

Start crop Point

End crop Point

0 -

10 -

0

250

500

750

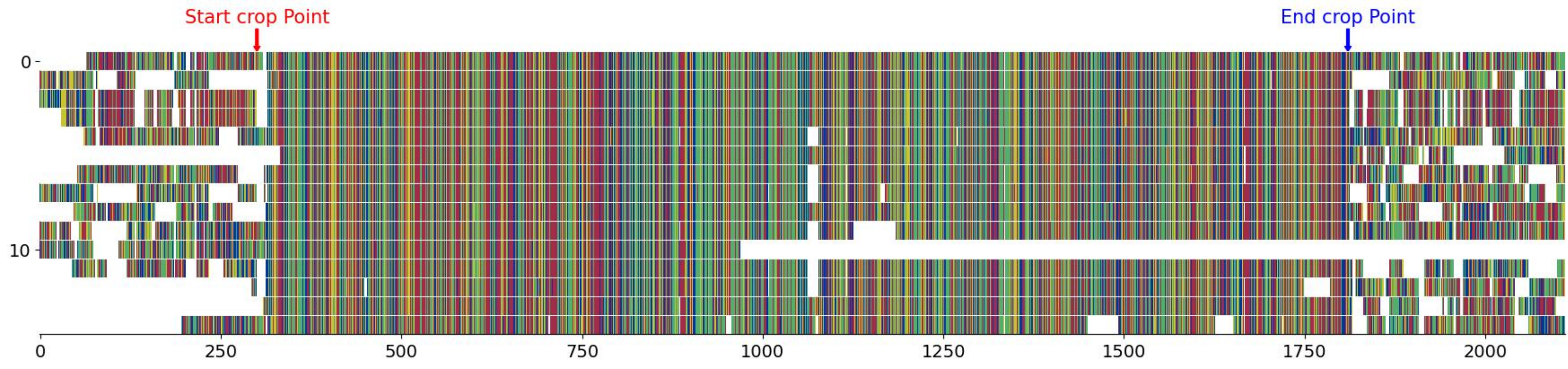
