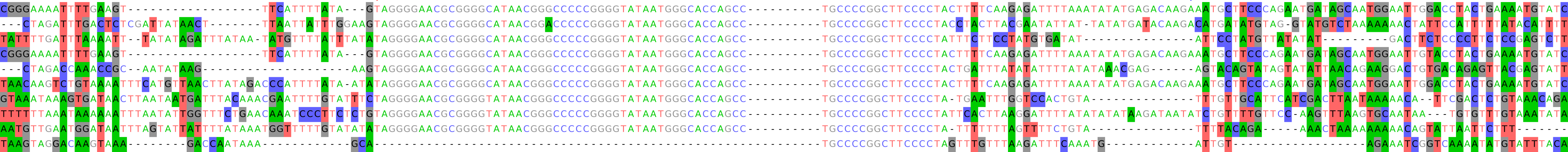


Start Crop Point

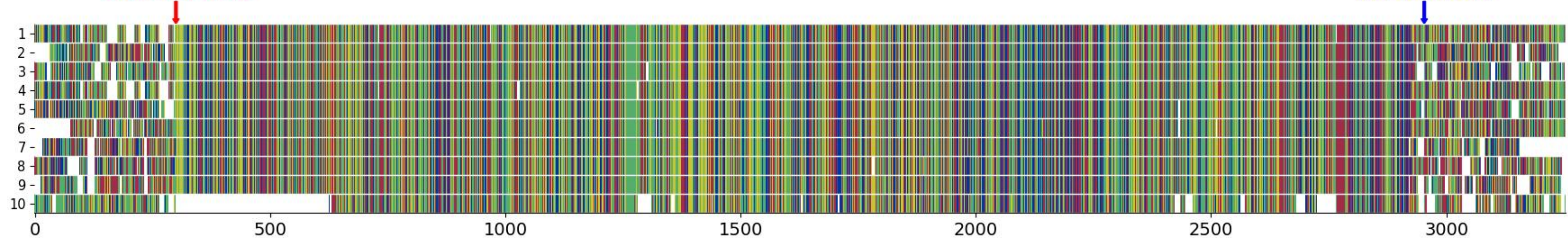
End Crop Point

MSA length = 2654



