

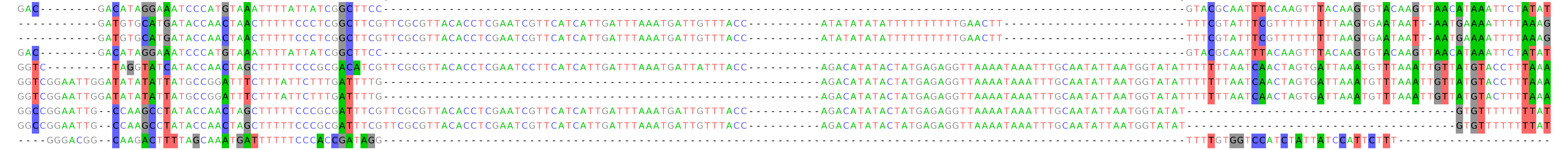
Start Crop Point



End Crop Point

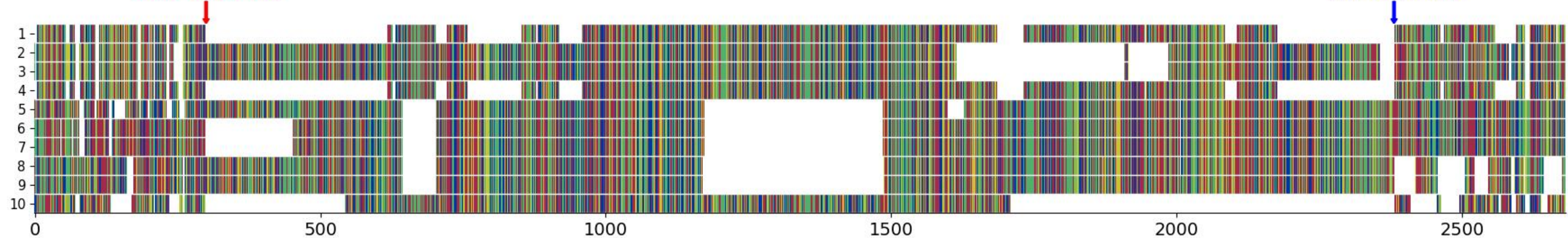


