

Start crop Point



End crop Point

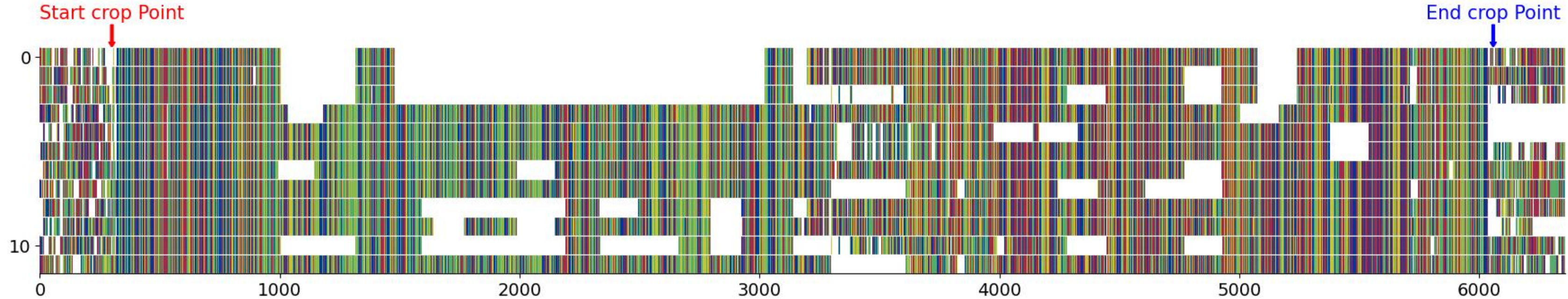


MSA length = 5760



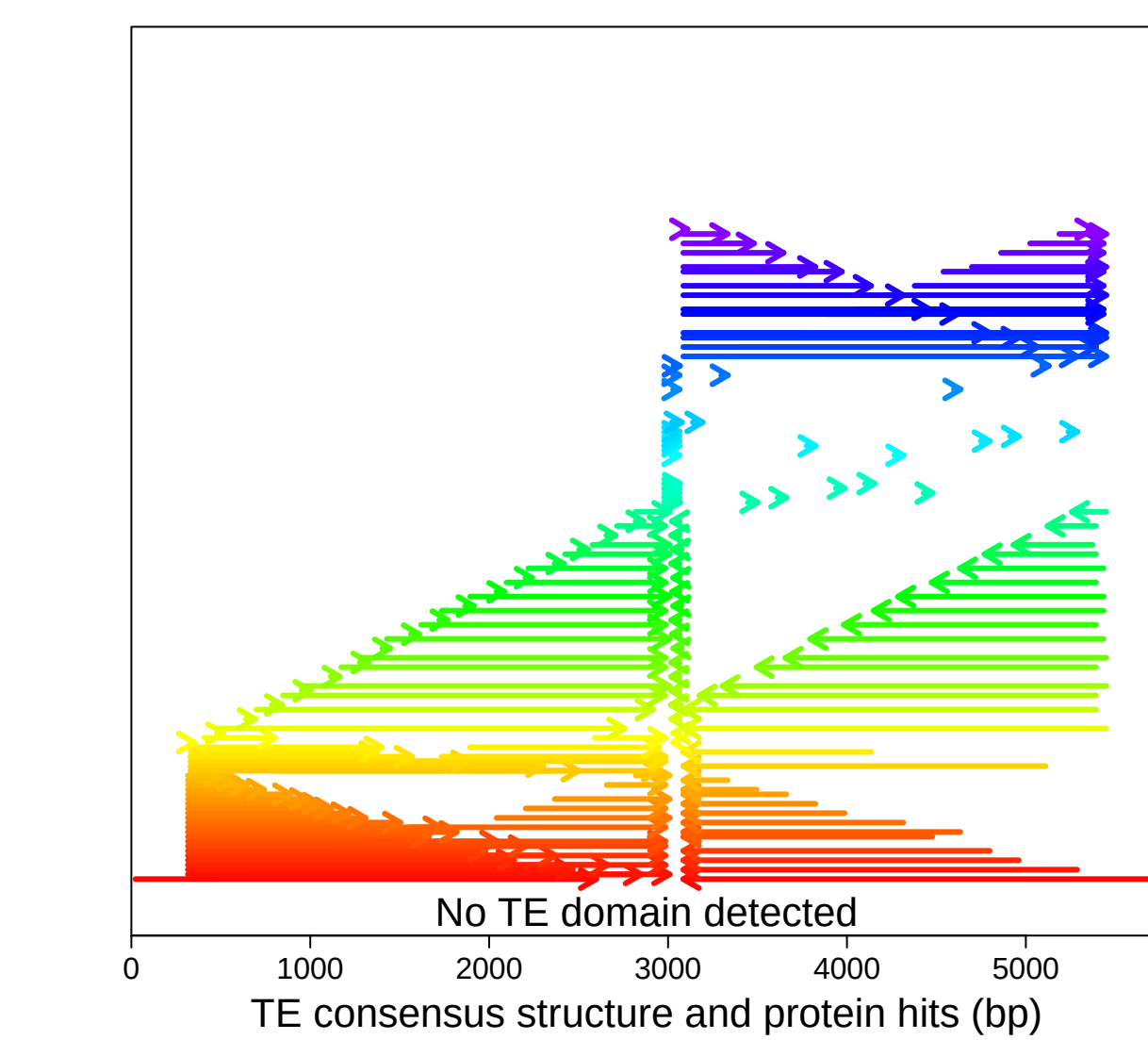
