

MSA length = 3642

Start crop Point

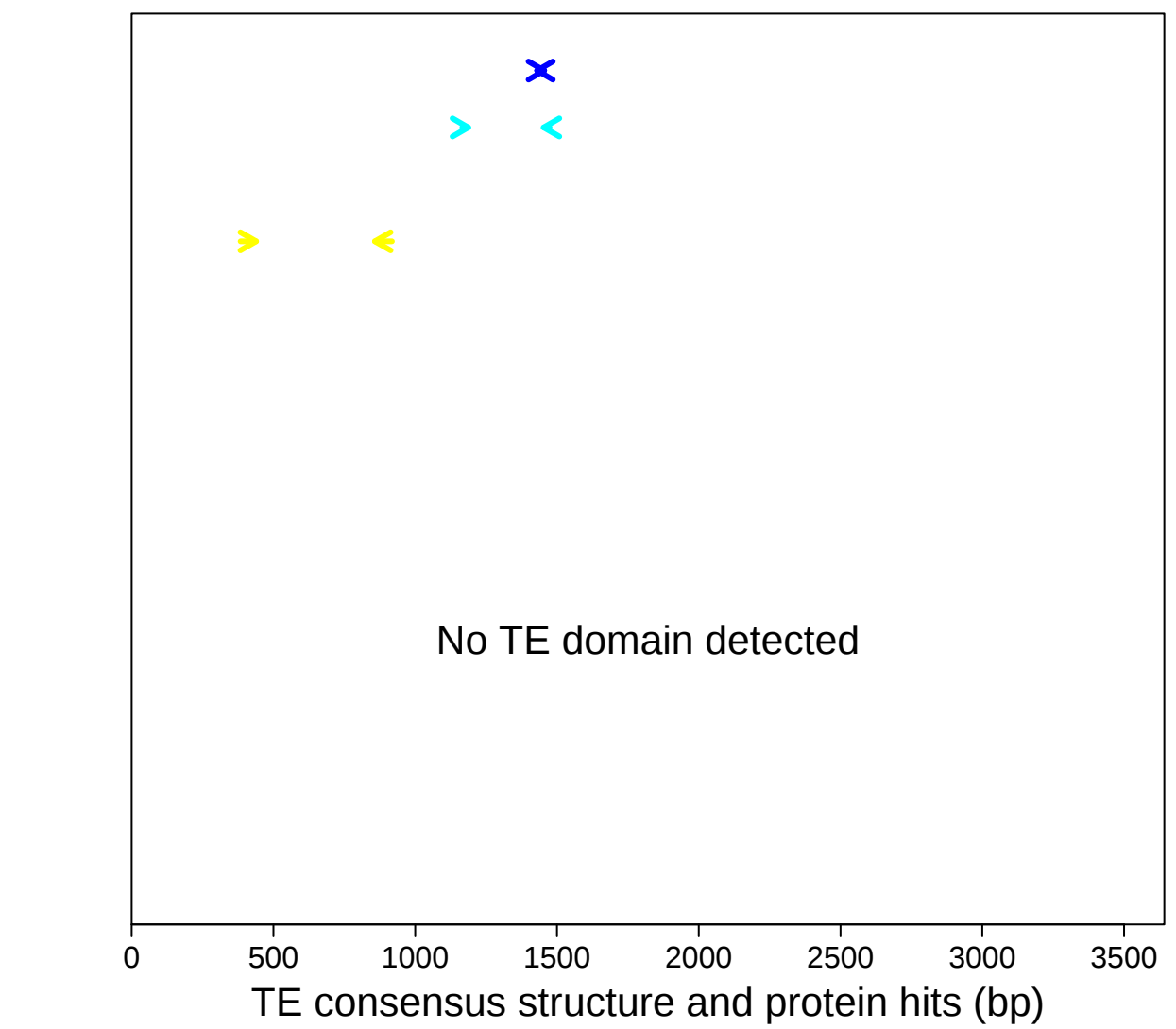
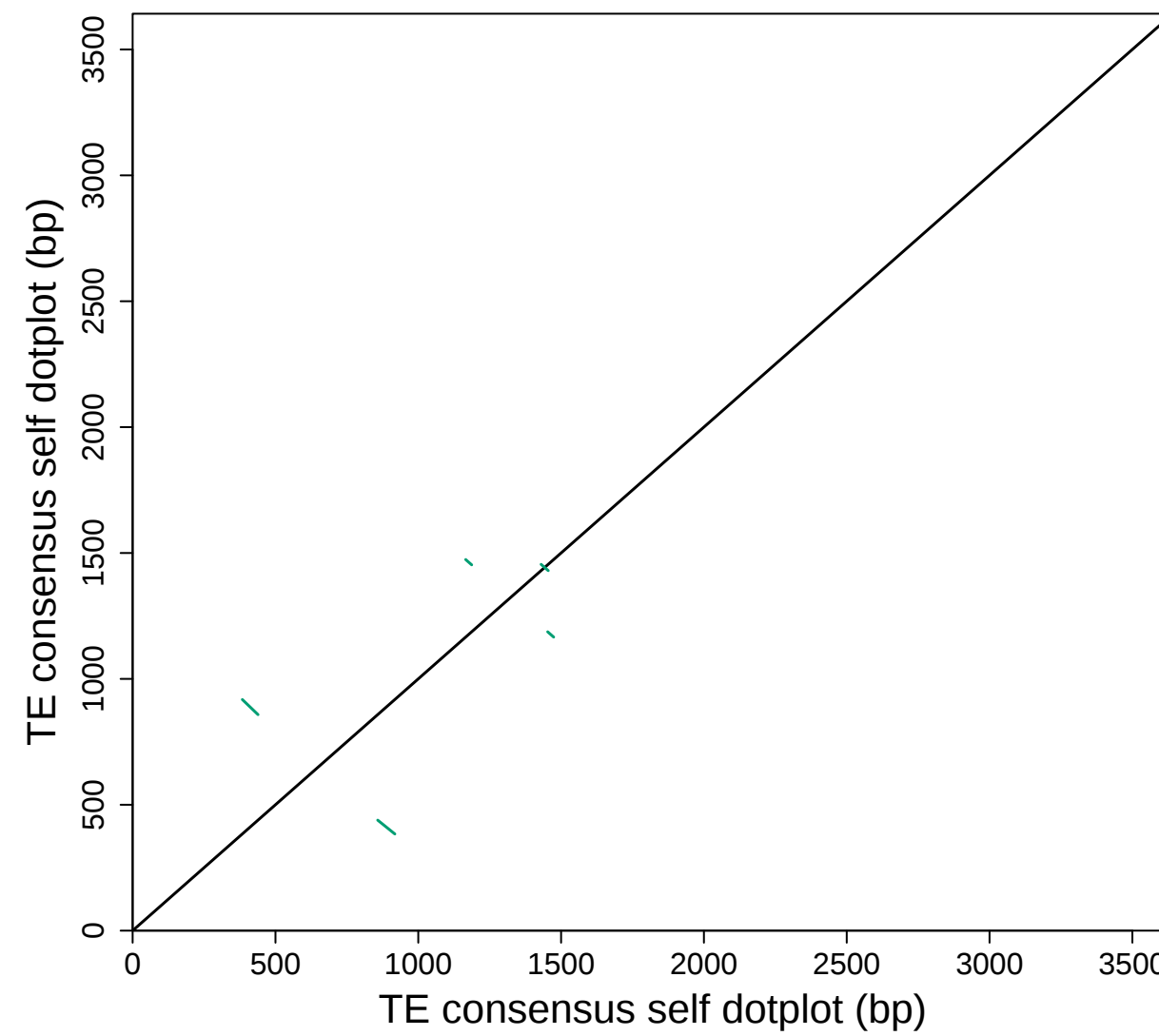