MSA length = 2082



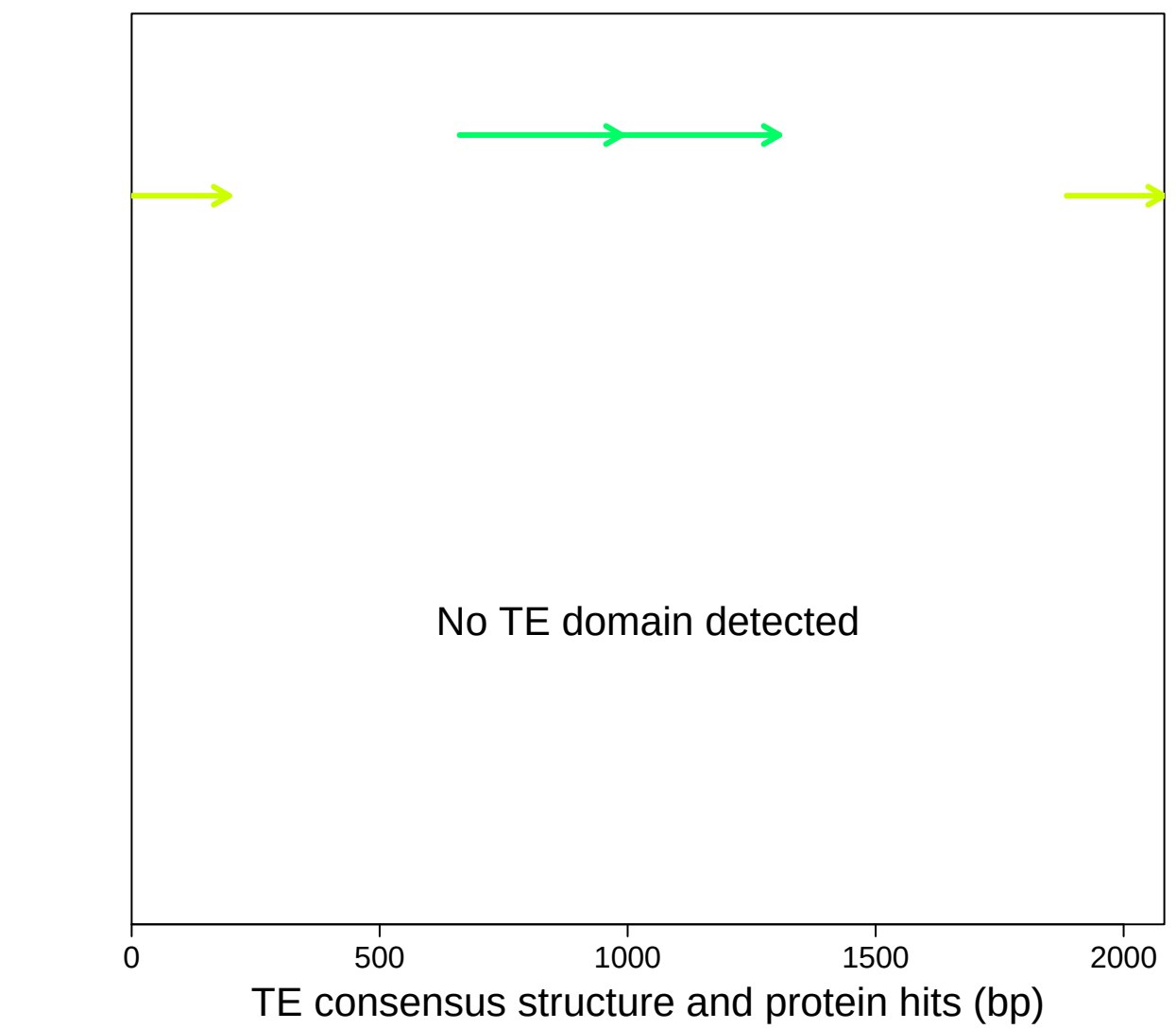
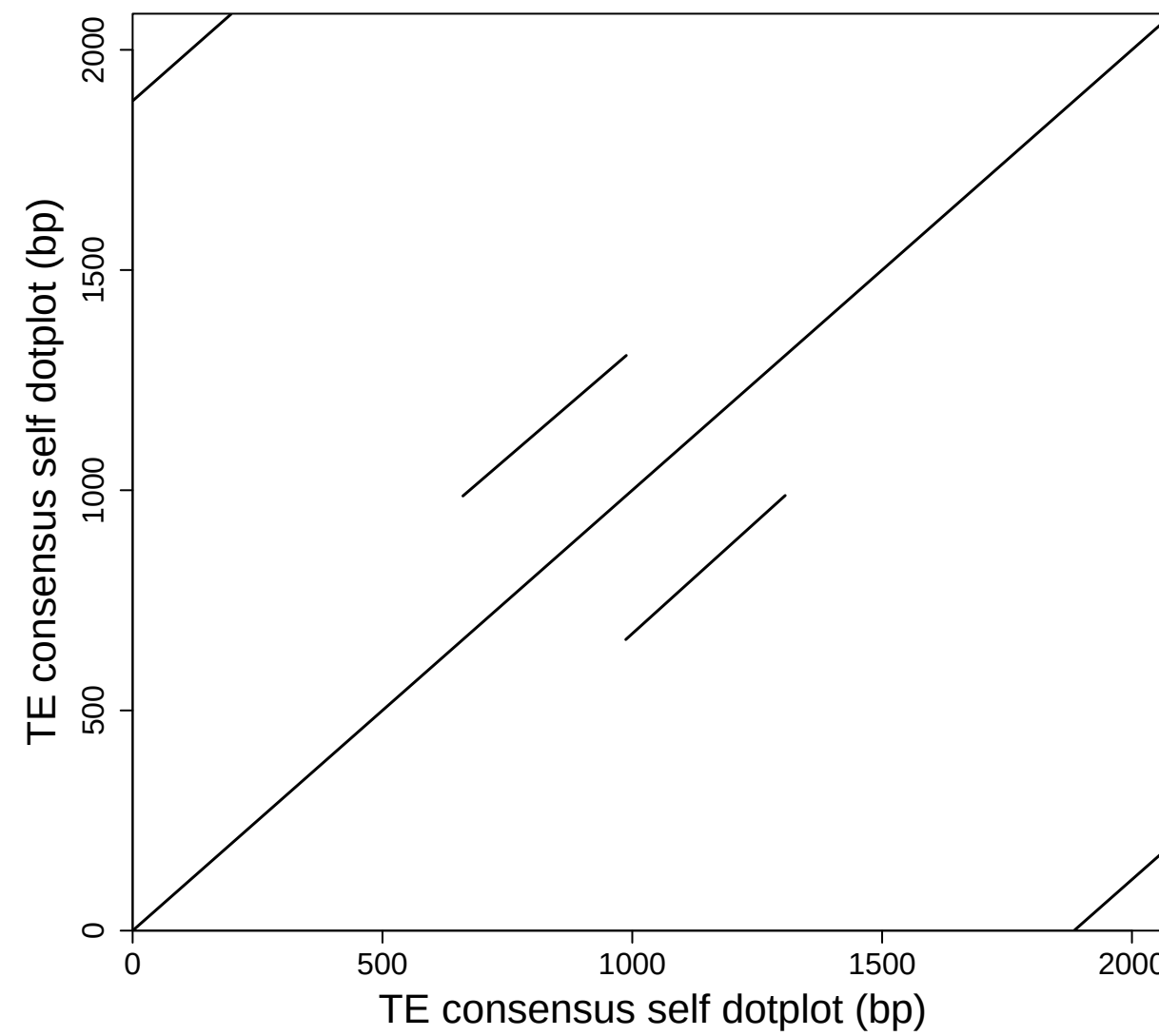
