

MSA length = 3884

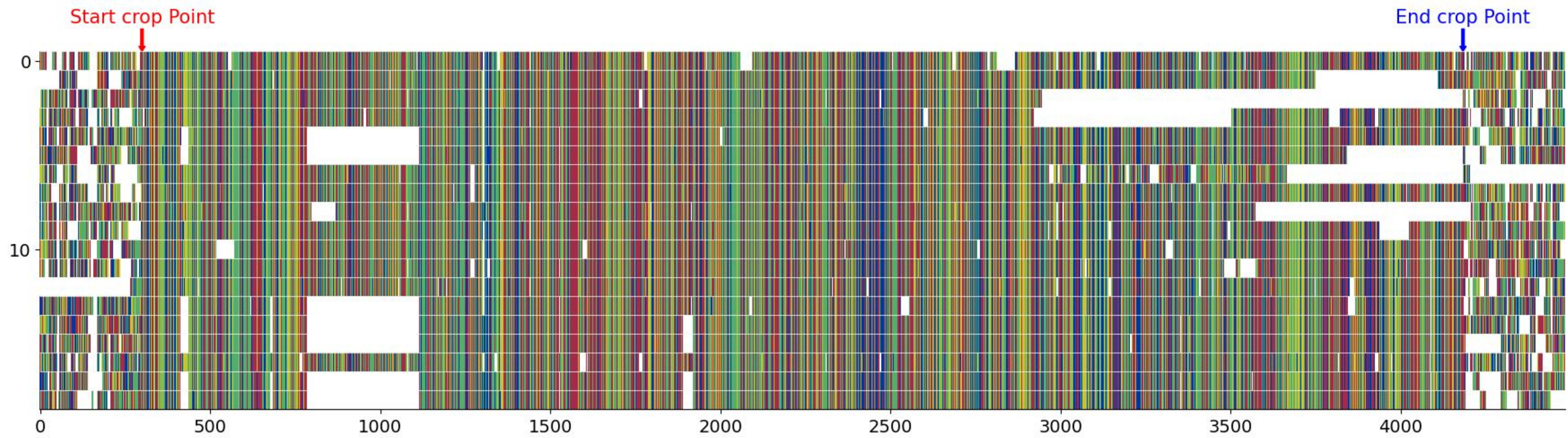
Start crop Point

End crop Point



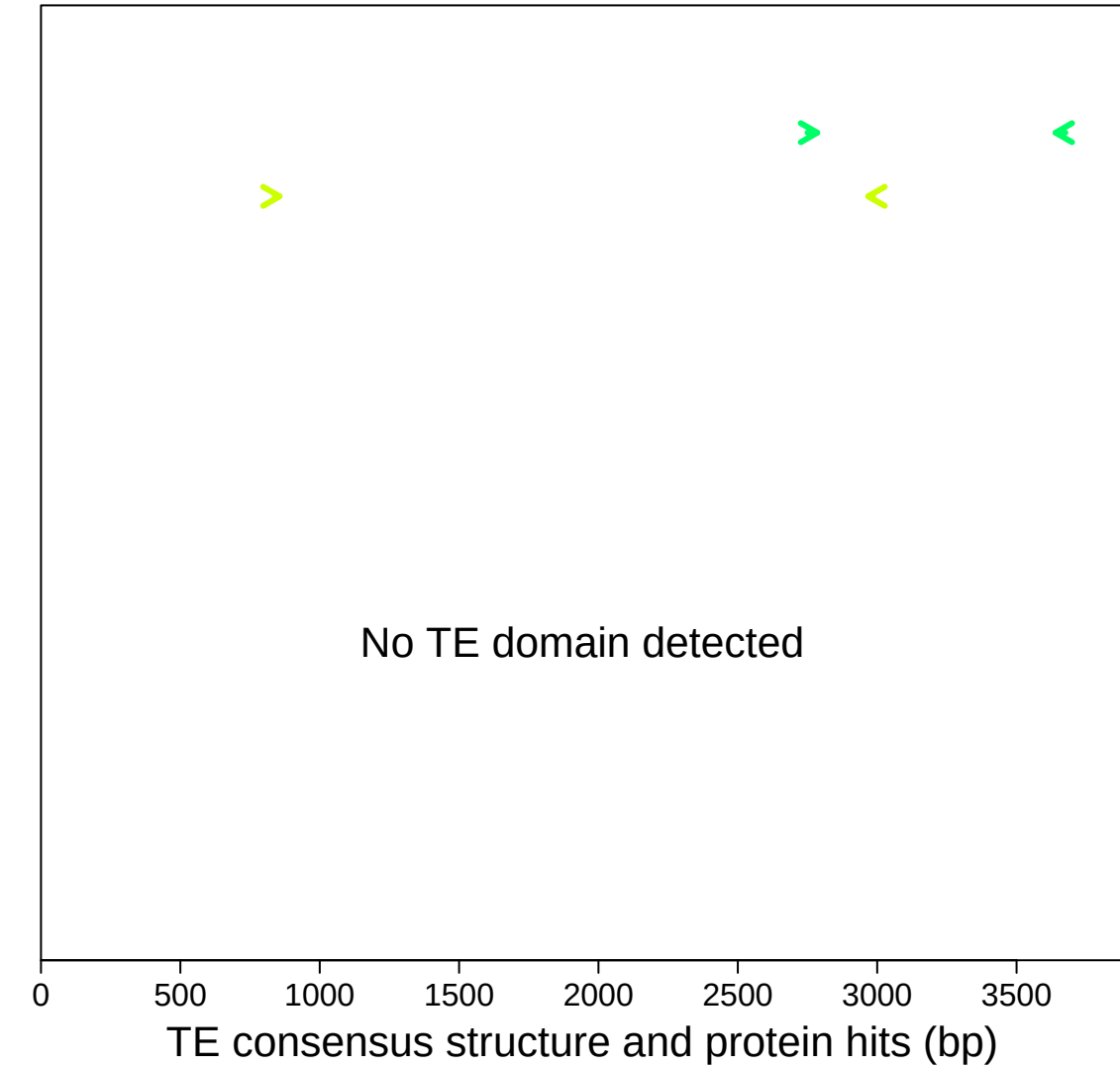
