

Start crop Point



End crop Point

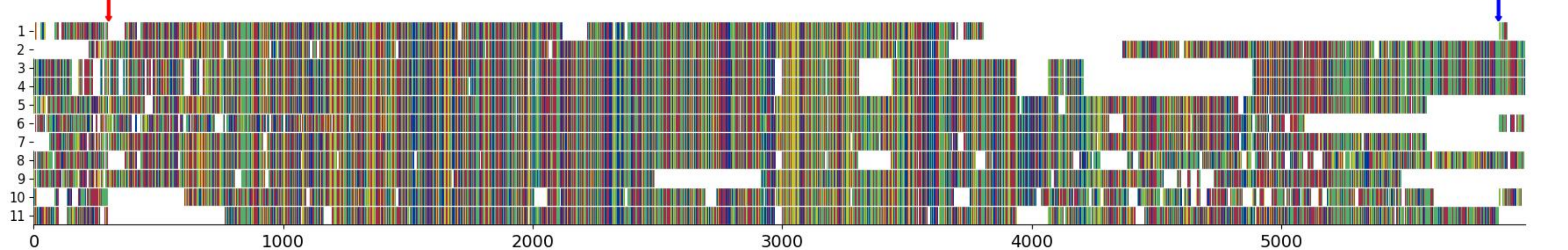


MSA length = 5572



