

Start crop Point



End crop Point



MSA length = 16385

TG TAGAAGAACCATTTC TAAAGCAT AATCATAACGGTCGCGTCGGTTTCGTGGTTCTGGAGCTG - - - - -

ATTAGCTCAAGTA - - - - -

TG TAGAAGAACCATTTC TAAAGCAT AATTATAACGGTCGCGTCGGTTTCGTGGTTCTGGAGCTG - - - - - AATTATGAAAAAAAC TTTAATGG AATGAAAC TTGGAAAAAGAAACCATTTTTTTTA

TG TAGAAGAACCATTTC TAAAGCAT AATTATAACGGTCGCGTCGGTTTCGTGGTTCTGGAGCTG - - - - - AATTATGAAAAAAAC TTTAATGG AATGAAAC TTGGAAAAAGAAACCATTTTTTTTA

- - - - - TTTCTAAAGCAT AATTATAACGGTCGCGTCGGTTTCGTGGTTCTGGAGCTG - - - - - AATTATGAAAAAAAC TTTAACGG AATGAAAC TTGGAAAAAGAAACCATTTTTTTTA

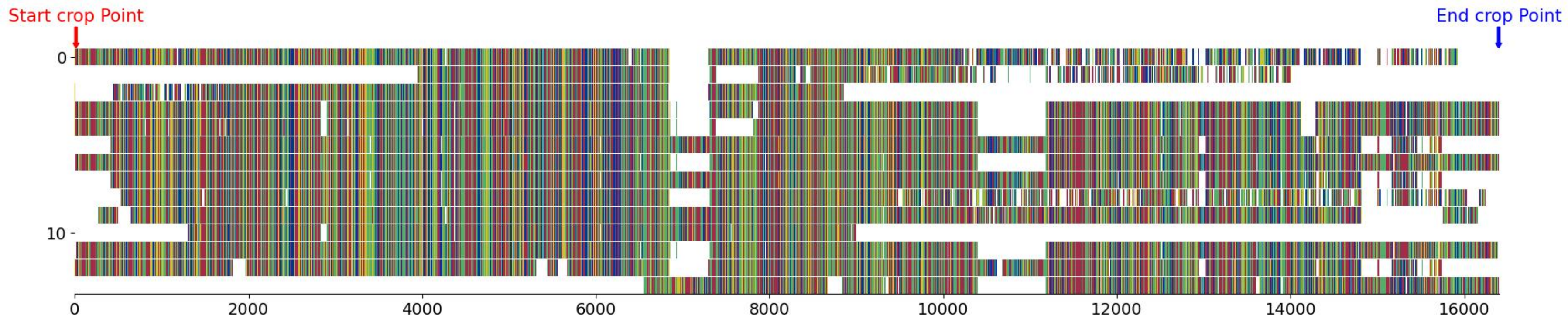
- - - - -

- - - - -

- - - - - TAAAGCAT AATTATAACGGTCGCGTCGGTTTCGTGGTTCTGGAGCTG - - - - - AATTATGAAAAAAAC TTTAATGG AATGAAAC TTGGAAAAAGAAACCATTT - - - - -