1000

1250

1500

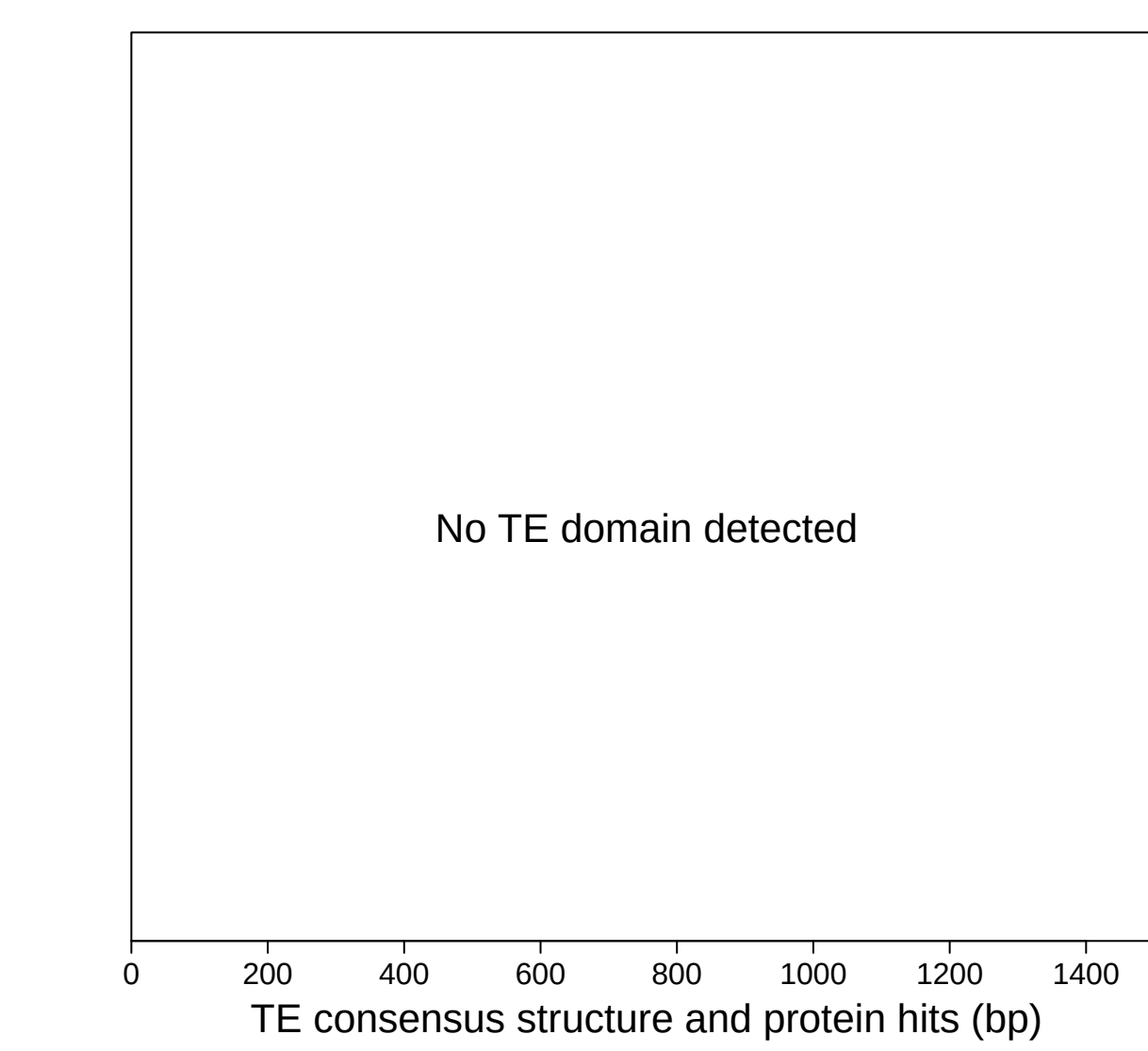
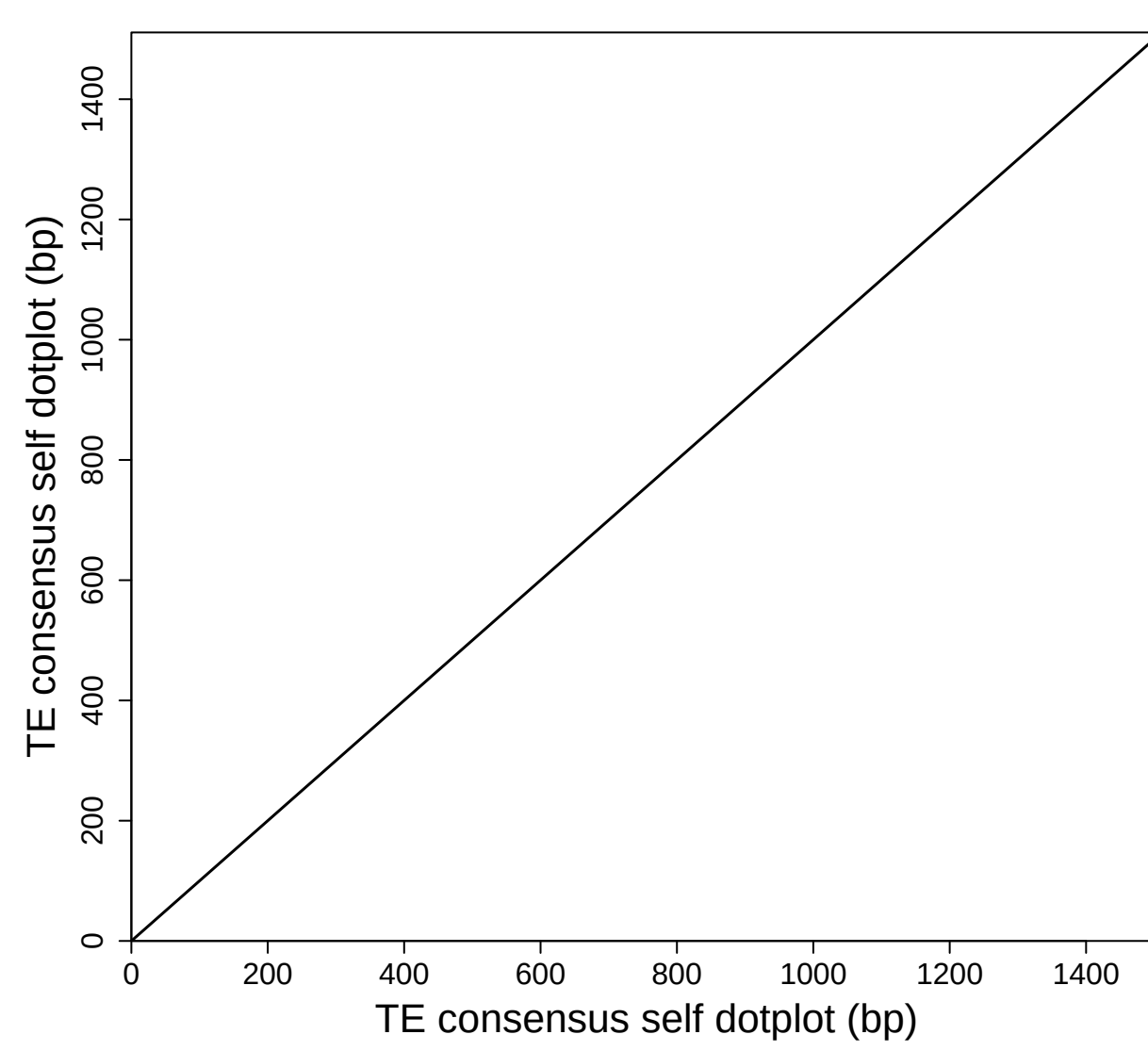
