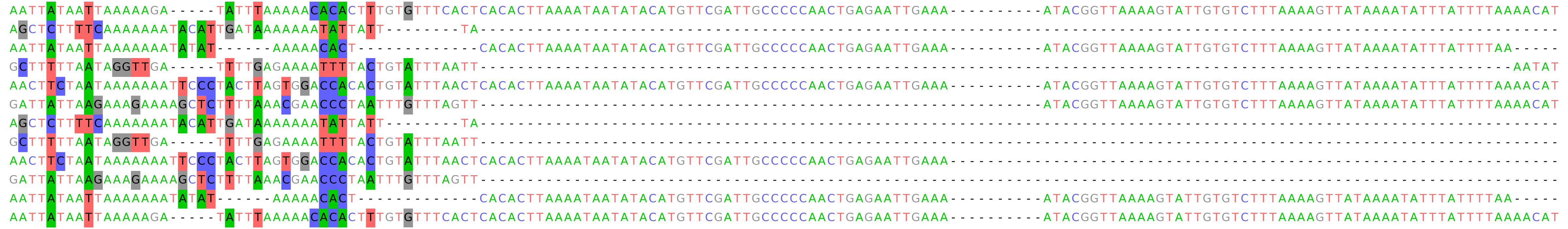
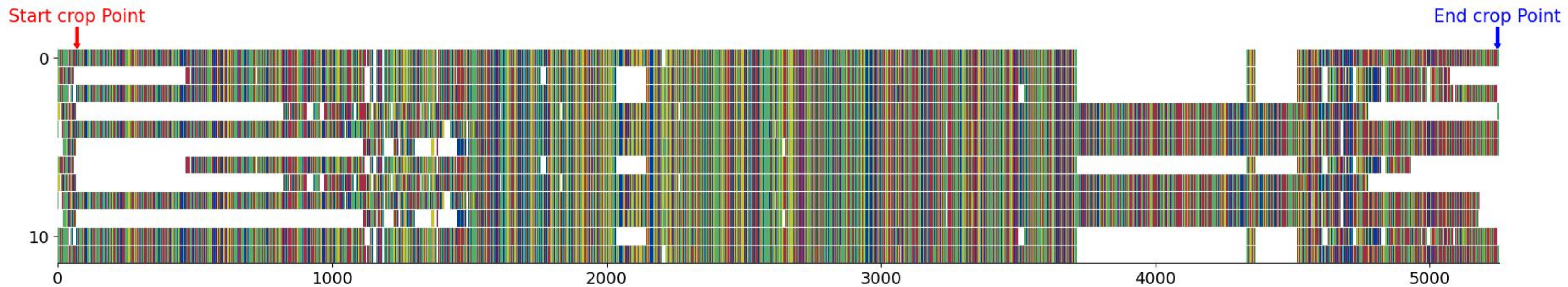


Start crop Point

End crop Point

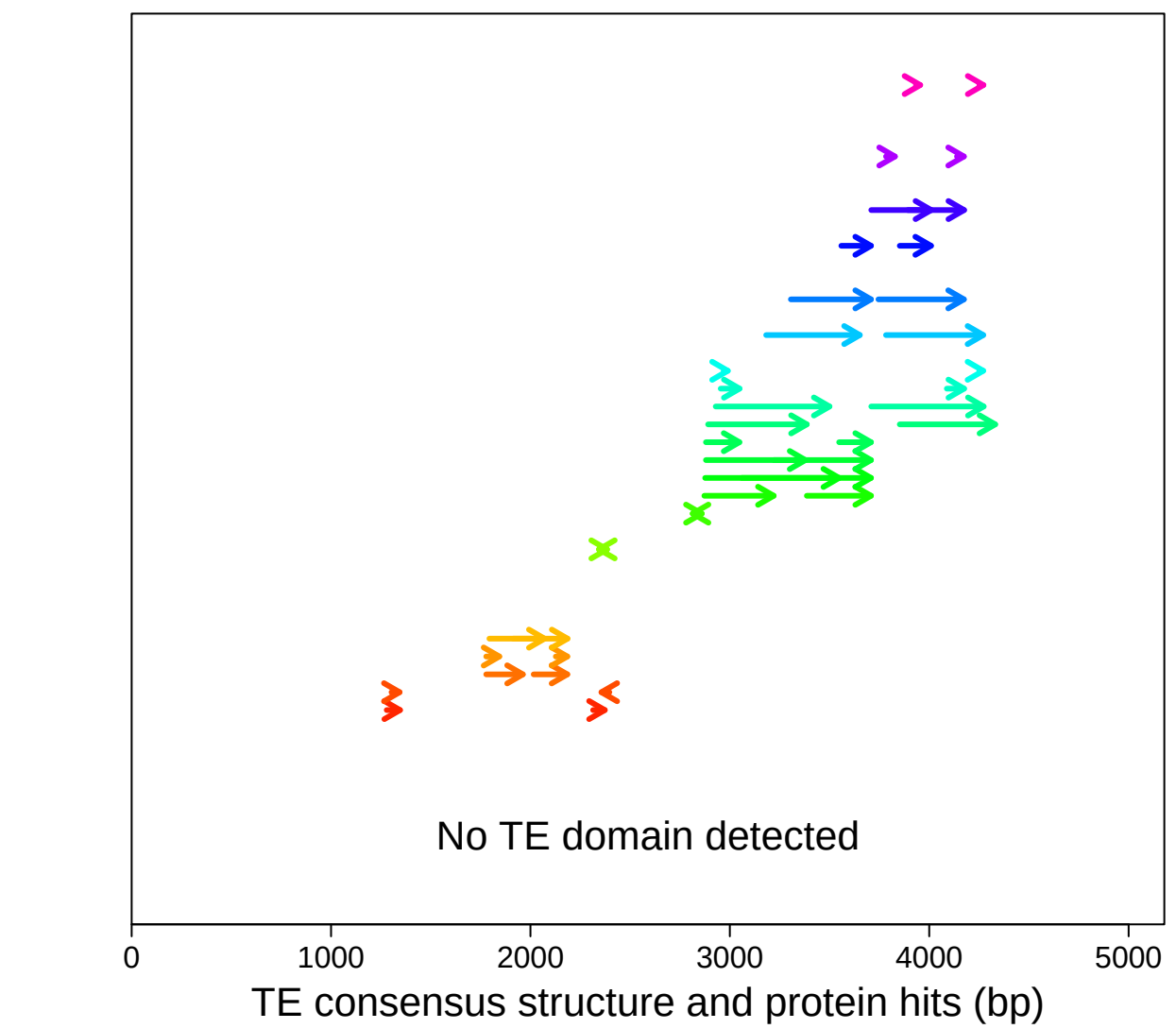
