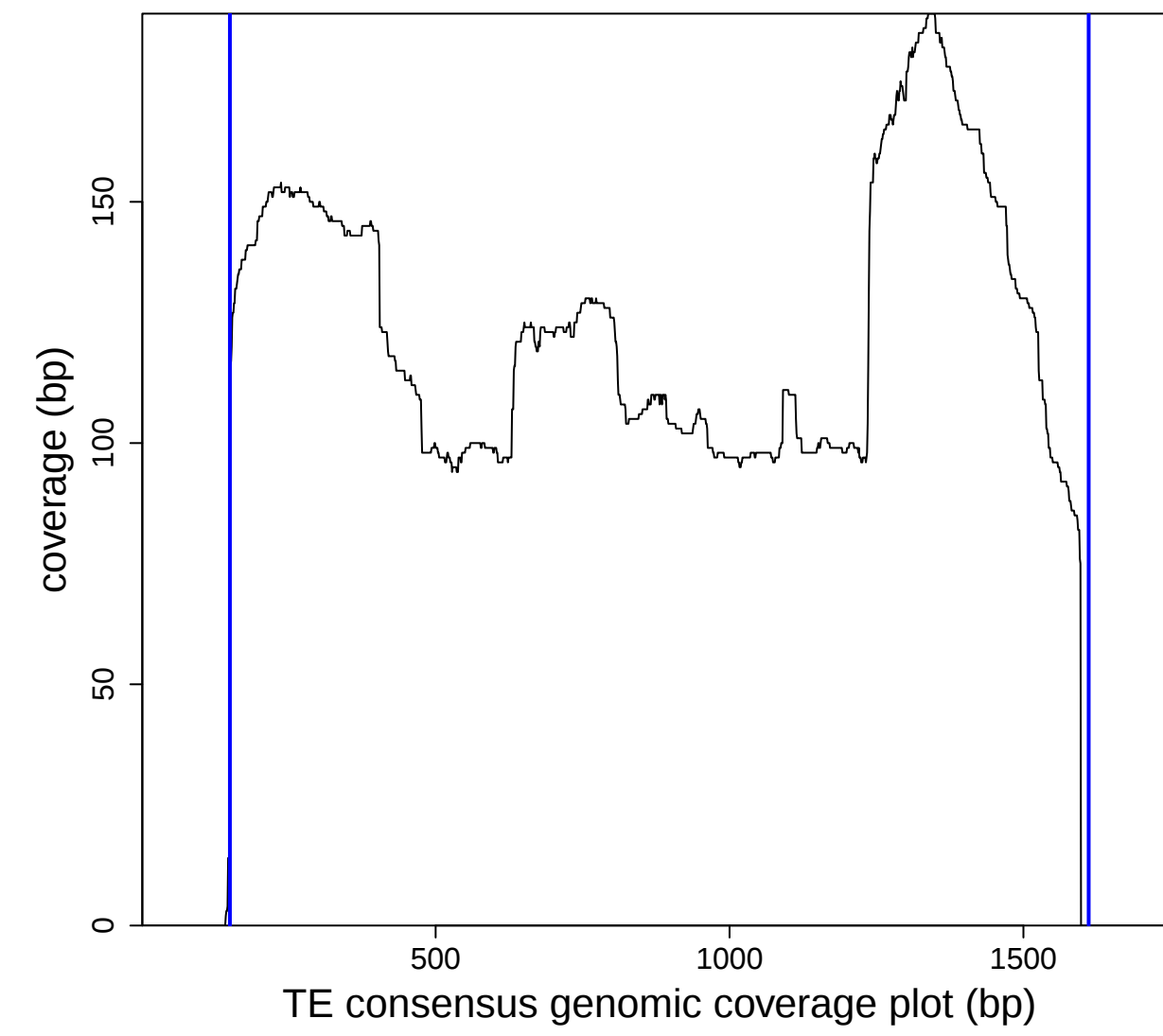
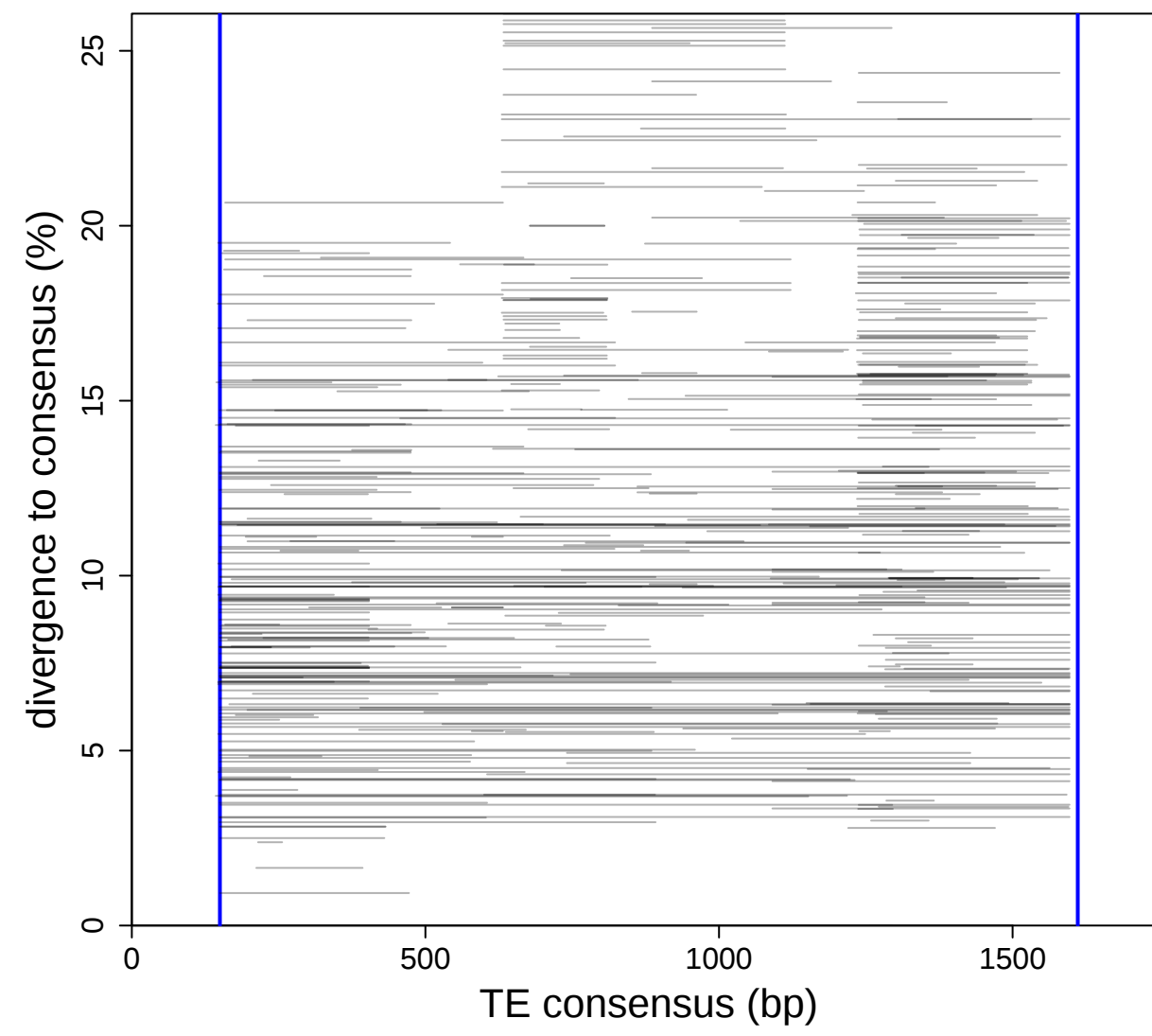
l.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcIn.fa_aln.fa_cl.fa_gs.fa
size: 1450bp; fragments: 421; full length: 25 (≥ 1305 bp)

After TEtrimmer 1450 bp

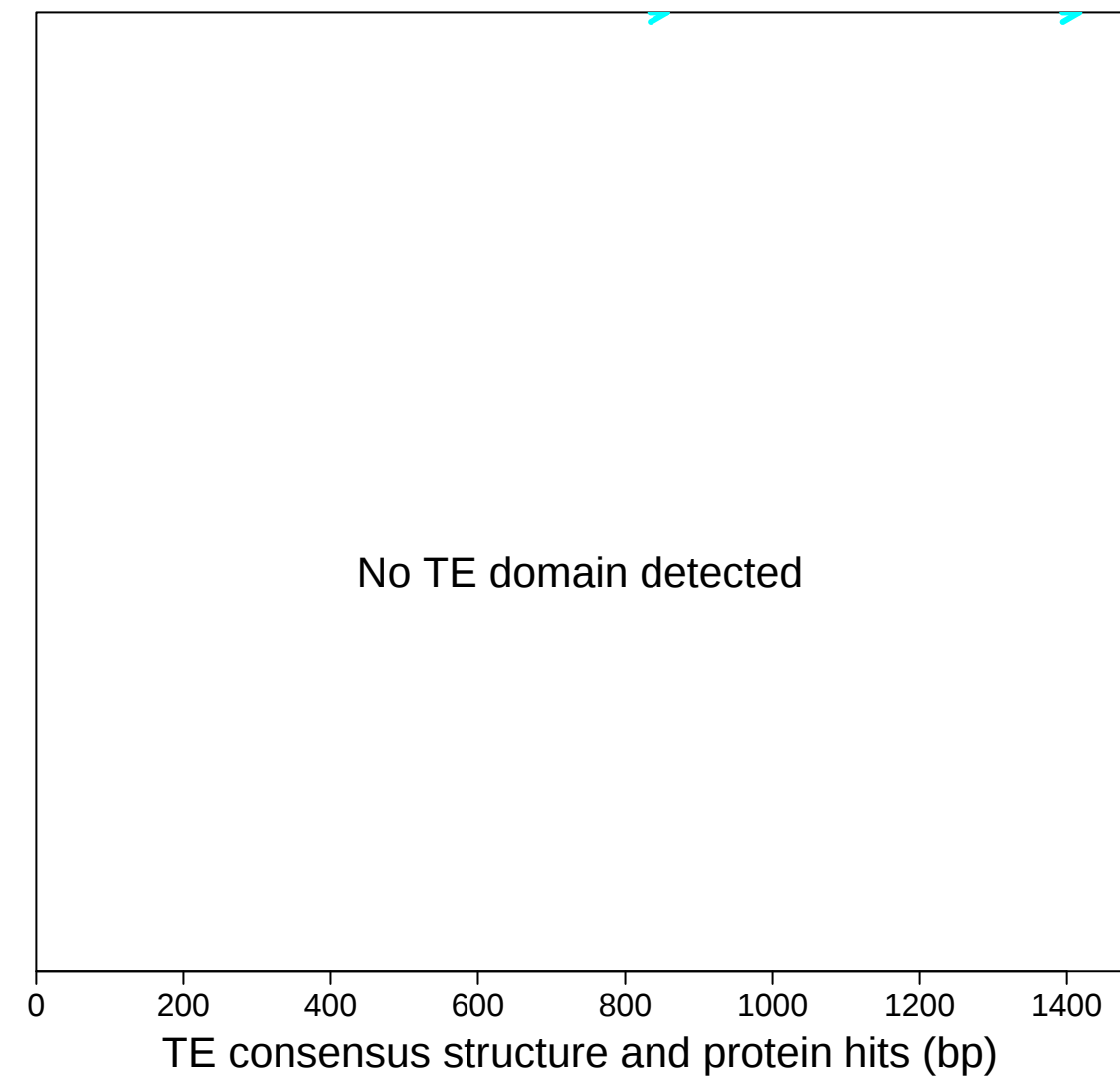
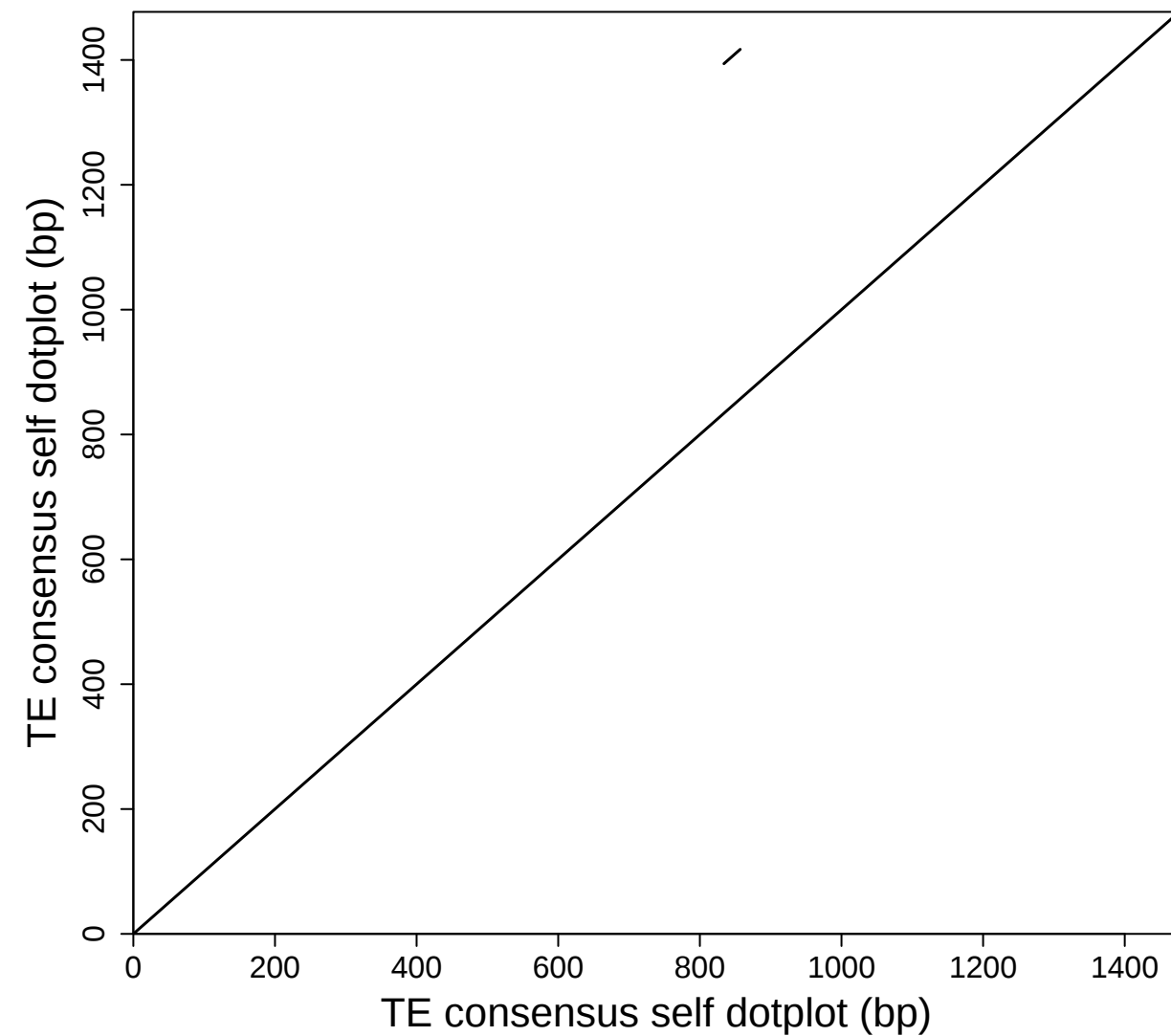
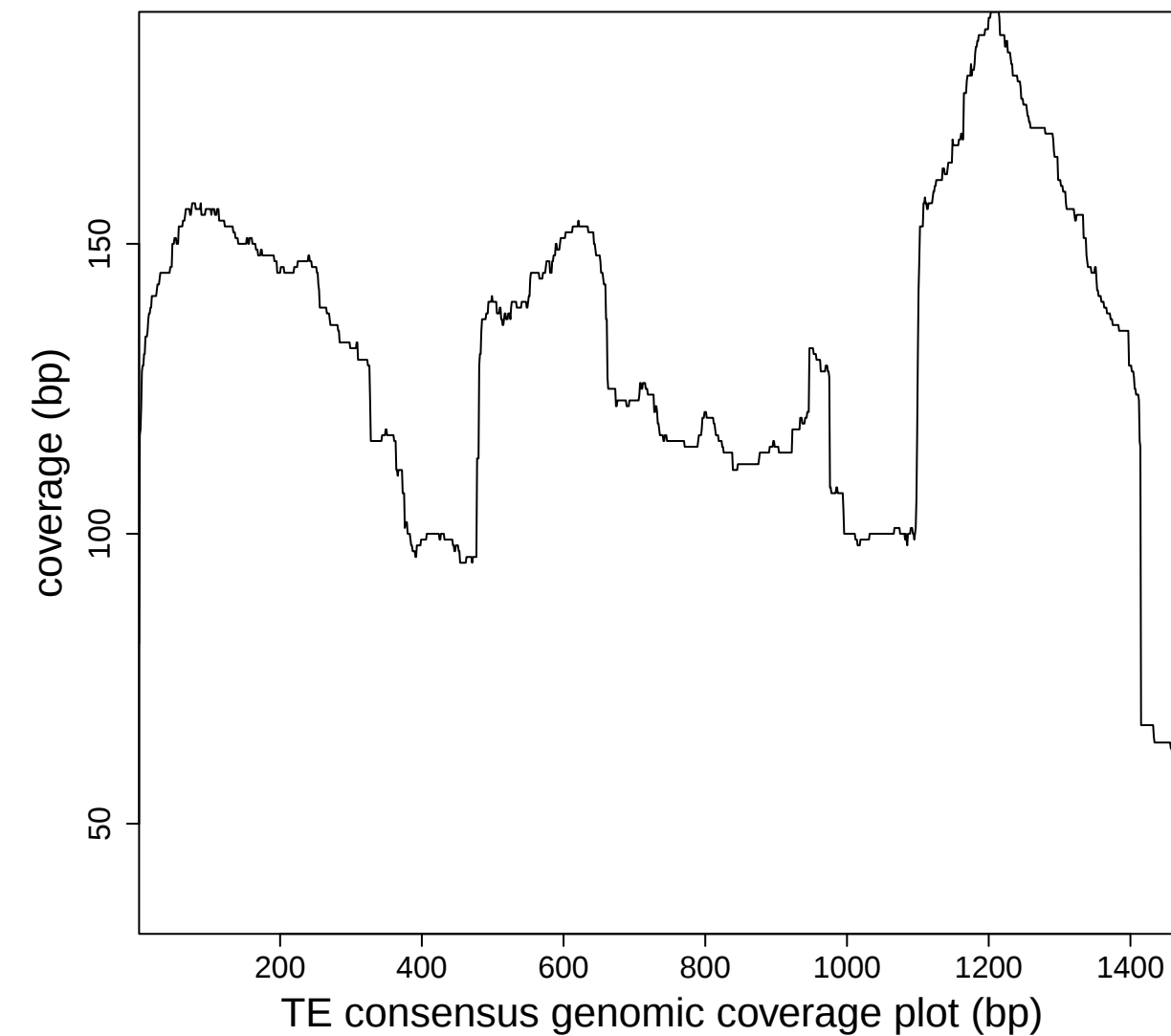
