

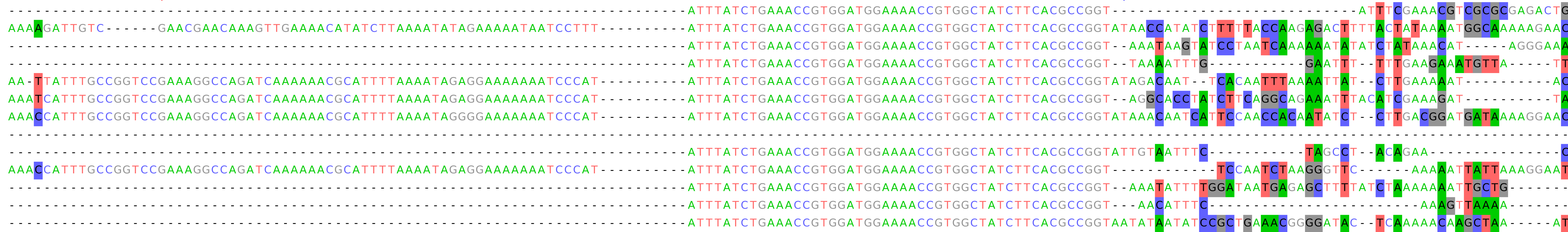
Start crop Point

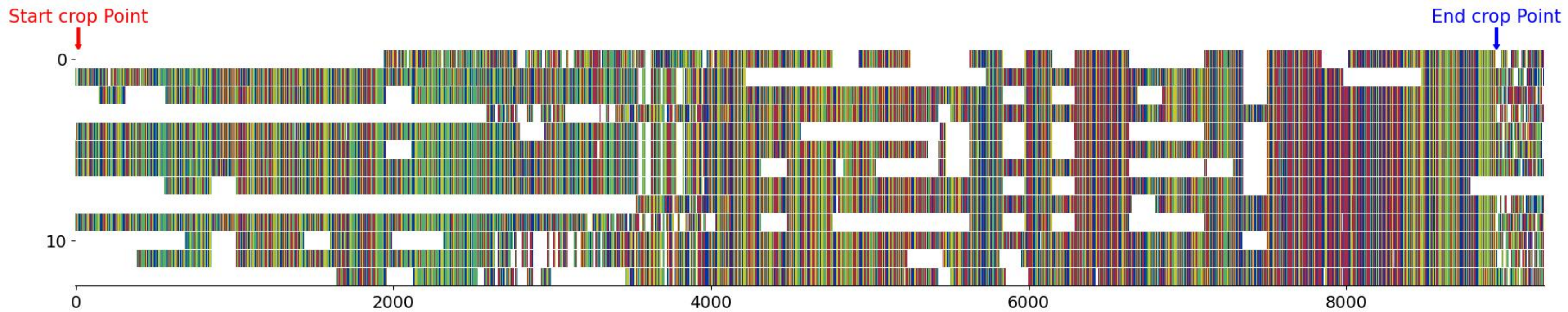


End crop Point



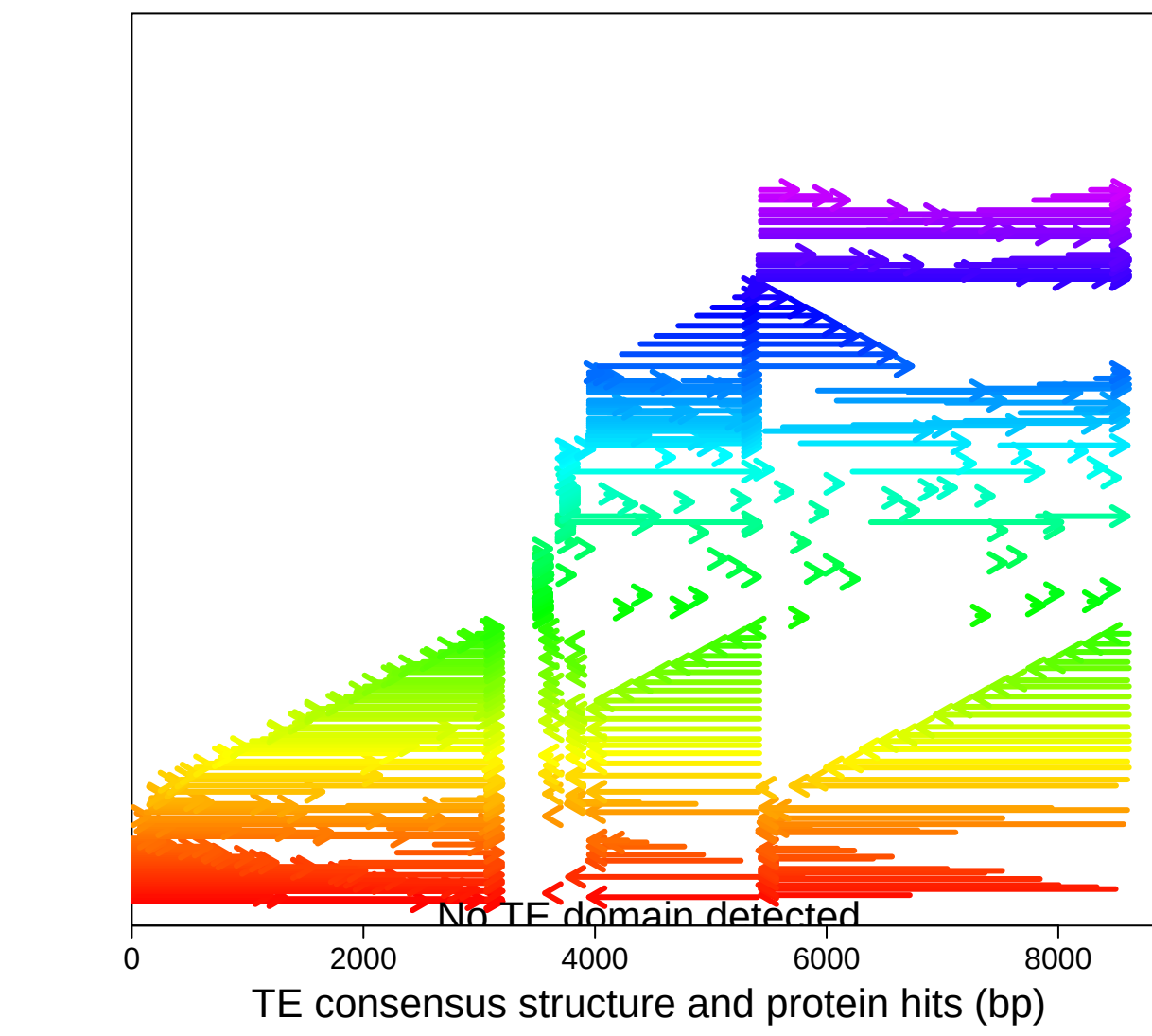
