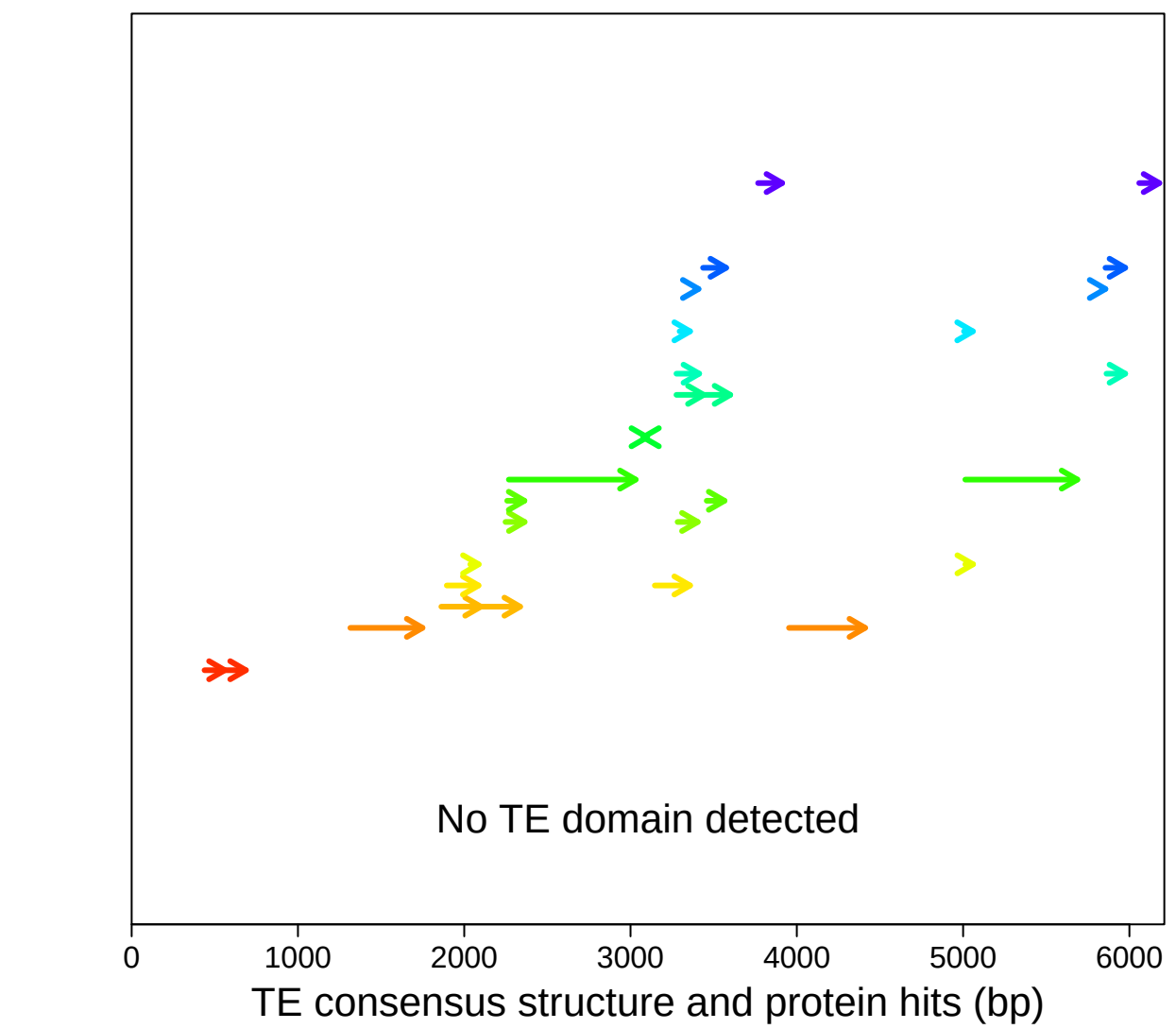
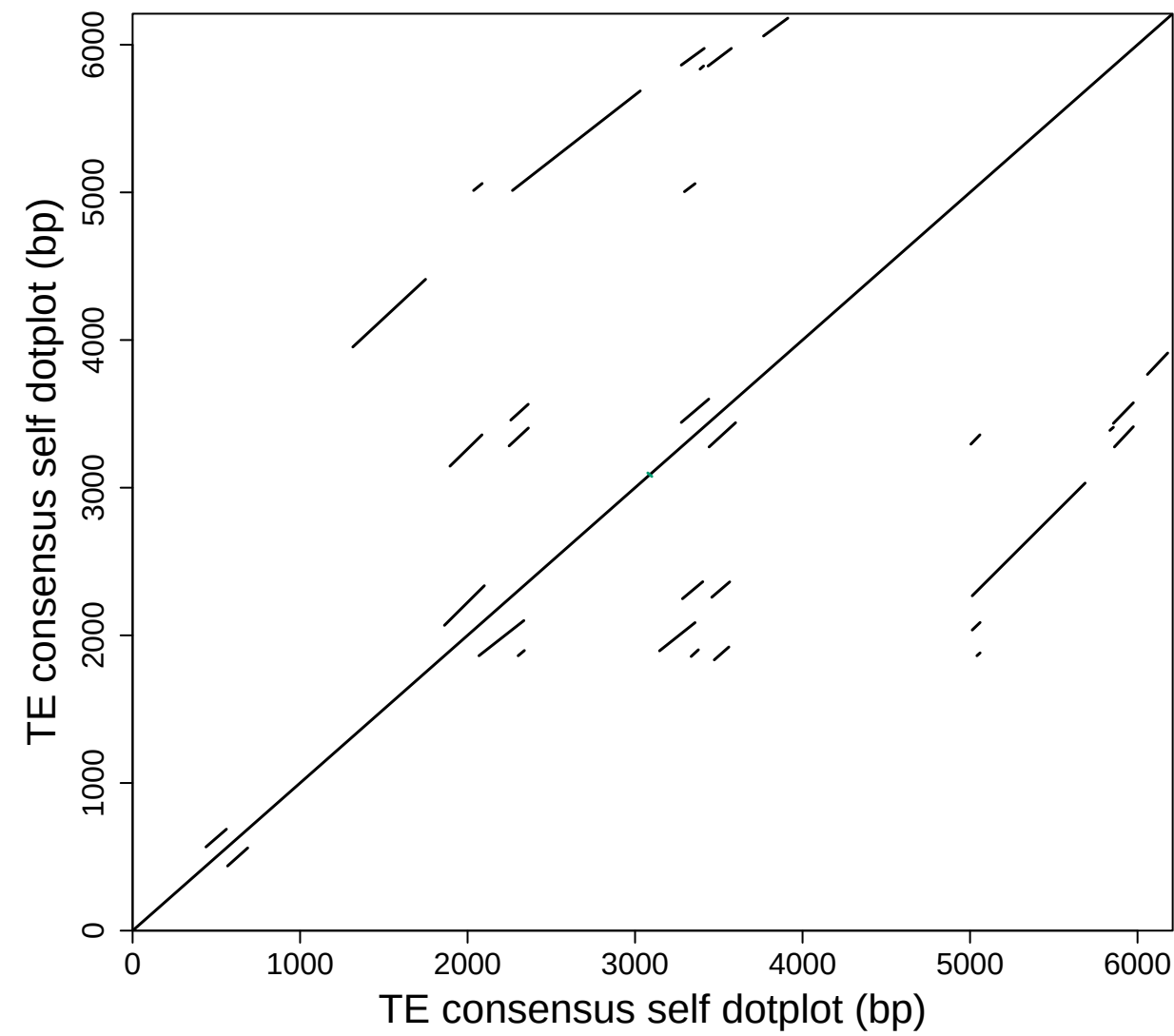
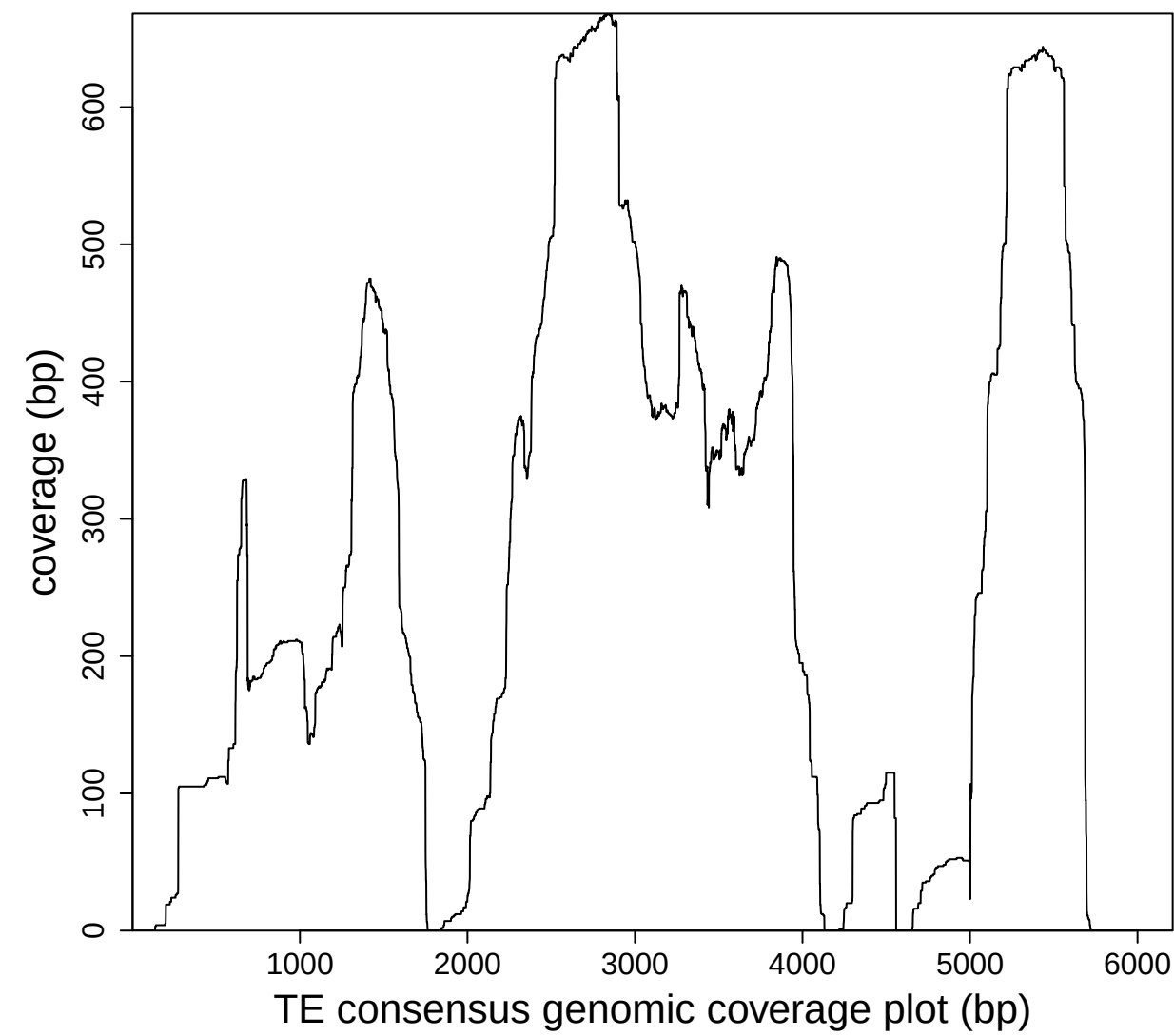
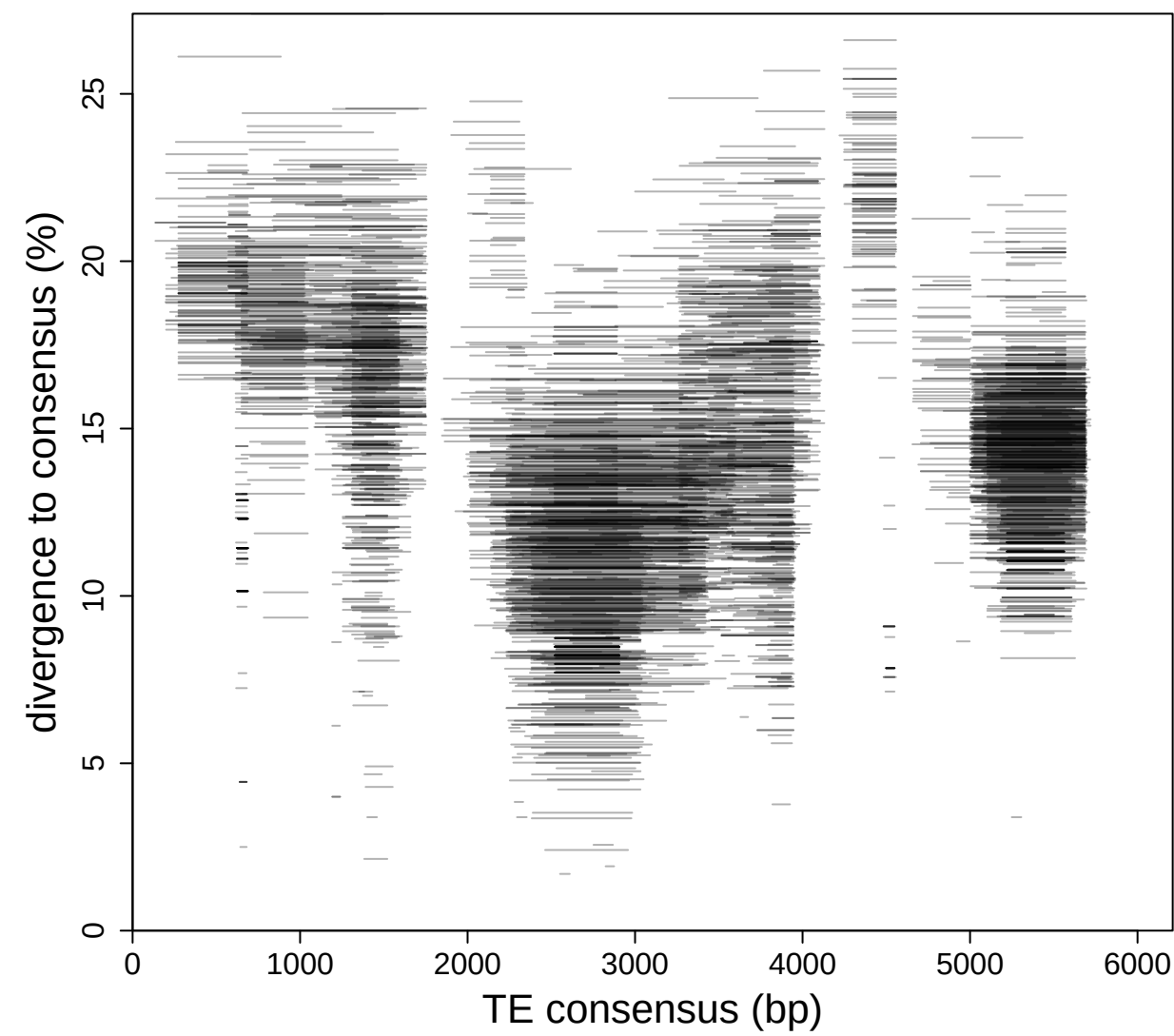