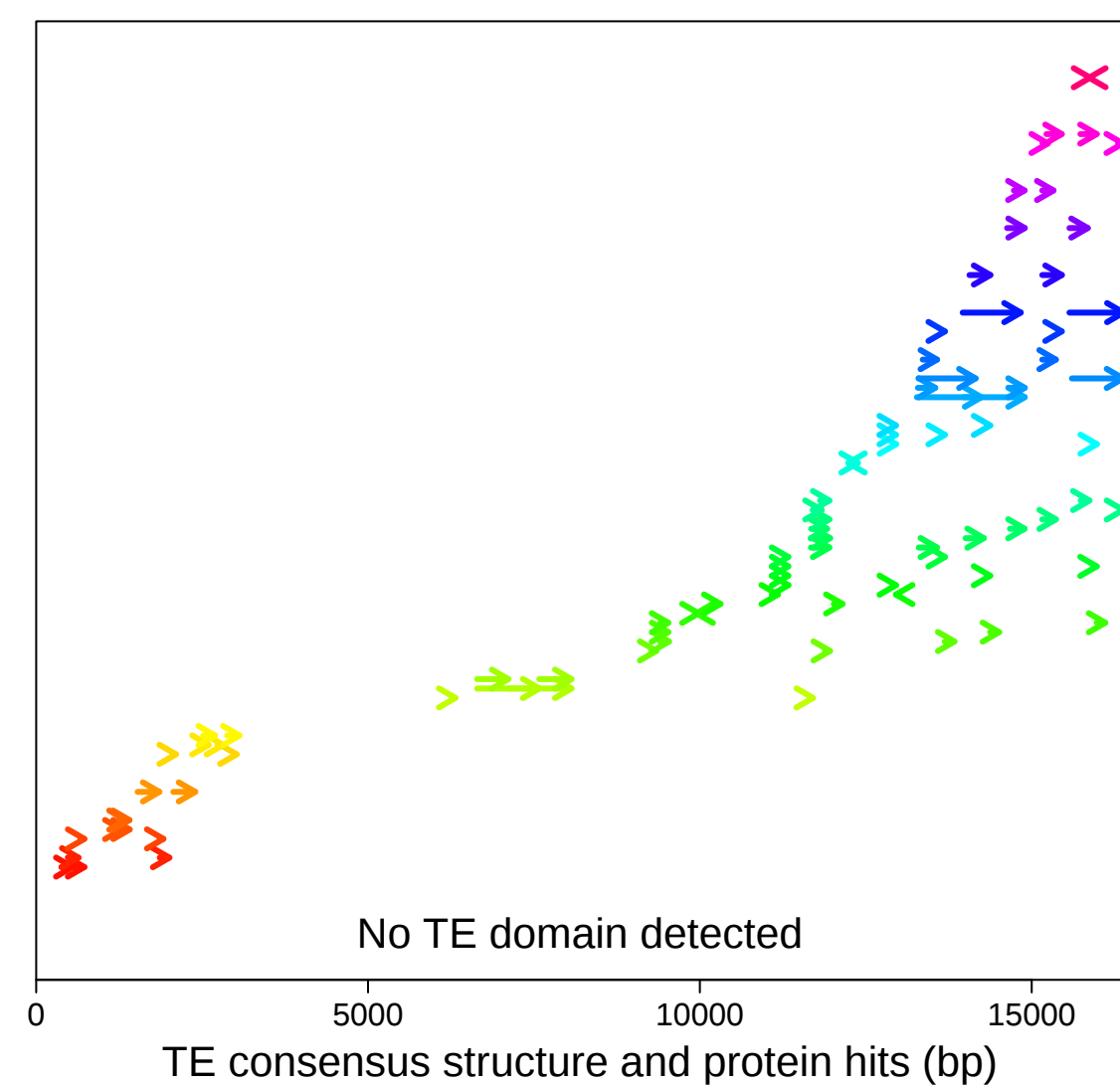
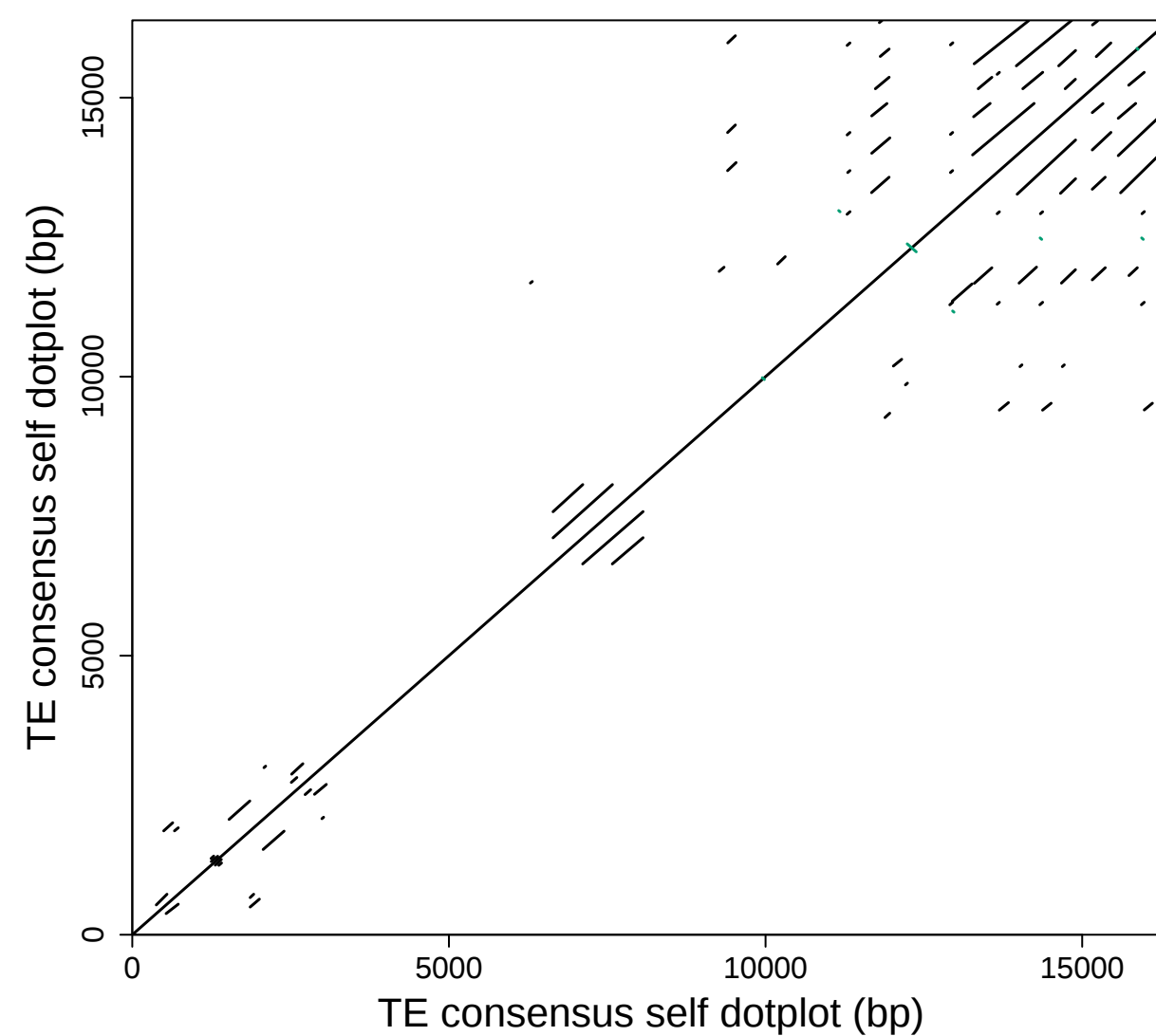
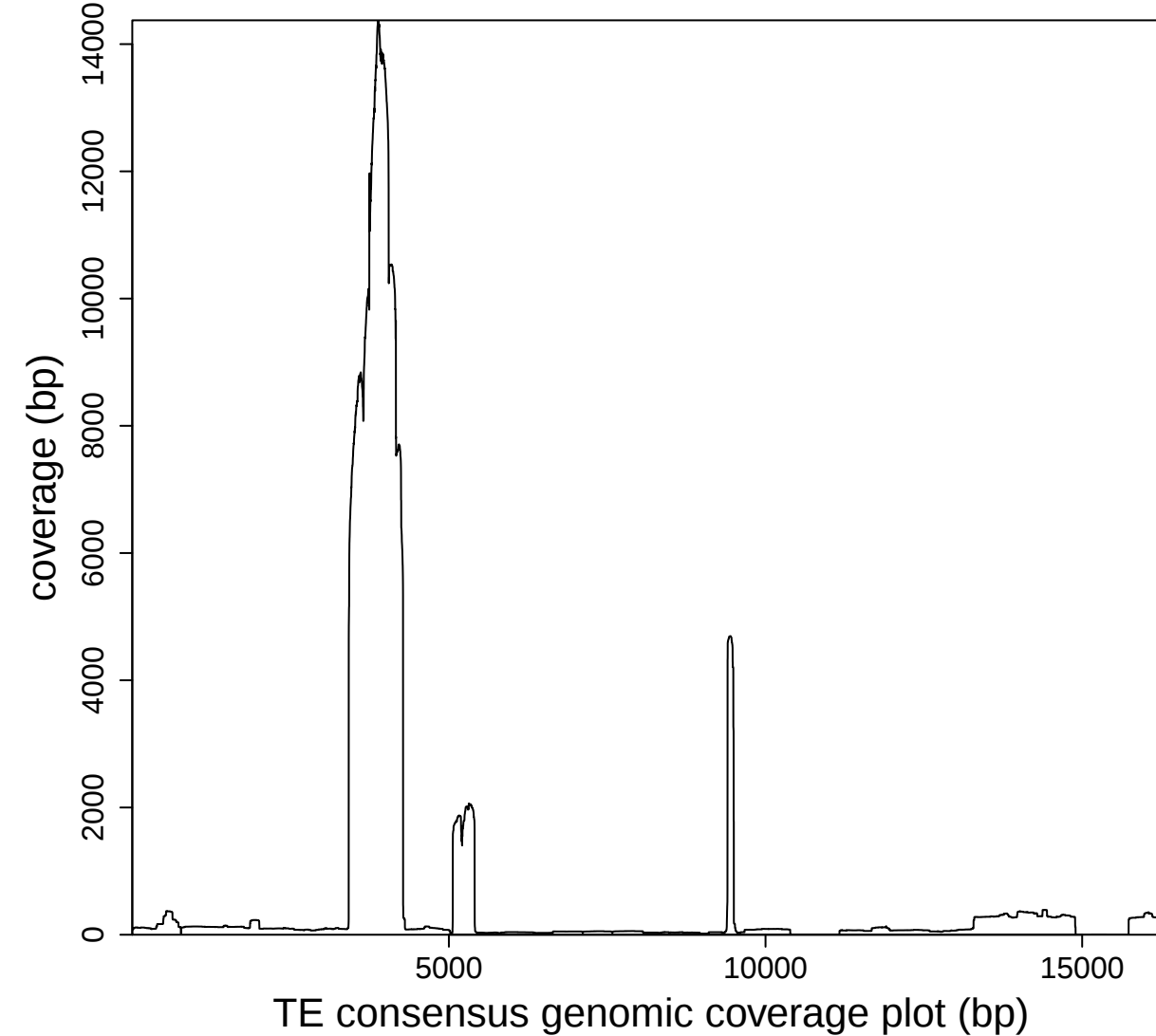