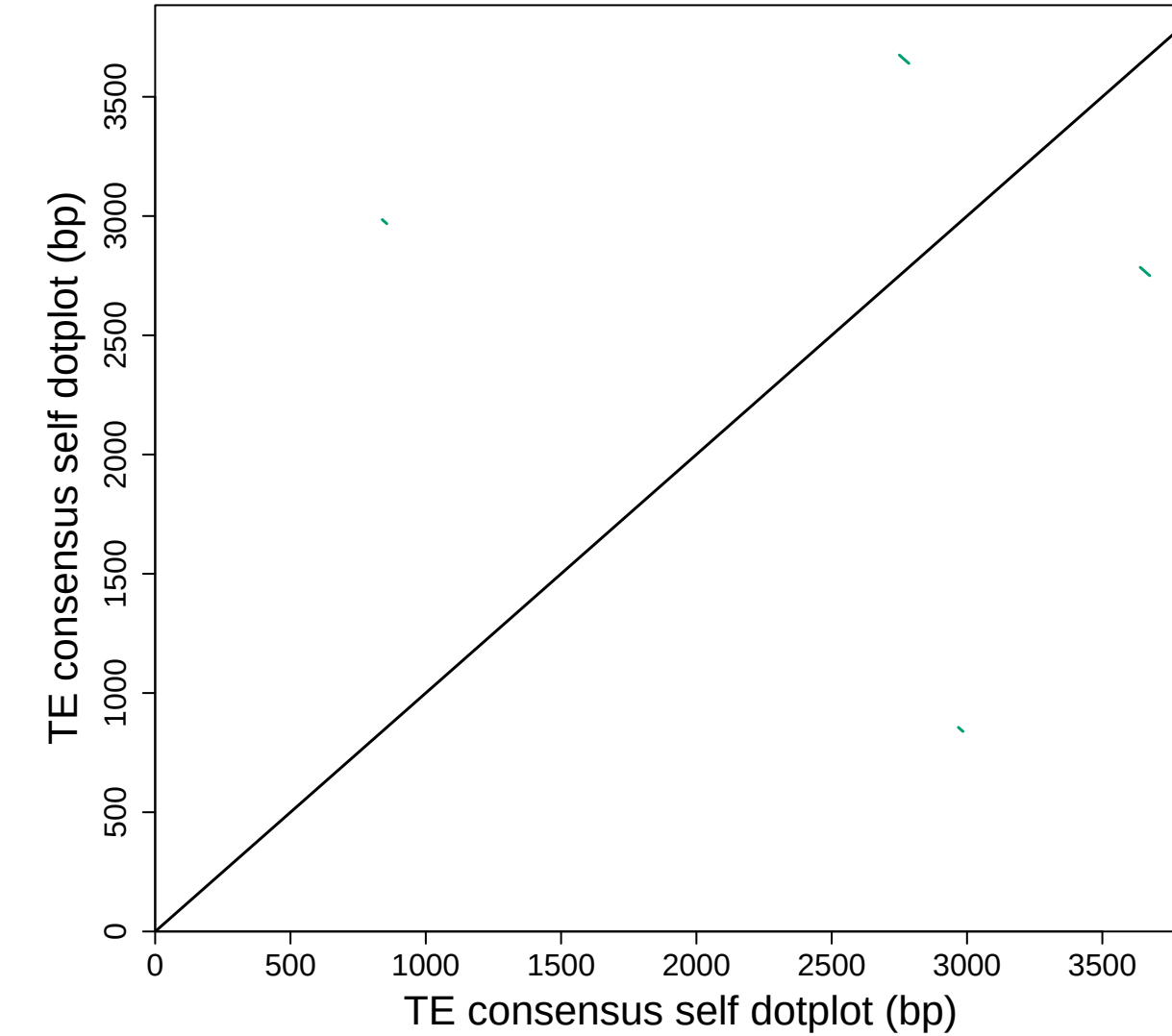
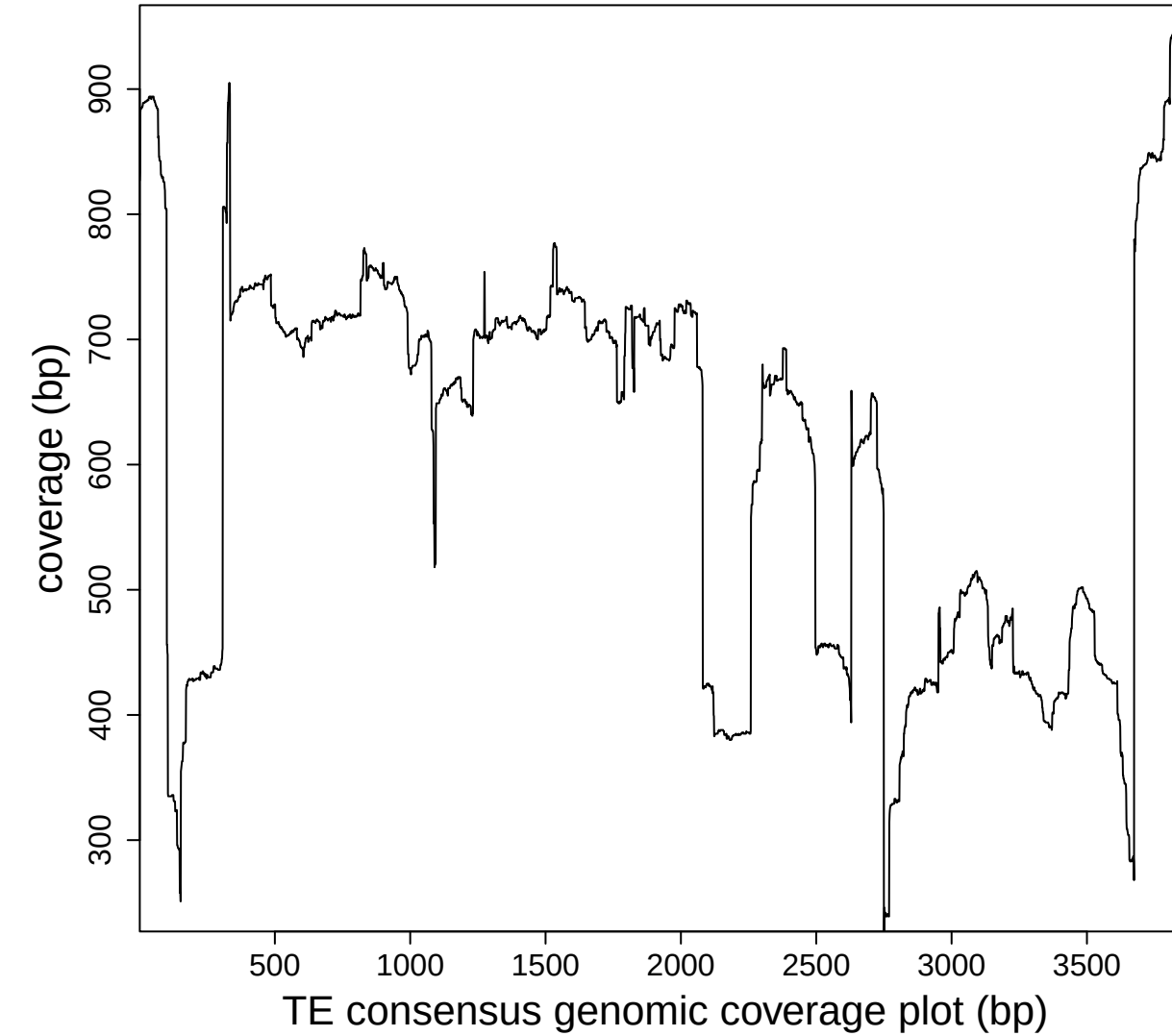
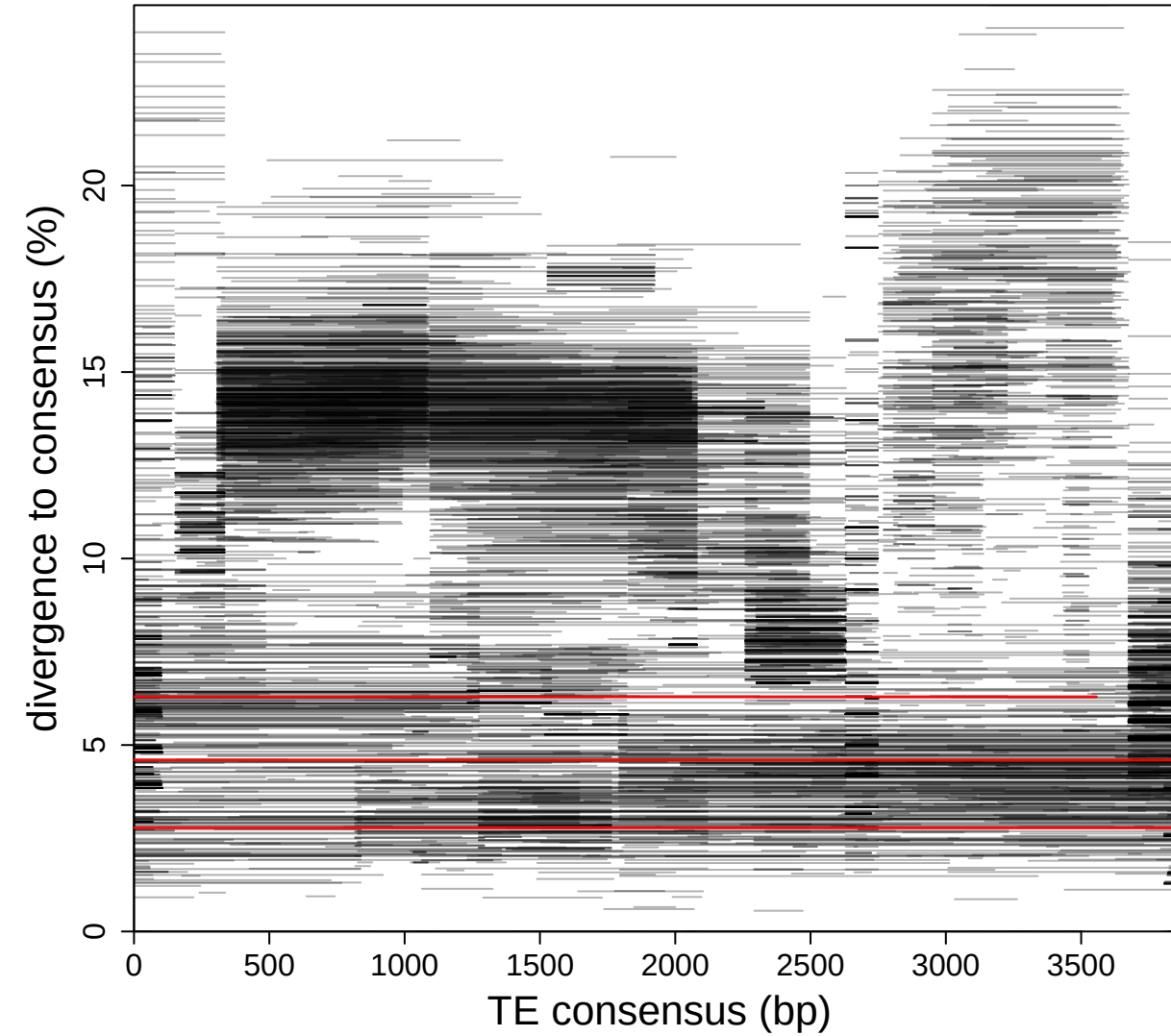
Start crop Point

