

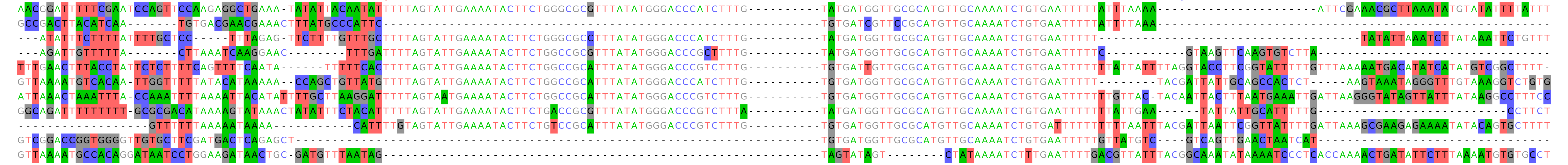
Start crop Point



End crop Point

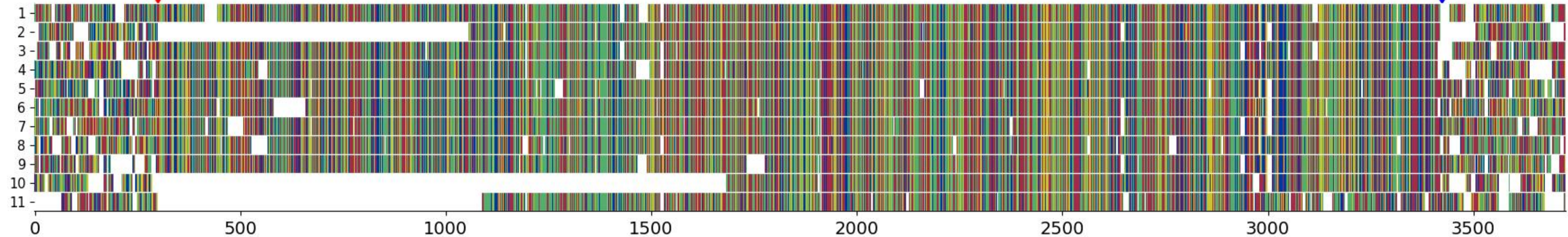


MSA length = 3125



