

Start crop Point



End crop Point



MSA length = 6885

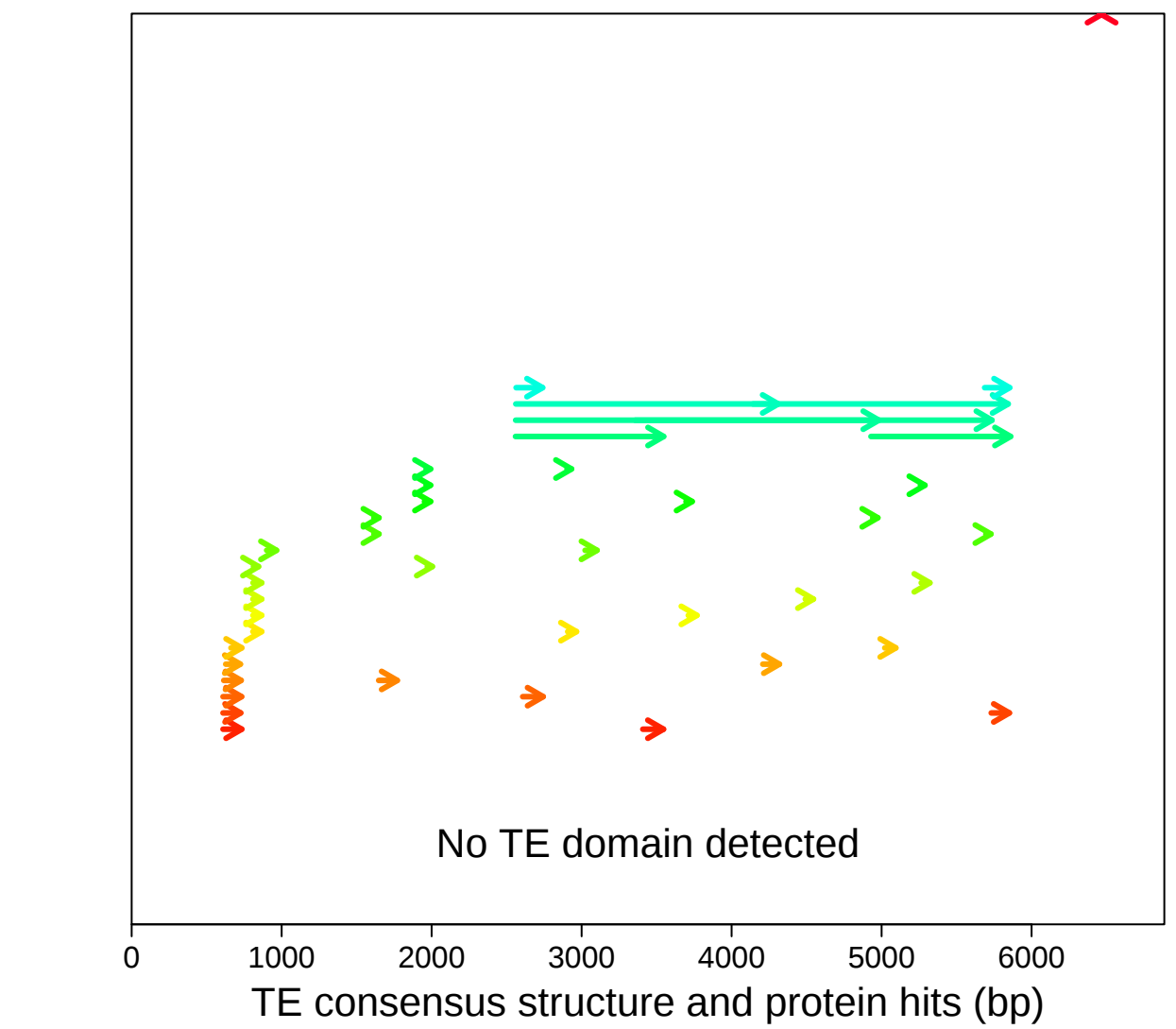