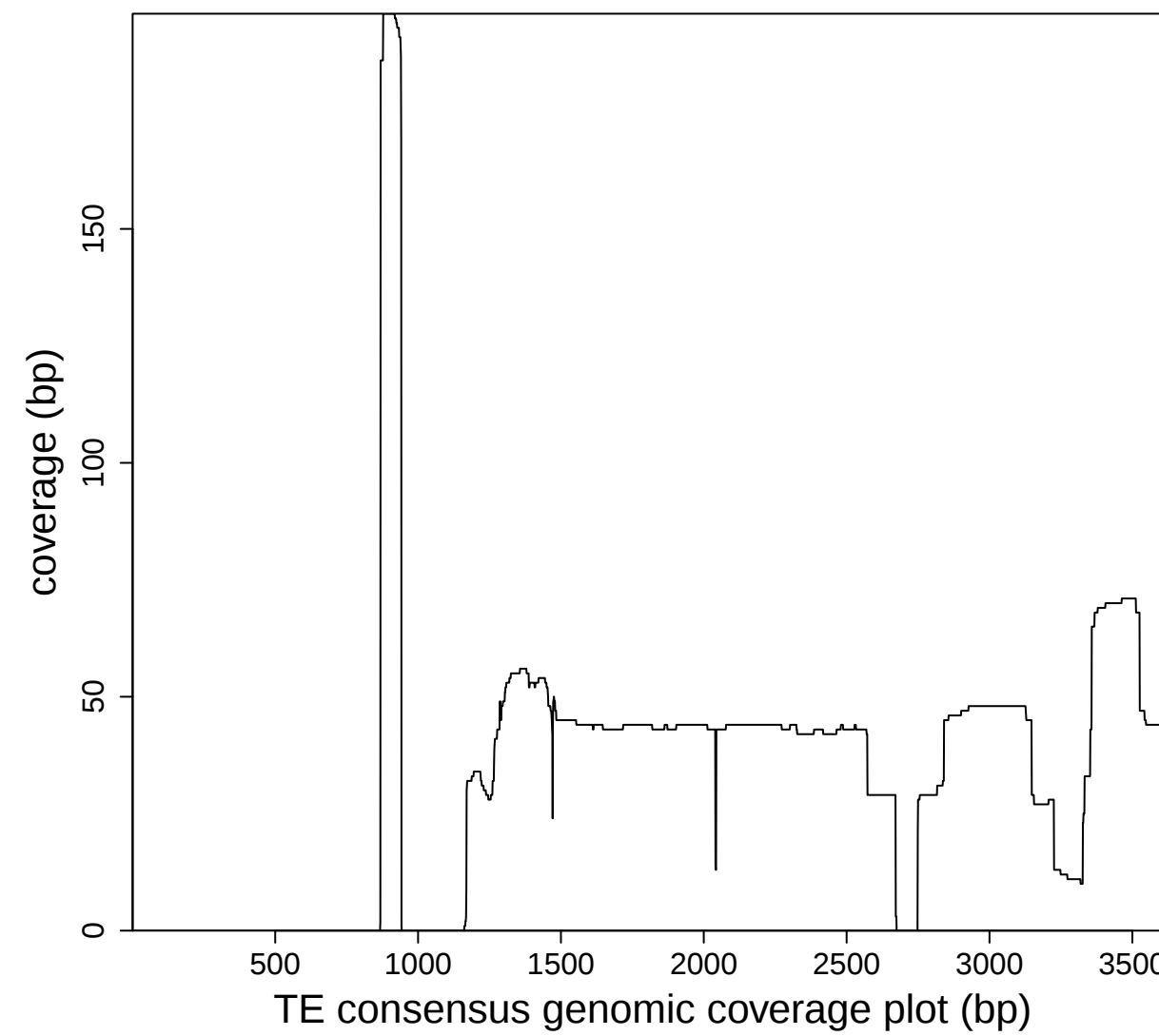
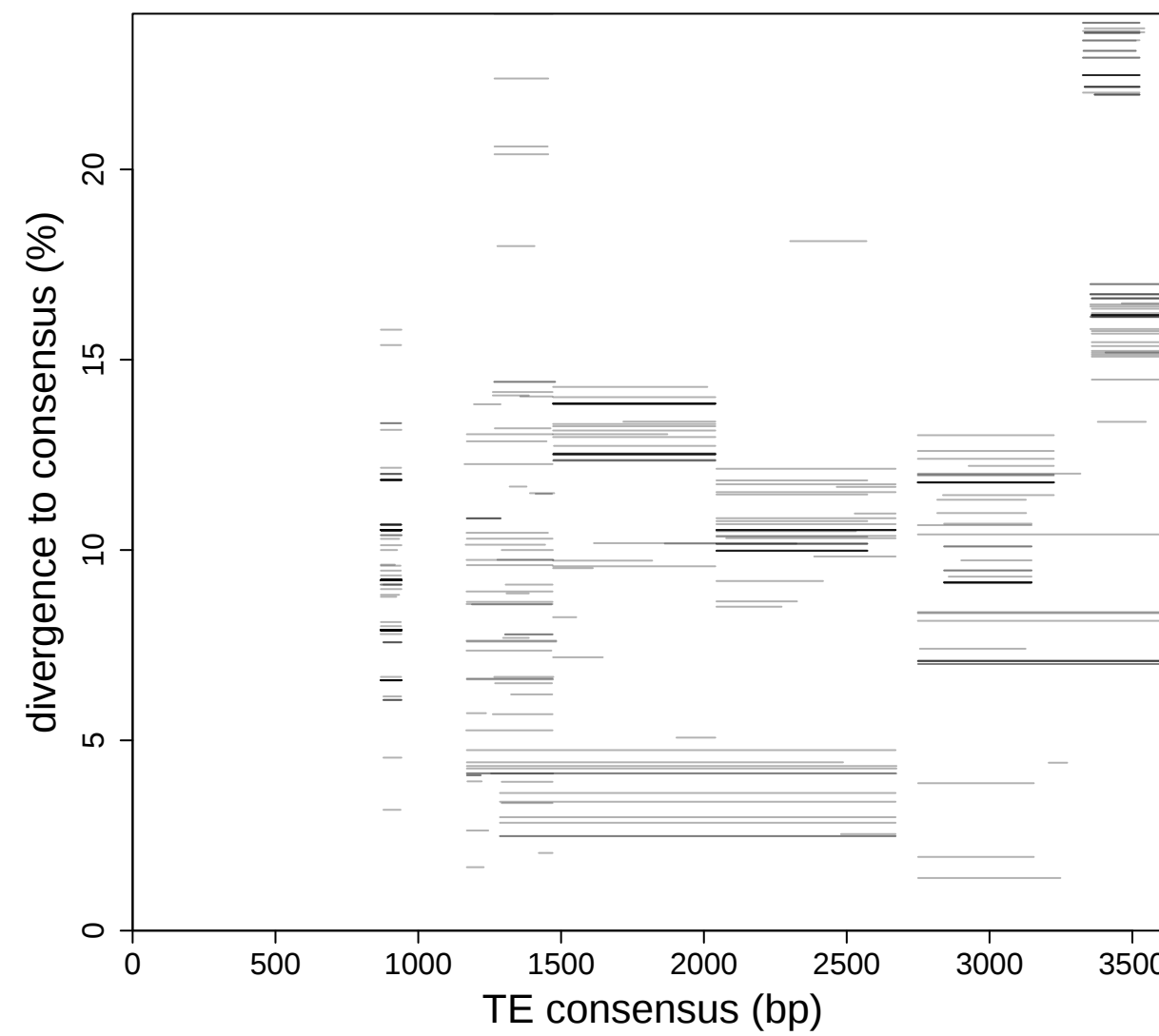
End crop Point





