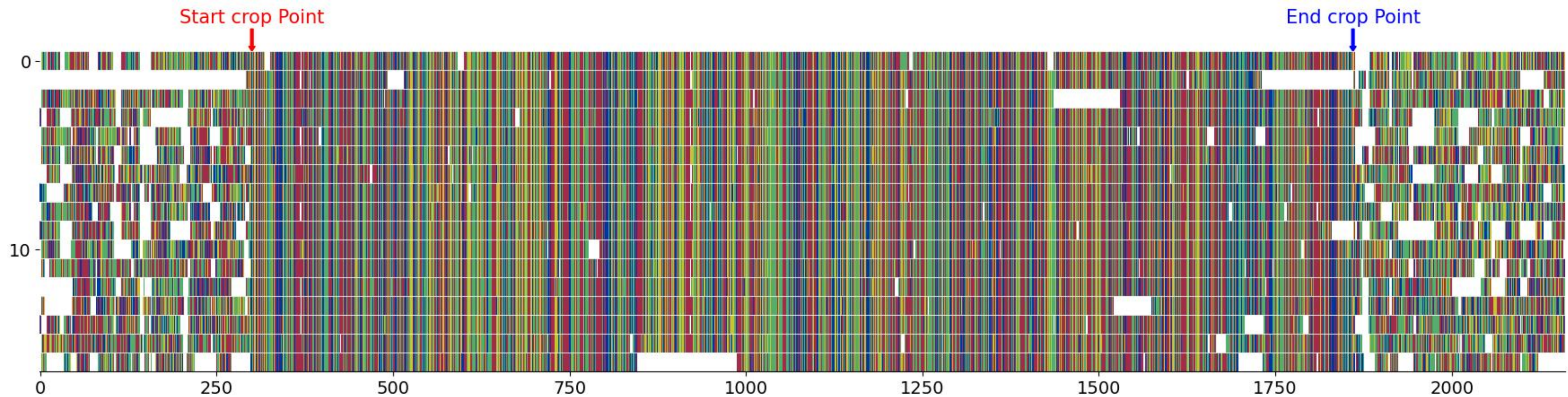


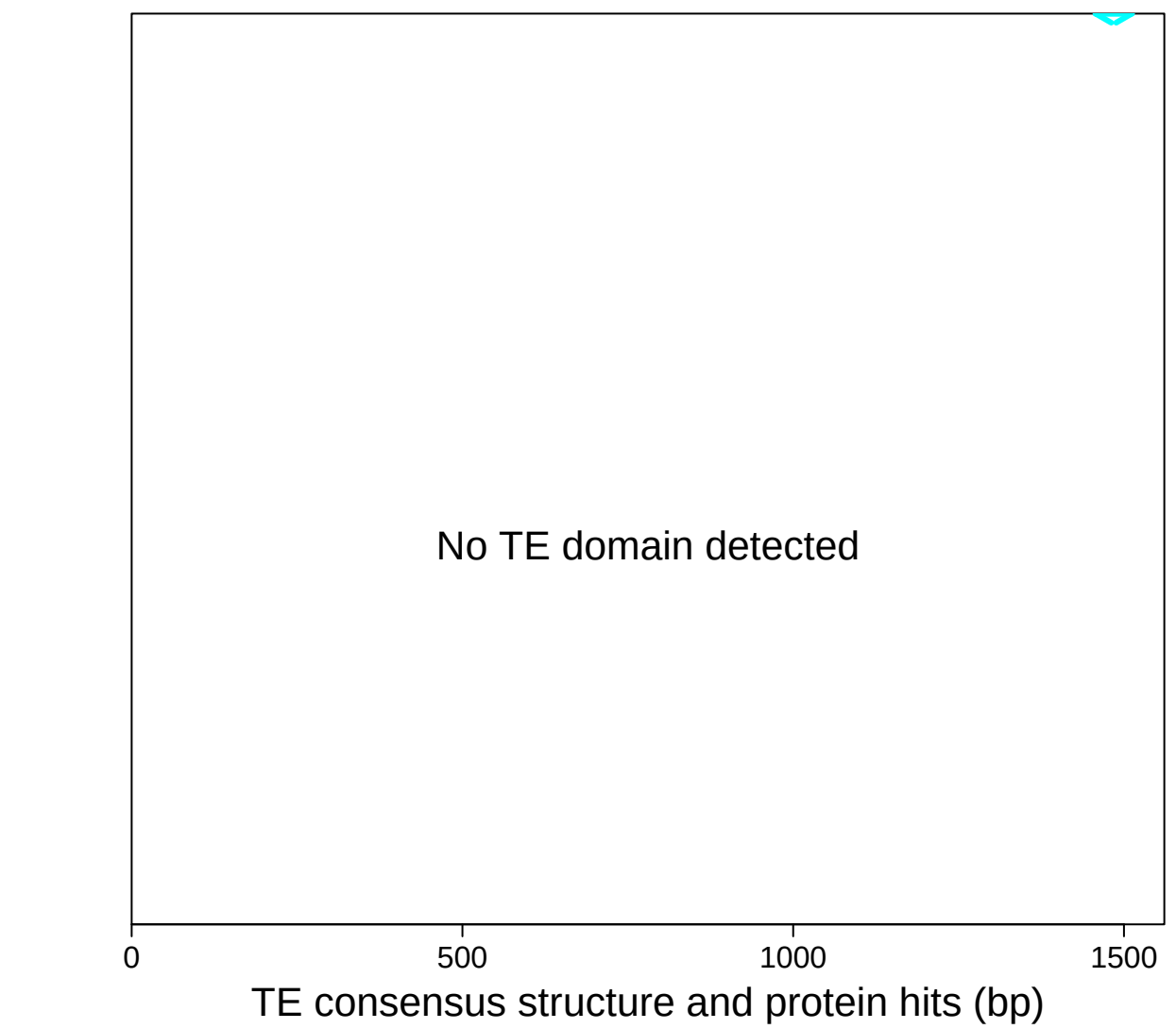
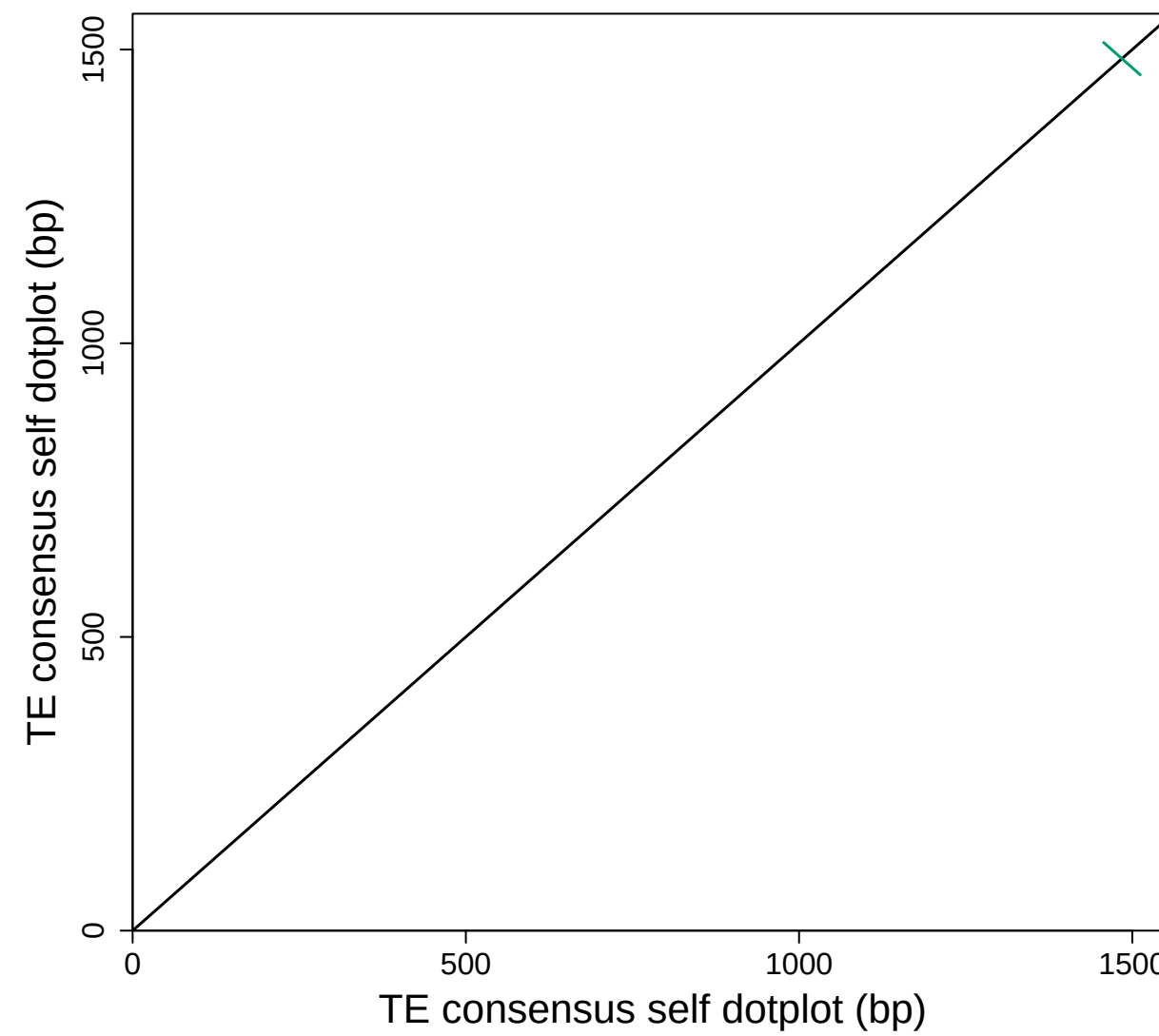
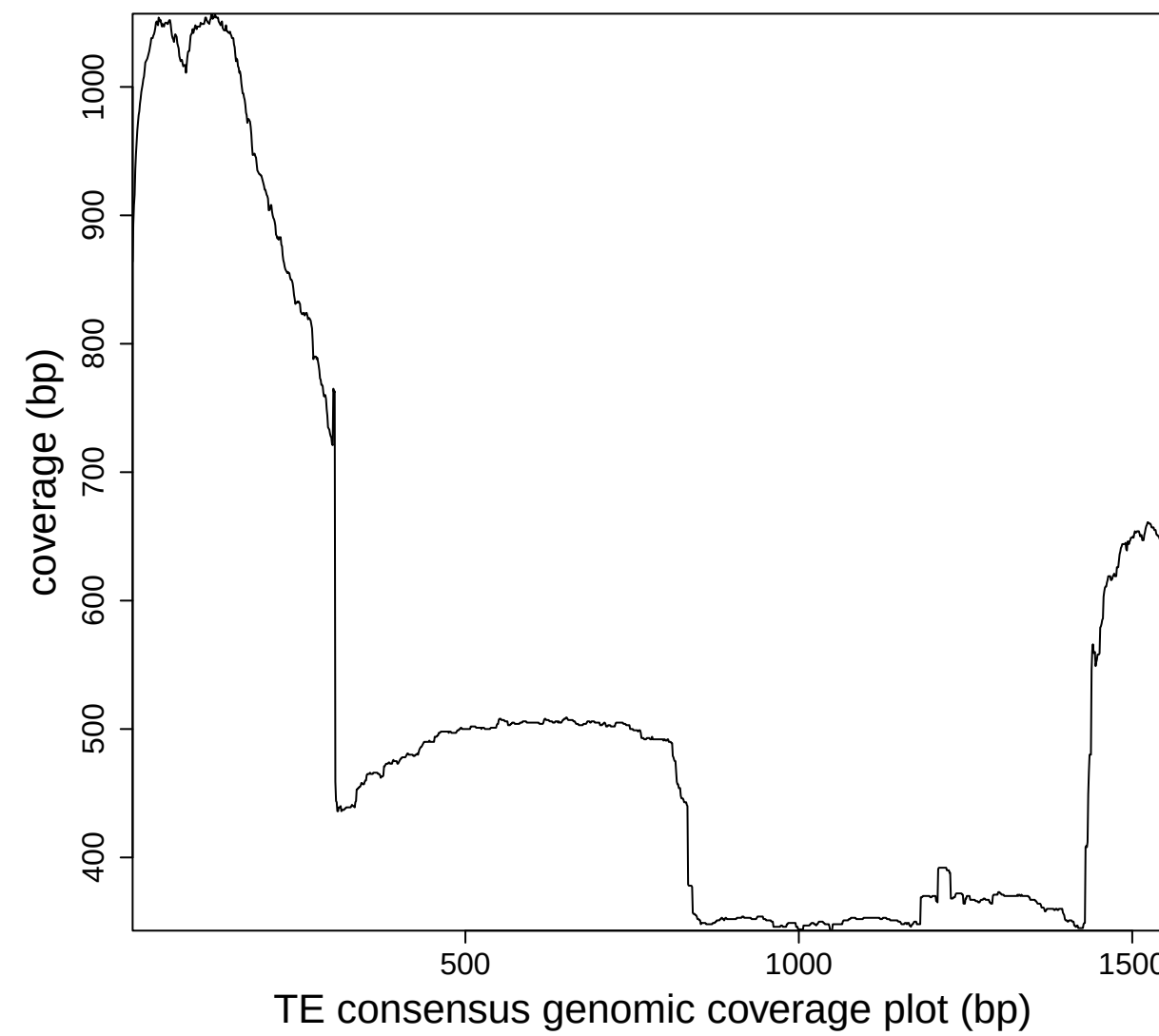
