

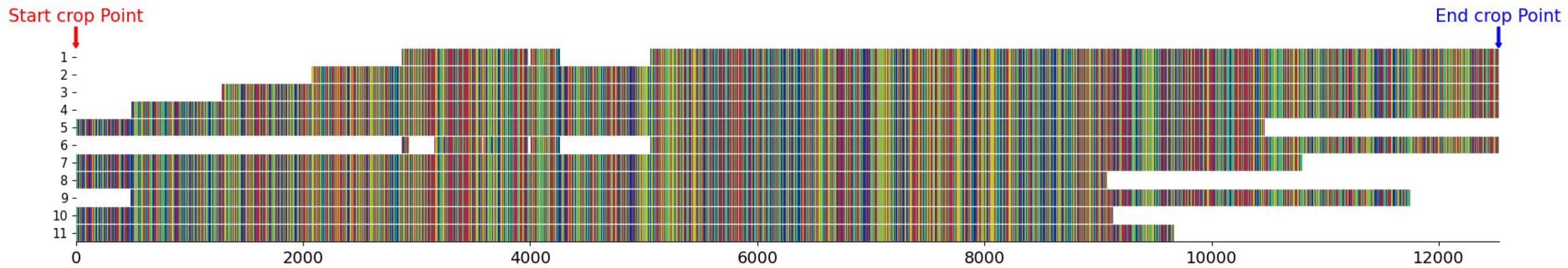
Start crop Point

End crop Point

MSA length = 12527

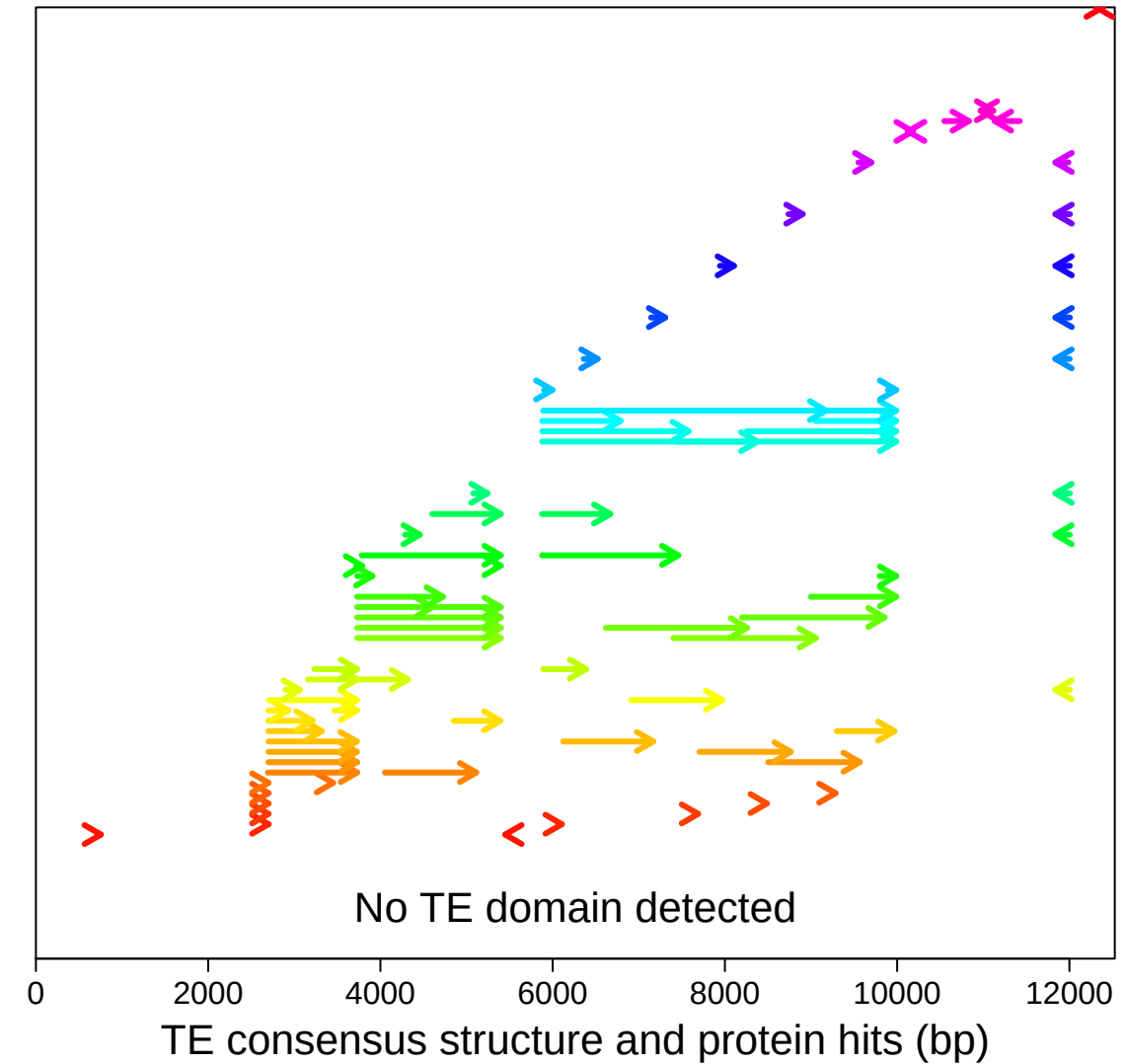
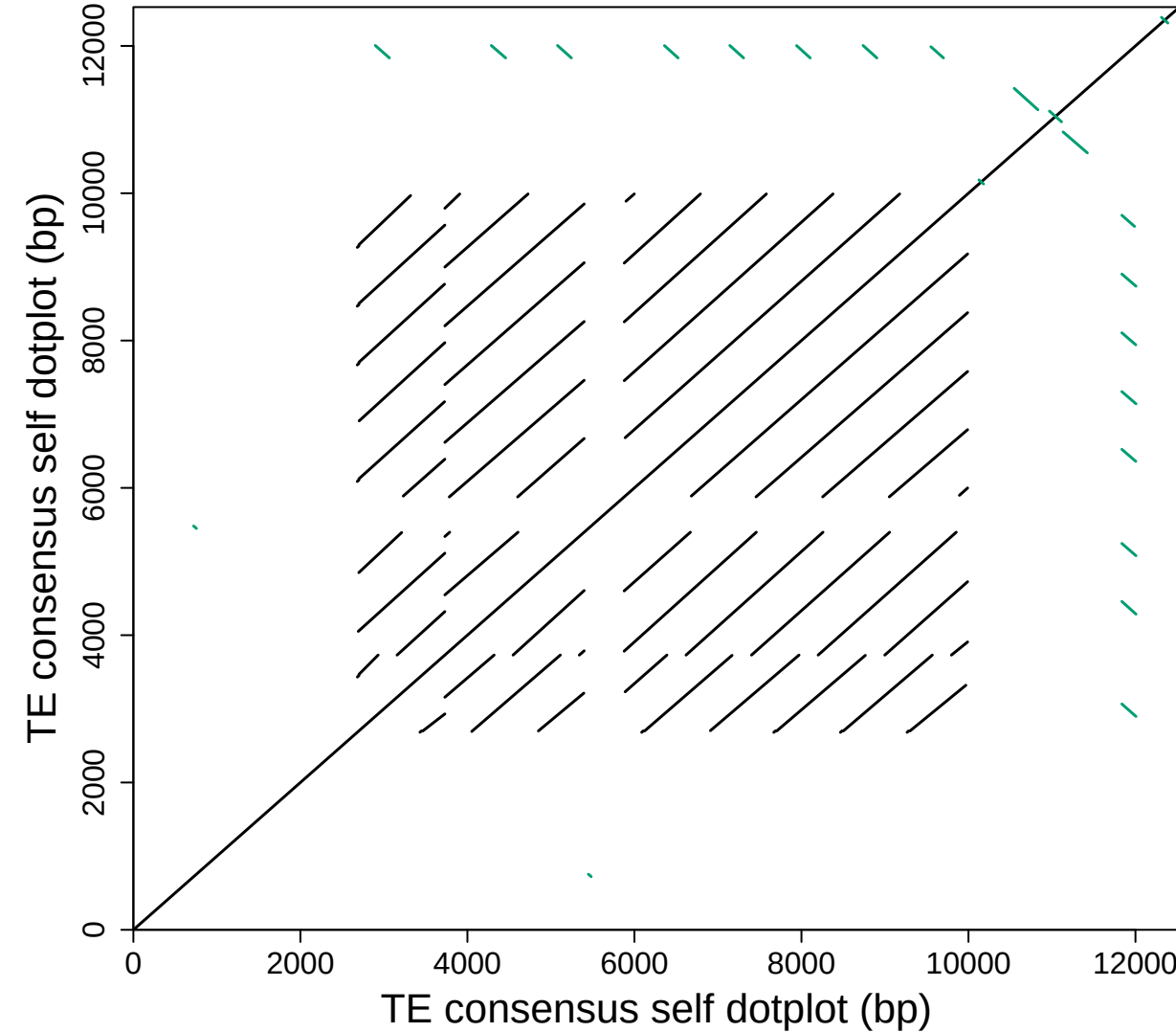
