

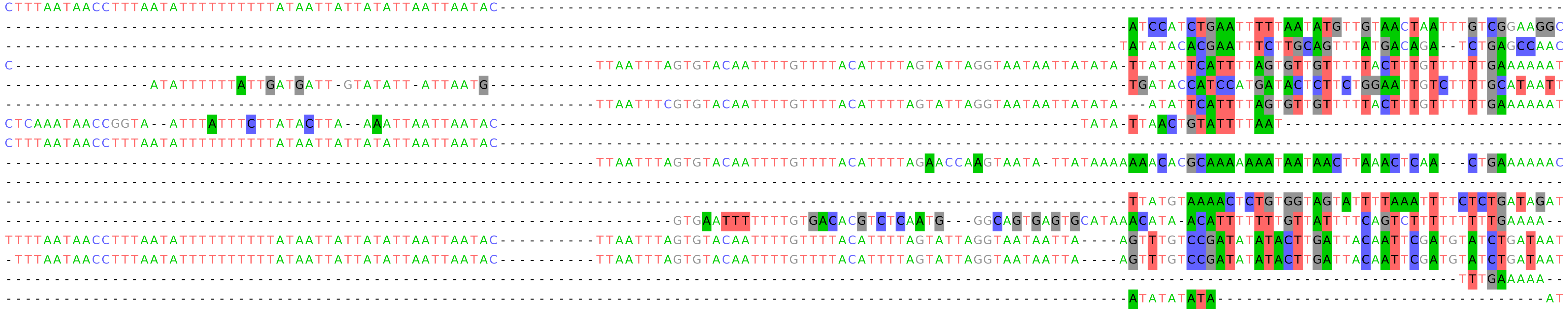
Start crop Point

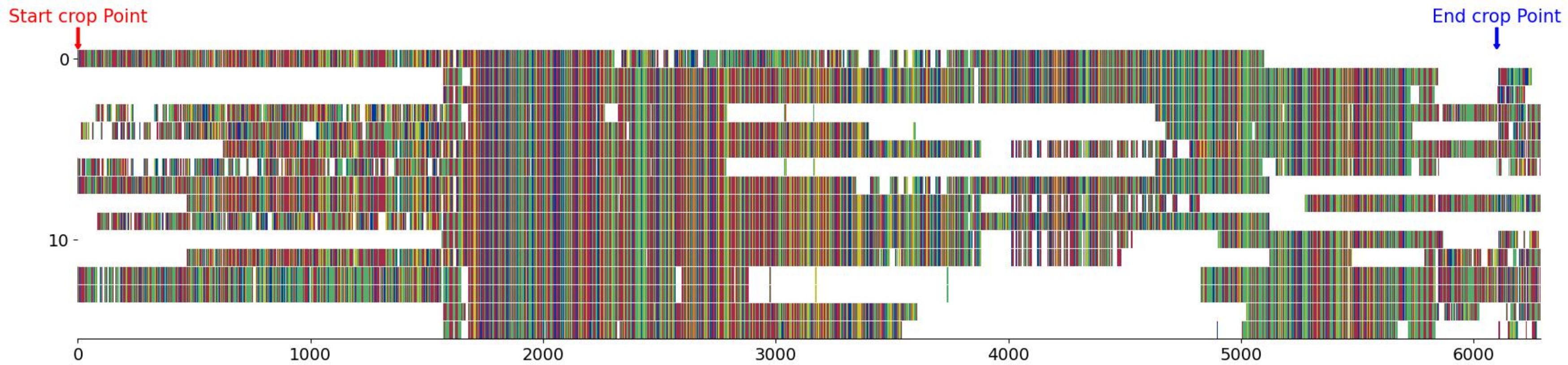