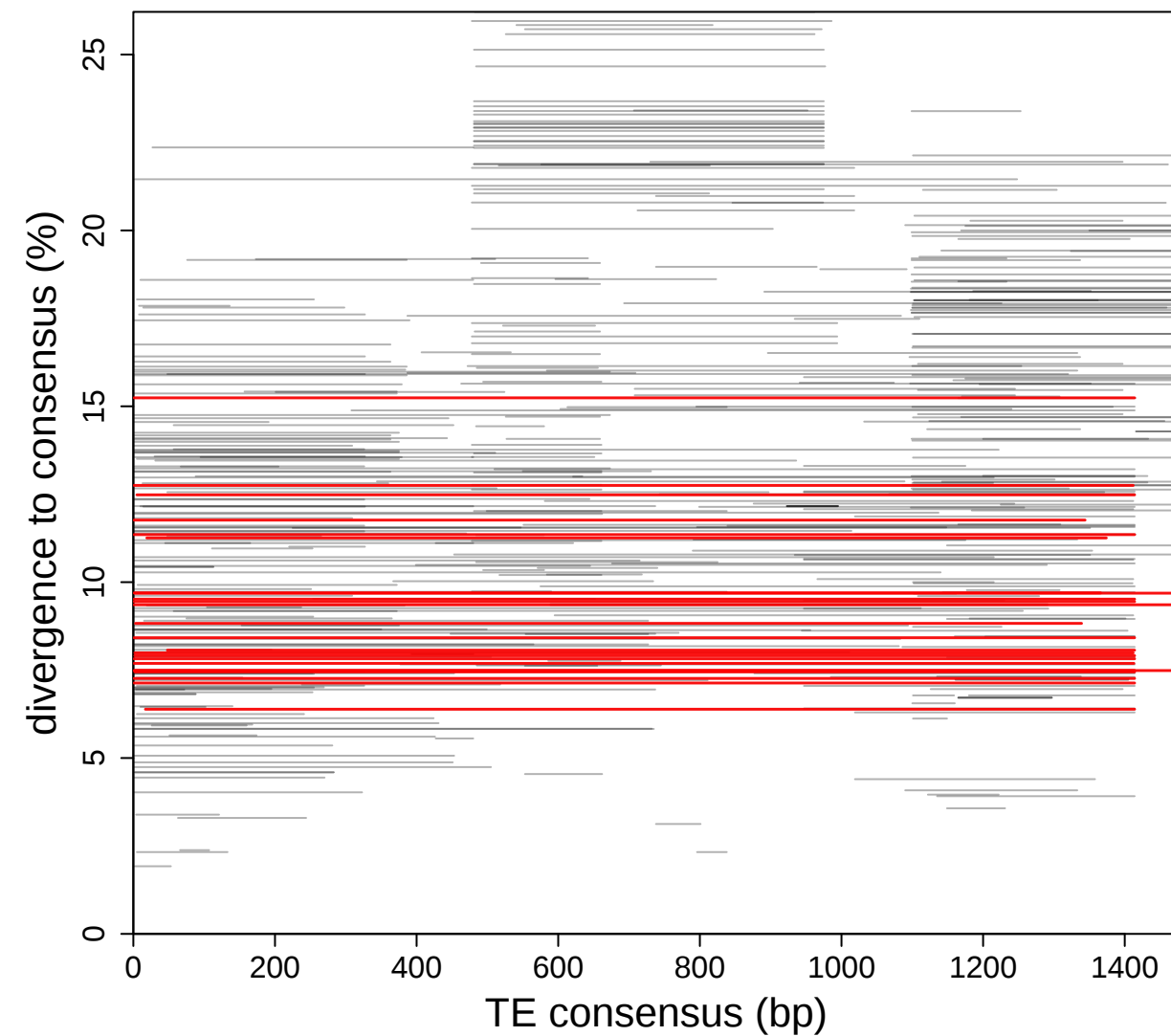


6.fasta.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa
size: 1761bp; fragments: 401; full length: 0 (>=1584.9bp)

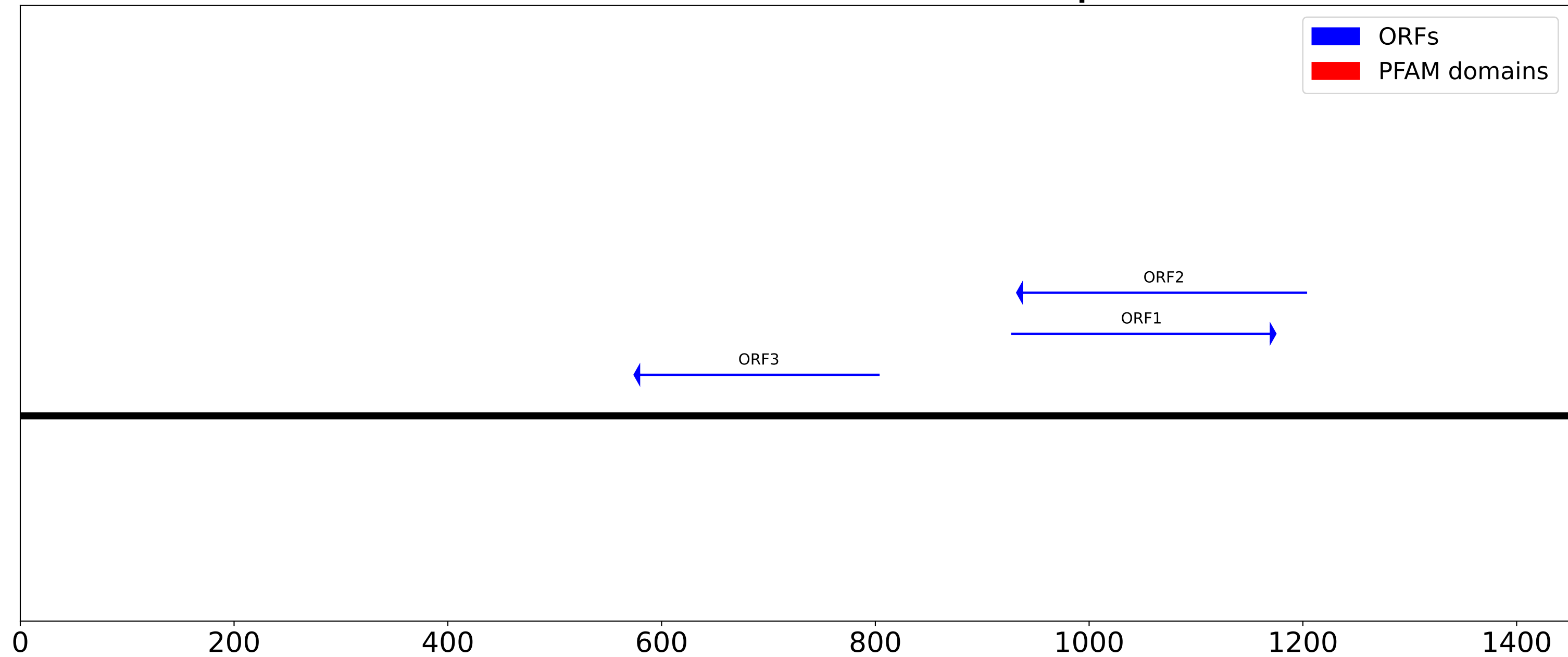
