

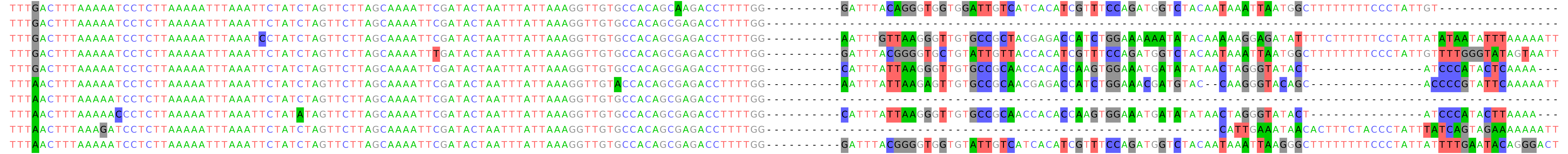
Start Crop Point



End Crop Point

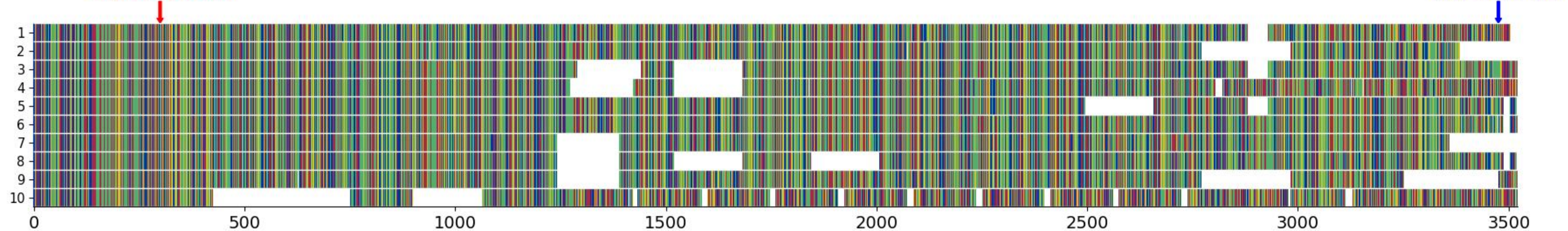


MSA length = 3177