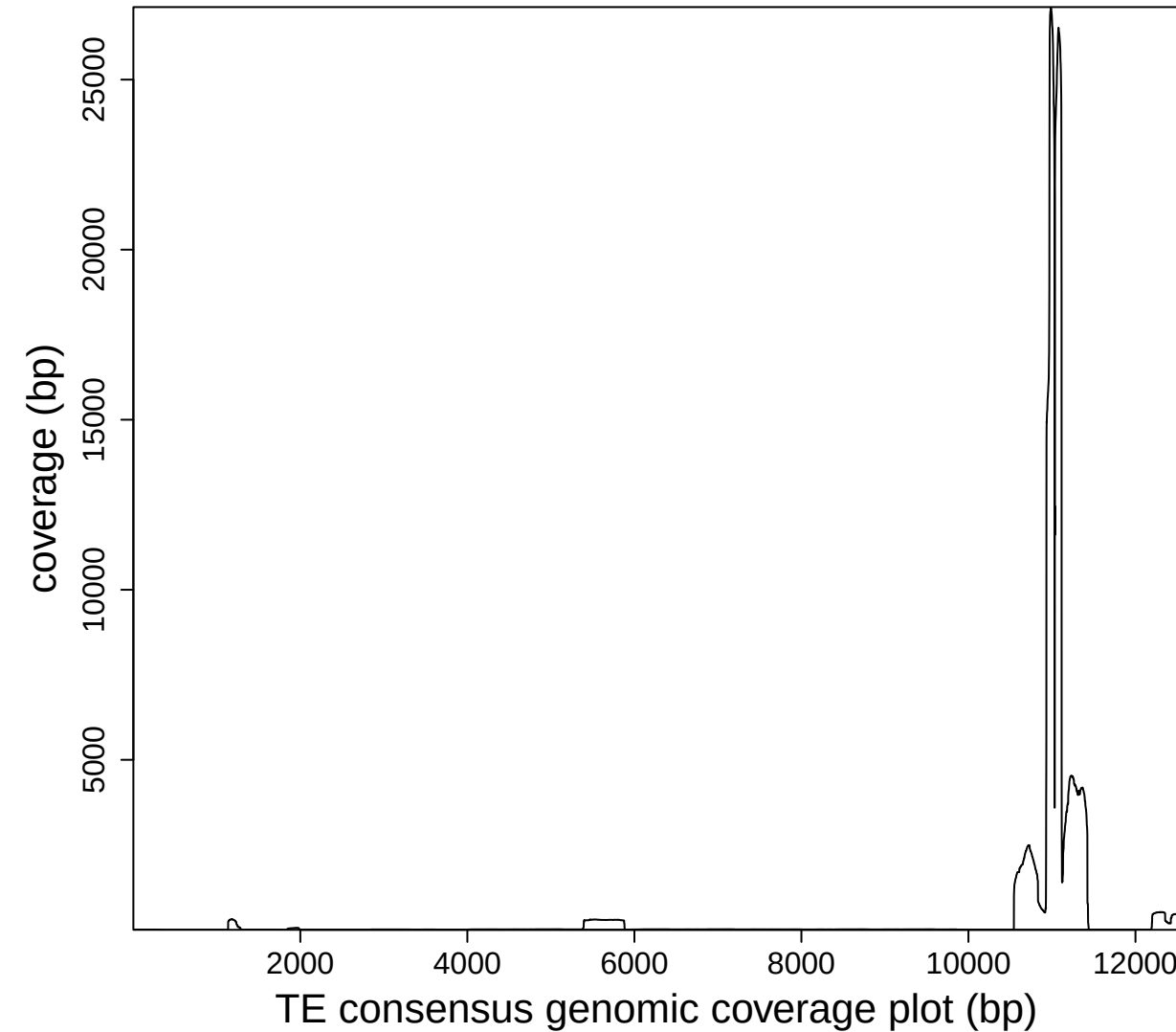
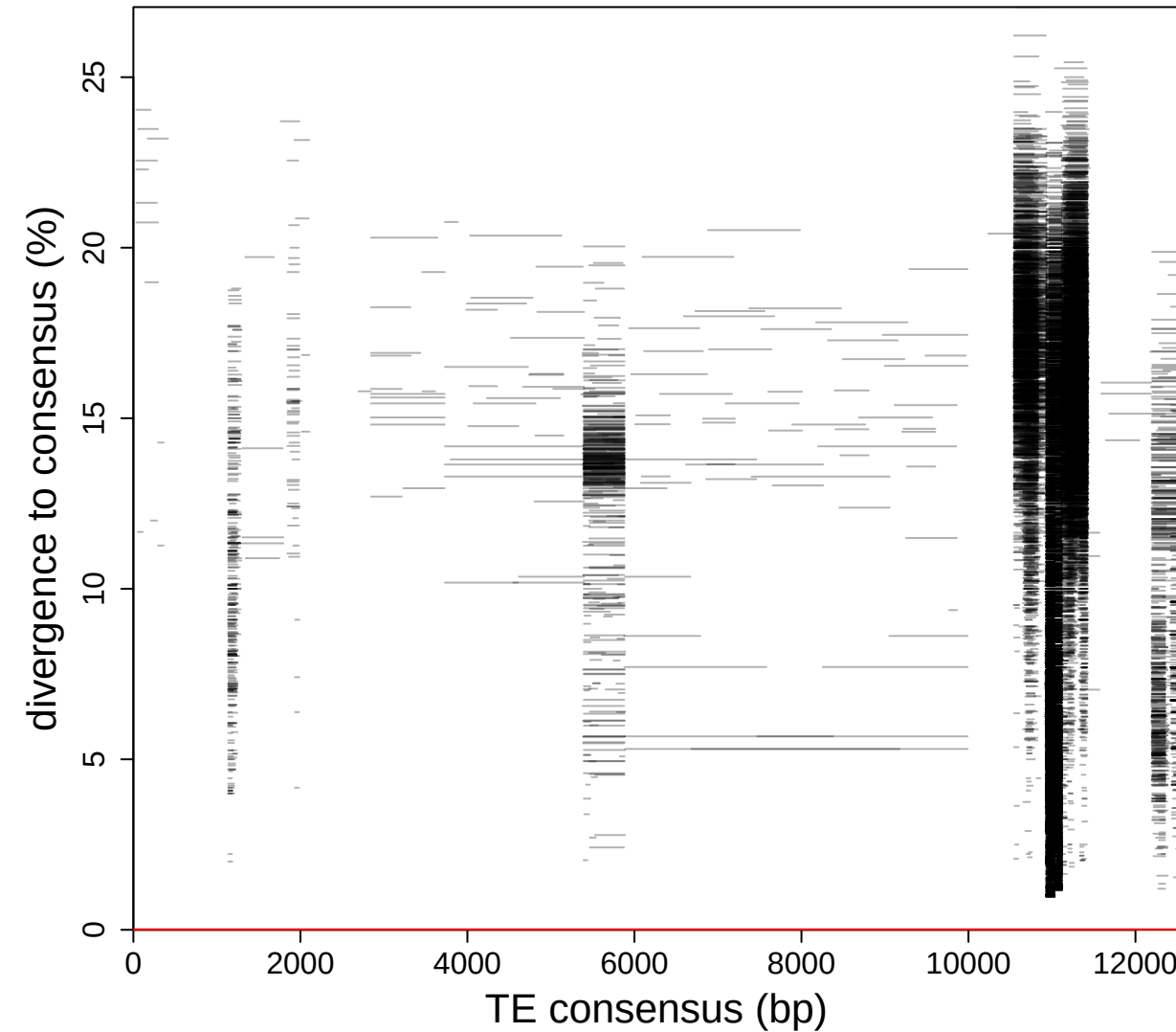
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-----TAATAAAATGTGTAATATGTGTAGTAAATATTATAAAGTCATTTTTTAGAA