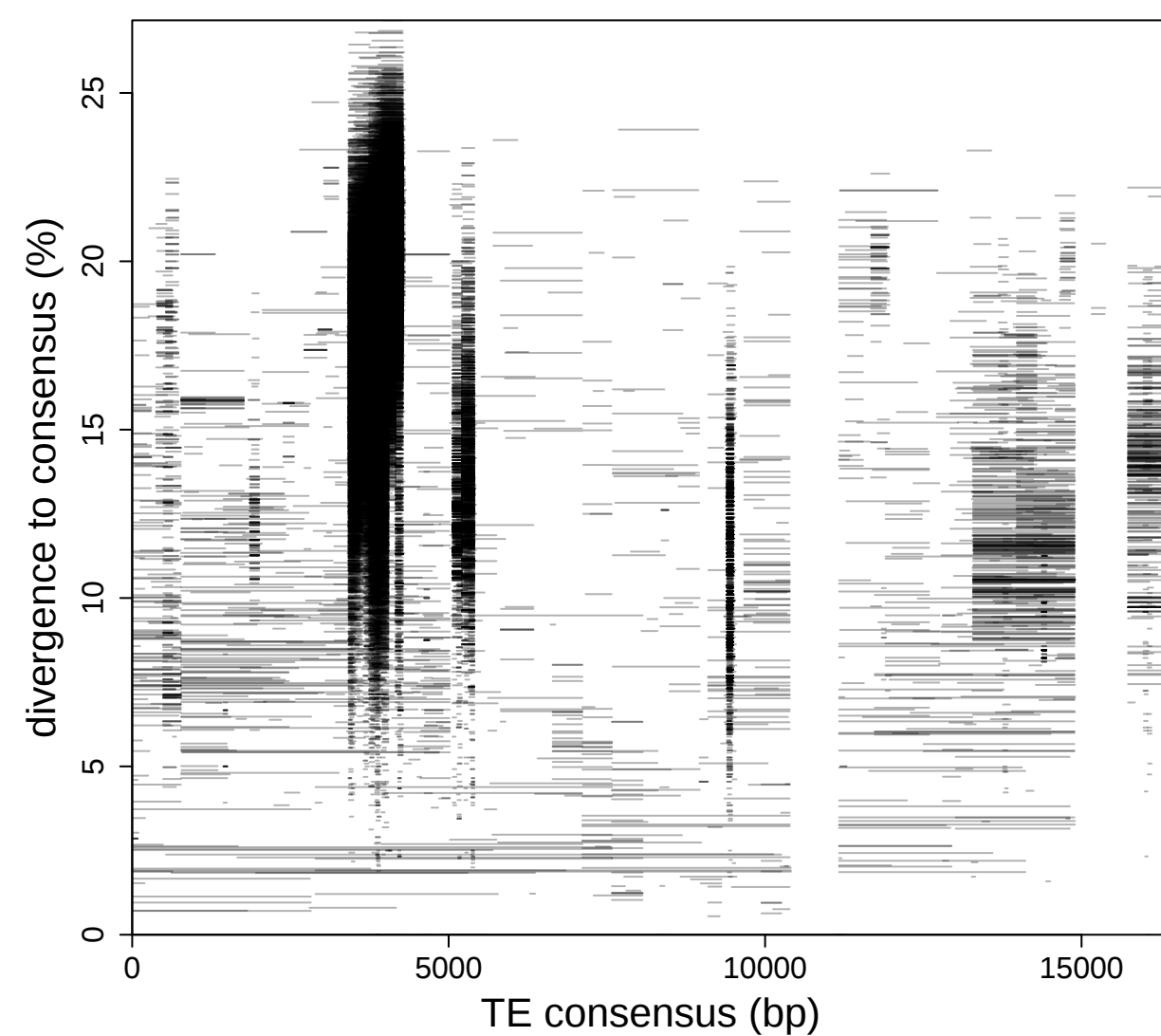
TG TAGAAGAACCATTTC TAAAGCAT AATTATAACGGTCGCGTCGGTTTCGTGGTTCTGGAGCTG - - - - -