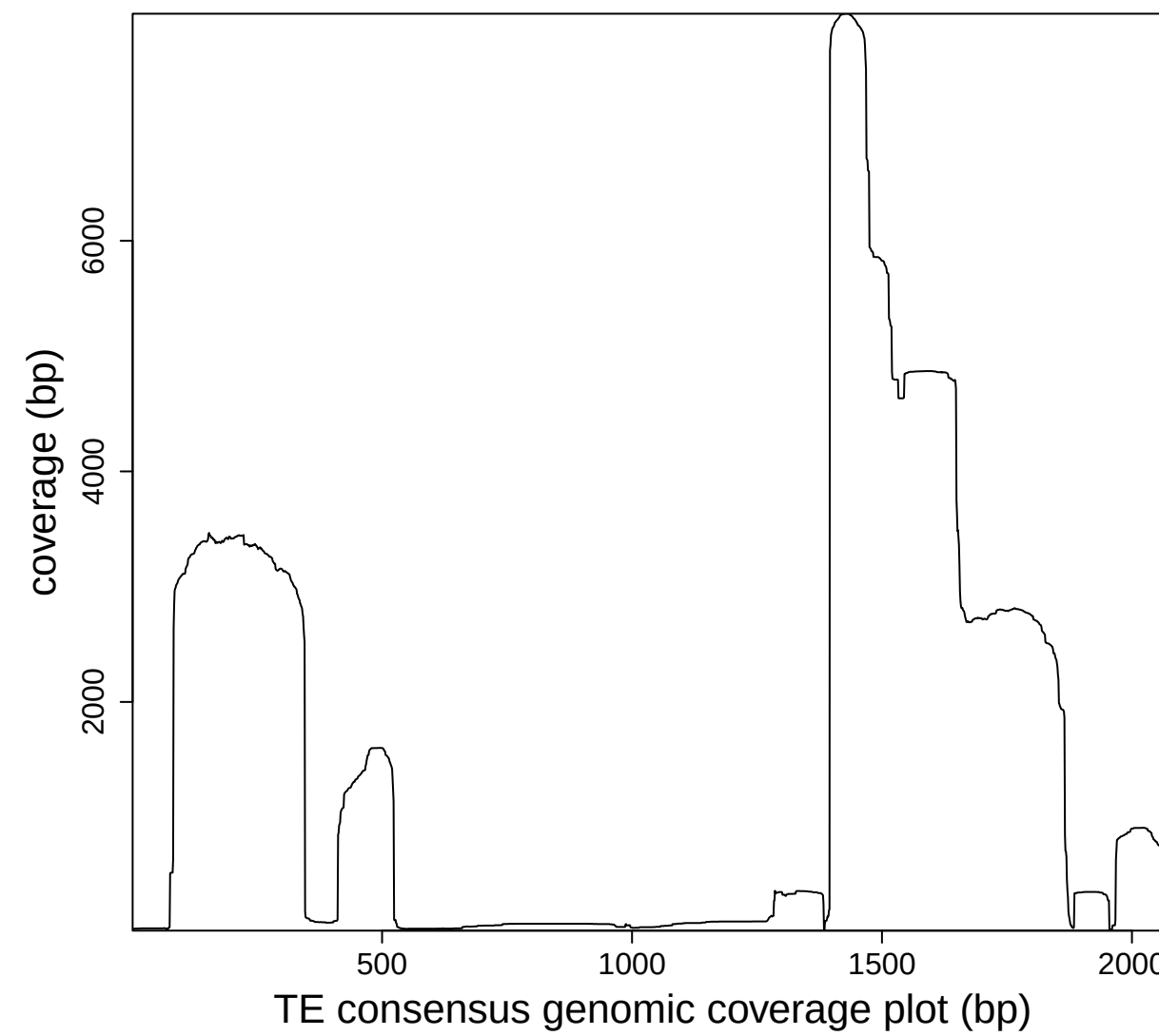
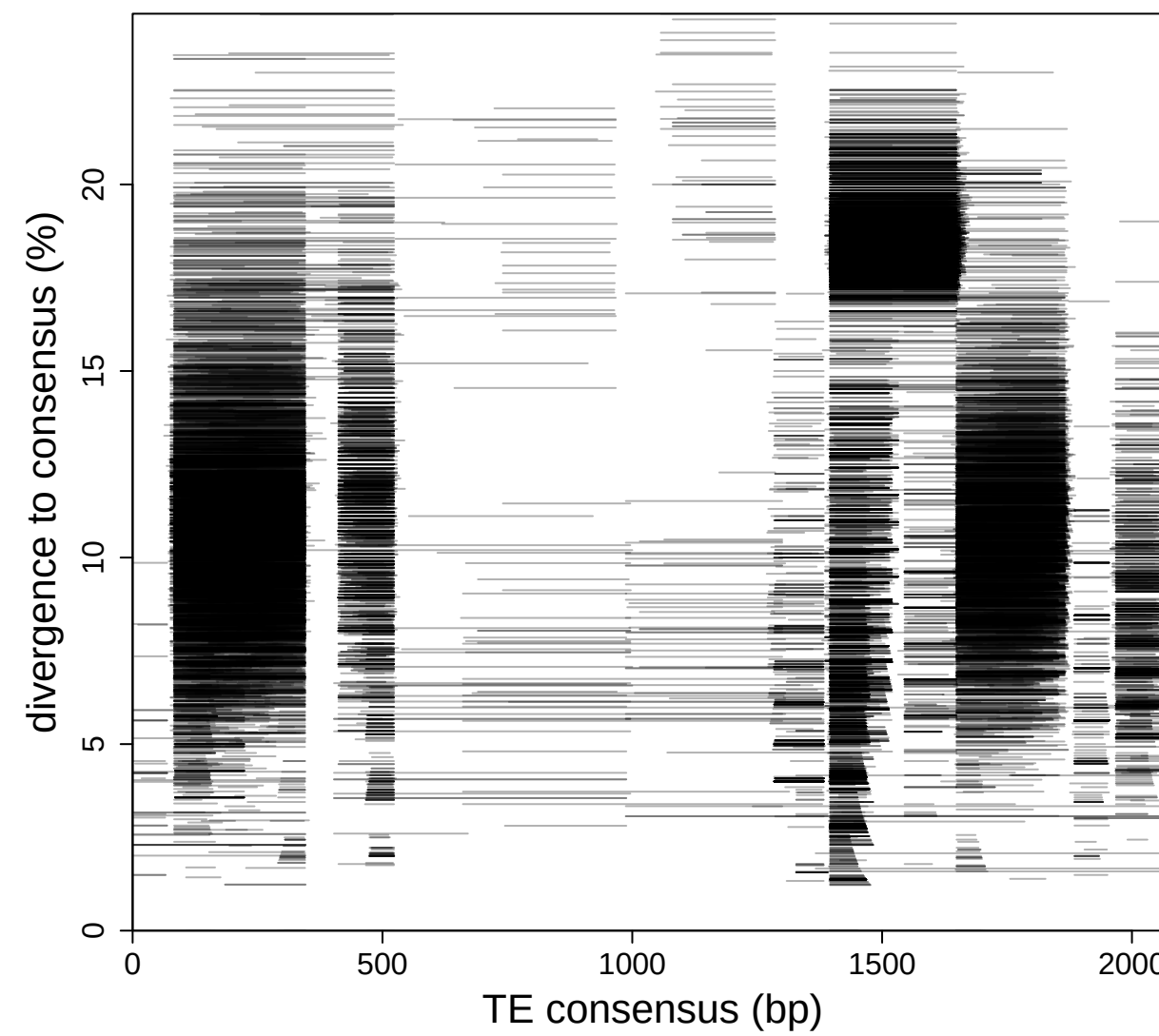
Start crop Point