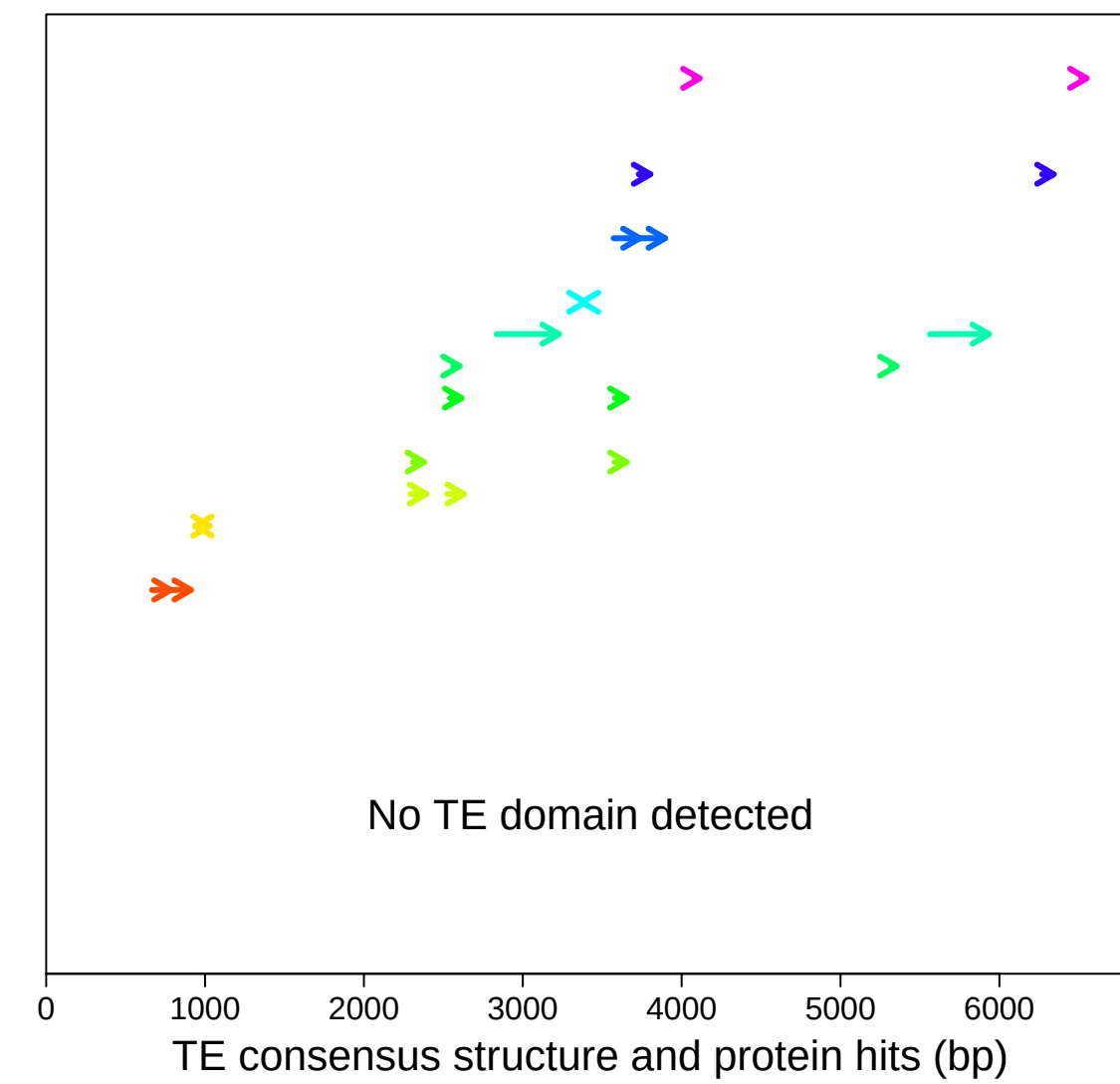
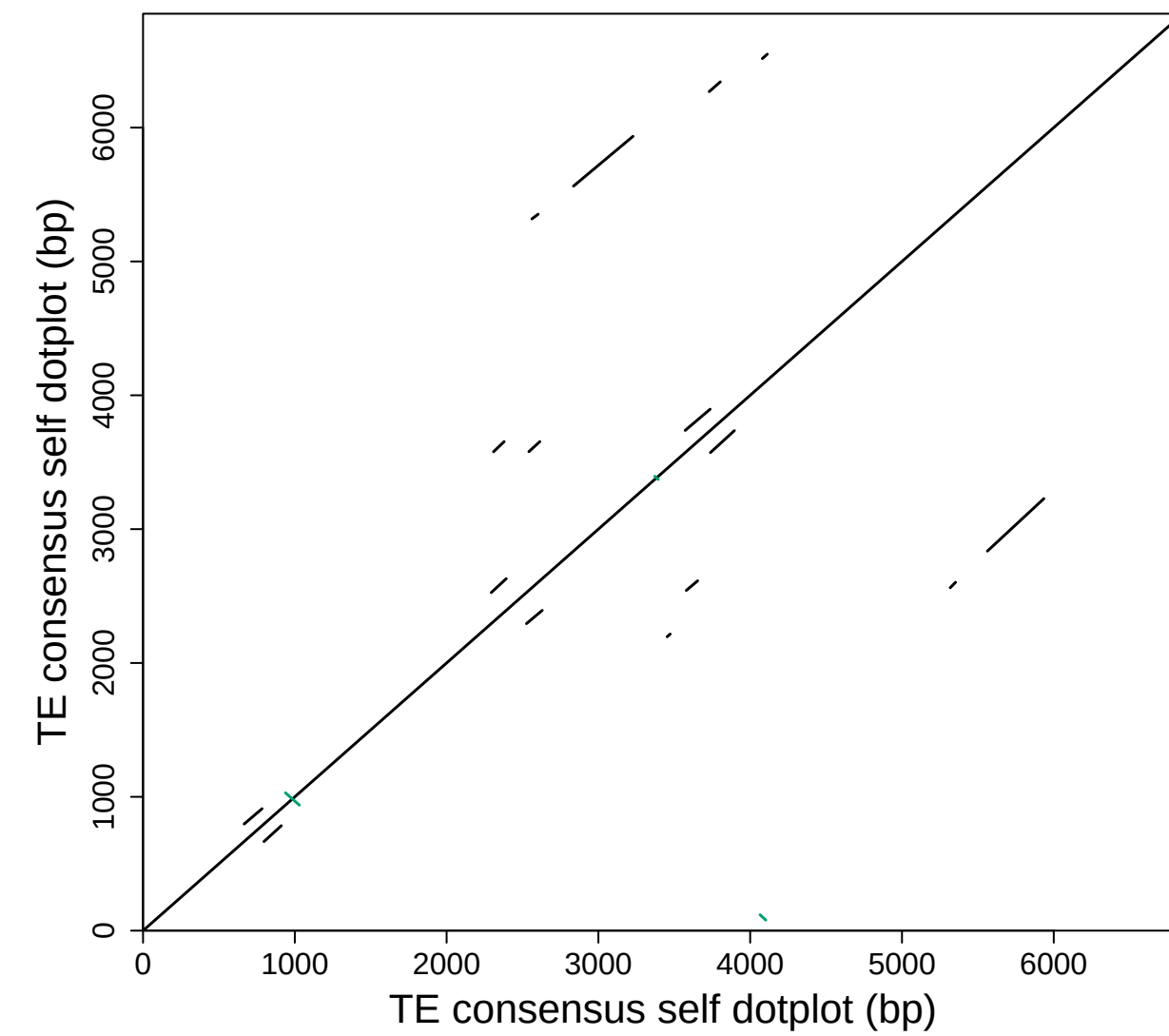
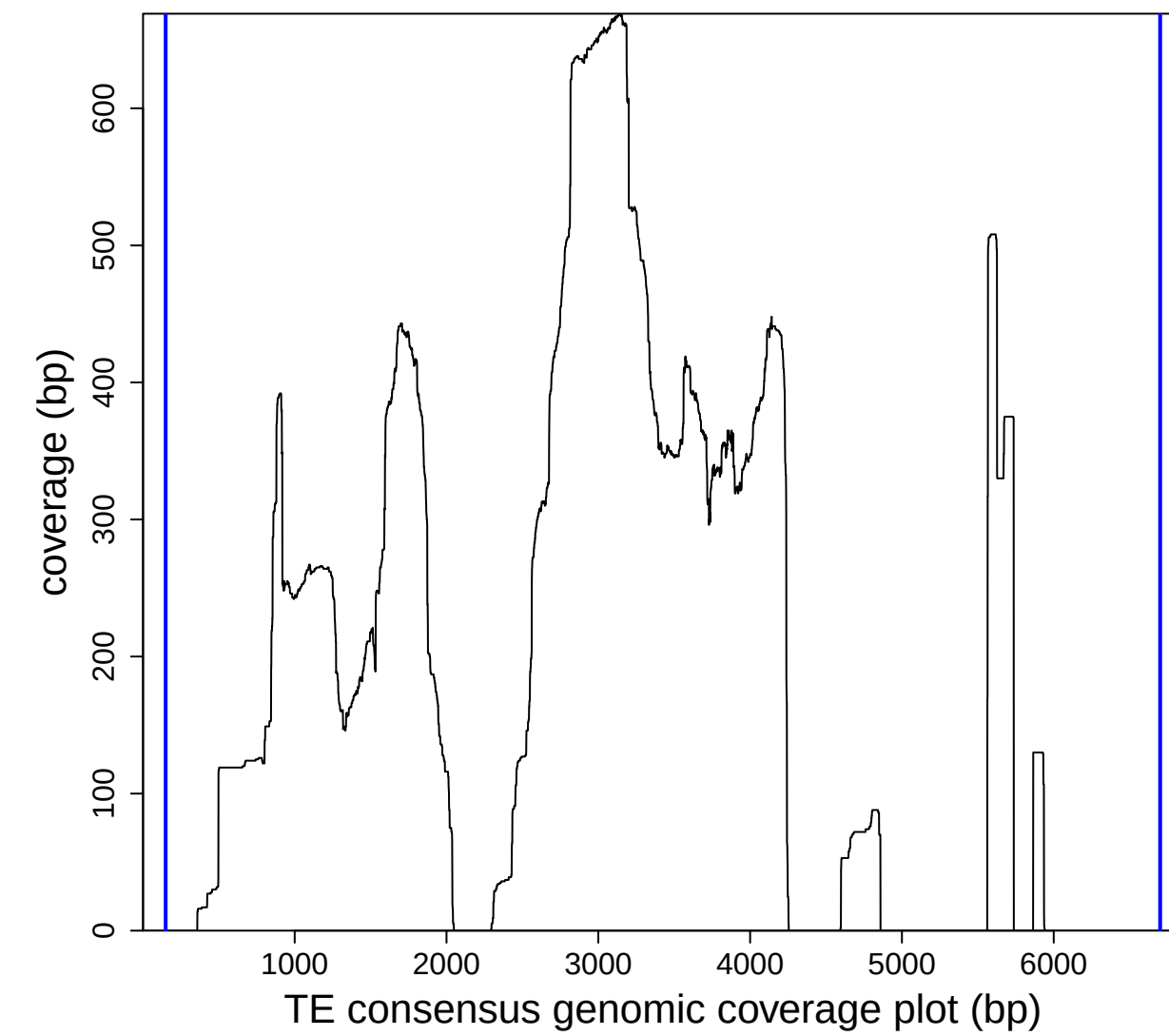
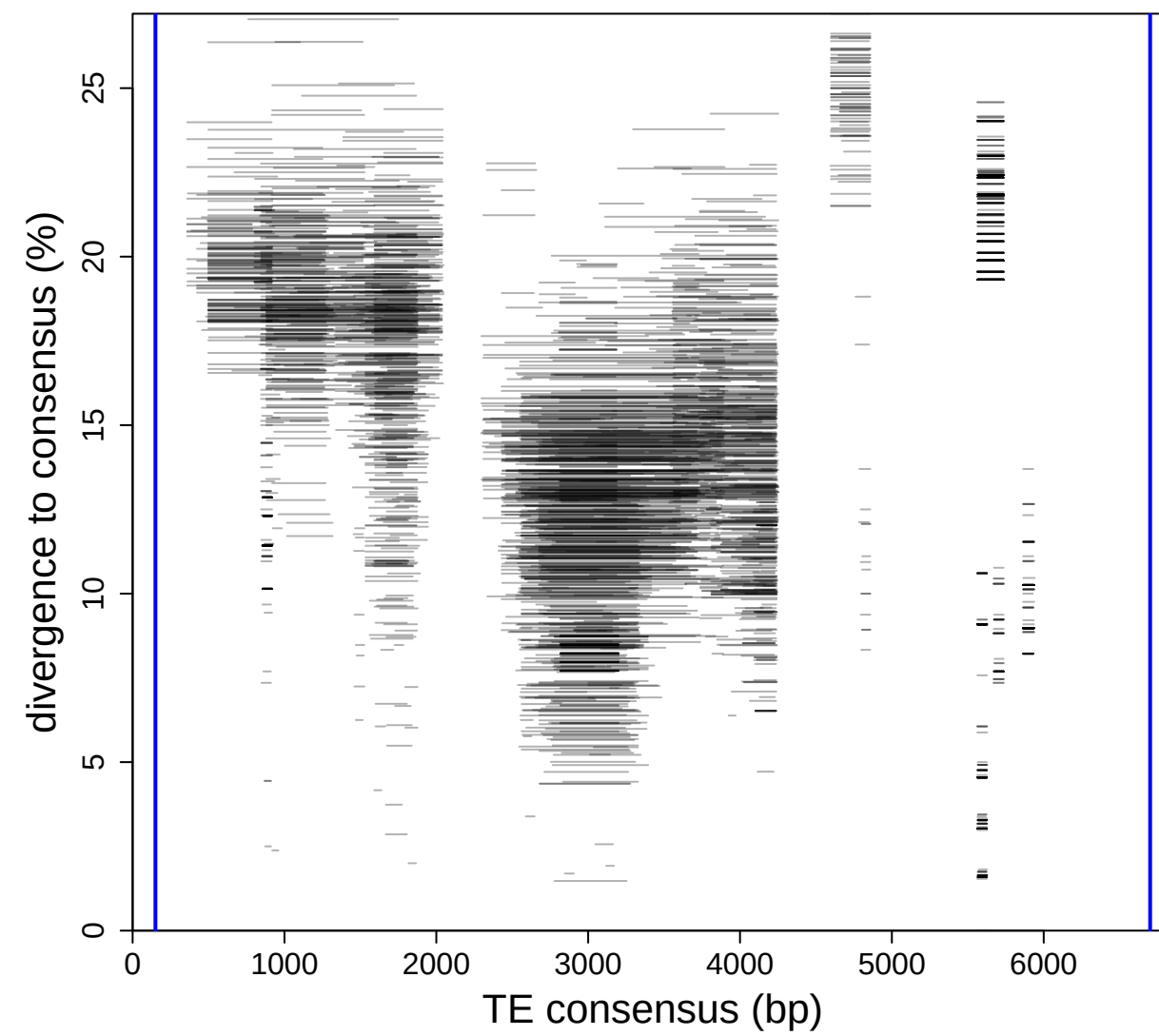
