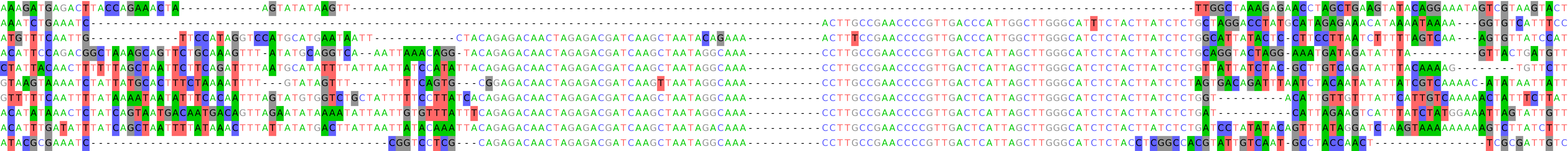


Start Crop Point

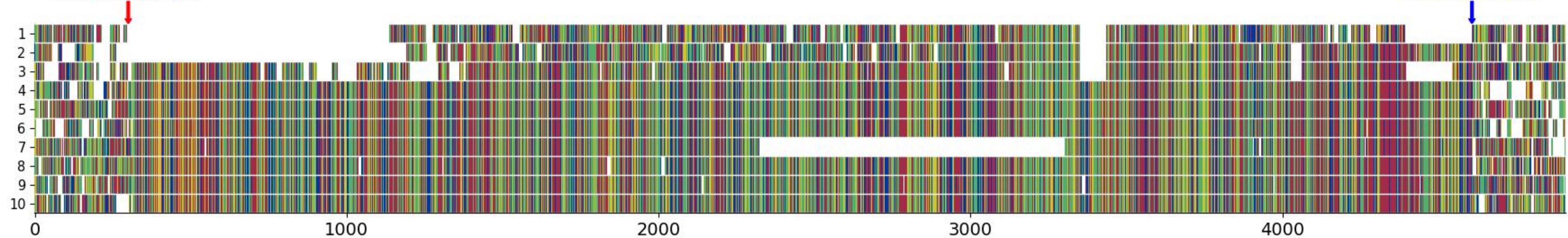
End Crop Point

MSA length = 4309



