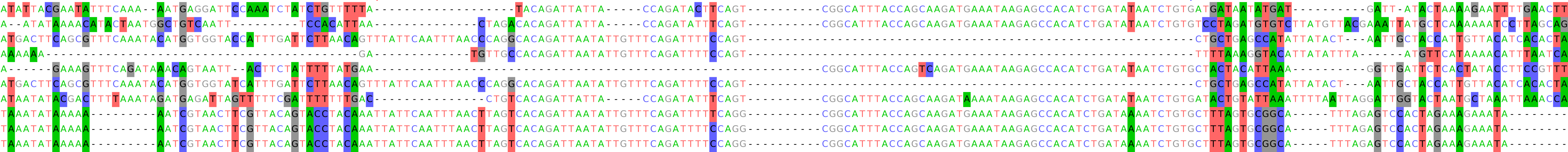


Start Crop Point

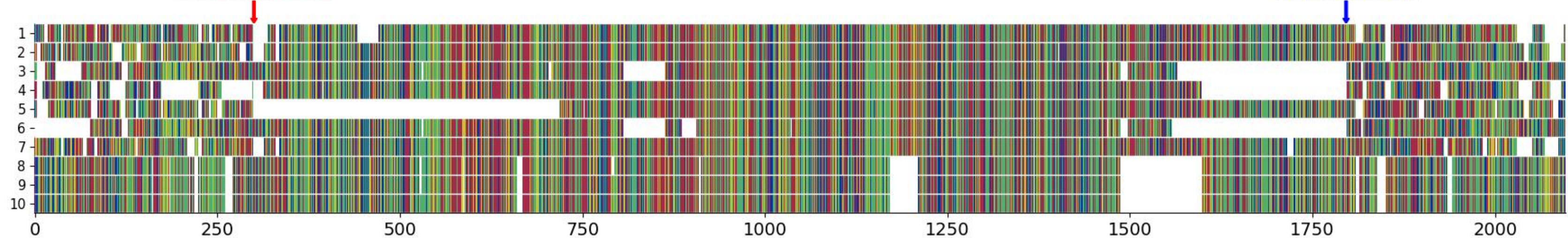
End Crop Point

MSA length = 1497



