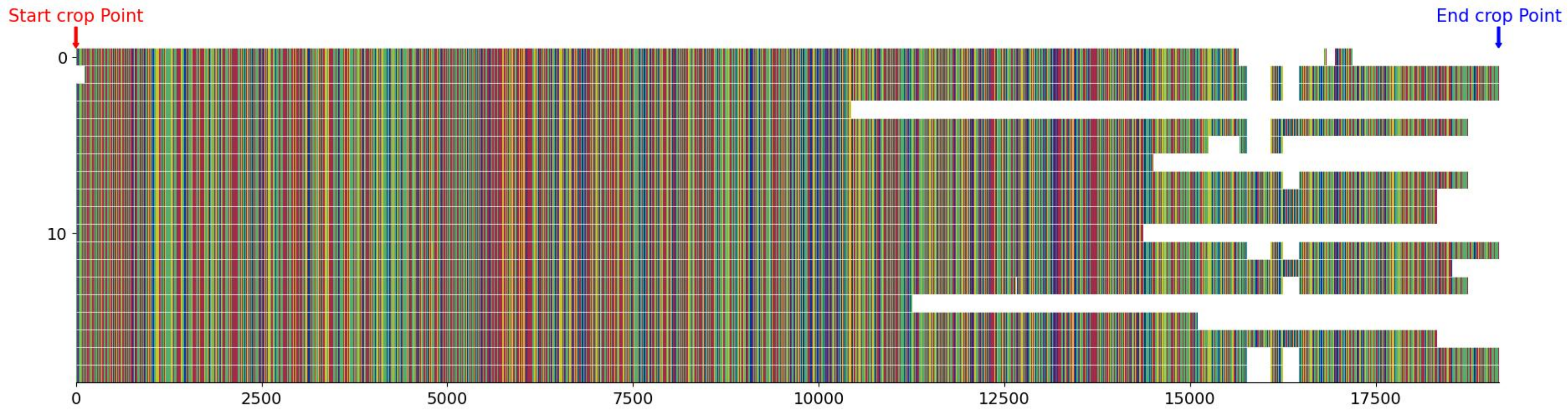




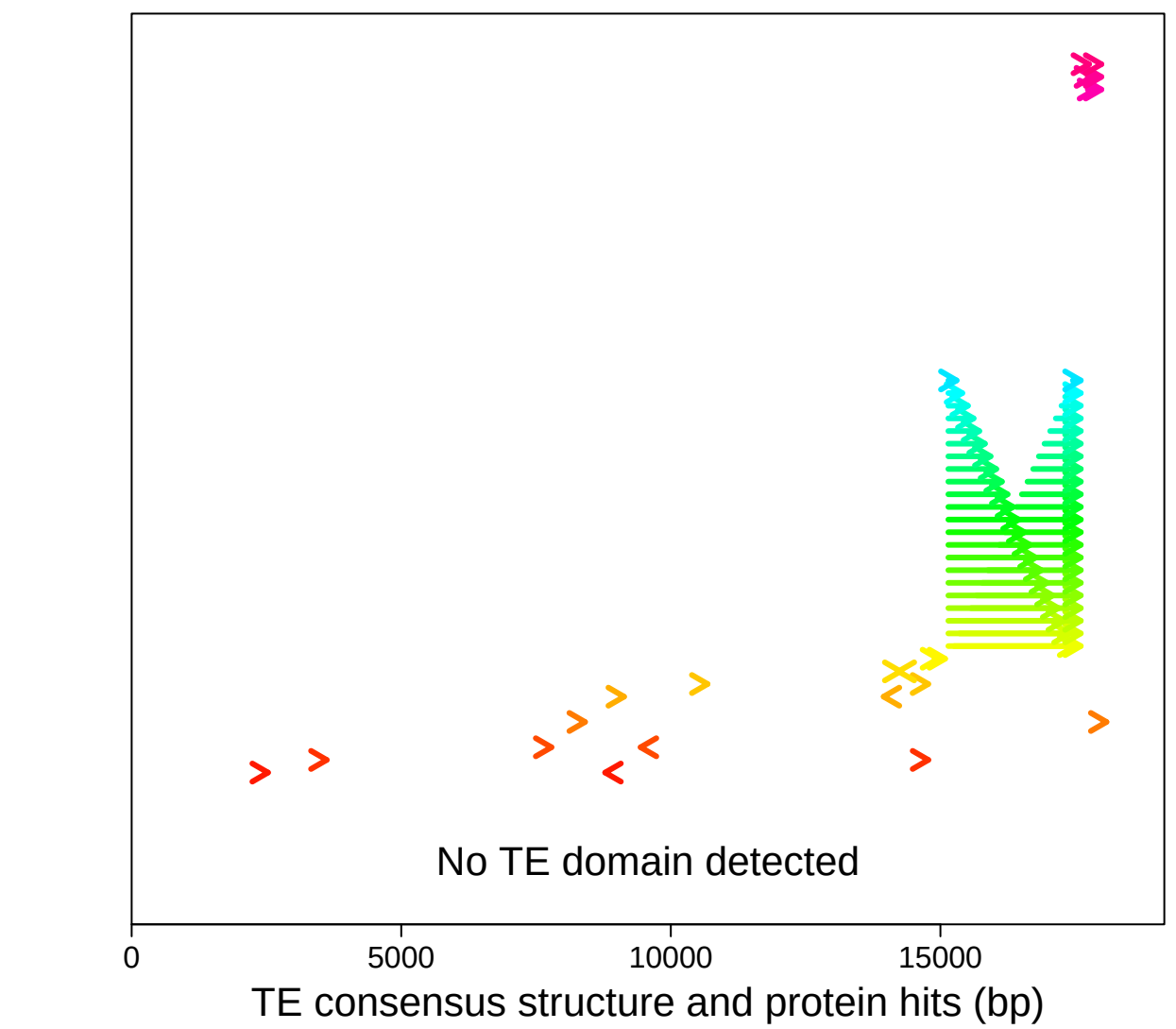
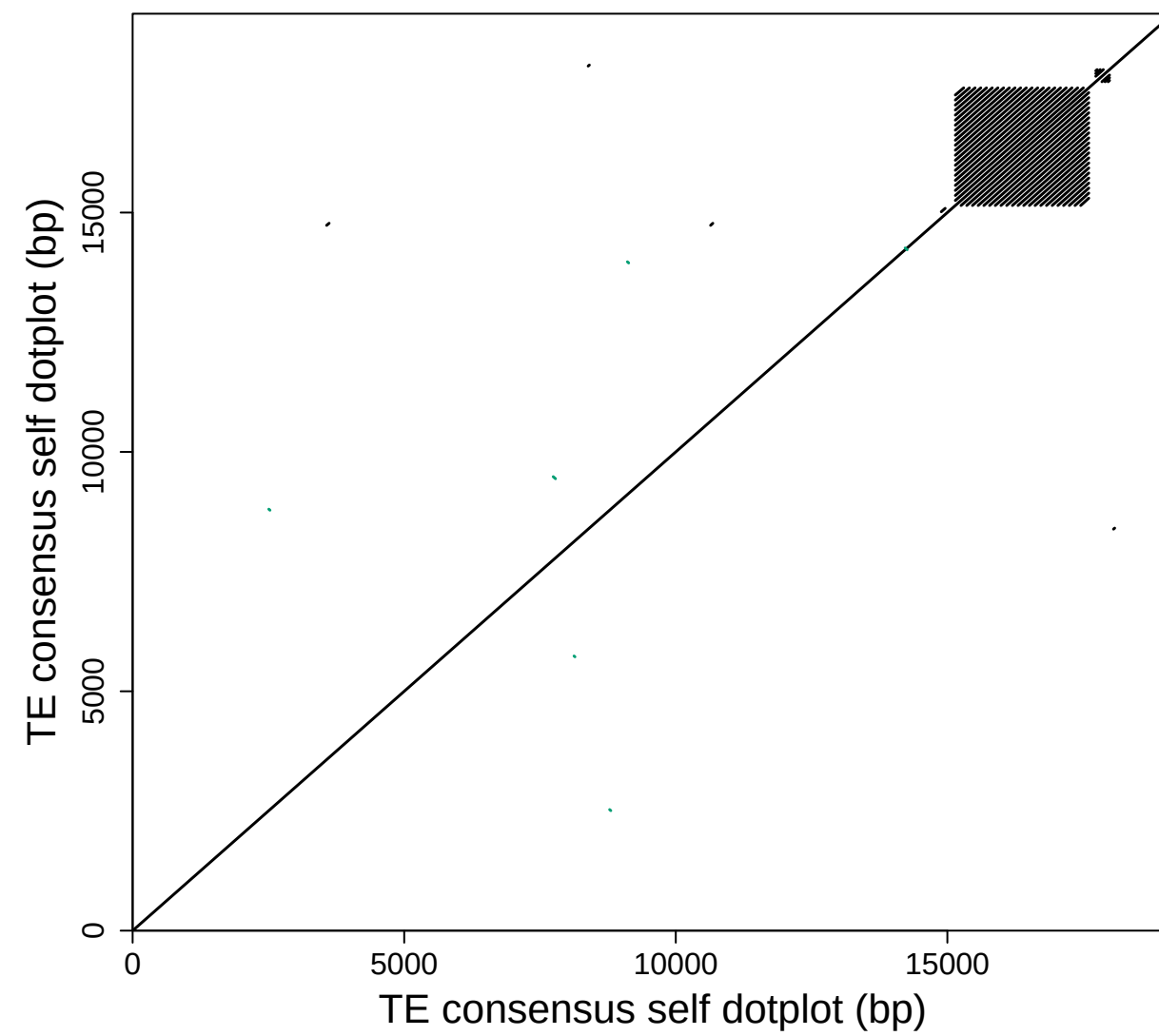
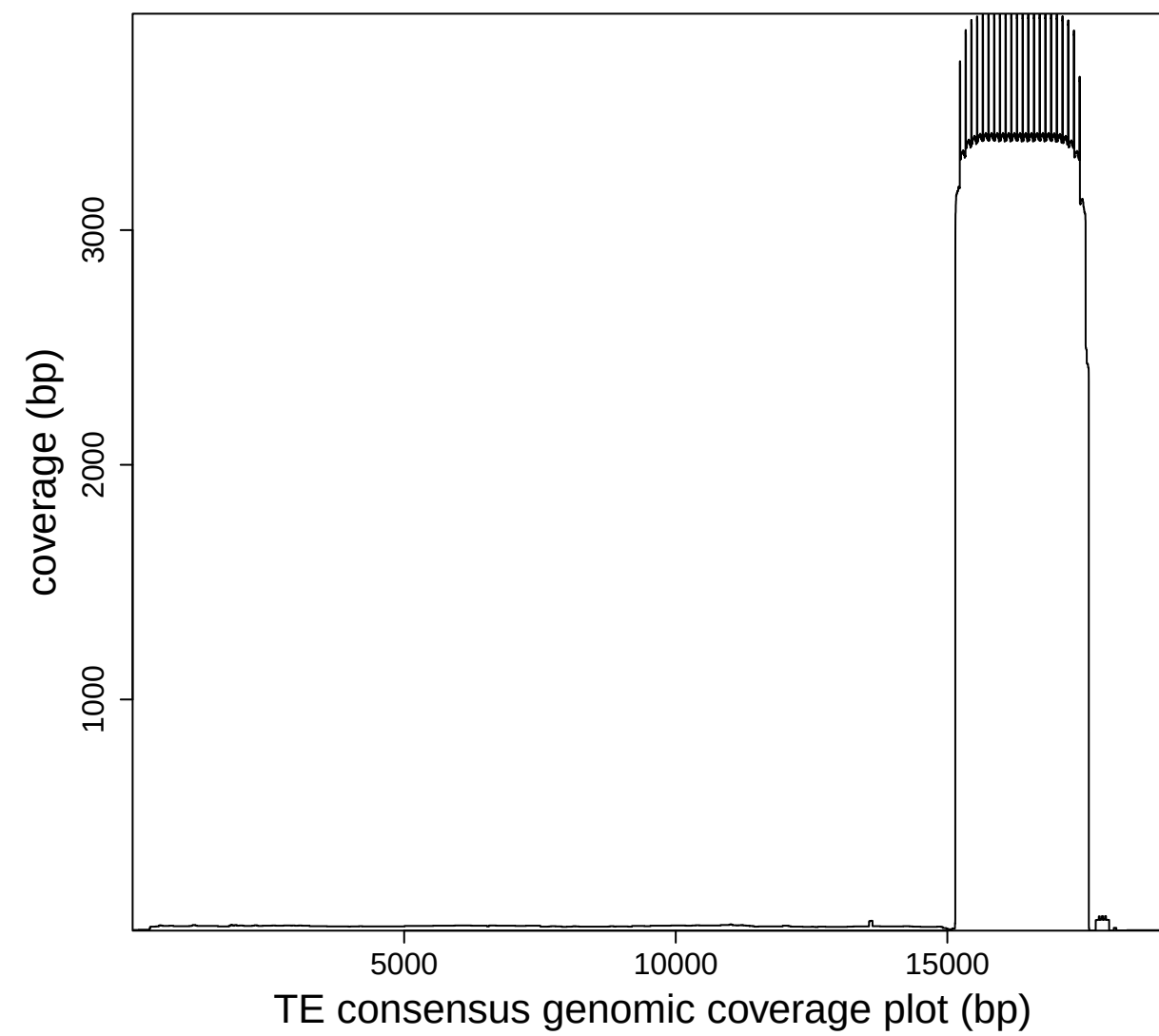