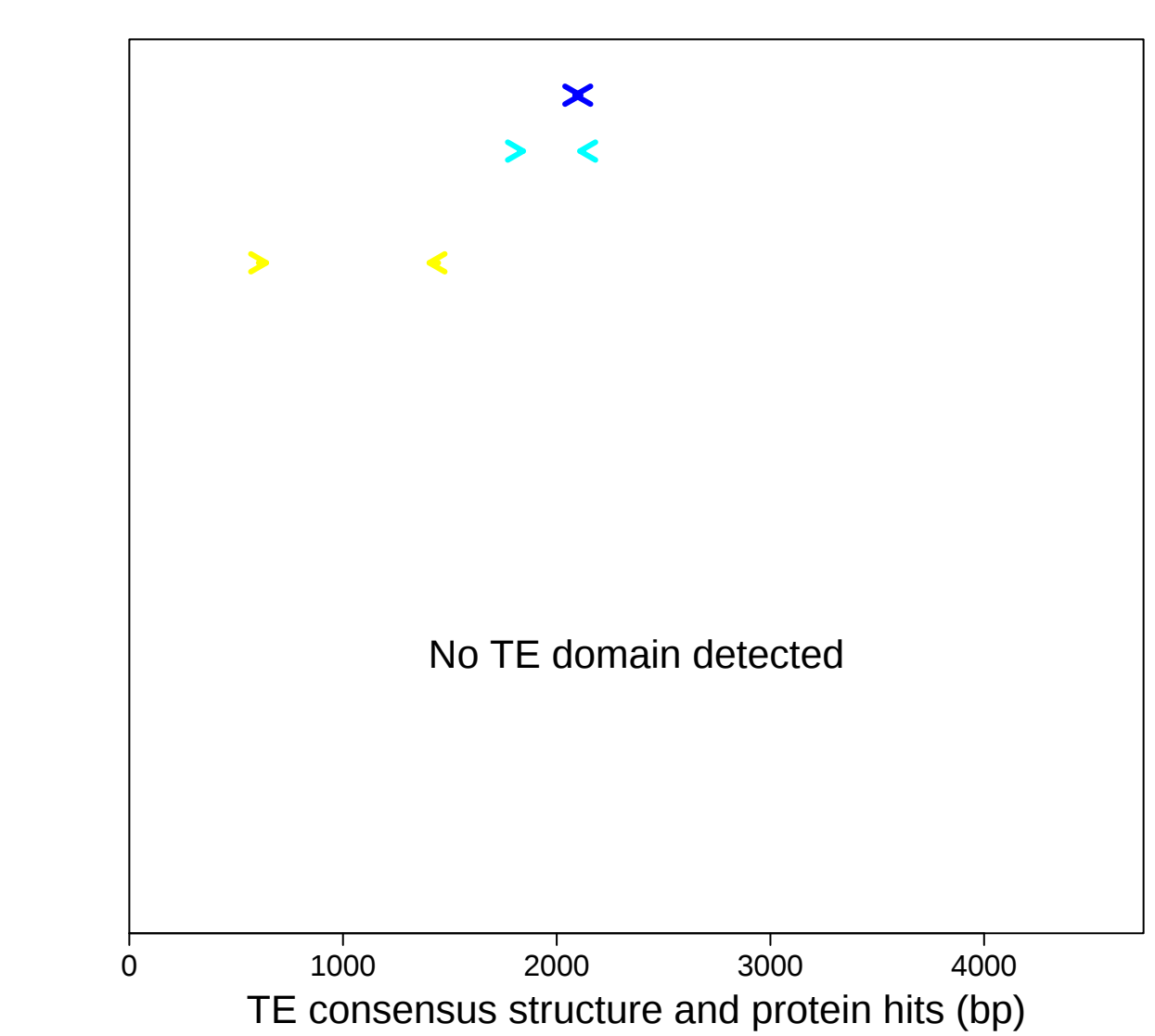
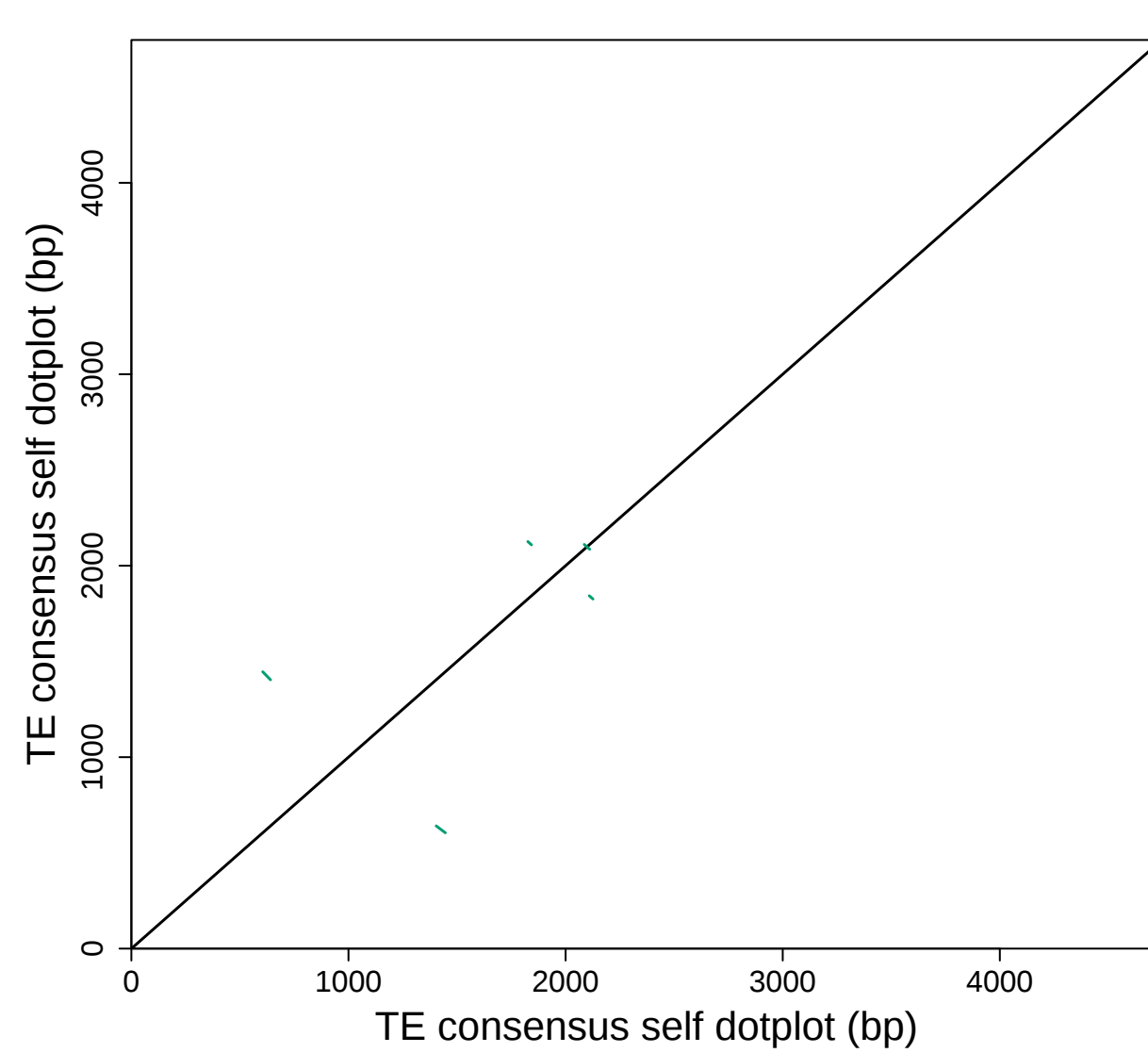