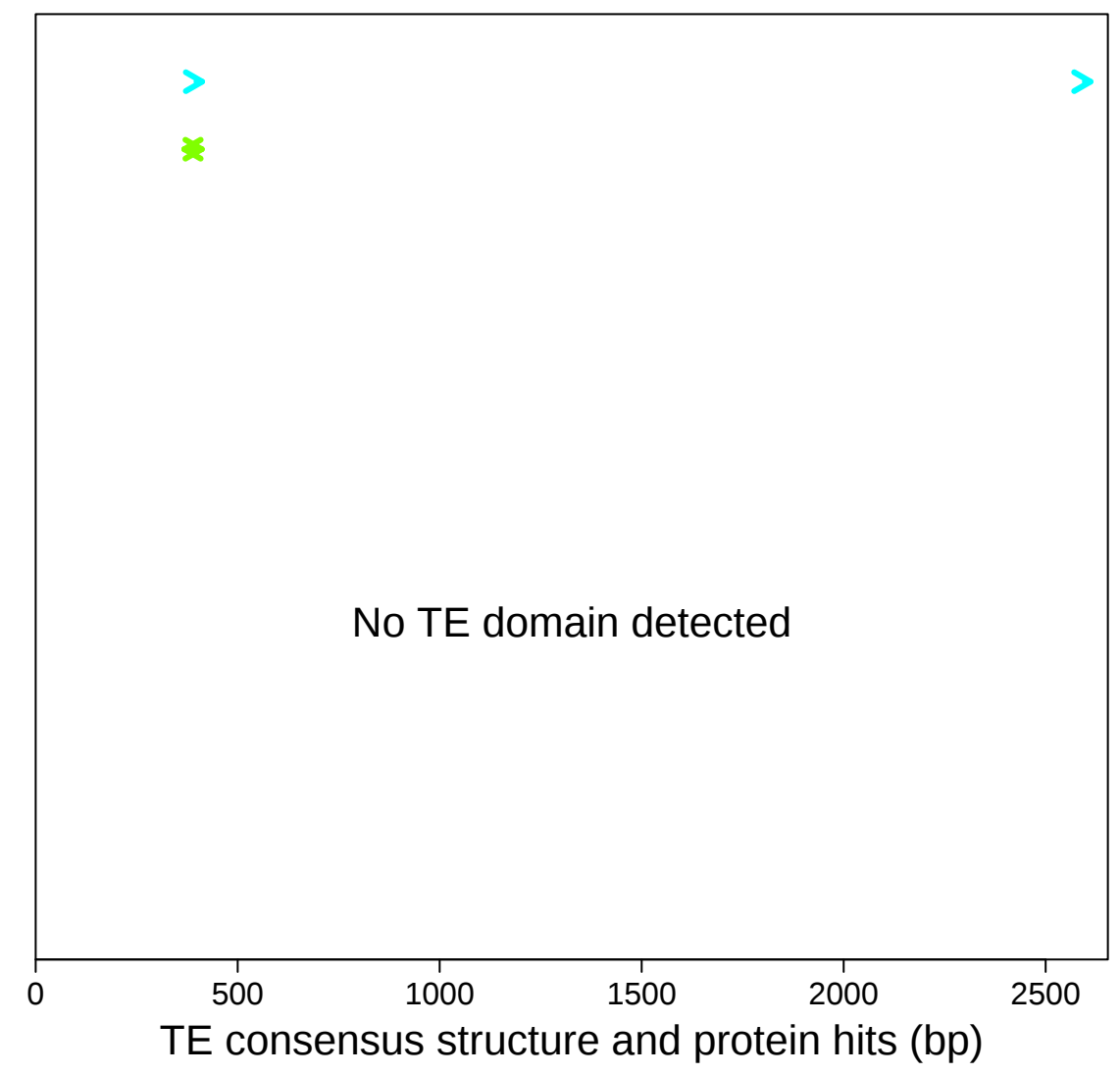
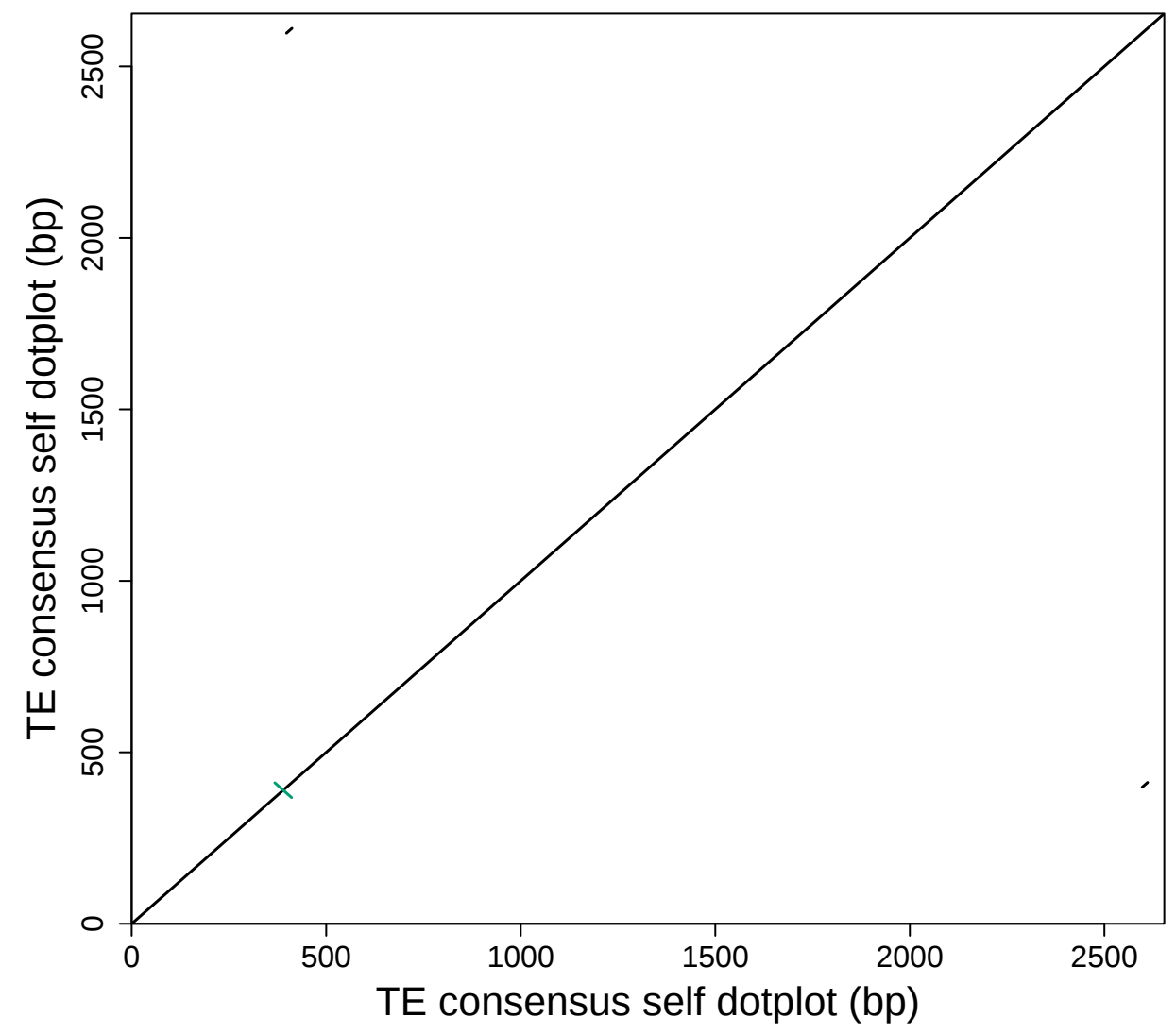
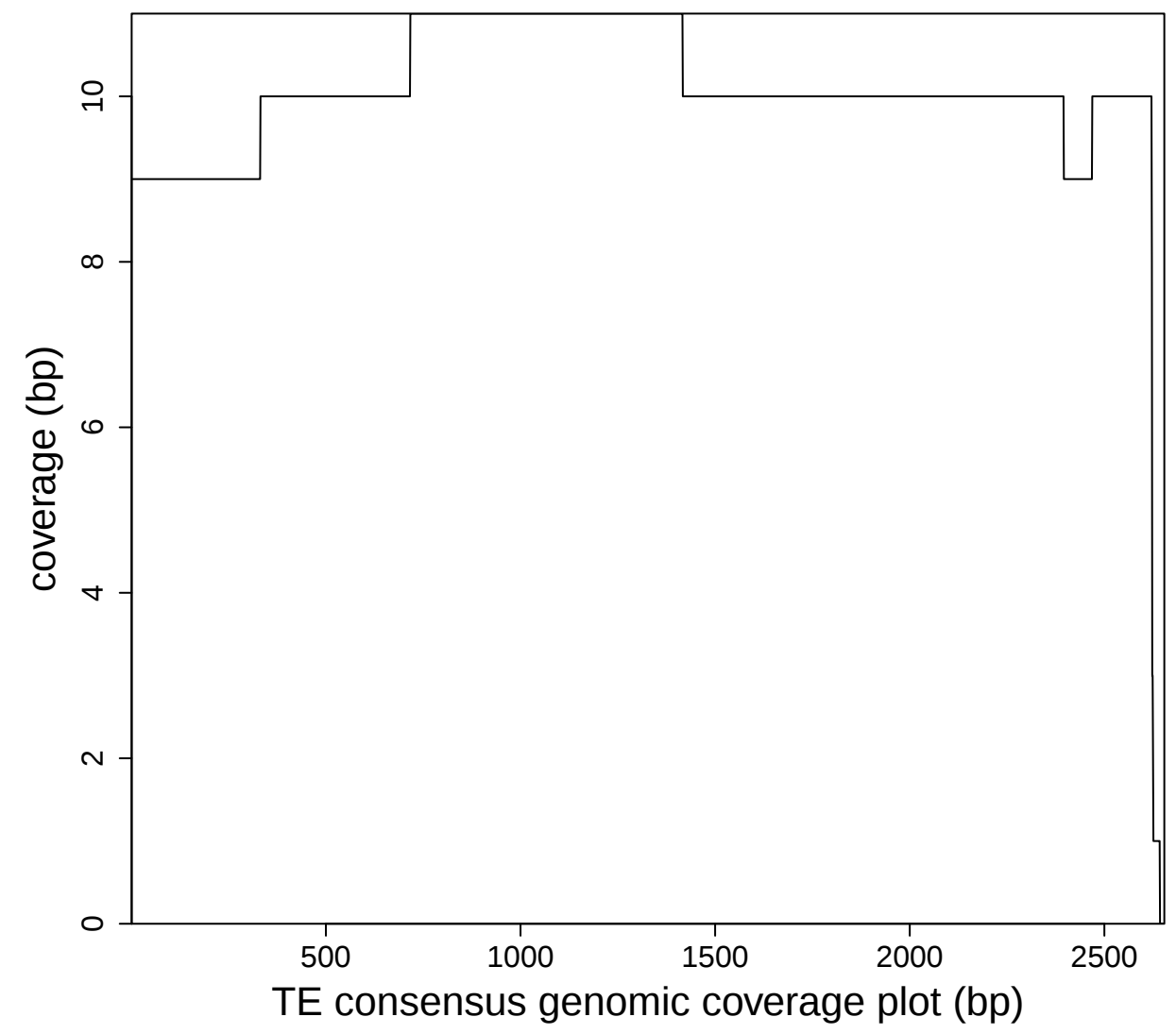
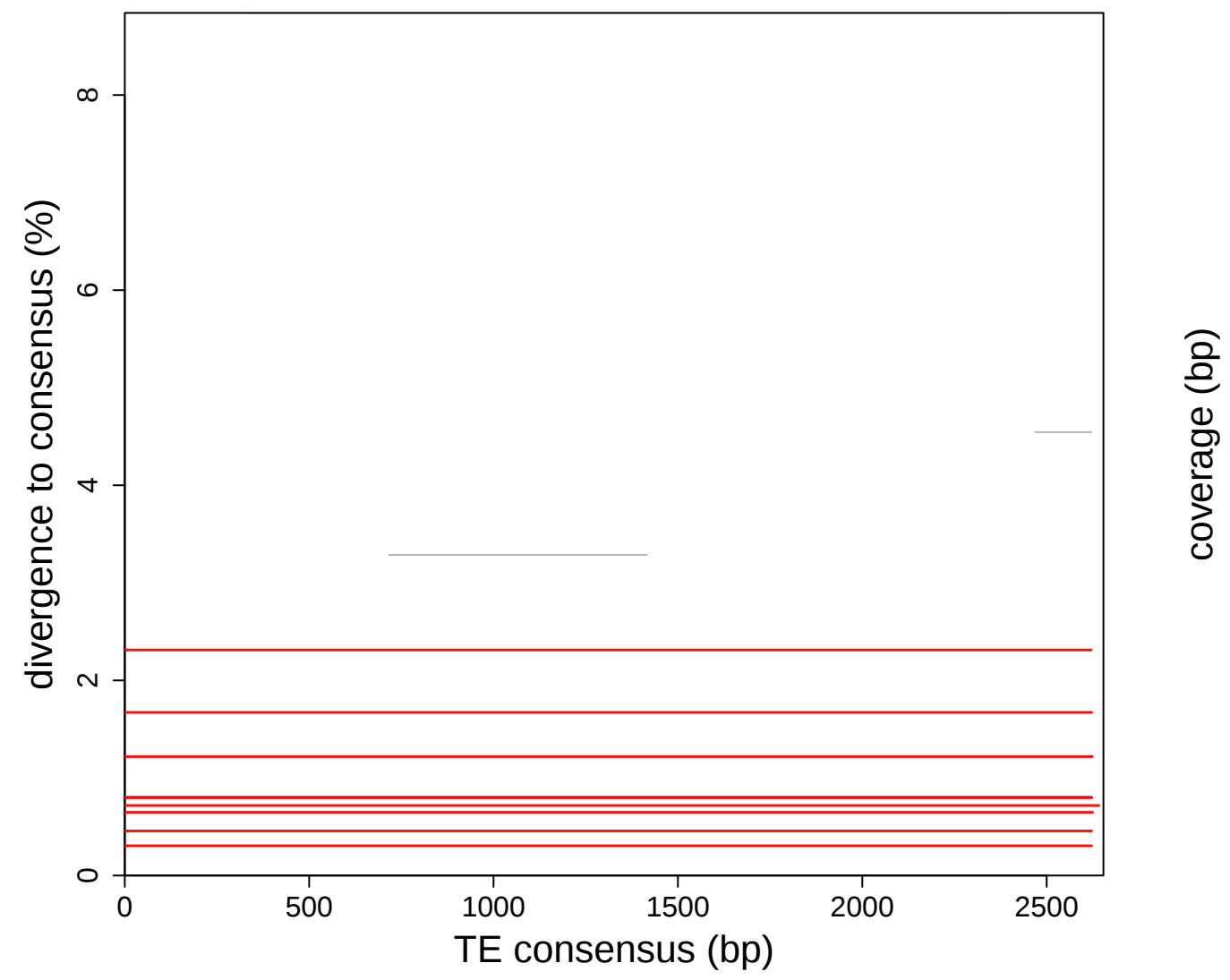