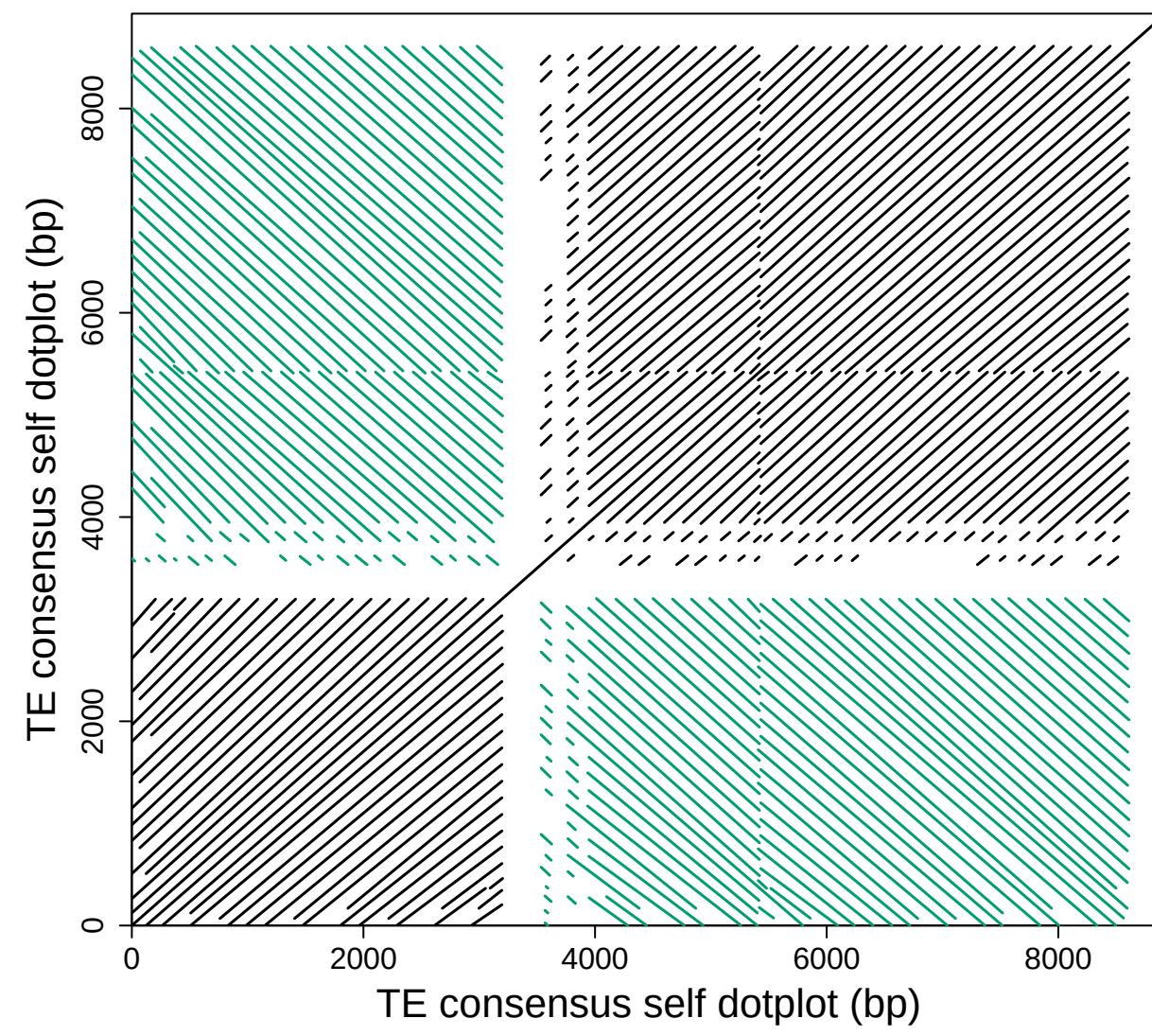
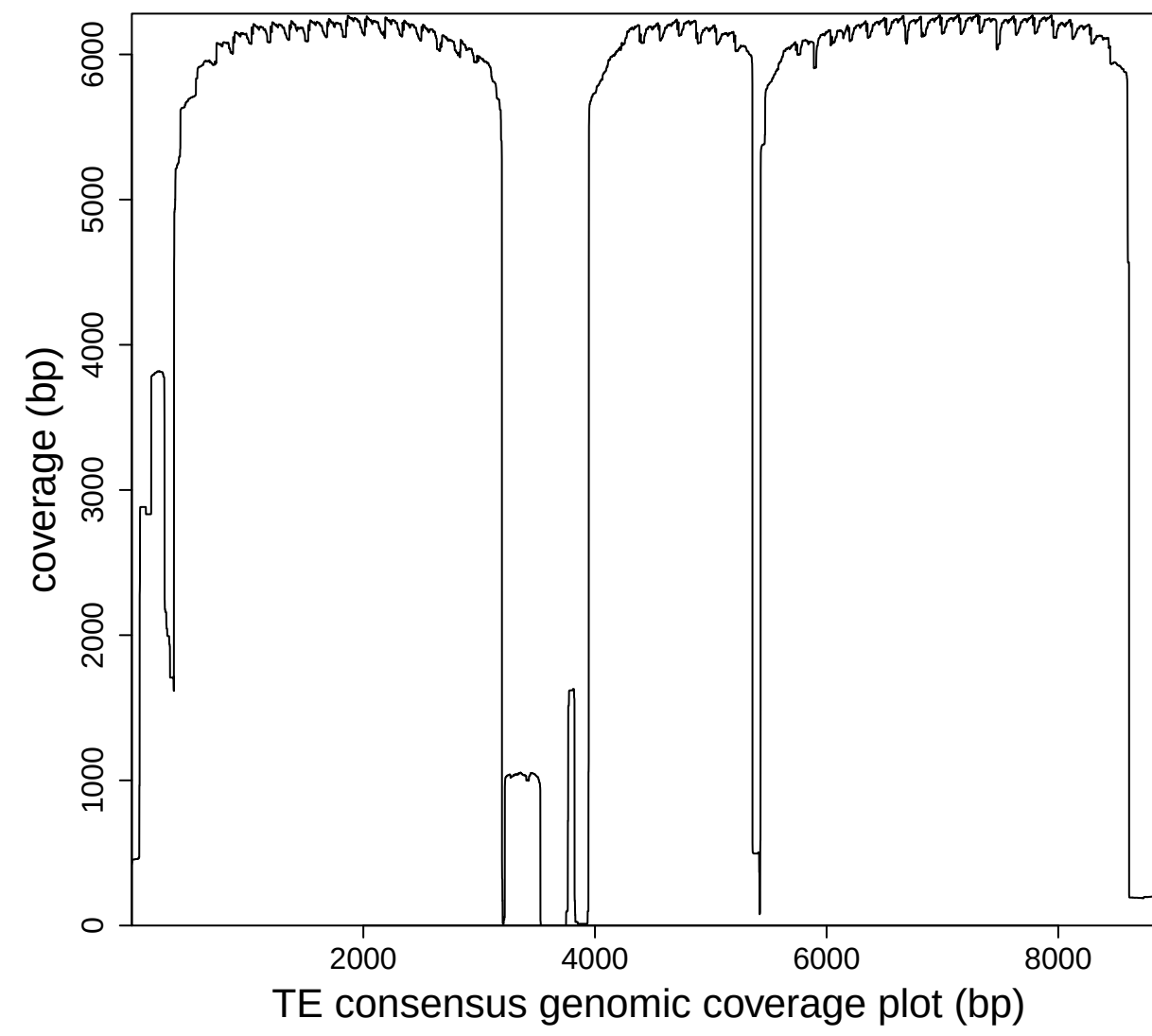
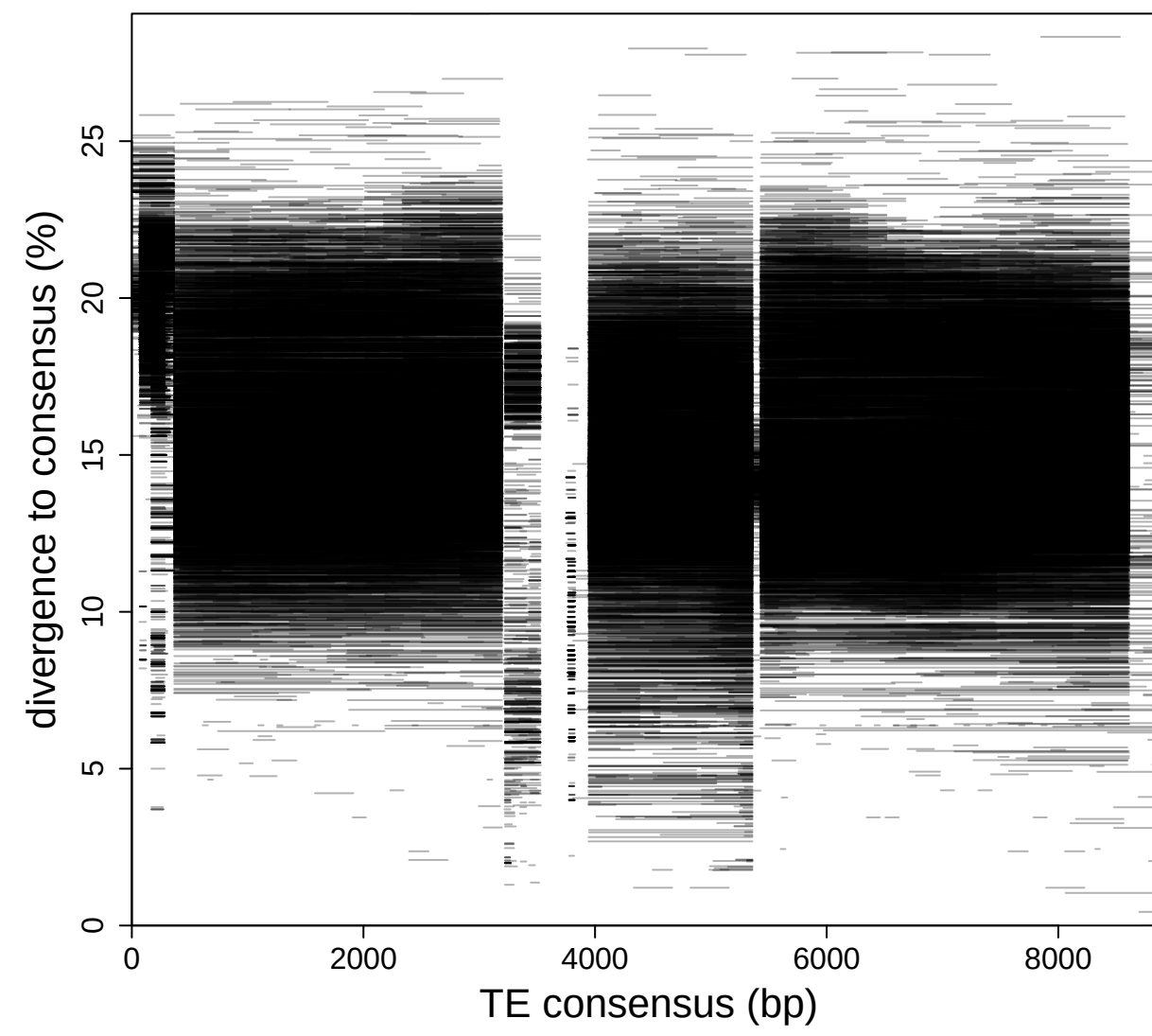
MSA length = 8929





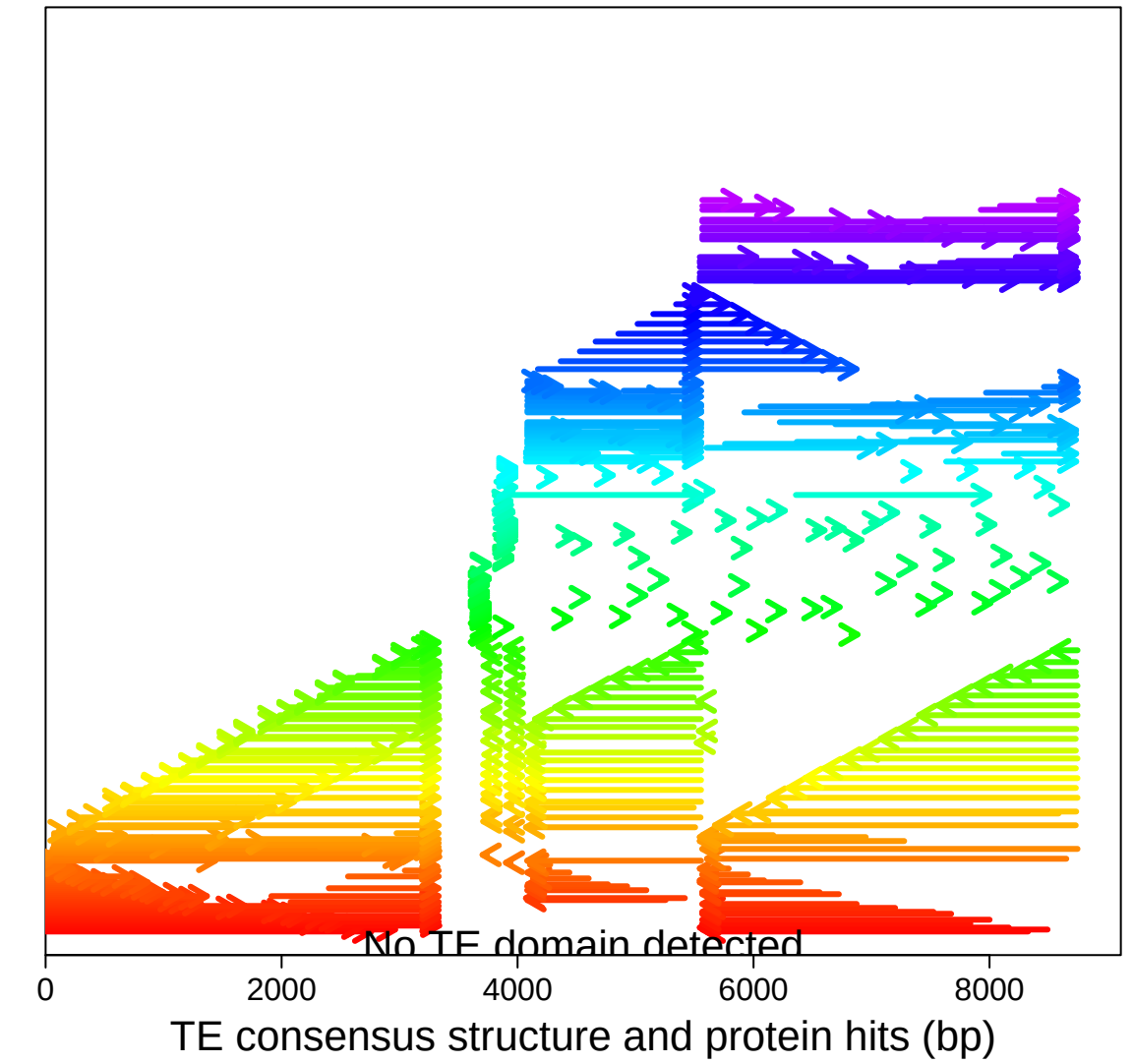