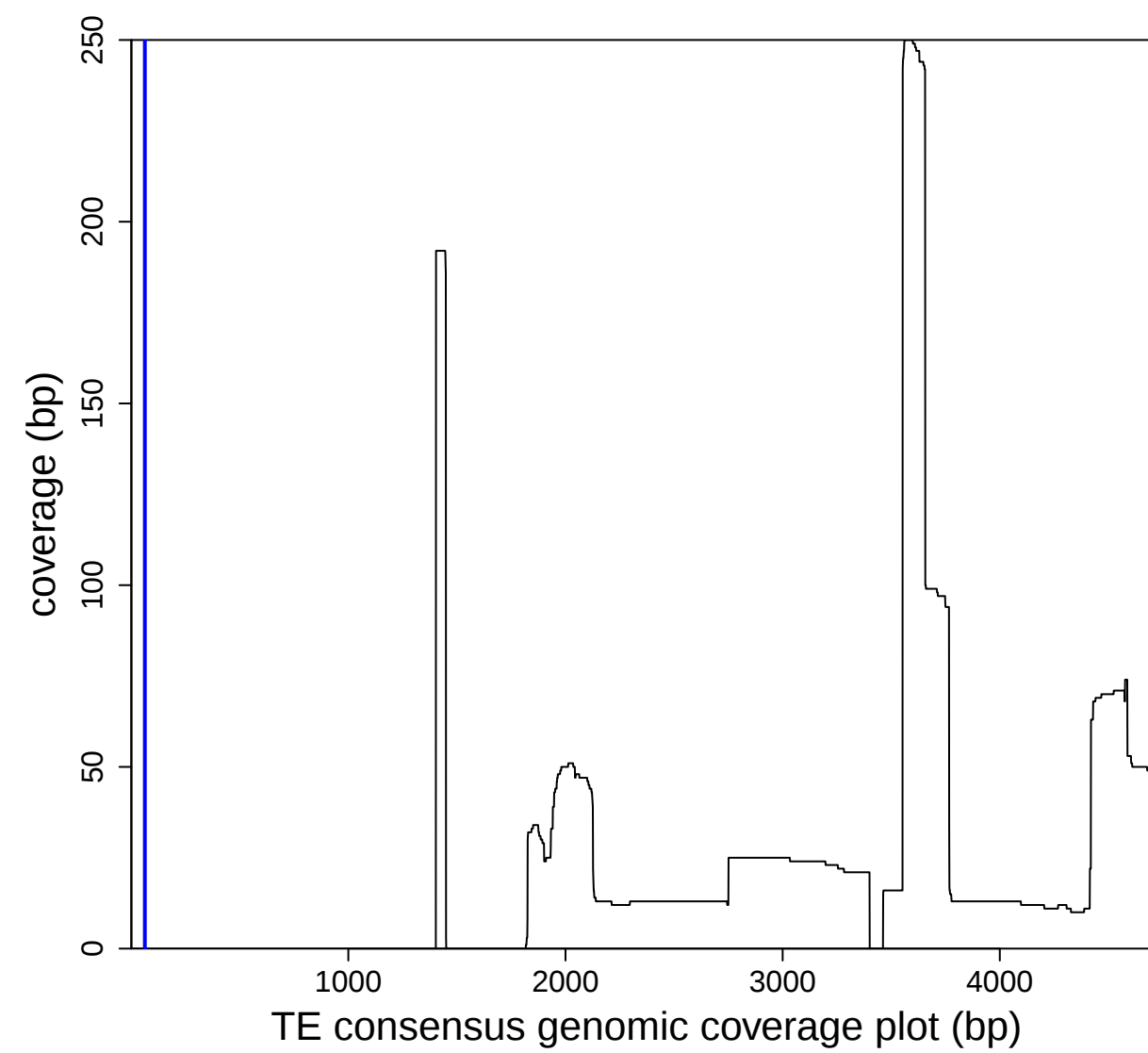
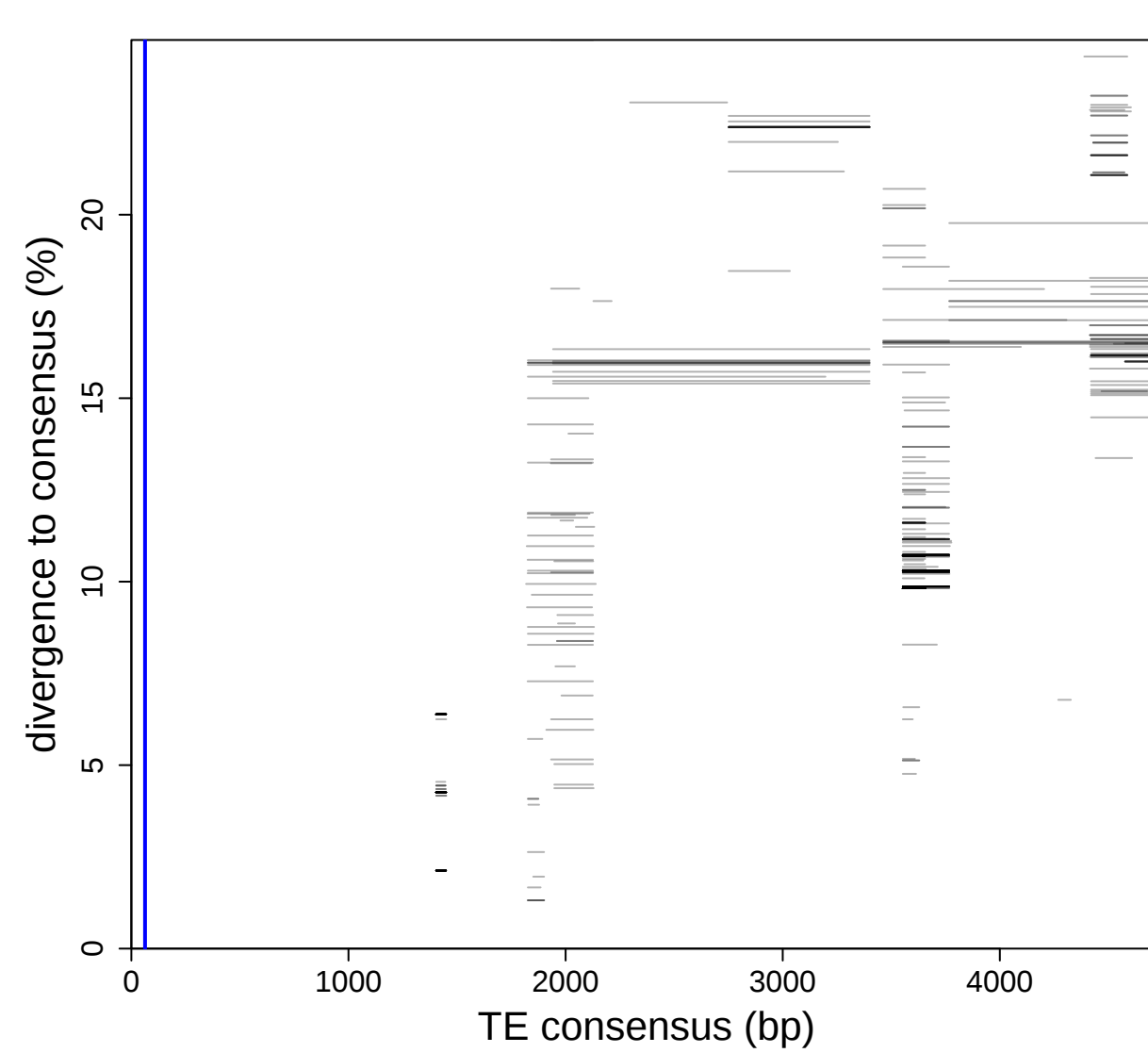
g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcIn.fa
size: 3642bp; fragments: 448; full length: 0 (>=3277.8bp)

