

Start crop Point



End crop Point

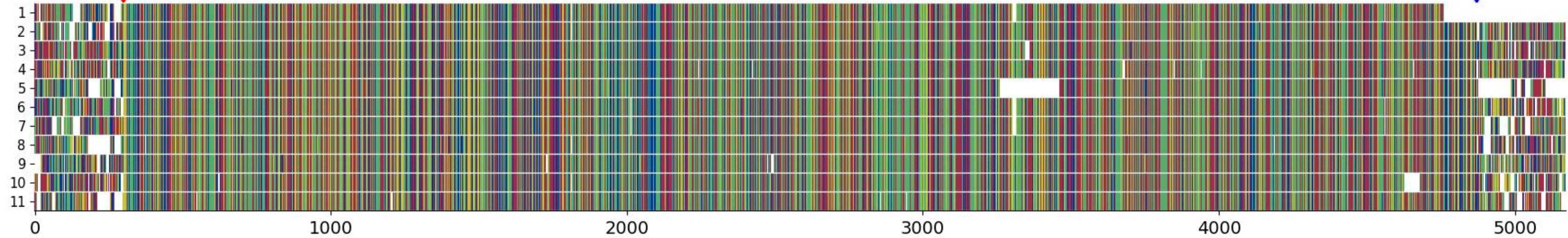


MSA length = 4572



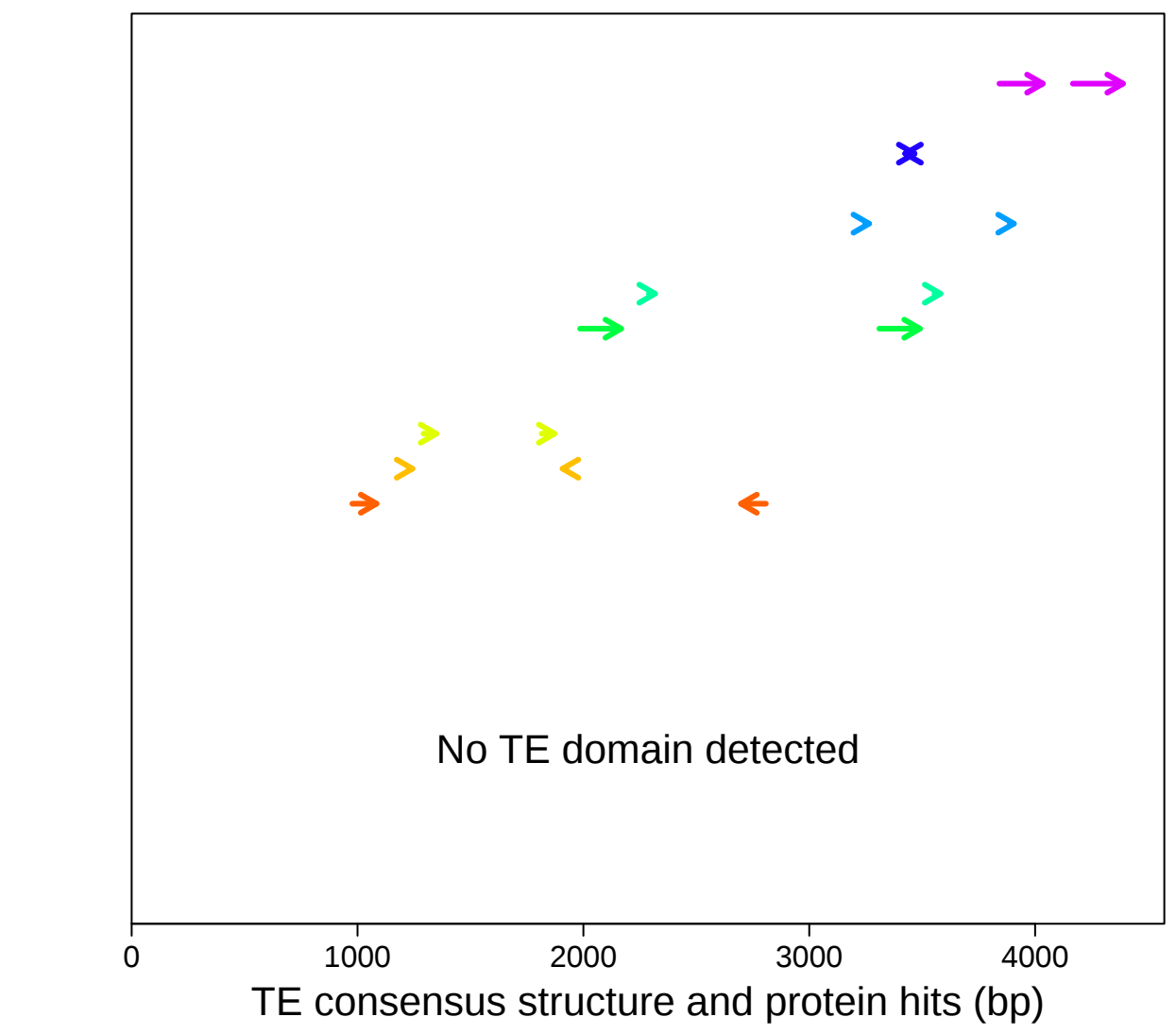