End crop Point



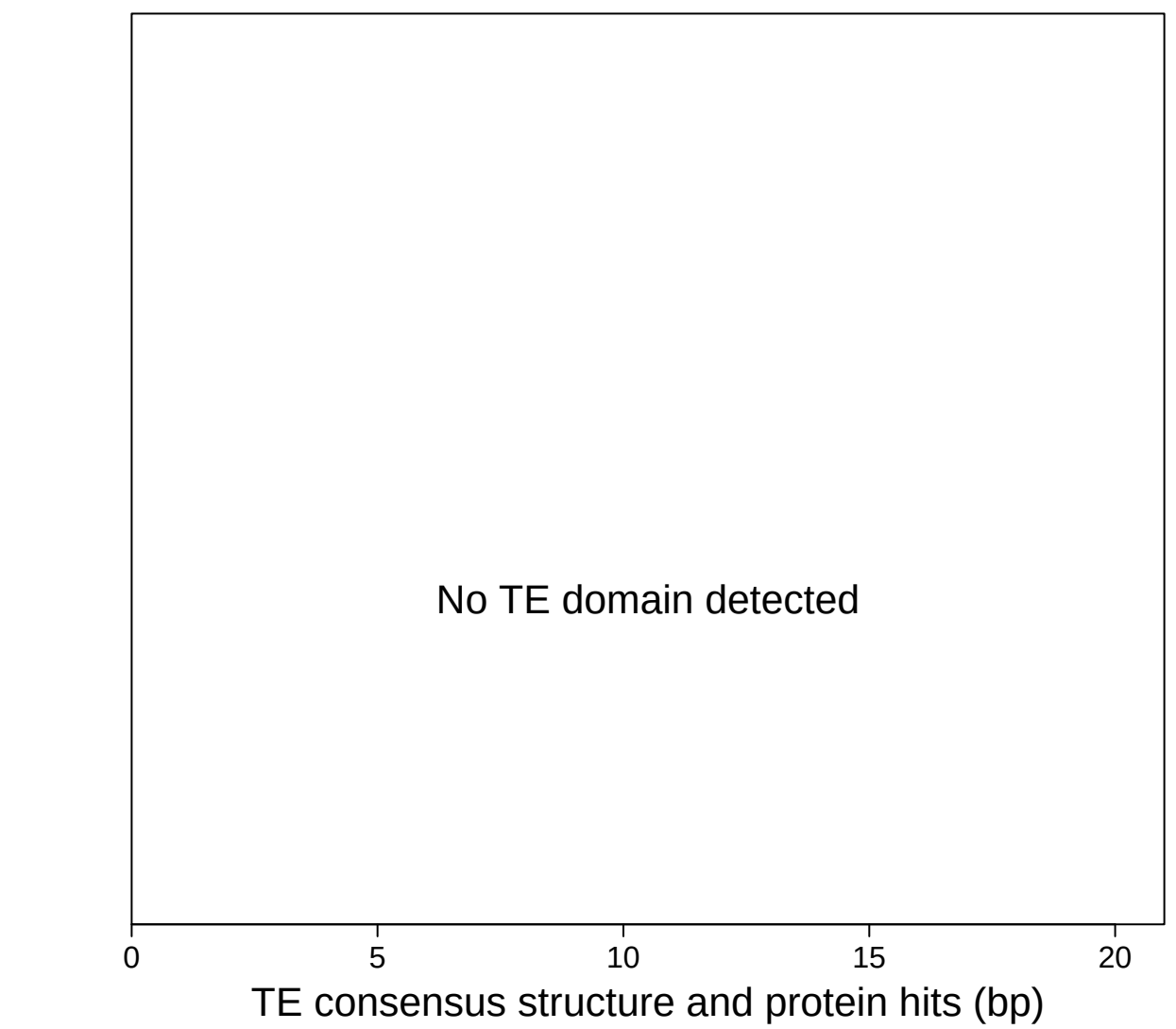
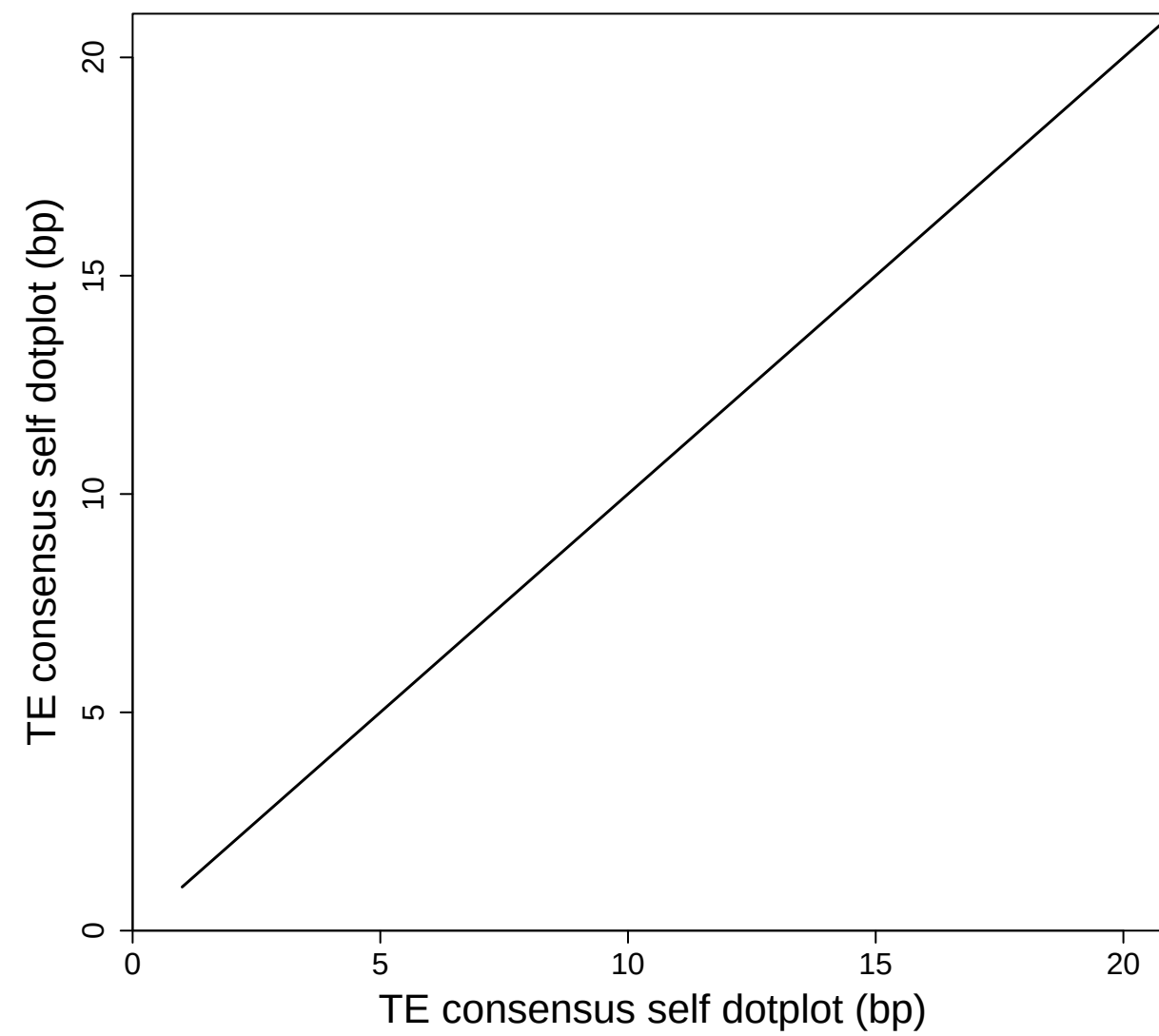
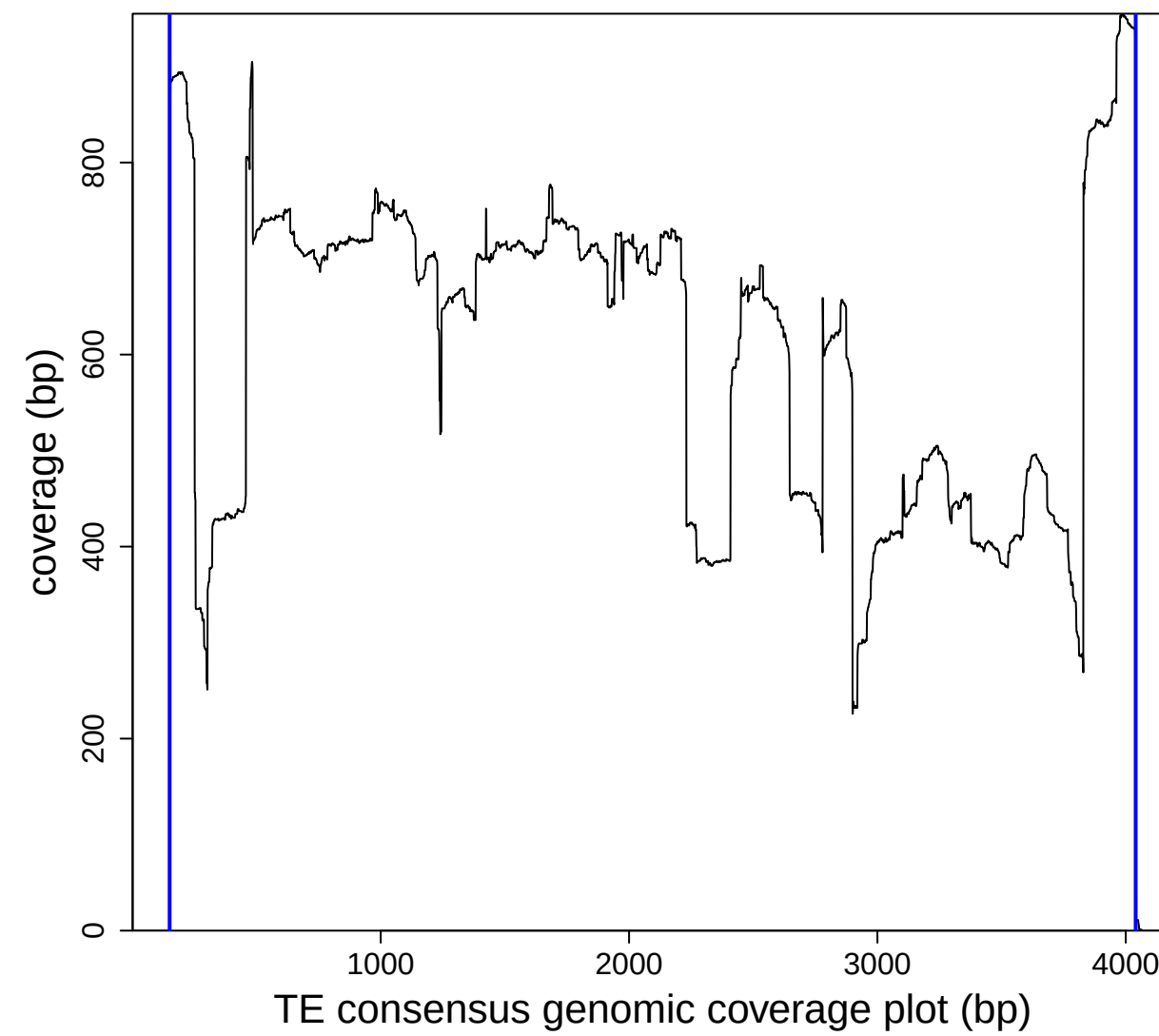