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GTGACCTTTACAGATCCACTGTGTTTCTCTTTTGATTCCCCACTCCTATT-----
-----TAATAAAATGTGTAATATGTGTAGTAAATATTATAAAGTCATTTTTTAGAA
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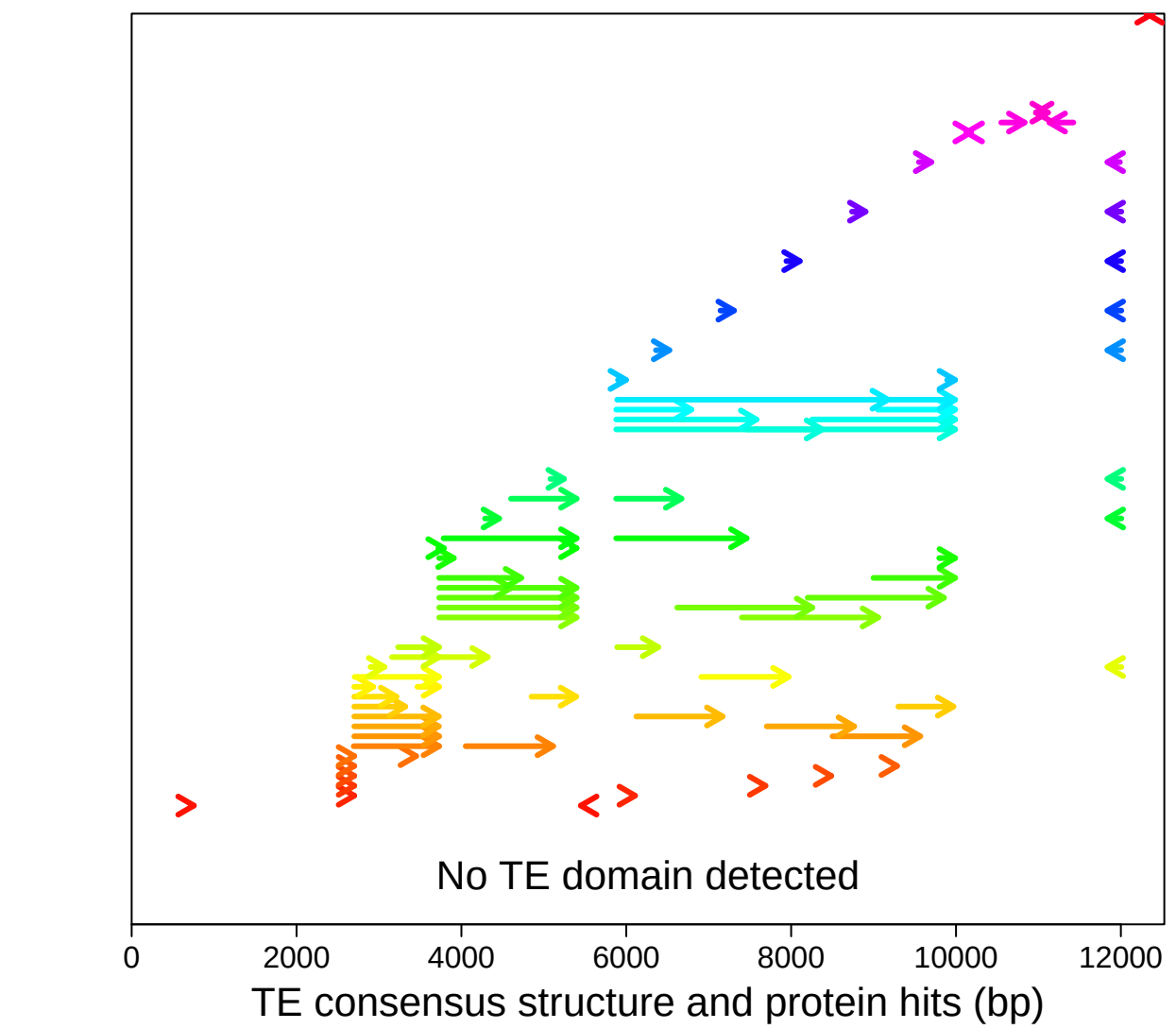
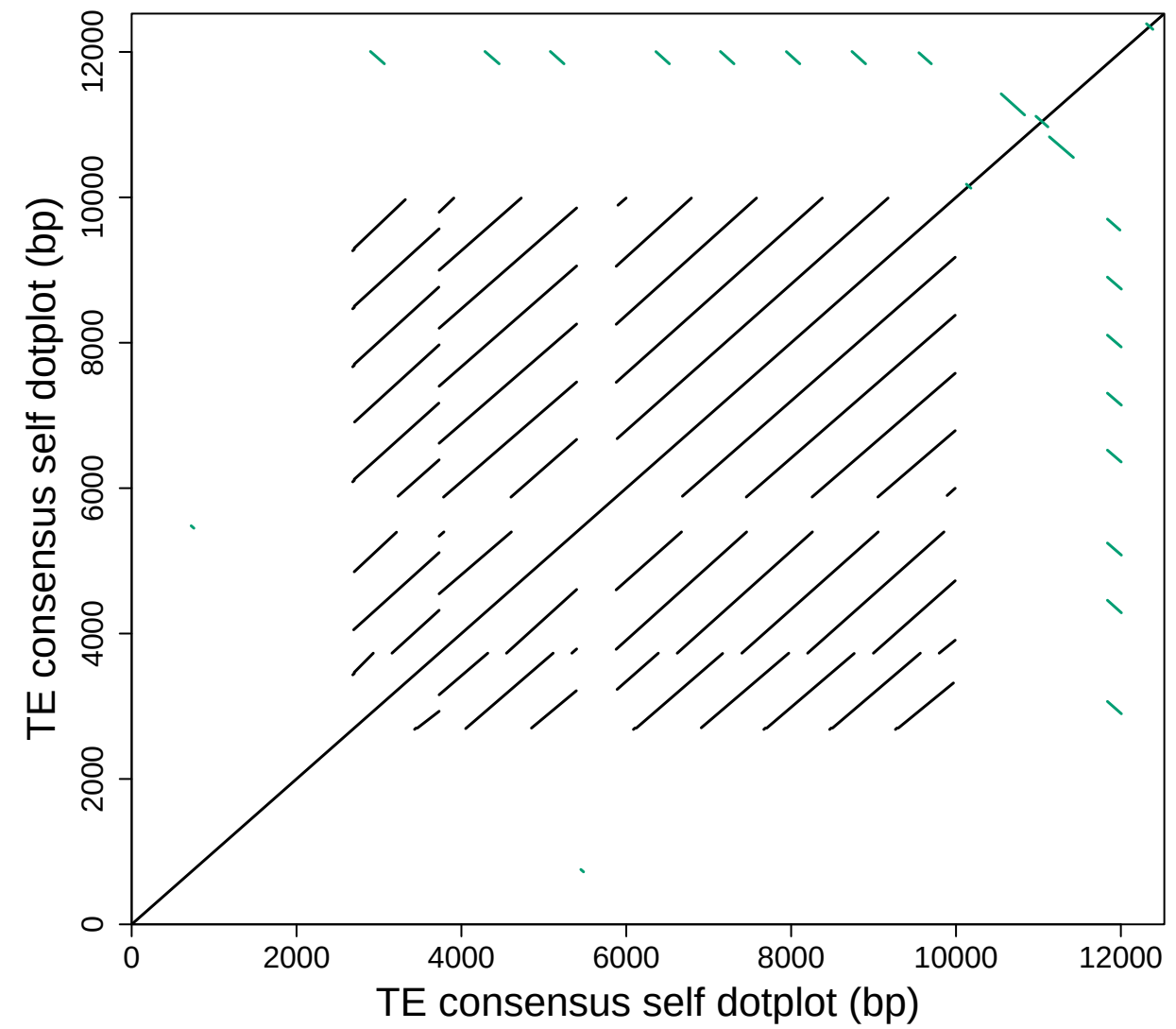
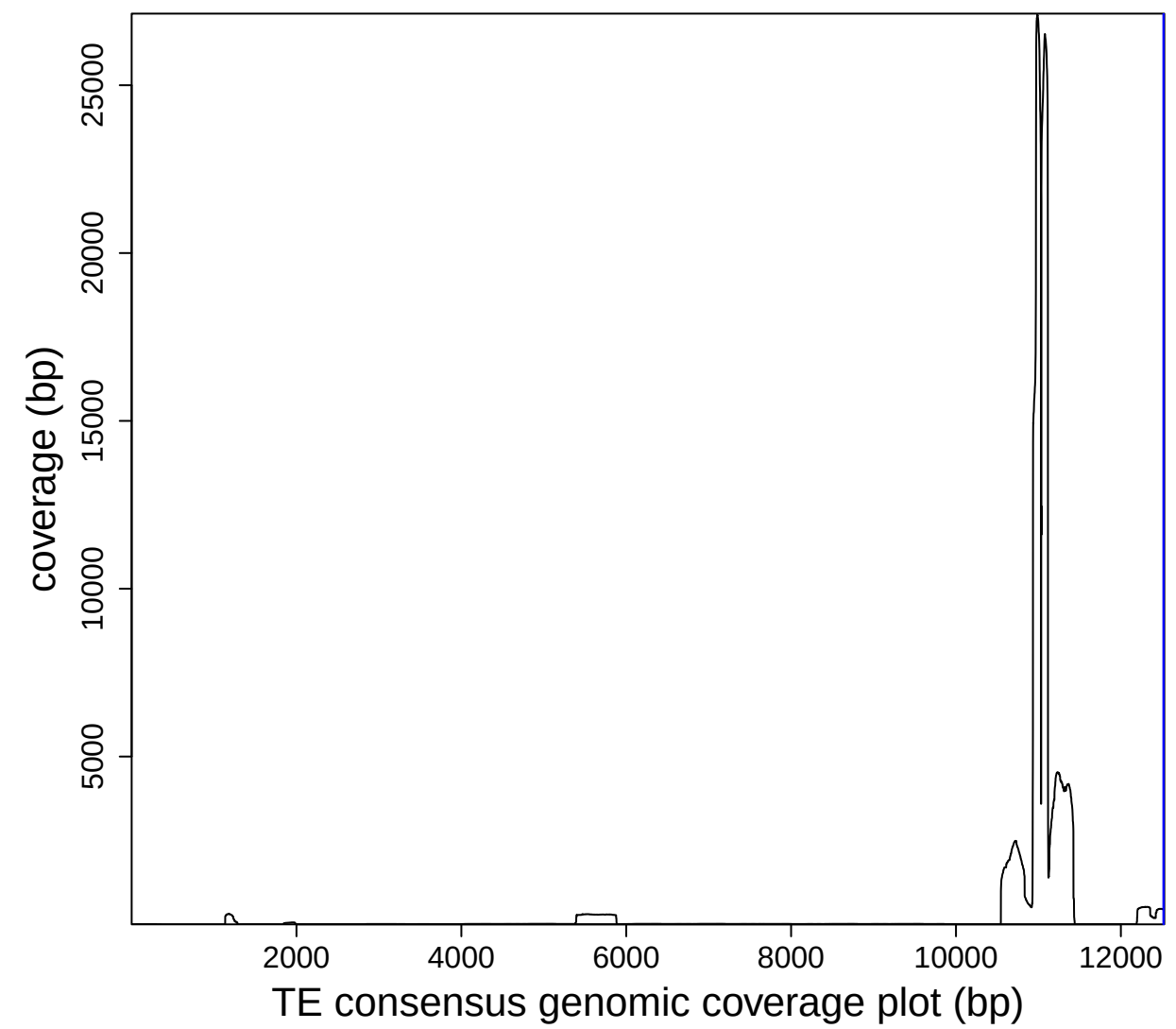