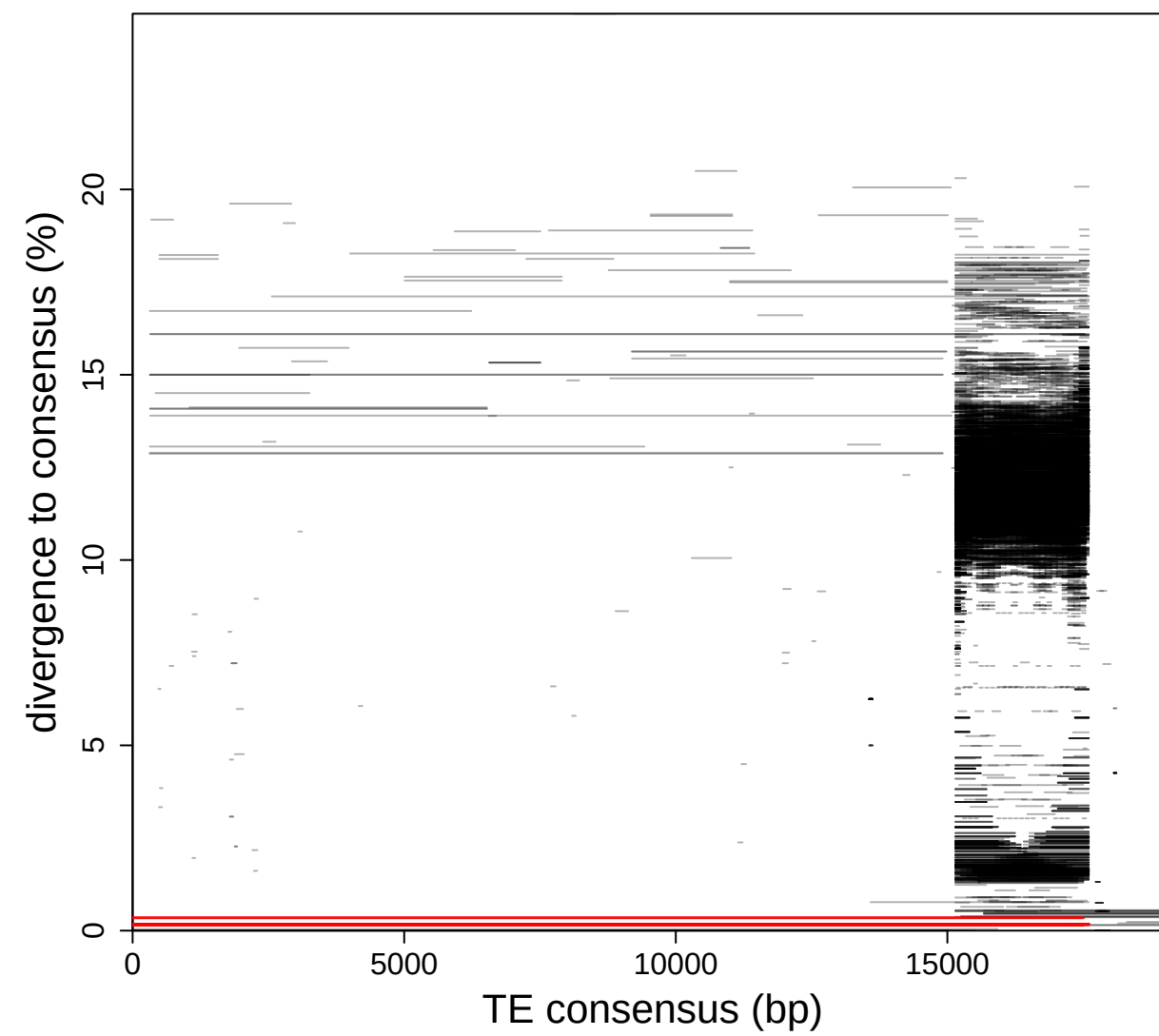
MSA length = 19153

[illegible]



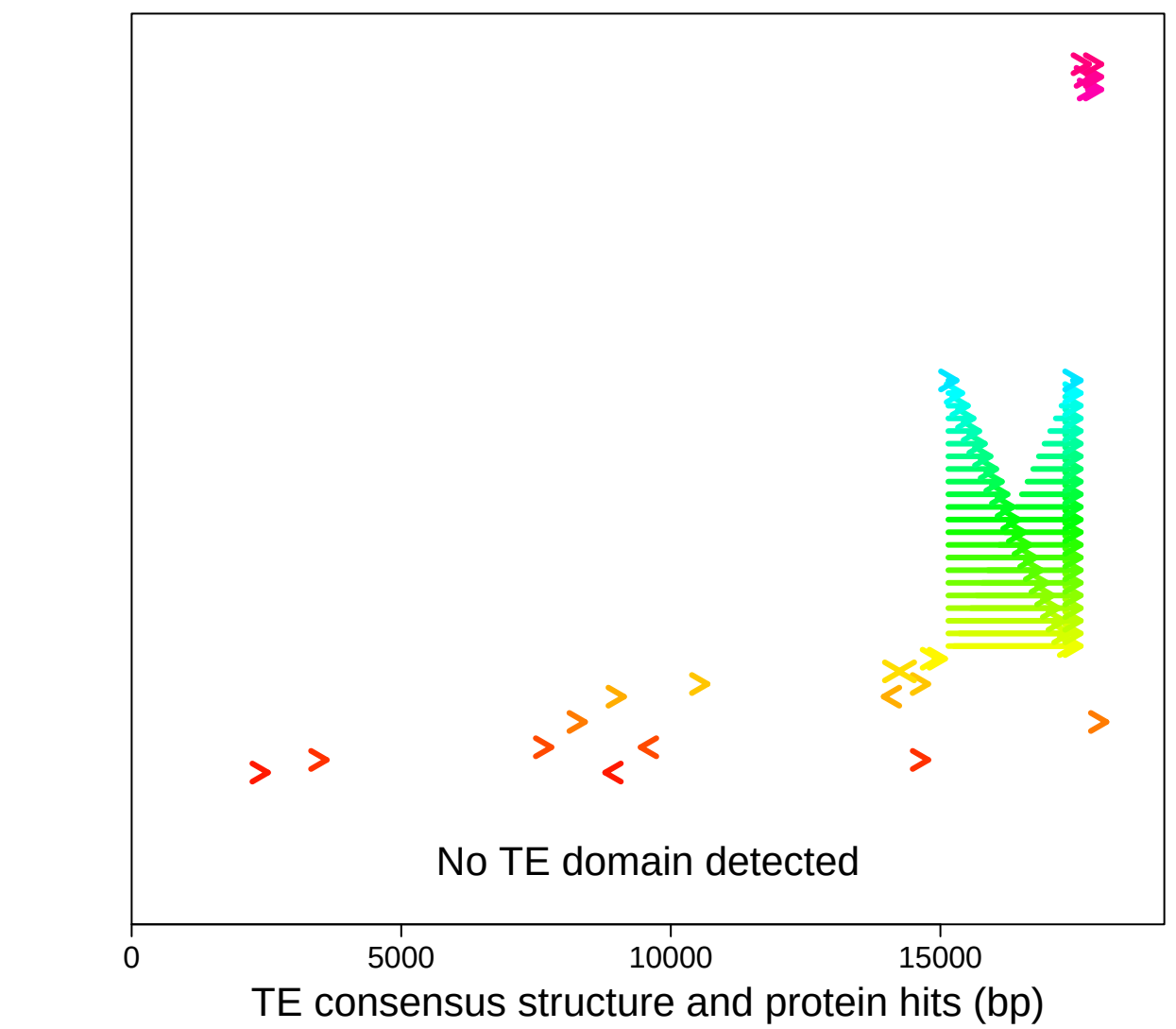
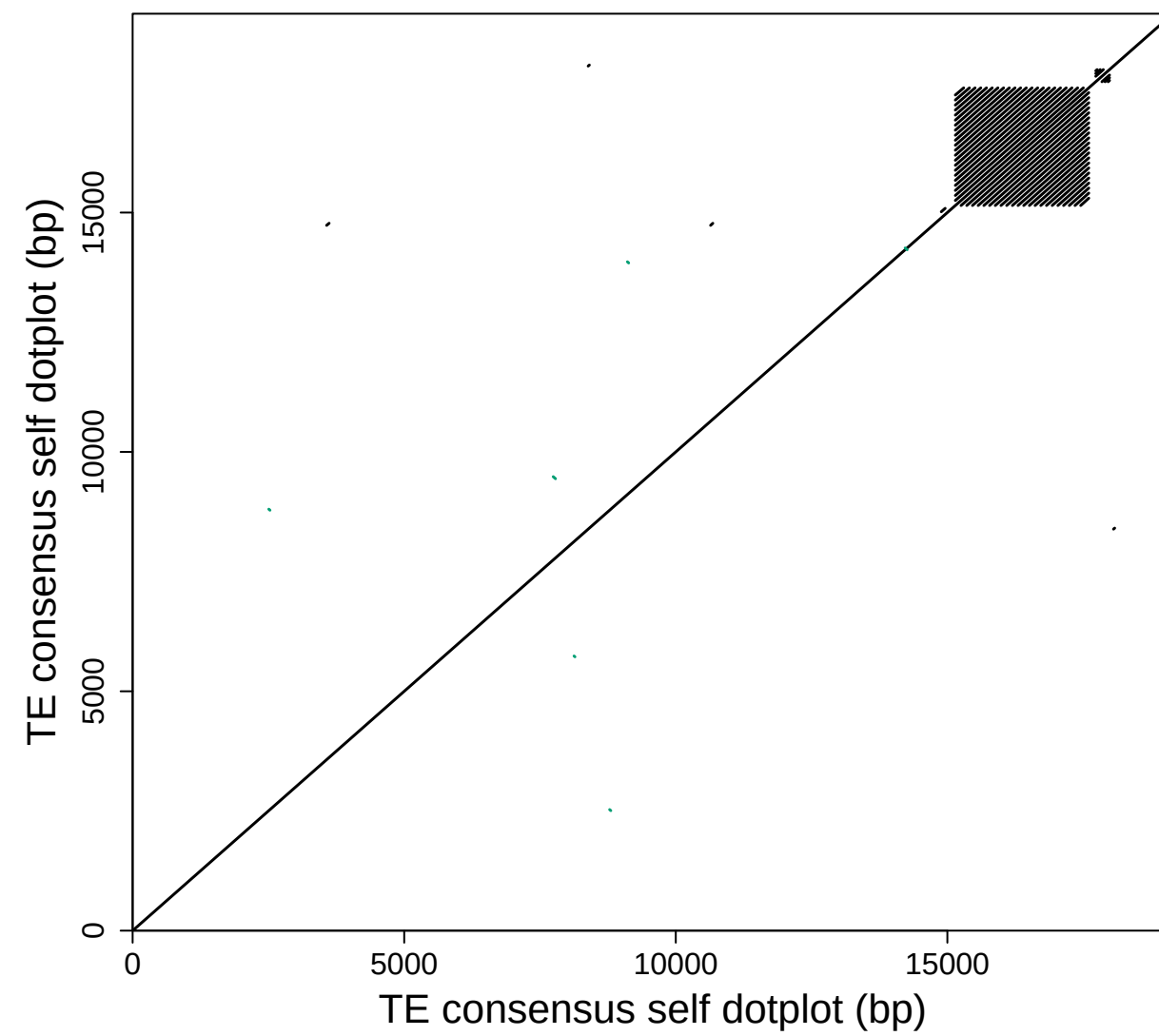