b.bed uf.bed g 2.bed fm 1.bed 0 0 bcln.fa aln.fa cl.fa gs.fa
size: 6210bp; fragments: 3030; full length: 0 (>=5589bp)

After TEtrimmer 6210 bp

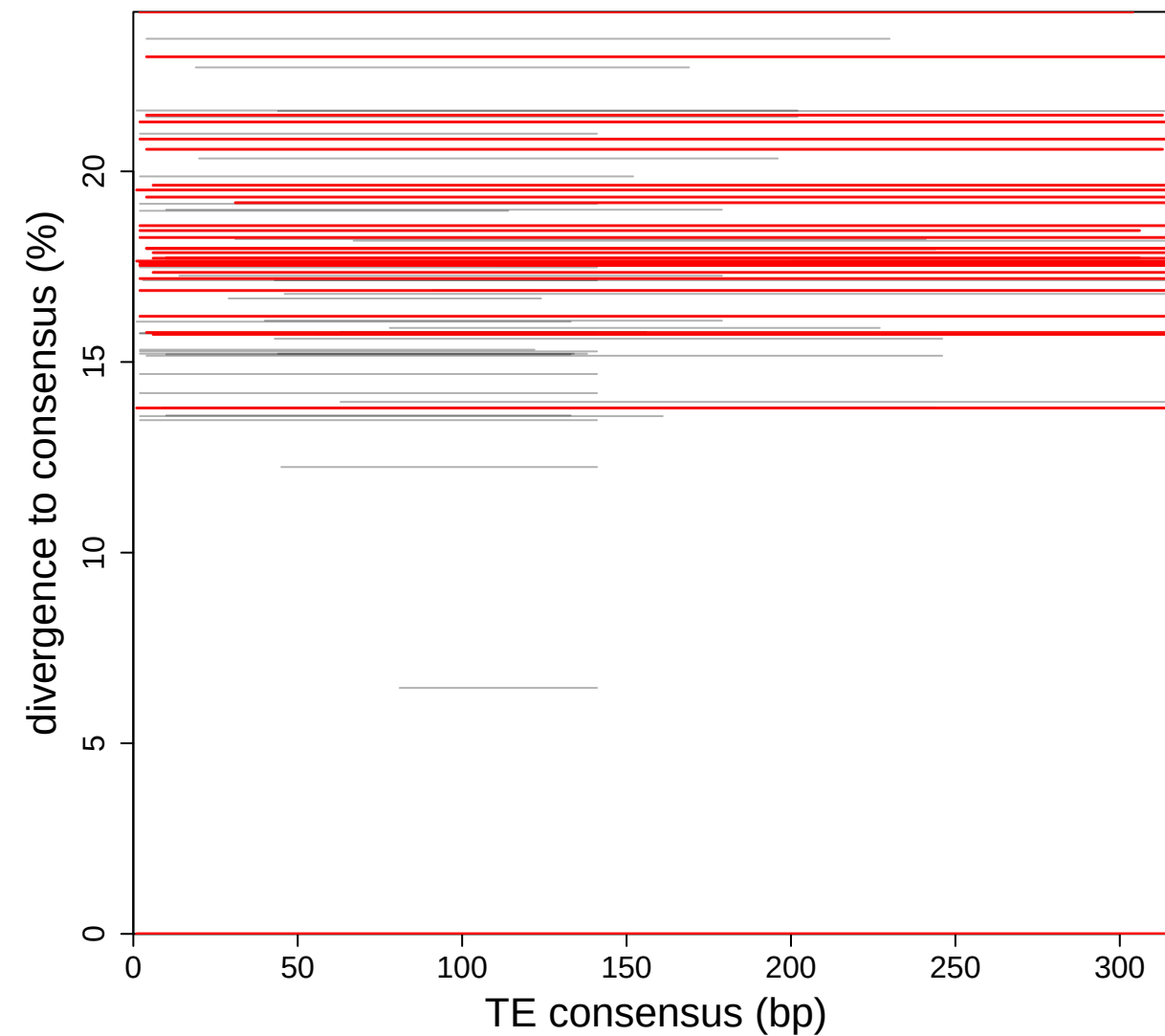


.fasta.b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa
size: 6851bp; fragments: 2879; full length: 0 (>=6165.9bp)

