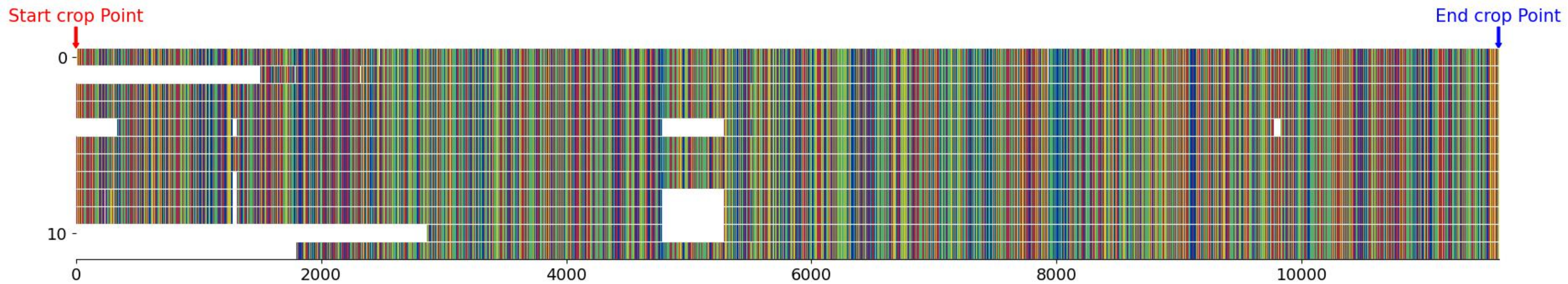




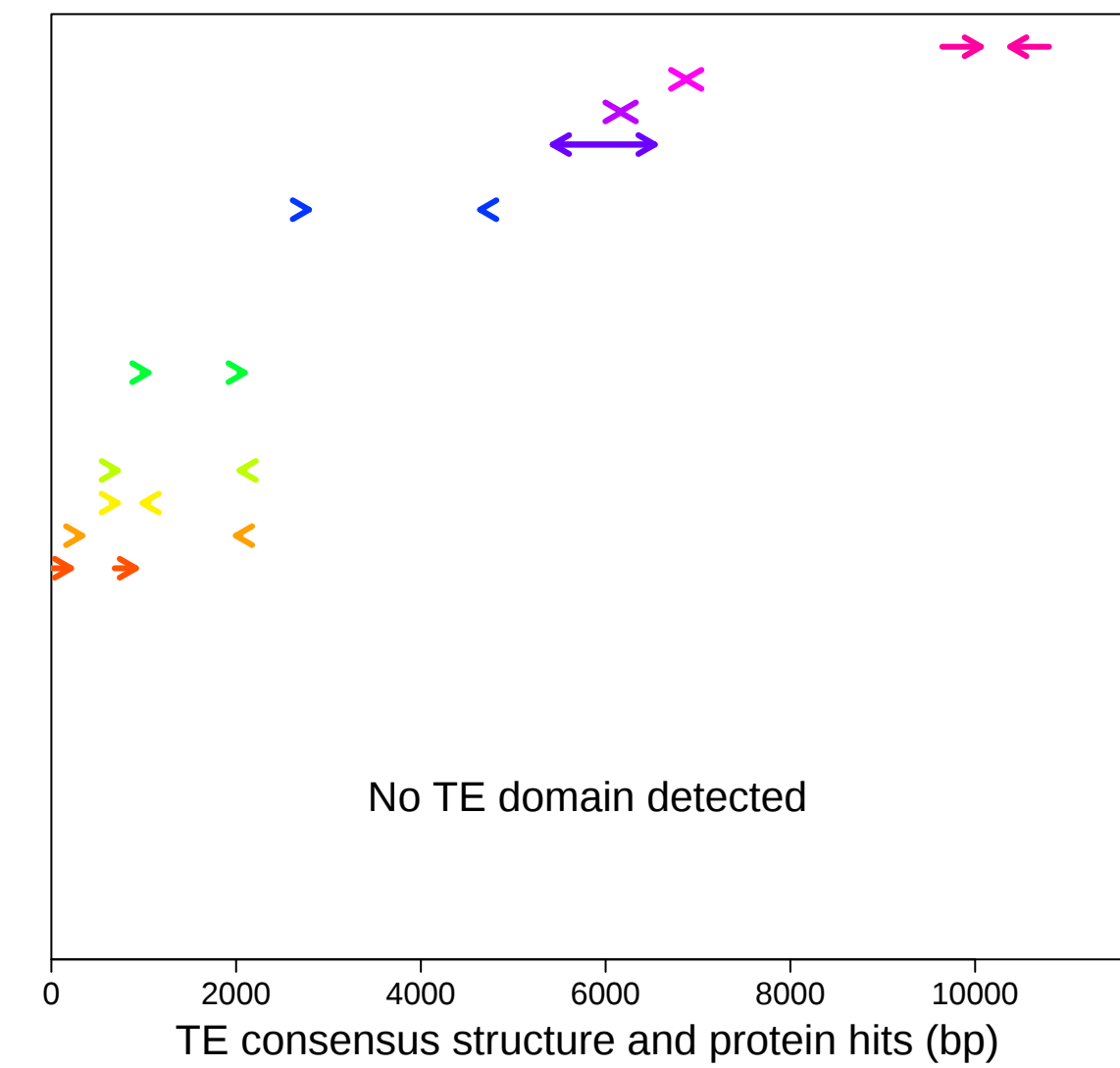
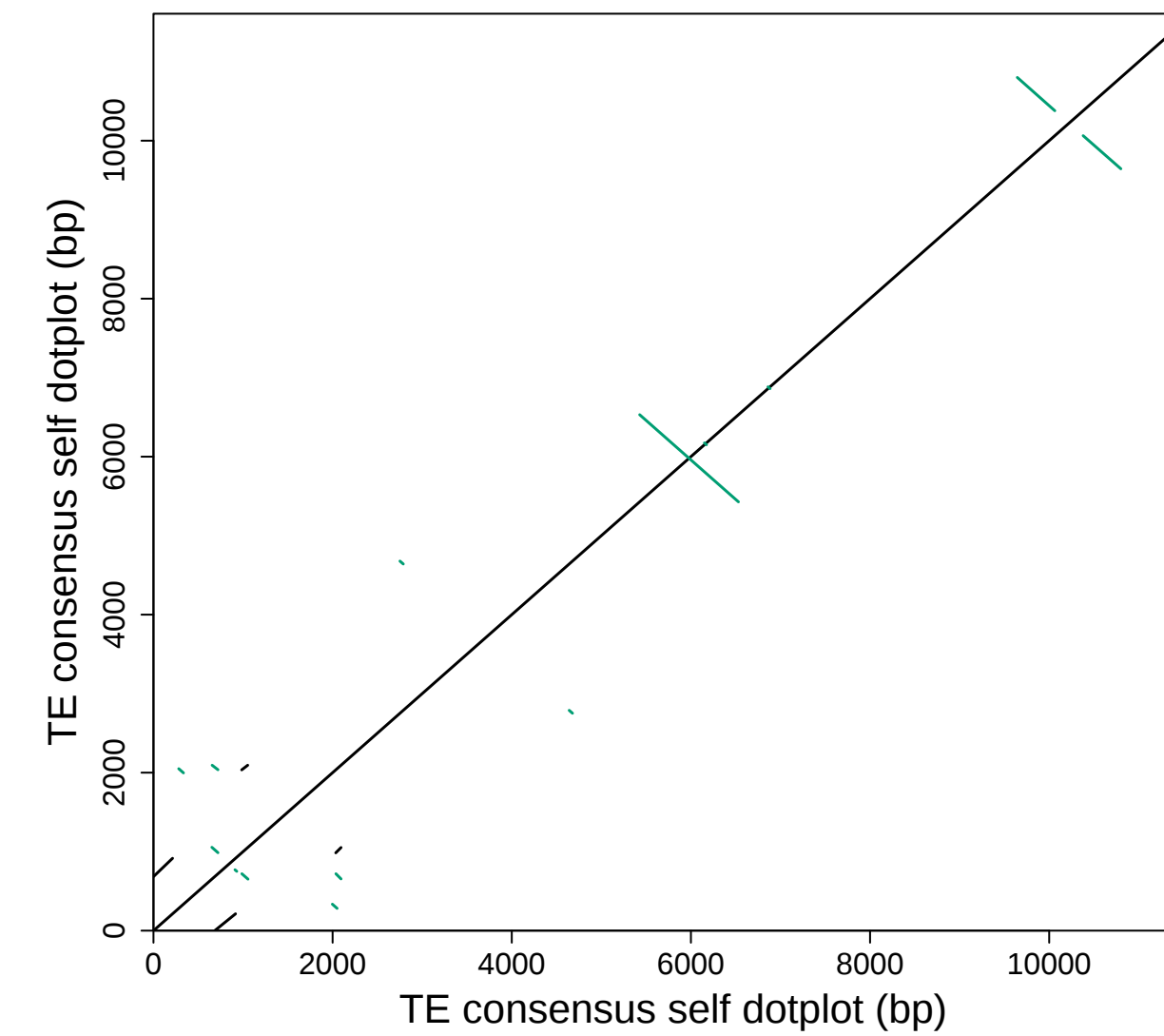
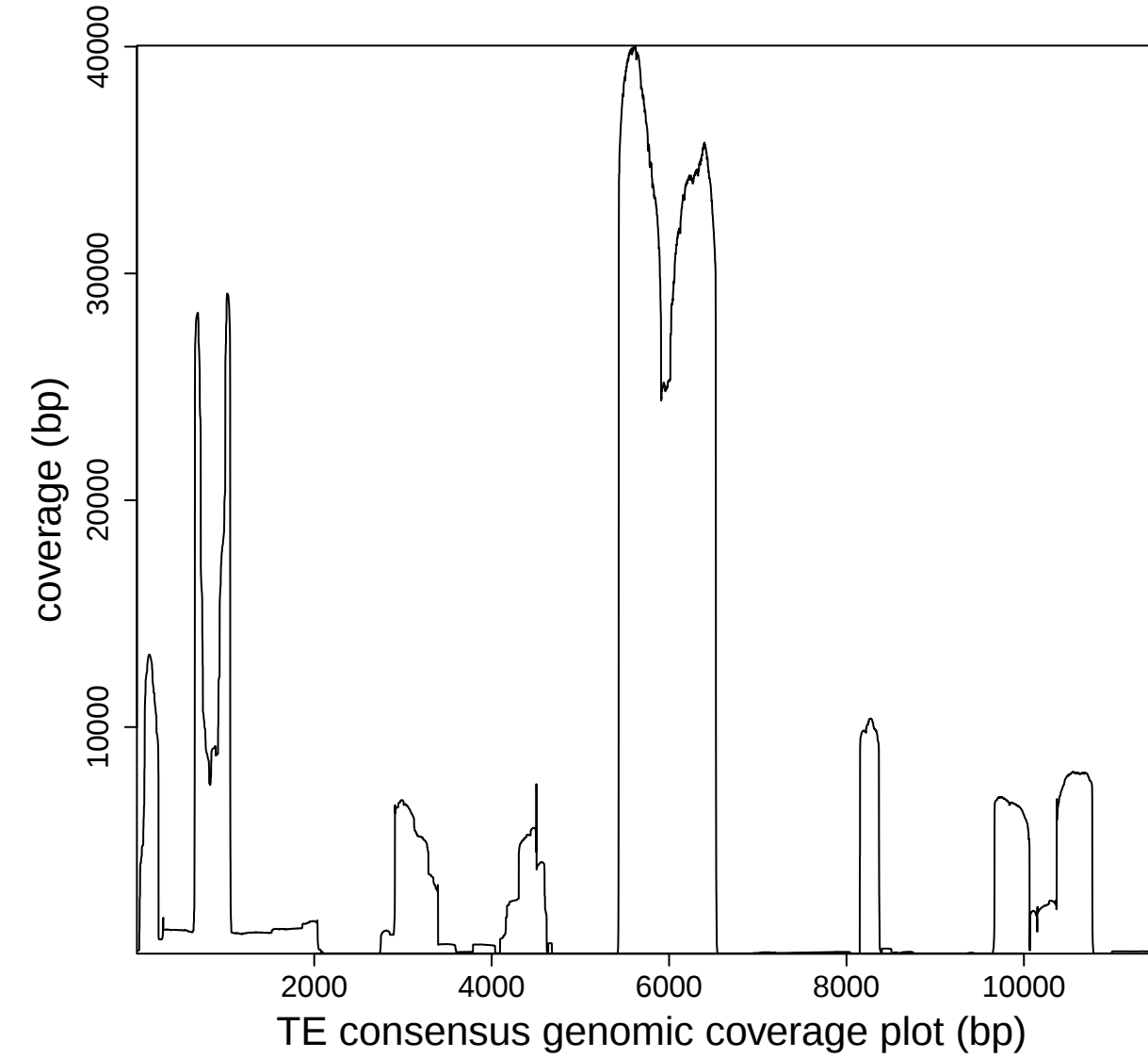
