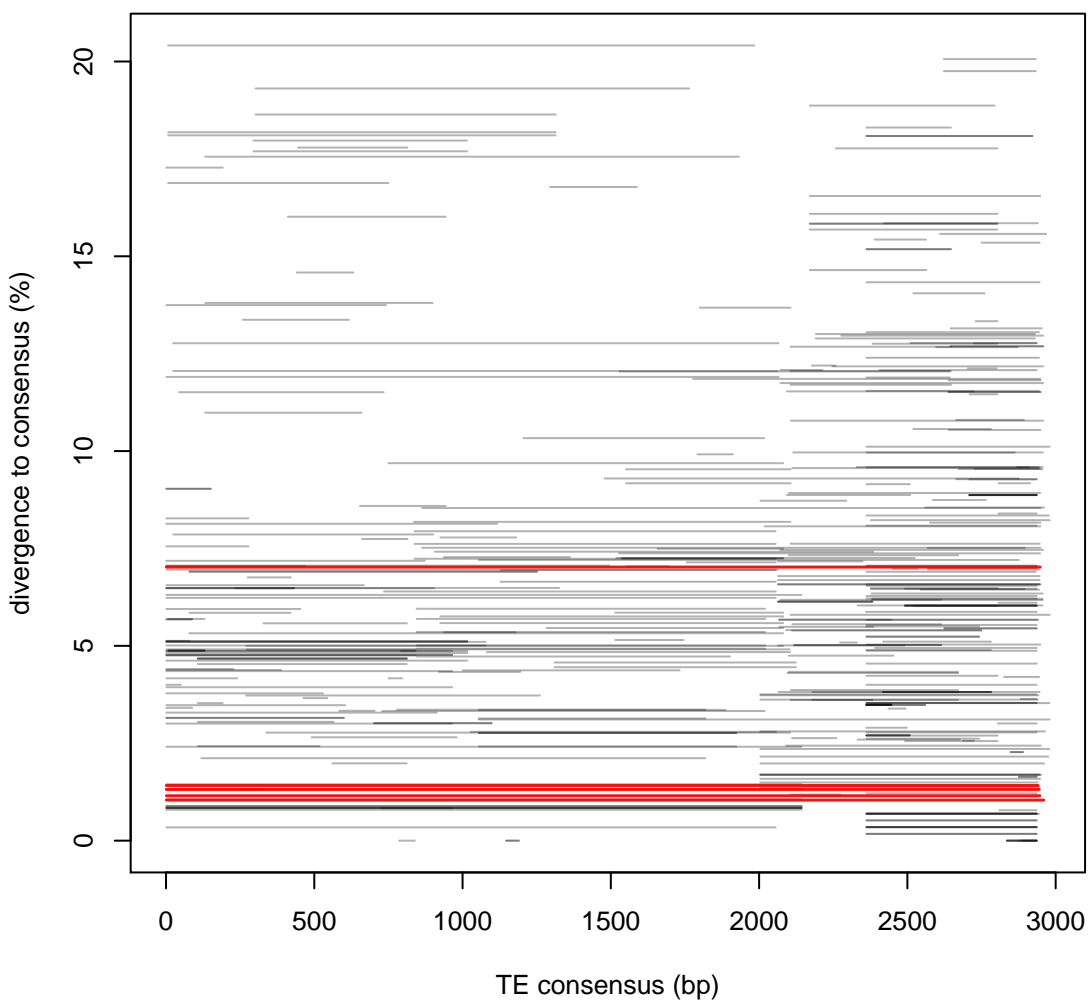
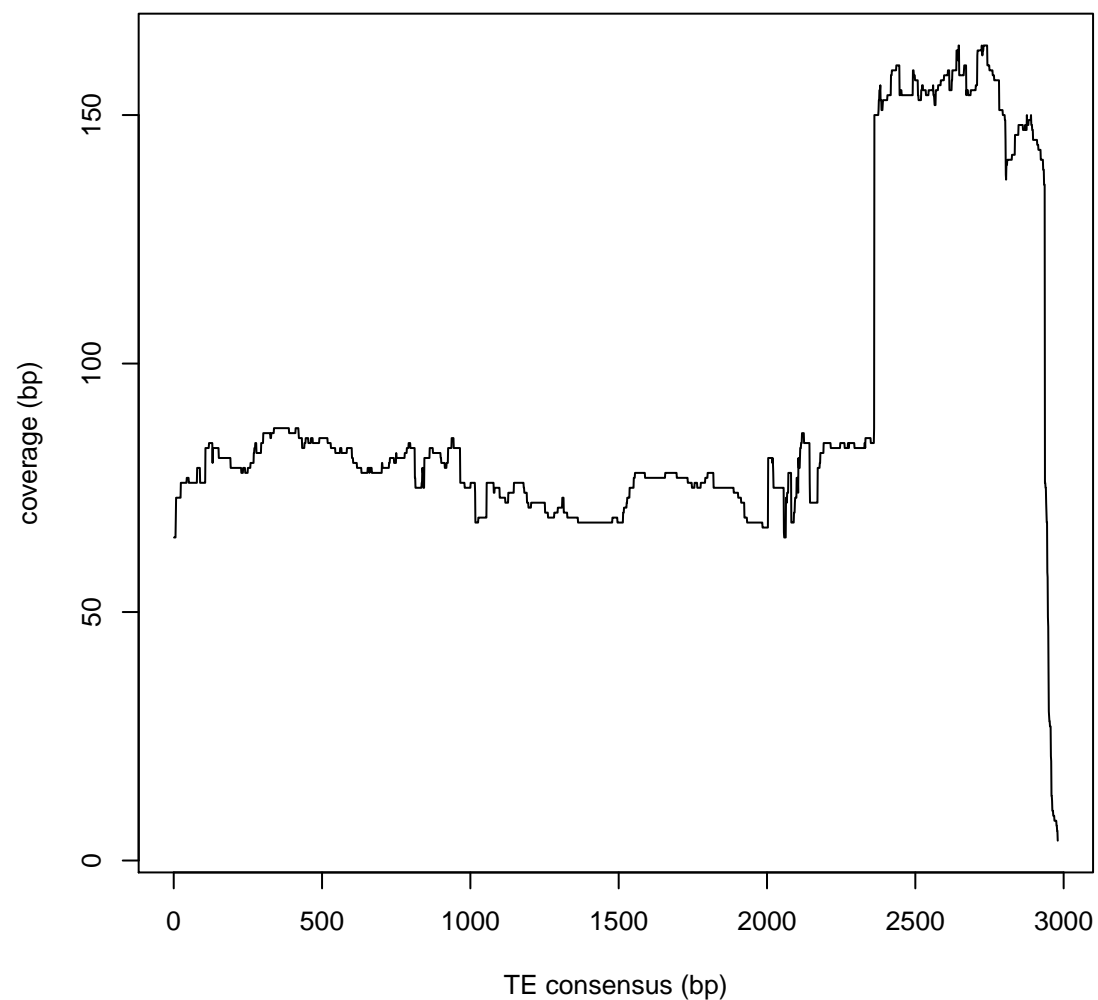


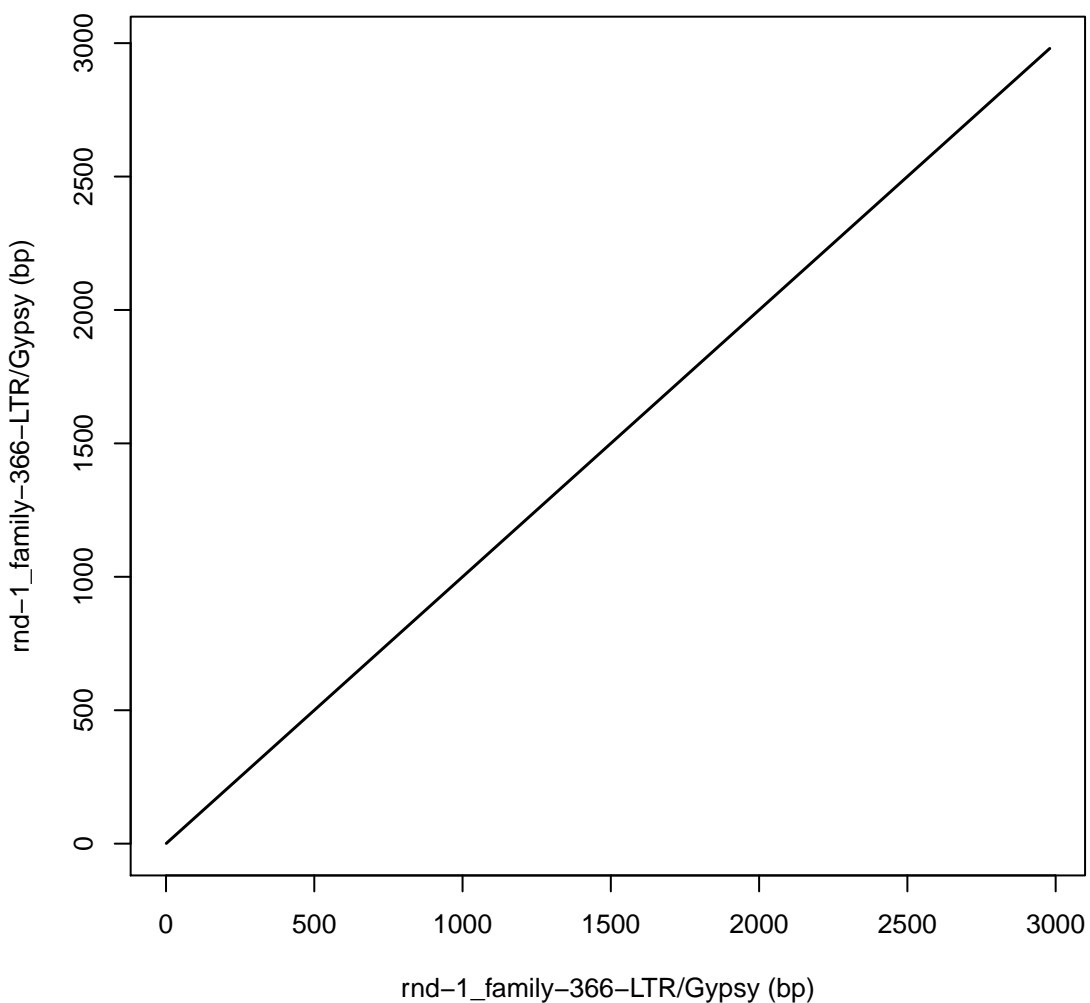
TE: rnd-1_family-366-LTR/Gypsy
consensus size: 2980bp; fragments: 401; full length: 5 (≥ 2682 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