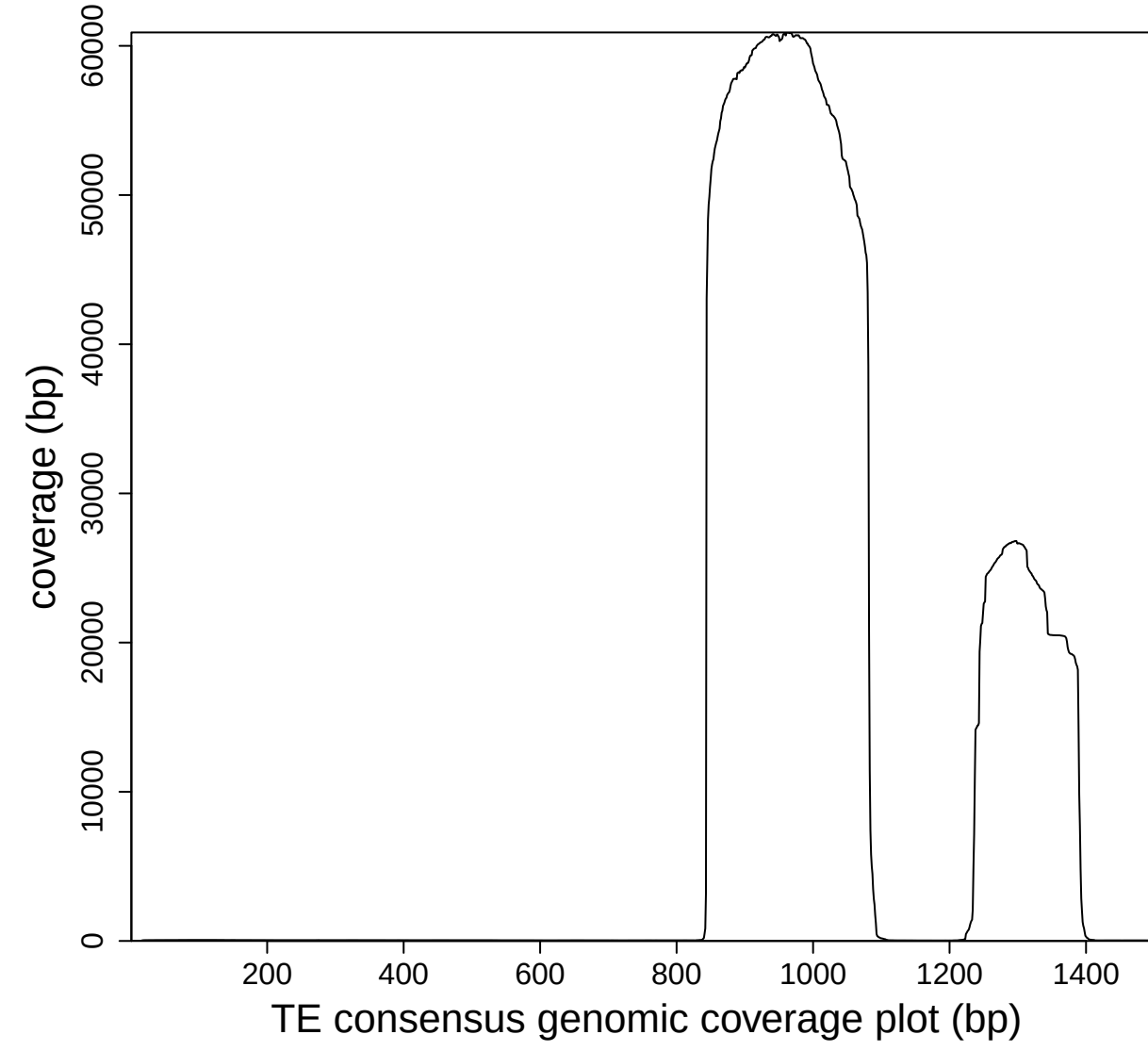
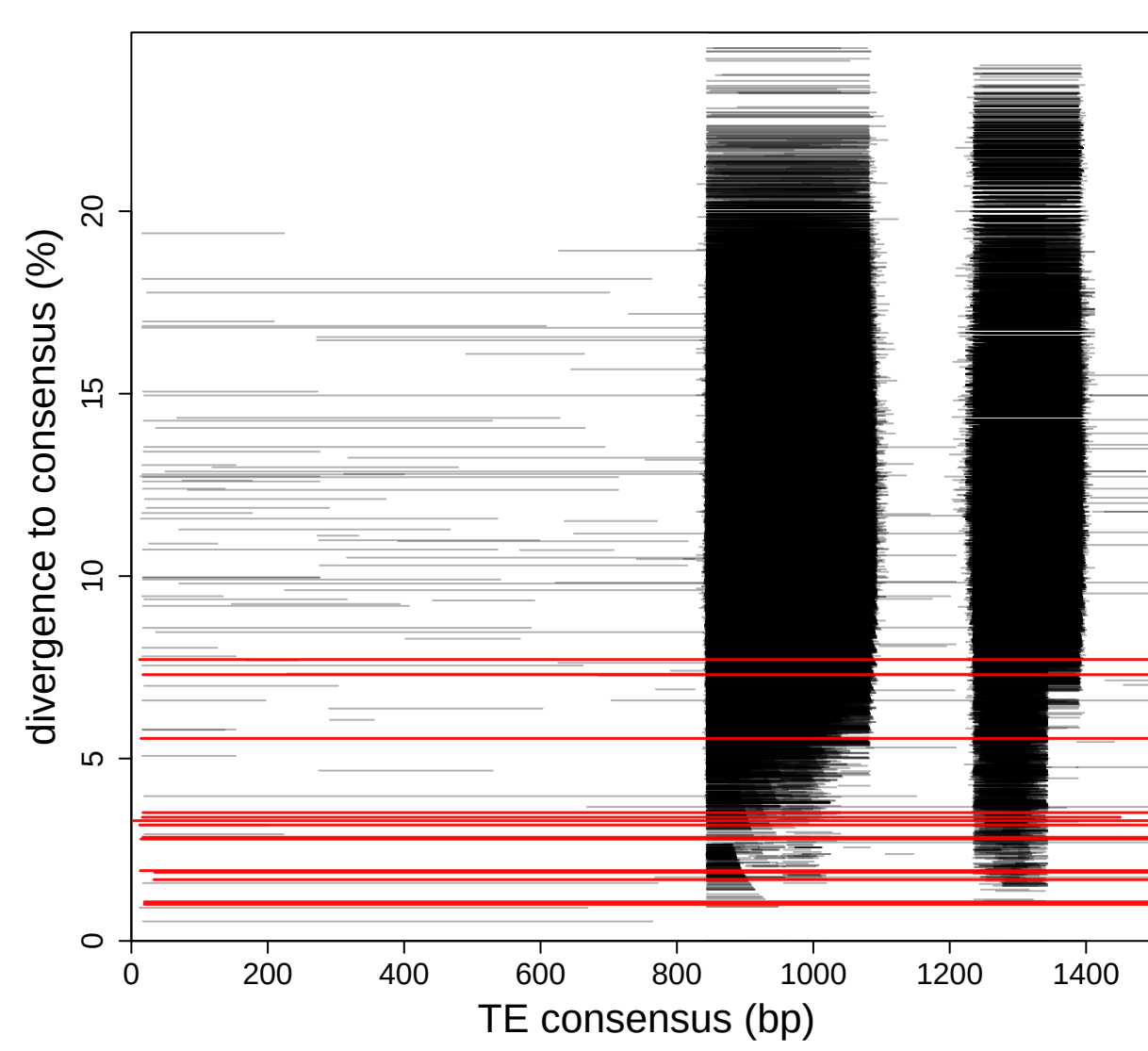
1750