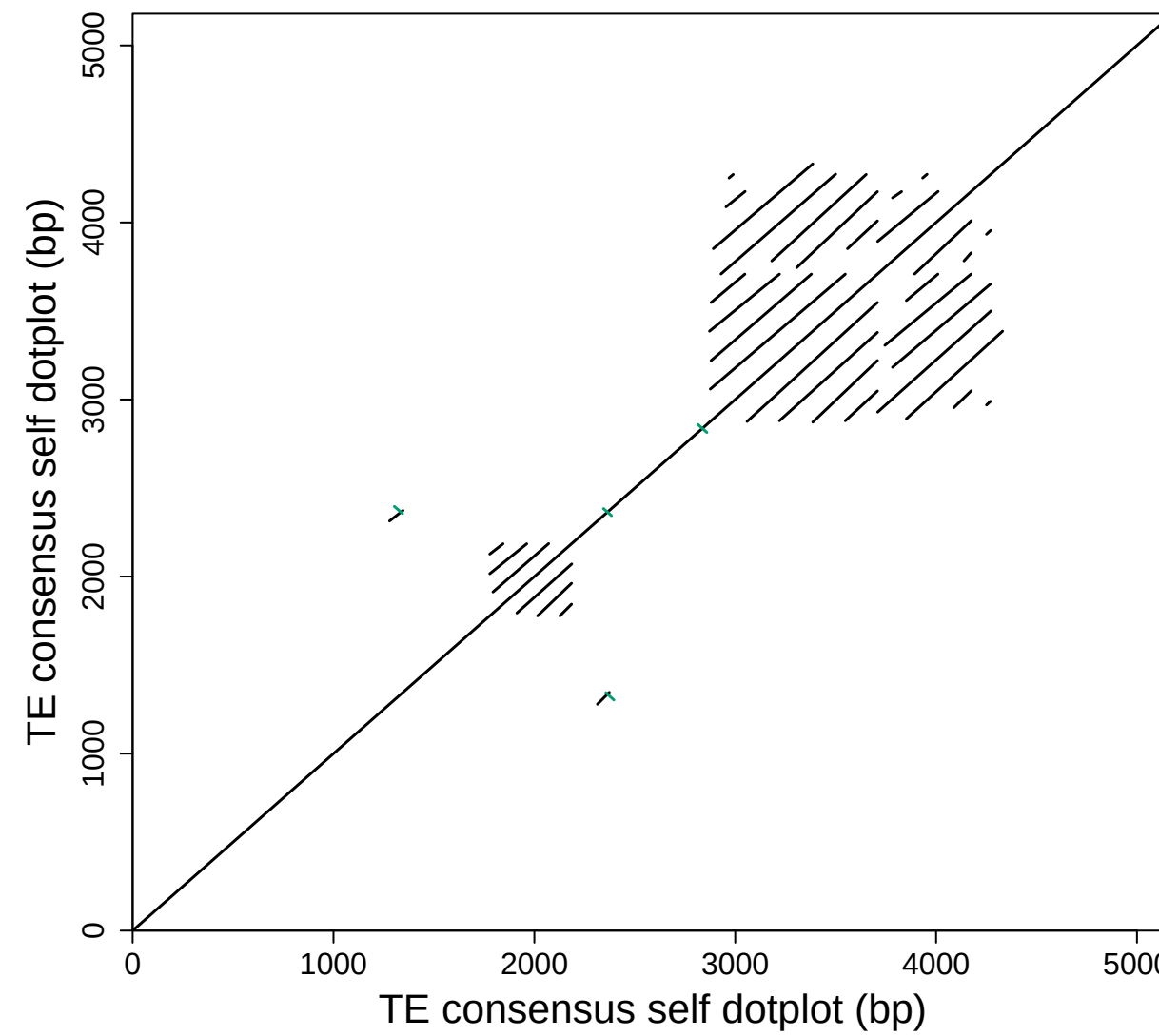
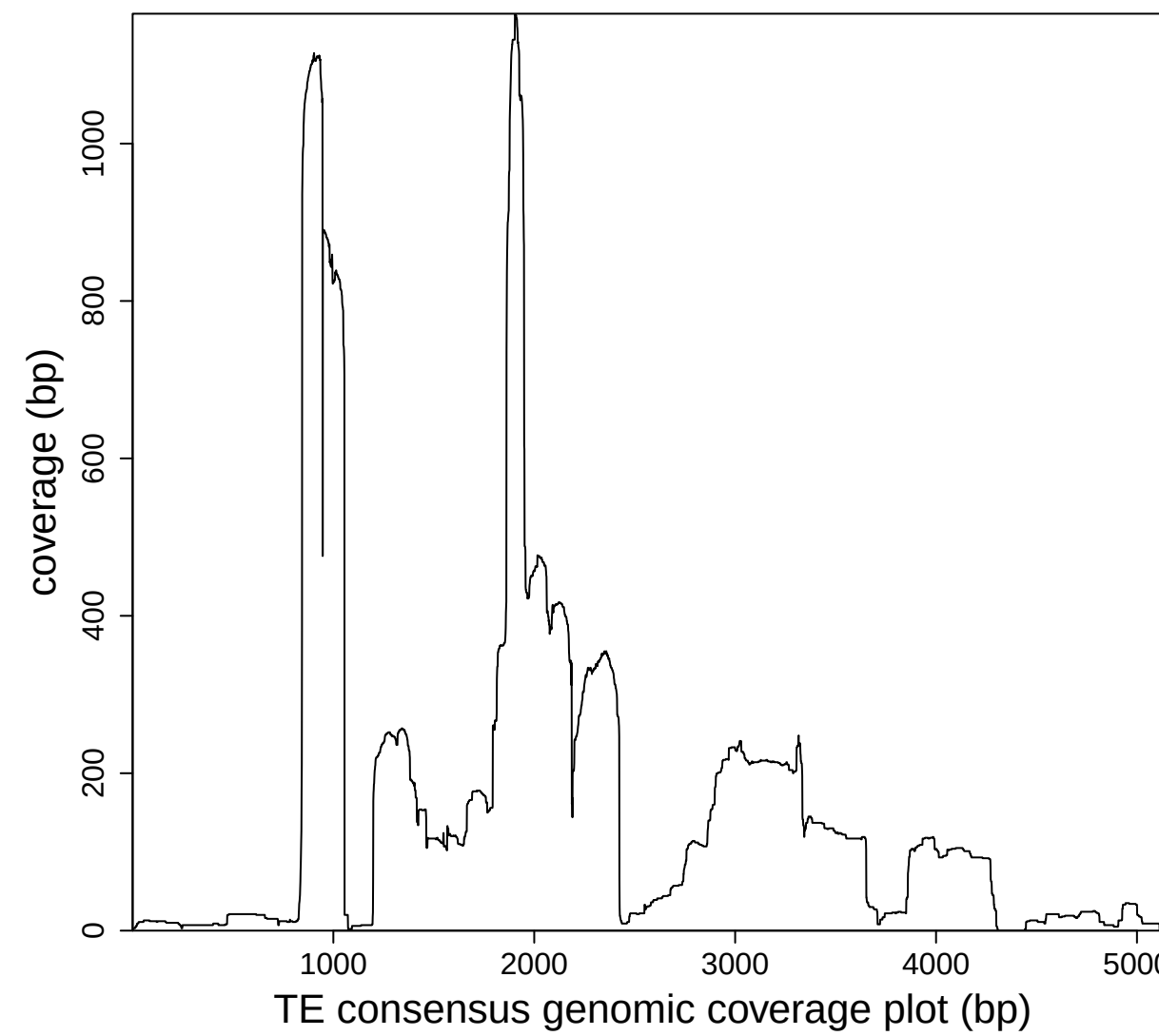
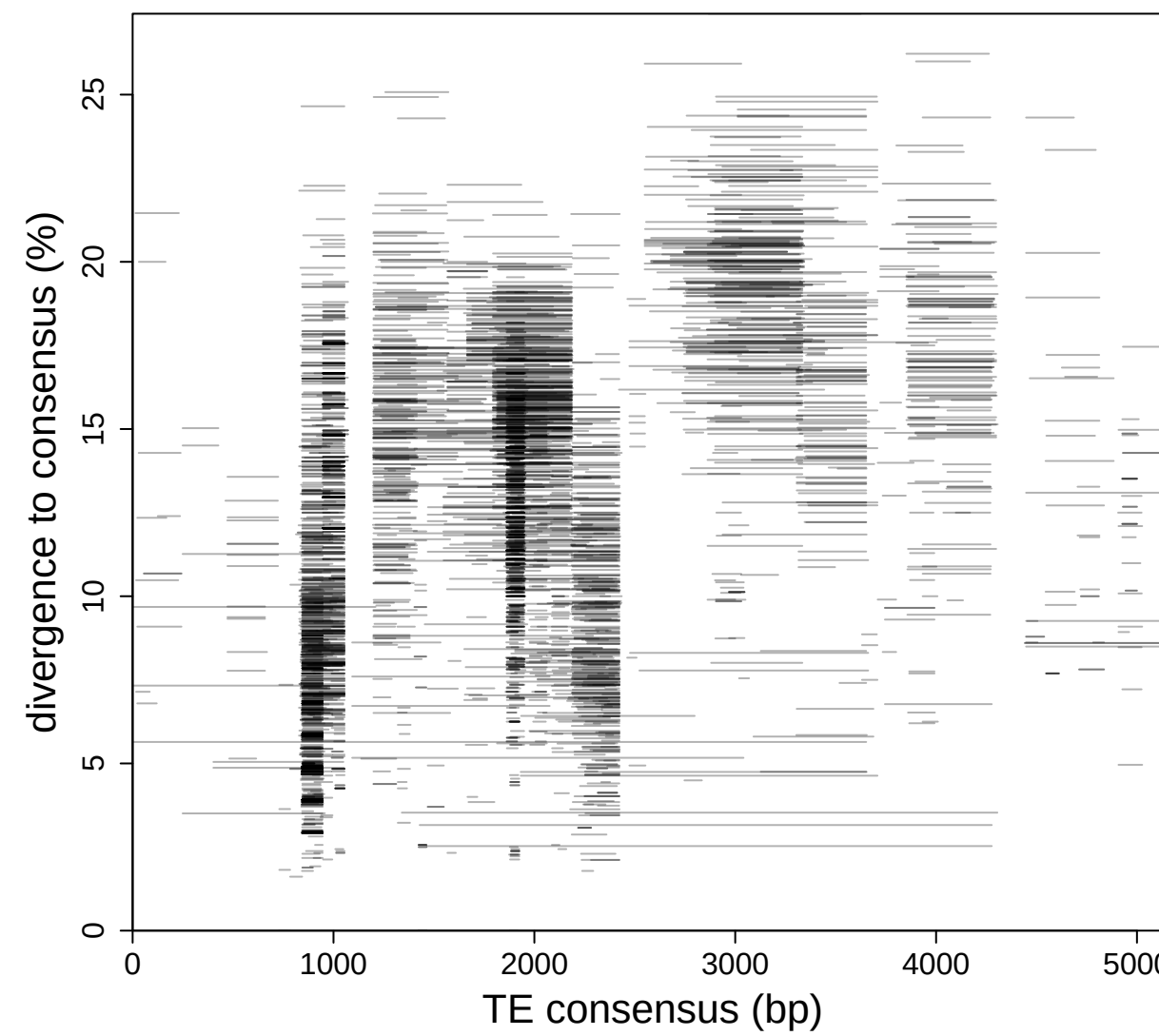
MSA length = 5179





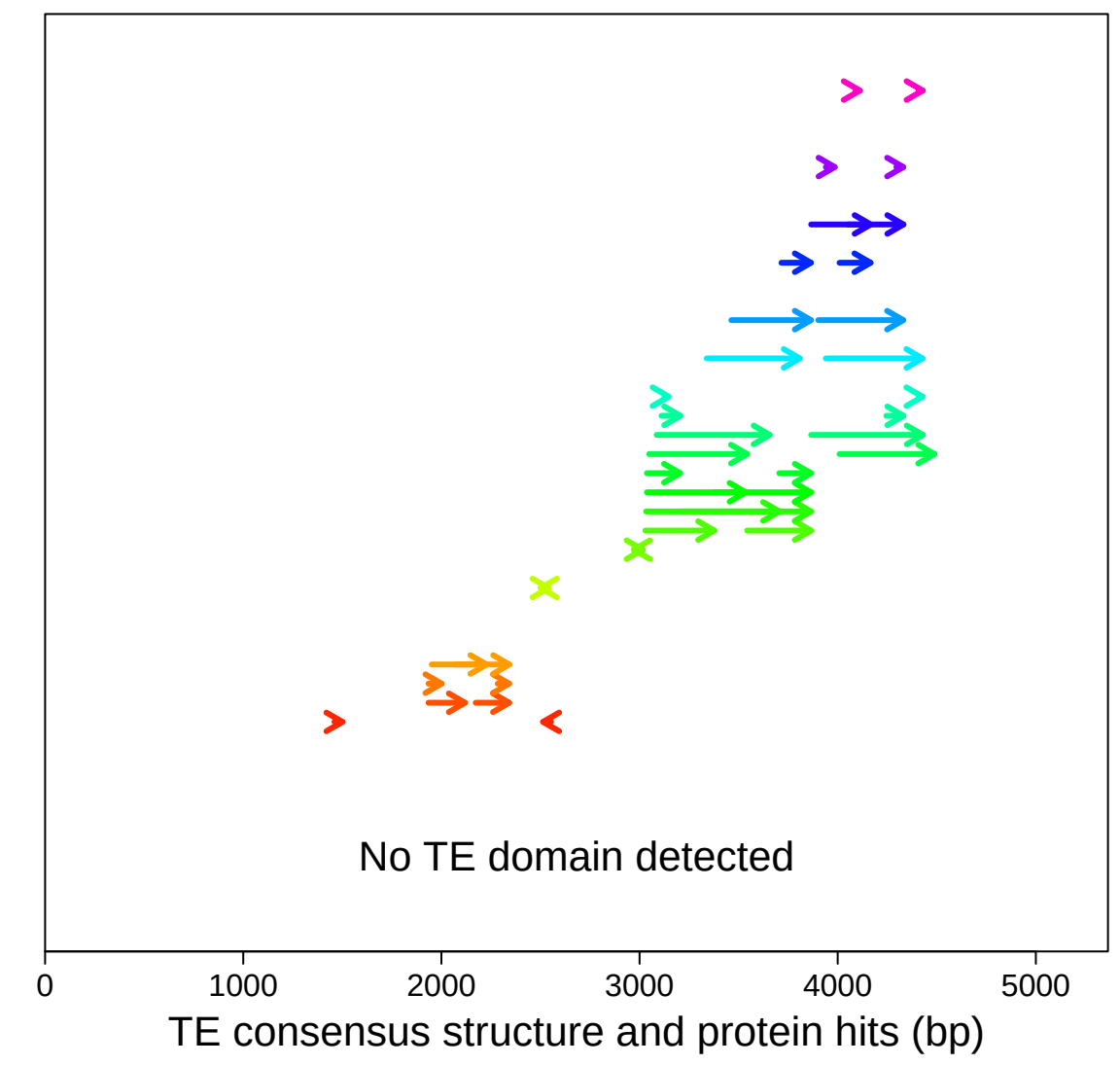