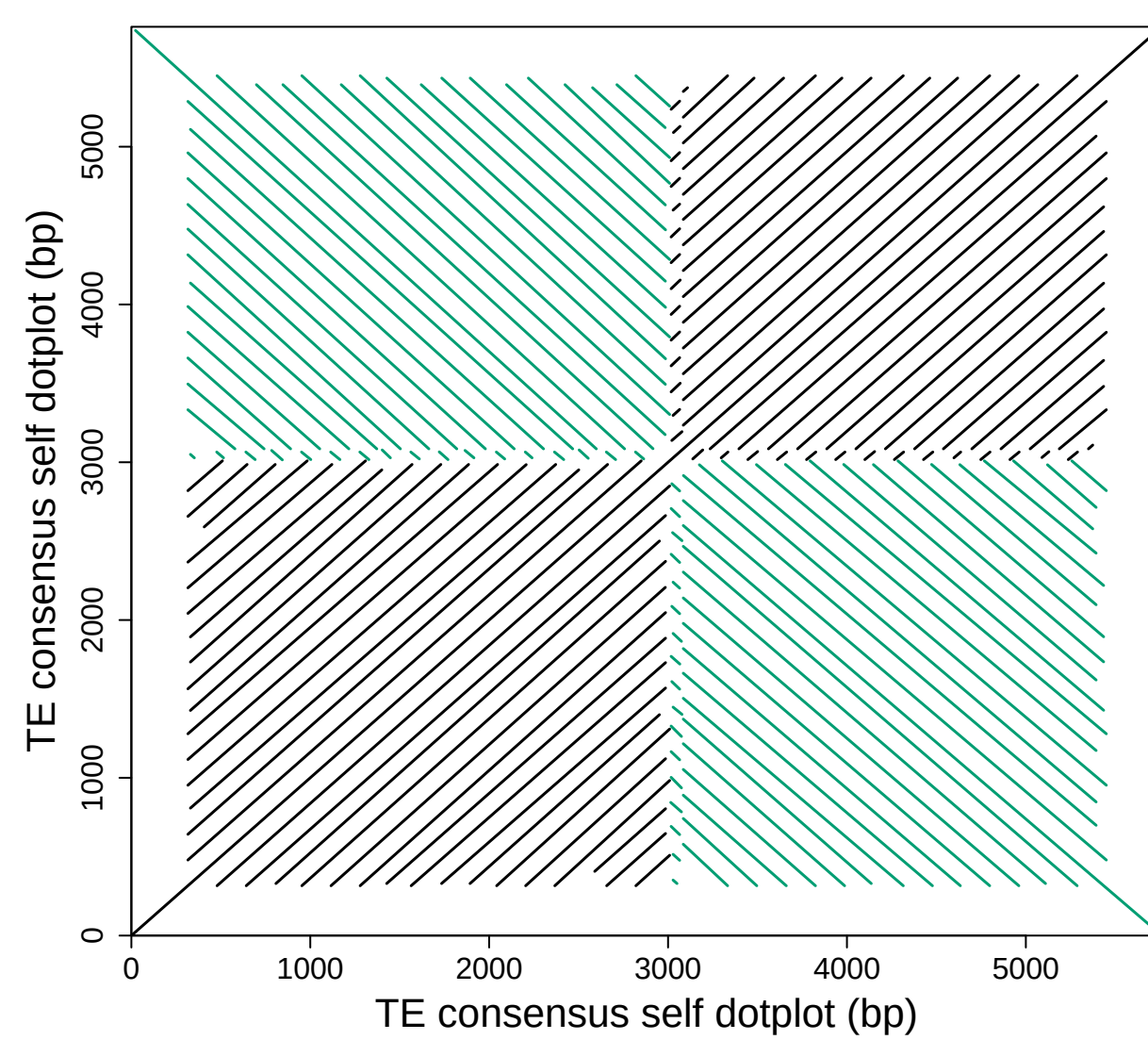
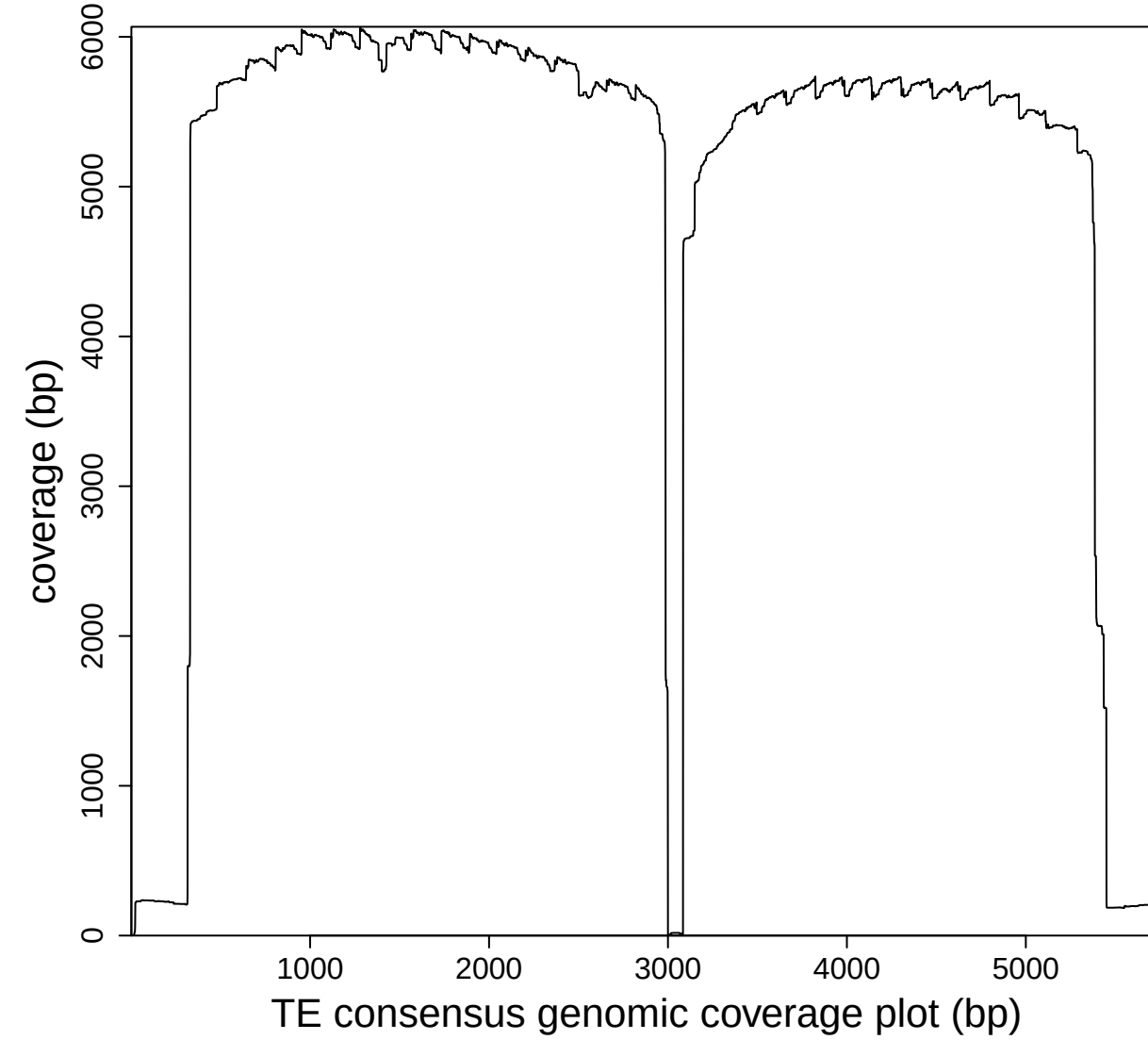
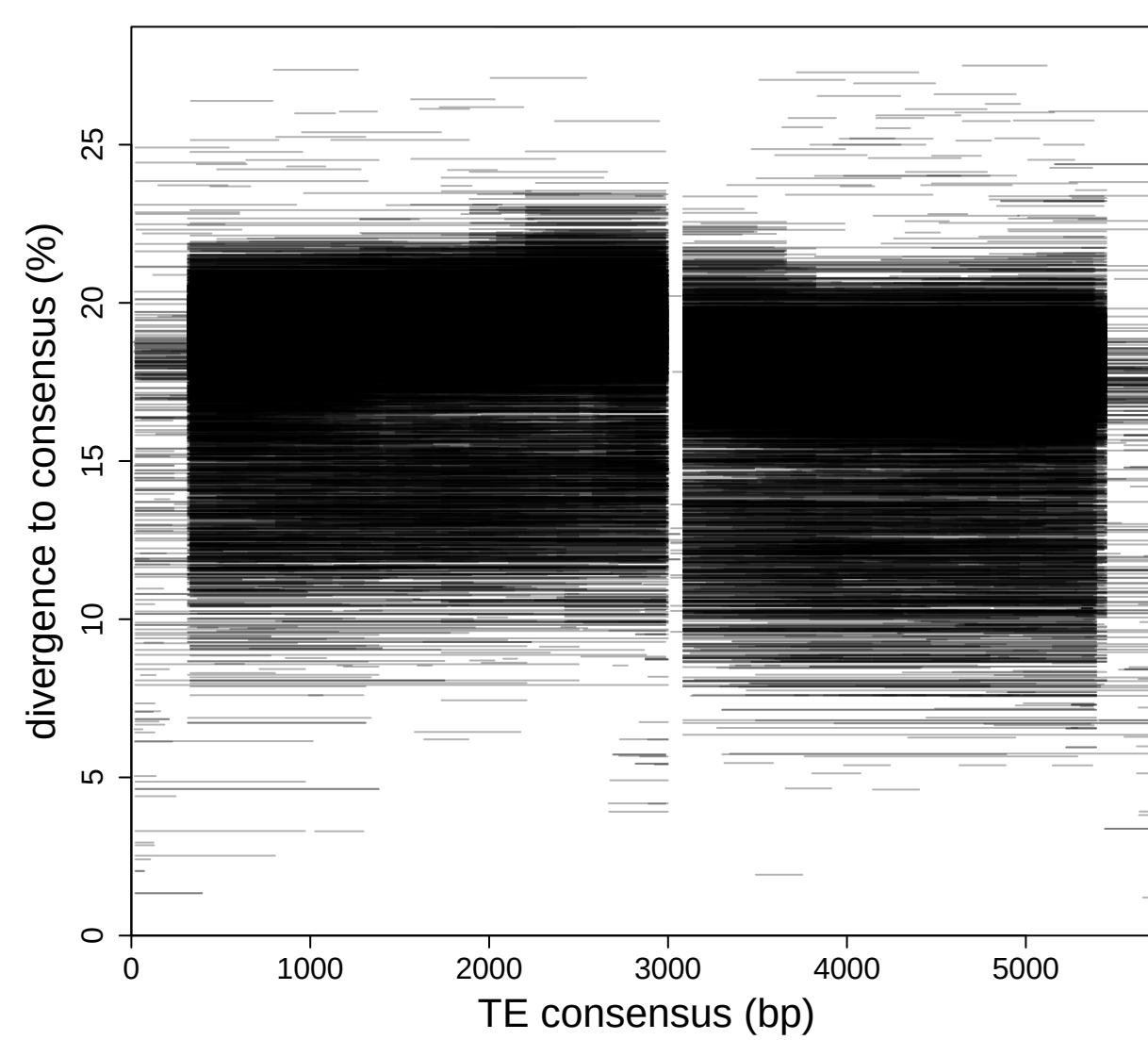
Start crop Point