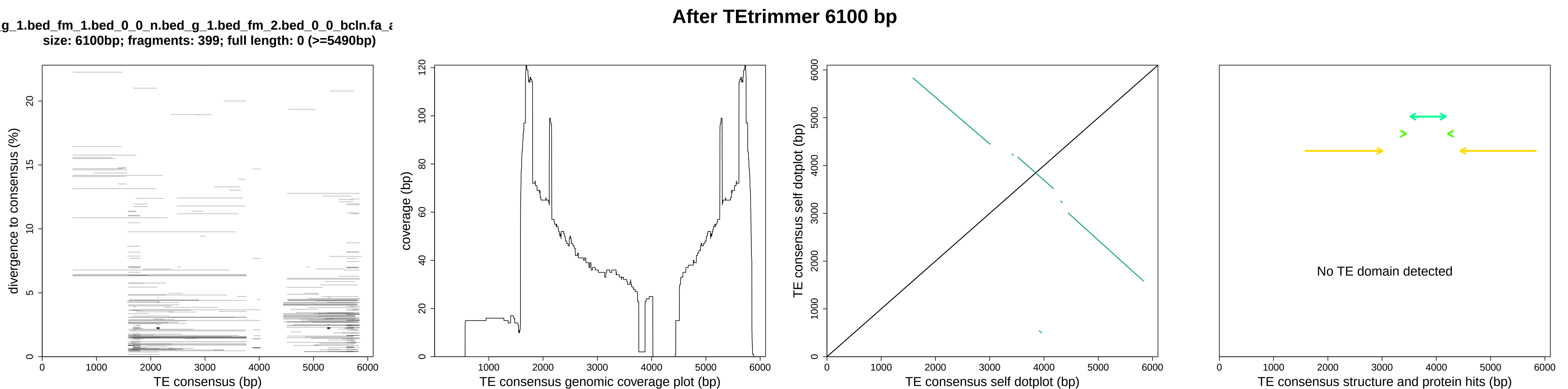
End crop Point



MSA length = 6100