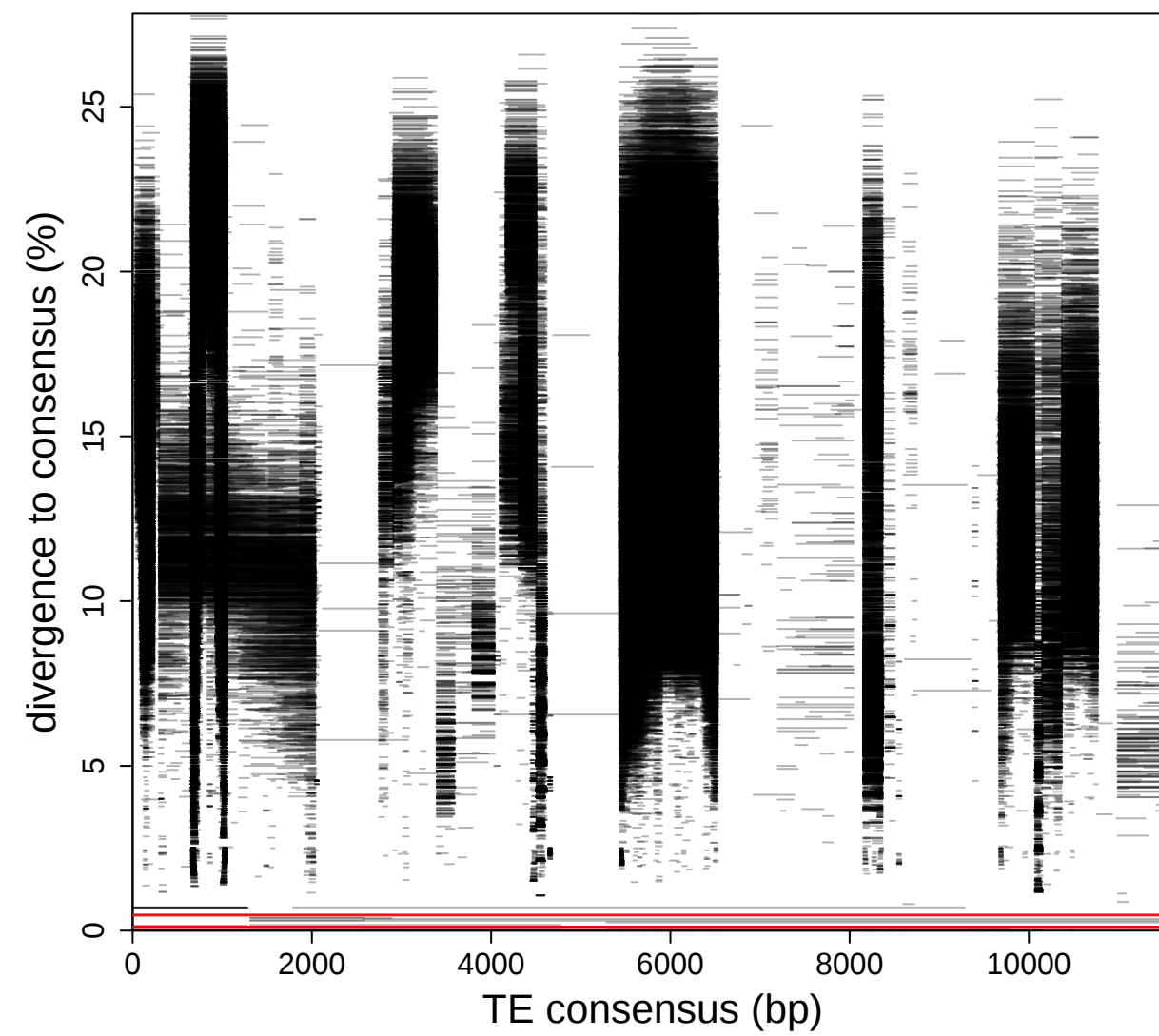
MSA length = 11607

[illegible]



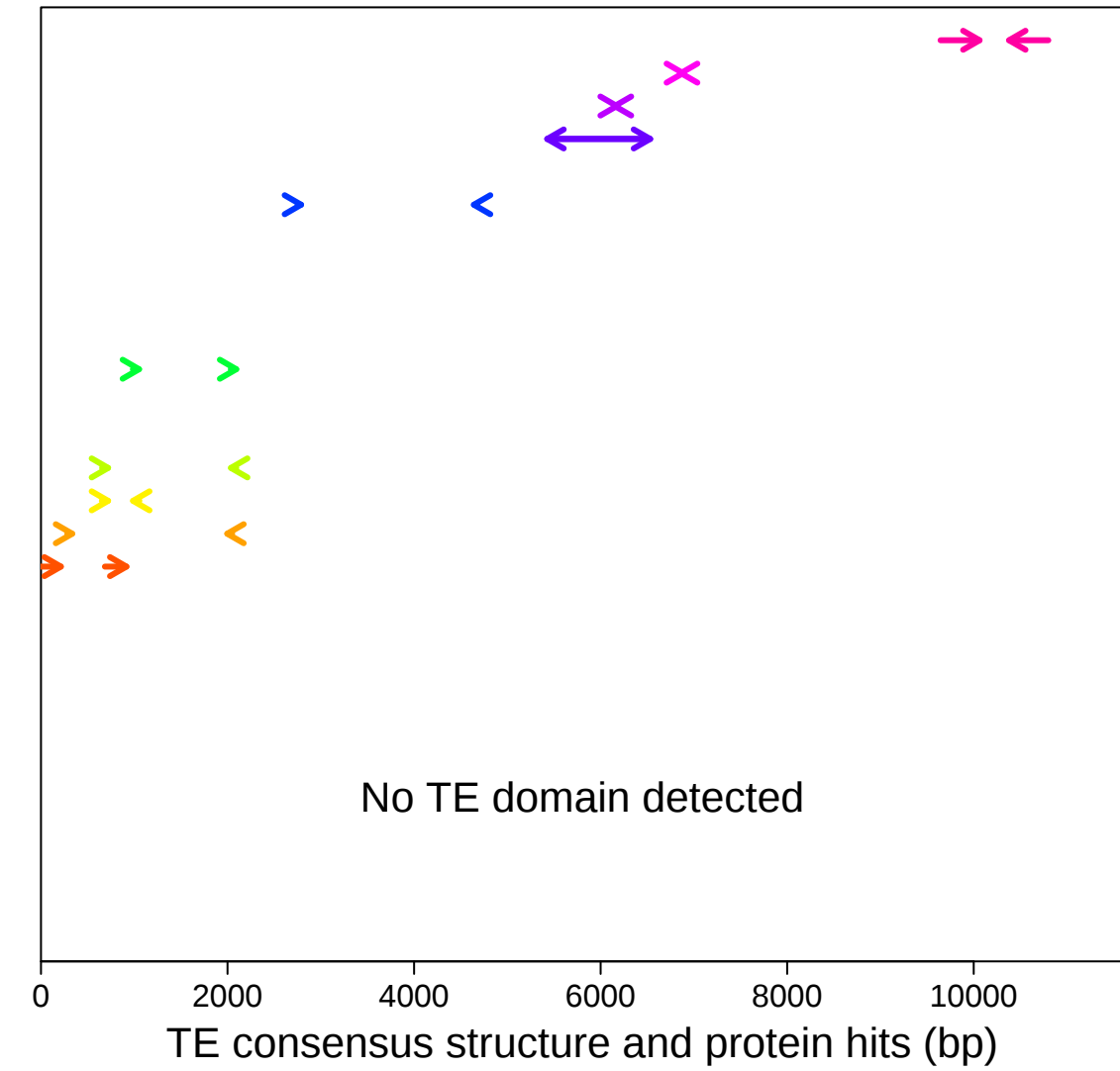
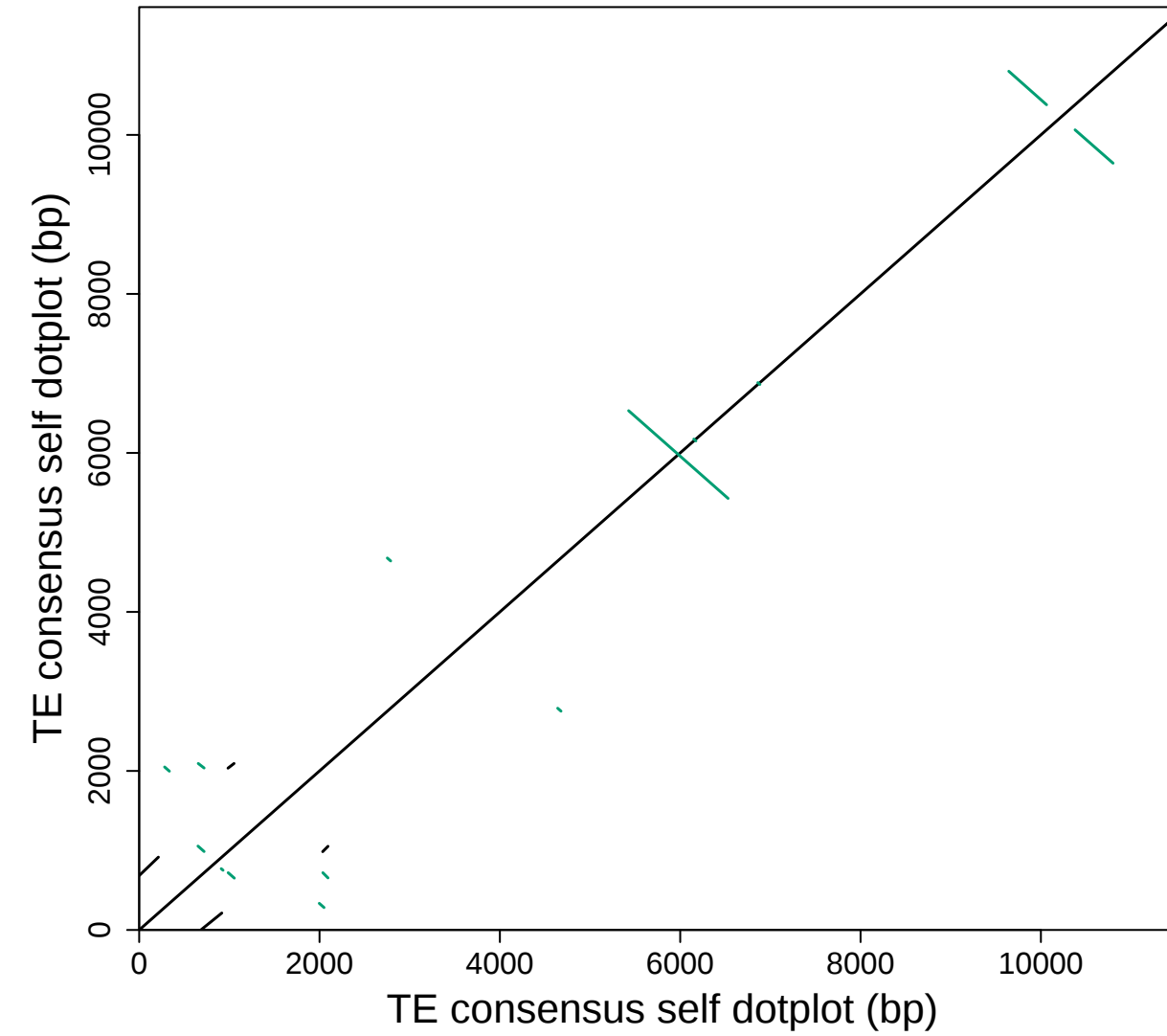