Start crop Point

End crop Point



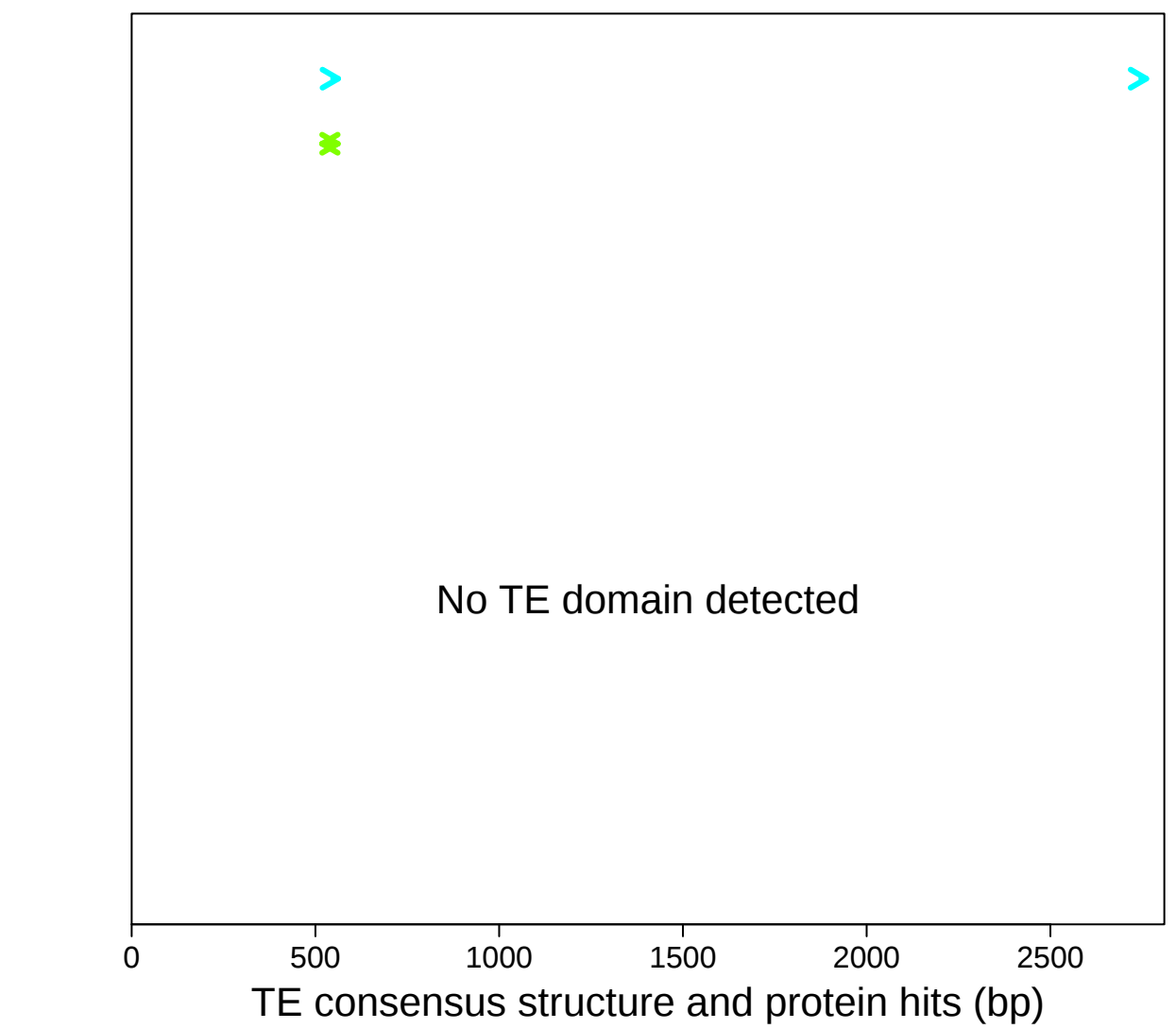
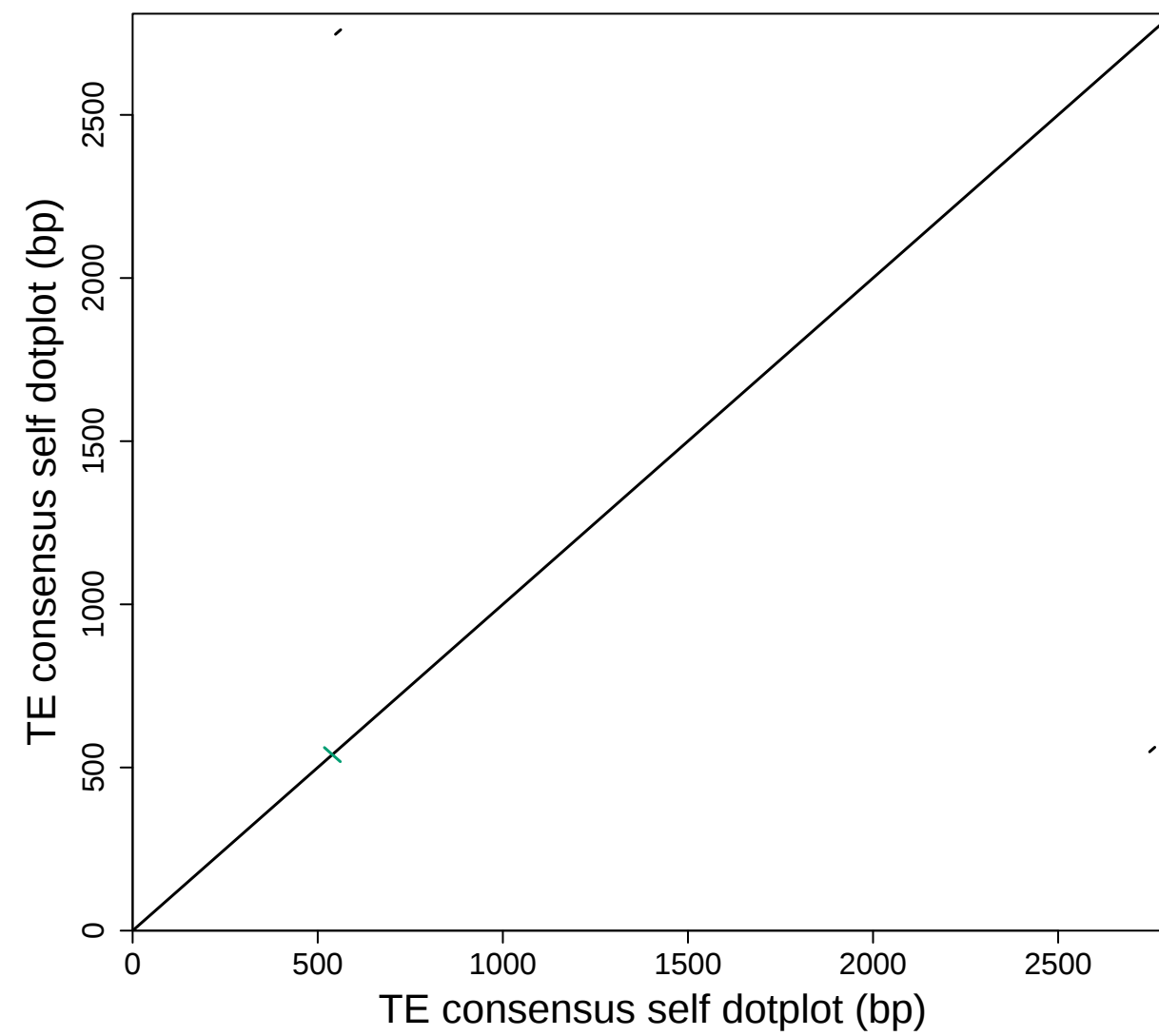
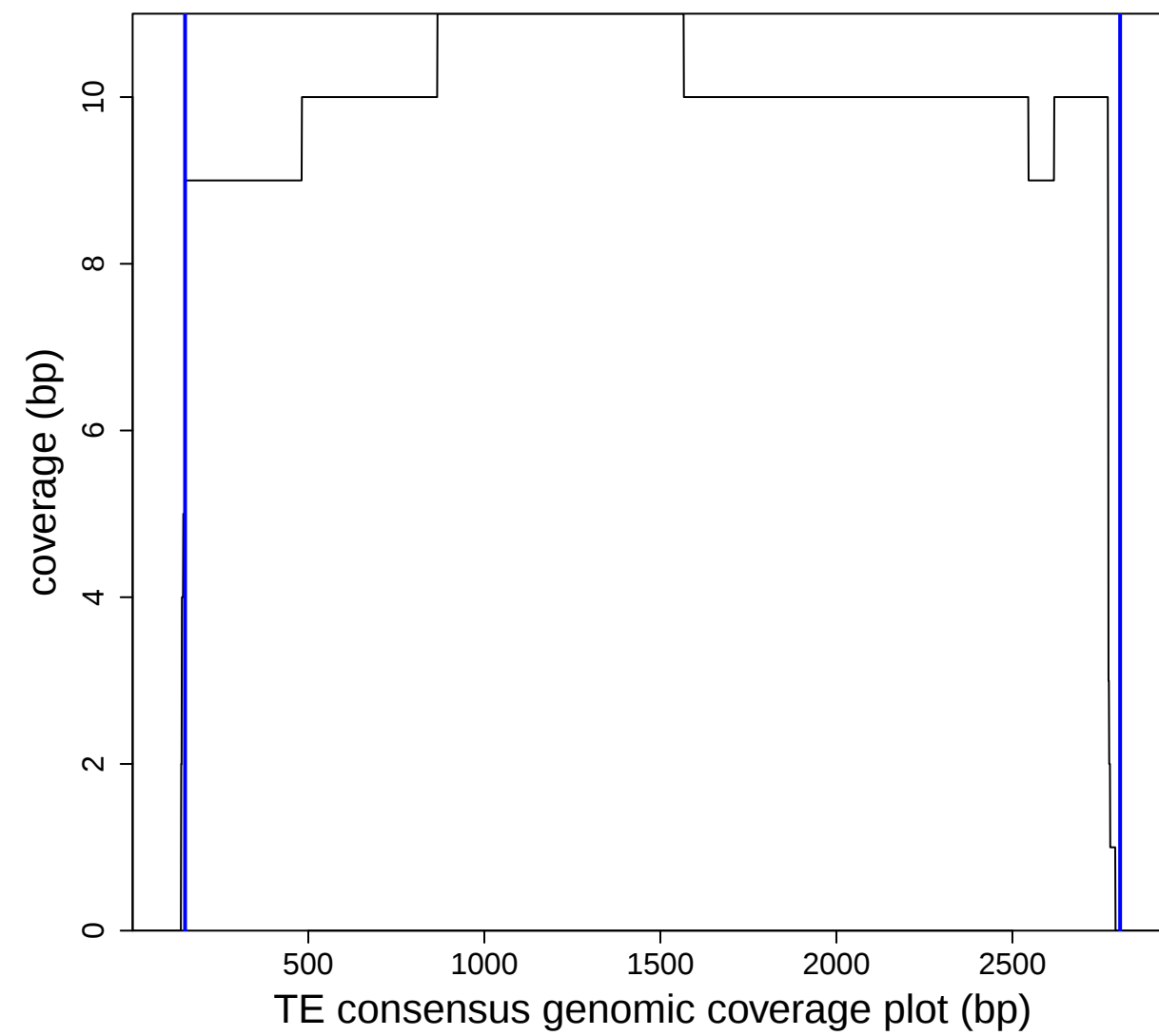
