

Start crop Point

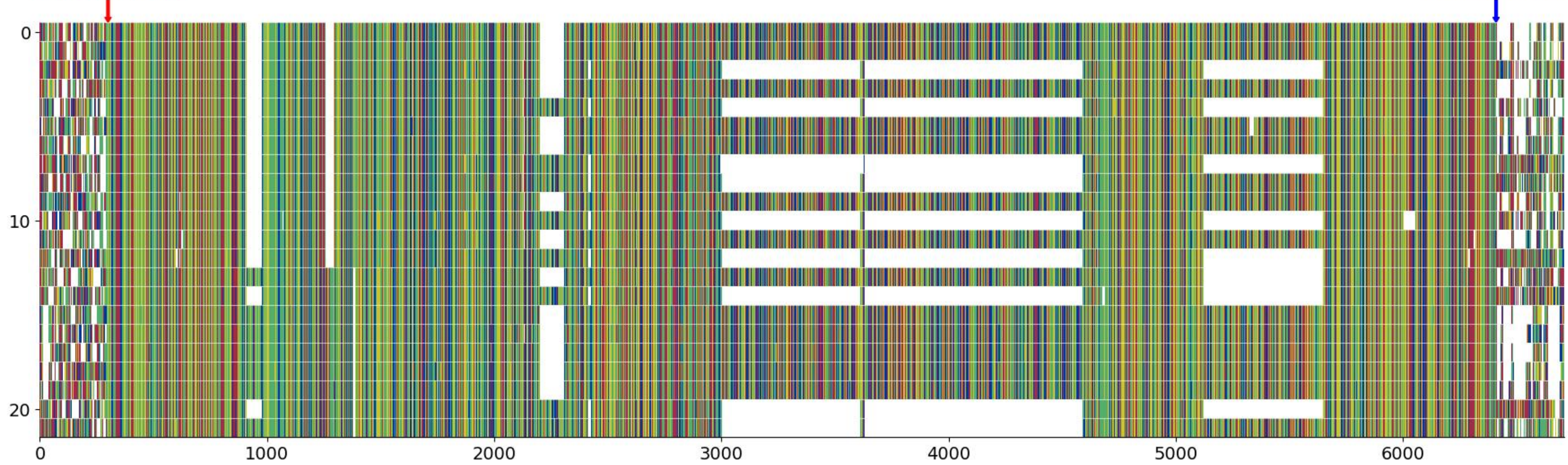
End crop Point

MSA length = 6111