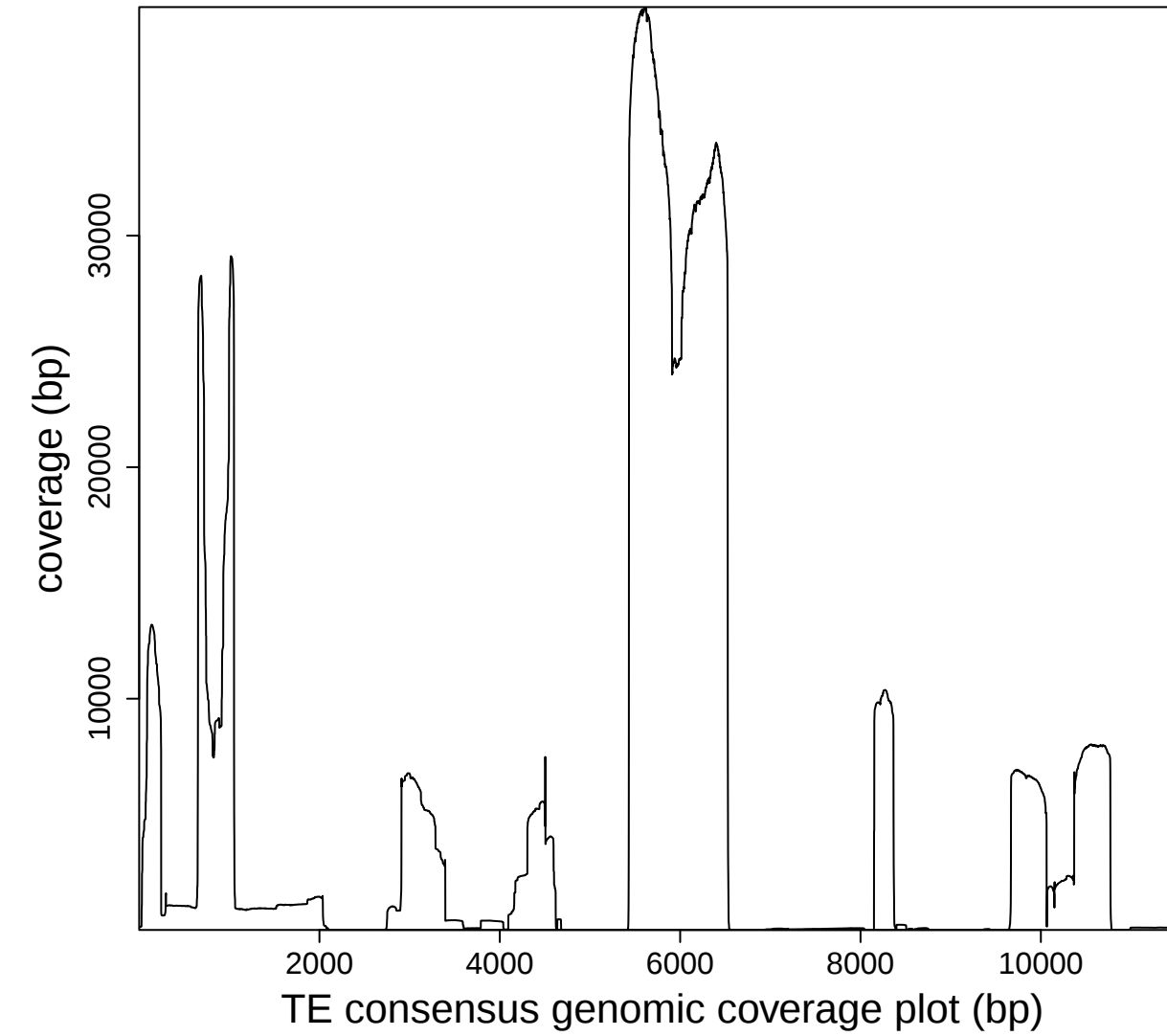
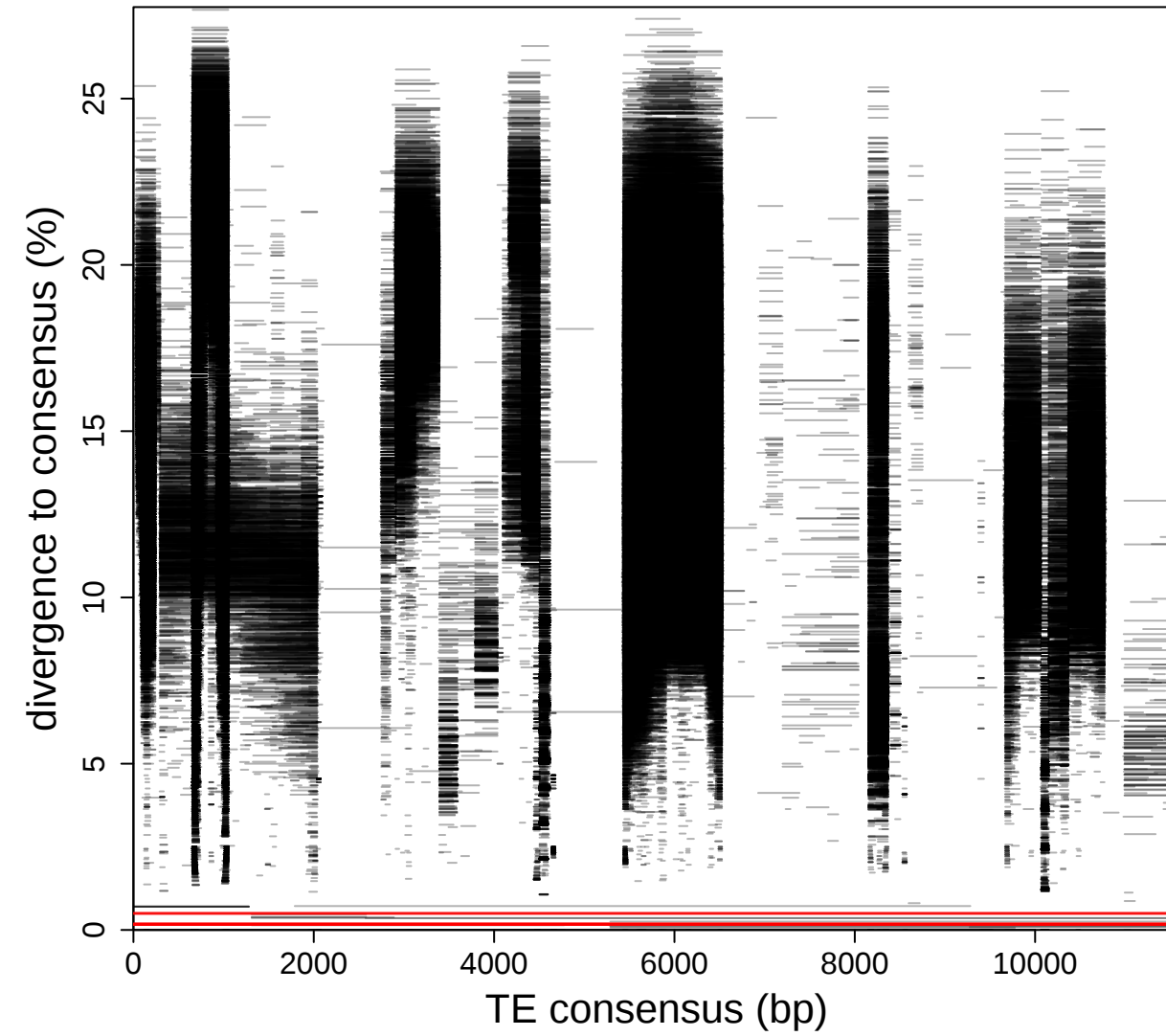
bed_uf.bed_g_3.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa_c
