

Start crop Point

End crop Point

0

10

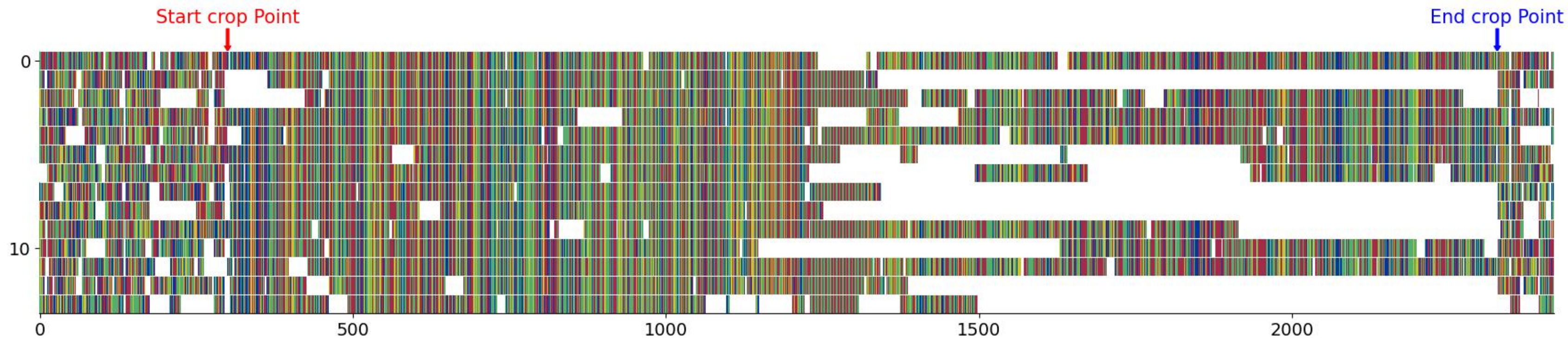
0

500

1000

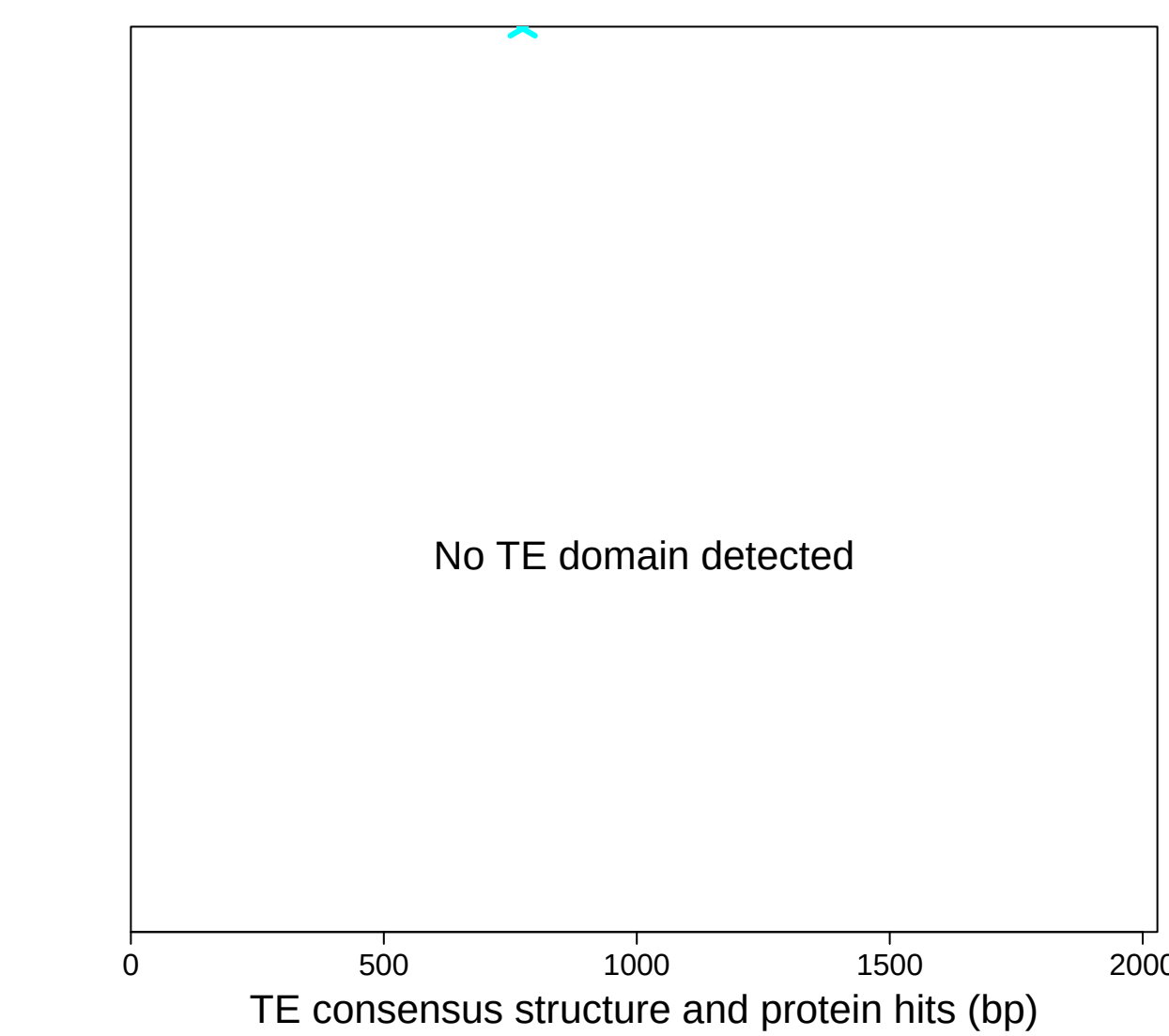