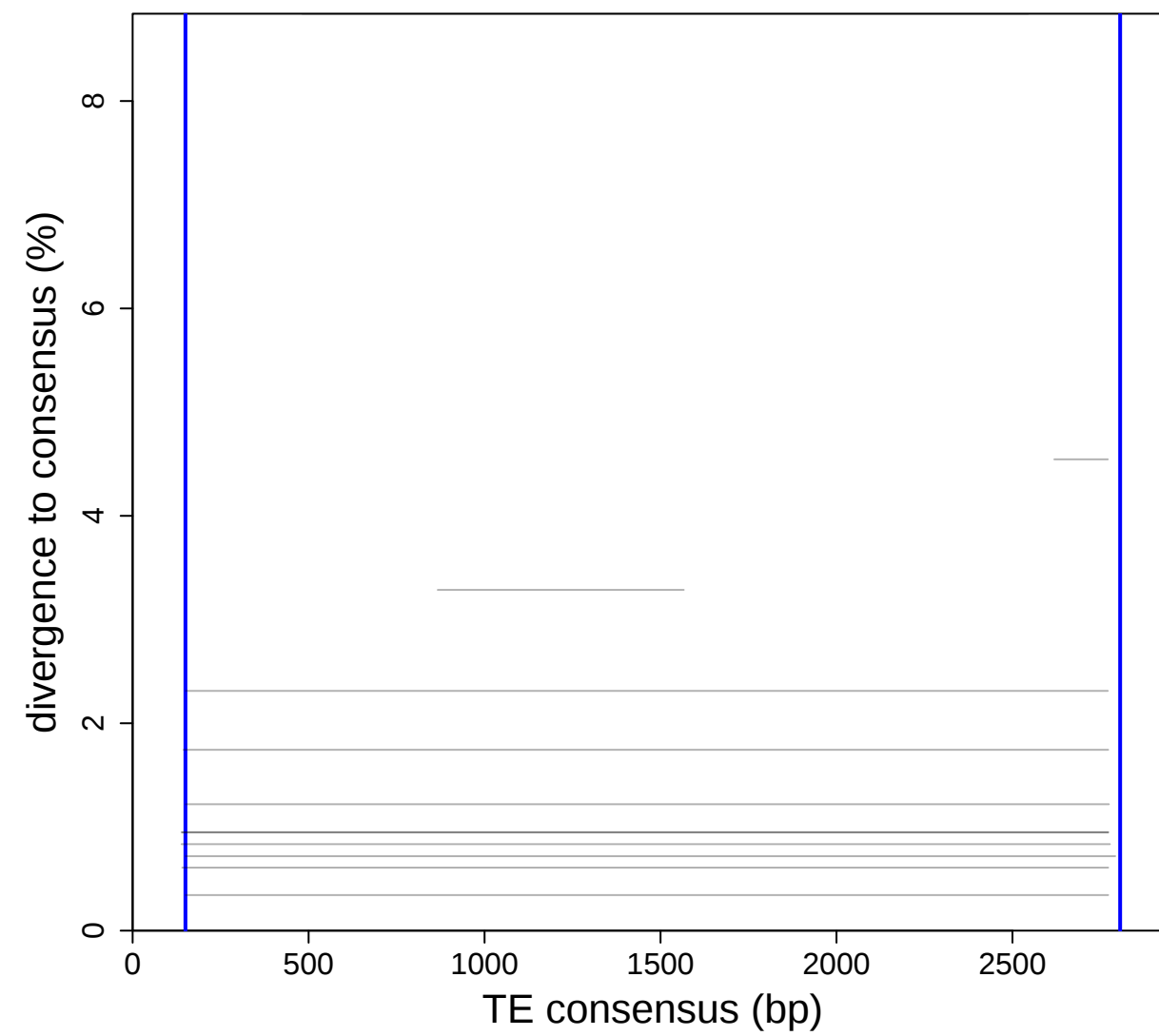
b.bed uf.bed g_1.bed fm_1.bed 0_0_bcln.fa aln.fa cl.fa gs.fa
size: 2654bp; fragments: 12; full length: 9 (>=2388.6bp)