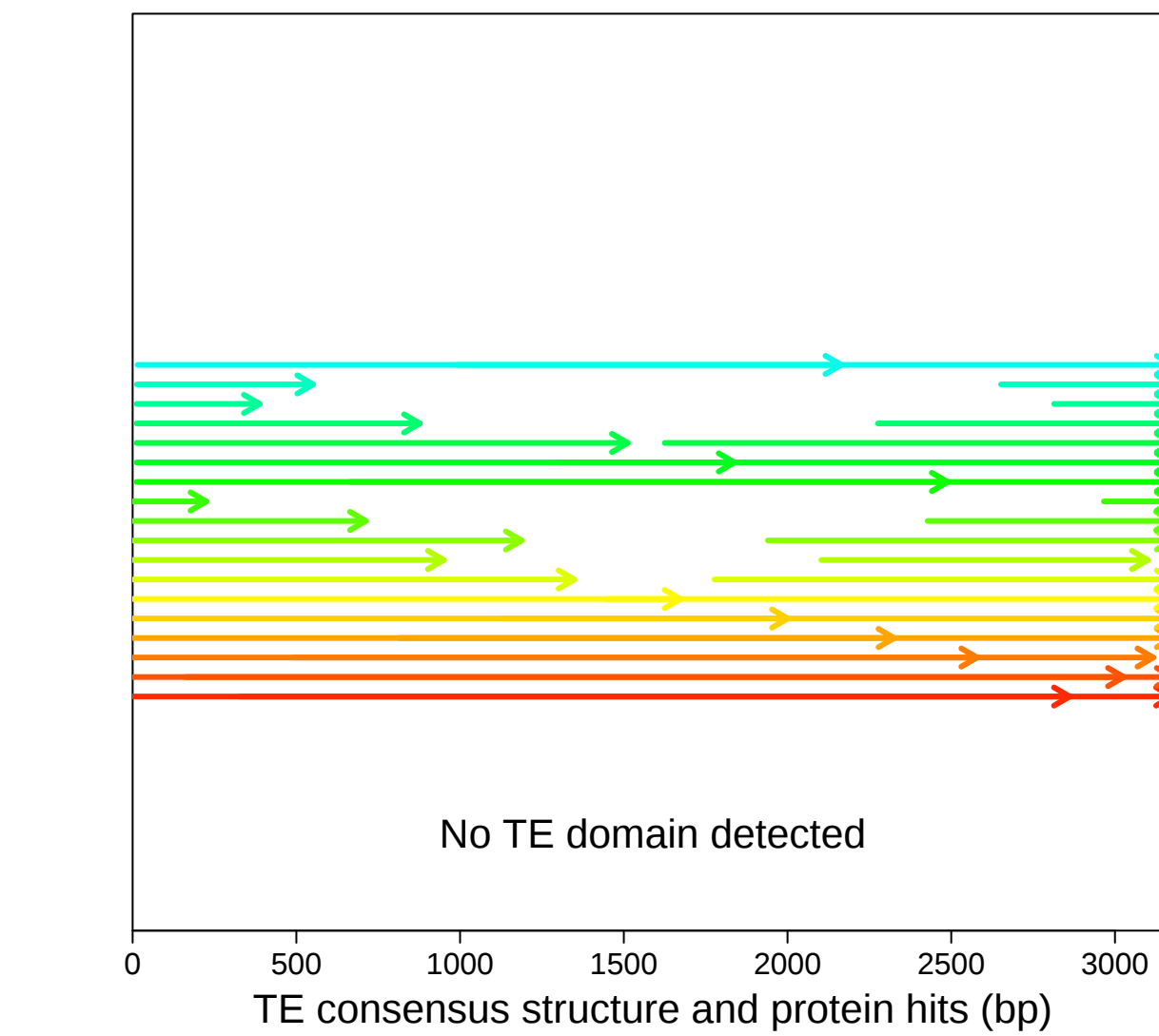
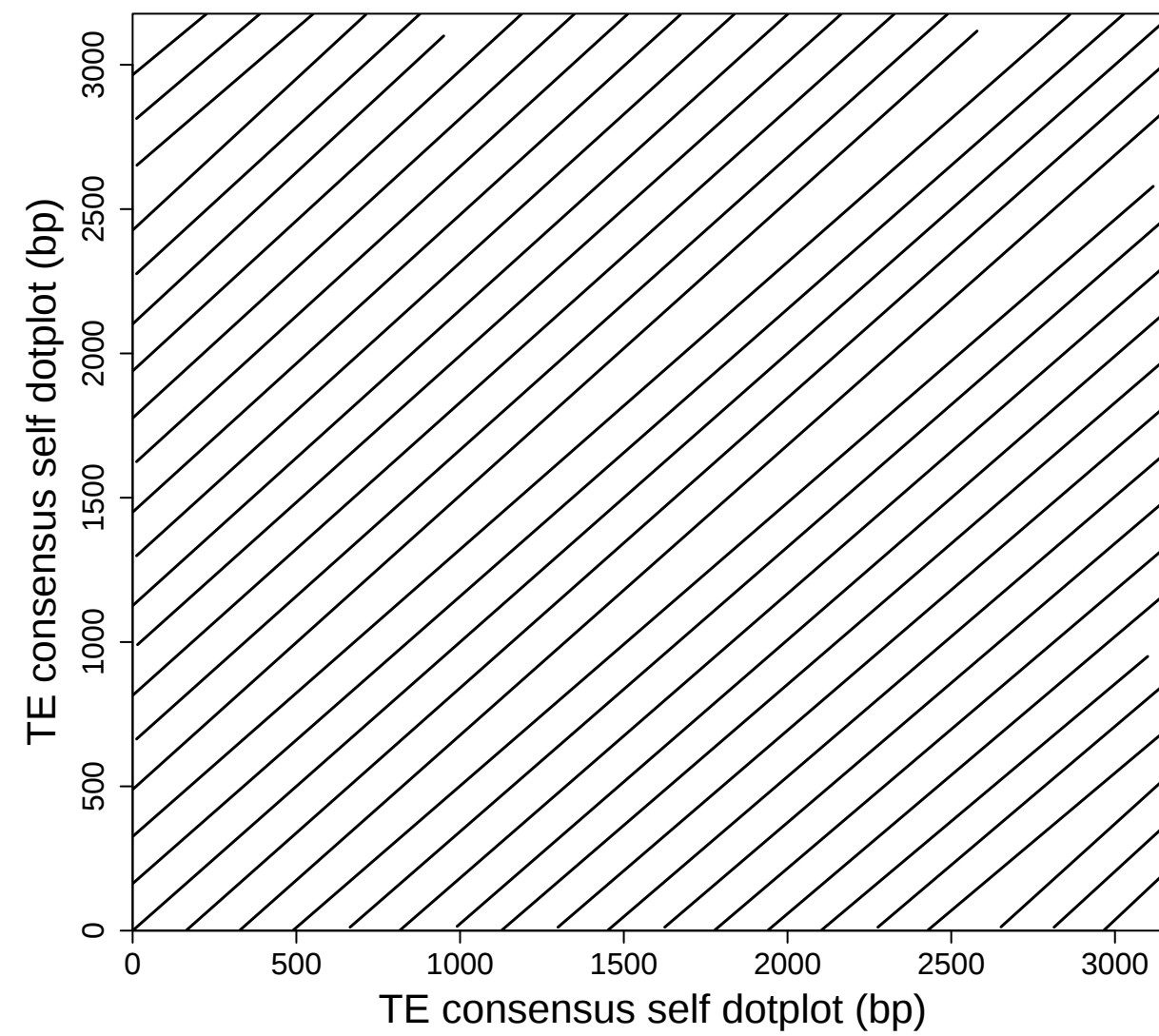
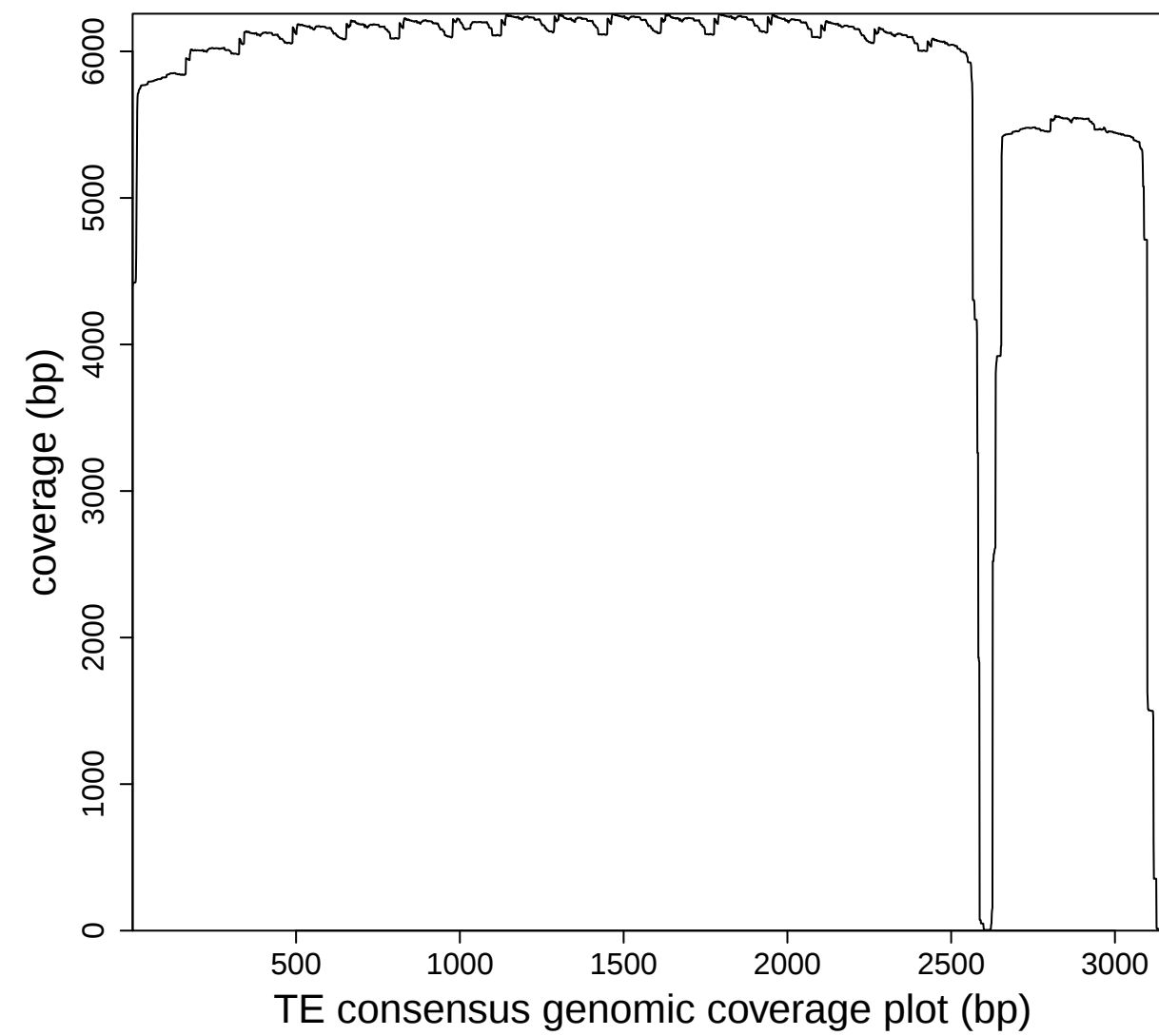
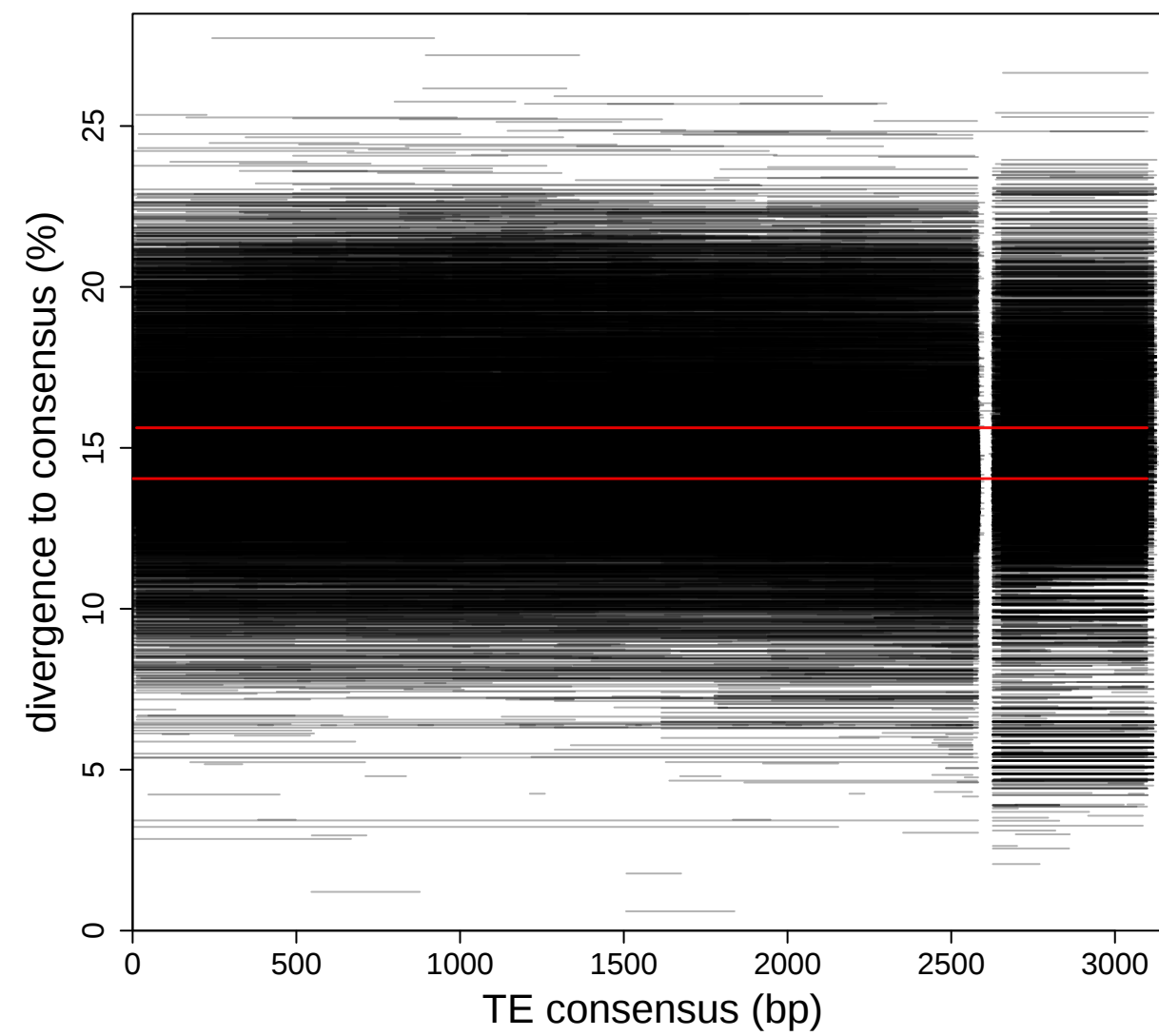