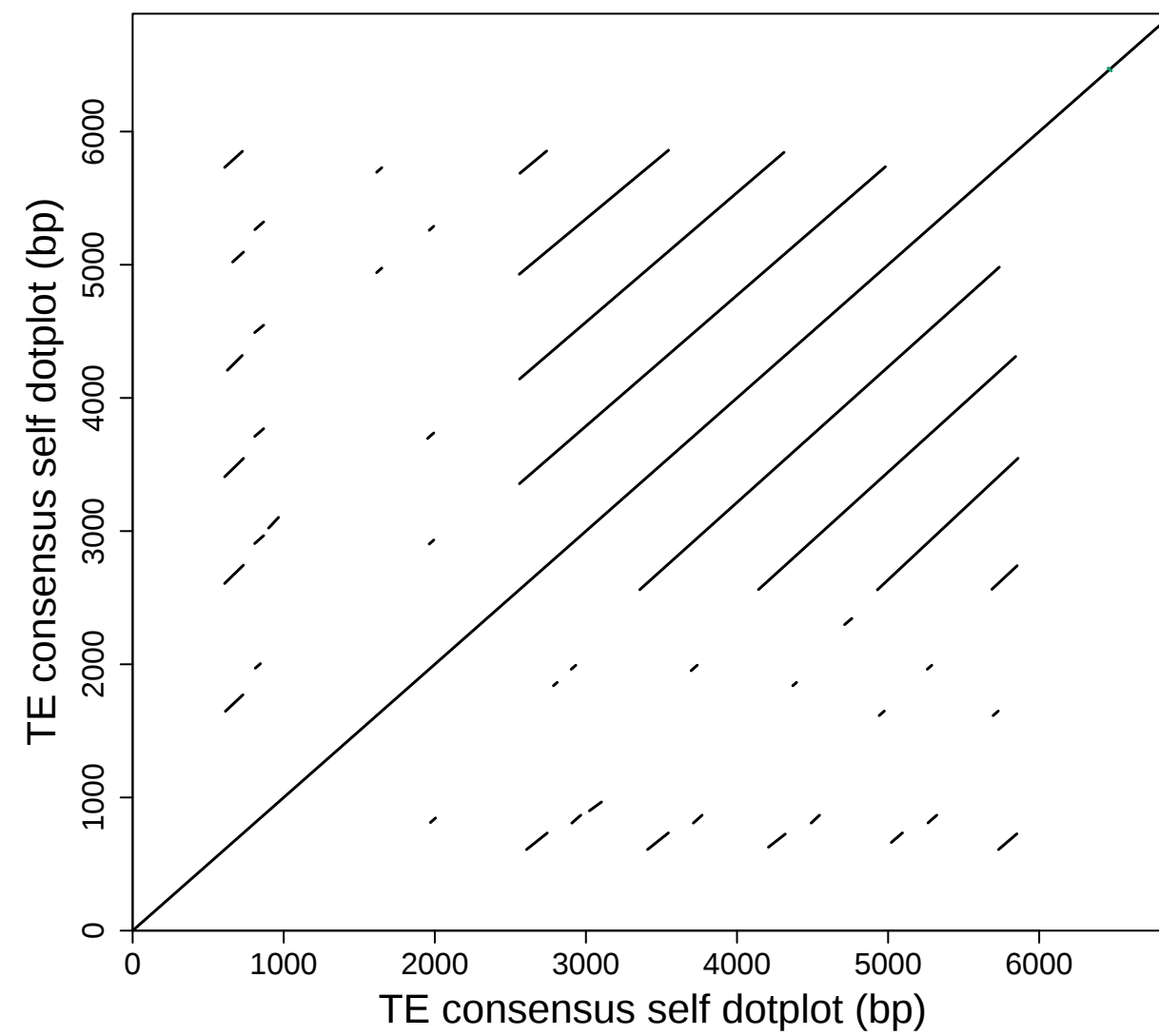
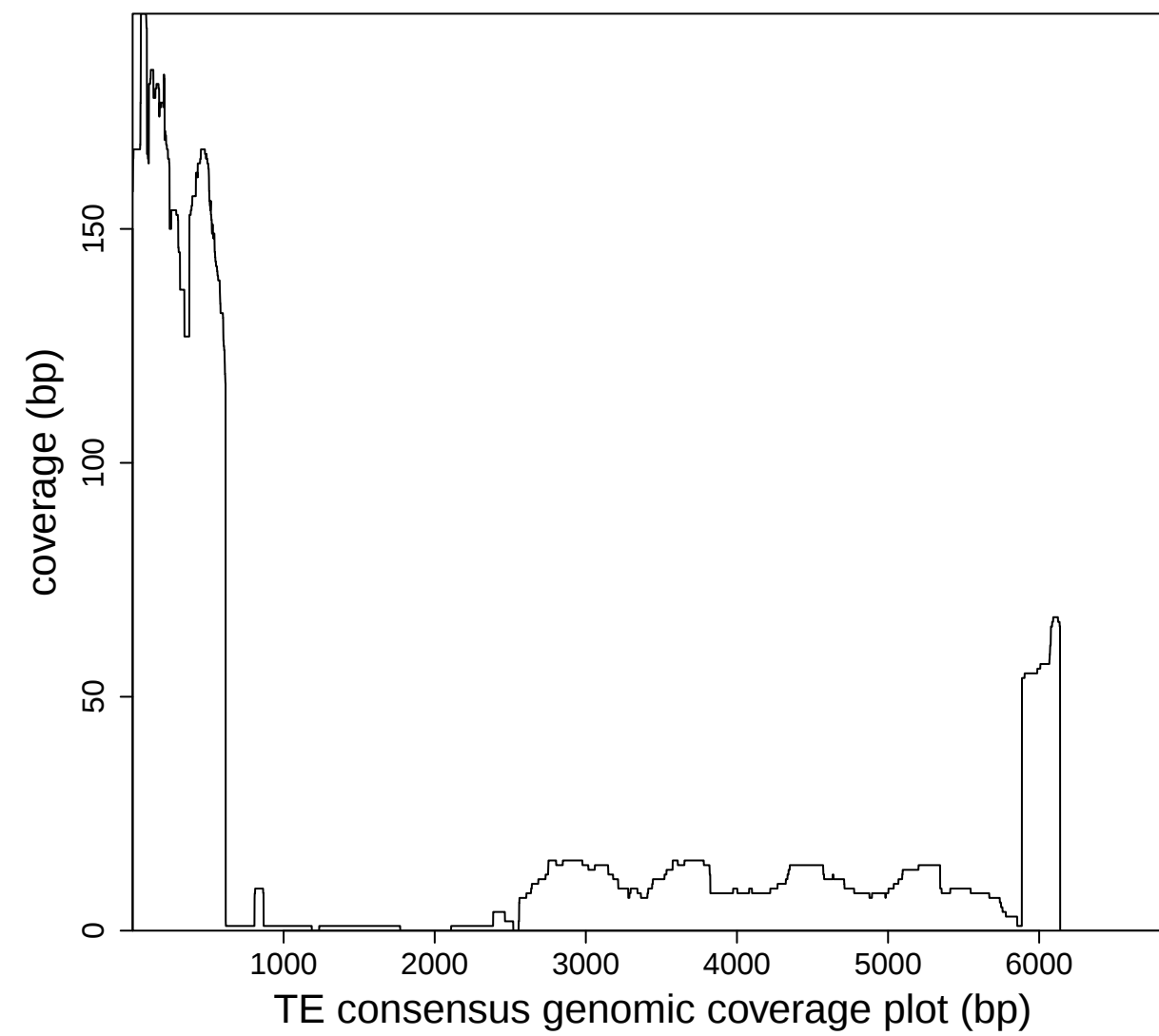
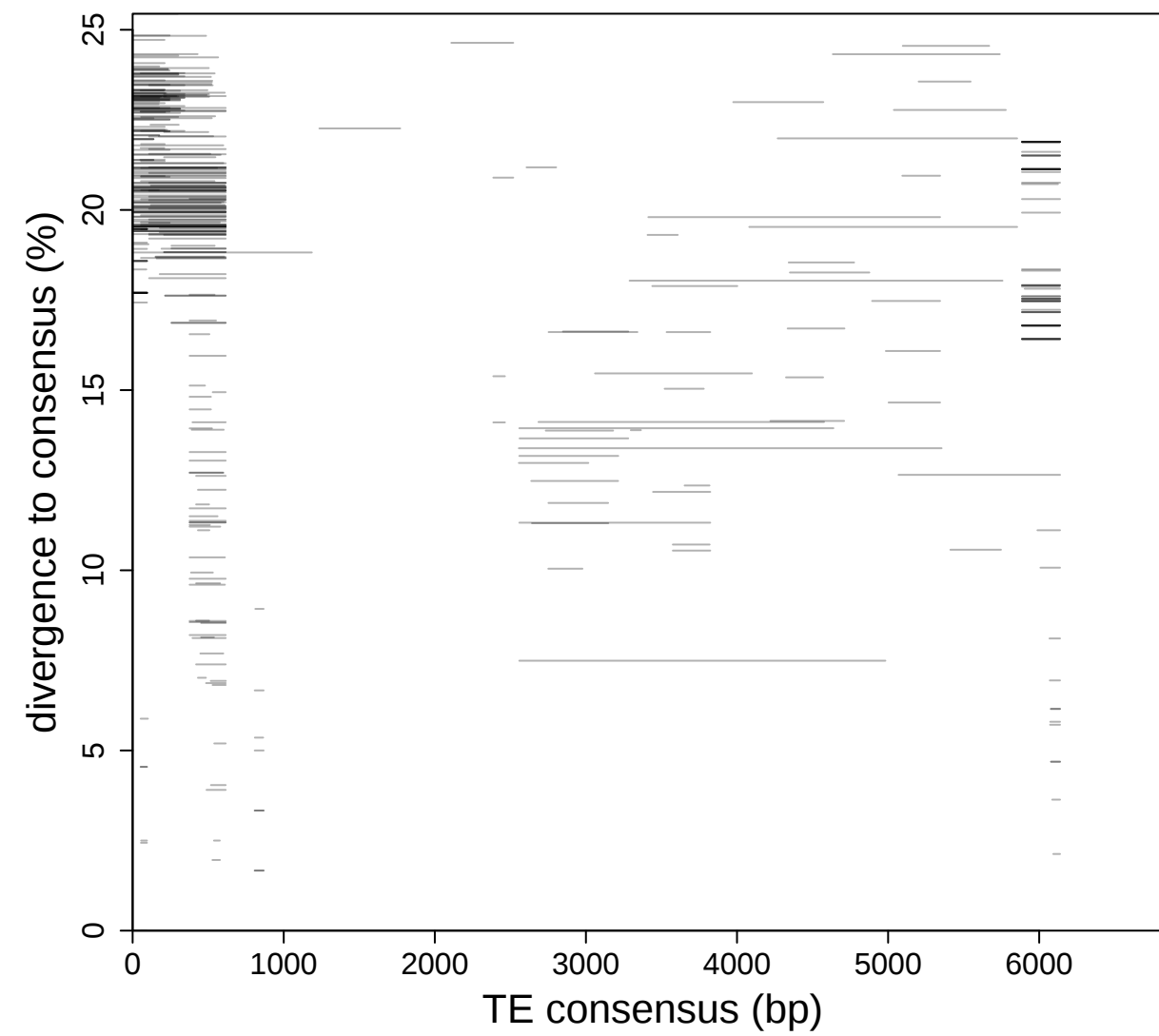