- - - - - AATTATGAAAAAAAC TTTAATGG AATGAAAC TTGGAAAAAGAAACCATTTTTTTTA



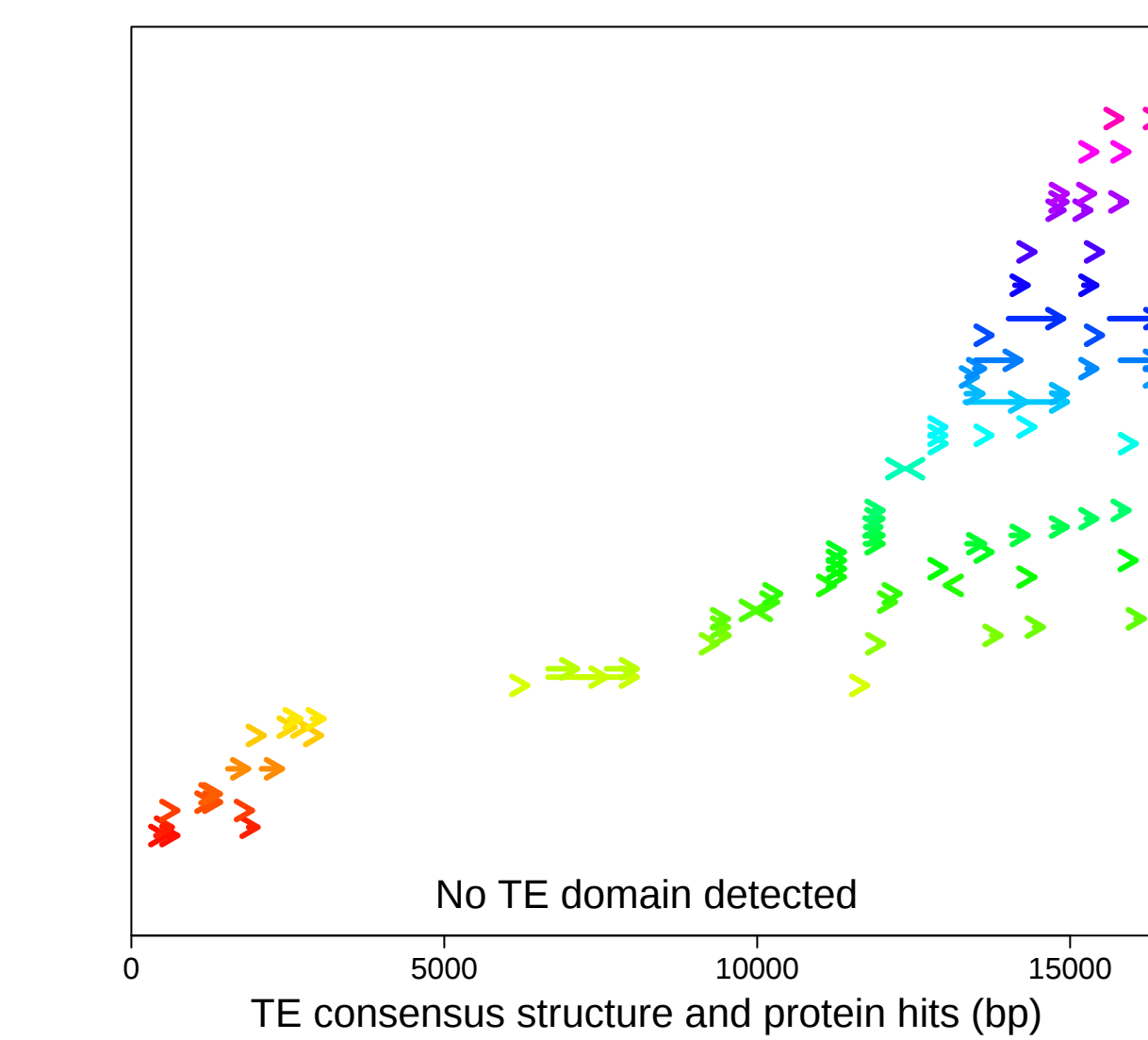
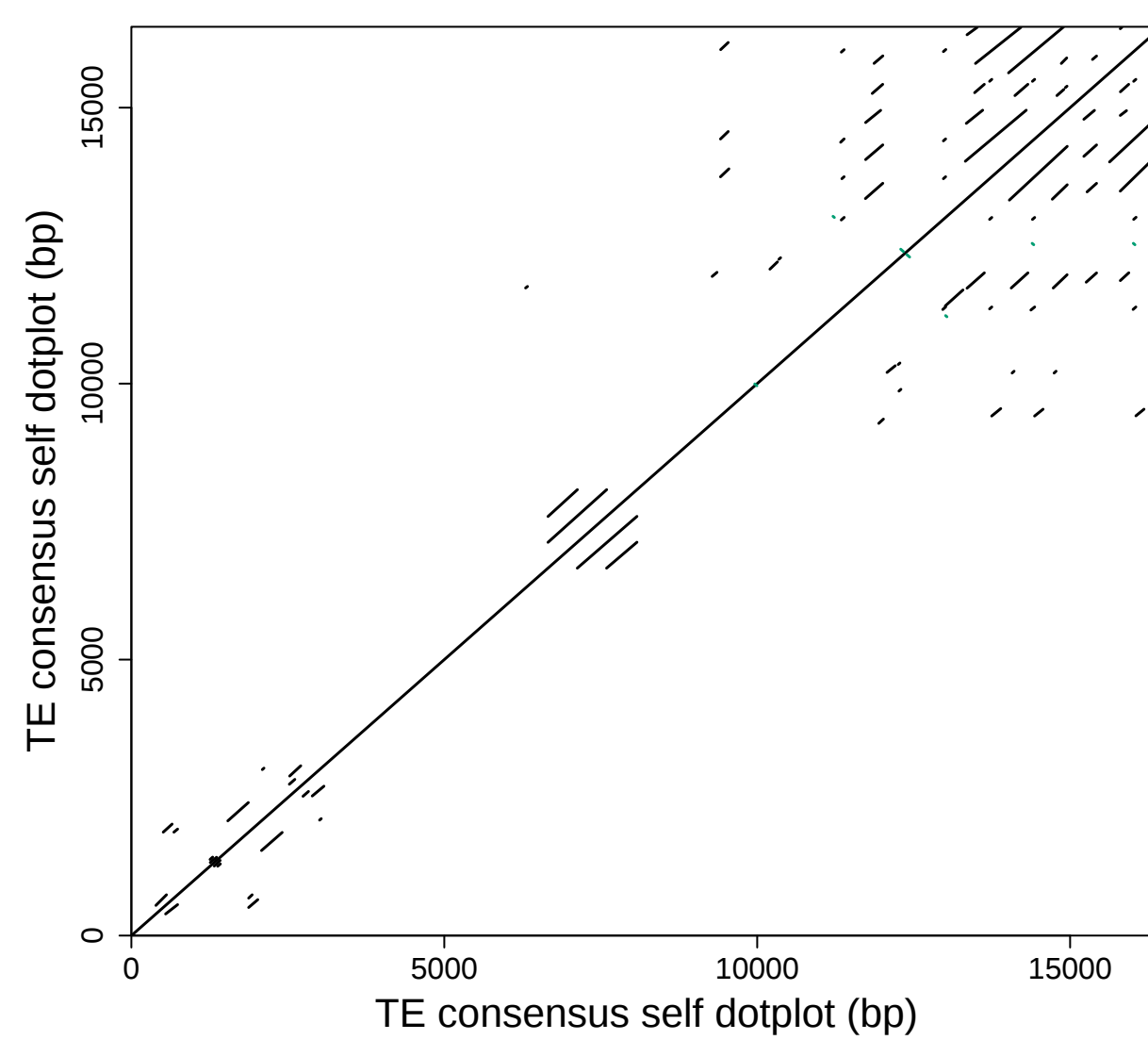
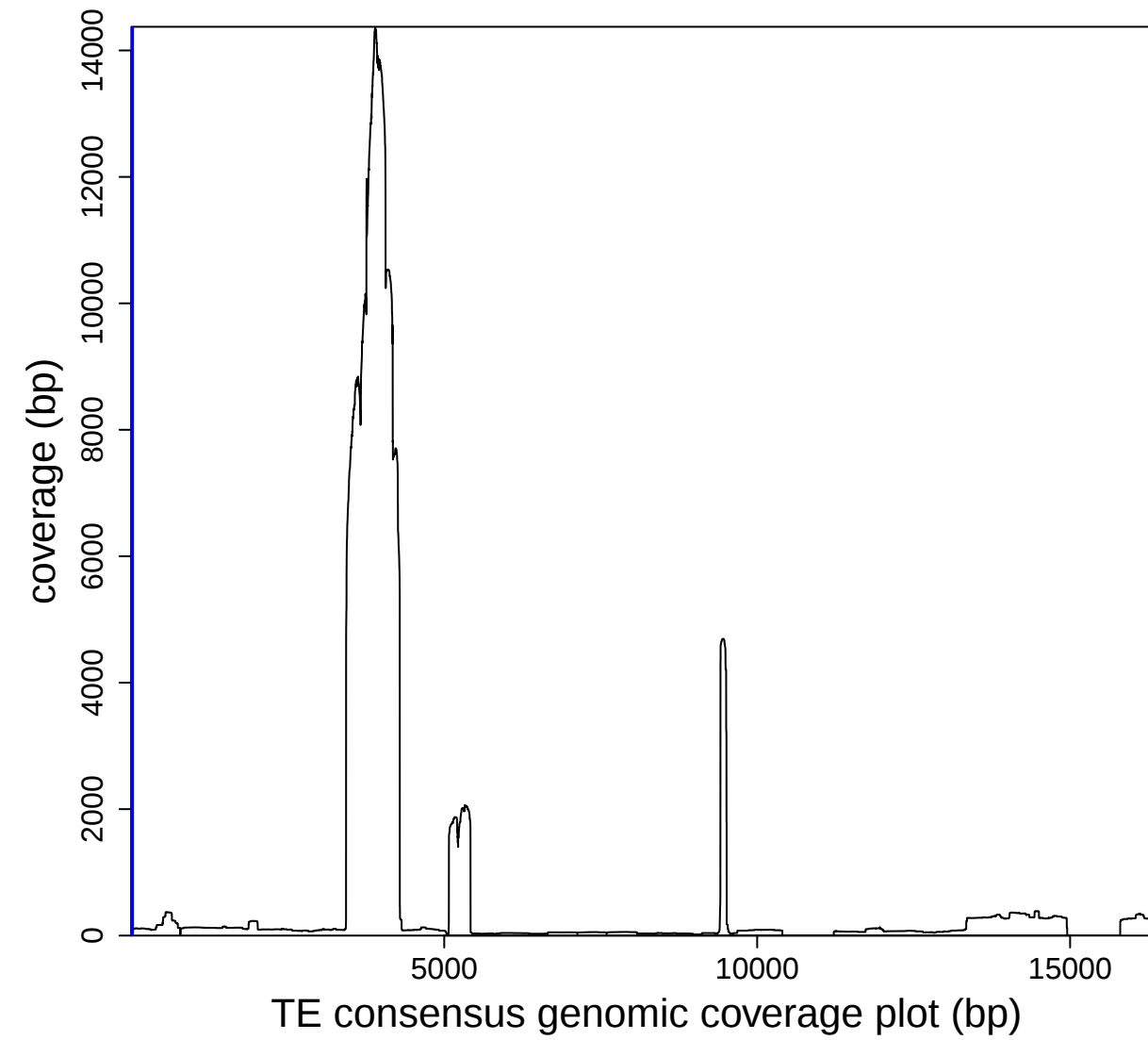