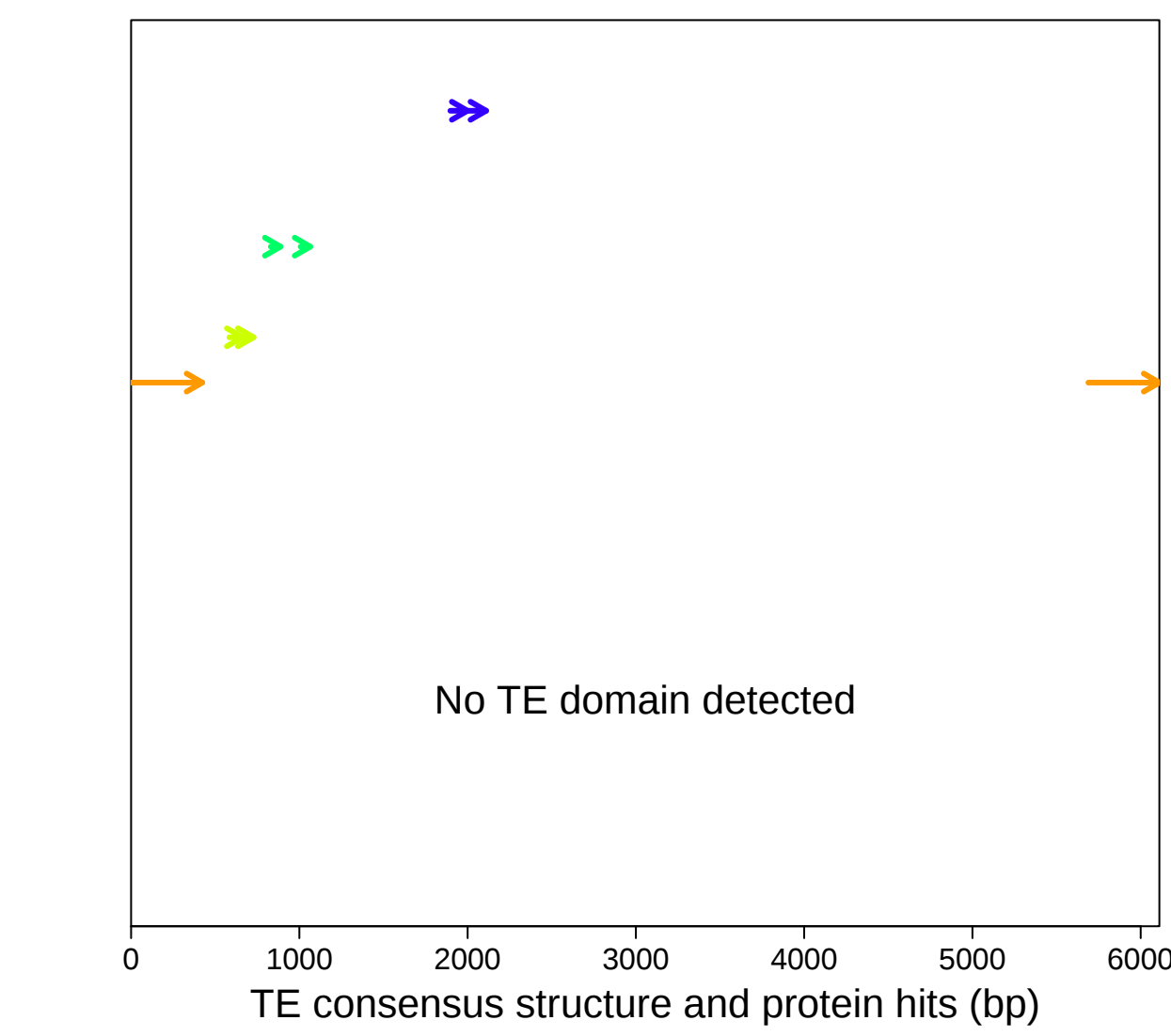
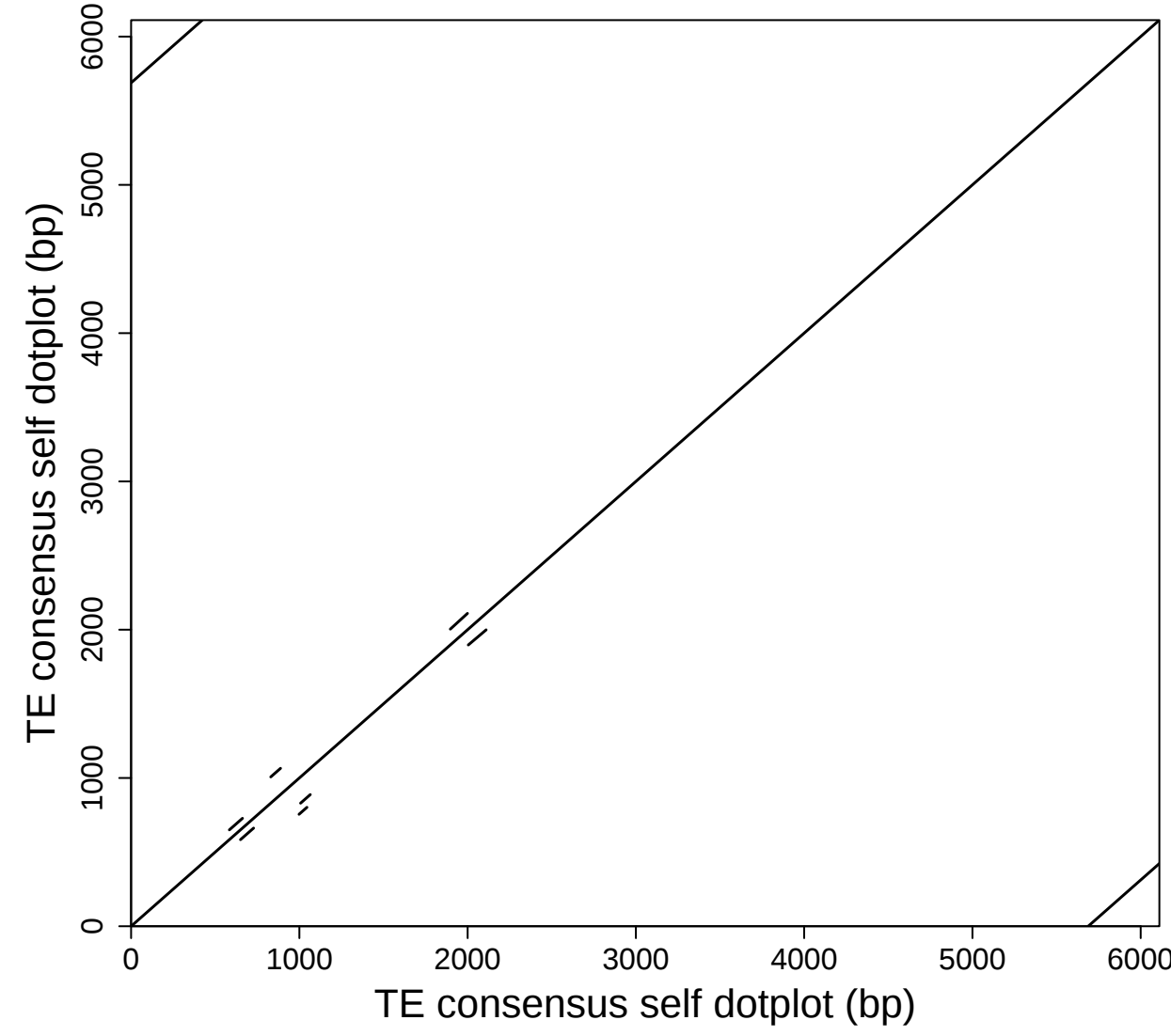