10 -



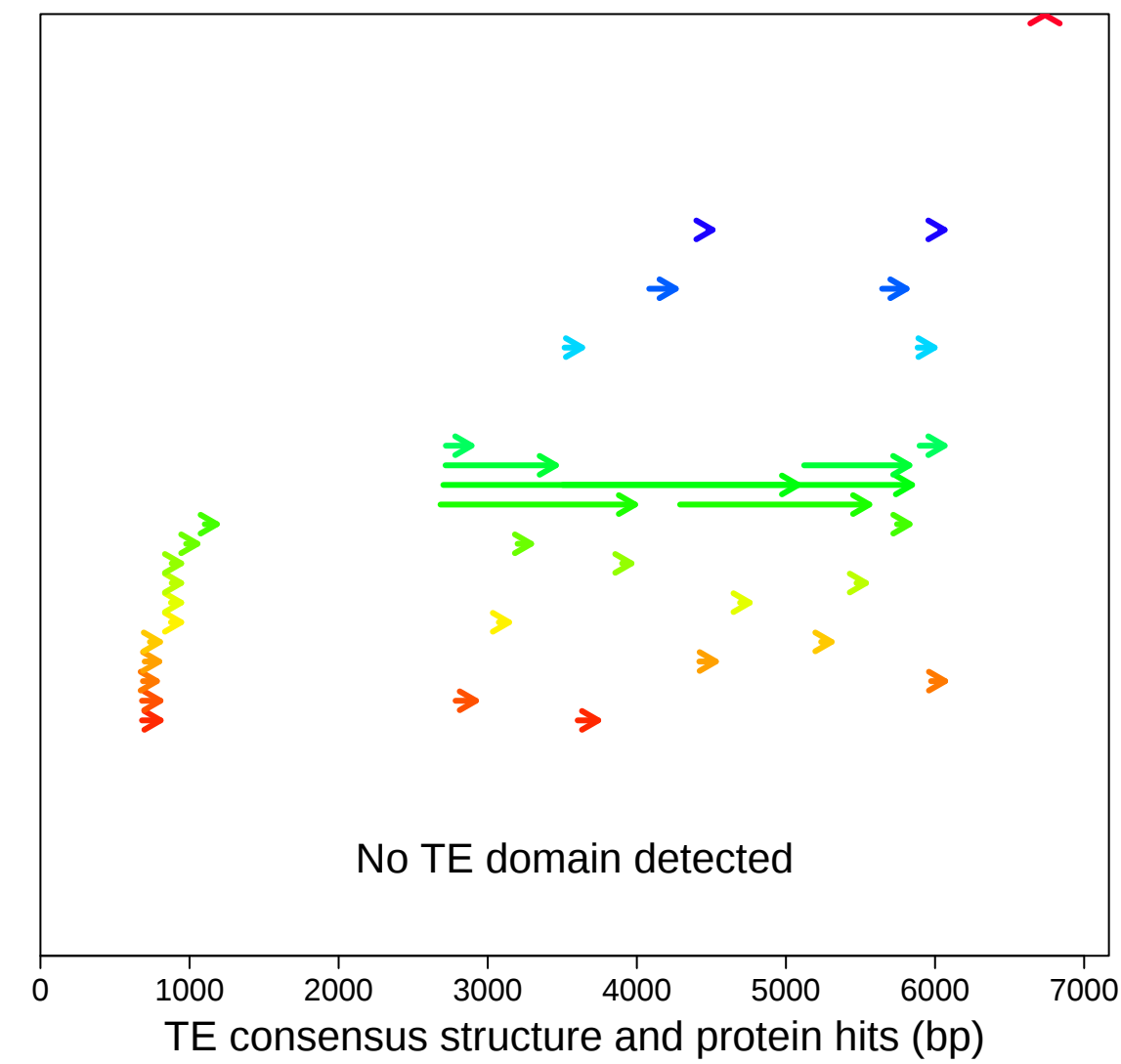
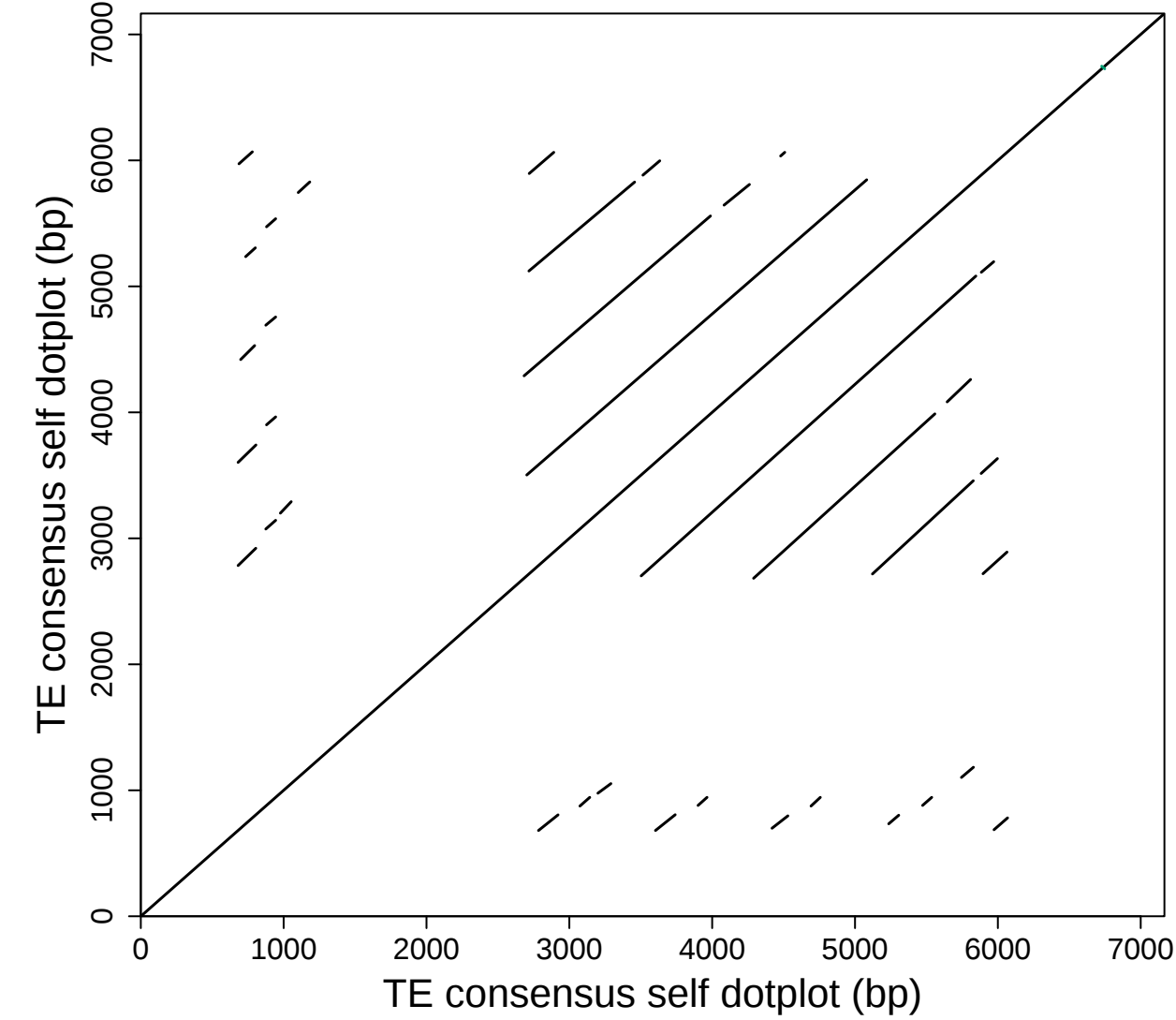