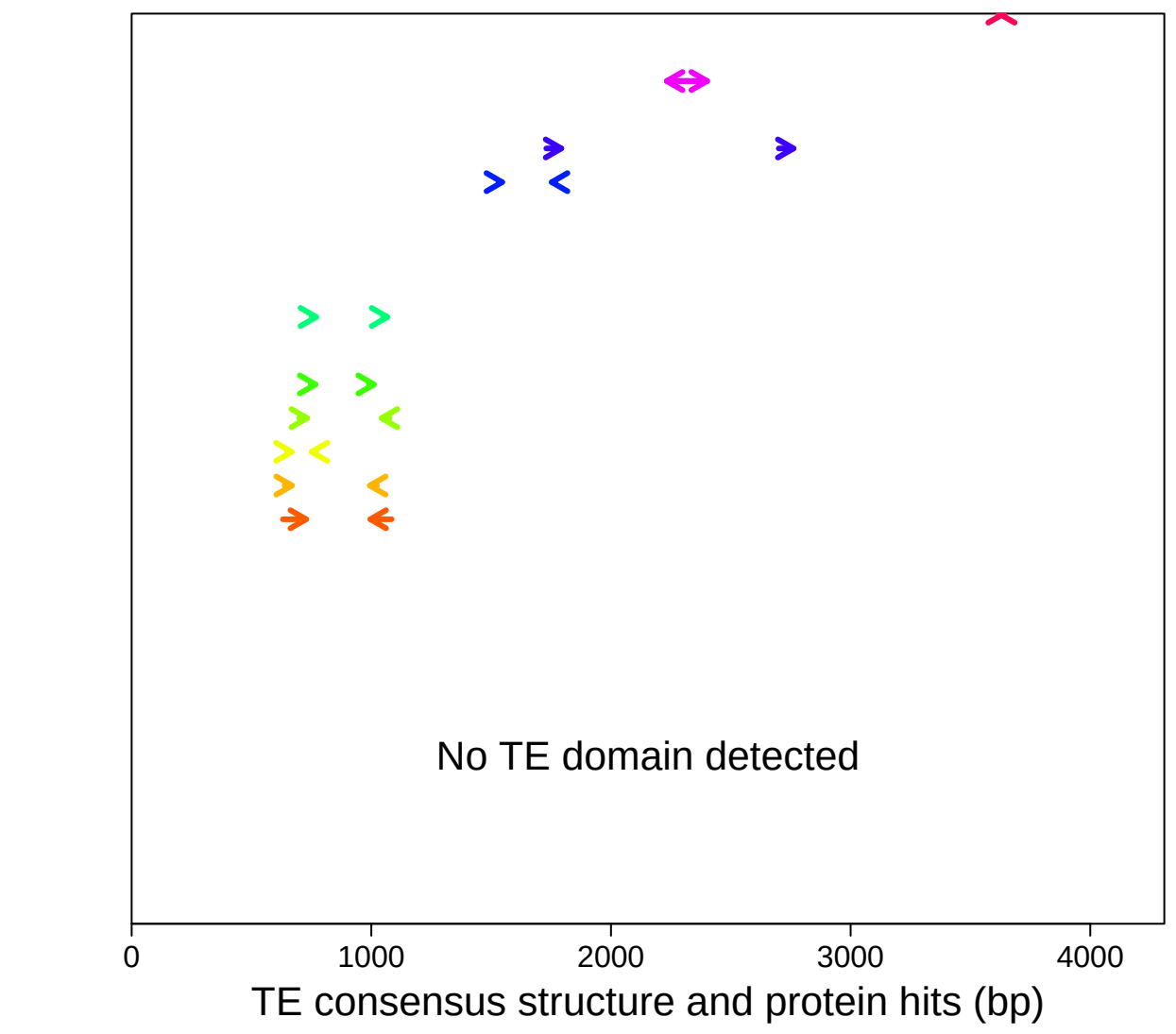
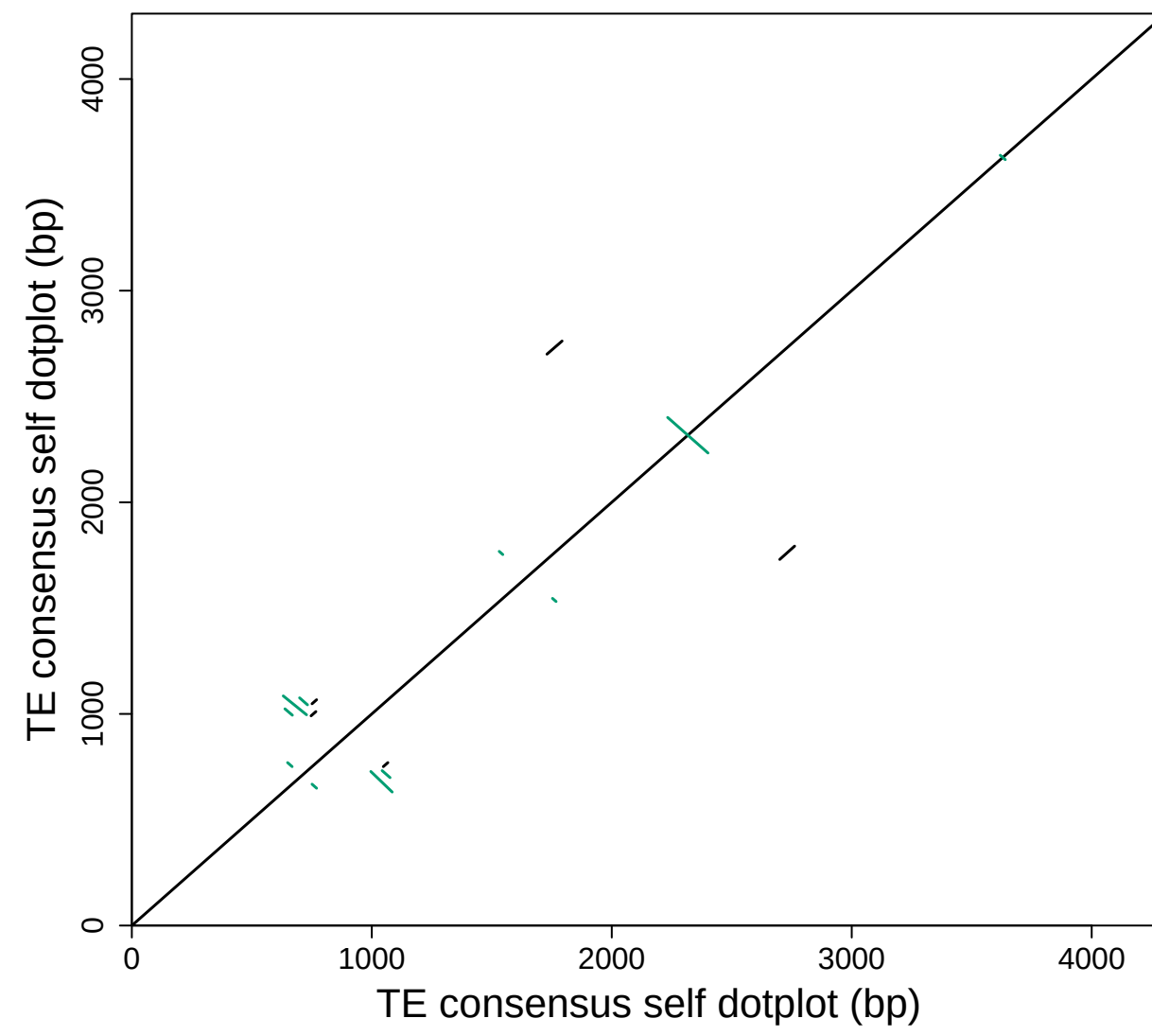
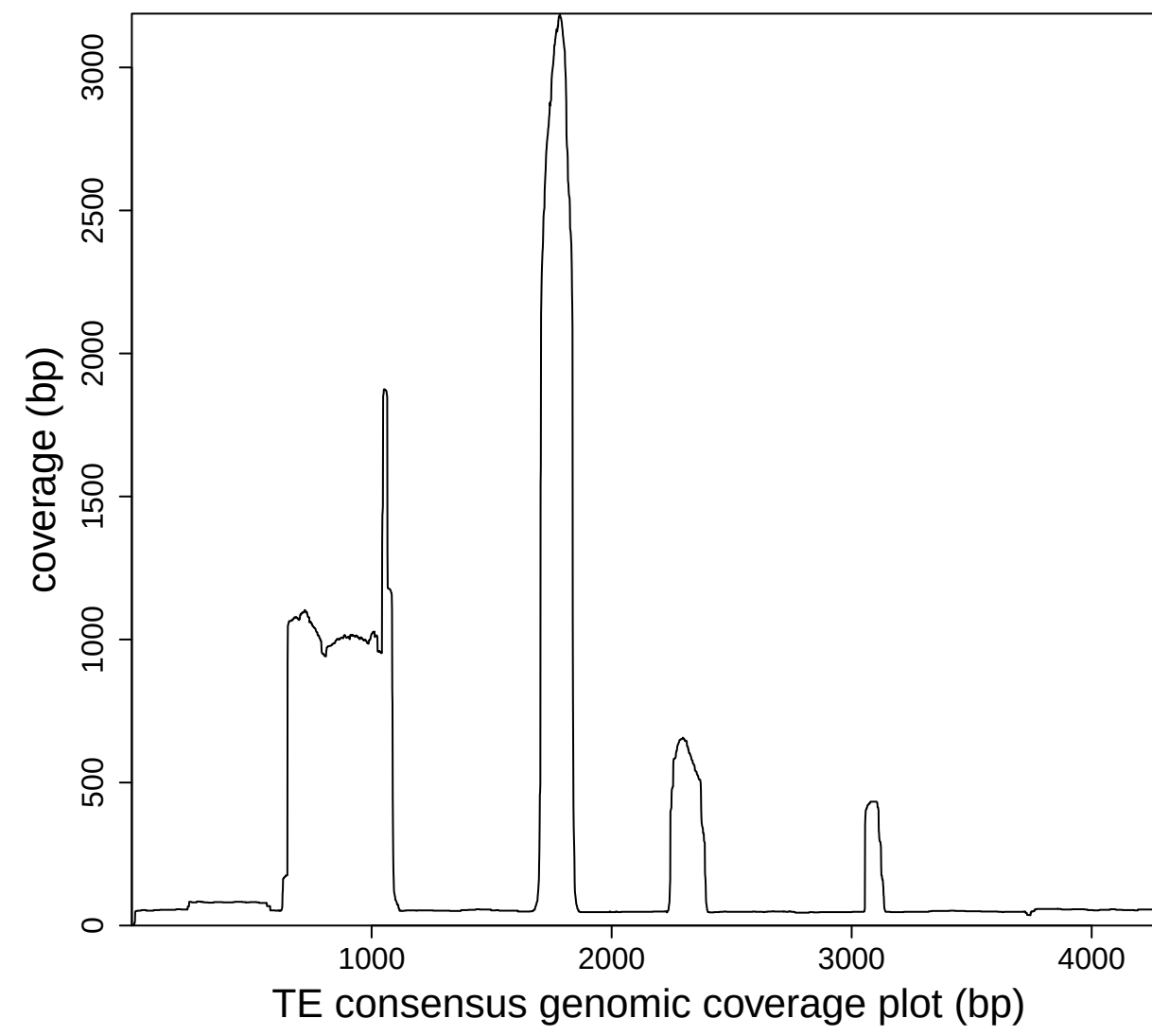
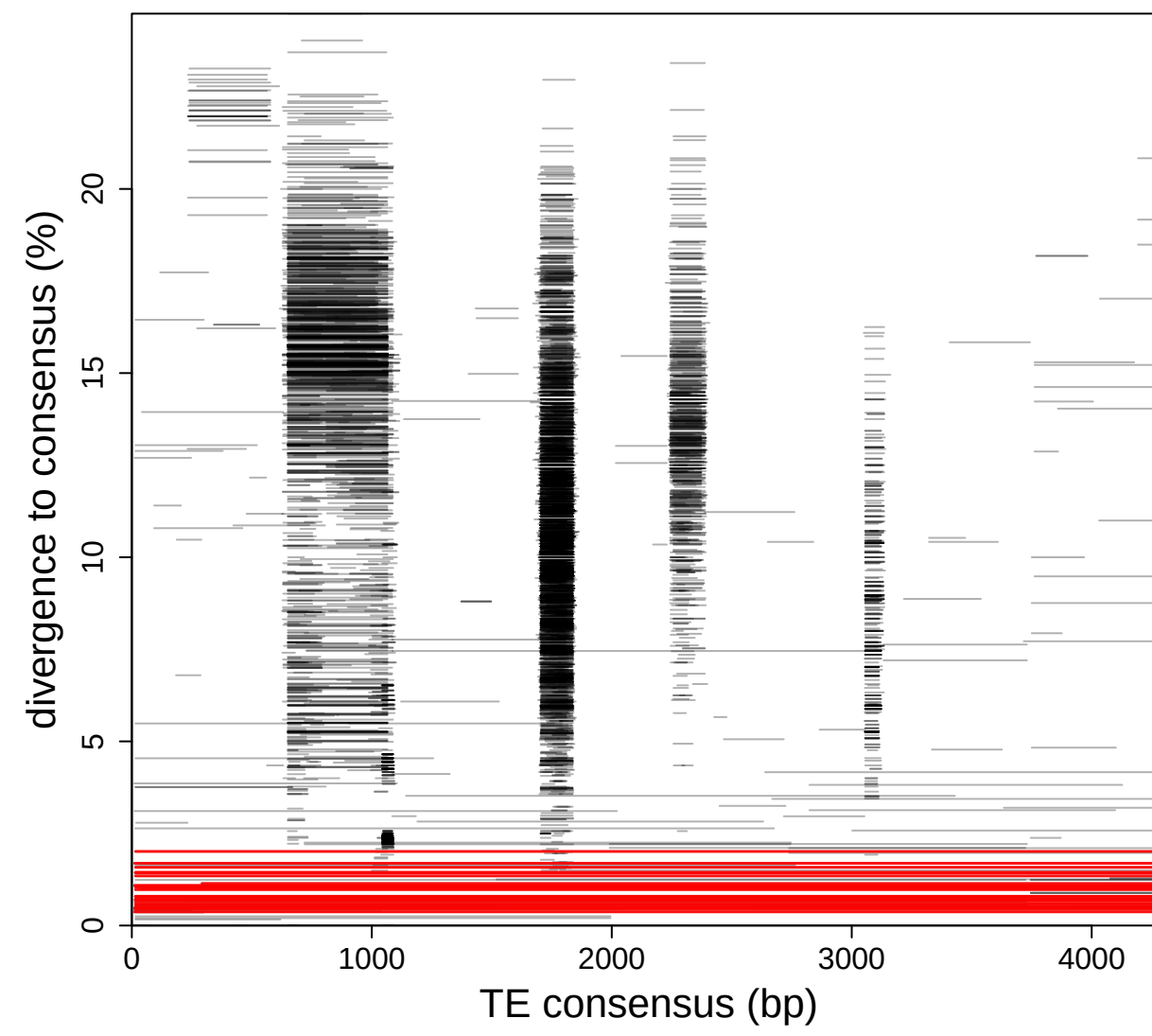