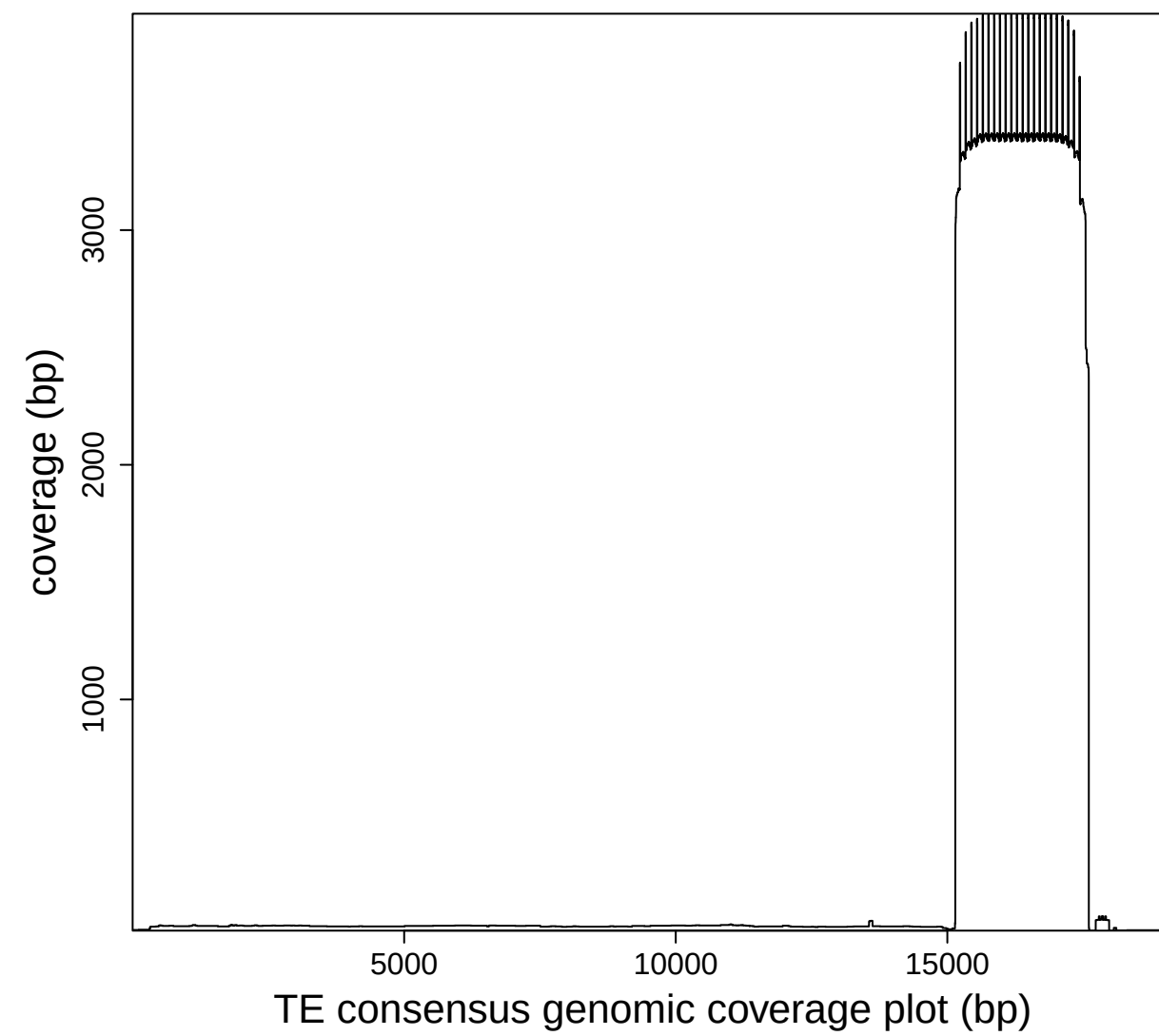
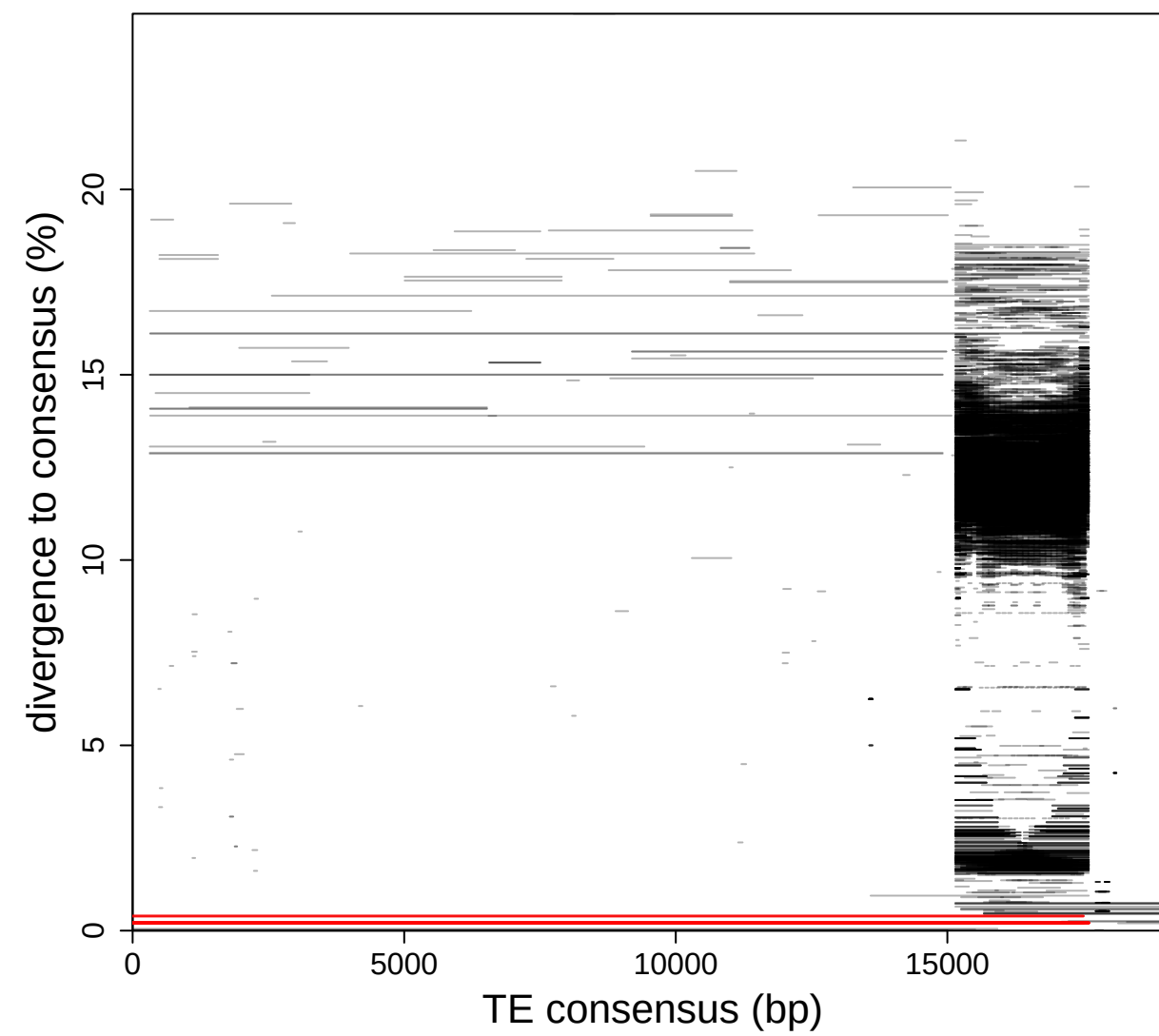
a.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.f