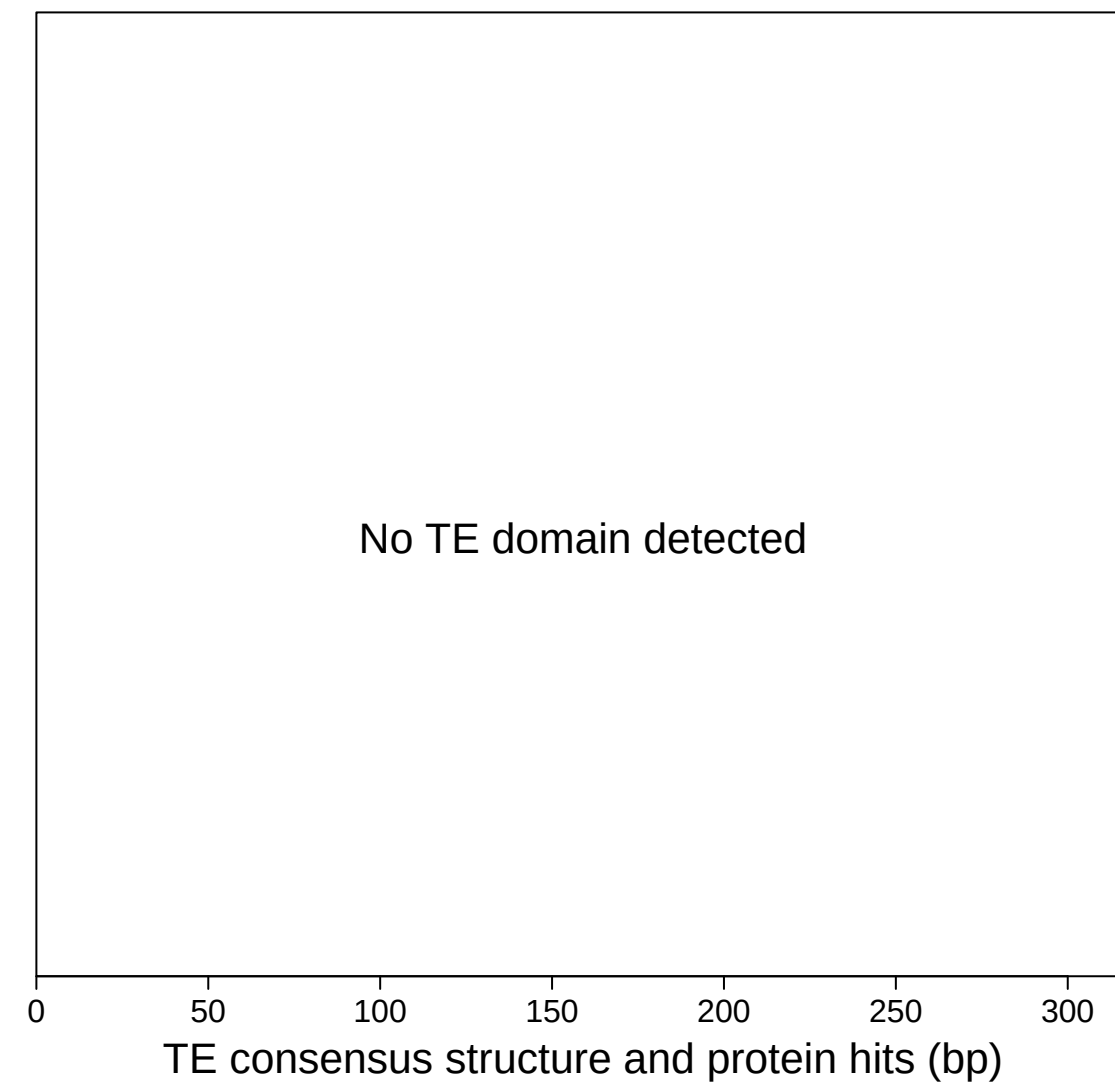
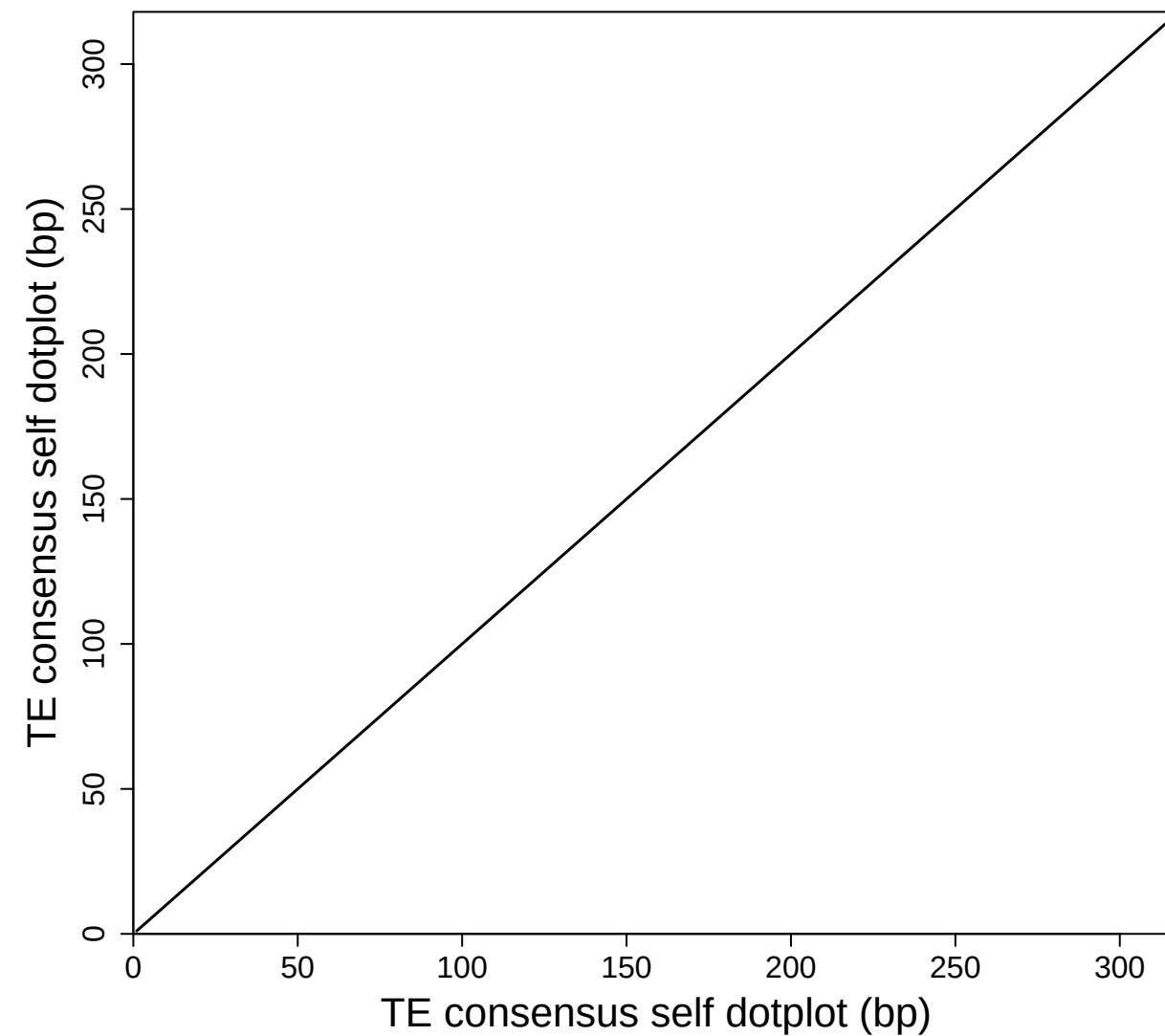
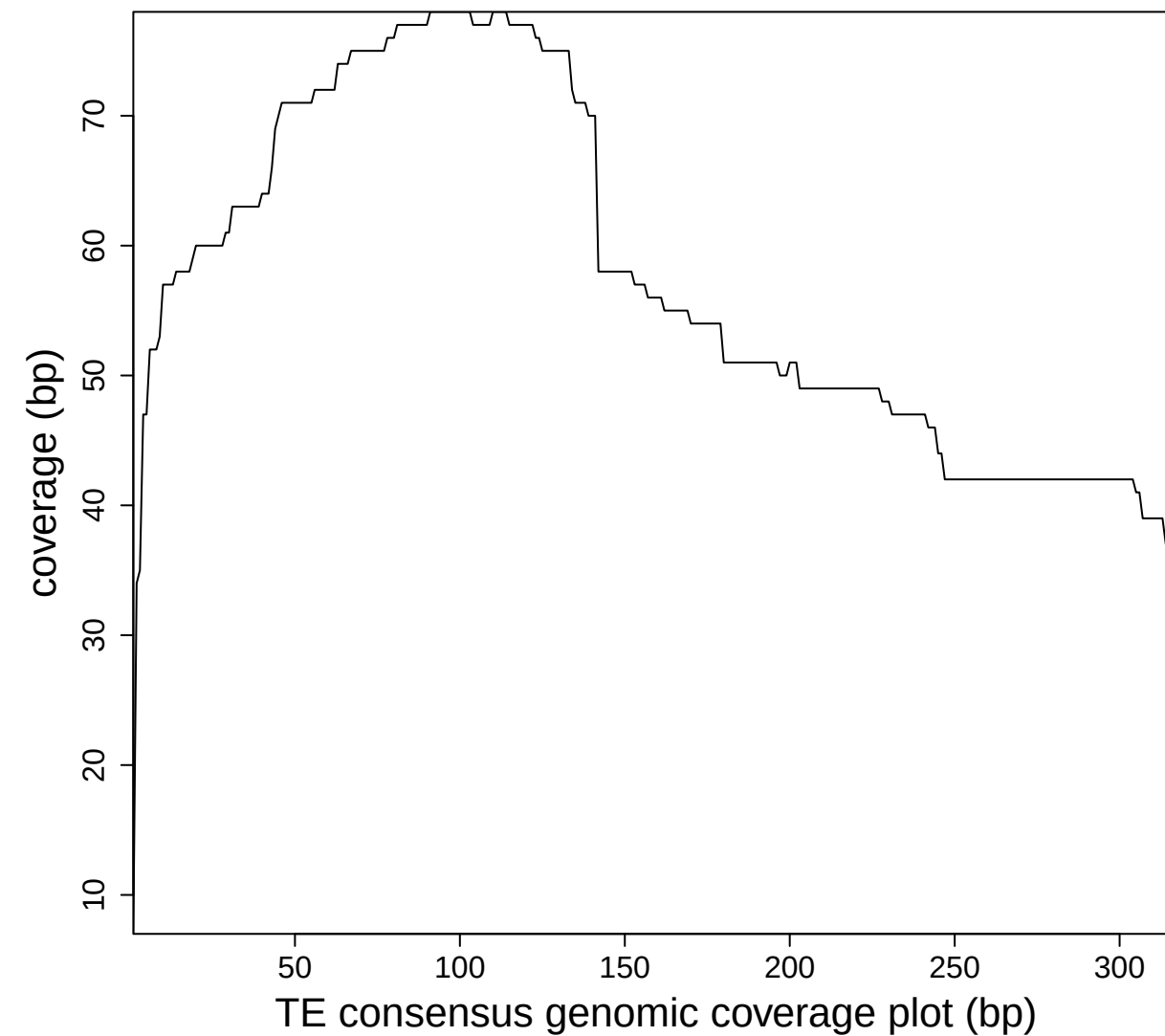