size: 11607bp; fragments: 190572; full length: 5 (>=10446.3bp)

After TEtrimmer 11607 bp



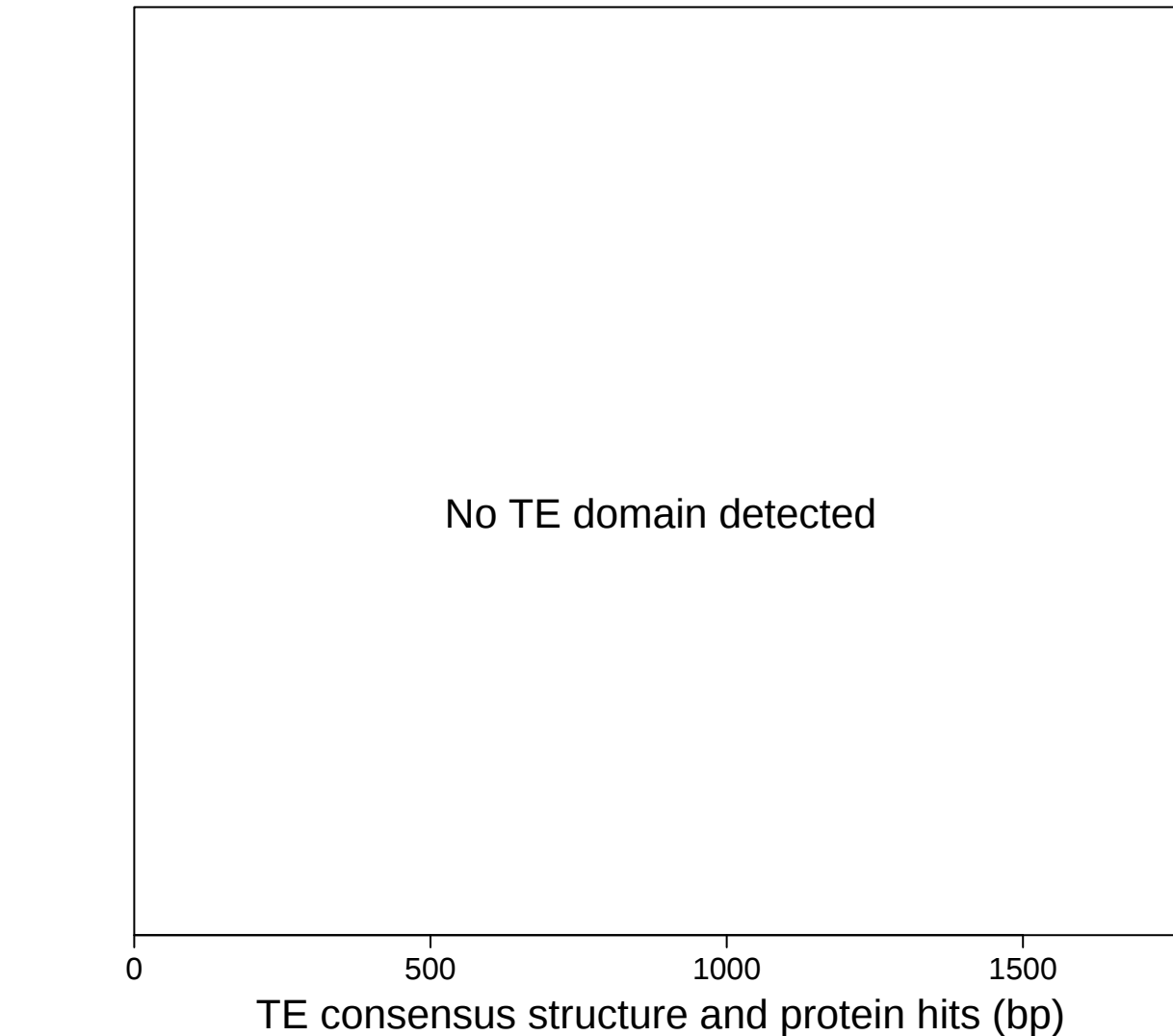
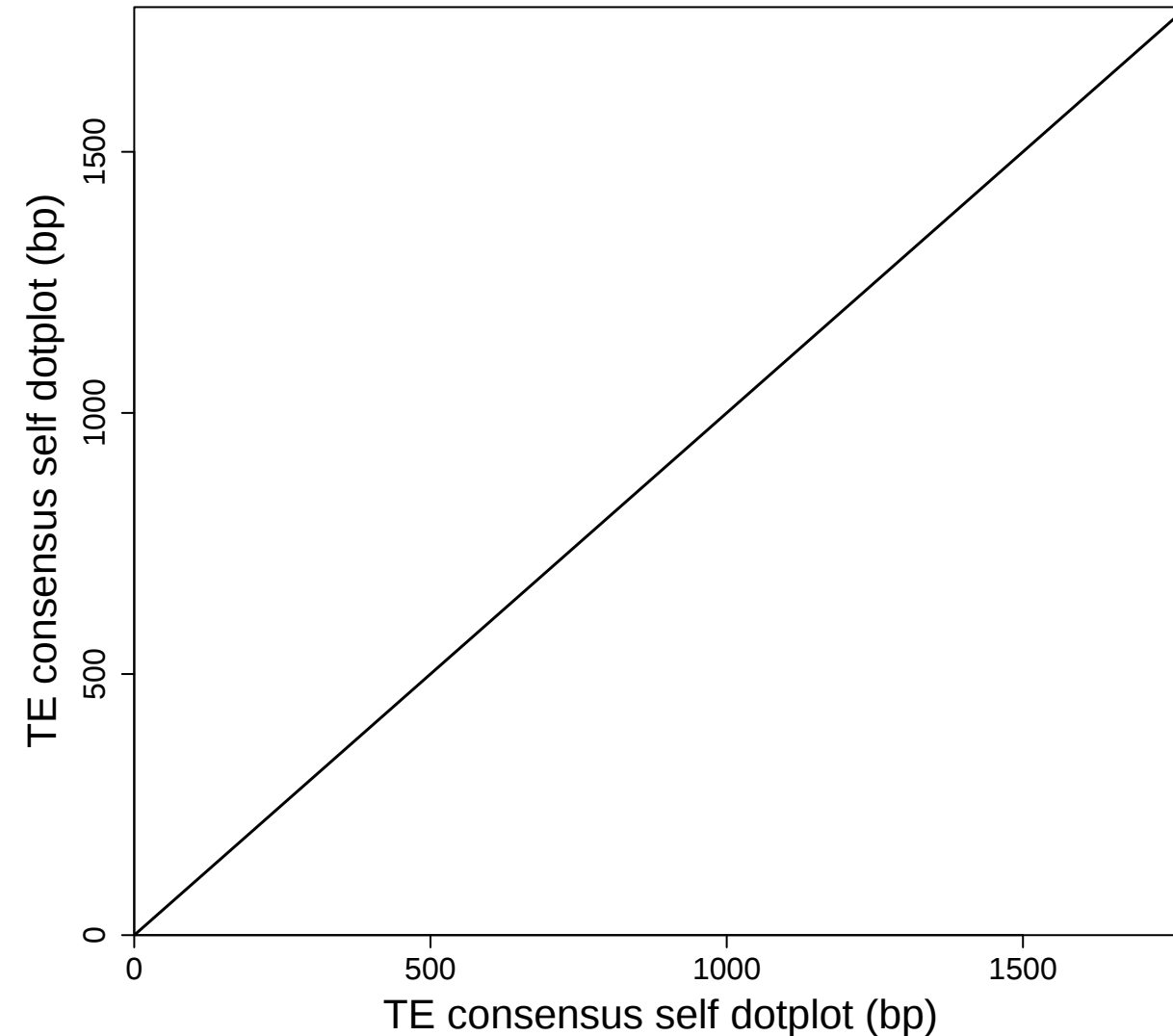