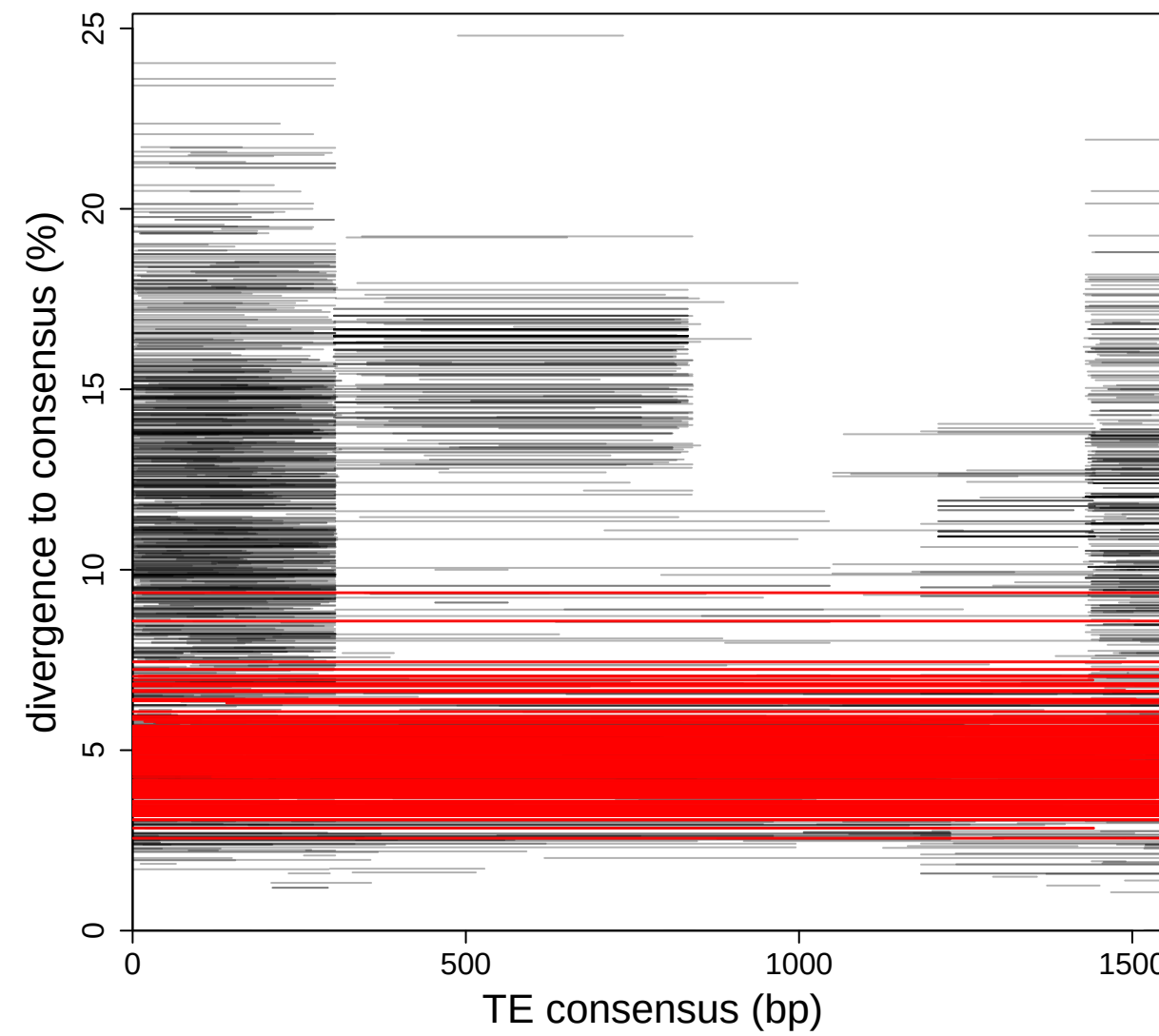
Start crop Point

End crop Point



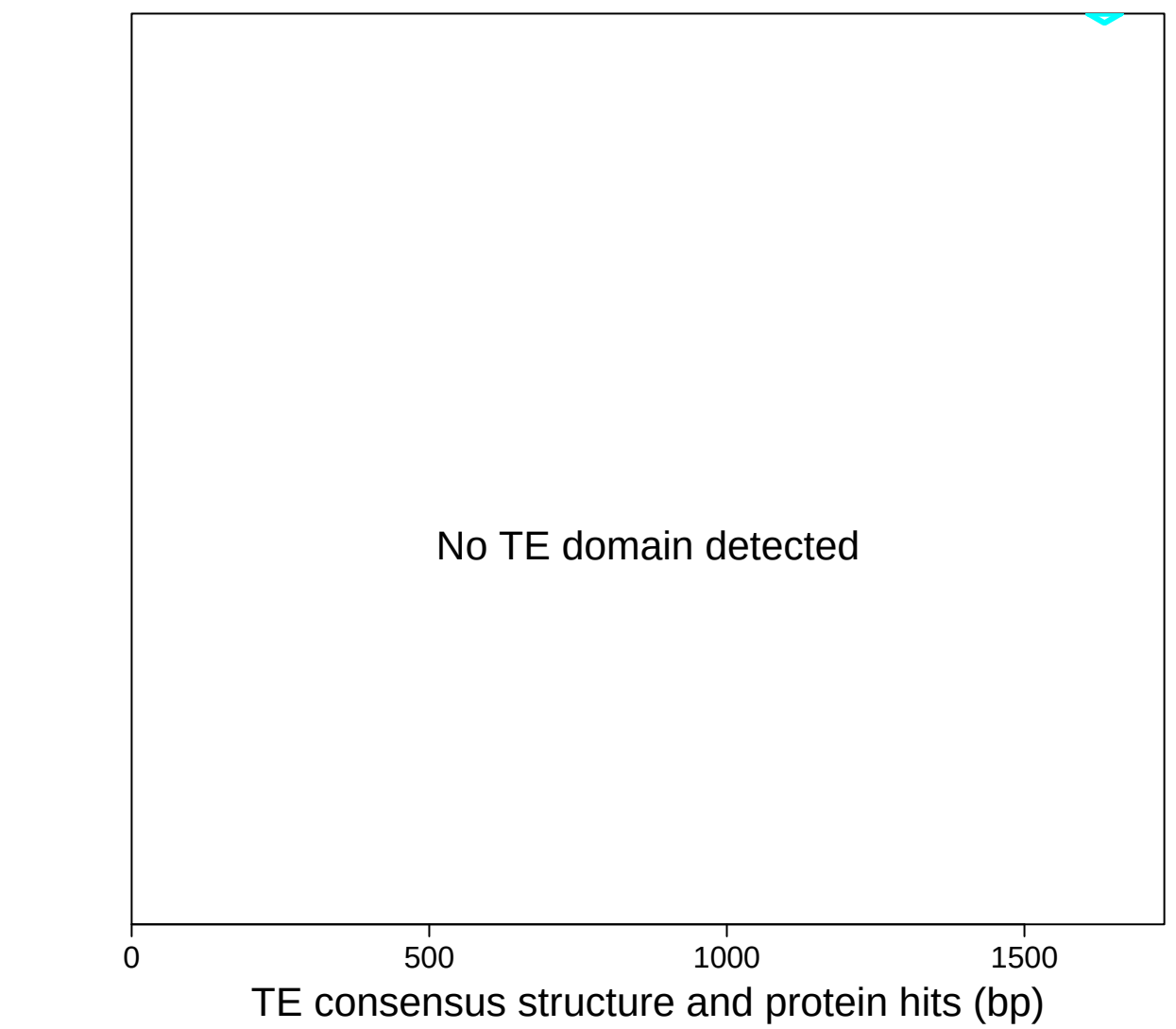
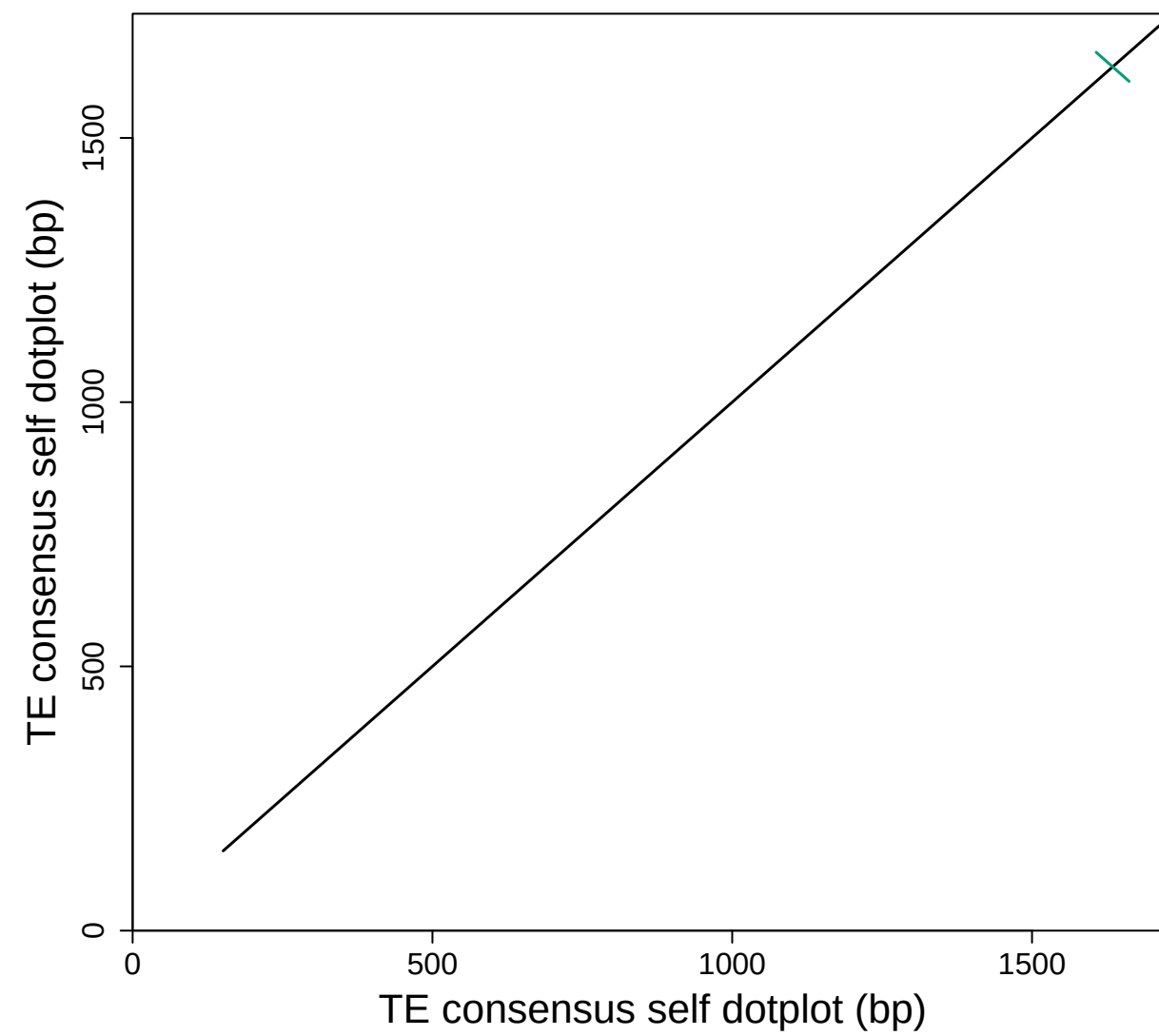