size: 19153bp; fragments: 16924; full length: 7 (>=17237.7bp)

After TEtrimmer 19153 bp



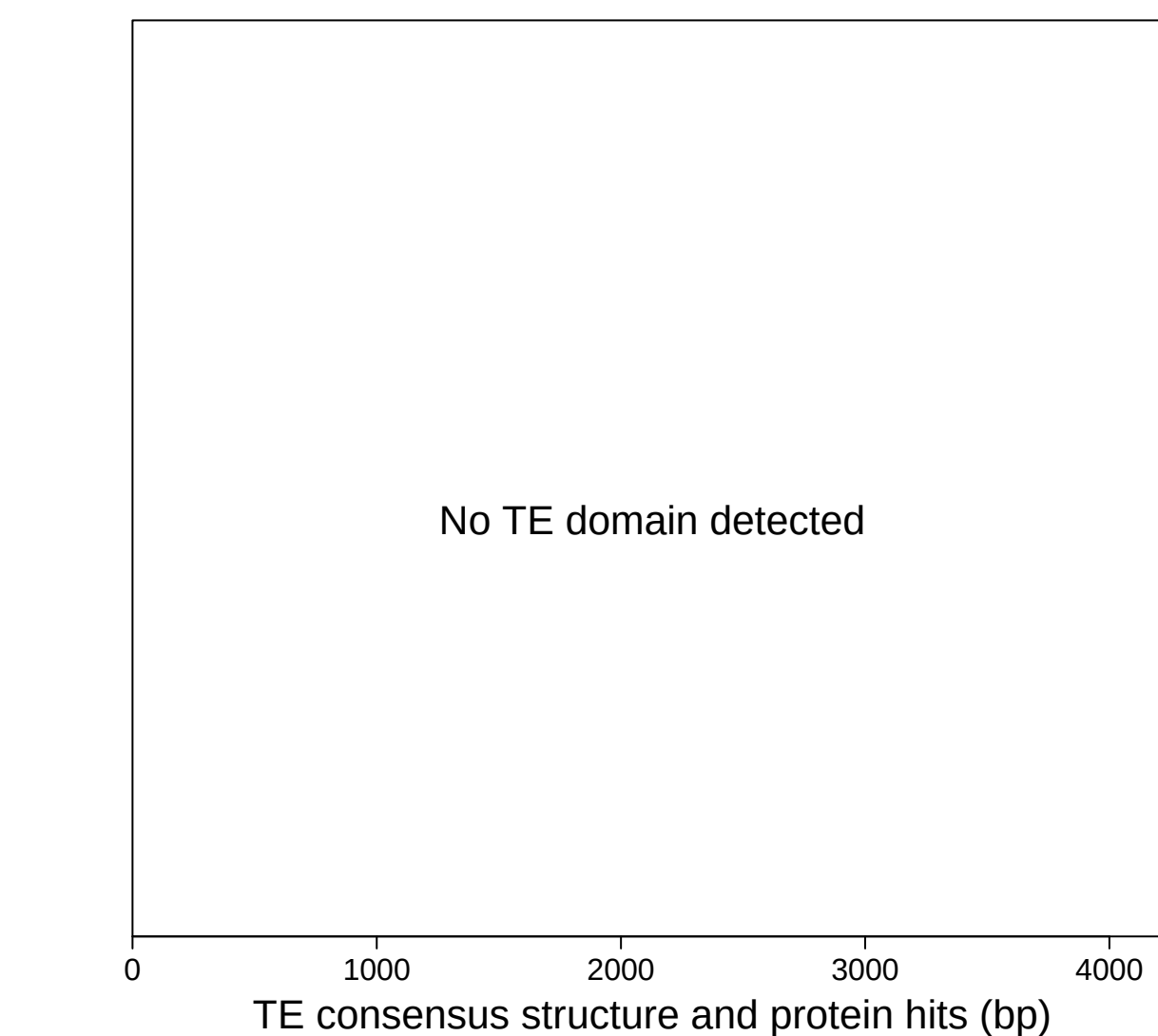
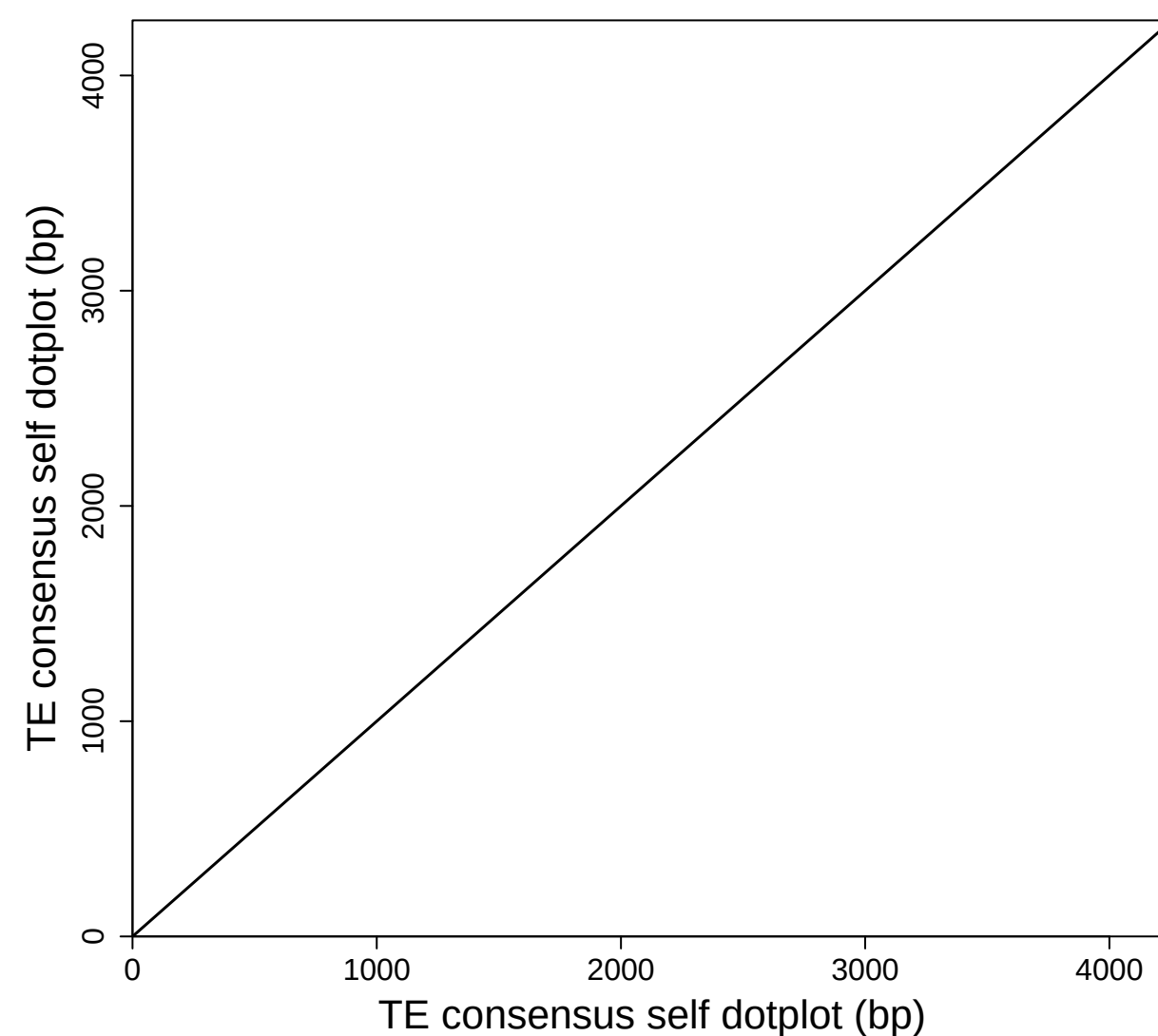