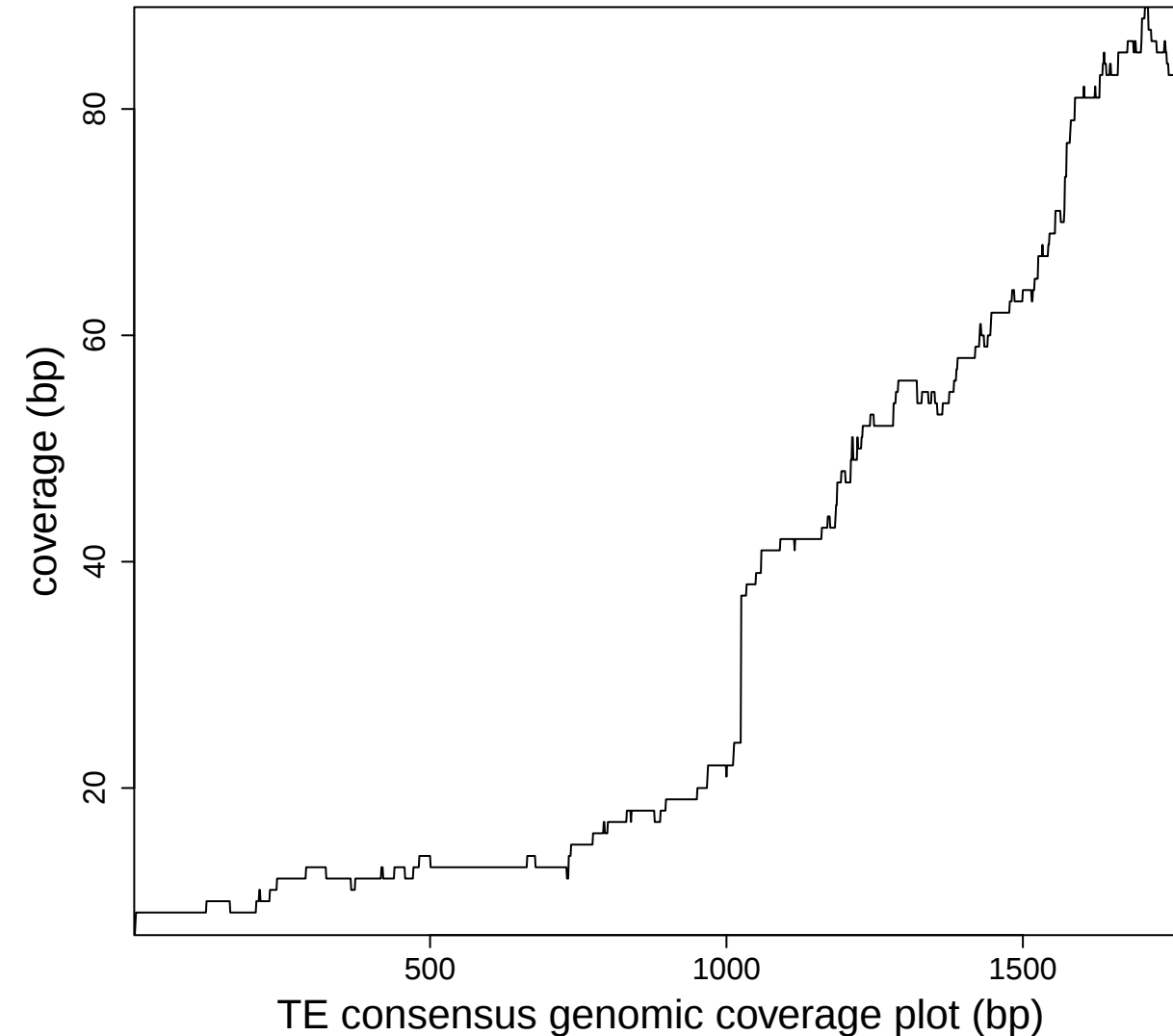
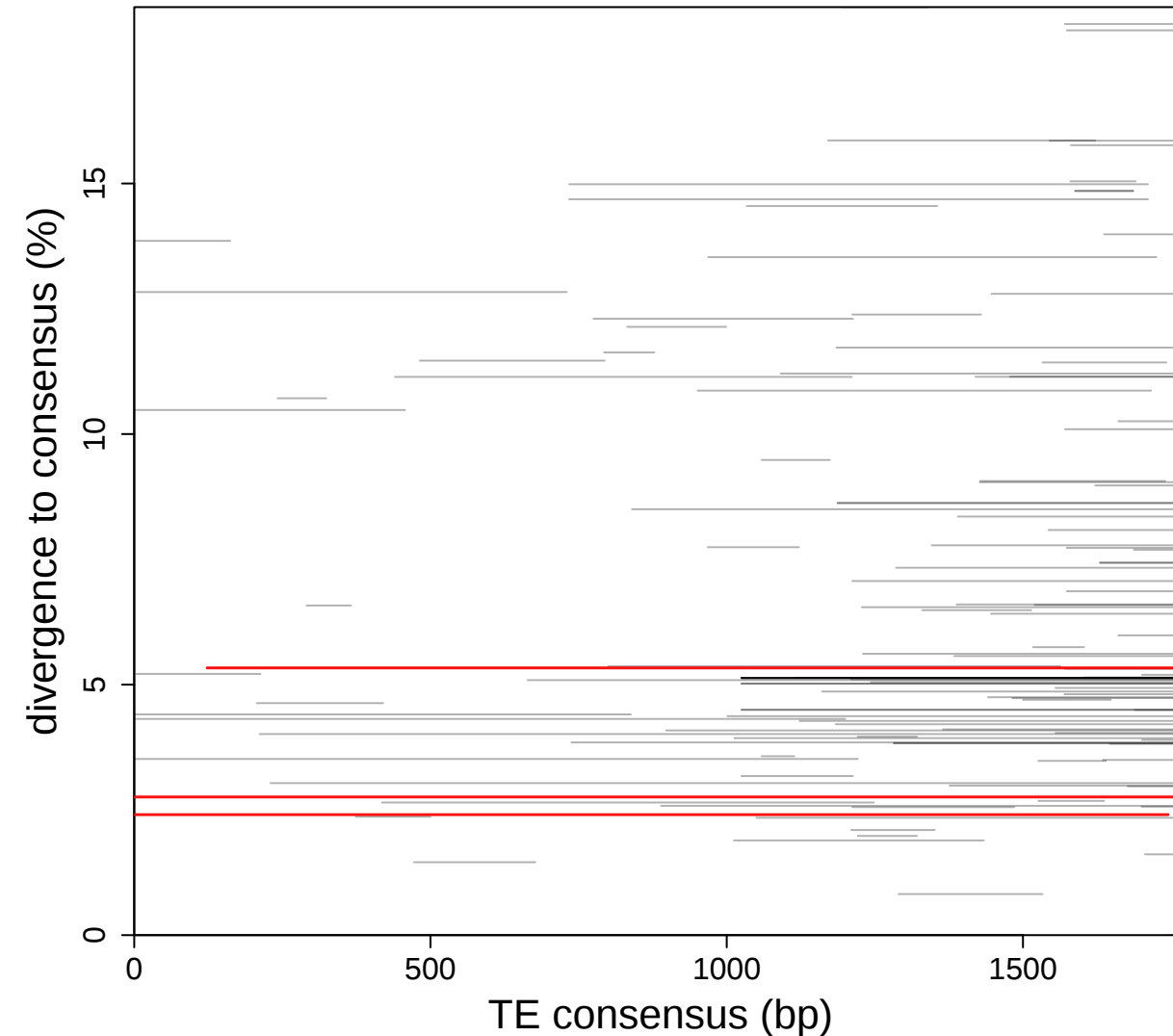
2.fasta.b.bed_uf.bed_g_3.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