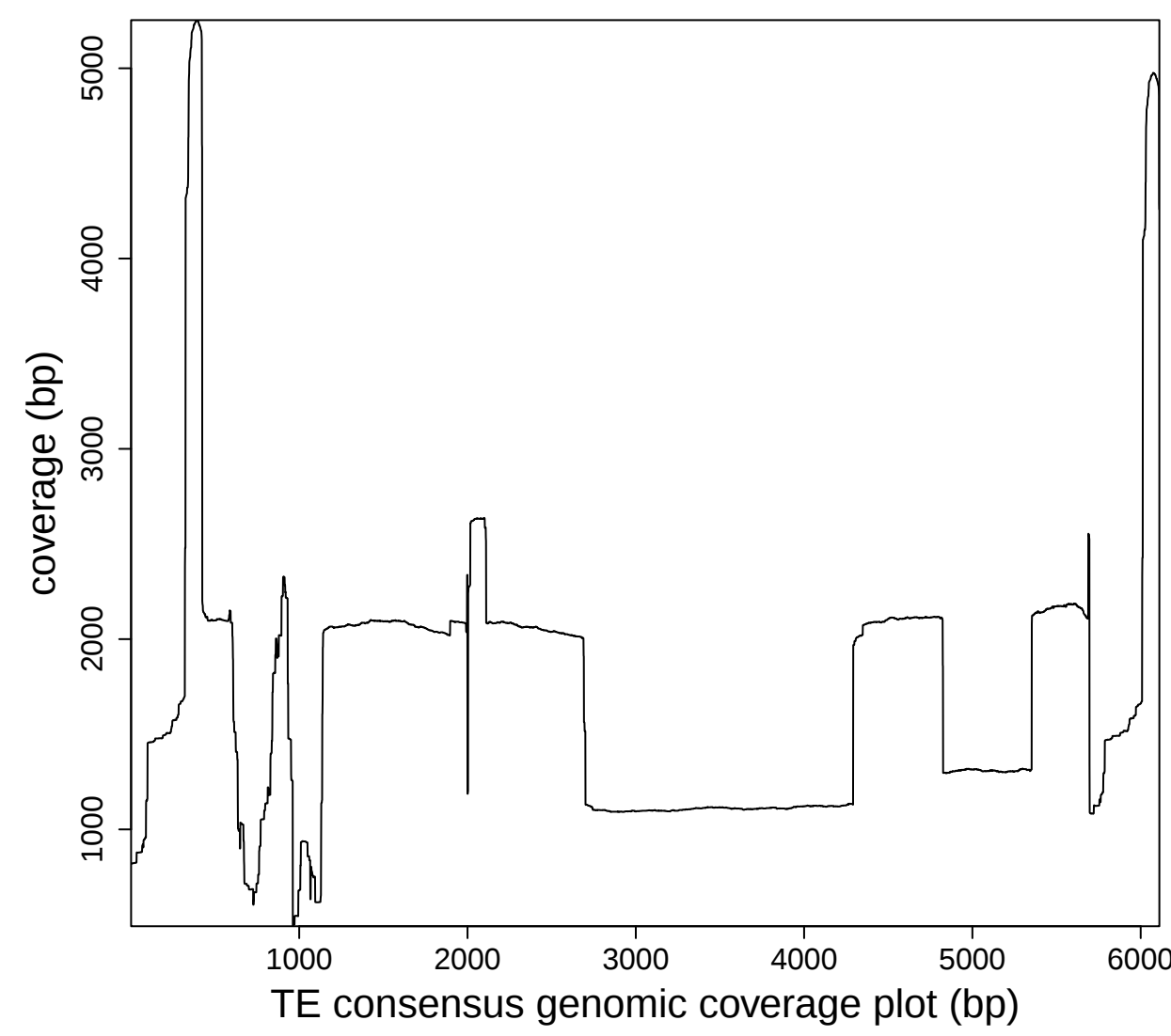
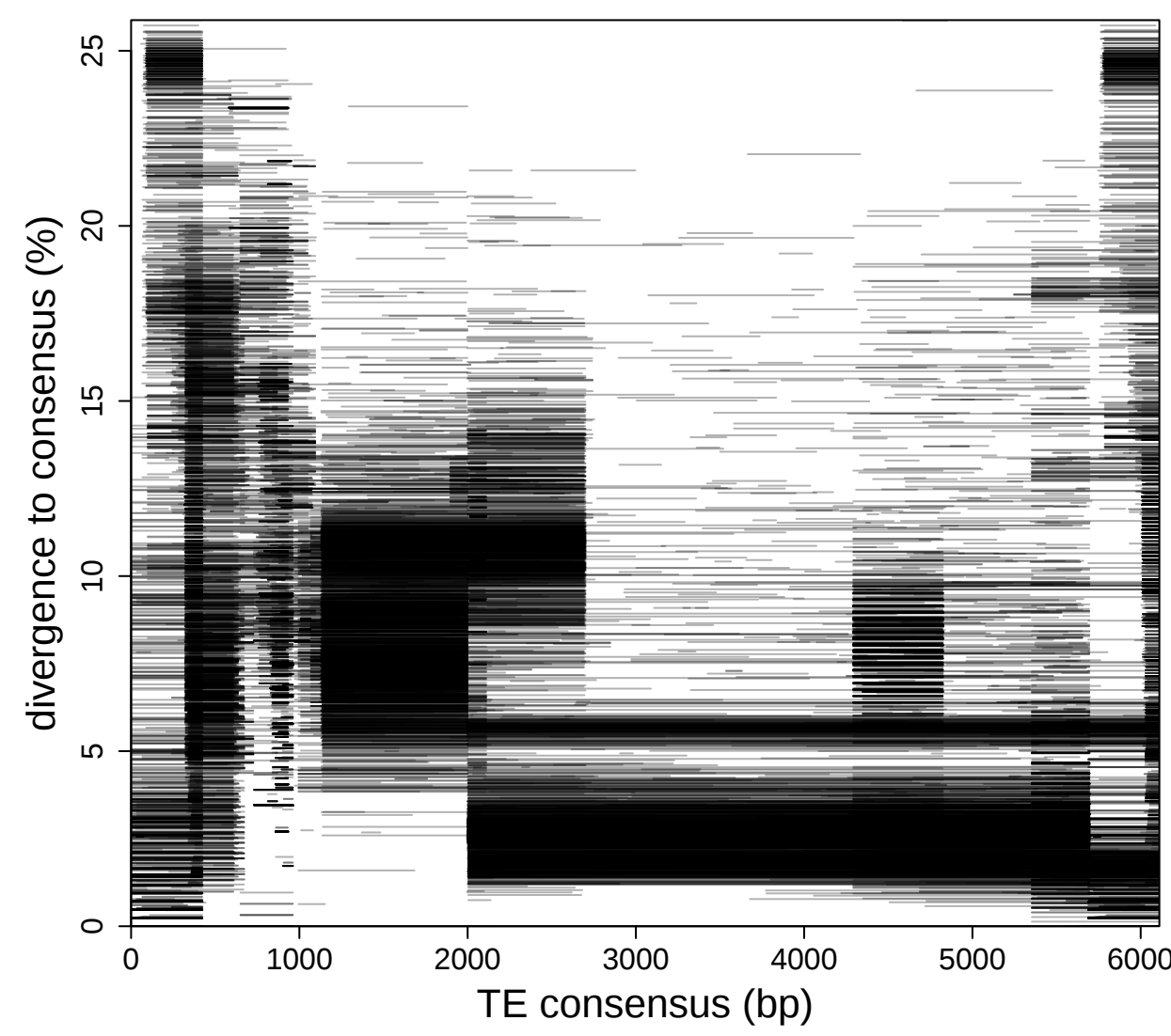
Start crop Point

