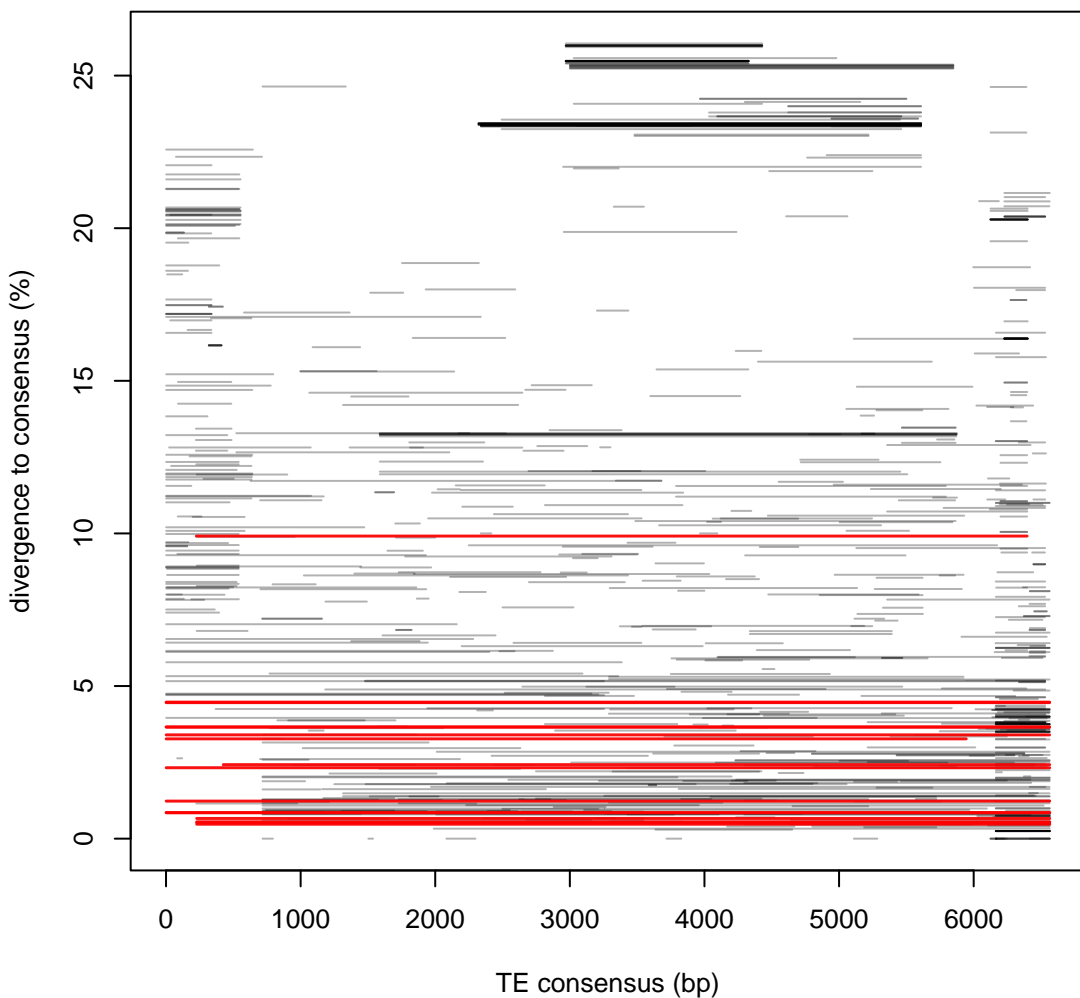
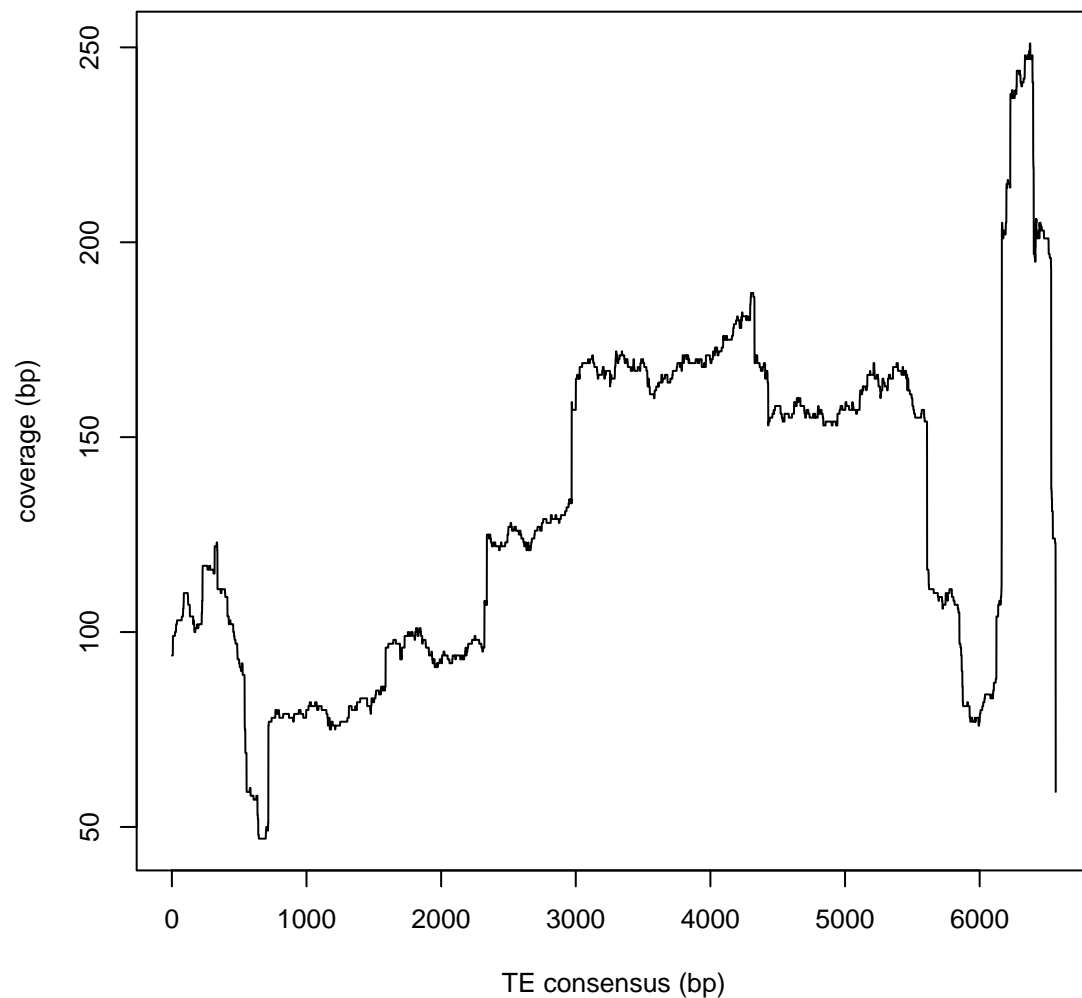


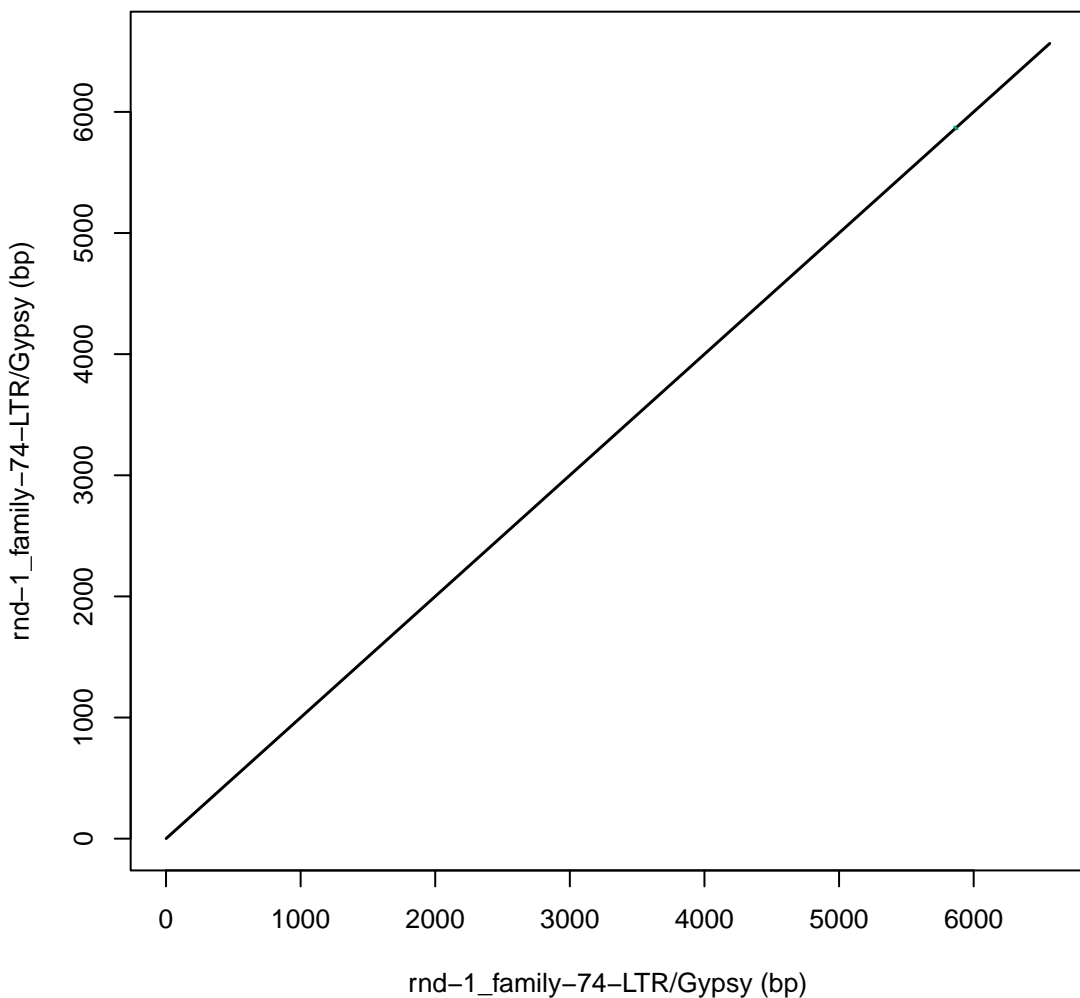
TE: rnd-1_family-74-LTR/Gypsy
consensus size: 6565bp; fragments: 788; full length: 16 (≥ 5908.5 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

