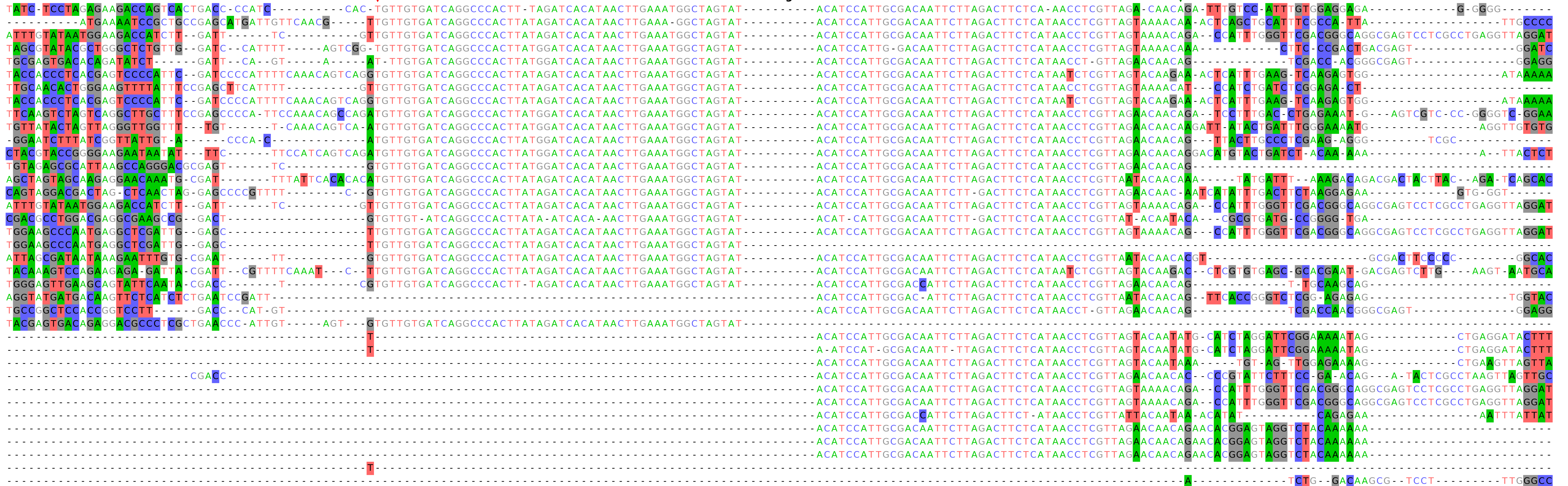
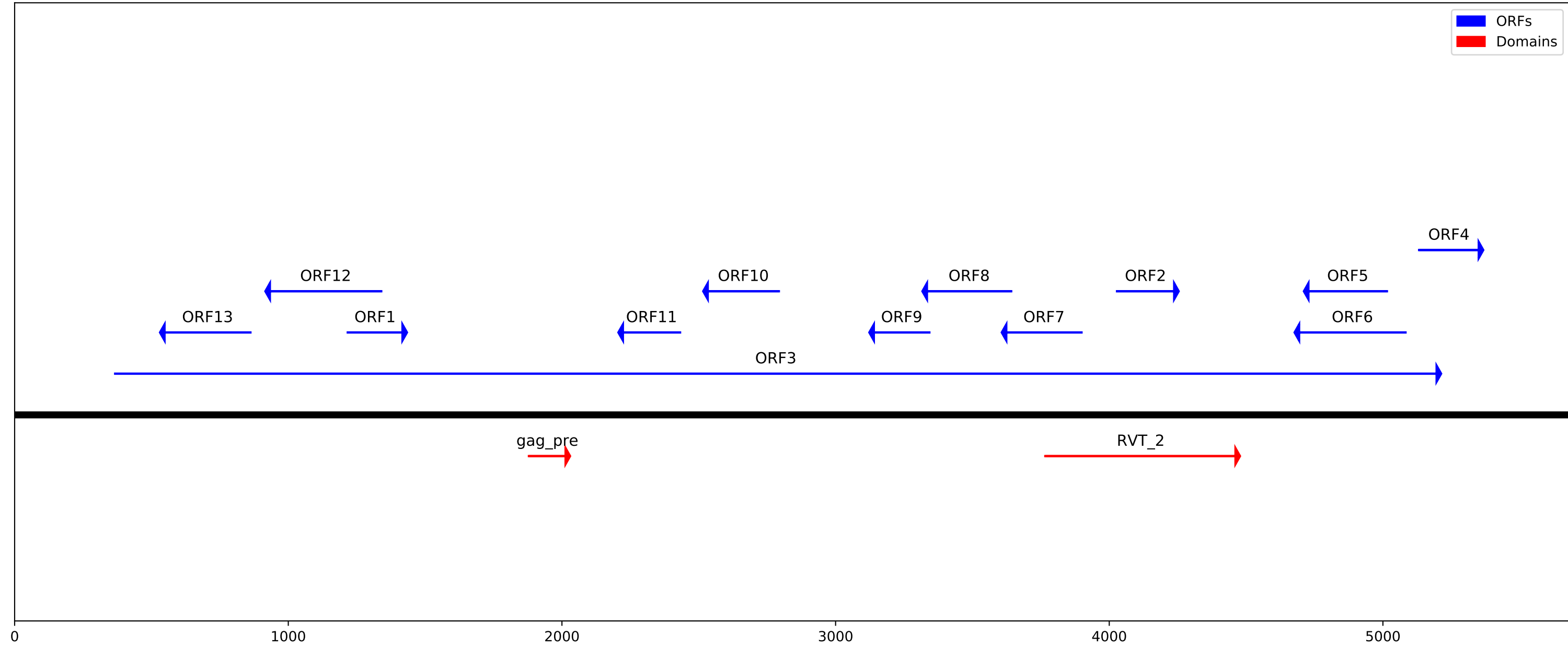