After TEtrimmer 3642 bp

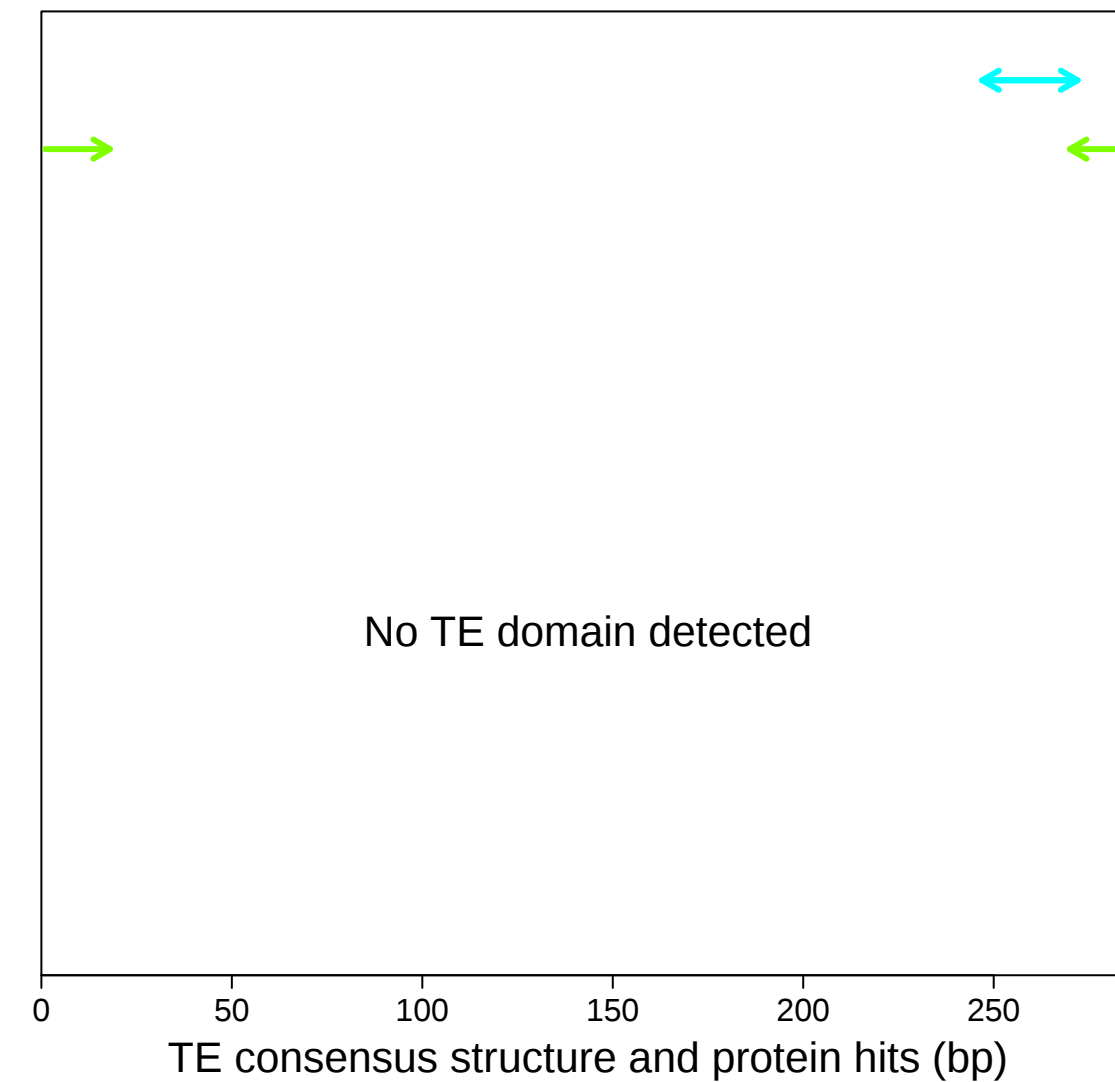
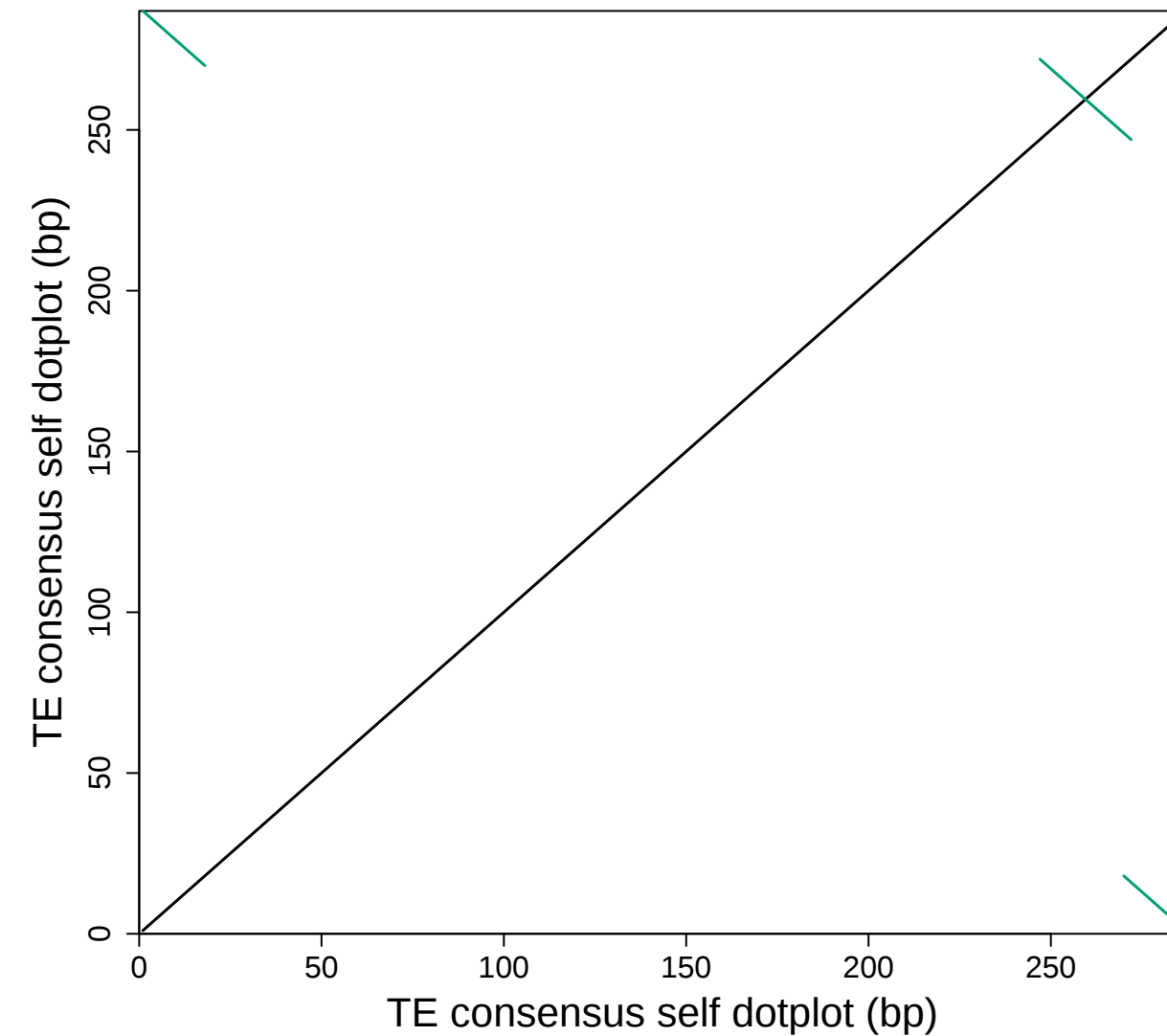