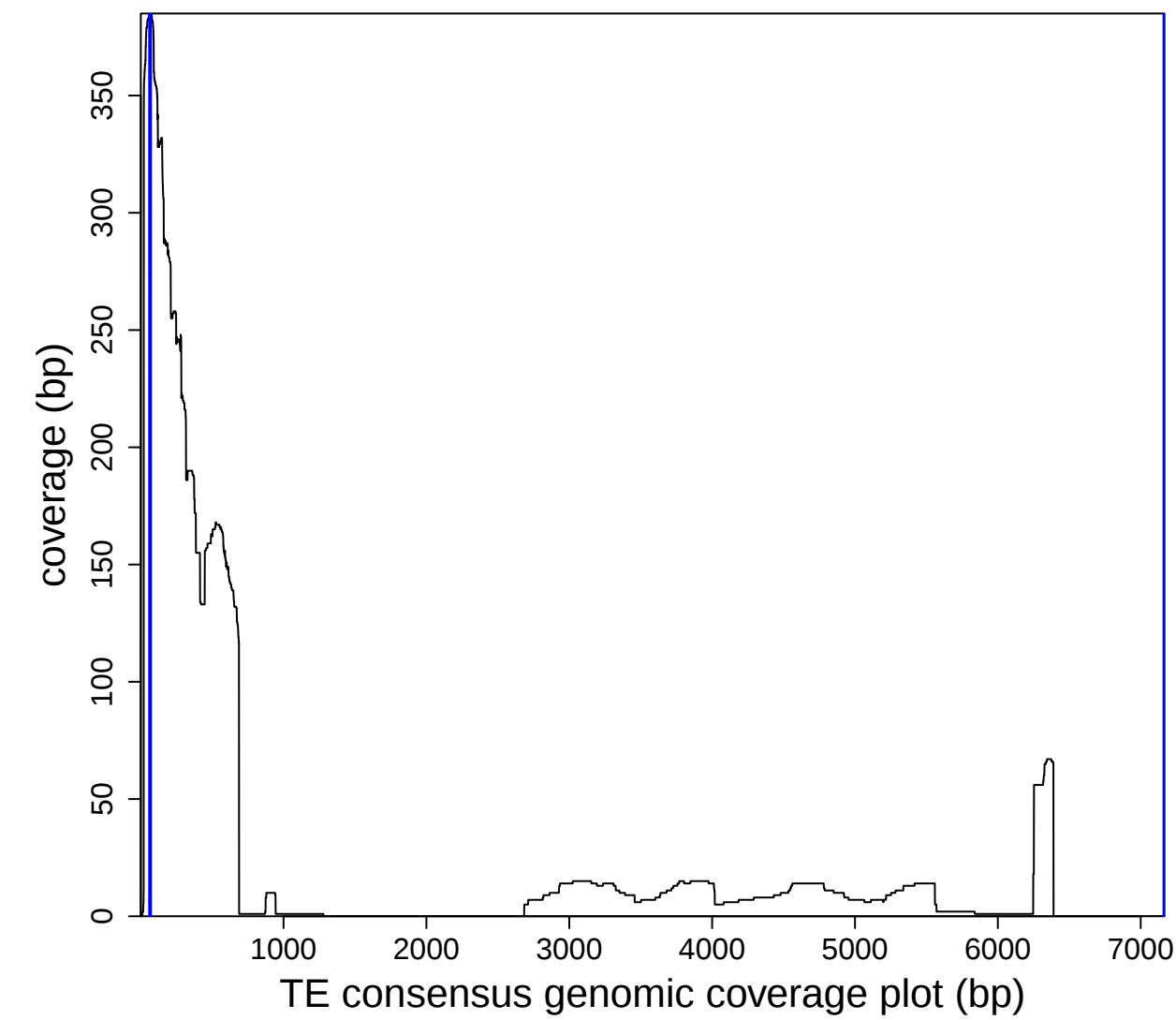
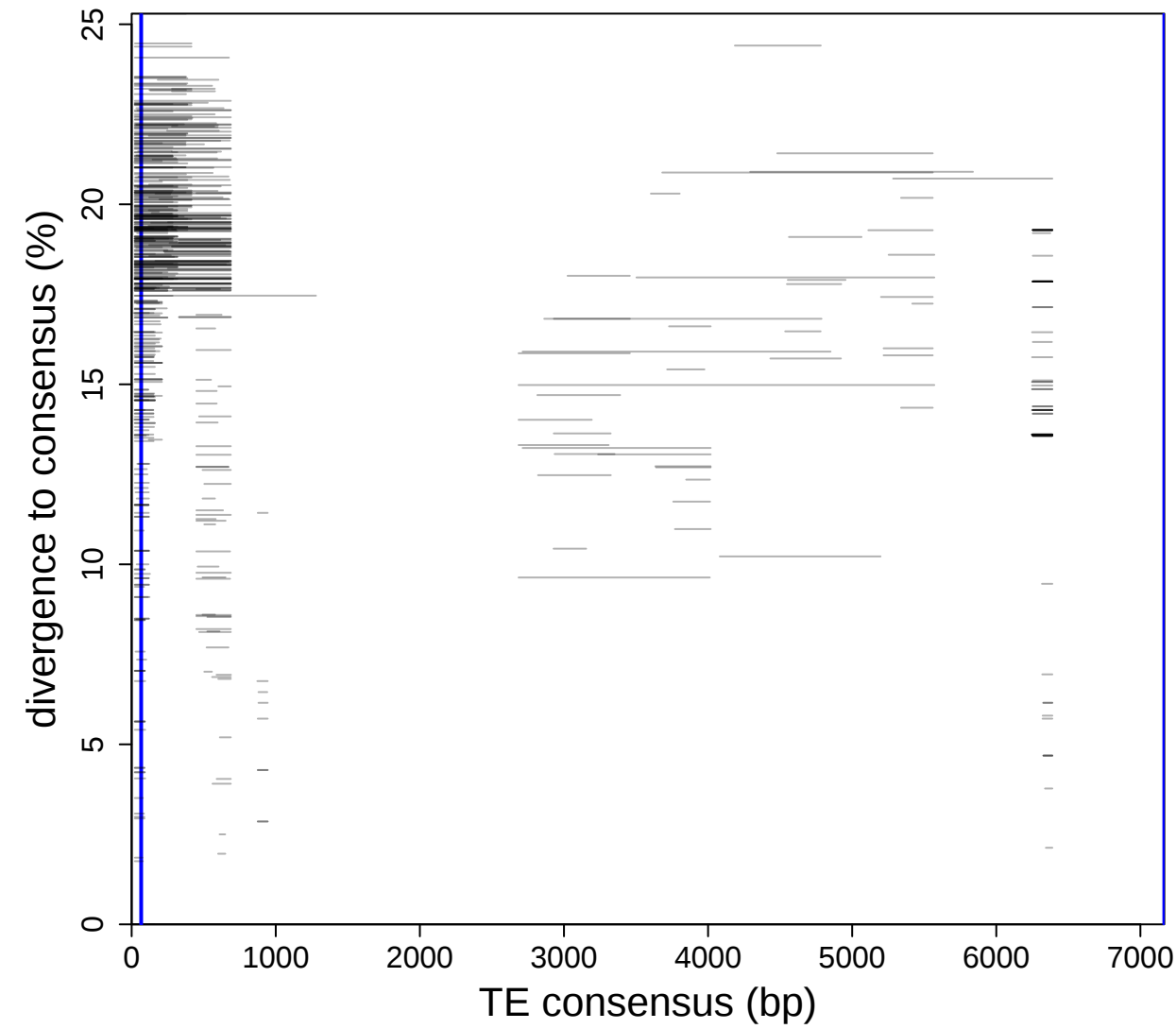
ta.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.f
size: 6885bp; fragments: 412; full length: 0 (>=6196.5bp)