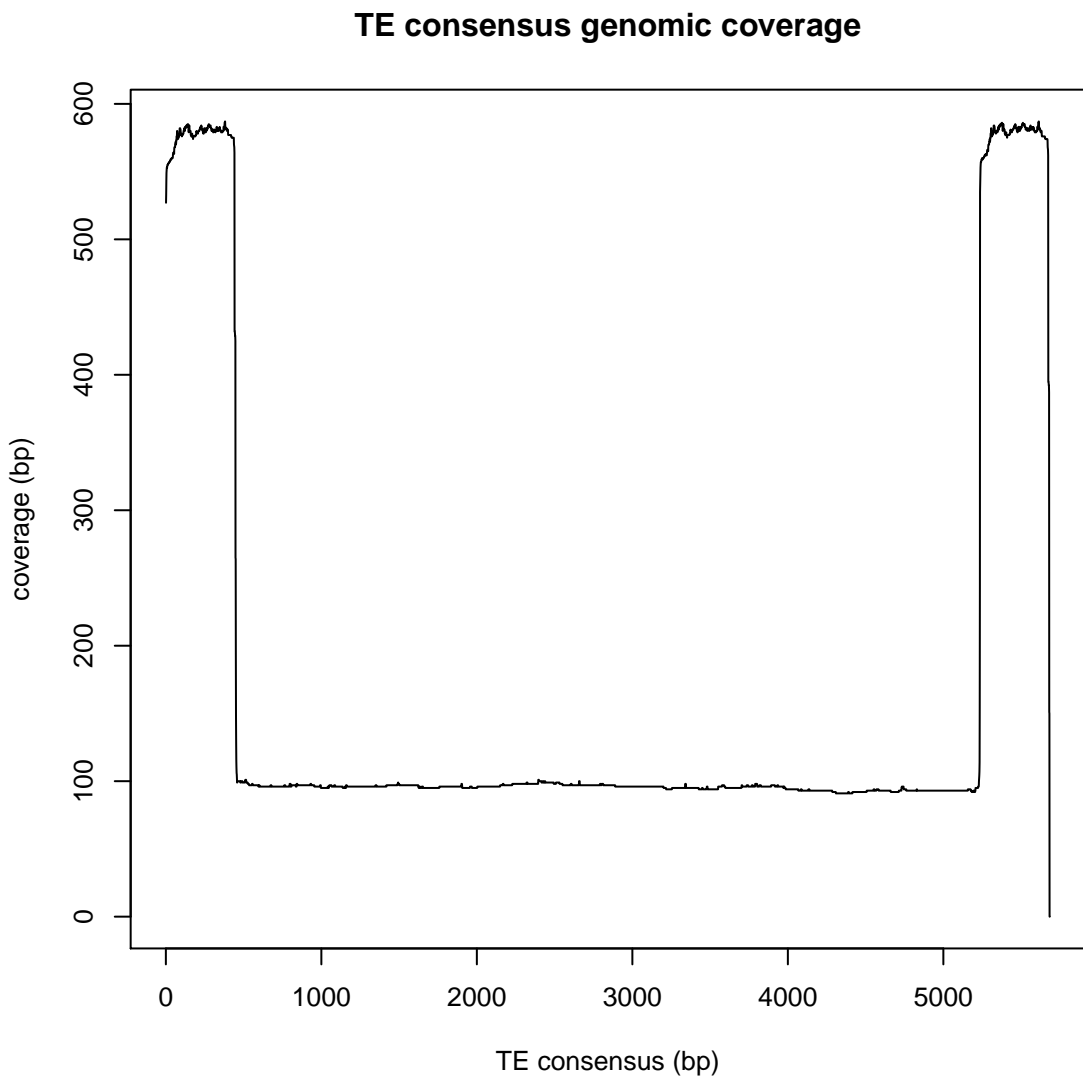