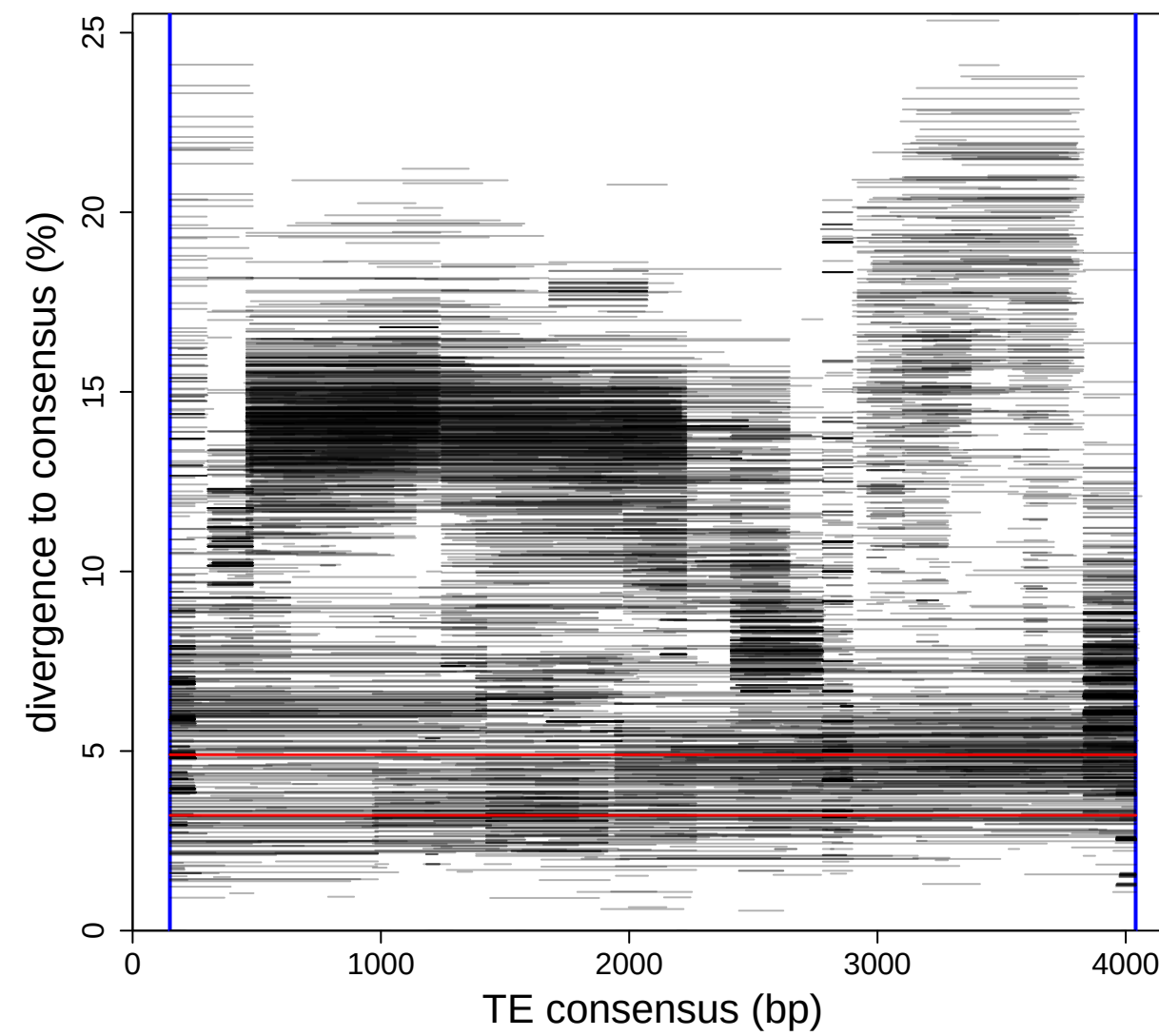
g_2.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.fa_a
size: 3884bp; fragments: 5285; full length: 3 (>=3495.6bp)