End crop Point



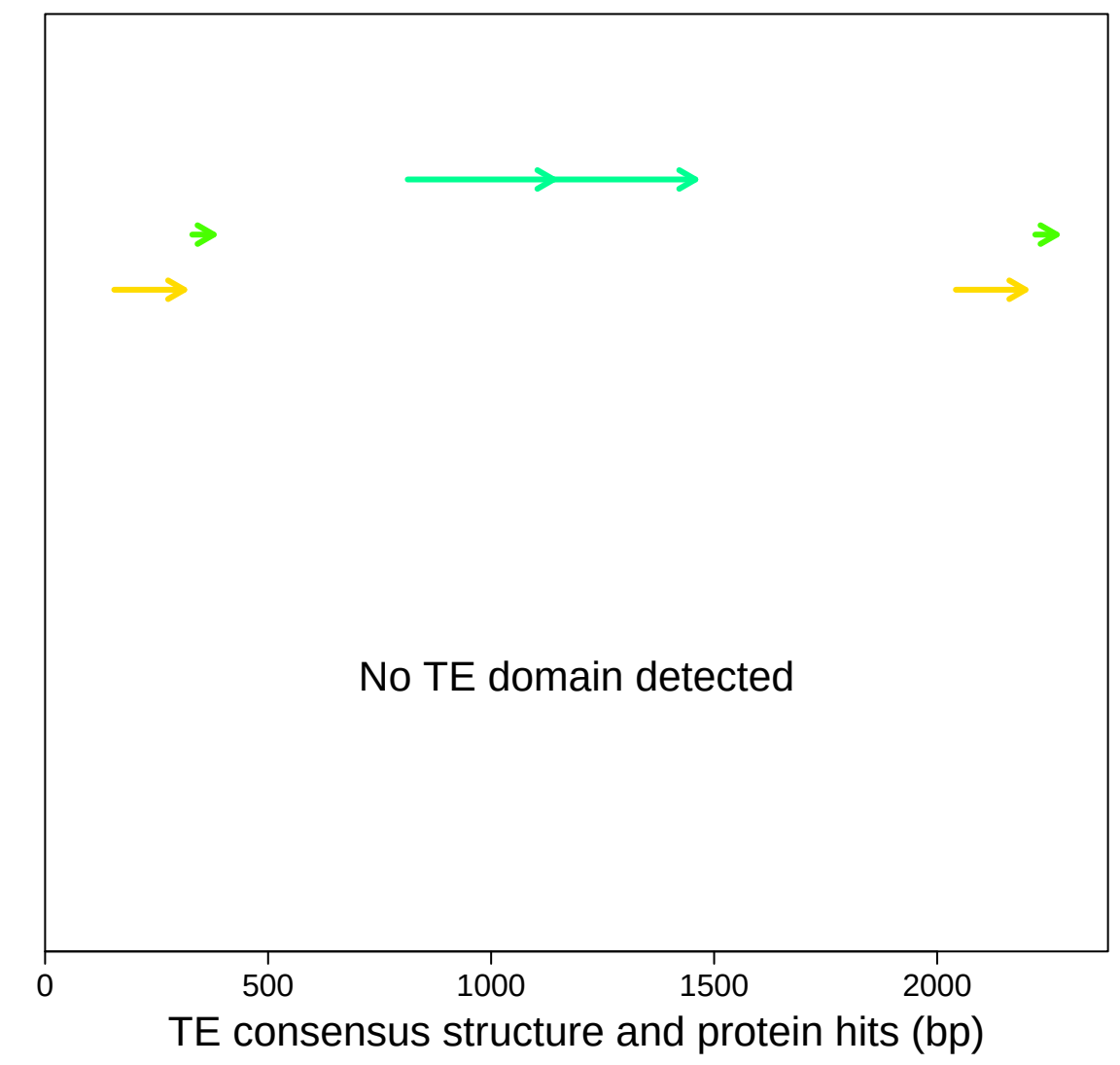
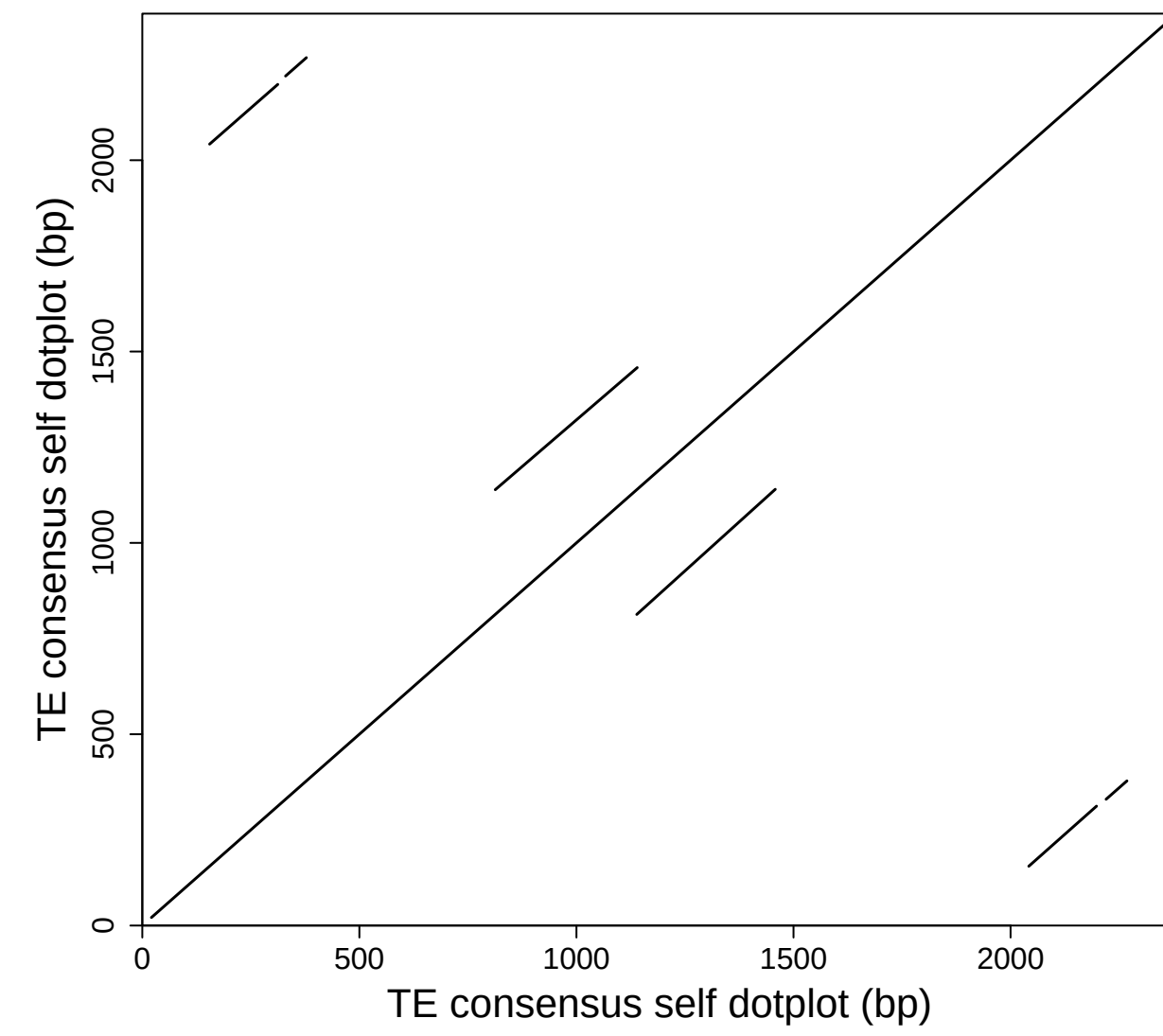
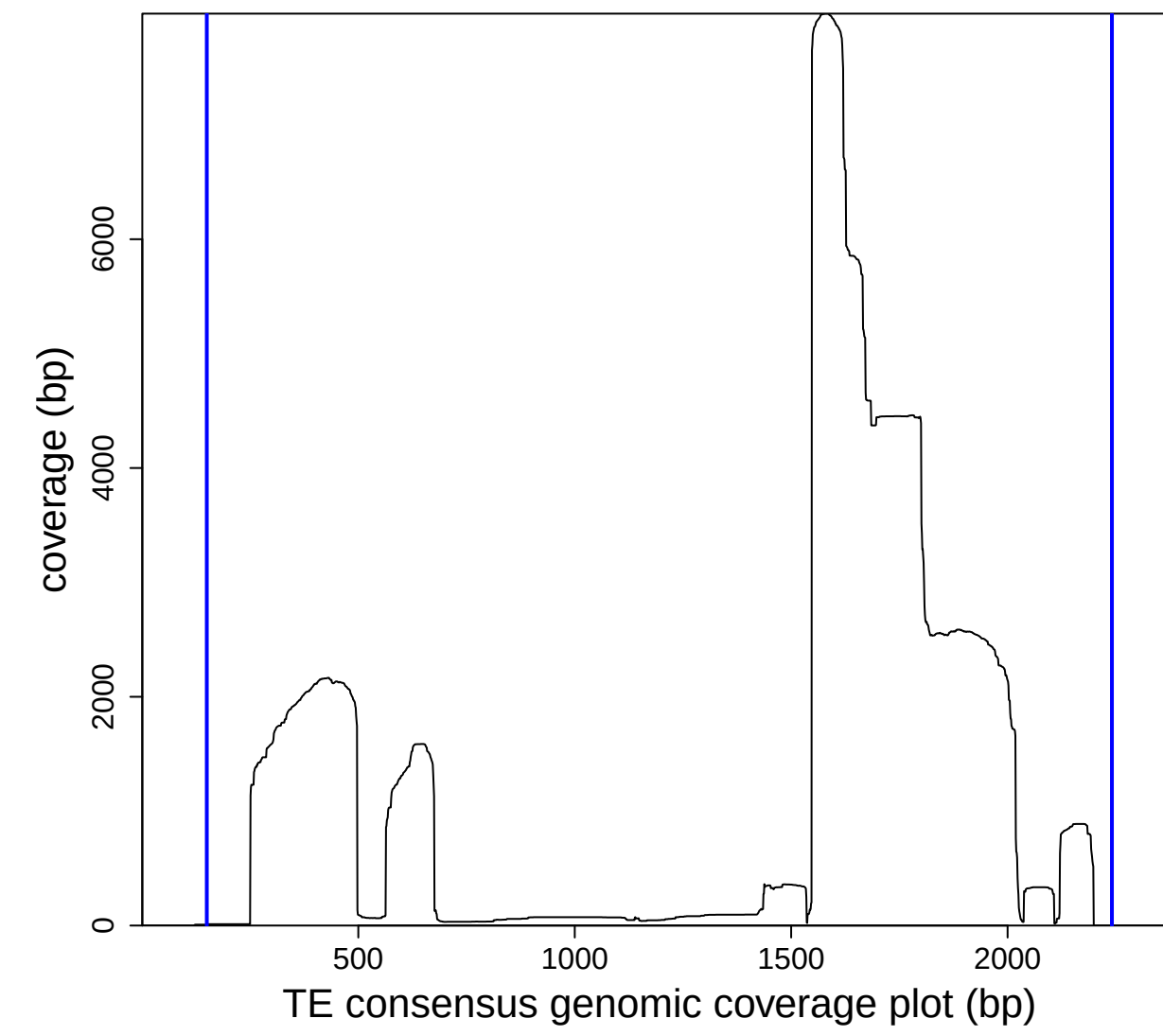