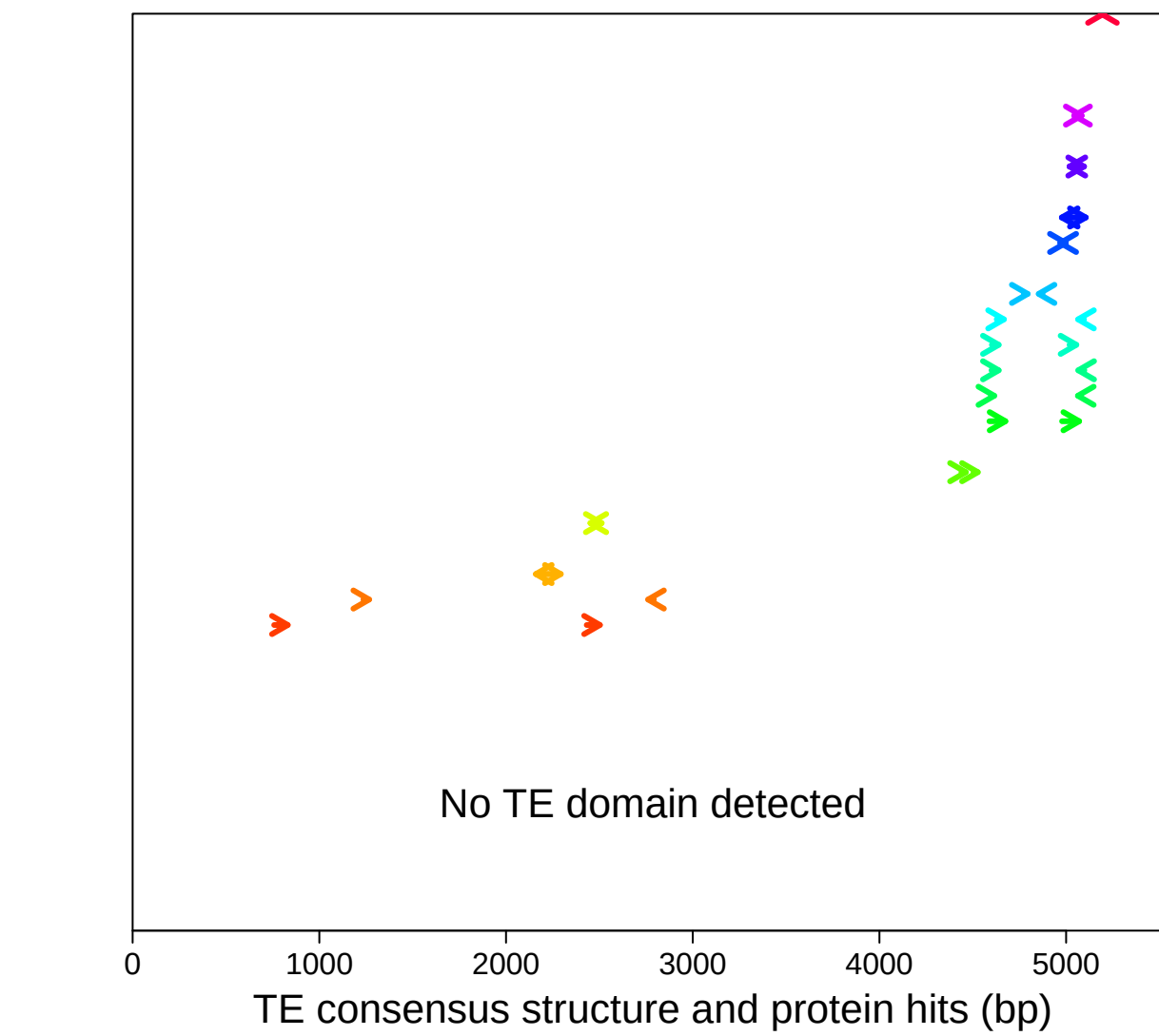
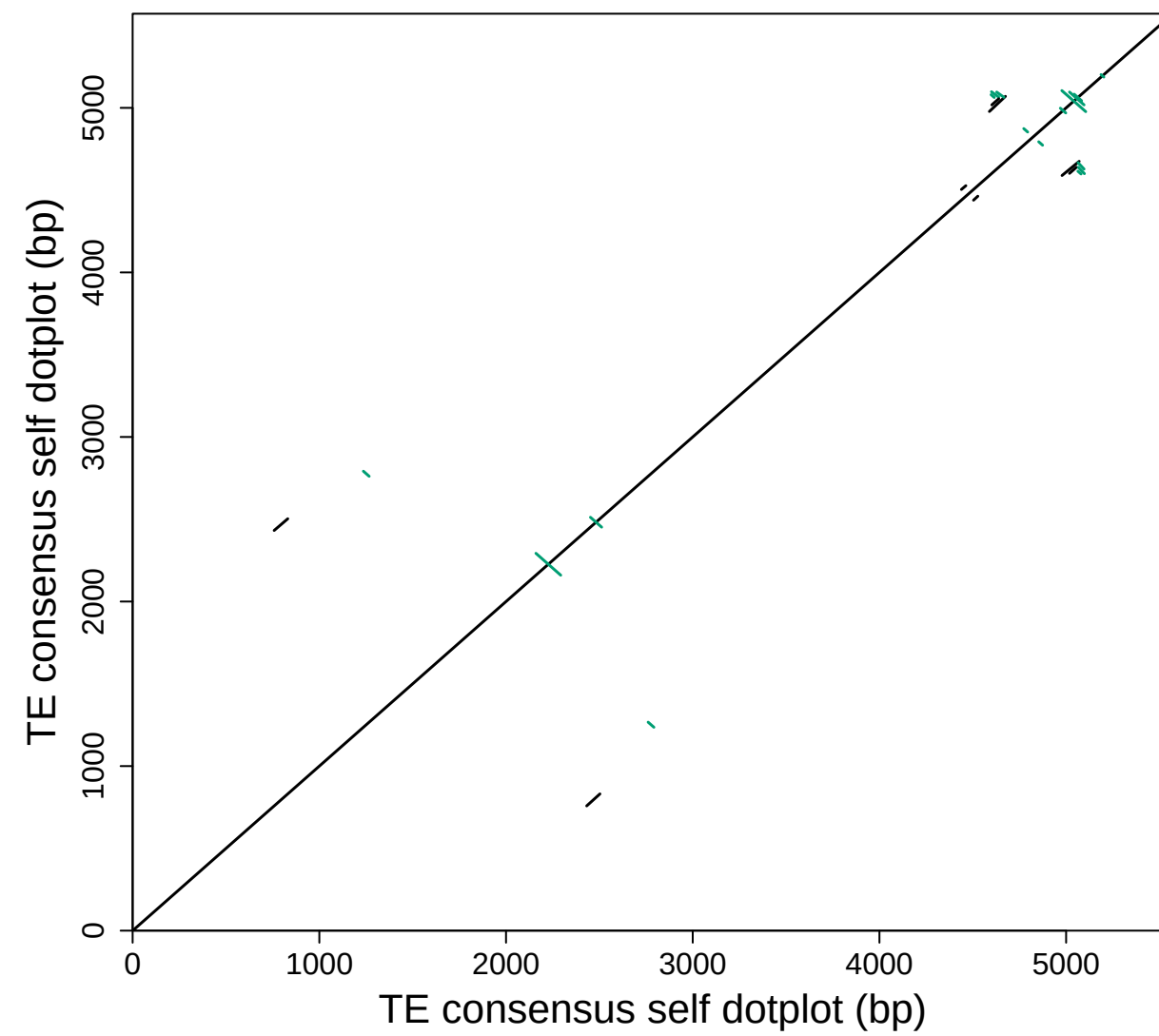
