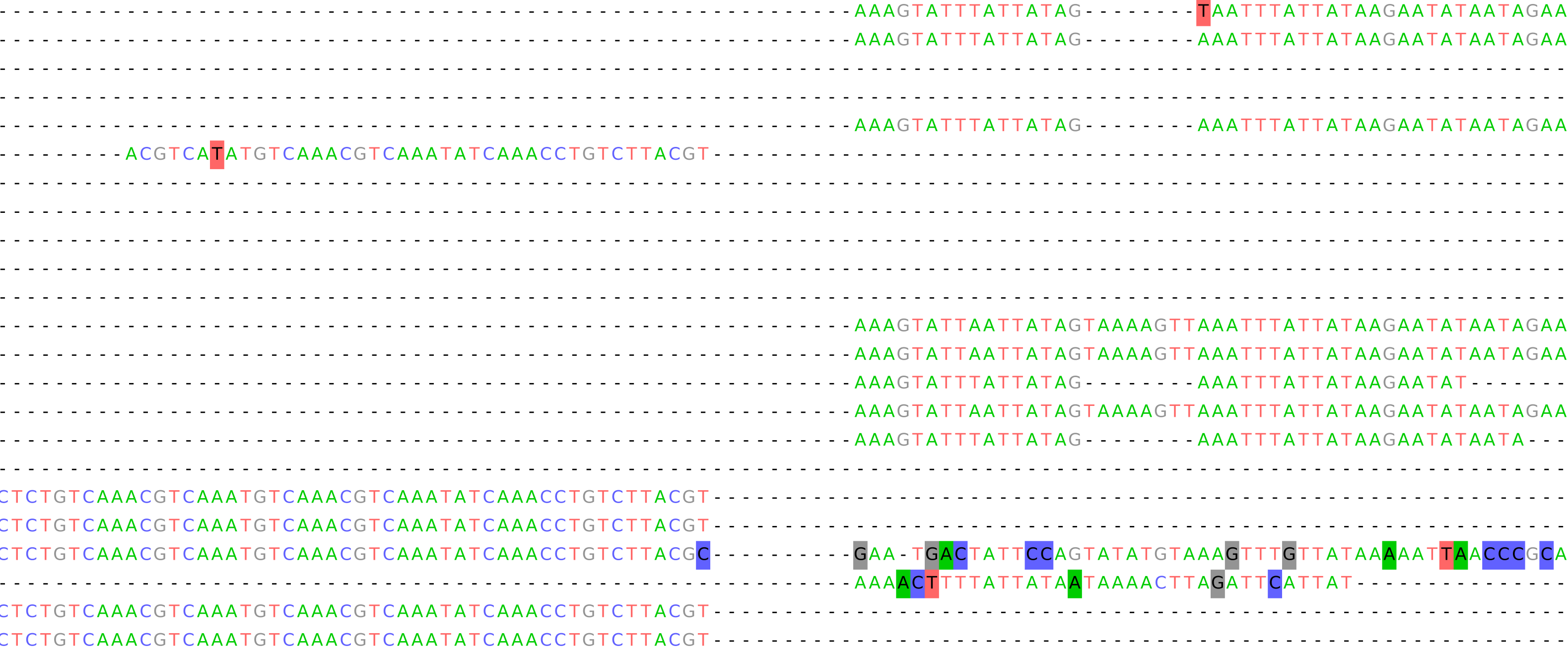


Start crop Point

End crop Point

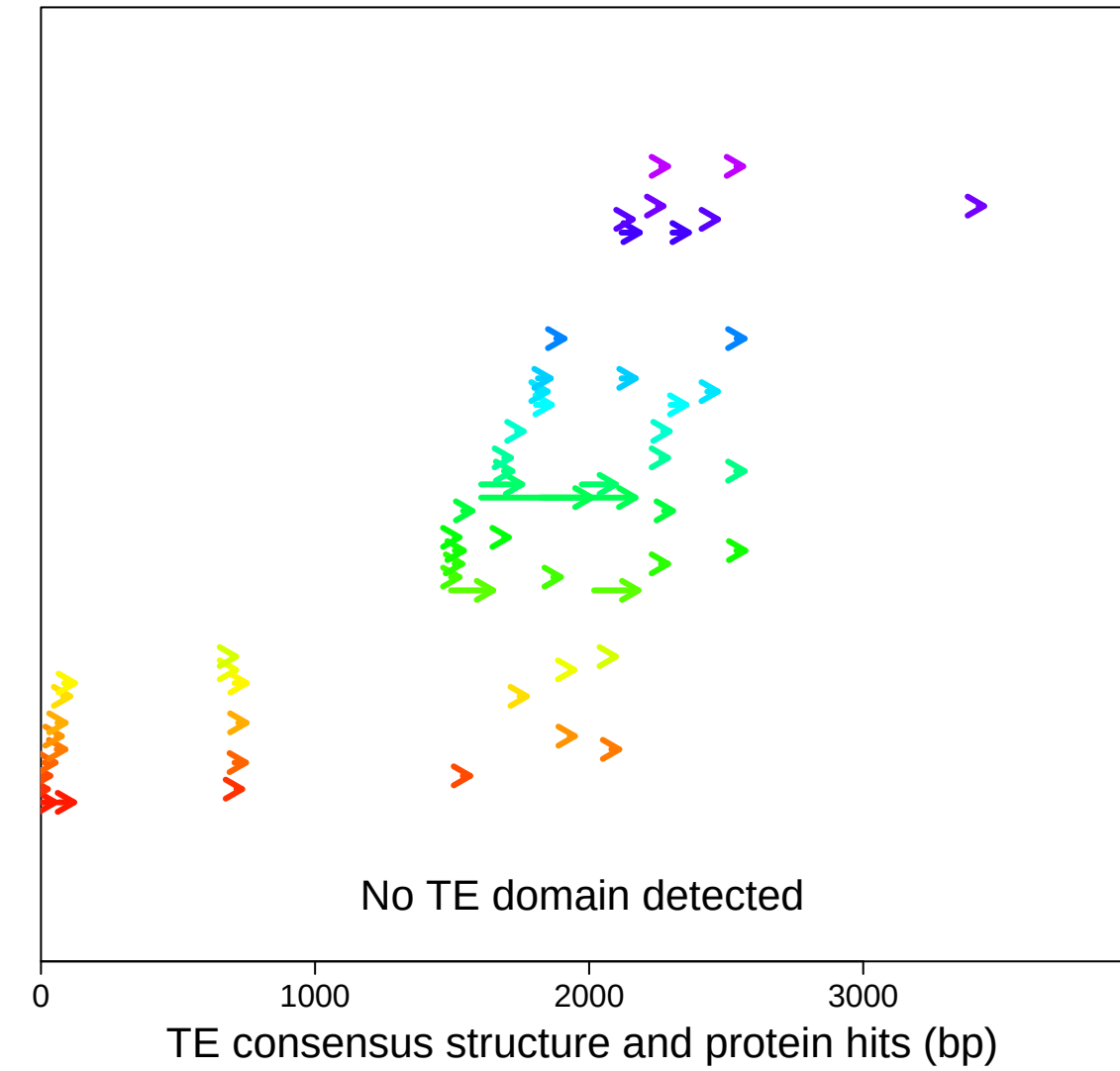
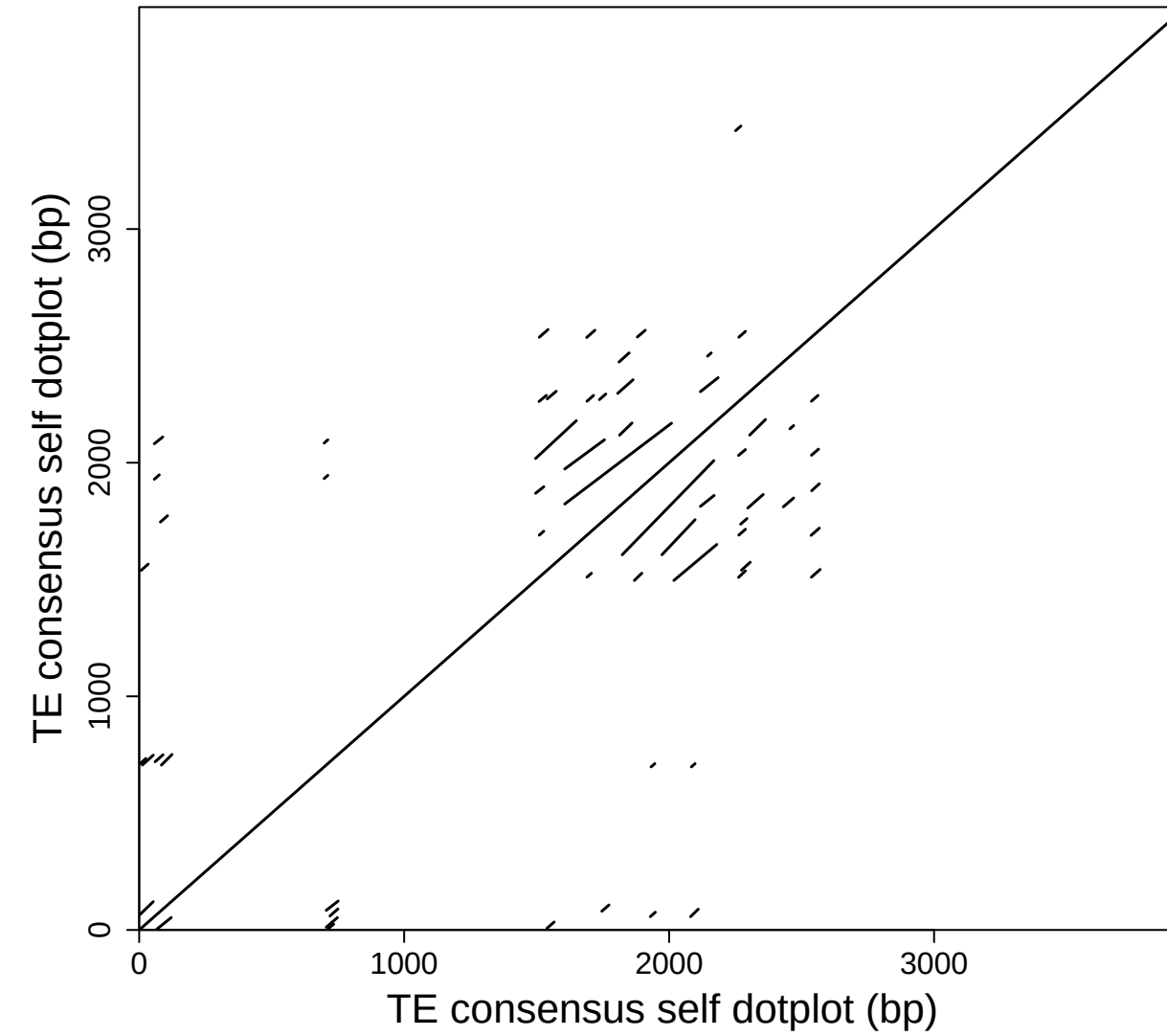
