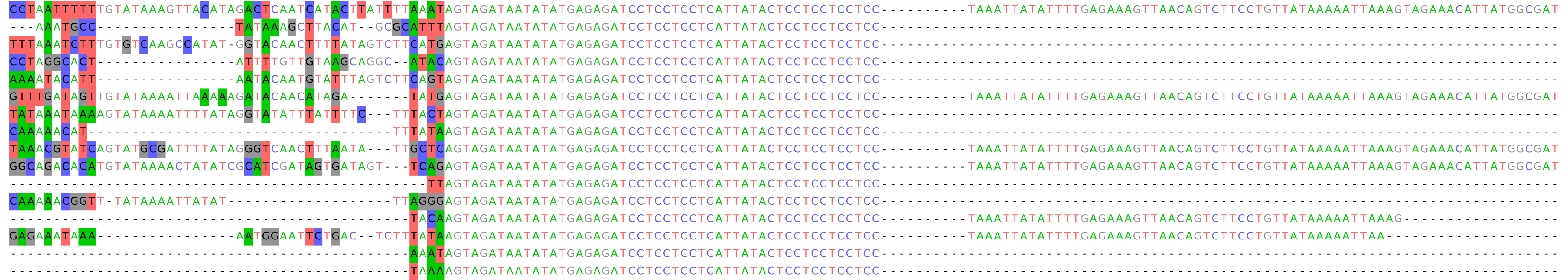
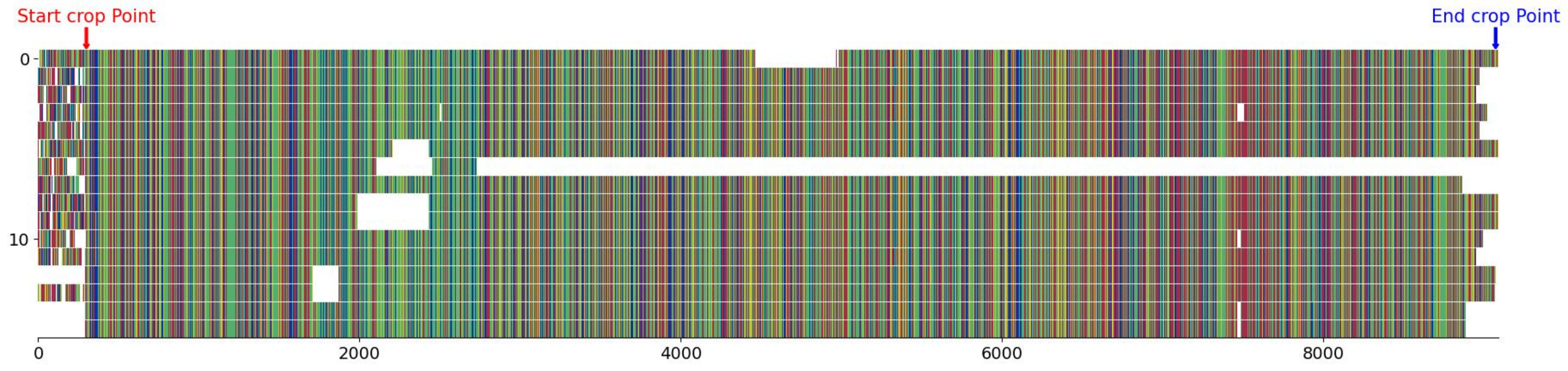


Start crop Point

End crop Point

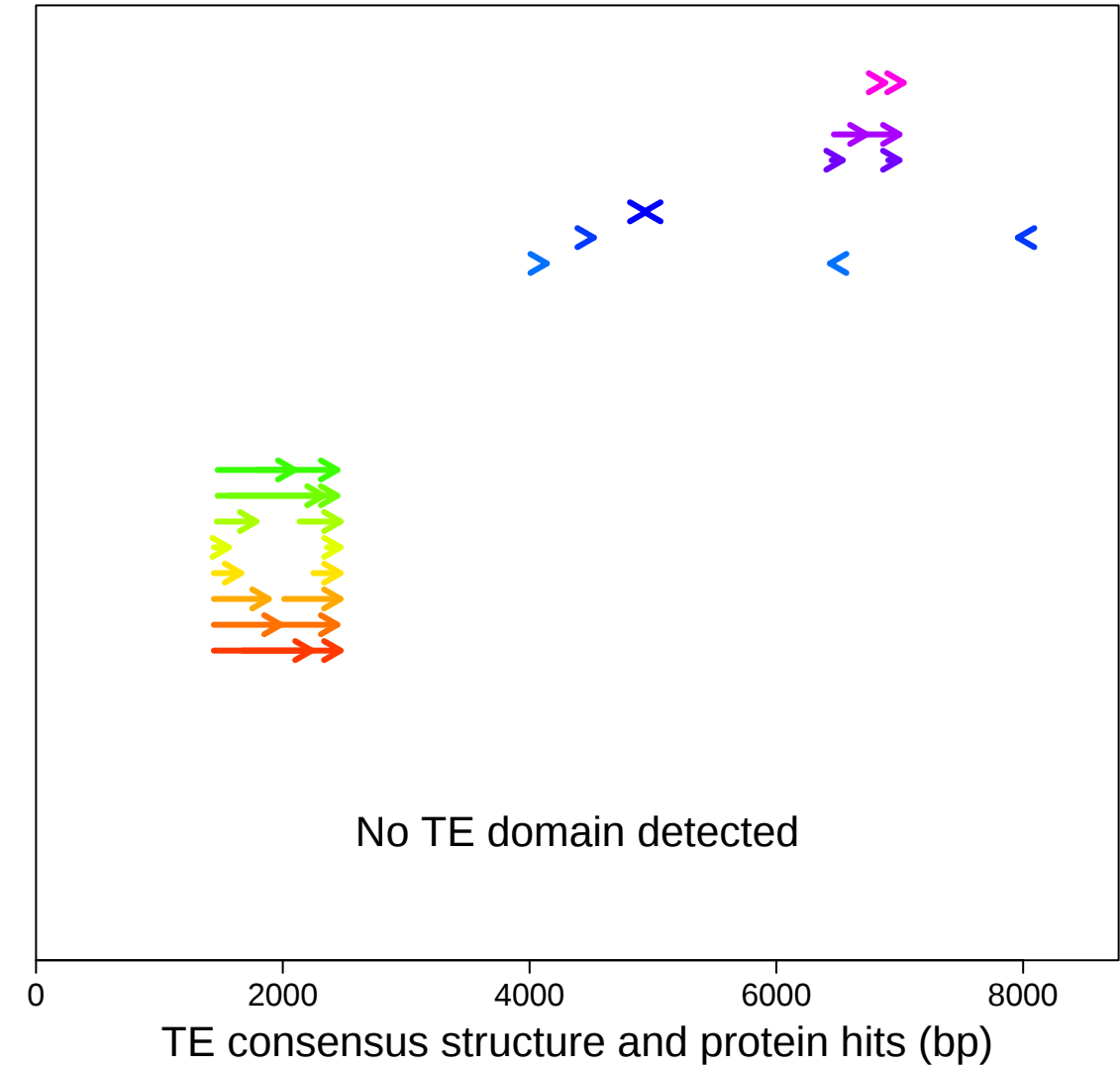