Start crop Point

End crop Point



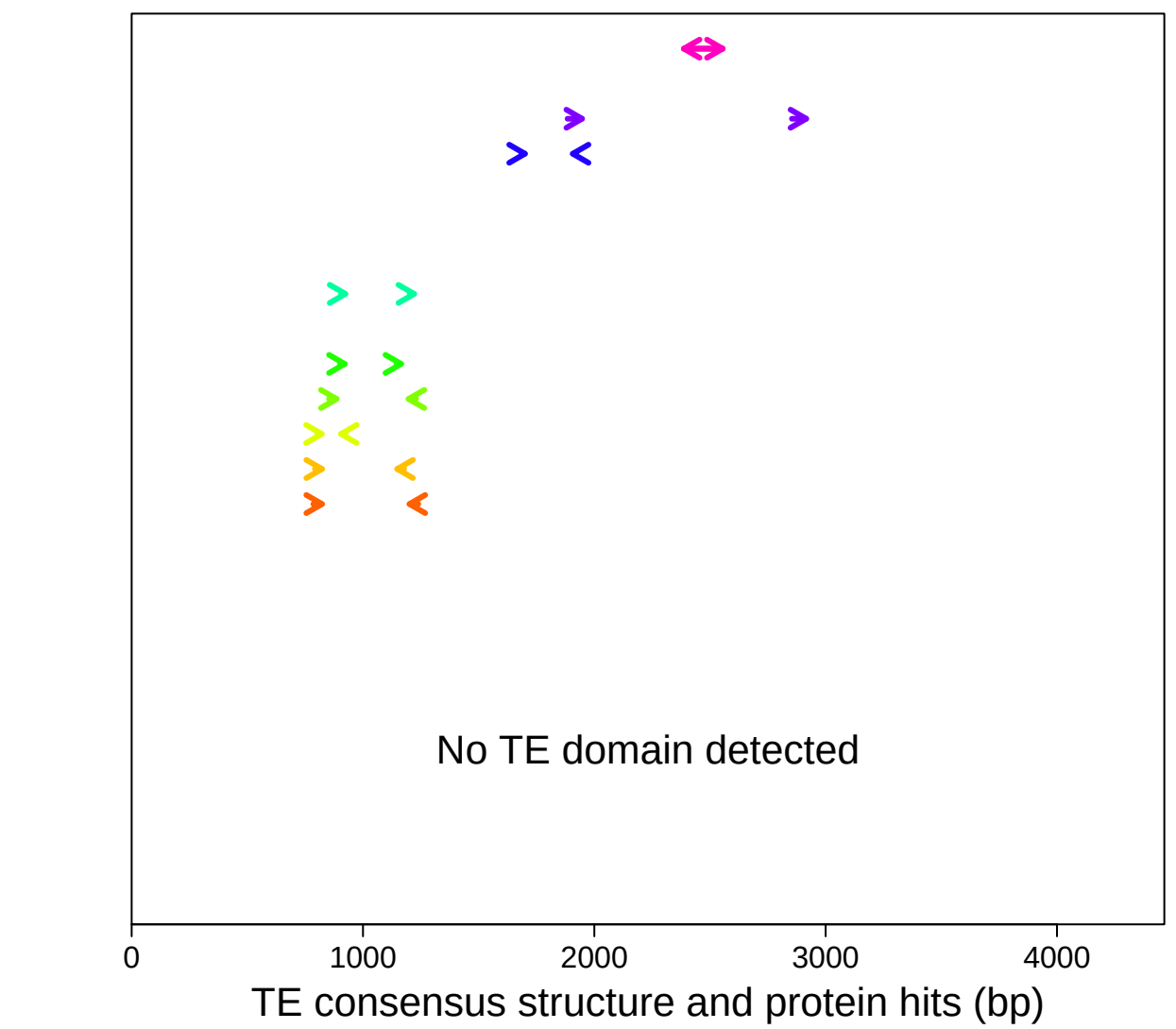
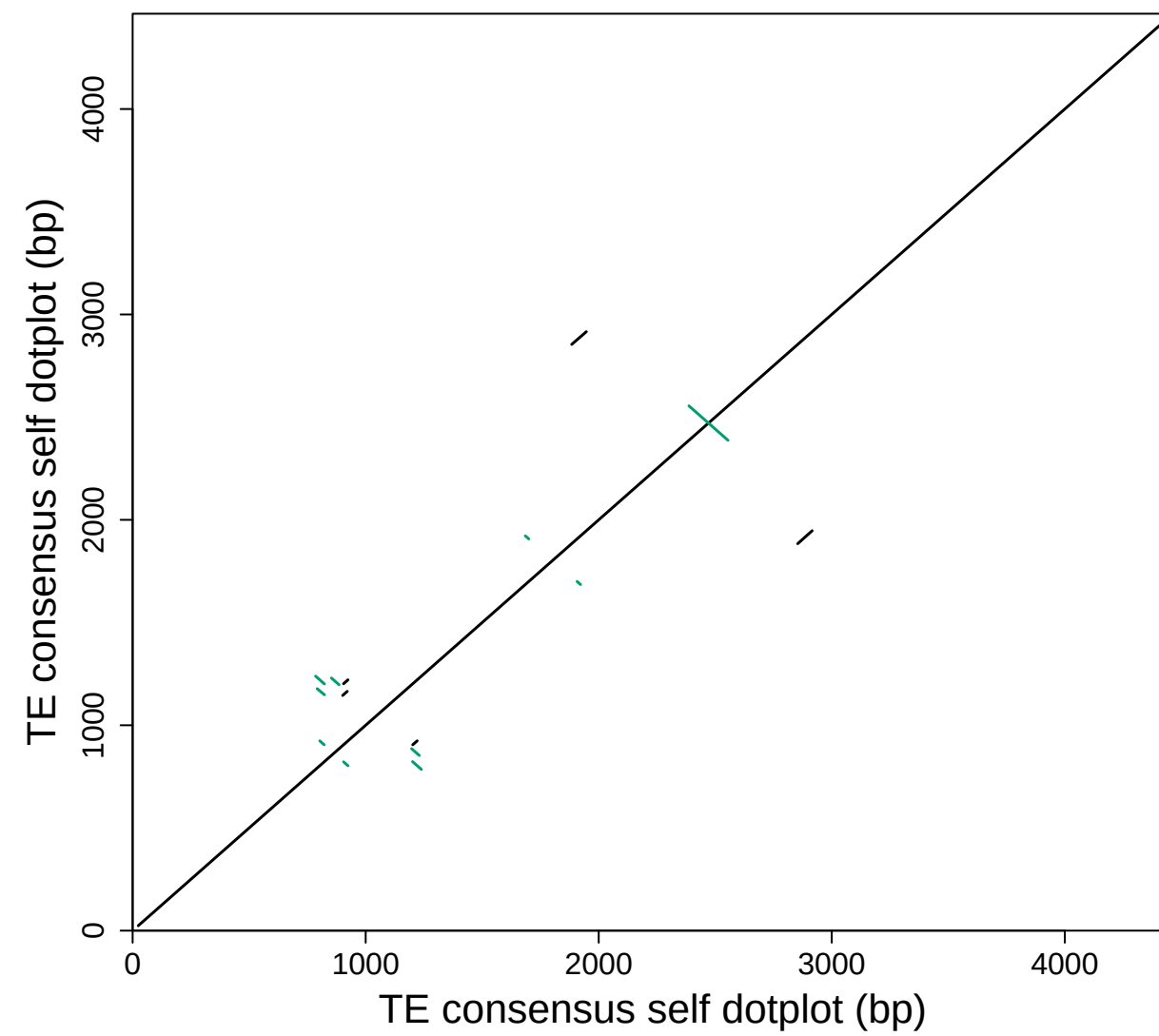
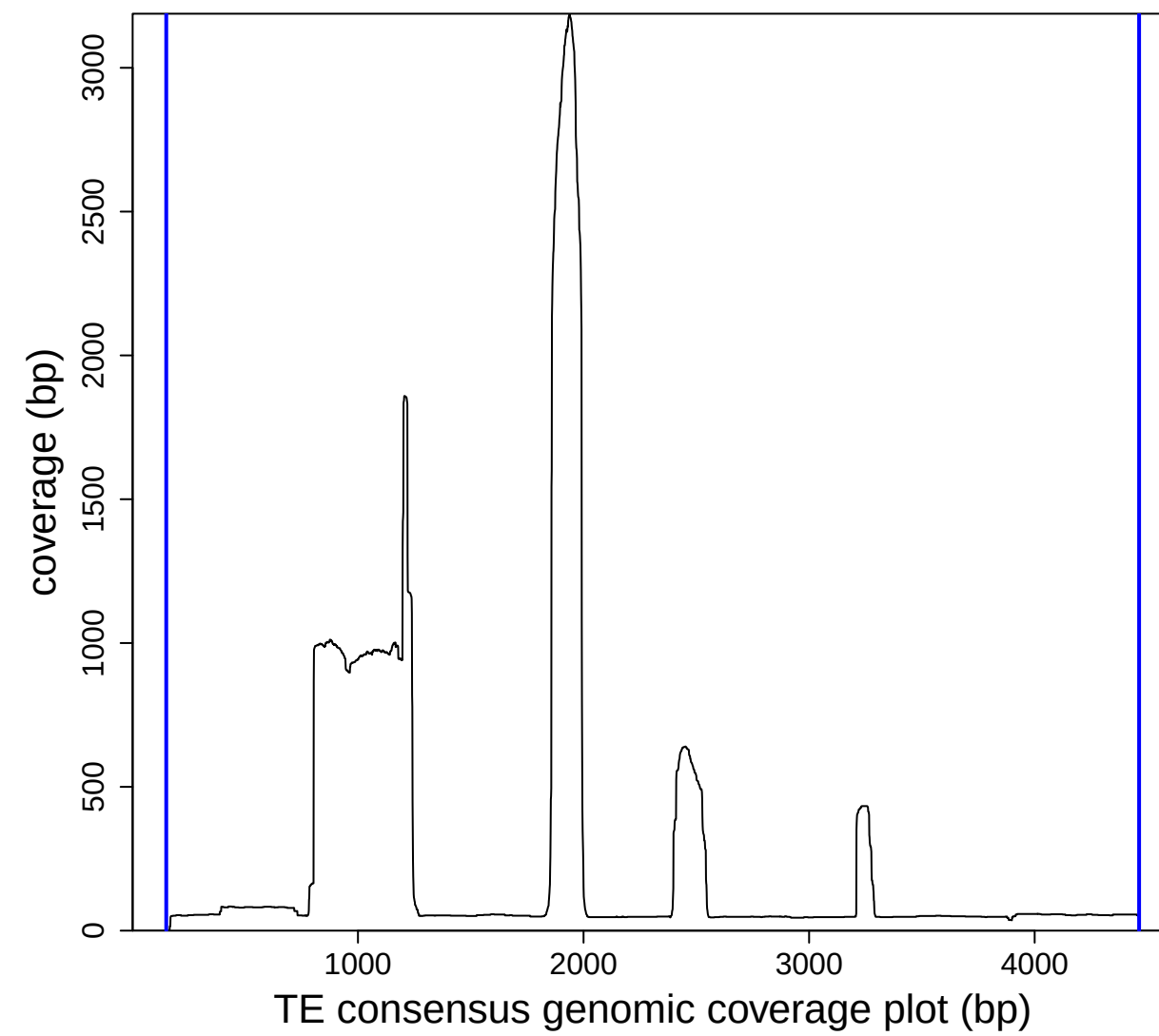