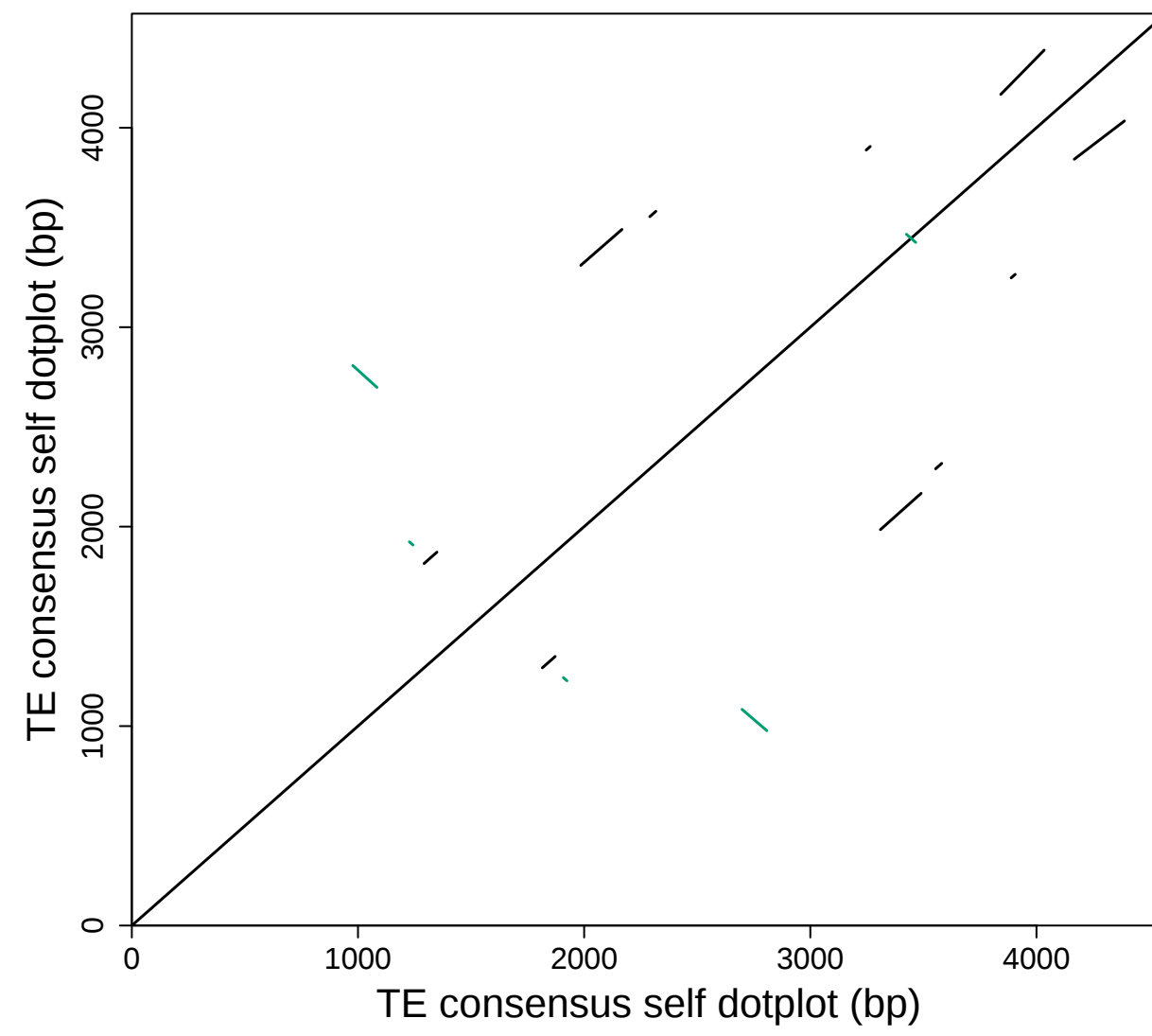
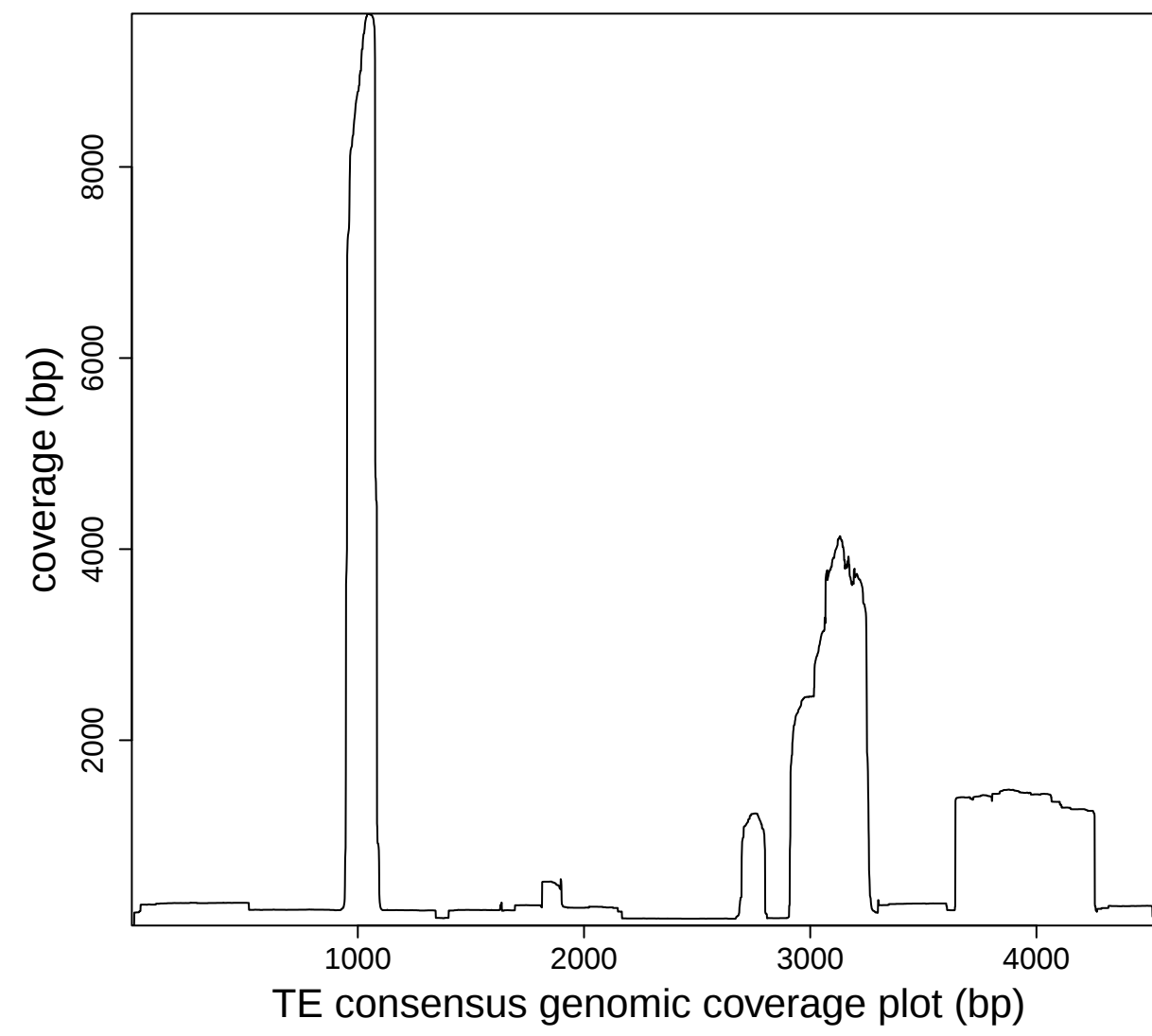
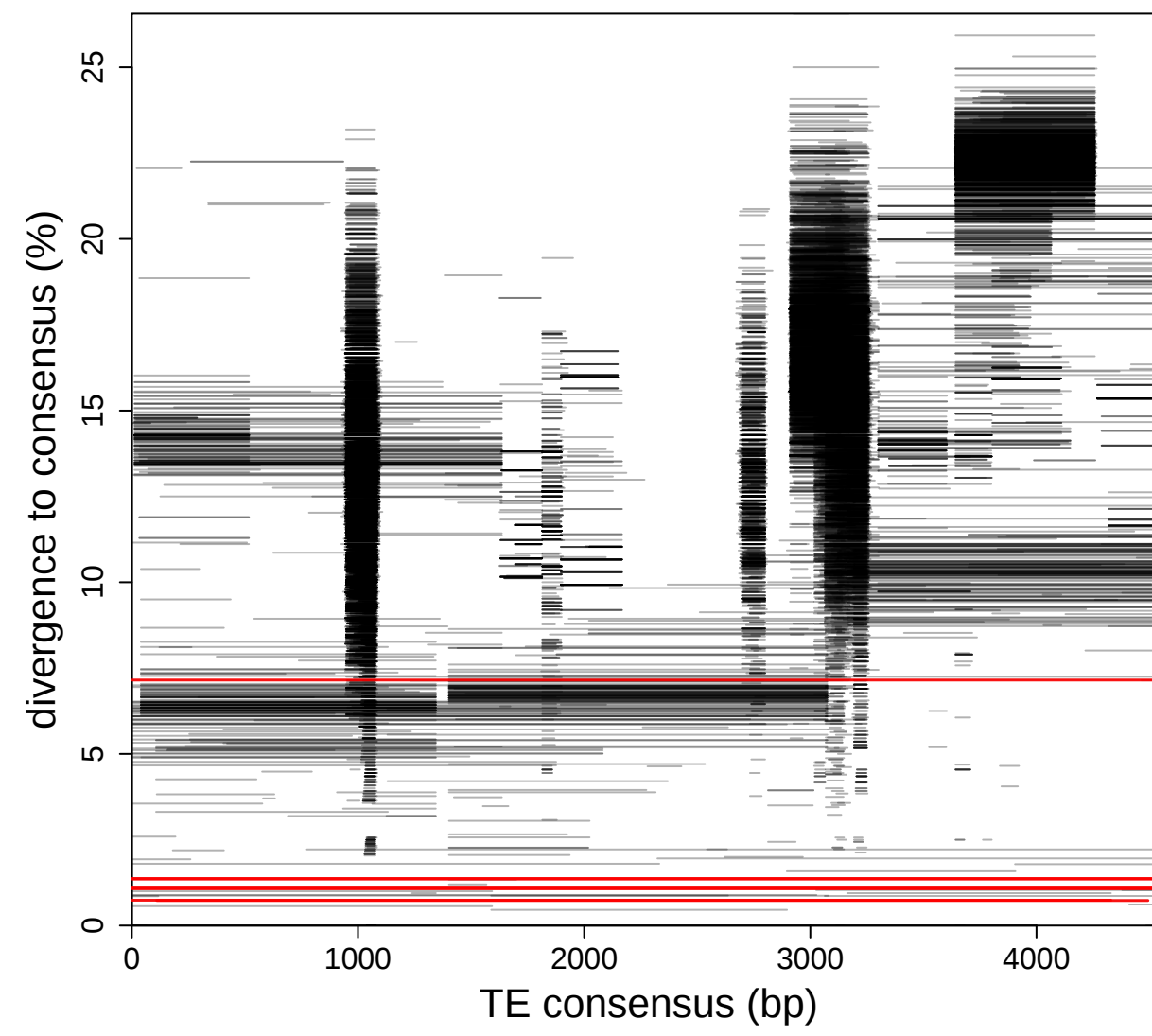
Start crop Point