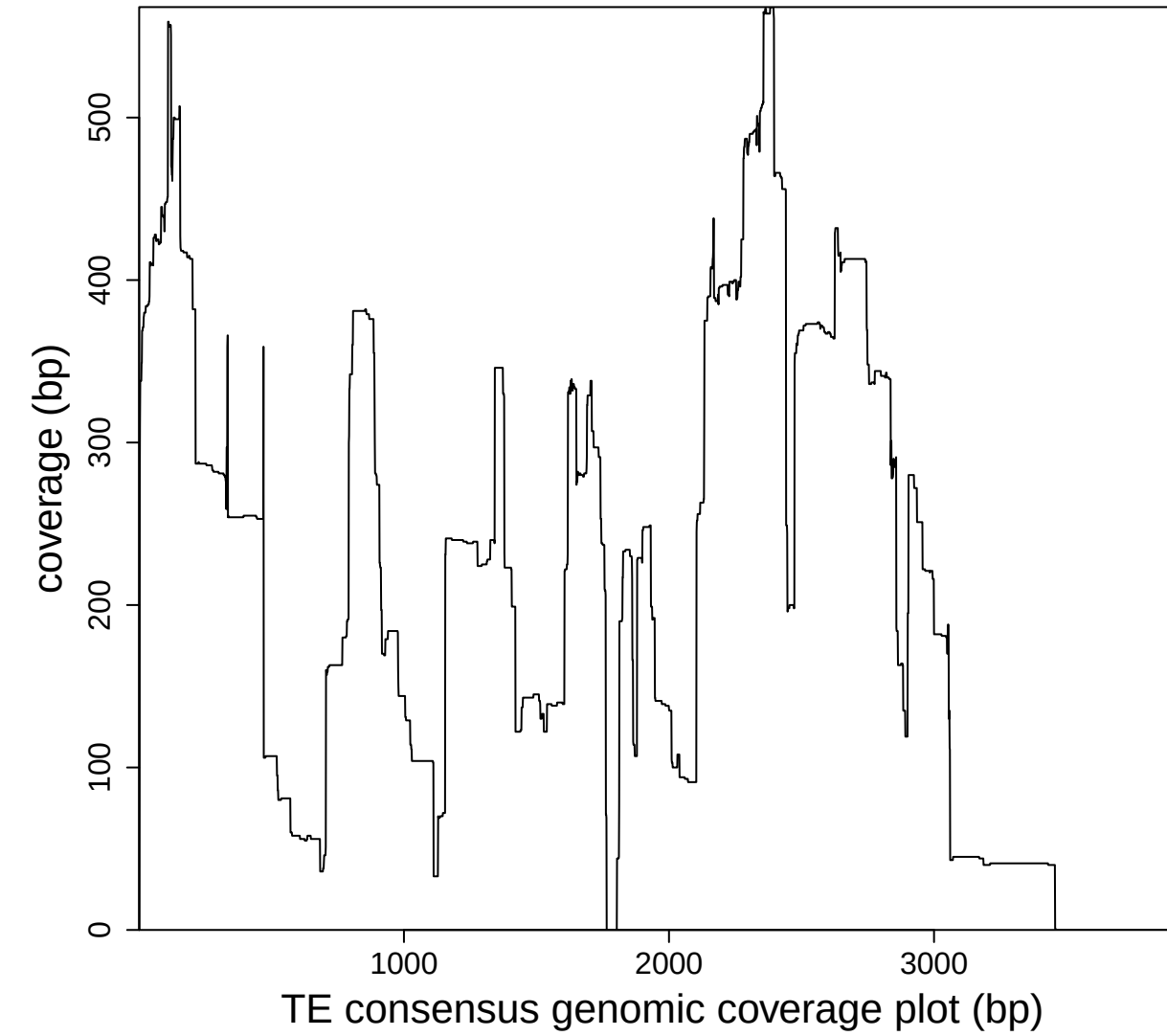
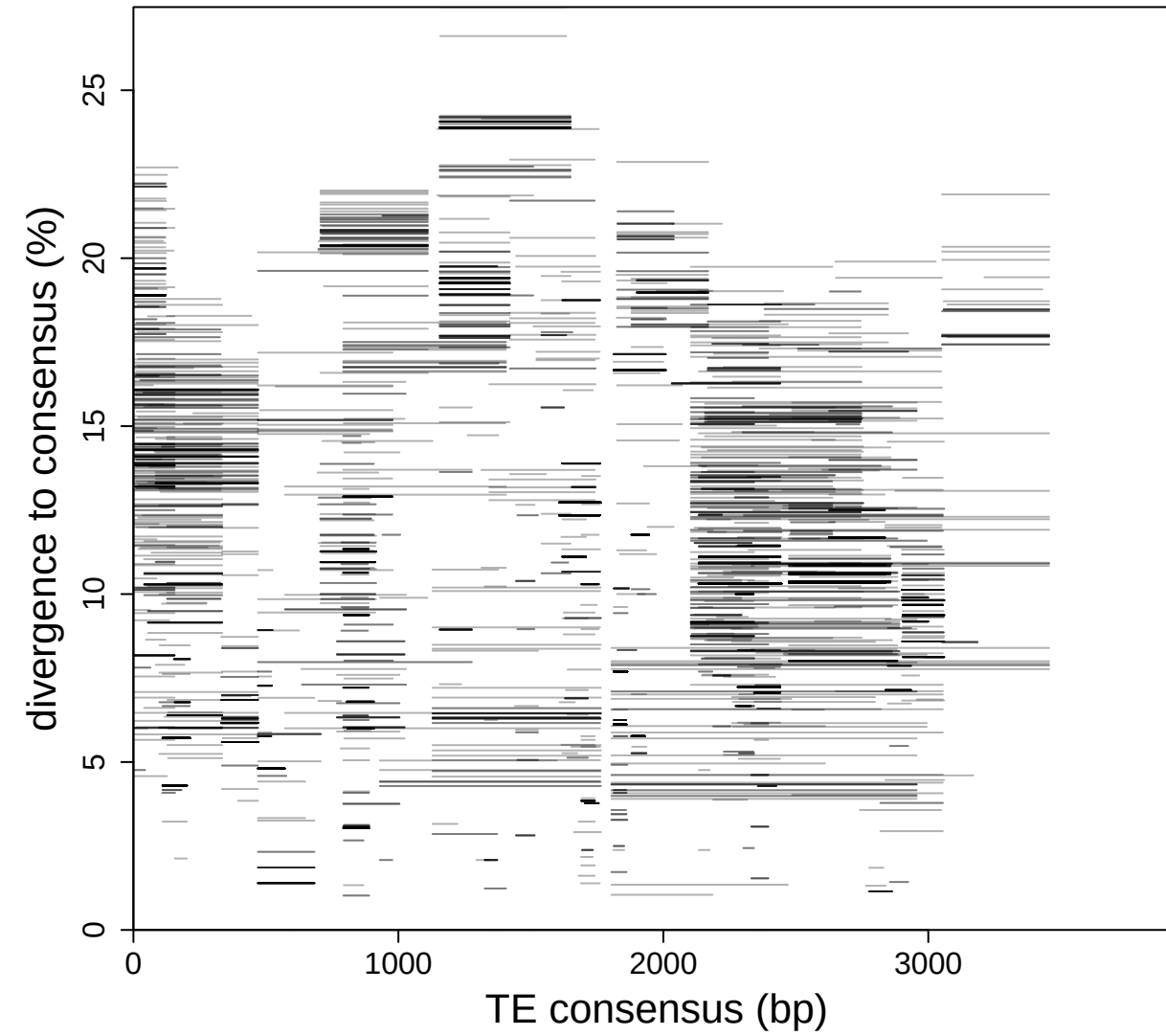
MSA length = 3950





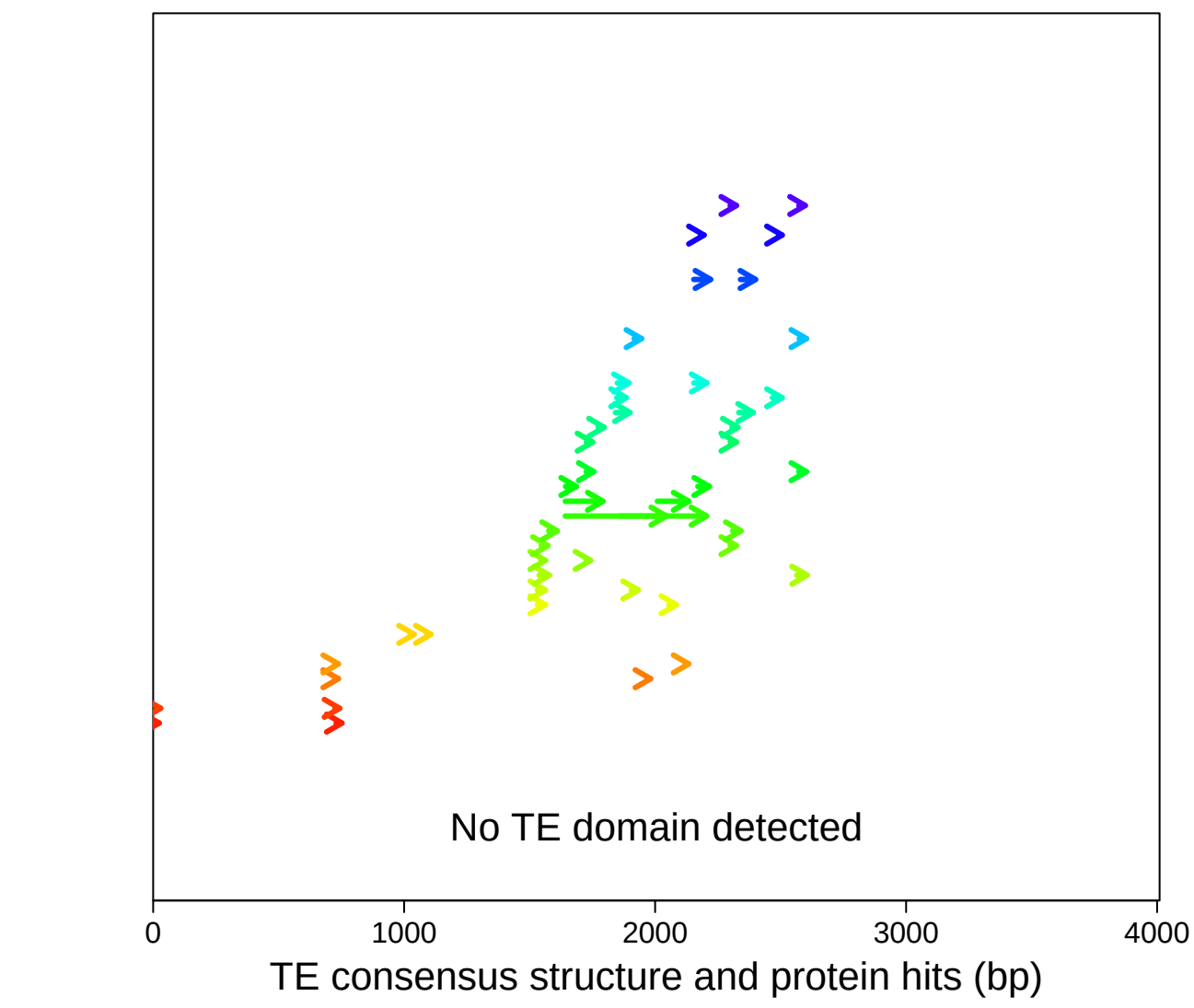
