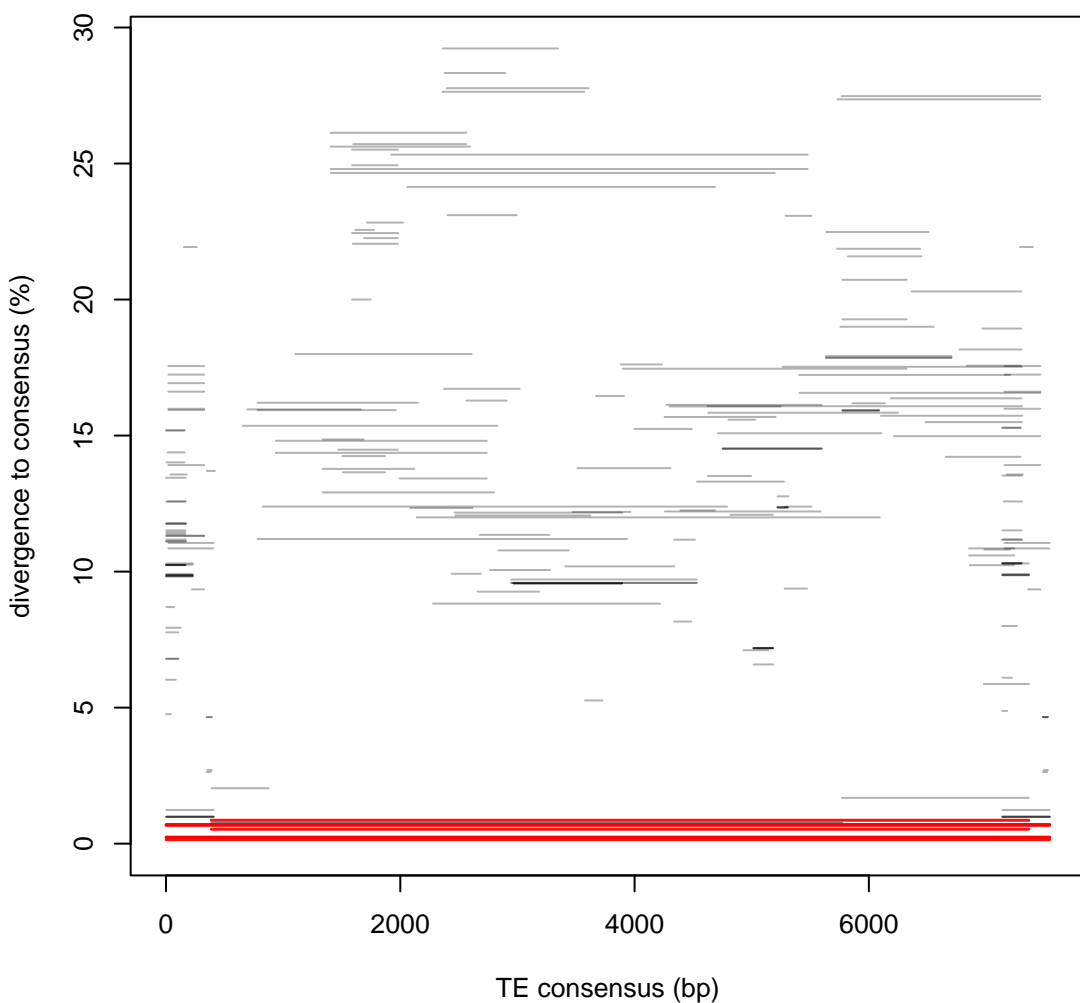
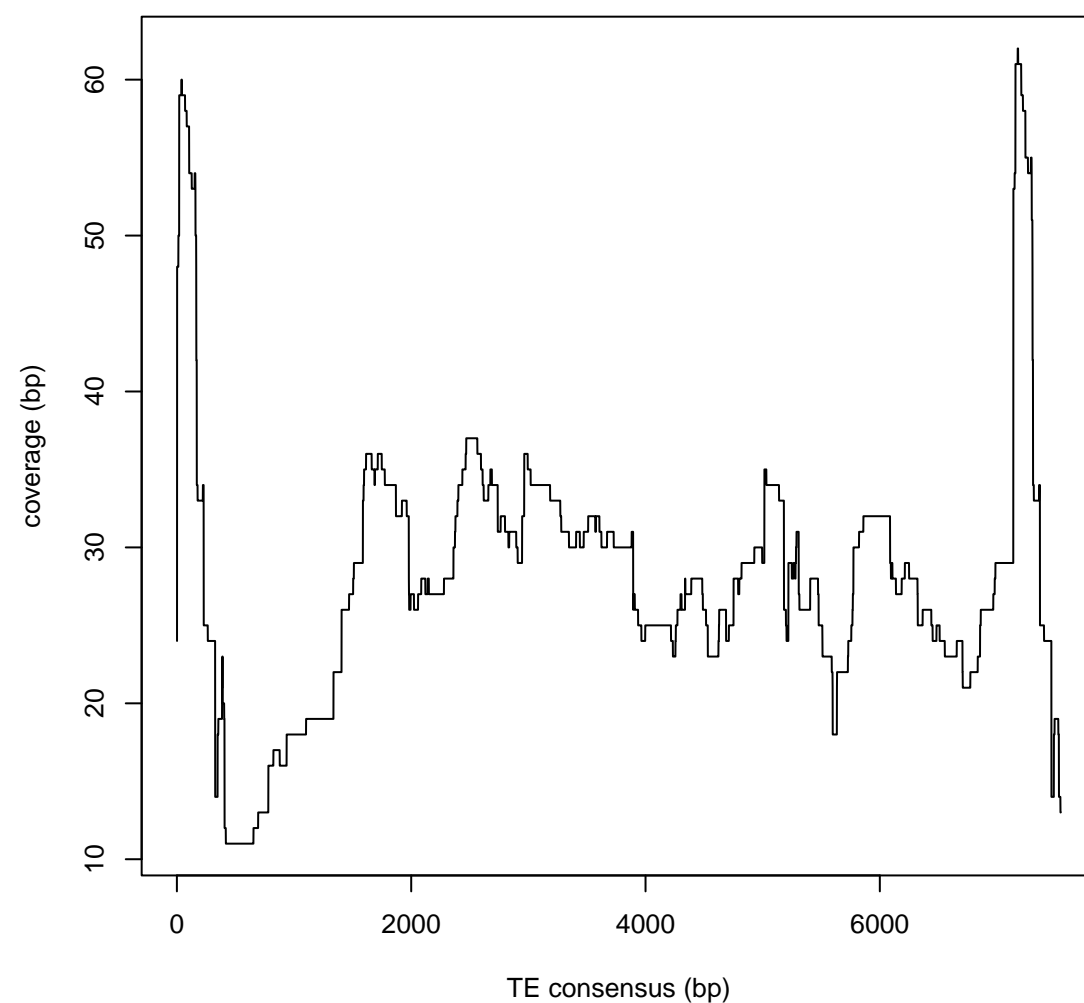


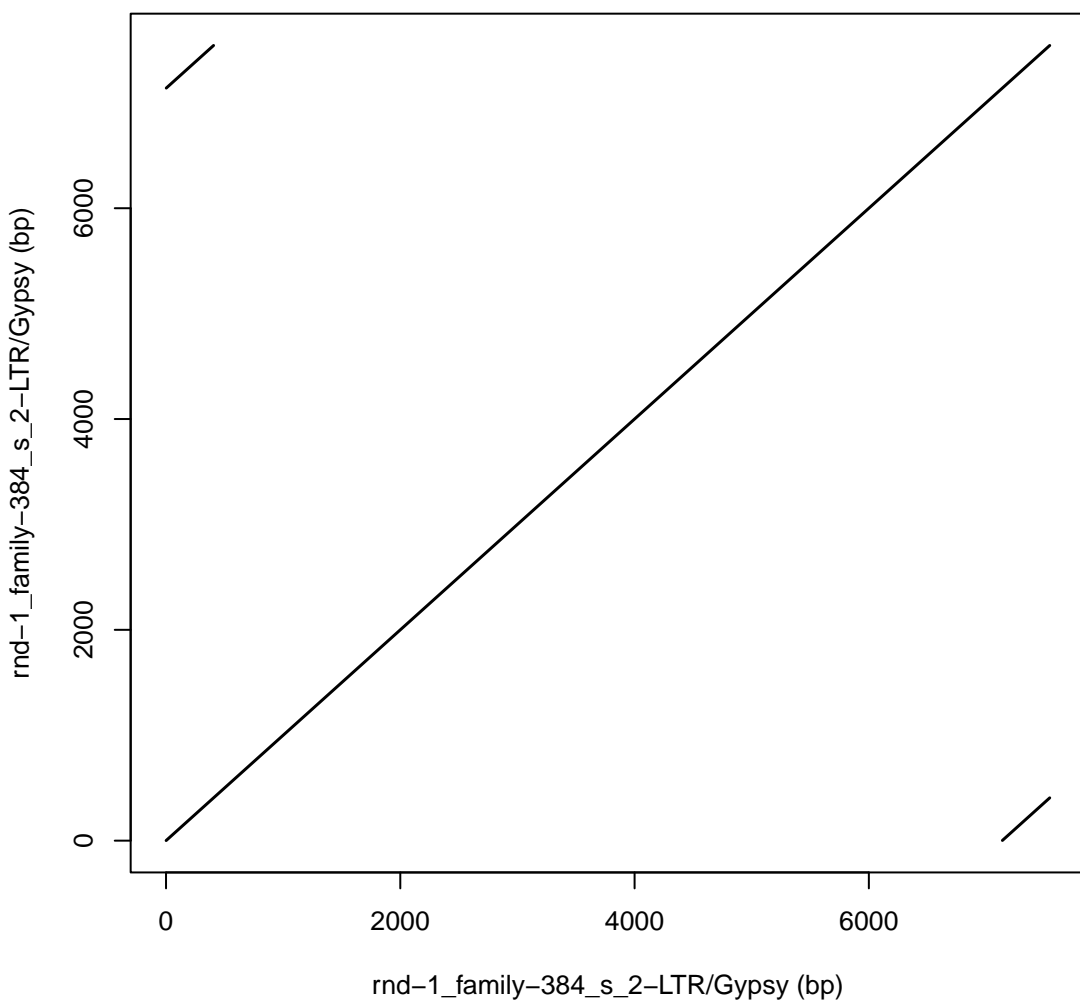
TE: rnd-1_family-384_s_2-LTR/Gypsy
consensus size: 7544bp; fragments: 240; full length: 8 (≥ 6789.6 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