End crop Point



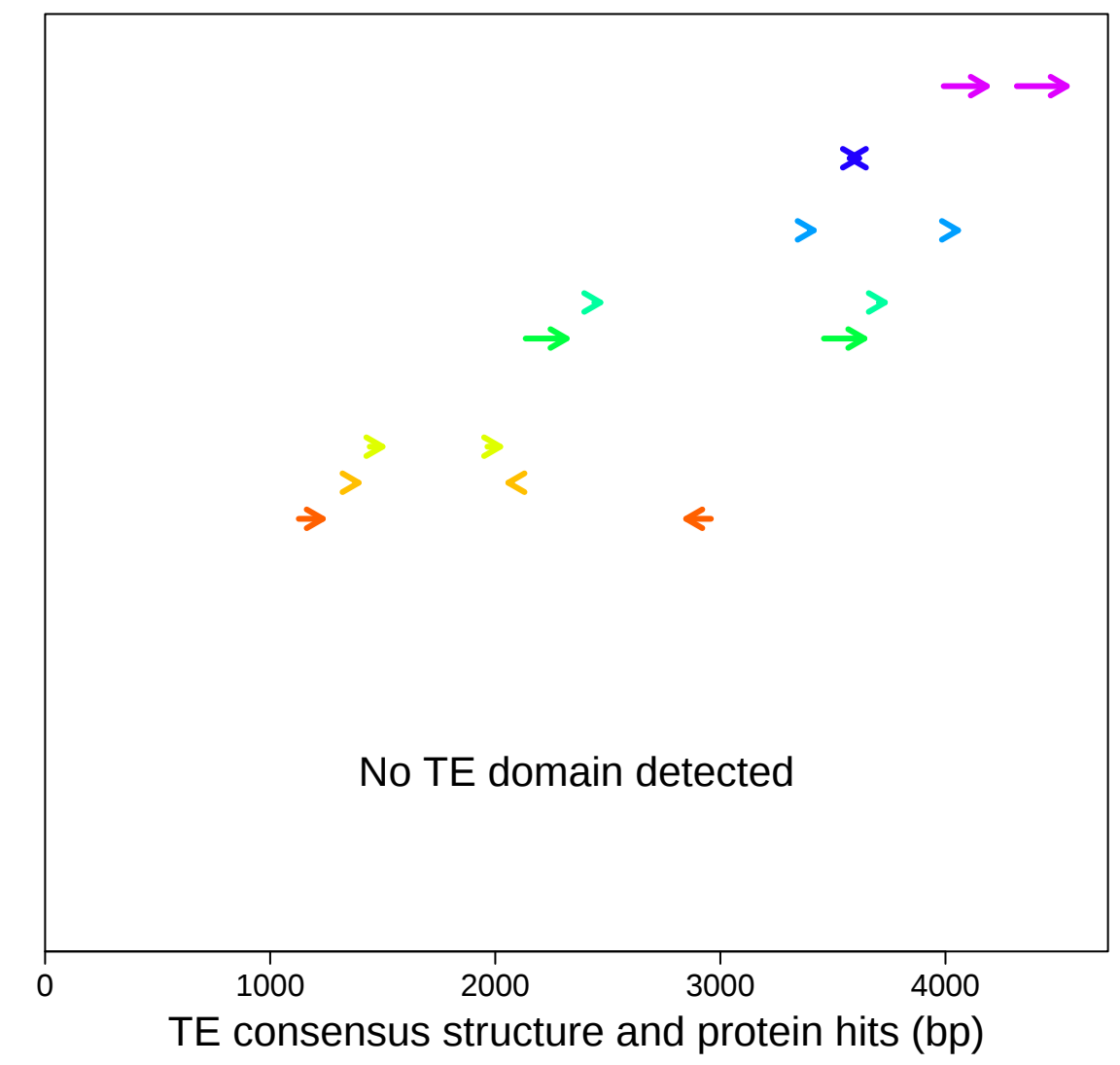
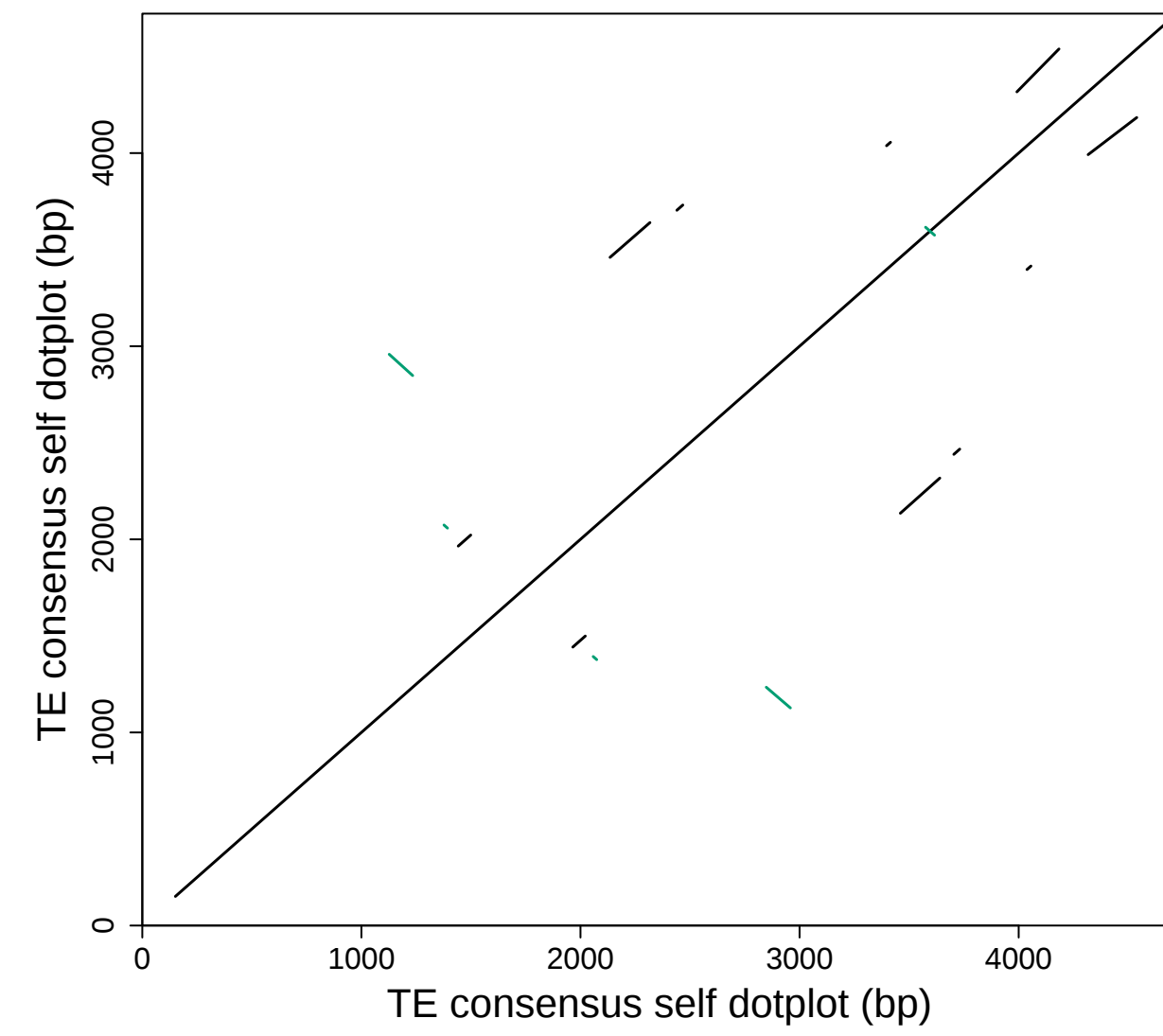
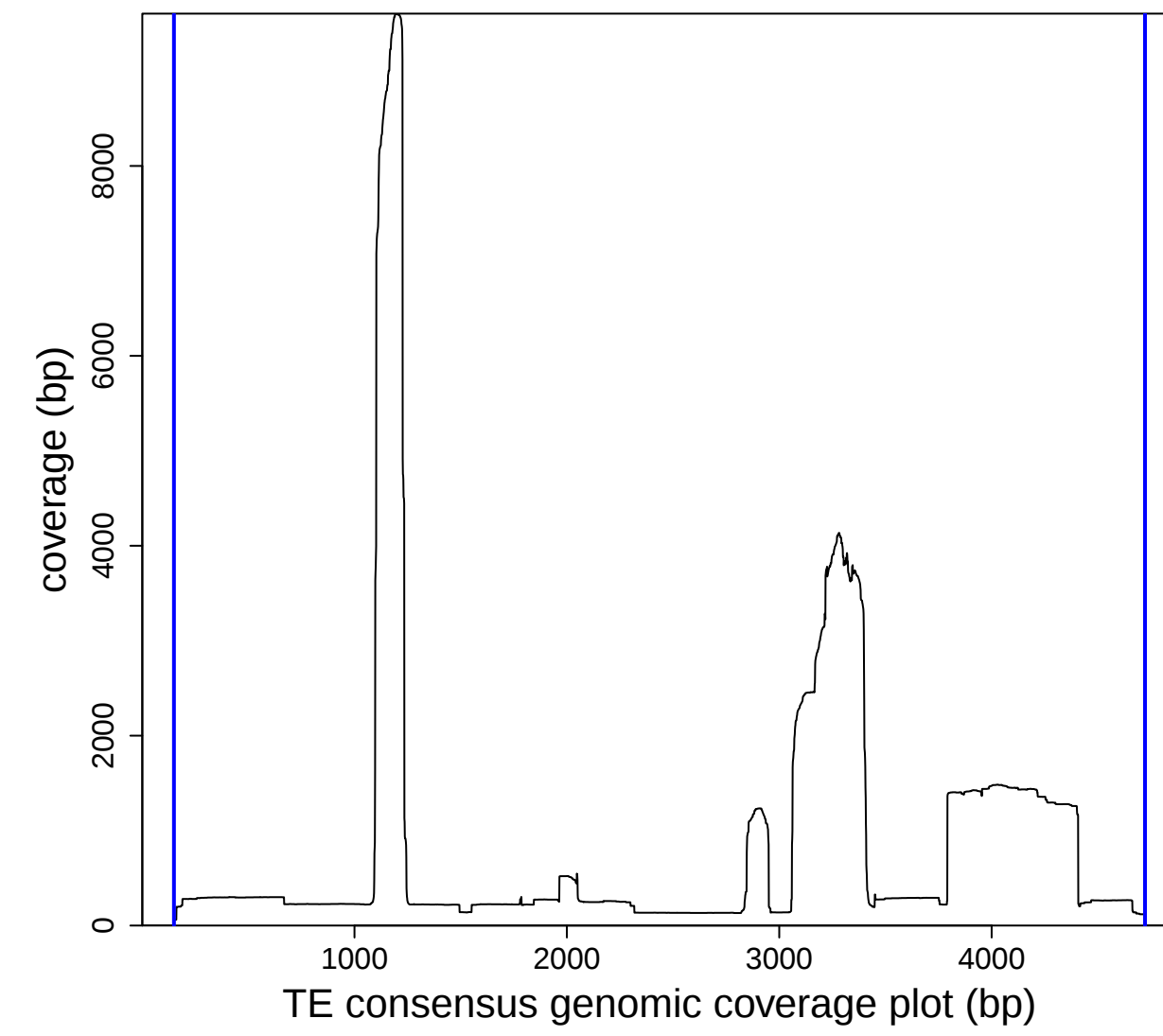