End crop Point



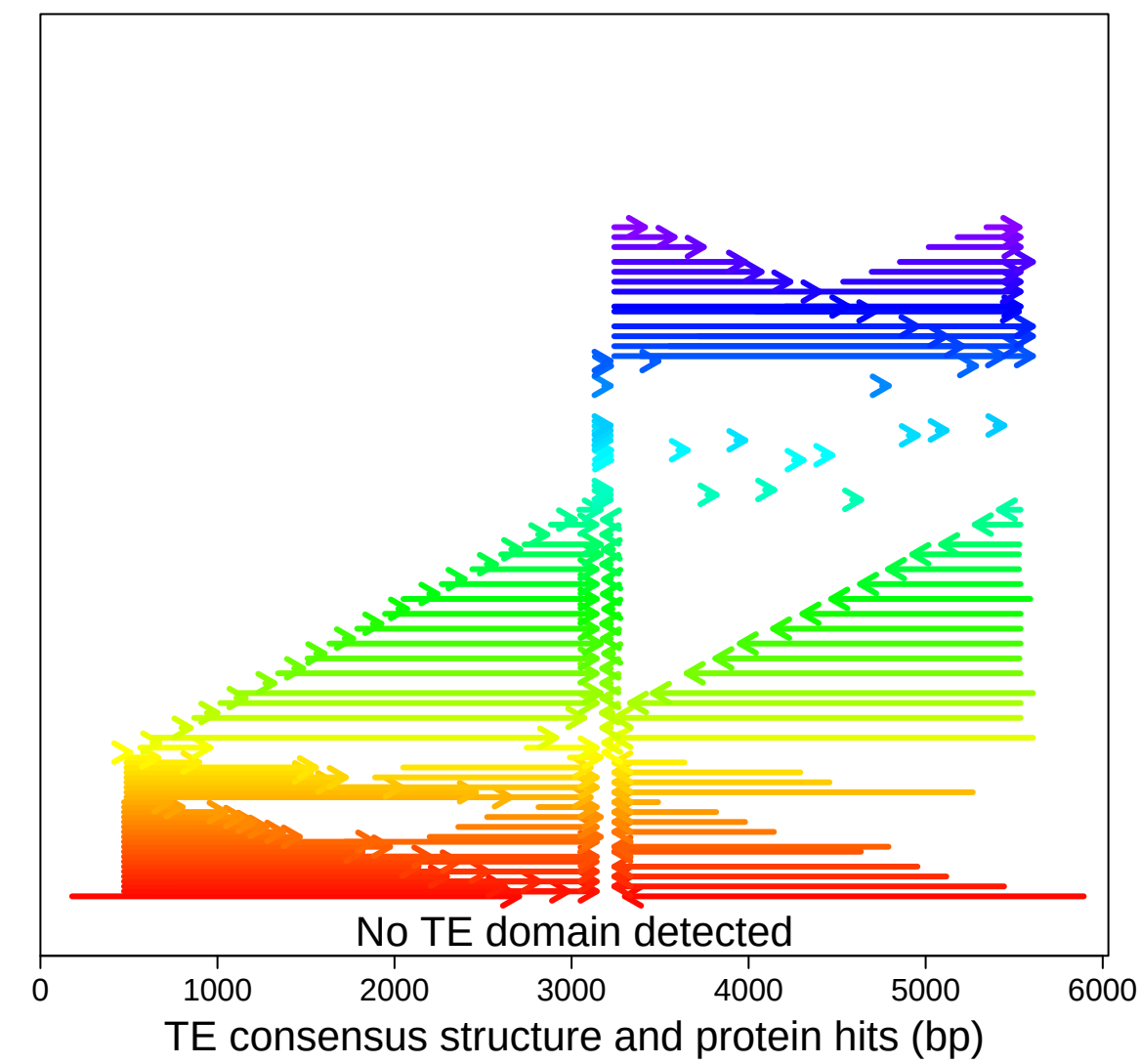
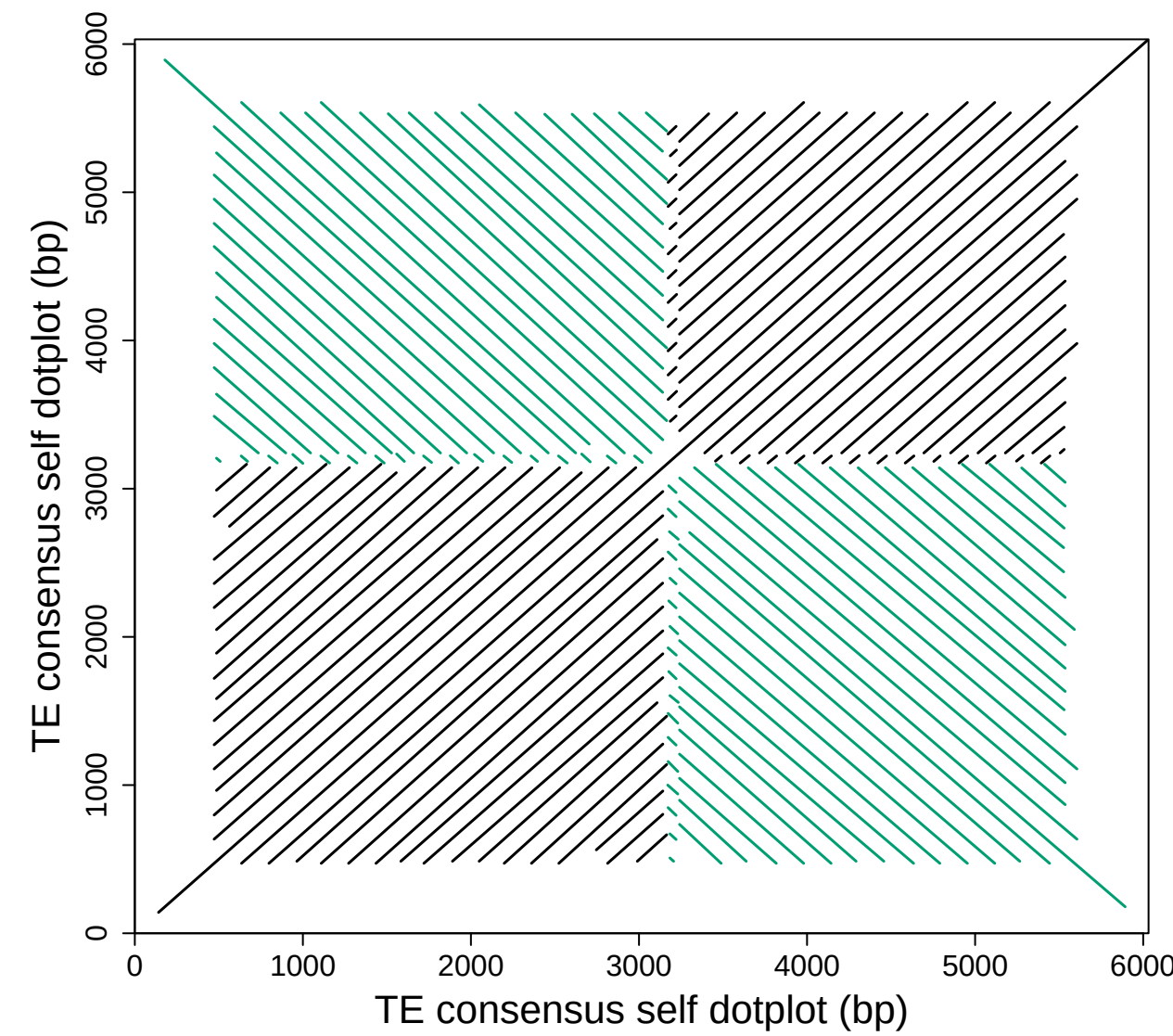
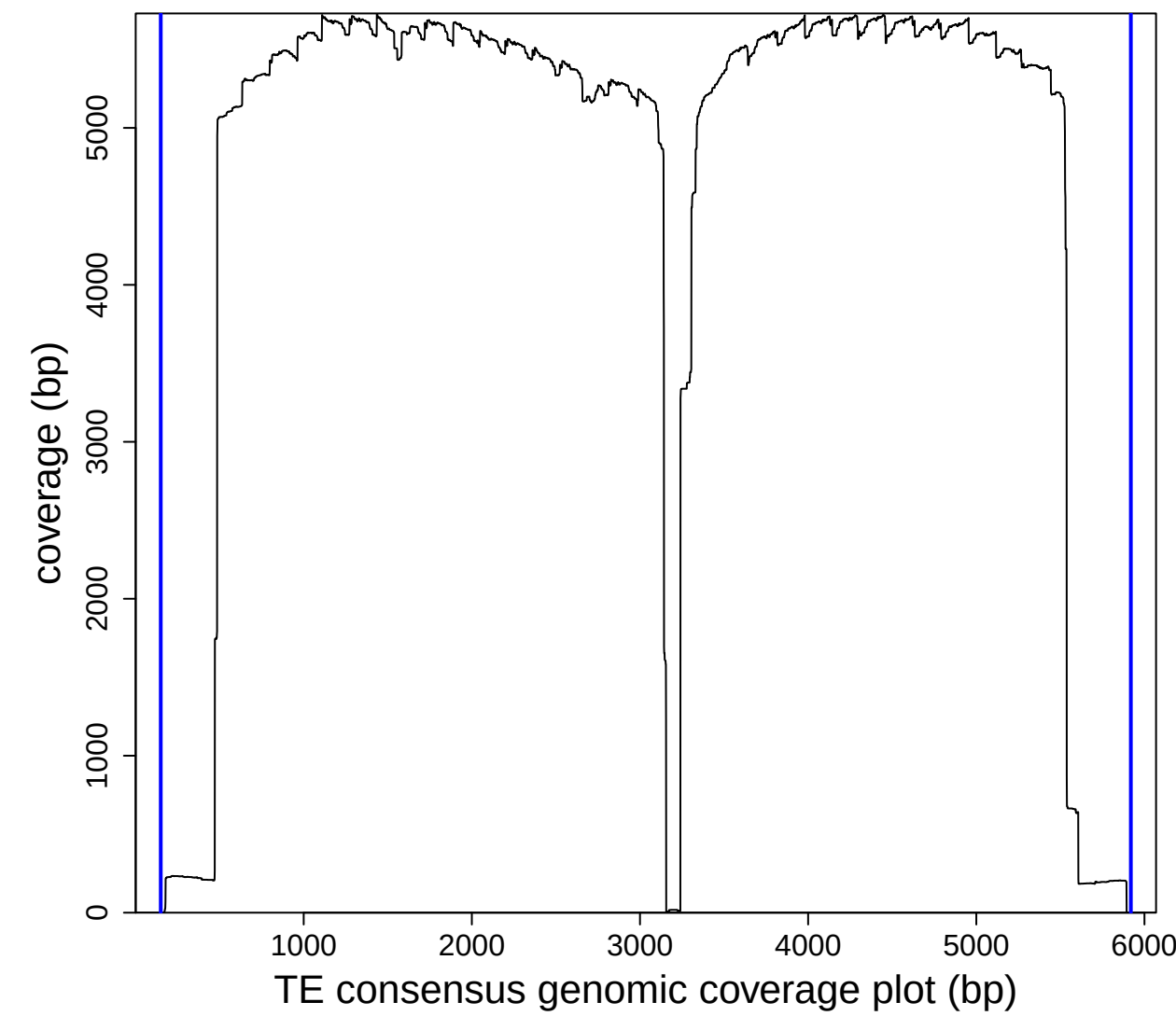
