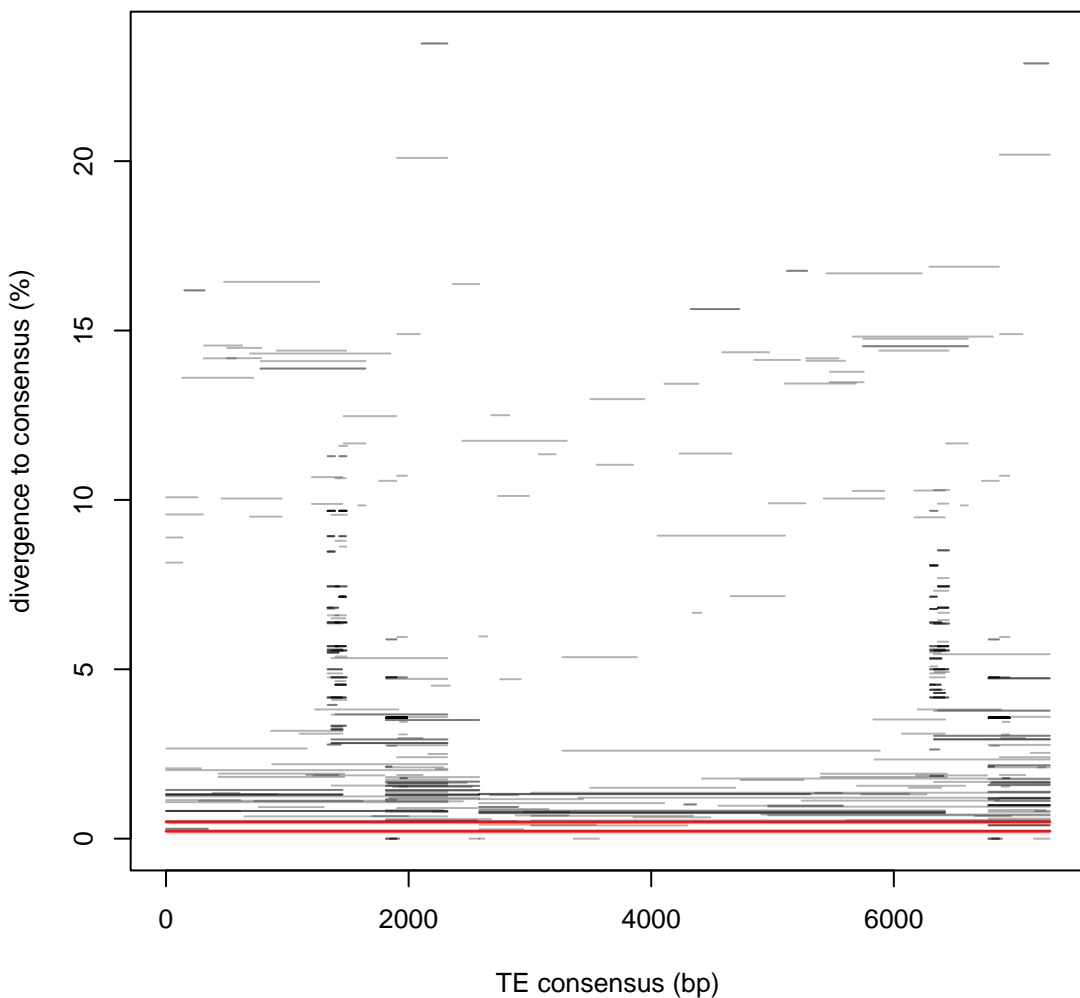
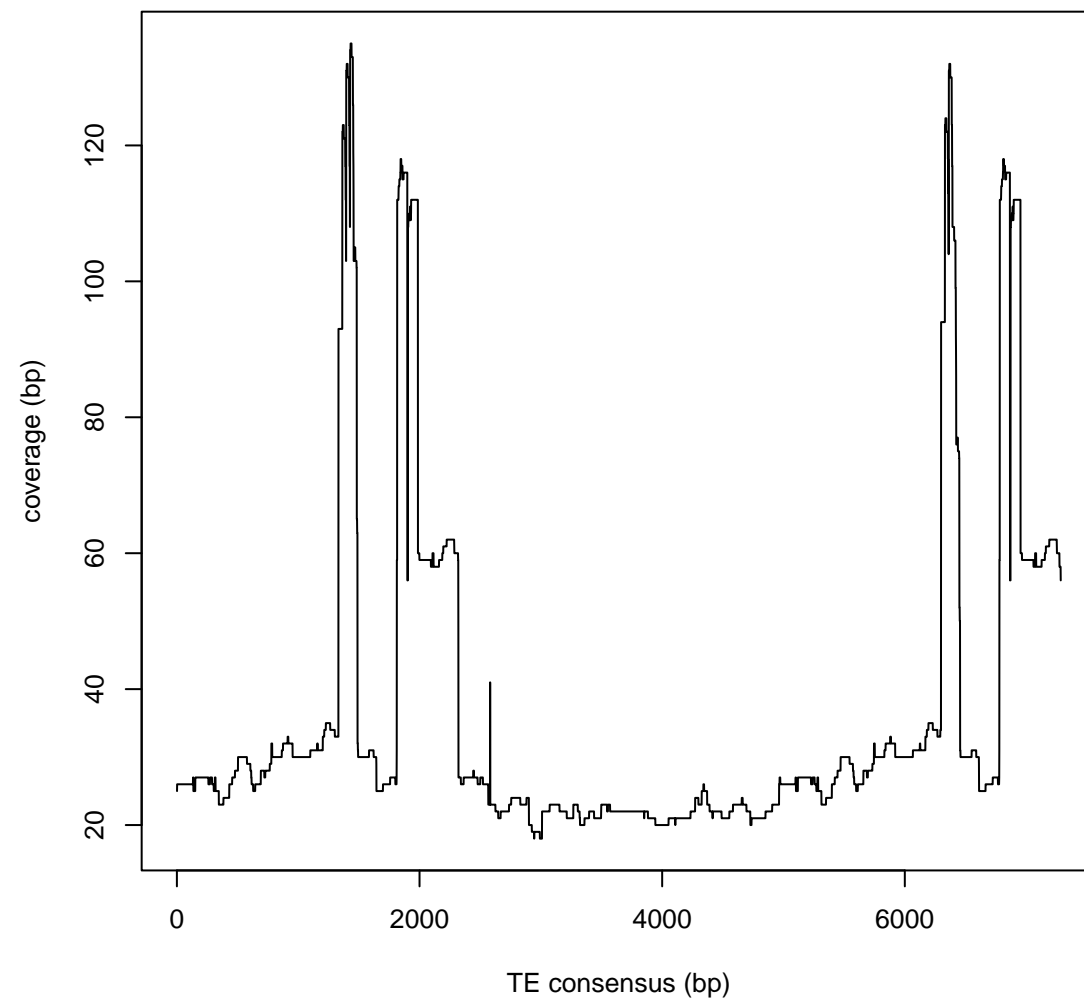


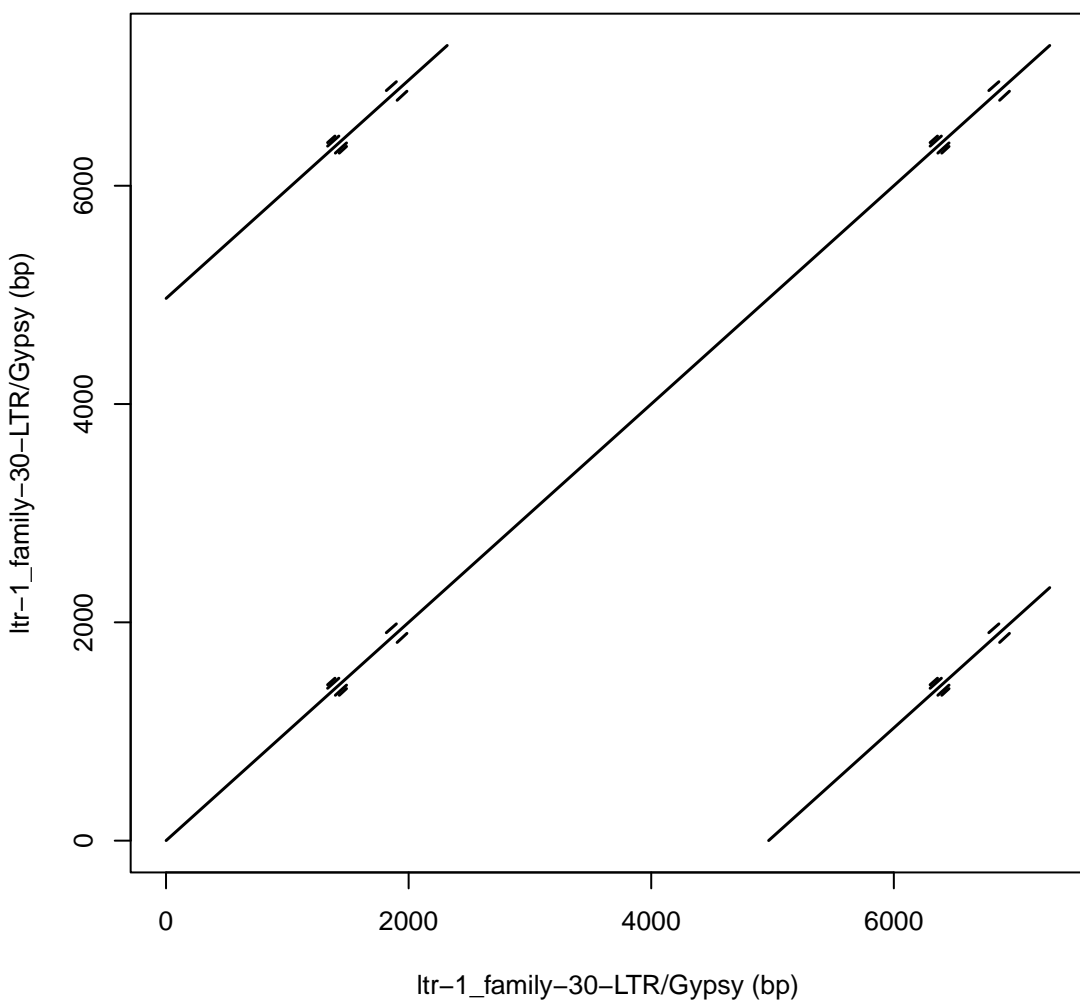
TE: ltr-1_family-30-LTR/Gypsy
consensus size: 7285bp; fragments: 735; full length: 2 (≥ 6556.5 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