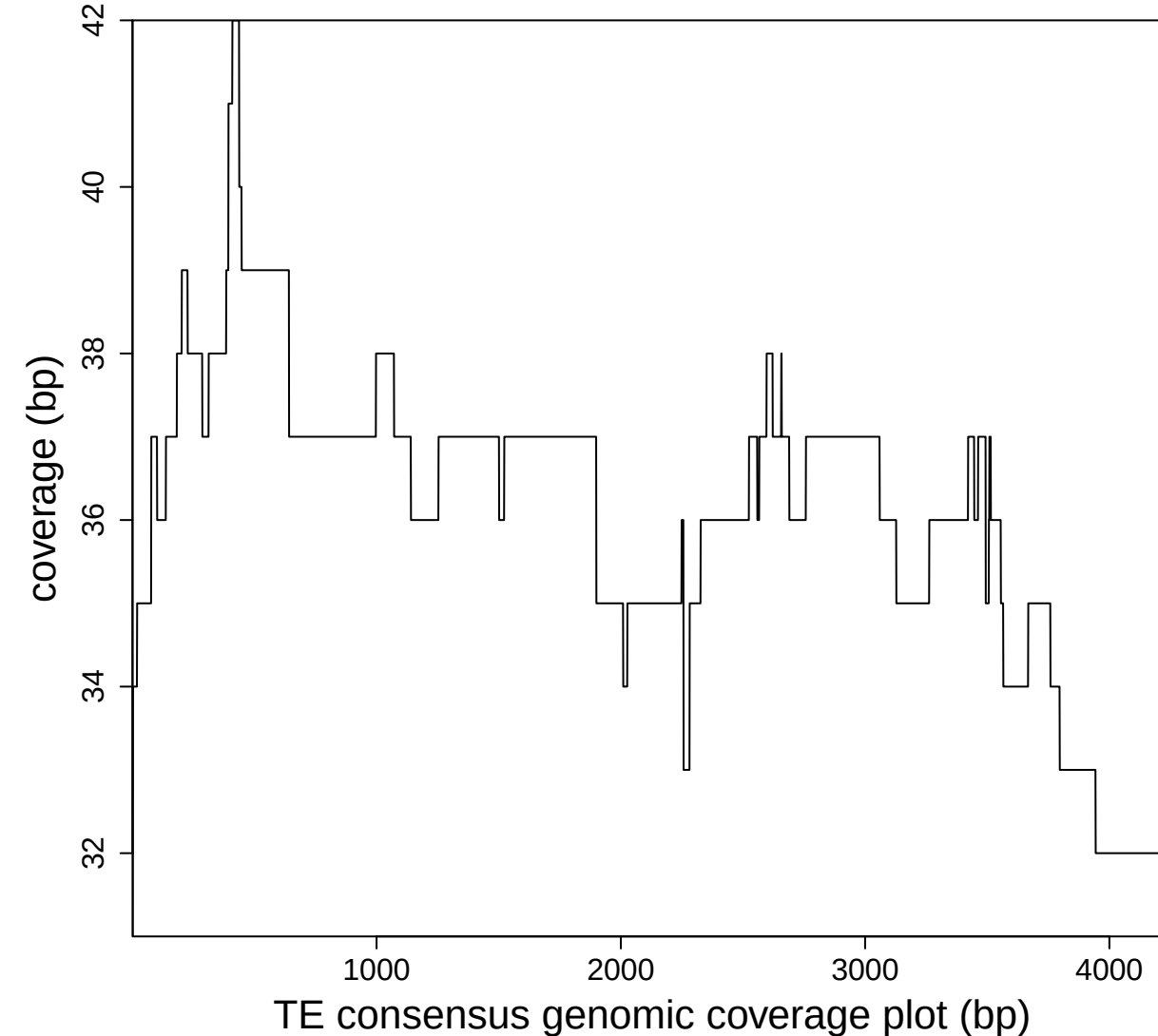
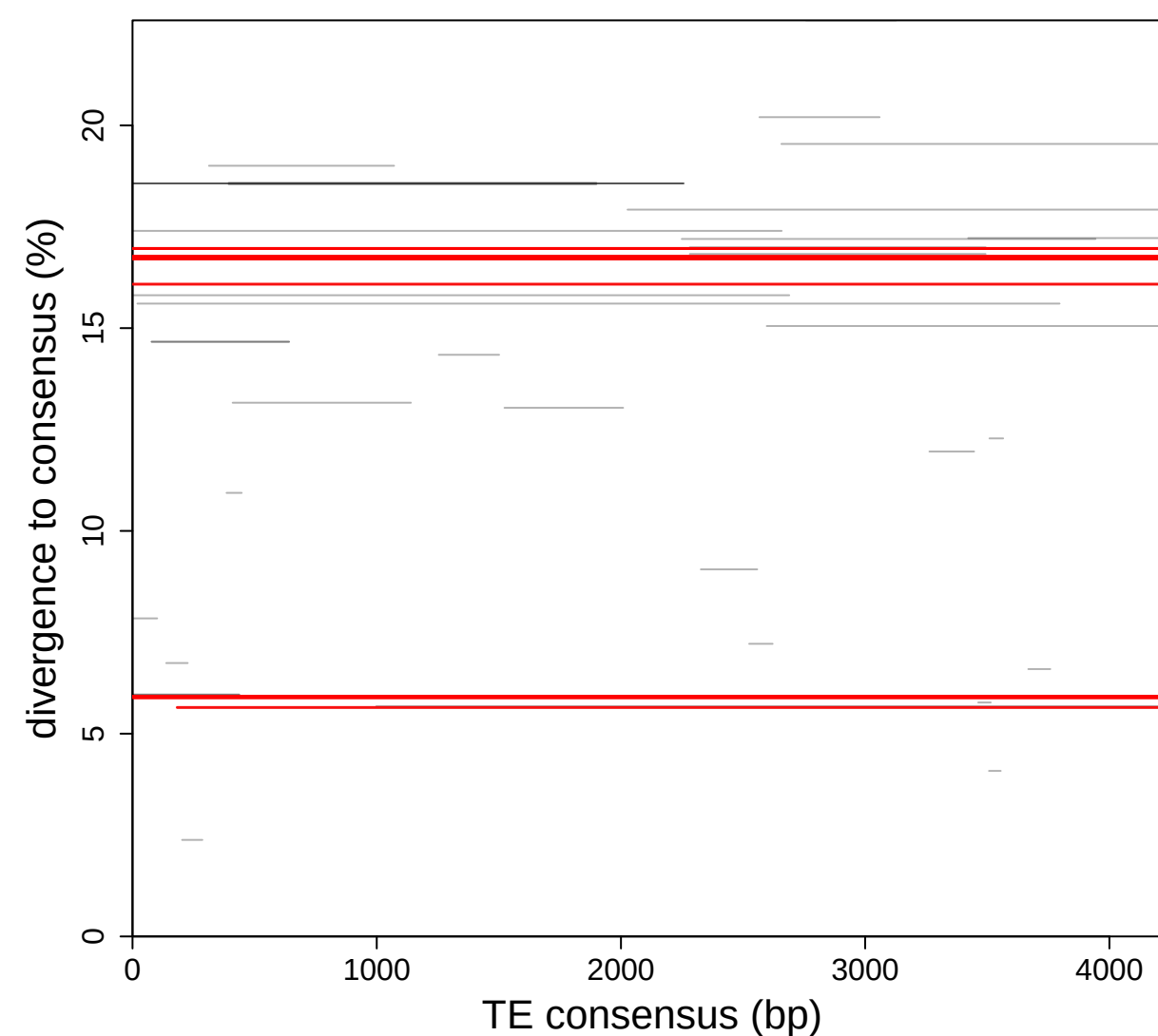
asta.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_c
size: 19153bp; fragments: 