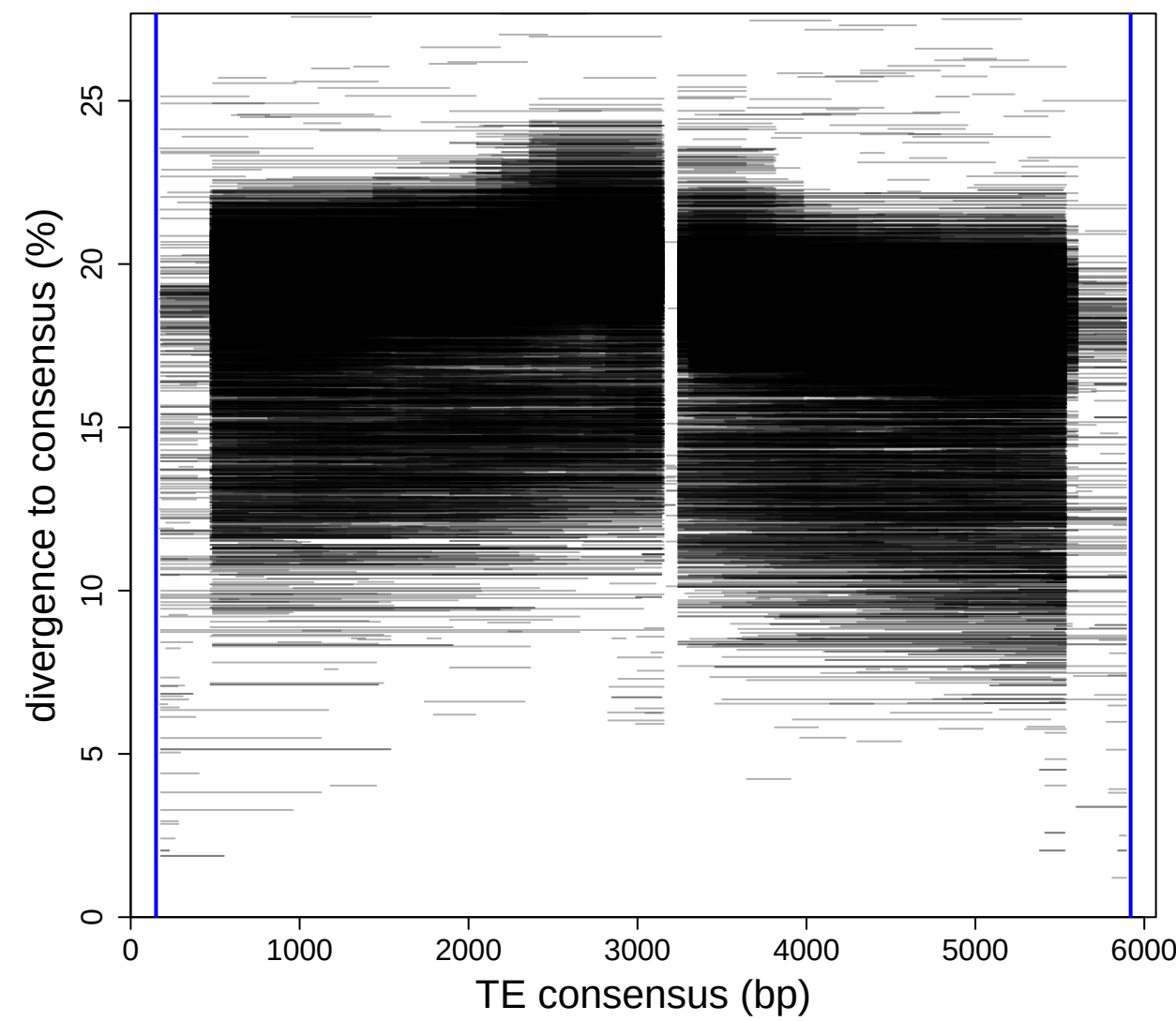
a.b.bed_uf.bed_g_3.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.f
size: 5760bp; fragments: 17202; full length: 0 (>=5184bp)

