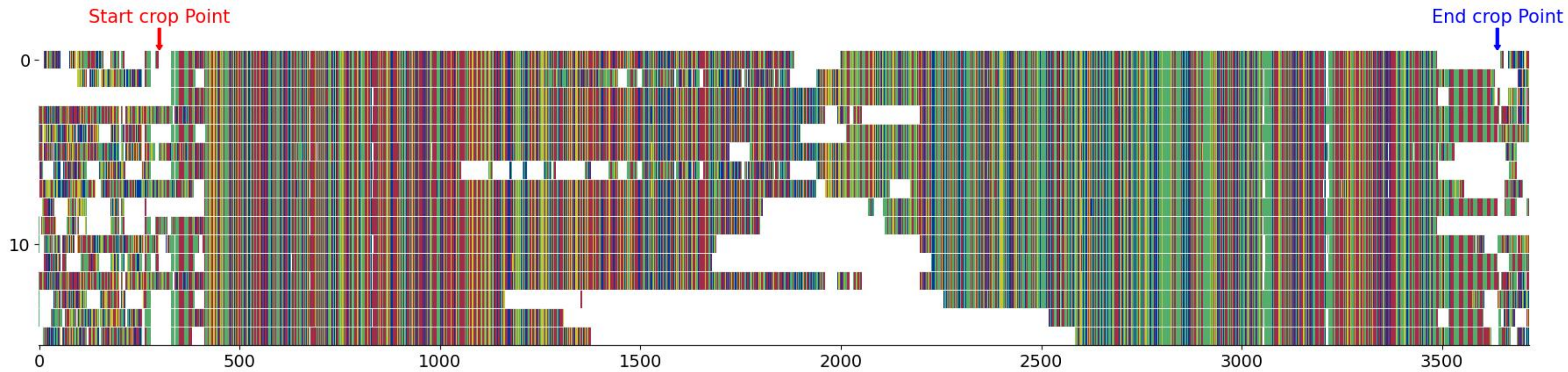


MSA length = 3339



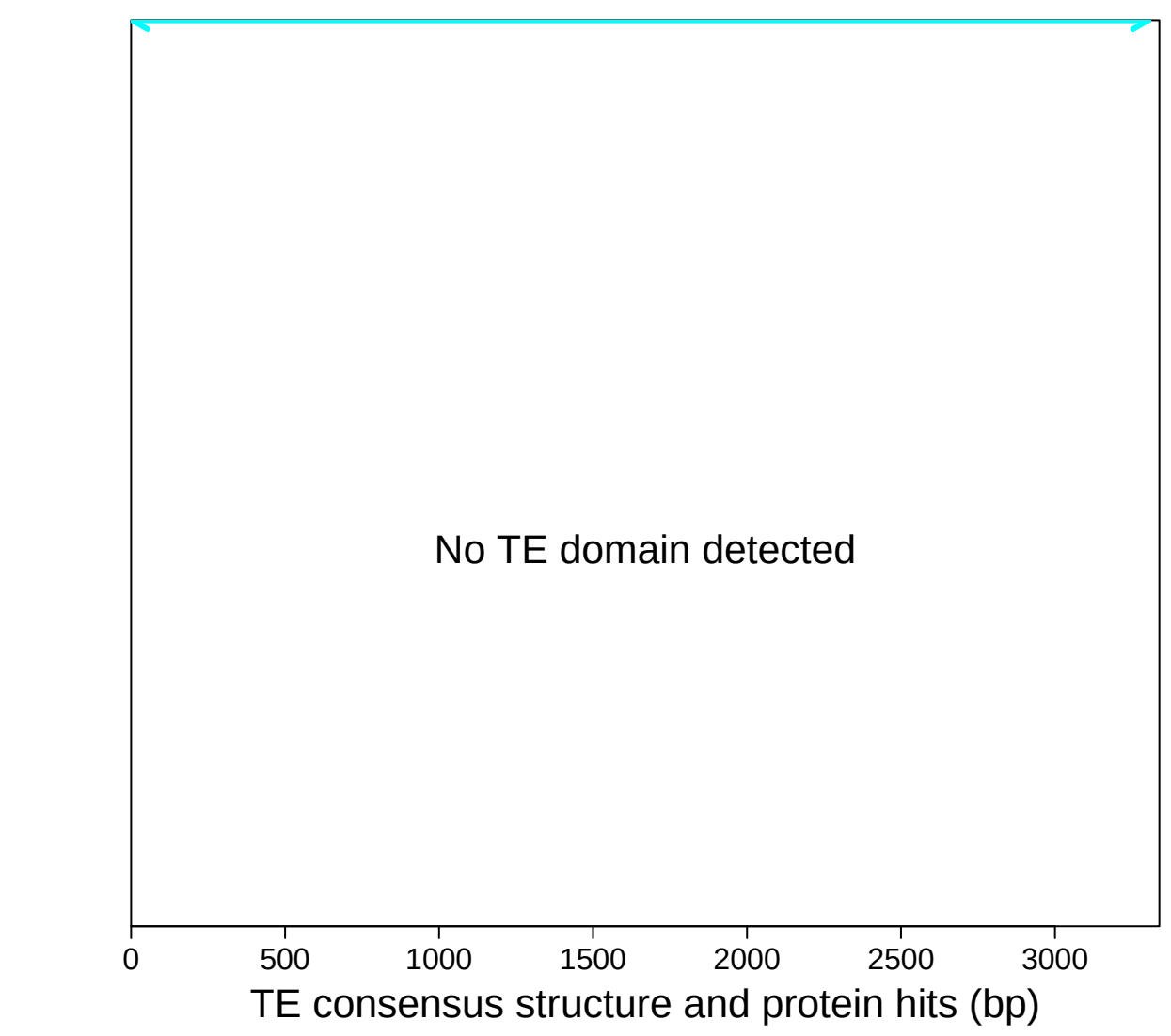
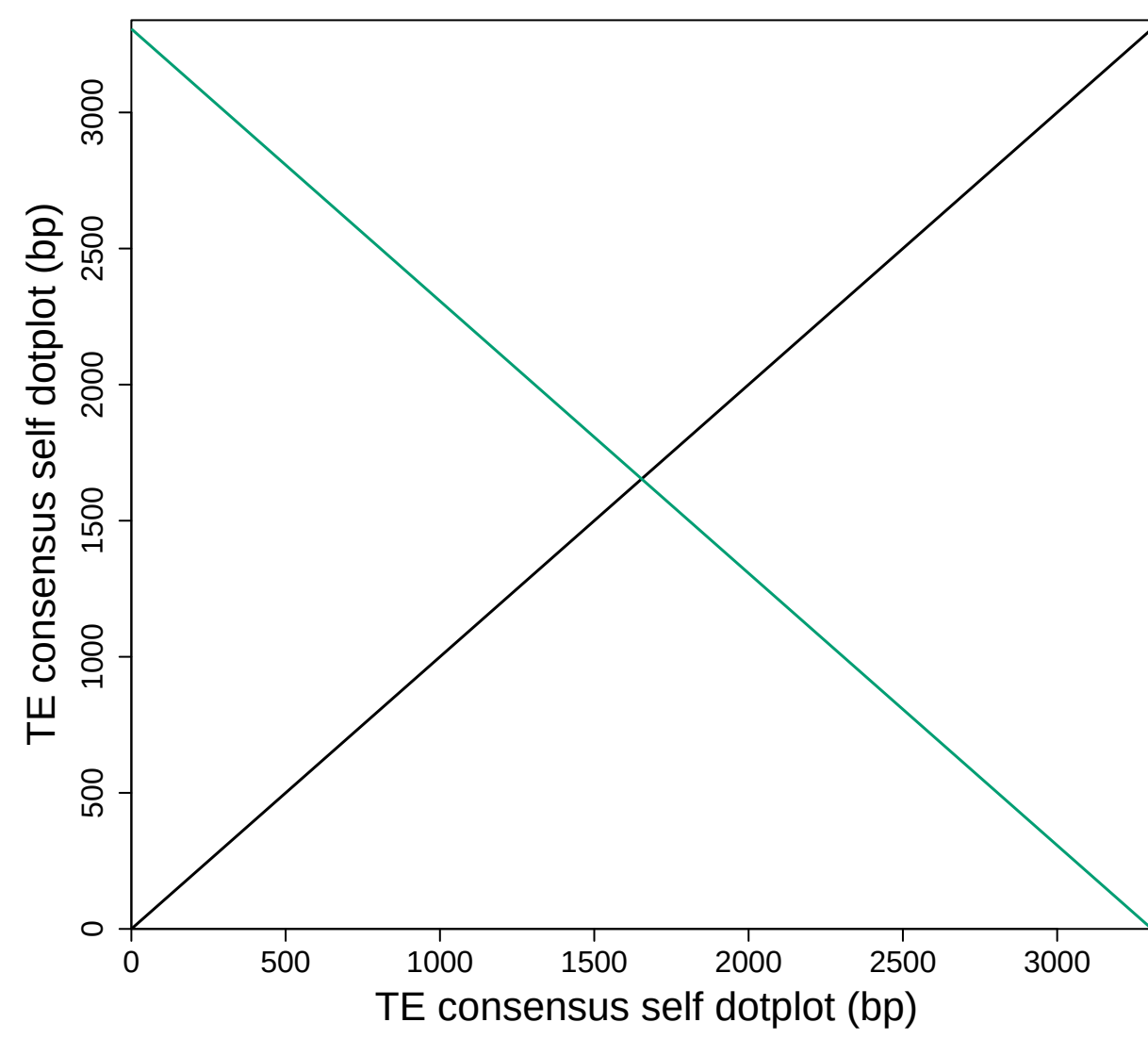