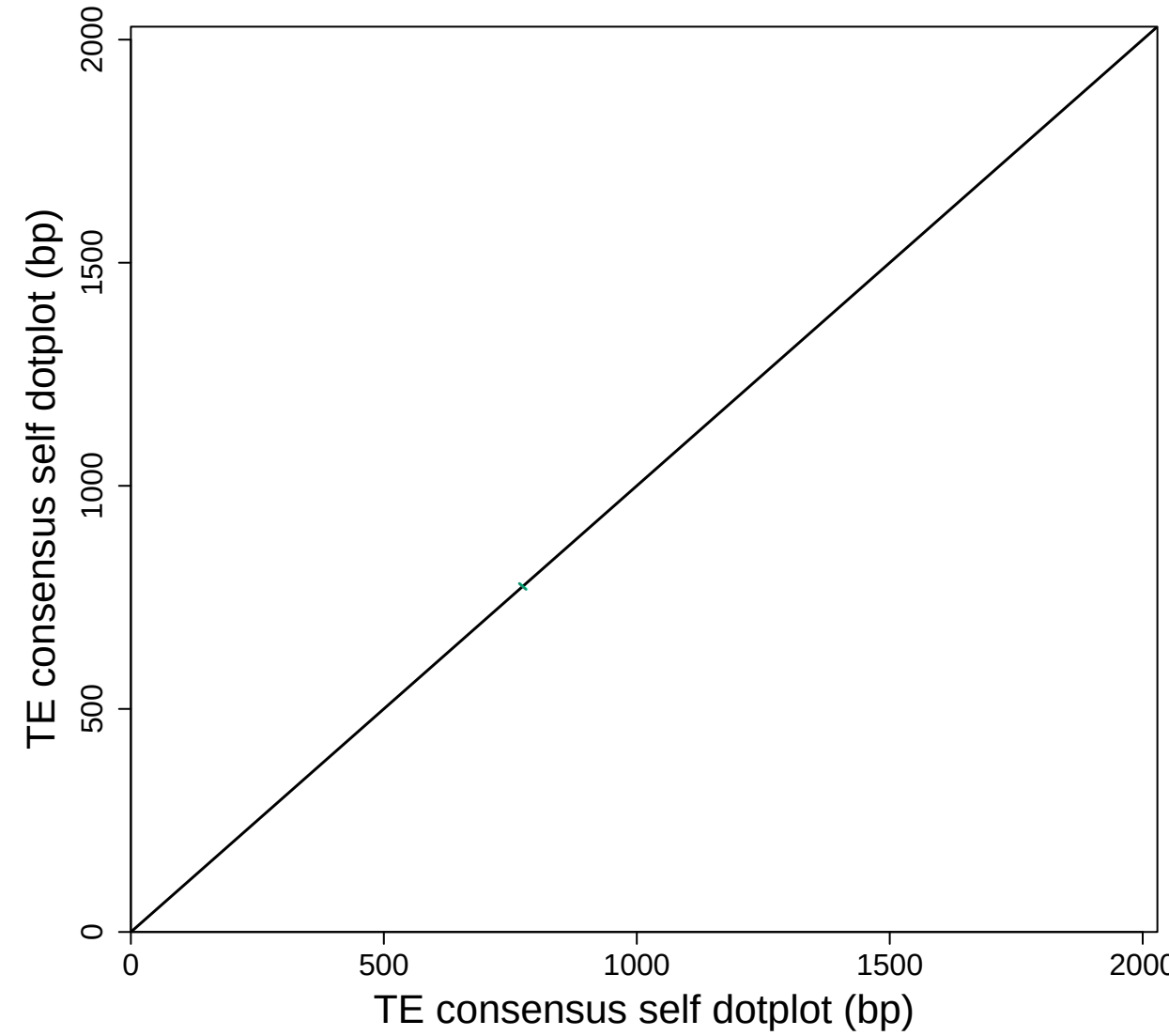
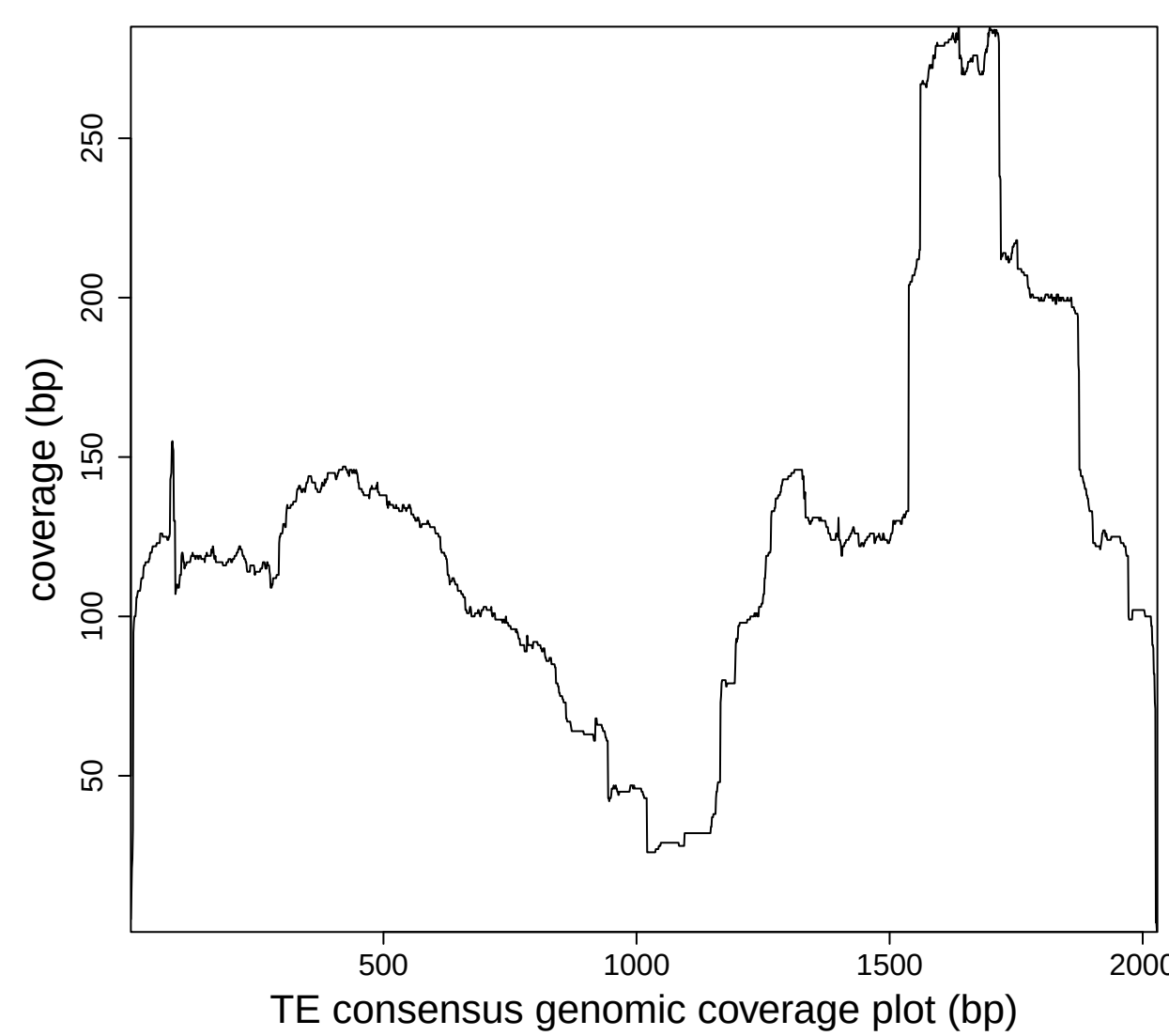
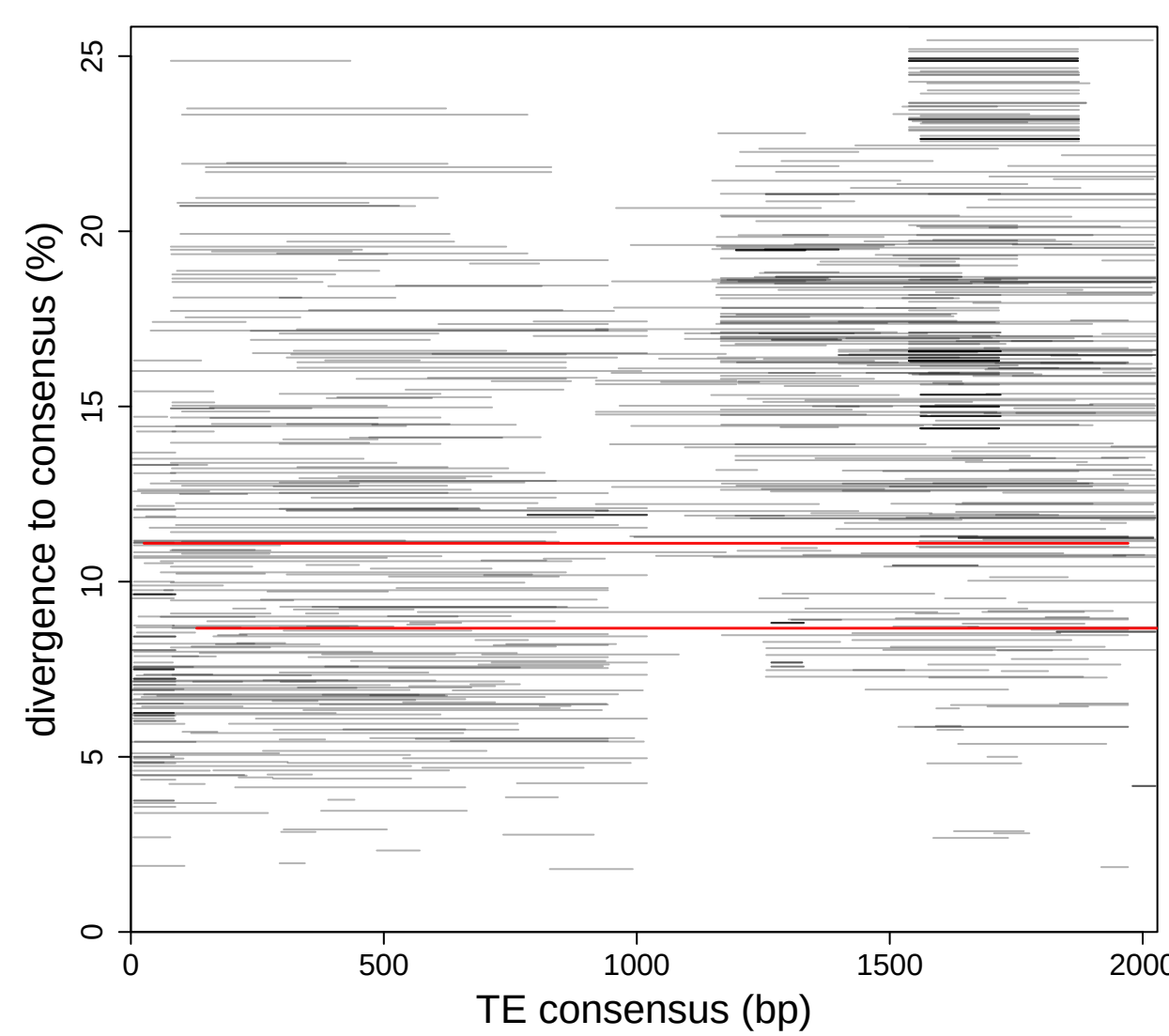
1500