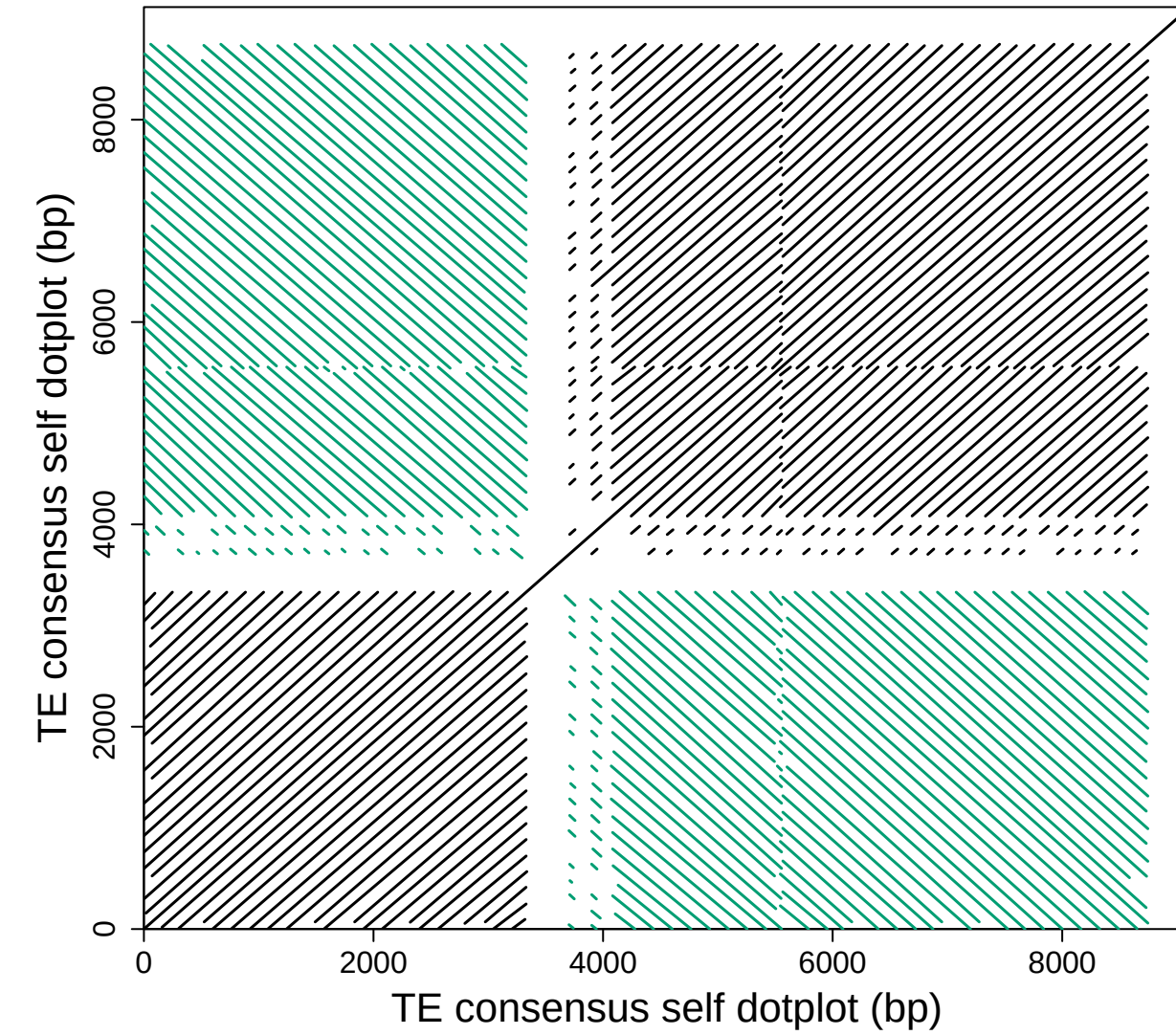
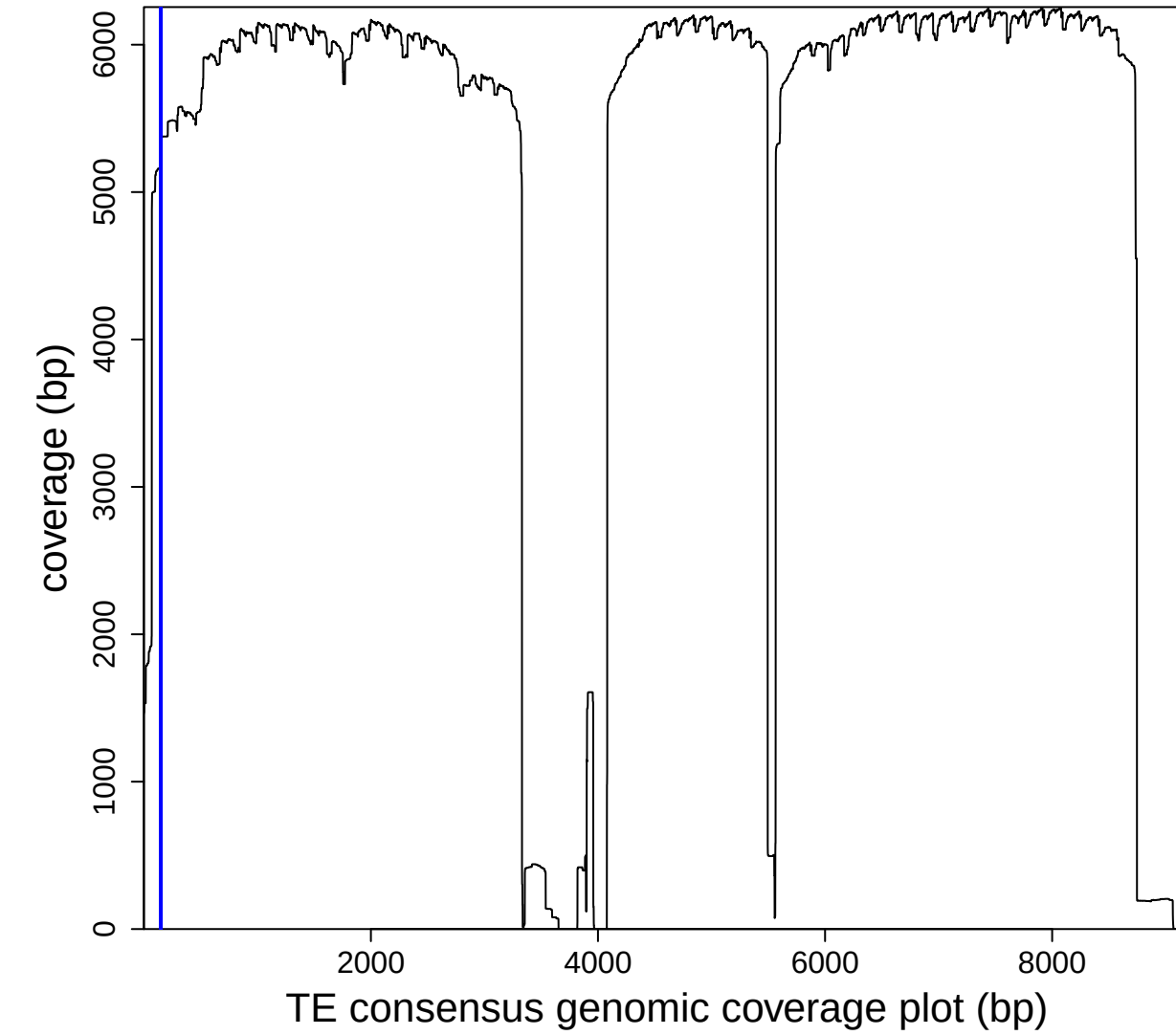
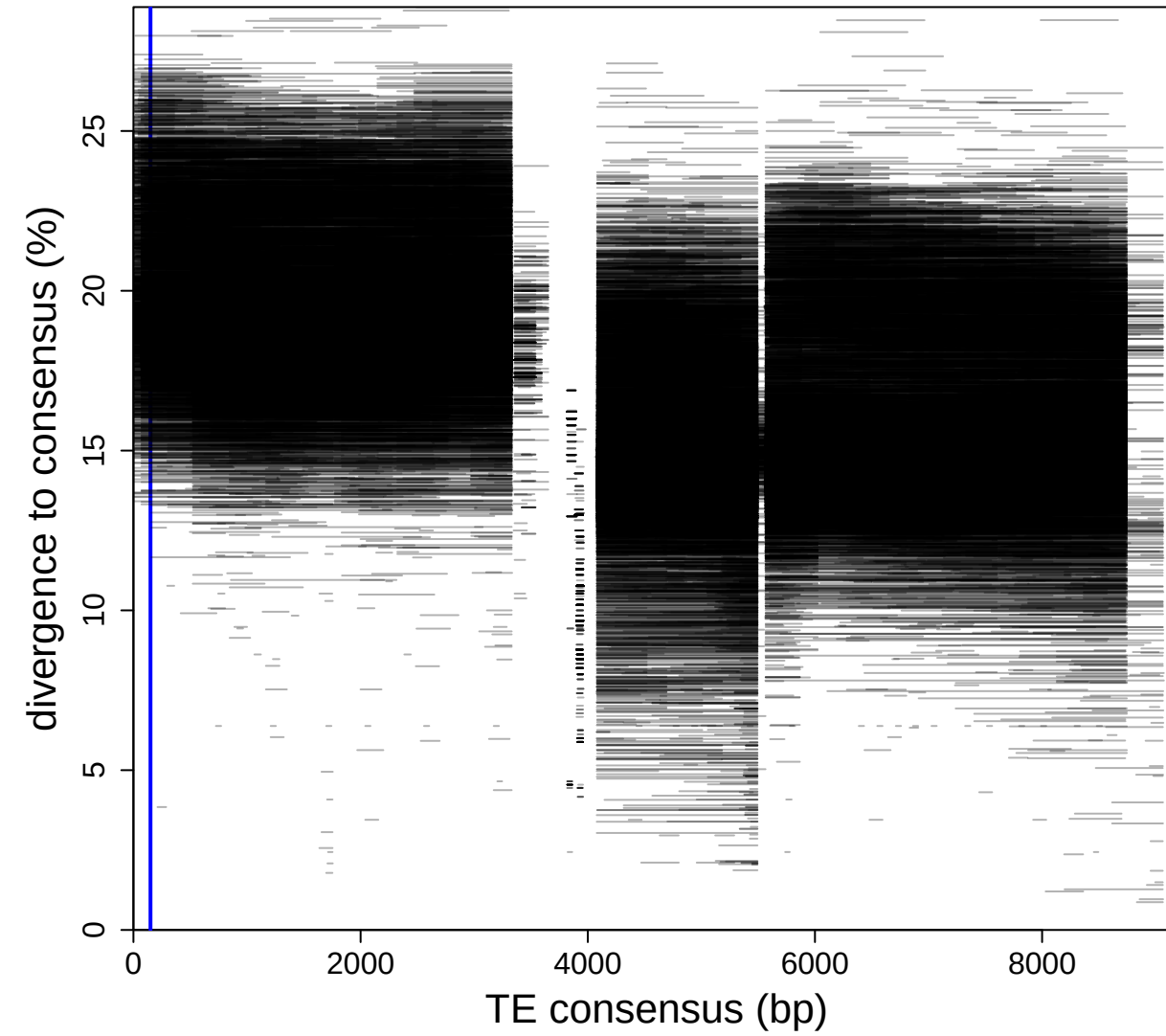
a.b