

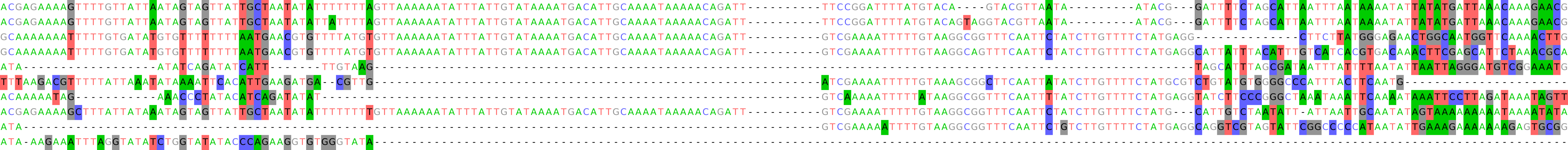
Start Crop Point



End Crop Point

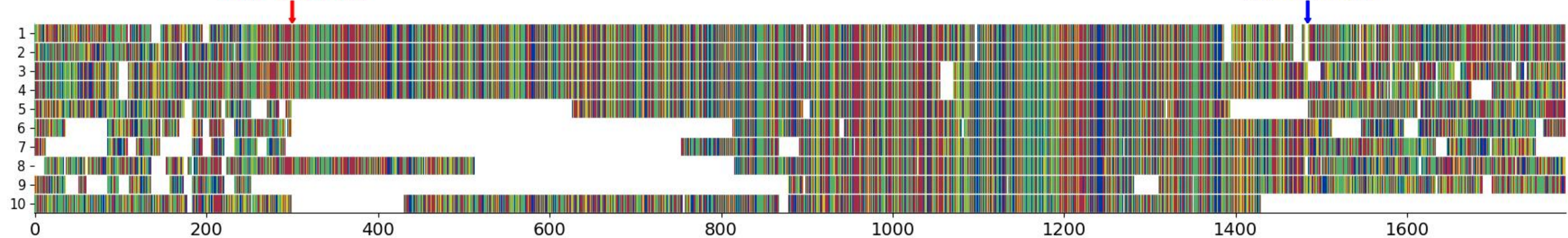