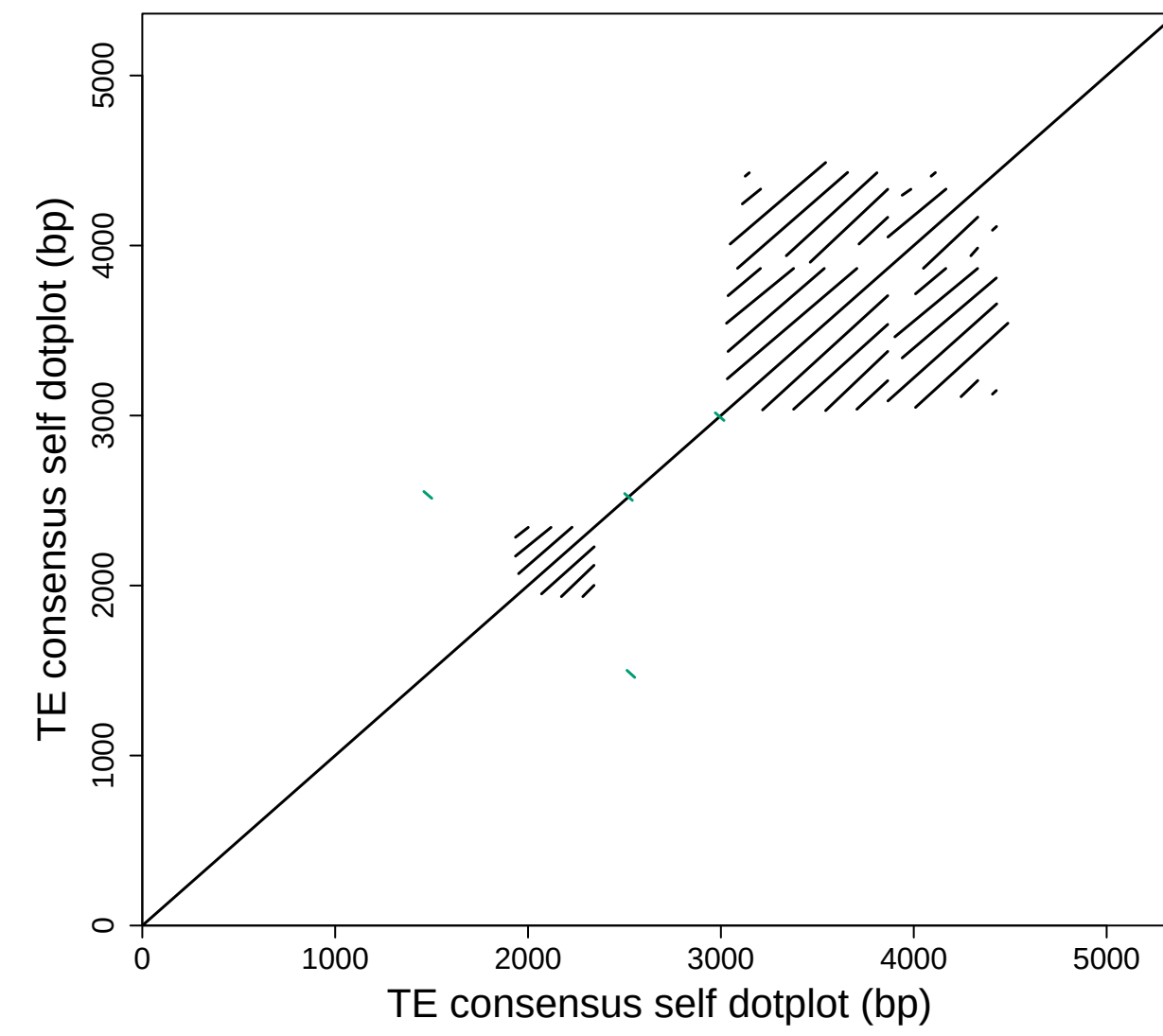
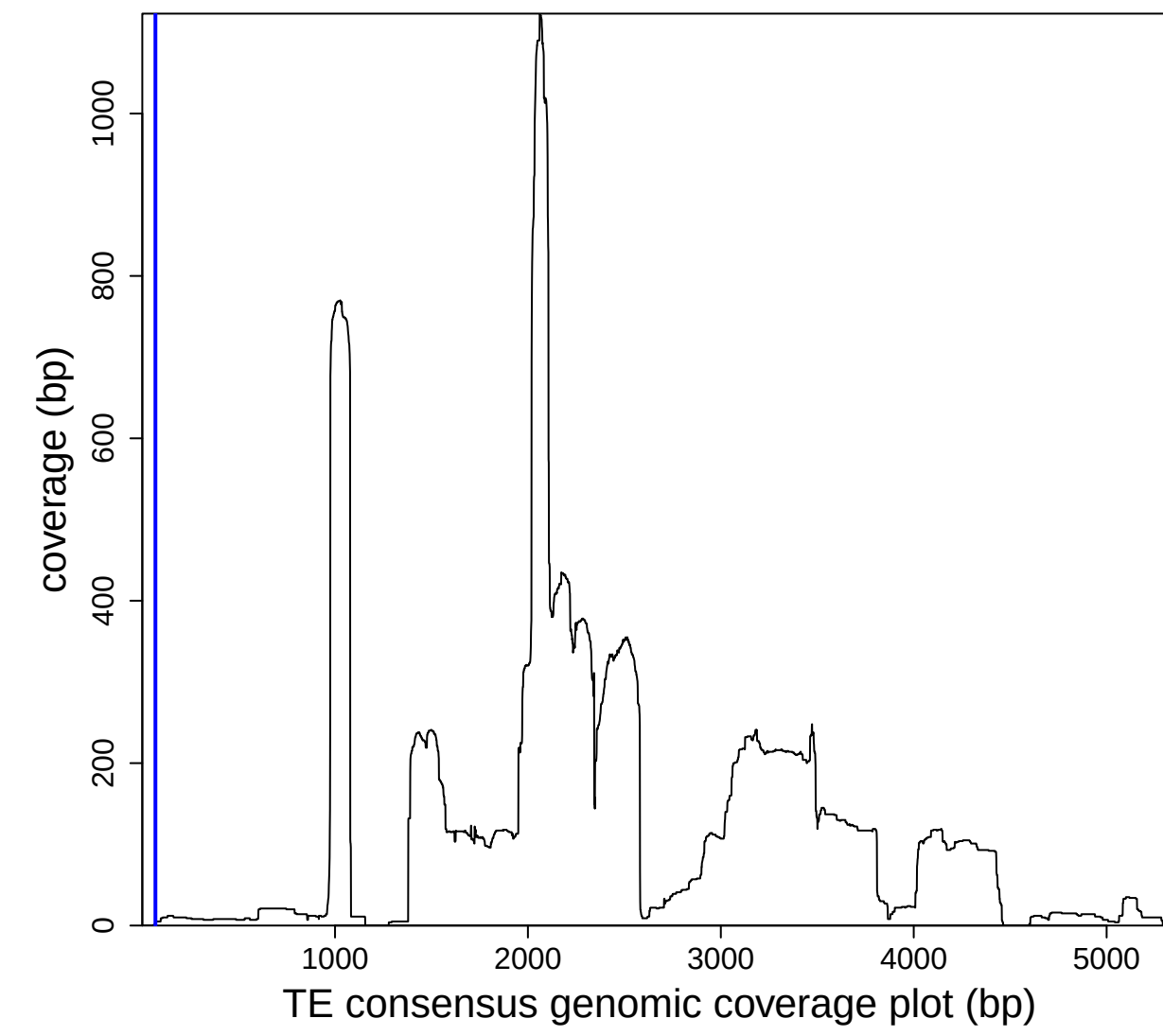
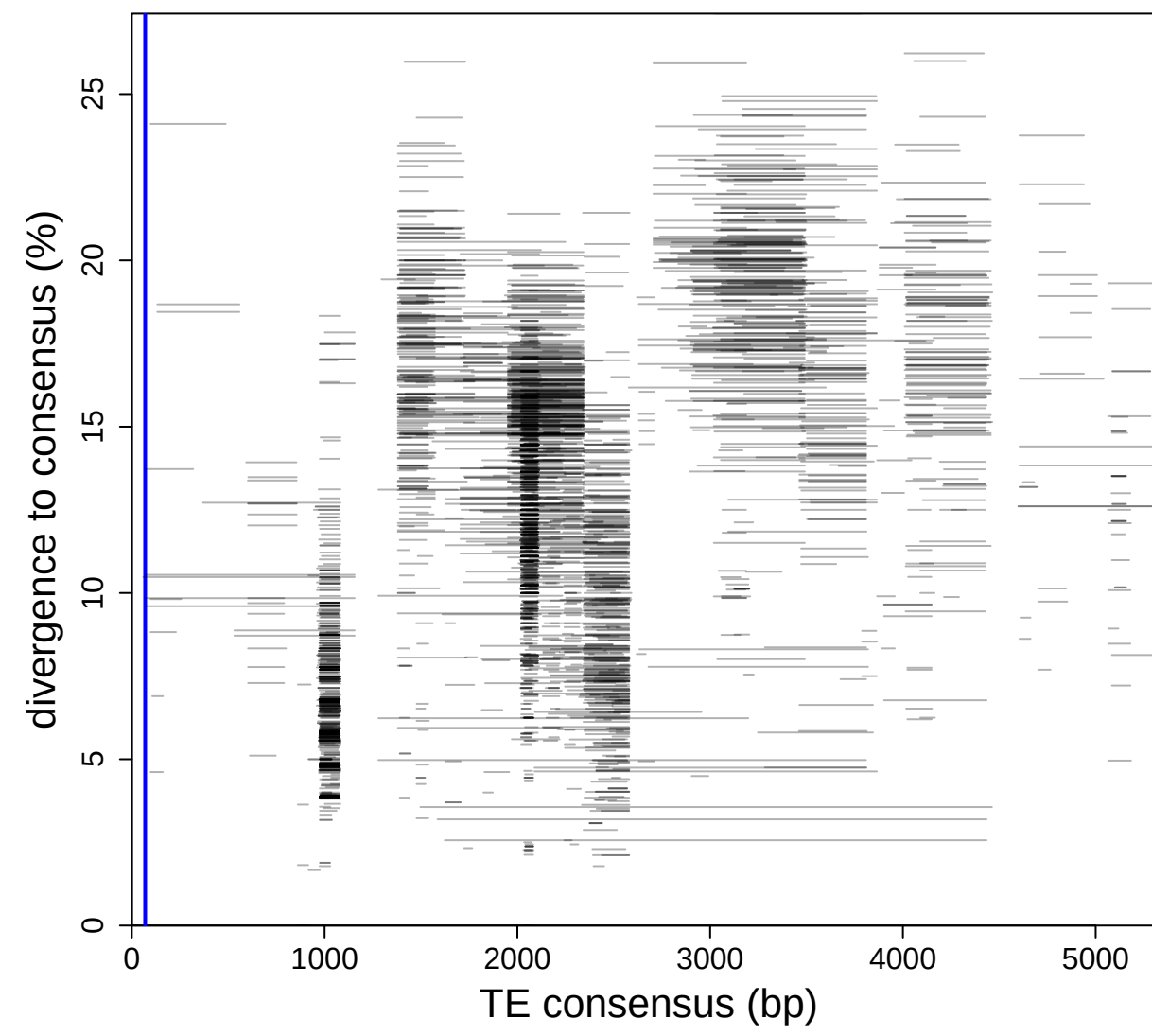
a.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.f