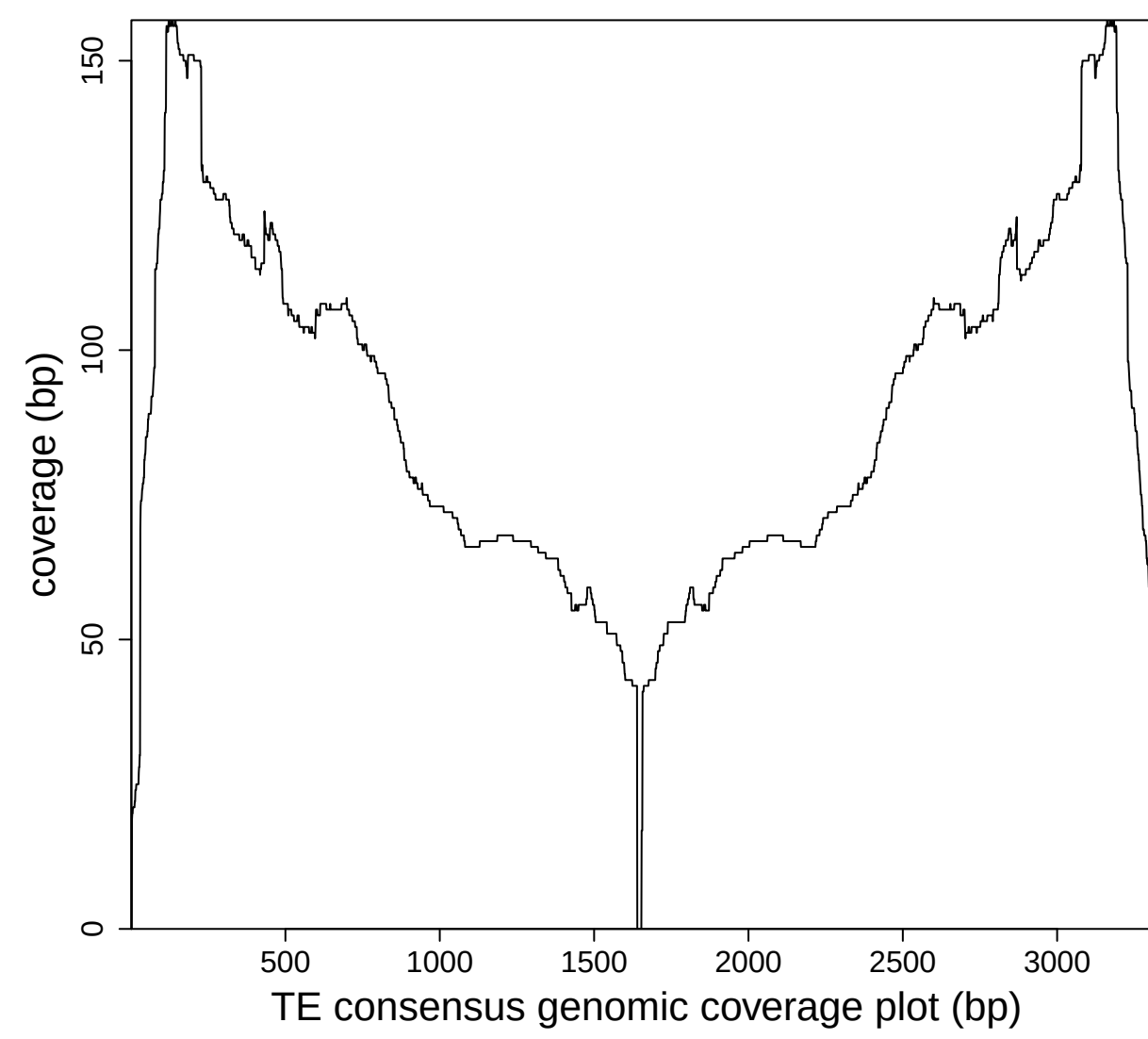
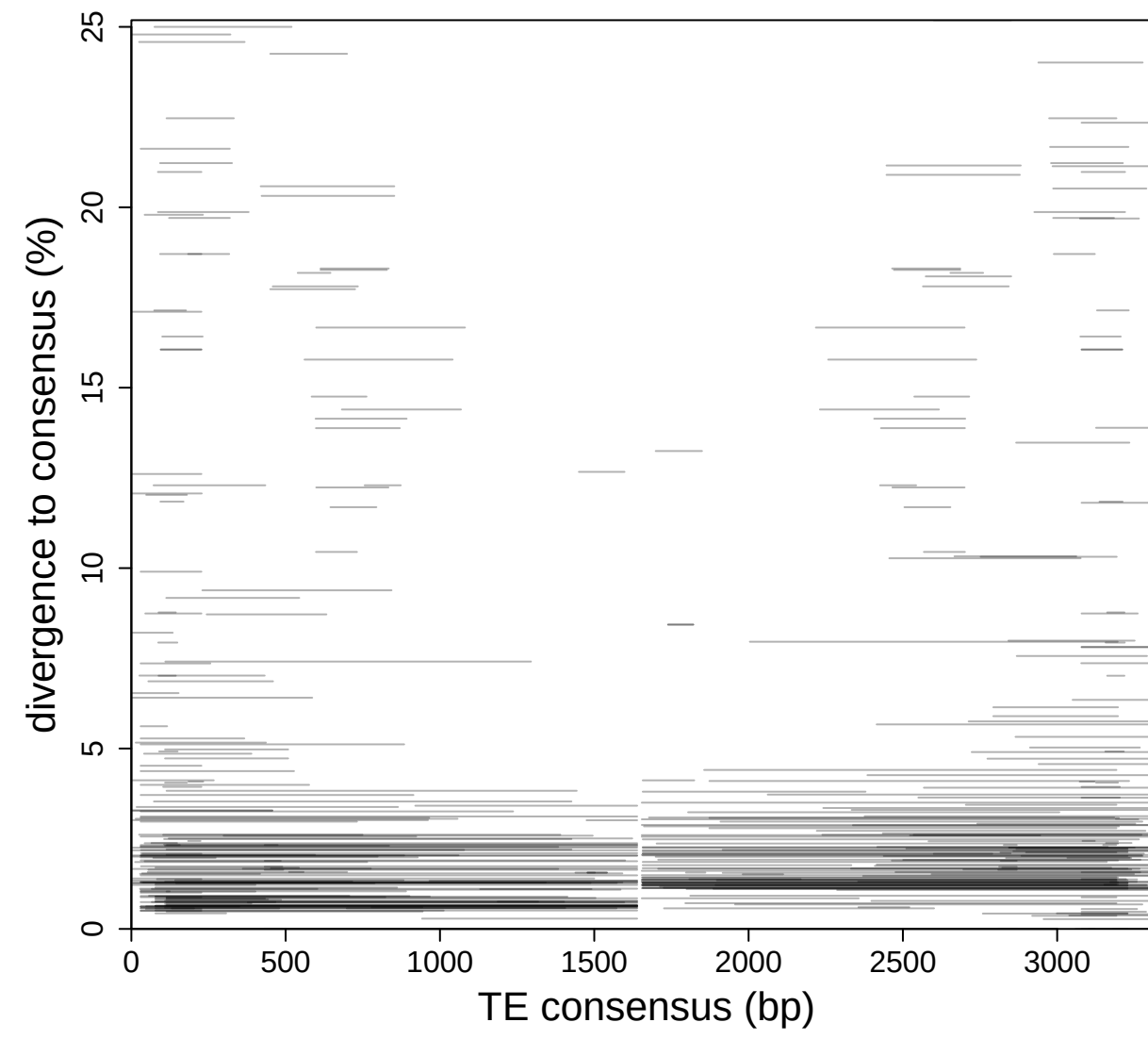
Start crop Point

End crop Point



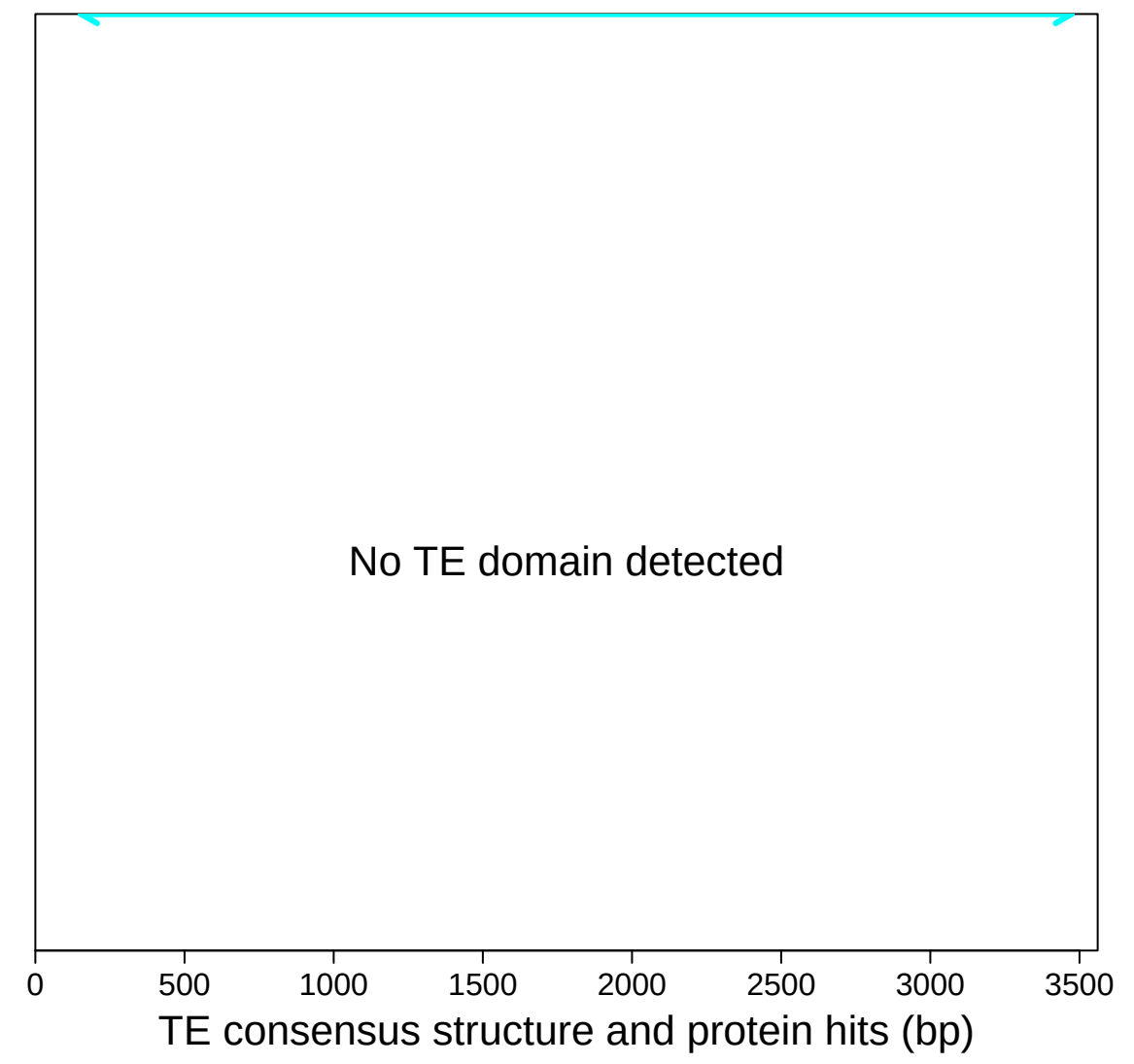