MSA length = 1185



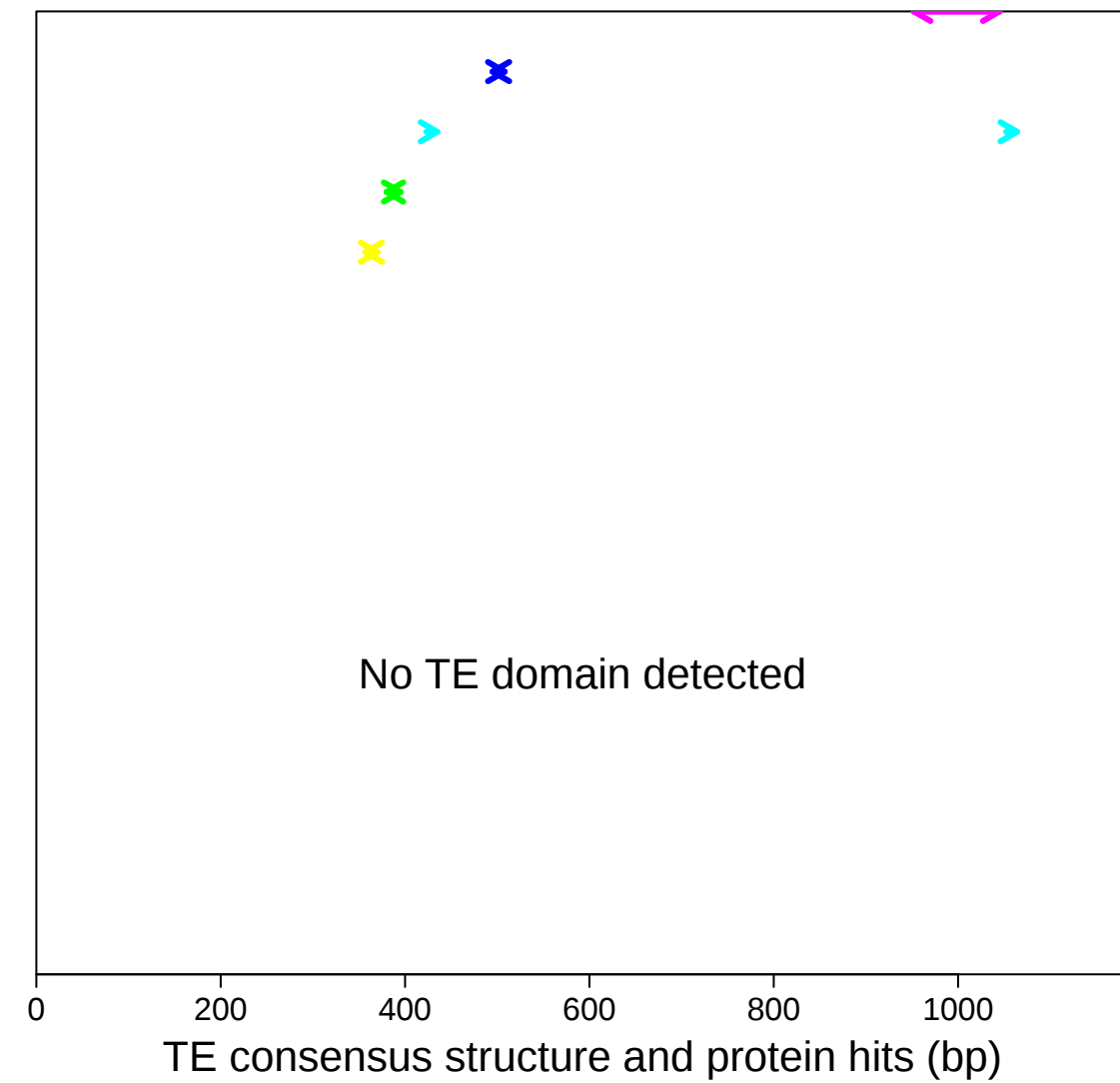
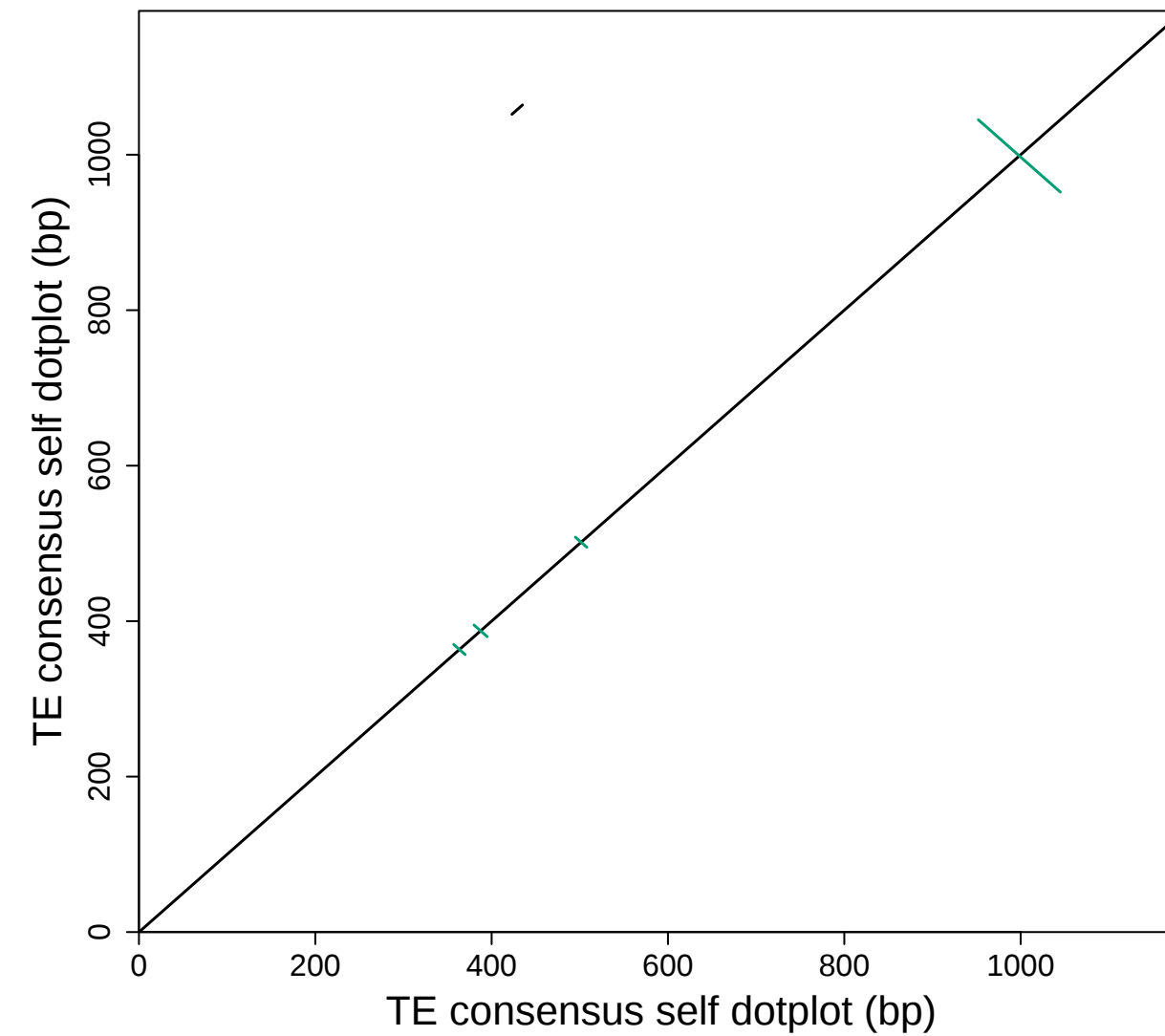