size: 5179bp; fragments: 4213; full length: 0 (>=4661.1bp)

After TEtrimmer 5179 bp



3.fasta.b.bed uf.bed g_1.bed fm_1.bed 0 0 bcln.fa aln.fa cl.f
size: 5364bp; fragments: 3221; full length: 0 (>=4827.6bp)

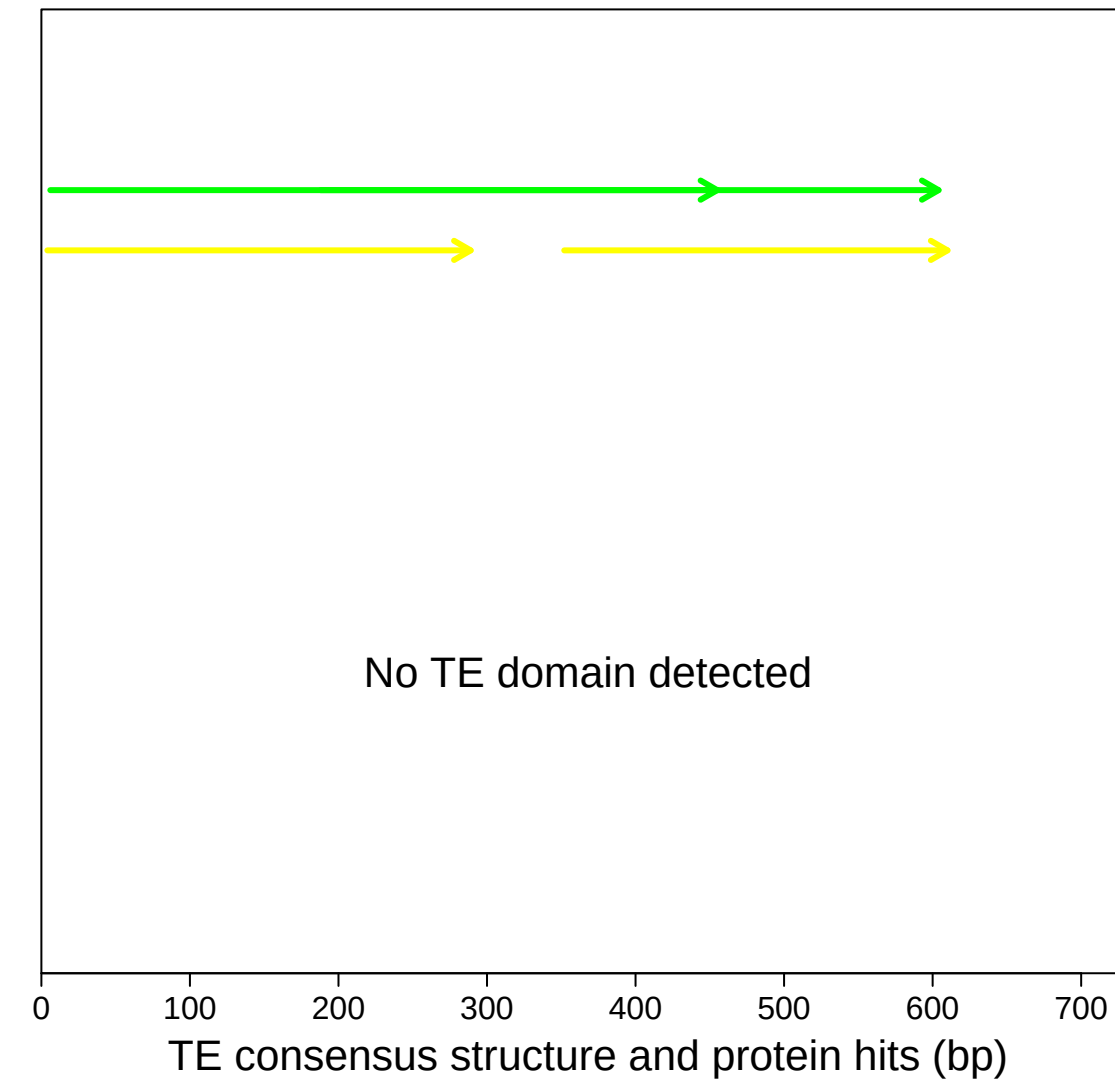
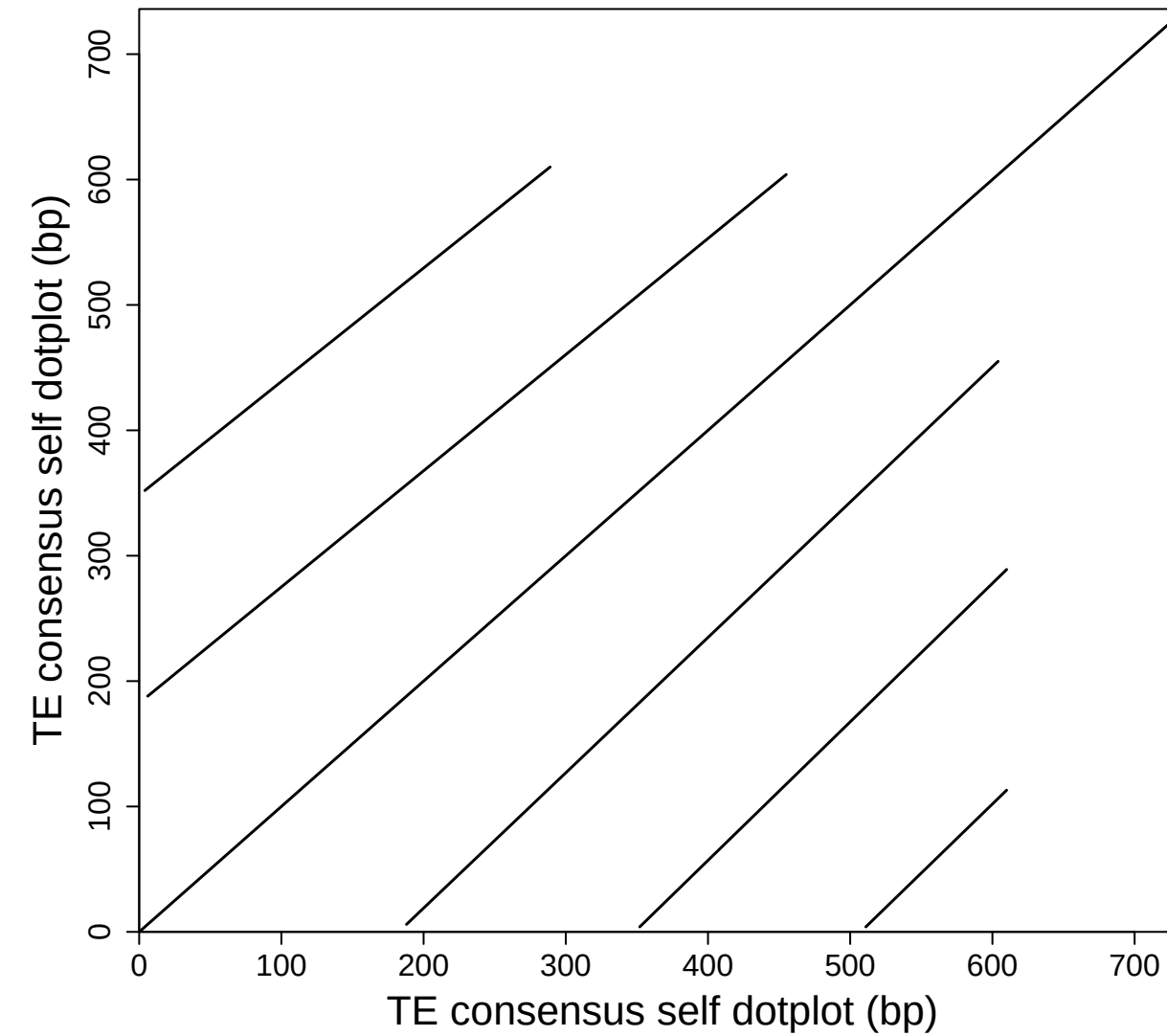
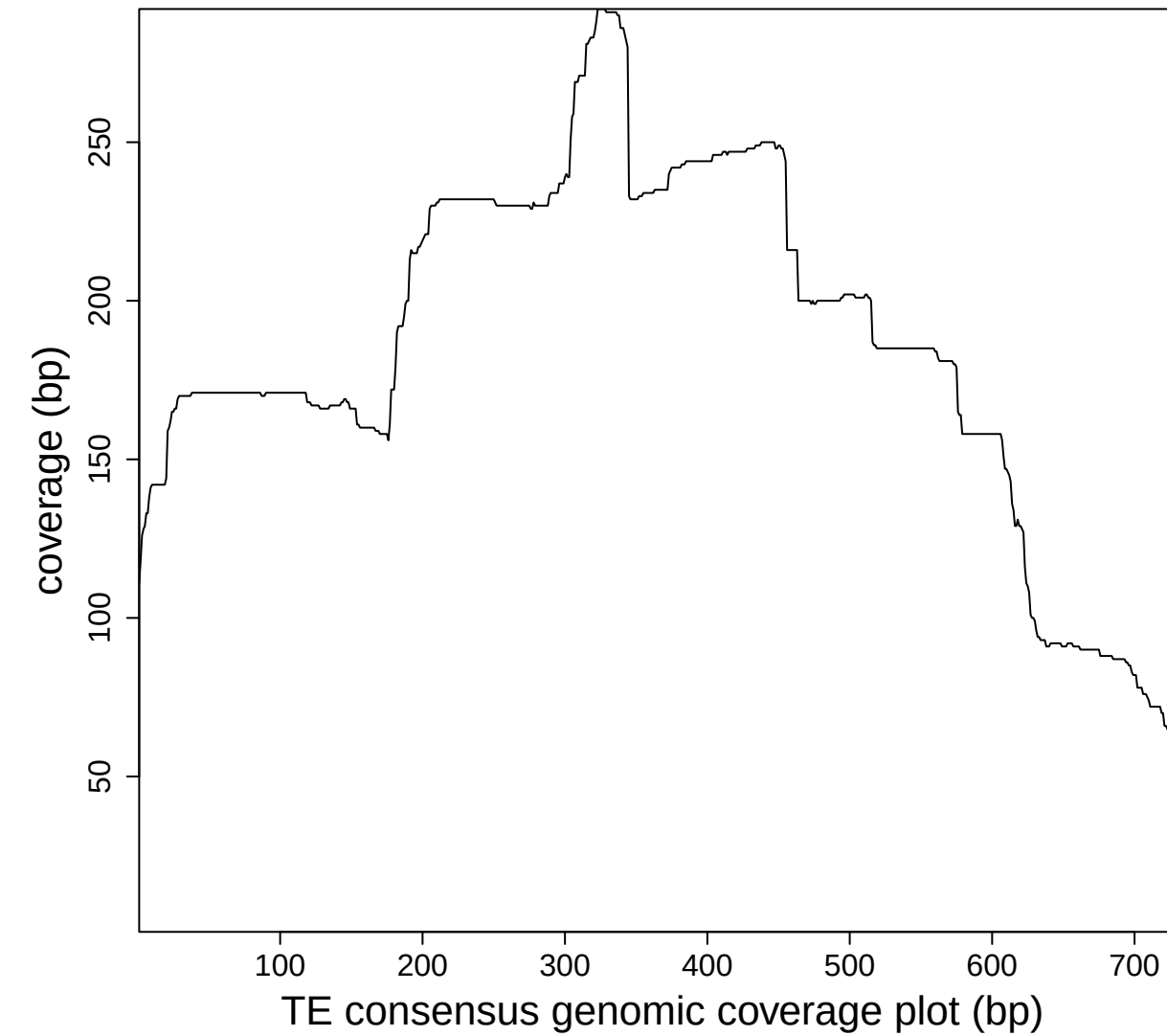
