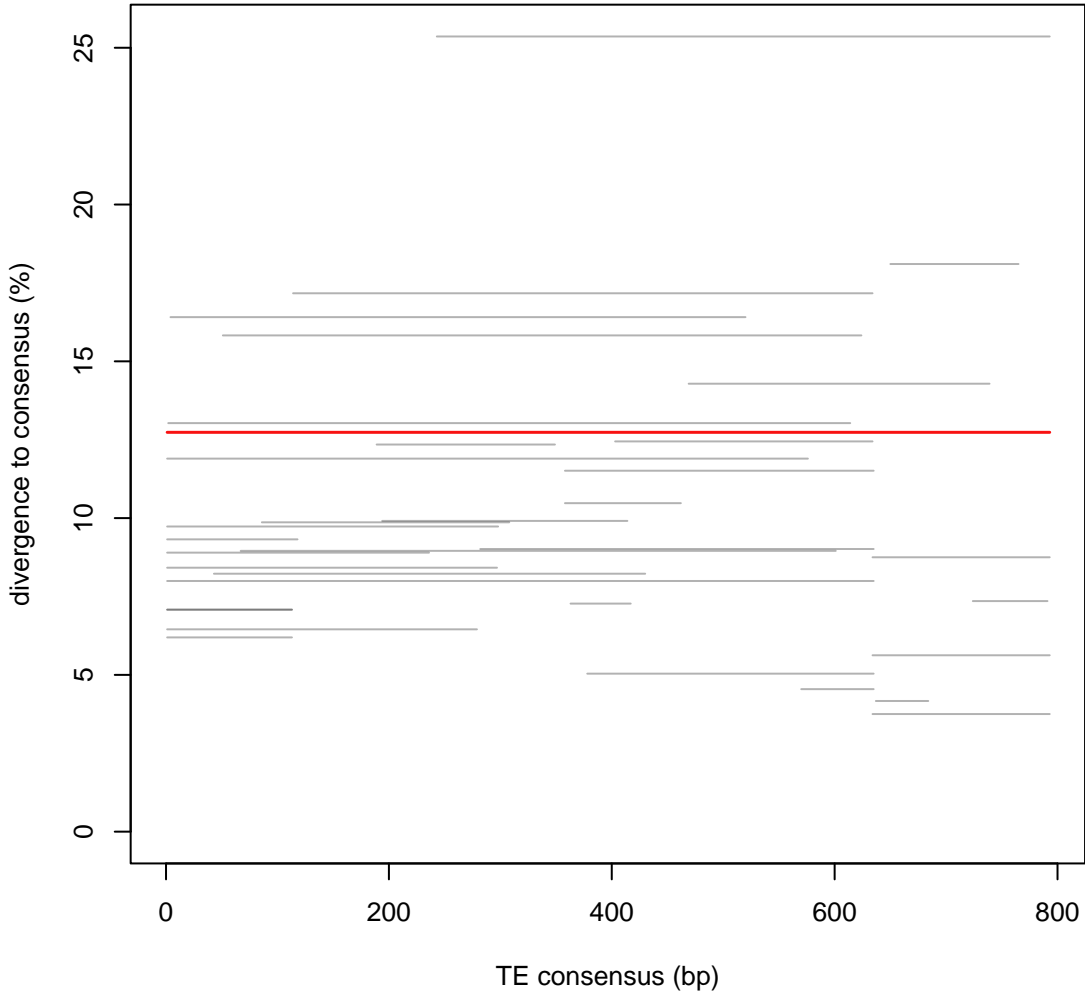
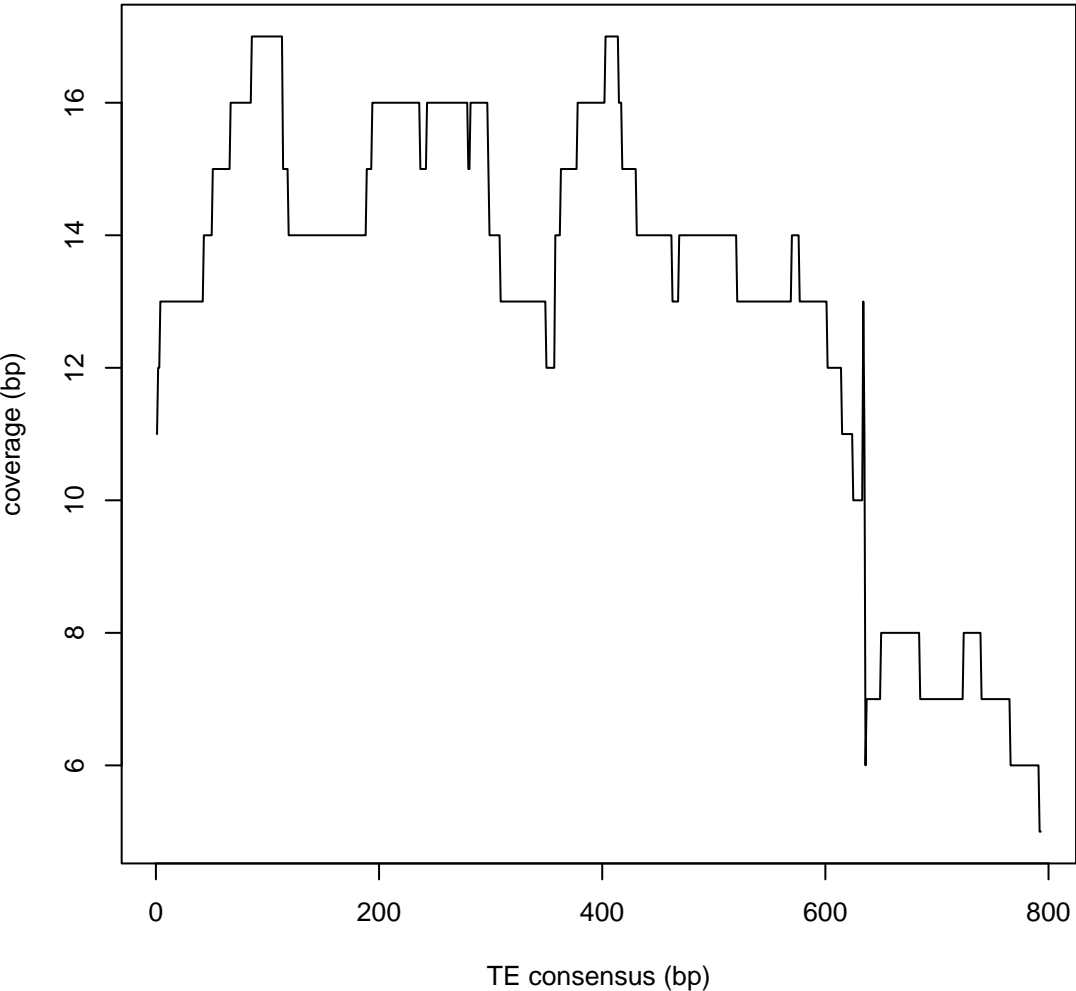


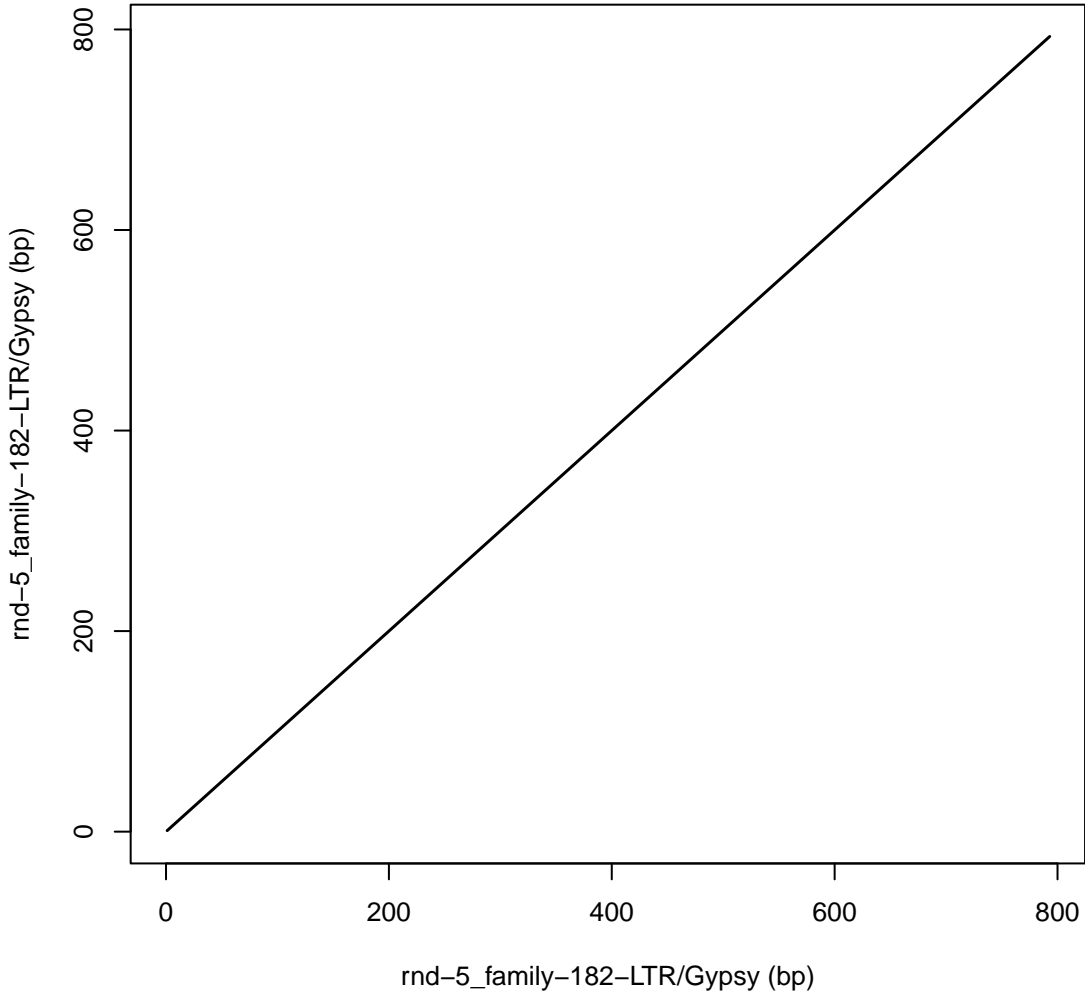
TE: rnd-5_family-182-LTR/Gypsy
consensus size: 793bp; fragments: 35; full length: 1 (≥ 713.7 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

