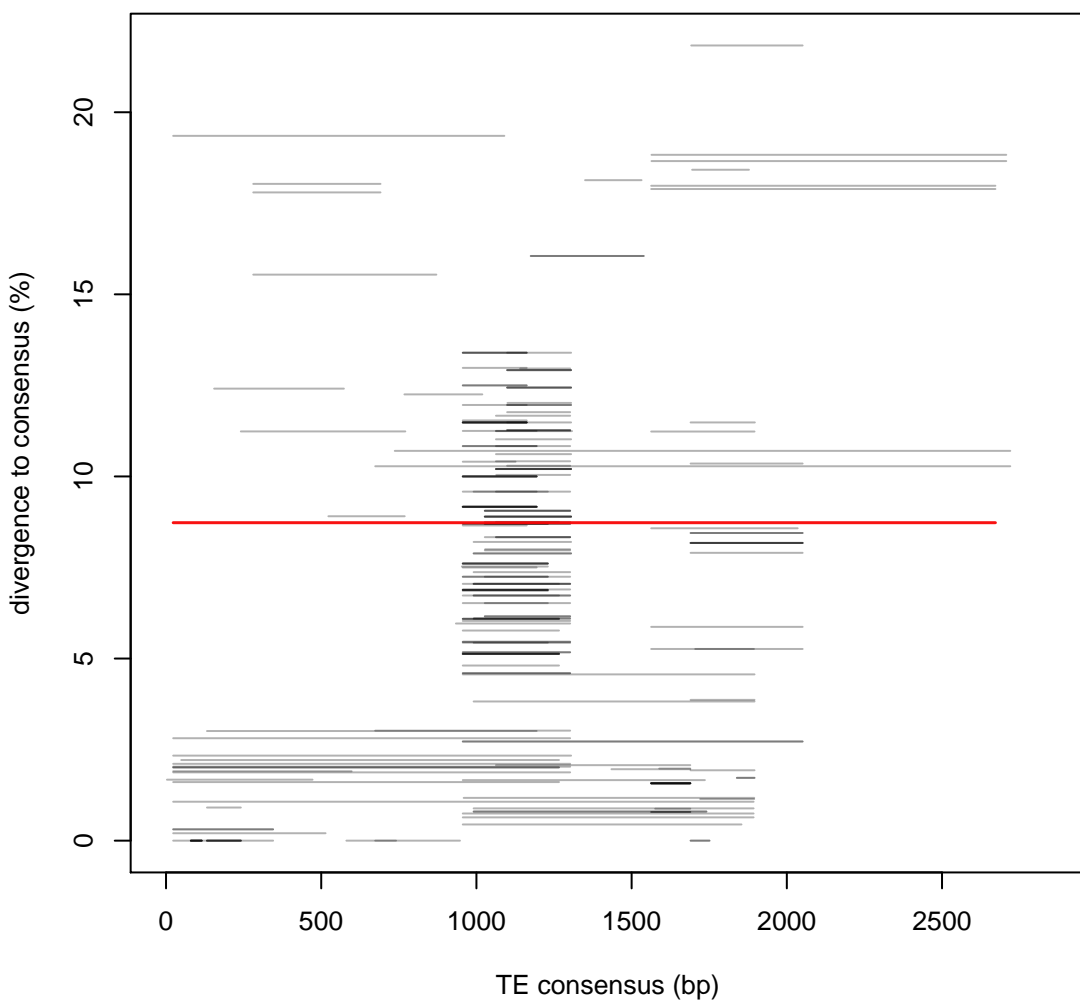
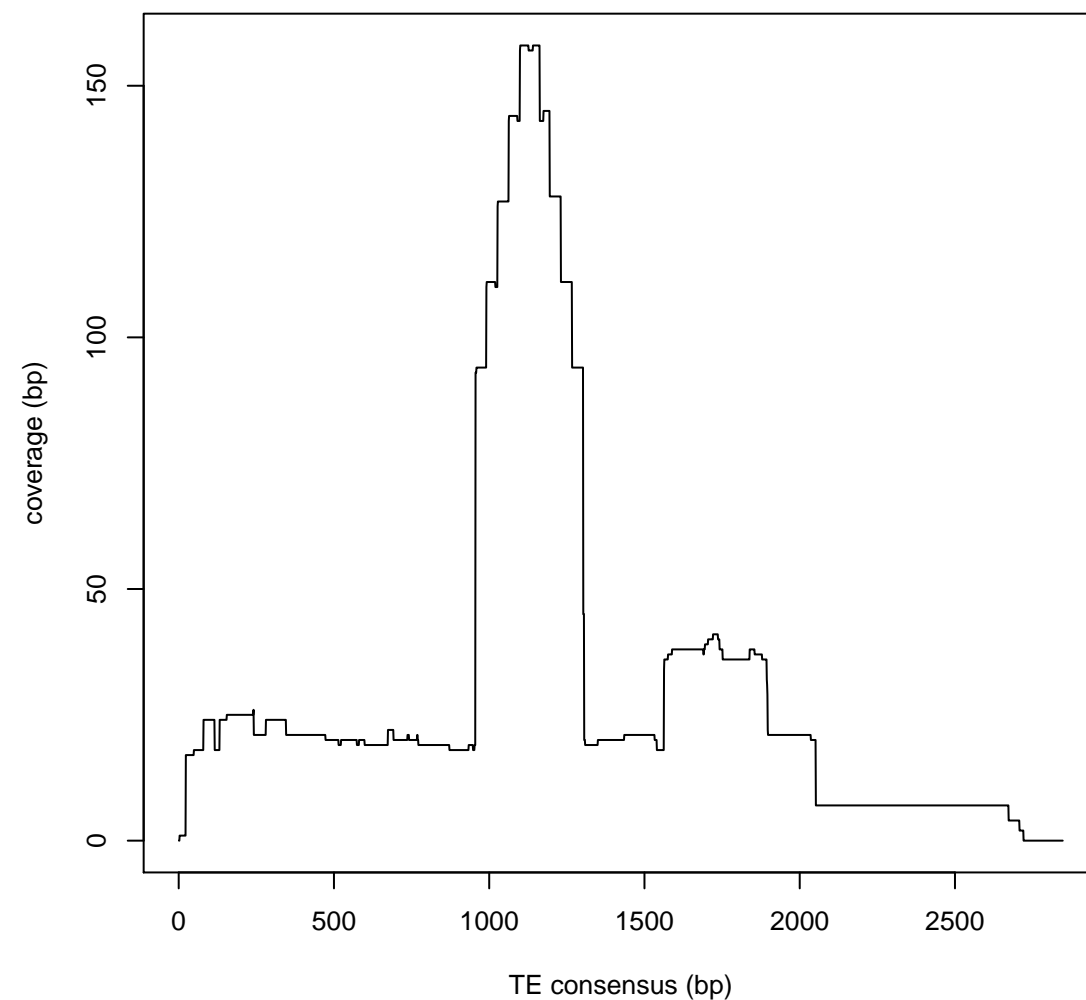


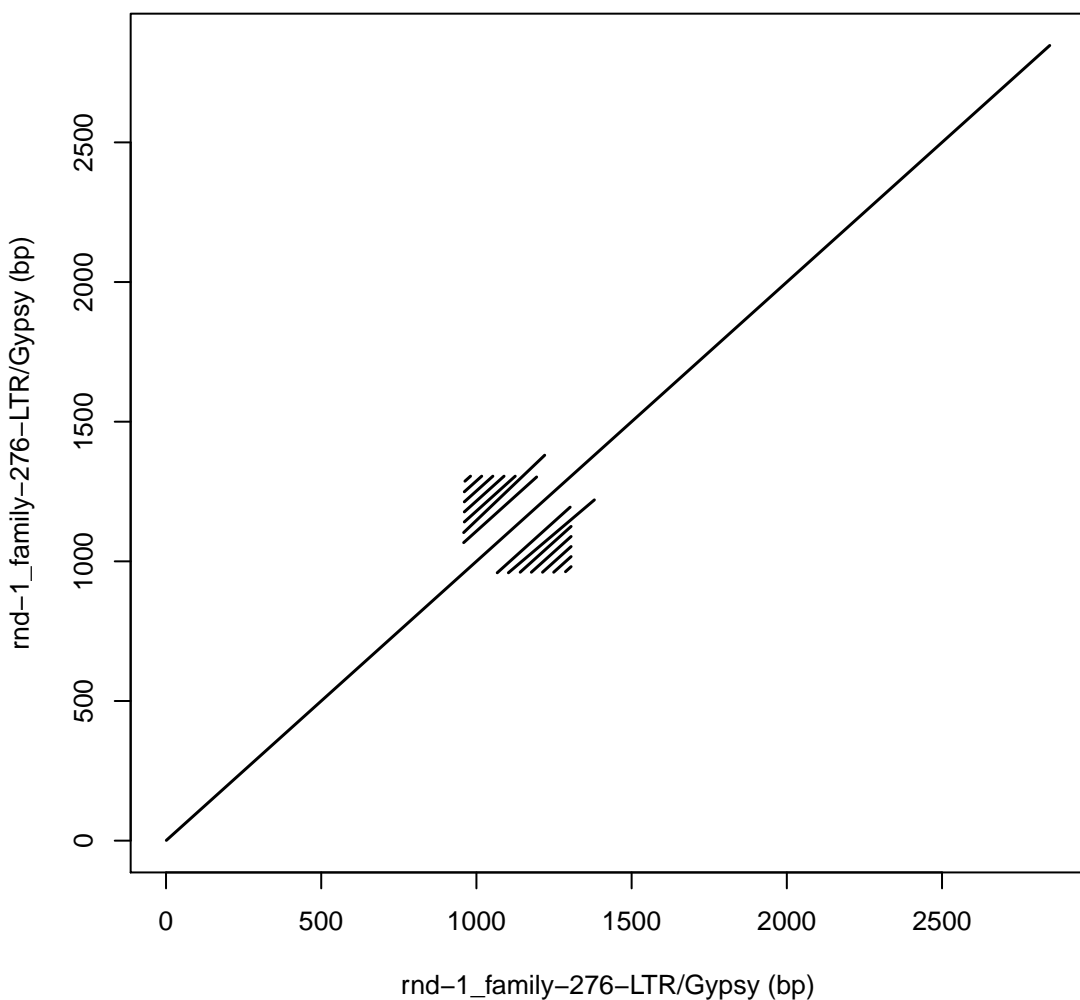
TE: rnd-1_family-276-LTR/Gypsy
consensus size: 2847bp; fragments: 229; full length: 1 (≥ 2562.3 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