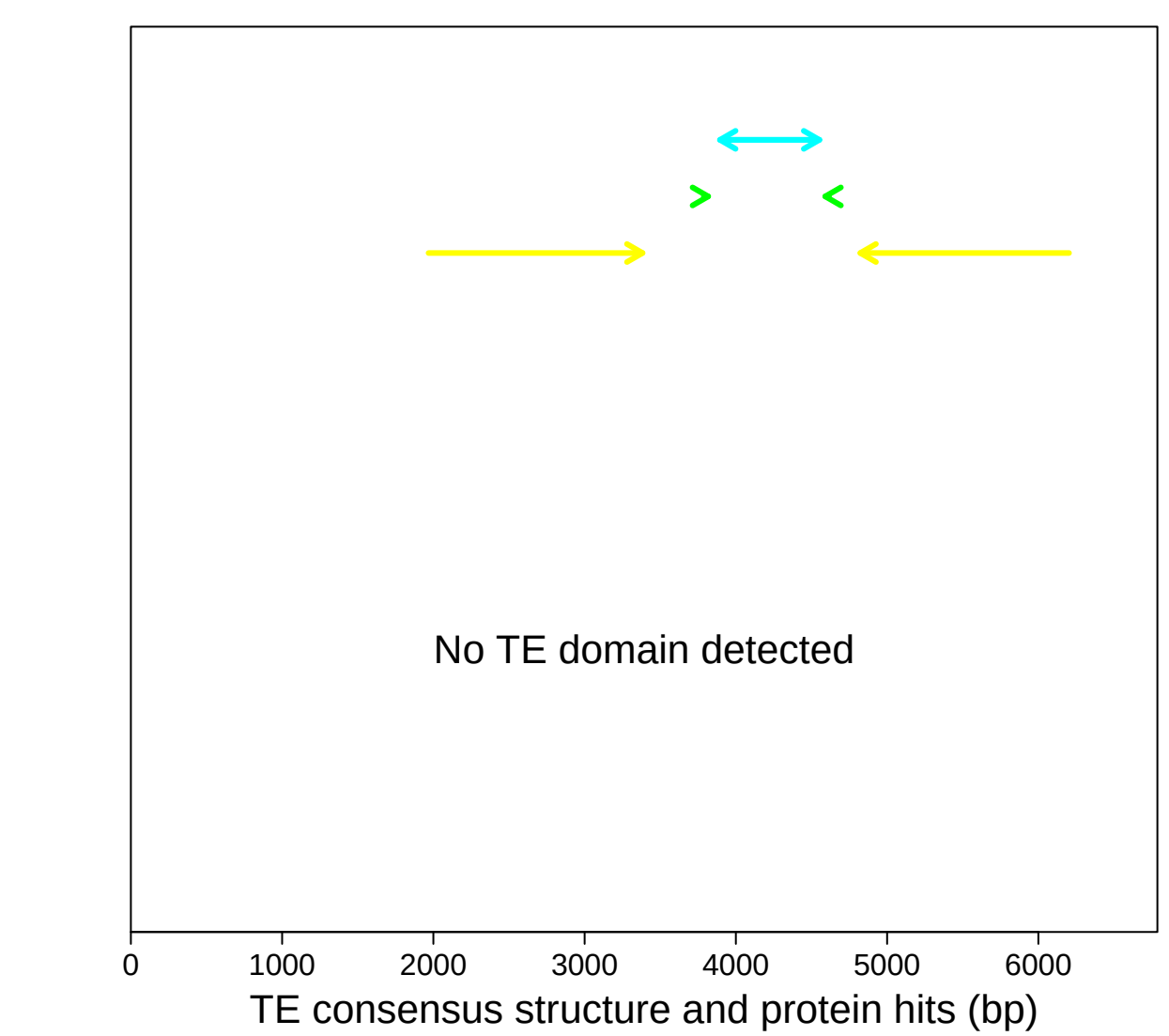
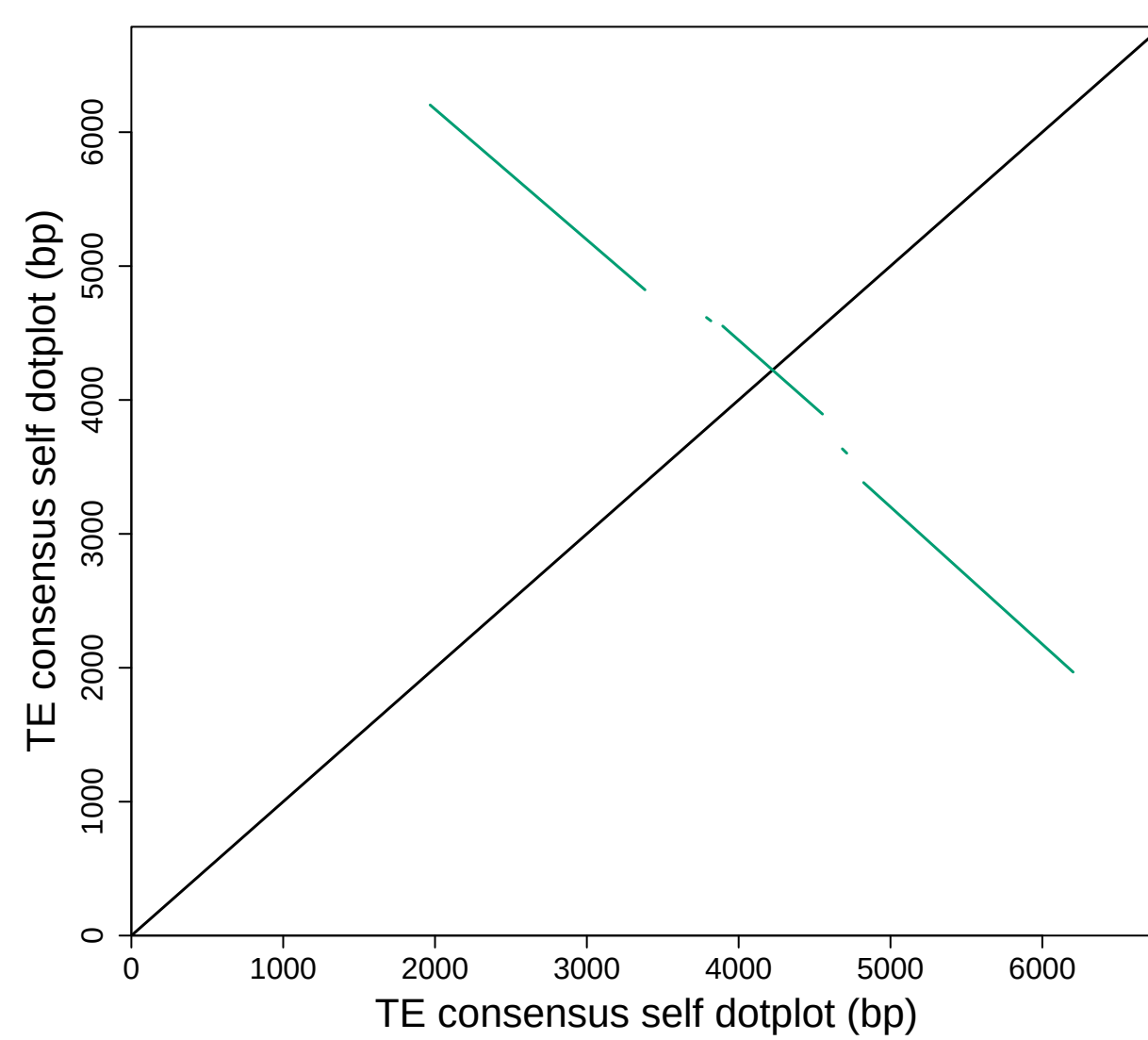