After TEtrimmer 6885 bp

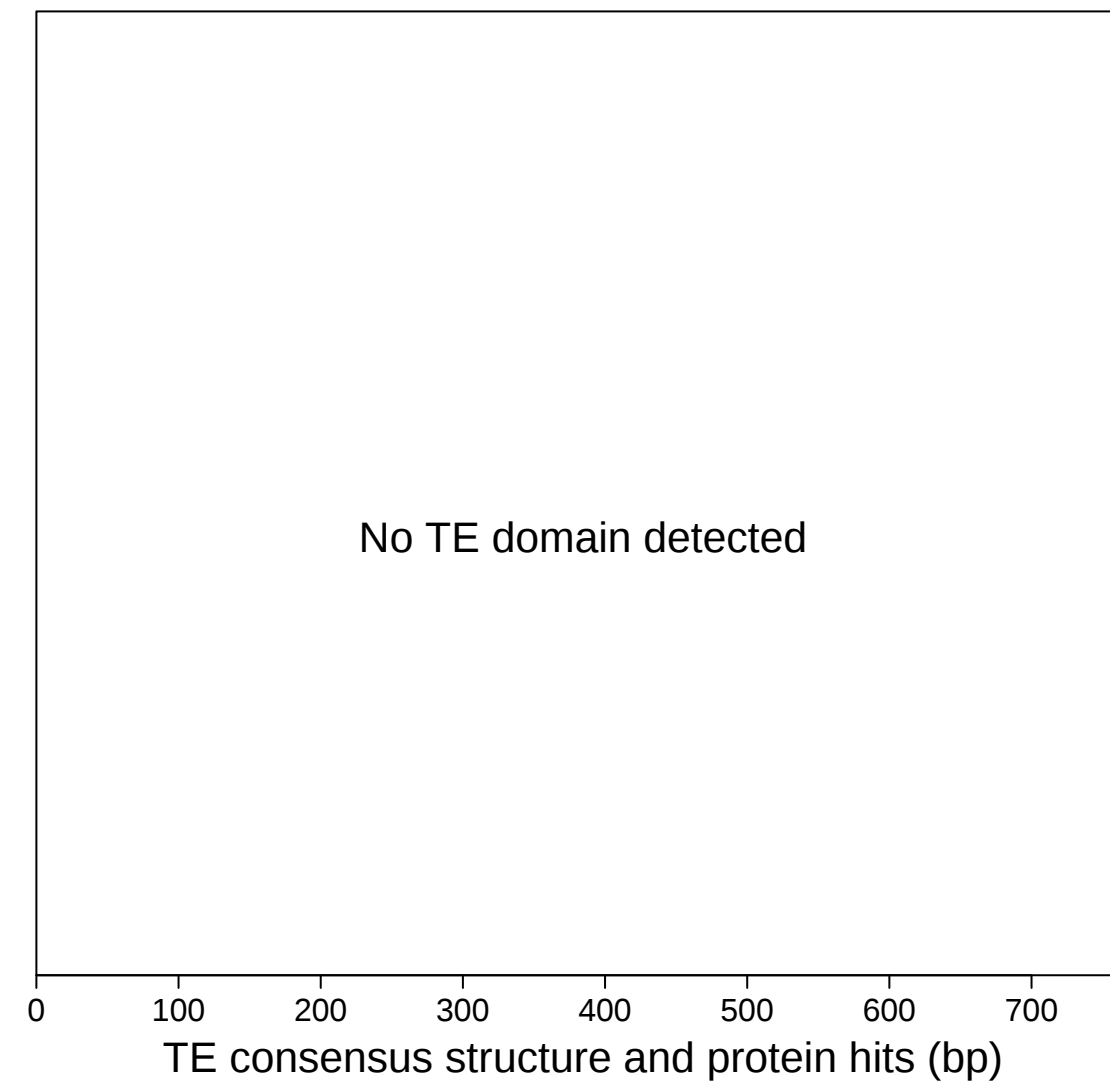
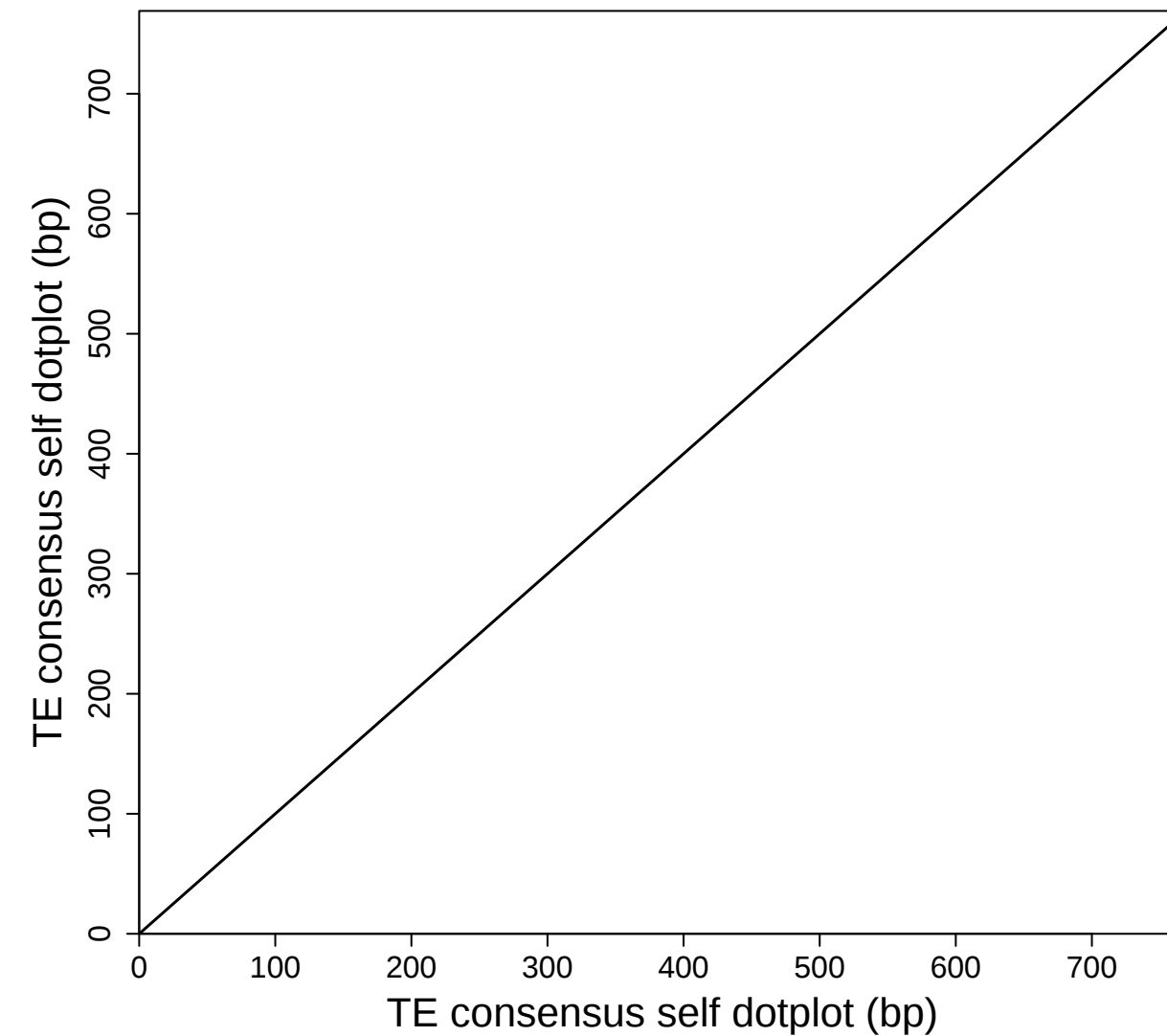