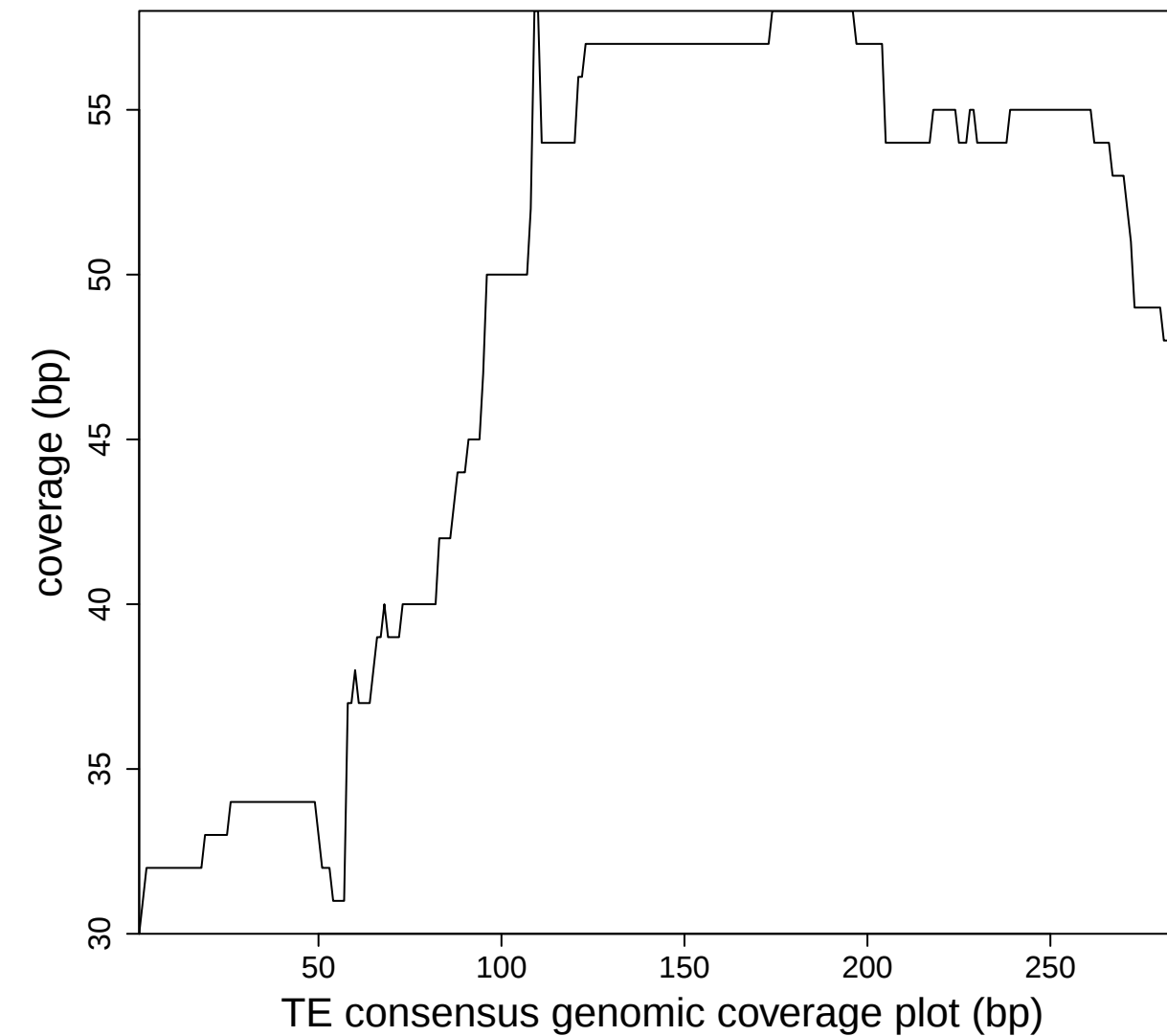
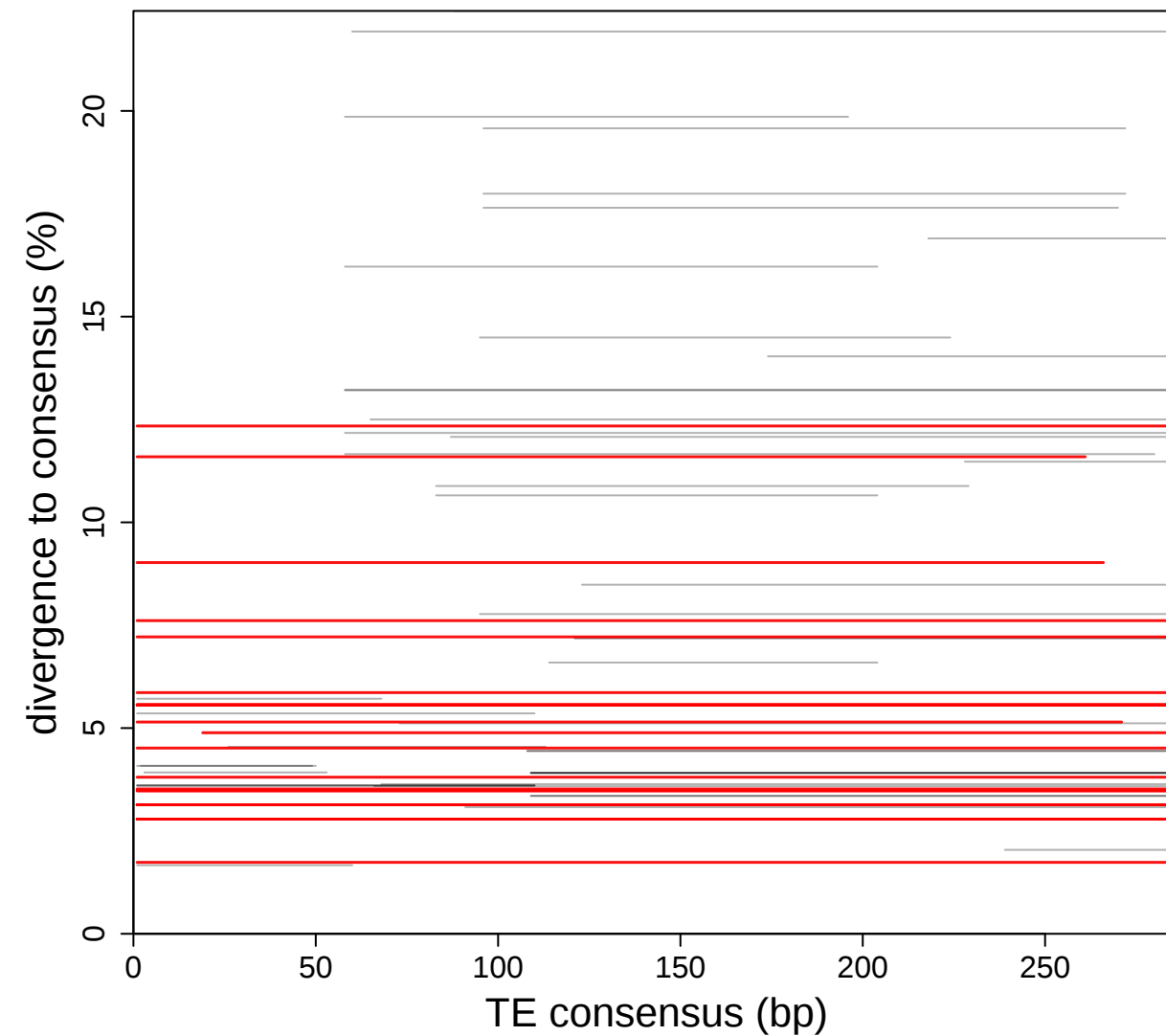