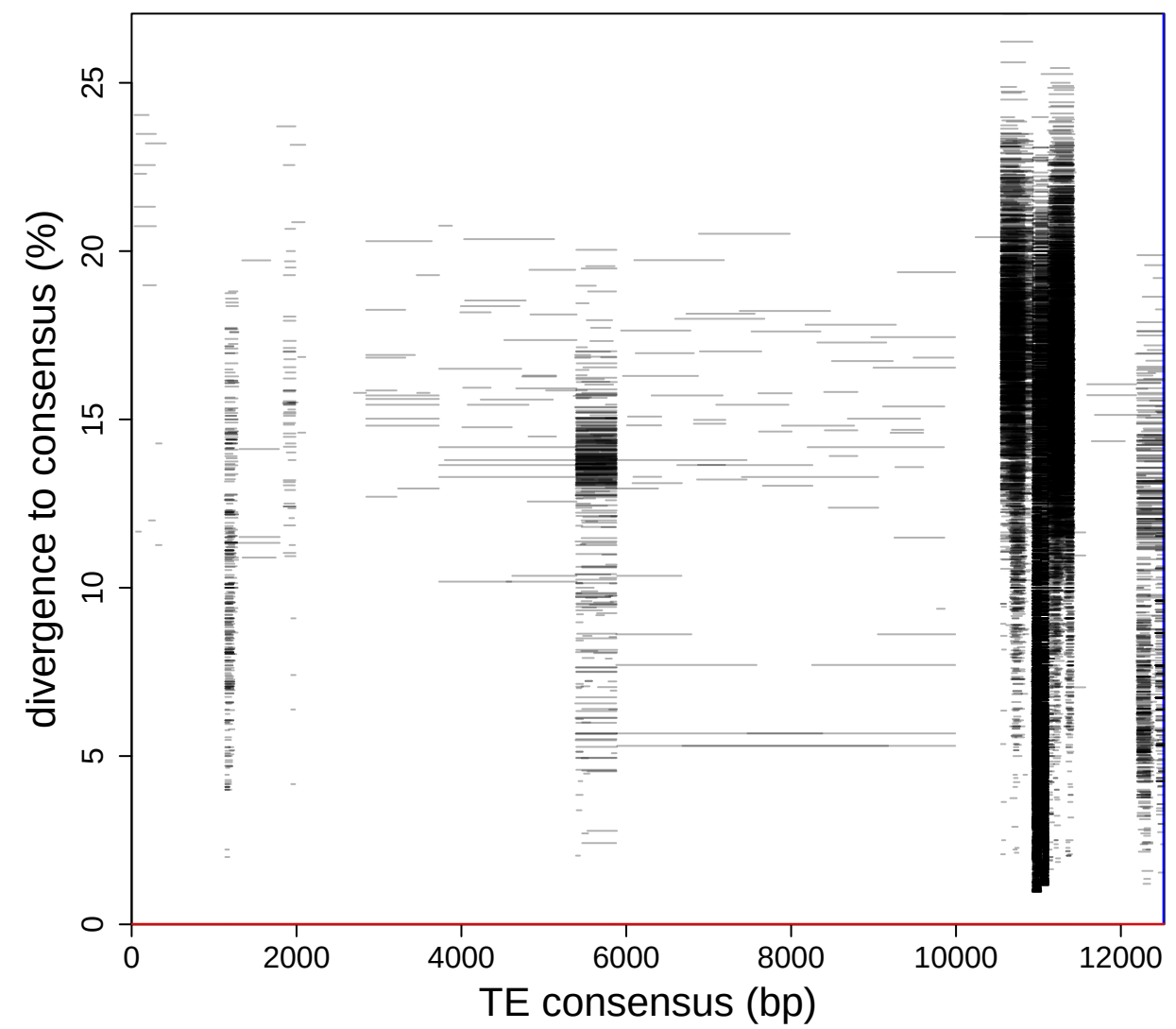
g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.fa_a
size: 12527bp; fragments: 61982; full length: 1 (>=11274.3bp)