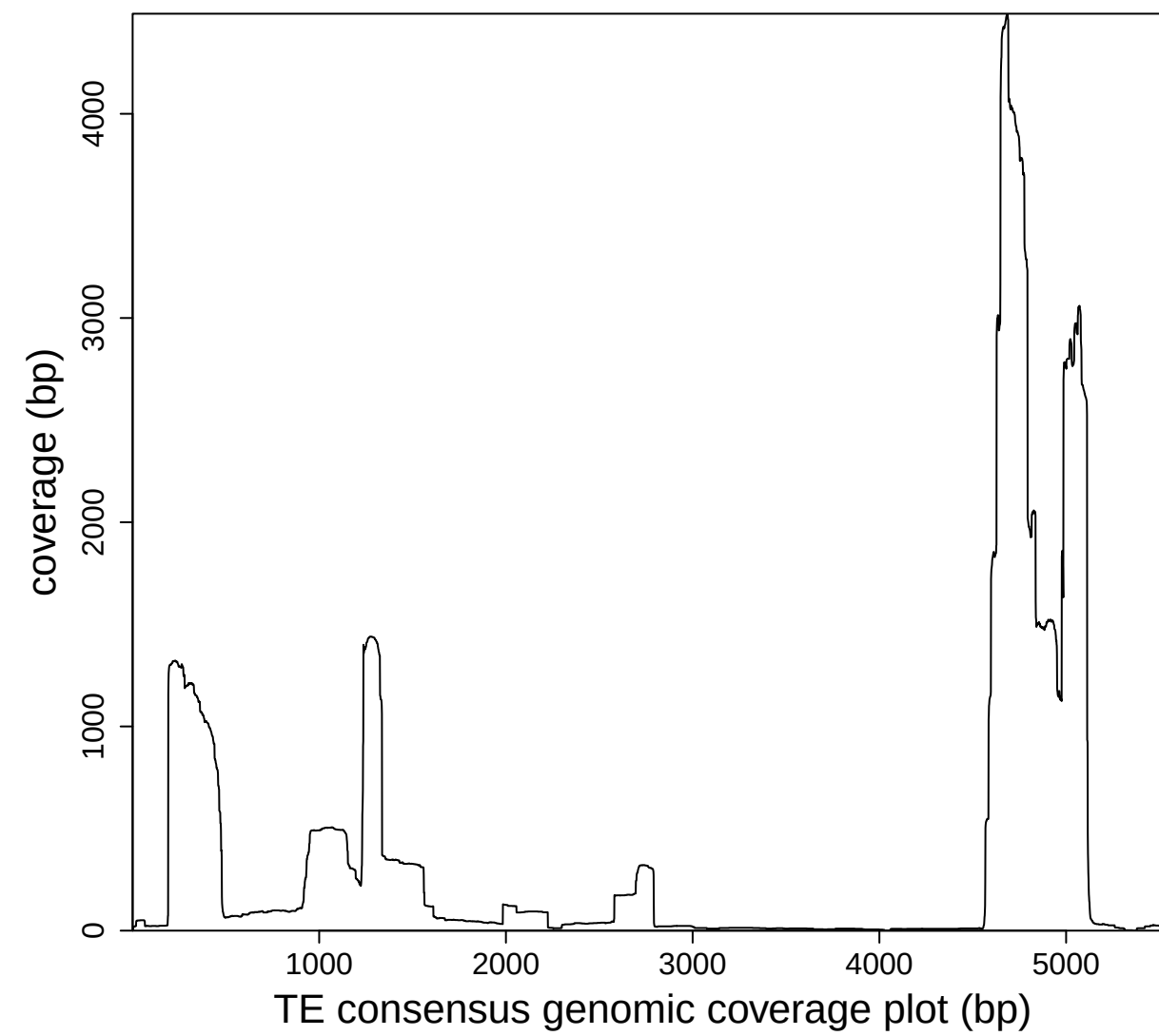
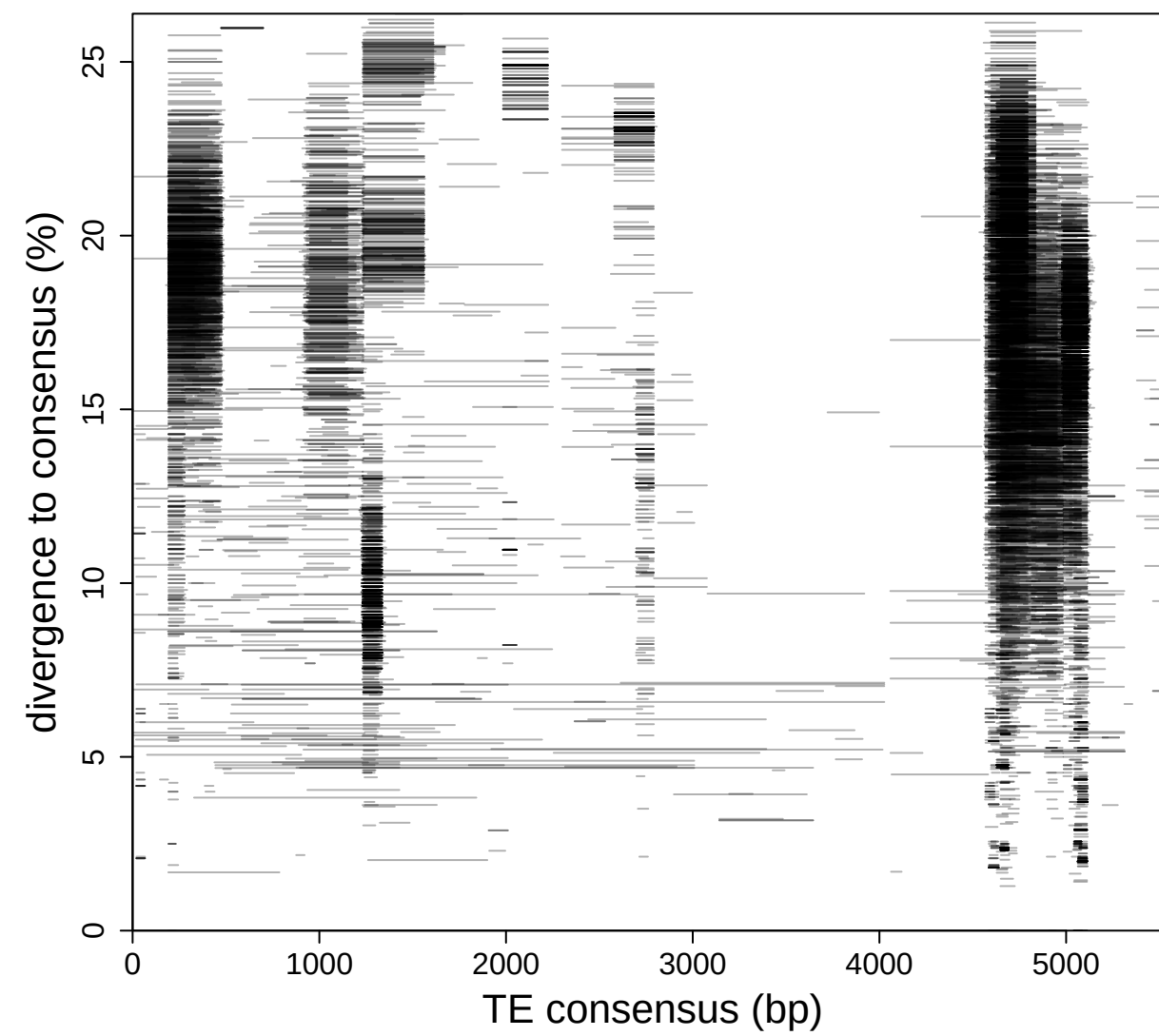
Start crop Point