After TEtrimmer 5760 bp

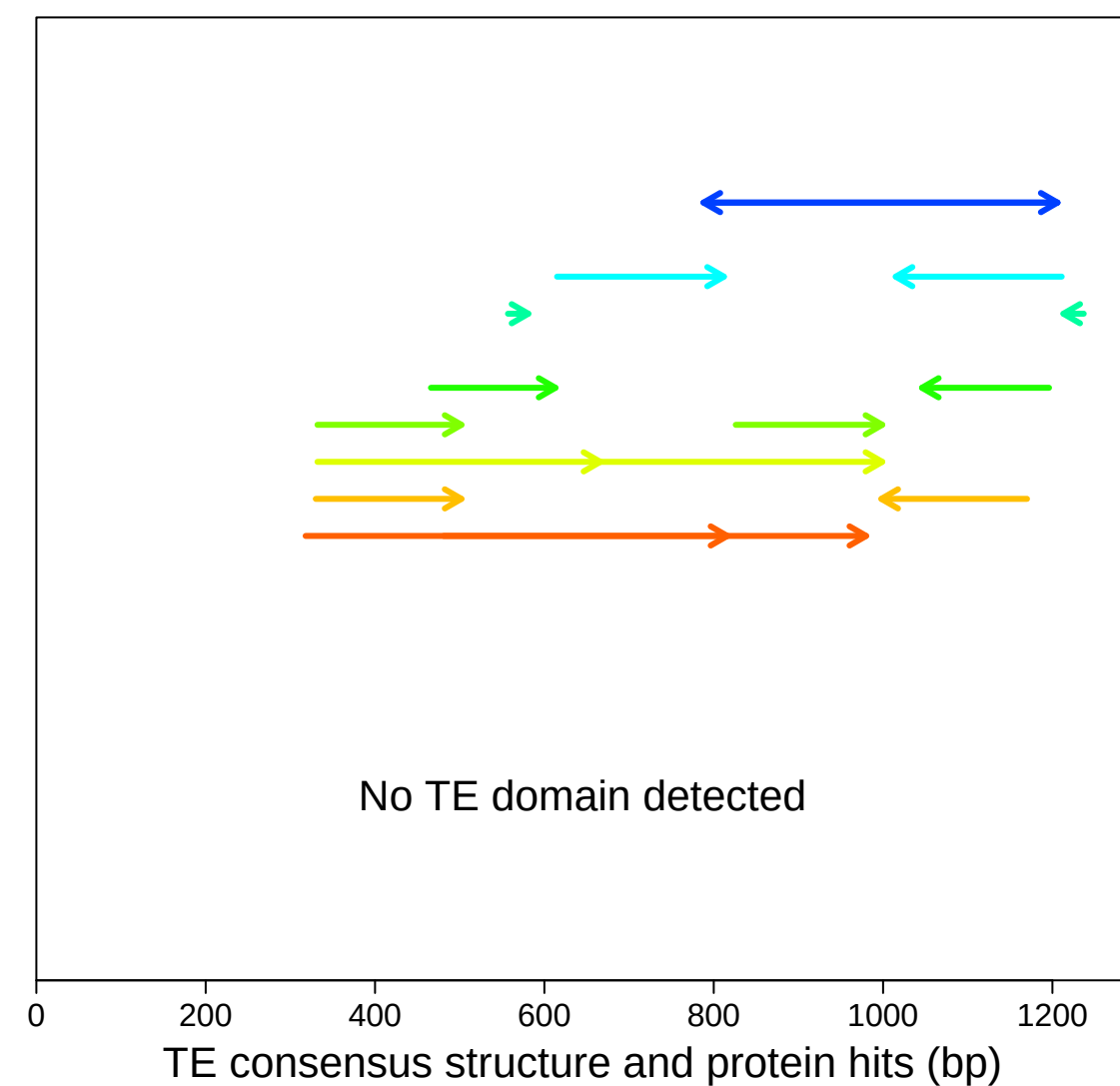
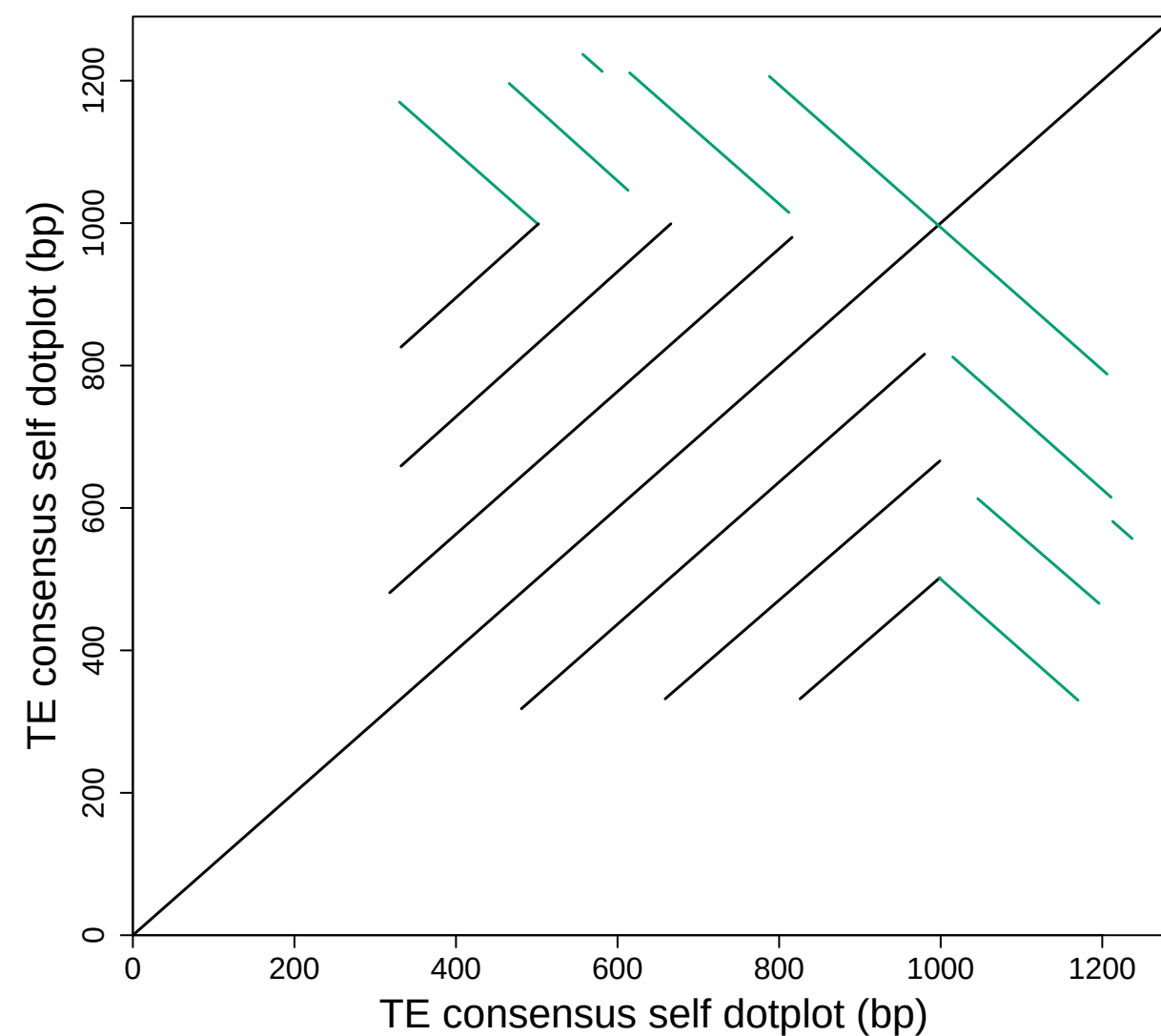