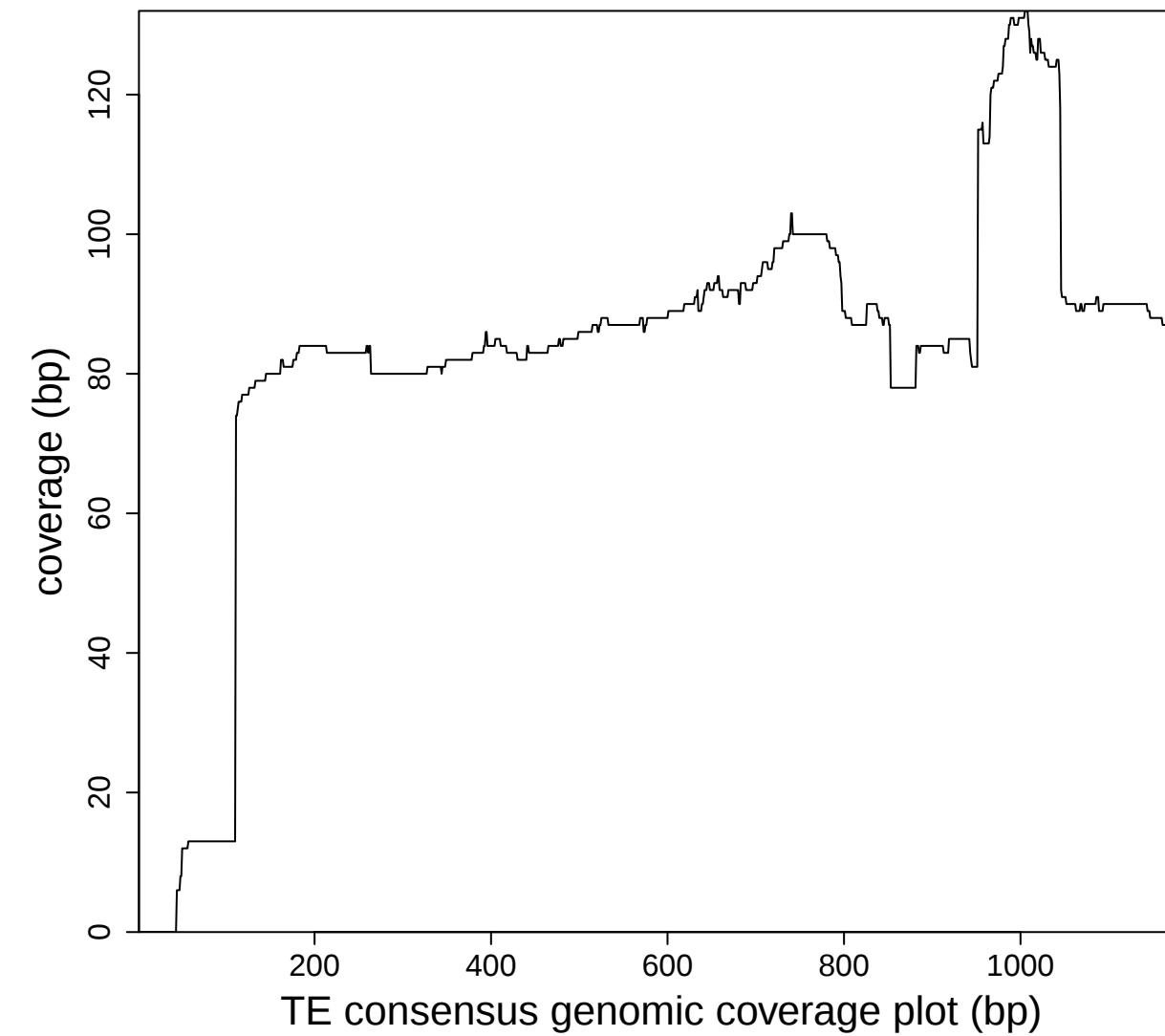
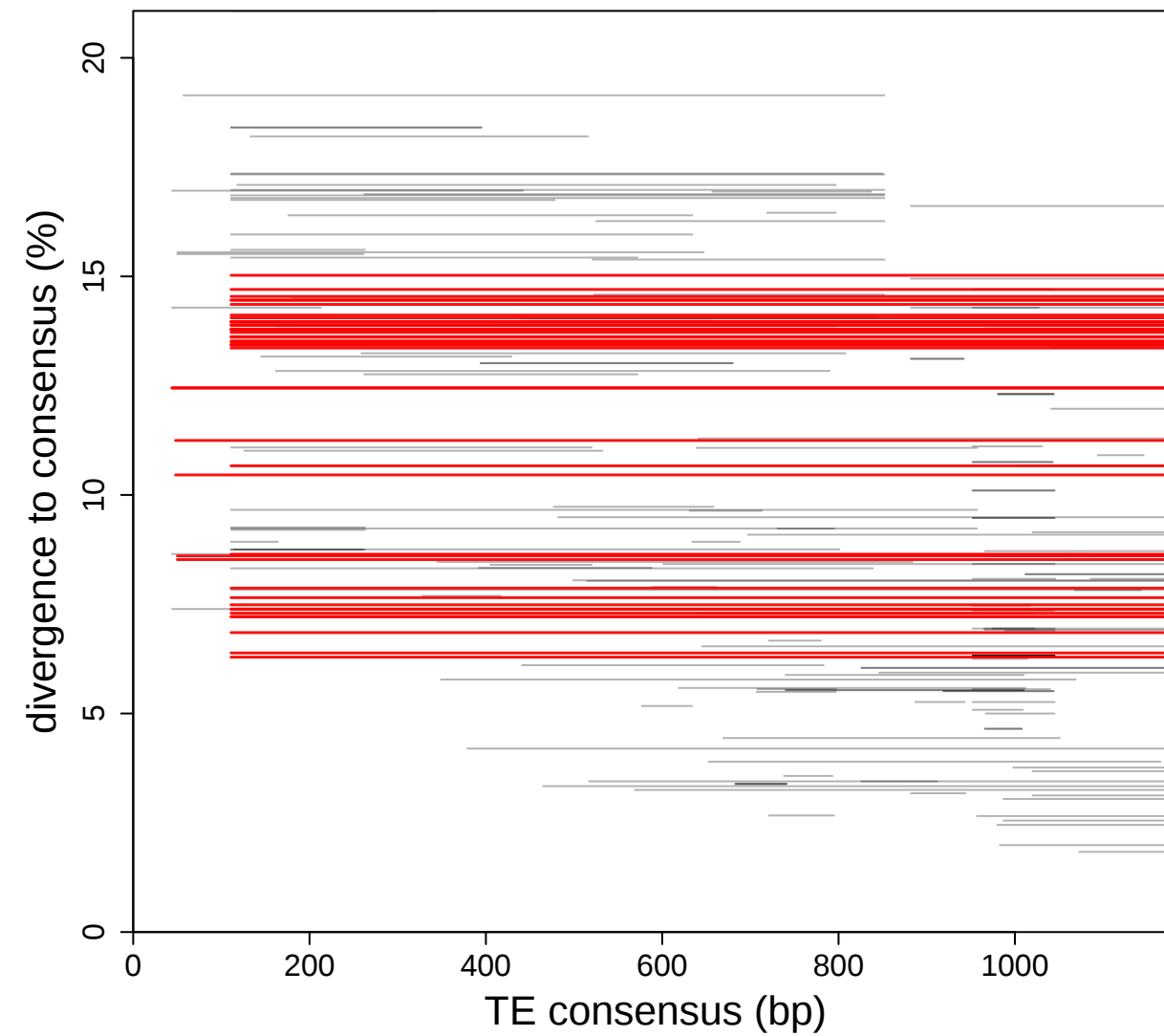
Start crop Point

