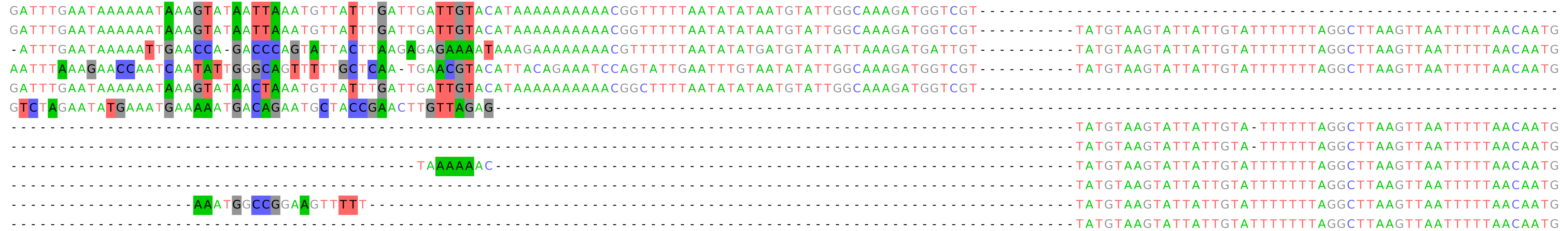
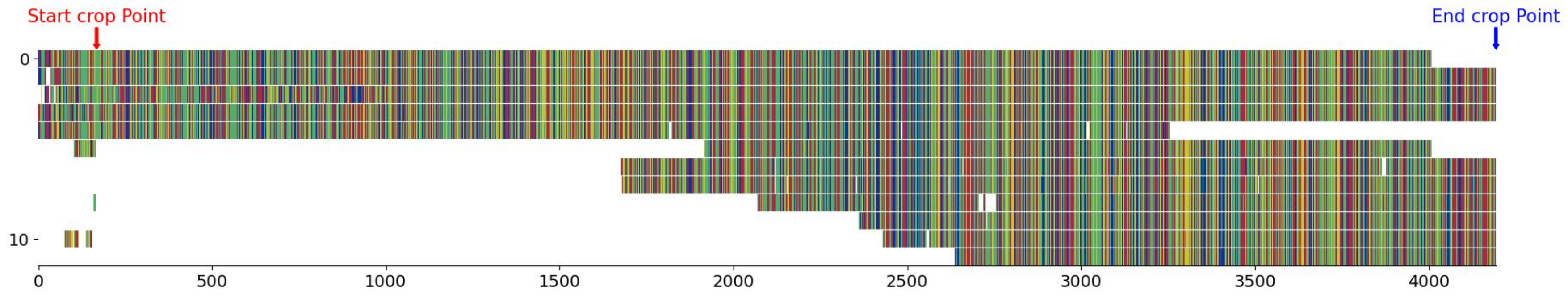


Start crop Point

End crop Point

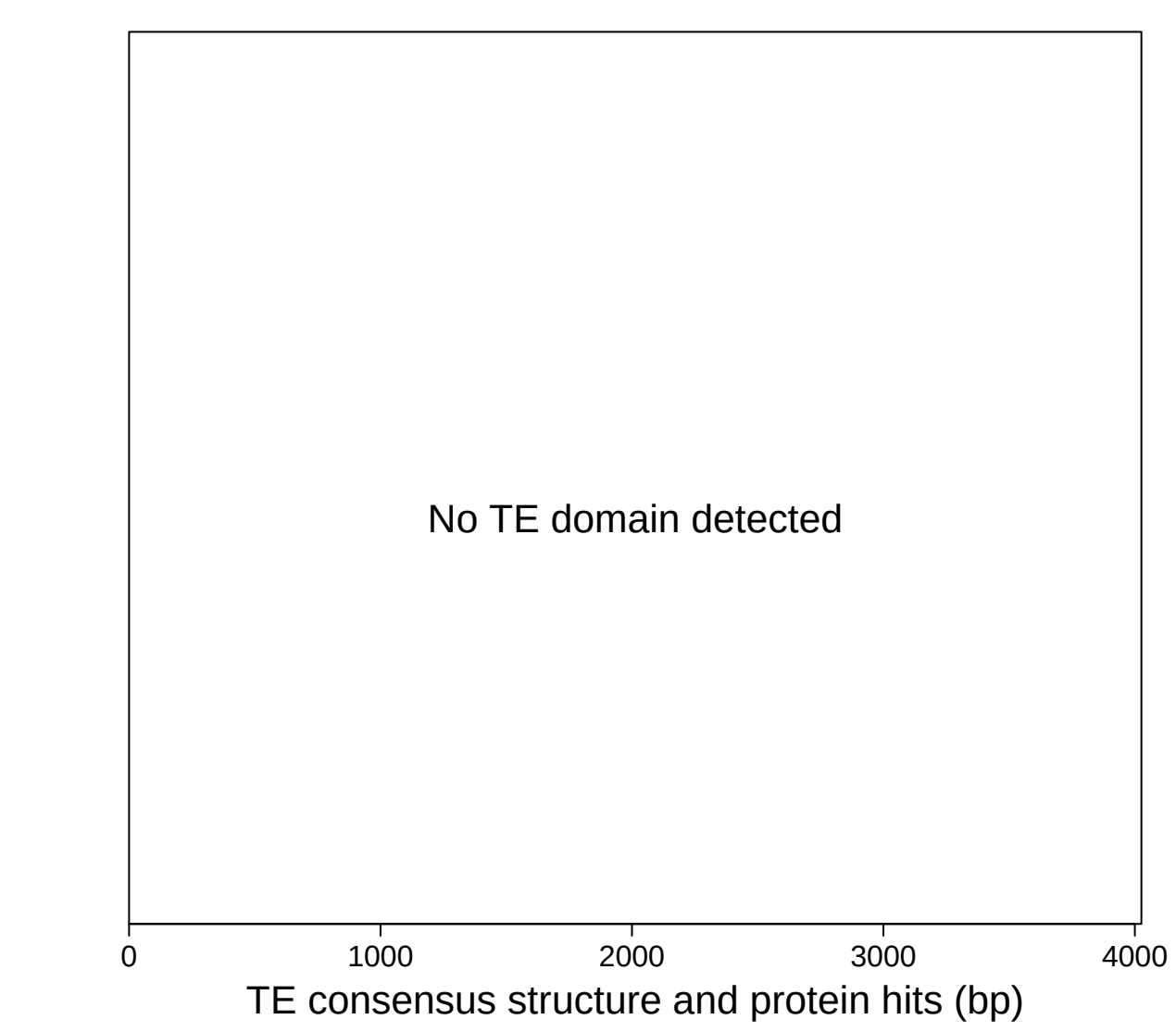