End crop Point



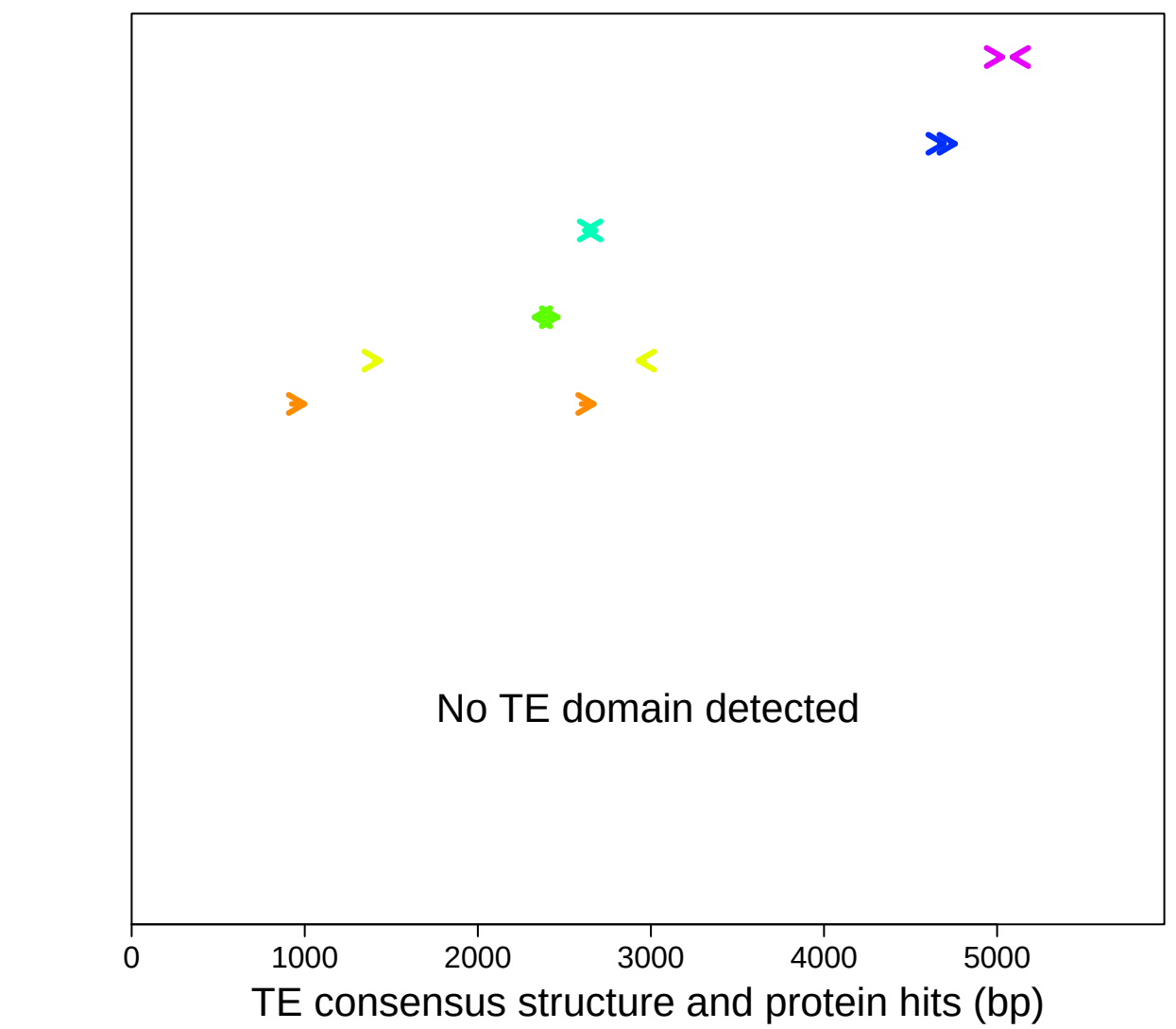
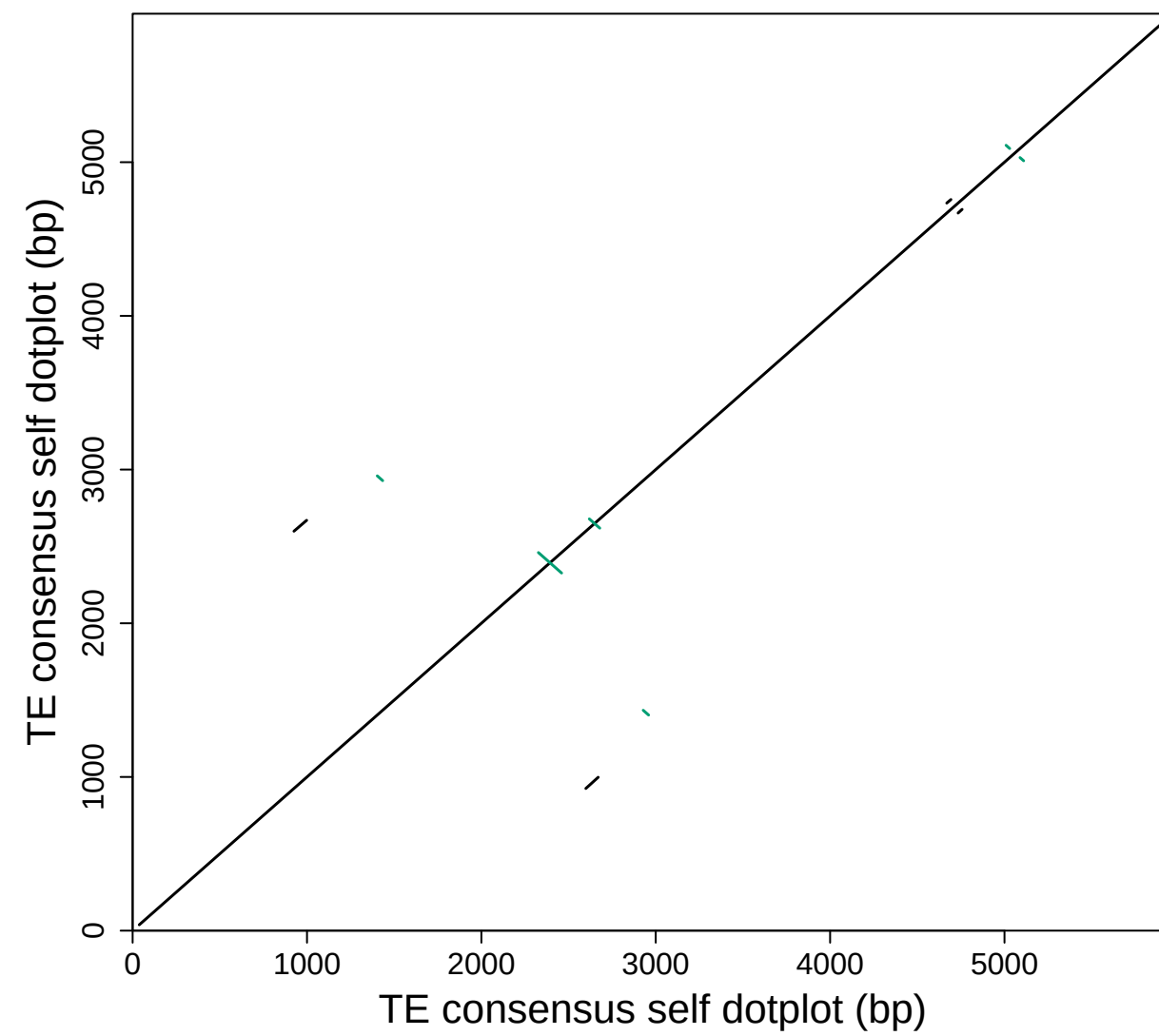
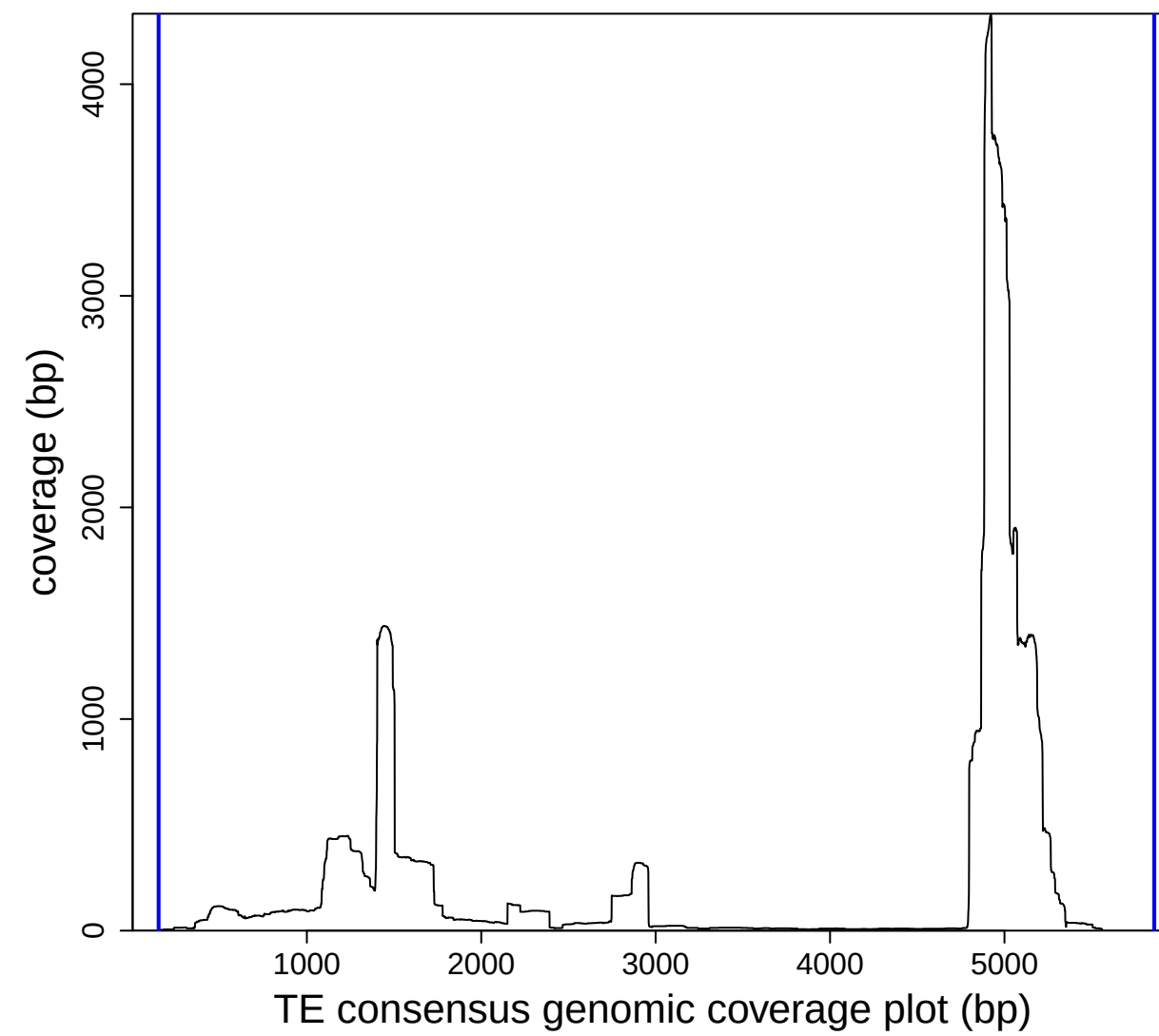