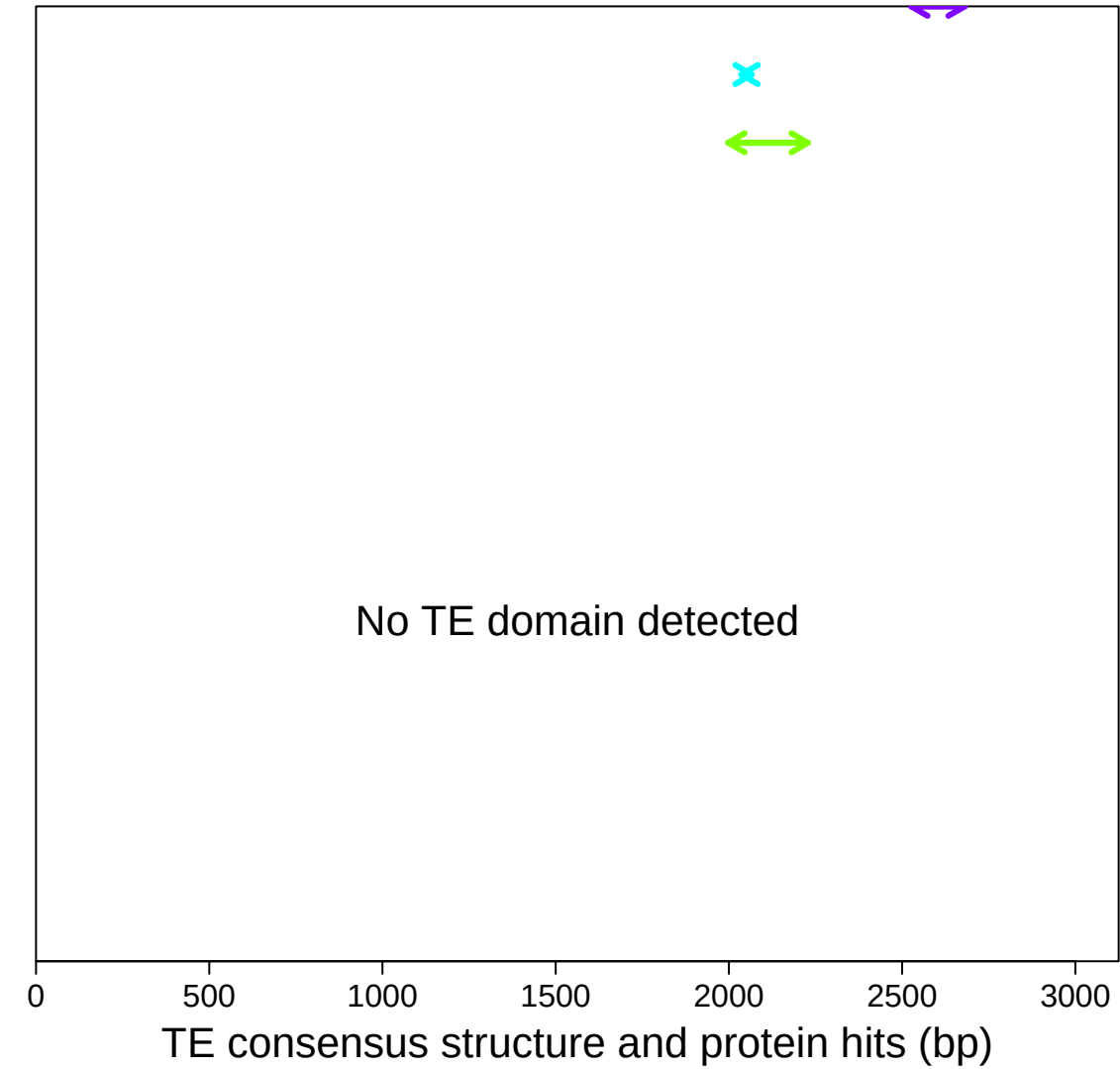
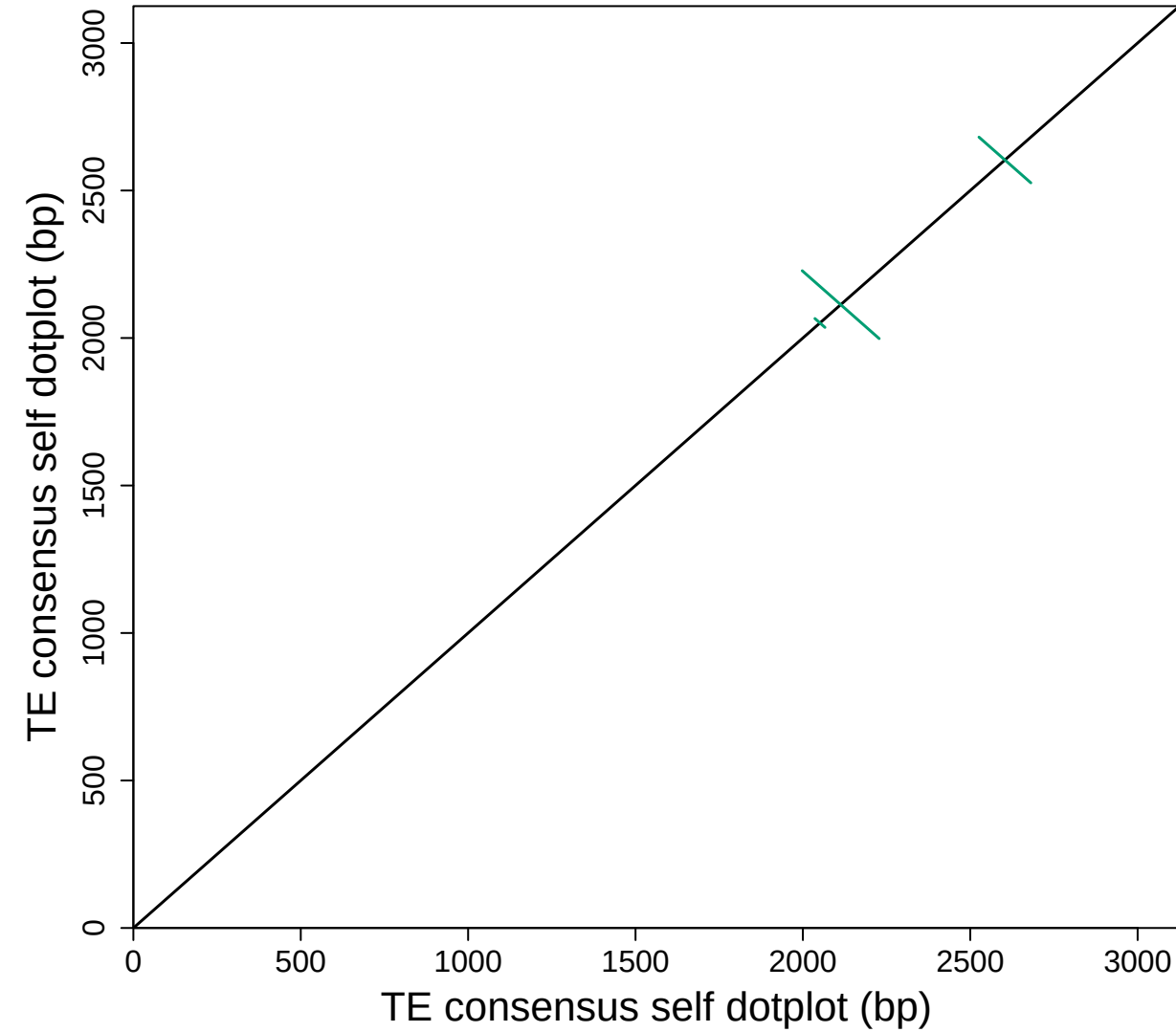
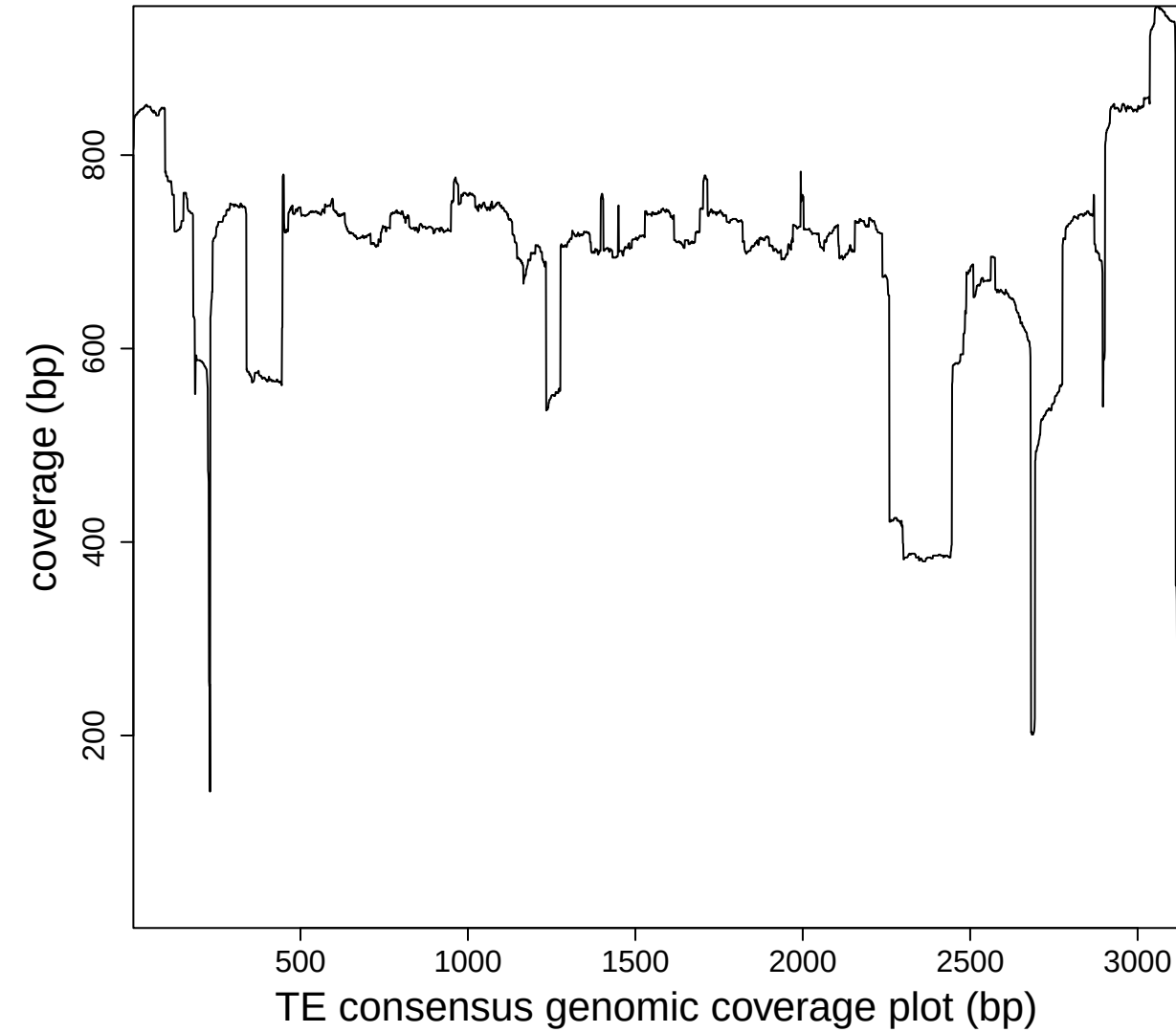
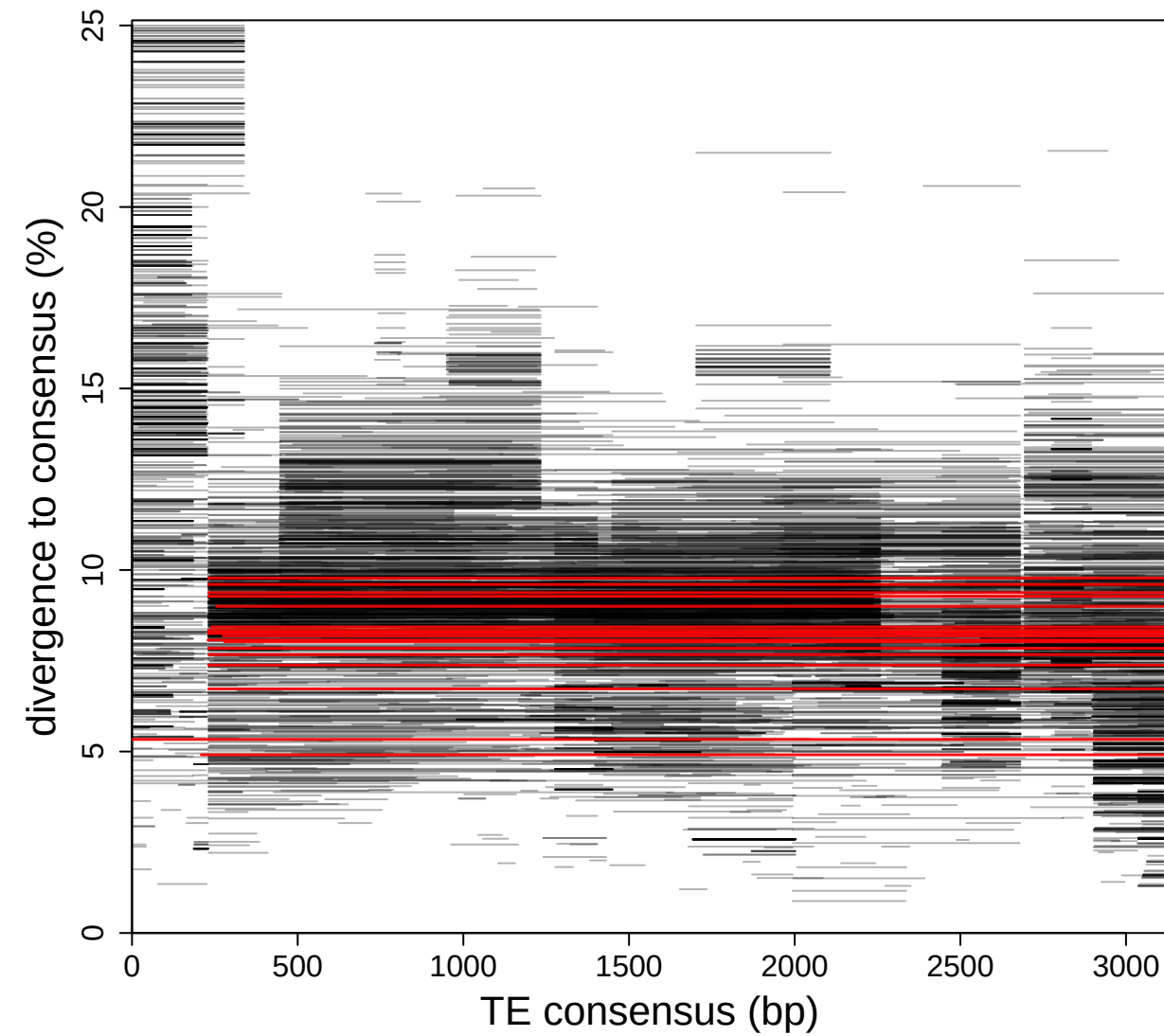
Start crop Point