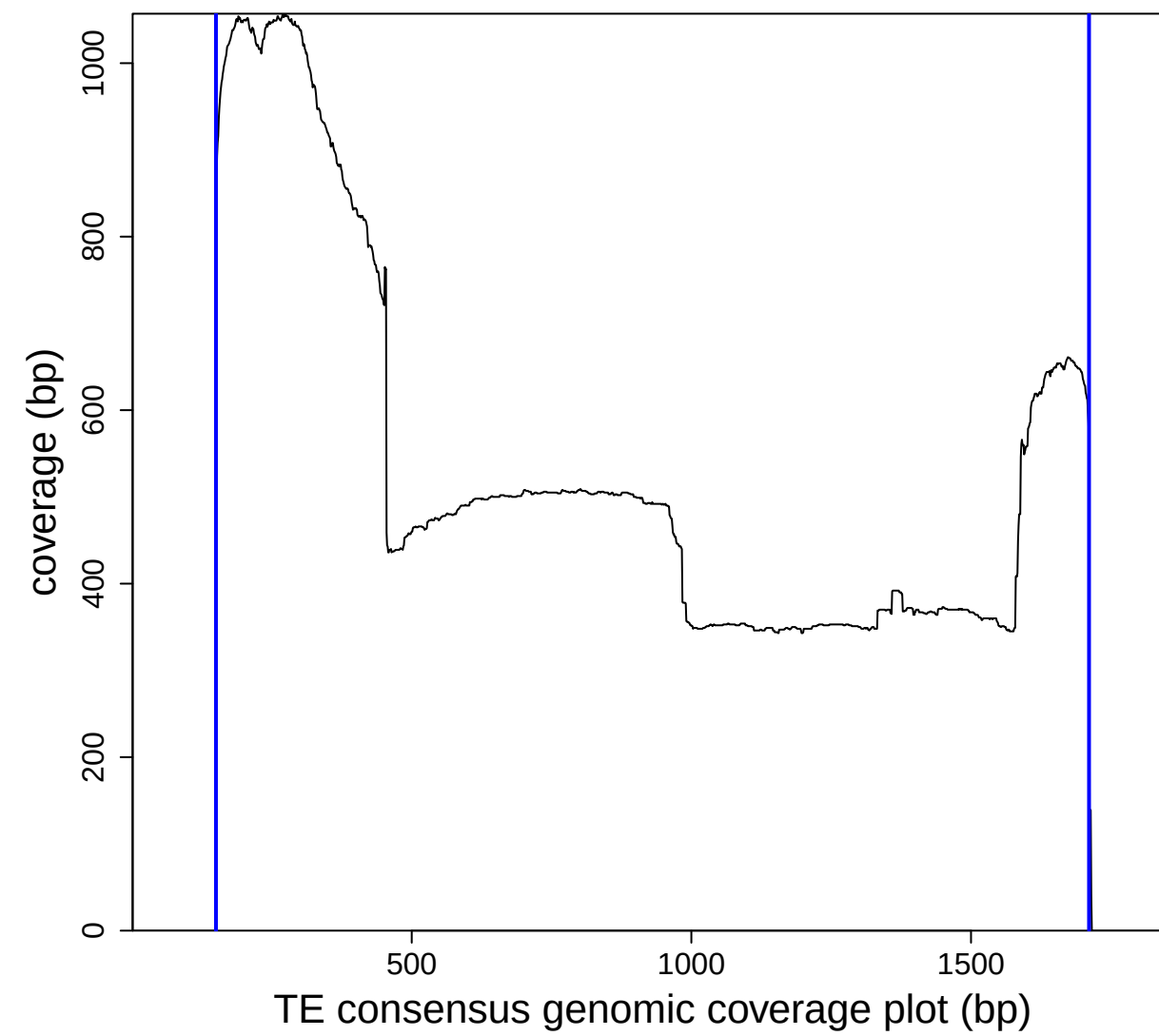
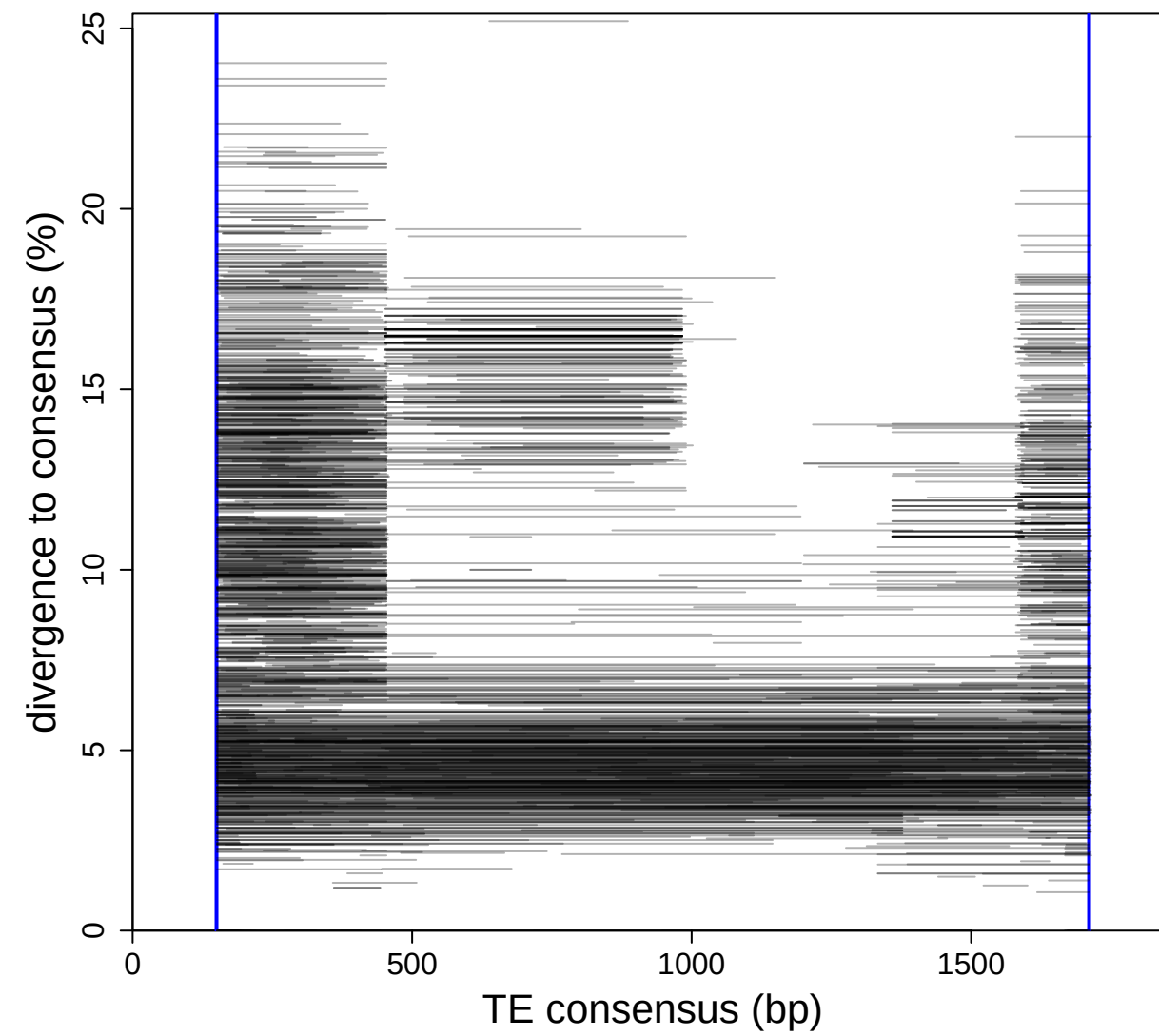
g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.fa_a