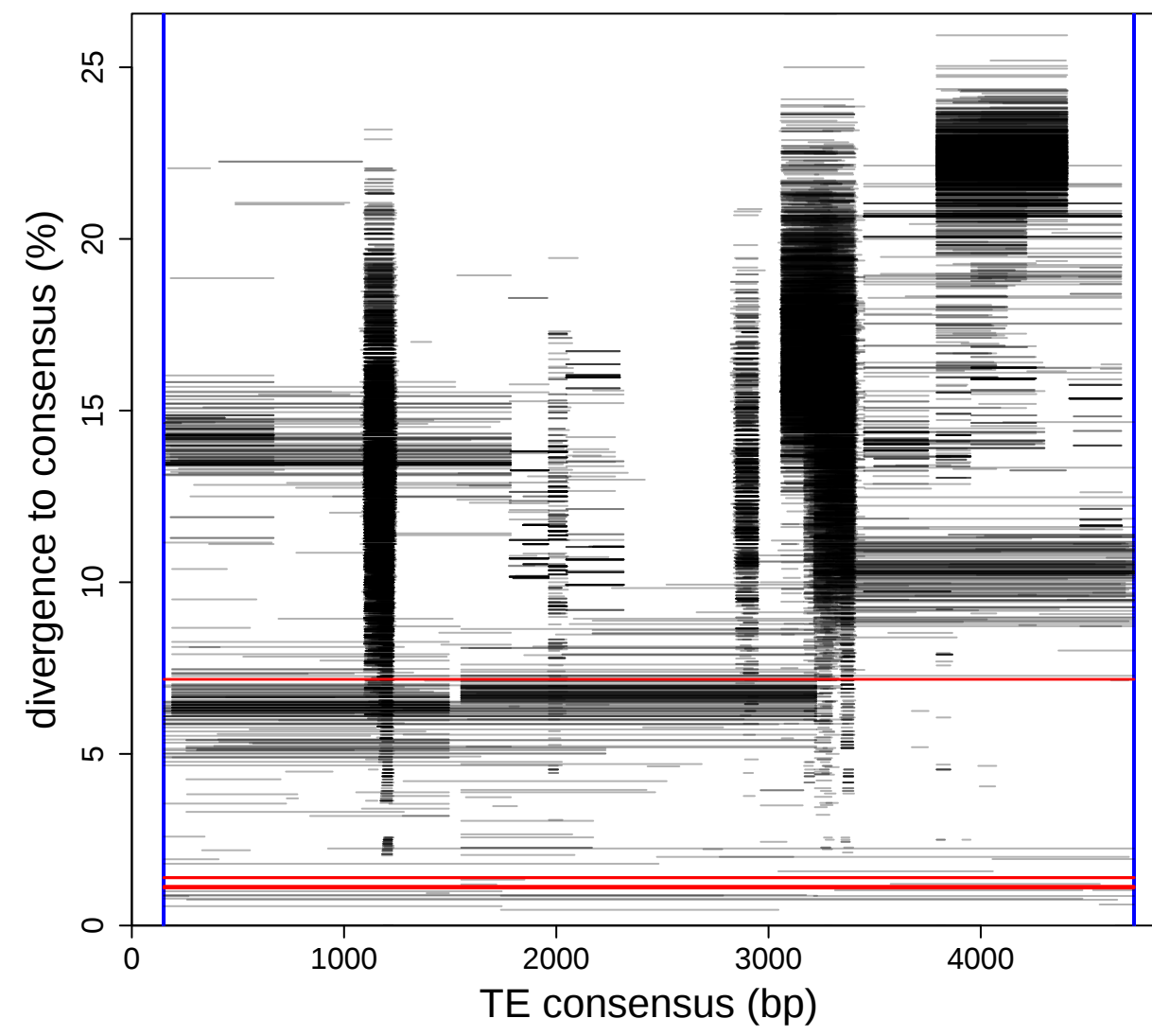
g_4.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.fa_3
size: 4572bp; fragments: 18344; full length: 9 (>=4114.8bp)