After TEtrimmer 2654 bp



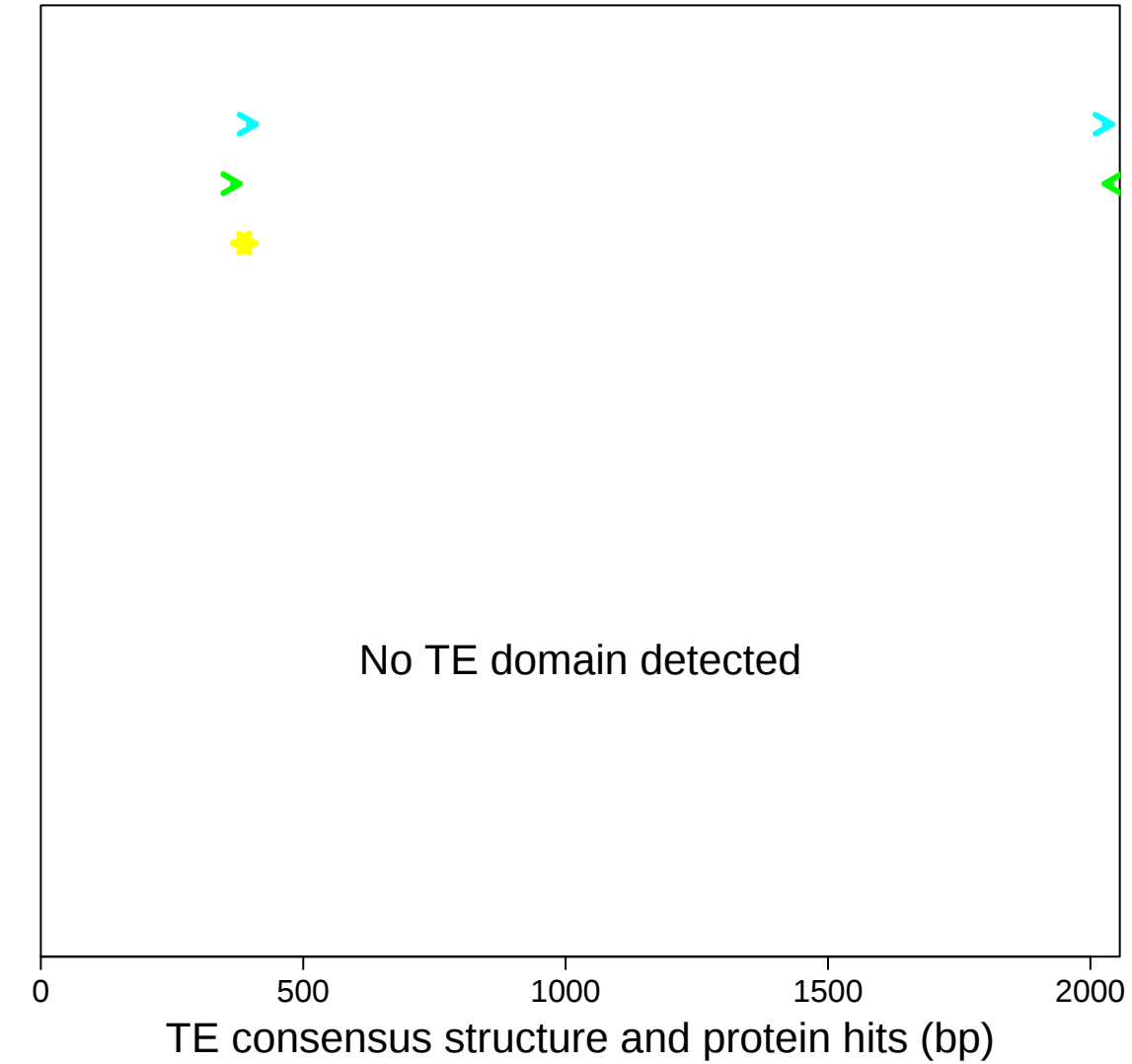
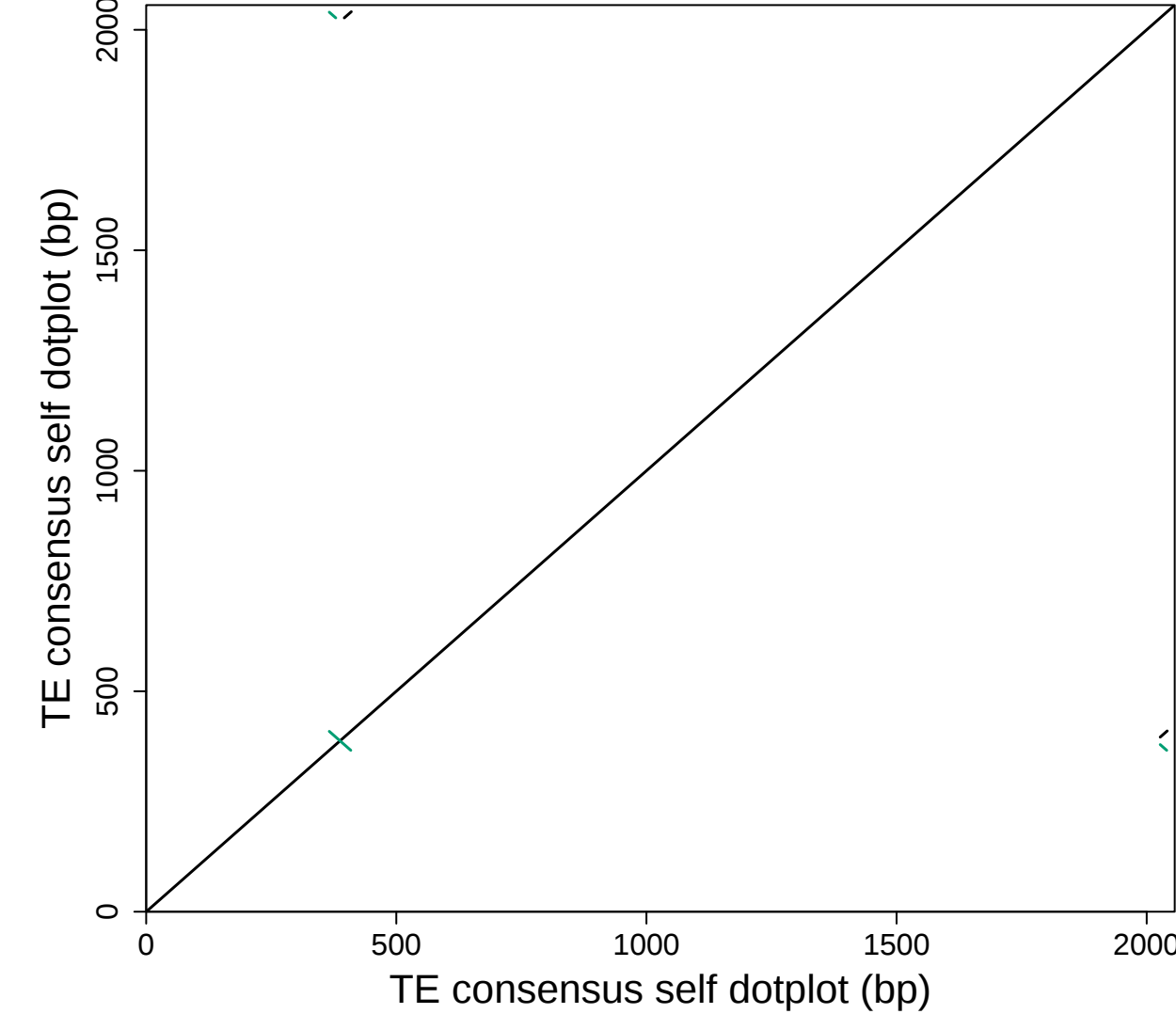
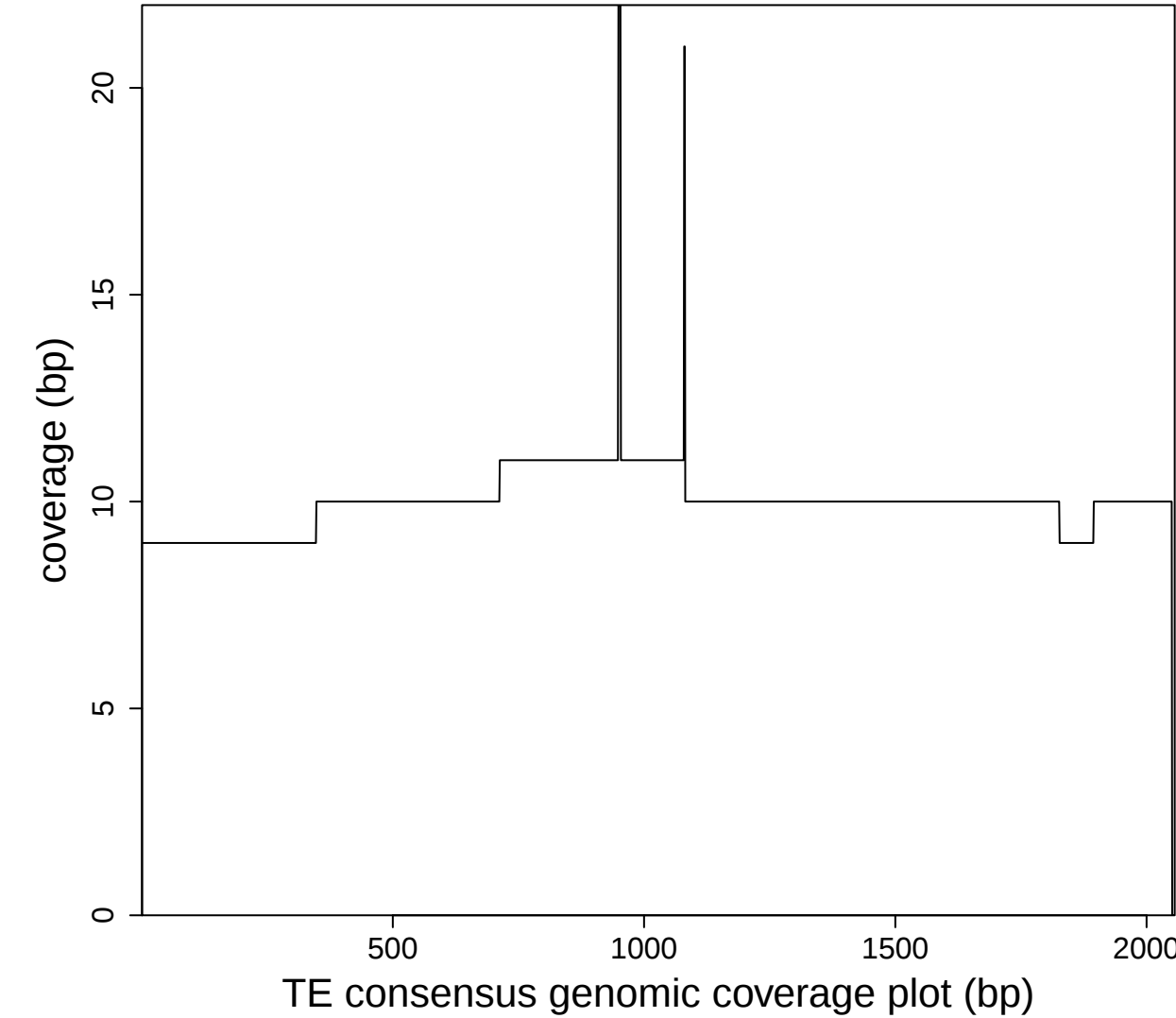