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.f
size: 8929bp; fragments: 34963; full length: 0 (>=8036.1bp)

After TEtrimmer 8929 bp

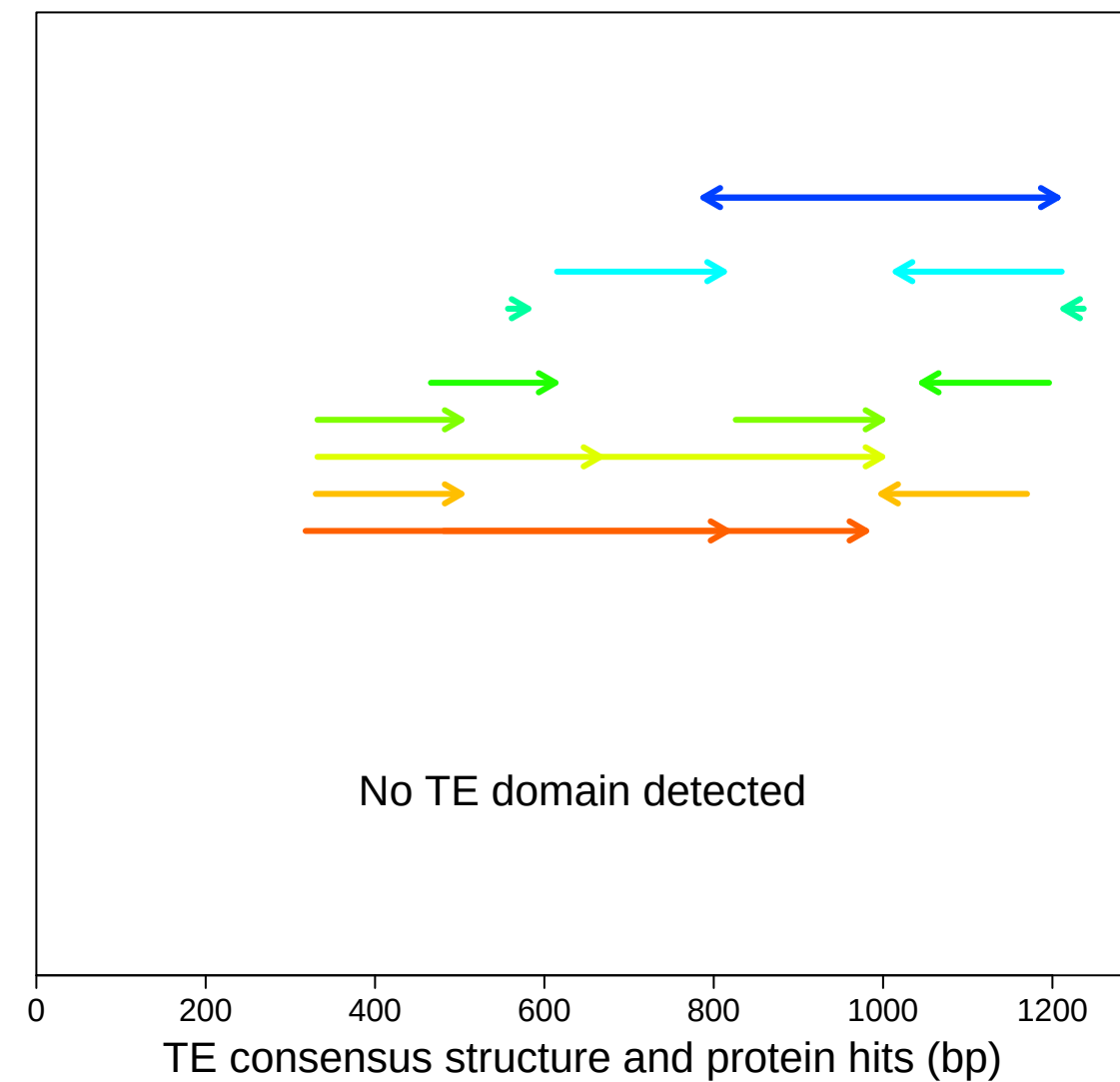