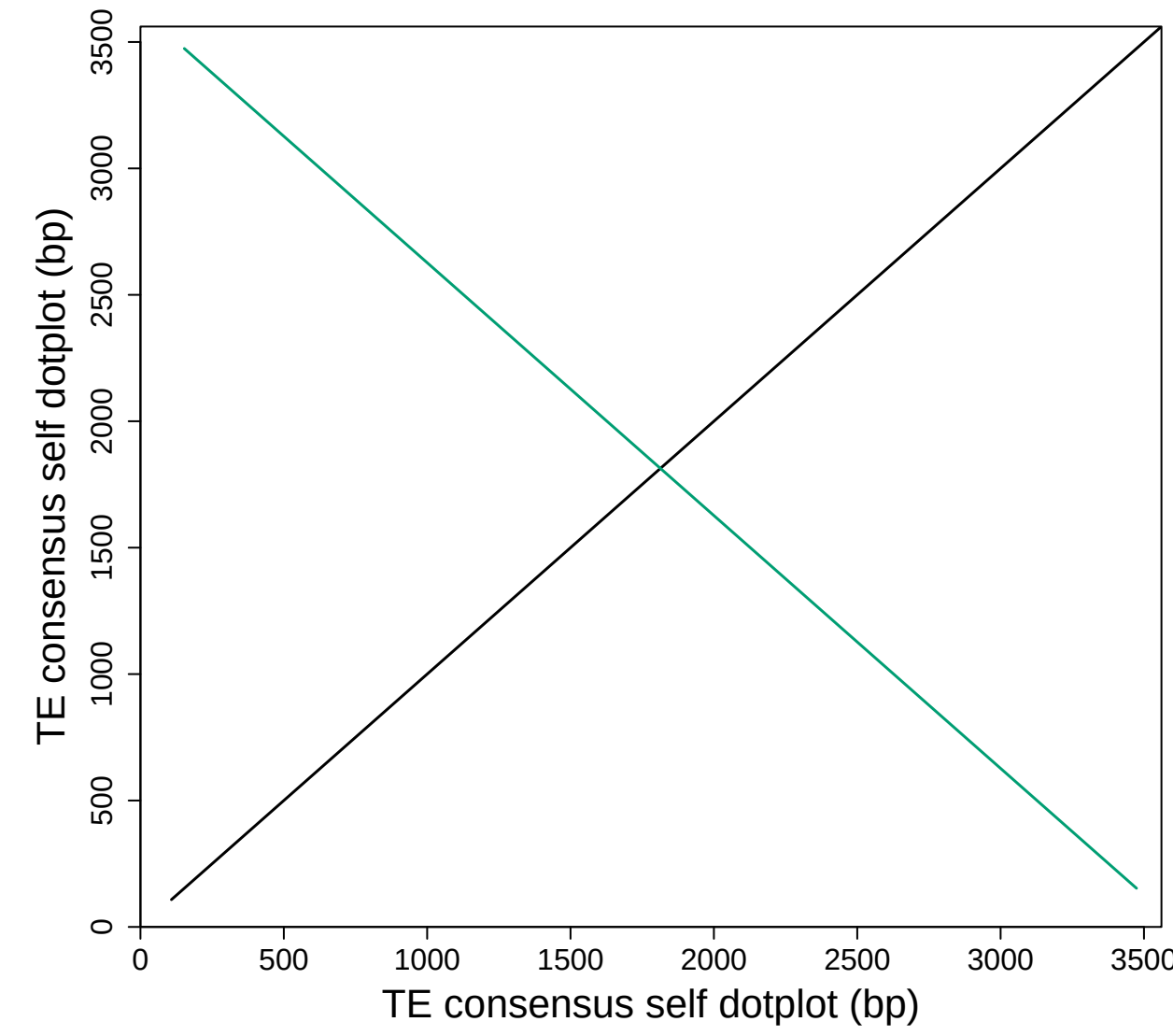
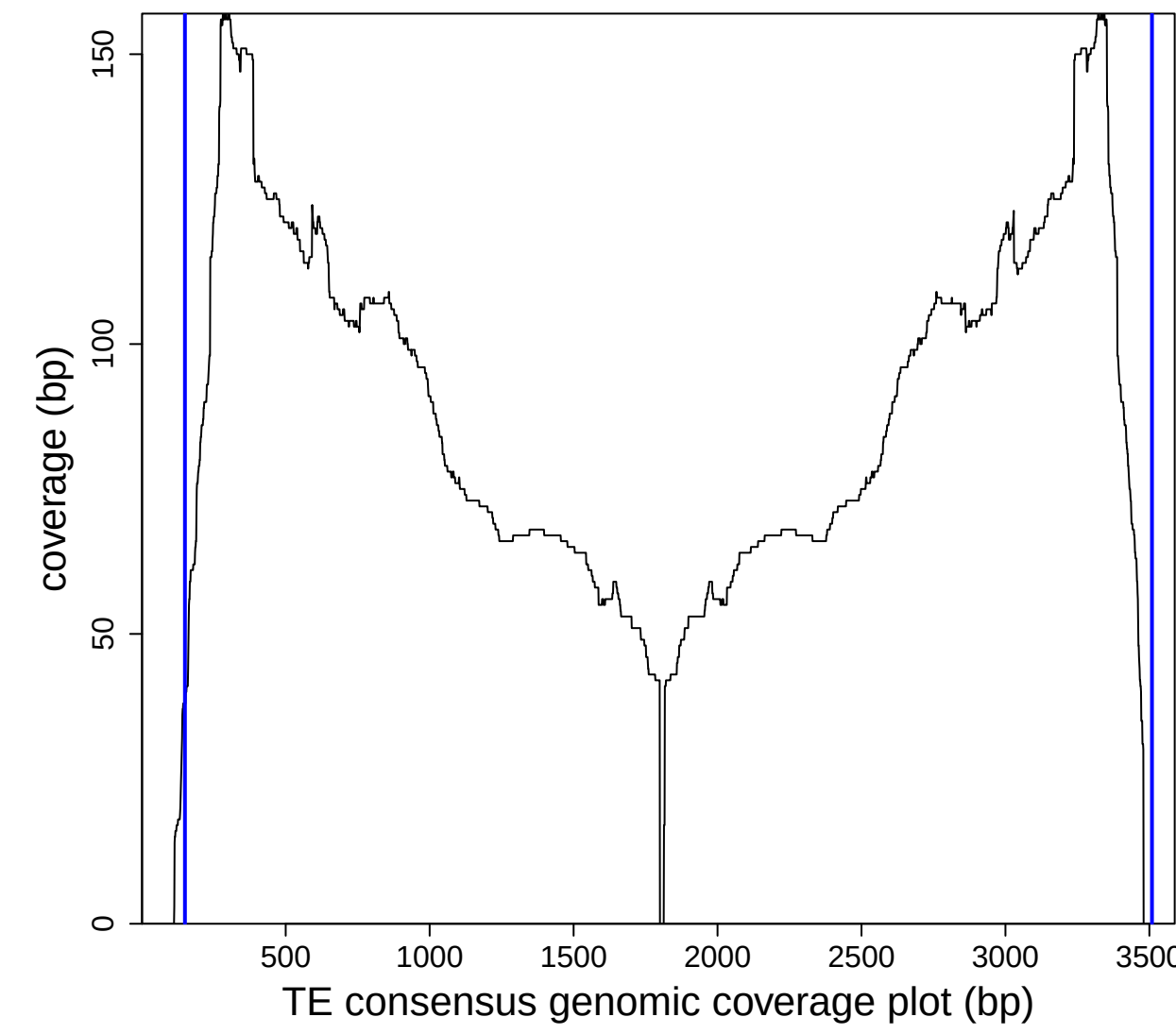
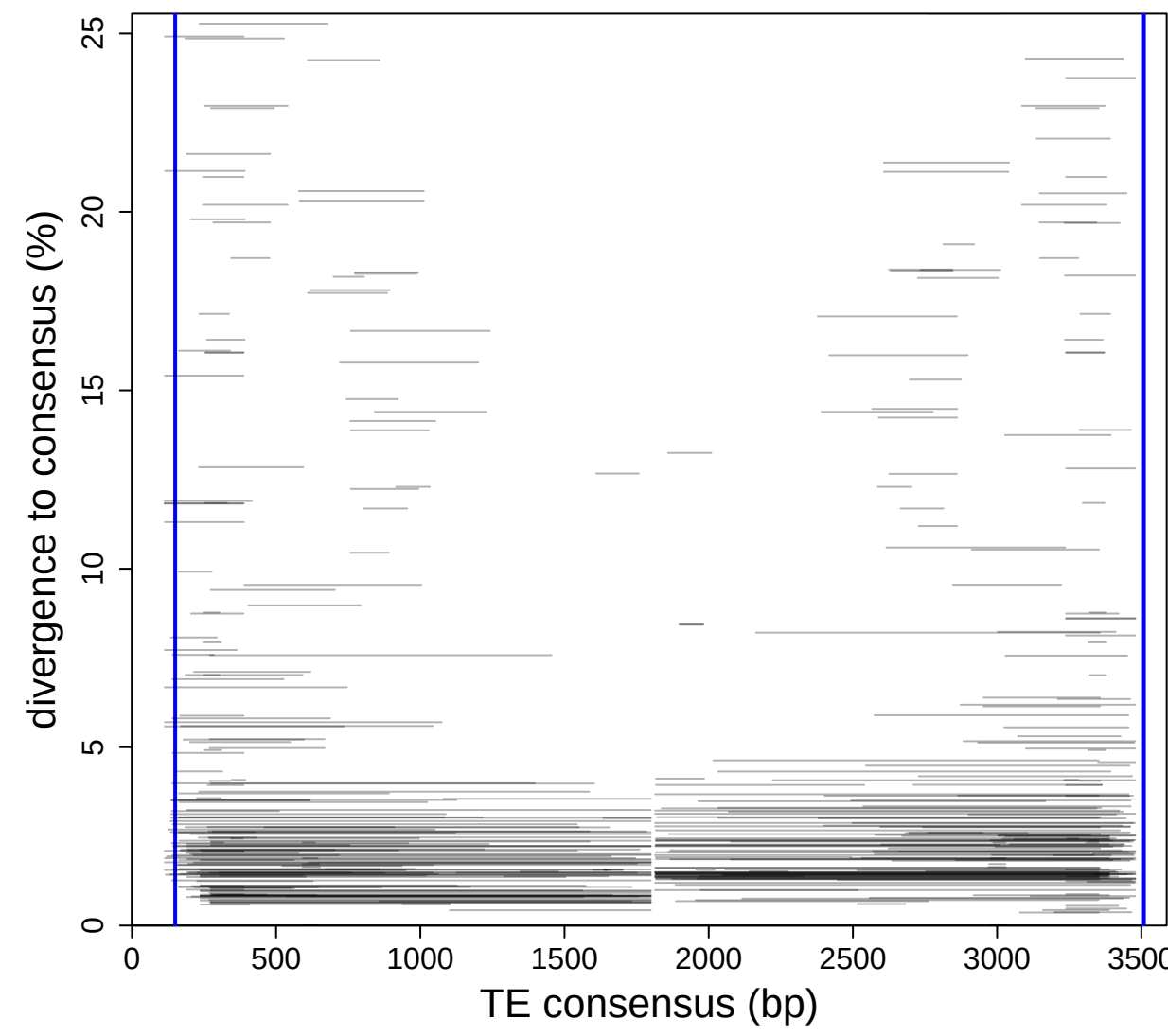
bed_fm_1.bed 0 0 n.bed g 1.bed fm_2.bed 0 0 bcln.fa aln.fa