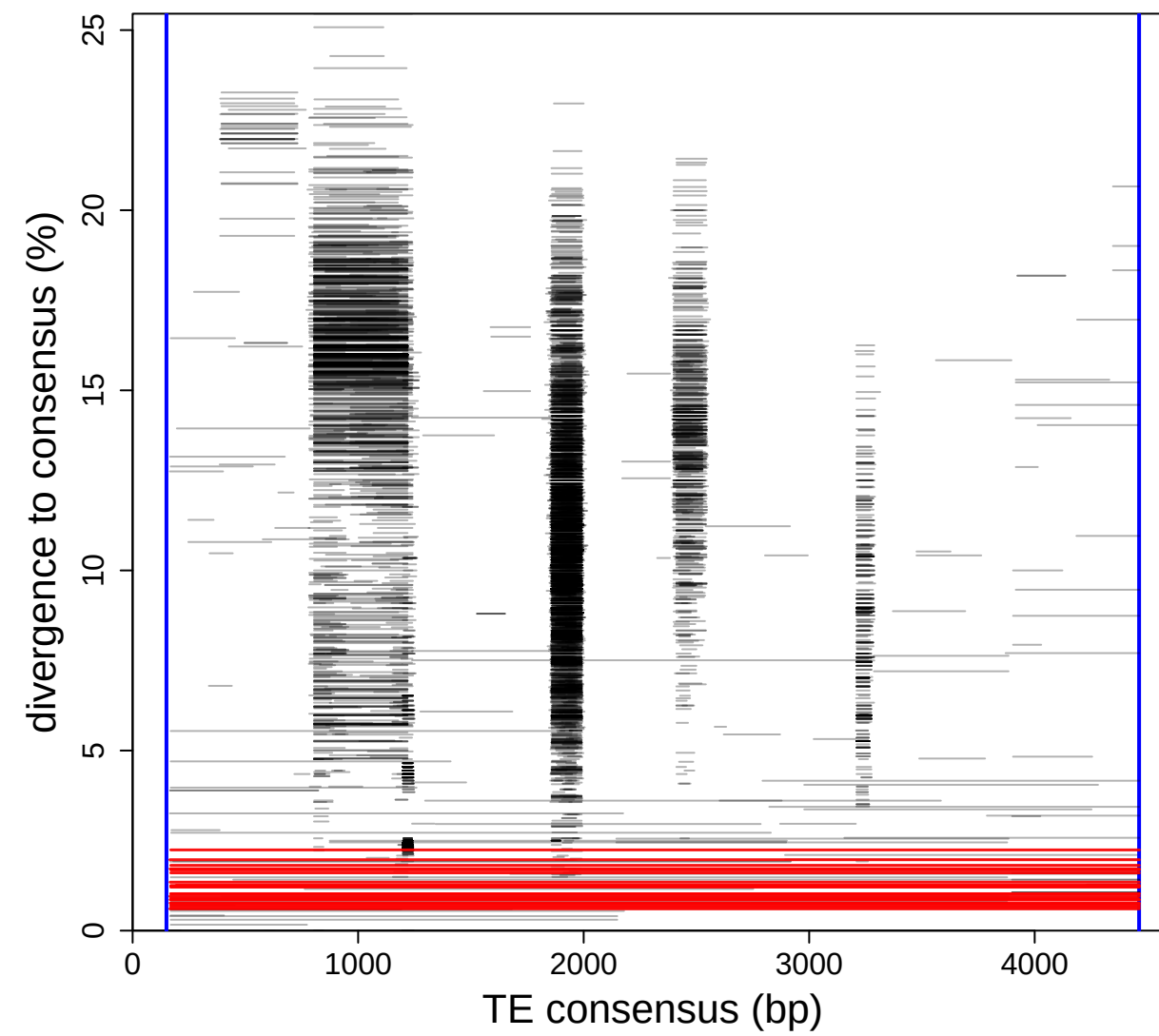
l.b.bed uf.bed g_3.bed fm_1.bed 0_0_bcln.fa aln.fa cl.fa gs.fa
size: 4309bp; fragments: 6660; full length: 24 (>=3878.1bp)