End crop Point



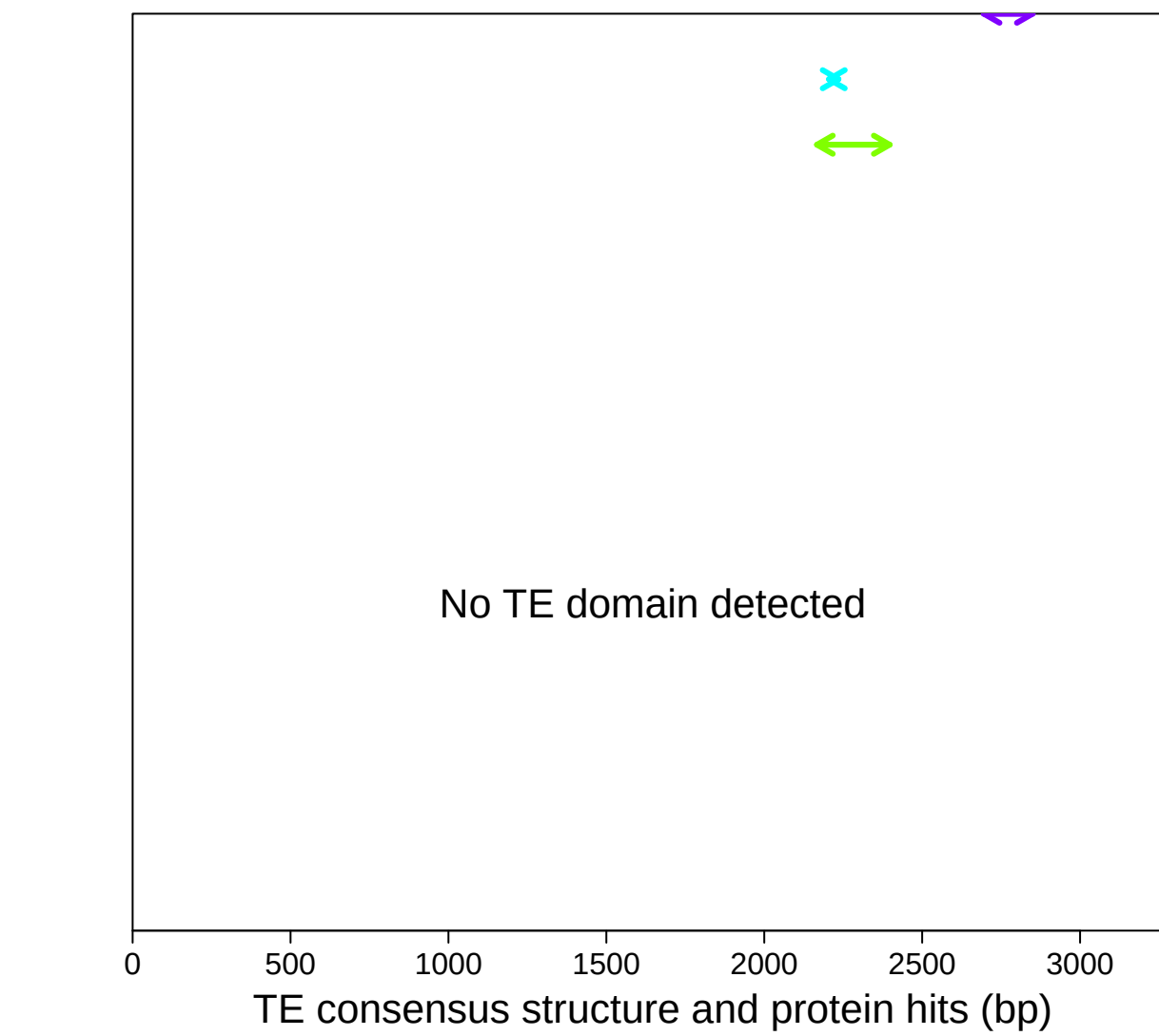
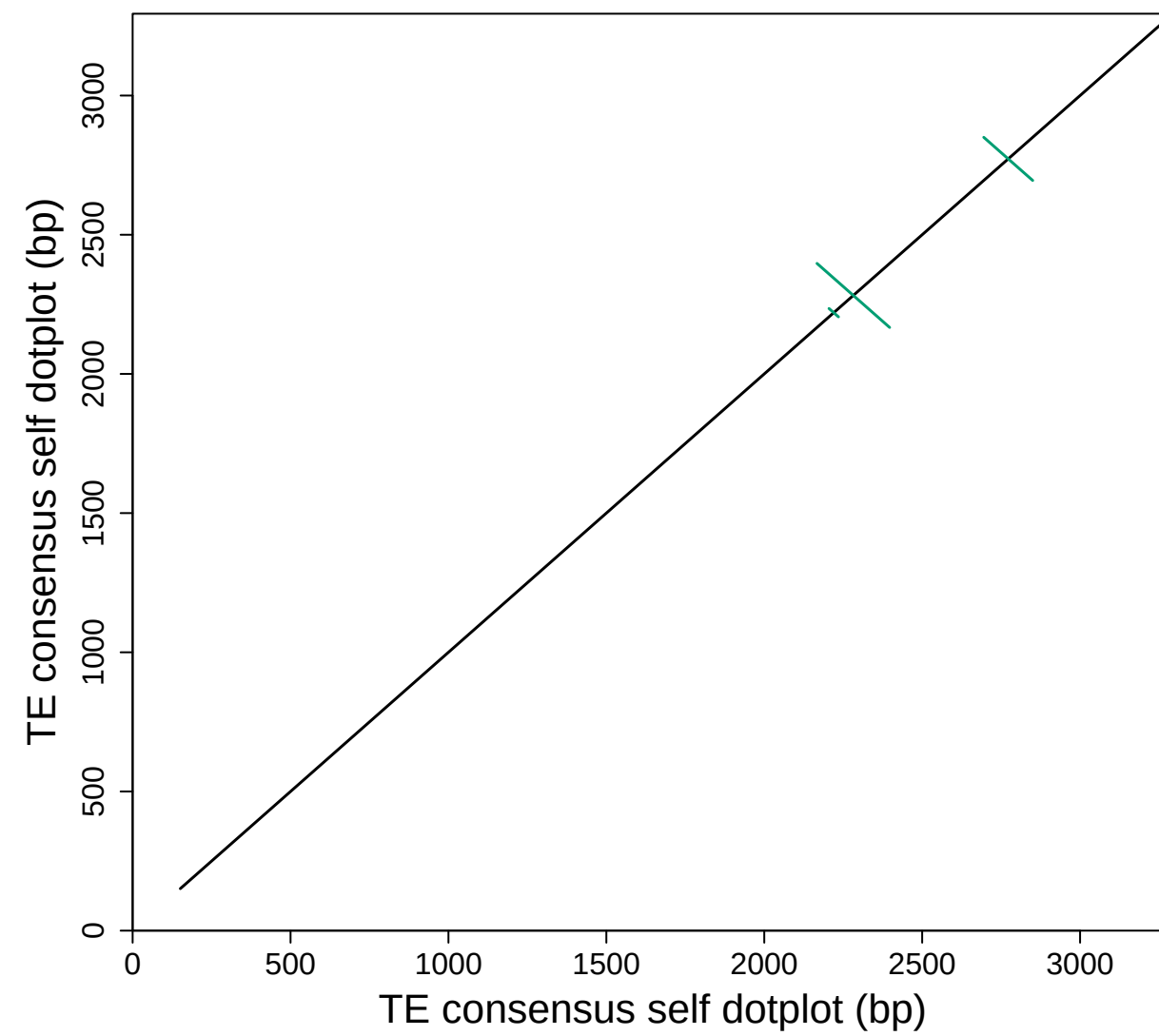
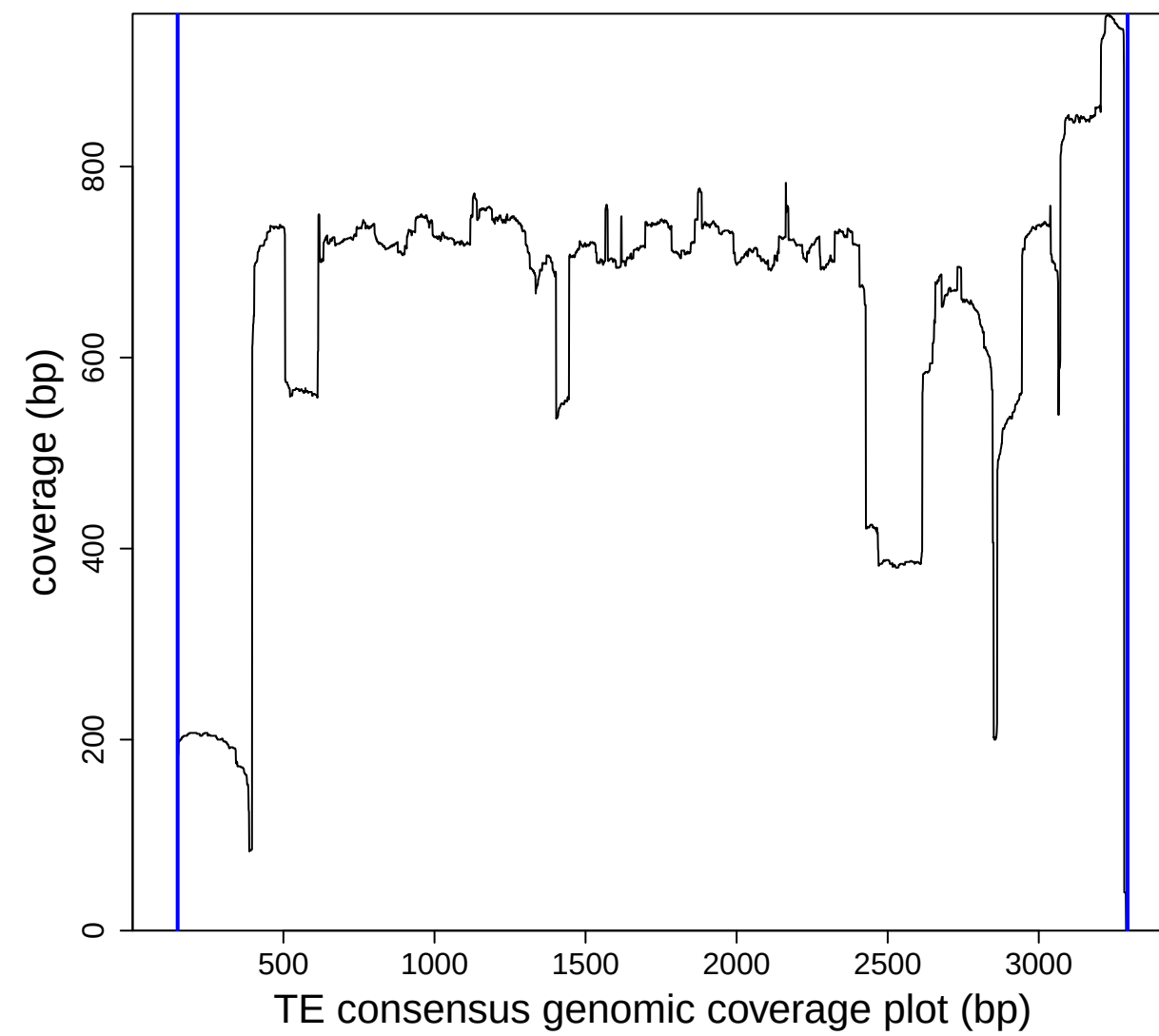