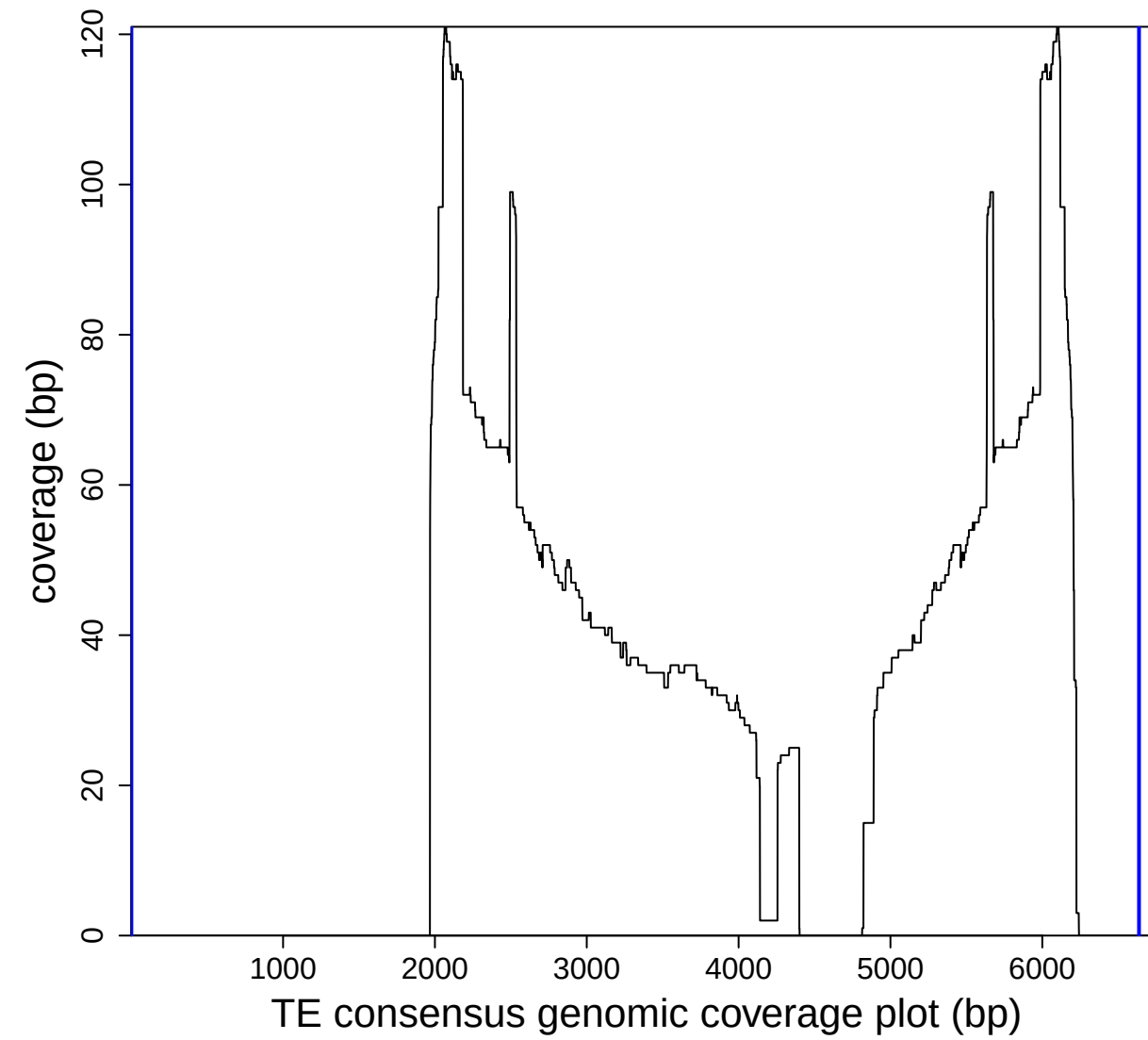
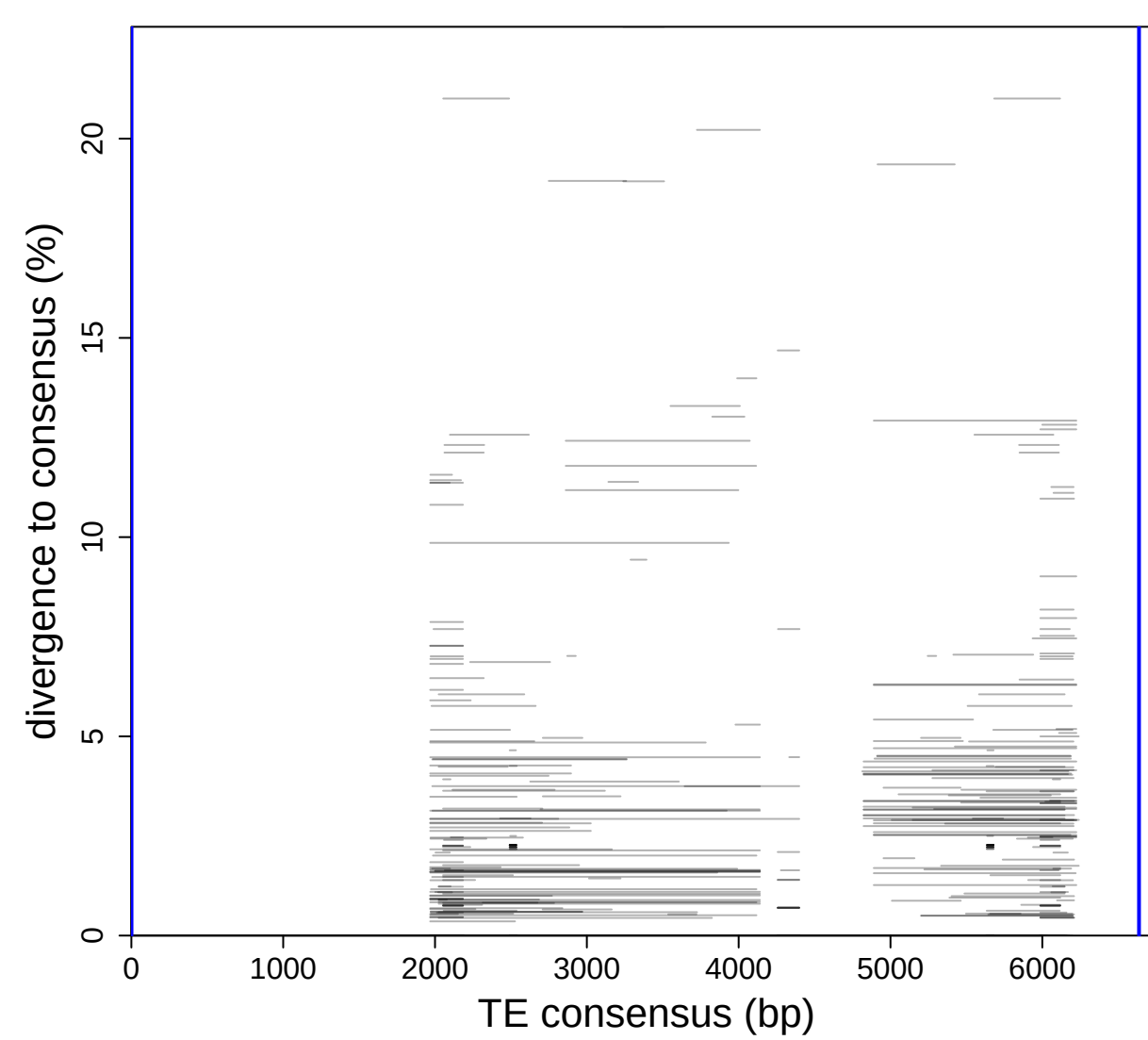