16914; full length: 7 (>=17237.7bp)

After TEtrimmer Extended plot Blue lines are boundaries

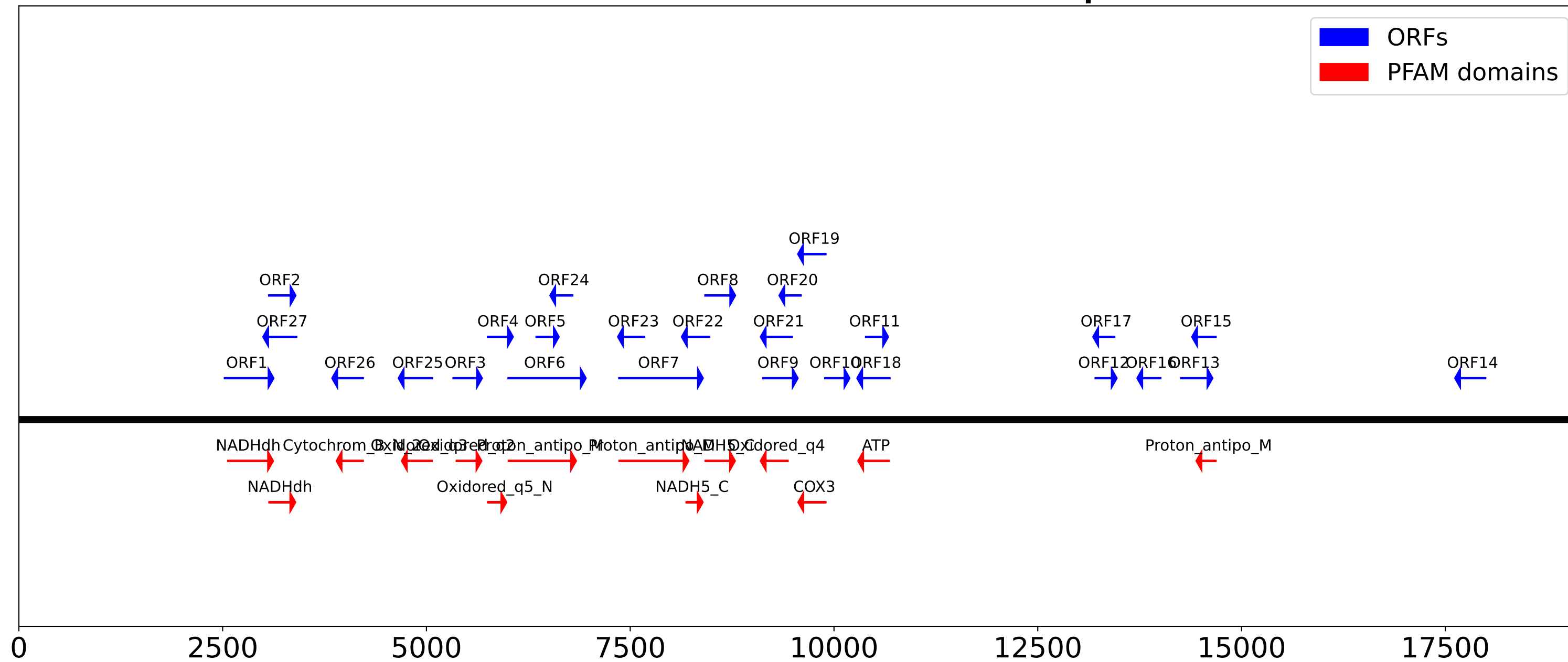


TE: rnd_5_family_4324
