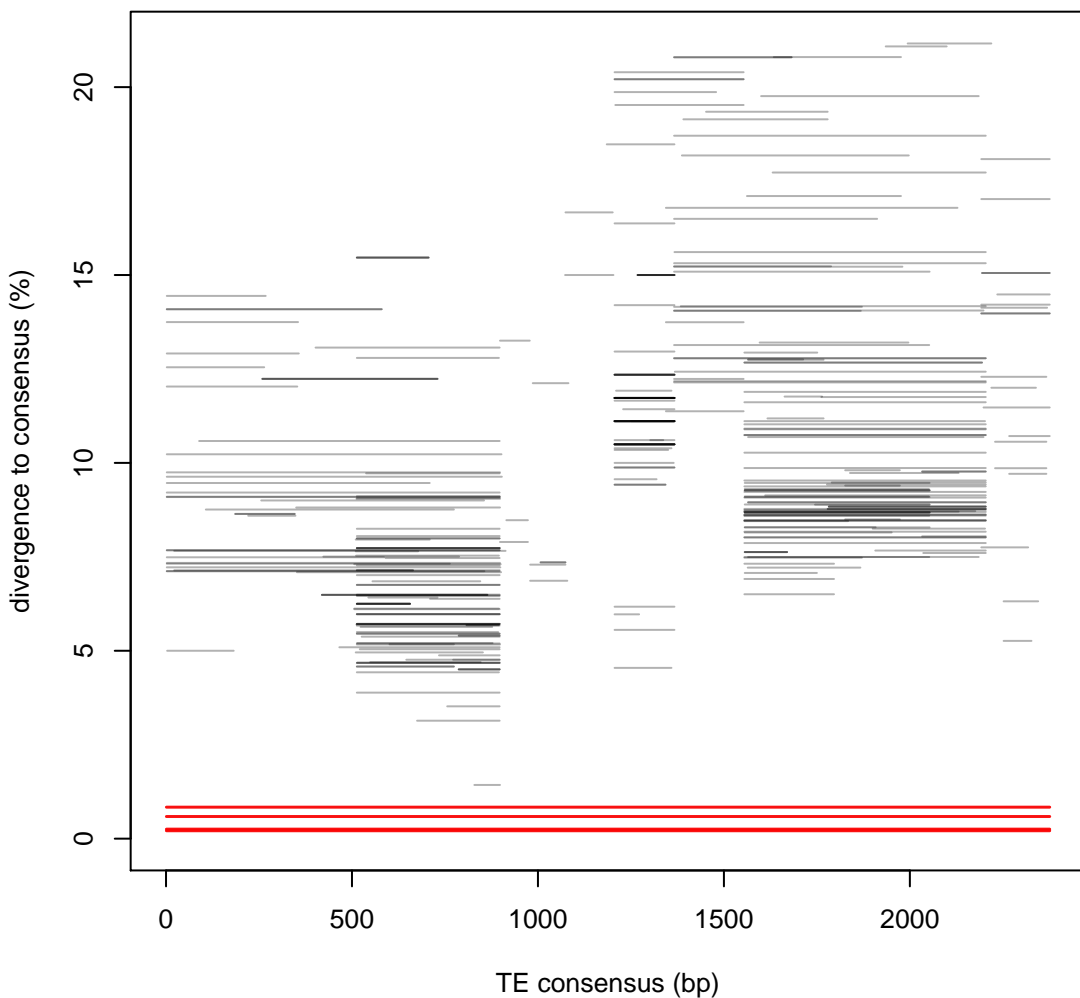
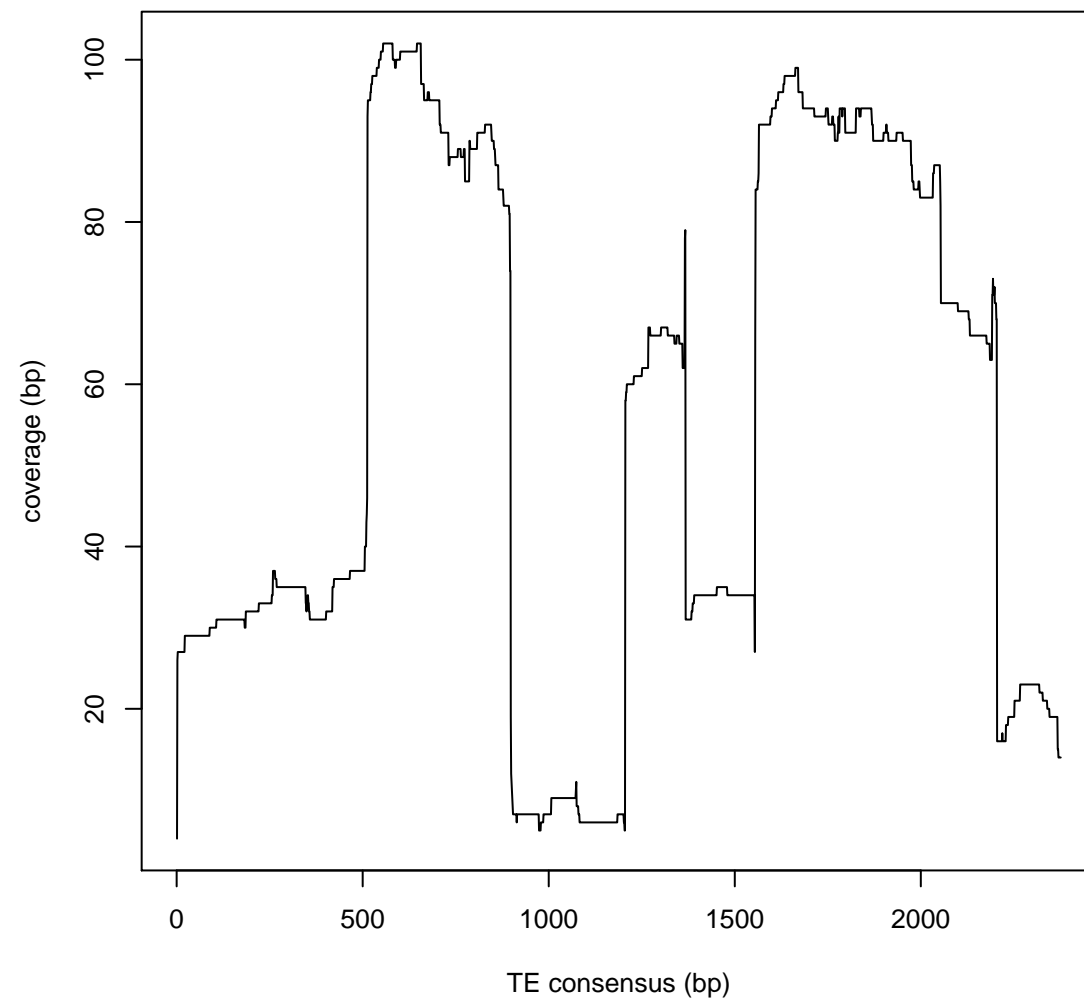


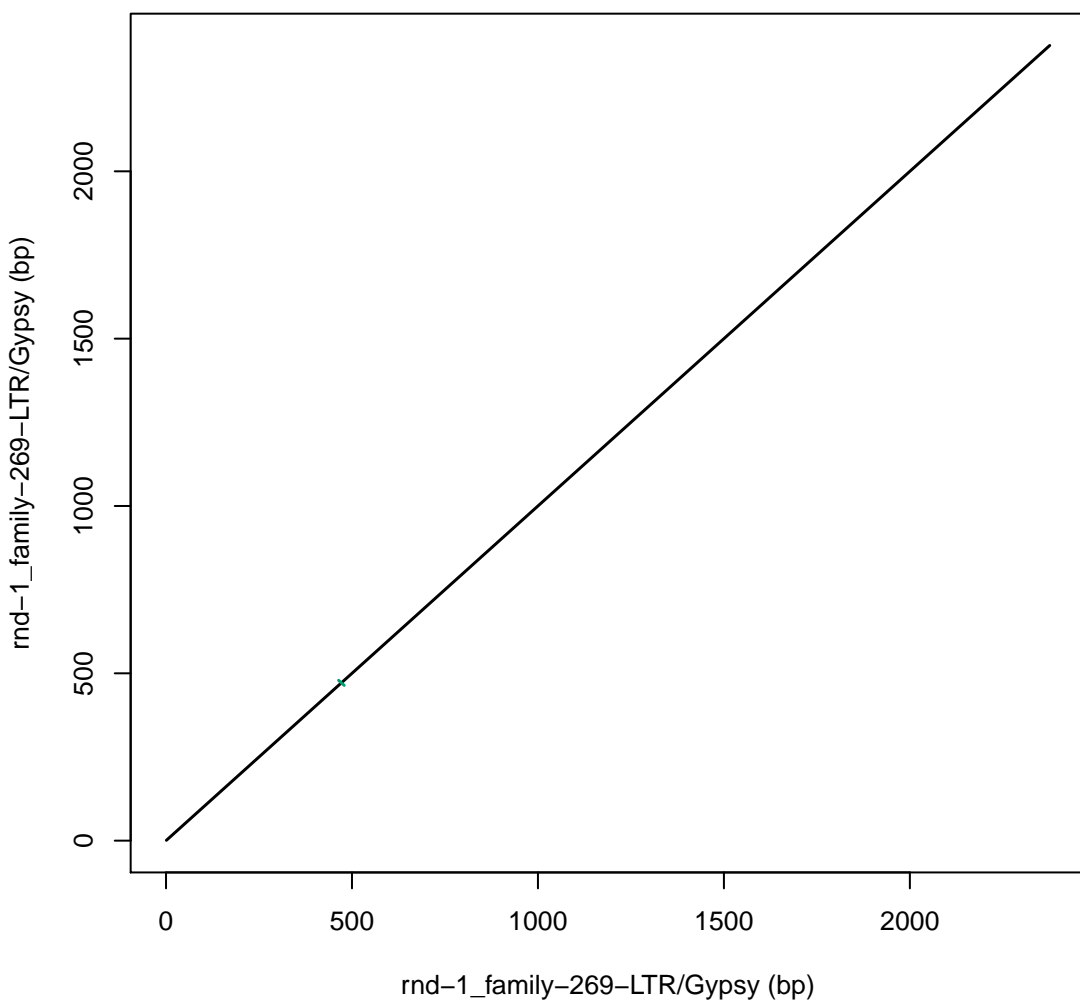
TE: rnd-1_family-269-LTR/Gypsy
consensus size: 2376bp; fragments: 339; full length: 4 (≥ 2138.4 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