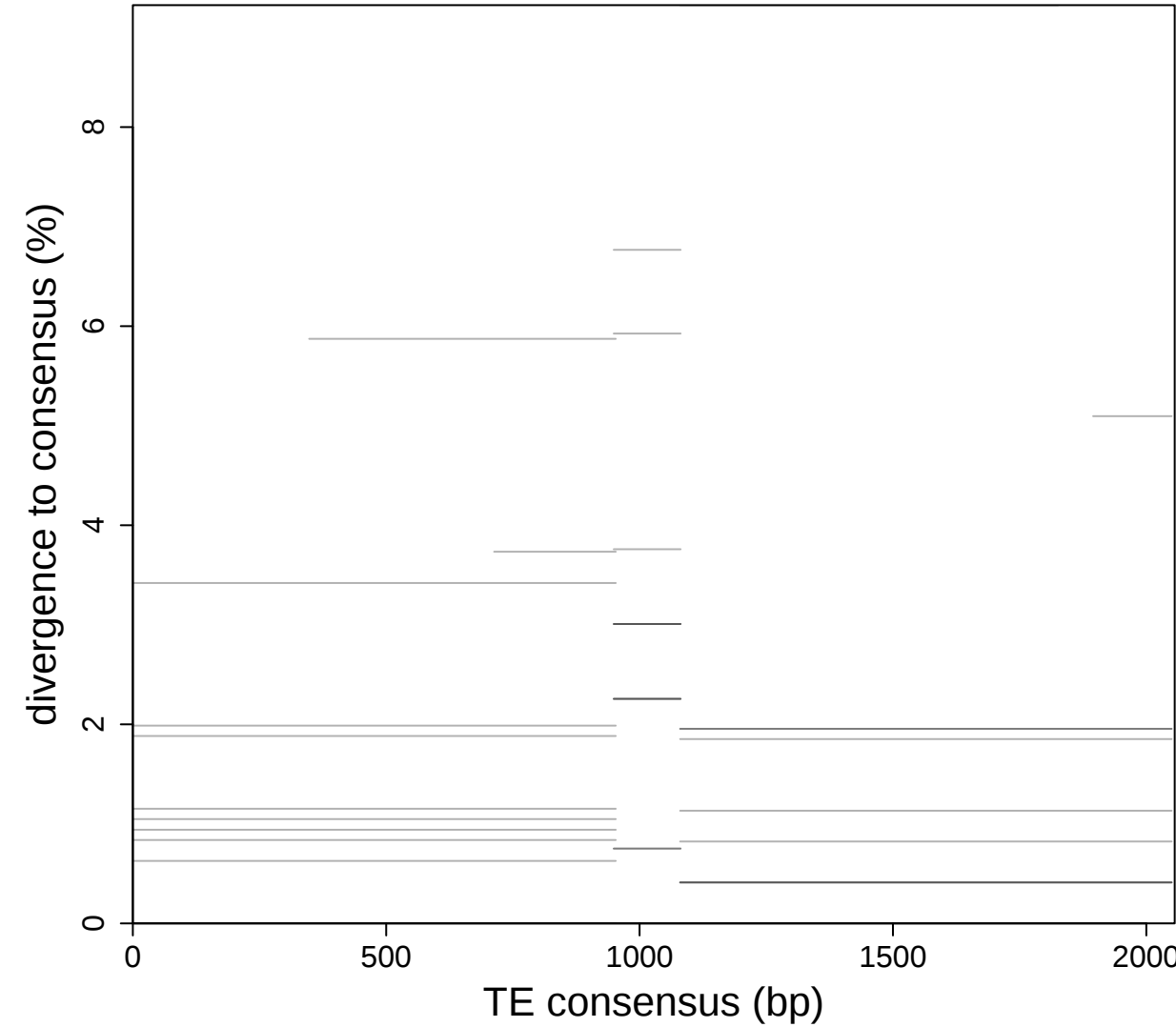
3.fasta.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa
size: 2956bp; fragments: 12; full length: 0 (>=2660.4bp)

