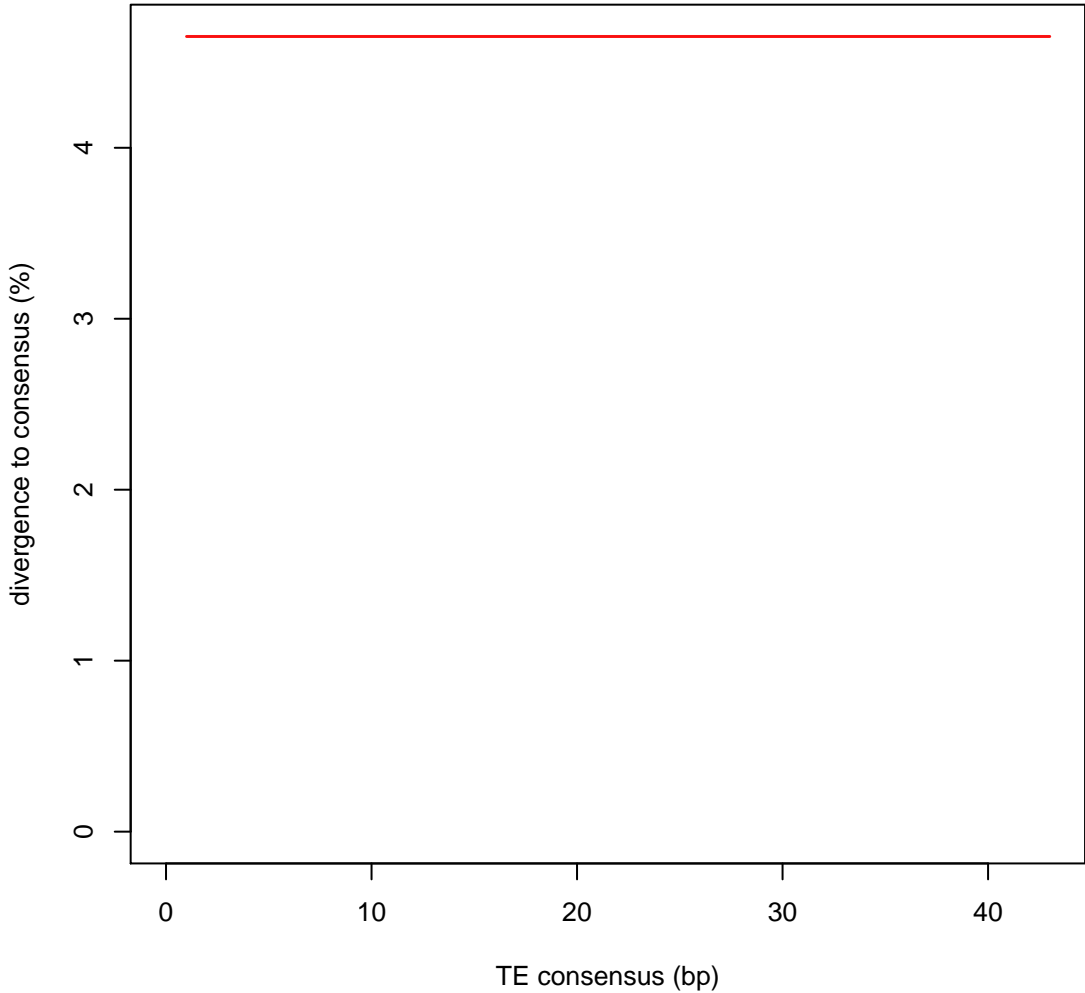
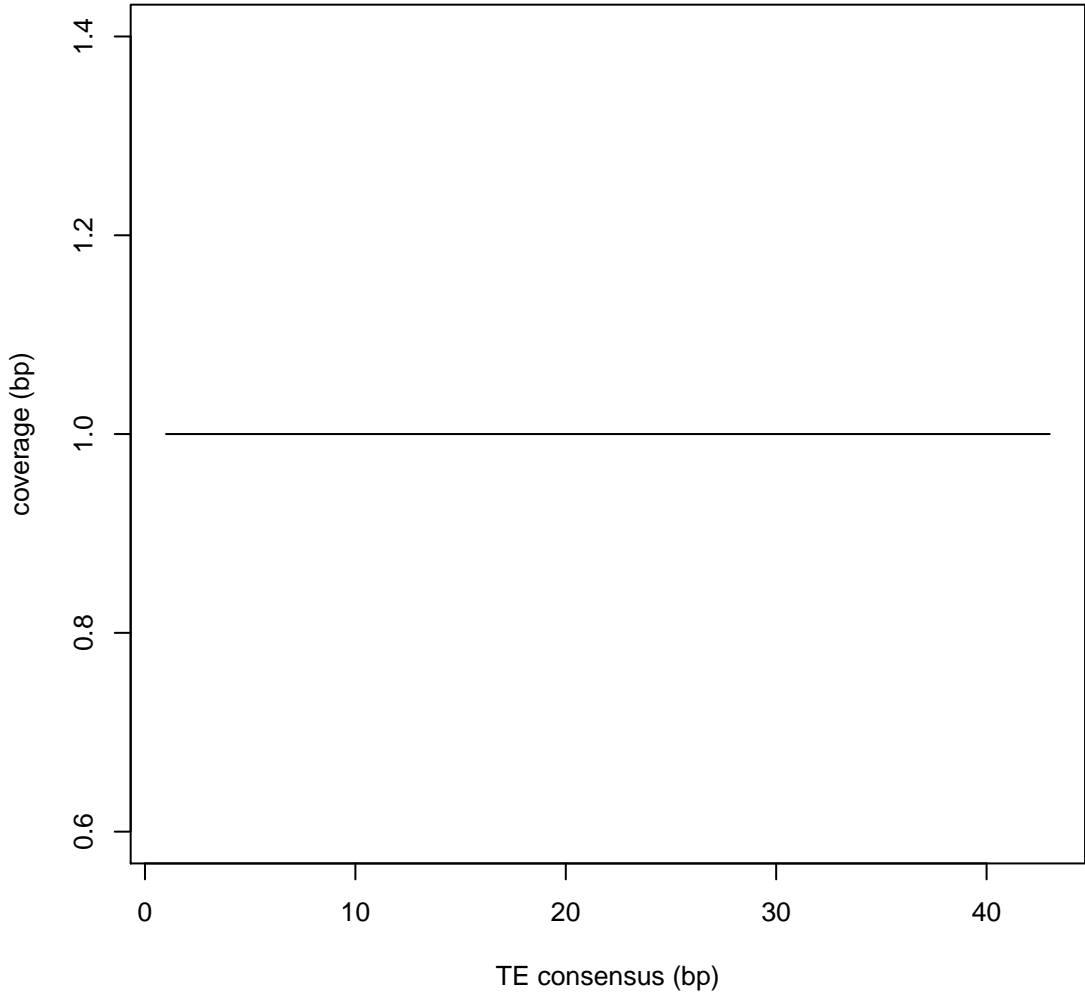


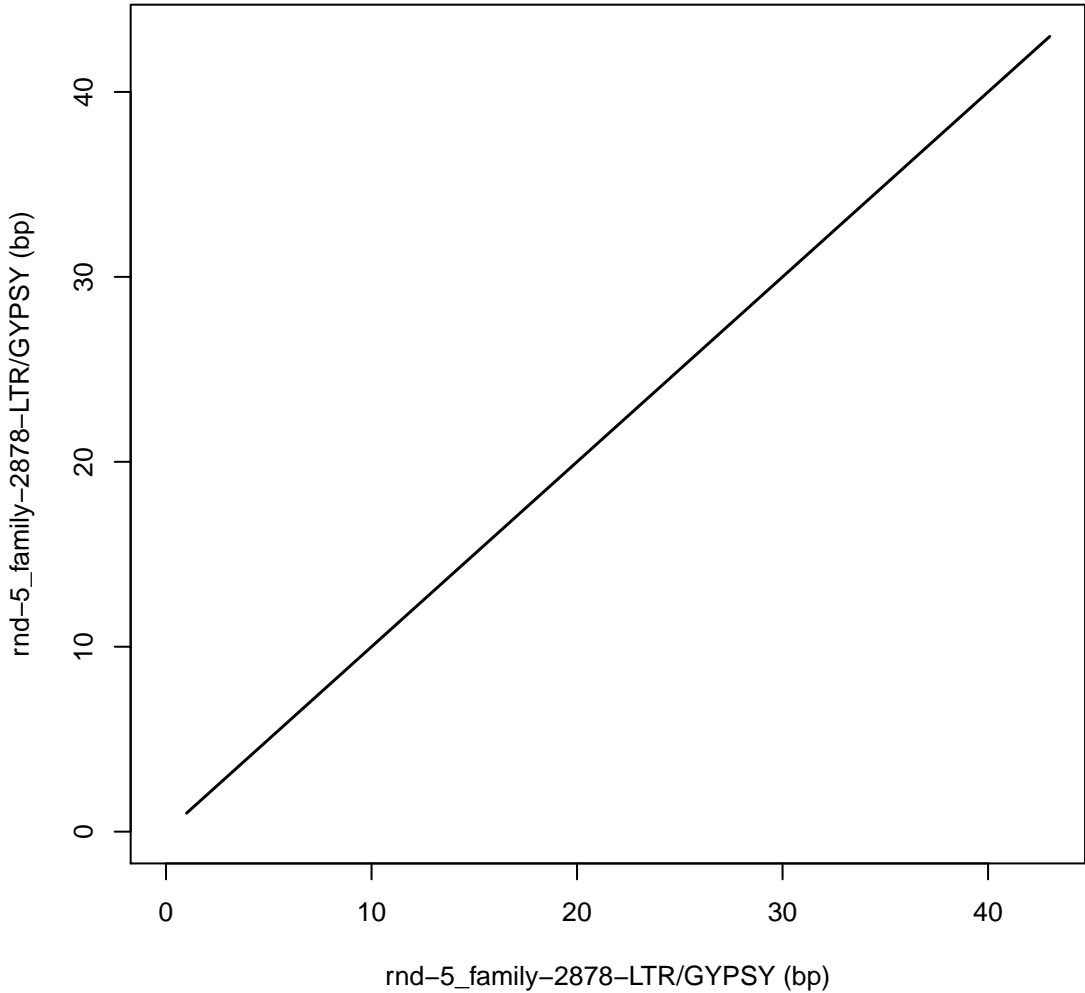
TE: rnd-5_family-2878-LTR/GYPSY
consensus size: 43bp; fragments: 1; full length: 1 (>=38.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