End crop Point



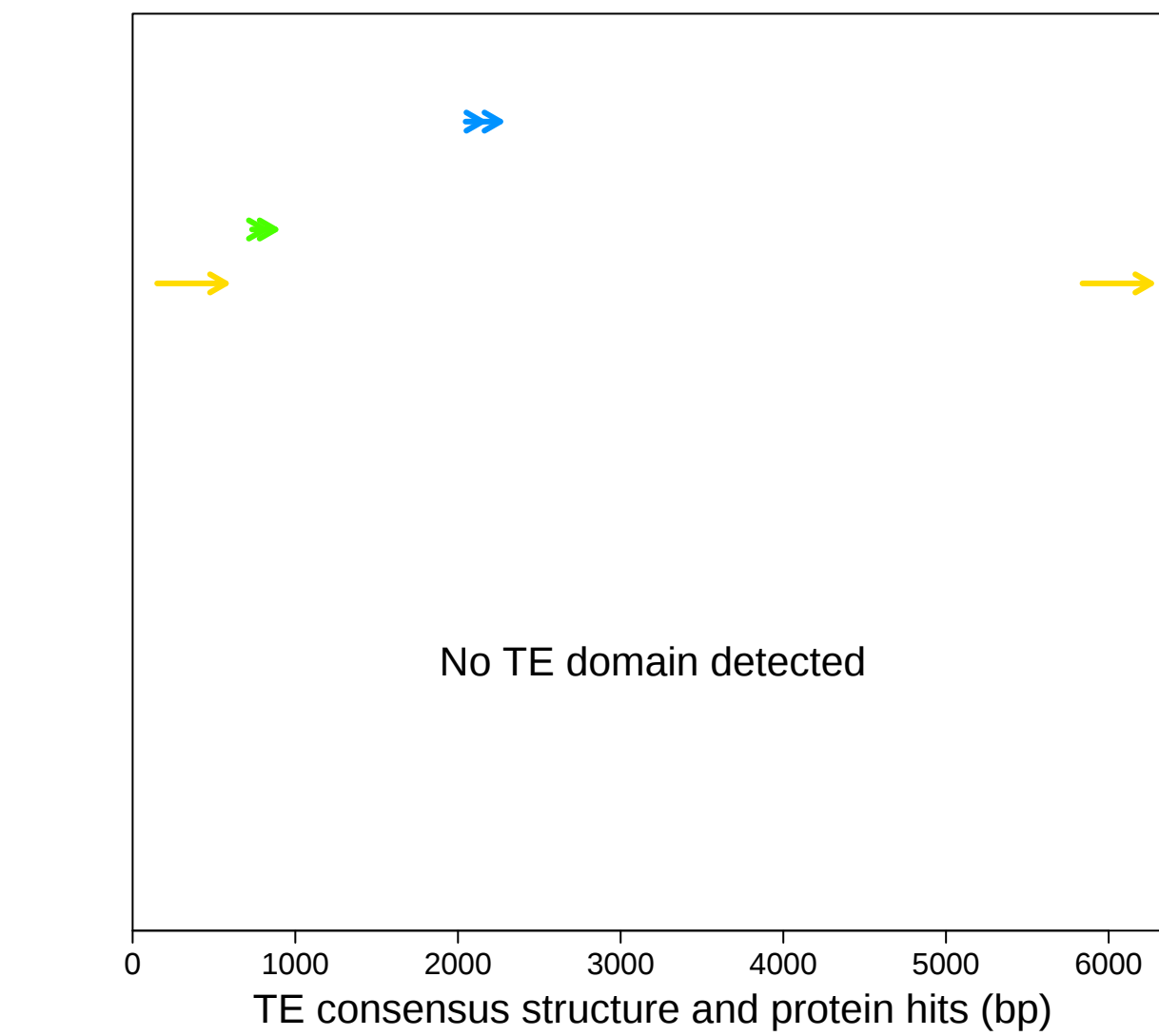
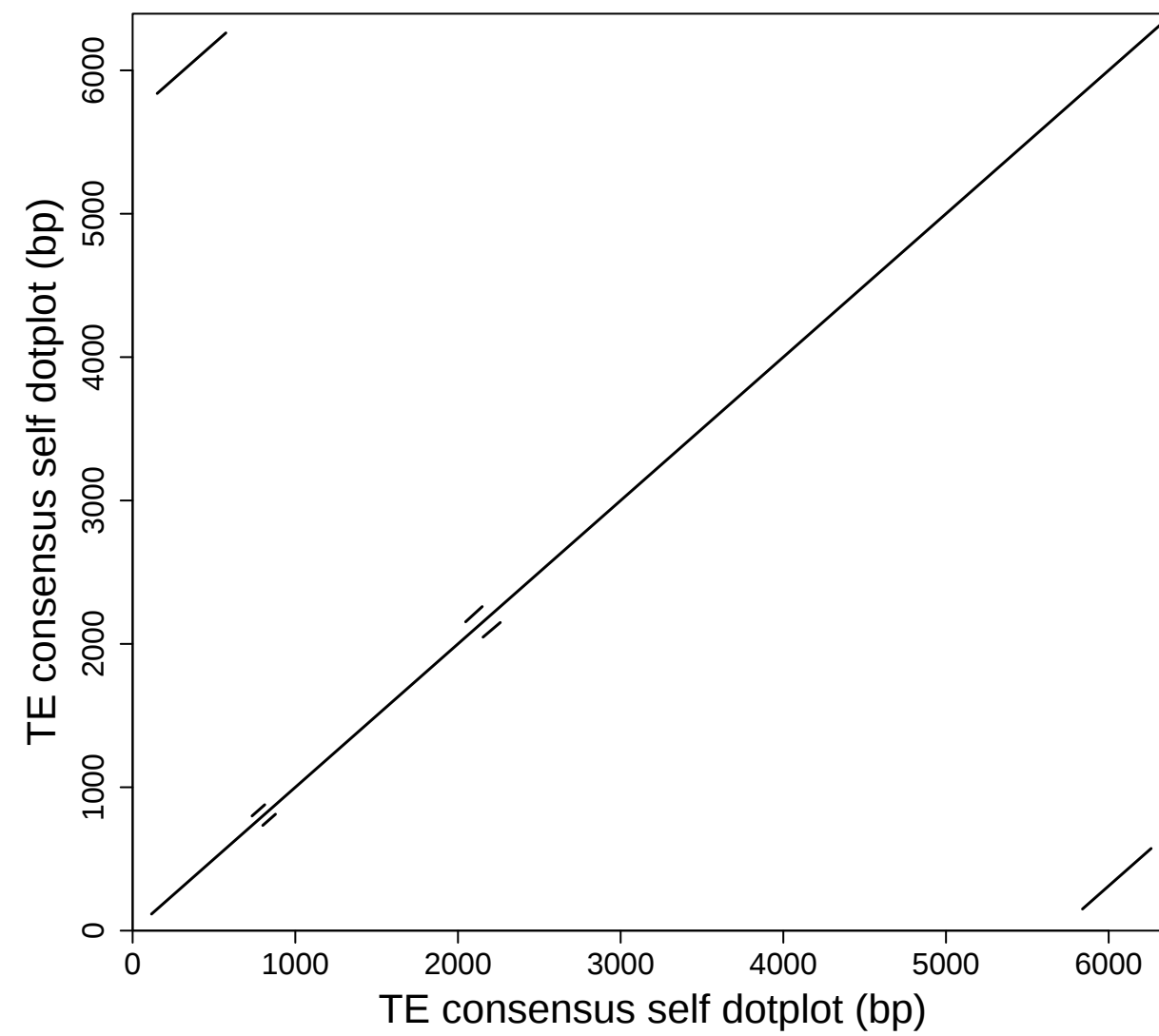
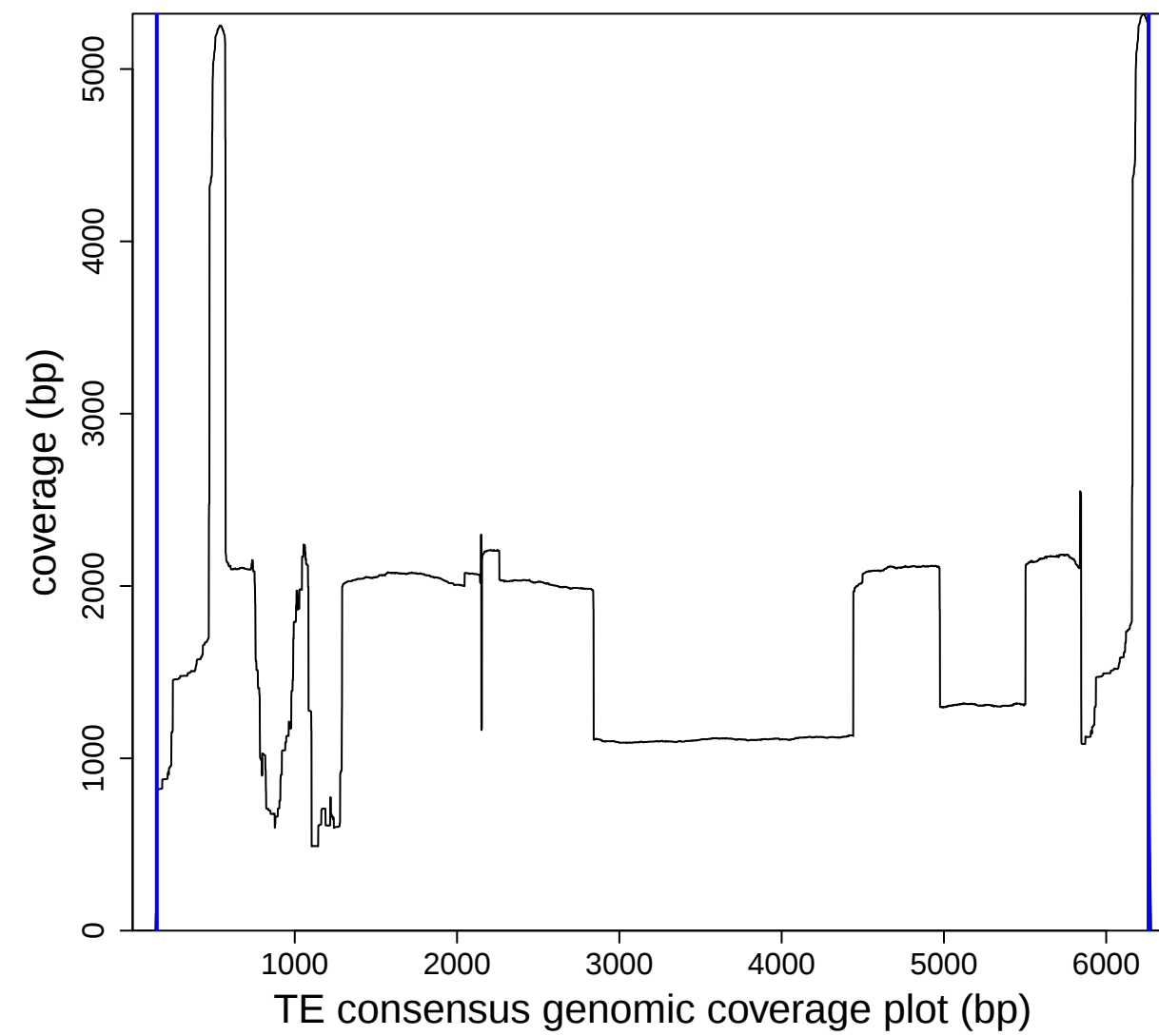