size: 4256bp; fragments: 64; full length: 27 (≥ 3830.4 bp)

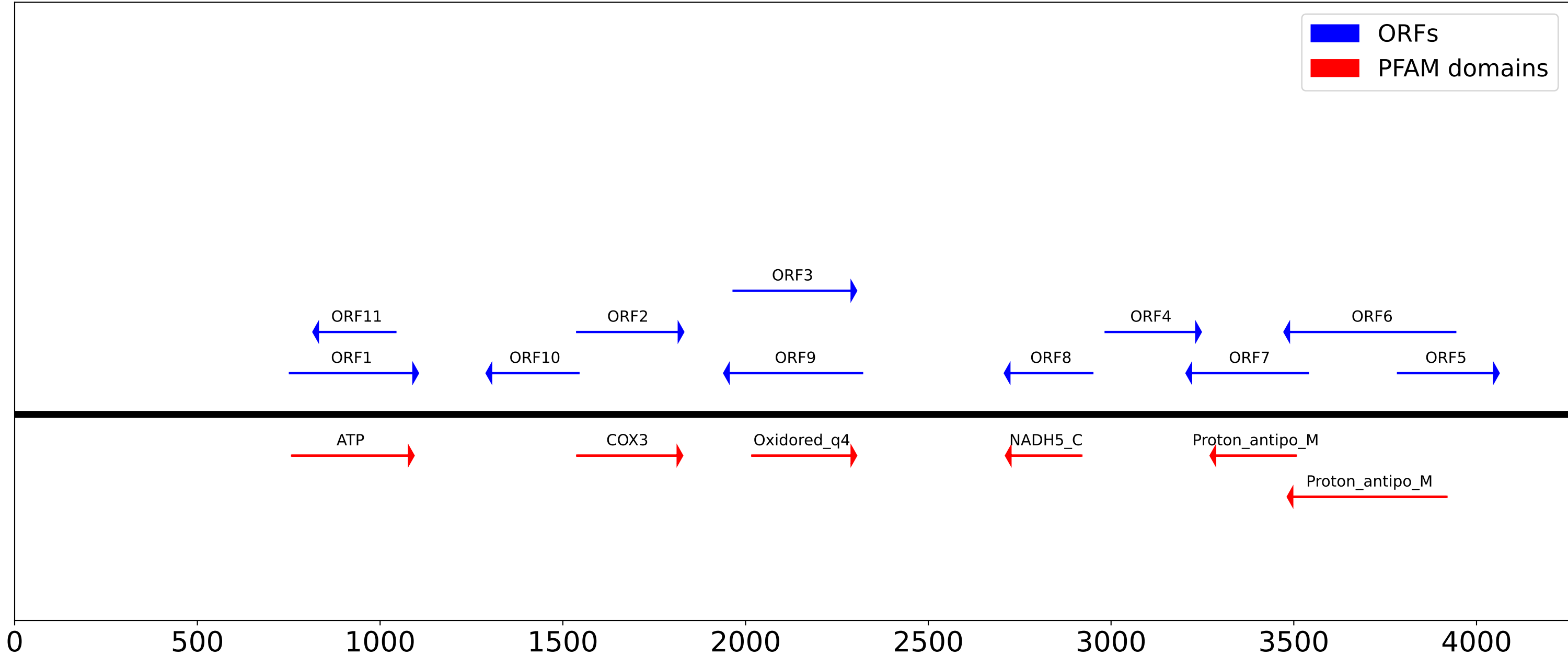
Before TEtrimmer 4256 bp



After TETrimmer ORF and PFAM domain plot