size: 3339bp; fragments: 421; full length: 0 (>=3005.1bp)

After TEtrimmer 3339 bp



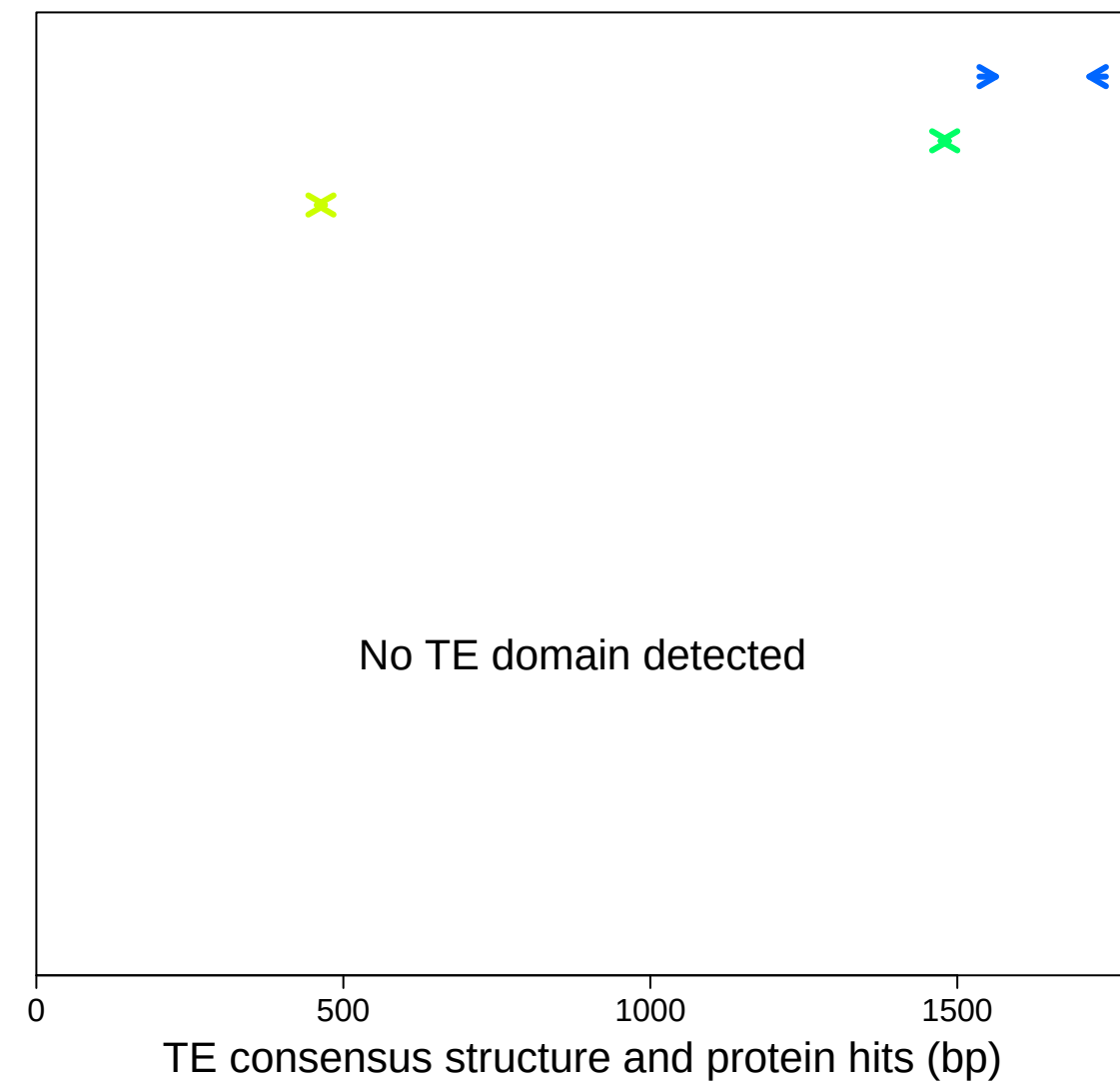