After TETrimmer 4572 bp



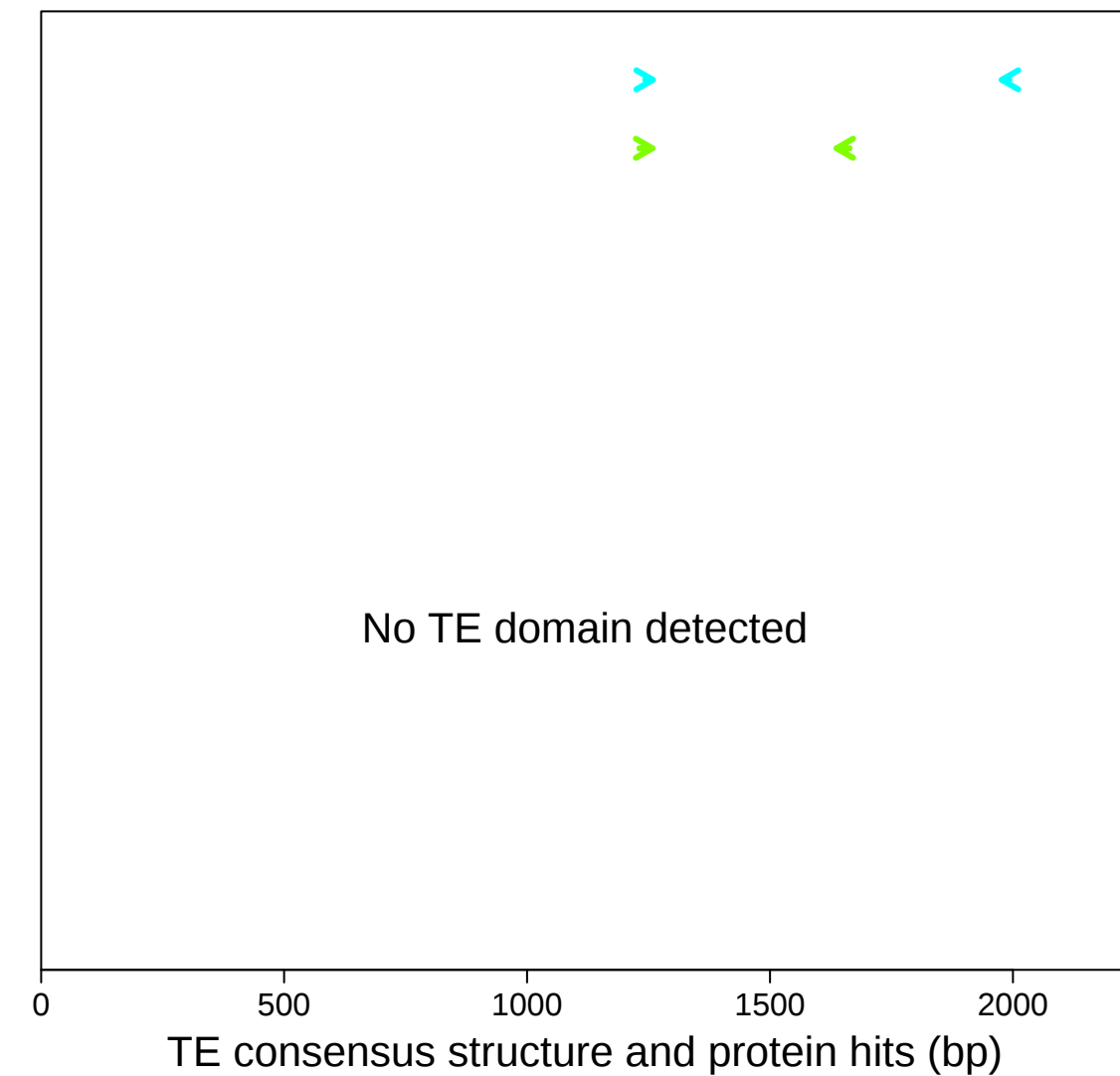
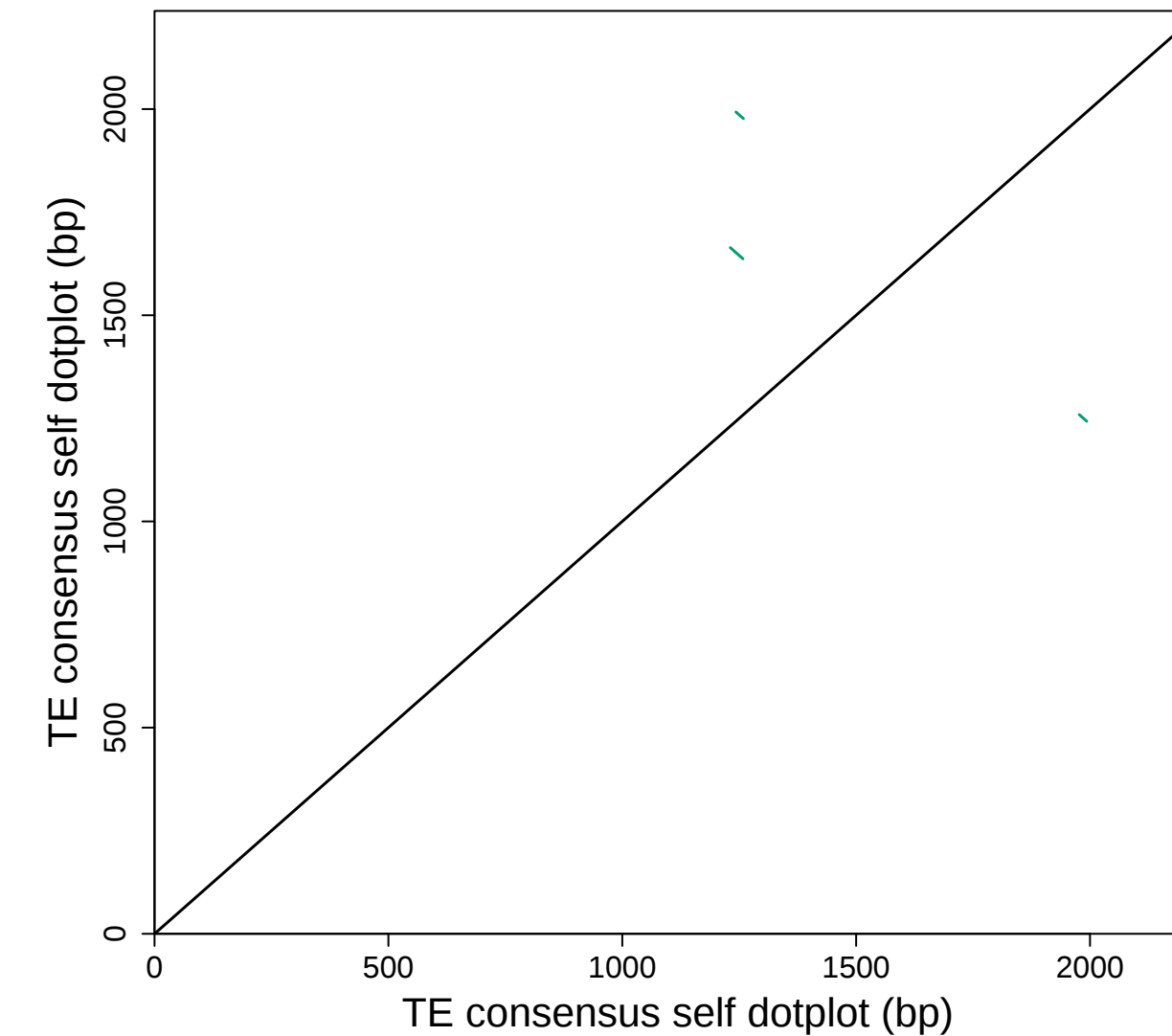
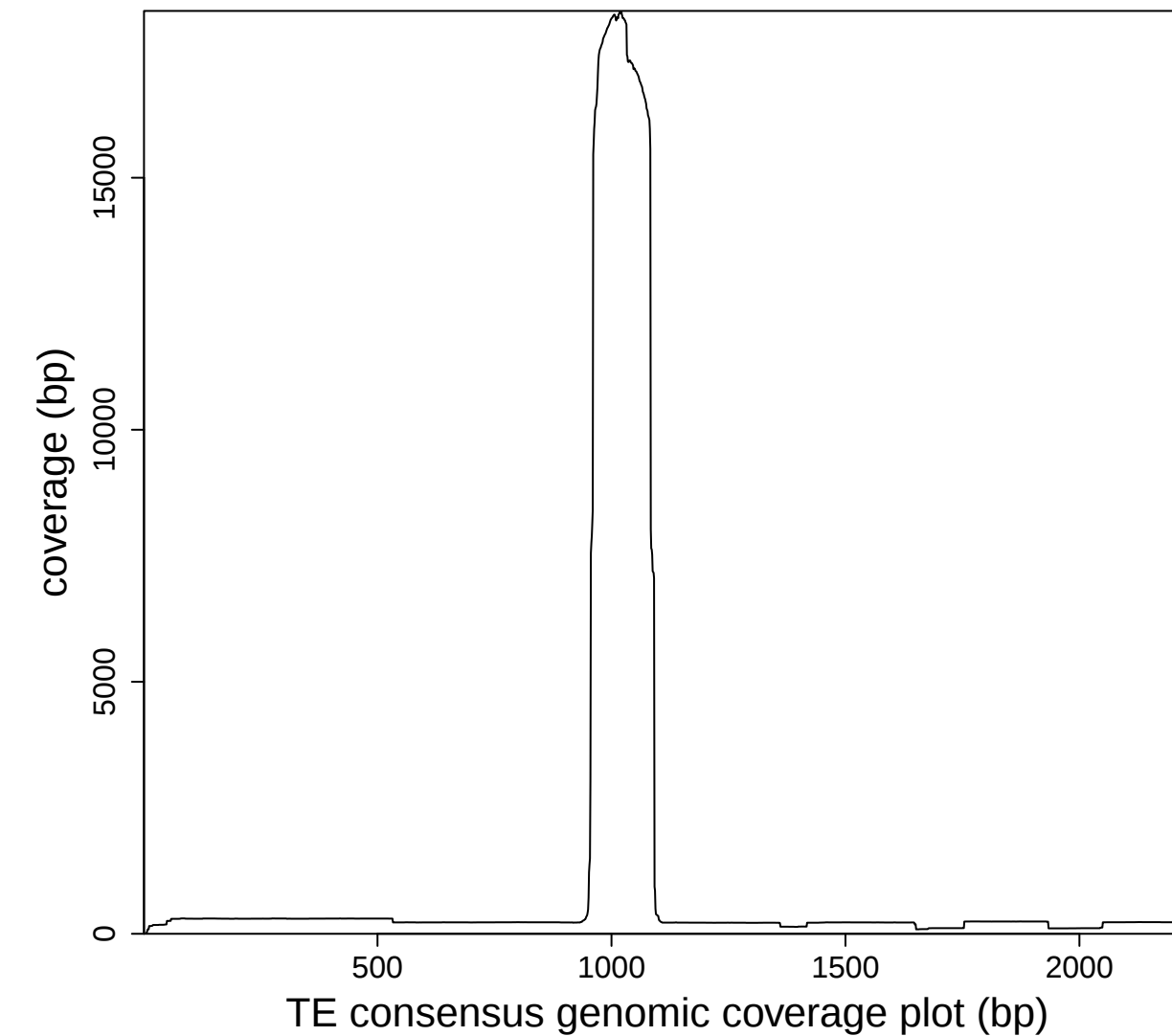