.f
size: 11607bp; fragments: 187971; full length: 5 (>=10446.3bp

After TEtrimmer Extended plot Blue lines are boundaries

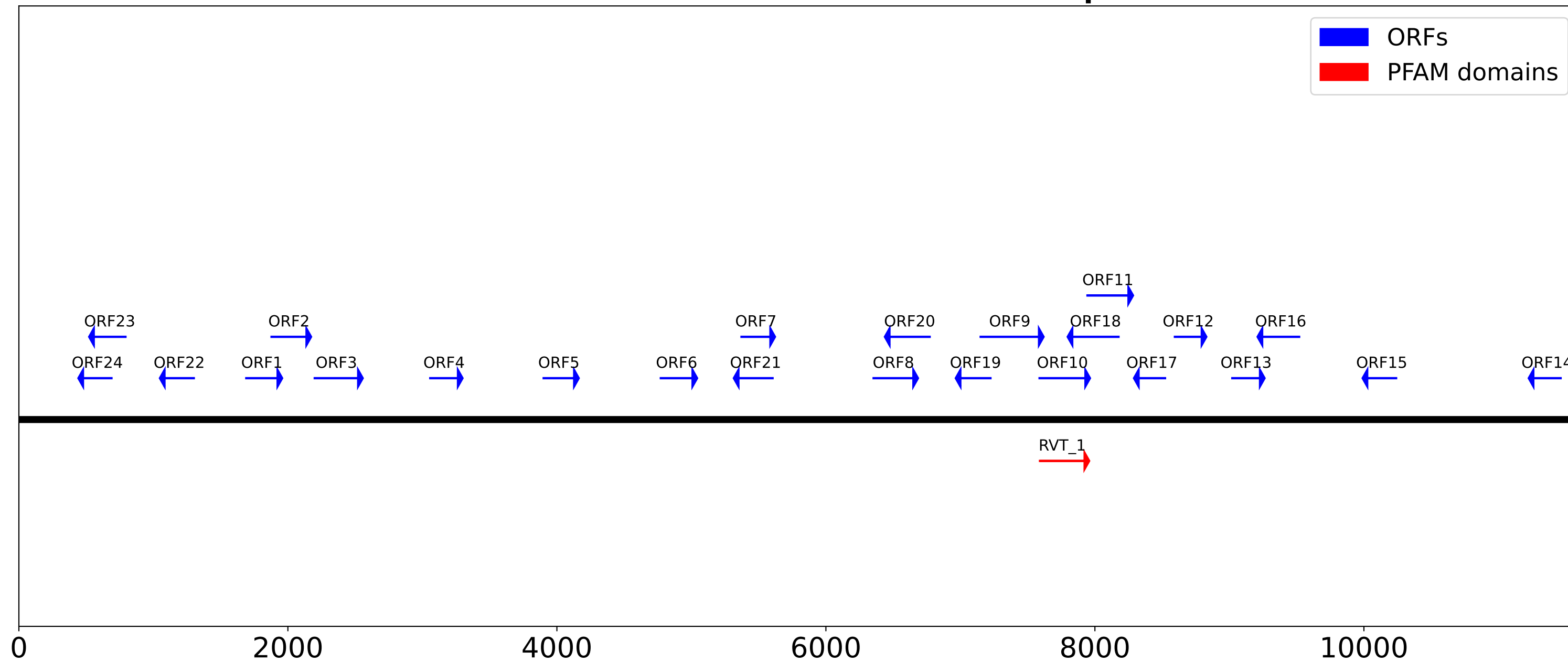


TE: rnd_5_family_2262
size: 1777bp; fragments: 131; full length: 3 (>=1599.3bp)

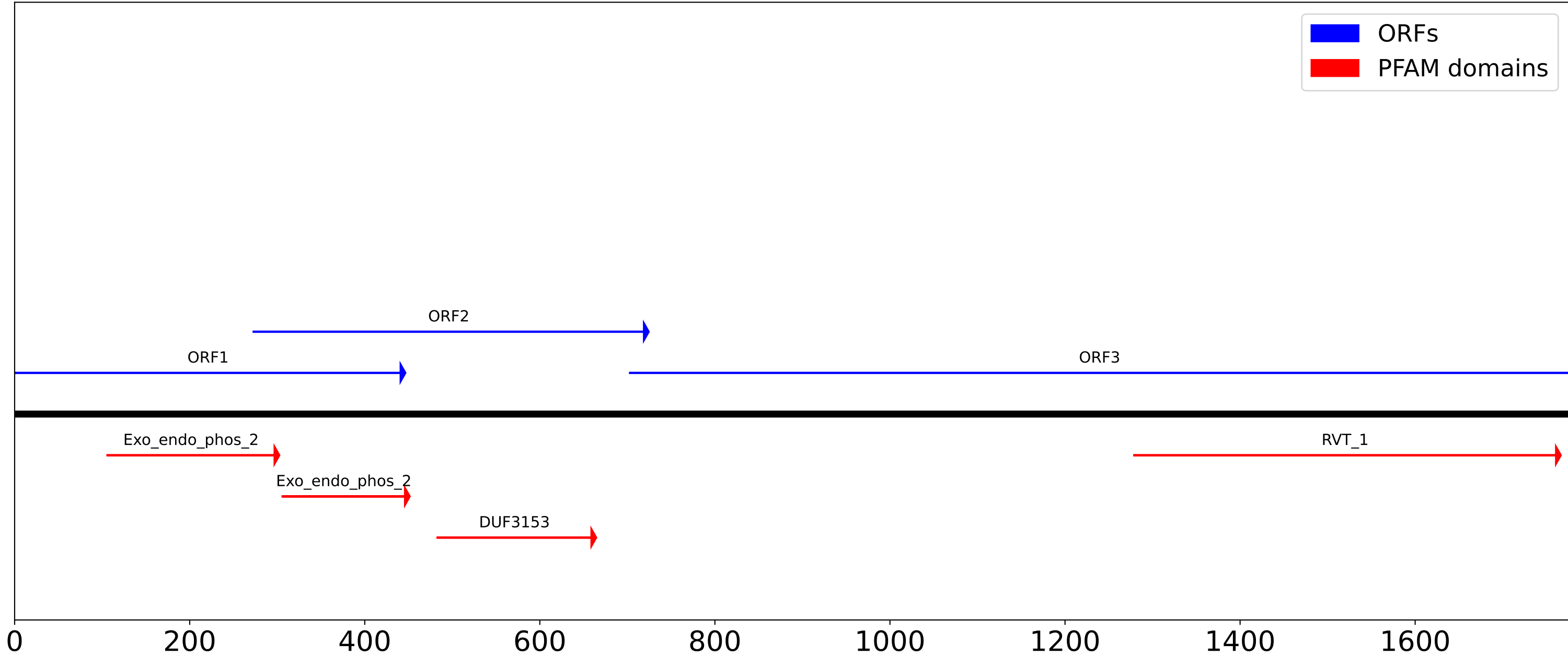