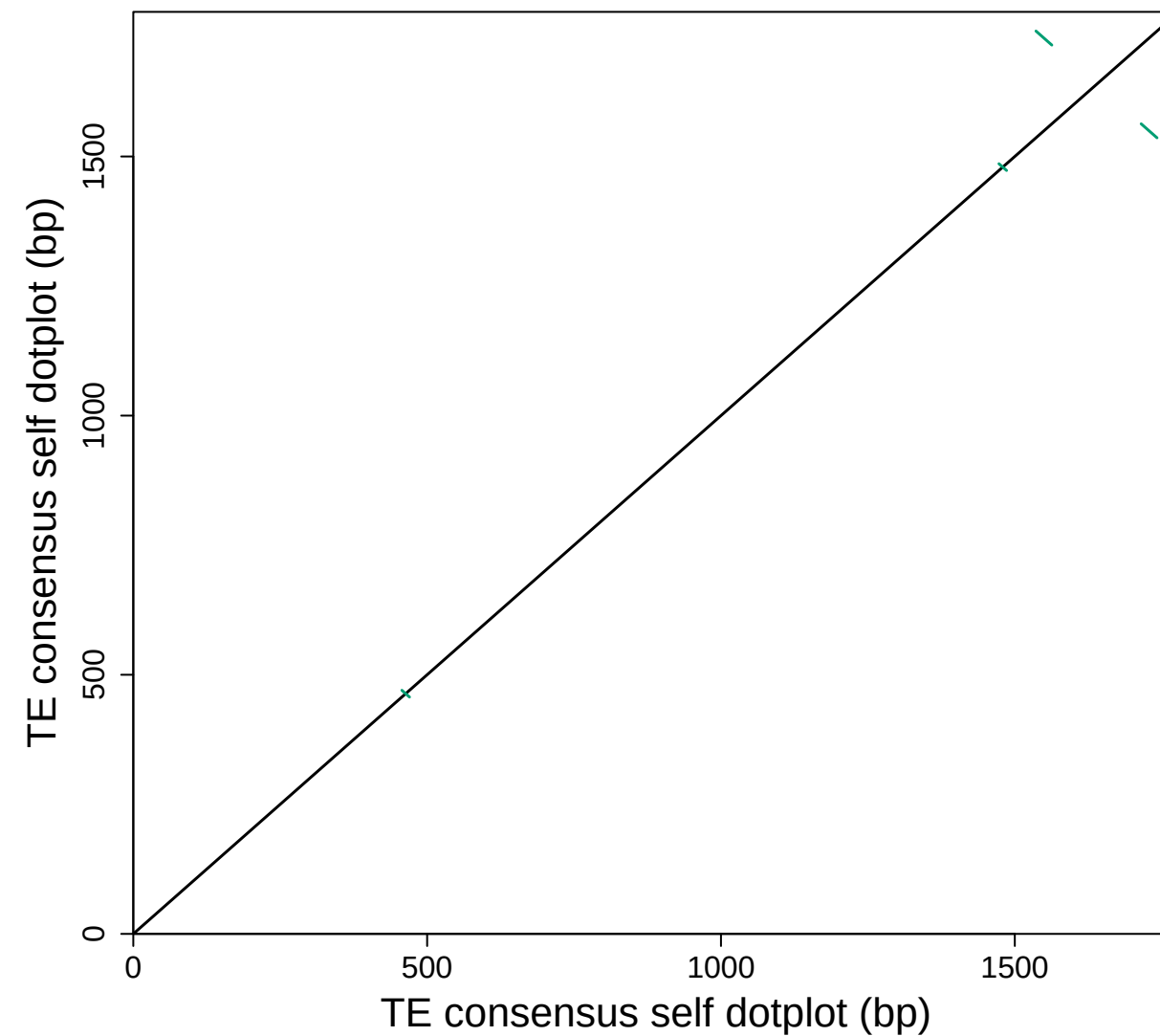
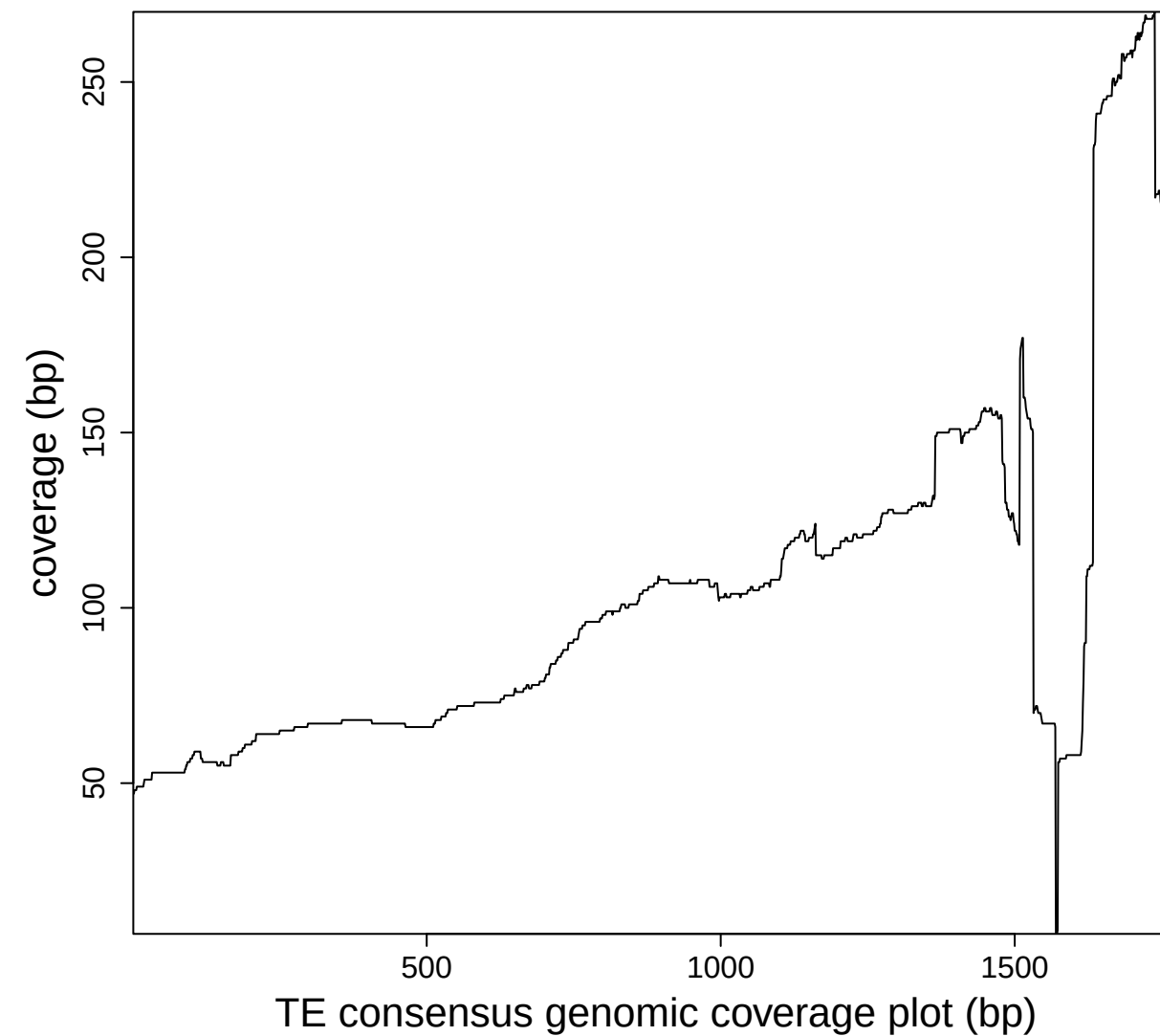
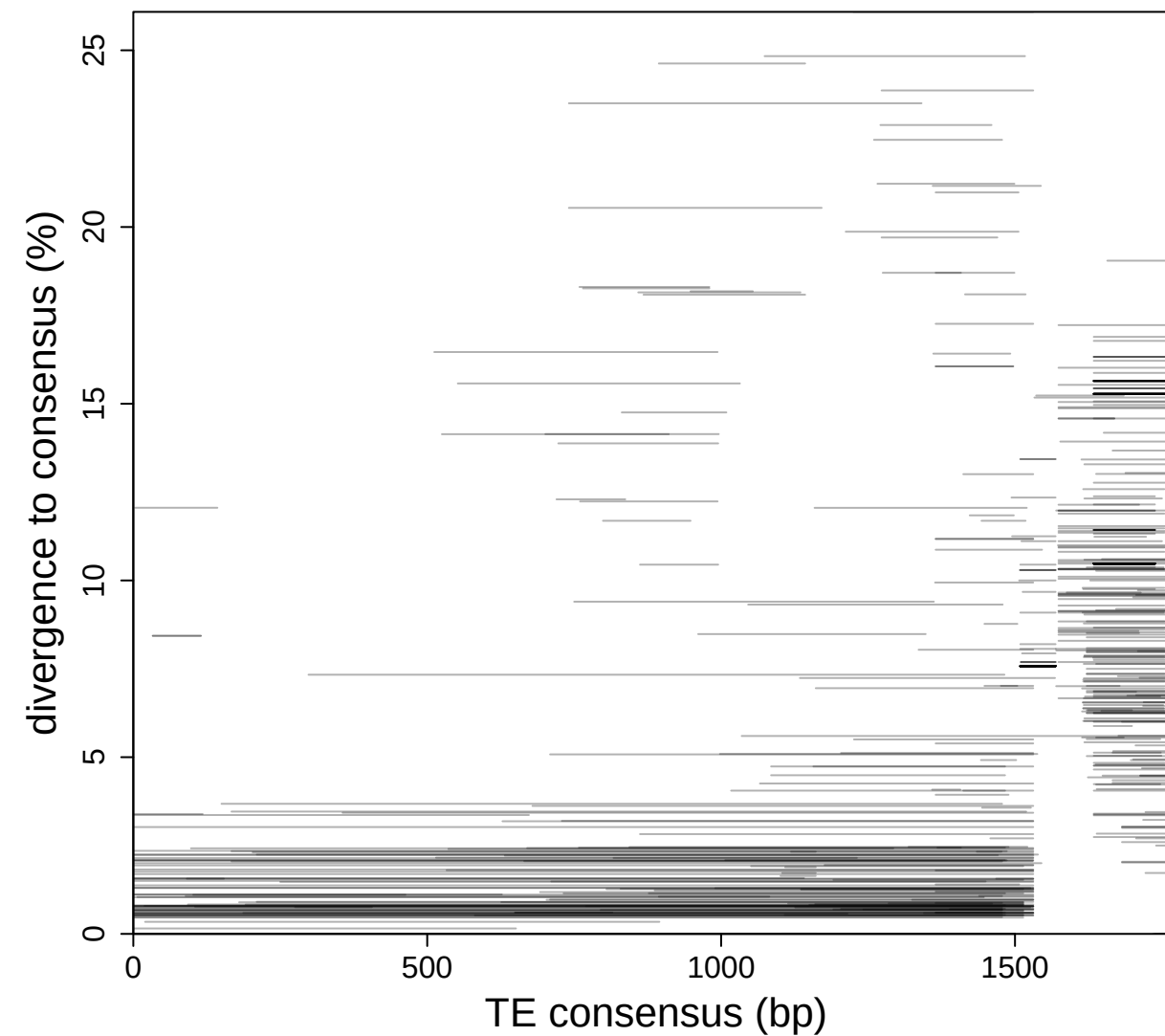
f.bed_g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcl
size: 3588bp; fragments: 421; full length: 0 (>=3229.2bp)

After TEtrimmer Extended plot Blue lines are boundaries

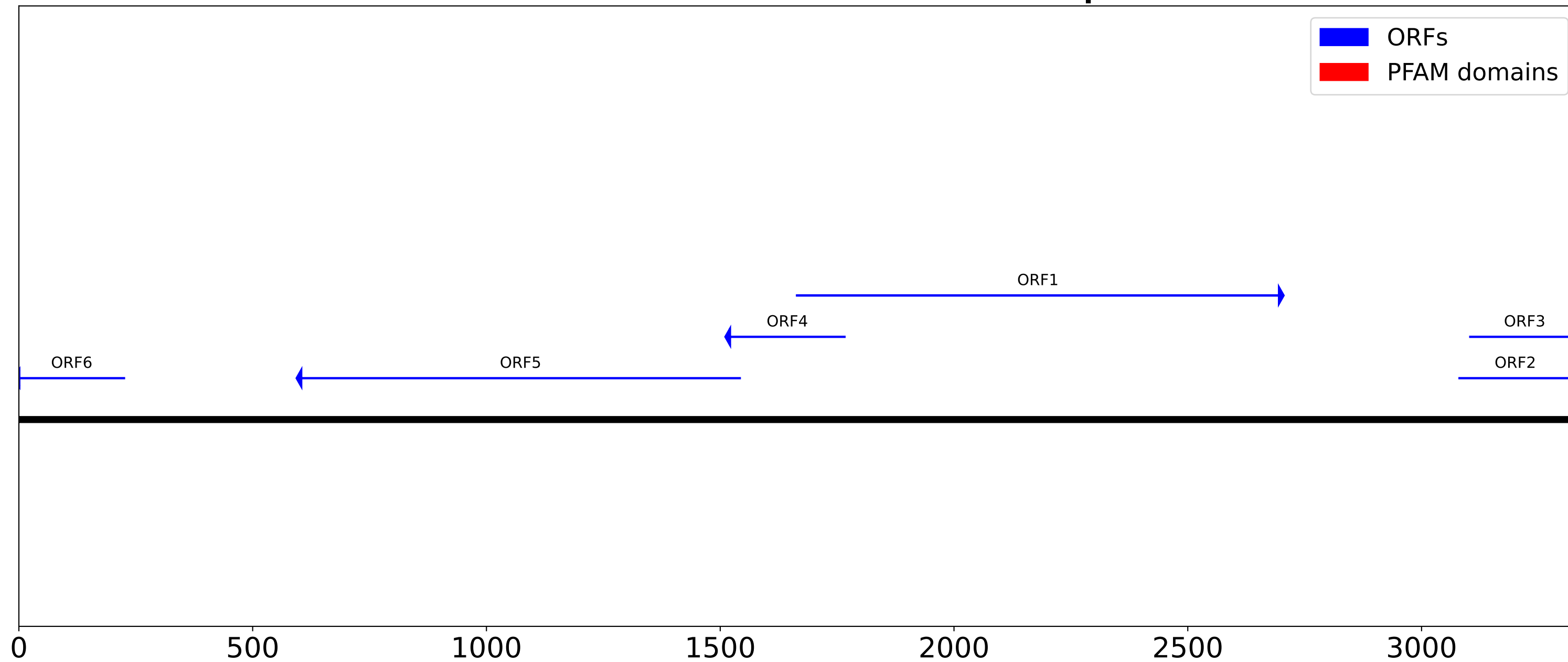


TE: rnd_5_family_983
size: 1779bp; fragments: 558; full length: 0 (>=1601.1bp)

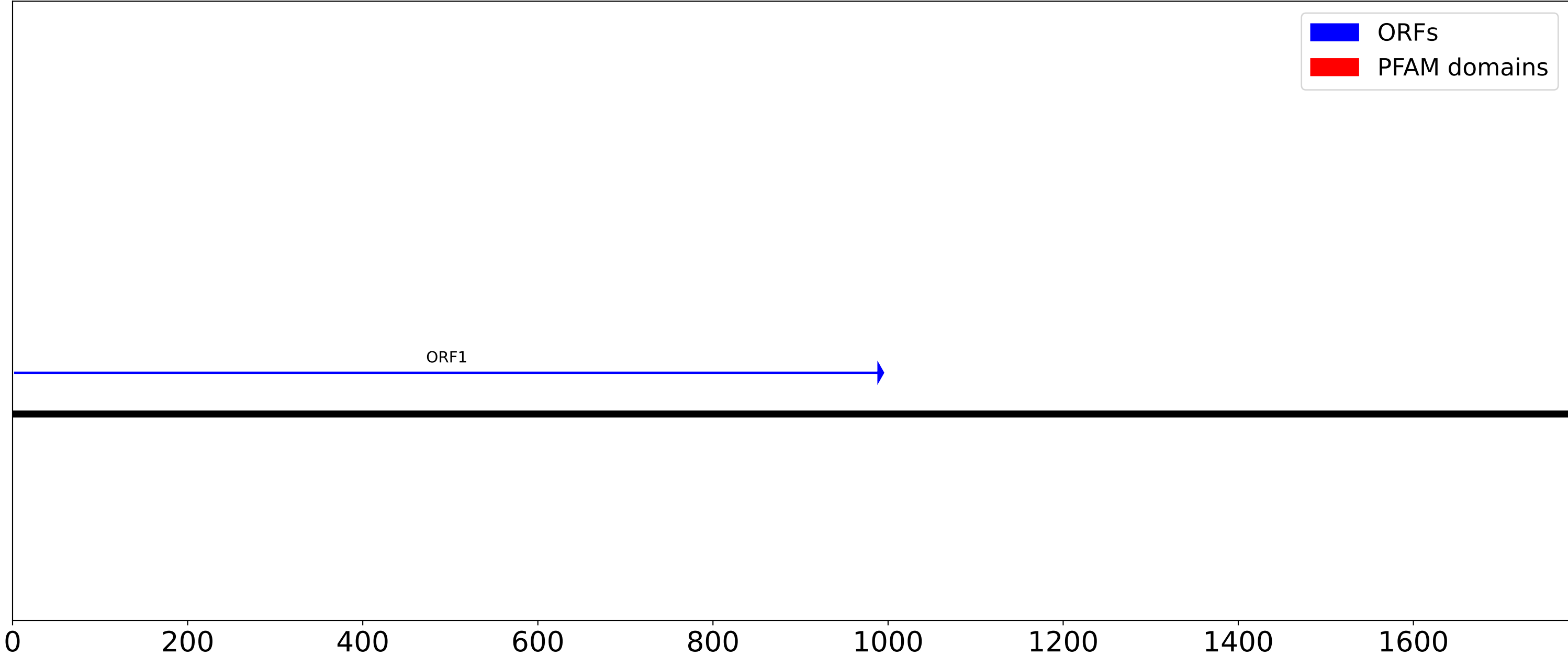