After TEtrimmer Extended plot Blue lines are boundaries



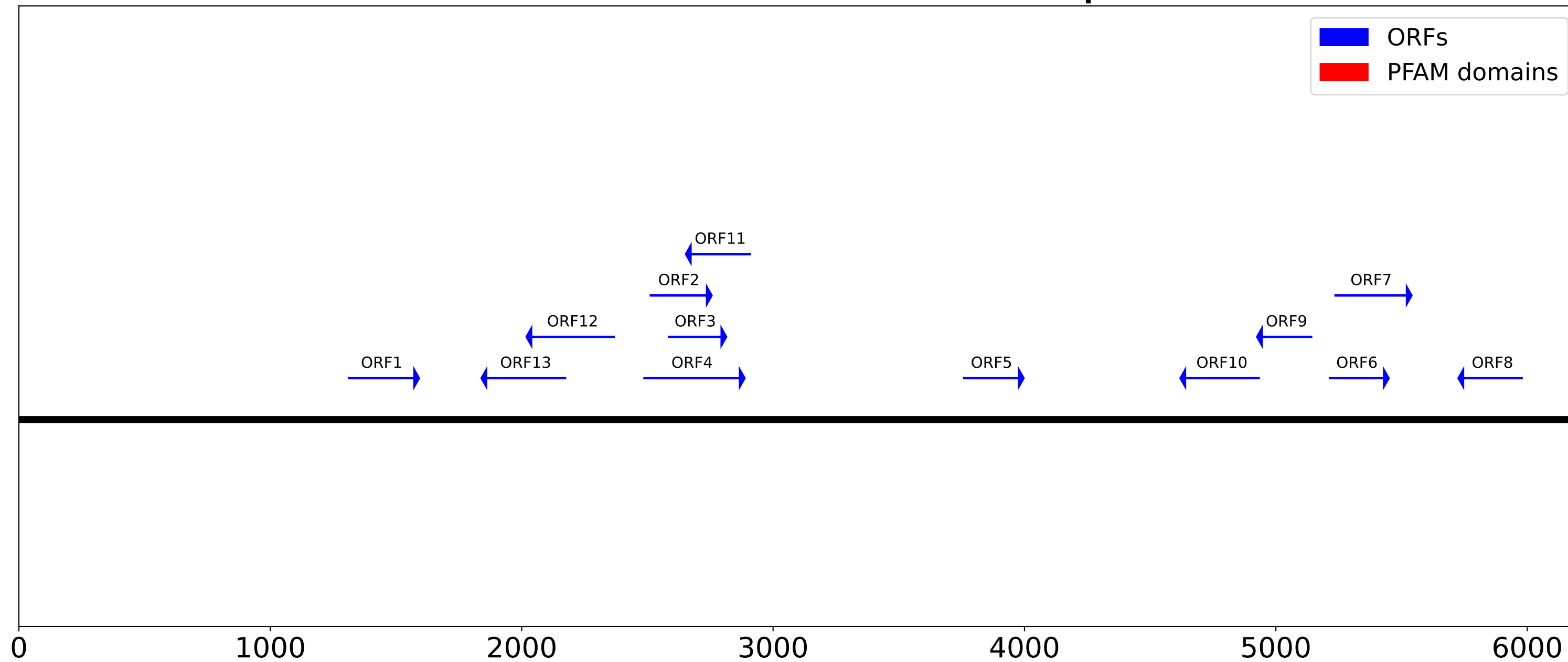
TE: ltr_1_family_226
size: 318bp; fragments: 80; full length: 32 (≥ 286.2 bp)