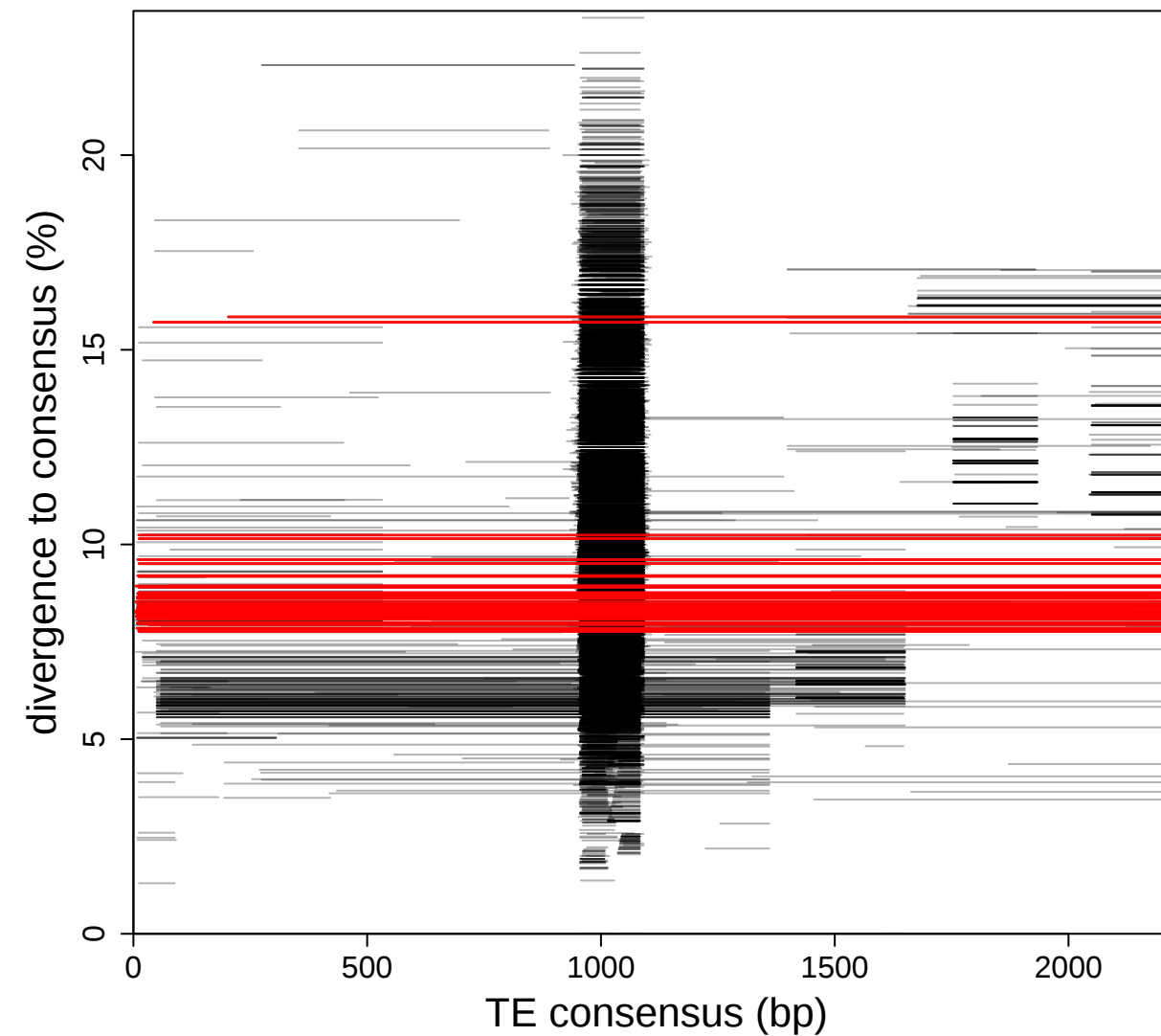
f.bed_g_4.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bc
size: 4872bp; fragments: 18343; full length: 7 (>=4384.8bp)