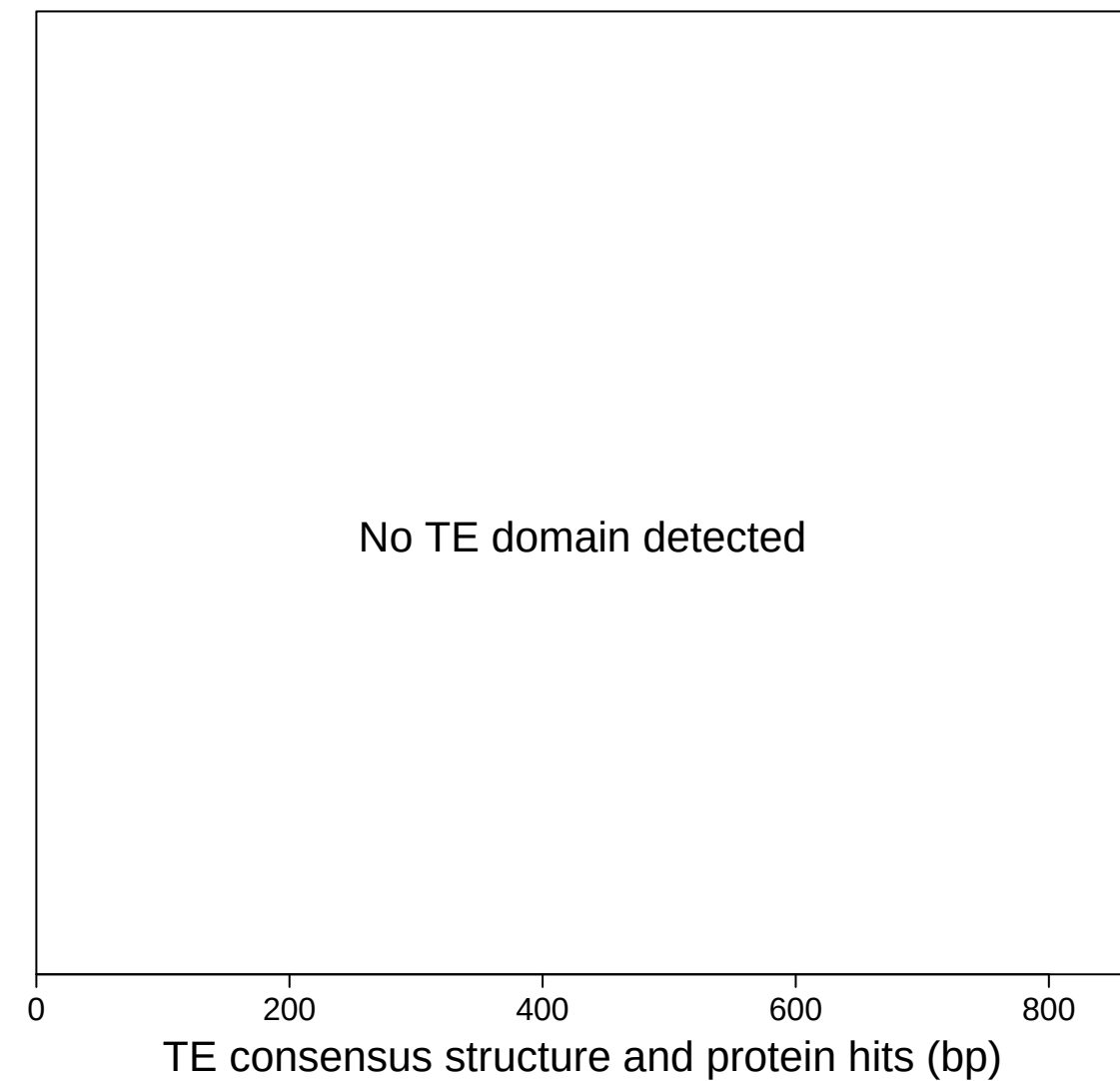