2000



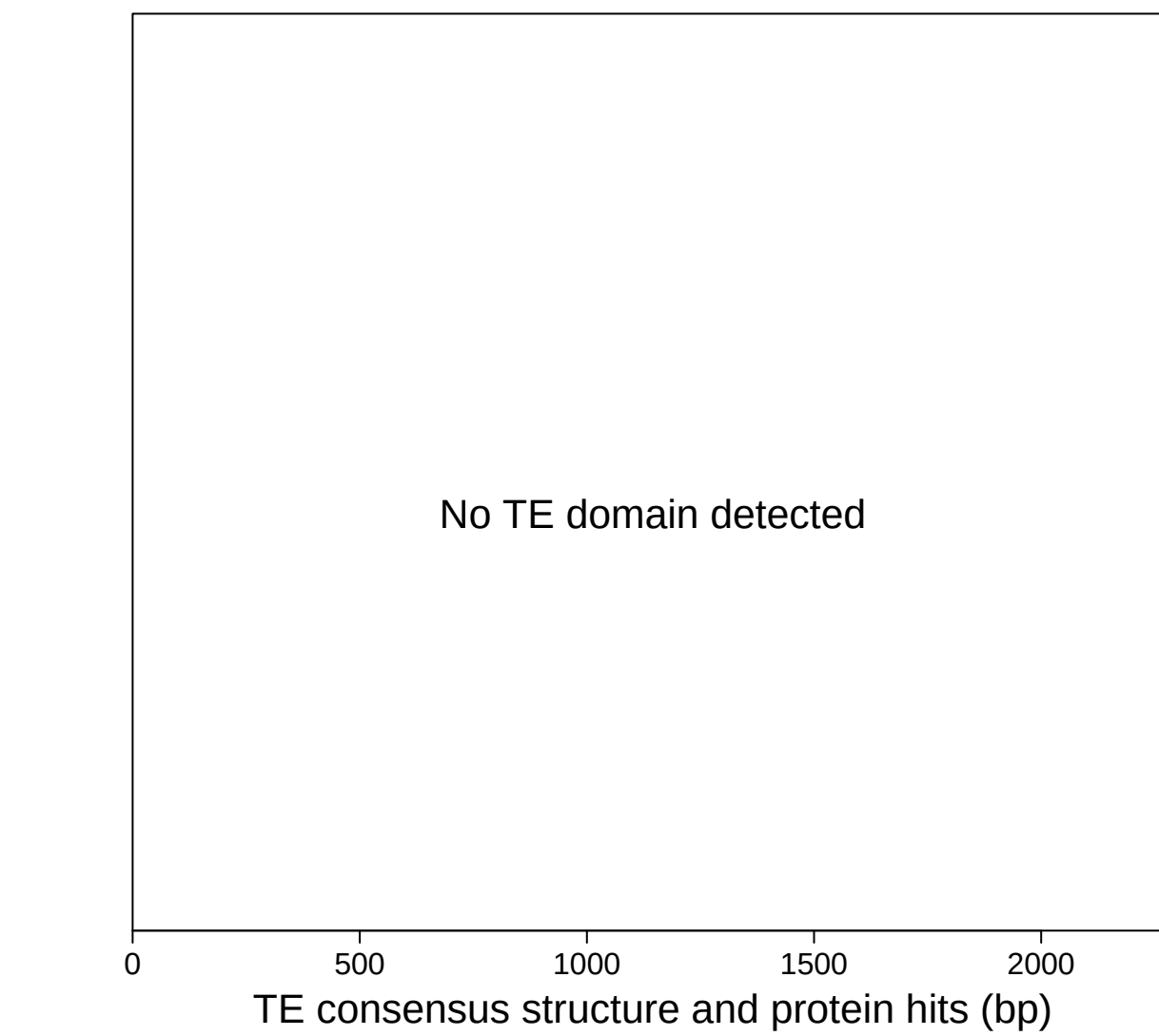
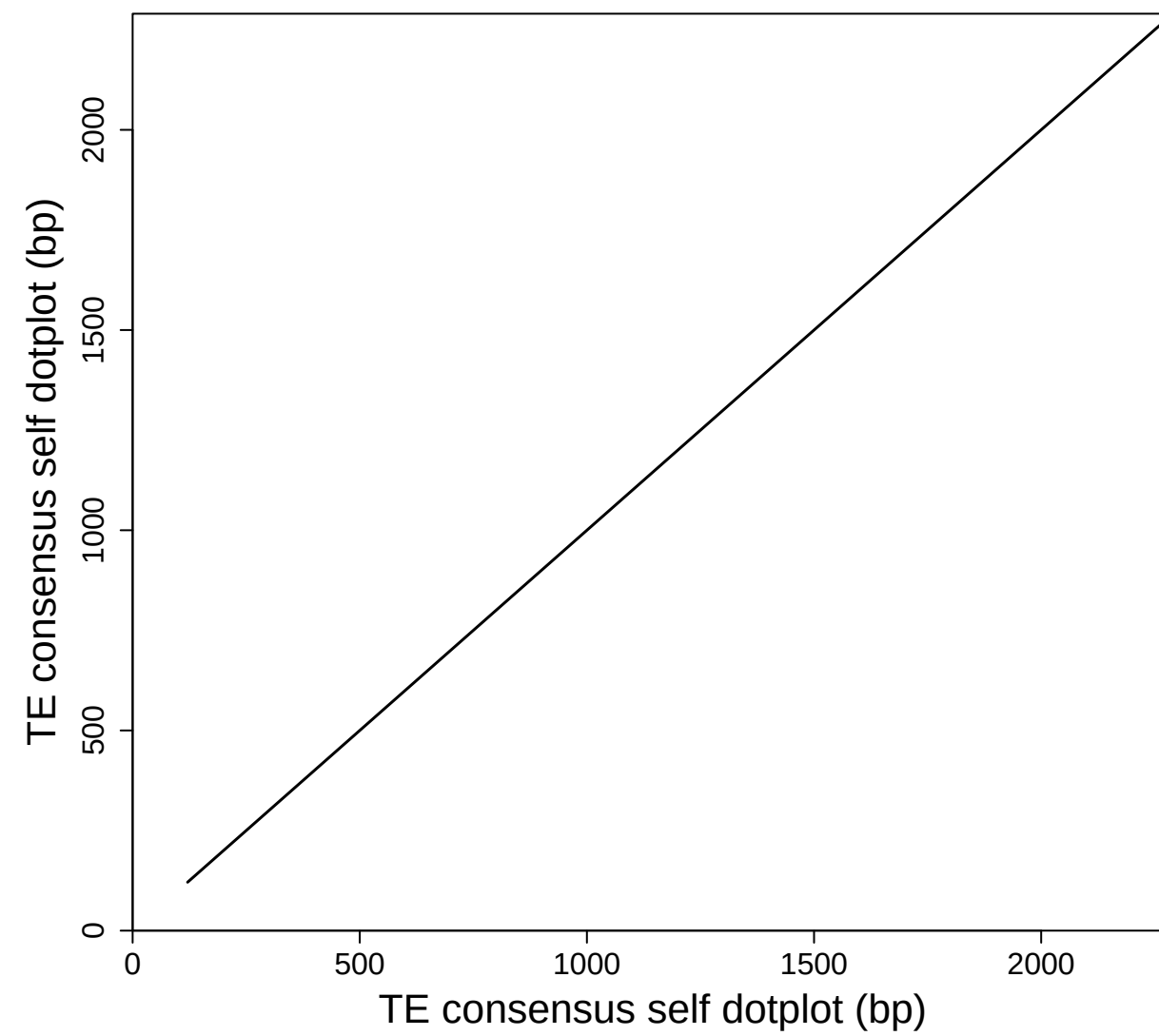
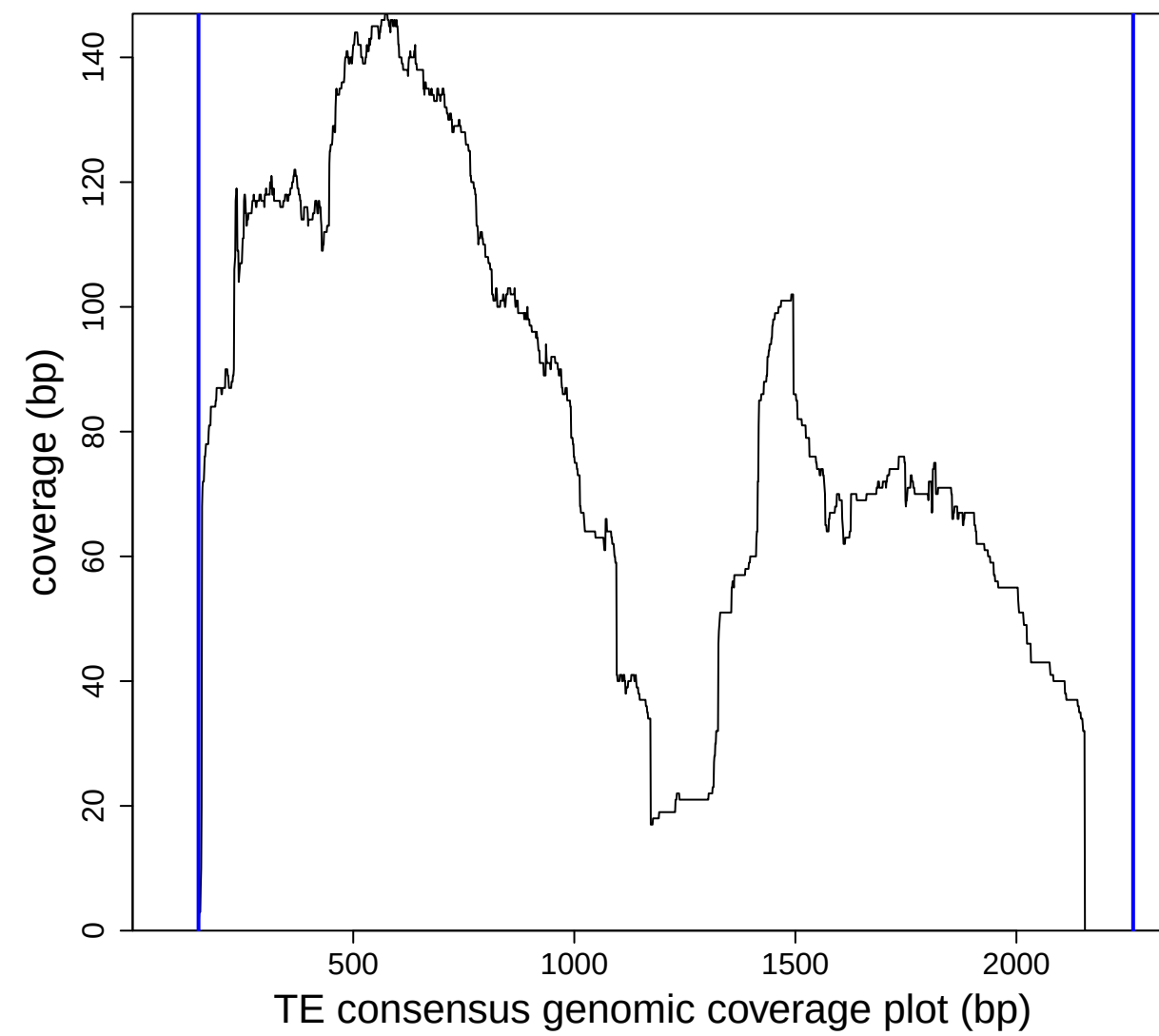