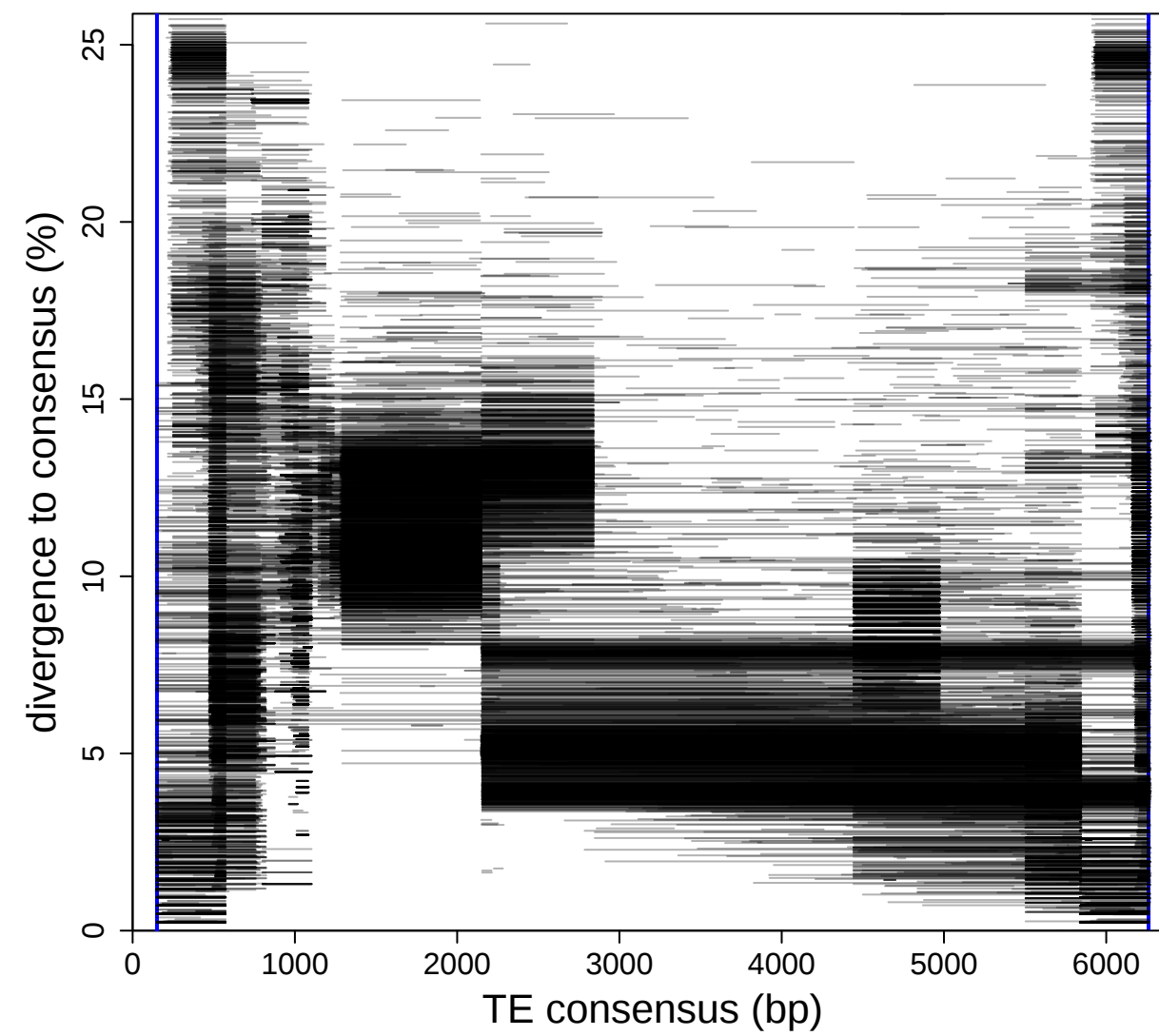
bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa_c
size: 6111bp; fragments: 18598; full length: 0 (>=5499.9bp)