2000



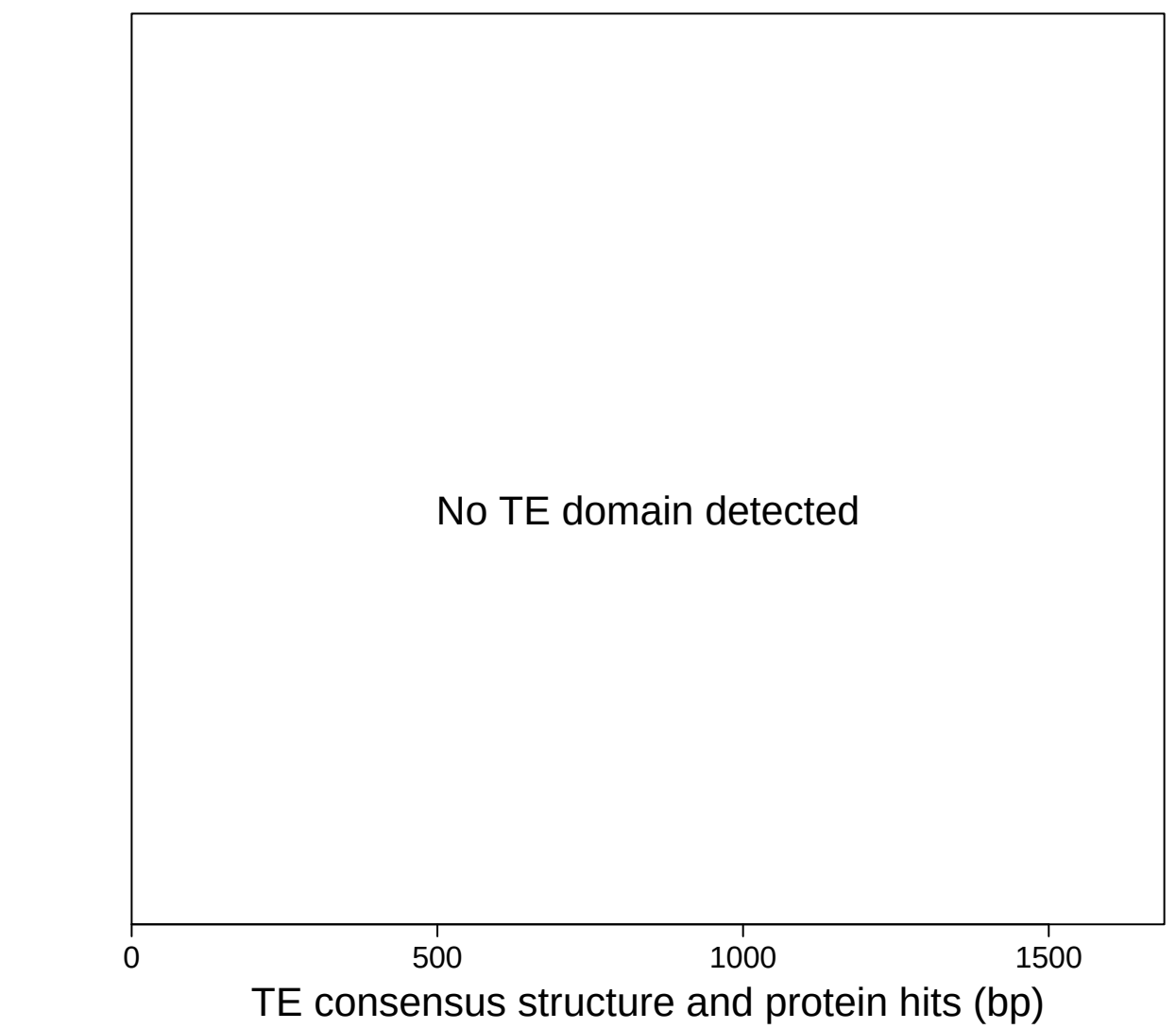
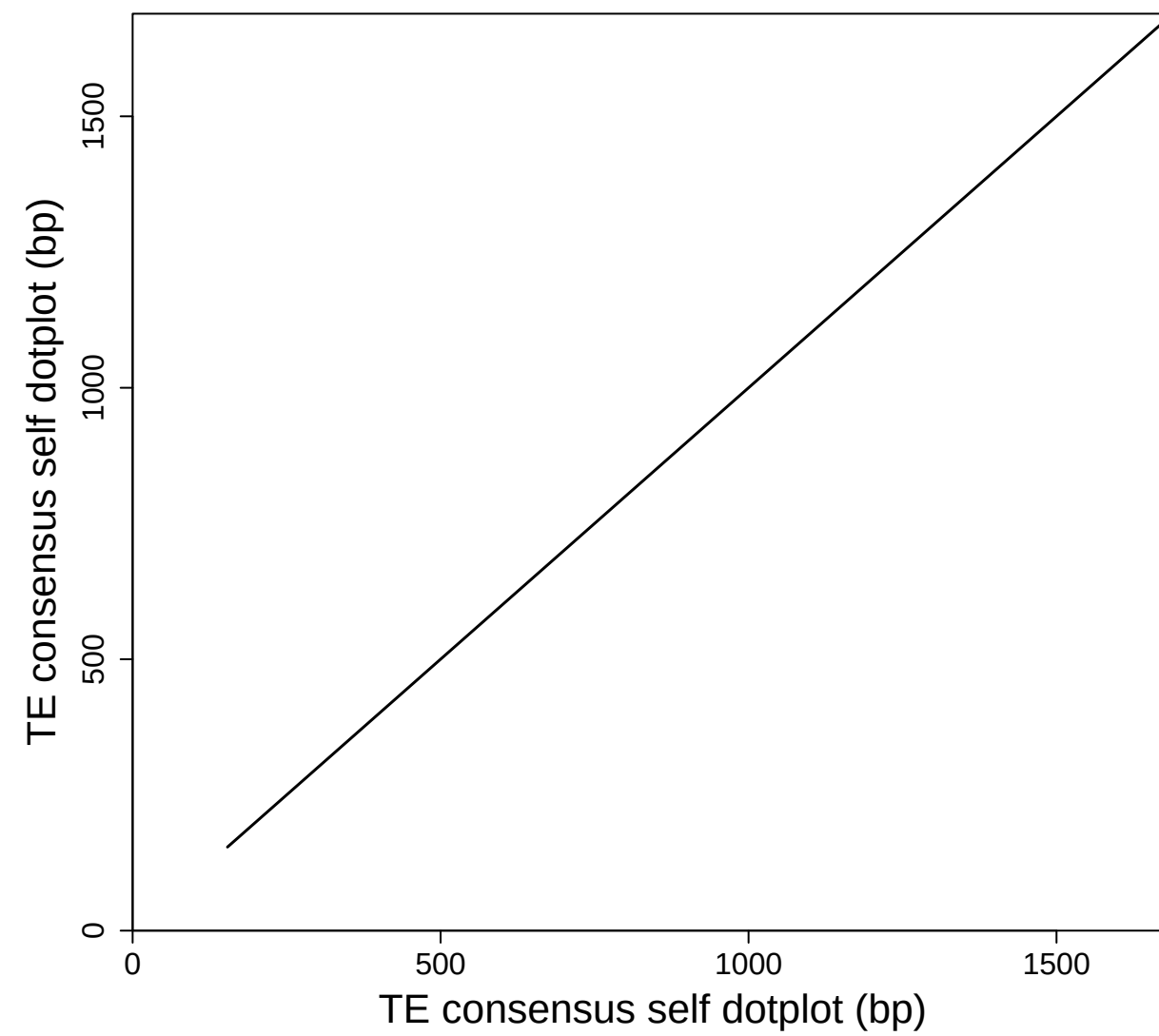
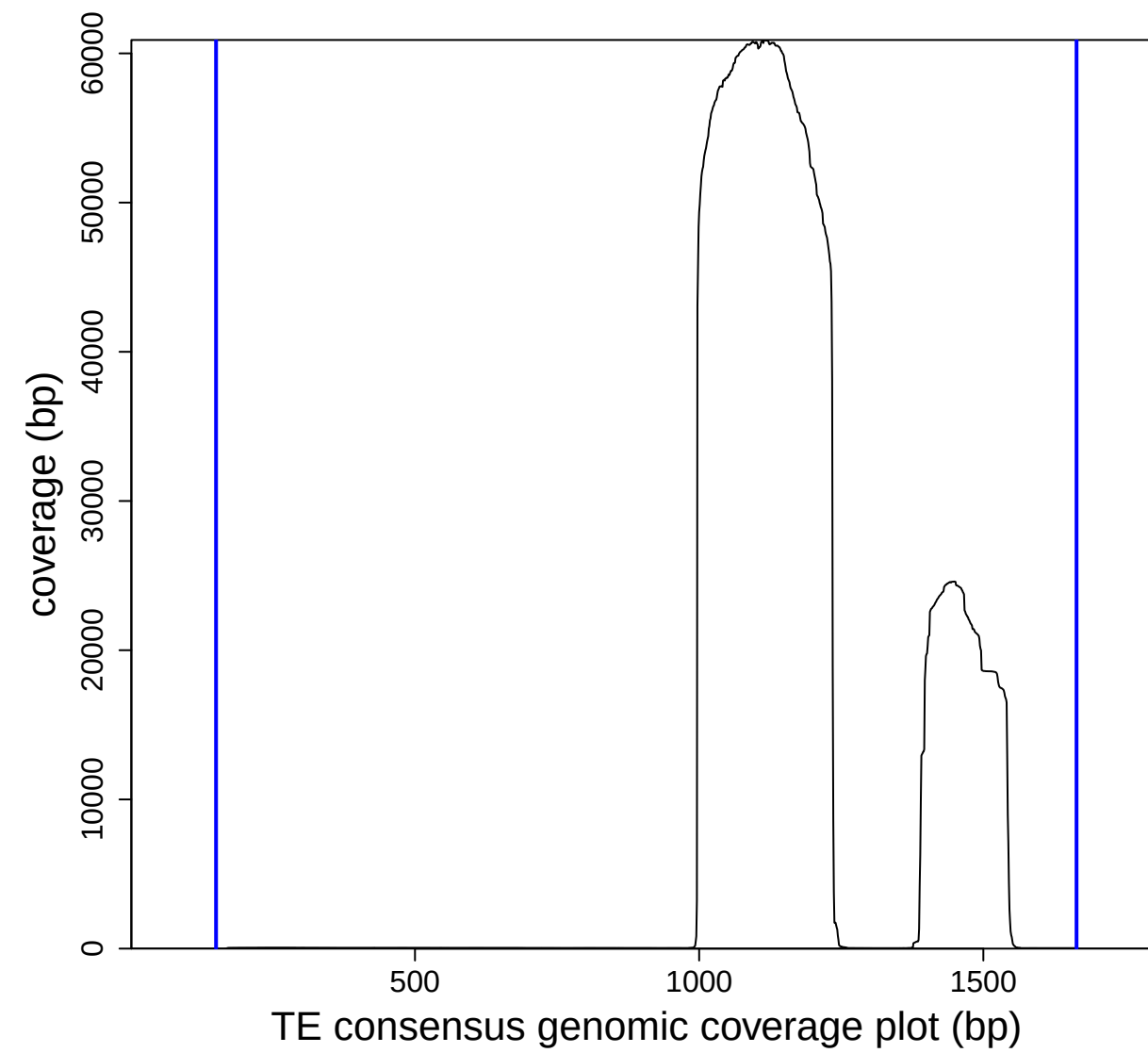