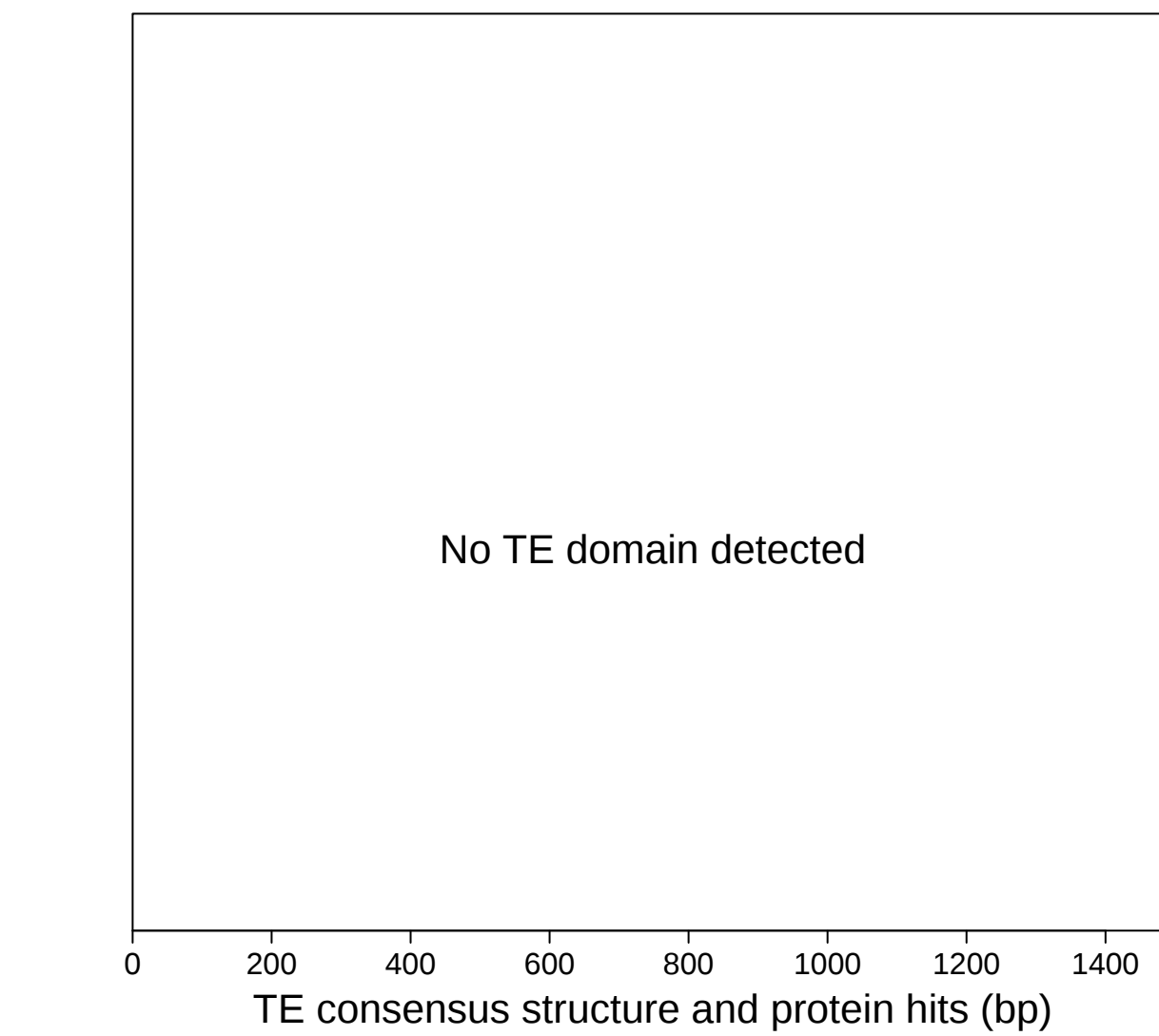
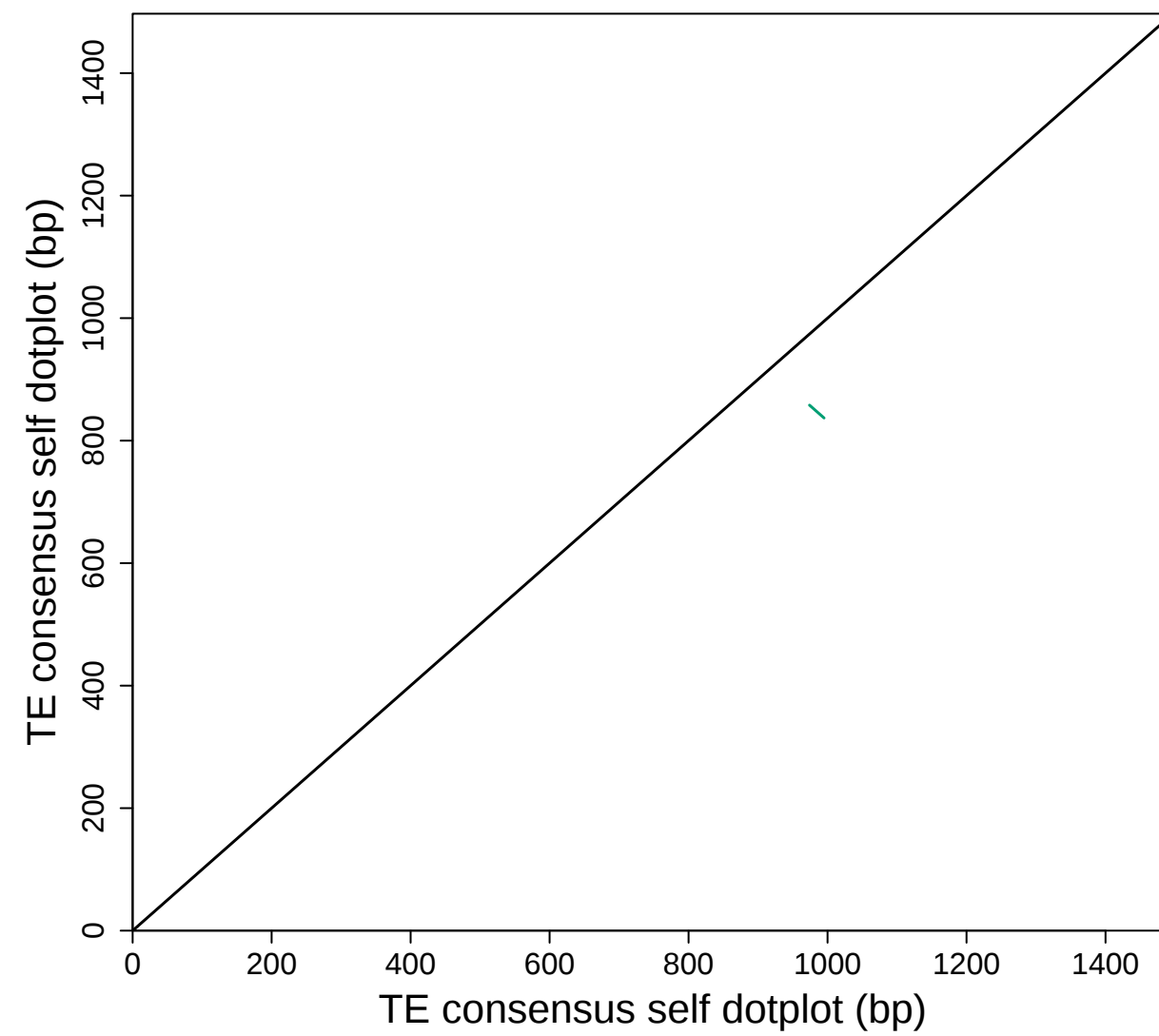
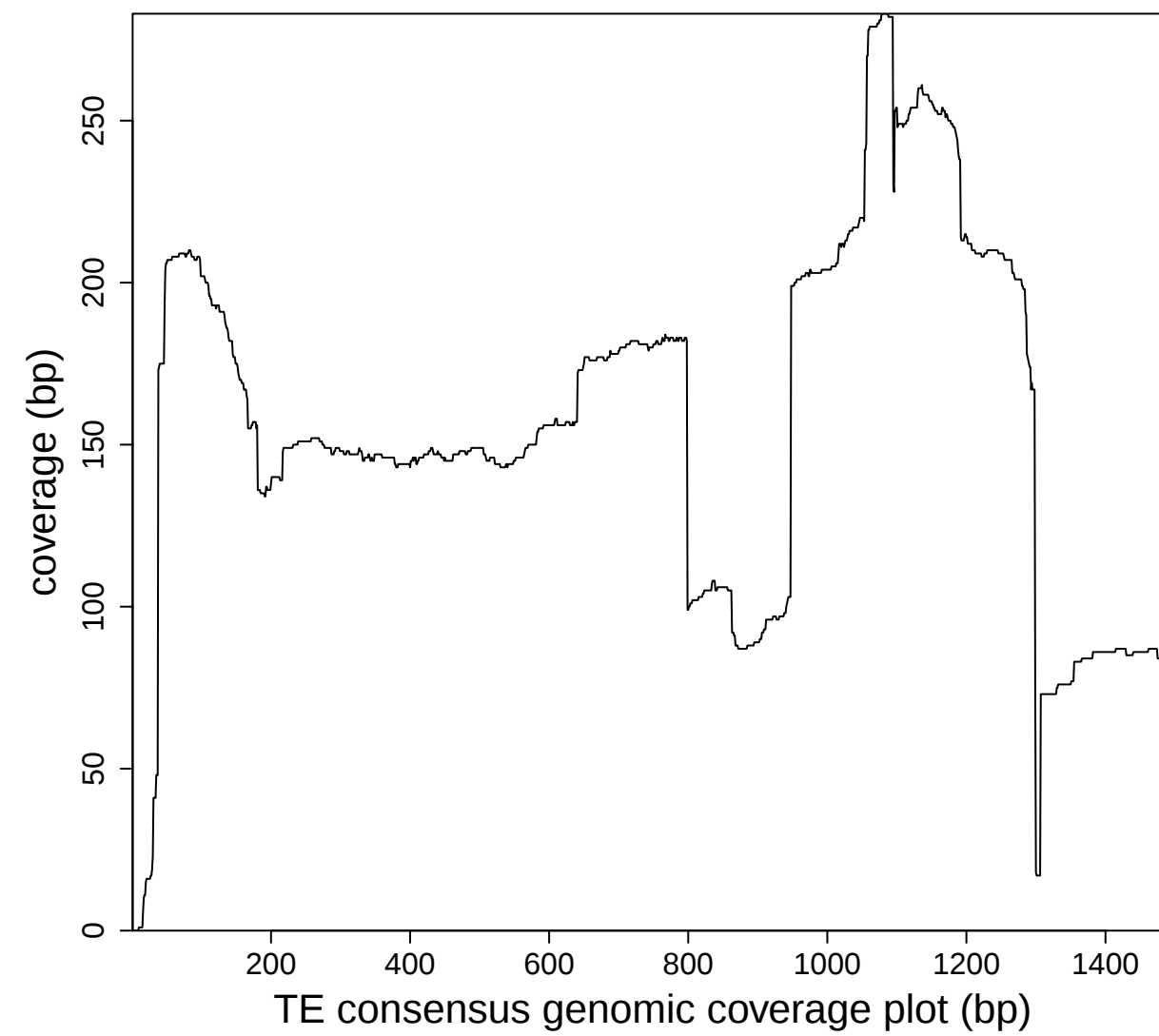
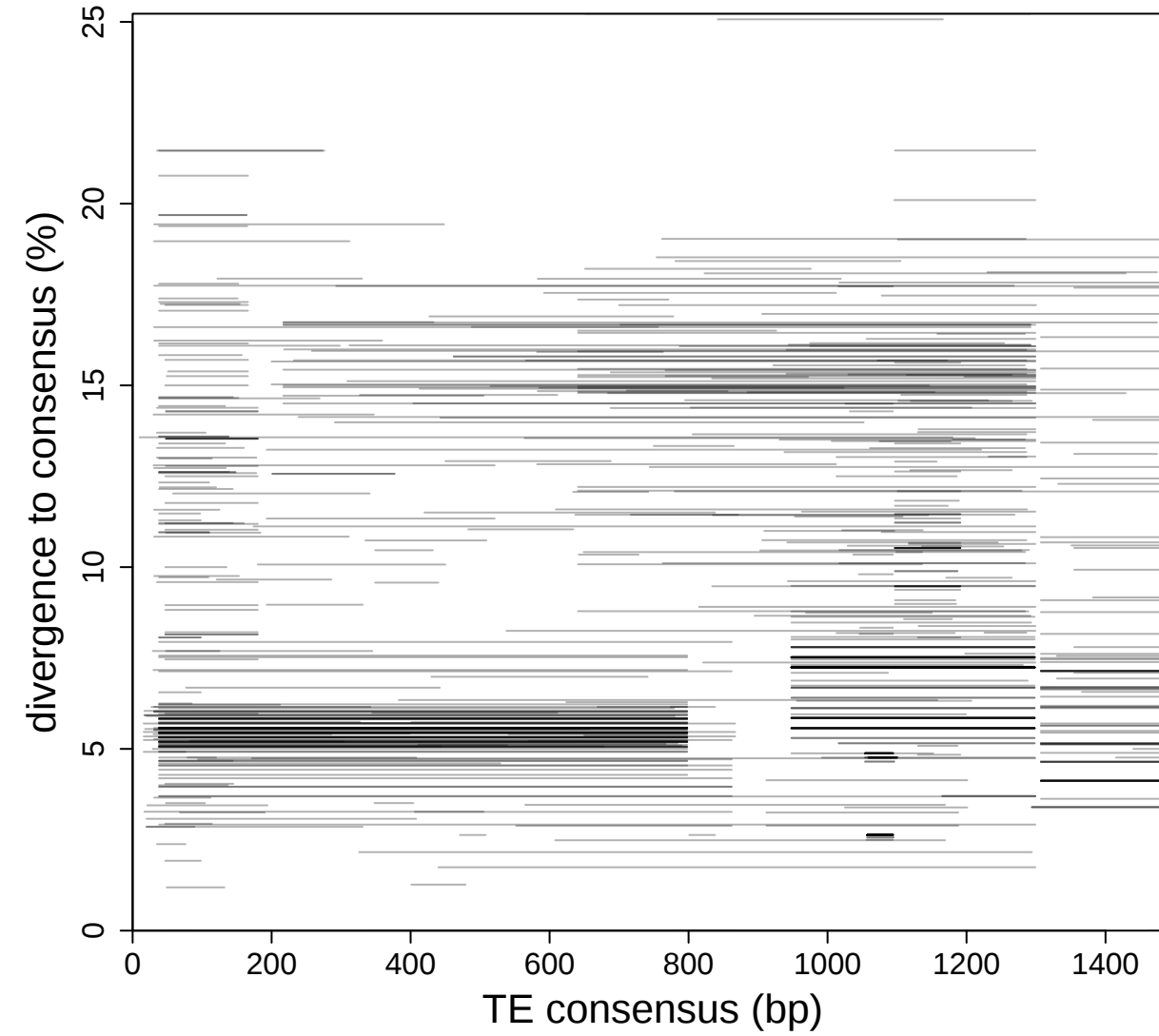
Start crop Point

