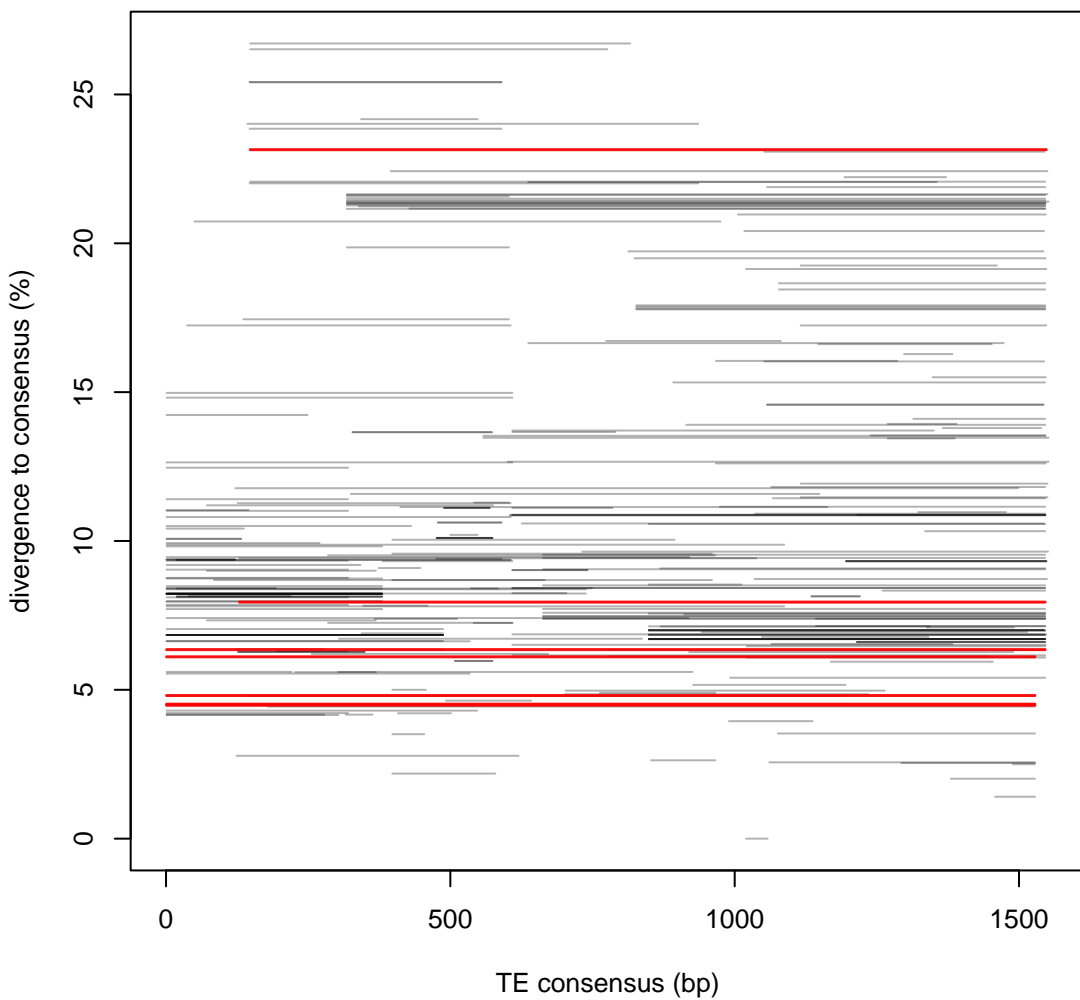
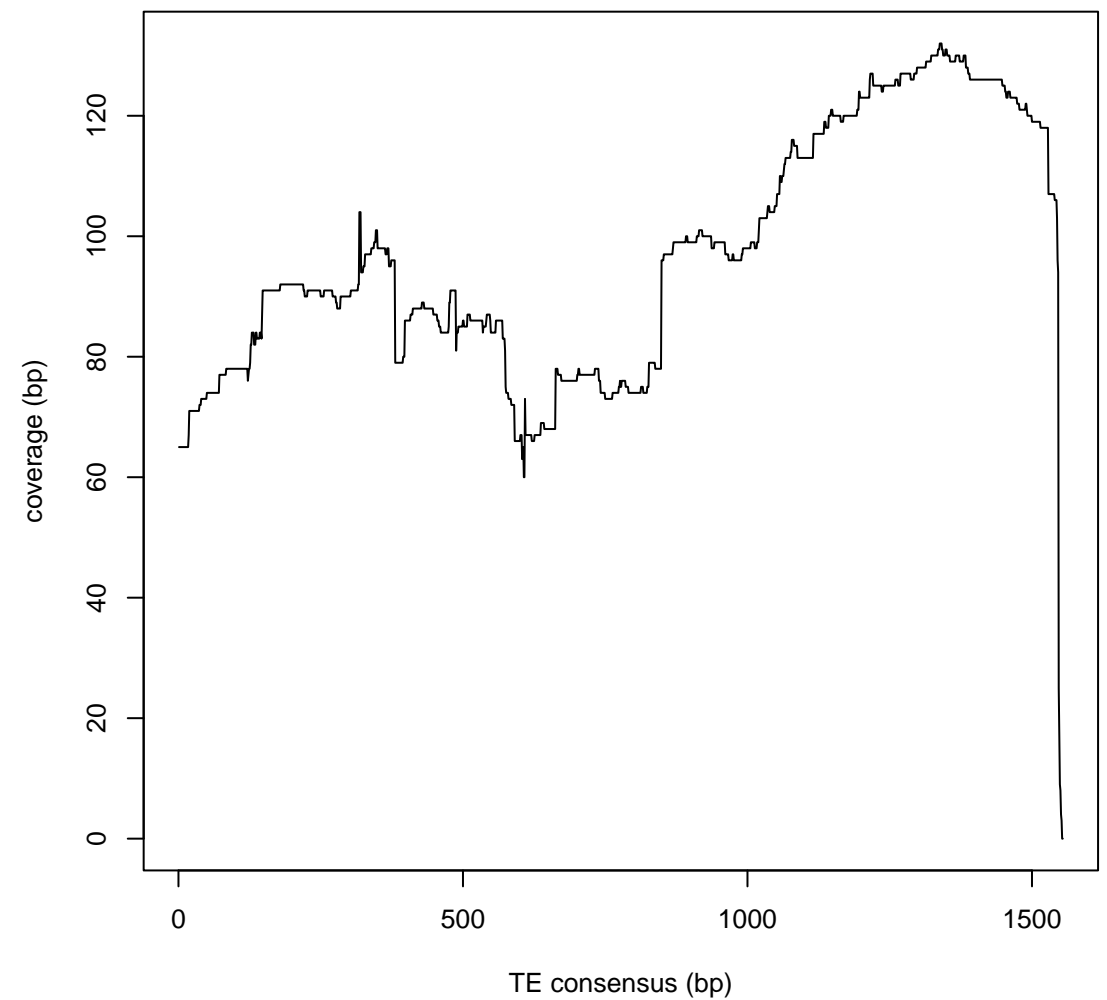


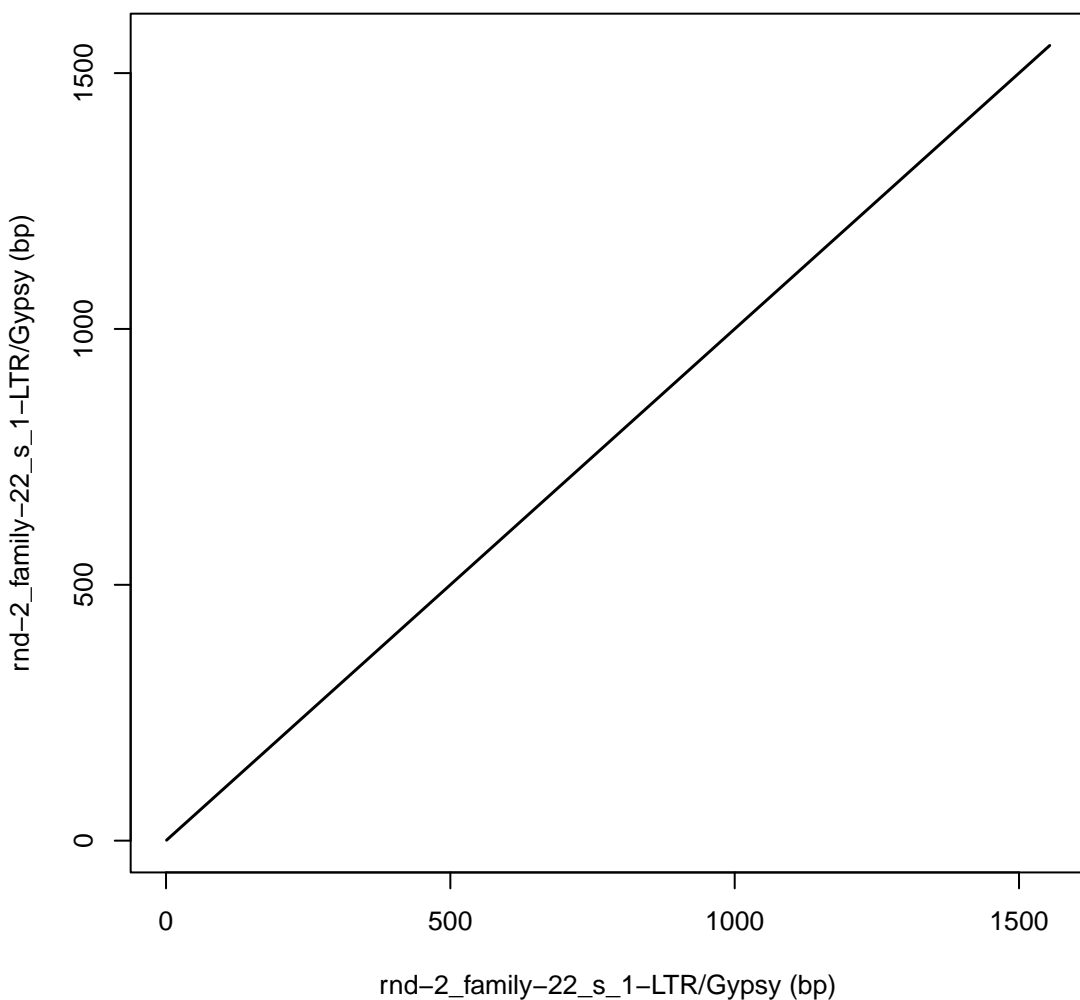
TE: rnd-2_family-22_s_1-LTR/Gypsy
consensus size: 1554bp; fragments: 288; full length: 7 (≥ 1398.6 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