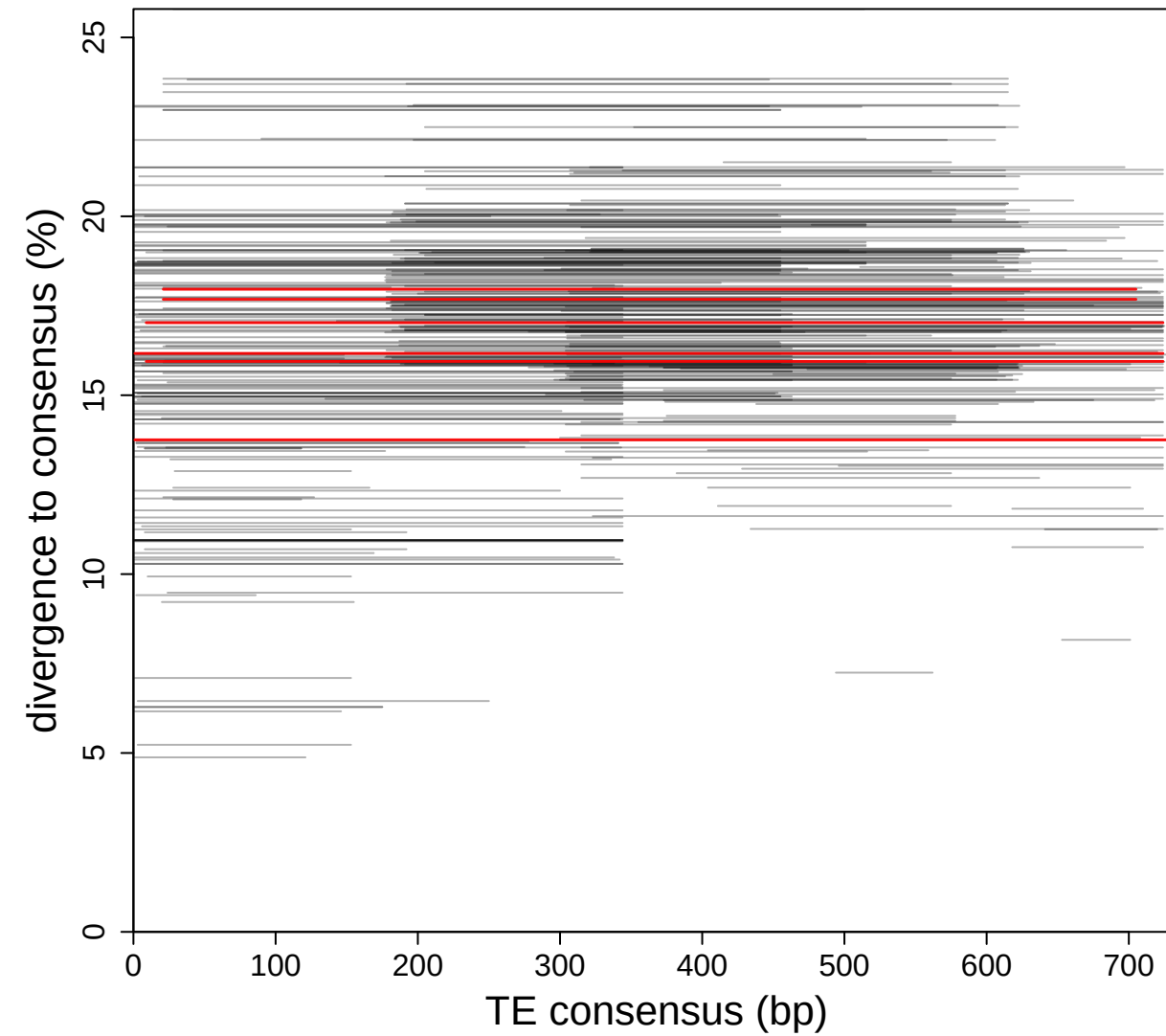
After TEtrimmer Extended plot Blue lines are boundaries



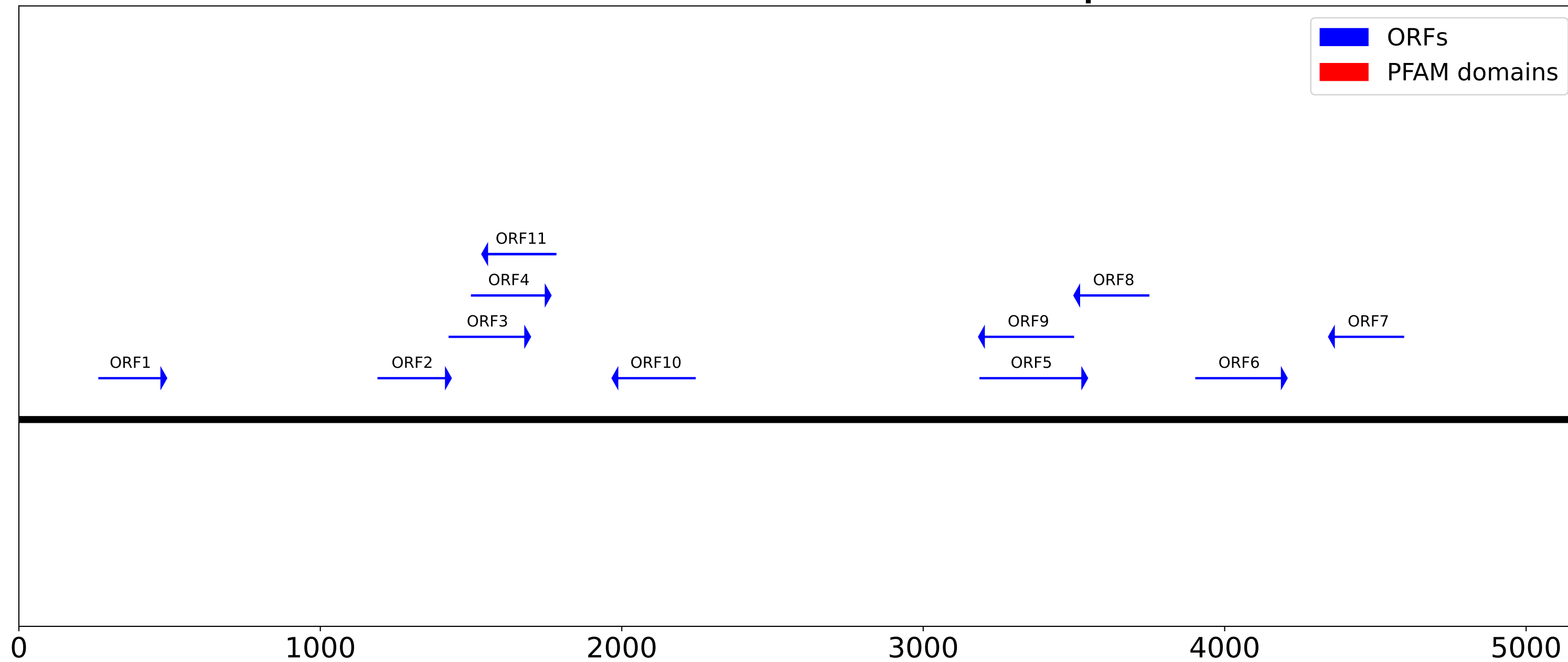