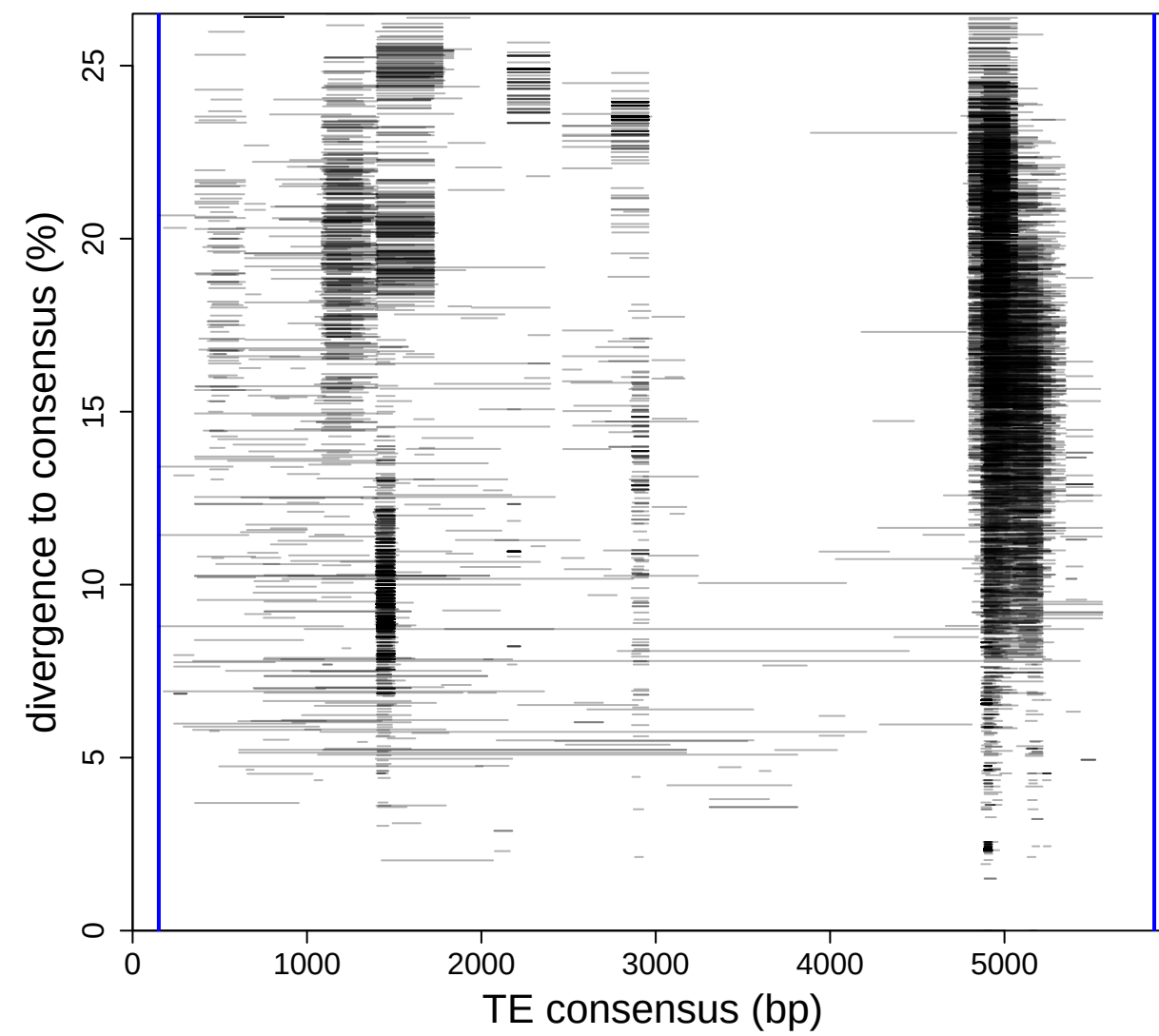
a.b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.f
size: 5572bp; fragments: 12052; full length: 0 (>=5014.8bp)