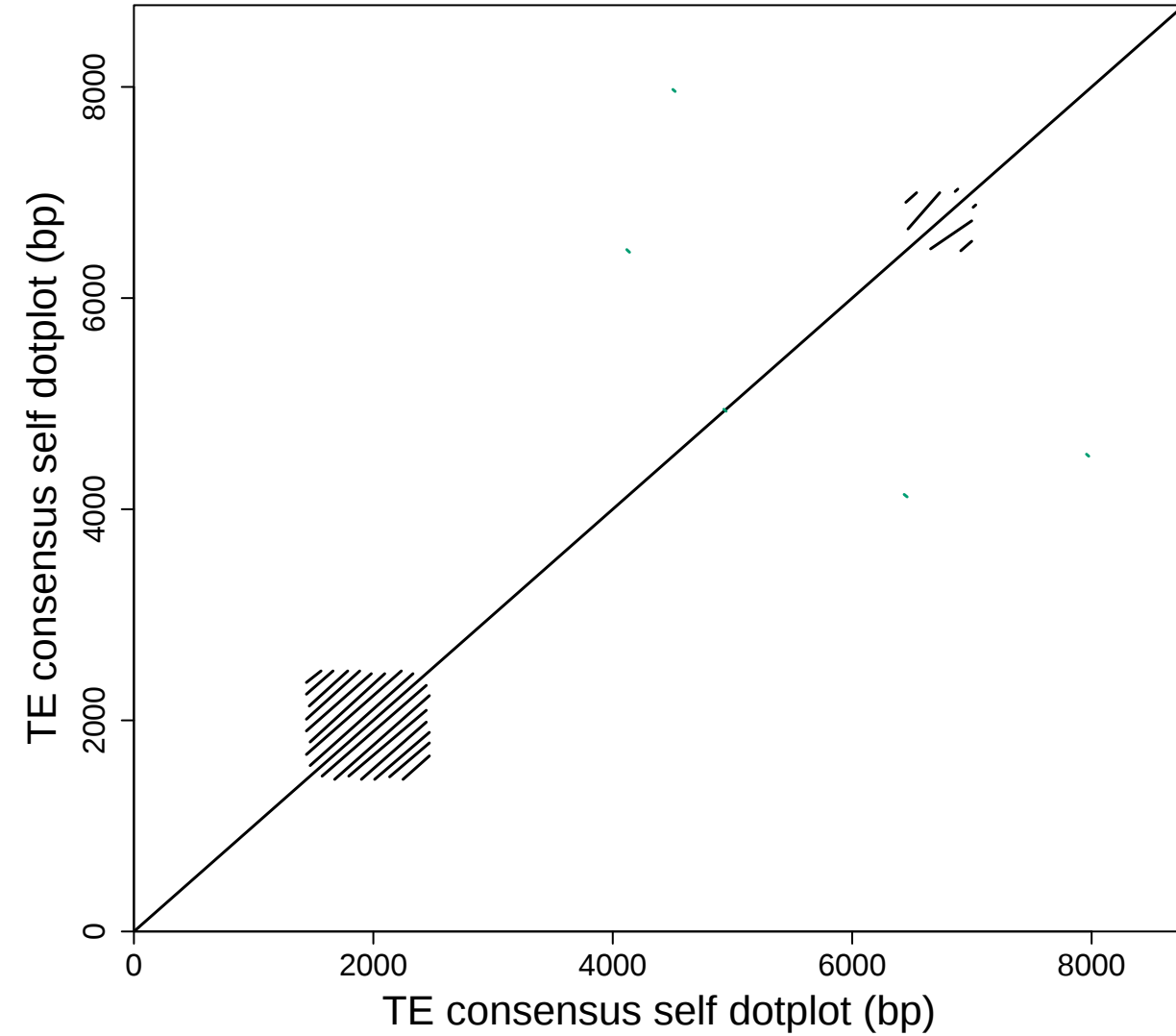
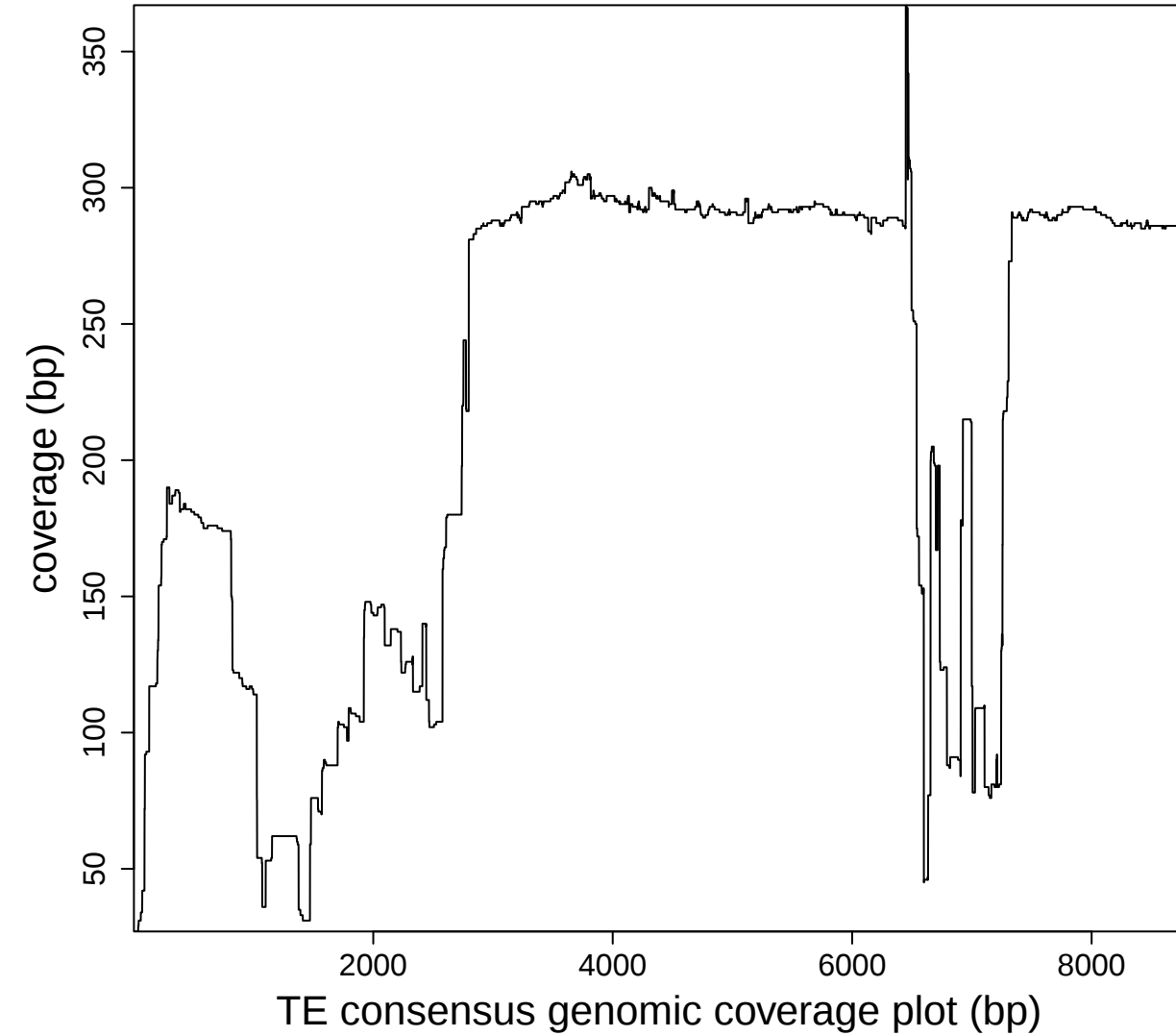
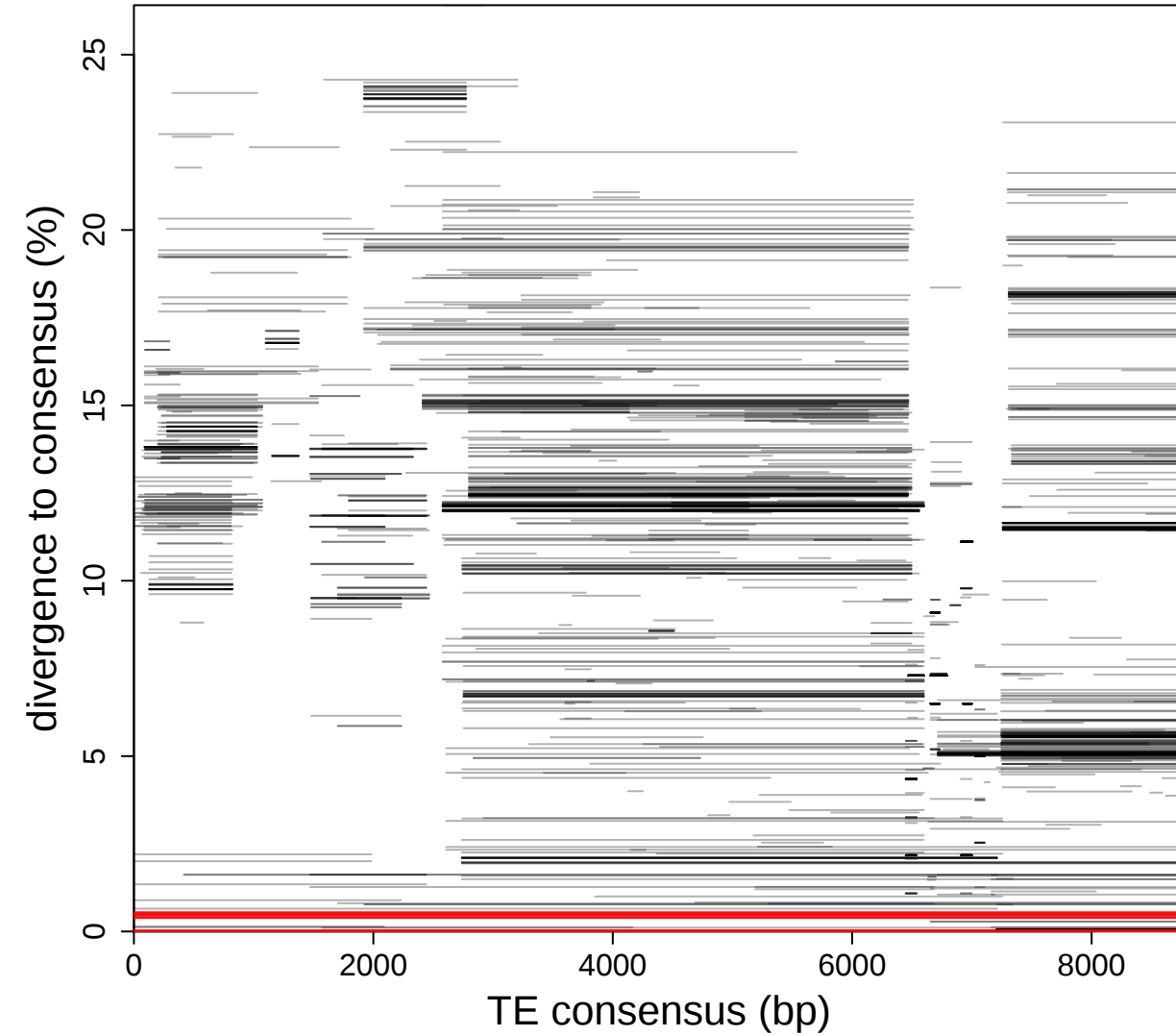
MSA length = 8774