End crop Point



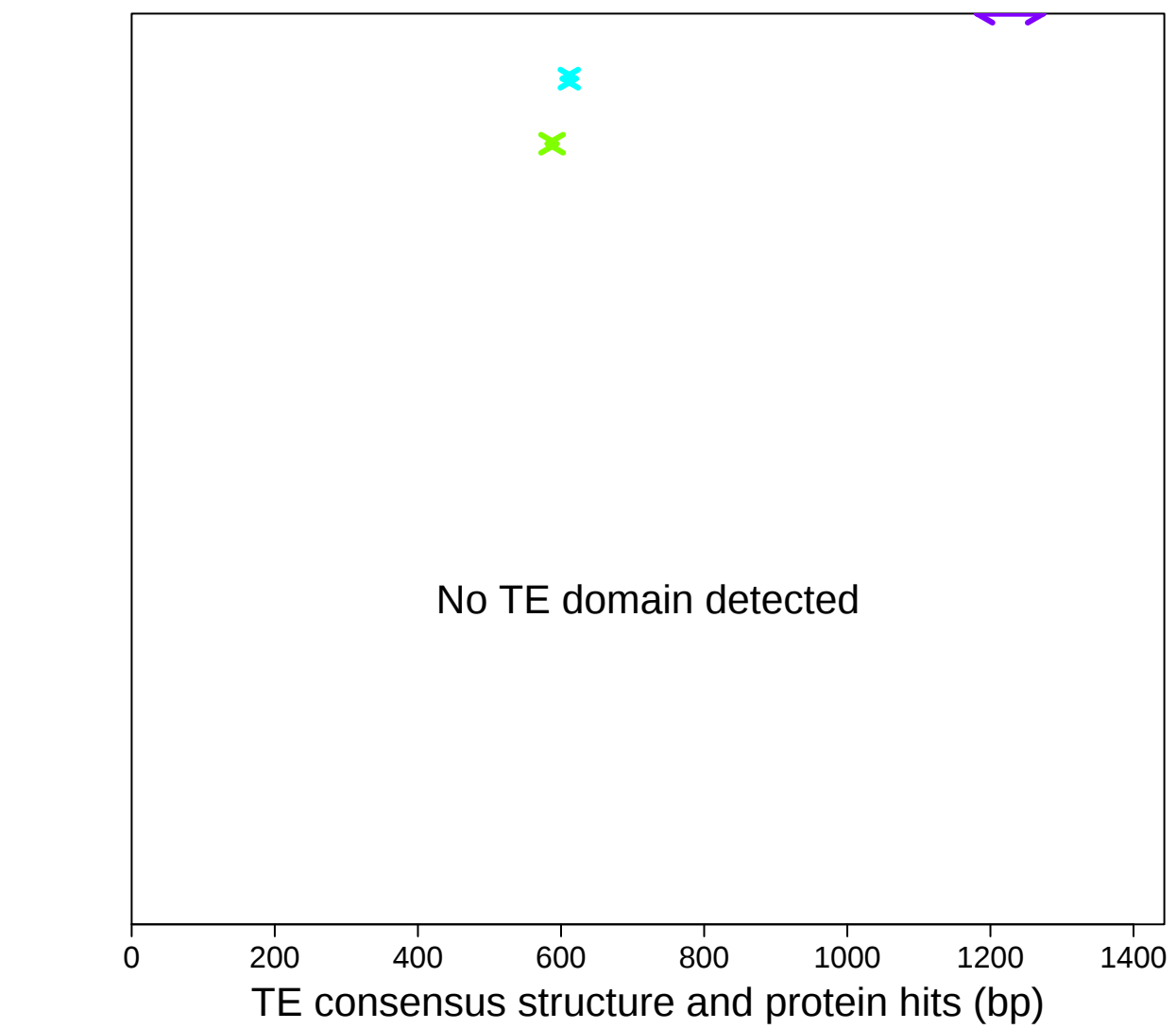
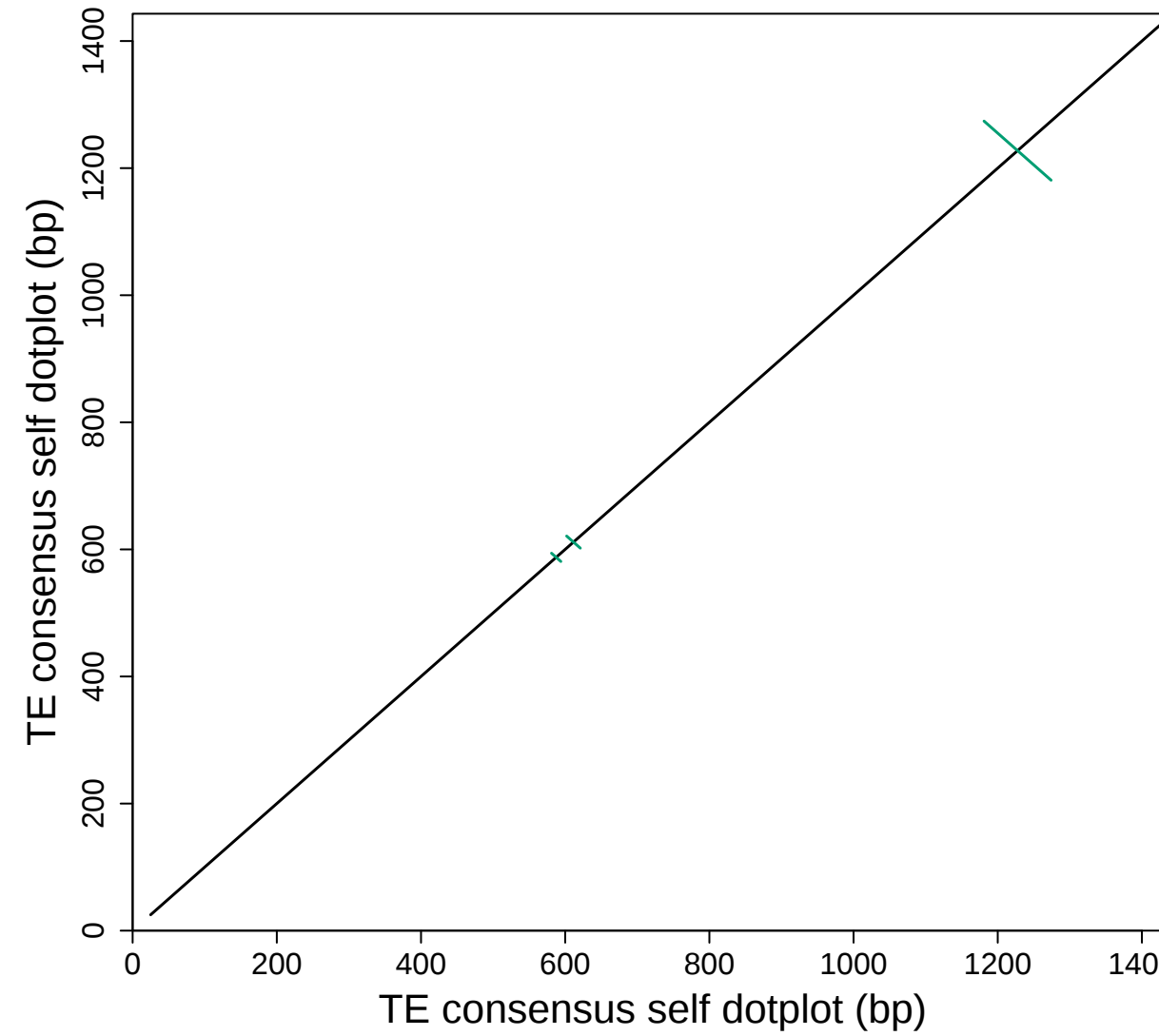
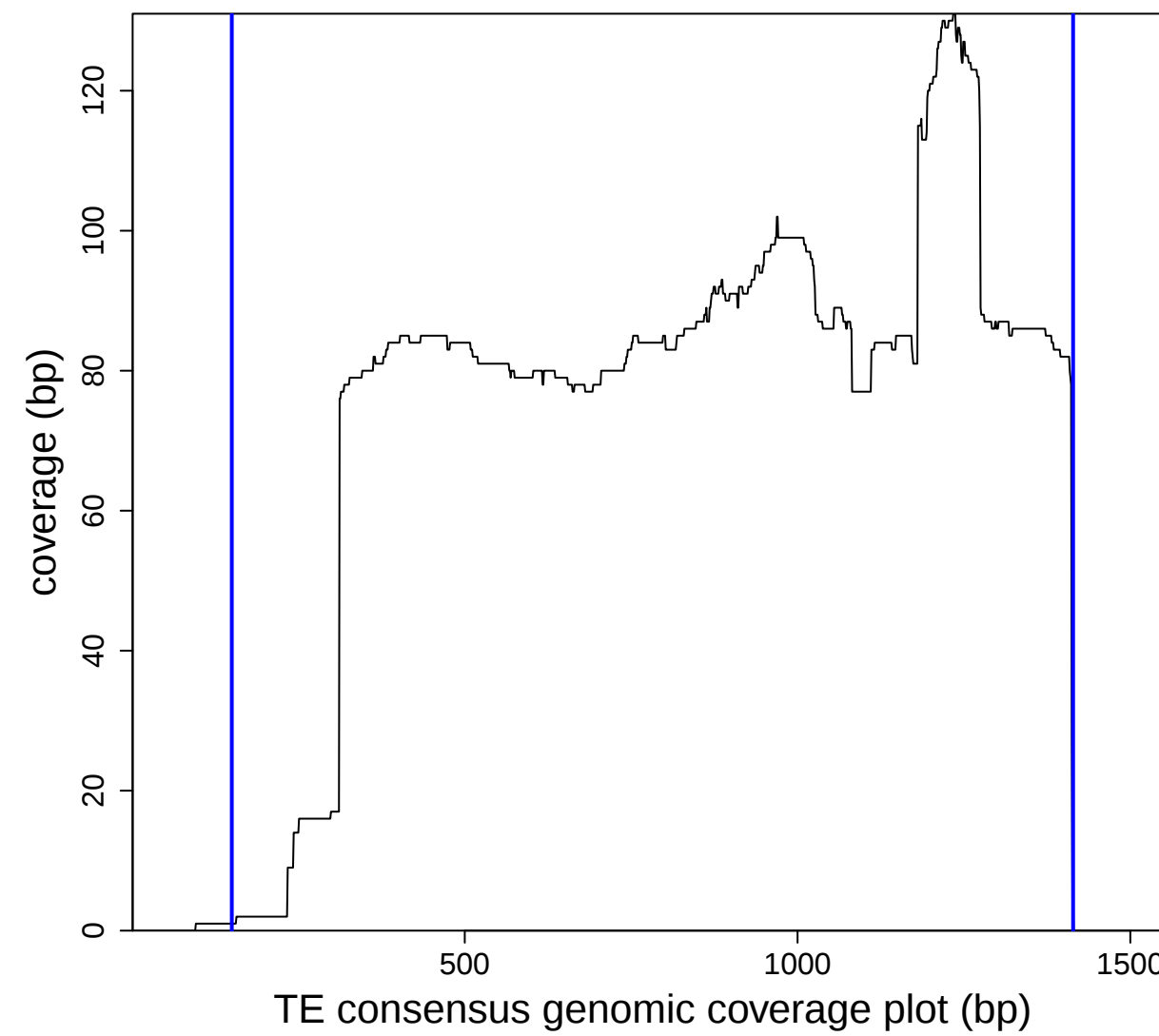