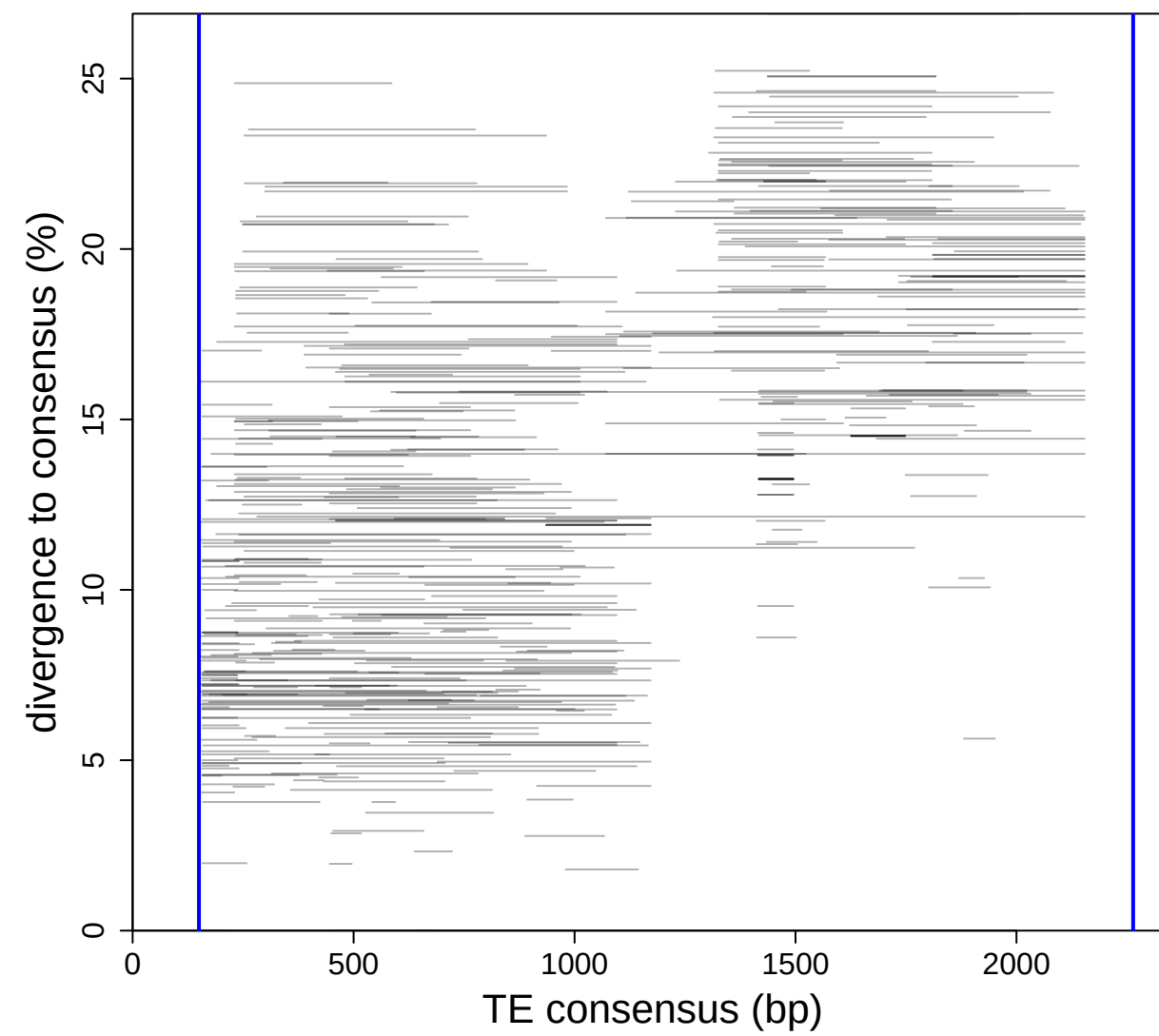
a.b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.f
size: 2029bp; fragments: 778; full length: 2 (>=1826.1bp)