End crop Point



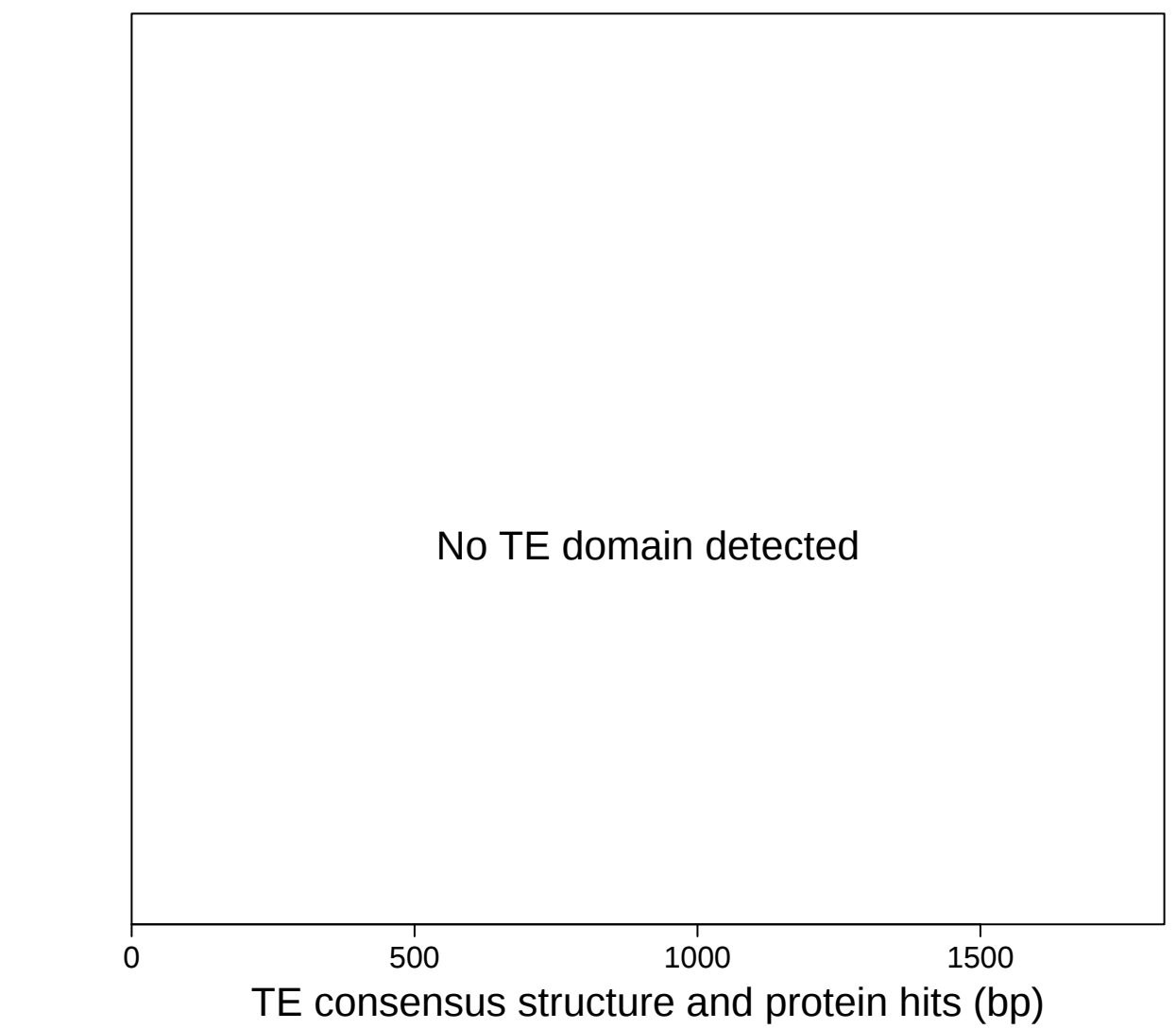
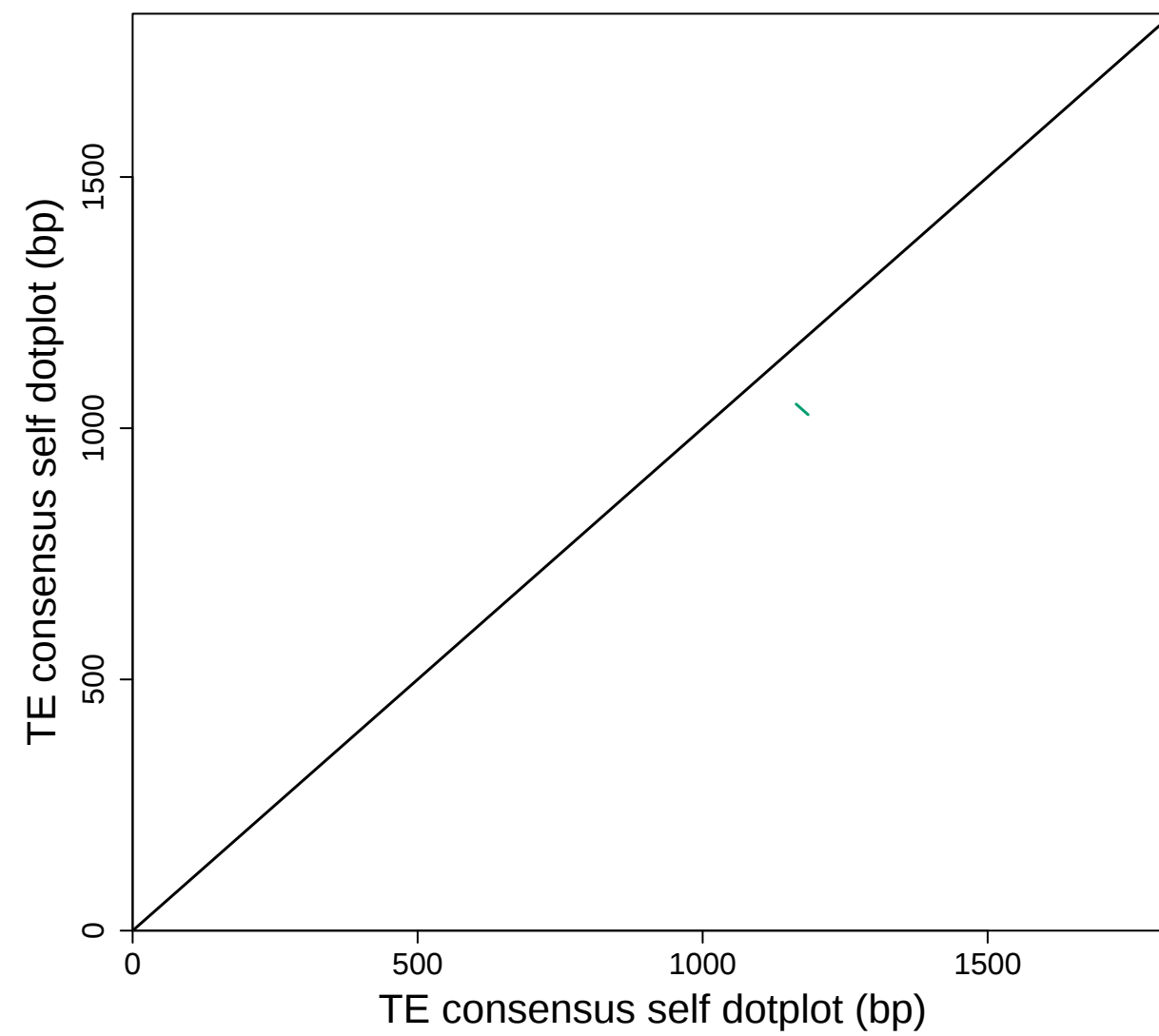
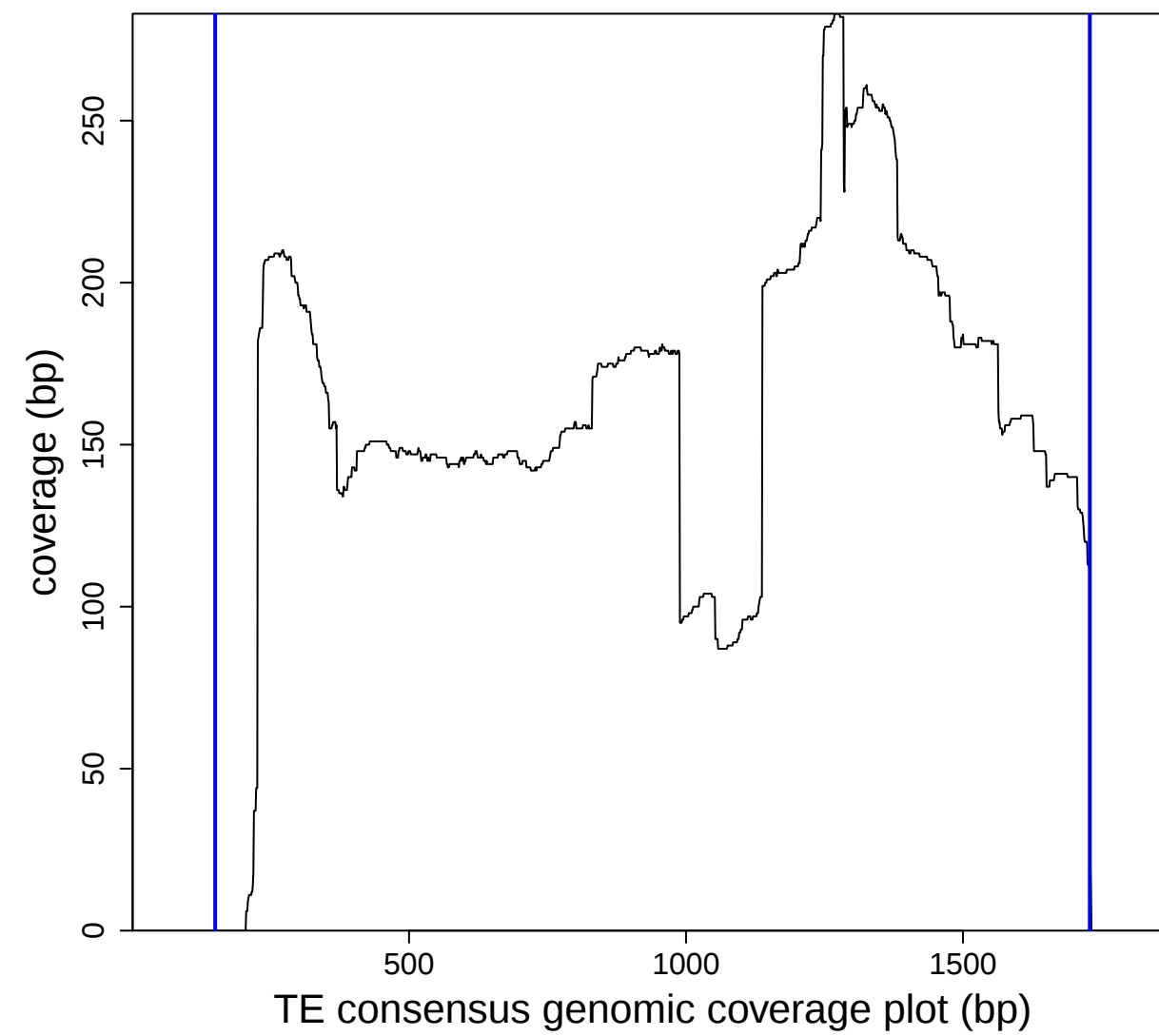