size: 1561bp; fragments: 2044; full length: 172 (>=1404.9bp)

After TEtrimmer 1561 bp

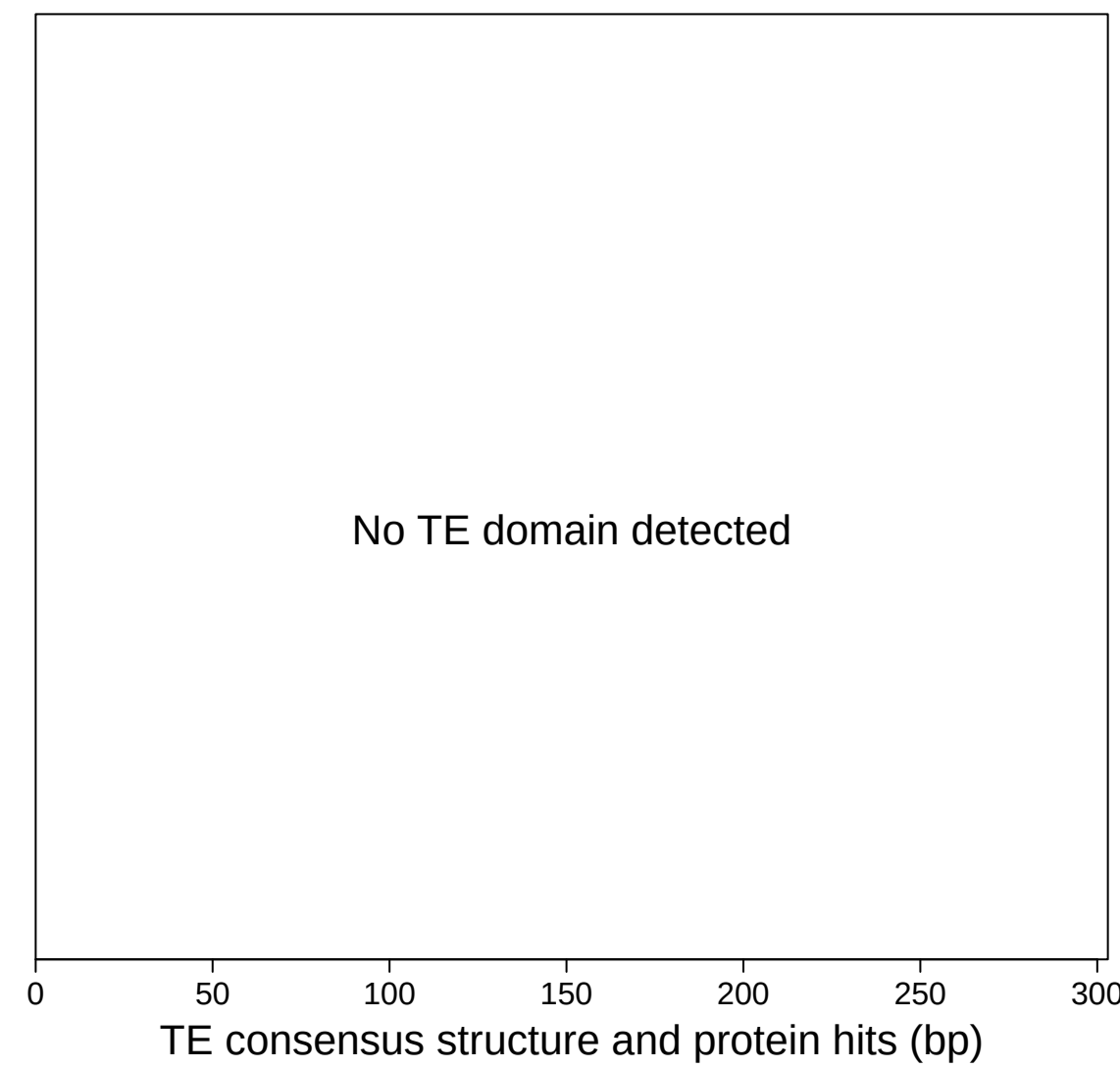
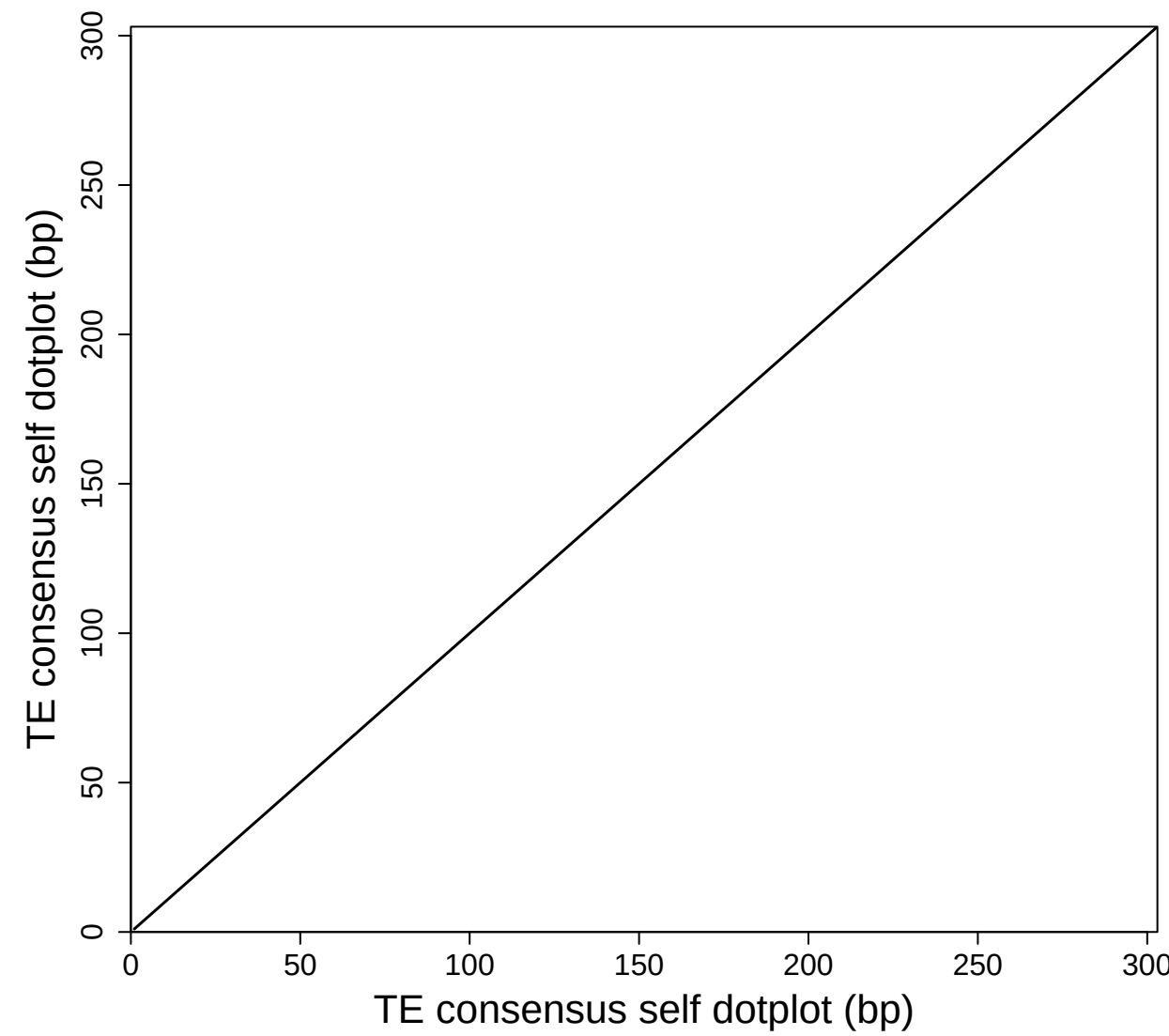
