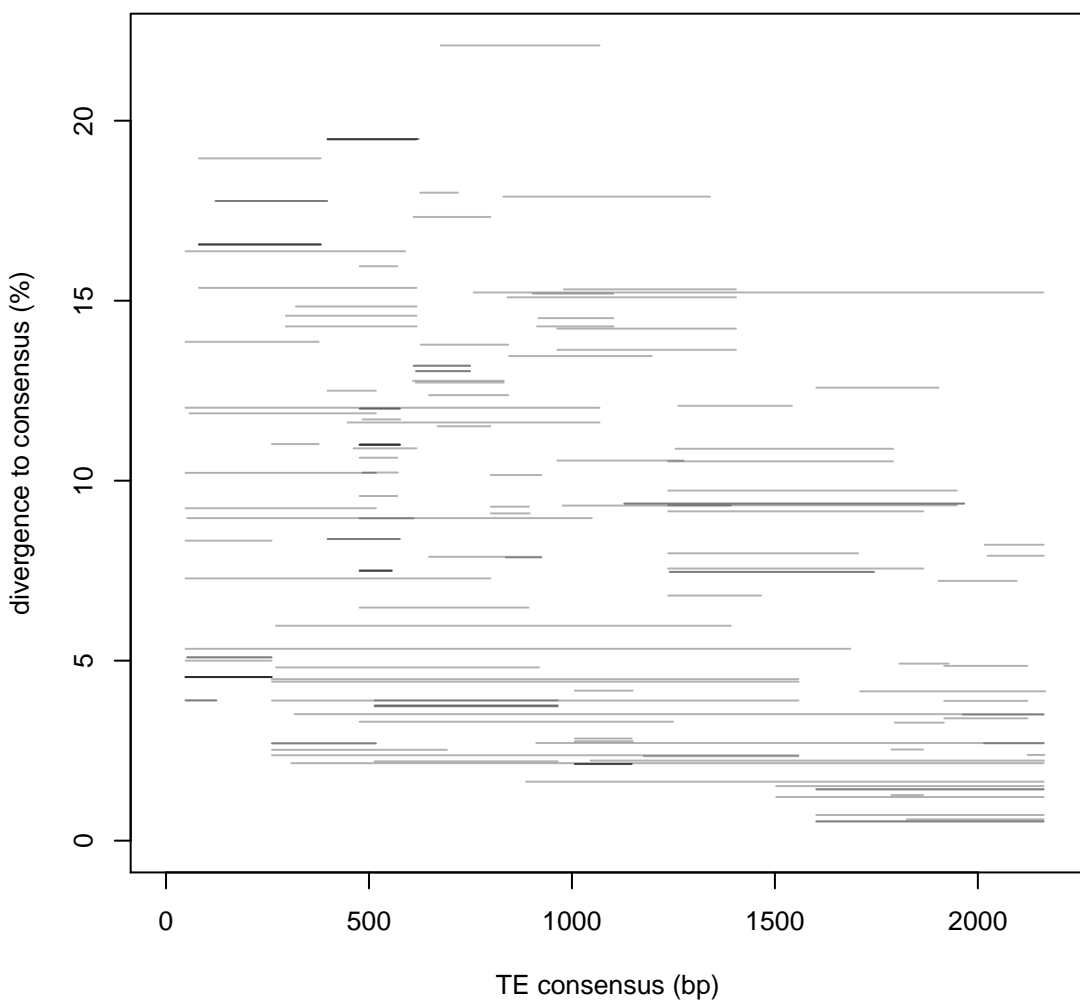
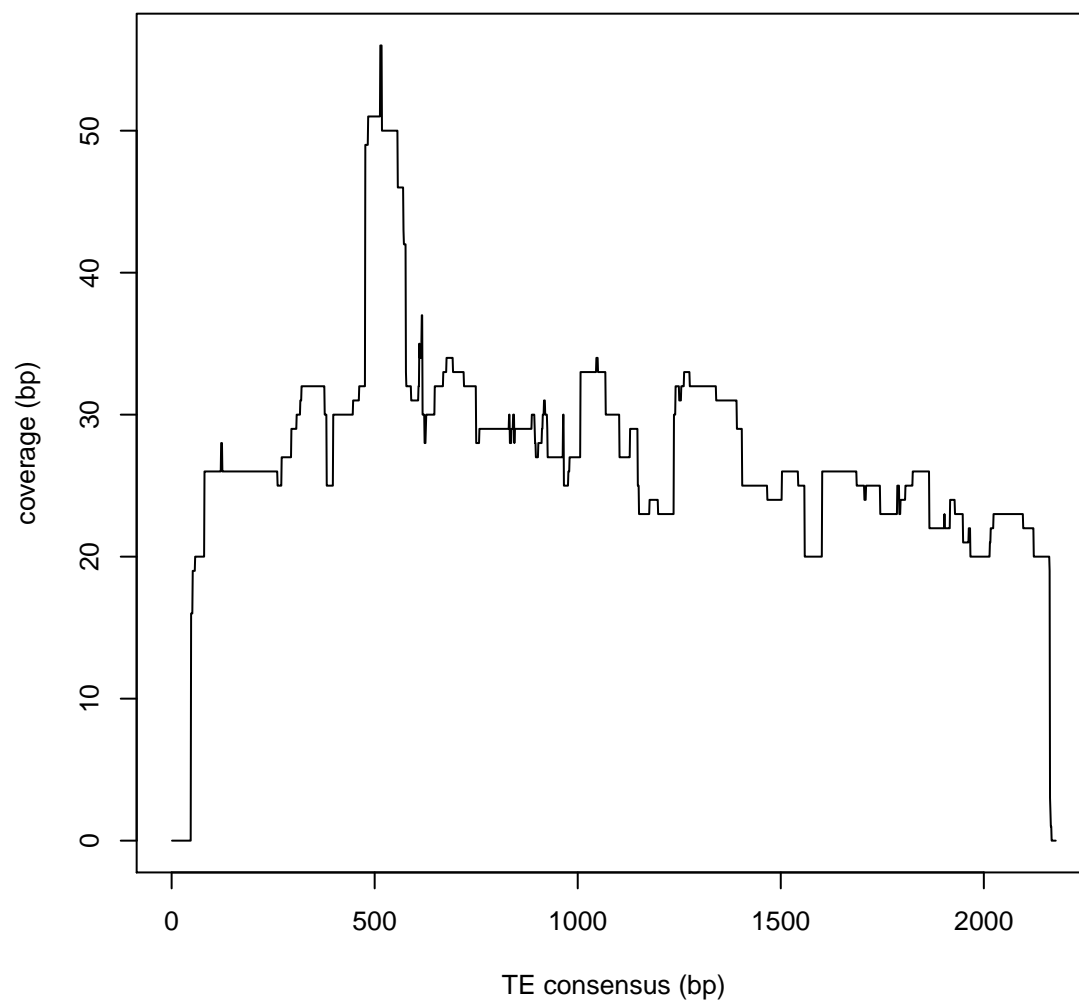


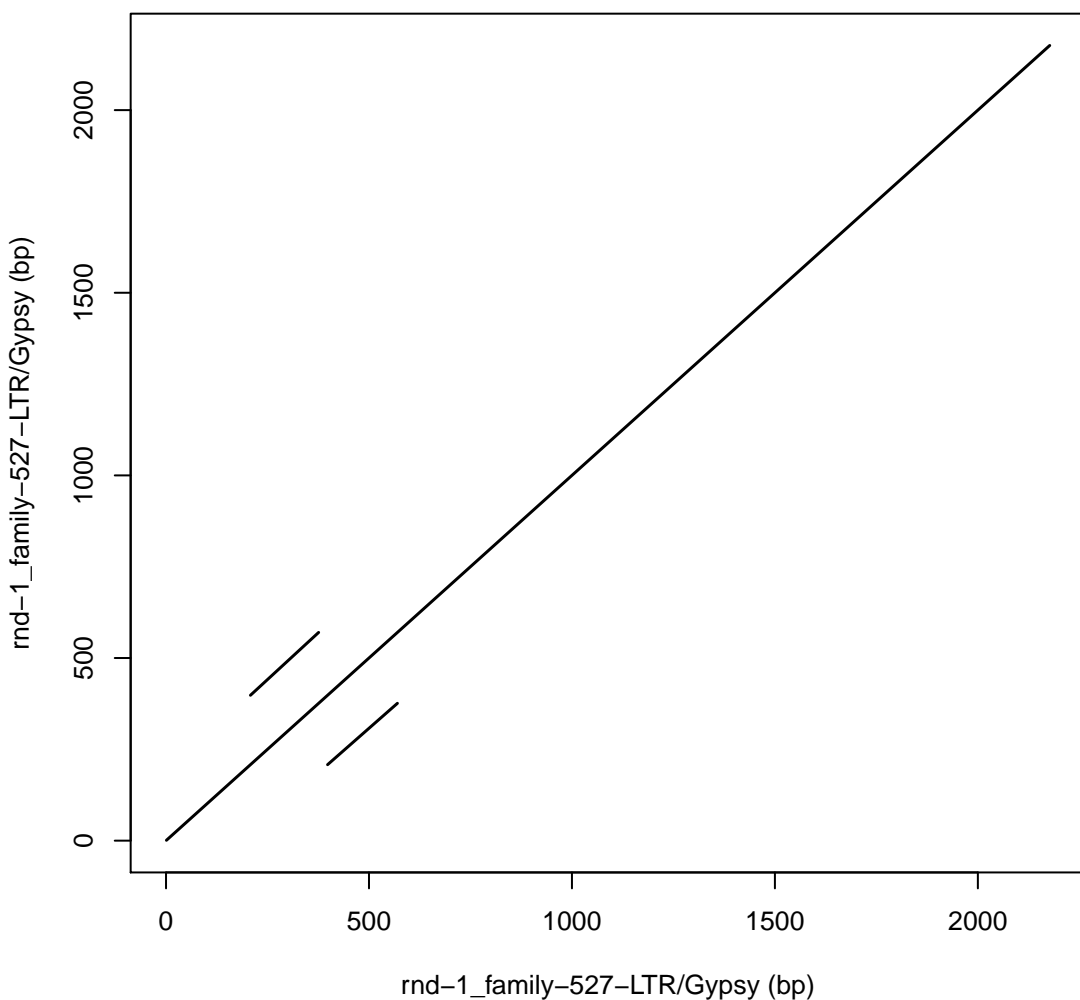
TE: rnd-1_family-527-LTR/Gypsy
consensus size: 2177bp; fragments: 151; full length: 0 (>=1959.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