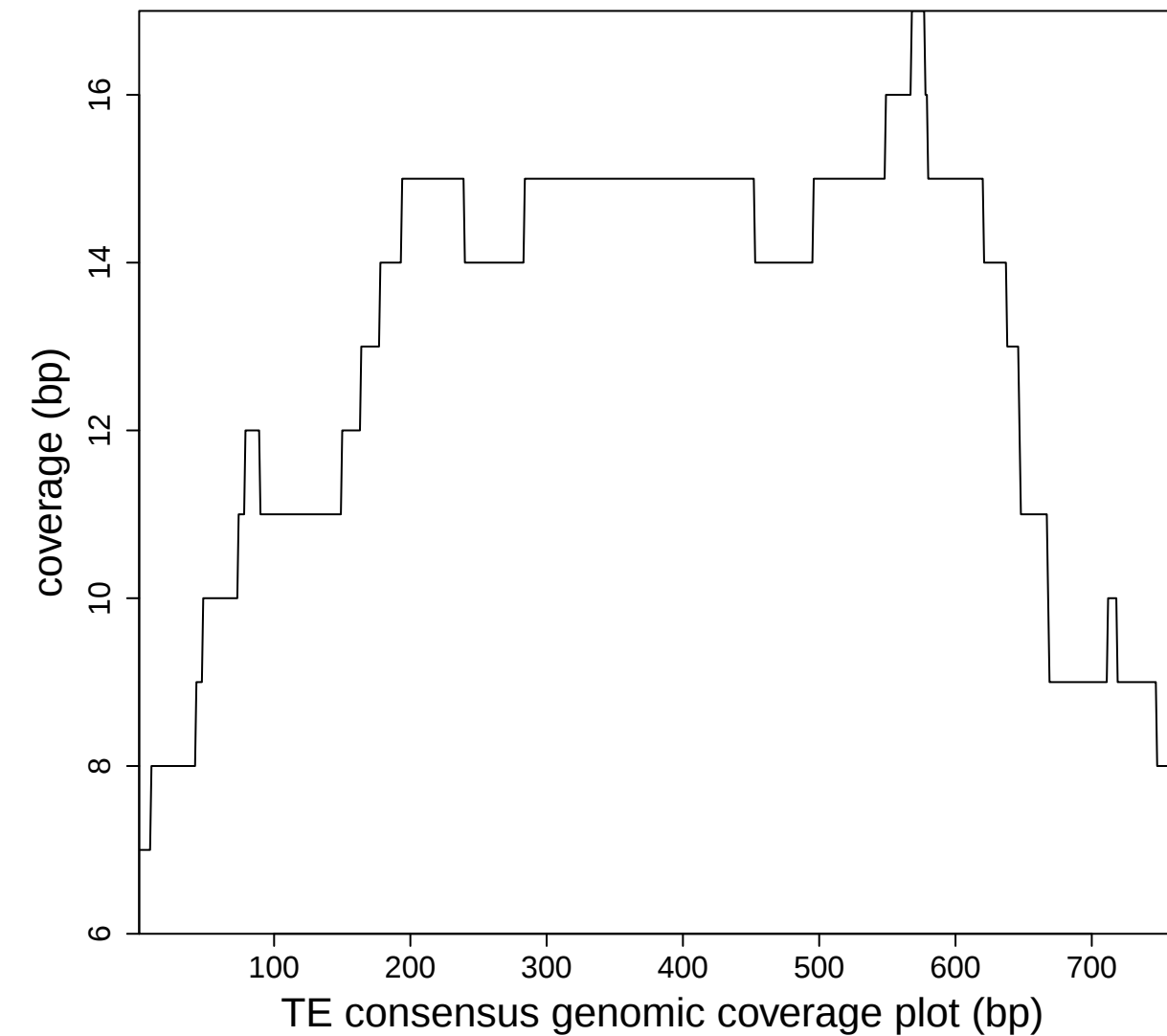
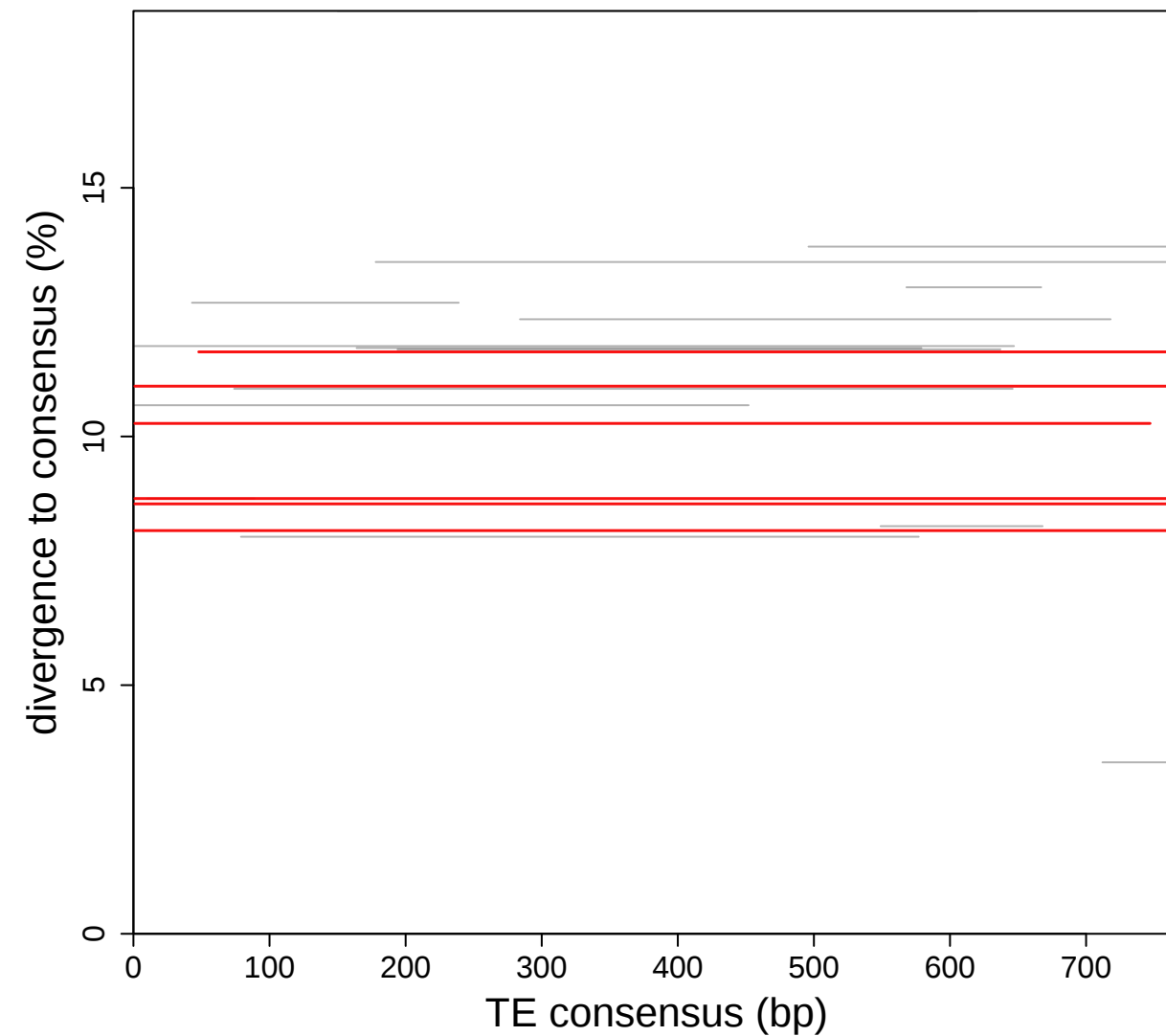