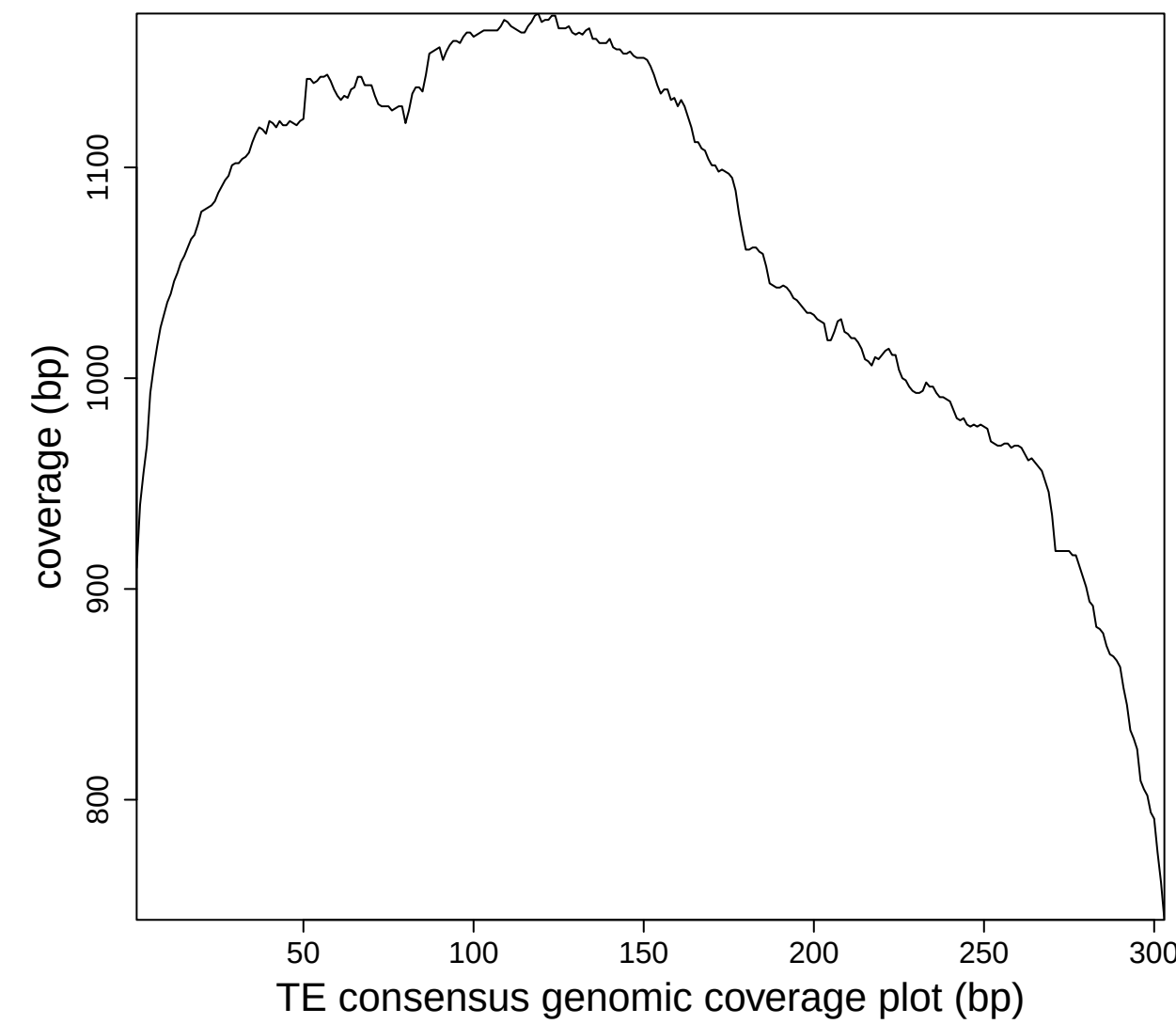
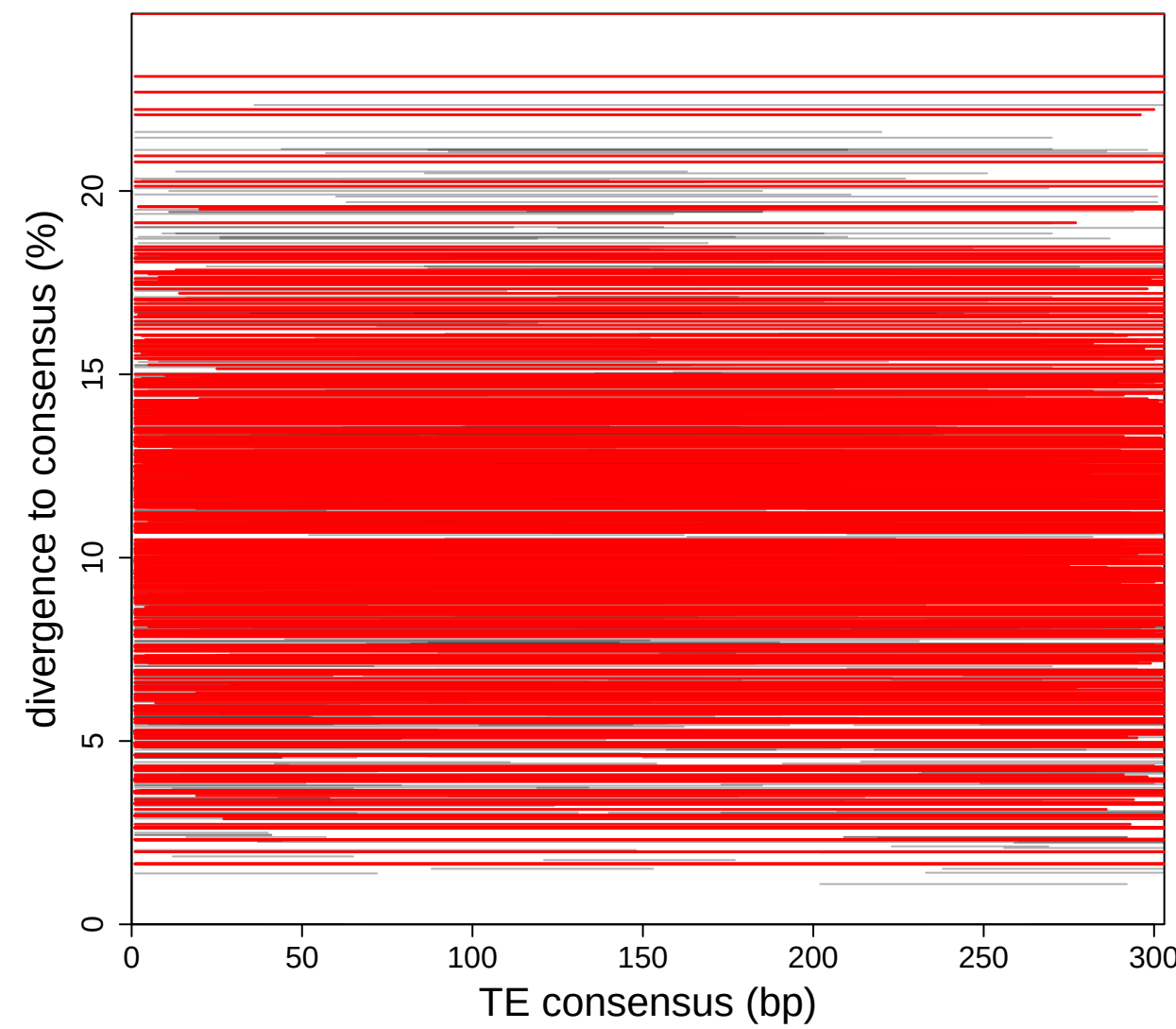


f.bed_g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcl
size: 1861bp; fragments: 2044; full length: 0 (>=1674.9bp)

After TEtrimmer Extended plot Blue lines are boundaries

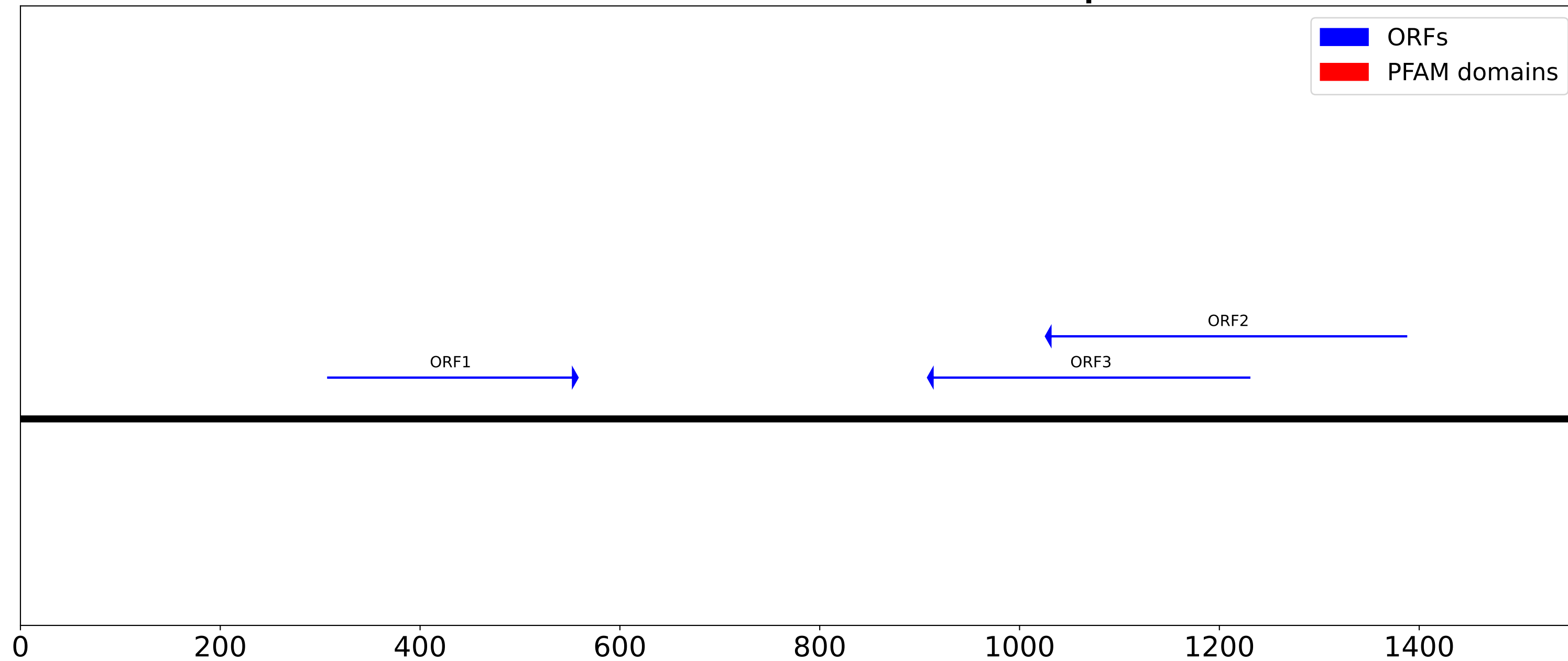