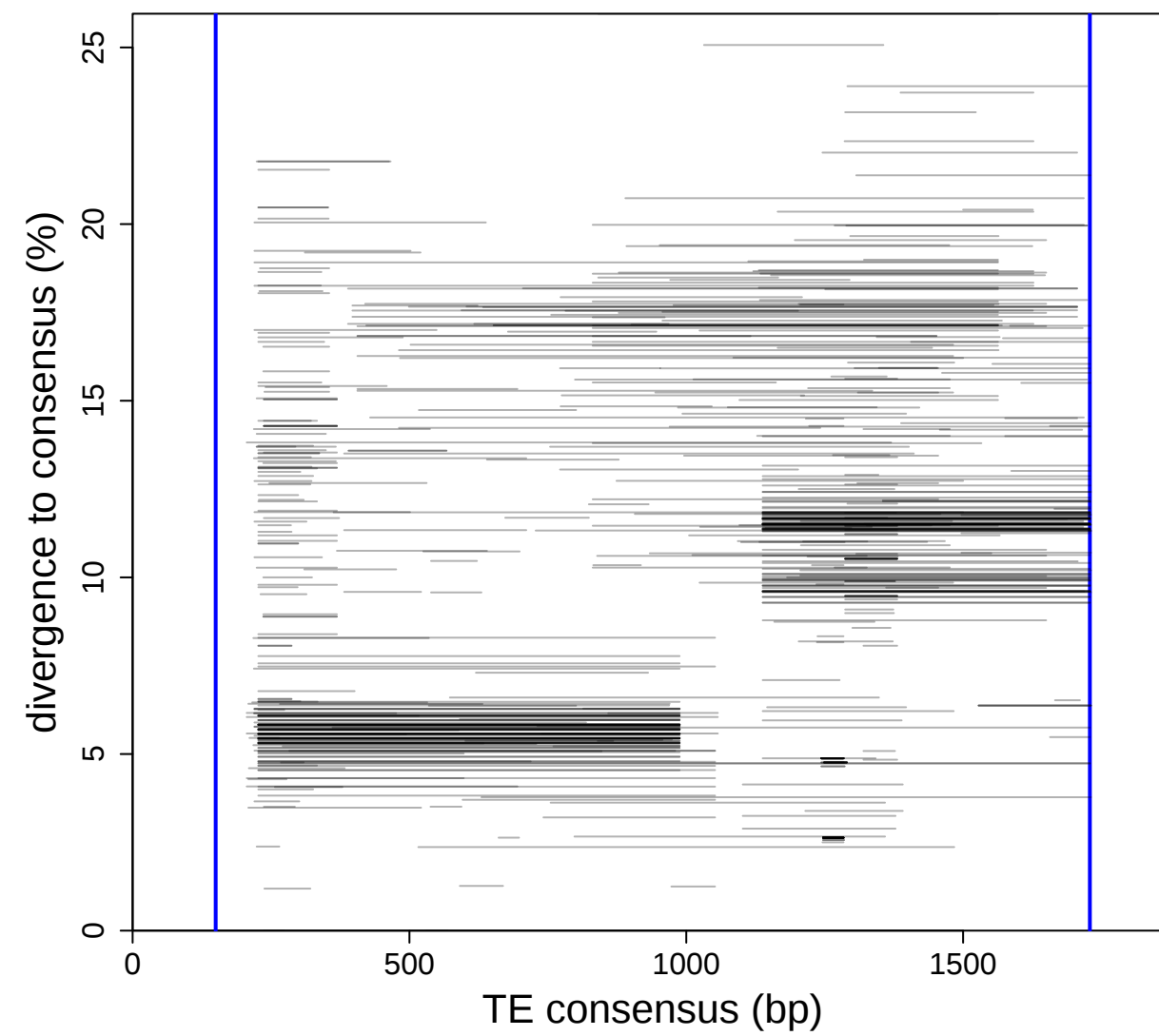
a.b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.f
size: 1497bp; fragments: 675; full length: 0 (>=1347.3bp)