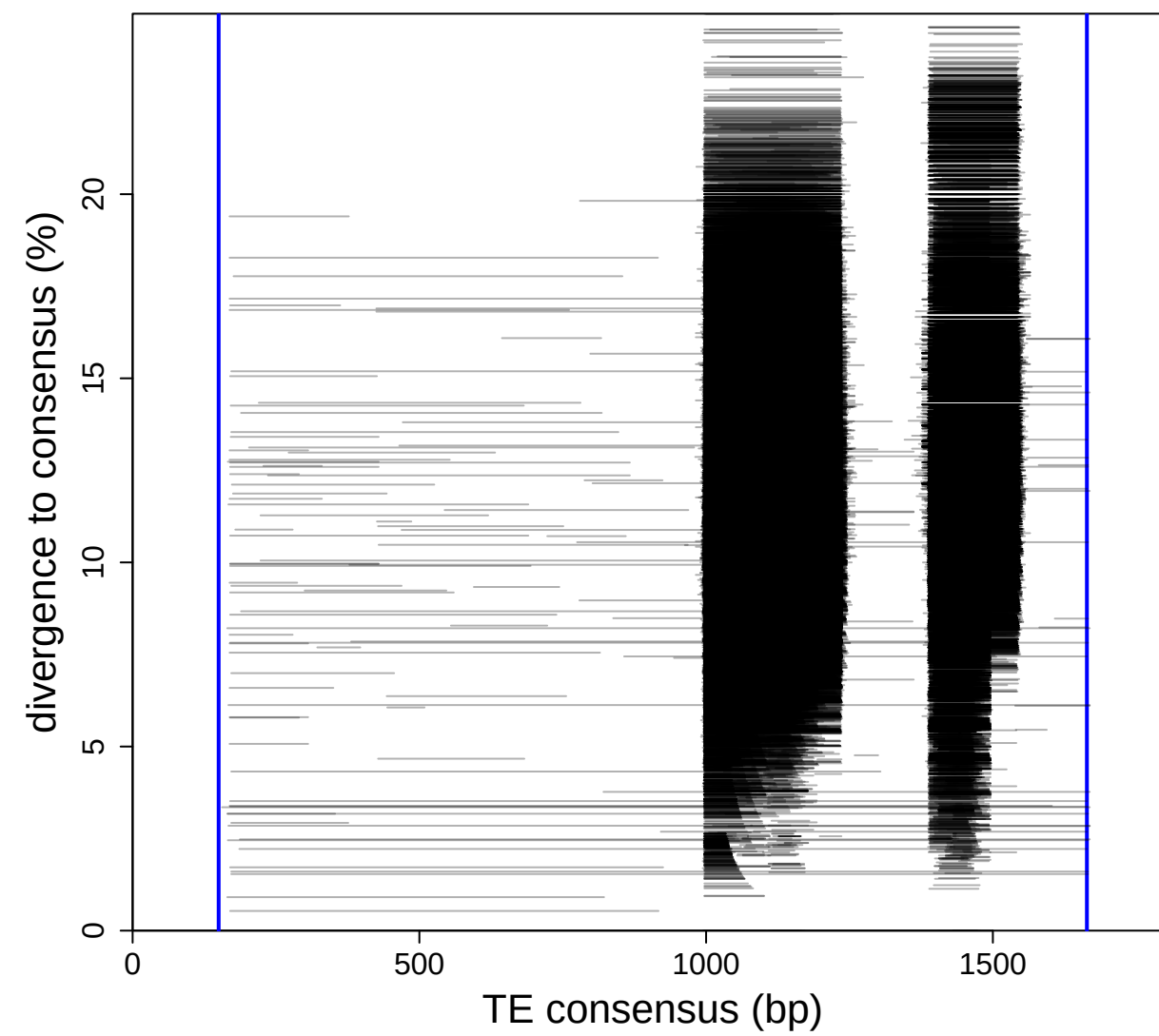
a.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.f
size: 1511bp; fragments: 92599; full length: 14 (>=1359.9bp)