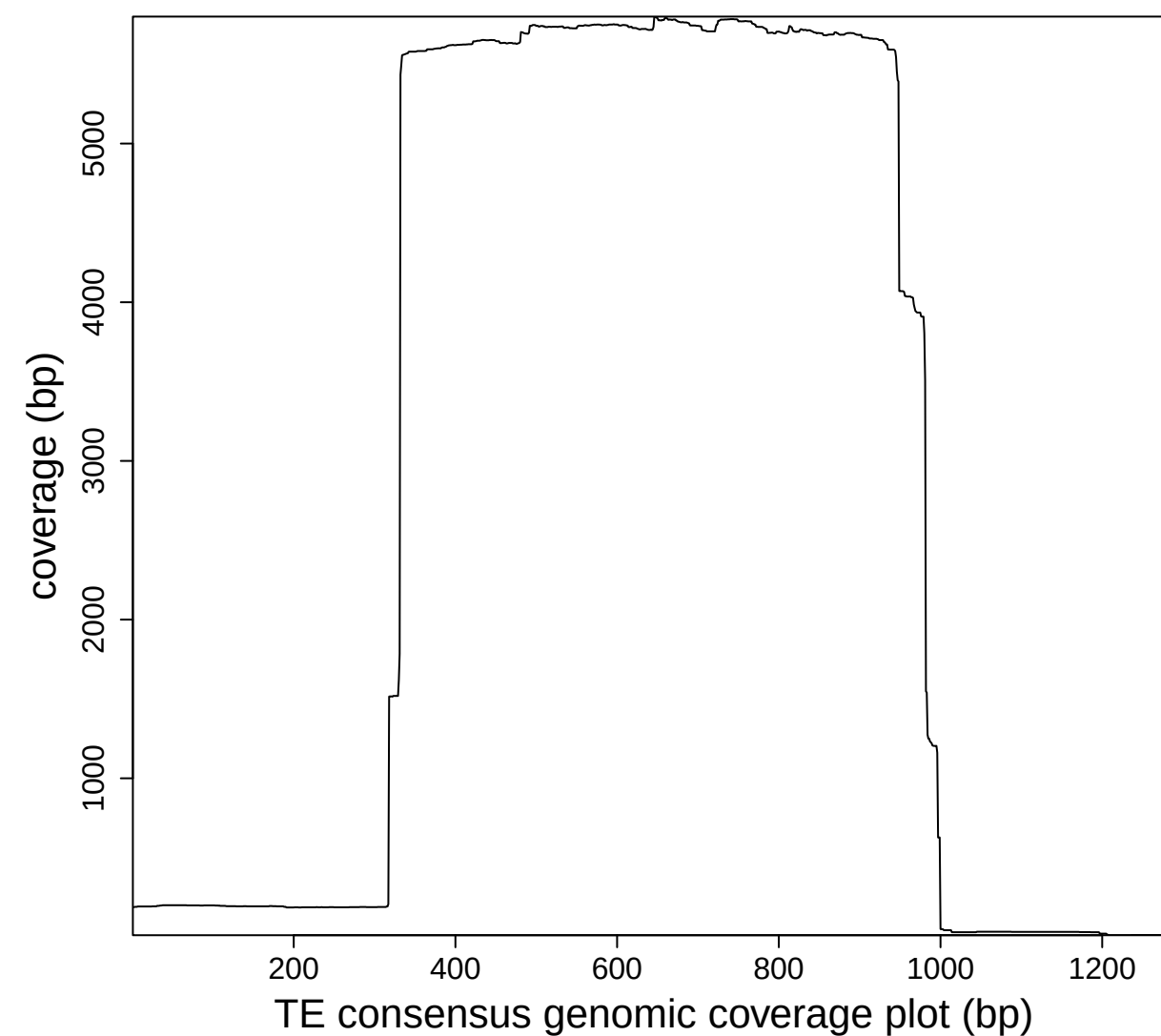
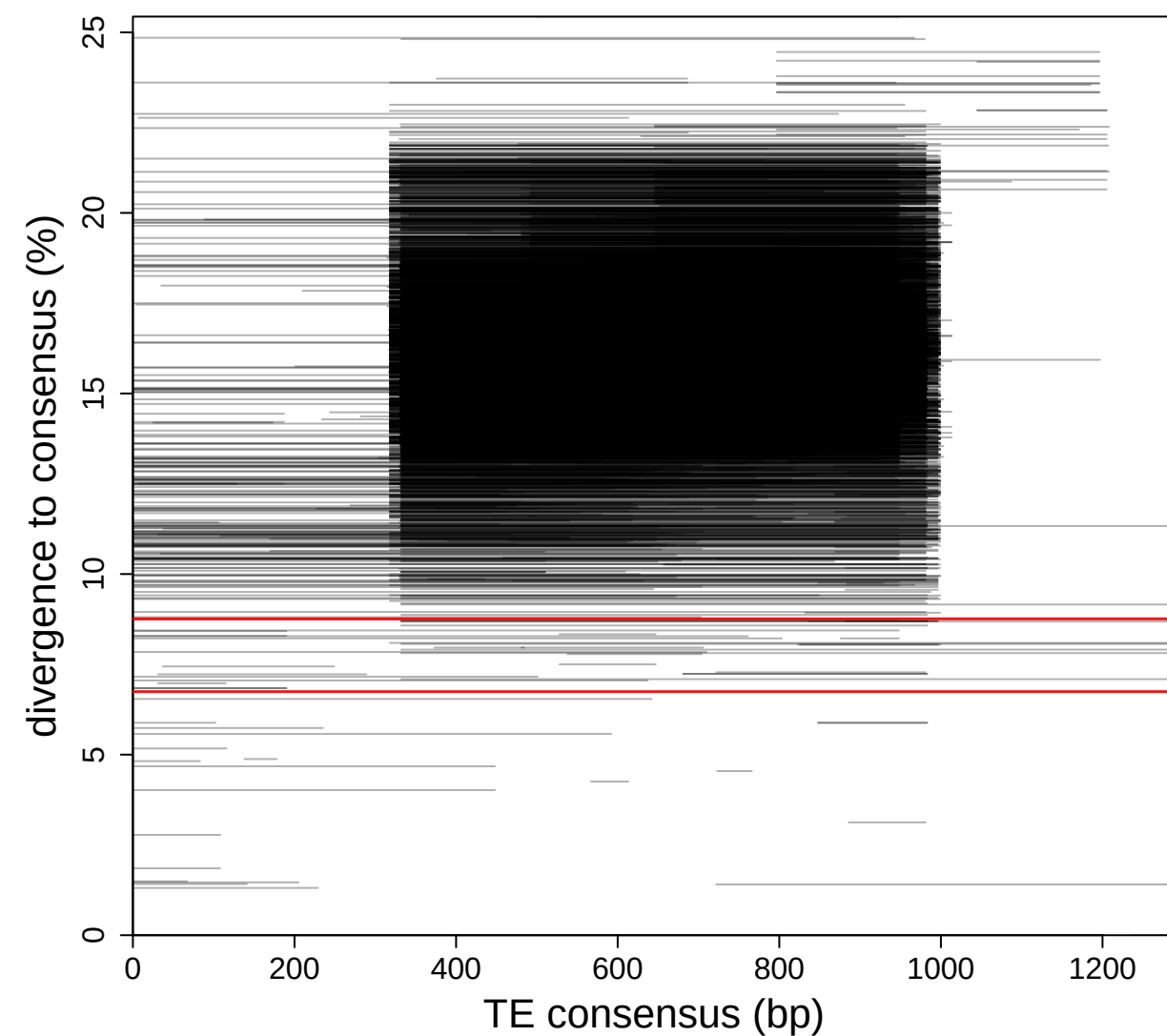