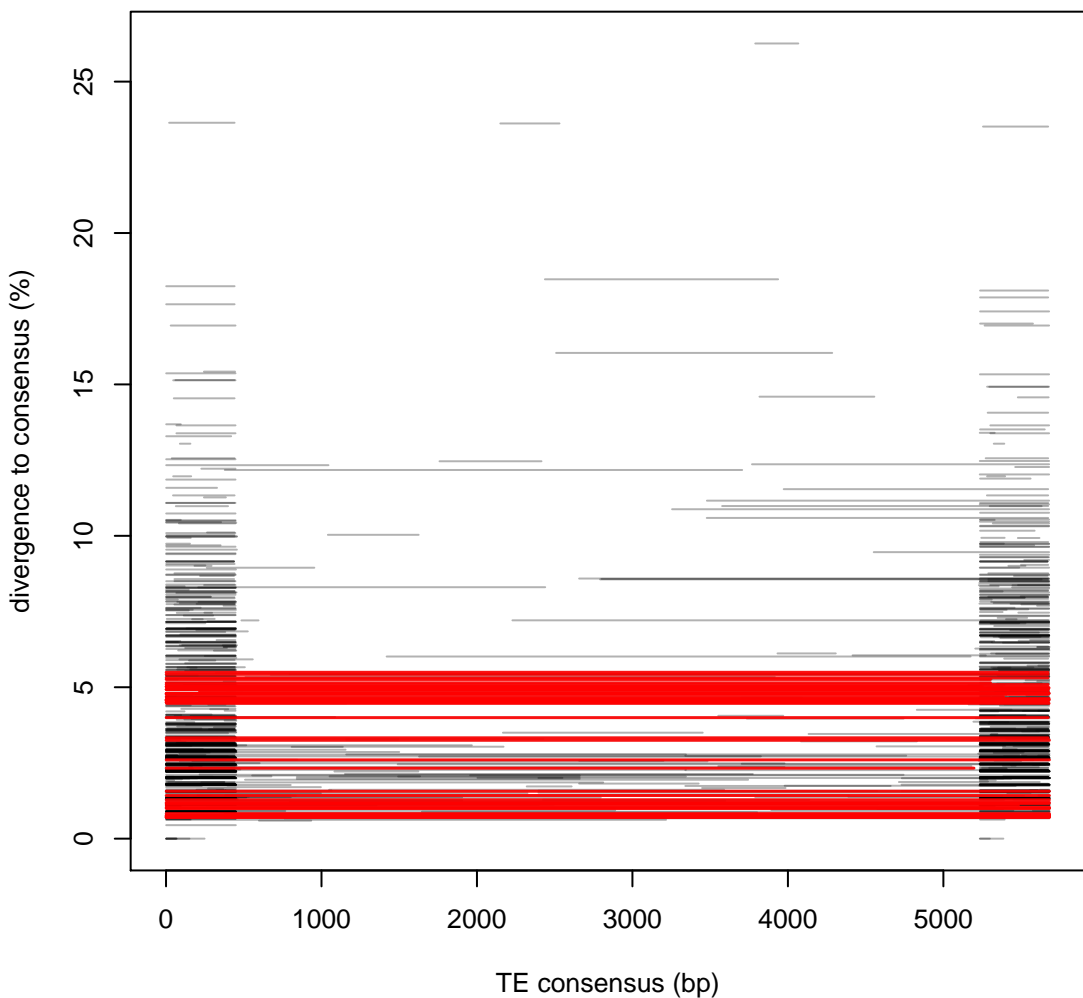
End crop Point