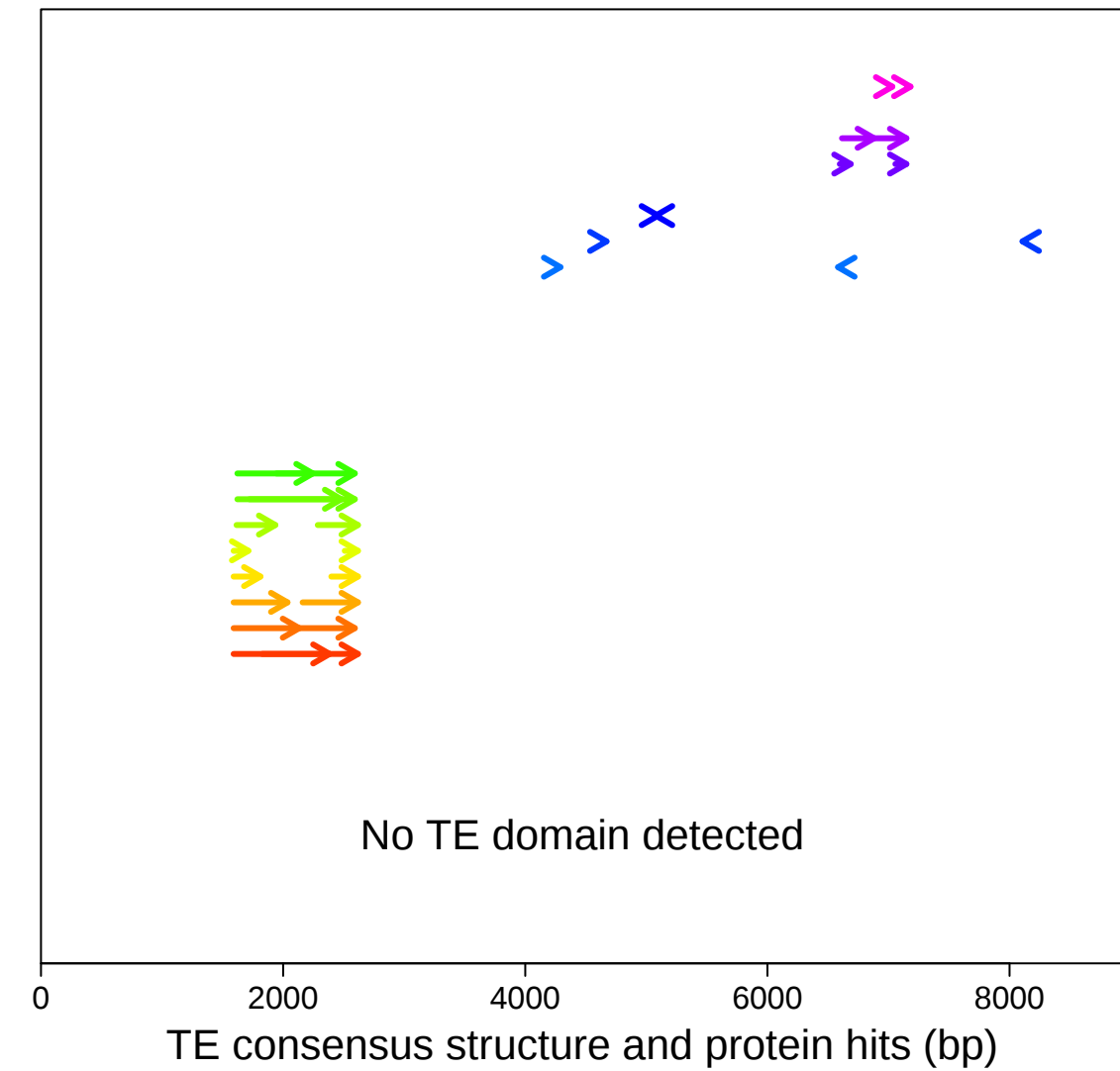
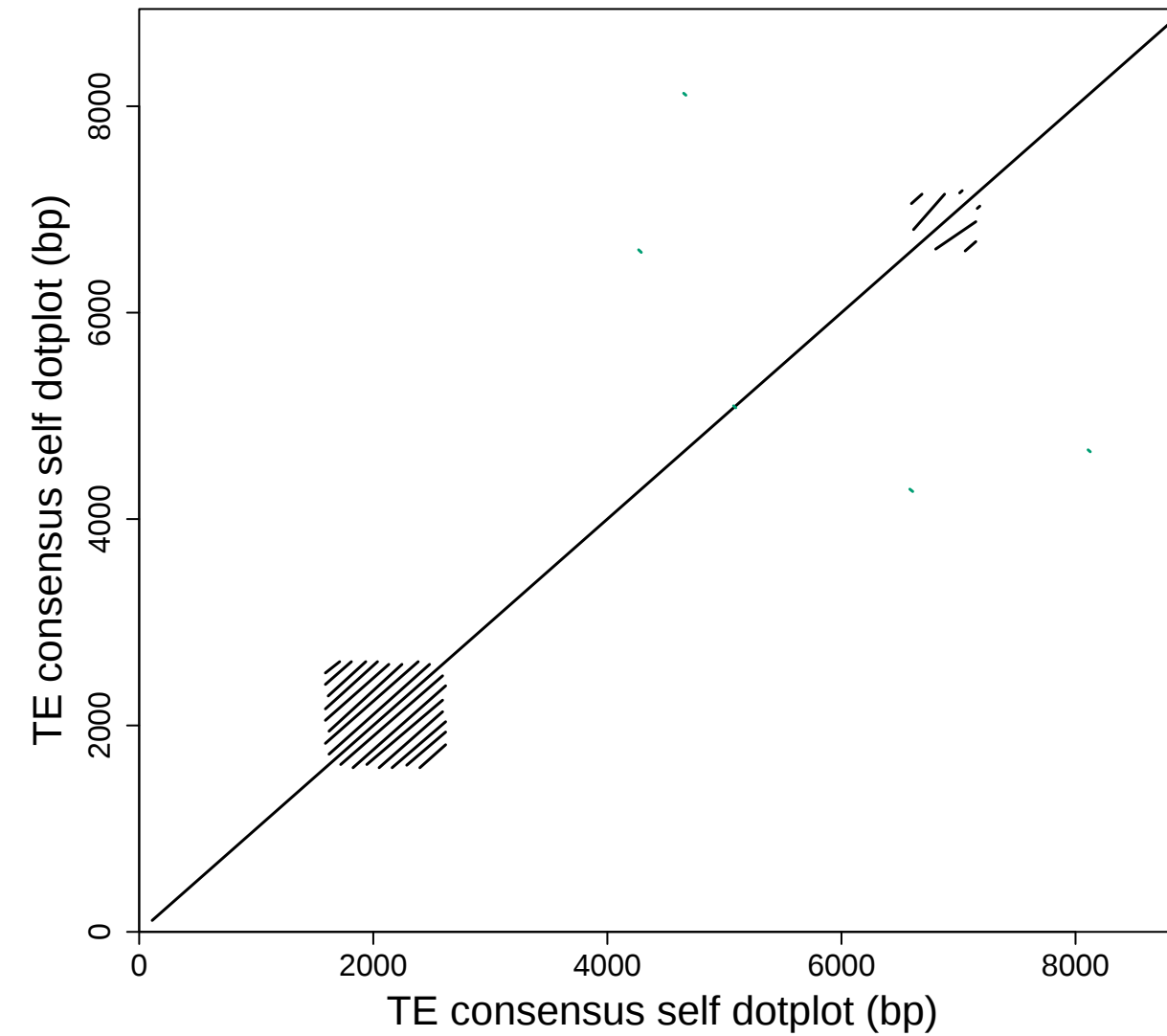
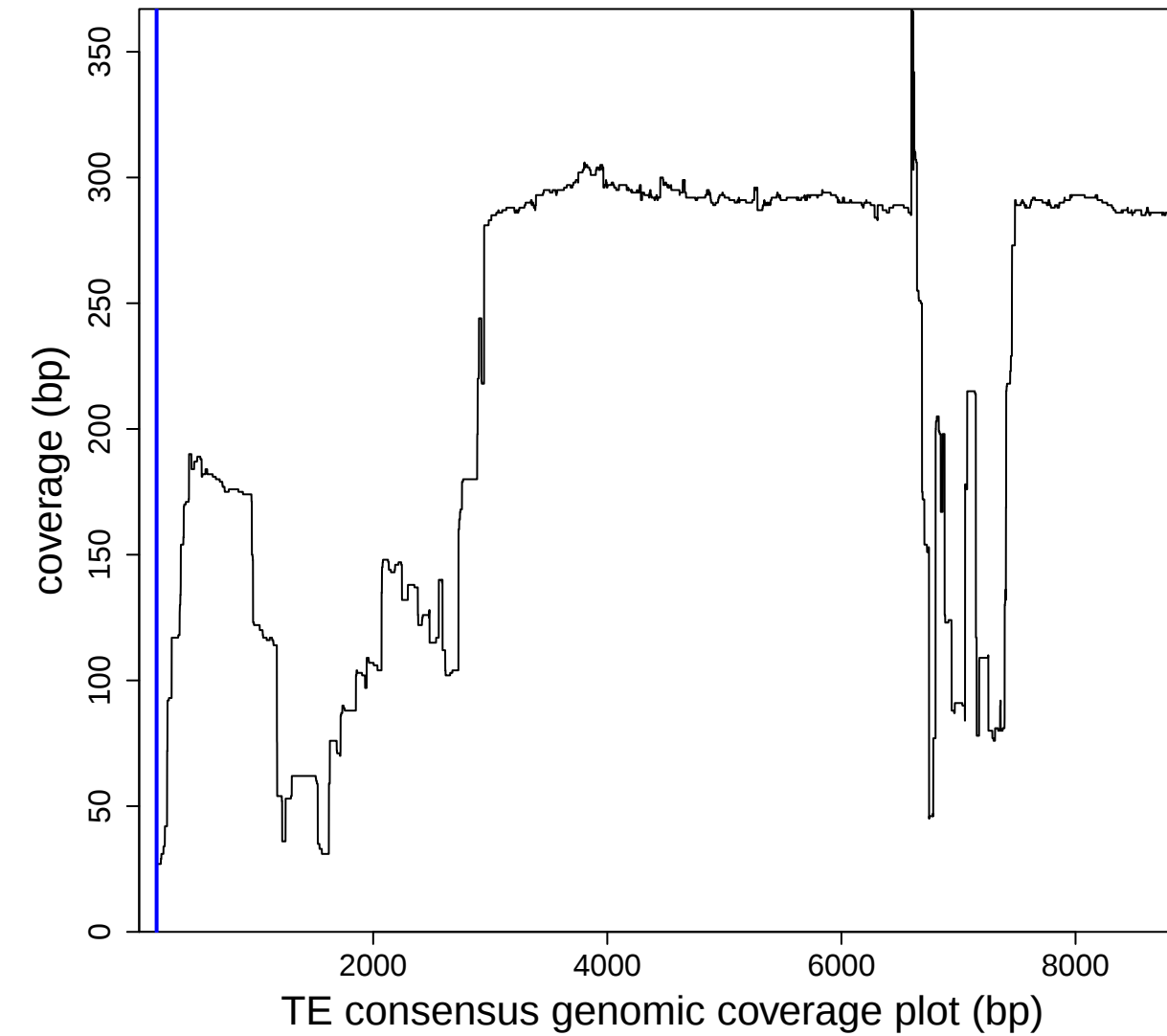
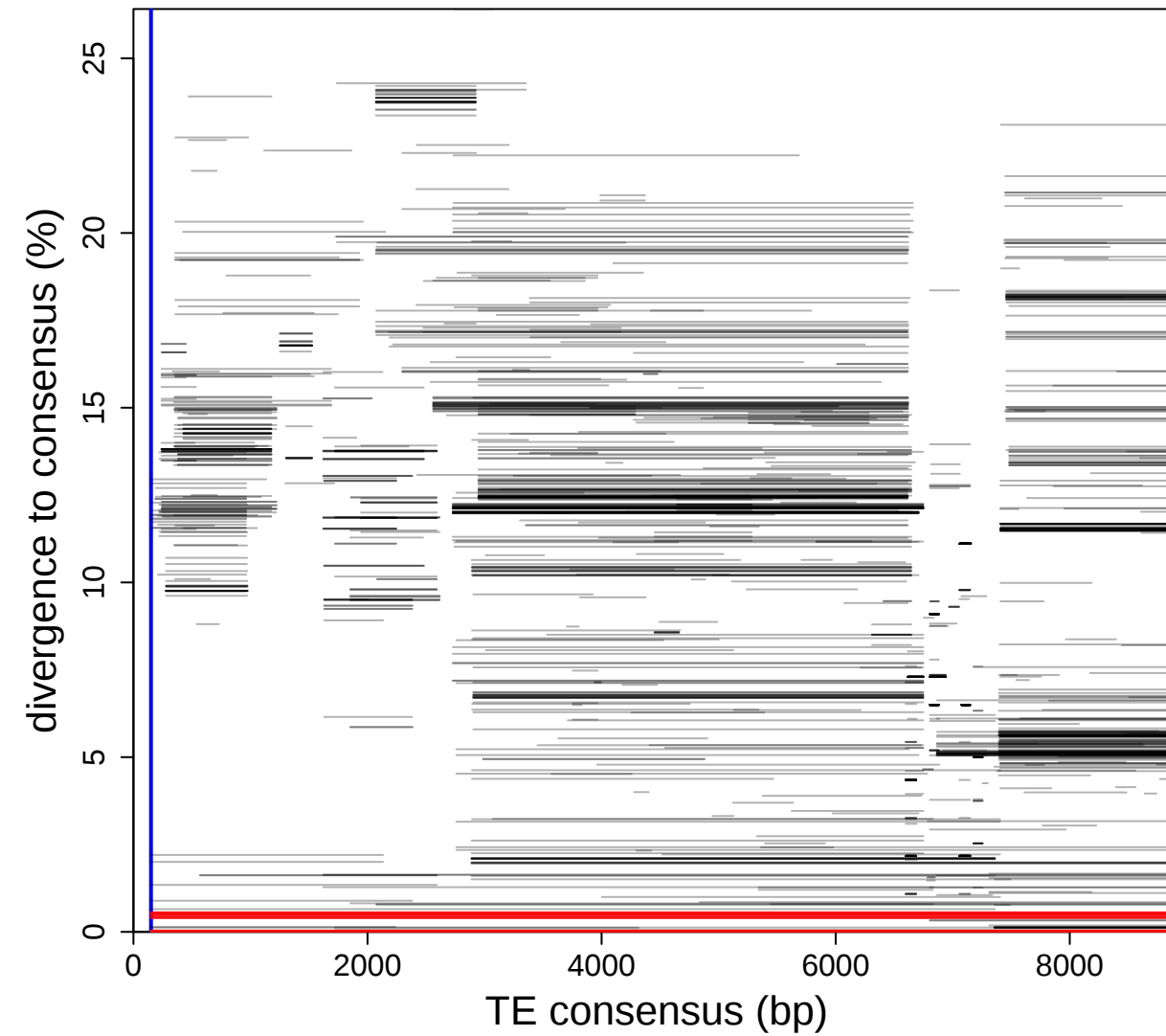