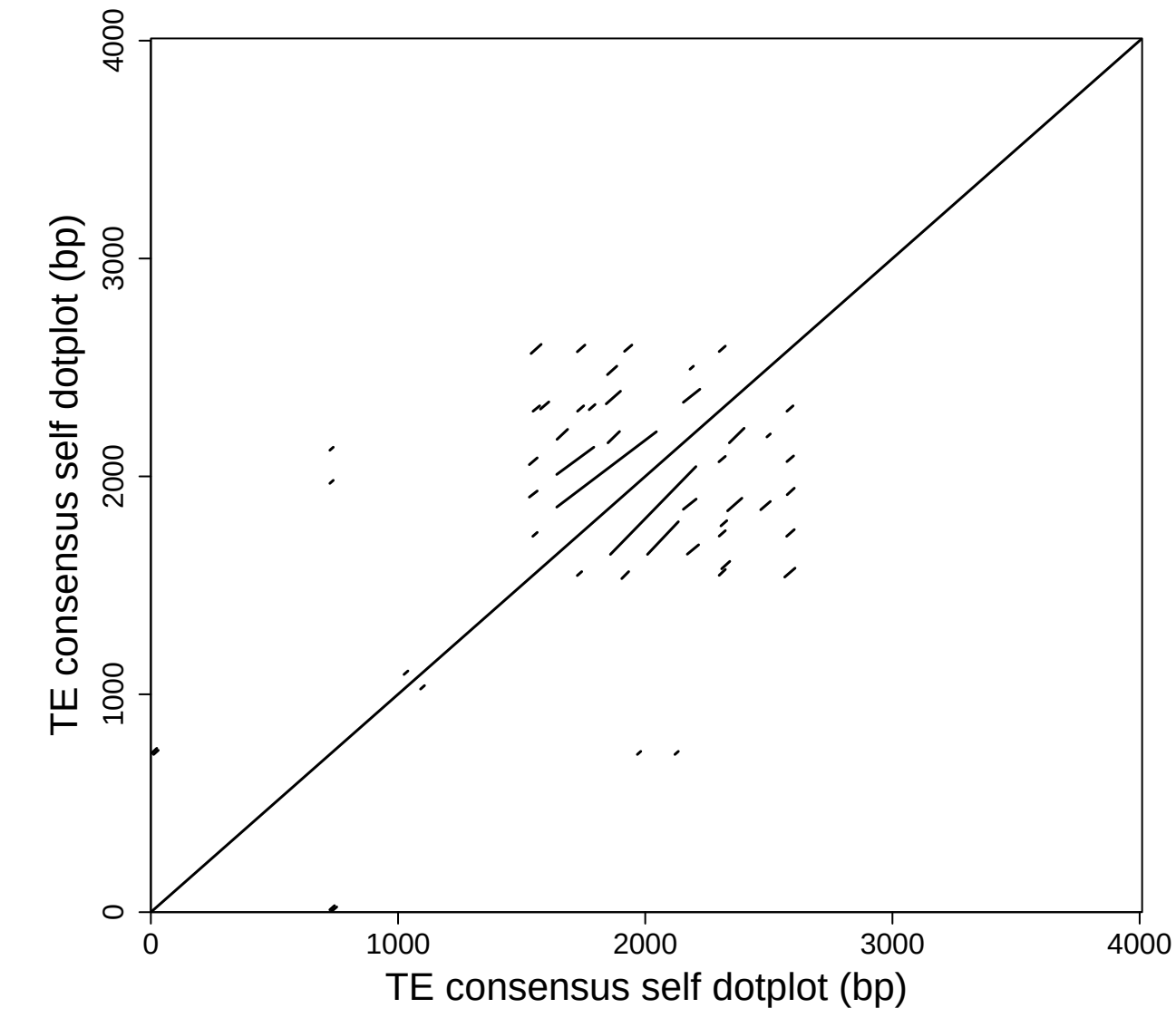
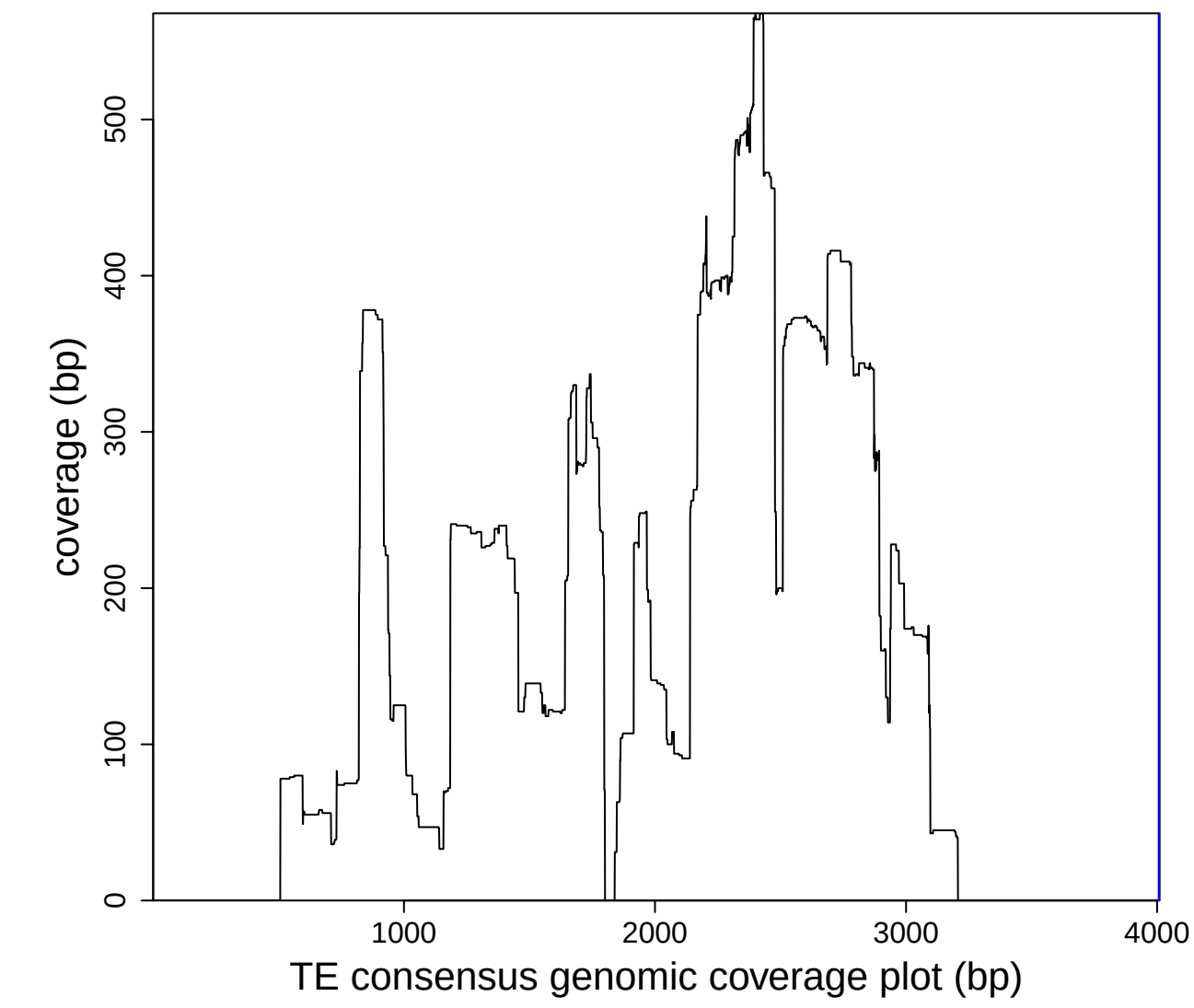
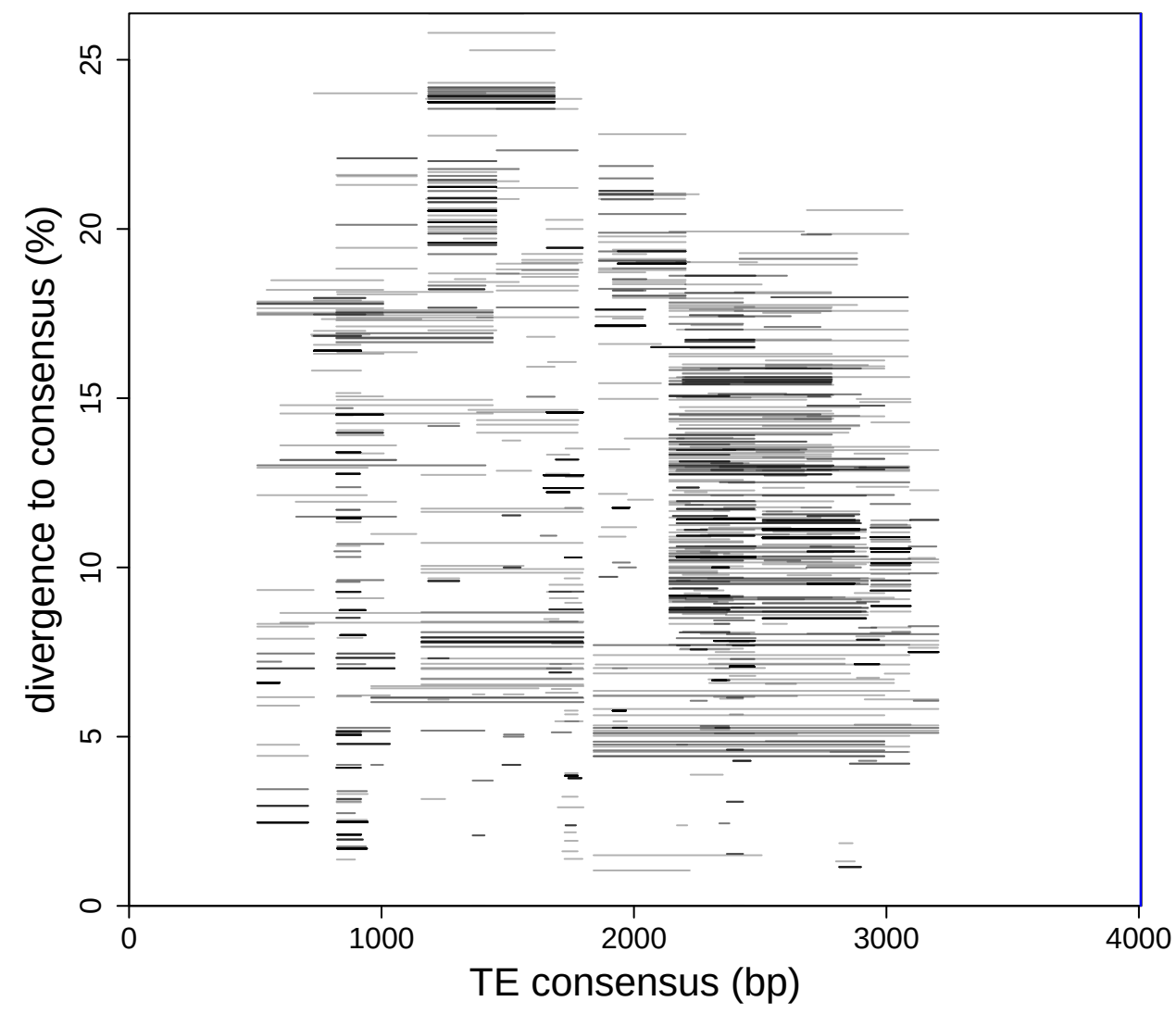