After TEtrimmer 1511 bp

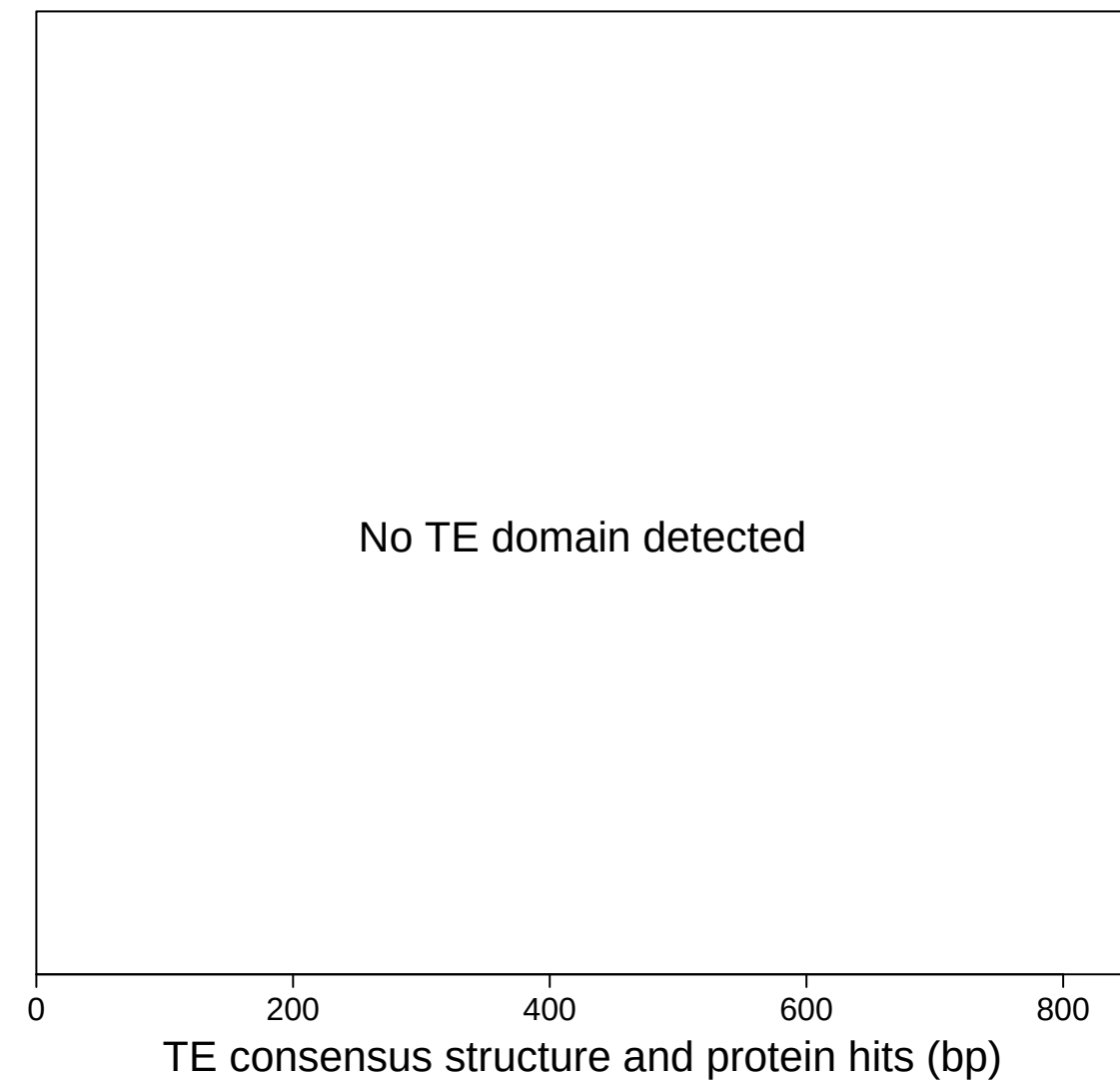
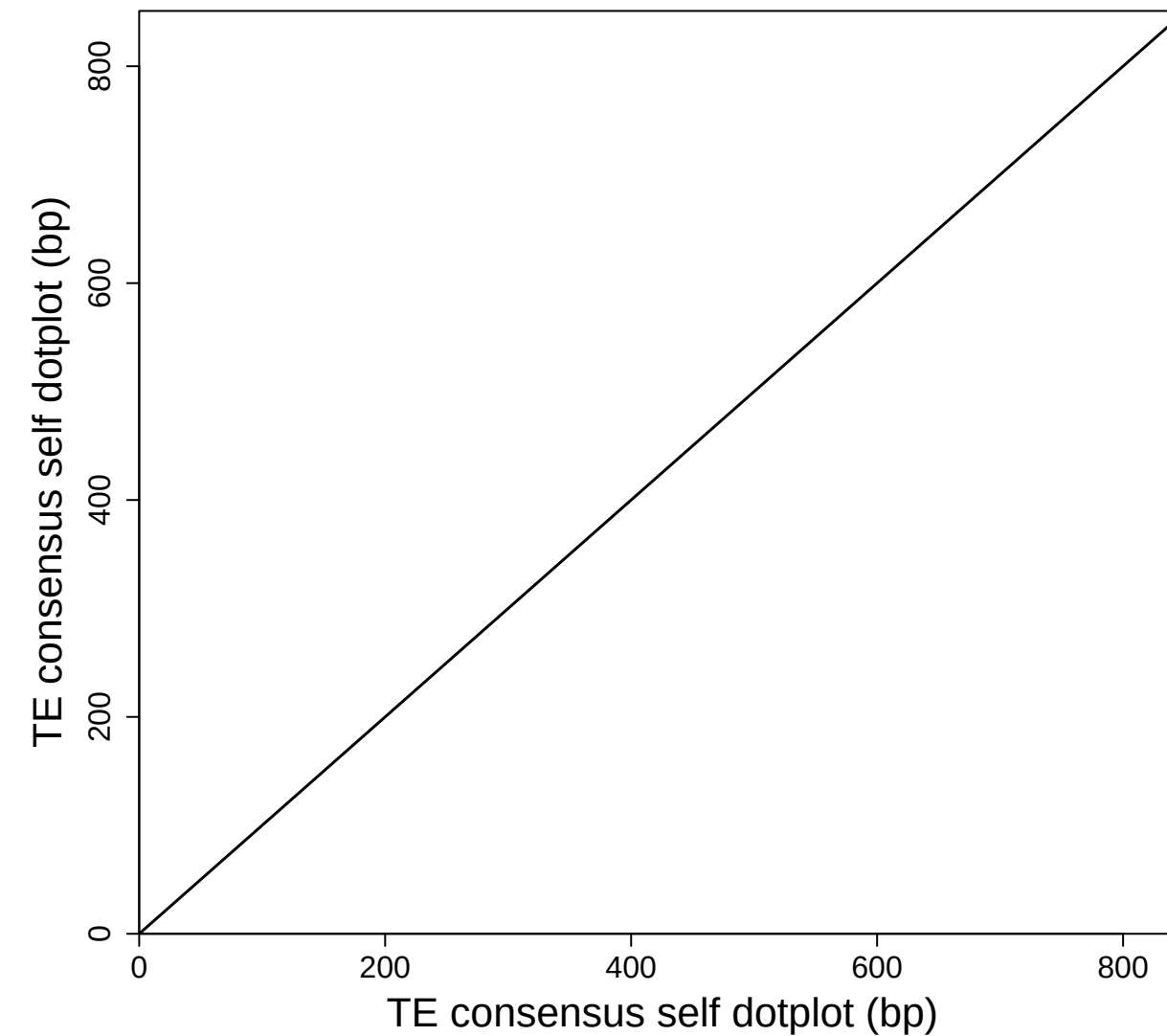