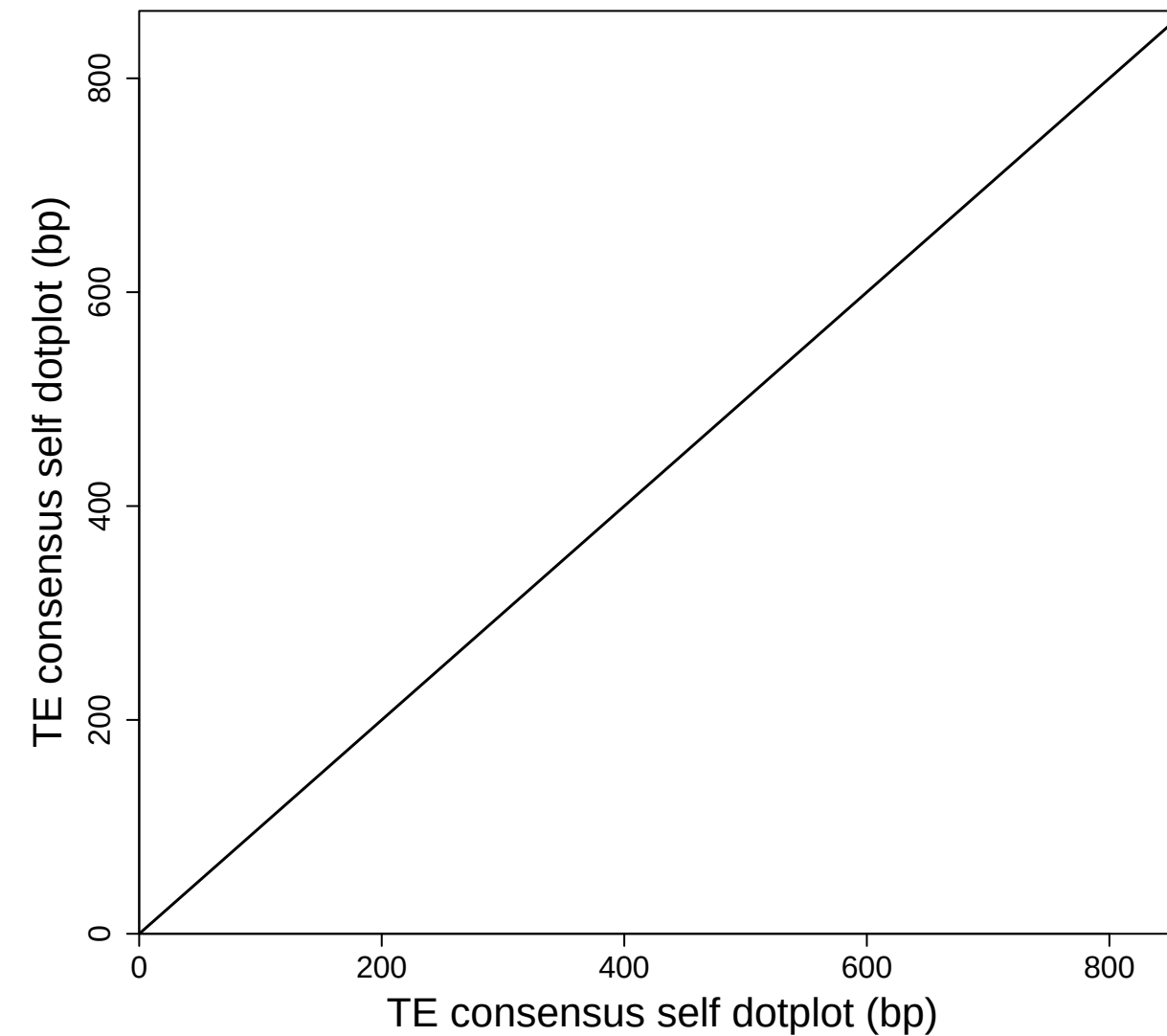
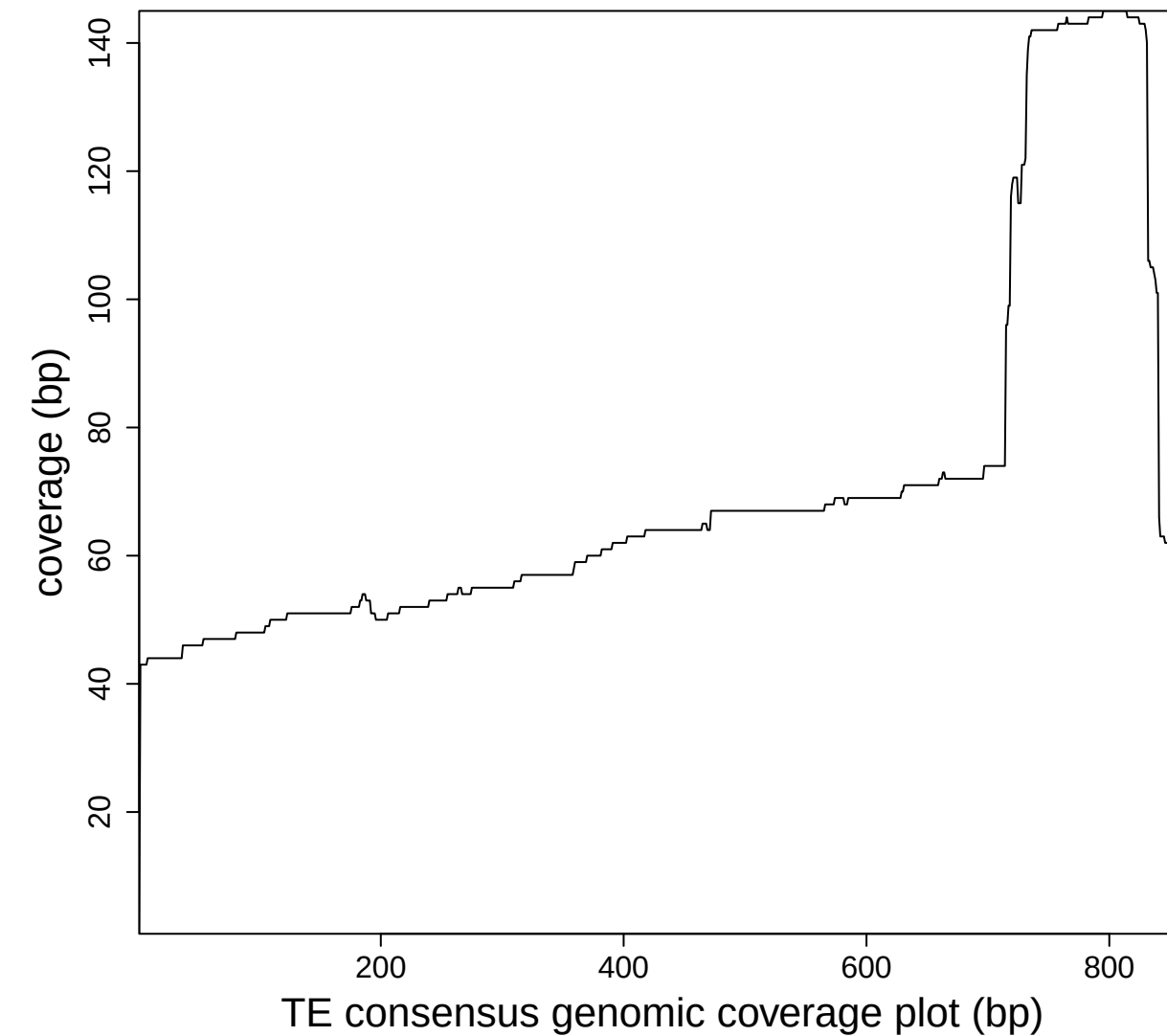