3.fasta.b.bed uf.bed g_3.bed fm_1.bed 0 0 bcln.fa aln.fa cl.f
size: 6069bp; fragments: 16553; full length: 0 (>=5462.1bp)

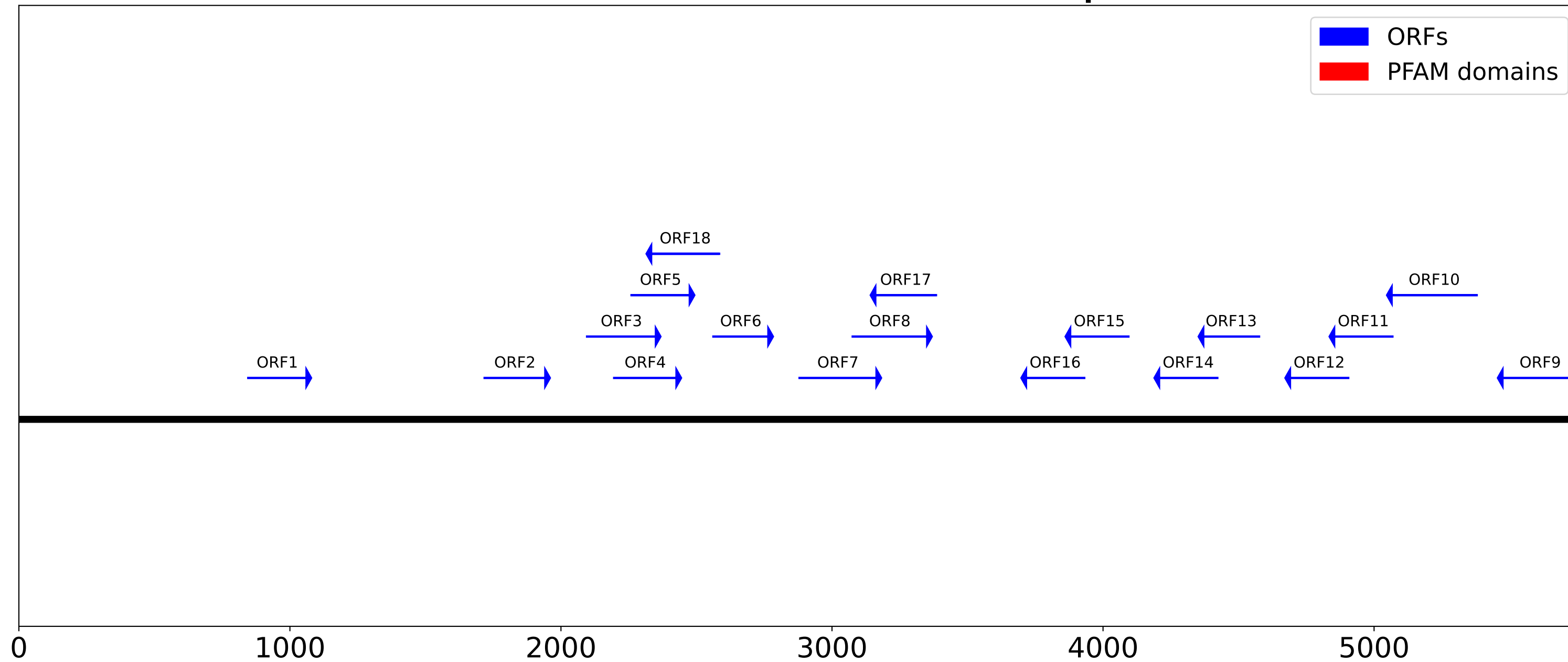
After TEtrimmer Extended plot Blue lines are boundaries