a.b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.f
size: 3950bp; fragments: 3500; full length: 0 (>=3555bp)

After TEtrimmer 3950 bp

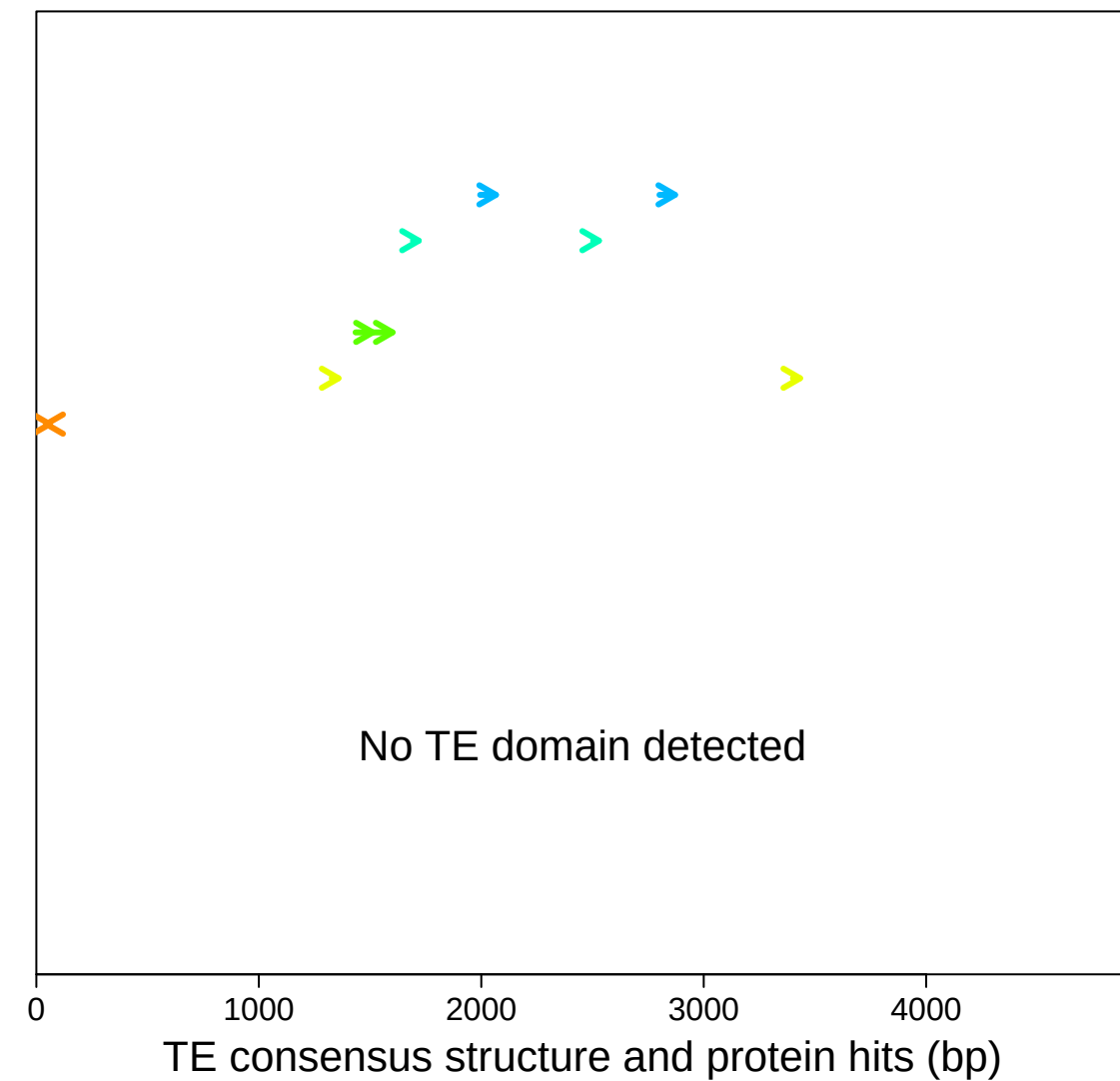
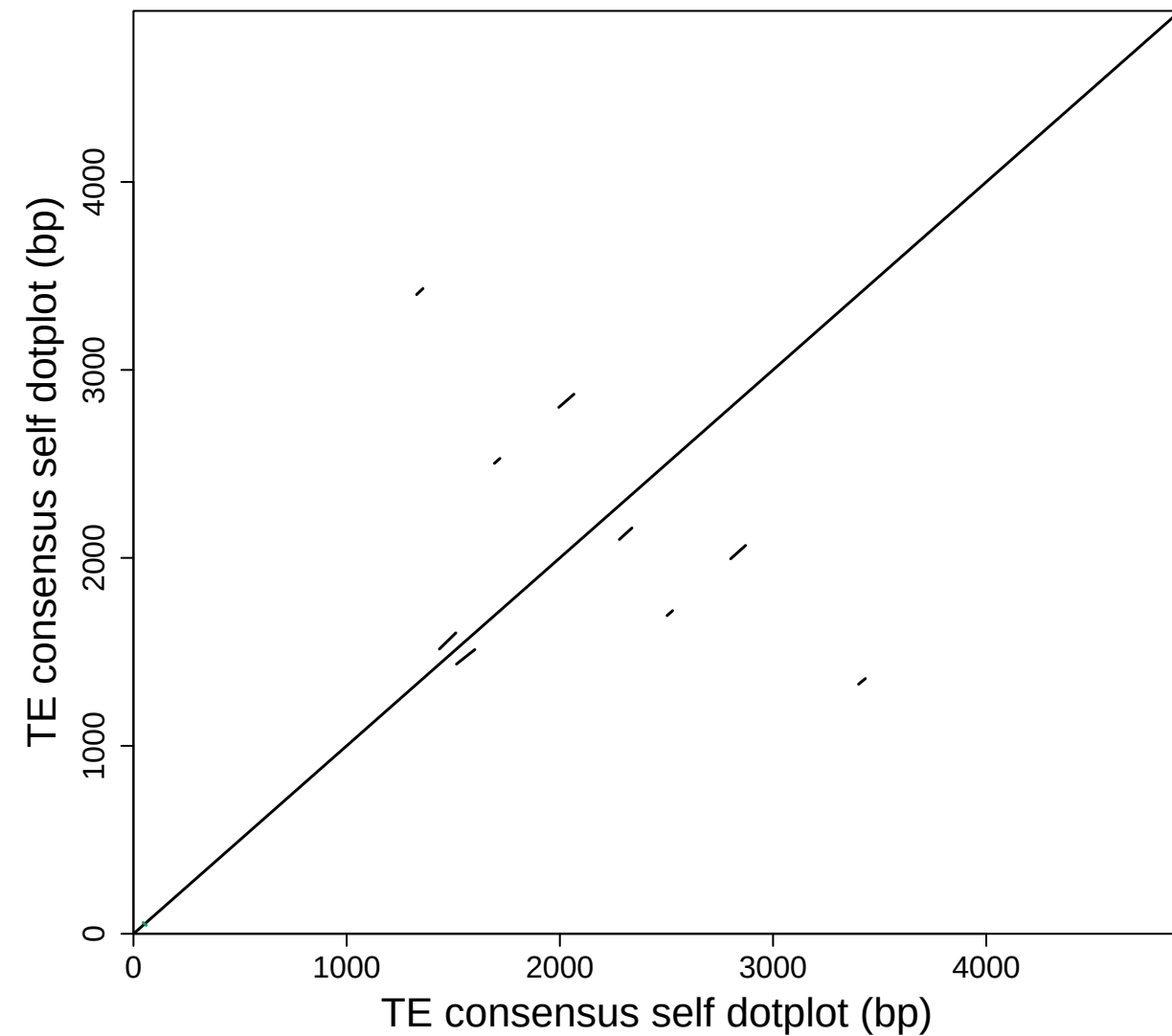
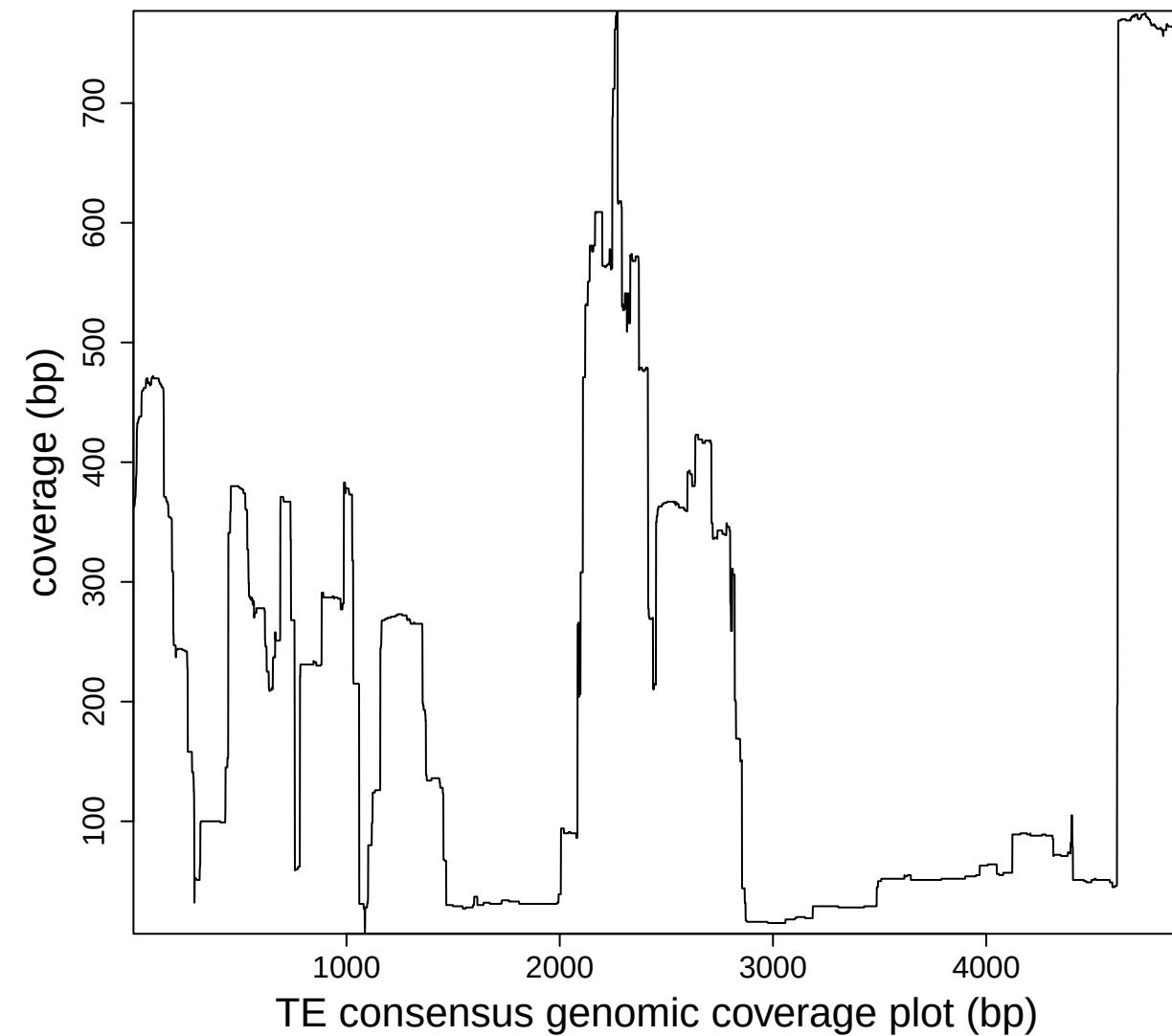
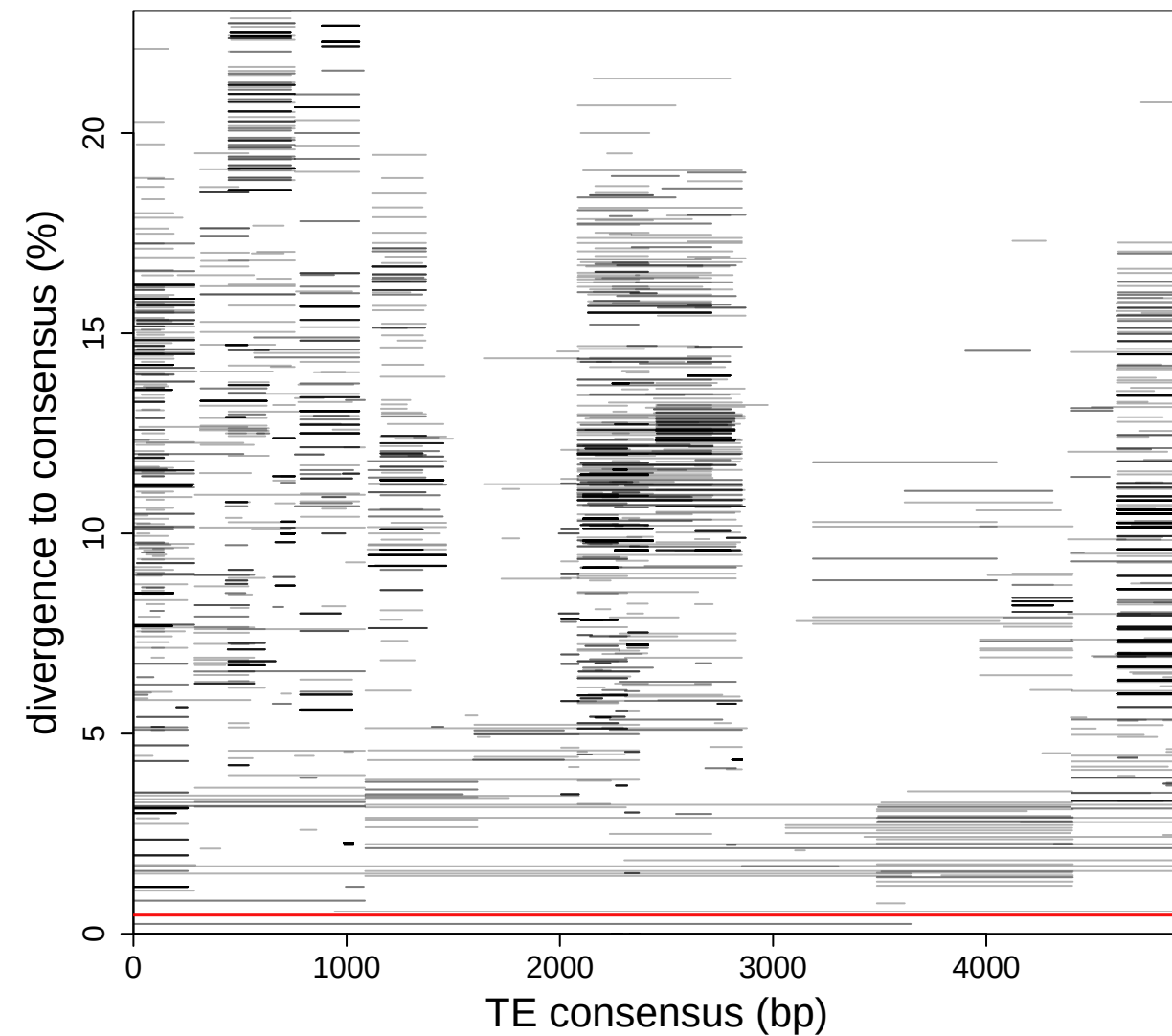