l.b.bed_uf.bed_g_5.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa
size: 8774bp; fragments: 1505; full length: 5 (≥ 7896.6 bp)

After TEtrimmer 8774 bp

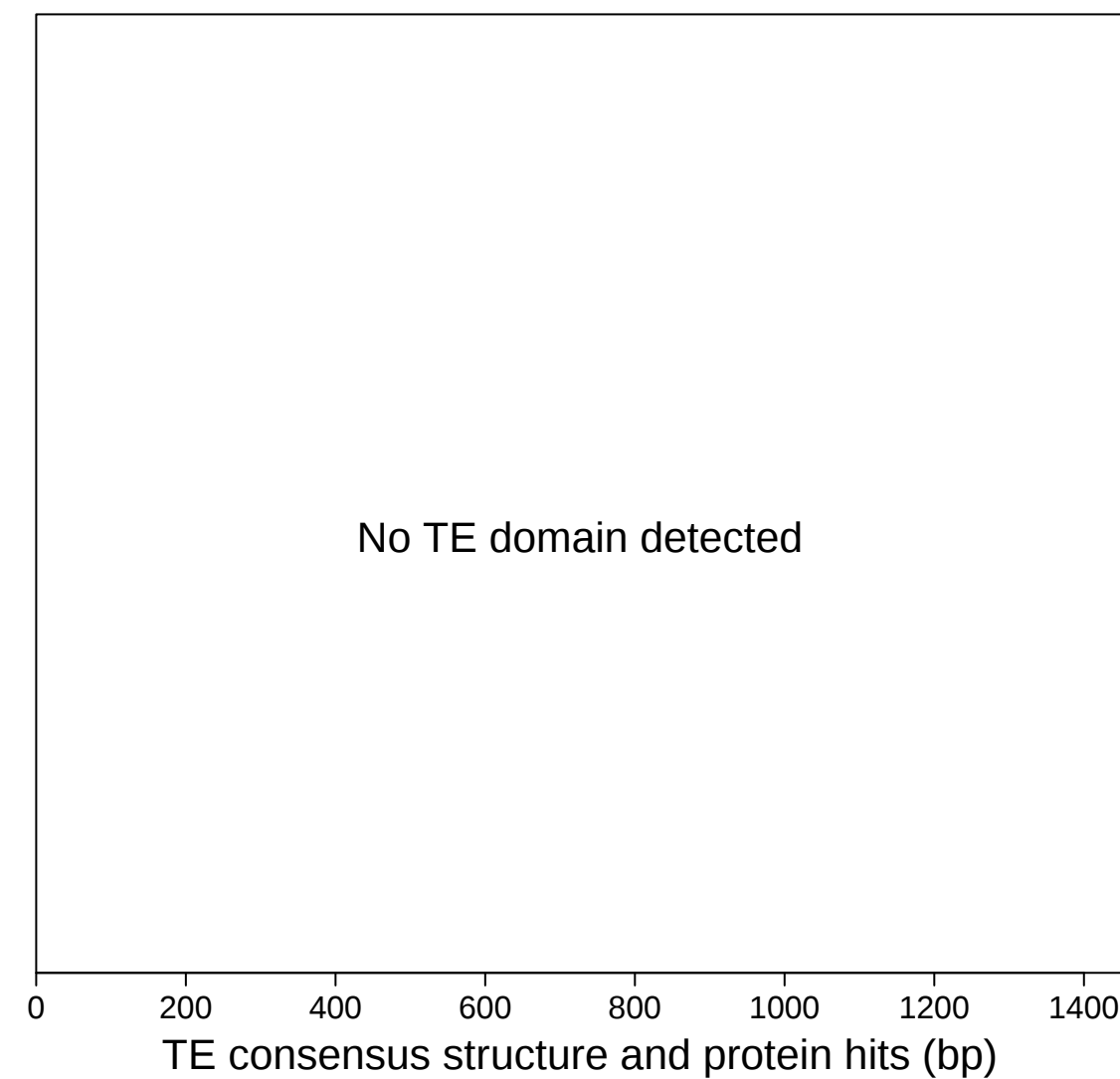
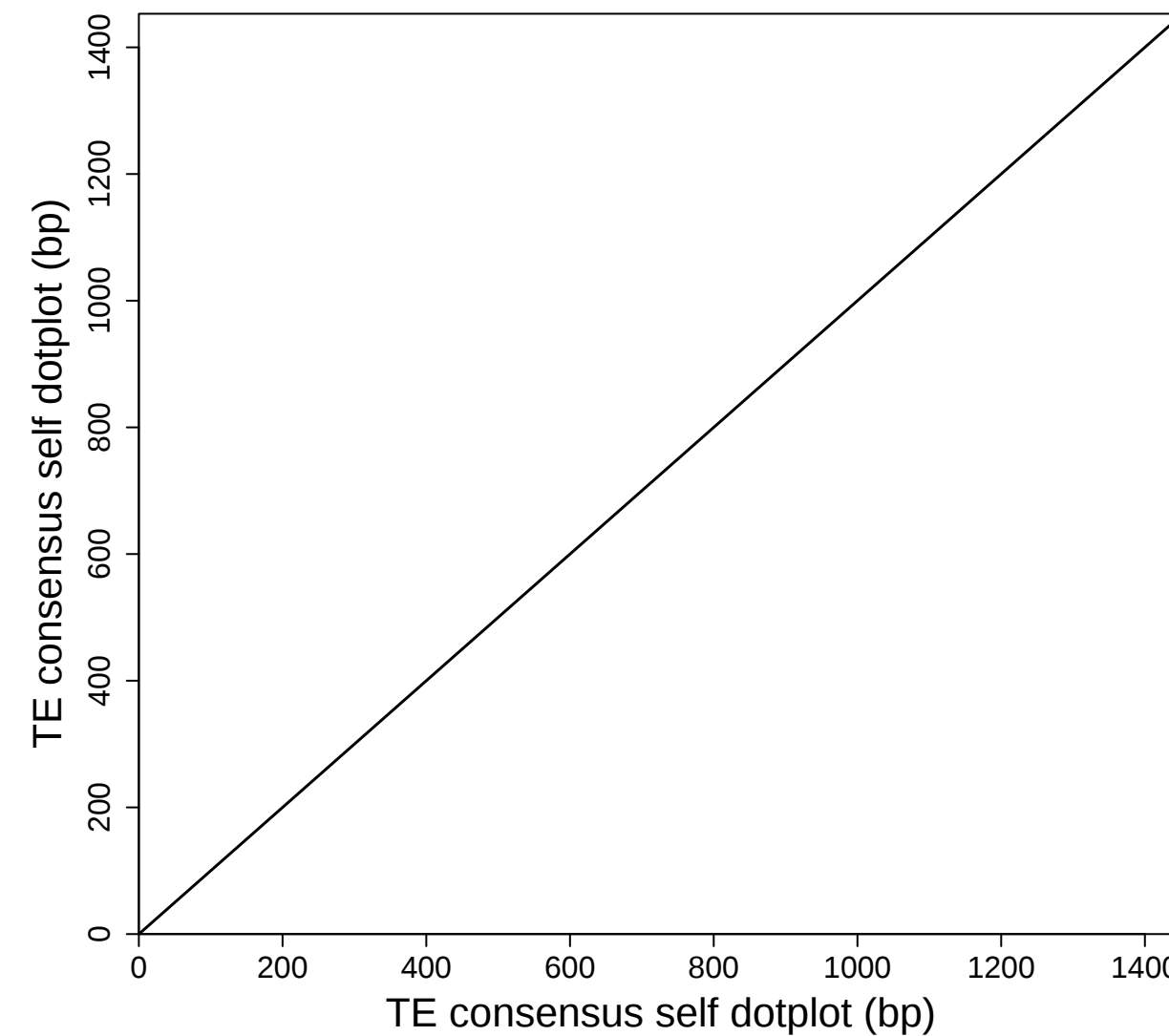
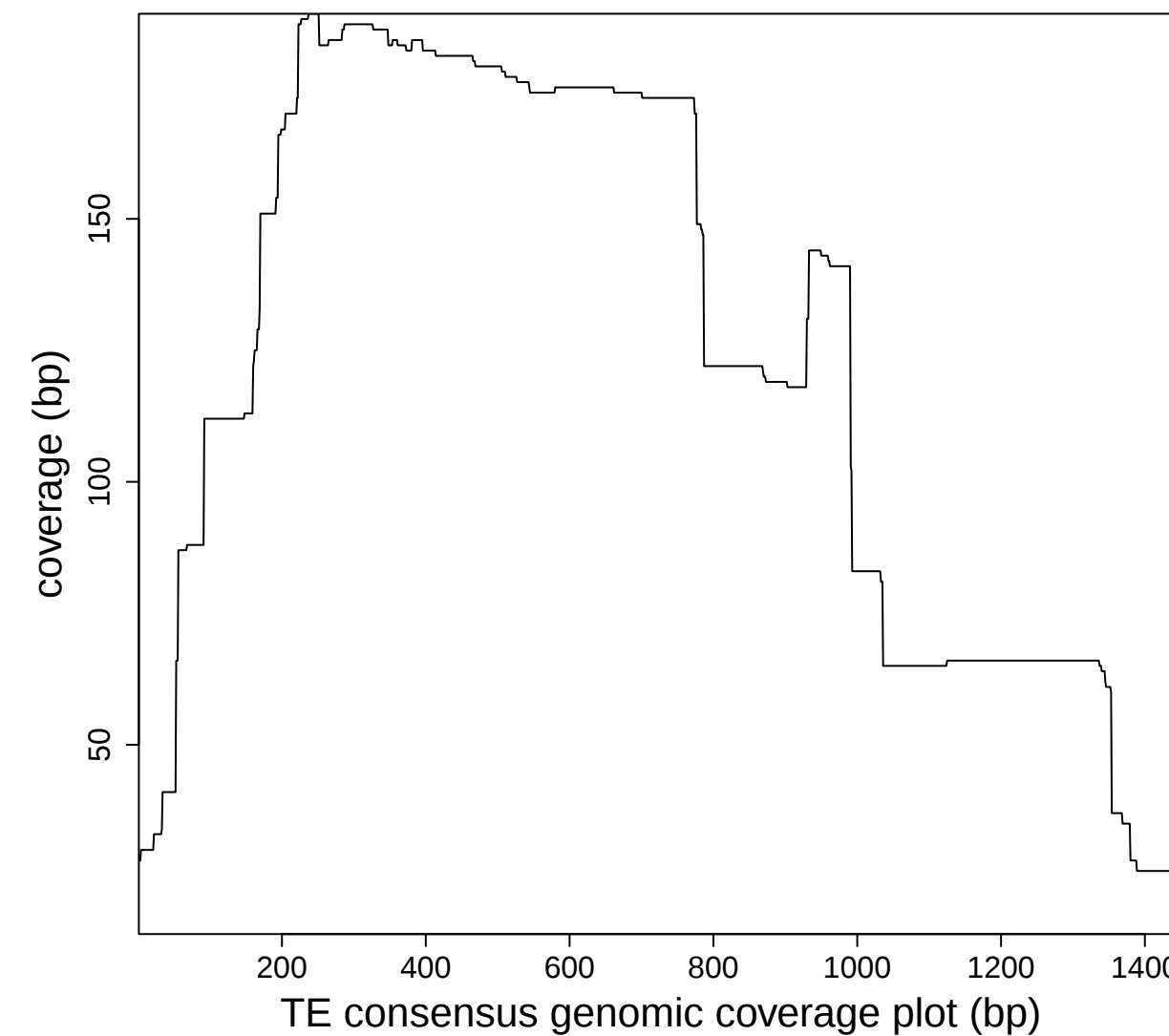
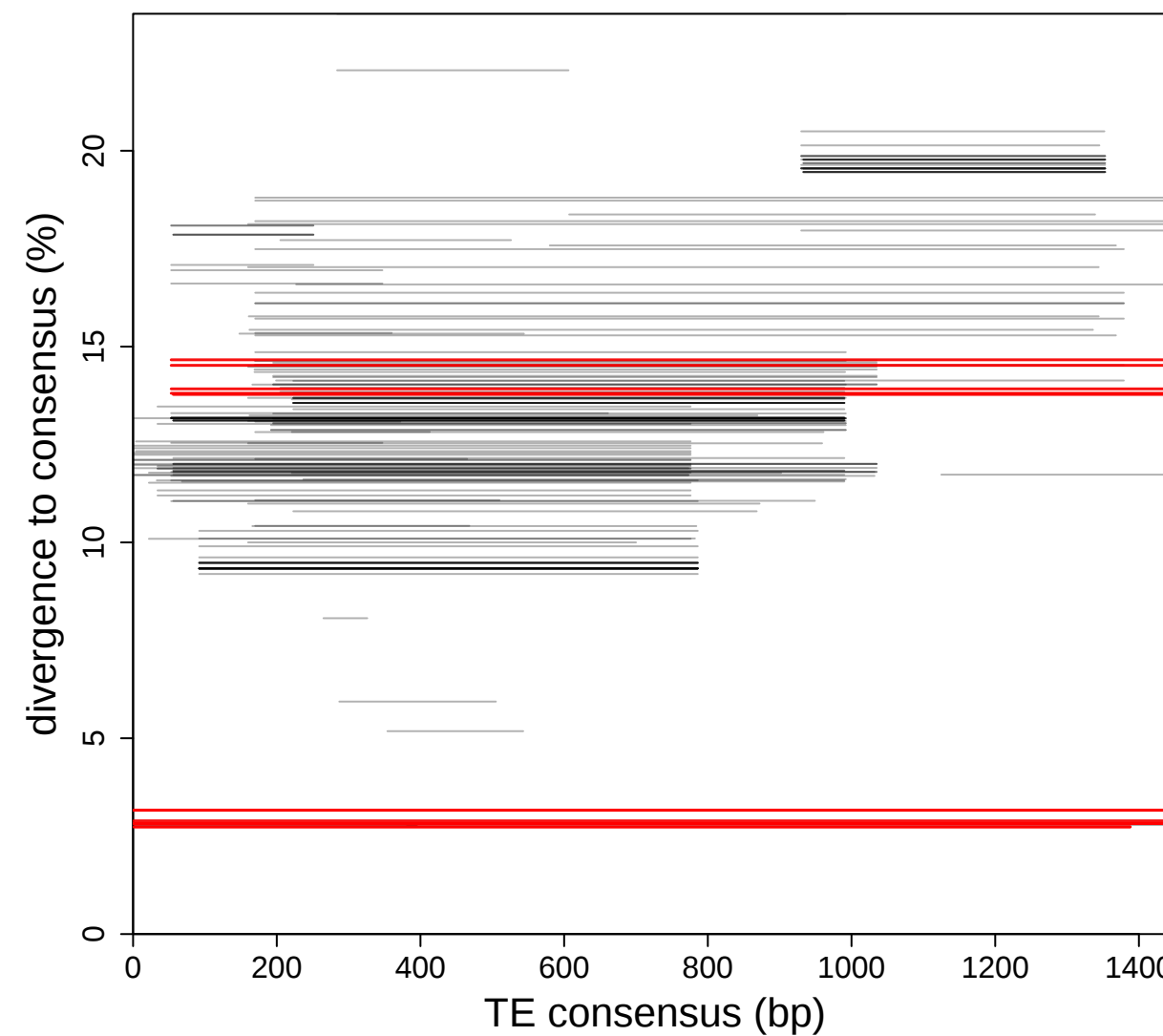