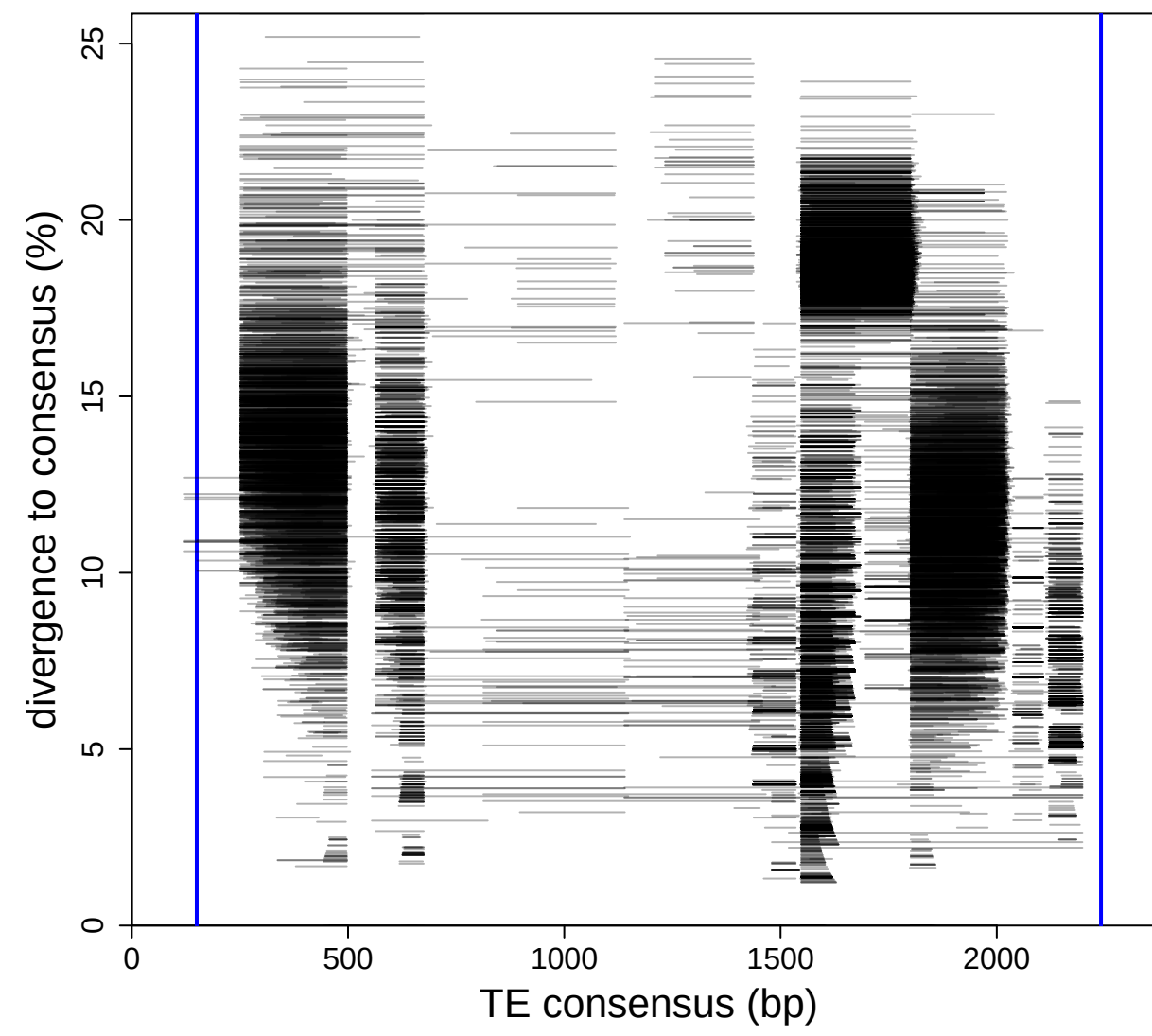
bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa_ce
size: 2082bp; fragments: 18428; full length: 0 (>=1873.8bp)