After TEtrimmer Extended plot Blue lines are boundaries

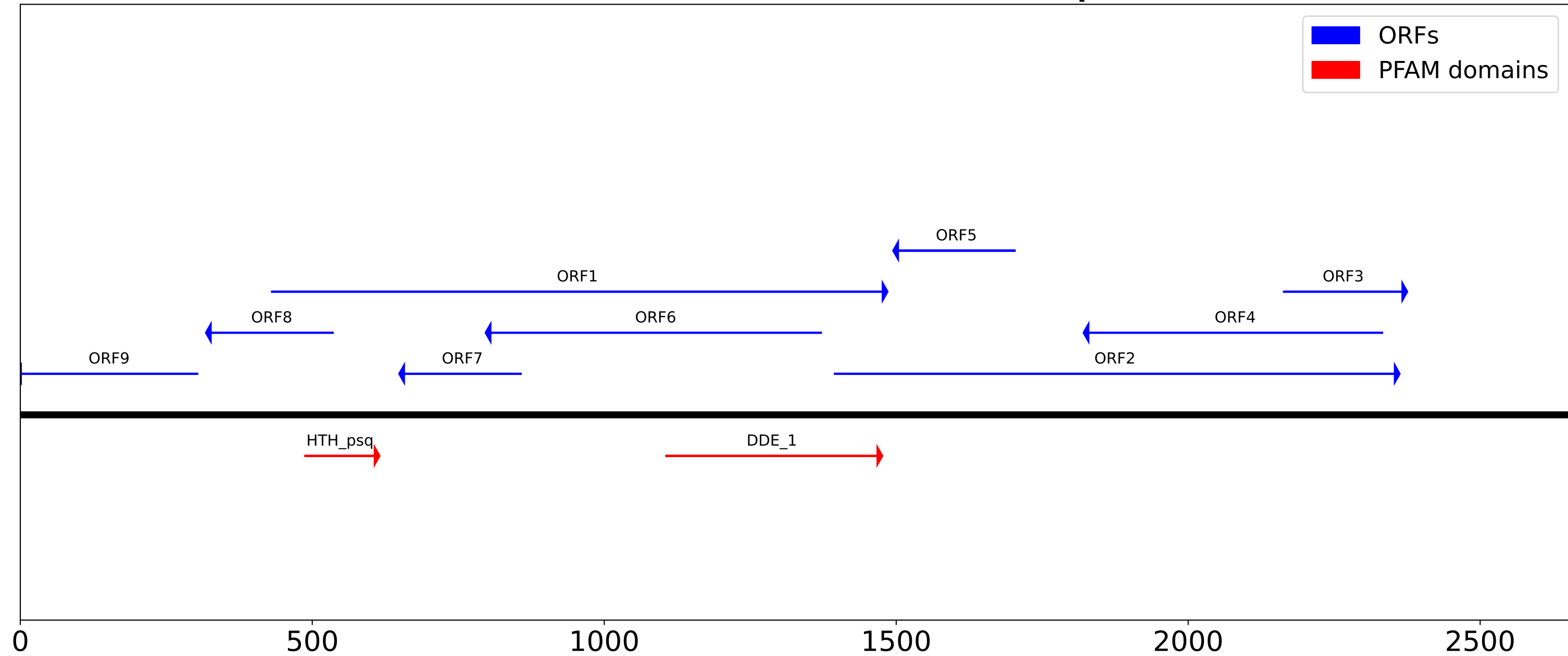


TE: ltr_1_family_963
size: 2056bp; fragments: 33; full