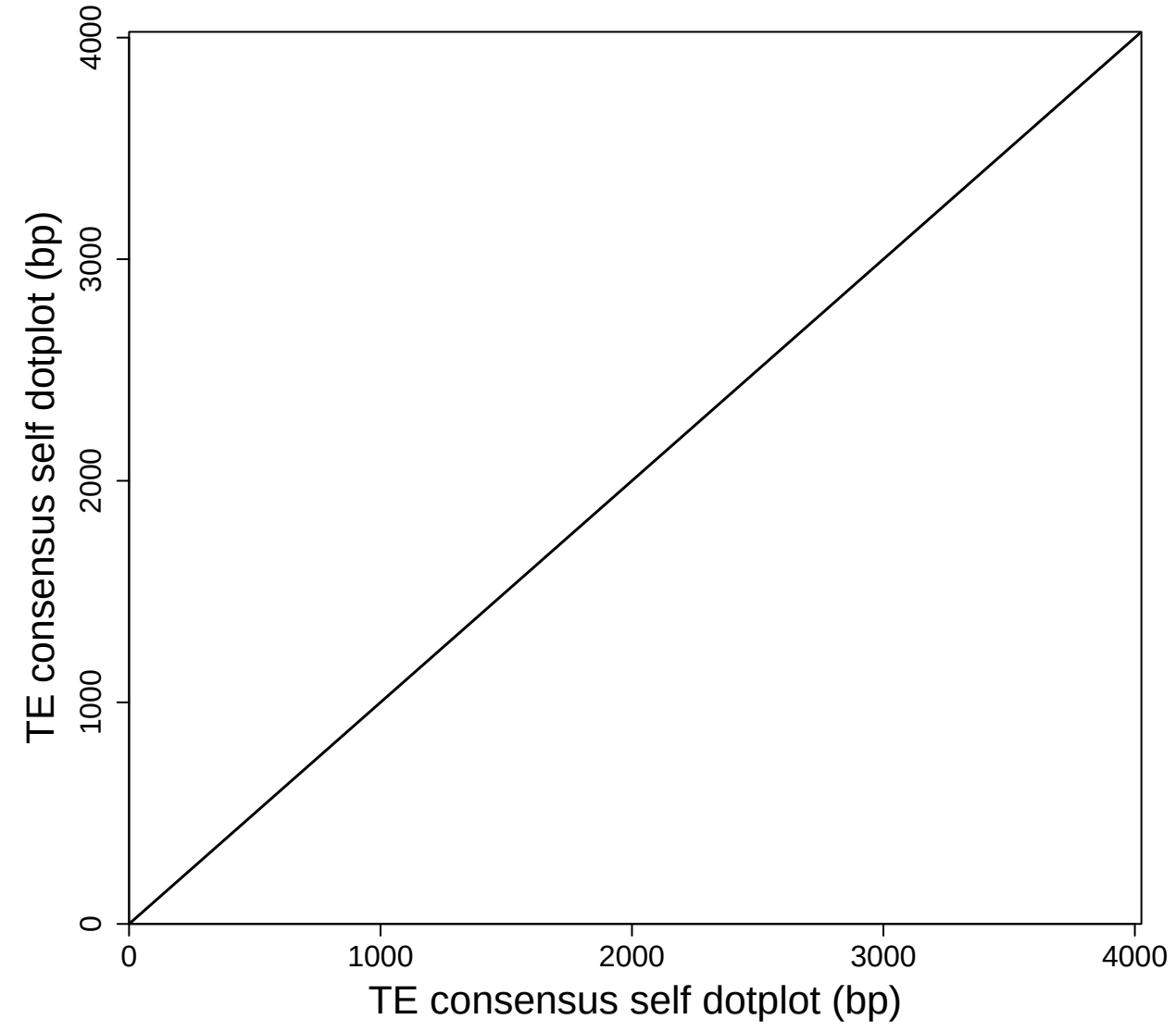
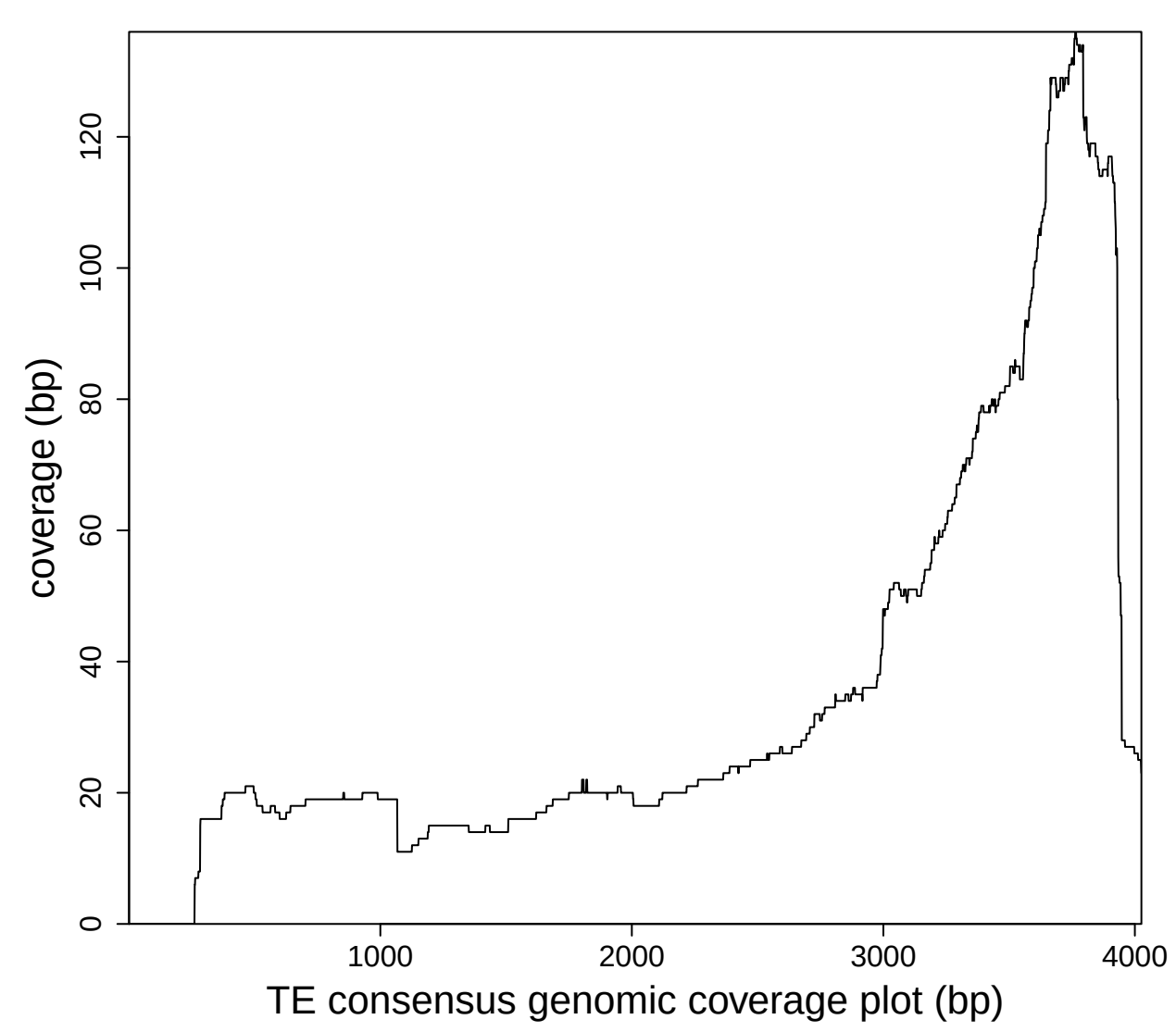
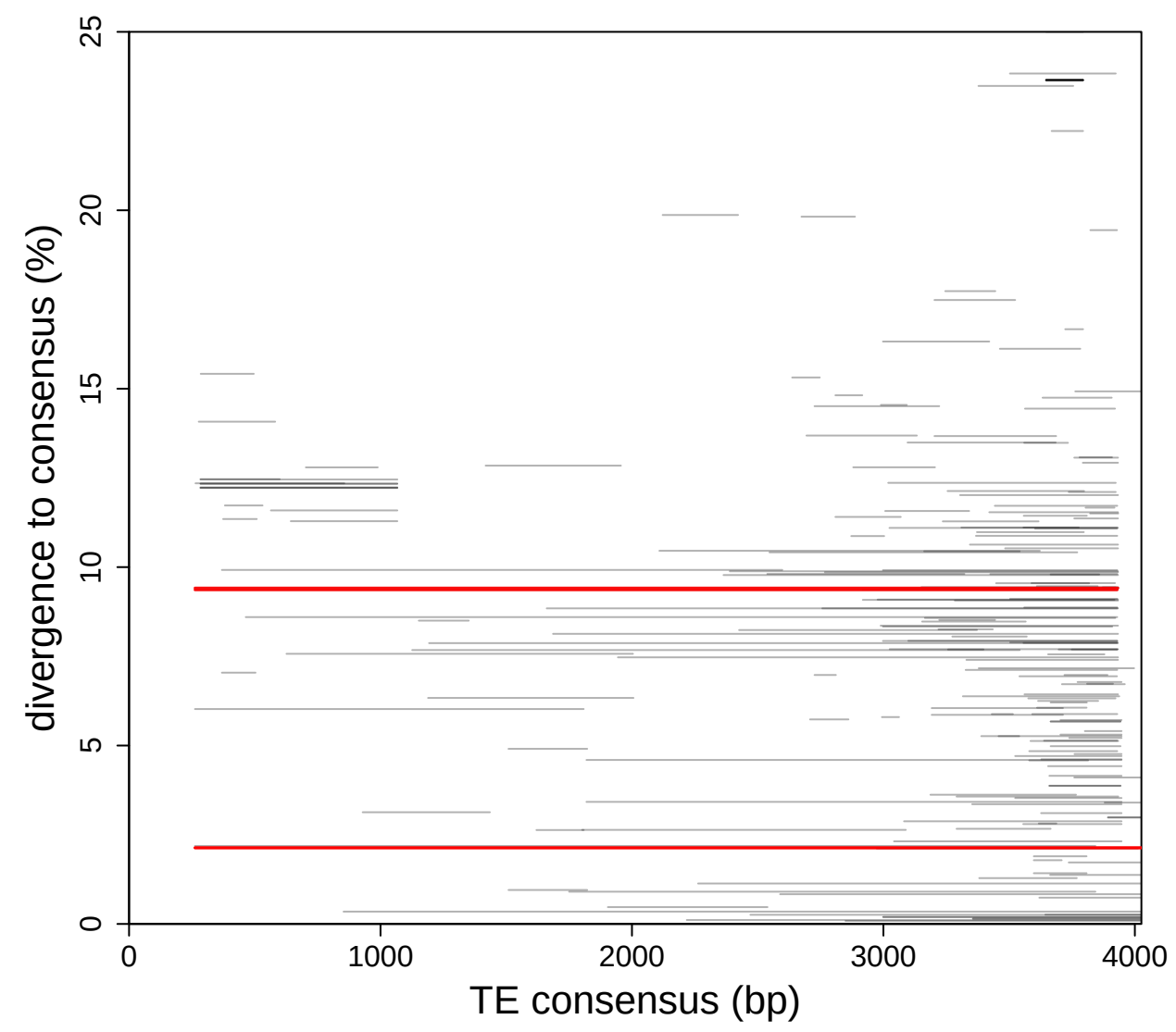
MSA length = 4026