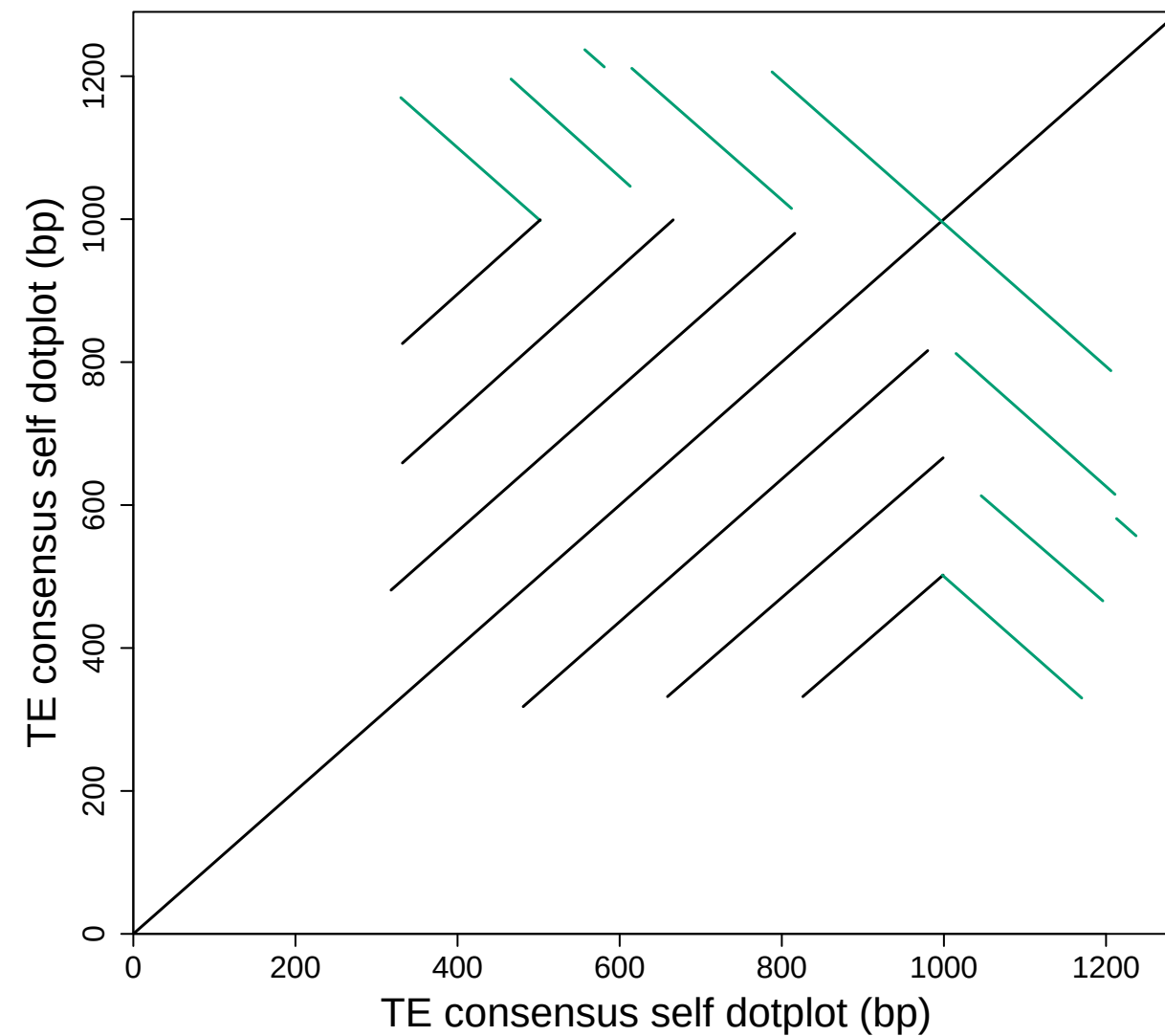
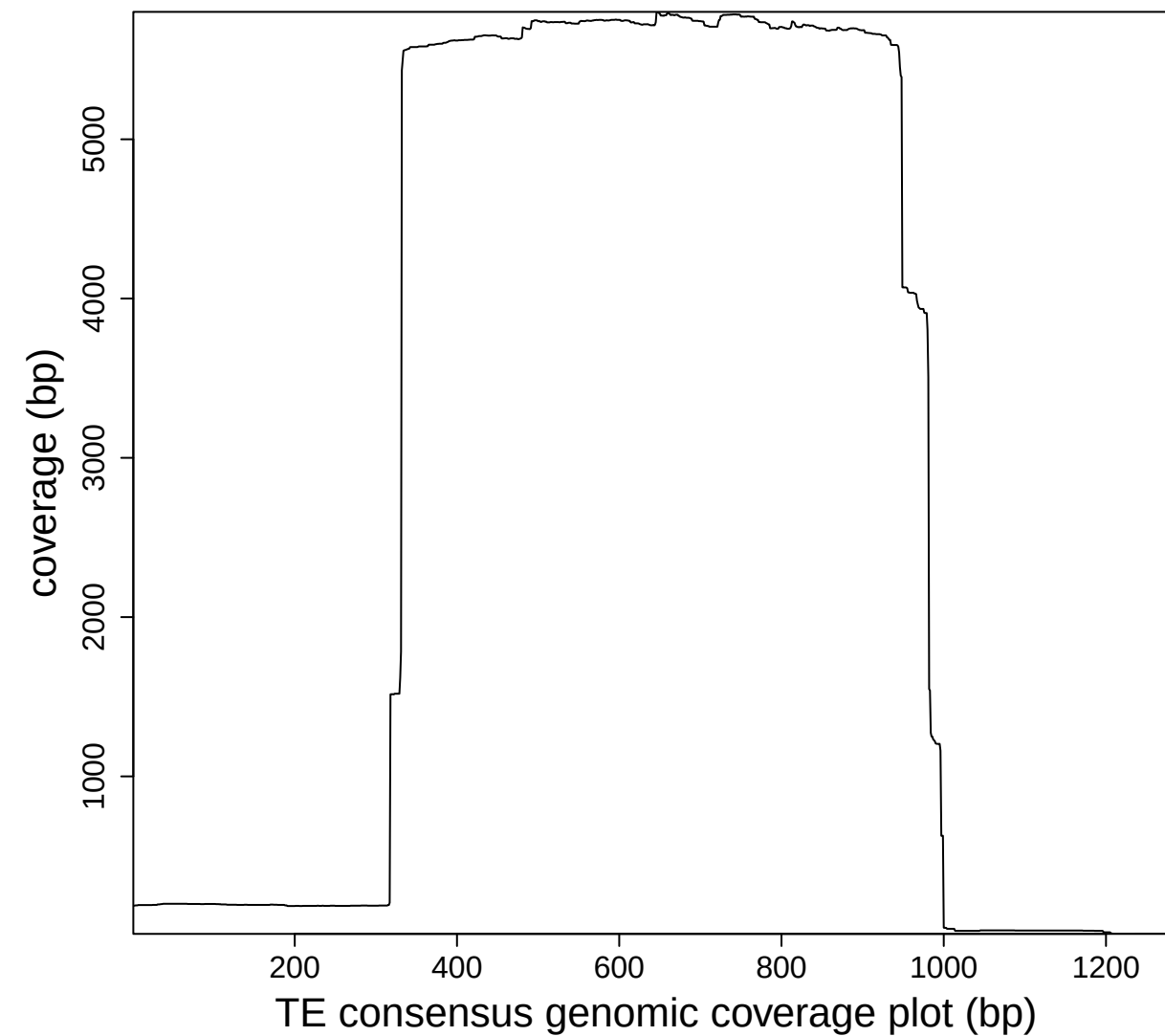
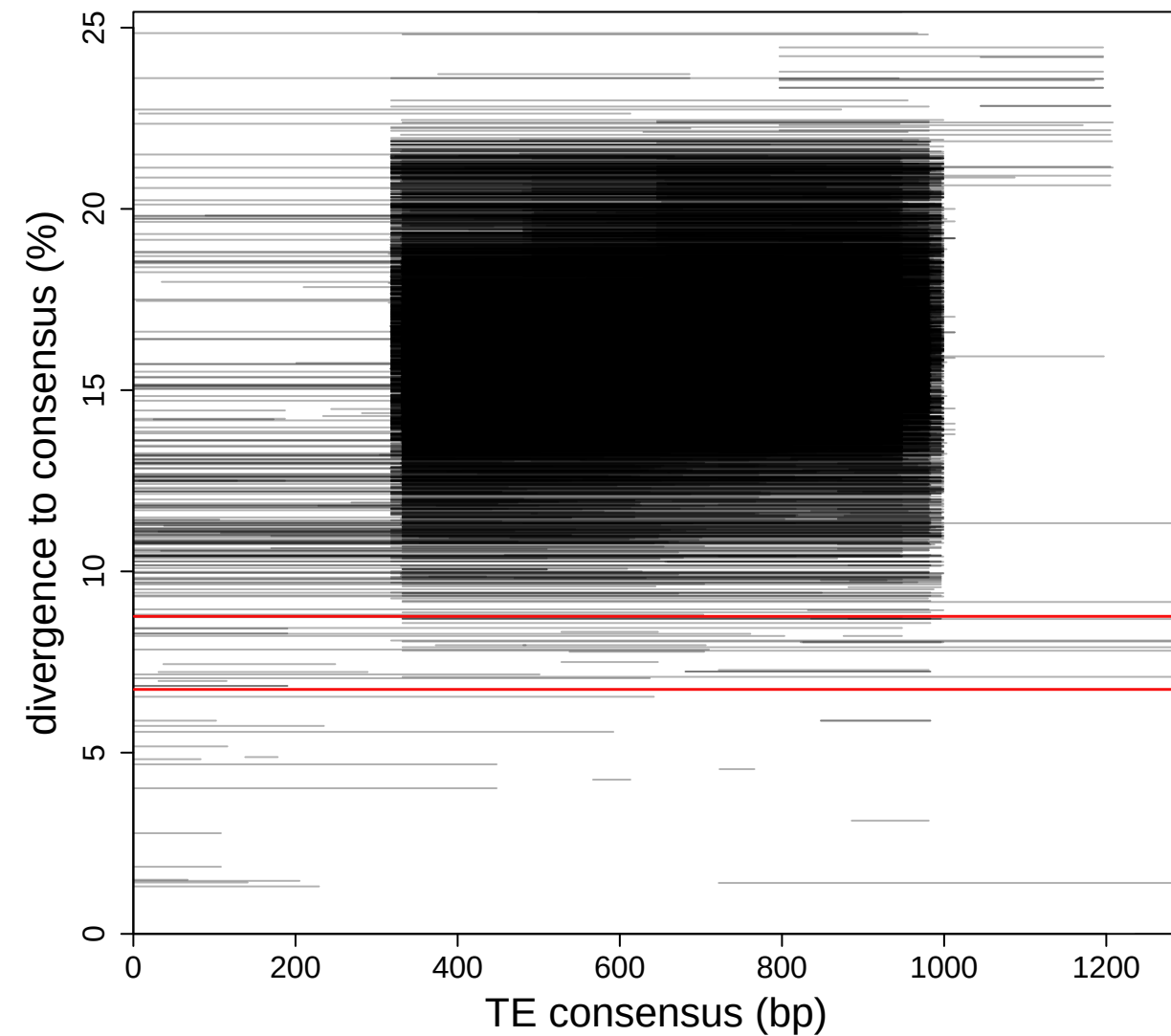


asta.b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_g