After TEtrimmer 3884 bp

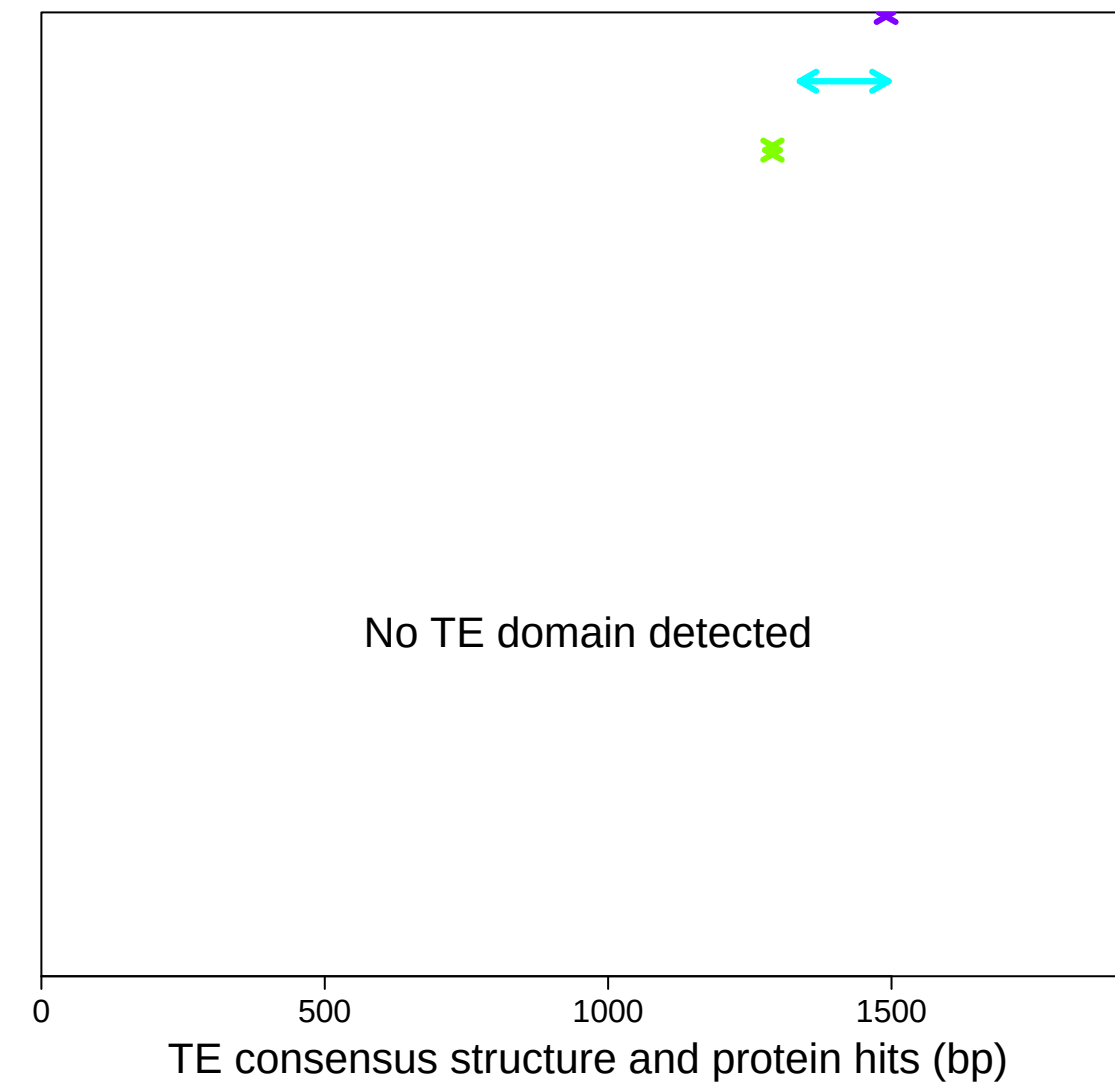
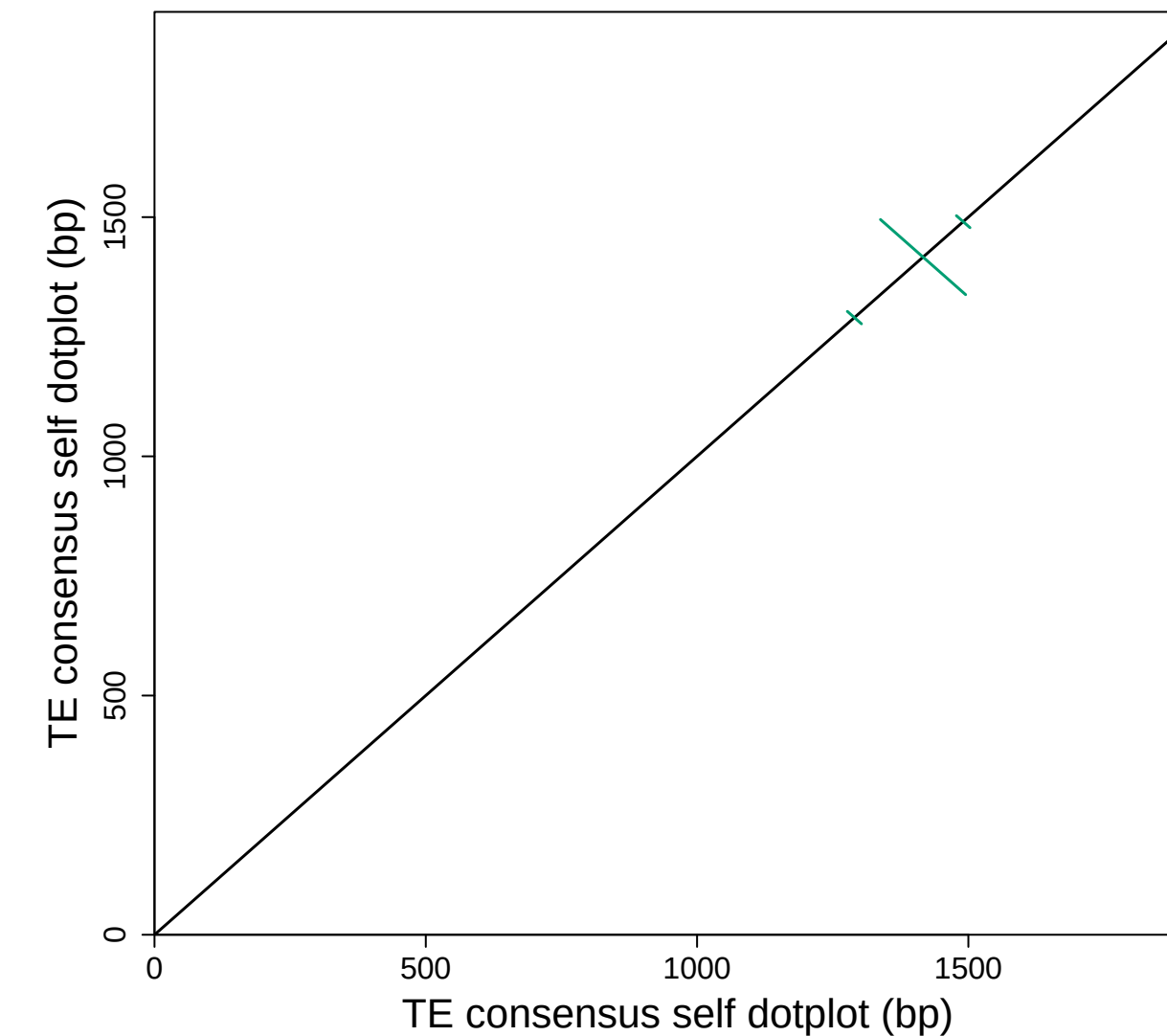