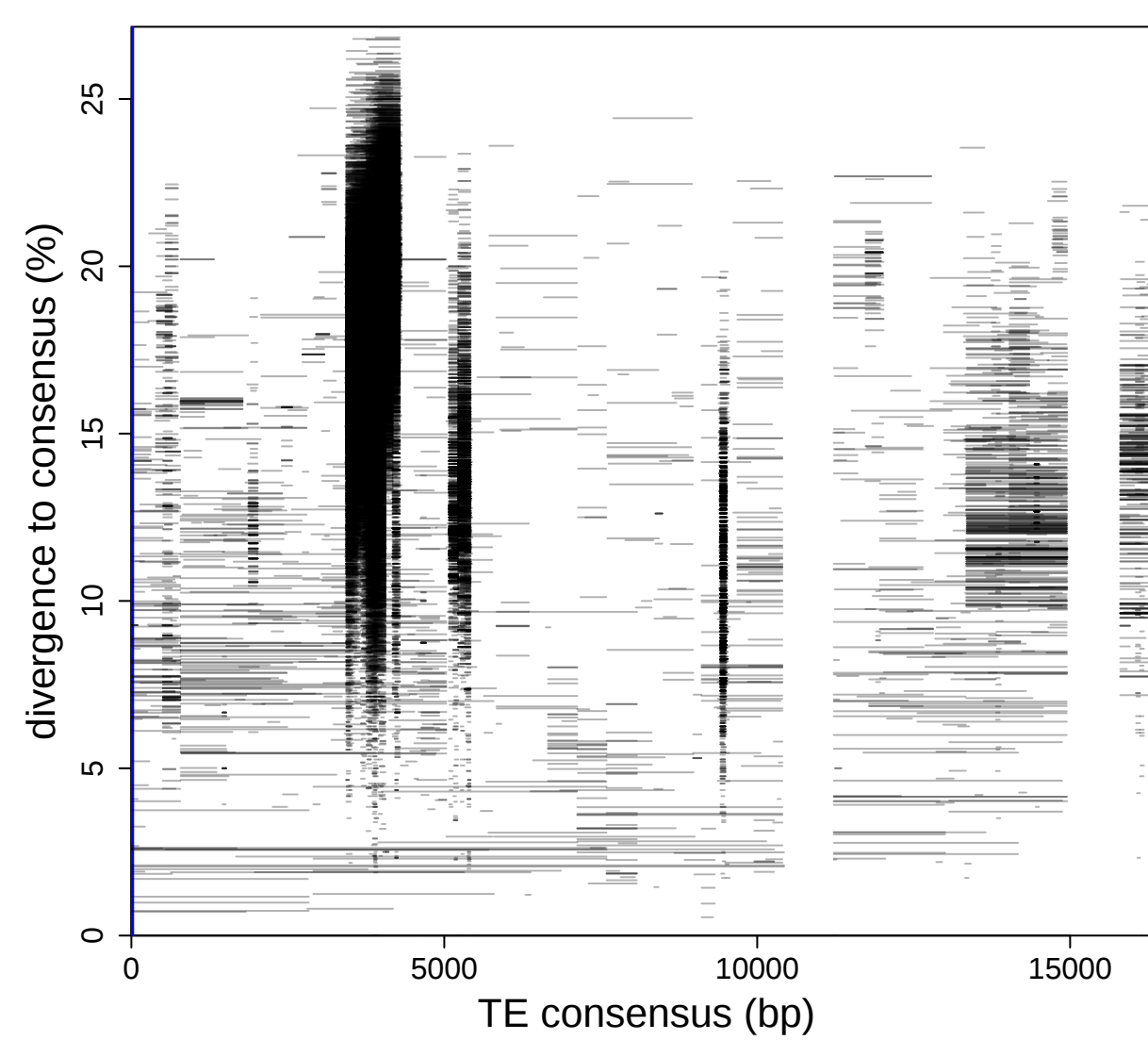
ta.b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.f
size: 16385bp; fragments: 33011; full length: 0 (>=14746.5bp)