Start crop Point

End crop Point



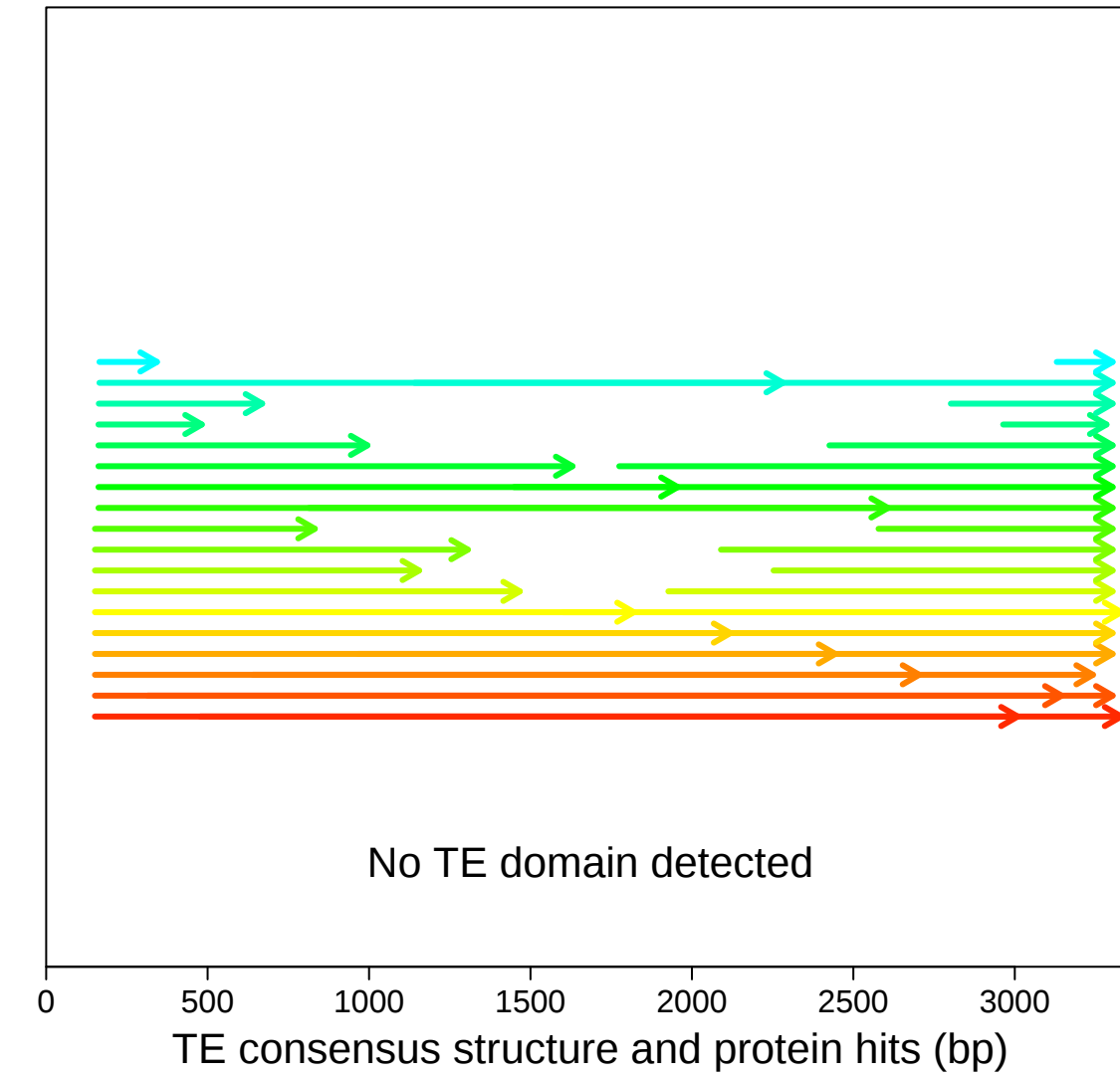
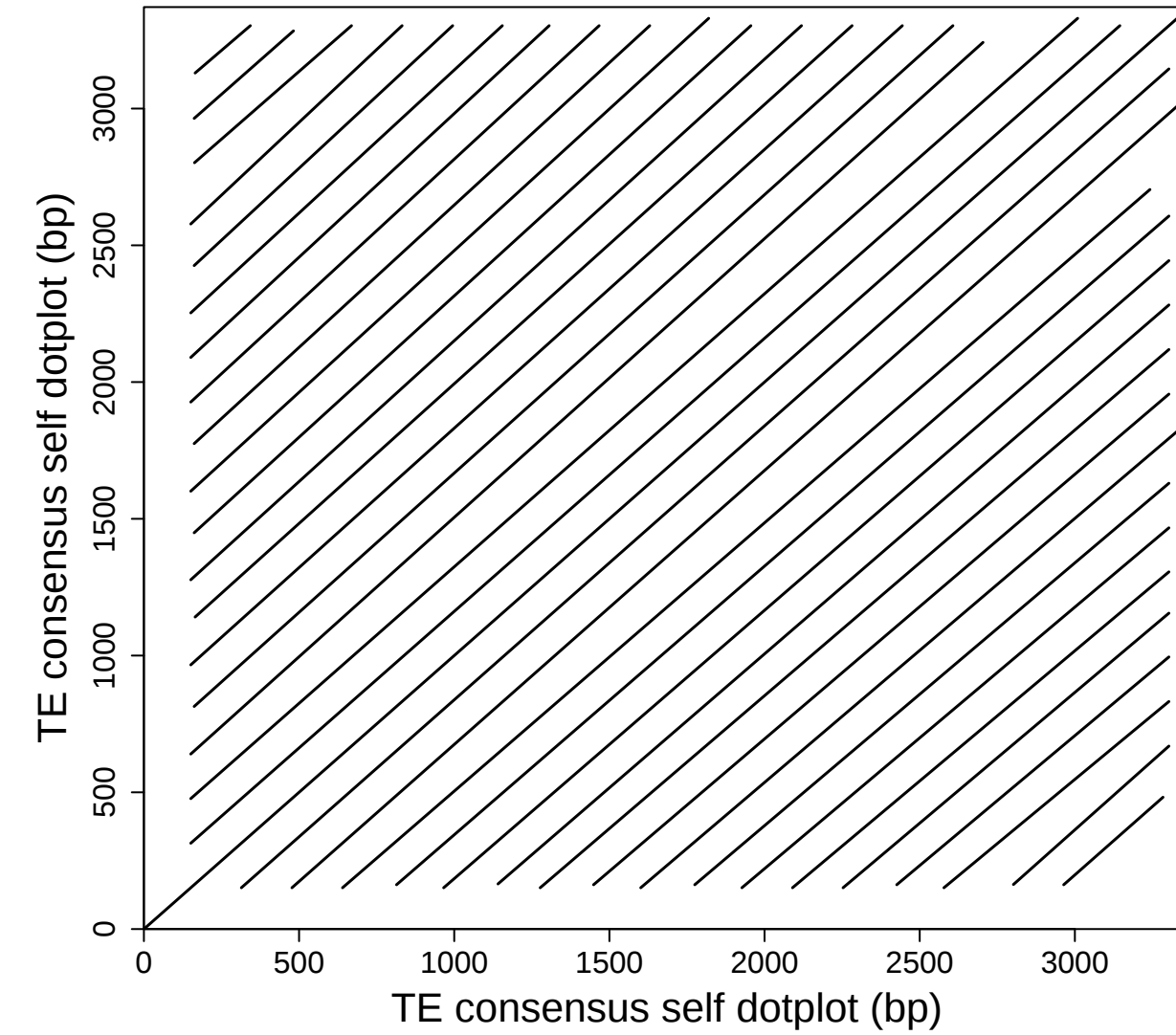
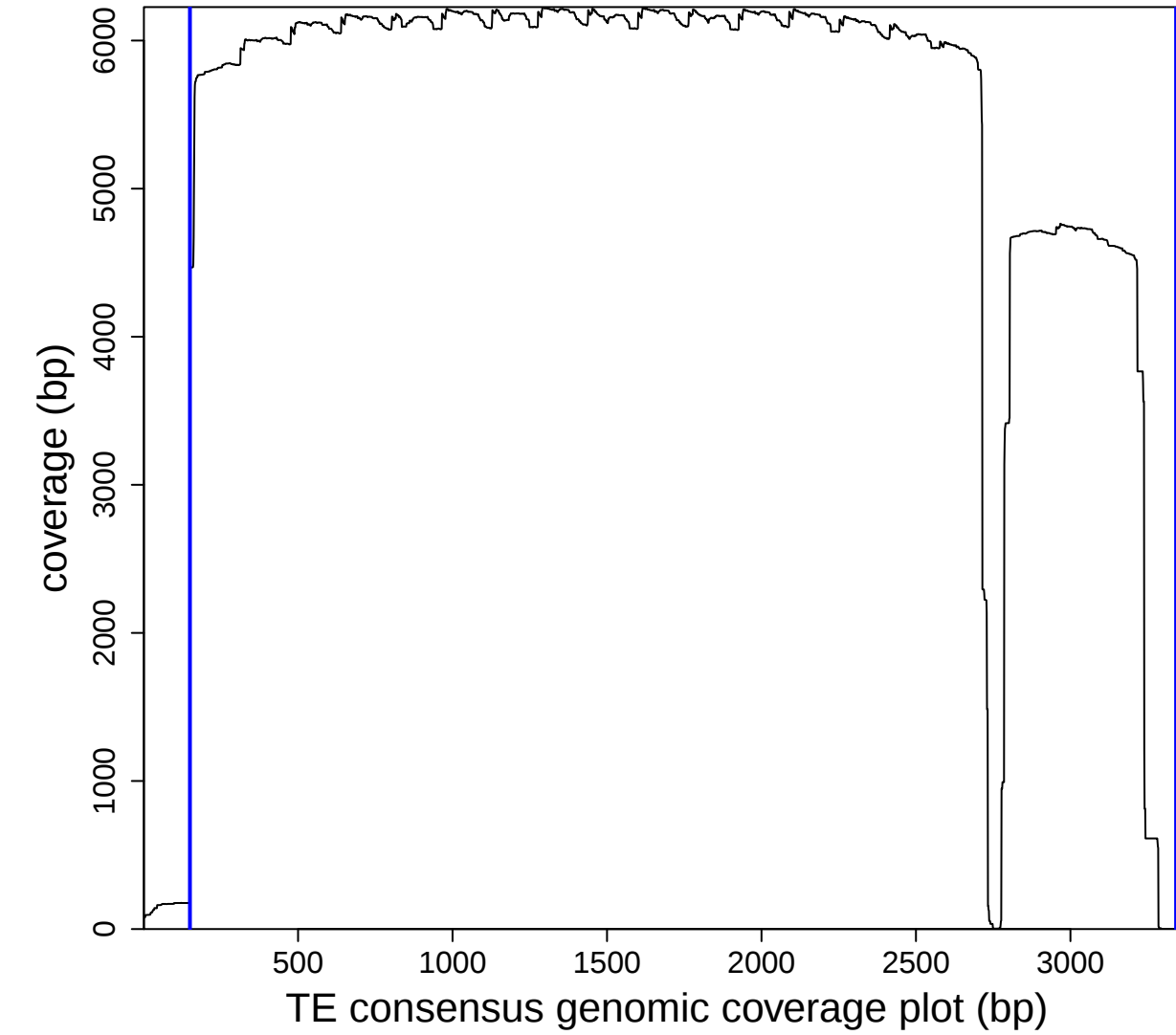