After TETrimmer 2082 bp

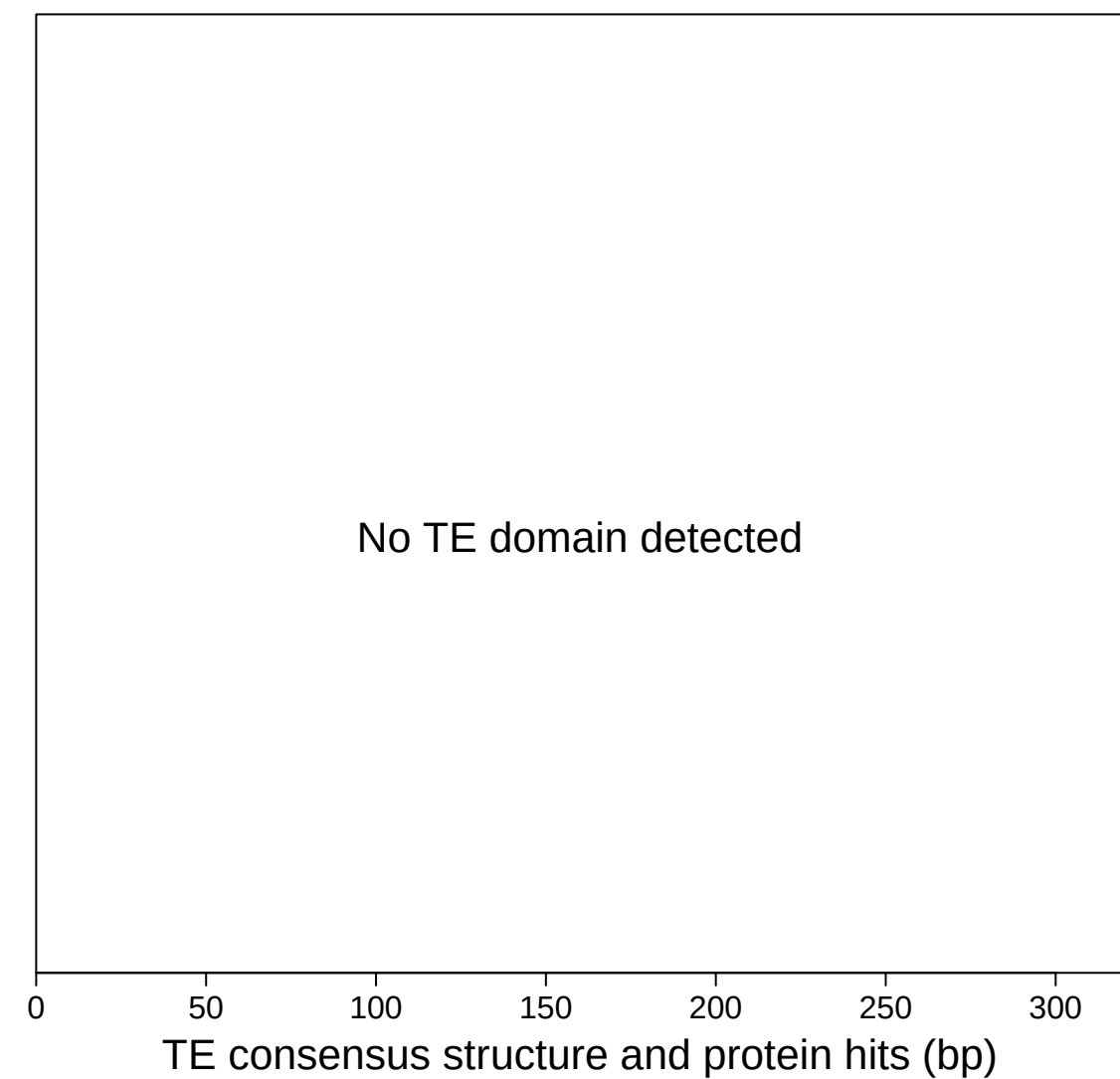
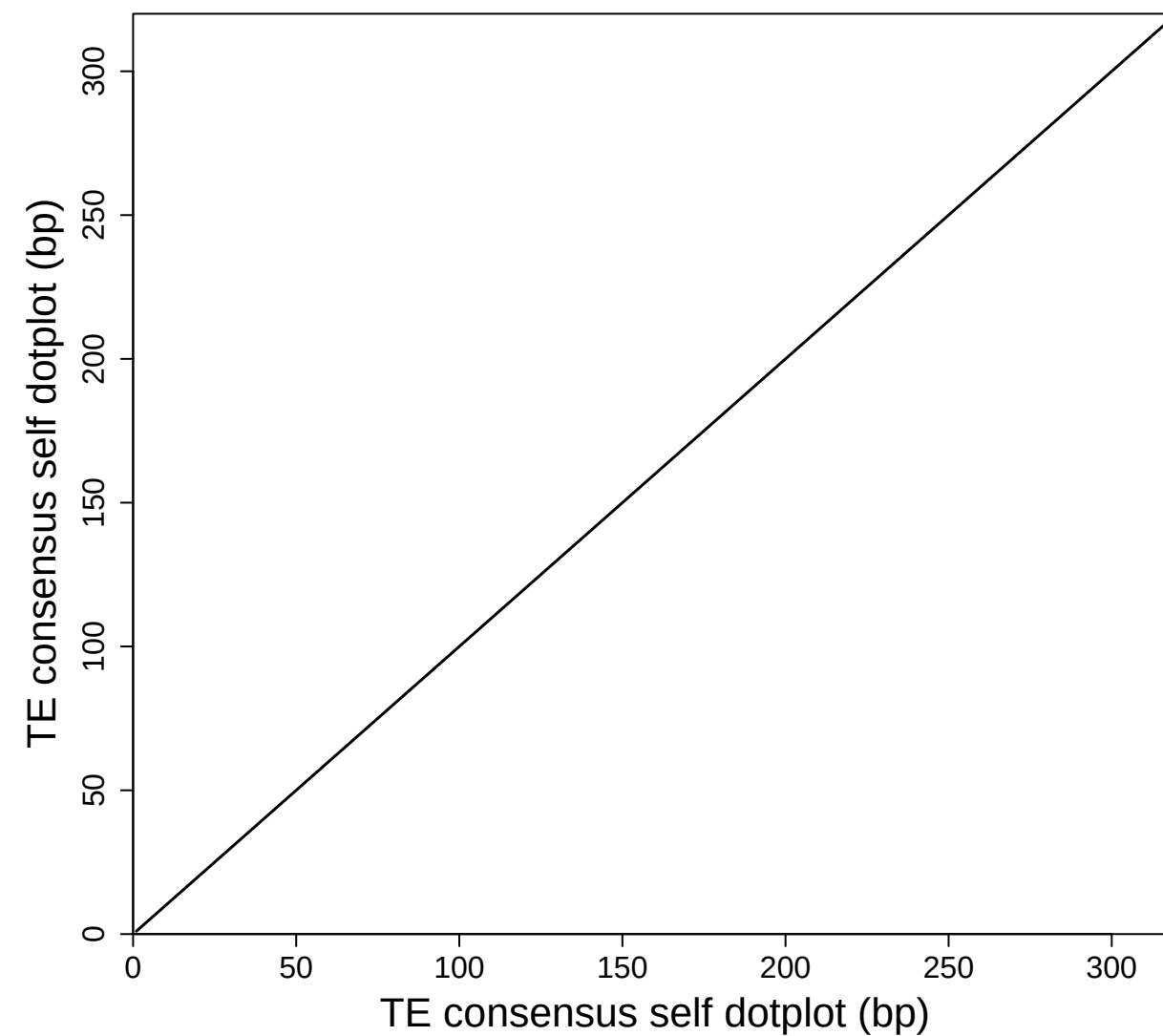