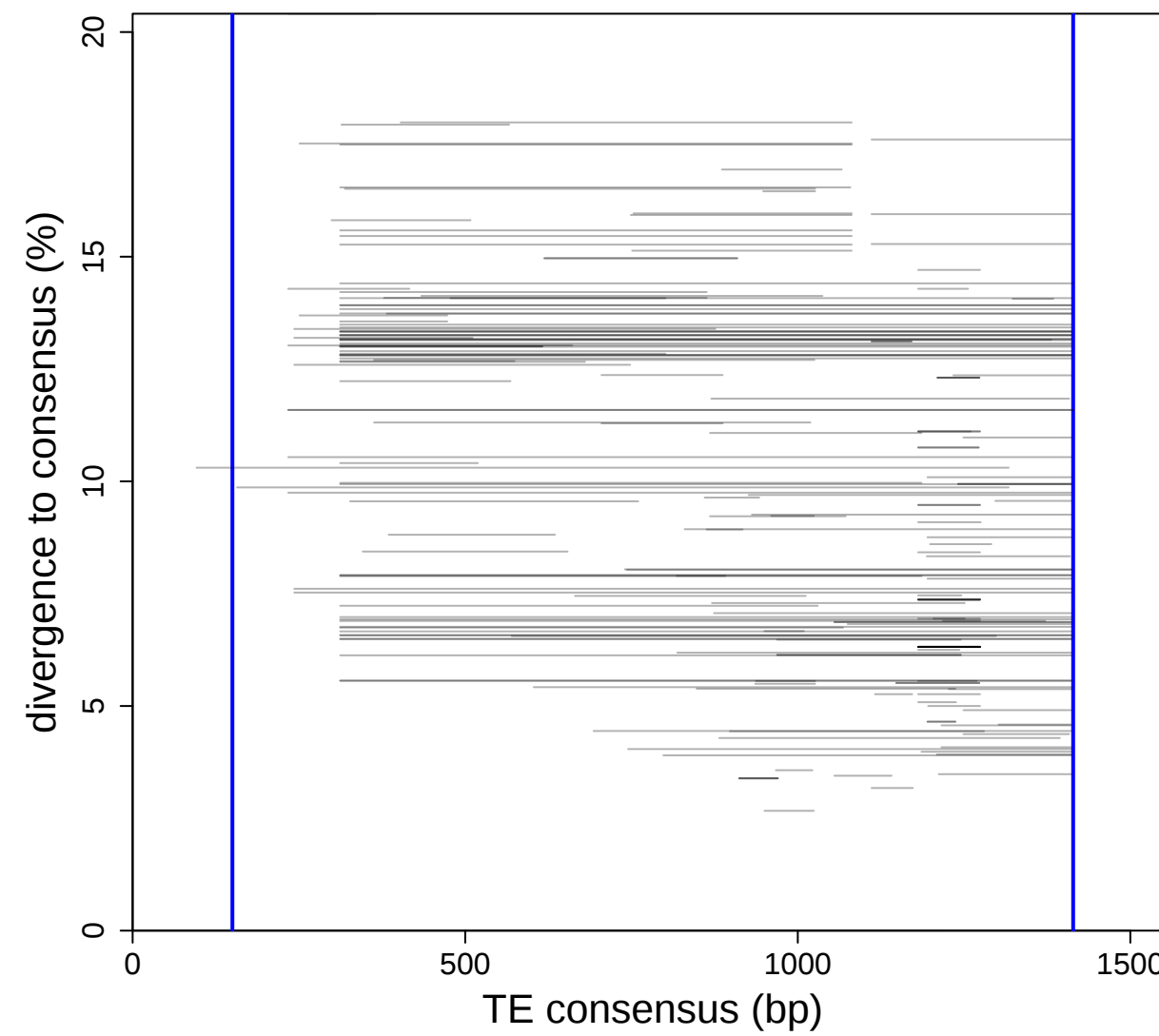
a.b.bed_uf.bed_g_3.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.f
size: 1185bp; fragments: 216; full length: 45 (≥ 1066.5 bp)