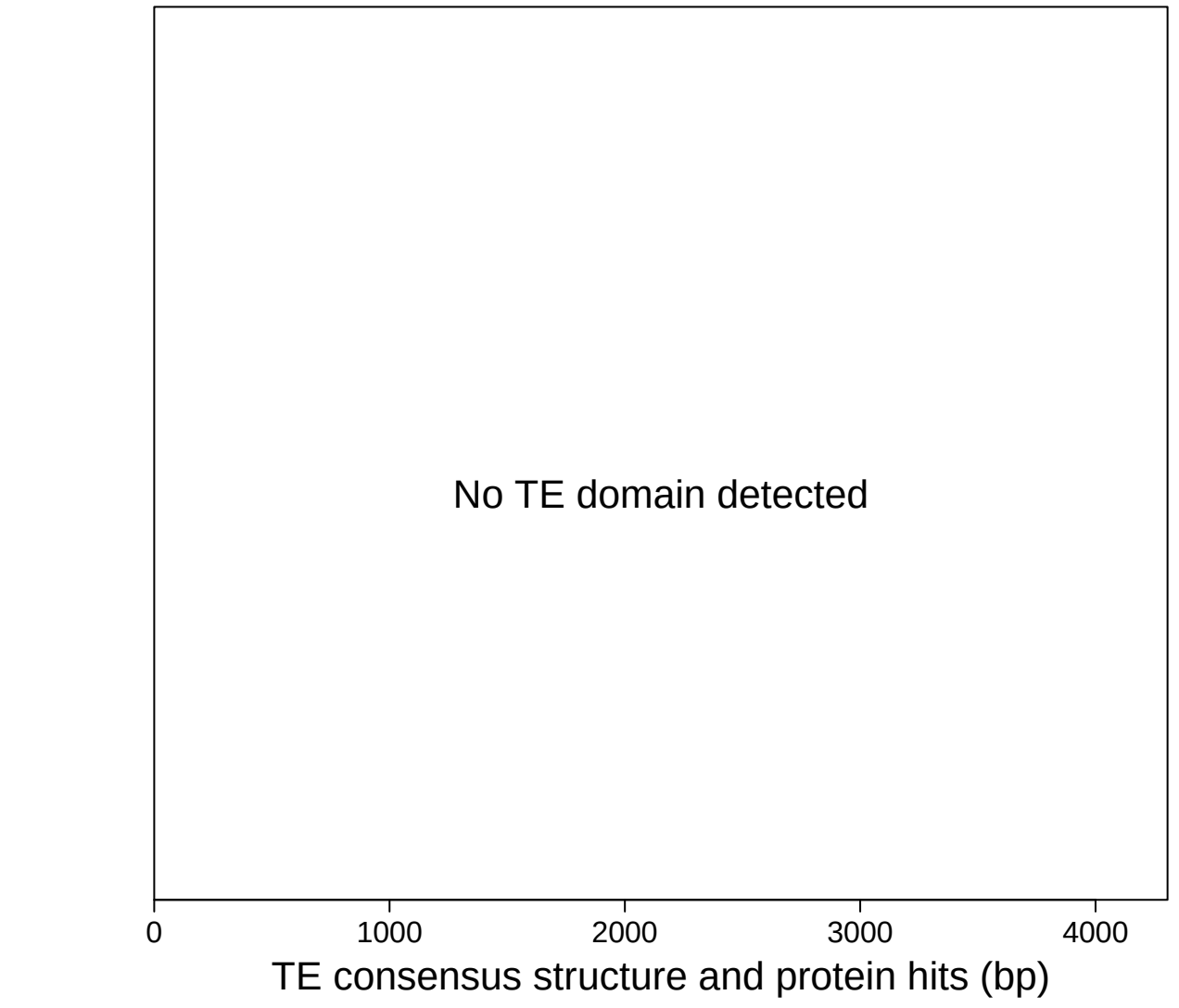
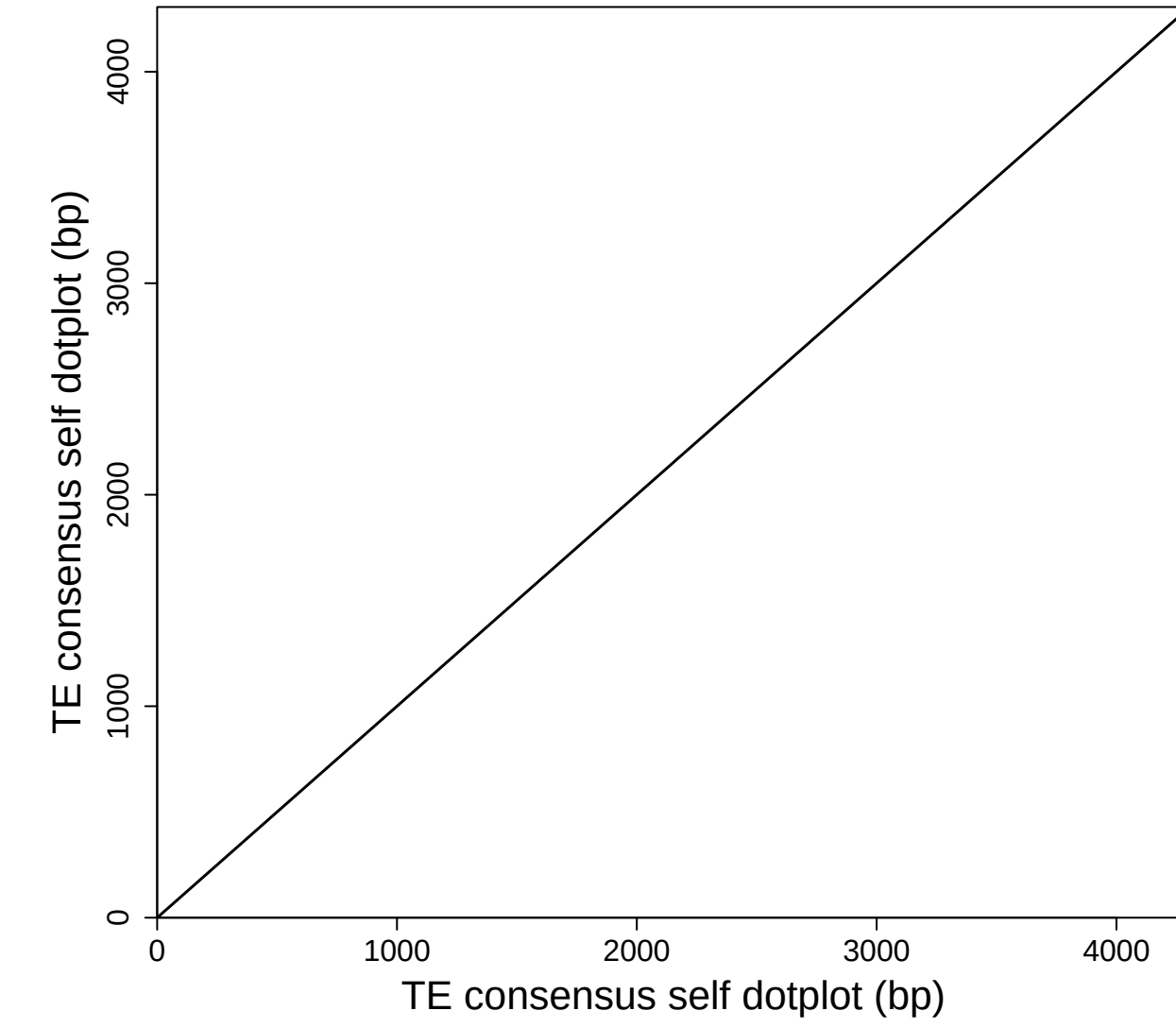
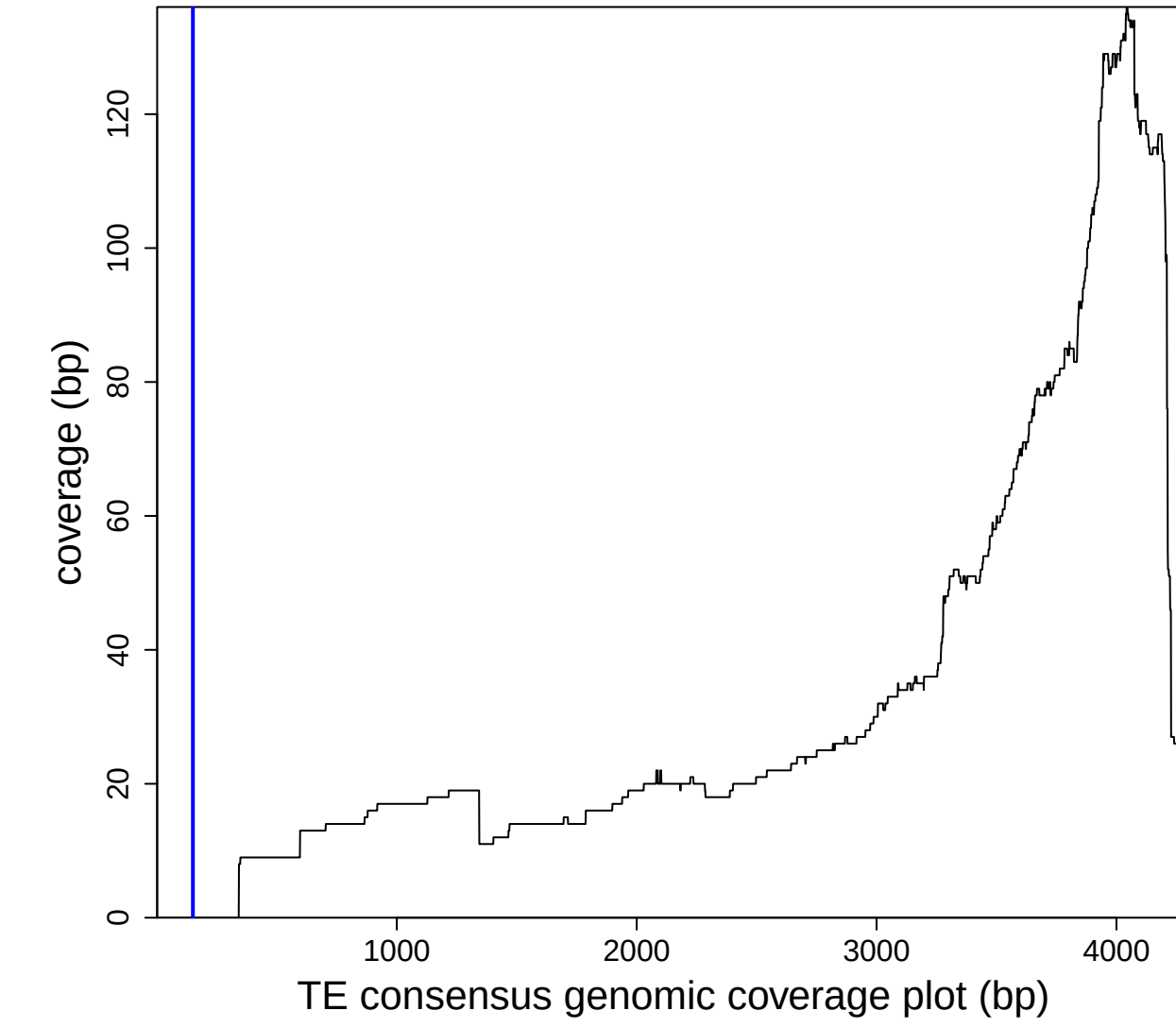