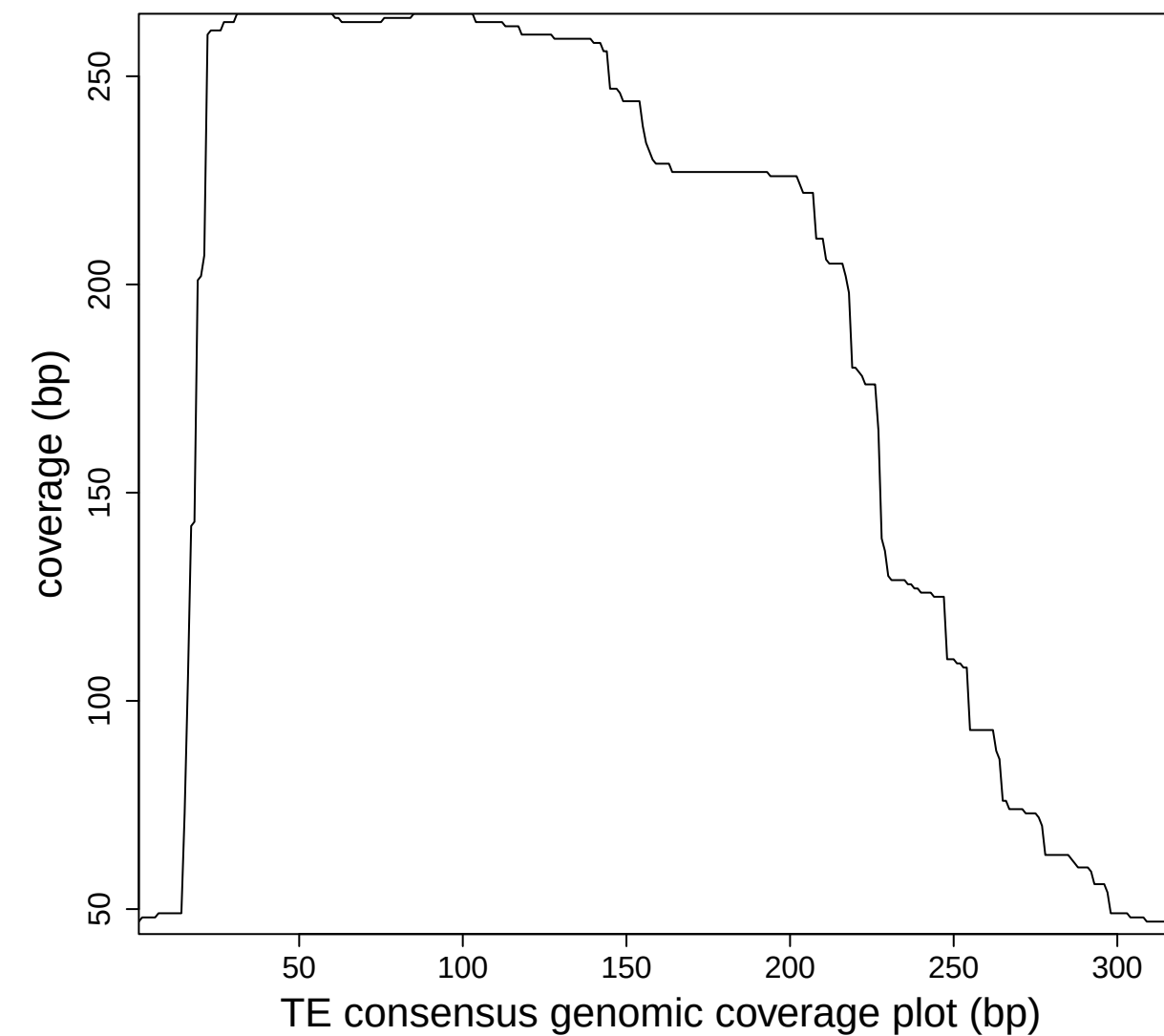
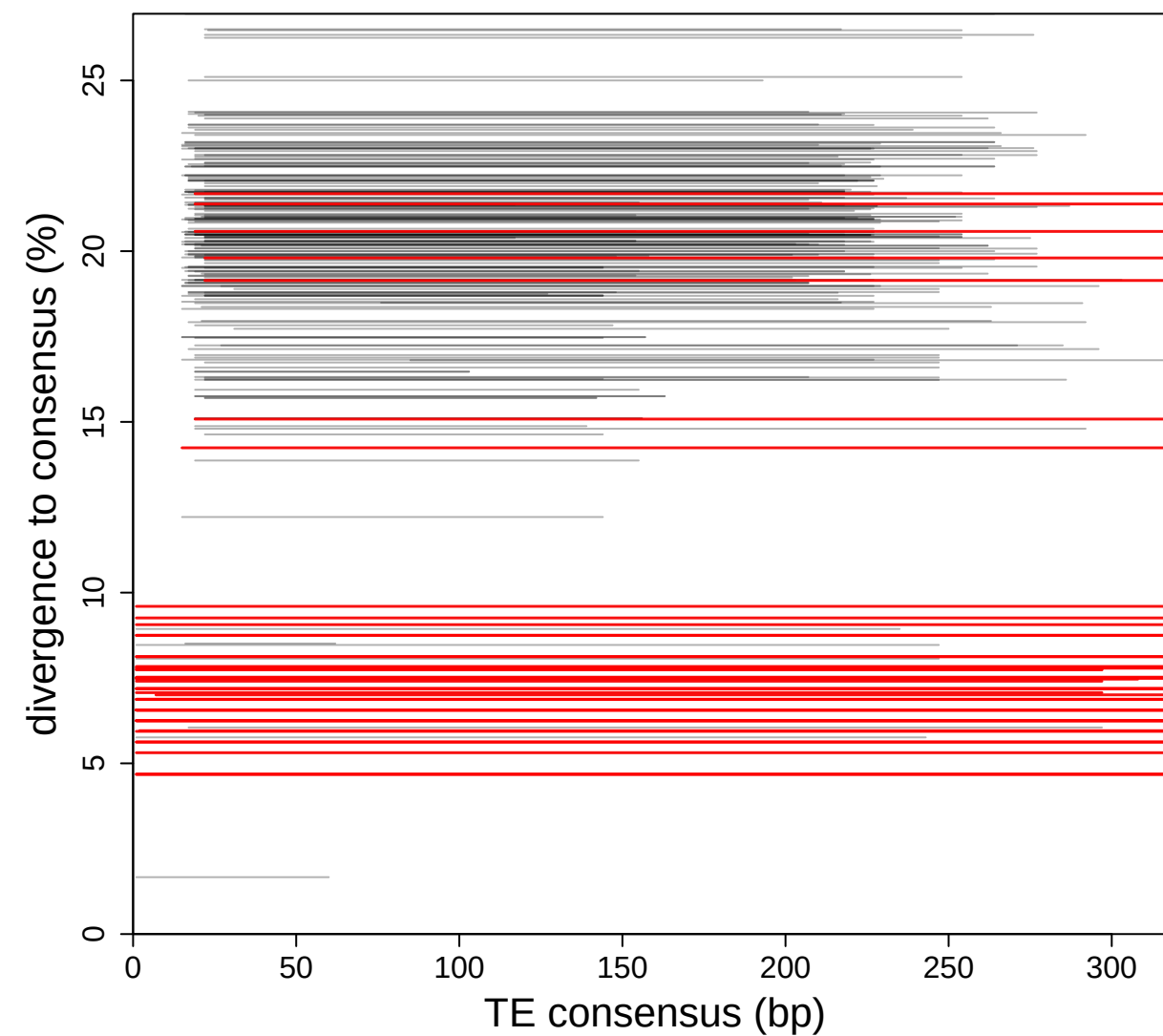