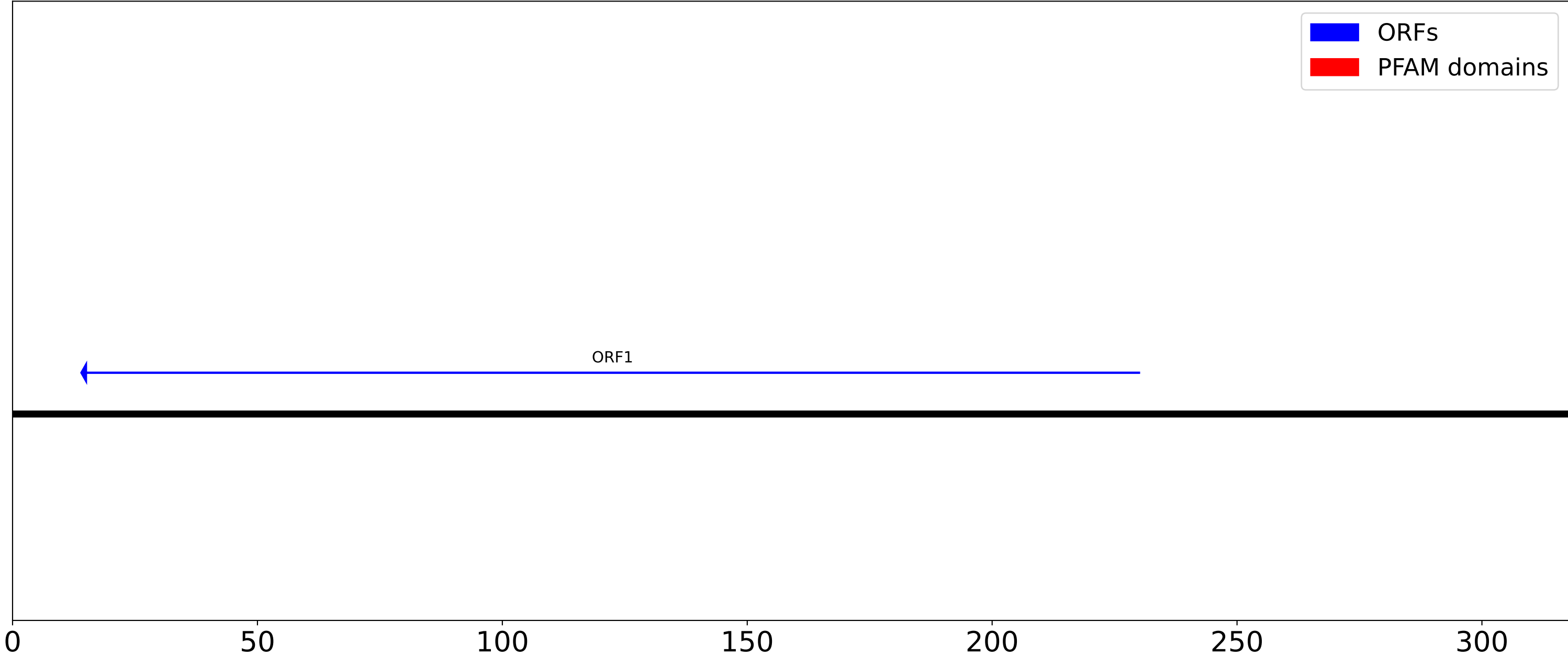
Before TEtrimmer 318 bp



After TEtrimmer ORF and PFAM domain plot



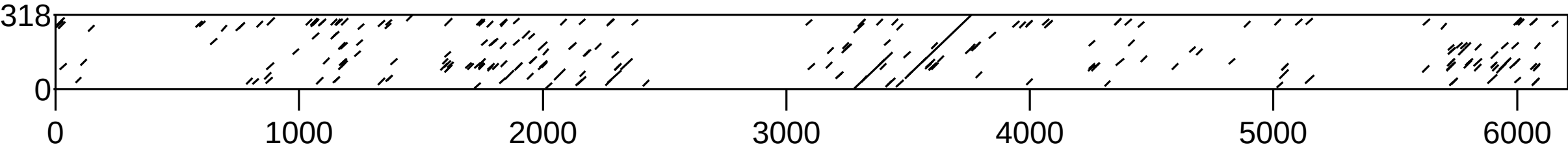
Before TEtrimmer ORF and PFAM domain plot



Dotplot

(windowsize = 25, threshold = 50.00 09/10/25)

consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)