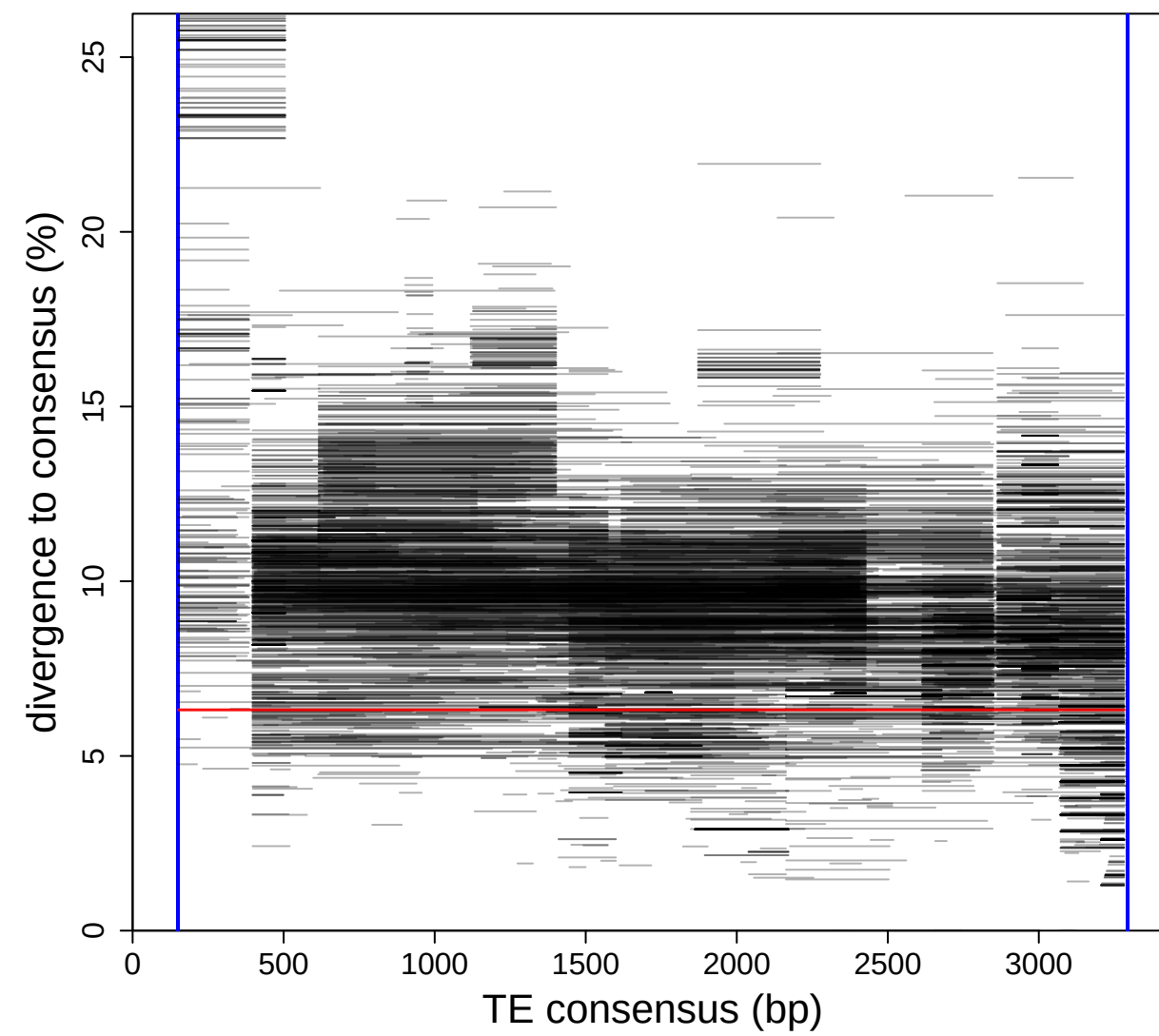
g_3.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.fa_a
size: 3125bp; fragments: 4677; full length: 18 (>=2812.5bp)