After TEtrimmer 4309 bp



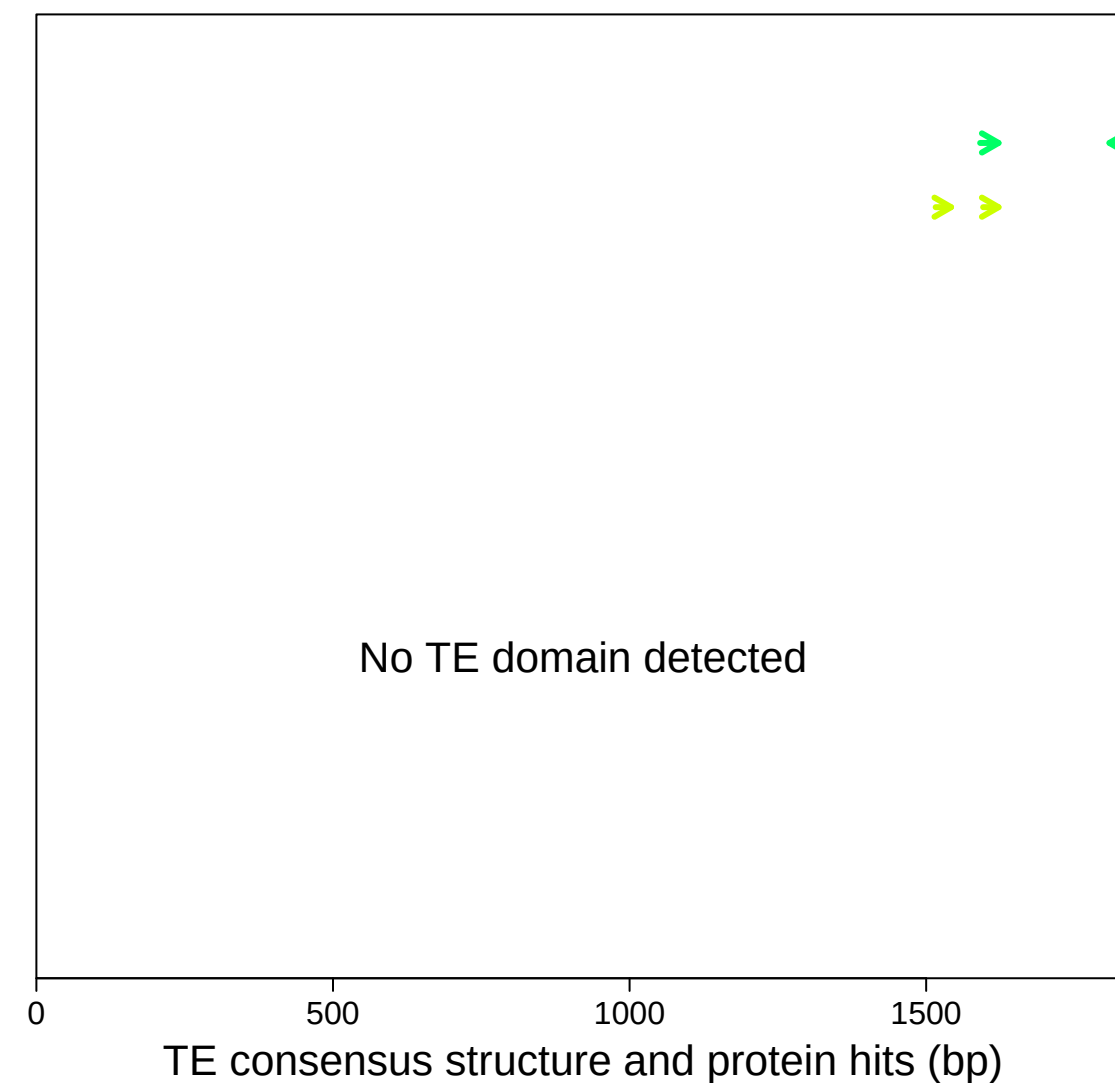
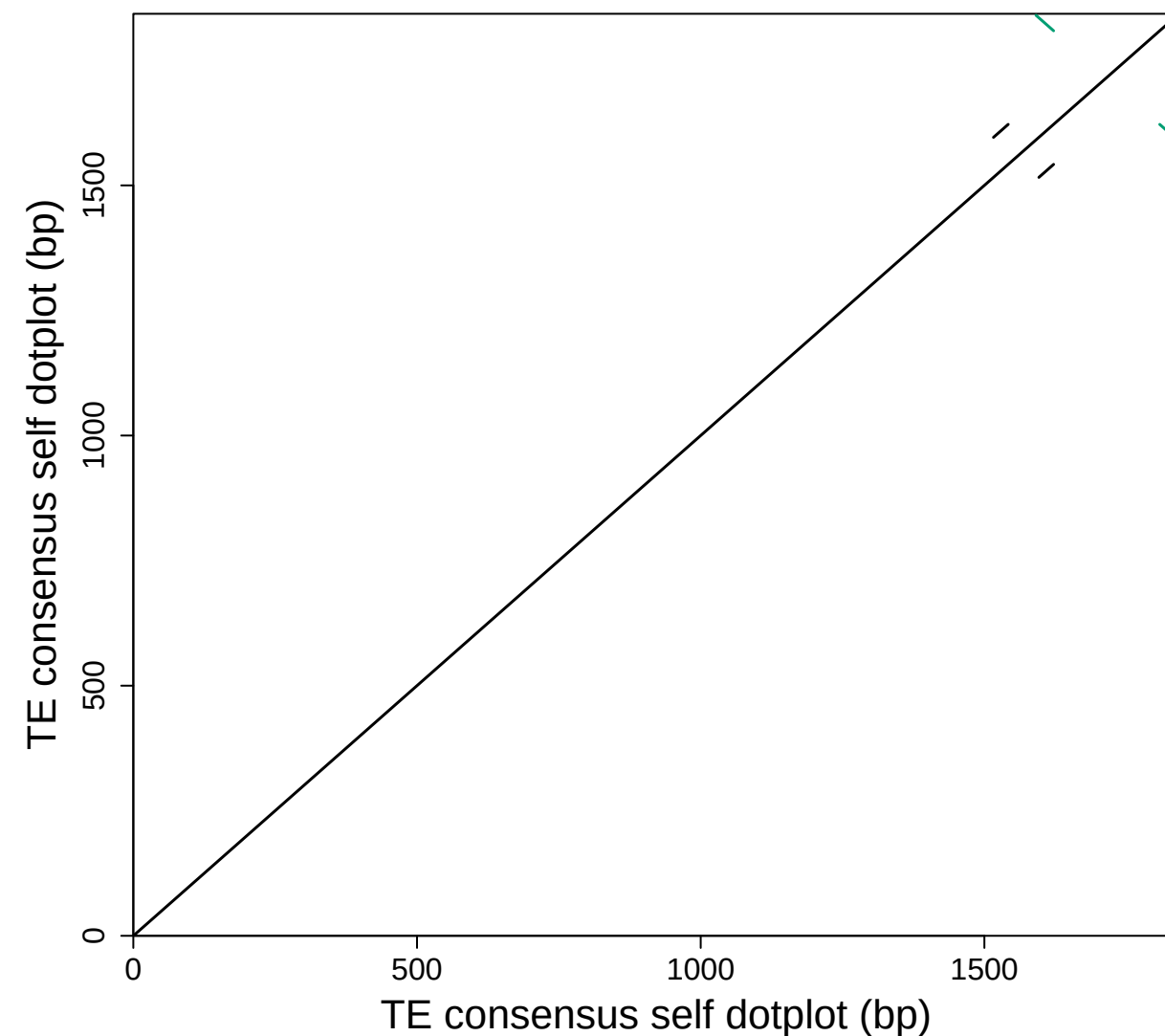
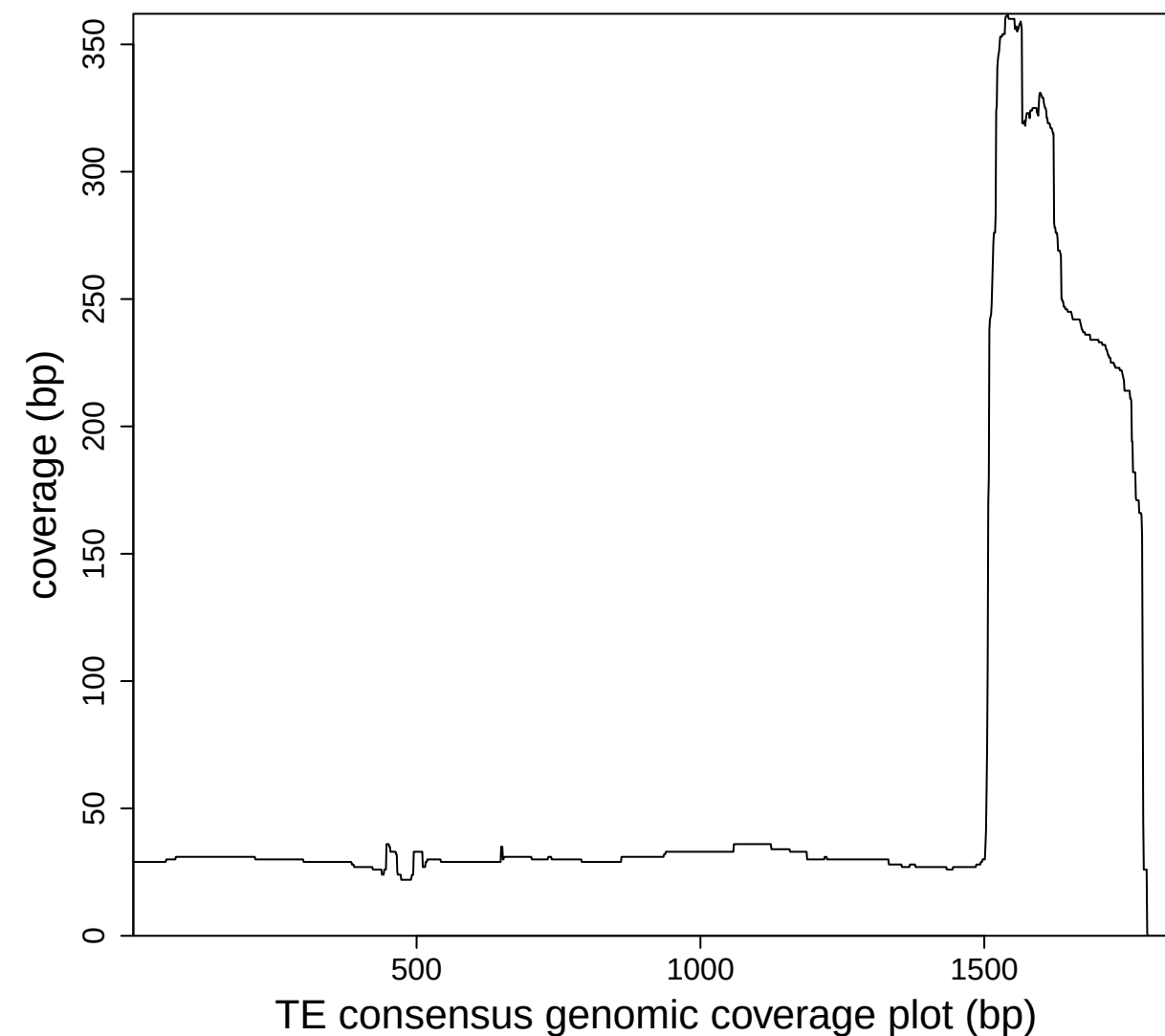