After TETrimmer 5572 bp

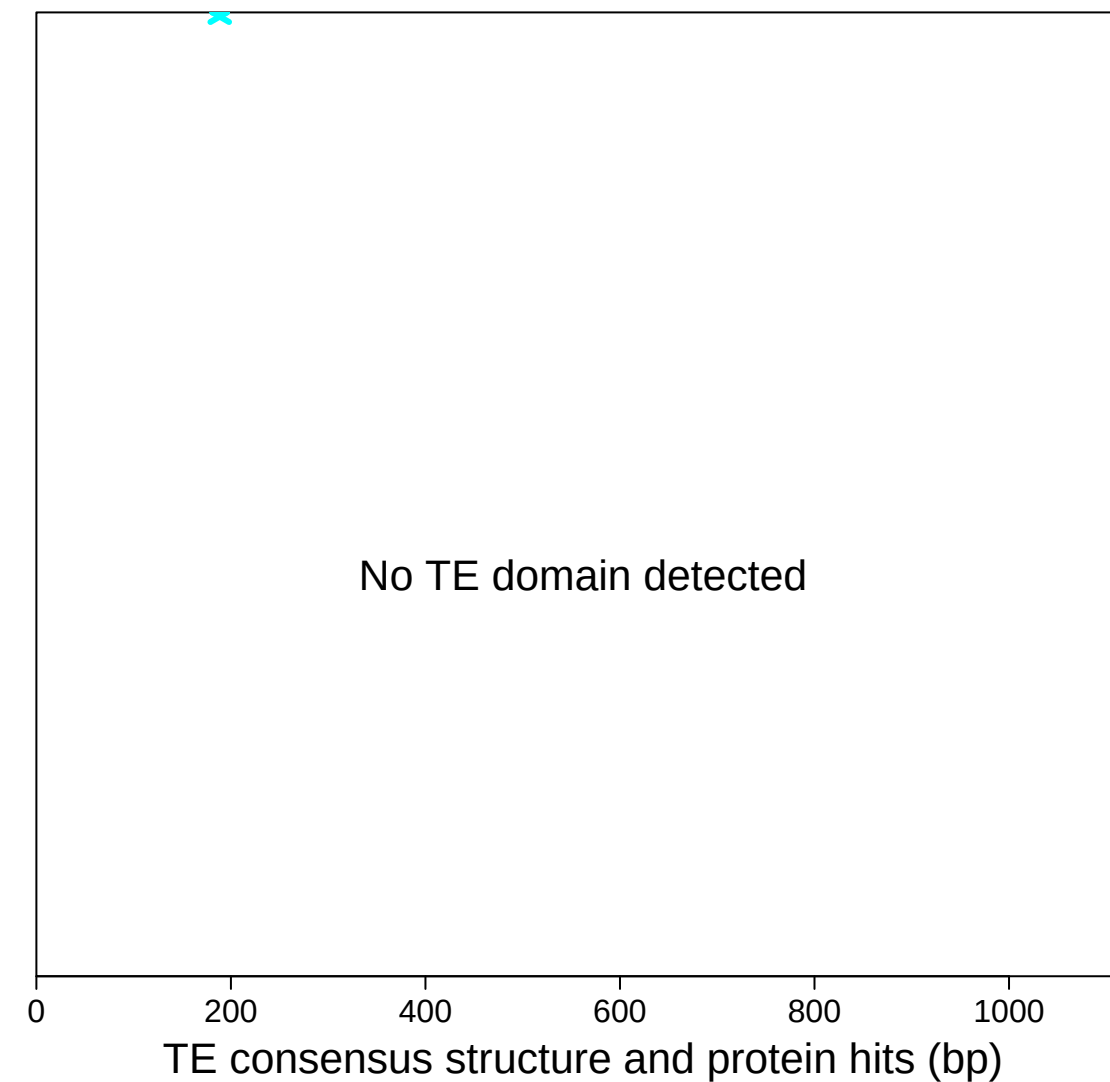
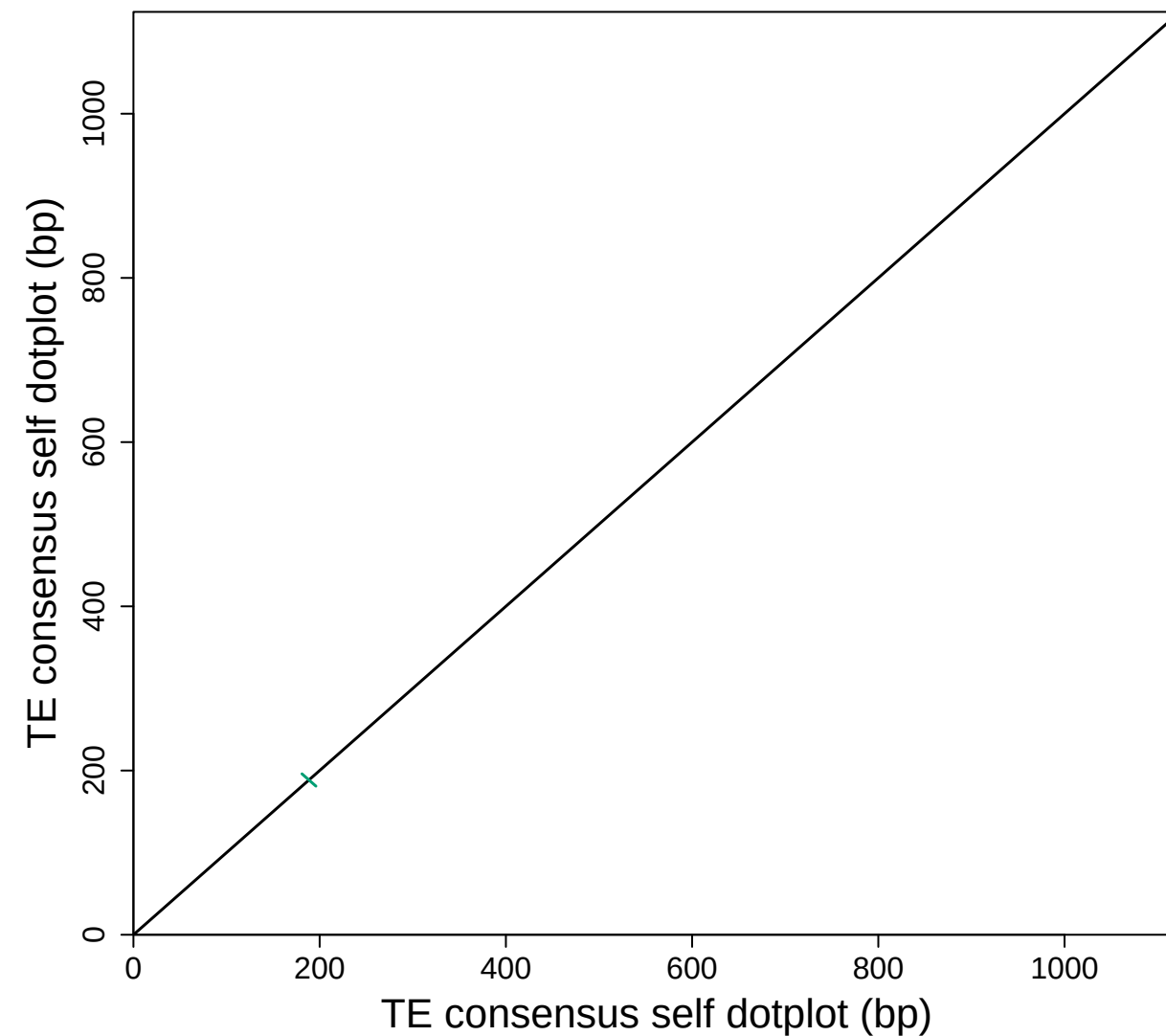