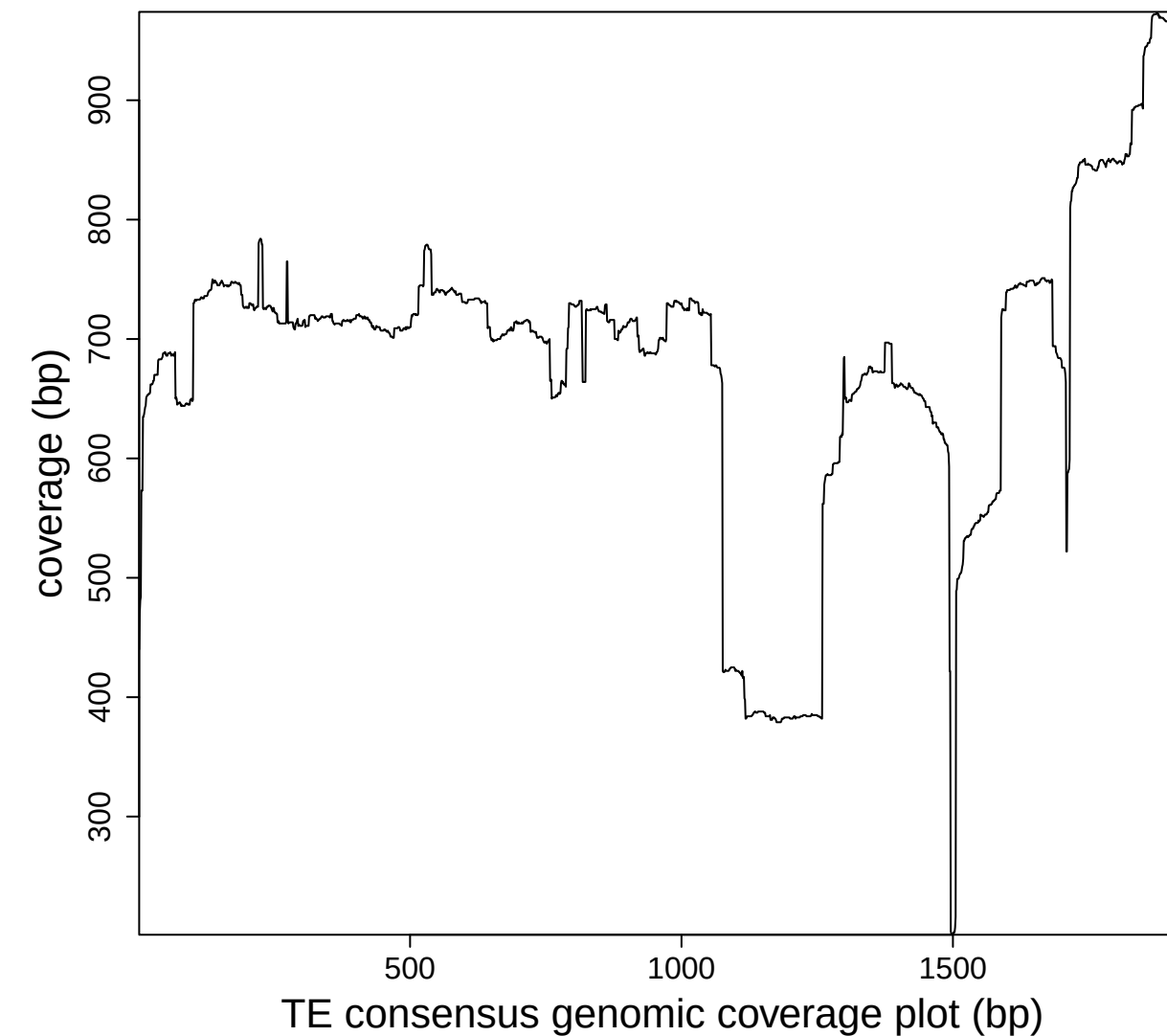
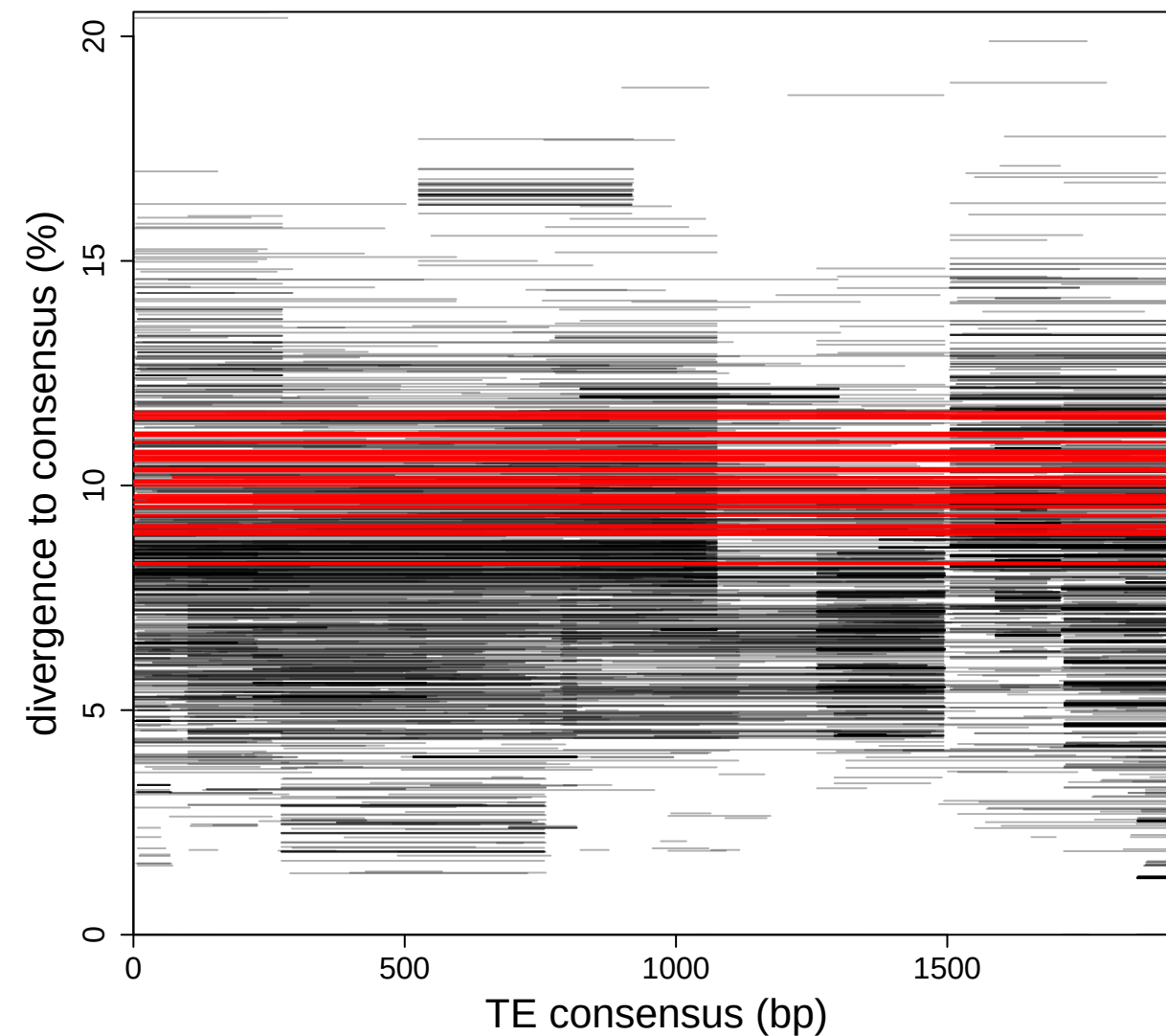