8.fasta.b.bed_uf.bed_g_5.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa
size: 8942bp; fragments: 1505; full length: 5 (>=8047.8bp)

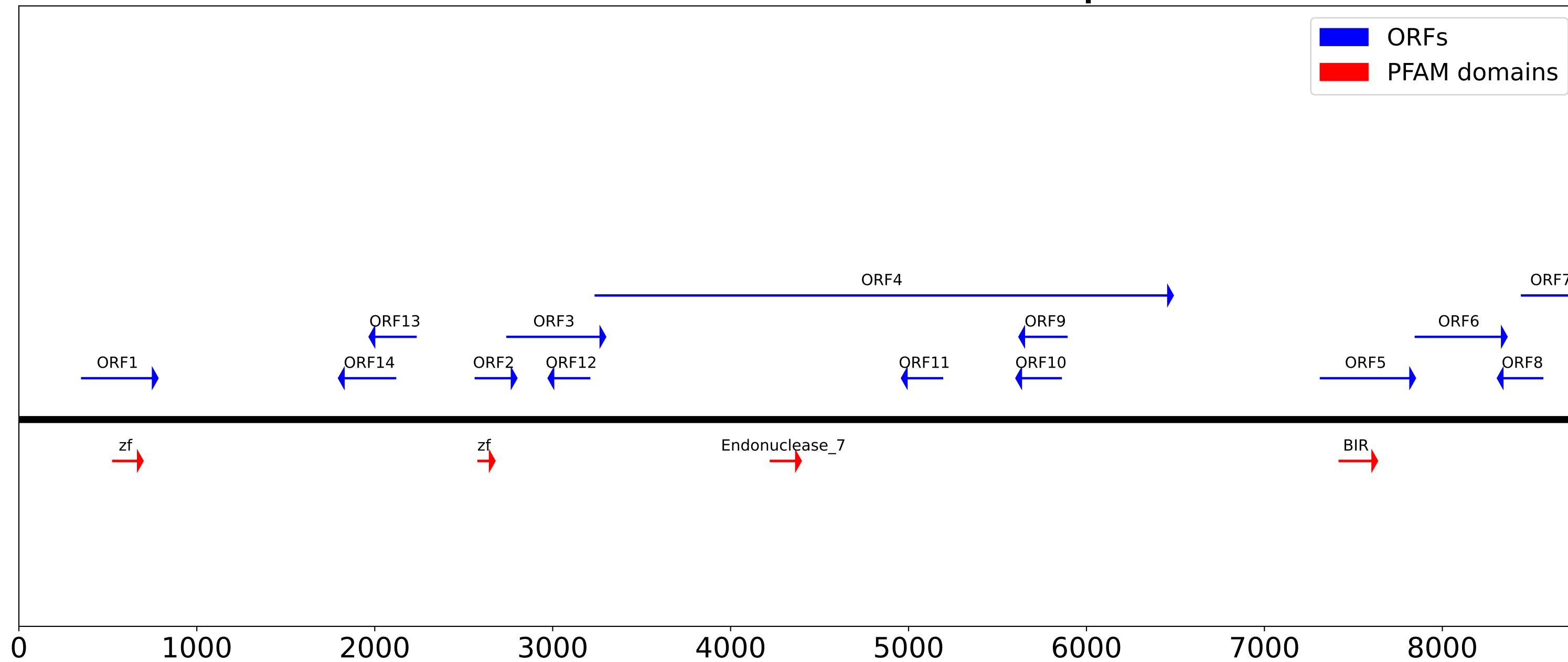