After TEtrimmer 12527 bp

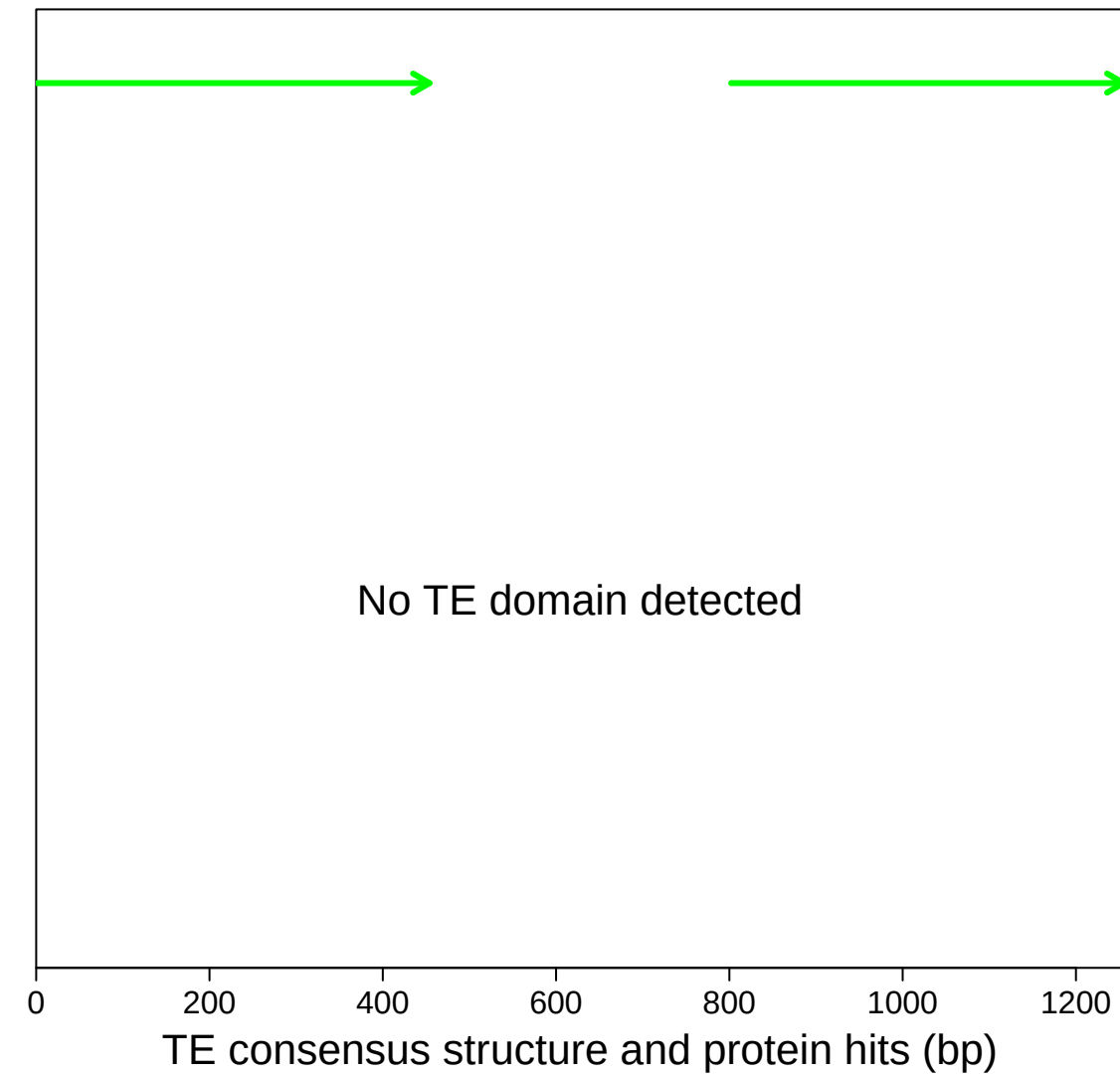
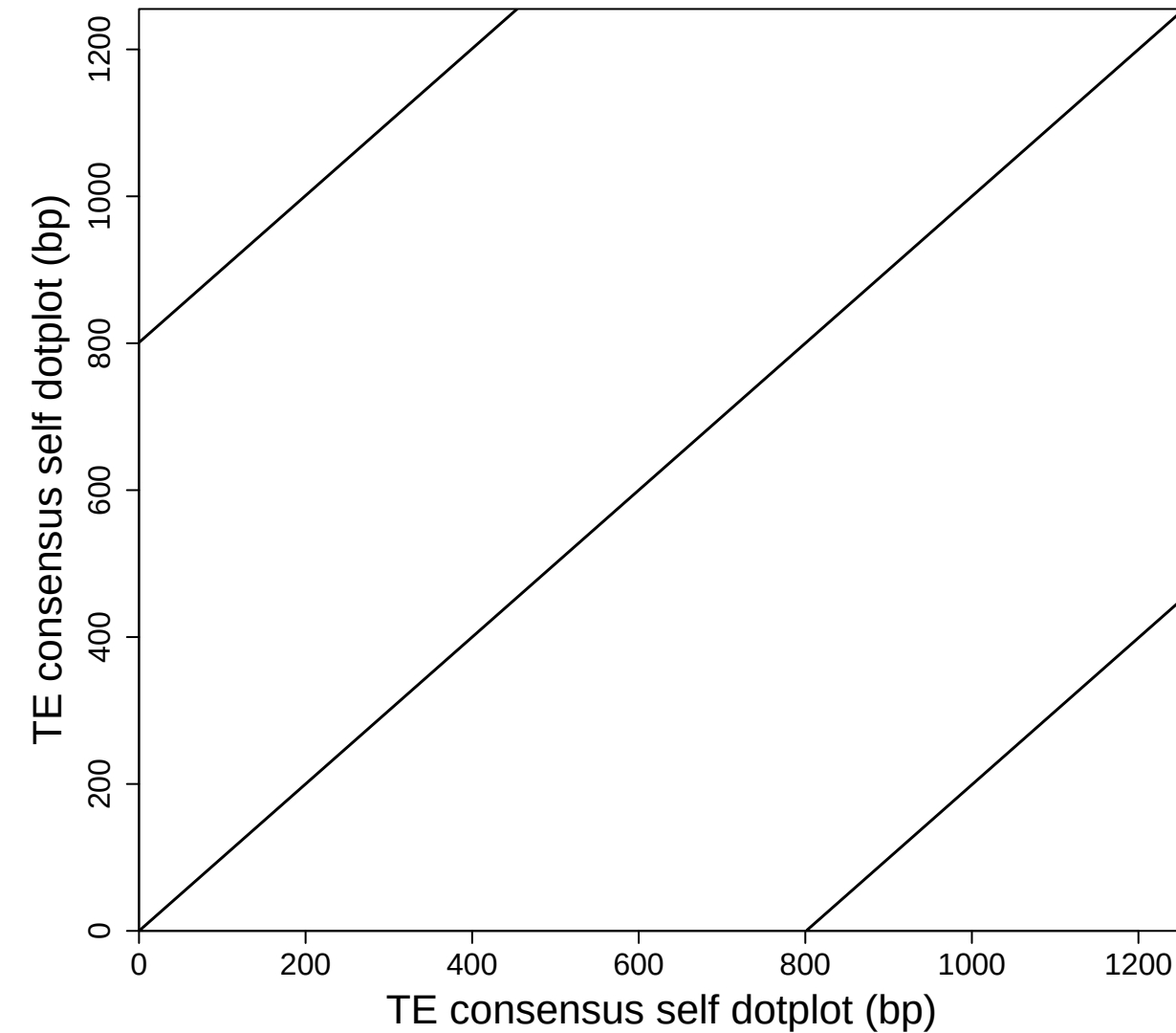