f.bed_g_2.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcl
size: 4190bp; fragments: 5224; full length: 2 (>=3771bp)

After TEtrimmer Extended plot Blue lines are boundaries

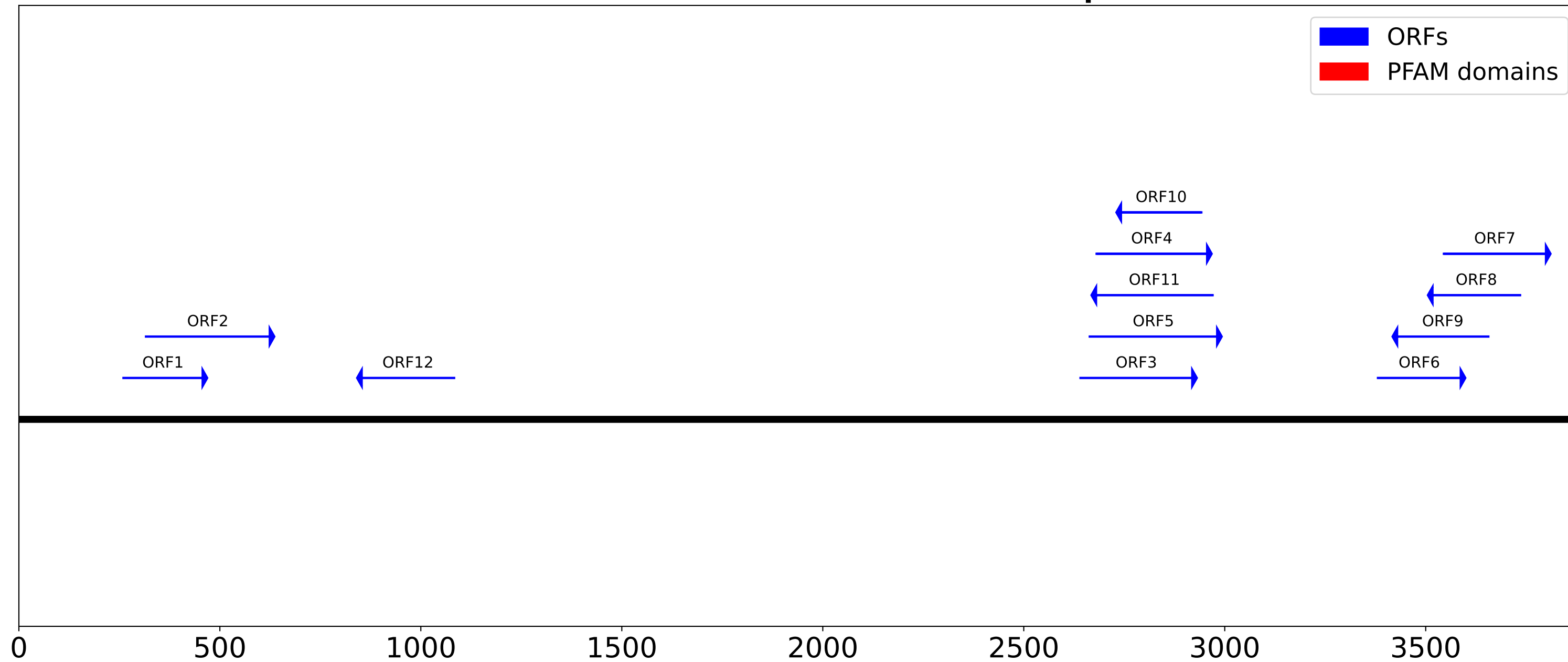


TE: rnd_4_family_557
size: 1929bp; fragments: 3075; full length: 42 (≥ 1736.1 bp)