After TEtrimmer Extended plot Blue lines are boundaries

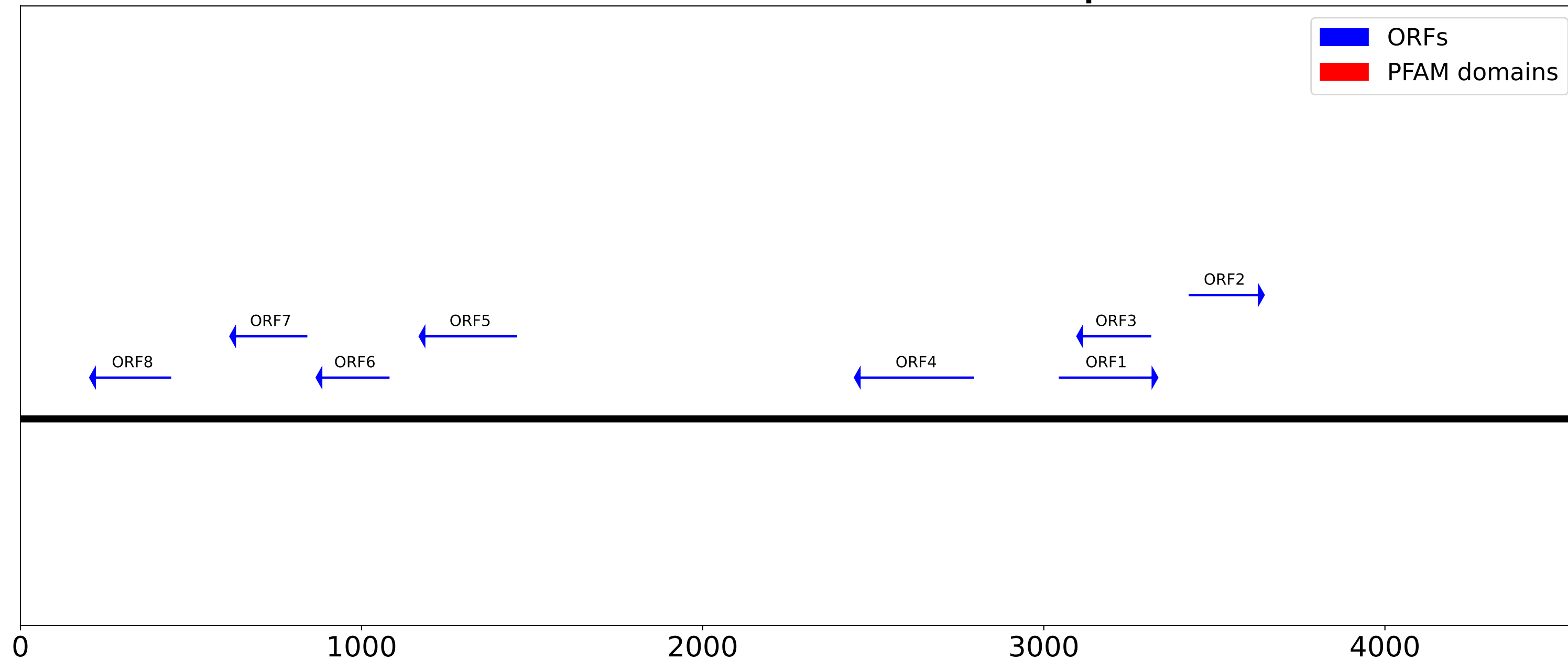


TE: rnd_5_family_3052
size: 2238bp; fragments: 19614; full length: 63 (≥ 2014.2 bp)