After TEtrimmer Extended plot Blue lines are boundaries



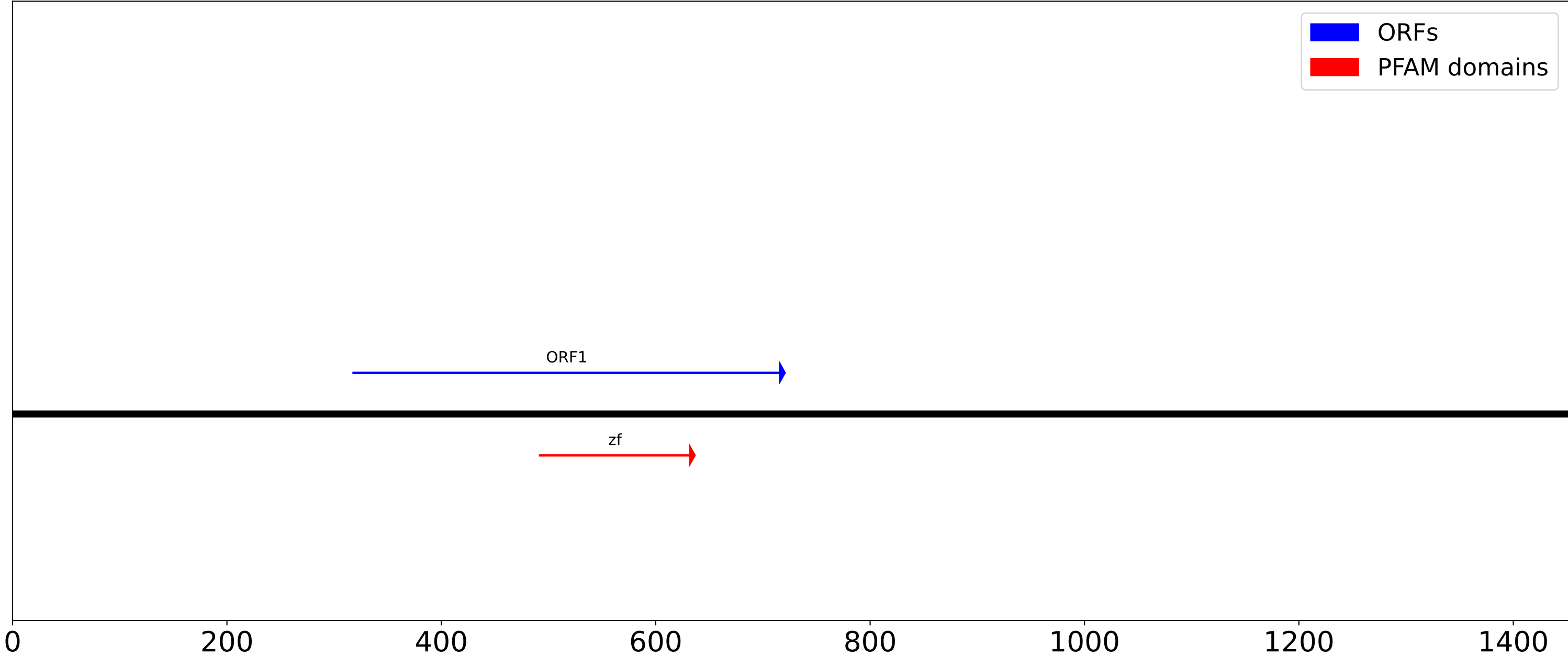
TE: rnd_5_family_588
size: 1453bp; fragments: 225; full length: 19 (≥ 1307.7 bp)