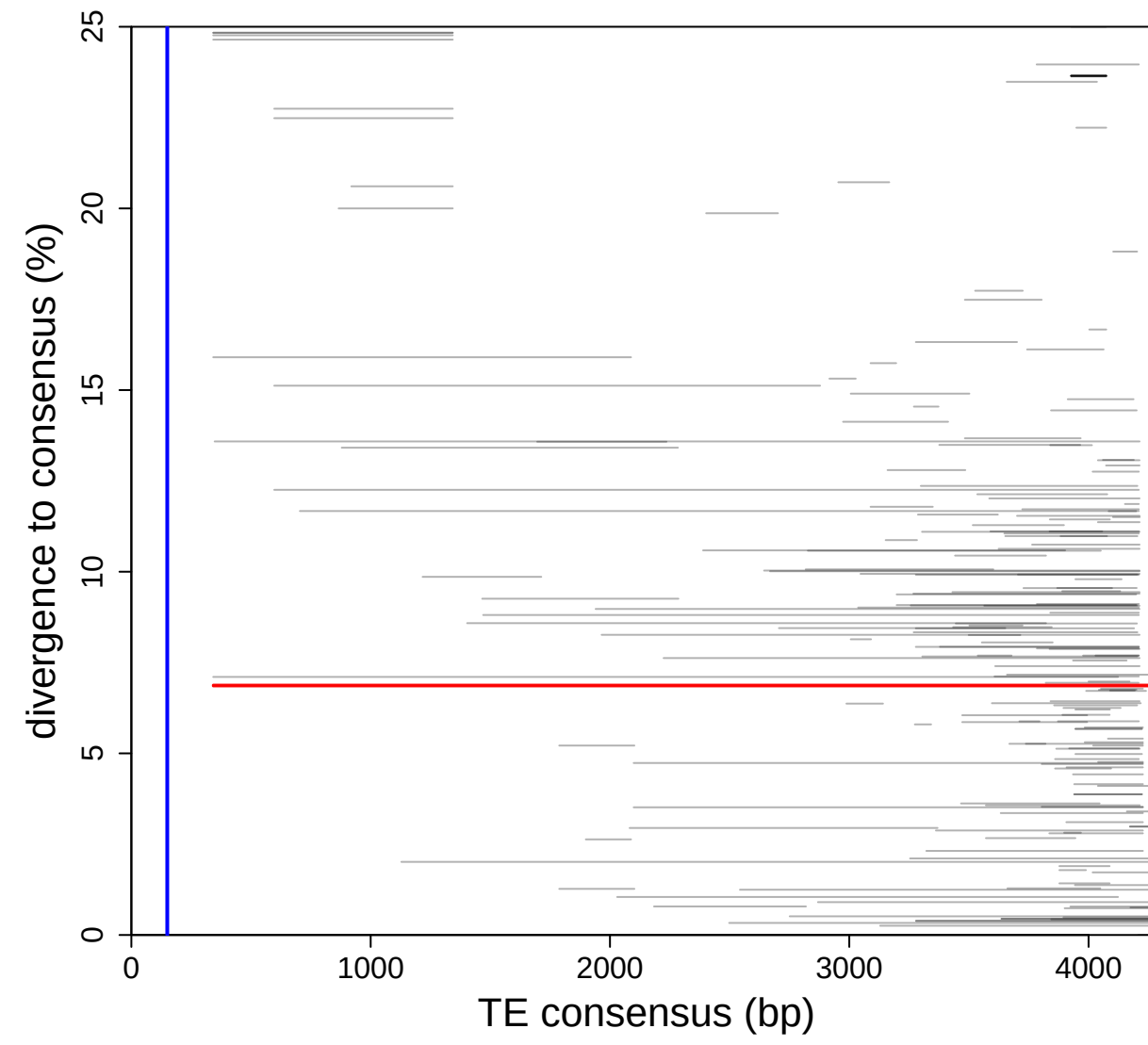
ed_uf.bed_g_4.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa_ce
size: 4026bp; fragments: 219; full length: 4 (>=3623.4bp)