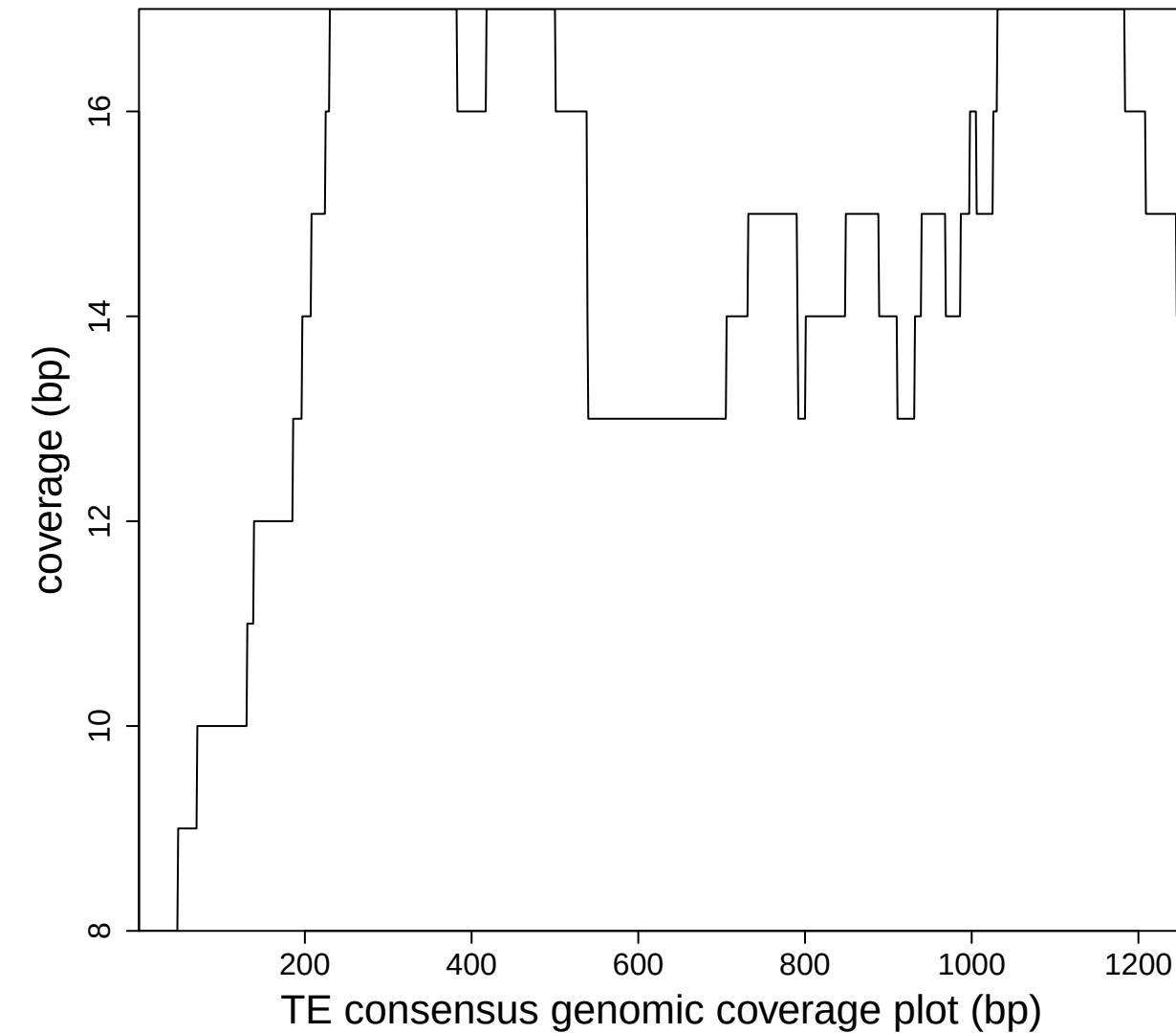
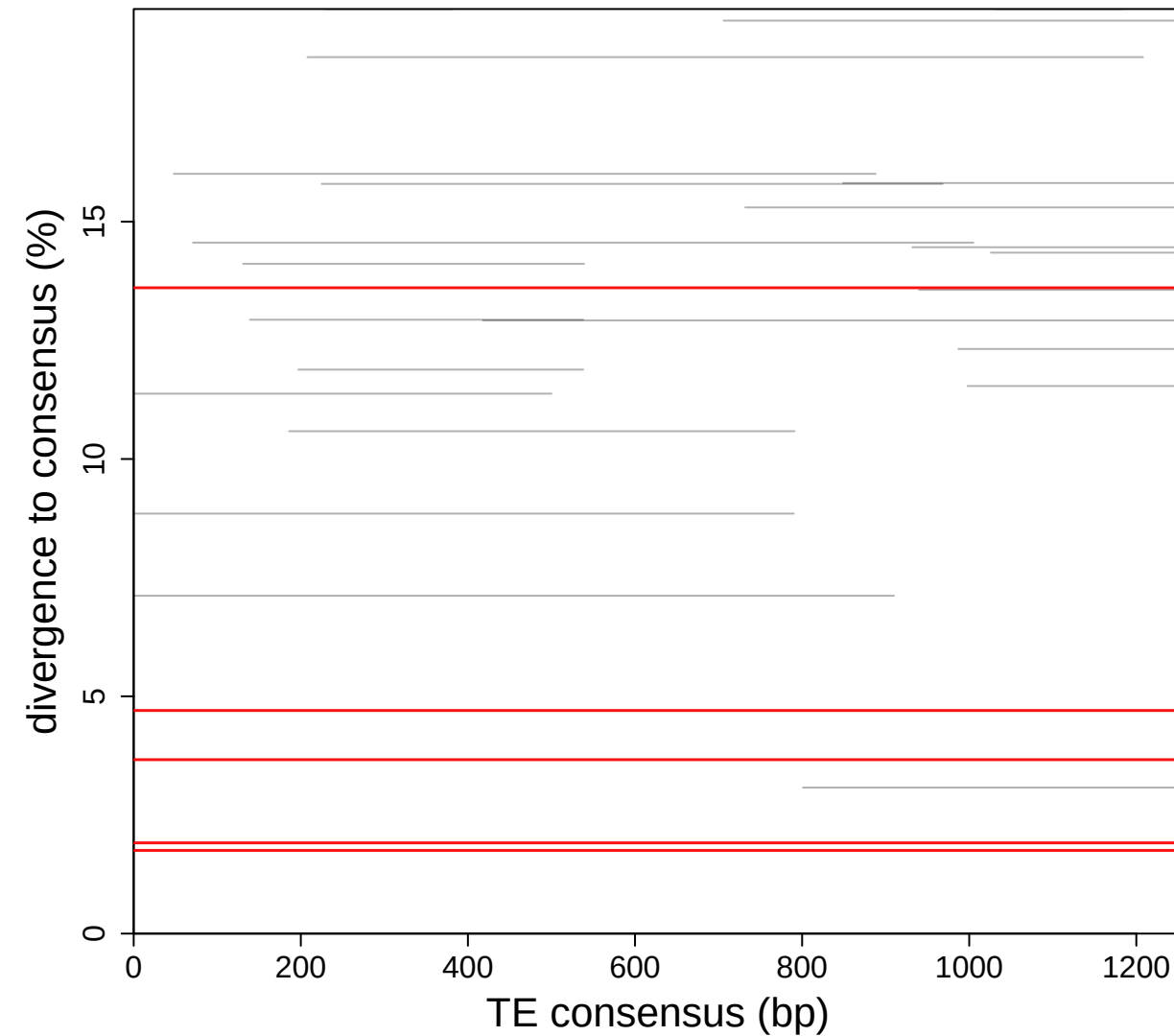