After TEtrimmer ORF and PFAM domain plot



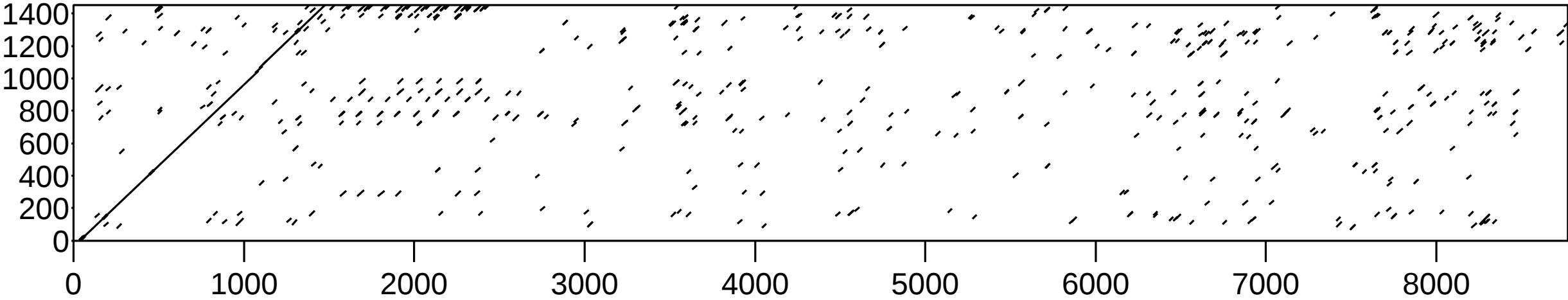
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)