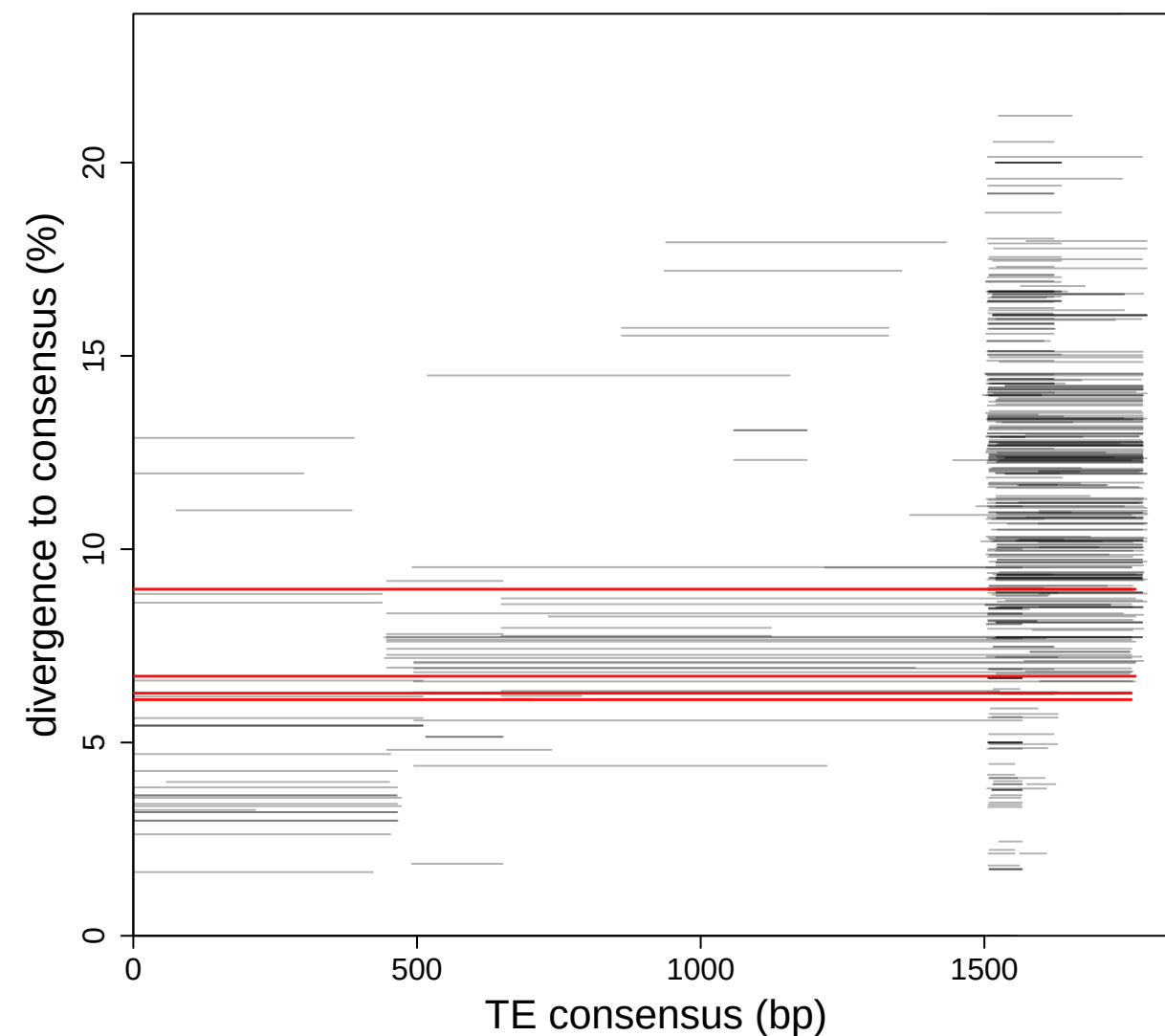
asta.b.bed uf.bed g_3.bed fm_1.bed 0_0_bcln.fa aln.fa cl.fa g
size: 4613bp; fragments: 6546; full length: 23 (>=4151.7bp)