Before TEtrimmer 1779 bp



After TEtrimmer ORF and PFAM domain plot



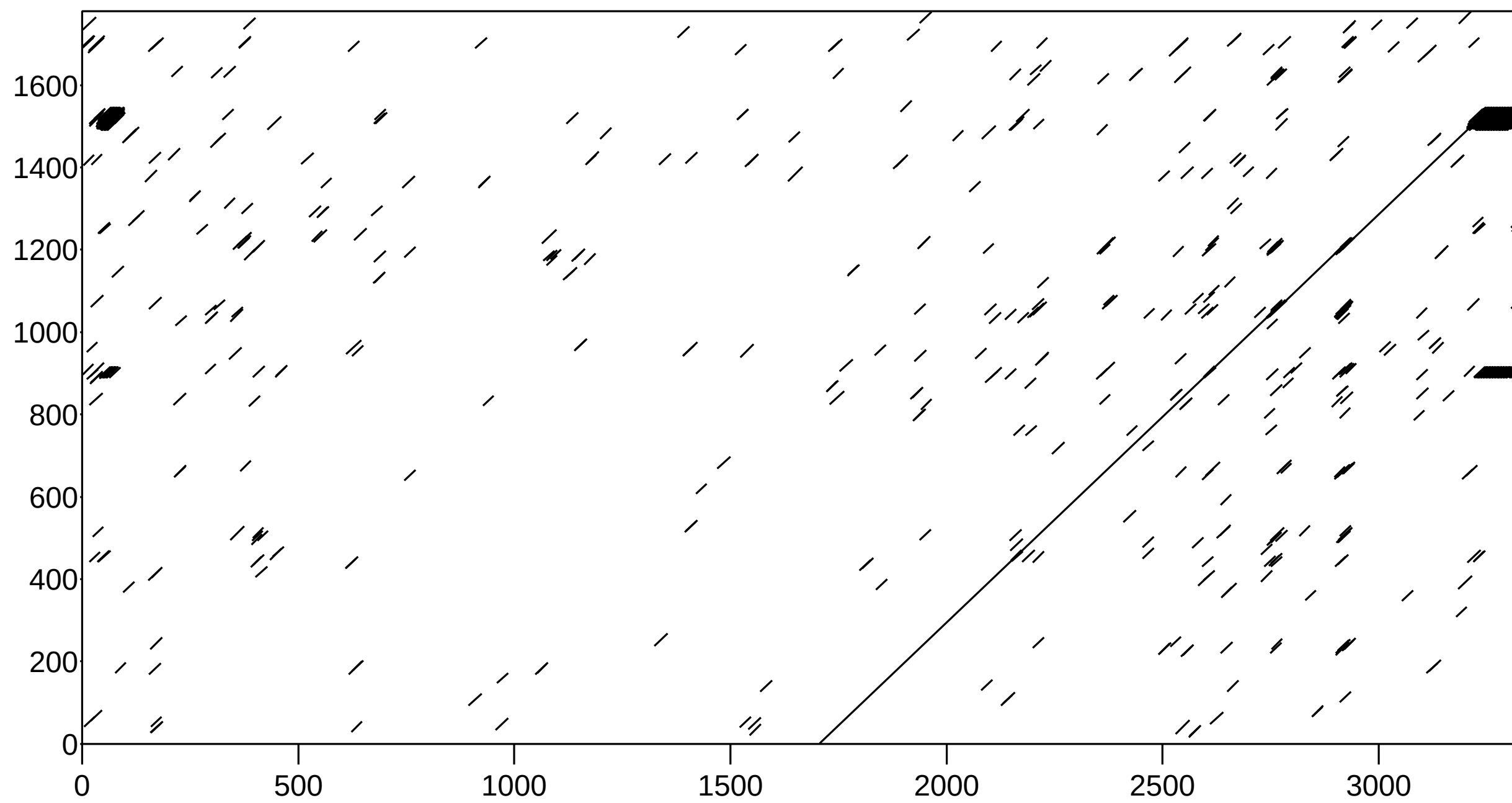
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)