After TEtrimmer 1497 bp

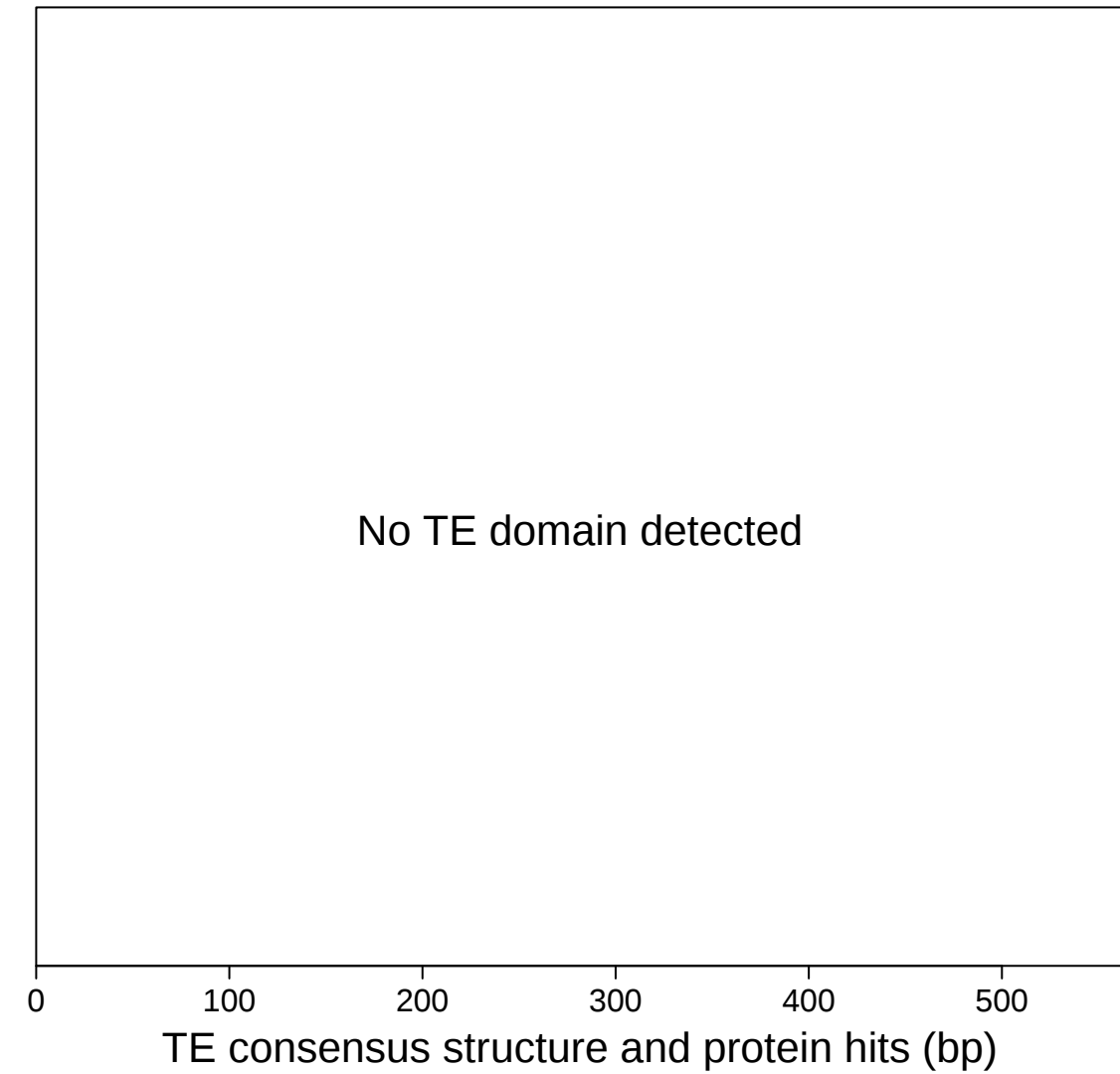
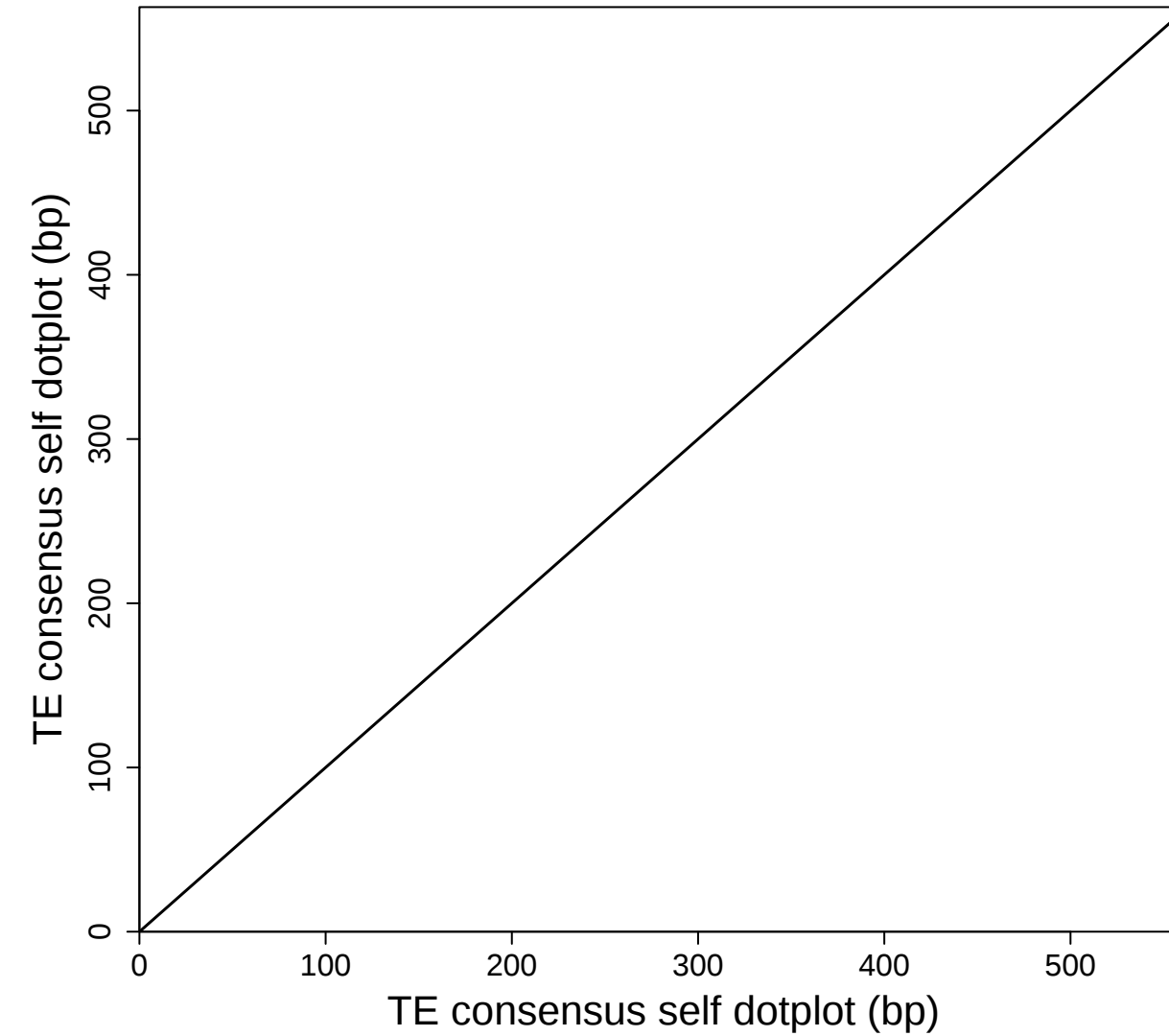
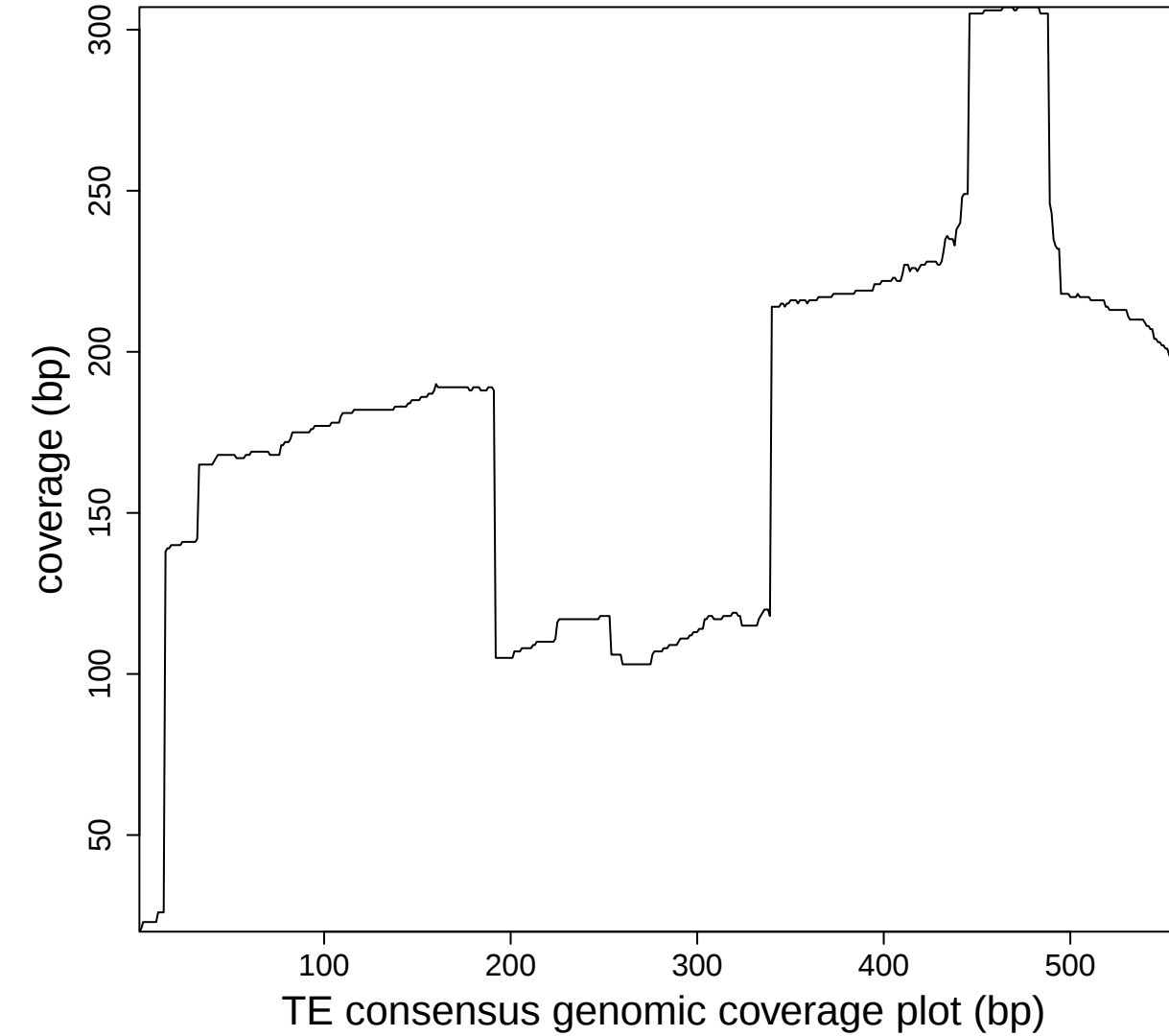