After TEtrimmer Extended plot Blue lines are boundaries



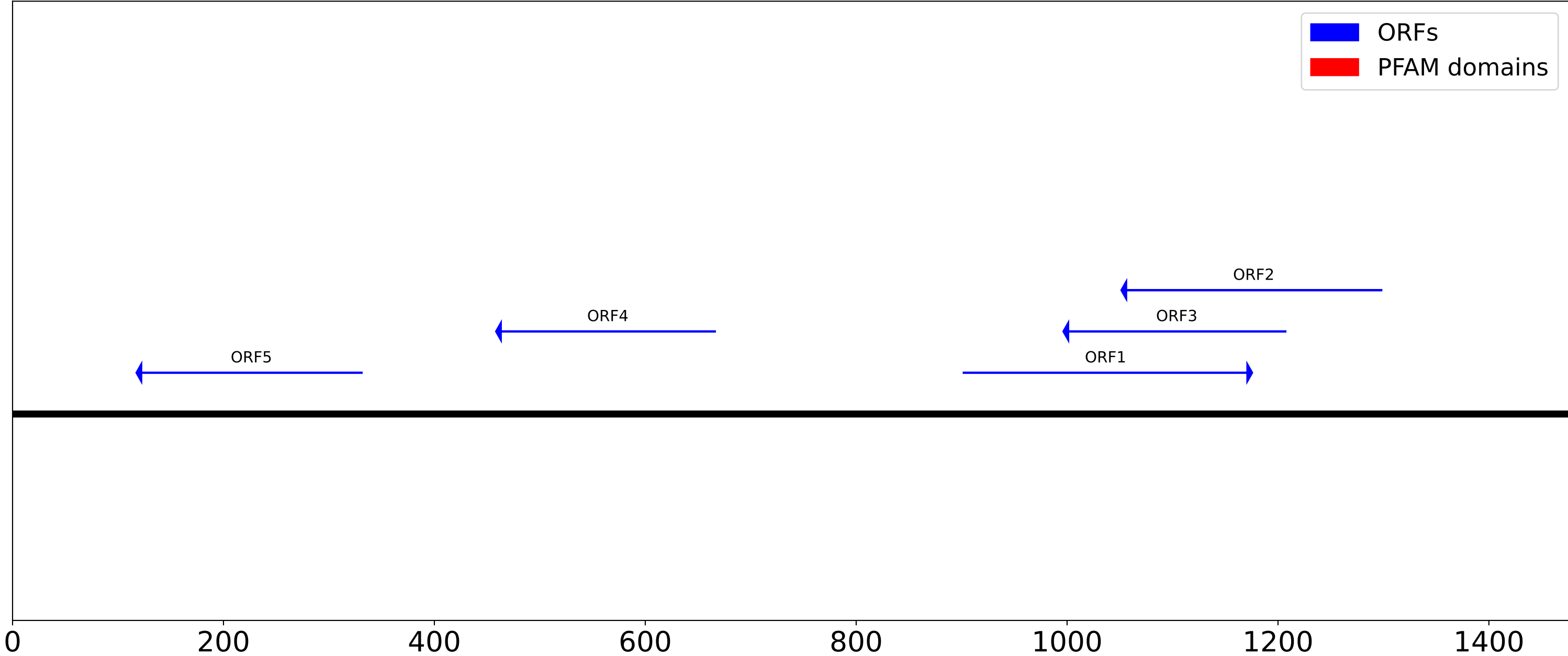
TE: rnd_5_family_106
size: 1477bp; fragments: 433; full length: 24 (≥ 1329.3 bp)



After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