After TEtrimmer 2029 bp

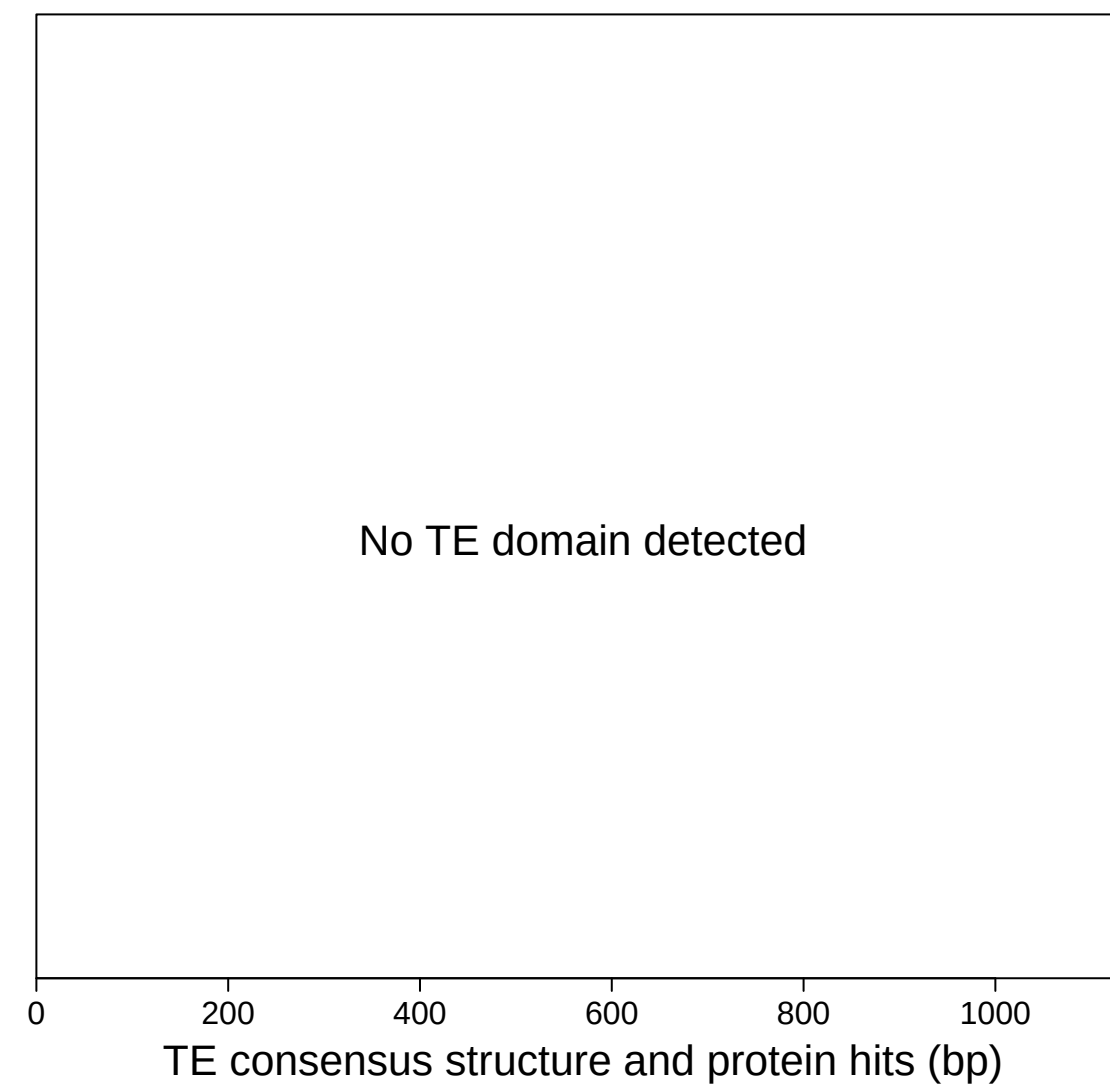
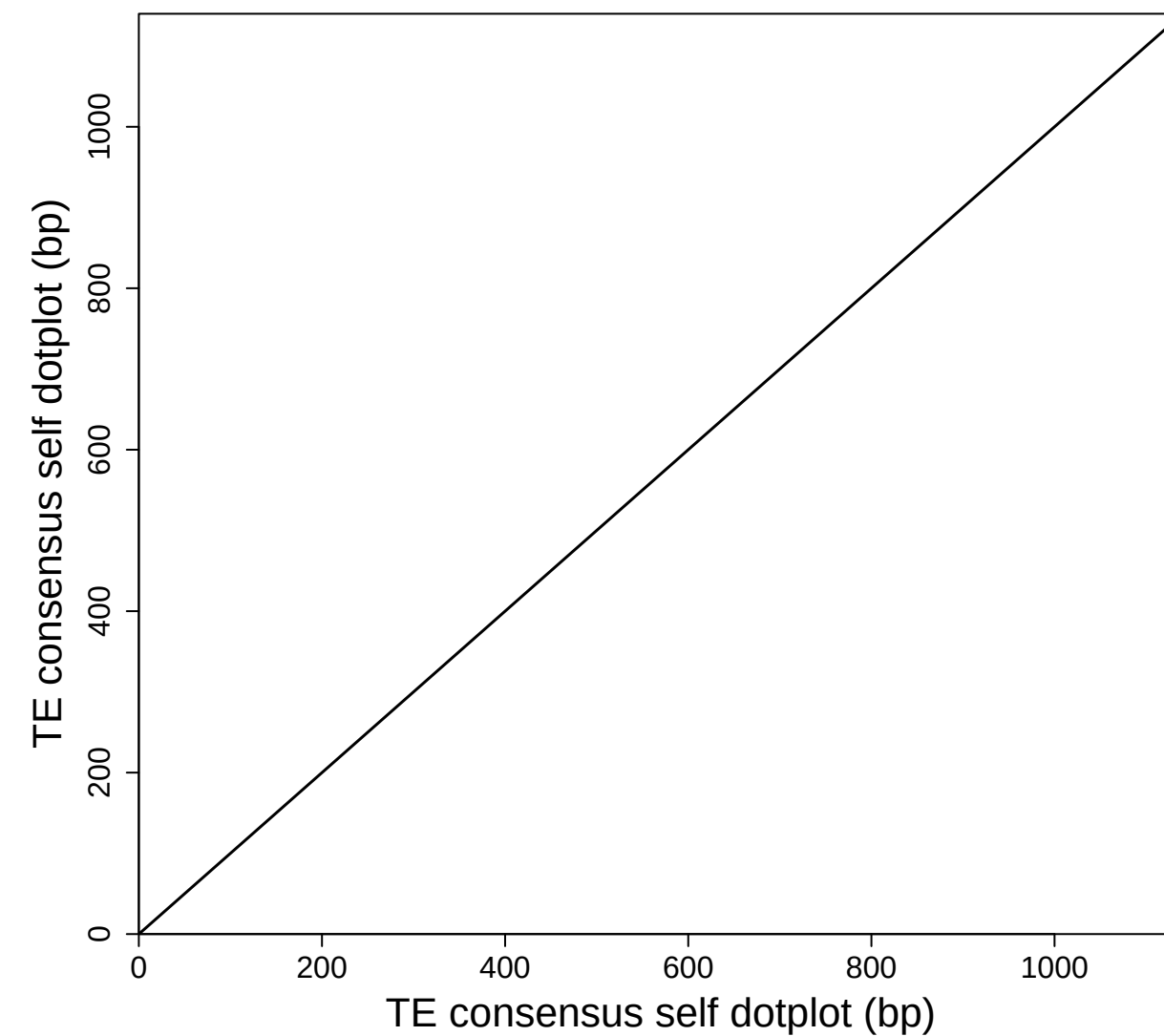