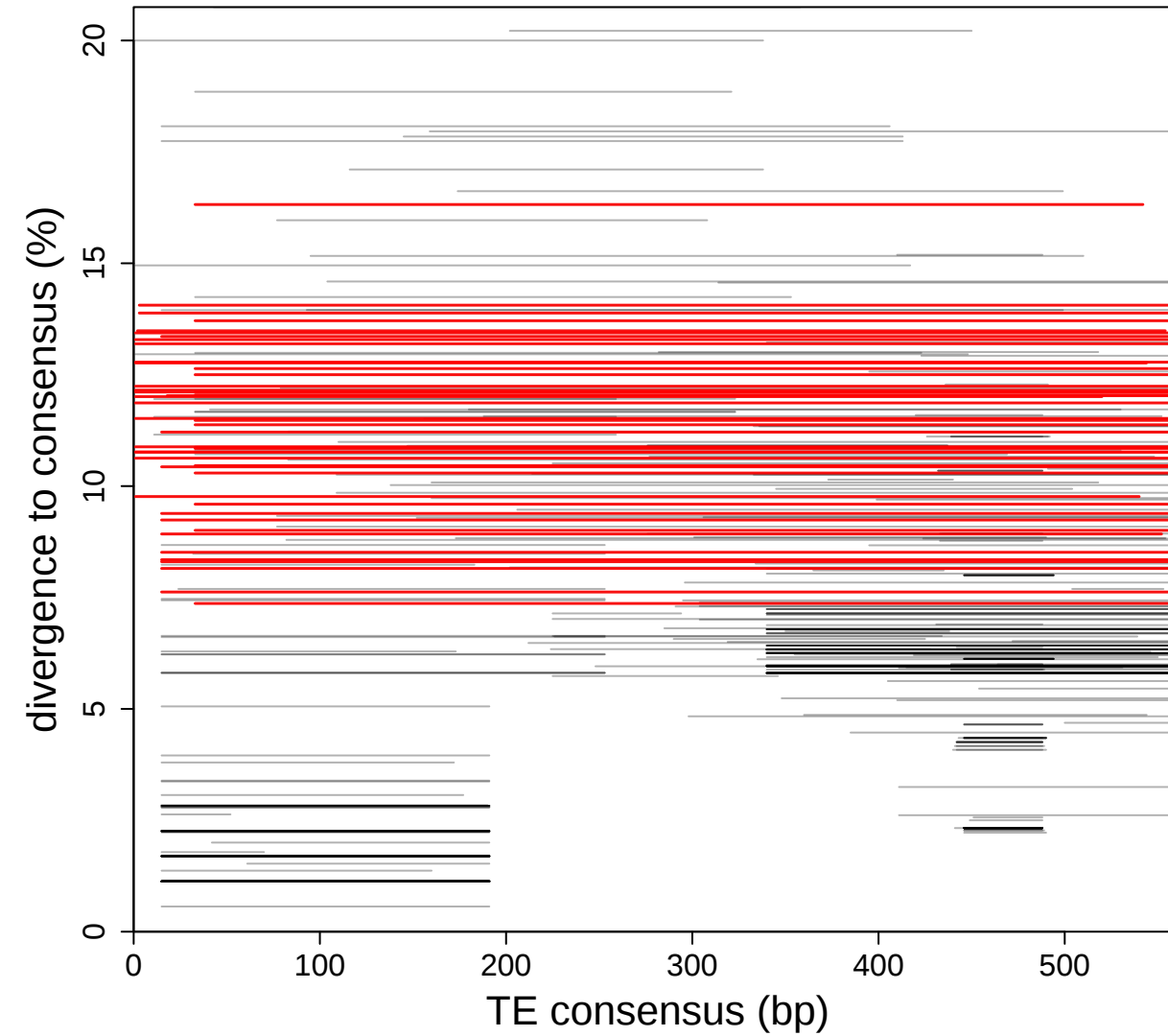


1.fasta.b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcIn.fa_aln.fa_cl.f
size: 1879bp; fragments: 613; full length: 0 (>=1691.1bp)