length: 0 (≥ 1850.4 bp)

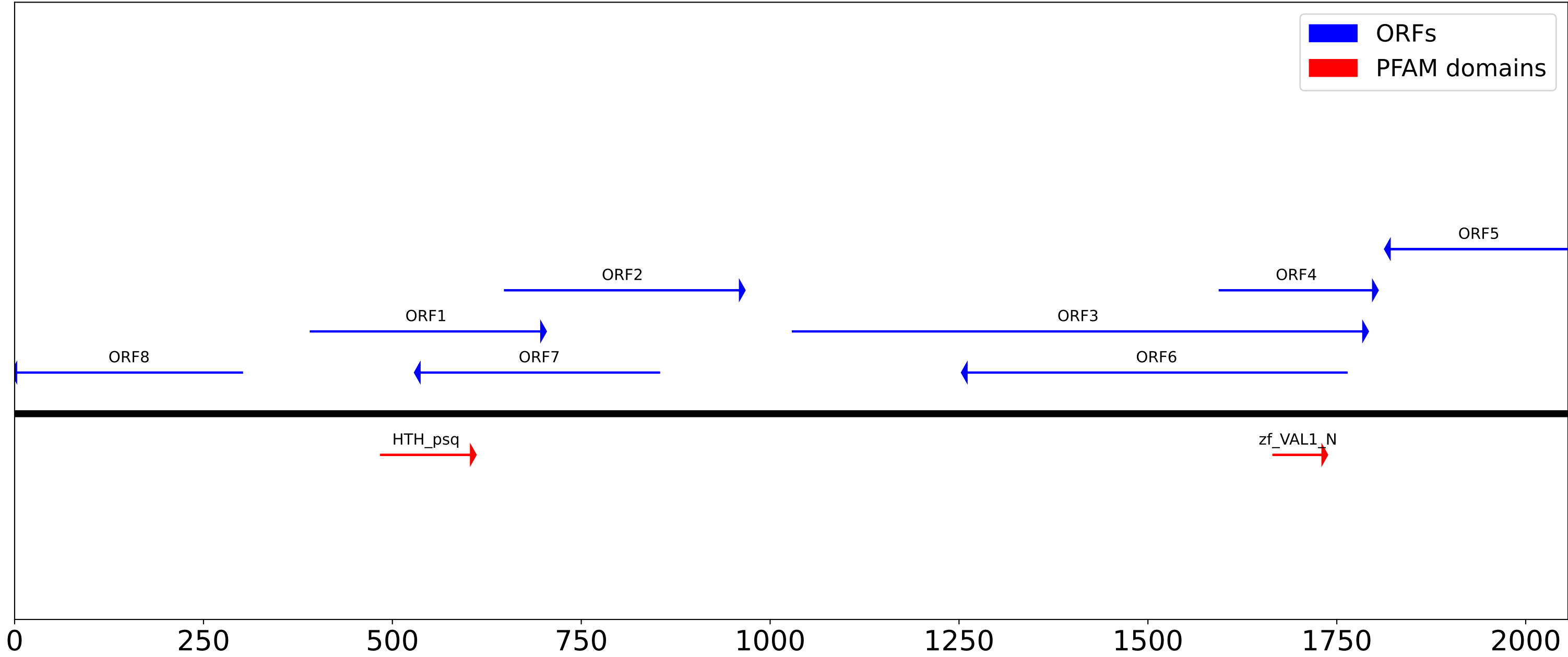
Before TEtrimmer 2056 bp



After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