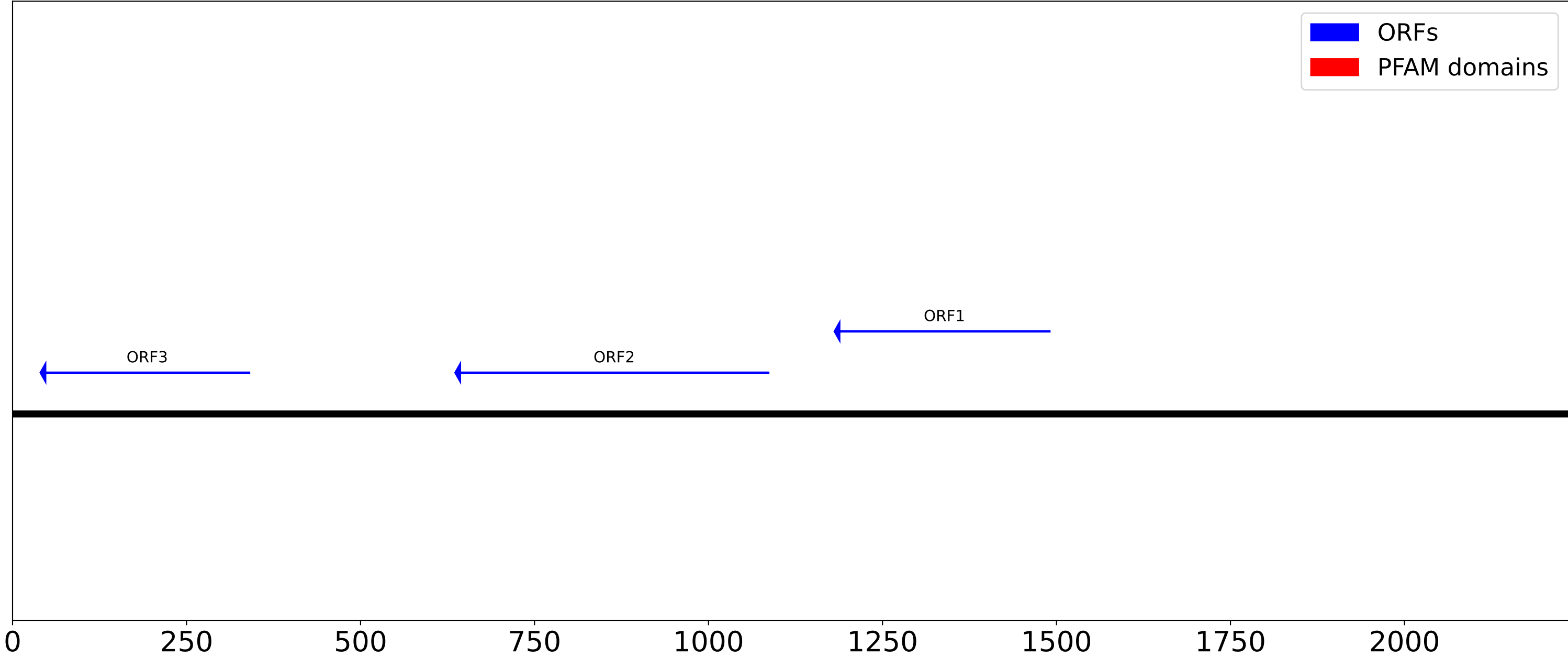
Before TEtrimmer 2238 bp



After TEtrimmer ORF and PFAM domain plot



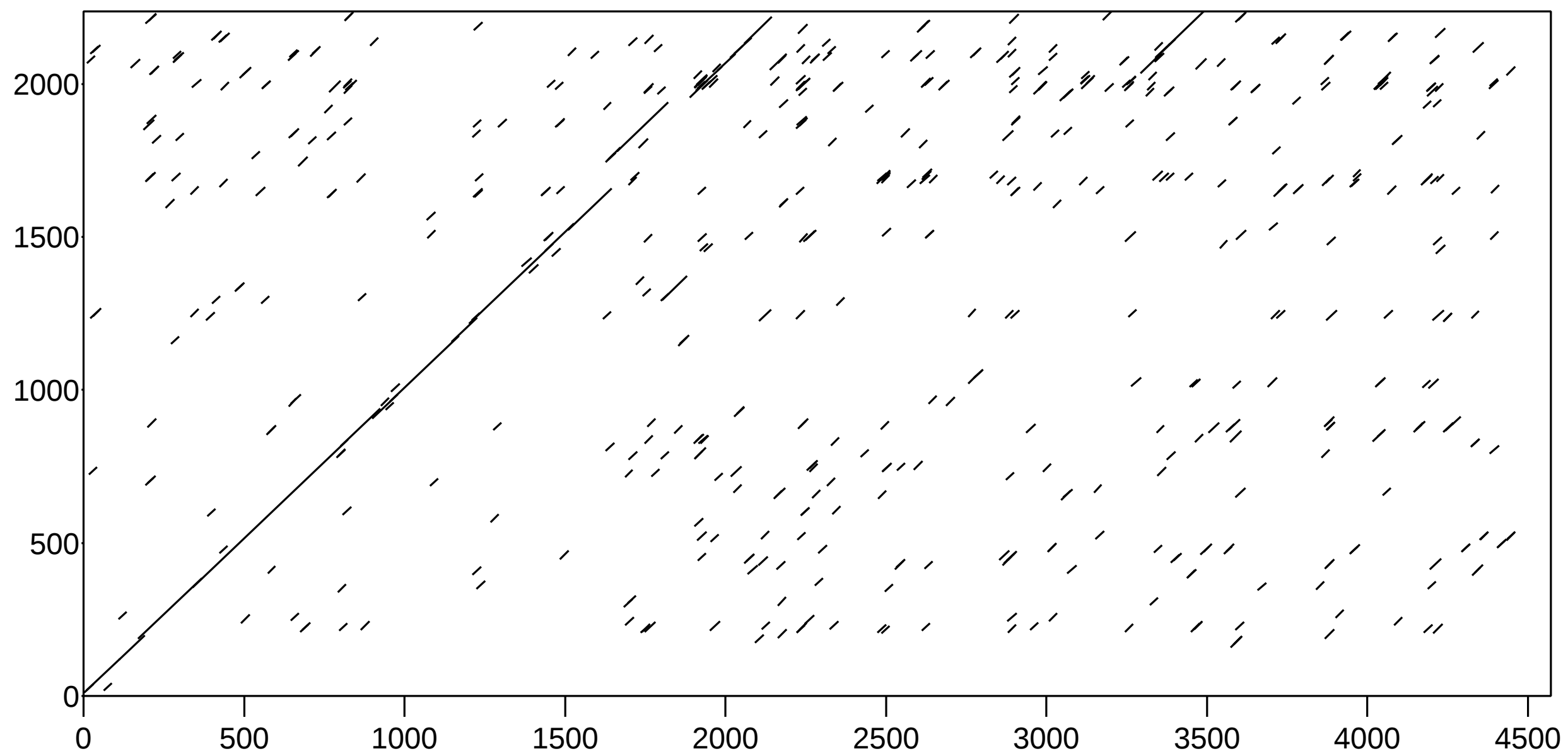
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)