Before TEtrimmer 1777 bp



After TEtrimmer ORF and PFAM domain plot



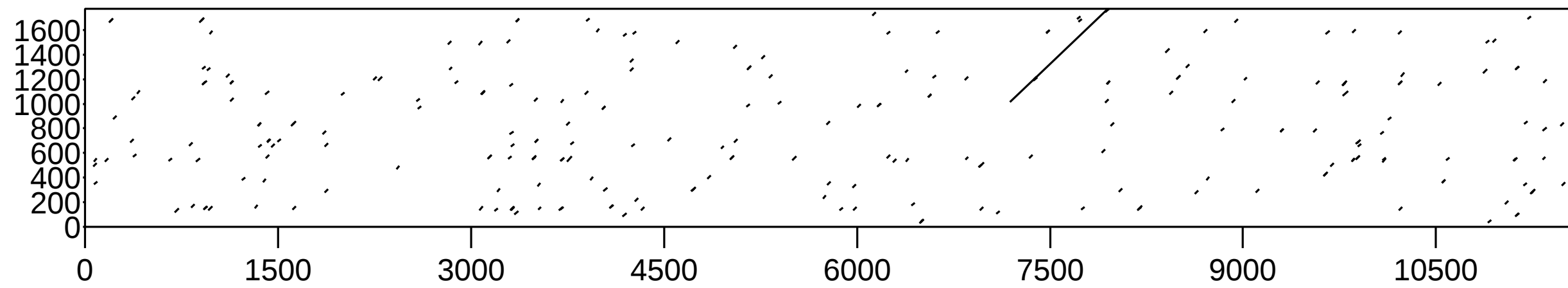
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)