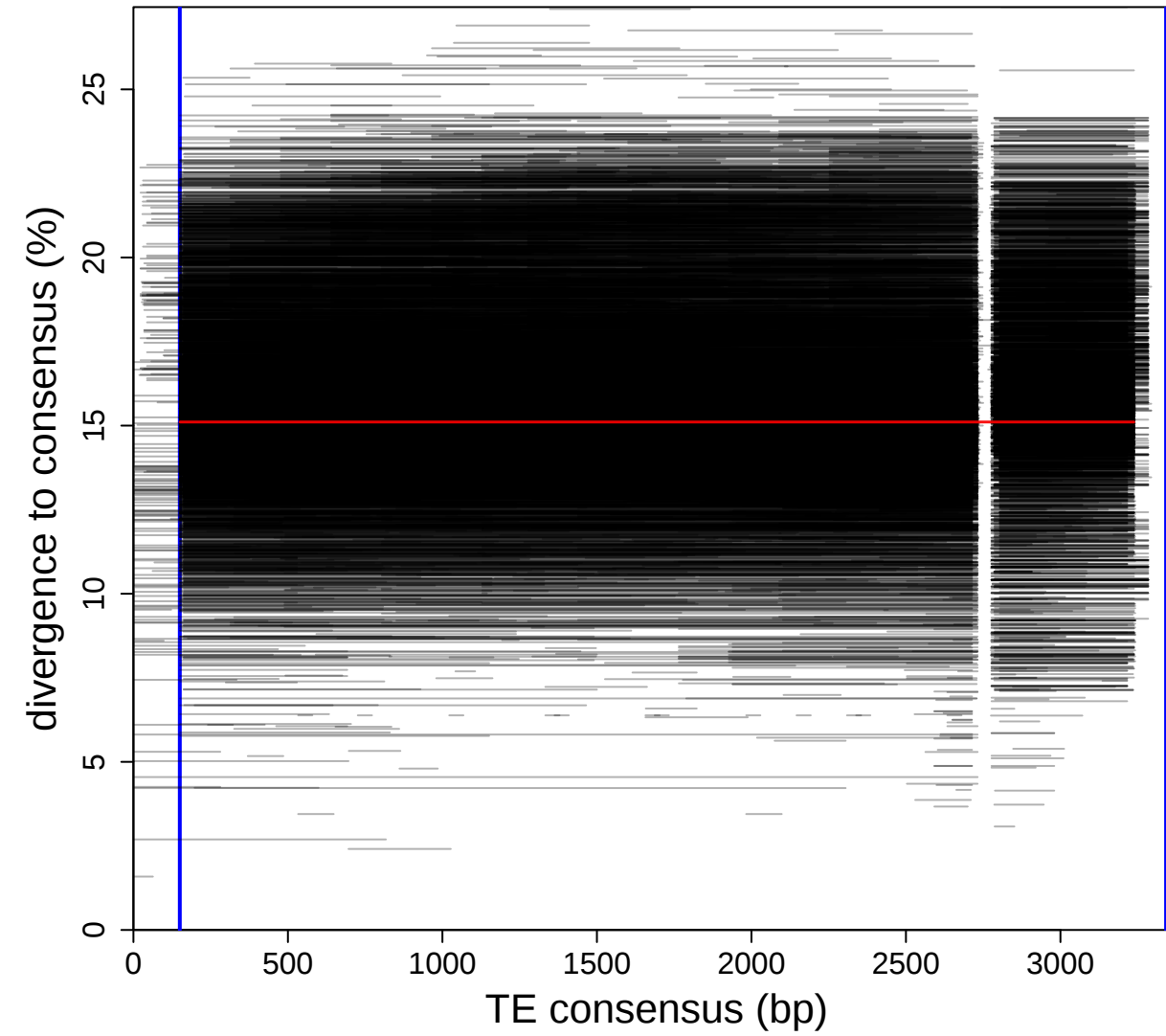
bed_uf.bed_g_5.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa_c
size: 3177bp; fragments: 15496; full length: 2 (>=2859.3bp)

