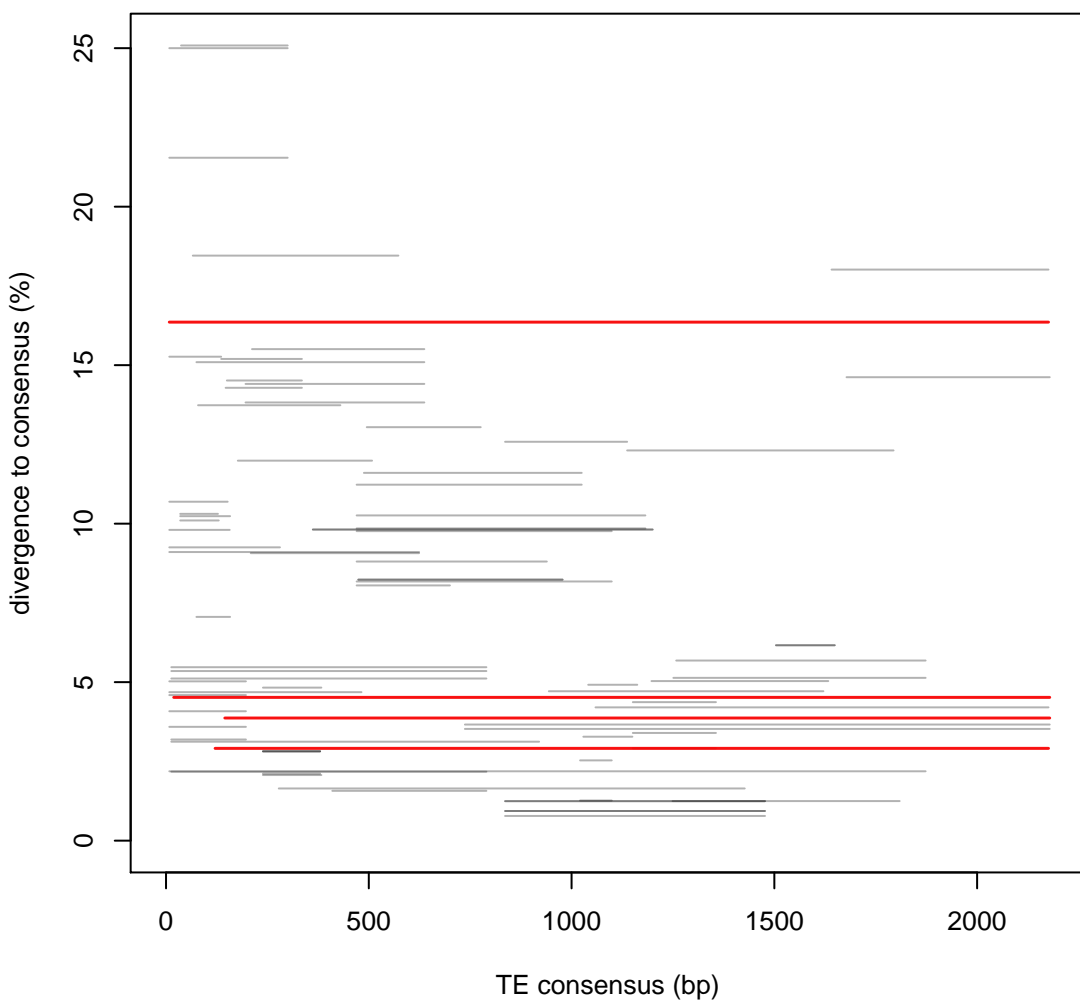
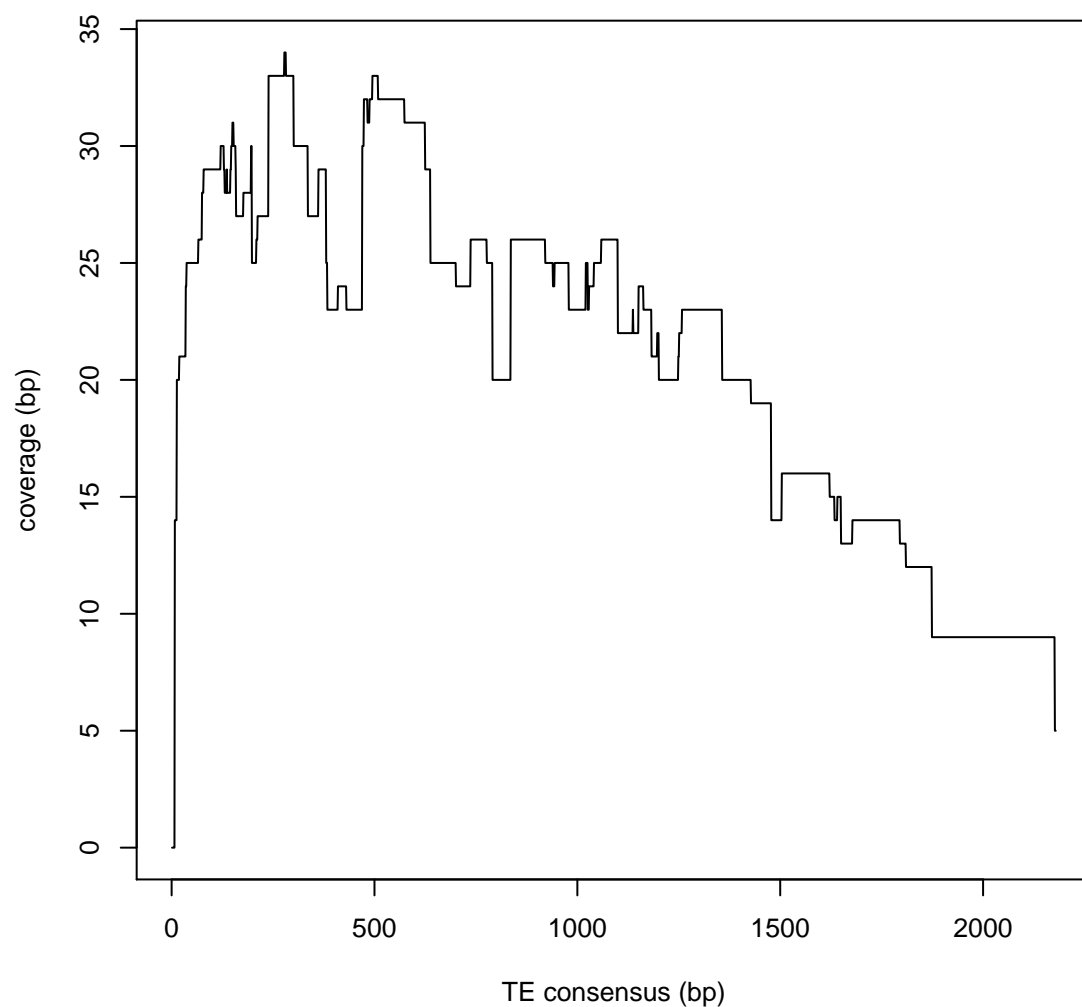


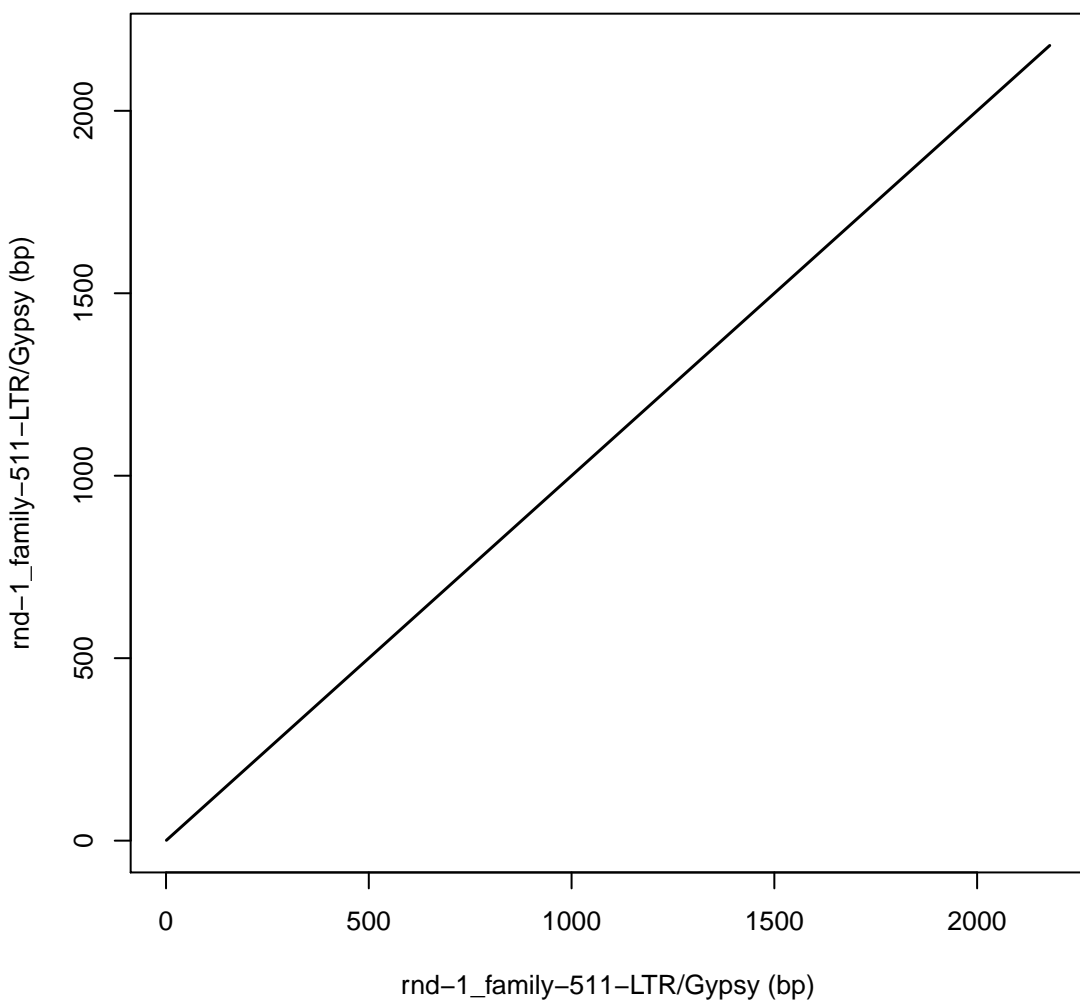
TE: rnd-1_family-511-LTR/Gypsy
consensus size: 2179bp; fragments: 86; full length: 4 (≥ 1961.1 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

