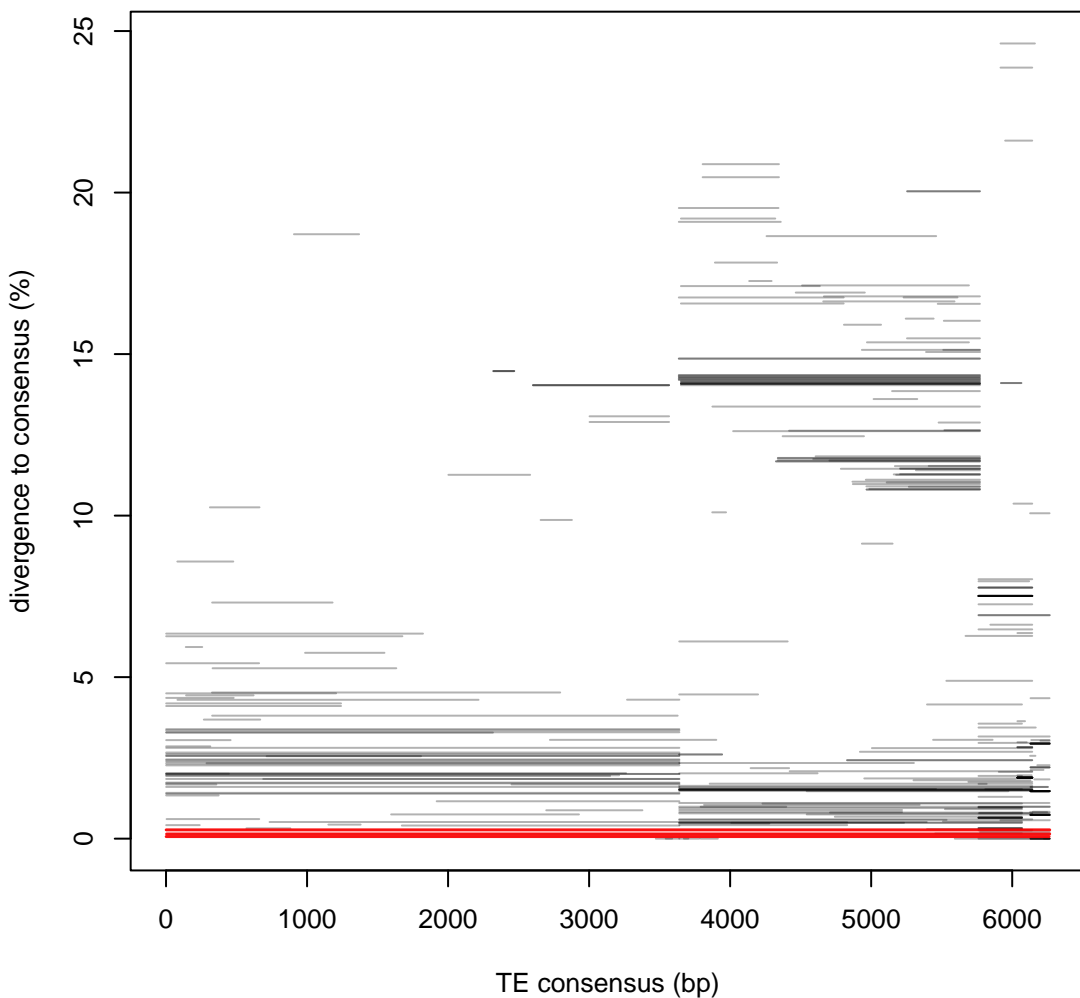
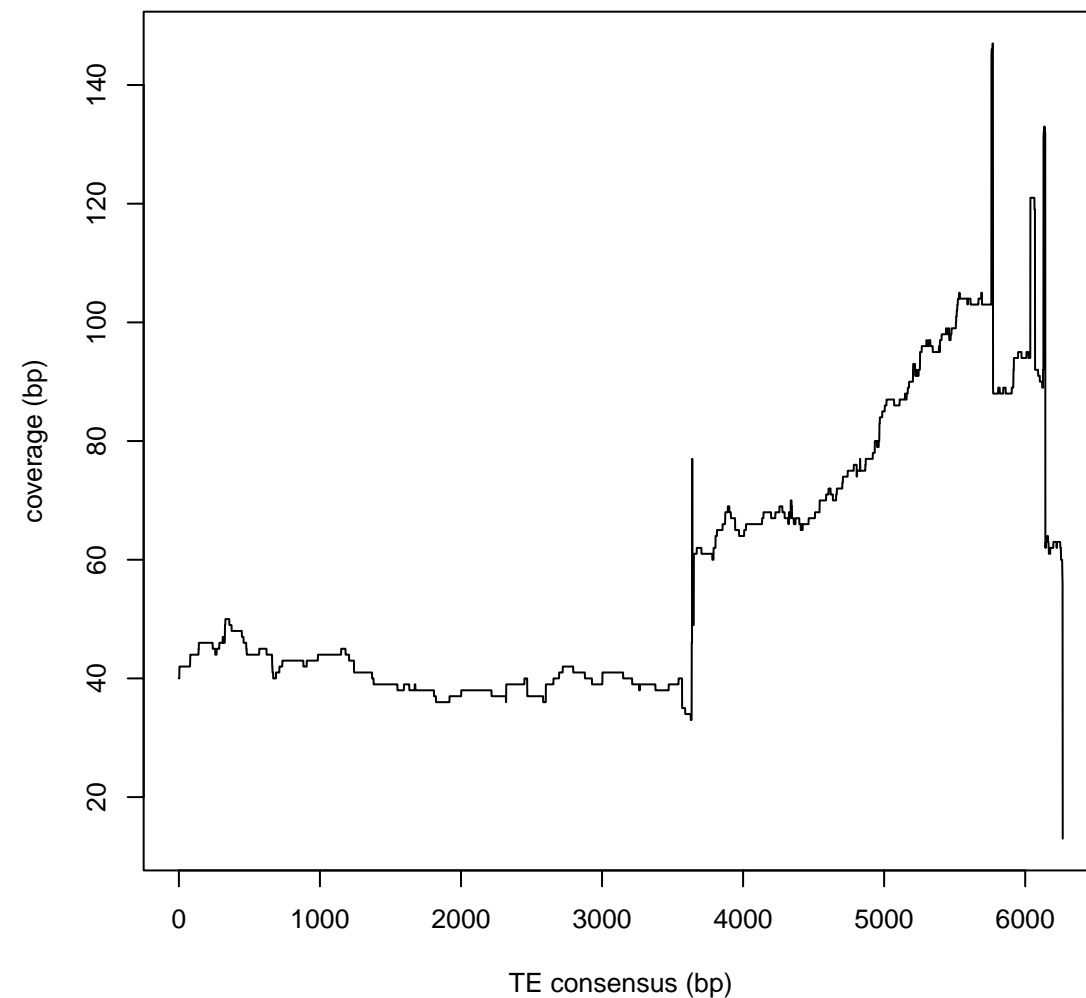


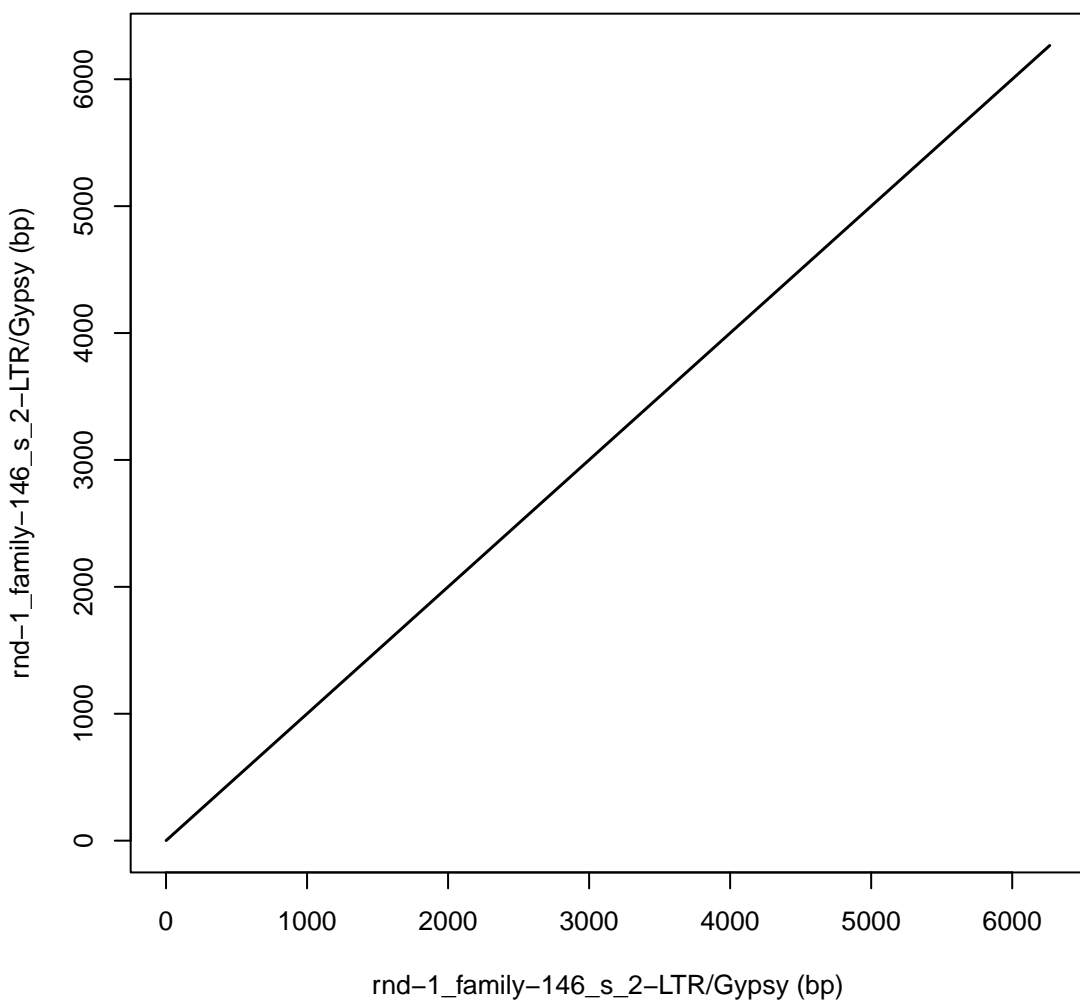
TE: rnd-1_family-146_s_2-LTR/Gypsy
consensus size: 6266bp; fragments: 353; full length: 3 (≥ 5639.4 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