After TEtrimmer 1185 bp

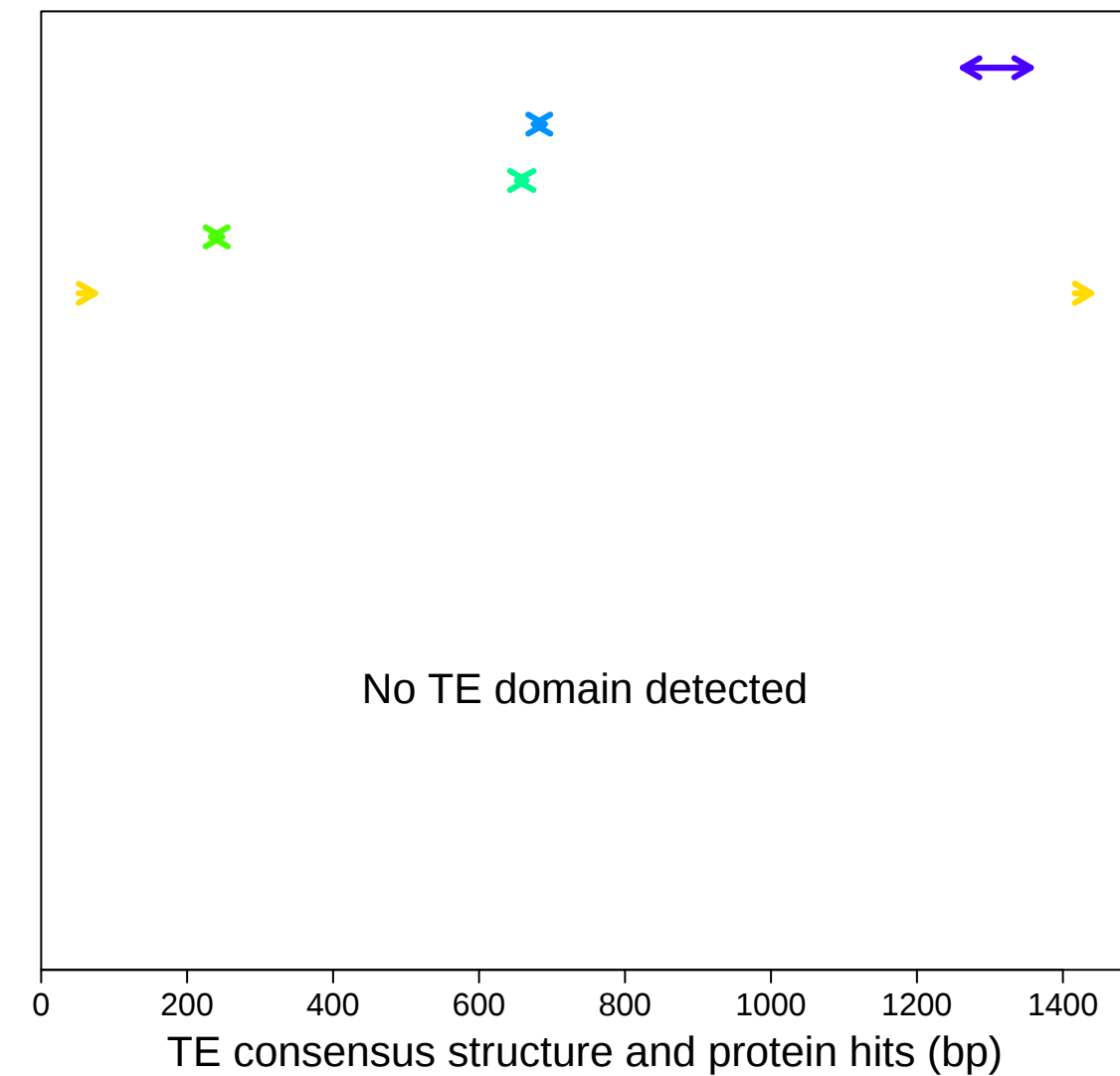
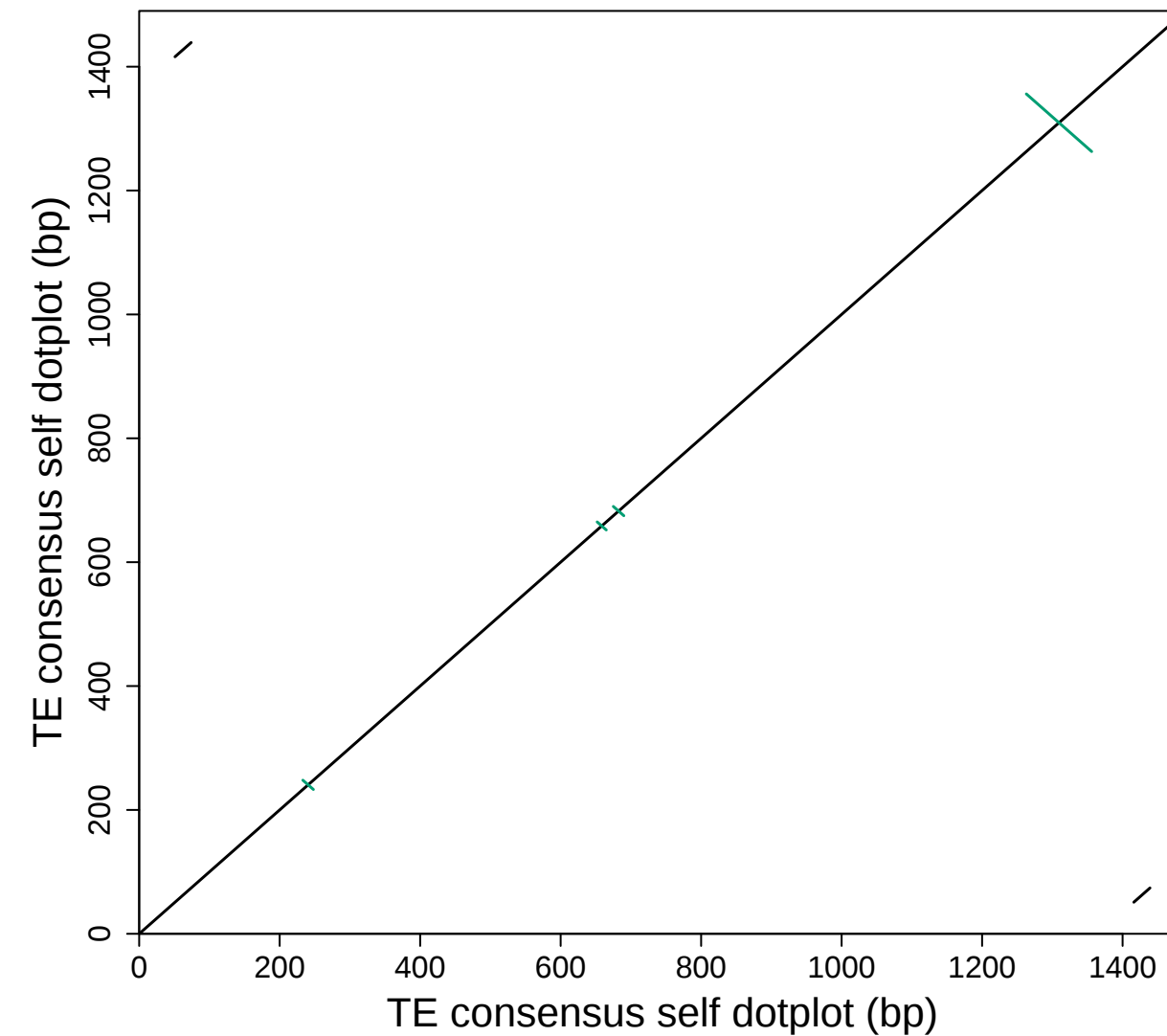