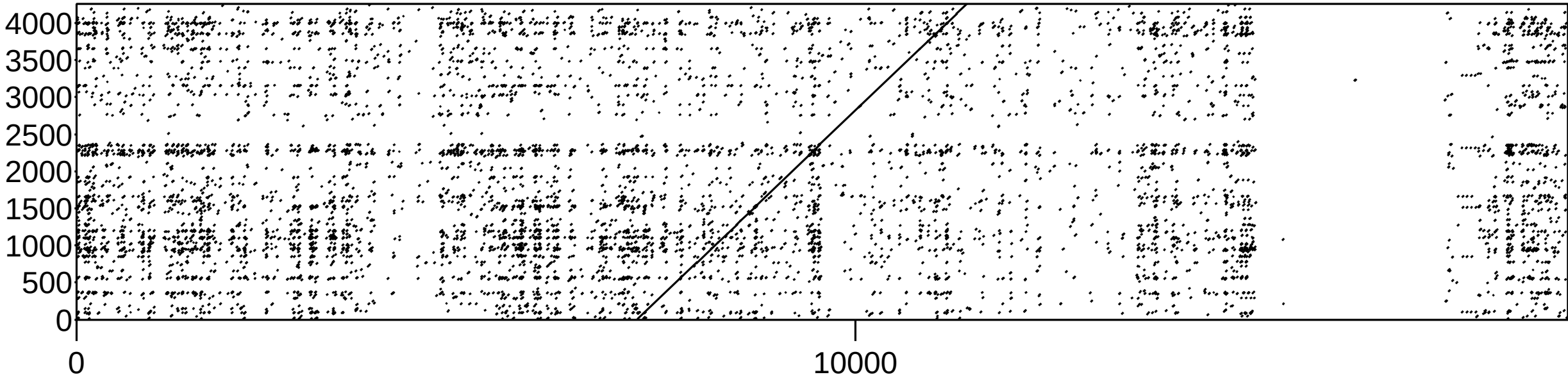
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

onsensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)