After TEtrimmer 16385 bp



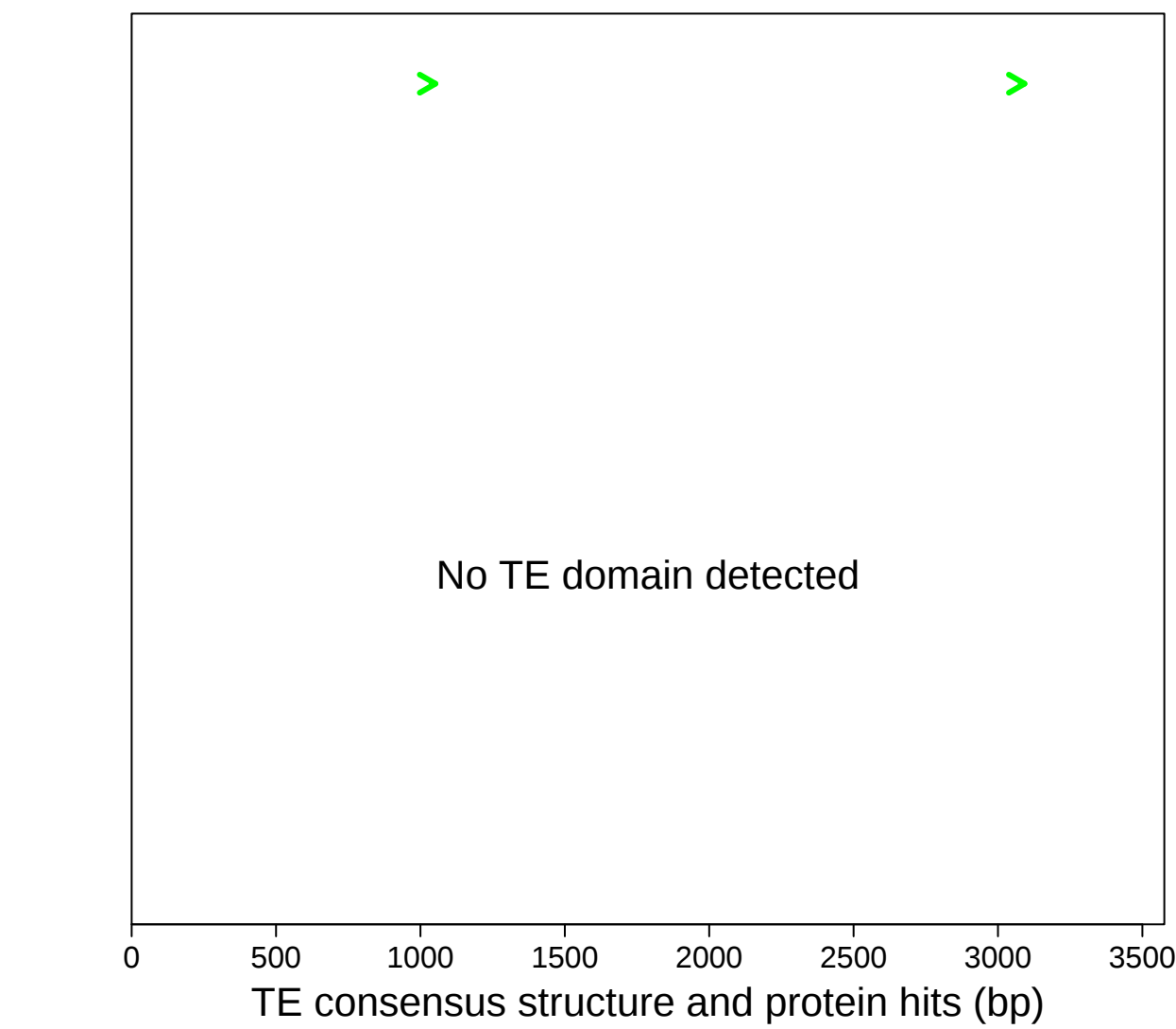
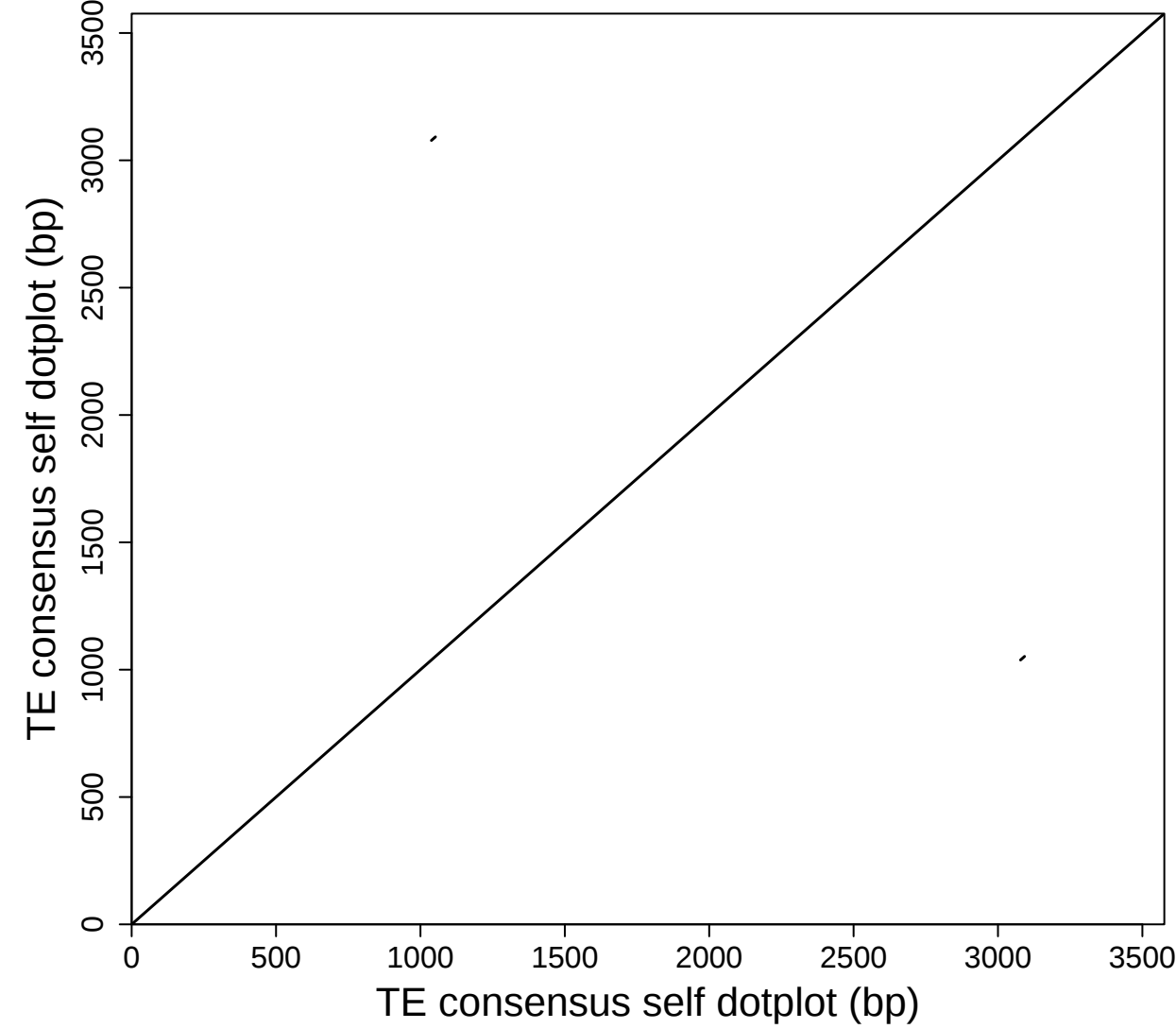
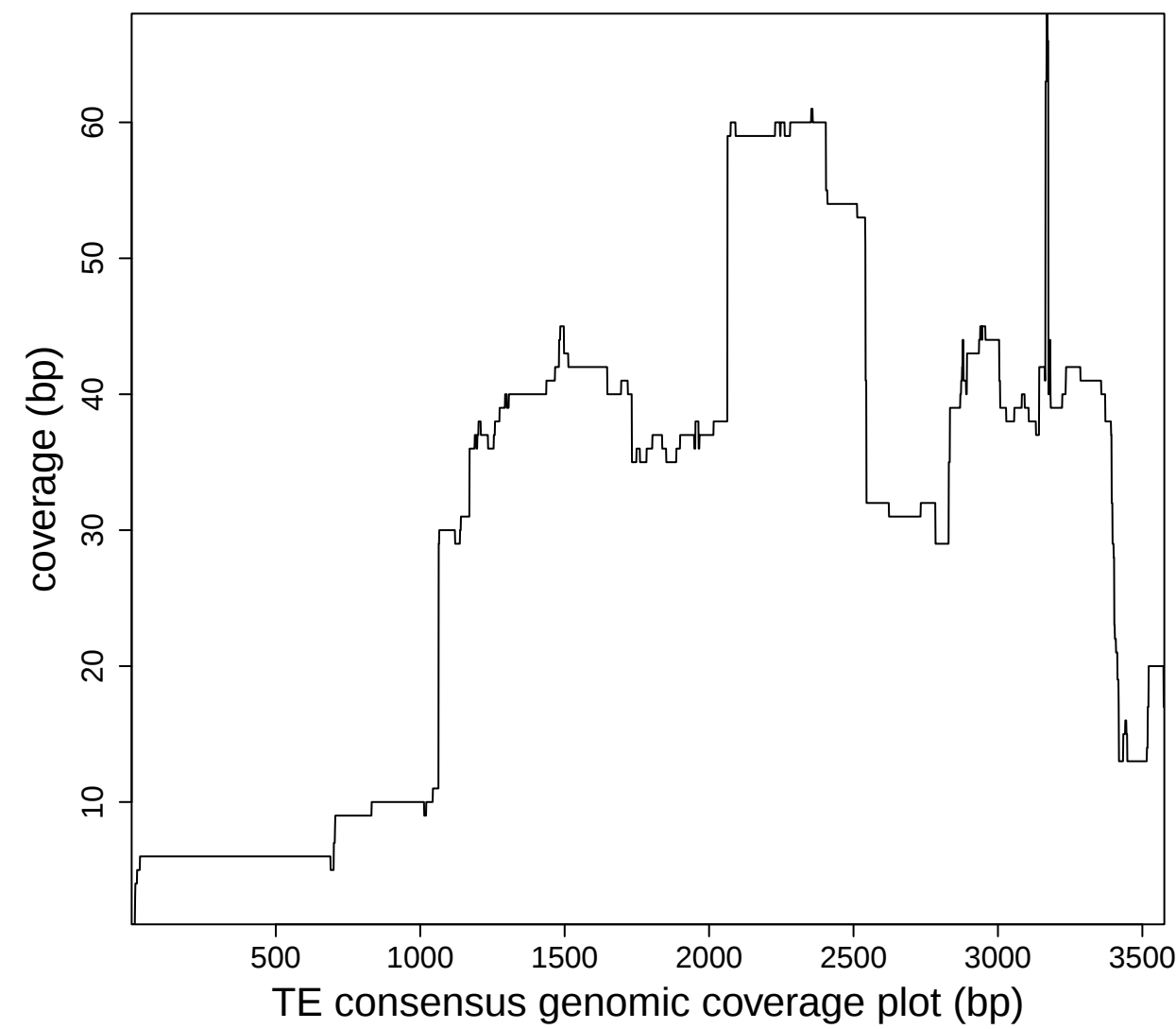