After TEtrimmer Extended plot Blue lines are boundaries

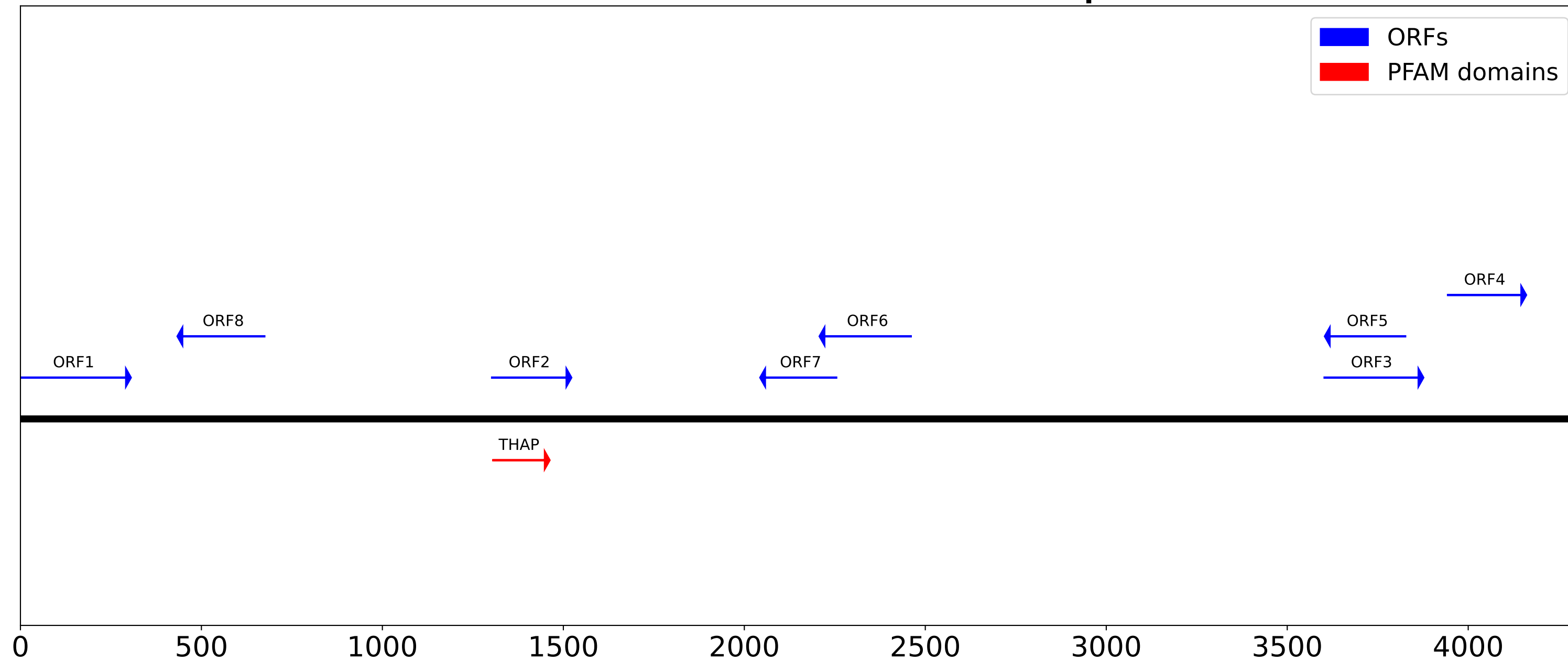


TE: rnd_4_family_278
size: 1843bp; fragments: 