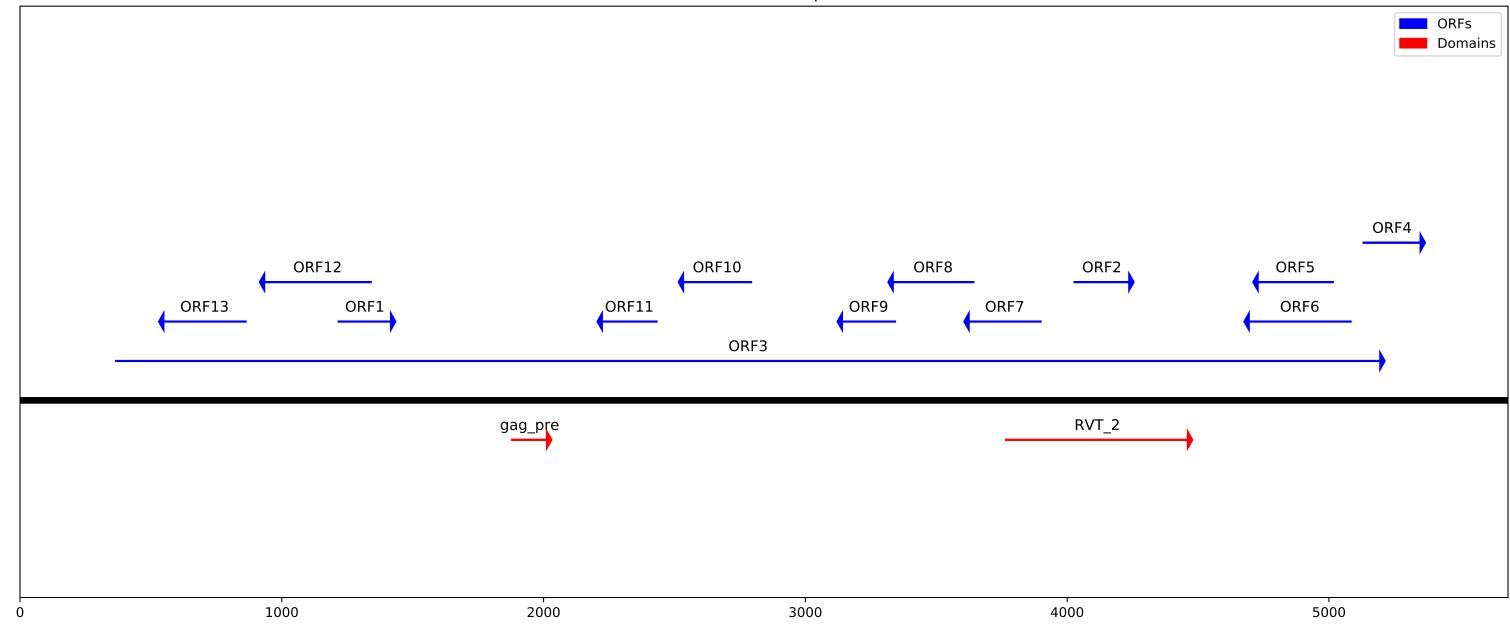
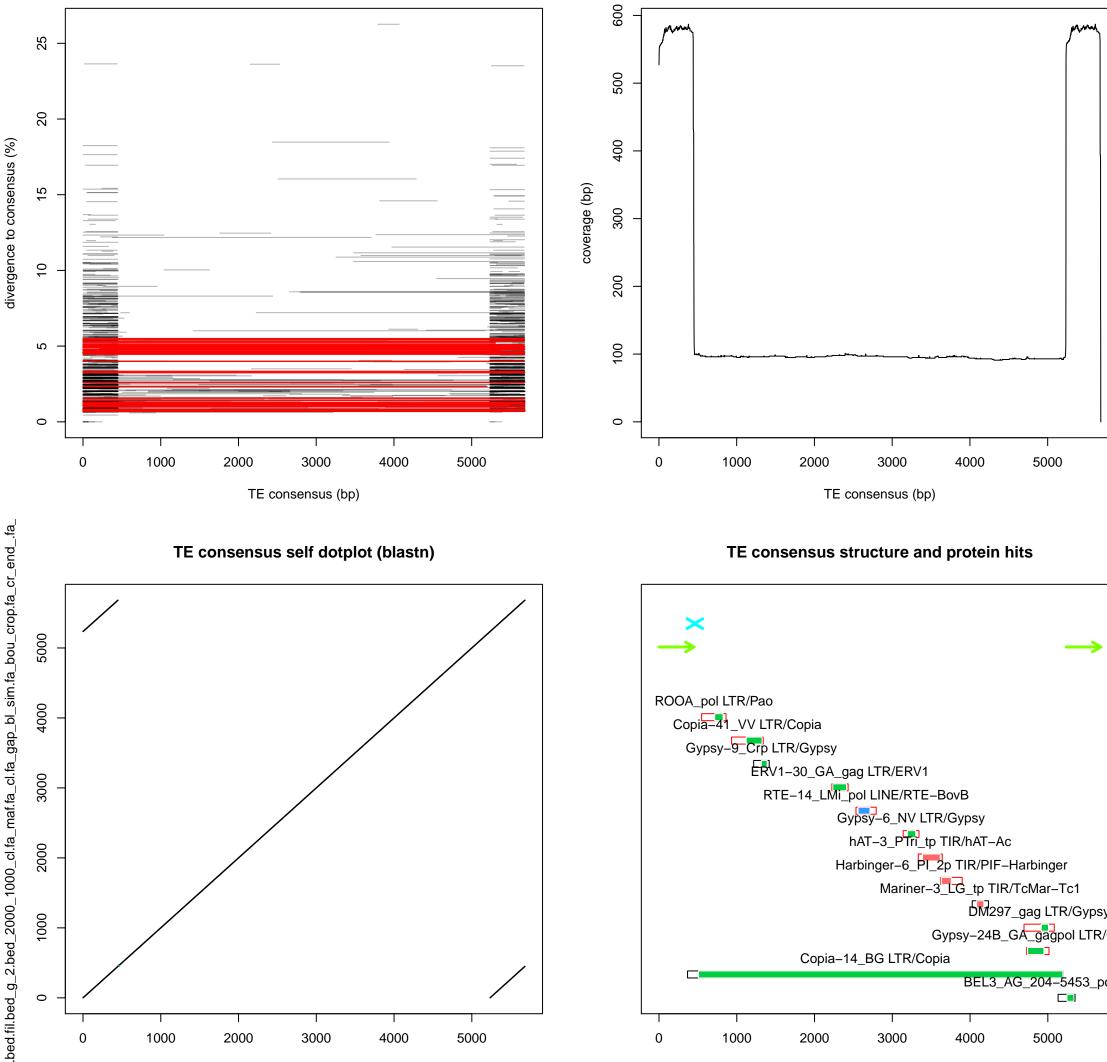
Start crop Point End crop Point MSA length = 5684 - CCAT<mark>C</mark> - - - - - - - - - CAC - TGTTGTGATCAGGCCCACTT - TAGATCACATAACTTGAAATGGCTAGTAT - - - - - - - - ACATCCATTGCGACAAT

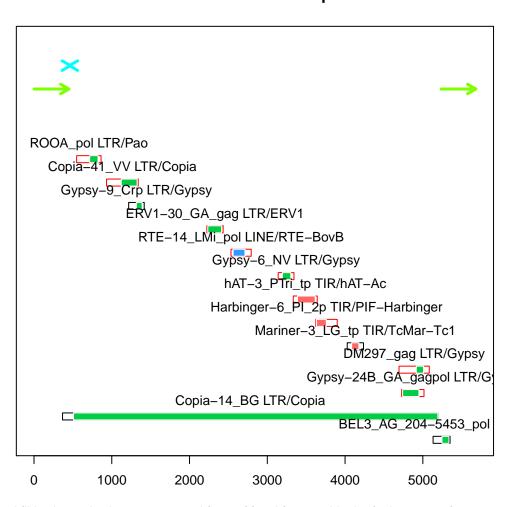


TE consensus genomic coverage



TE consensus self dotplot (blastn)

TE consensus structure and protein hits



e⊈uniq.bed.fil.bed_g_2.bed_2000_1000_cl.fa_maf.fa_cl.fa_gap_bl_sim.fa_bou_crop.fa_cr_end_.fæd.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