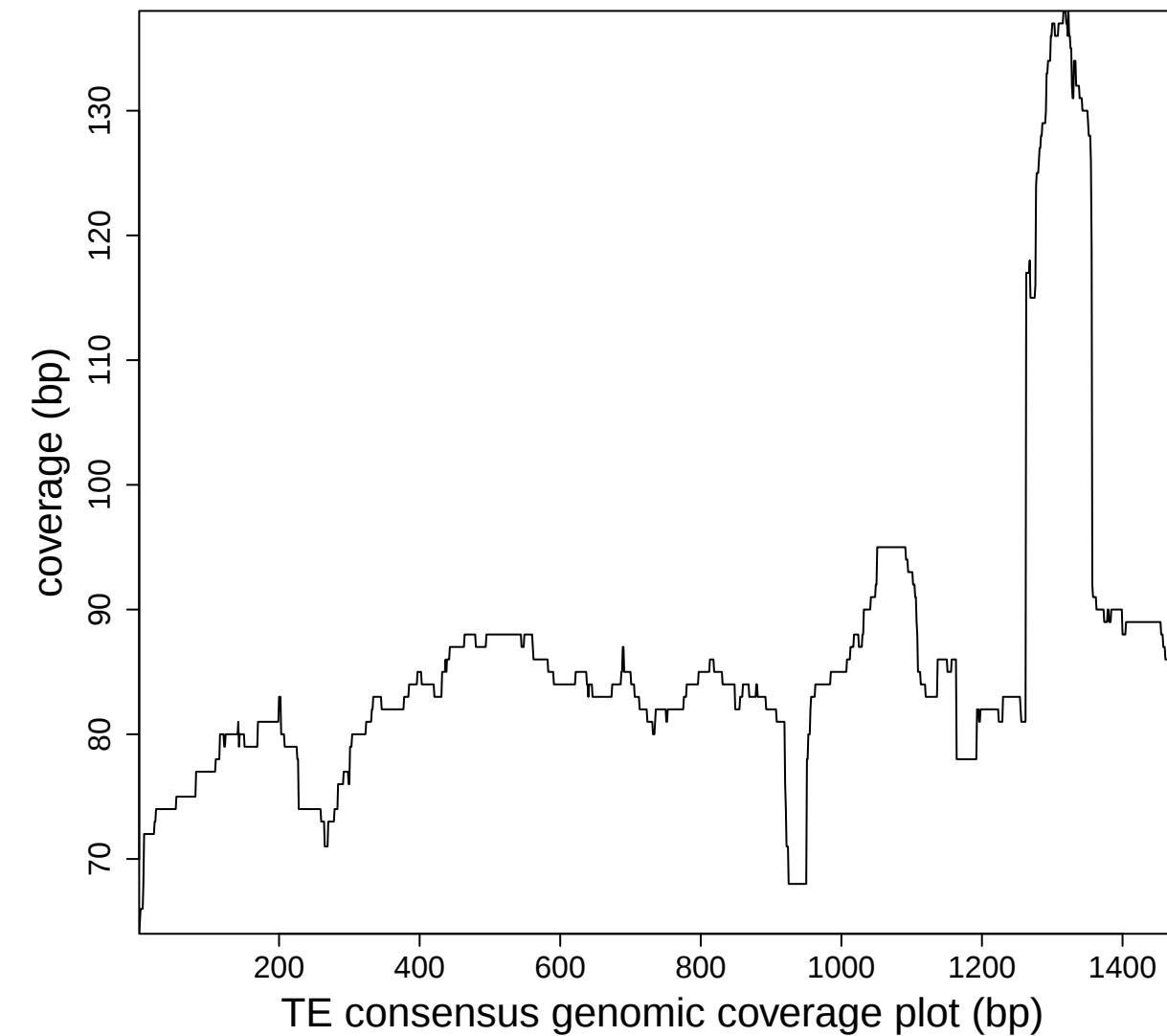
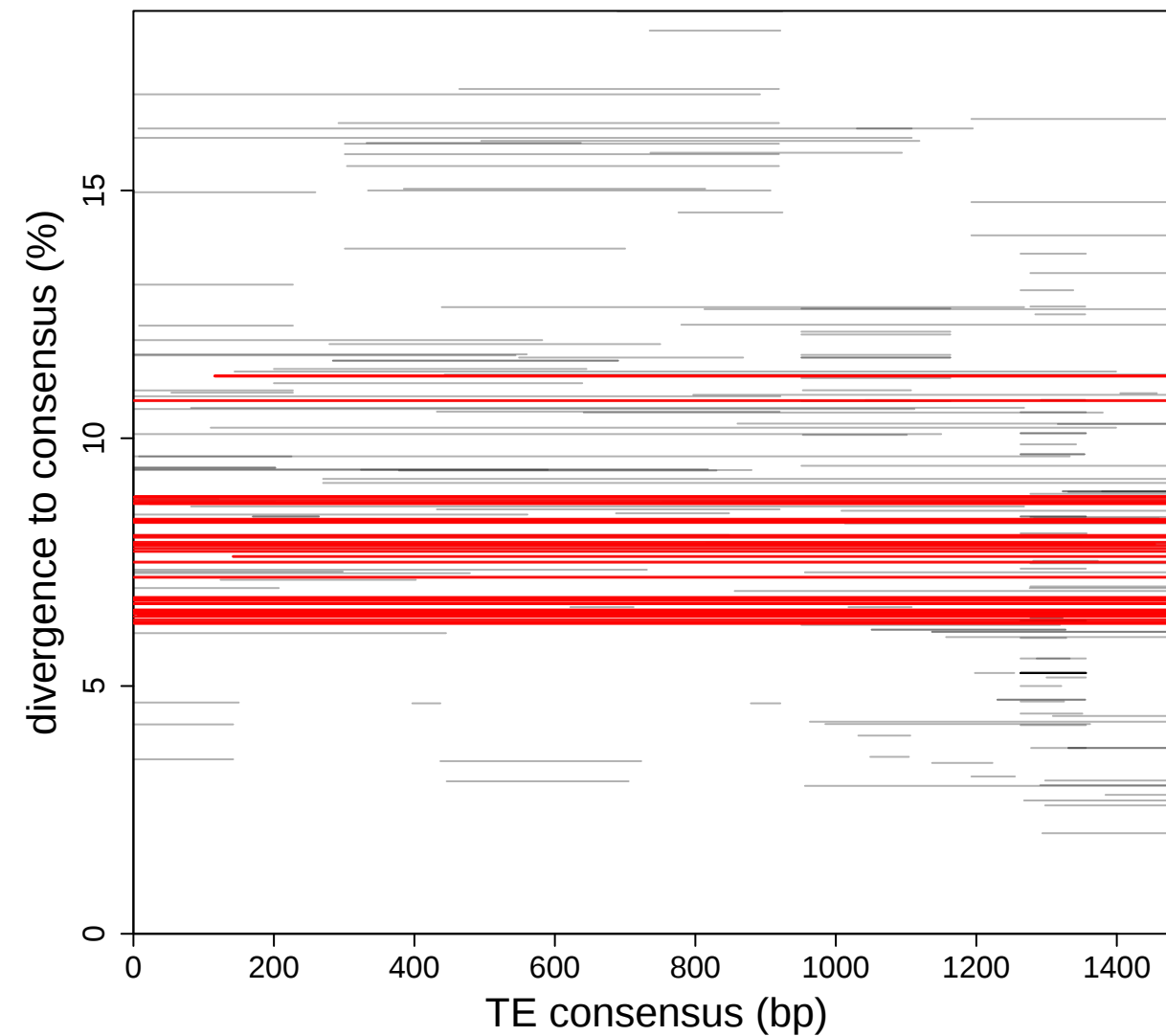