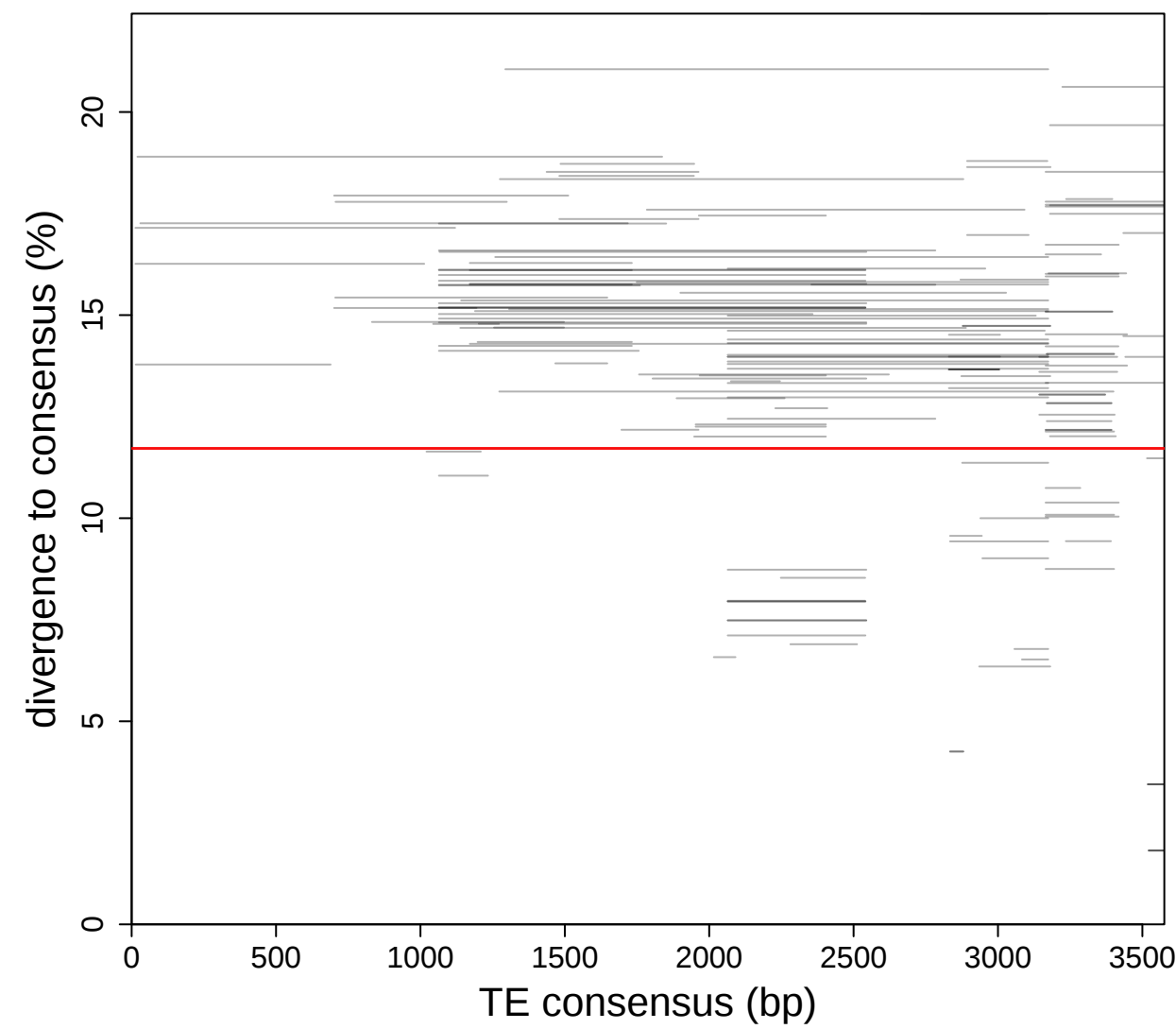
07.fasta.b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.f
size: 16465bp; fragments: 32992; full length: 0 (>=14818.5bp)