8.fasta.b.bed uf.bed g_2.bed fm_1.bed 0_0_bcln.fa aln.fa cl.f
size: 4010bp; fragments: 2393; full length: 0 (>=3609bp)

After TEtrimmer Extended plot Blue lines are boundaries

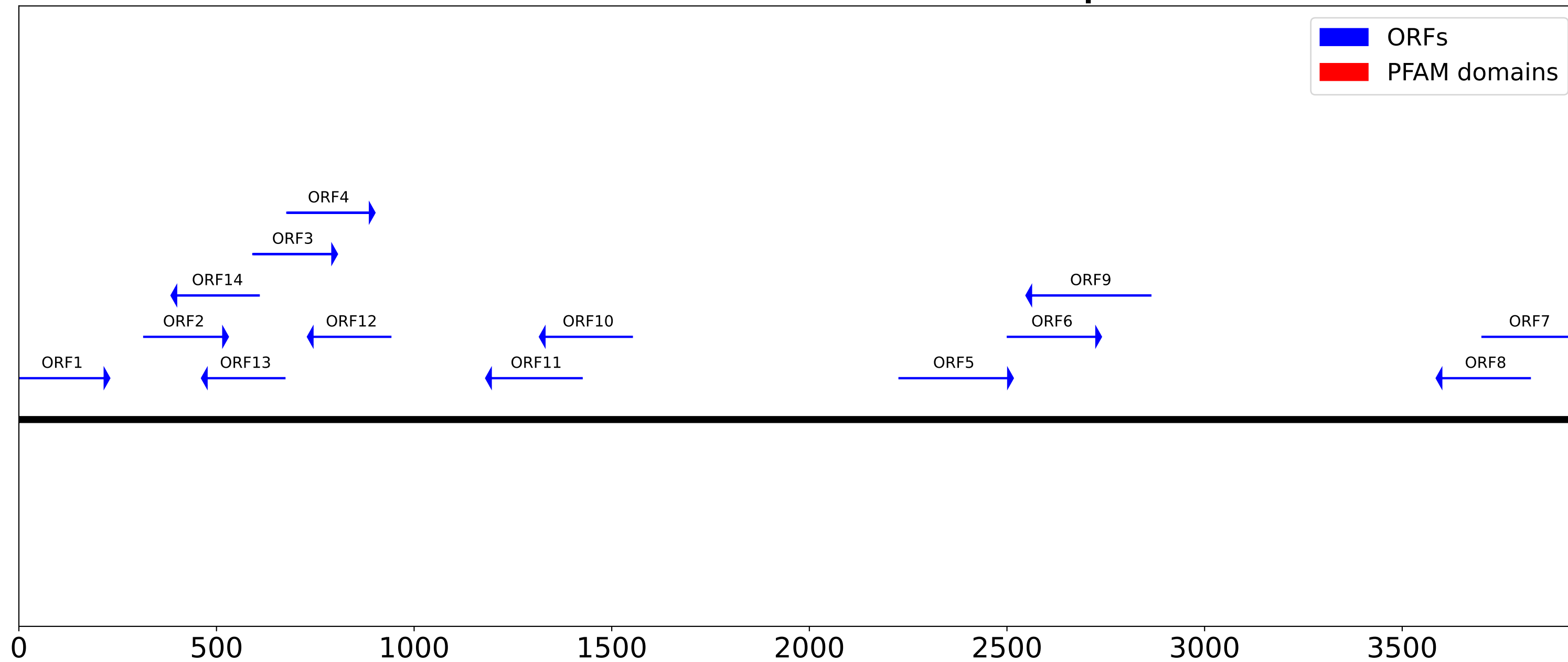


TE: rnd_5_family_3948
size: 4910bp; fragments: 3997; full length: 1 (>=4419bp)