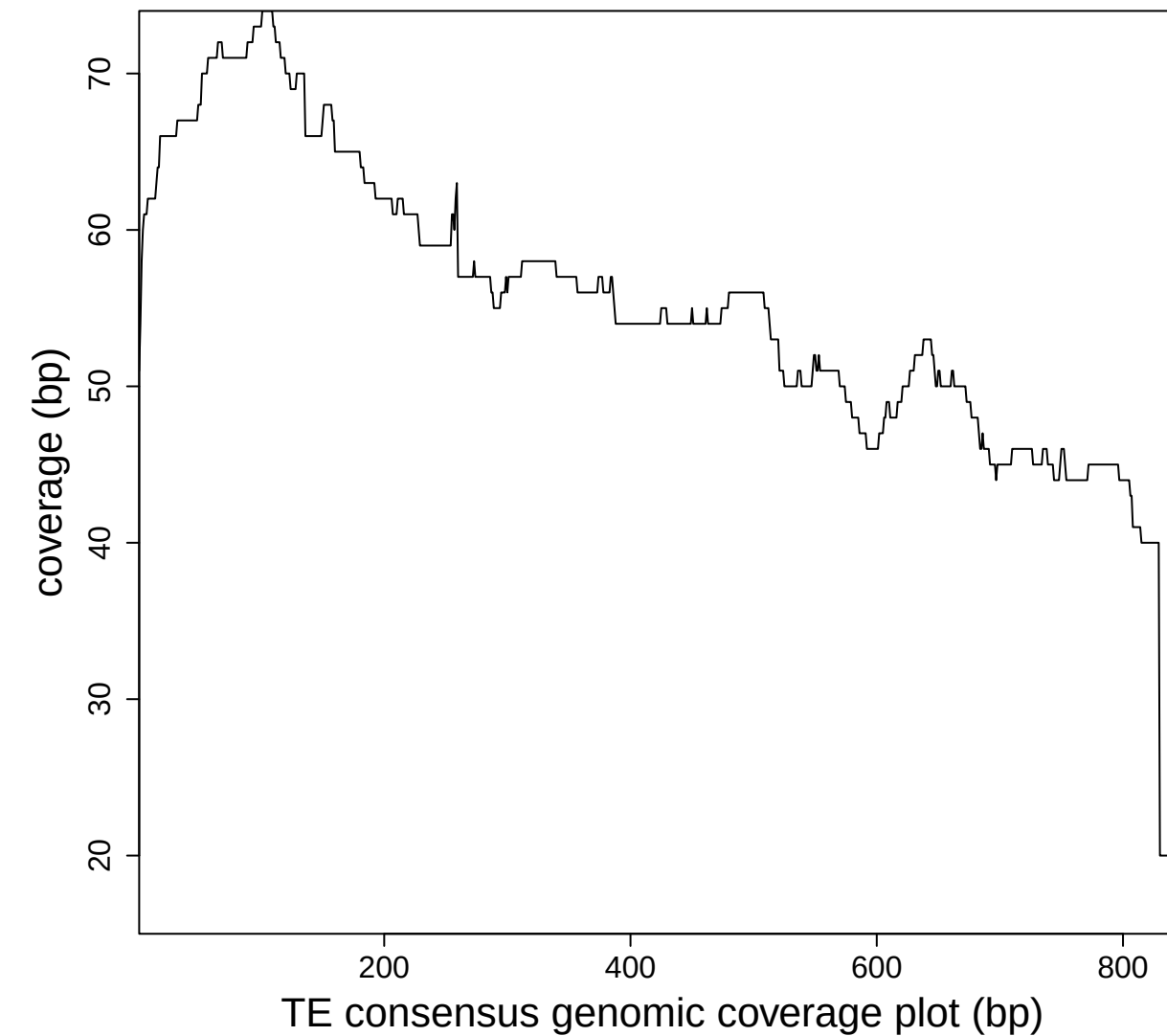
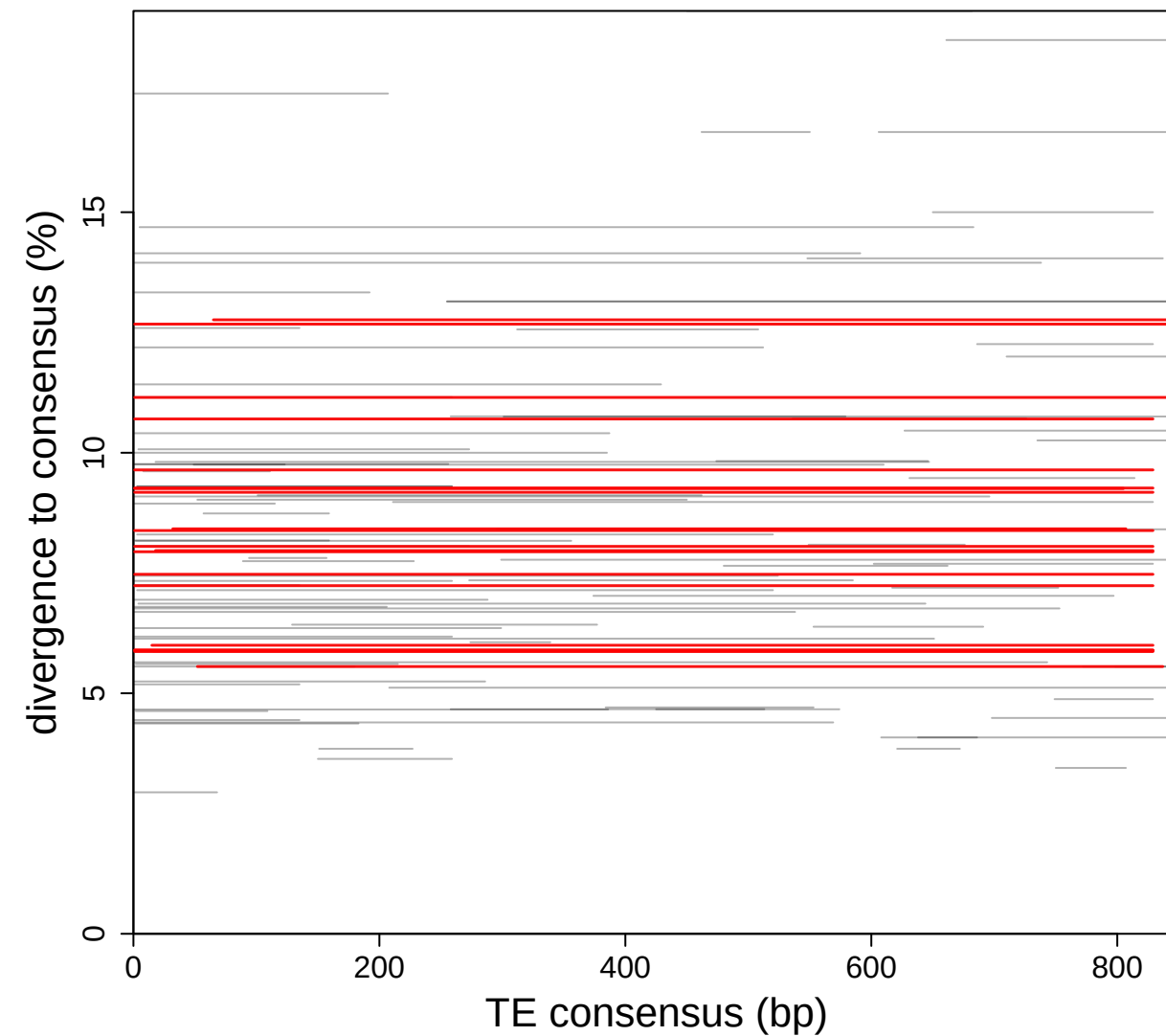