After TEtrimmer 3125 bp

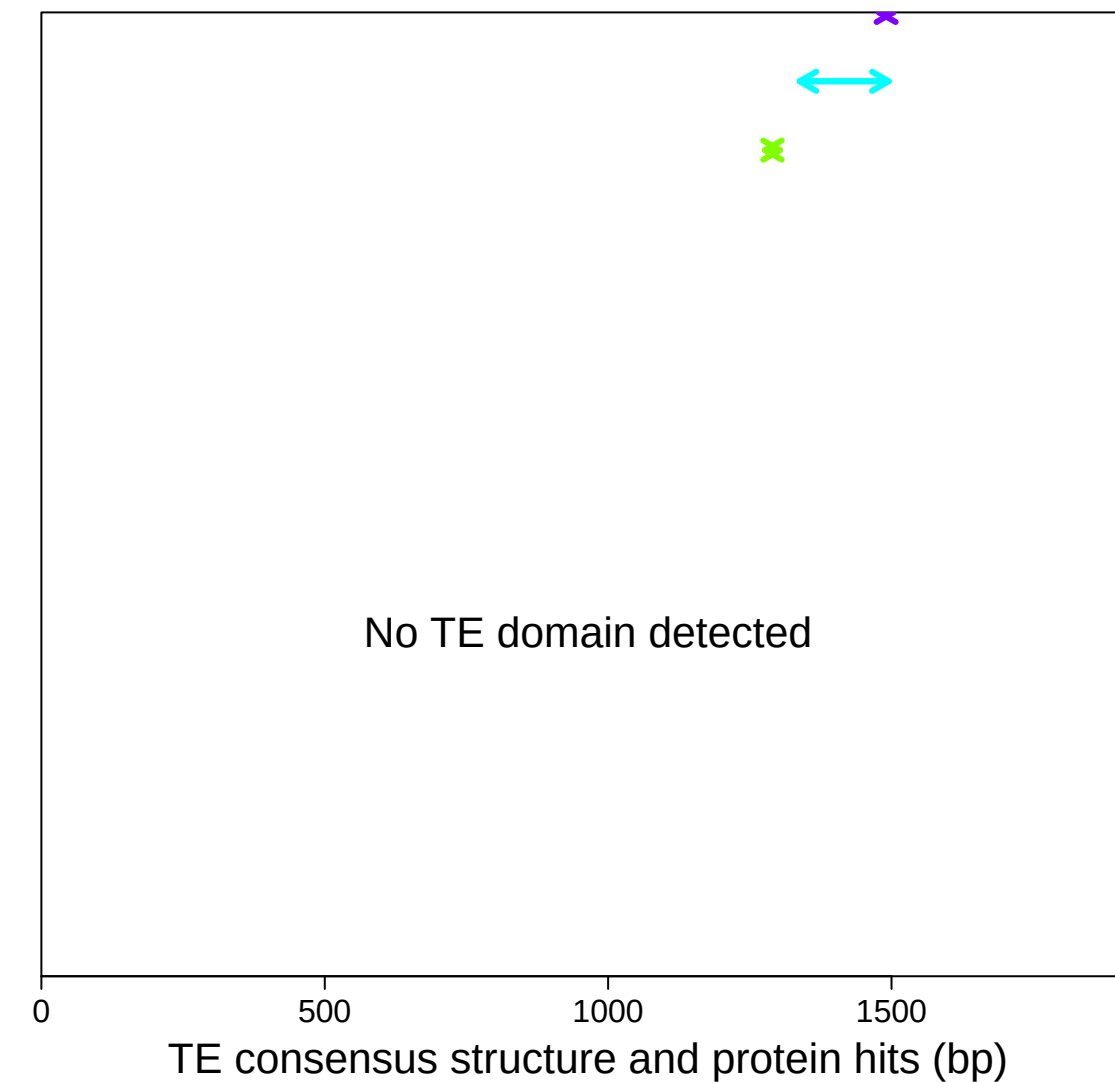
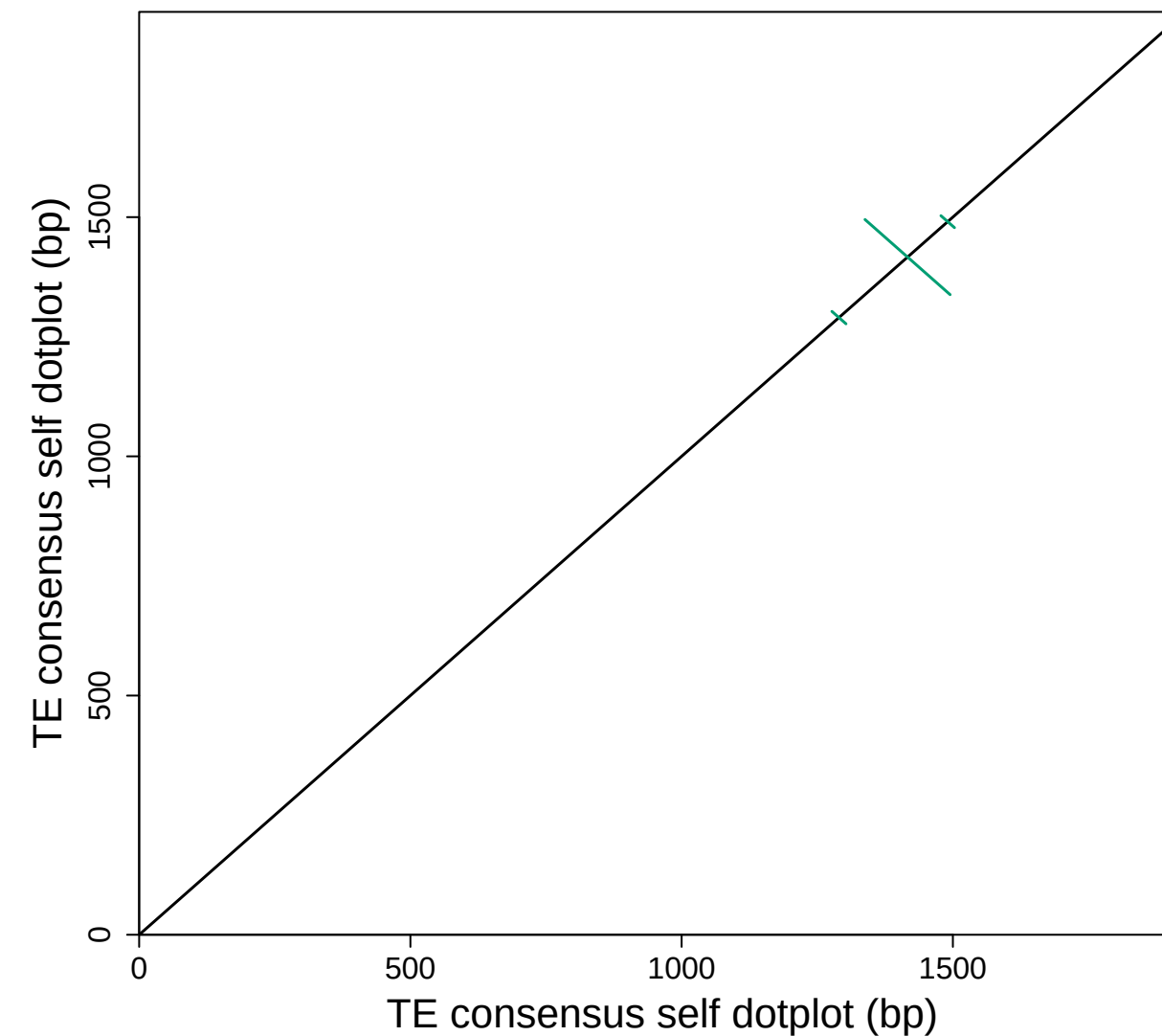