After TETrimmer 6111 bp



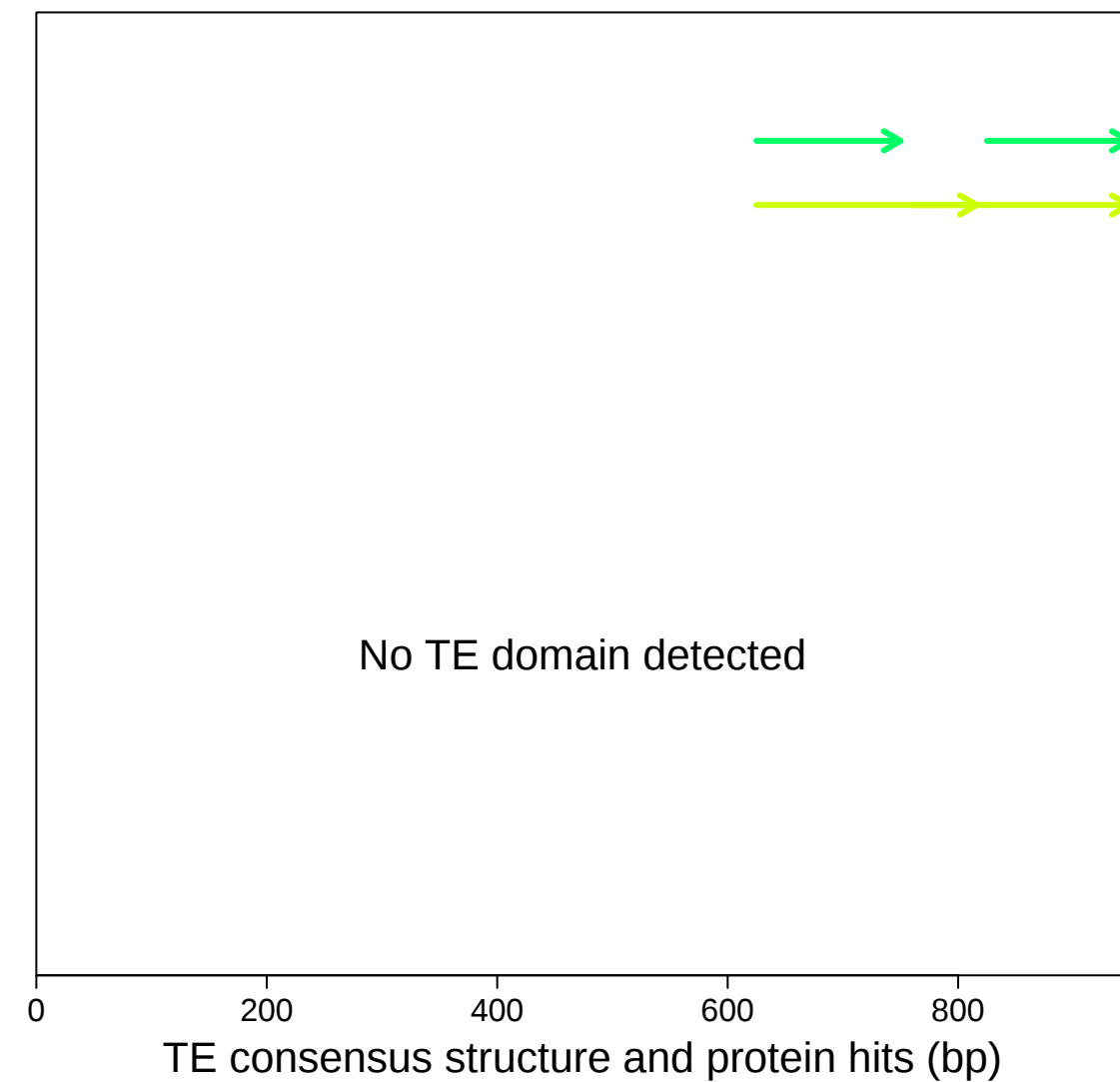
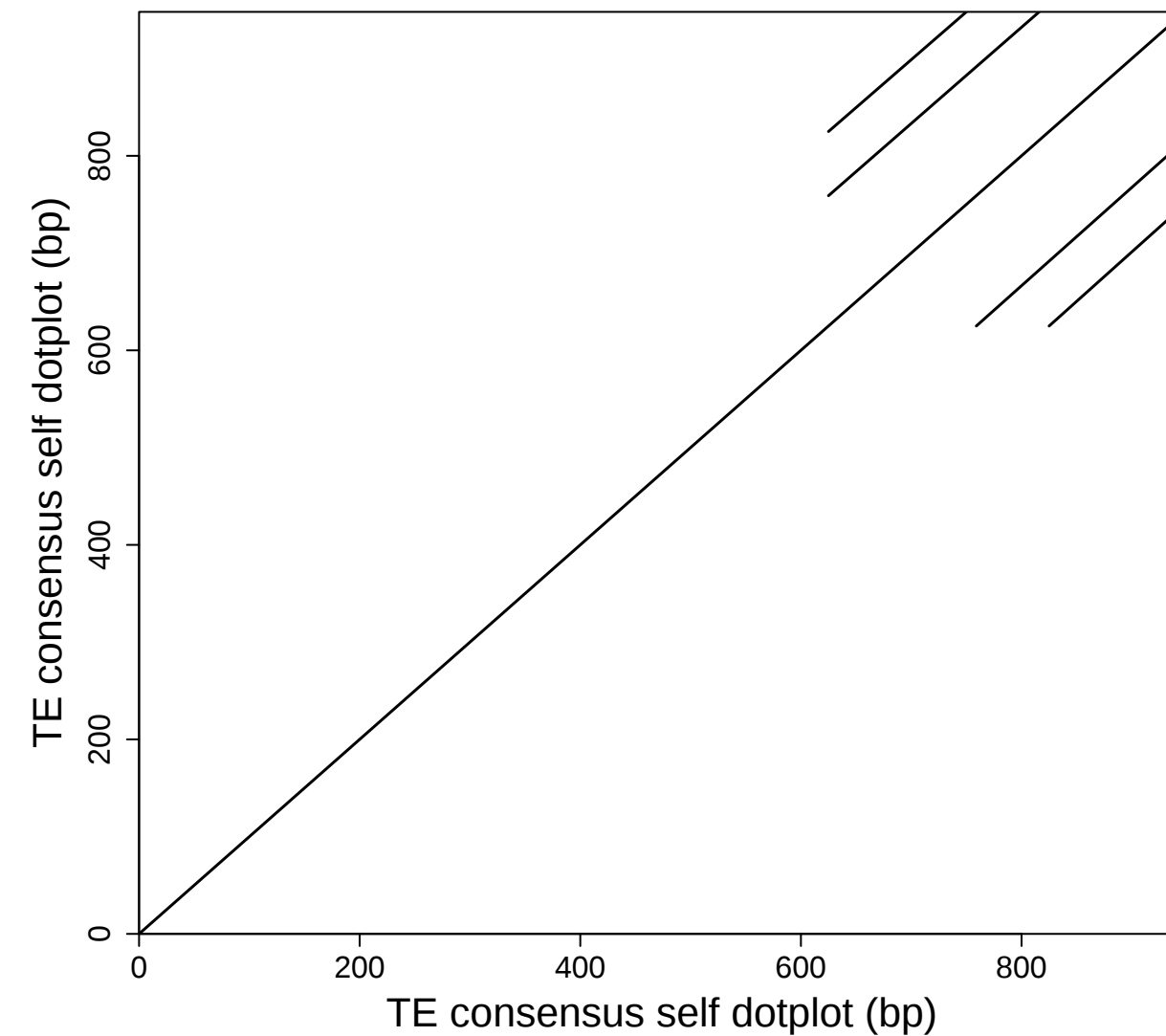
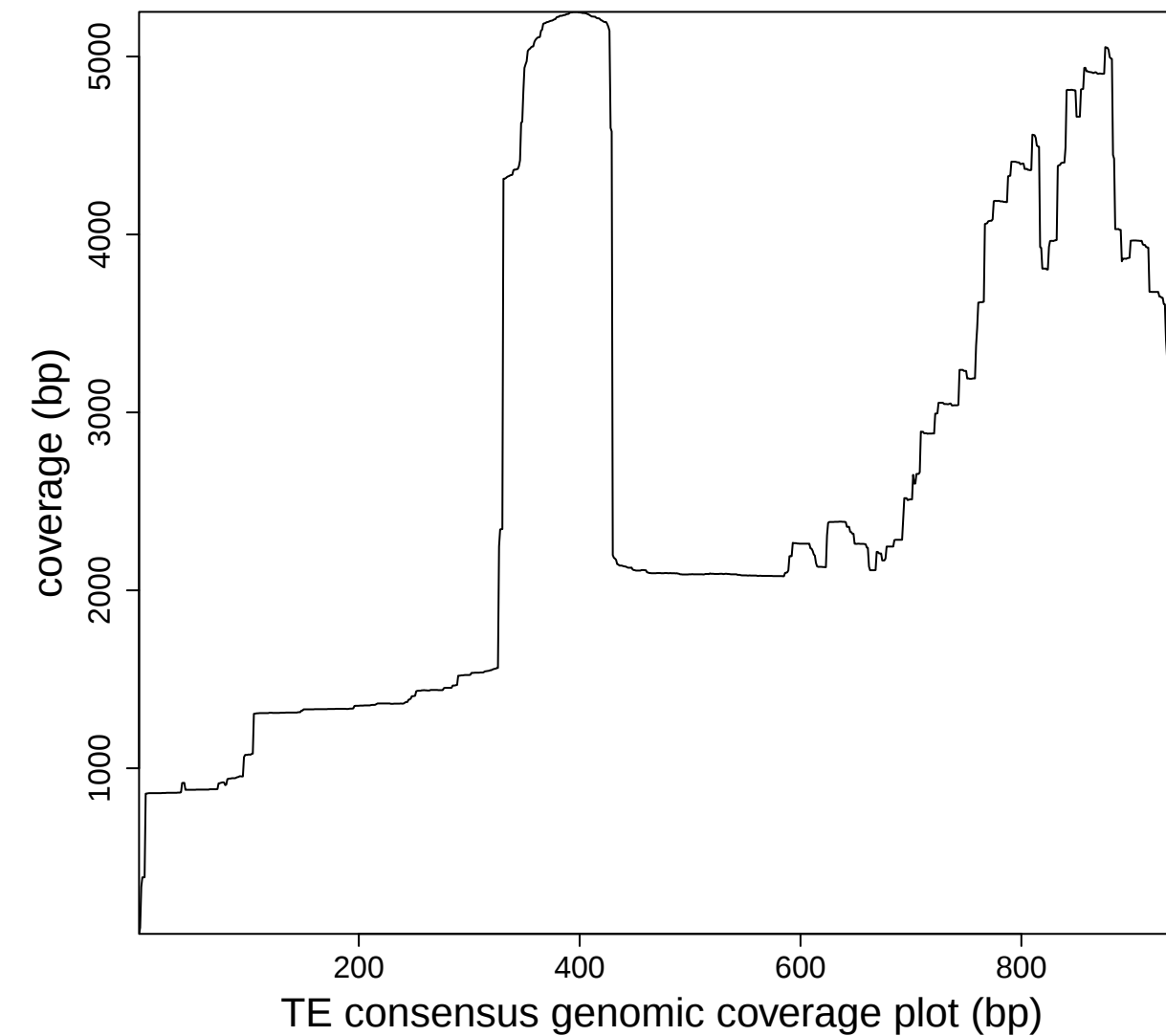