size: 9214bp; fragments: 30692; full length: 0 (>=8292.6bp)

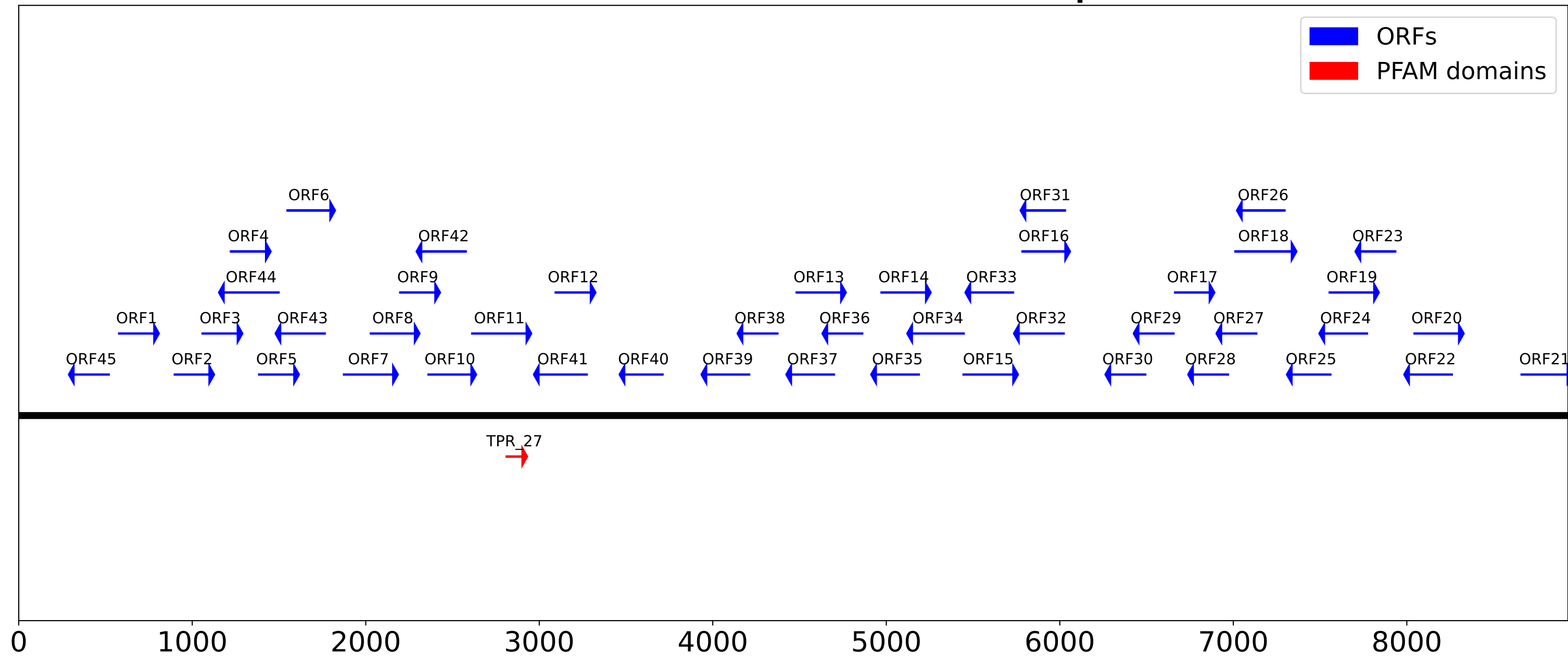
After TEtrimmer Extended plot Blue lines are boundaries