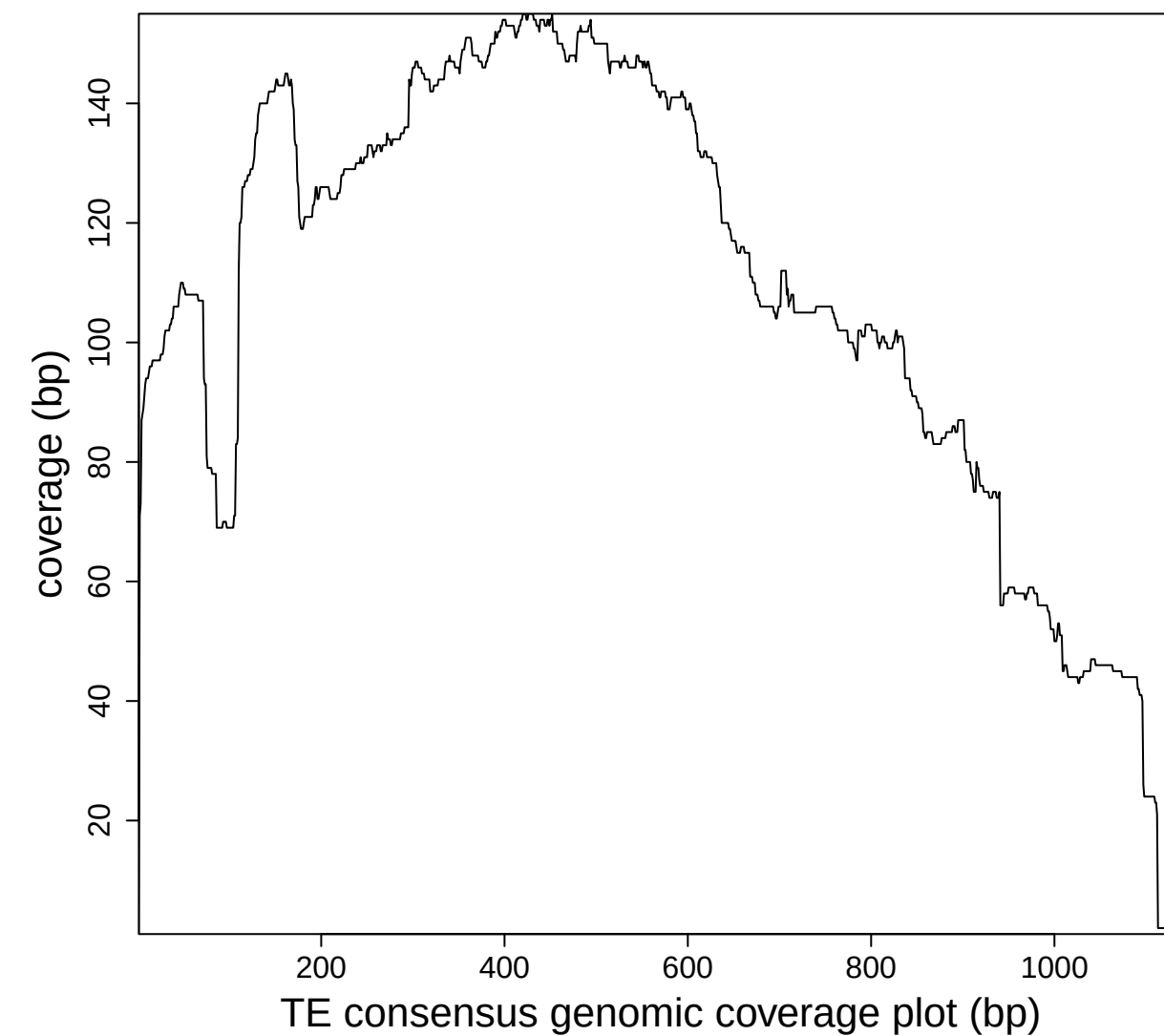
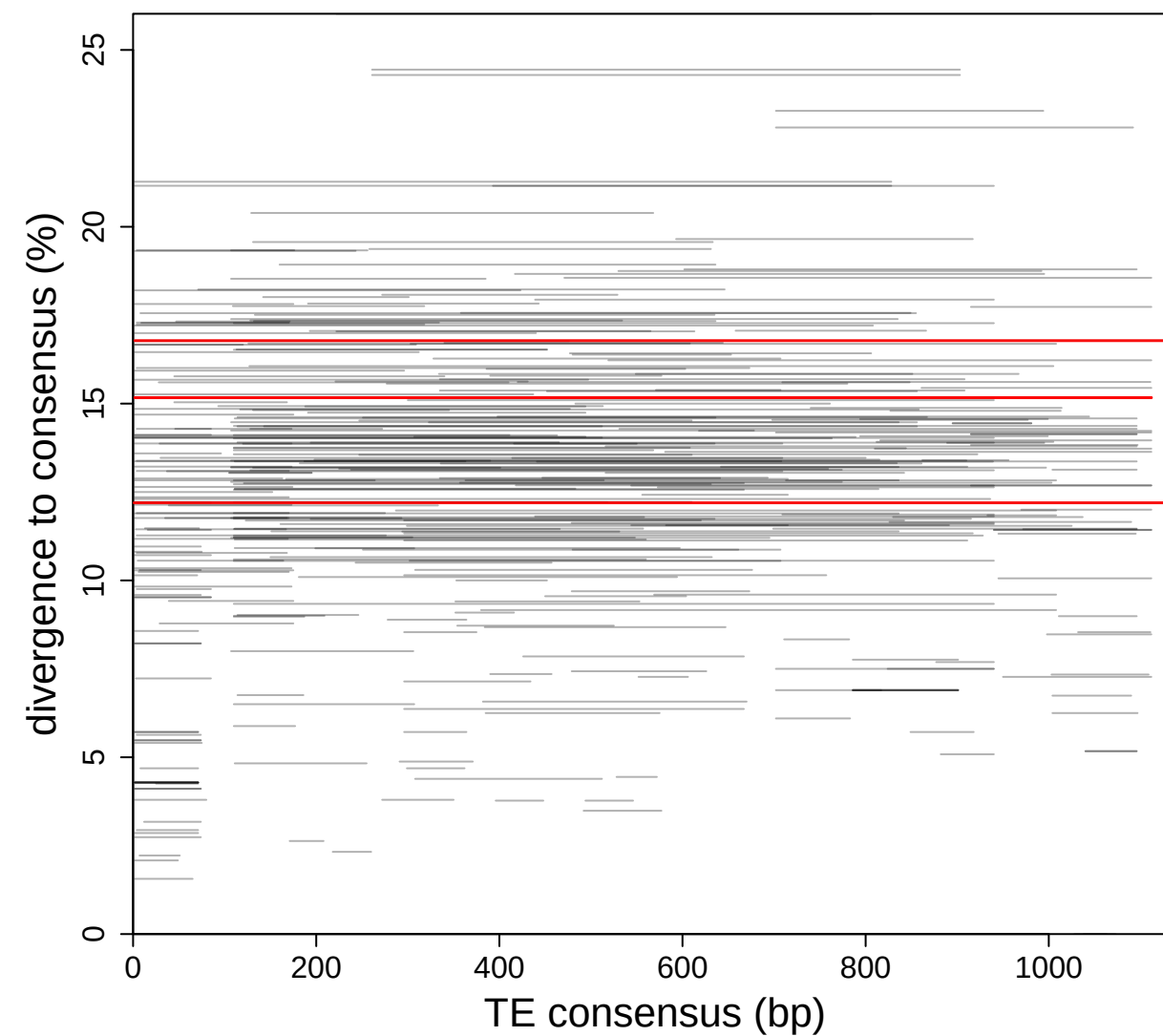