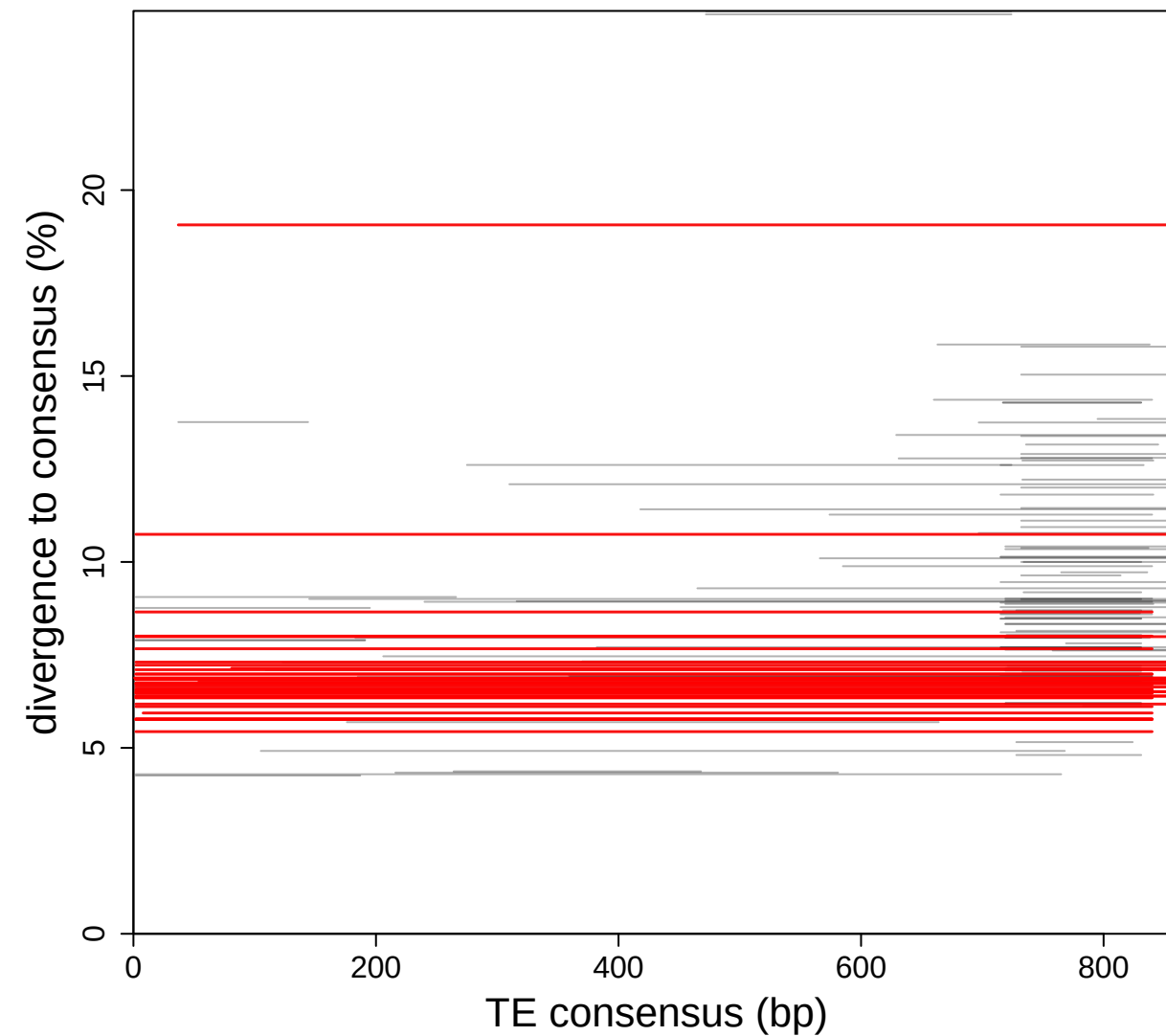