After TEtrimmer Extended plot Blue lines are boundaries

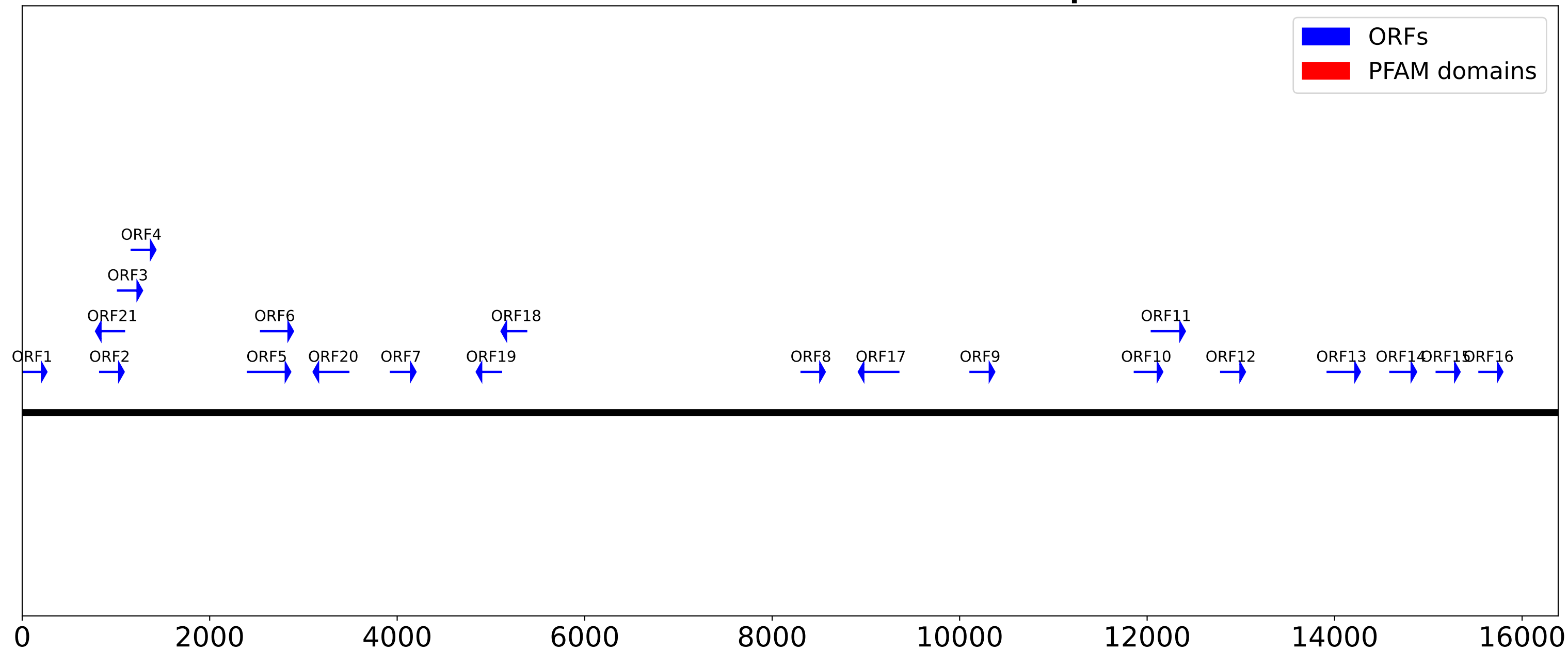


TE: rnd_5_family_13307
size: 3576bp; fragments: 166; full length: 1 (>=3218.4bp)

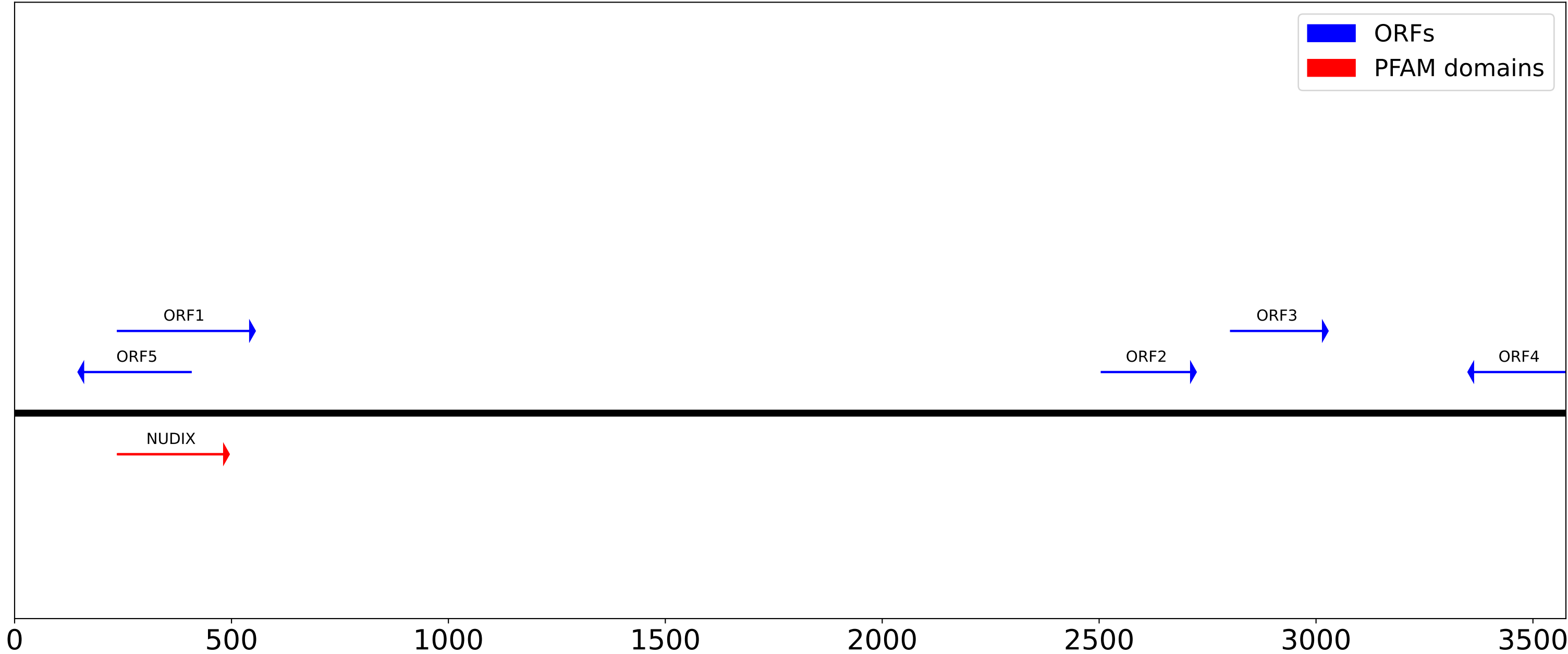
Before TEtrimmer 3576 bp



After TEtrimmer ORF and PFAM domain plot



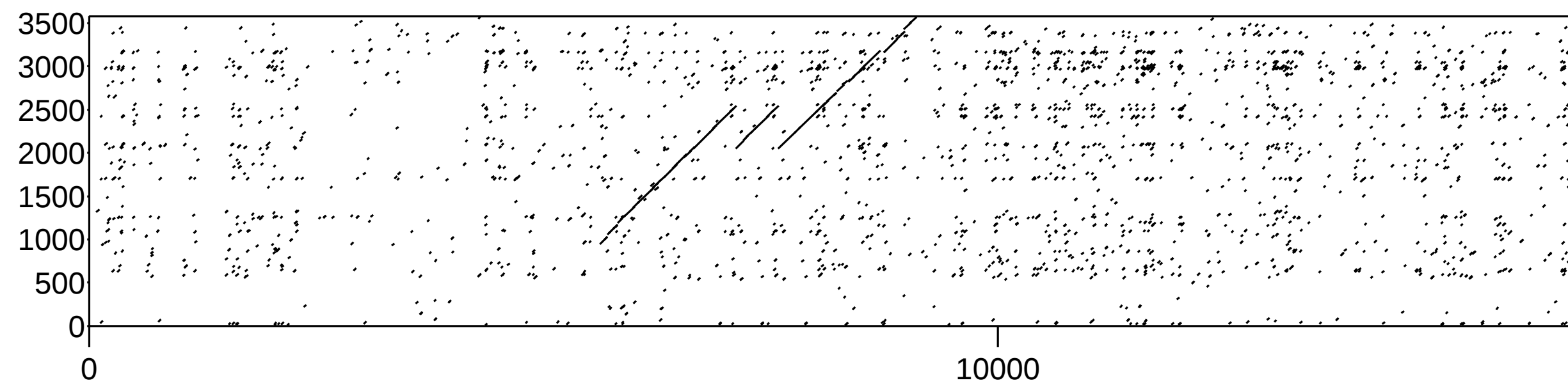
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

onsensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)