6.fasta.b.bed_uf.bed_g_3.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.f
size: 1564bp; fragments: 207; full length: 0 (>=1407.6bp)

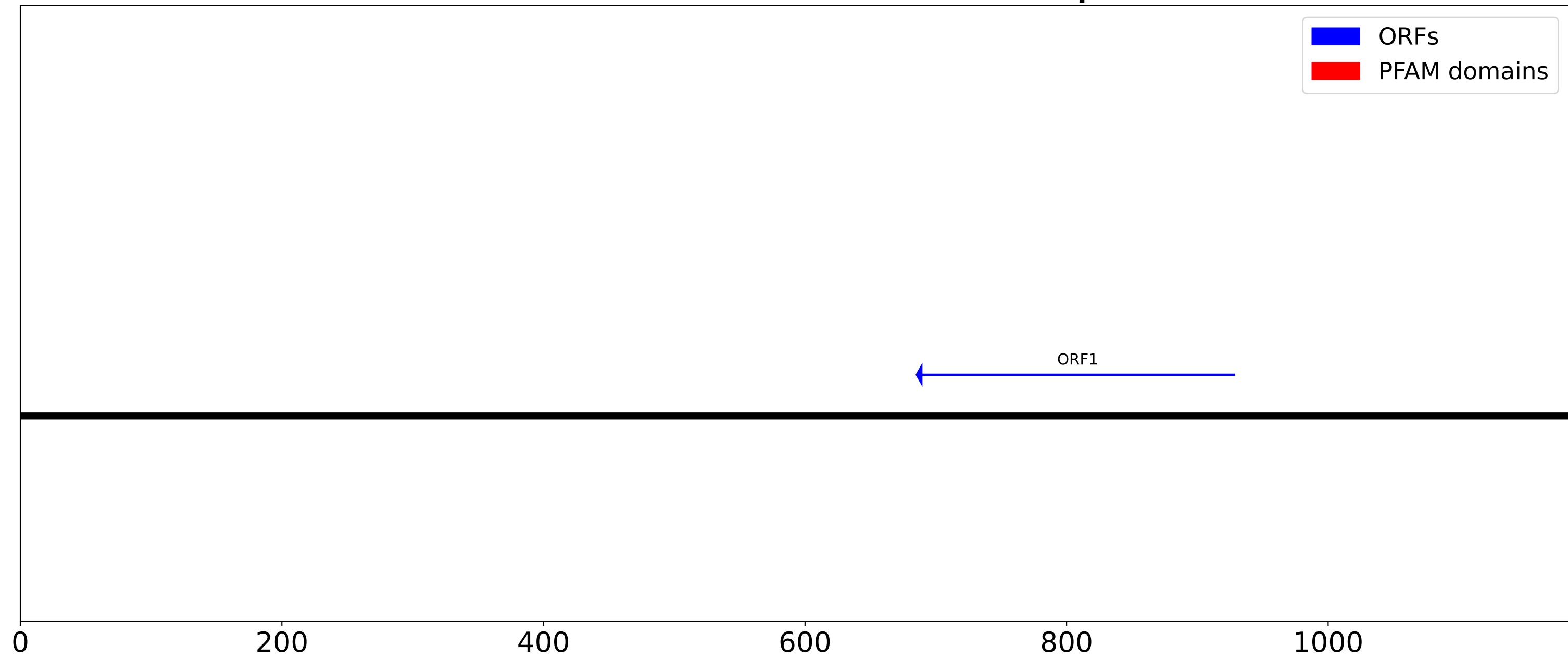
After TEtrimmer Extended plot Blue lines are boundaries



TE: rnd_5_family_3556
size: 1490bp; fragments: 237; full length: 44 (≥ 1341 bp)



After TEtrimmer ORF and PFAM domain plot



Dotplot

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