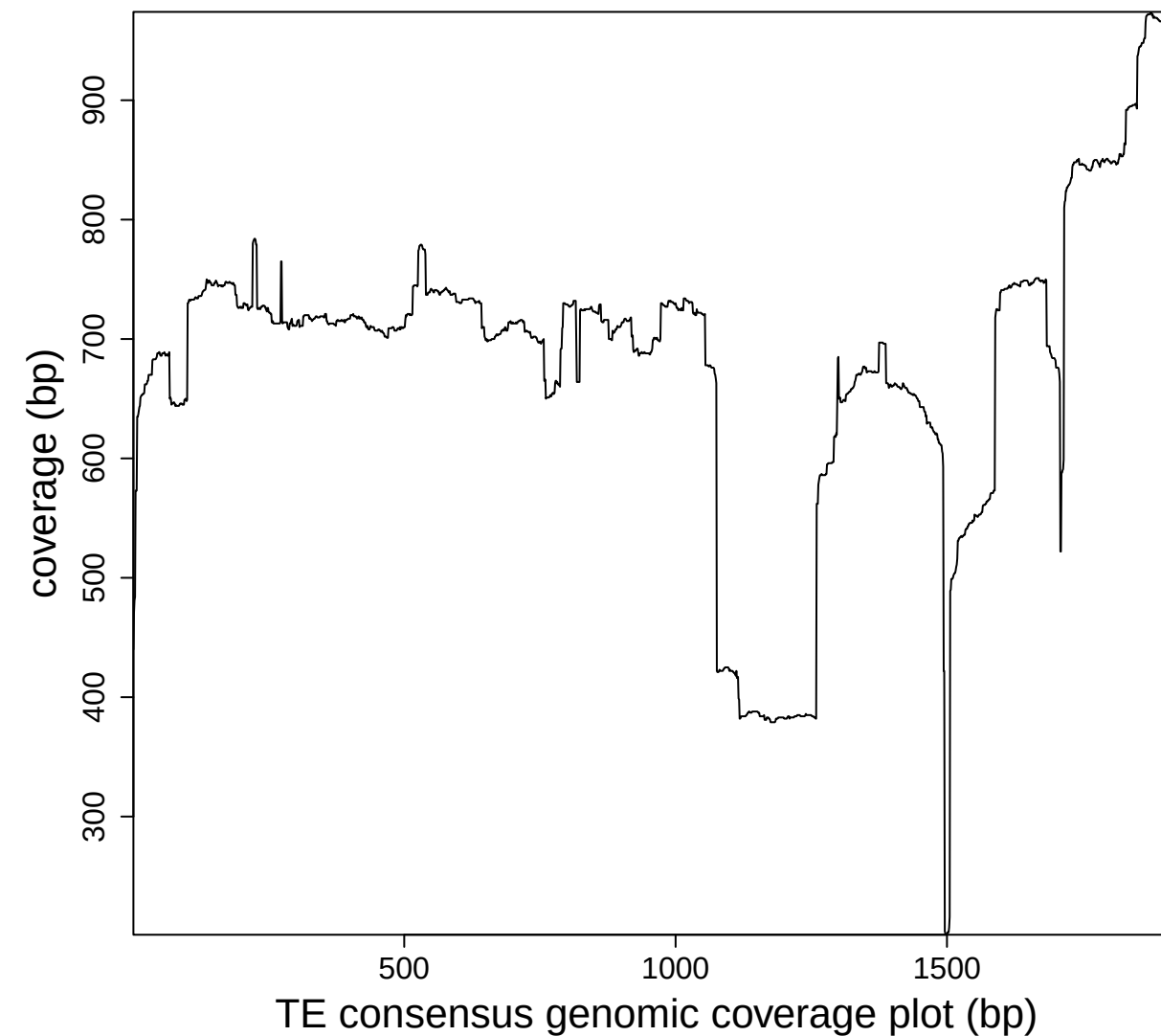
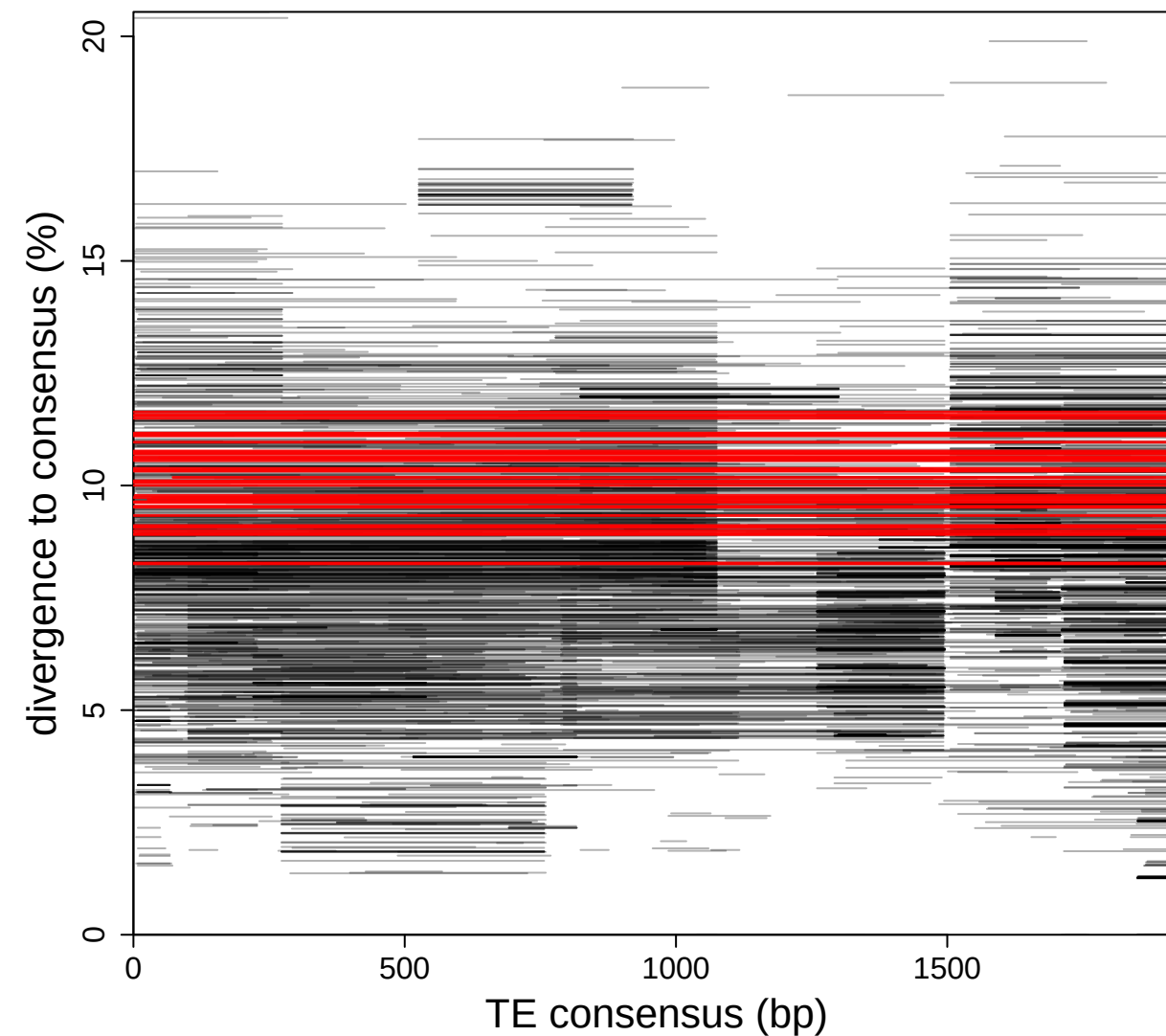