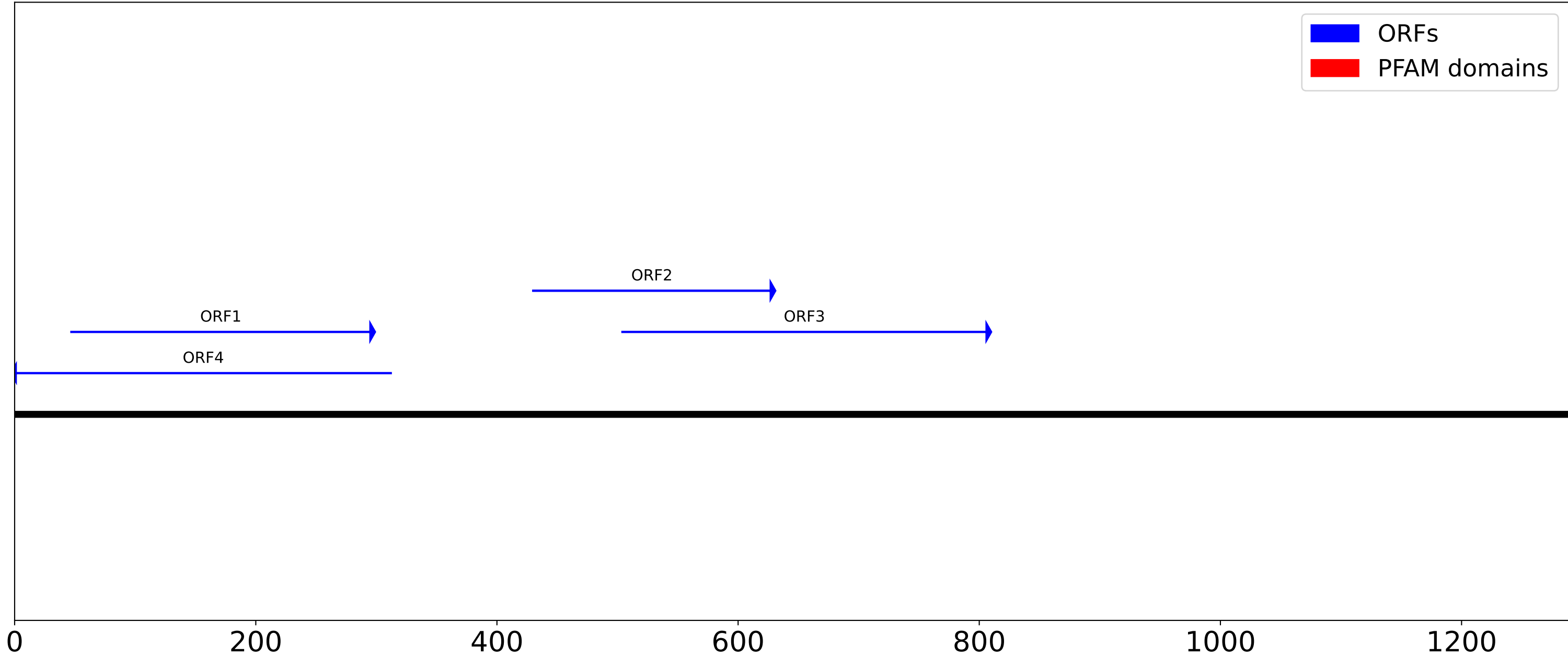
TE: rnd_5_family_8603
size: 1290bp; fragments: 6250; full length: 2 (≥ 1161 bp)



After TEtrimmer ORF and PFAM domain plot