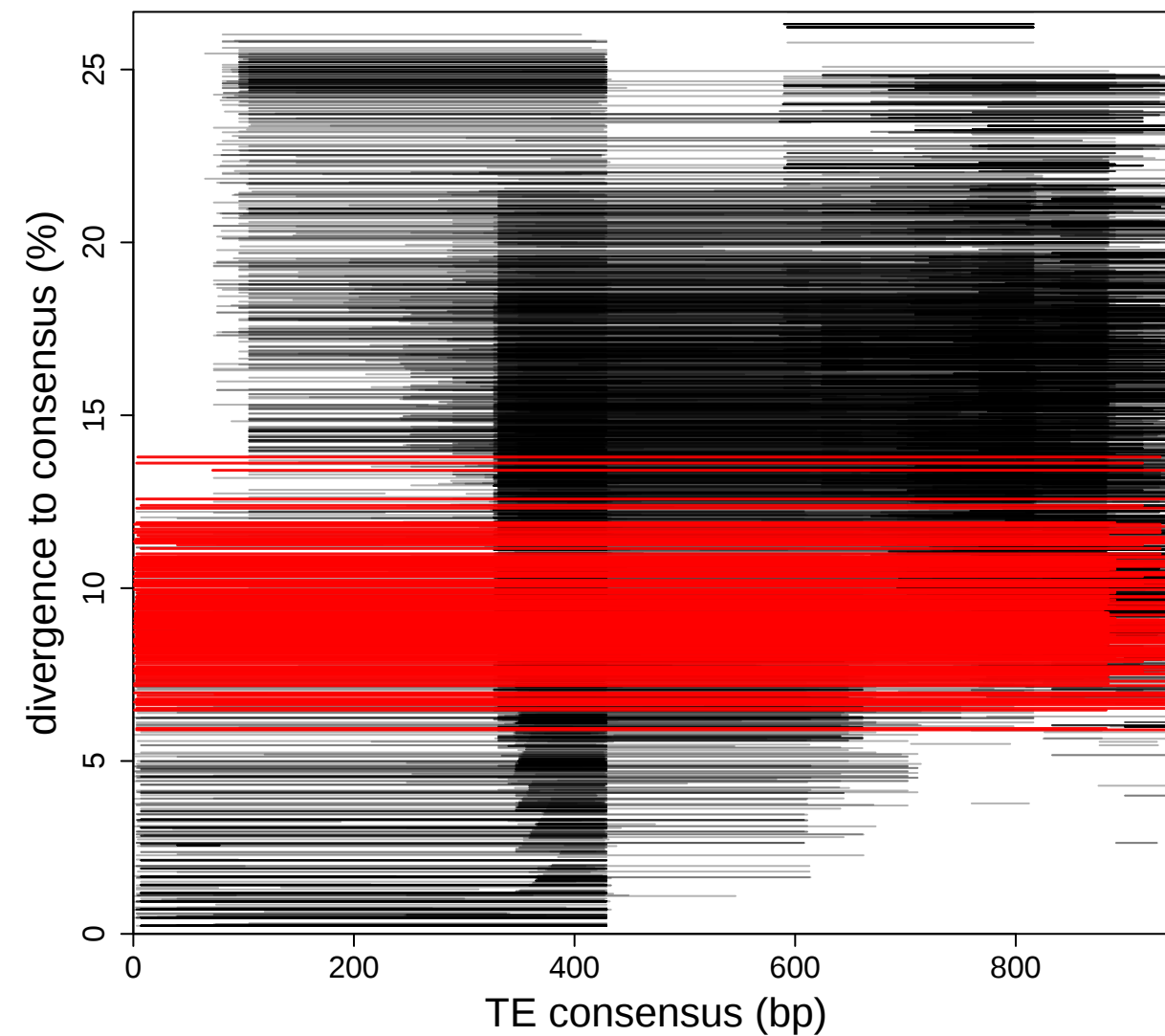
ly_1353.fasta.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.f
size: 6411bp; fragments: 18112; full length: 0 (>=5769.9bp)