TE: rnd_1_family_962
size: 303bp; fragments: 1412; full length: 691 (≥ 272.7 bp)



Before TEtrimmer 303 bp

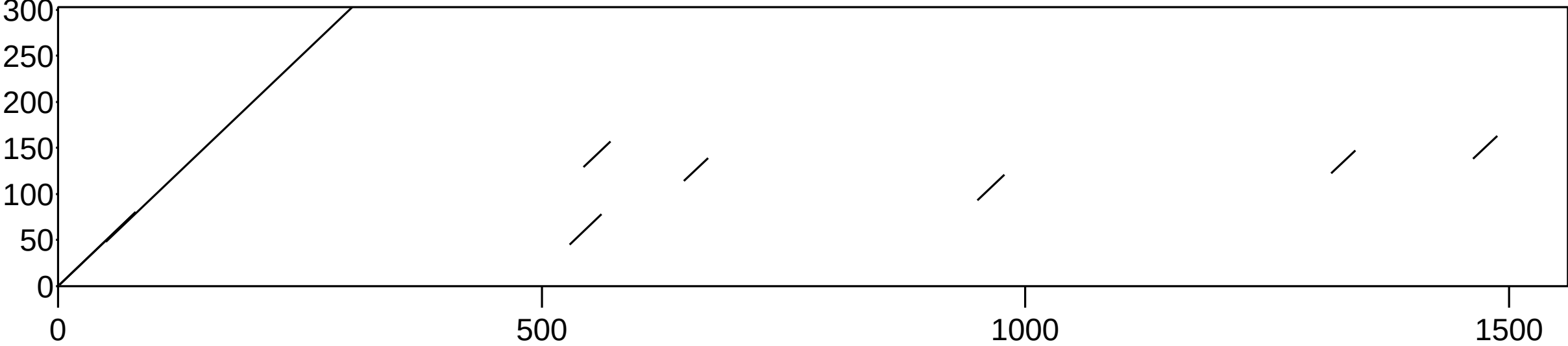
After TEtrimmer ORF and PFAM domain plot



Dotplot

(windowsize = 25, threshold = 50.00 09/10/25)

nsensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)