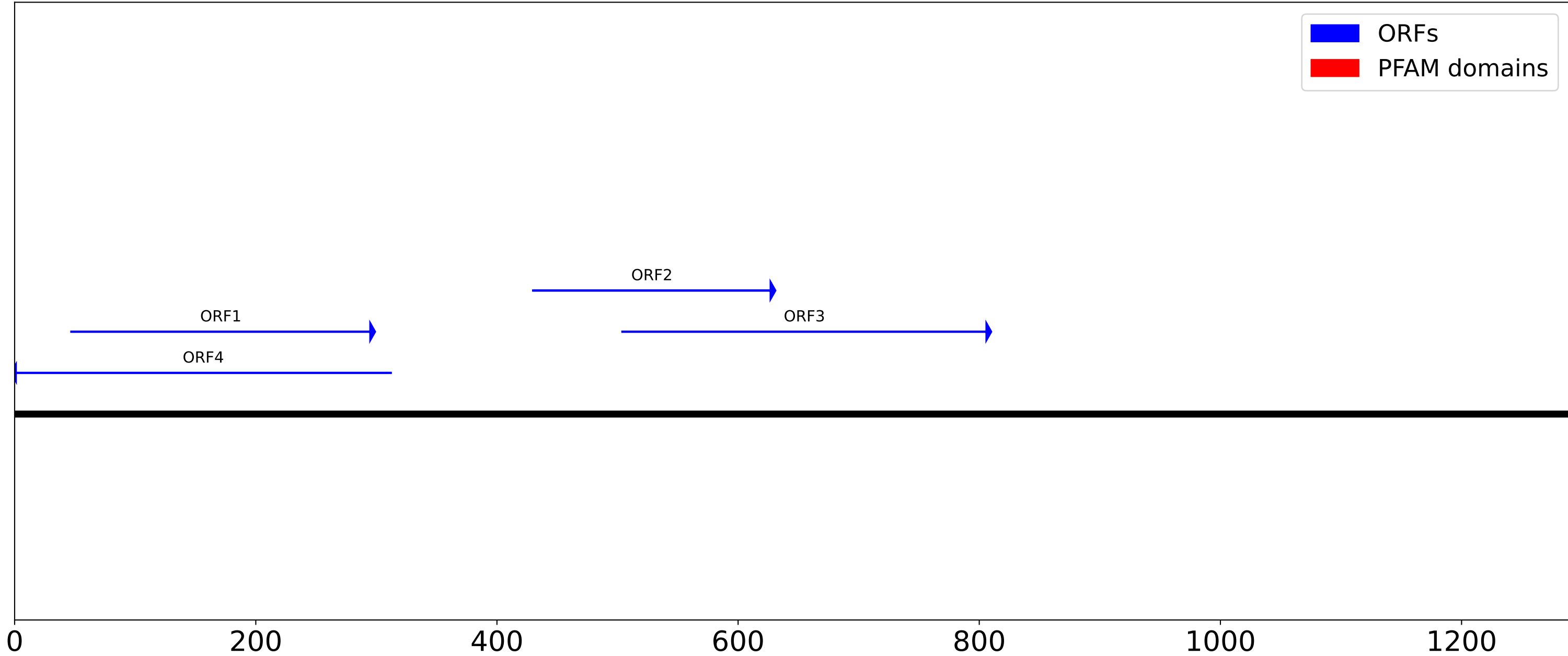
TE: rnd_5_family_8603
size: 1290bp; fragments: 6250; full length: 2 (≥ 1161 bp)



After TEtrimmer ORF and PFAM domain plot



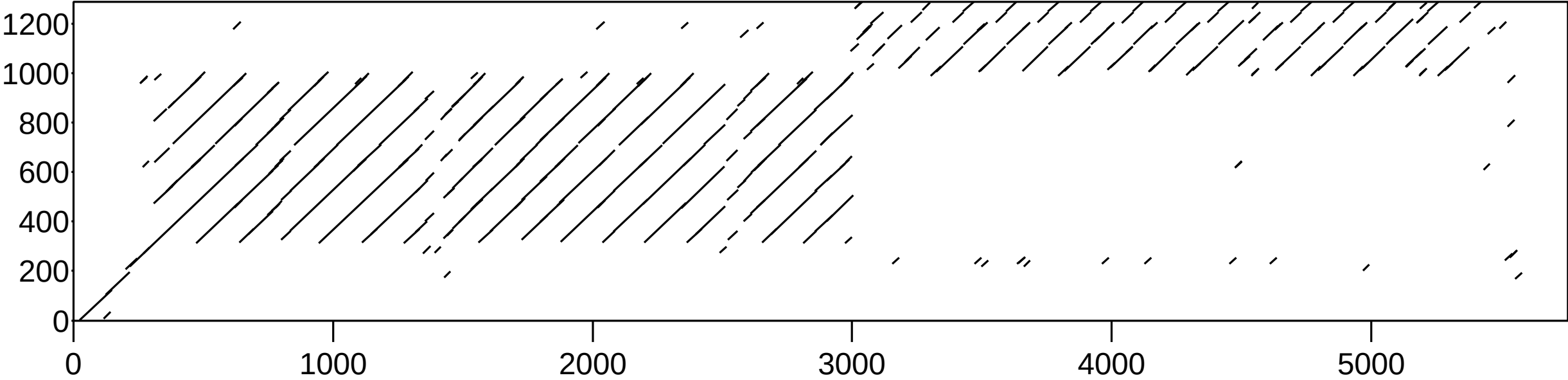
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

onsensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)