2.fasta.b.bed uf.bed g_2.bed fm_1.bed 0 0 bcln.fa aln.fa cl.f
size: 2354bp; fragments: 441; full length: 0 (>=2118.6bp)

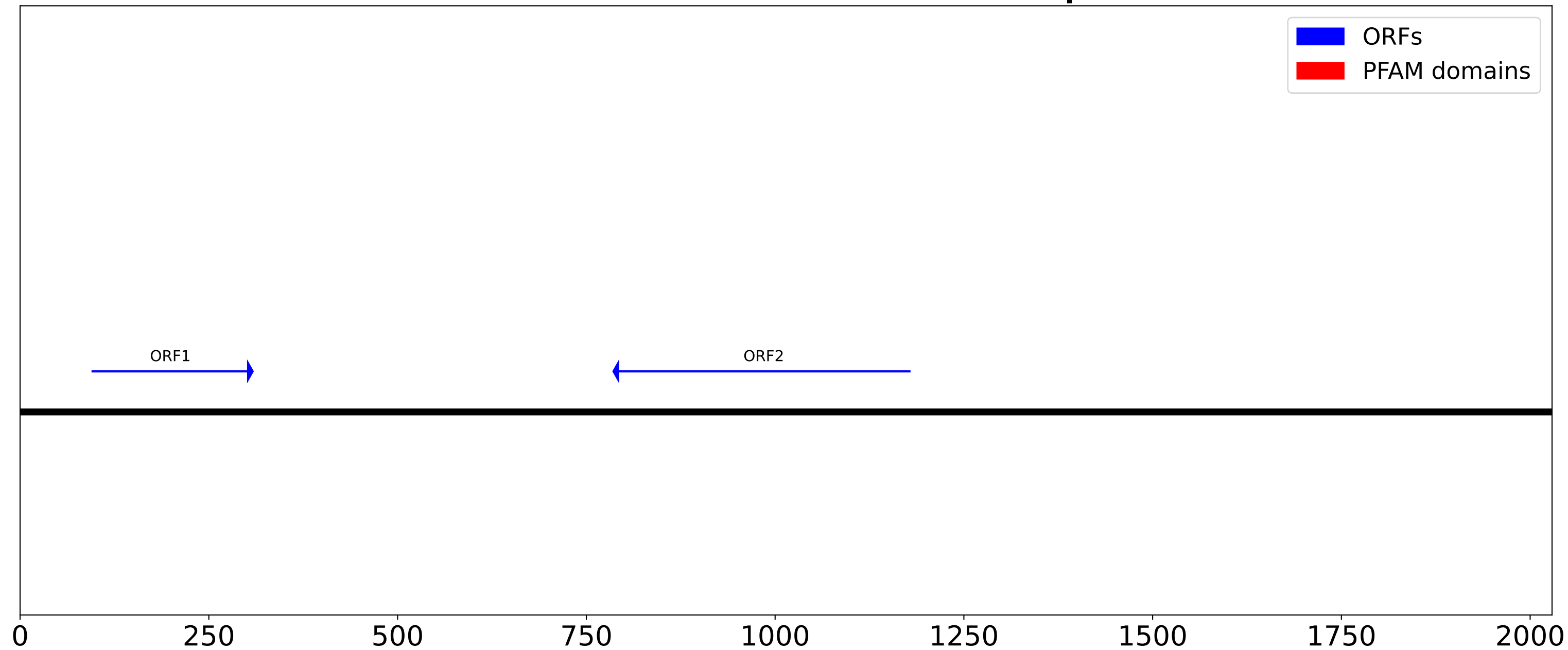
After TEtrimmer Extended plot Blue lines are boundaries