_947.fasta.b.bed uf.bed g 1.bed fm 1.bed 0 0 bcln.fa aln.fa c
size: 2391bp; fragments: 16155; full length: 0 (>=2151.9bp)

After TEtrimmer Extended plot Blue lines are boundaries

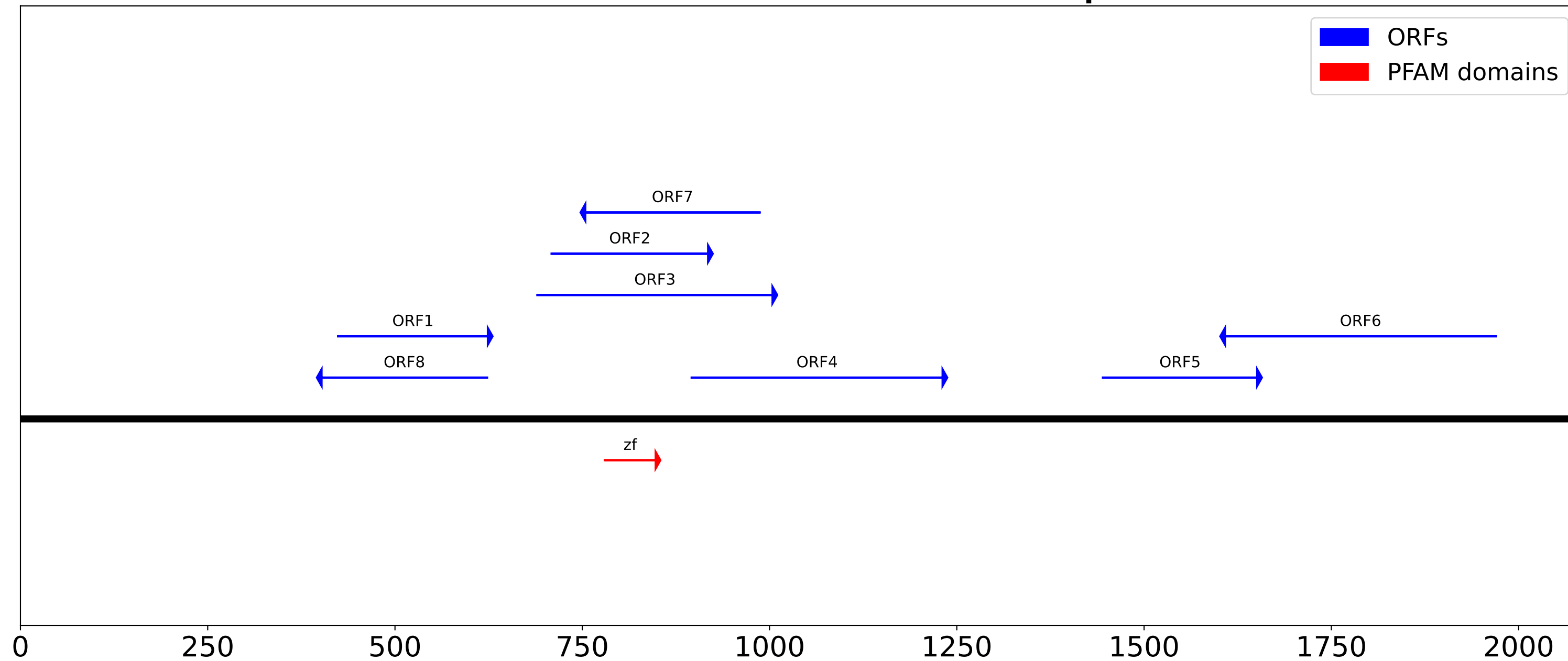


TE: rnd_1_family_947
size: 320bp; fragments: 267; full length: 51 (≥ 288 bp)

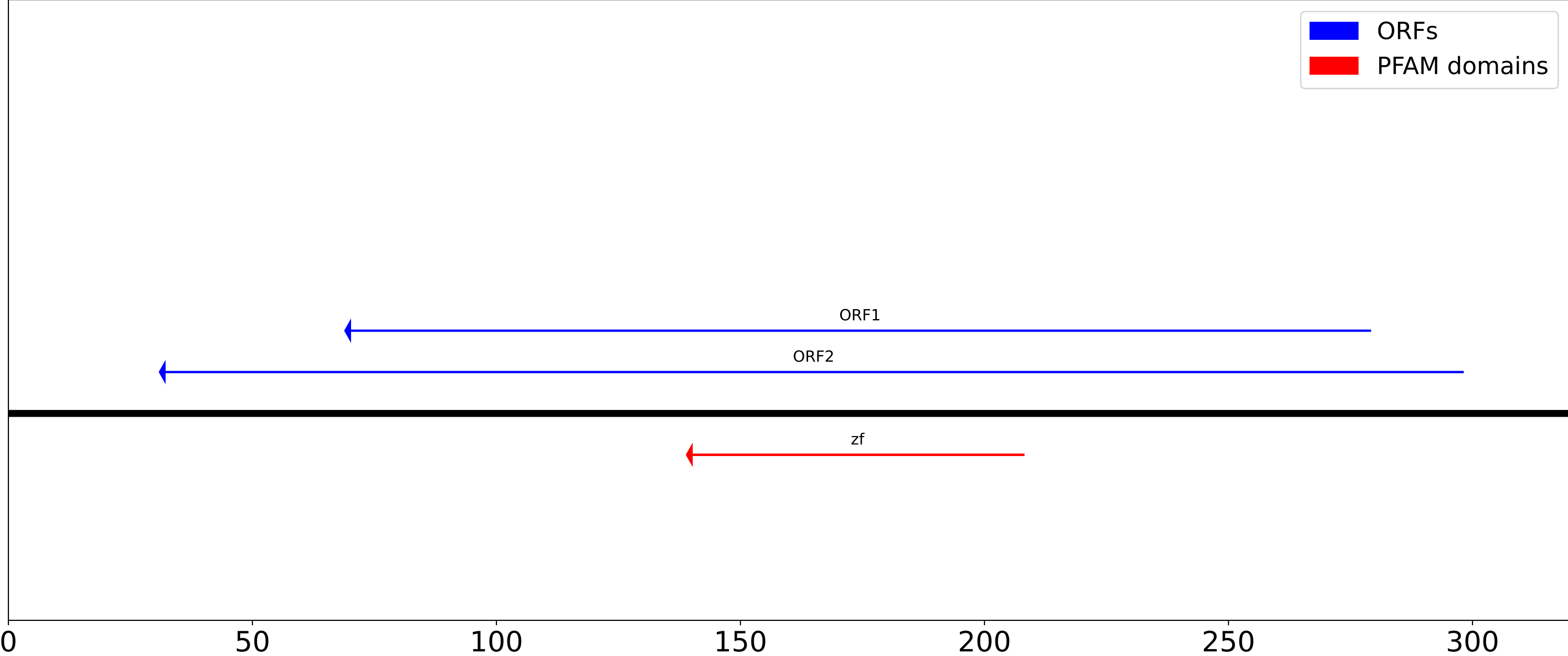


Before TEtrimmer 320 bp

After TEtrimmer ORF and PFAM domain plot



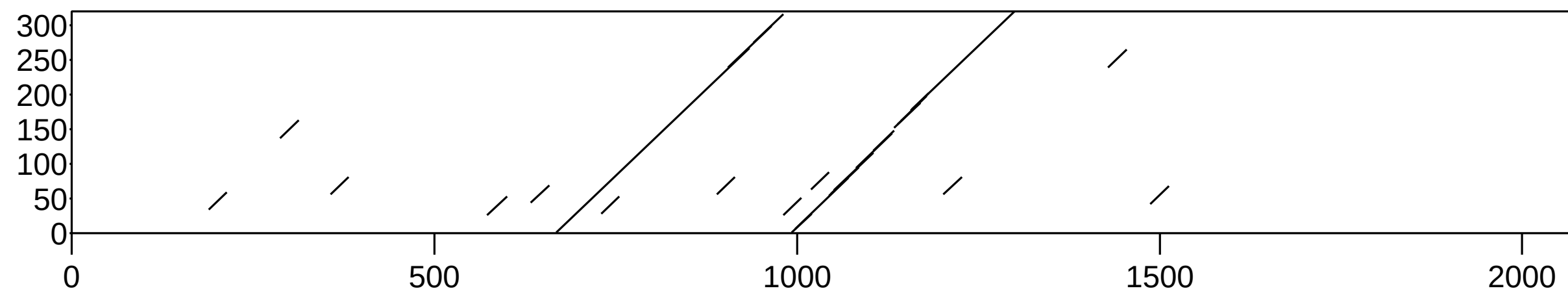
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)