Before TEtrimmer 736 bp

TE: rnd_4_family_1373

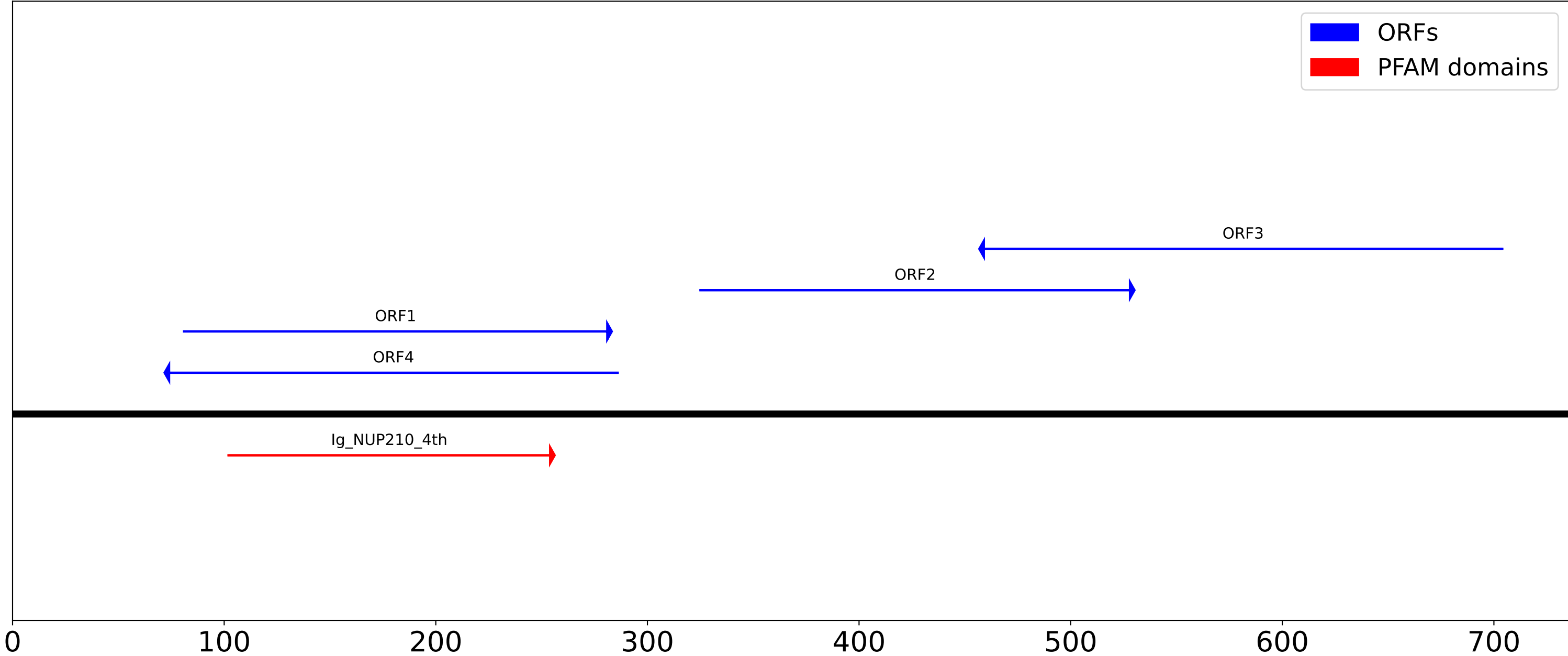
size: 736bp; fragments: 351; full length: 6 (≥ 662.4 bp)



After TEtrimmer ORF and PFAM domain plot