After TEtrimmer Extended plot Blue lines are boundaries

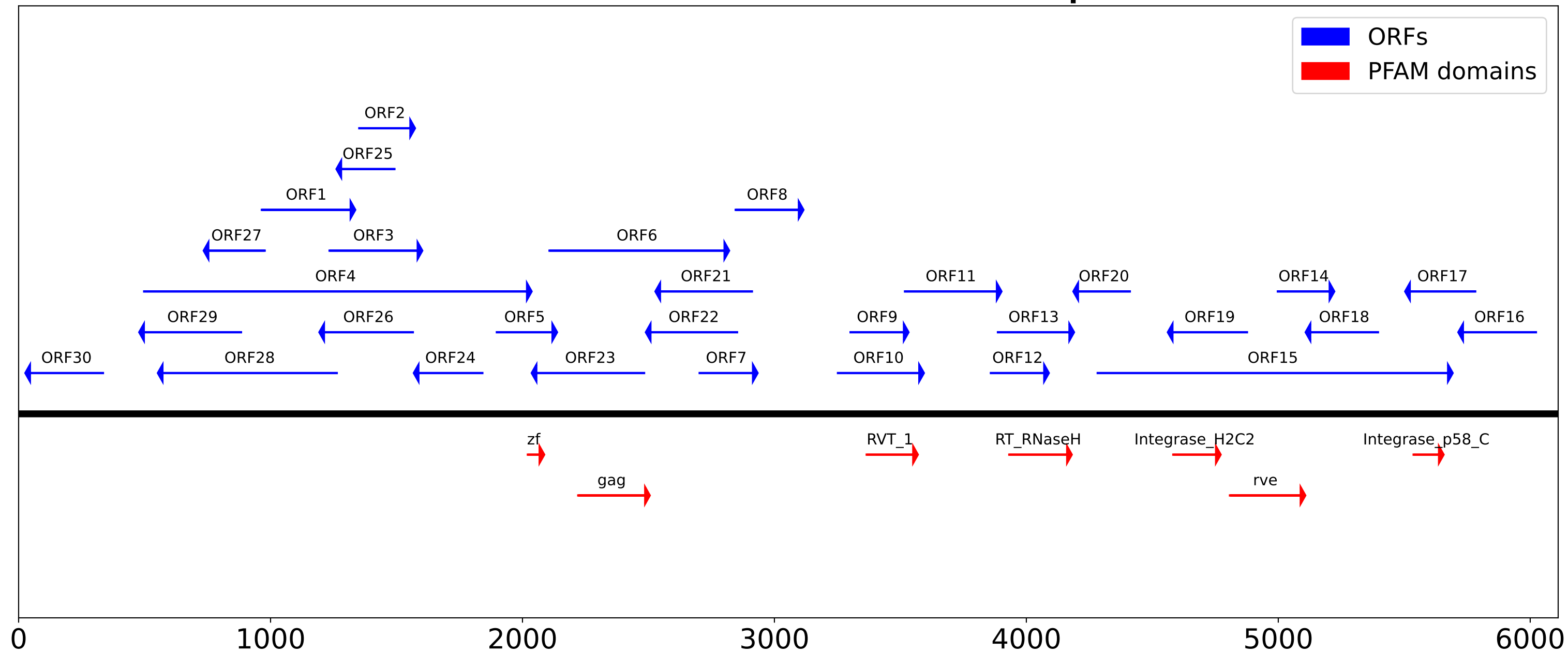


TE: rnd_5_family_1353
size: 948bp; fragments: 10200; full length: 222 (≥ 853.2 bp)

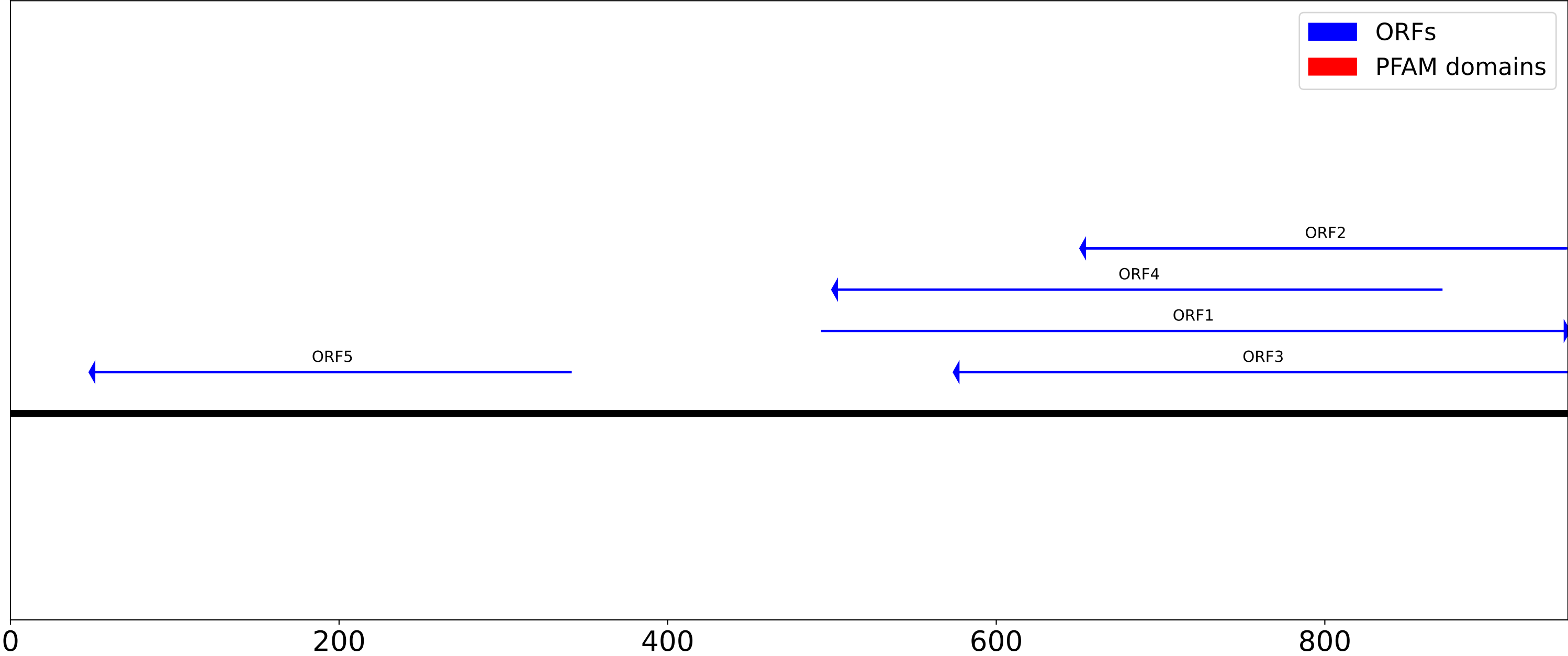
Before TEtrimmer 948 bp



After TETrimmer ORF and PFAM domain plot



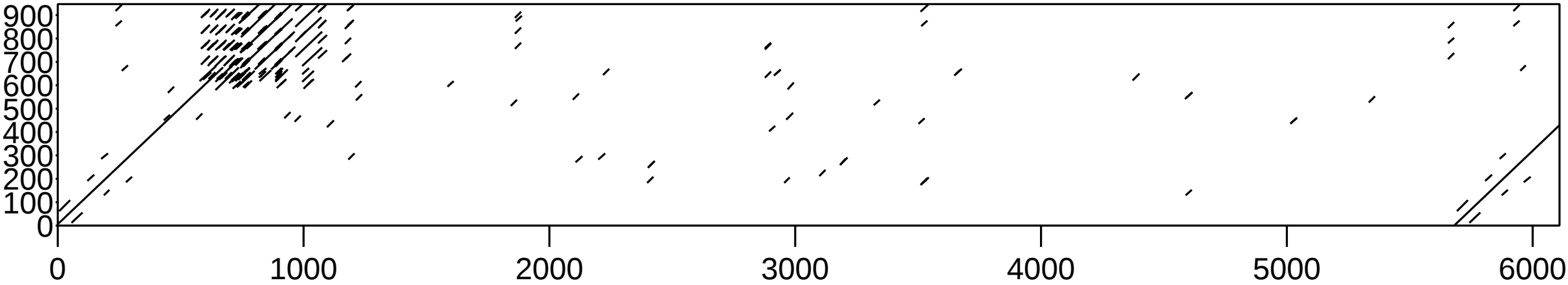
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

1sensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)