After TEtrimmer Extended plot Blue lines are boundaries

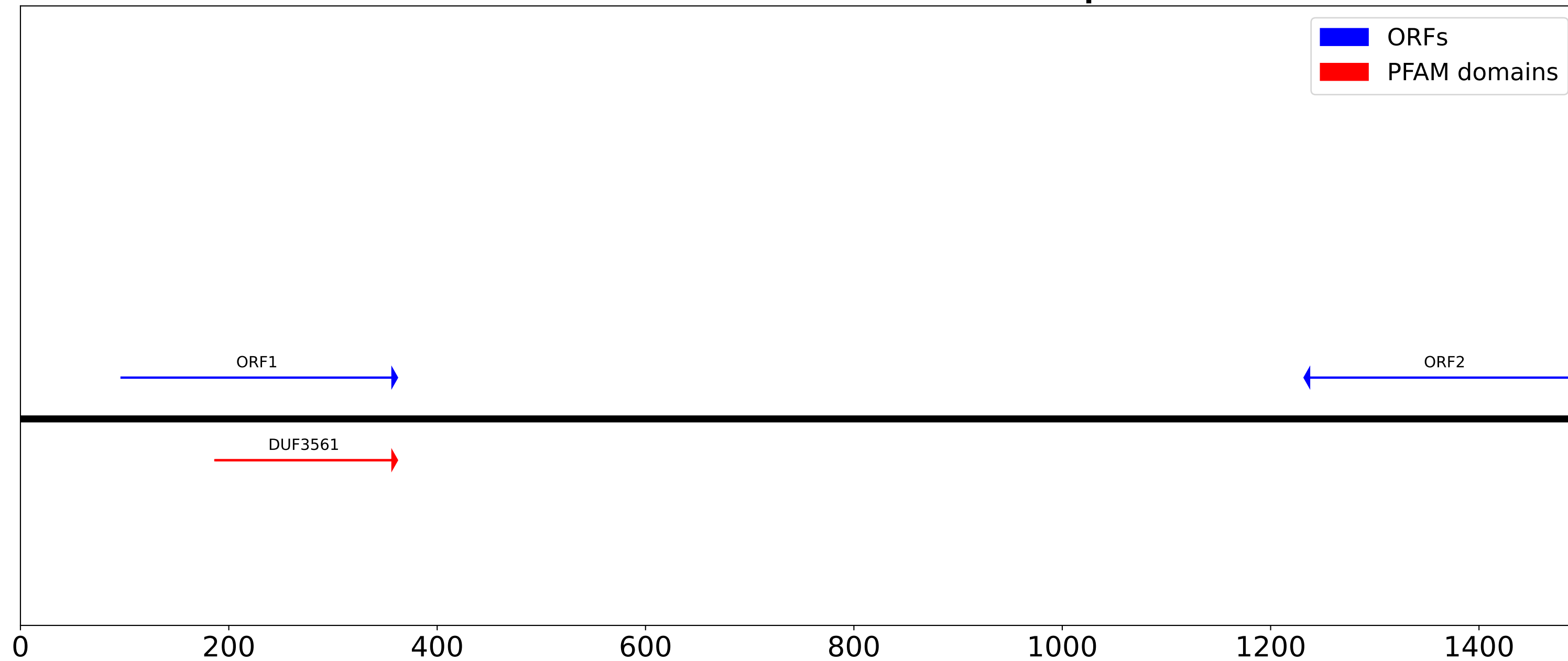


TE: rnd_5_family_8201
size: 563bp; fragments: 443; full