5.fasta.b.bed uf.bed g_1.bed fm_1.bed 0_0_bcln.fa aln.fa cl.f
size: 1814bp; fragments: 90292; full length: 0 (>=1632.6bp)

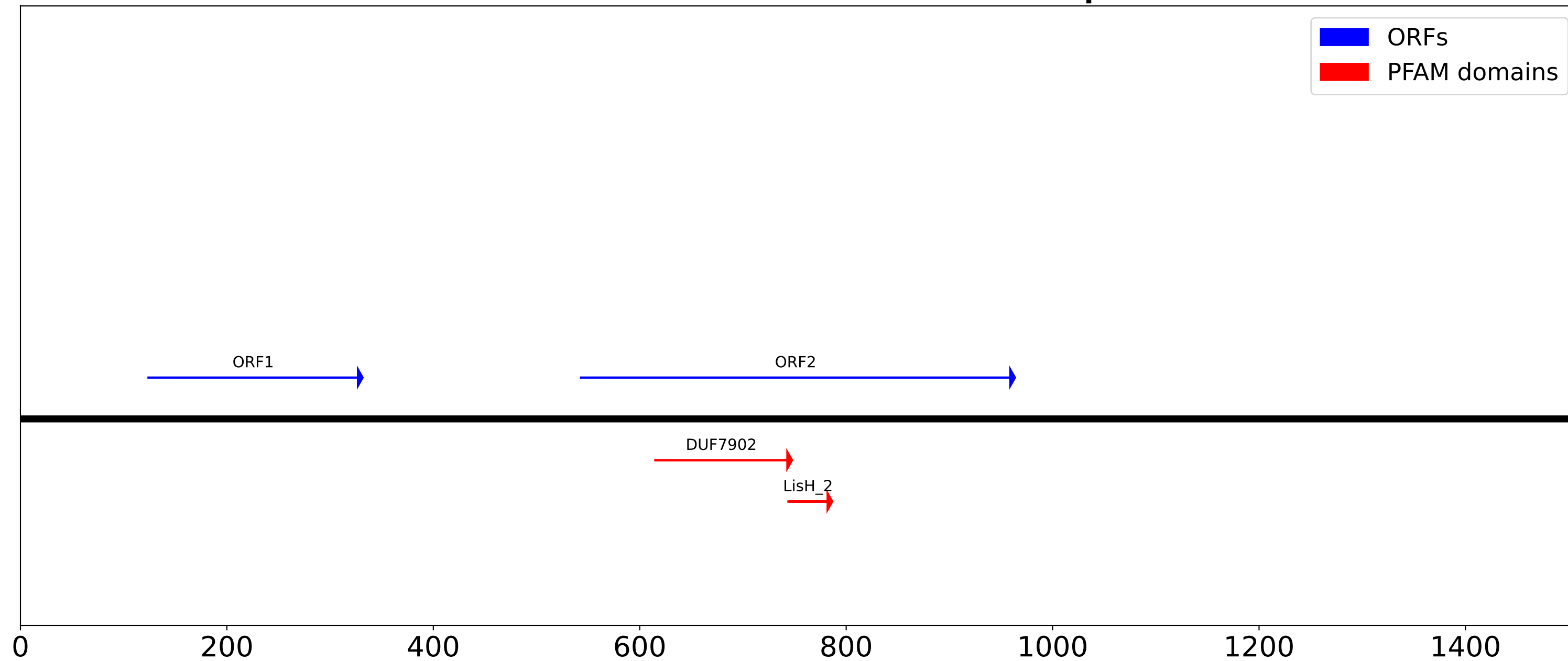
After TEtrimmer Extended plot Blue lines are boundaries



TE: rnd_5_family_9515
size: 851bp; fragments: 120; full length: 21 (≥ 765.9 bp)



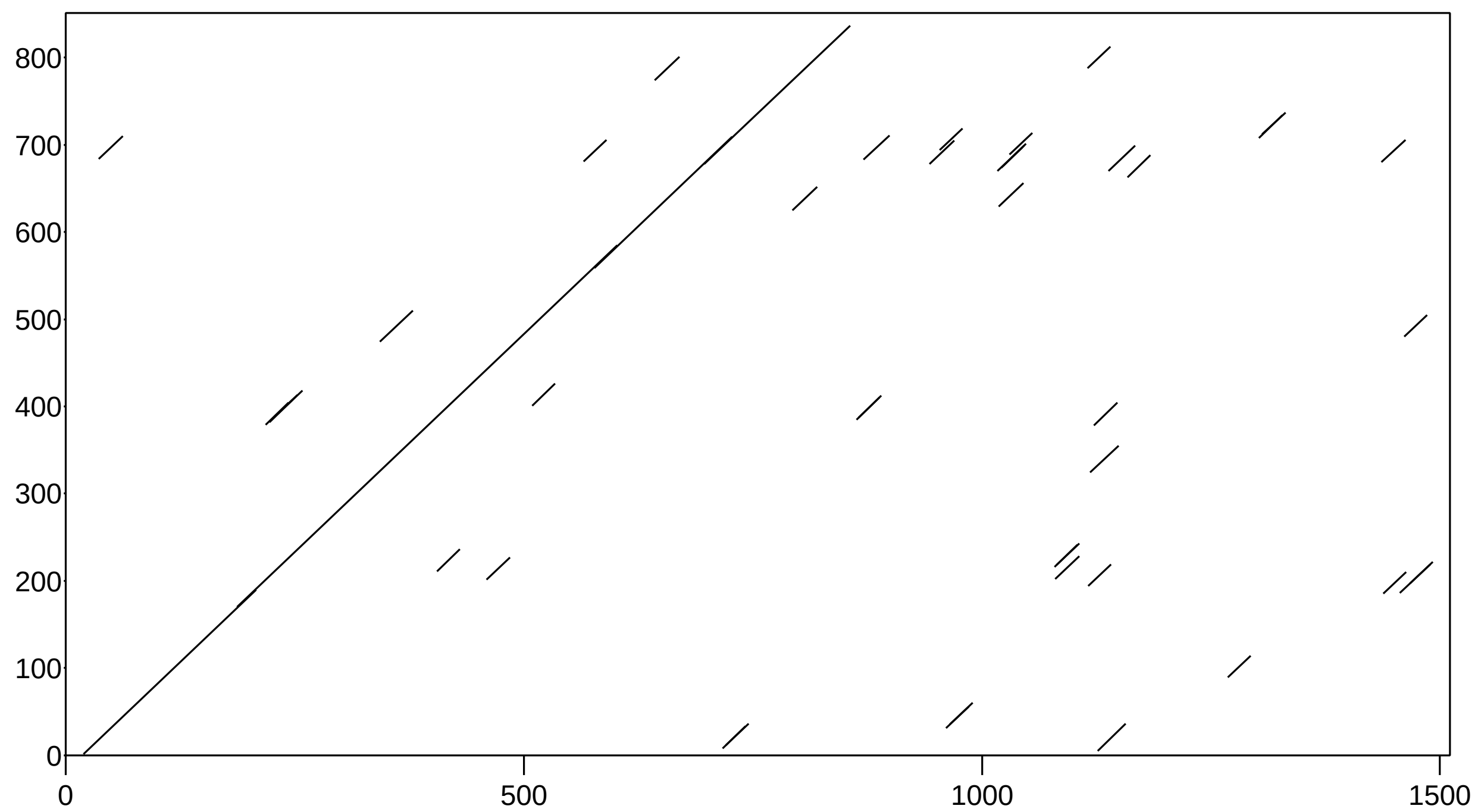
After TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)