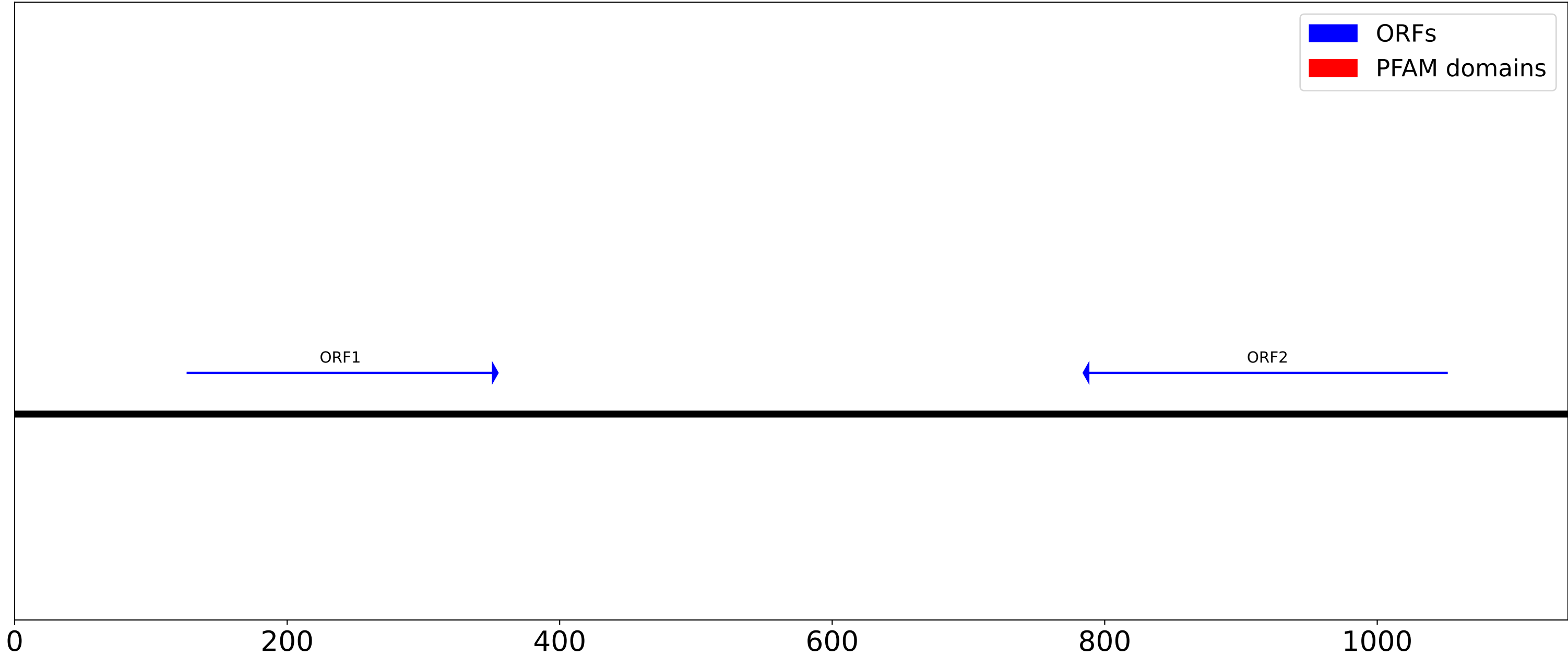
TE: rnd_5_family_3862
size: 1140bp; fragments: 368; full length: 4 (≥ 1026 bp)



After TEtrimmer ORF and PFAM domain plot



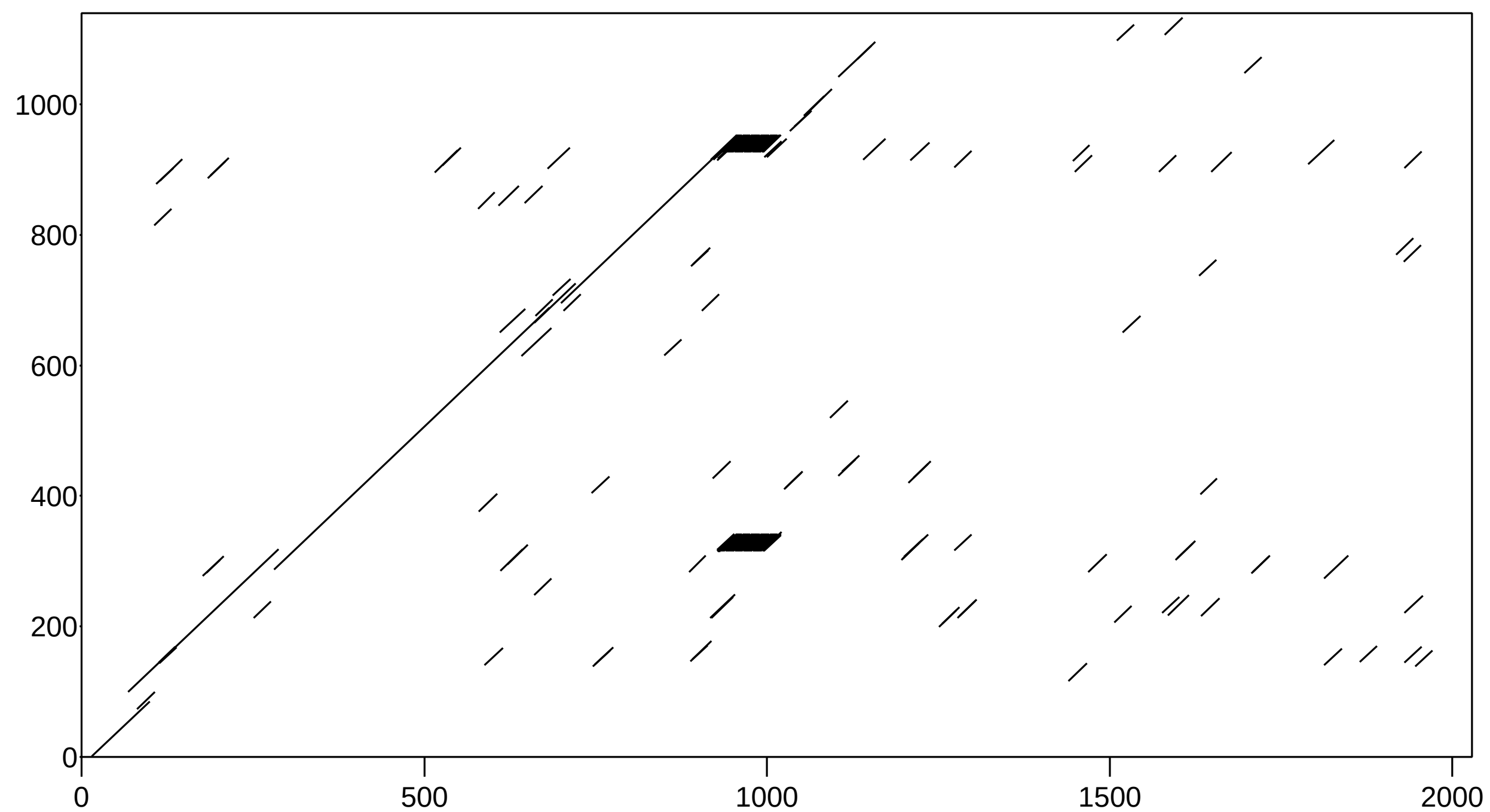
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)