length: 47 (≥ 506.7 bp)



Before TEtrimmer 563 bp

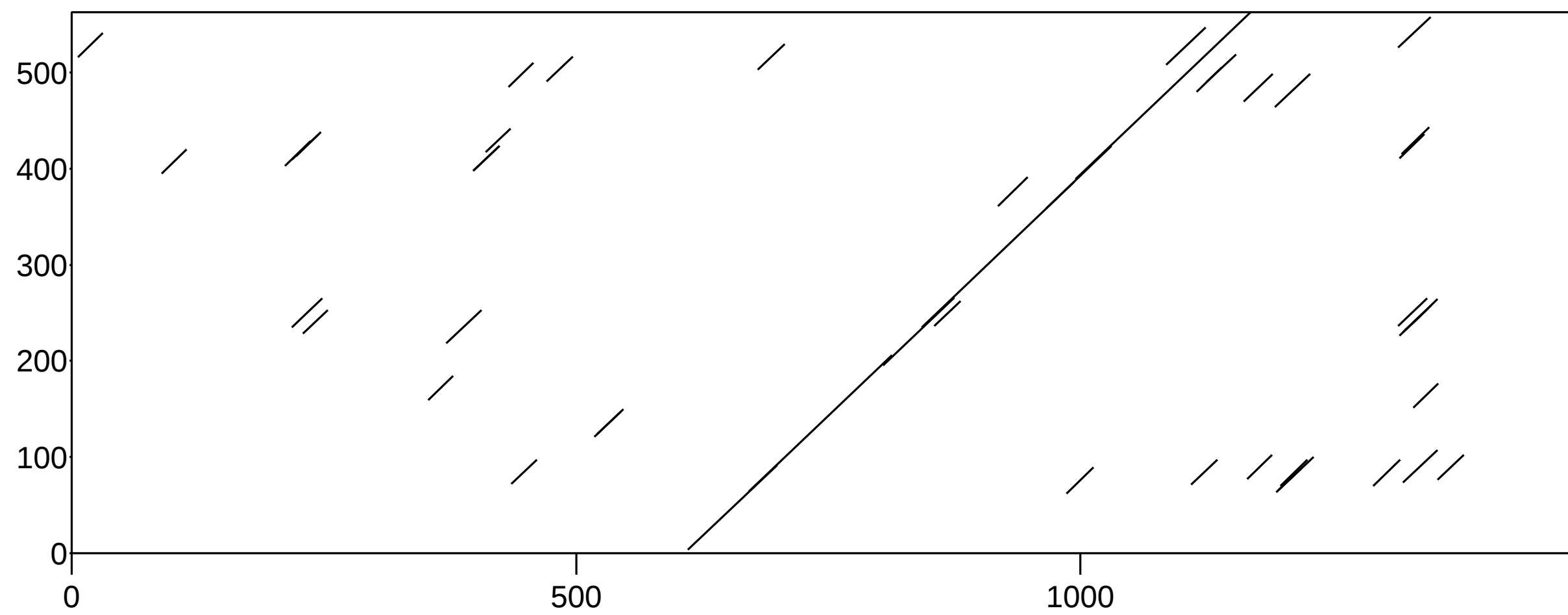
After TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)