After TEtrimmer 4026 bp

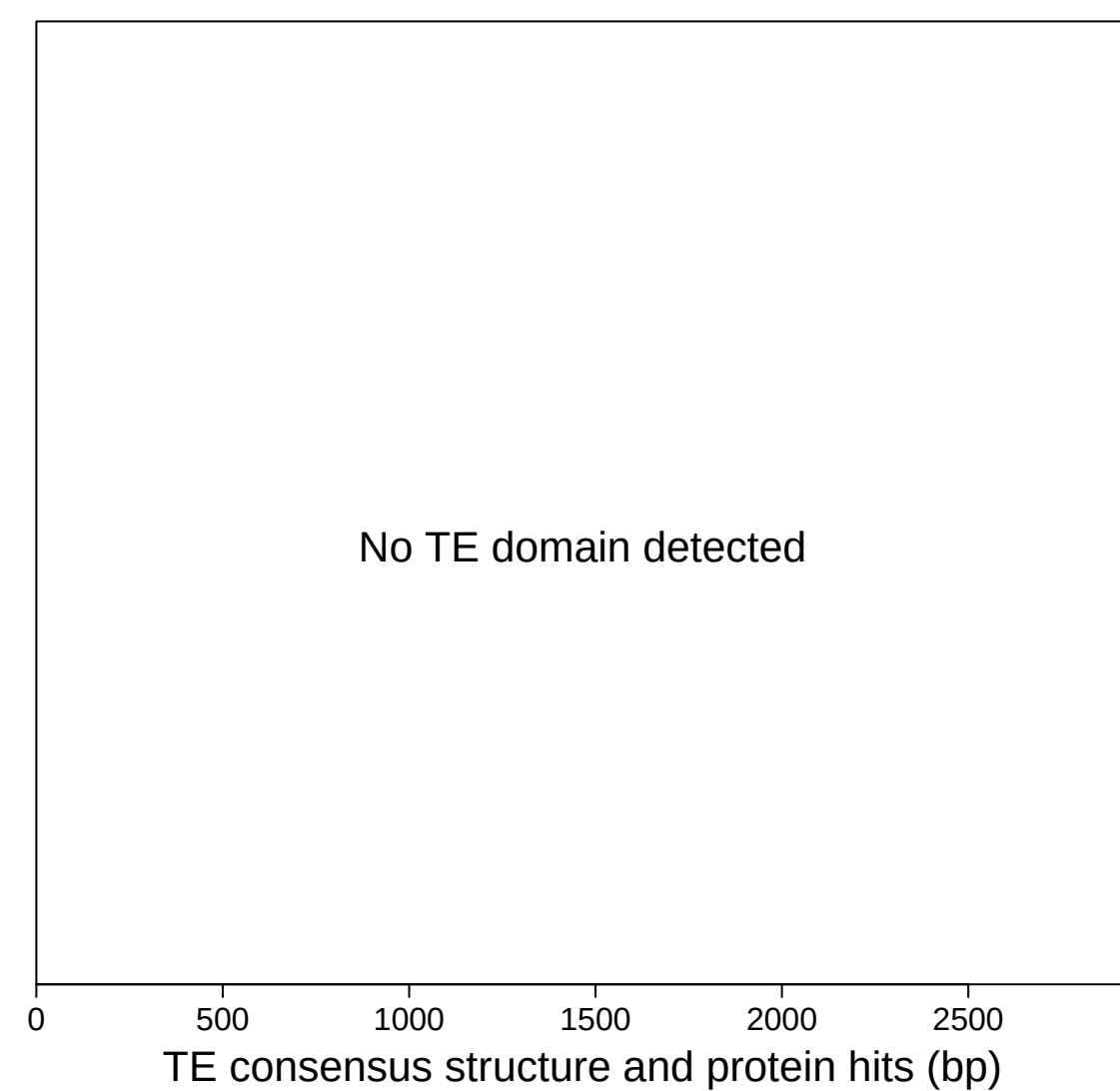
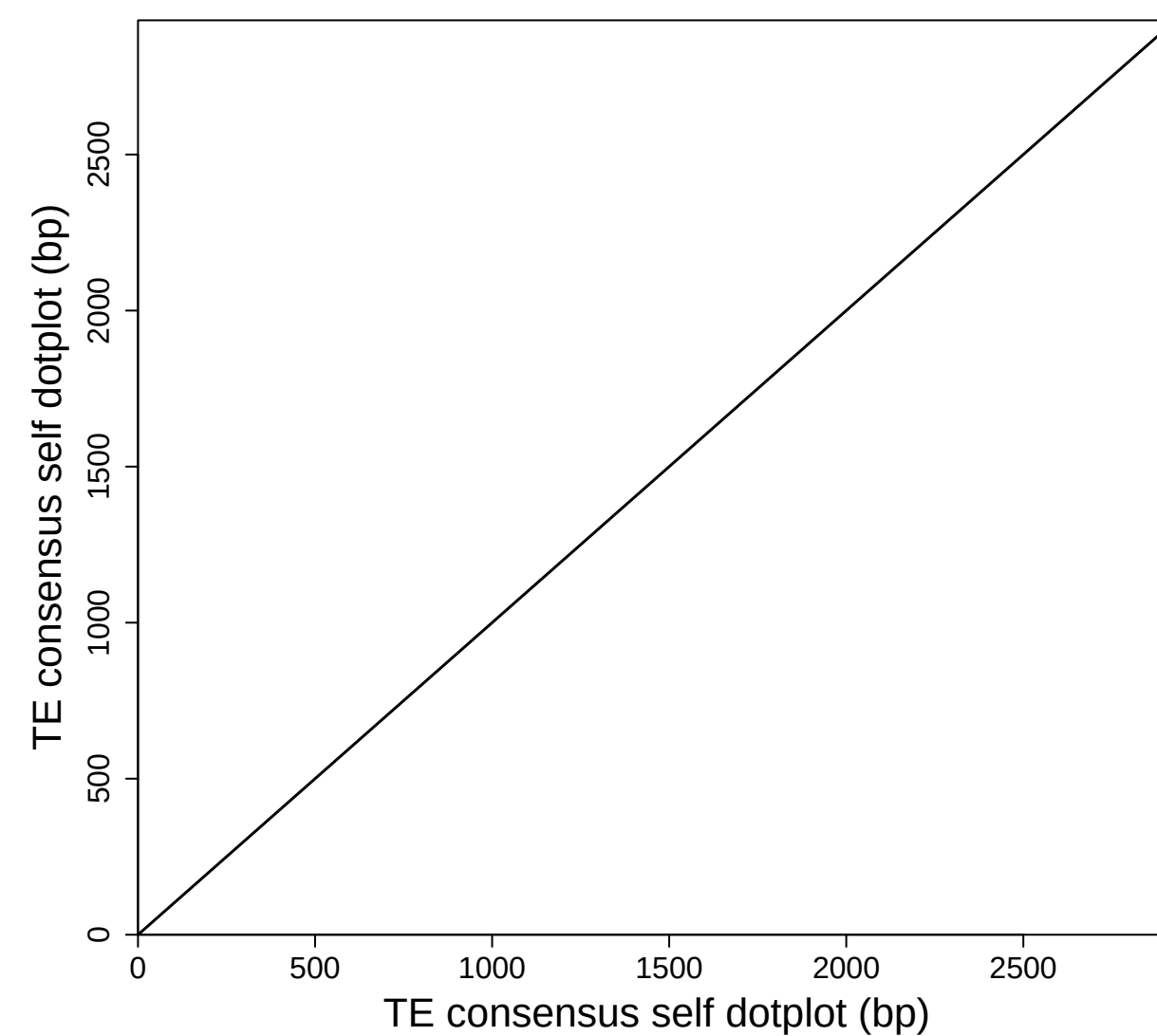
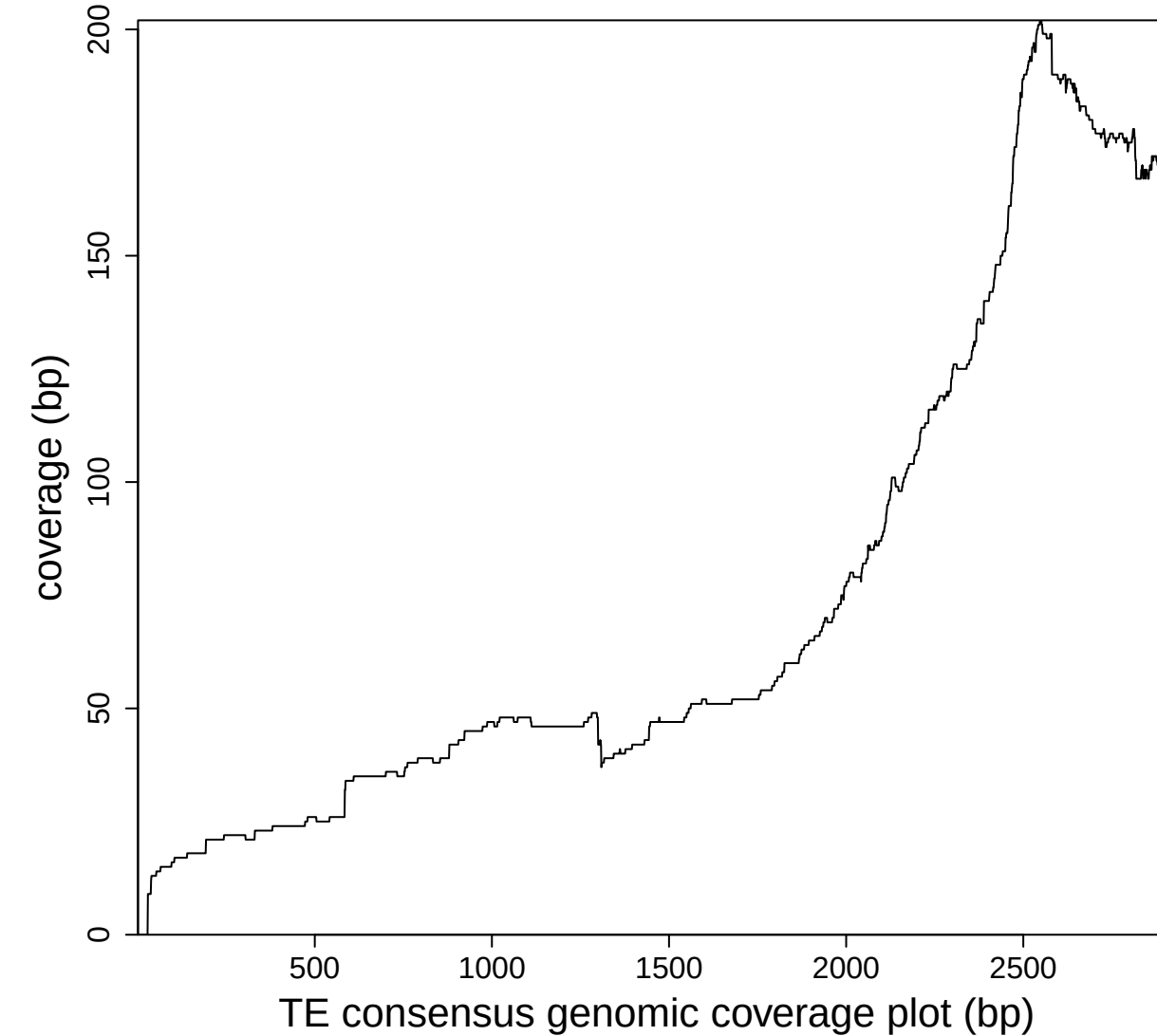