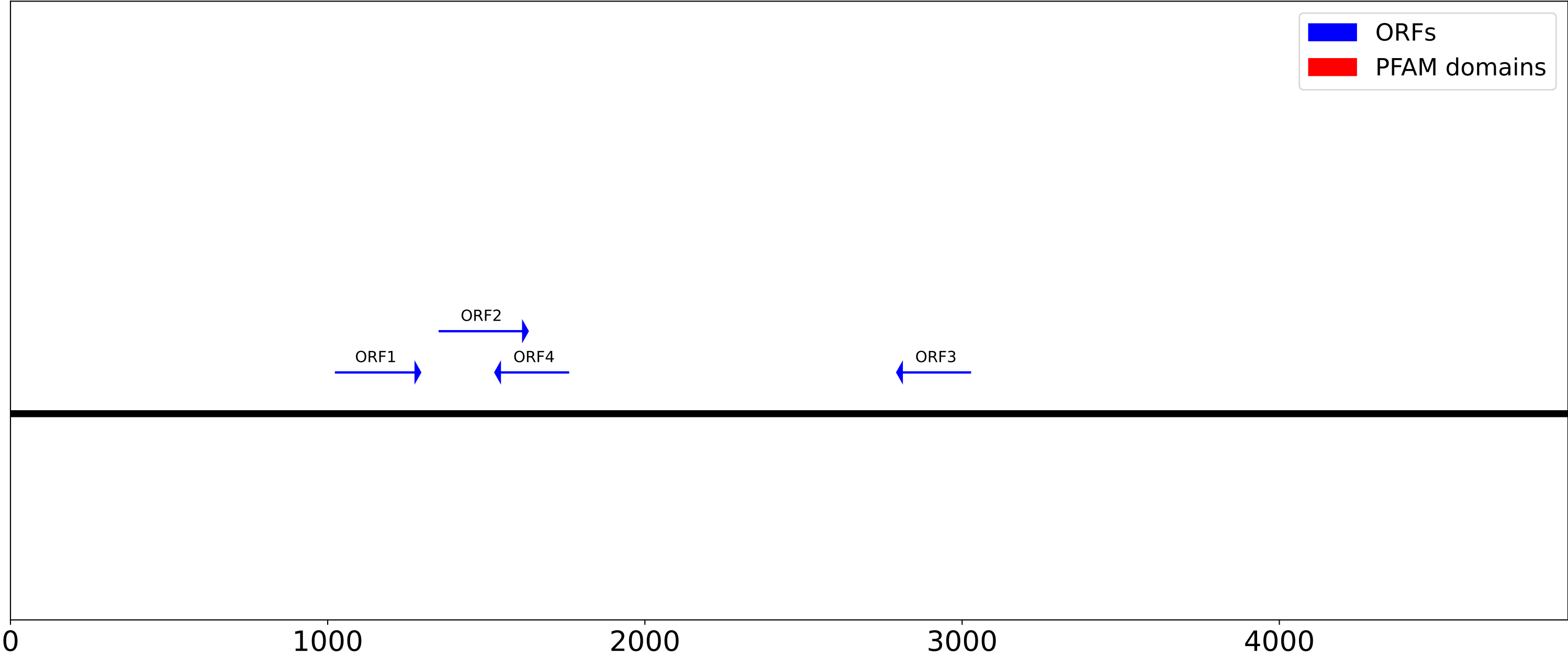


Before TETrimmer 4910 bp

After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