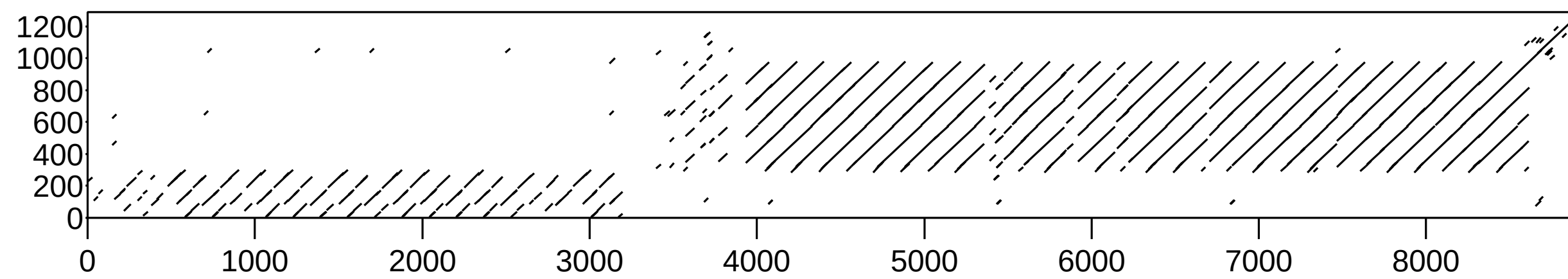
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

isensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)