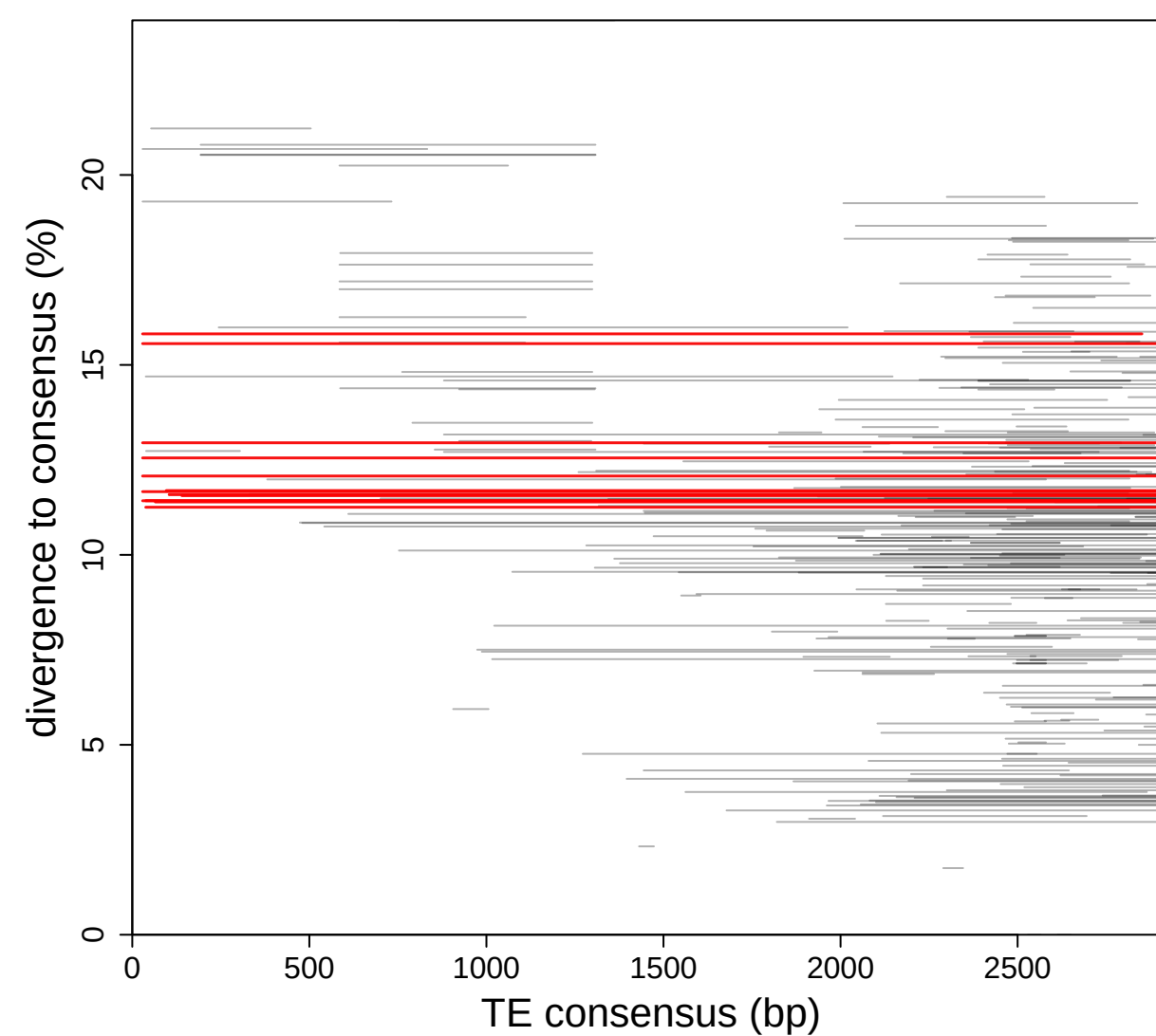


3.fasta.b.bed uf.bed g 4.bed fm 1.bed 0 0 bcln.fa aln.fa cl.fa
size: 4306bp; fragments: 210; full length: 2 (>=3875.4bp)