After TEtrimmer 3177 bp

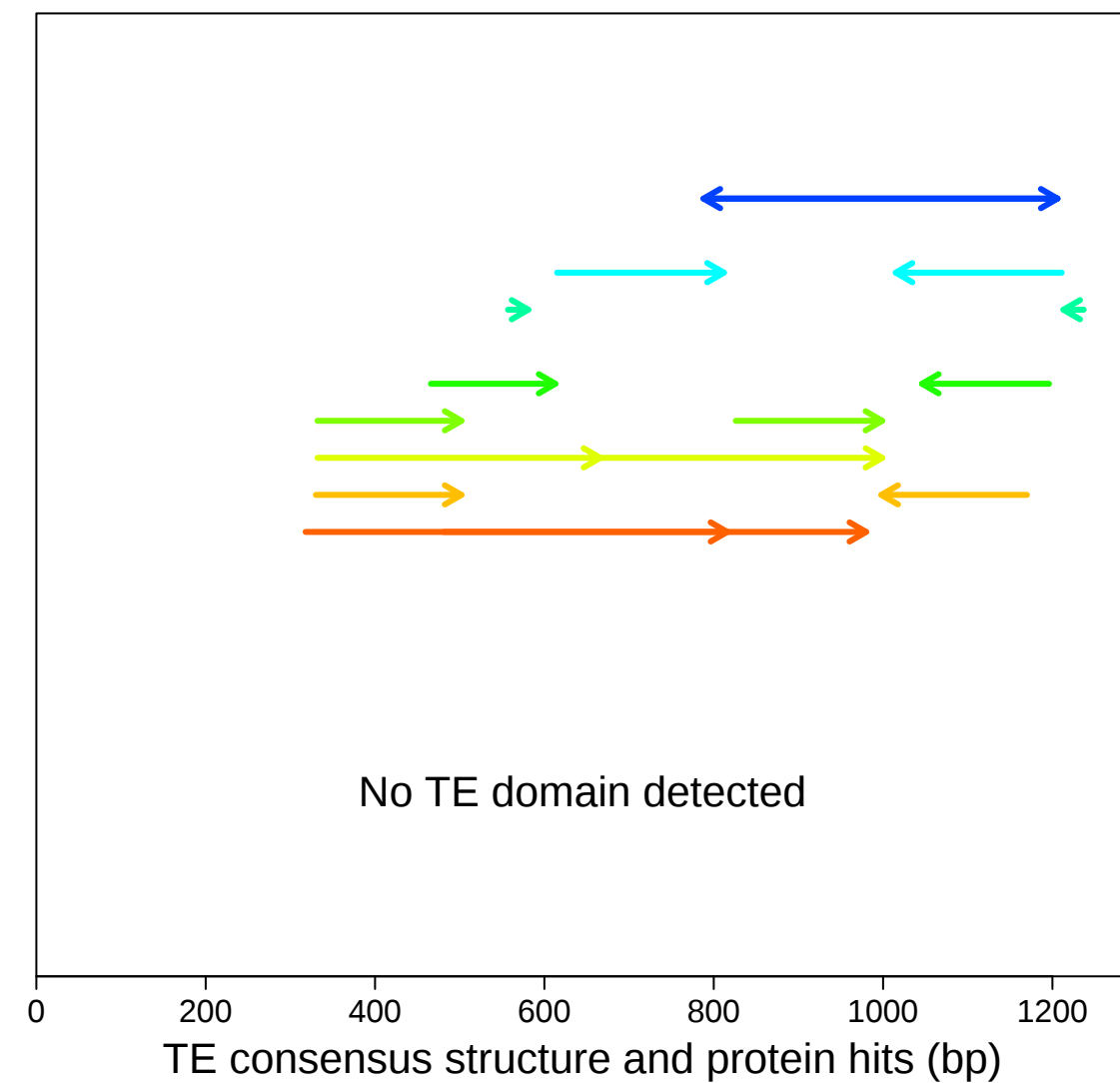
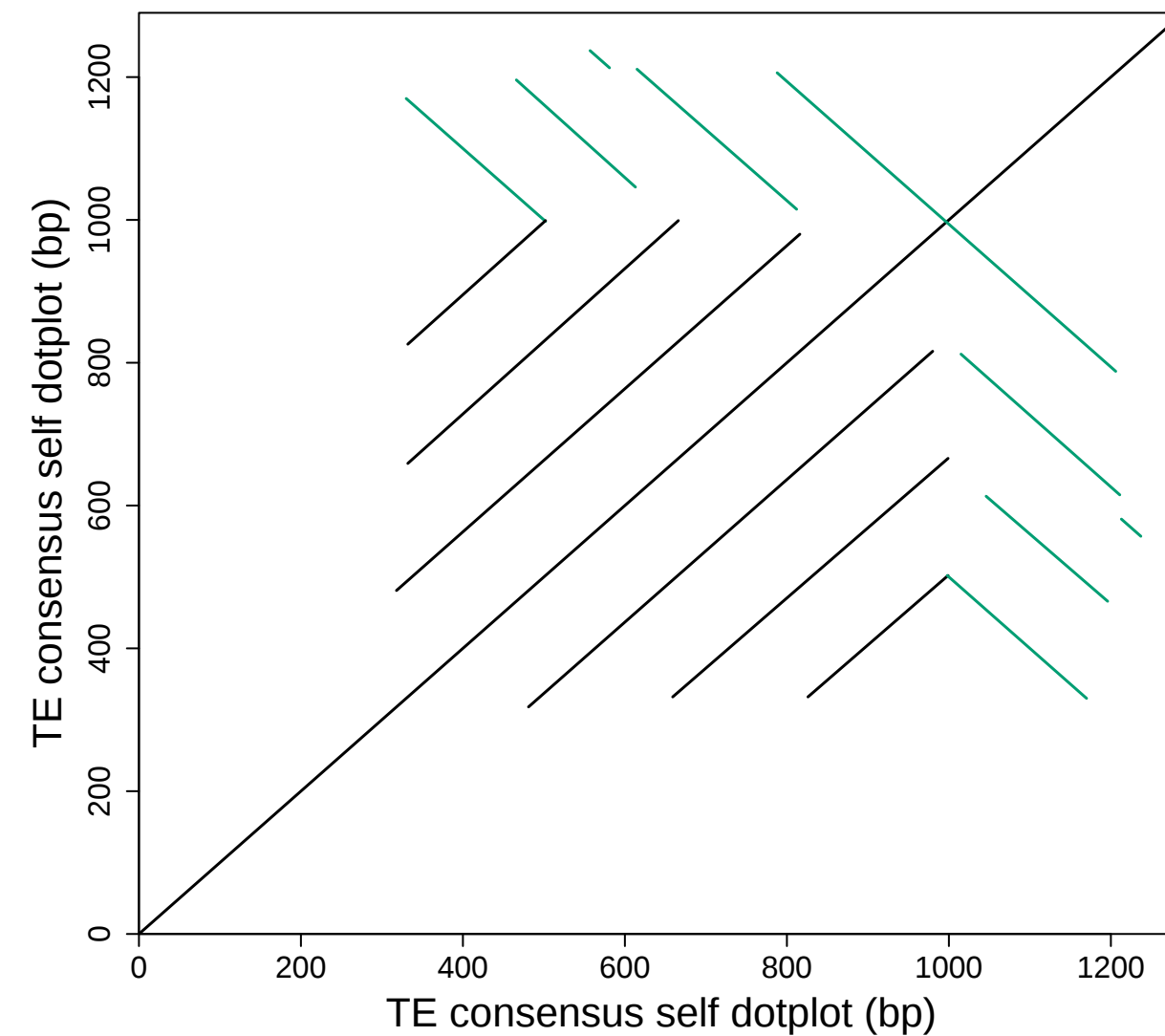
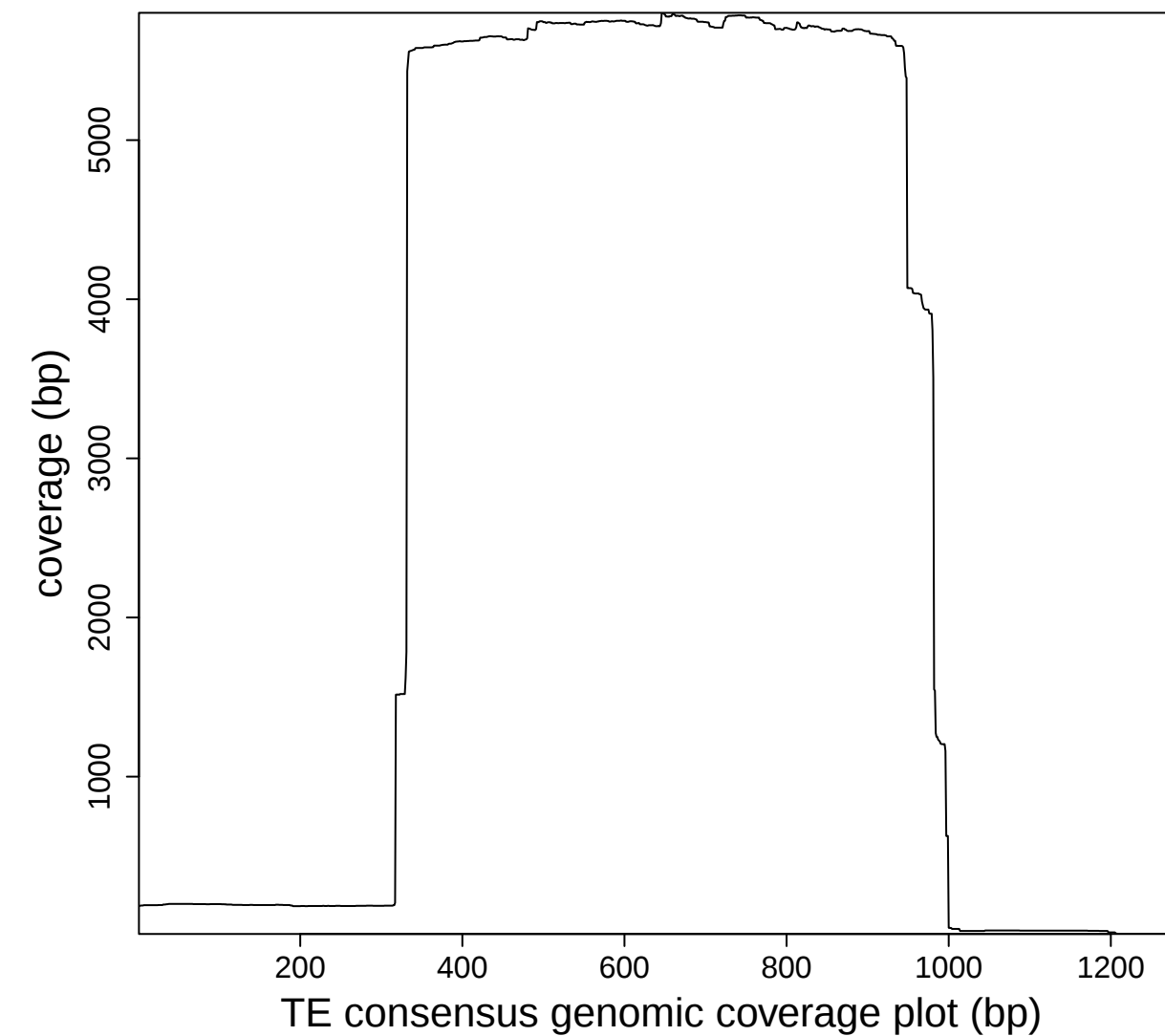