uf.bed_g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_b
size: 4746bp; fragments: 593; full length: 0 (>=4271.4bp)

After TEtrimmer Extended plot Blue lines are boundaries

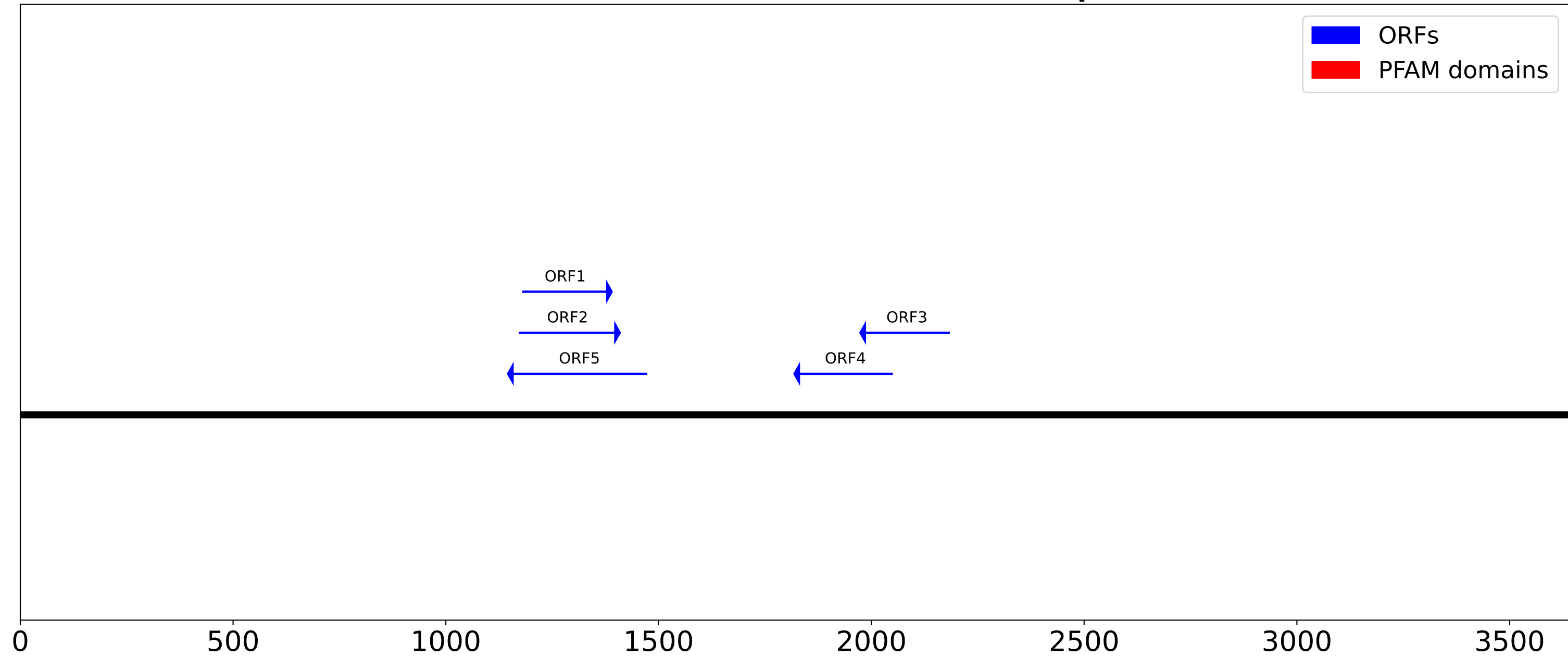


TE: rnd_5_family_12892
size: 287bp; fragments: 71; full length: 24 (≥ 258.3 bp)



Before TEtrimmer 287 bp

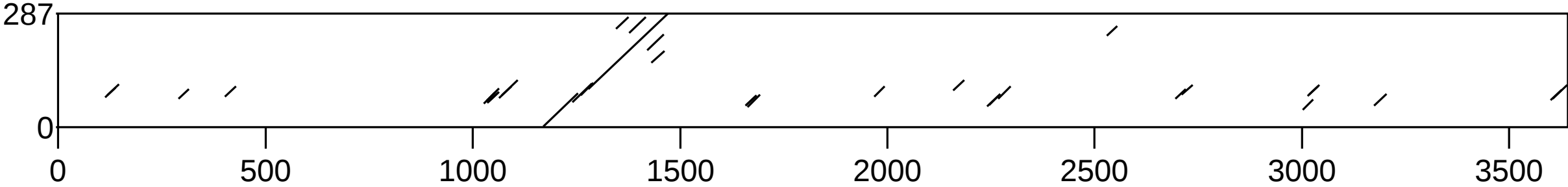
After TEtrimmer ORF and PFAM domain plot



Dotplot

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

ensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)