68.fasta.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.1
size: 7166bp; fragments: 589; full length: 0 (>=6449.4bp)

After TEtrimmer Extended plot Blue lines are boundaries

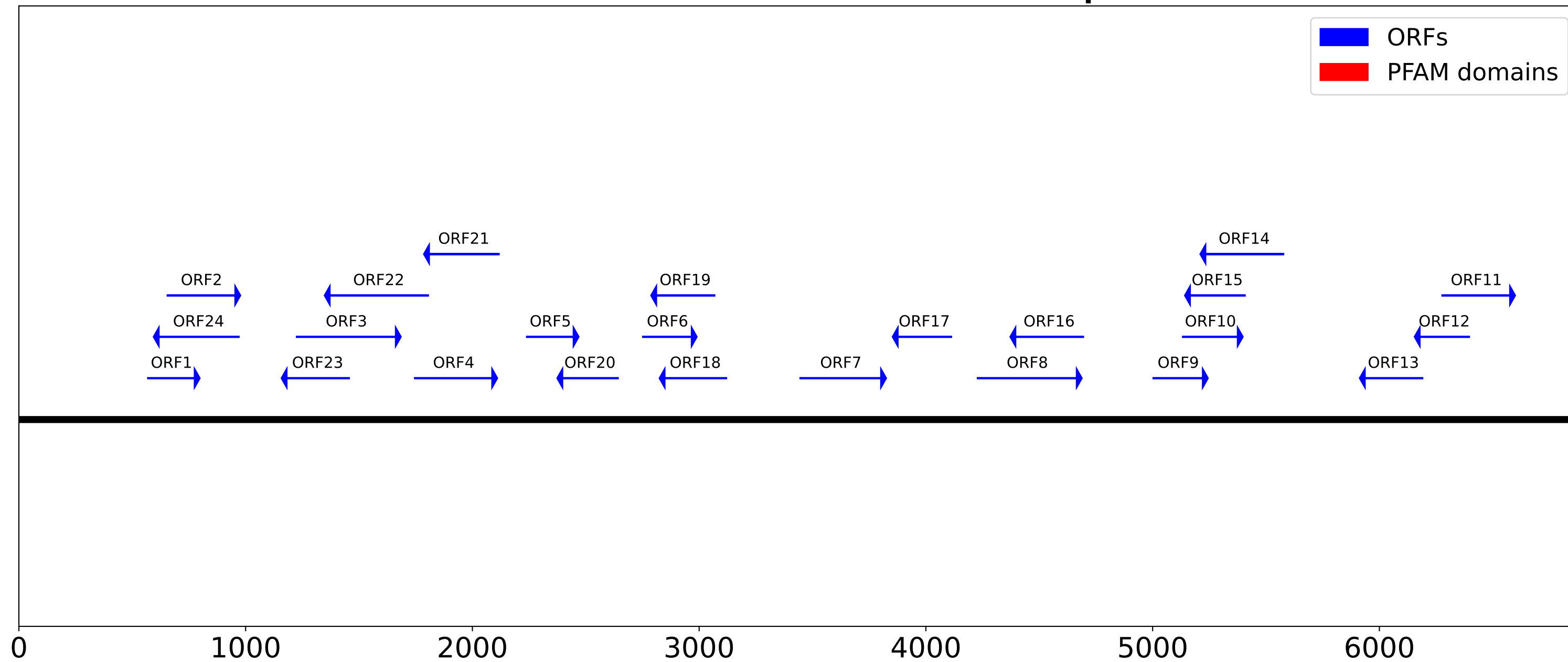


TE: rnd_5_family_27368
size: 769bp; fragments: 21; full length: 6 (≥ 692.1 bp)

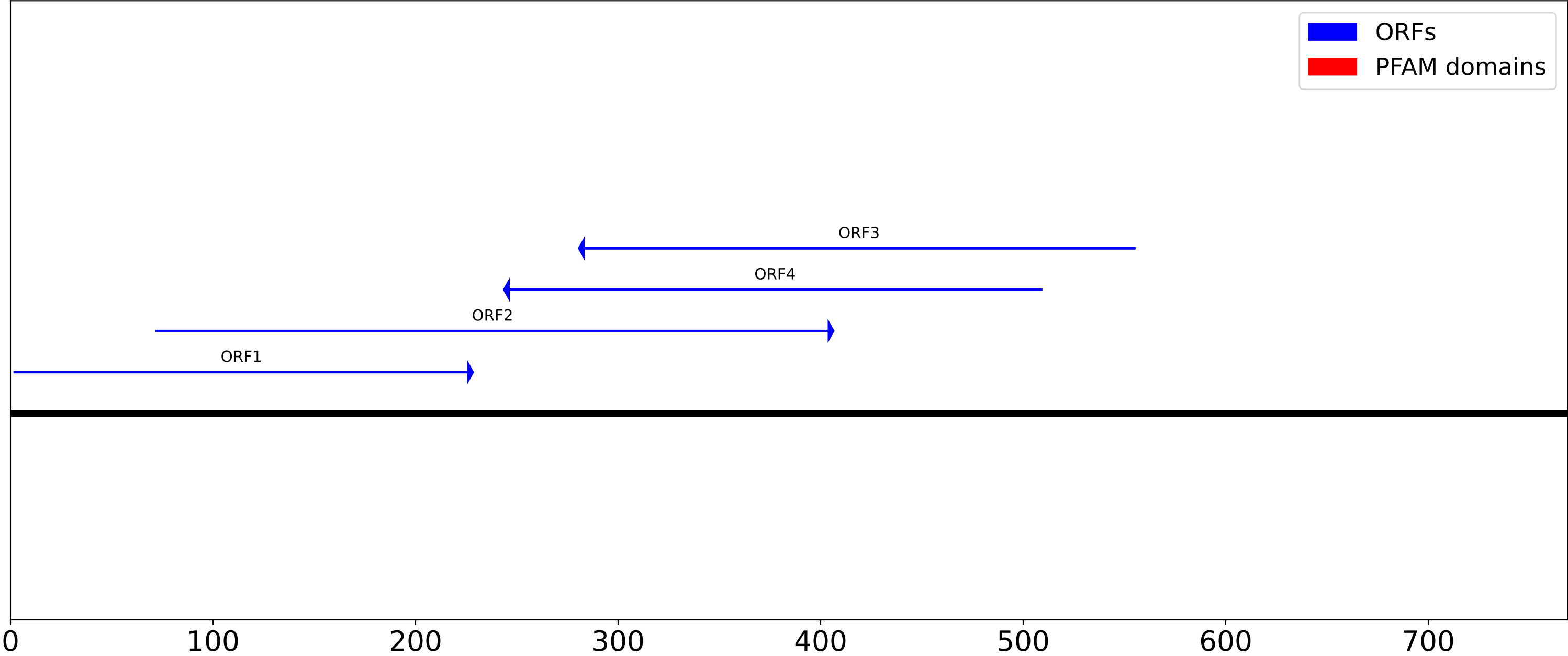


Before TEtrimmer 769 bp

After TEtrimmer ORF and PFAM domain plot



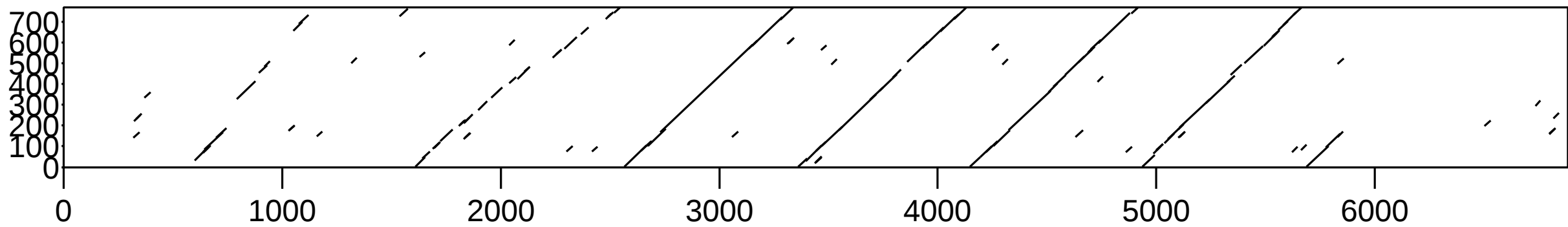
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)