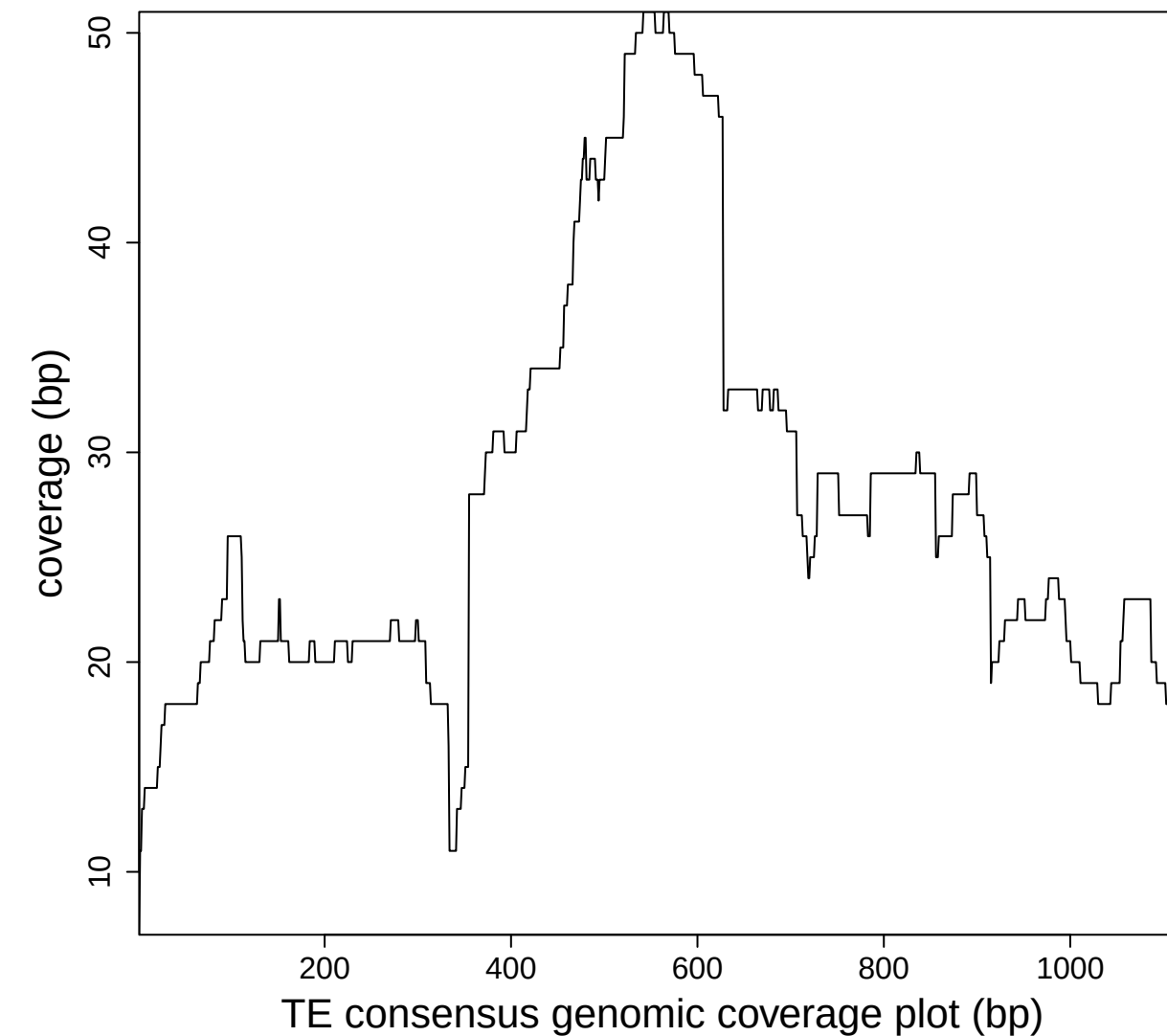
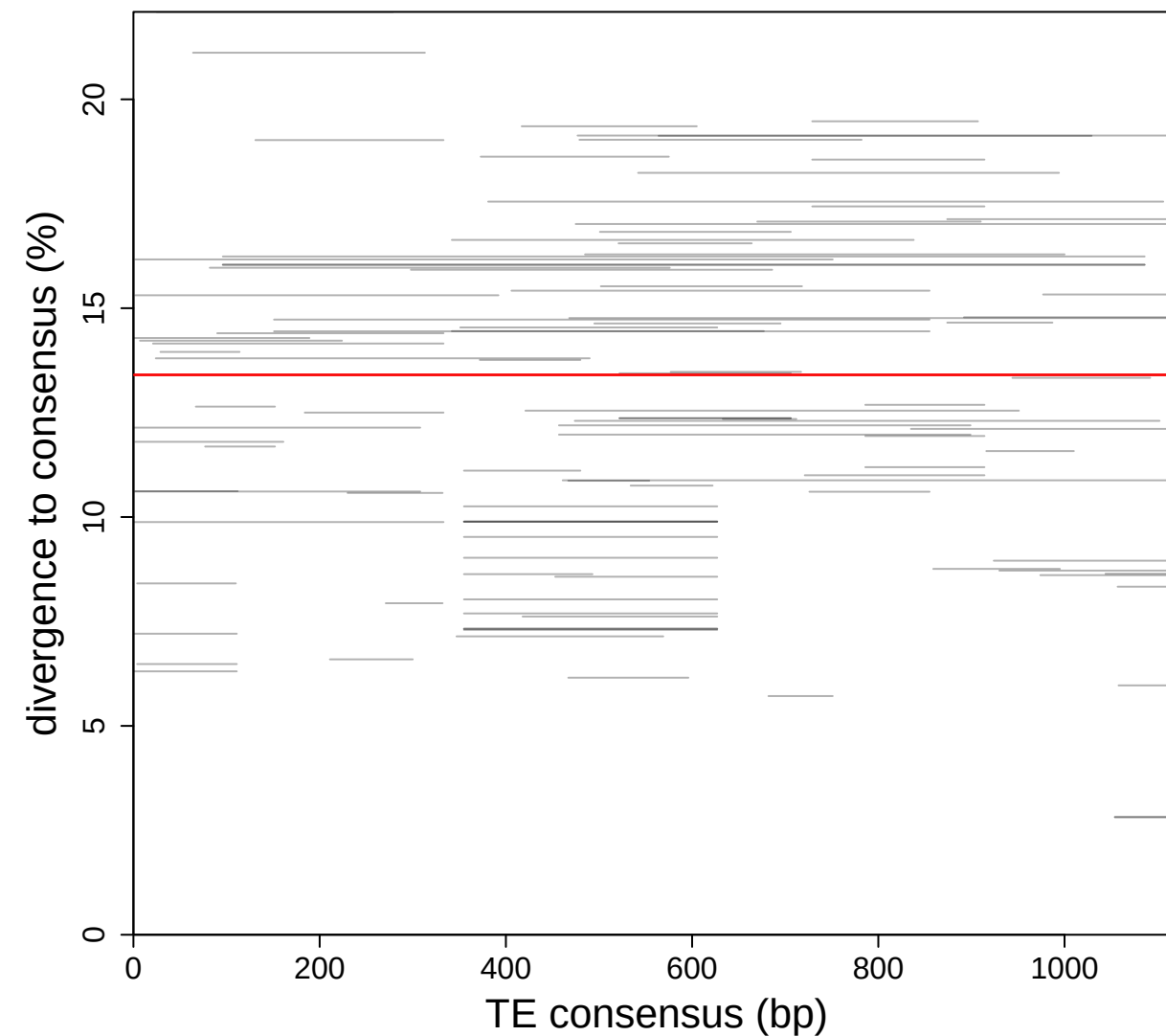