f.bed_g_3.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcl
size: 3444bp; fragments: 3947; full length: 1 (>=3099.6bp)

After TEtrimmer Extended plot Blue lines are boundaries

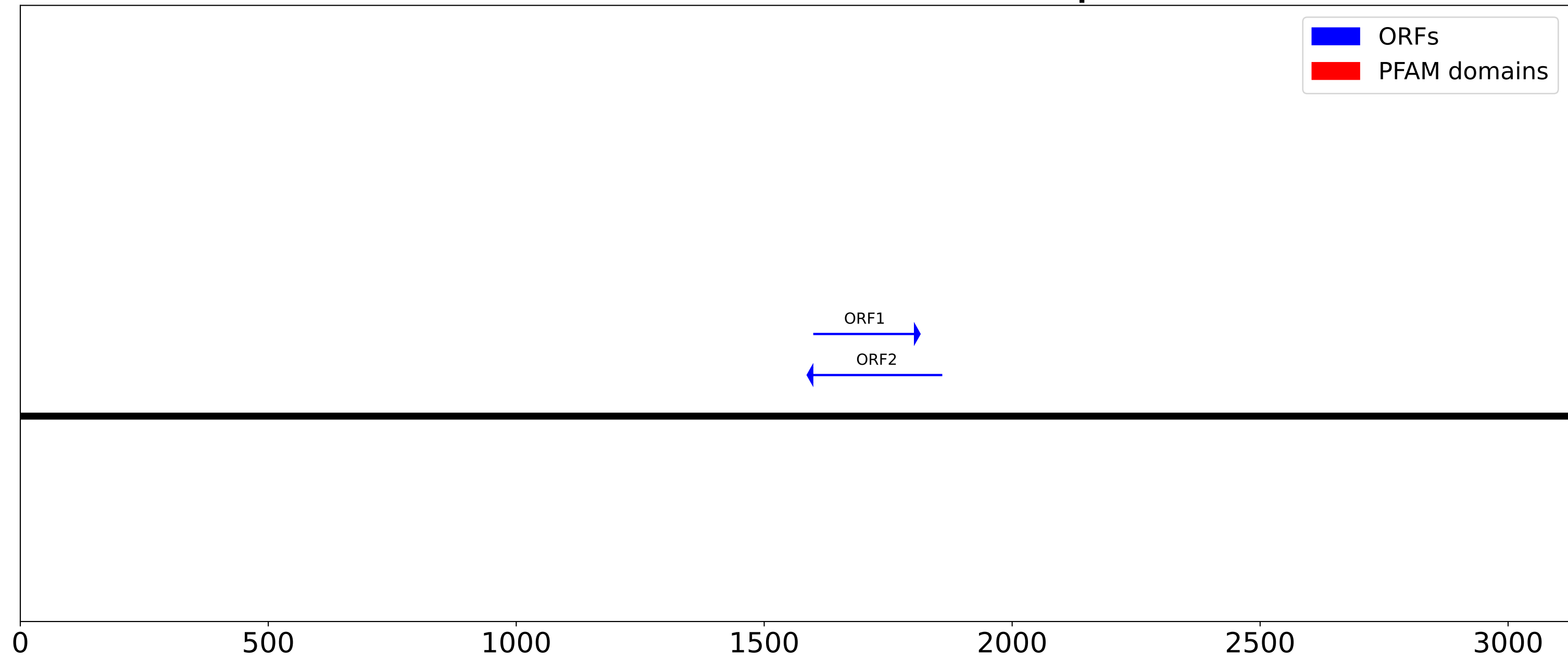


TE: rnd_4_family_557
size: 1929bp; fragments: 3075; full length: 42 (≥ 1736.1 bp)

