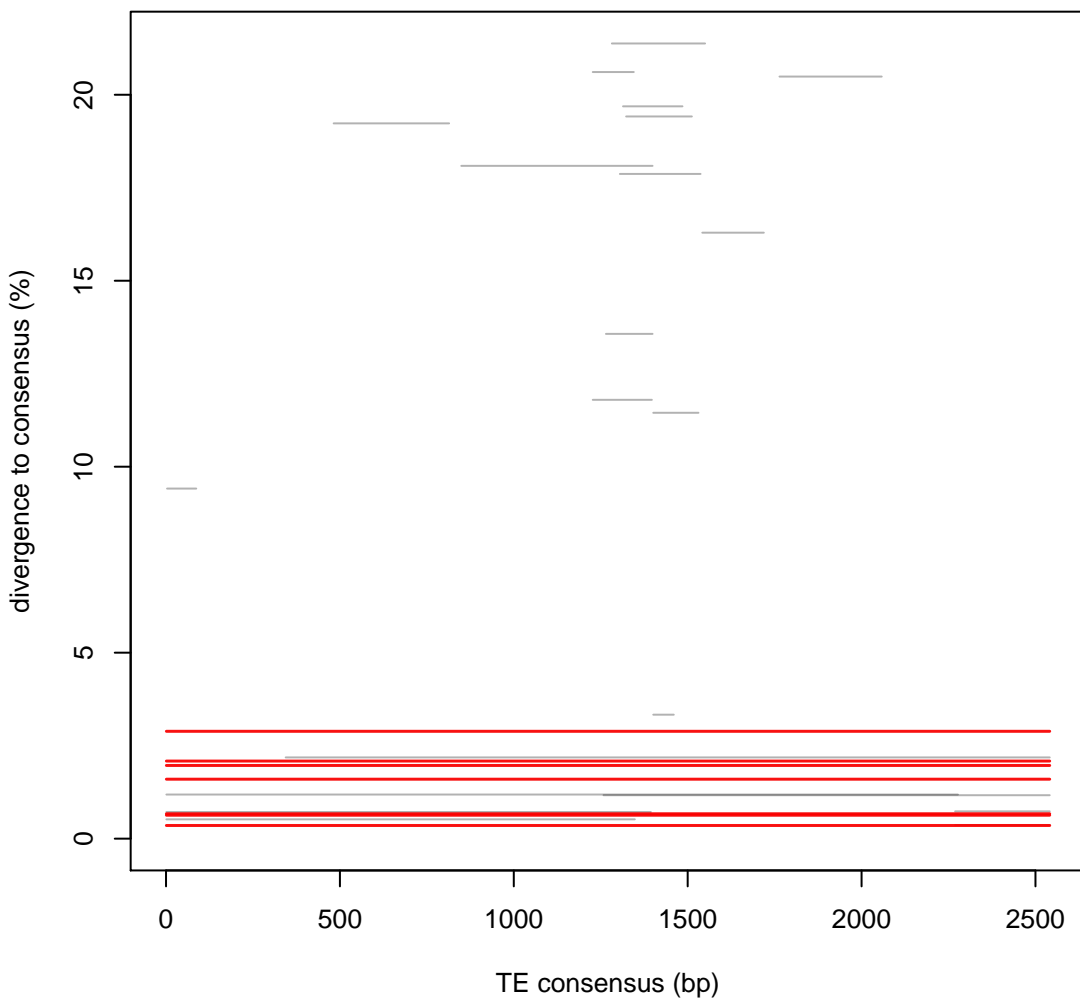
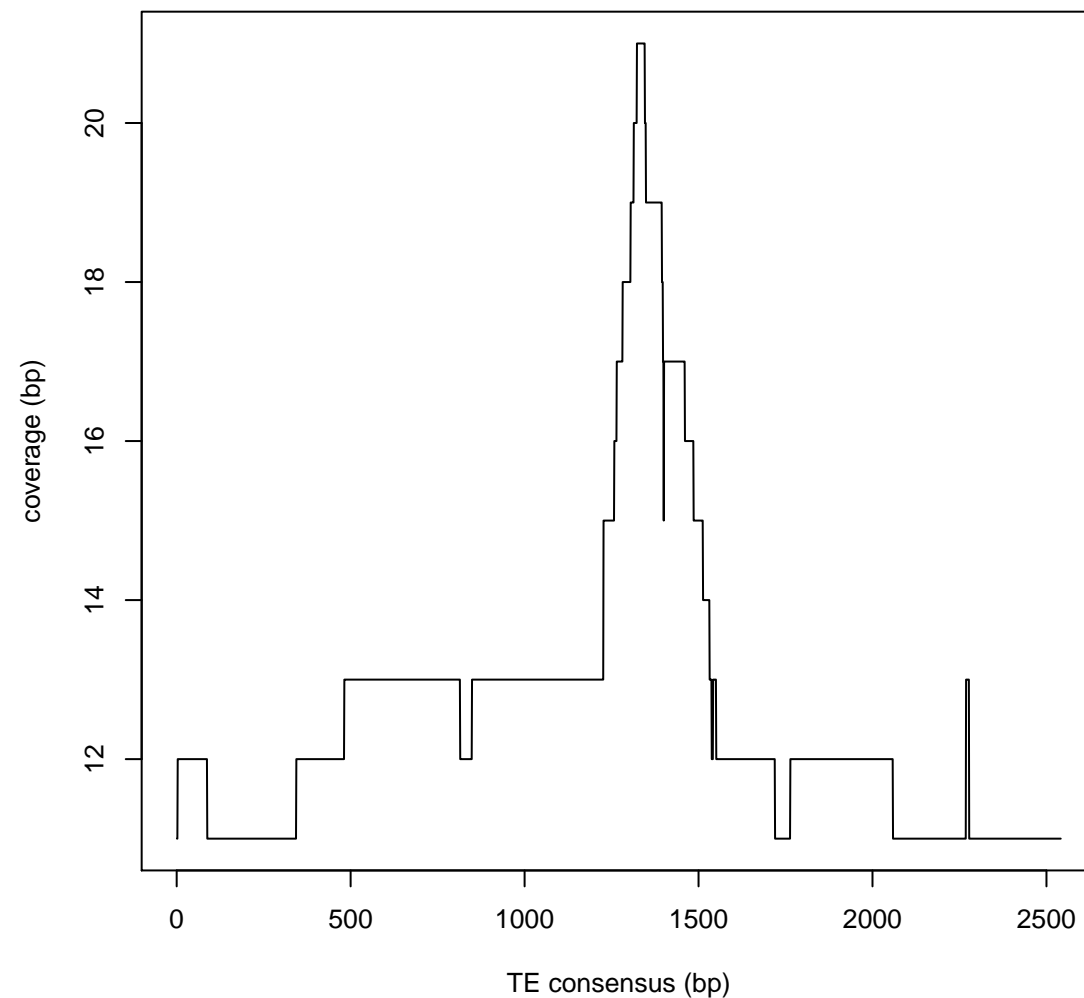


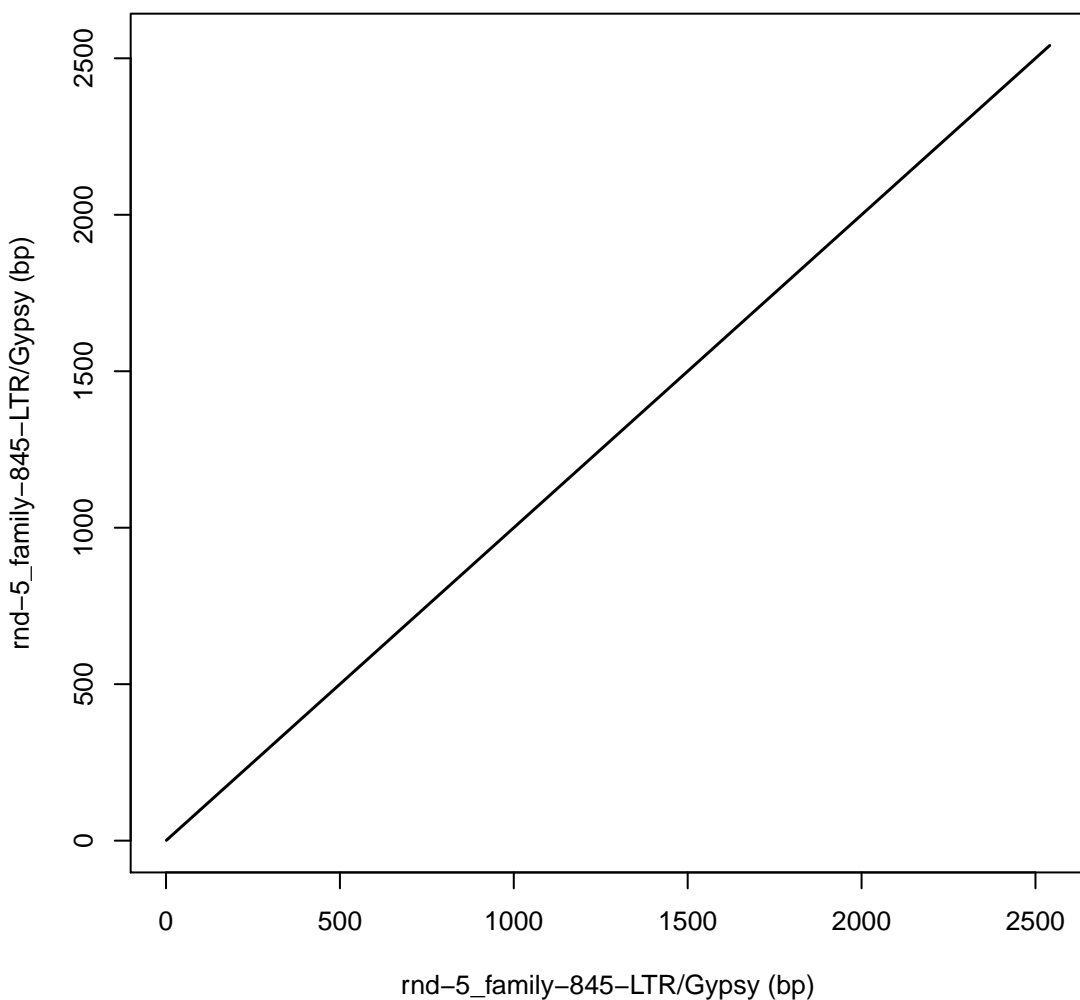
TE: rnd-5_family-845-LTR/Gypsy
consensus size: 2541bp; fragments: 29; full length: 7 (≥ 2286.9 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