0.fasta.b.bed uf.bed g_2.bed fm_1.bed 0_0_bcln.fa aln.fa cl.f
size: 5966bp; fragments: 7582; full length: 0 (>=5369.4bp)

After TEtrimmer Extended plot Blue lines are boundaries

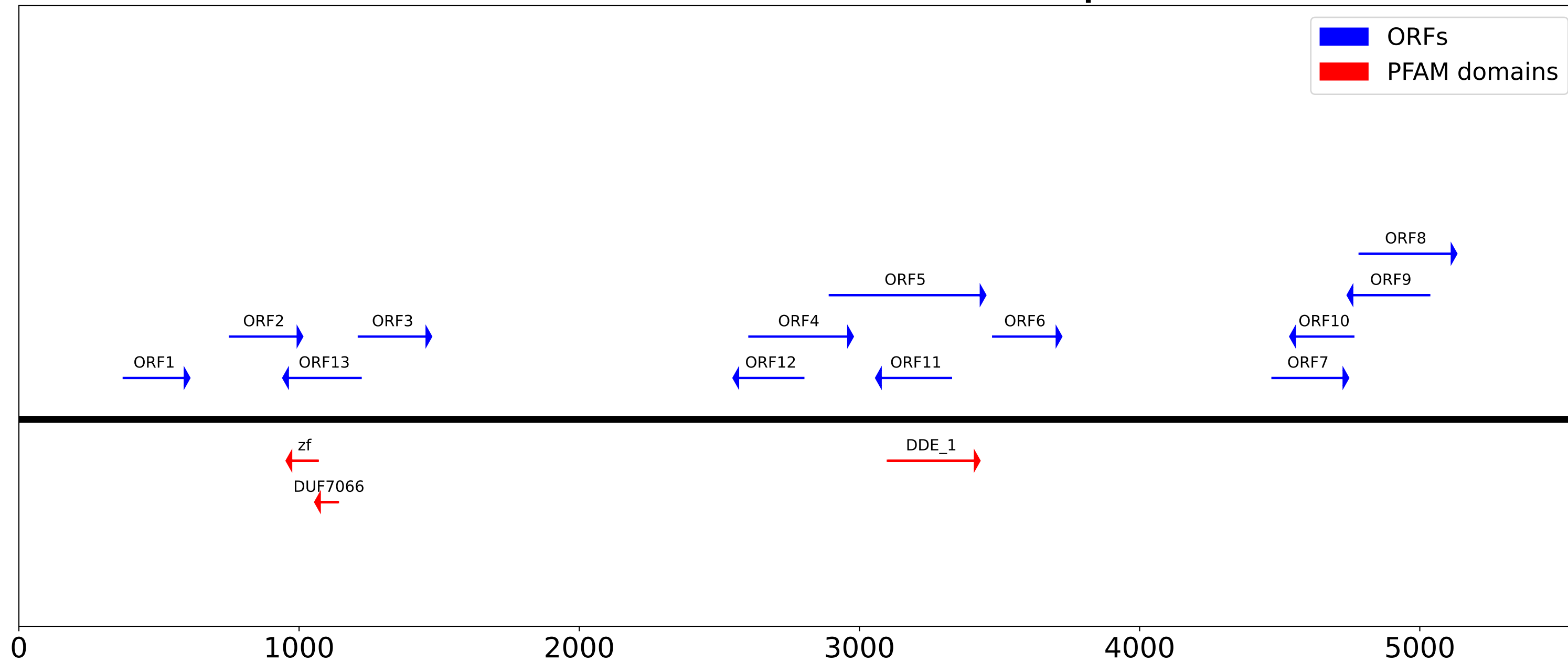


TE: rnd_5_family_3000
size: 1124bp; fragments: 108; full length: 1 (≥ 1011.6 bp)

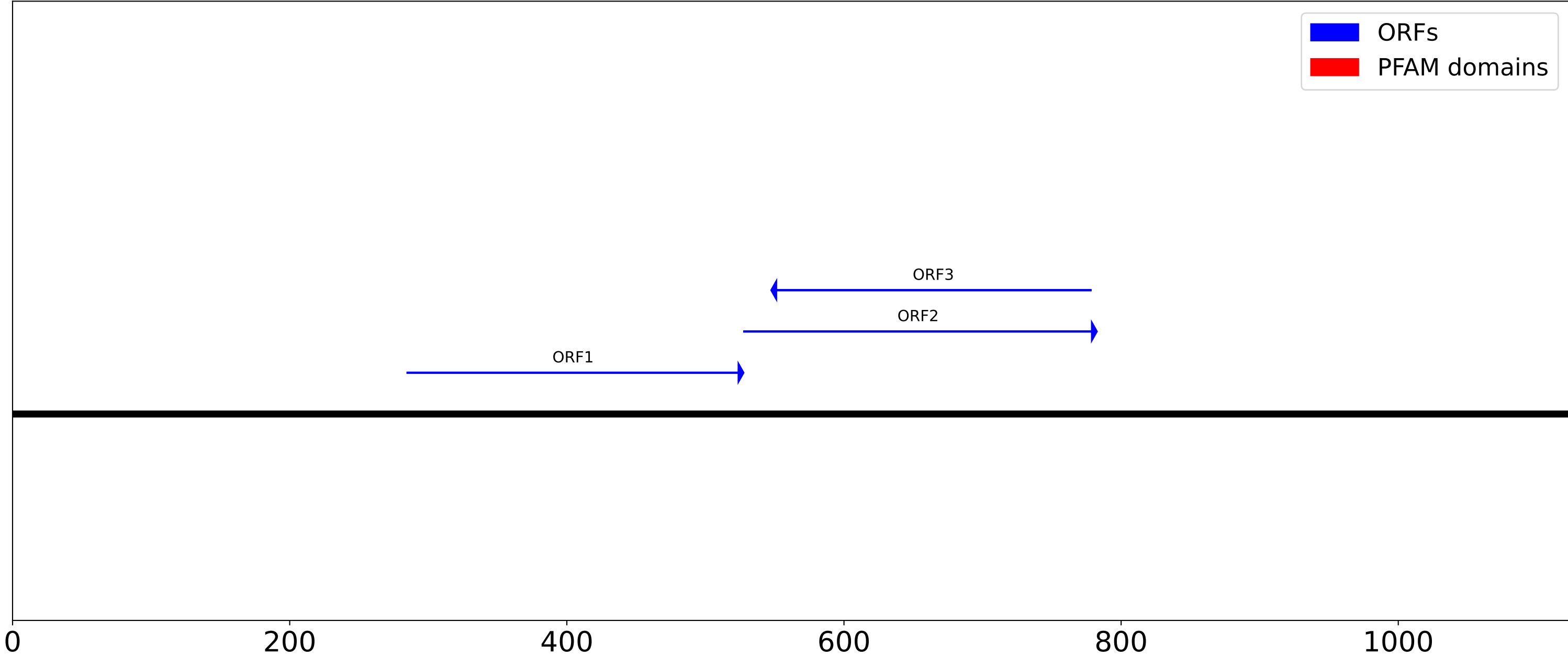


Before TEtrimmer 1124 bp

After TEtrimmer ORF and PFAM domain plot



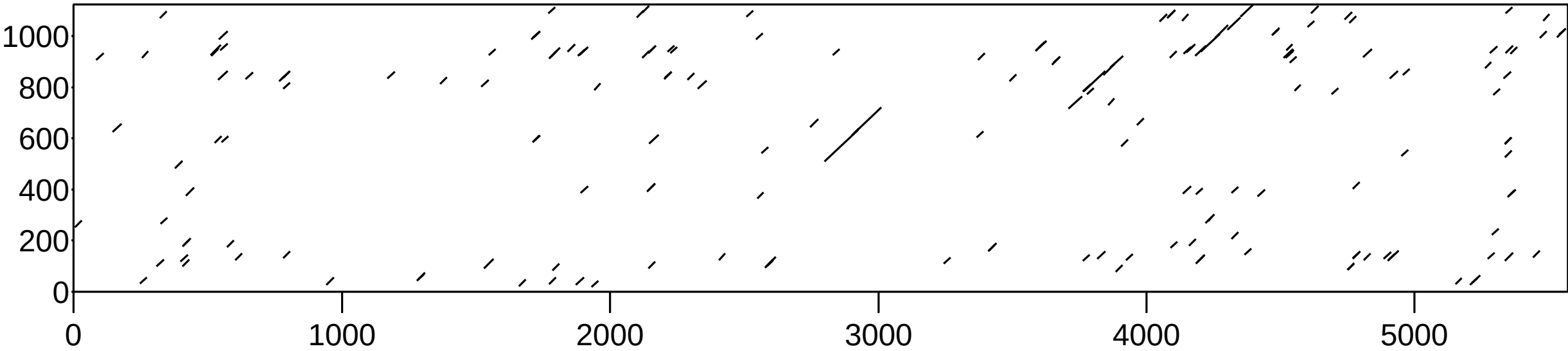
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)