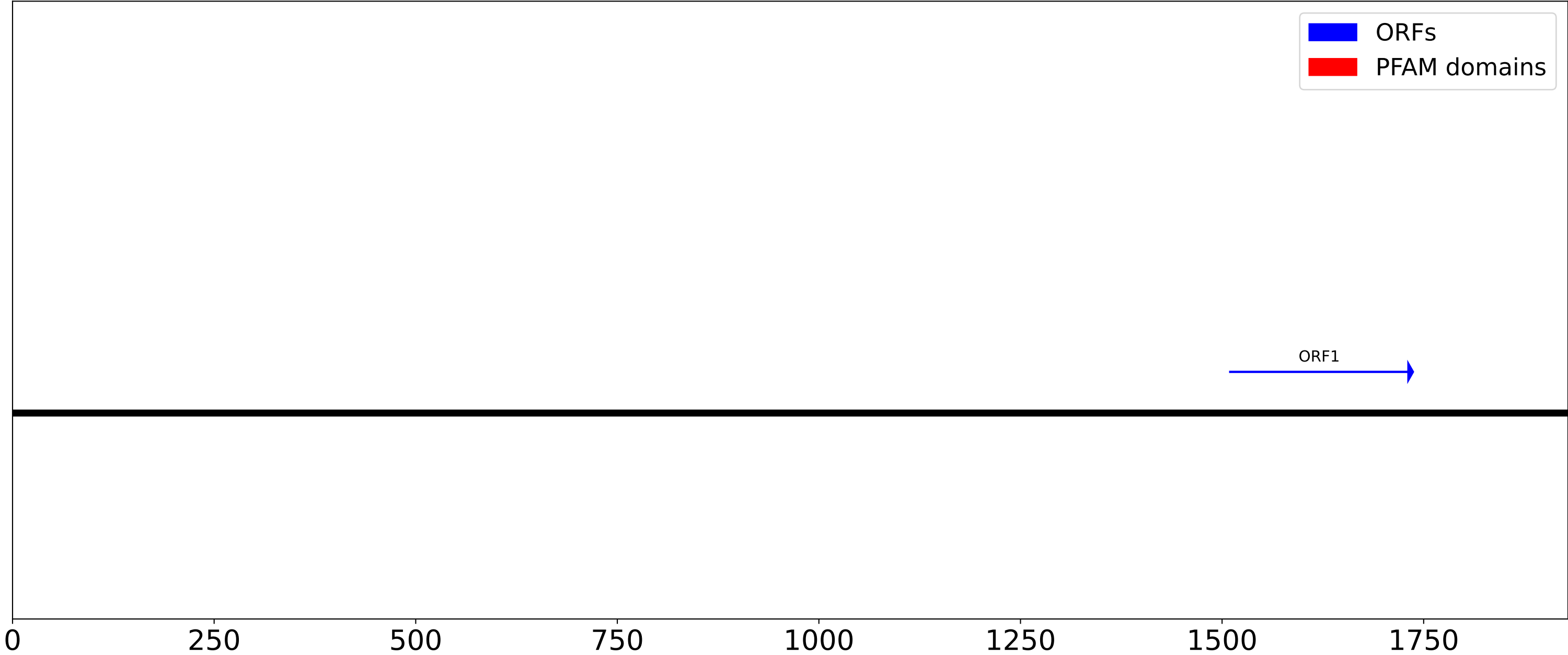


Before TEtrimmer 1929 bp

After TEtrimmer ORF and PFAM domain plot



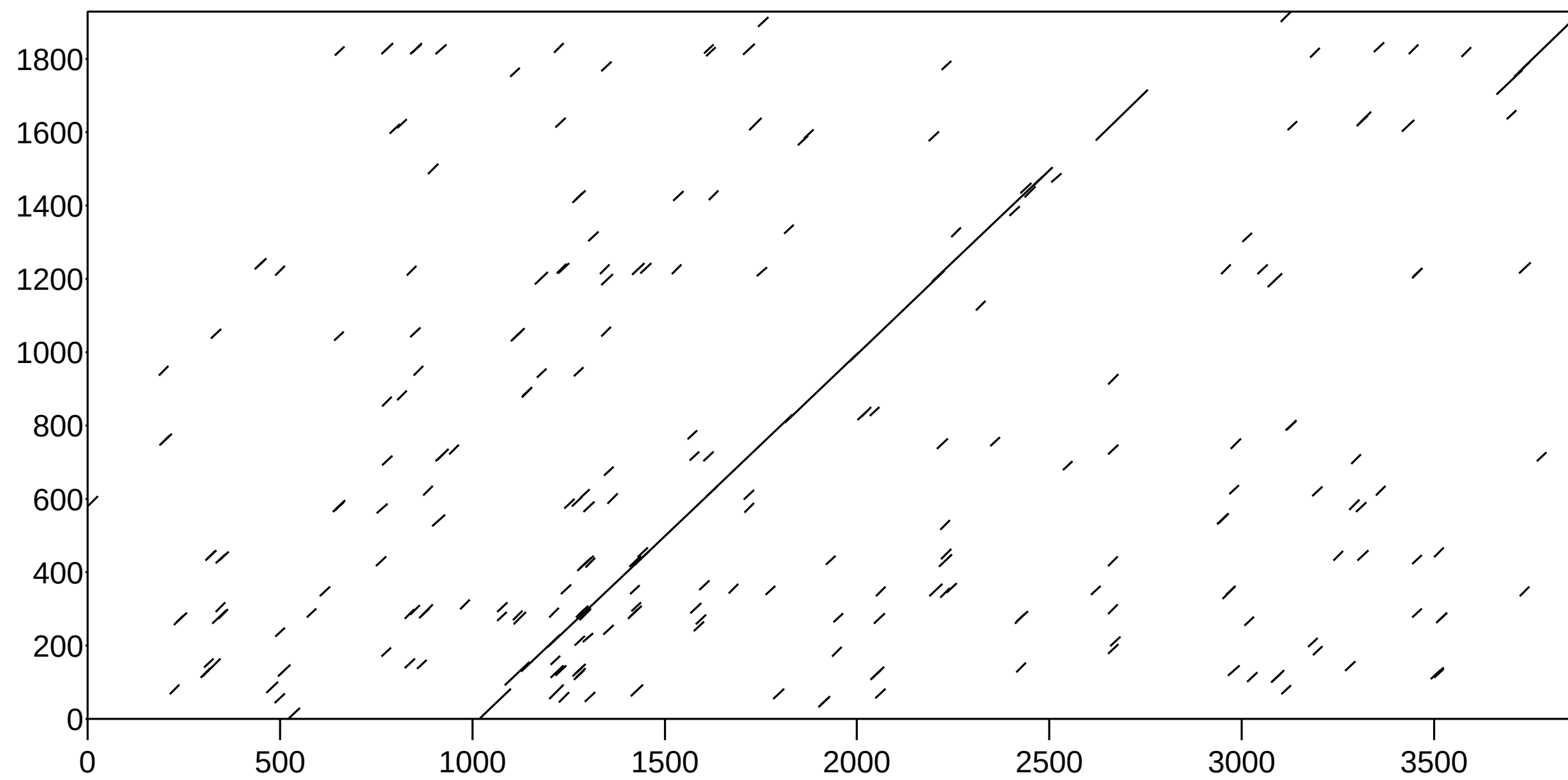
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)