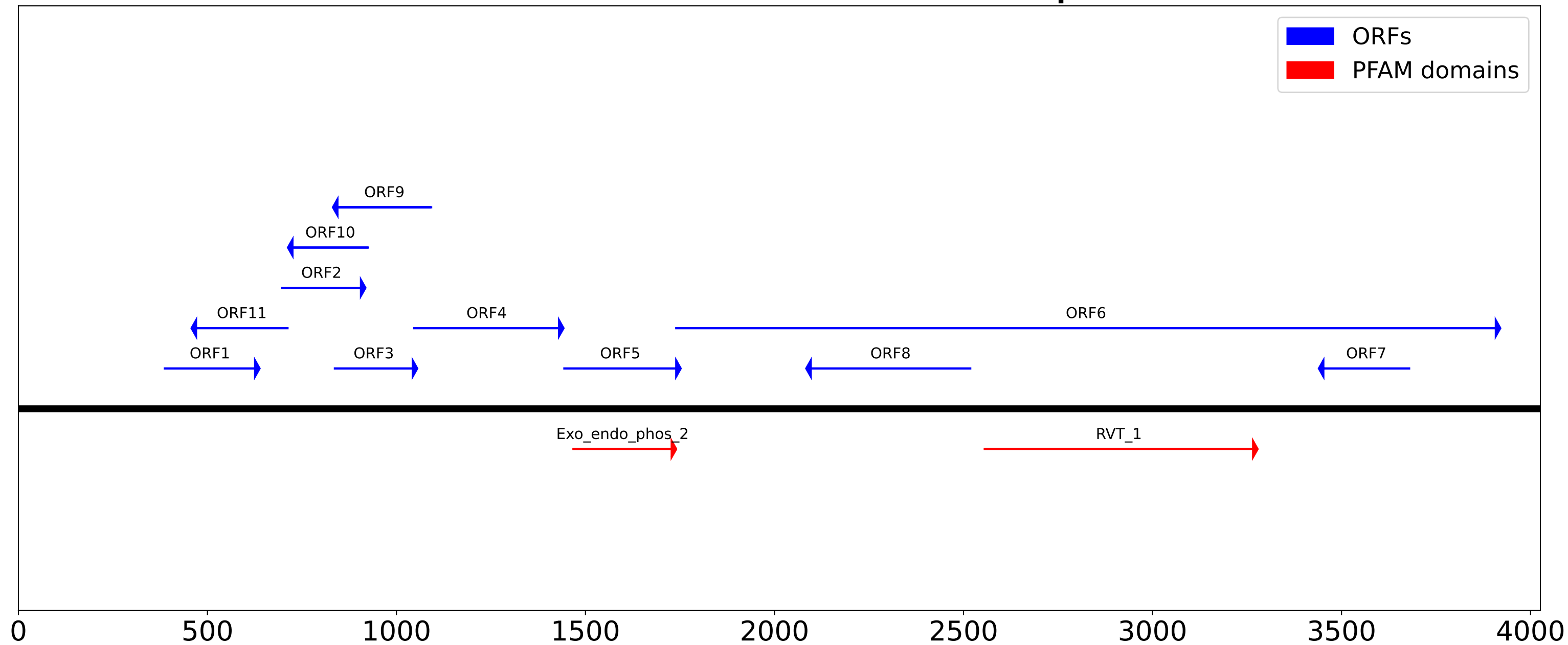
After TEtrimmer Extended plot Blue lines are boundaries



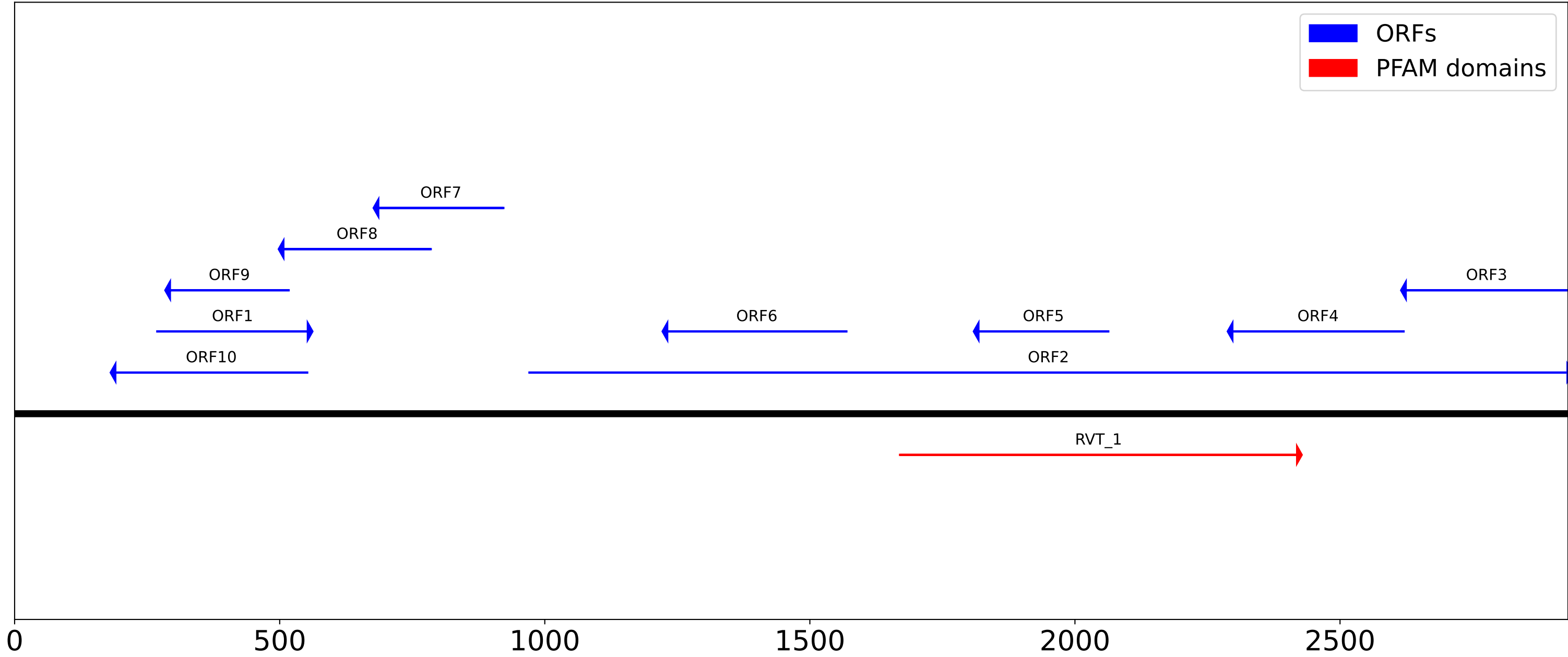
TE: rnd_5_family_383
size: 2930bp; fragments: 296; full length: 13 (>=2637bp)



After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