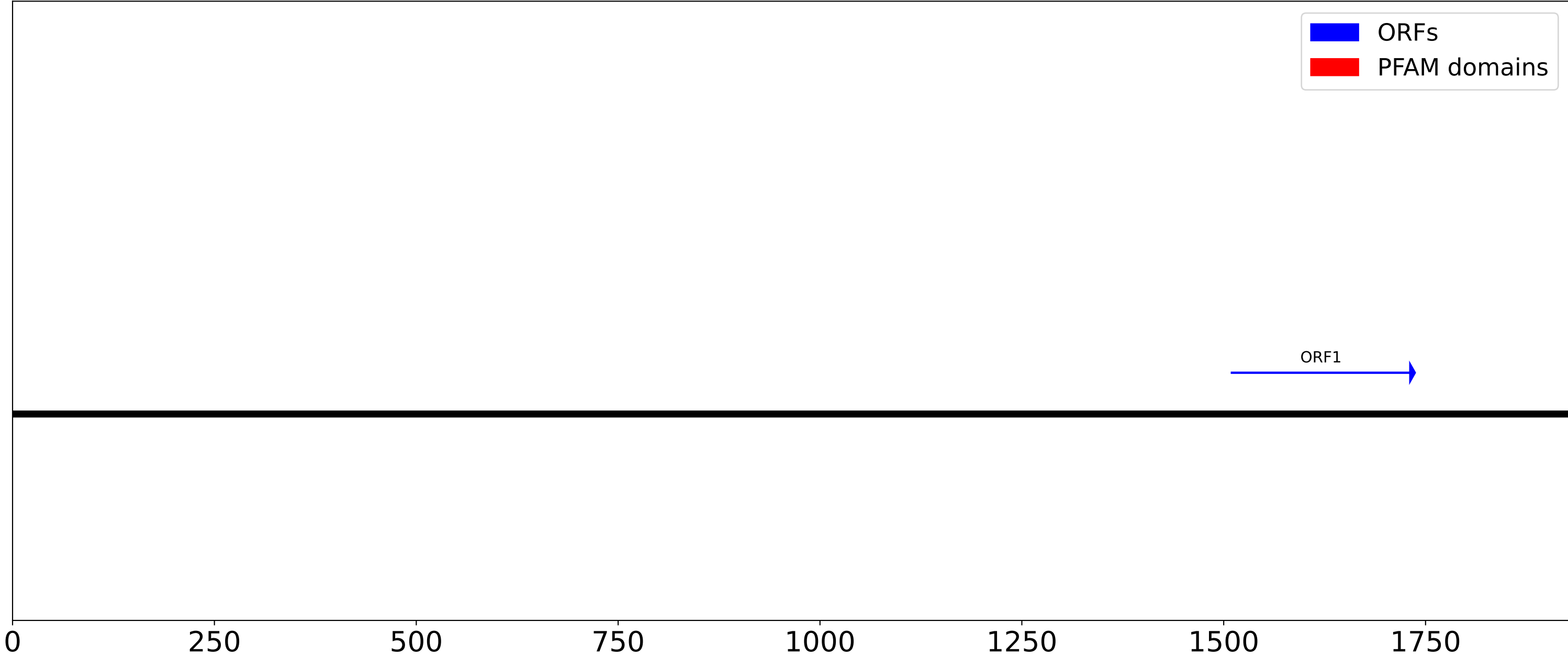


Before TEtrimmer 1929 bp

After TEtrimmer ORF and PFAM domain plot



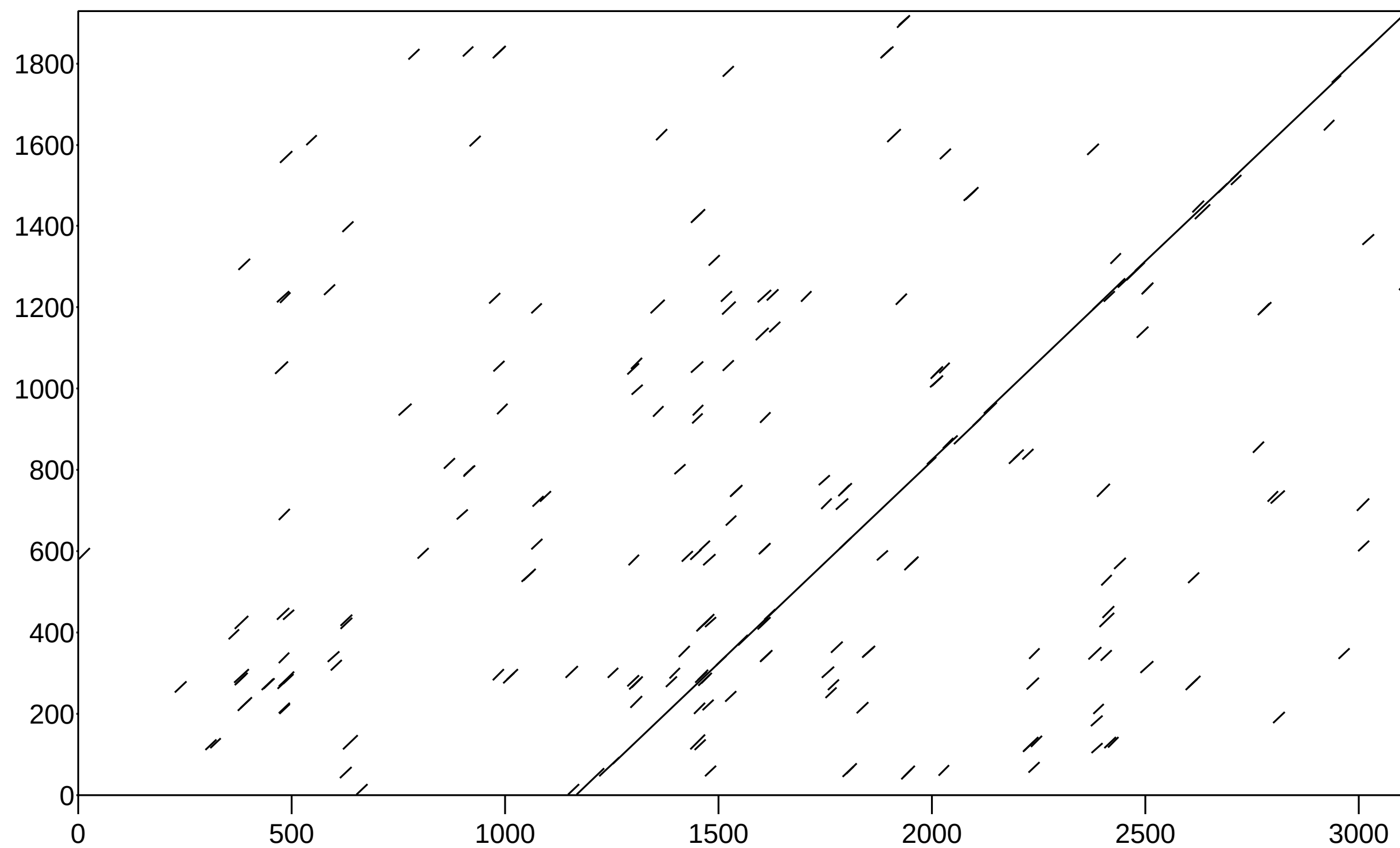
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)