f.bed_g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcl
size: 12527bp; fragments: 61982; full length: 1 (>=11274.3bp)

After TEtrimmer Extended plot Blue lines are boundaries

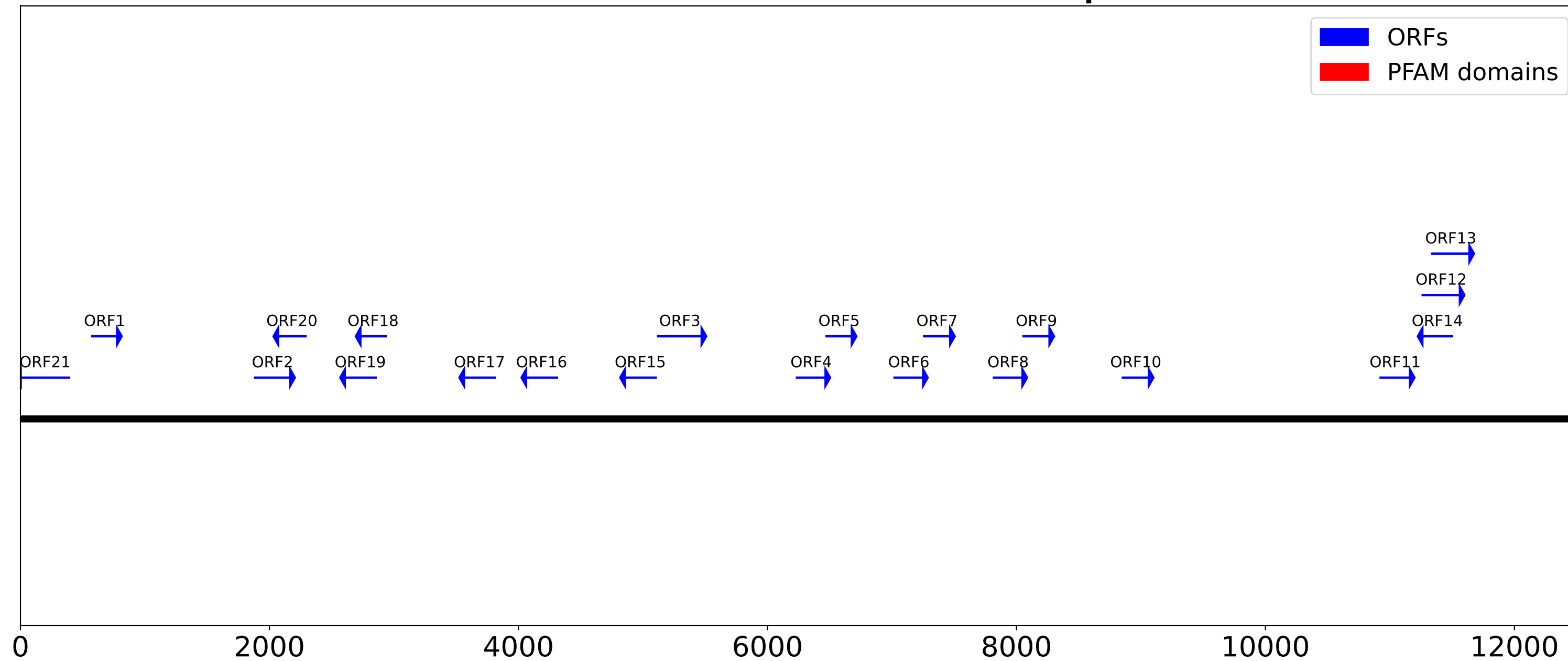


TE: rnd_1_family_819
size: 1255bp; fragments: 28; full length: 5 (≥ 1129.5 bp)

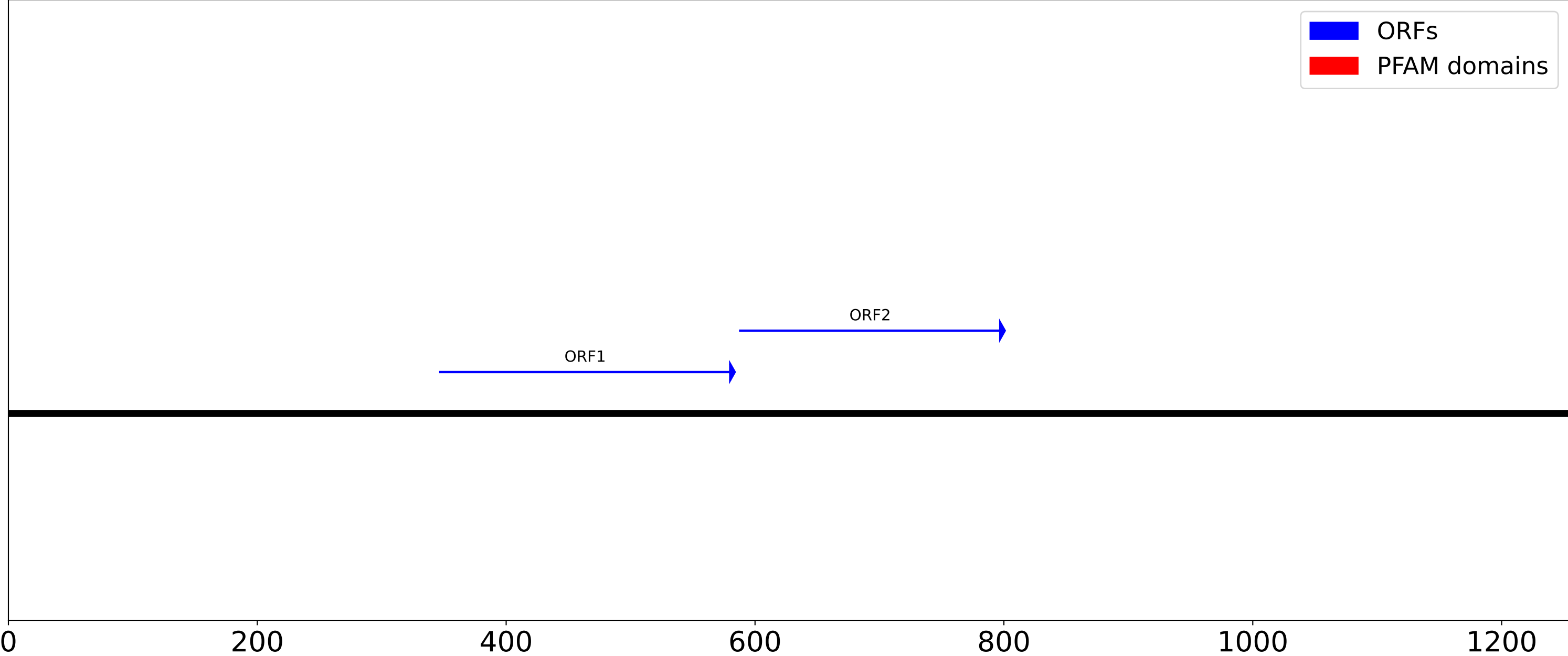


Before TETrimmer 1255 bp

After TEtrimmer ORF and PFAM domain plot



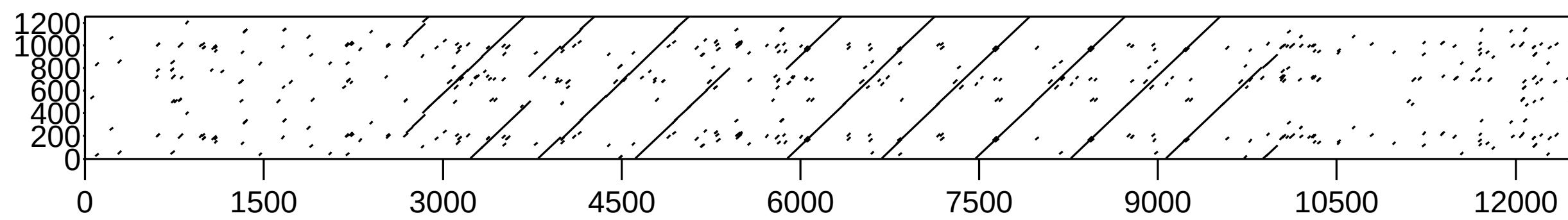
Before TEtrimmer ORF and PFAM domain plot



Dotplot

(windowsize = 25, threshold = 50.00 09/10/25)

consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)