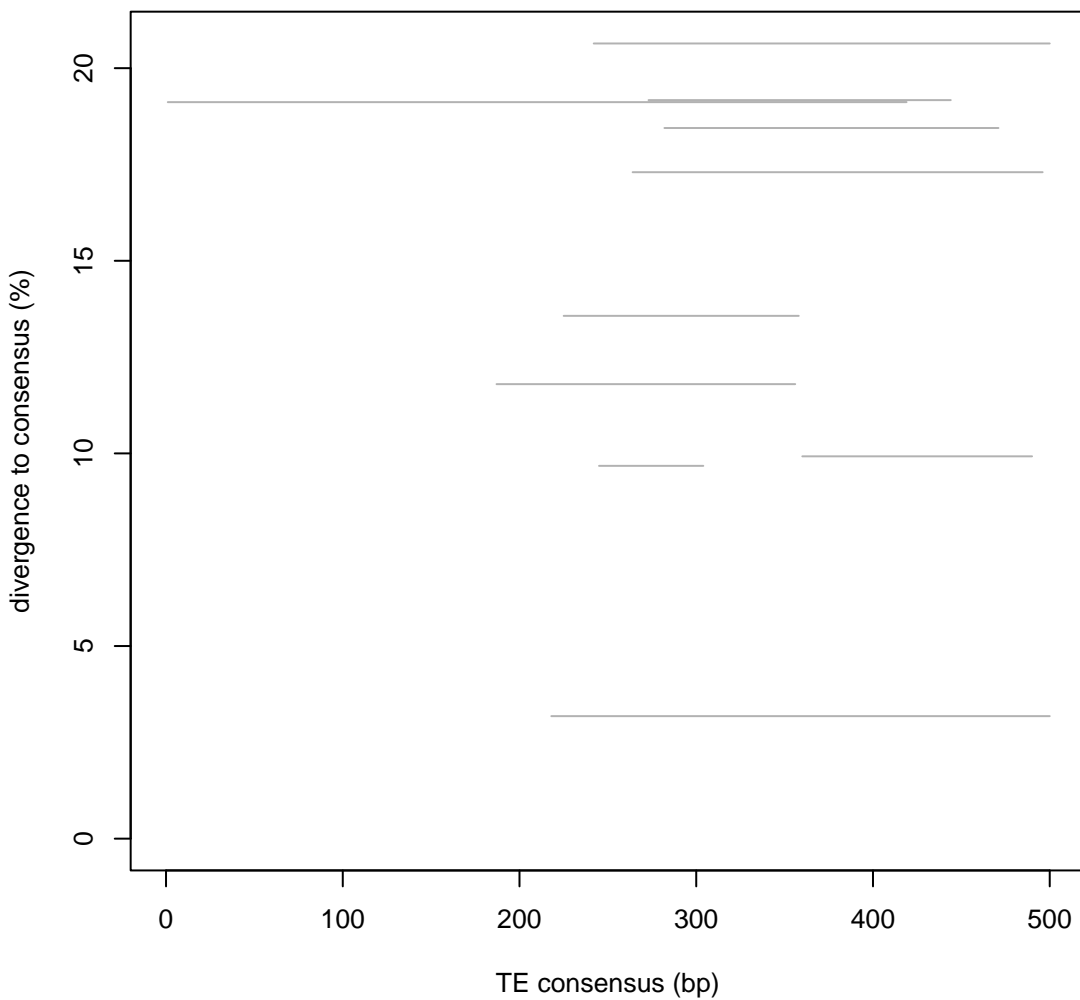
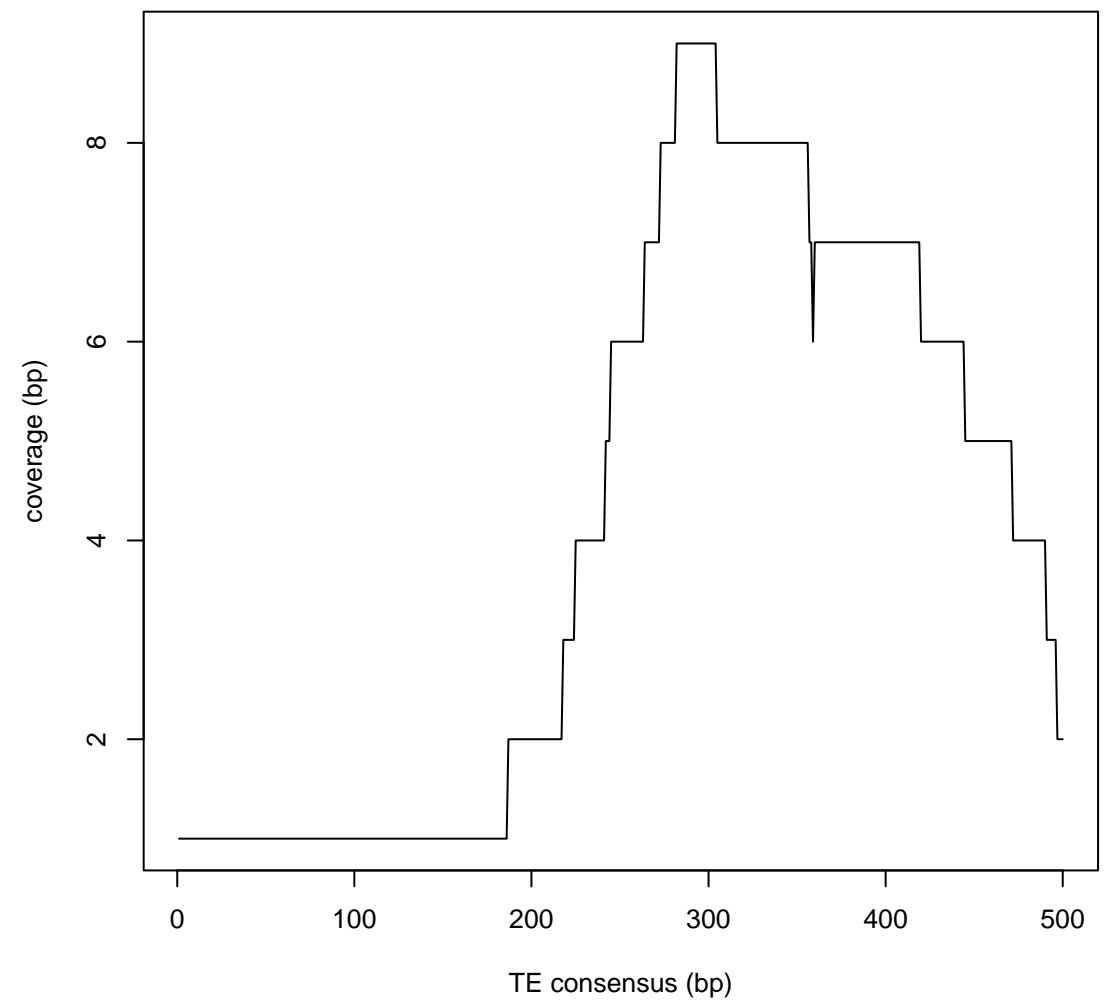


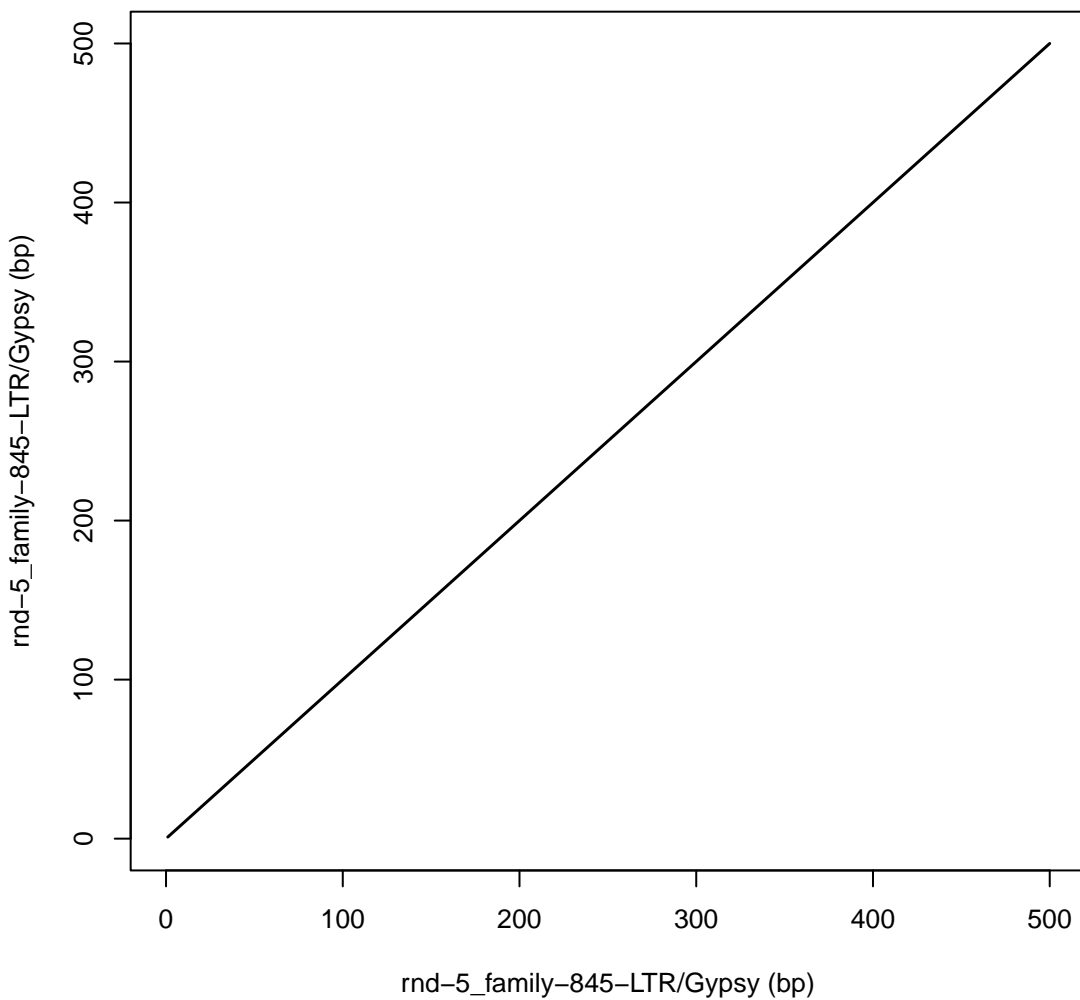
TE: rnd-5_family-845-LTR/Gypsy
consensus size: 500bp; fragments: 10; full length: 0 (>=450bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