f.bed_g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bc
size: 6787bp; fragments: 388; full length: 0 (>=6108.3bp)

After TEtrimmer Extended plot Blue lines are boundaries

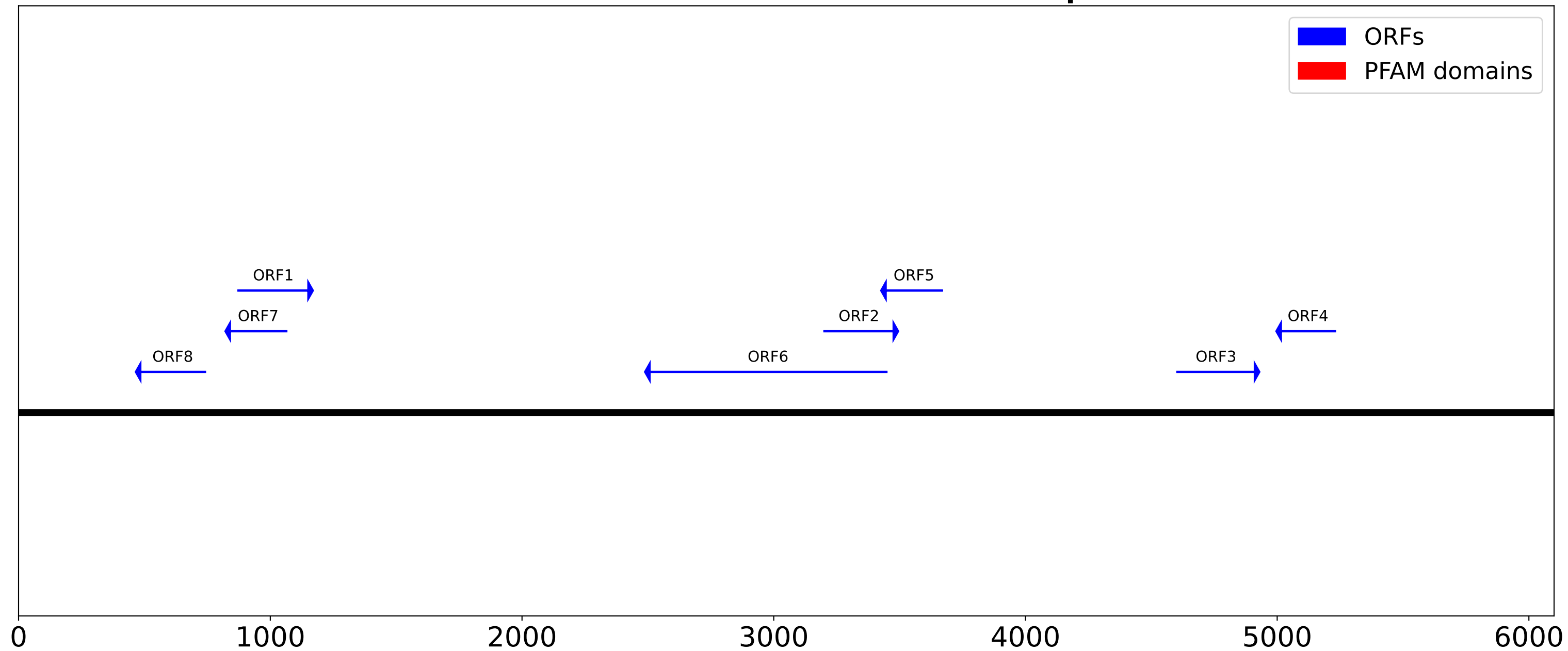


TE: rnd_5_family_2561
size: 863bp; fragments: 160; full length: 41 (≥ 776.7 bp)



Before TEtrimmer 863 bp

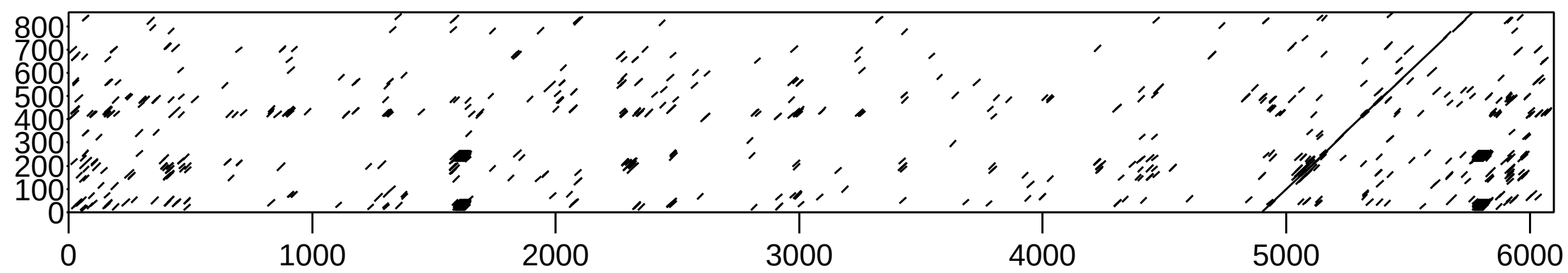
After TEtrimmer ORF and PFAM domain plot



Dotplot

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

isensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)