440; full length: 4 (≥ 1658.7 bp)

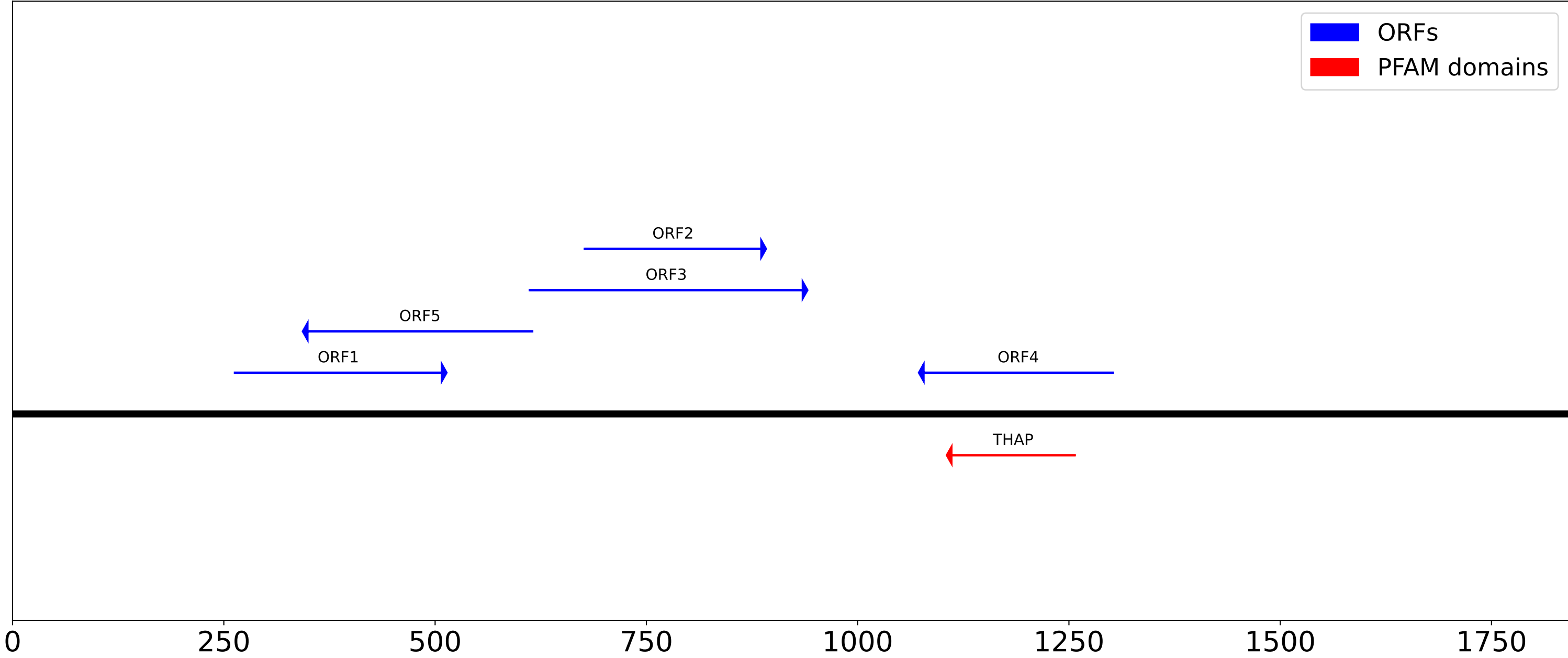
Before TETrimmer 1843 bp



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)