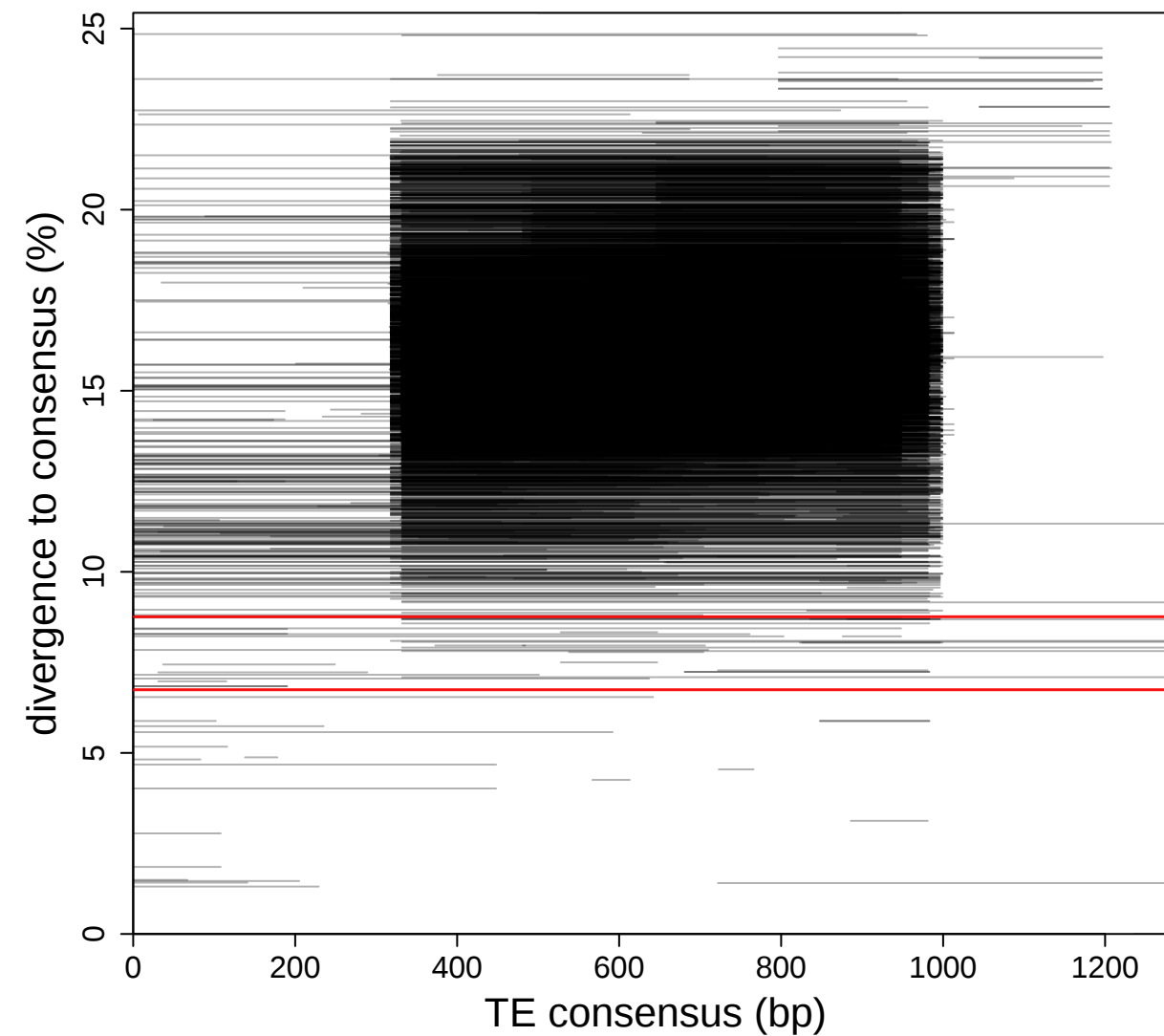


ly_8603.fasta.b.bed_uf.bed_g_5.bed_fm_1.bed_0_0_bcln.fa_aln.f
size: 3387bp; fragments: 14480; full