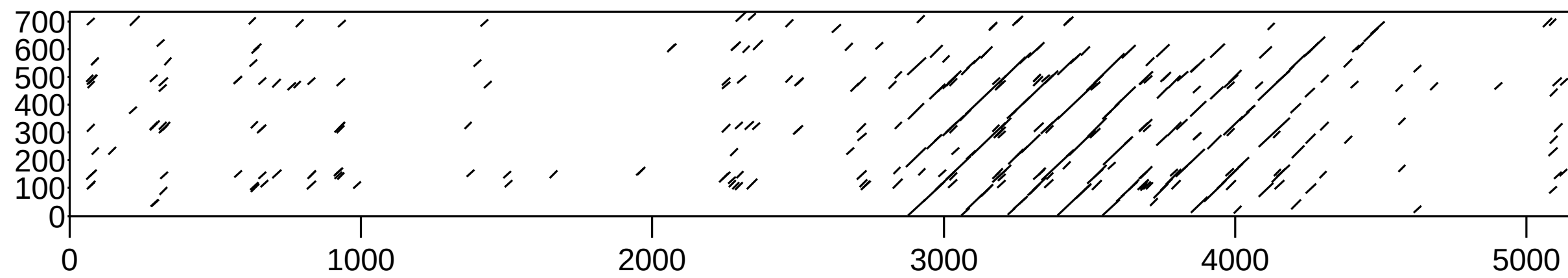
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

isensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)