Orf and domain plot

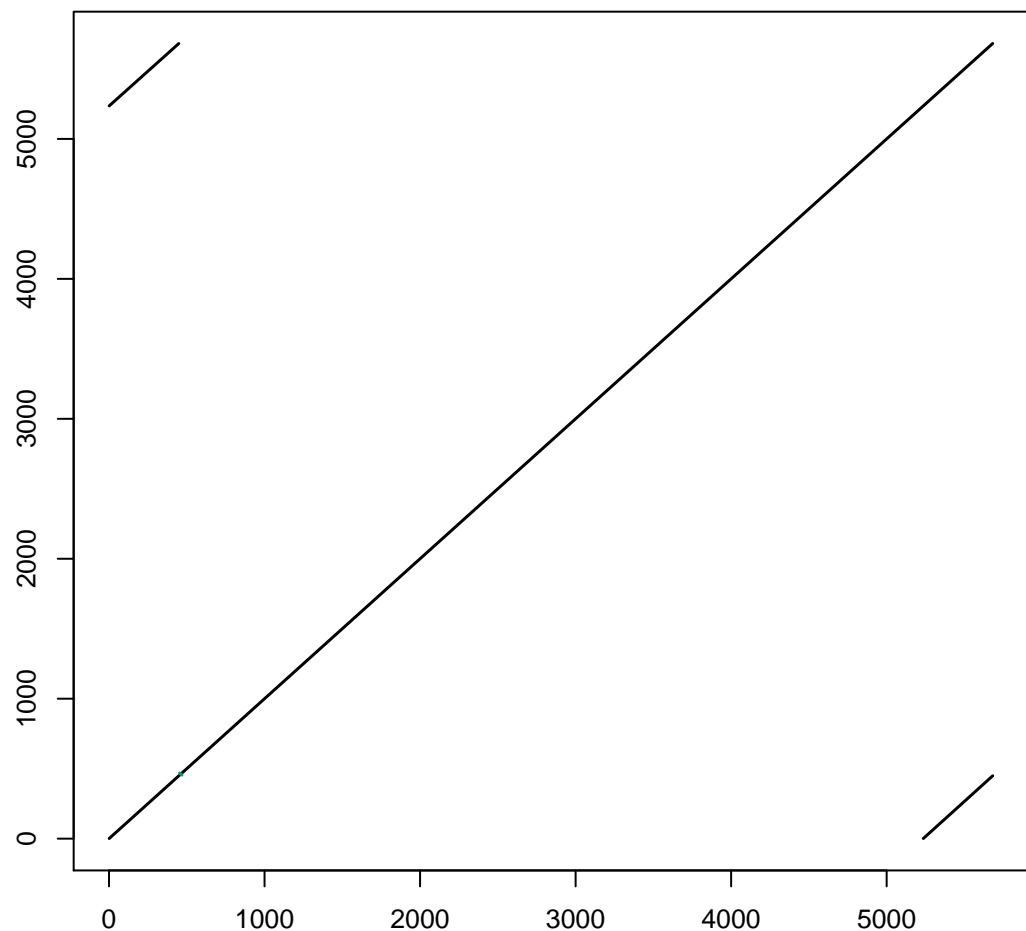


last.bed.uniq.bed.fil.bed_g_2.bed_2000_1000_cl.fa_maf.fa_cl.fa_gap_bl_sim.fa_bou_crop.fa_cr_end_
consensus size: 5684bp; fragments: 1359; full length: 49 (>=5115.6bp)



d.uniq.bed.fil.bed_g_2.bed_2000_1000_cl.fa_maf.fa_cl.fa_gap_bl_sim.fa_bou_crop.fa_cr_end_.fa

TE consensus self dotplot (blastn)



TE consensus structure and protein hits