length: 1 (>=3048.3bp)

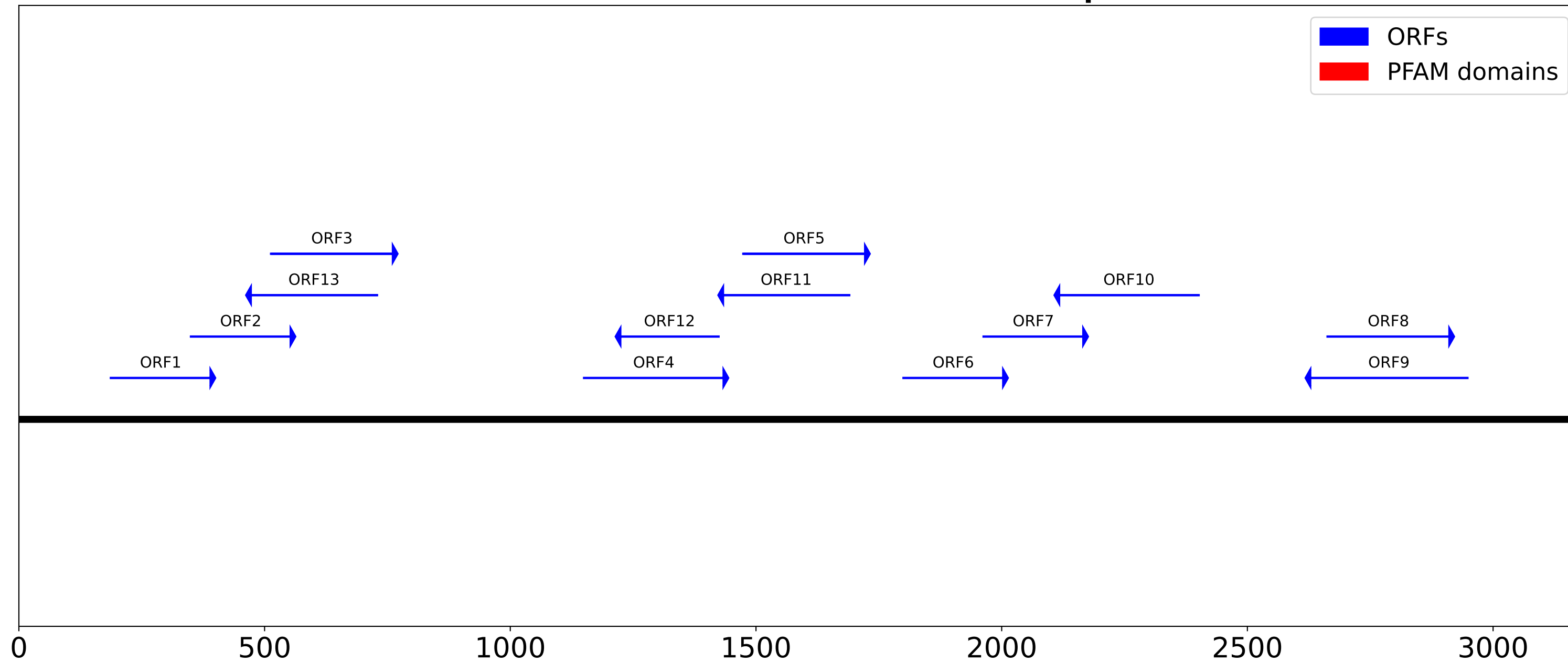
After TEtrimmer Extended plot Blue lines are boundaries



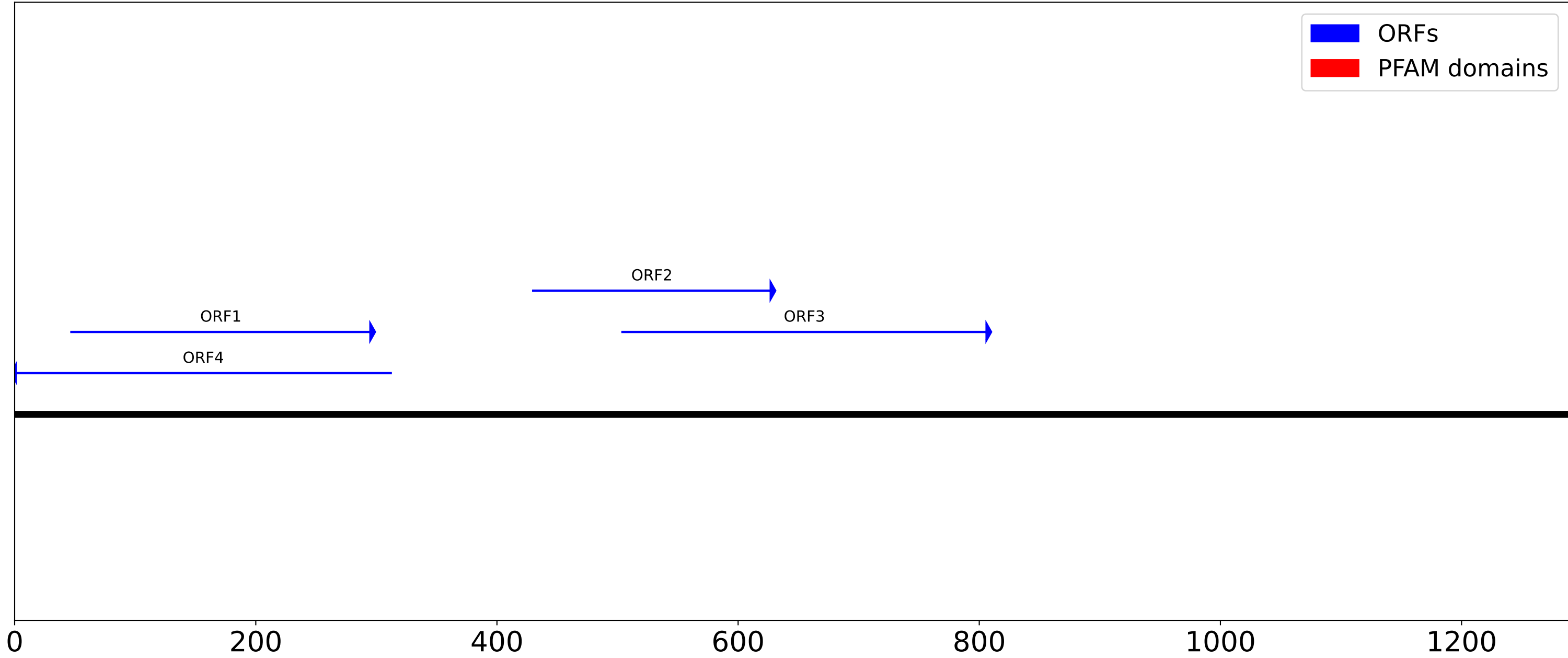
TE: rnd_5_family_8603
size: 1290bp; fragments: 6250; full length: 2 (≥ 1161 bp)



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)

