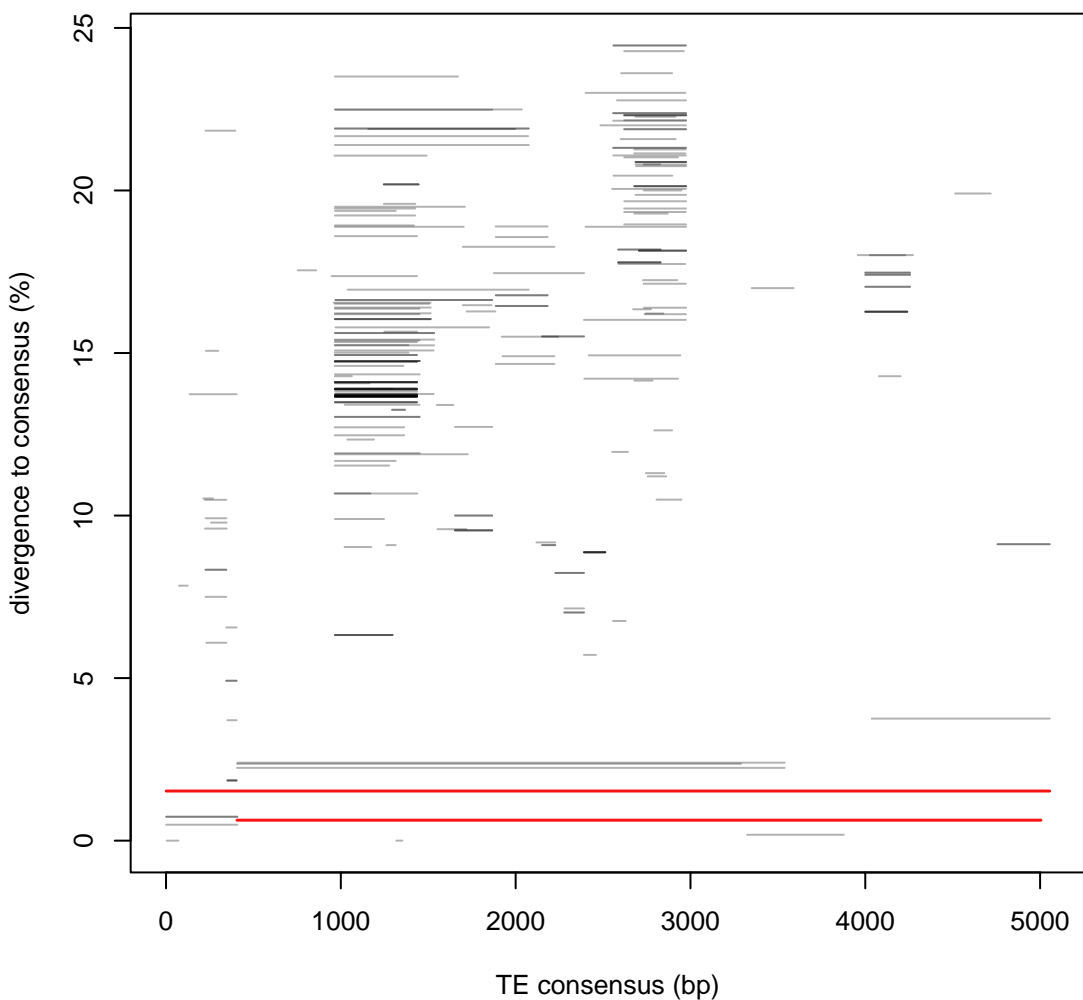
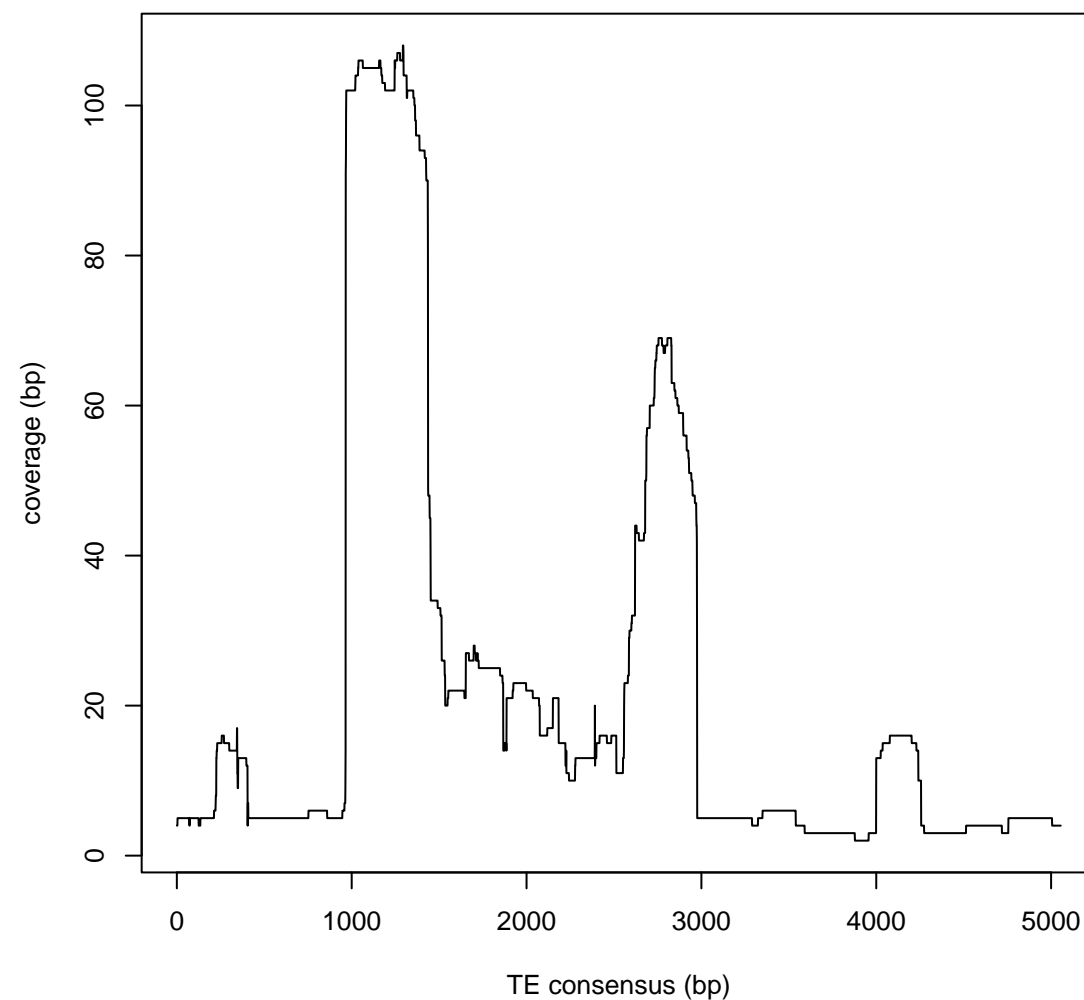


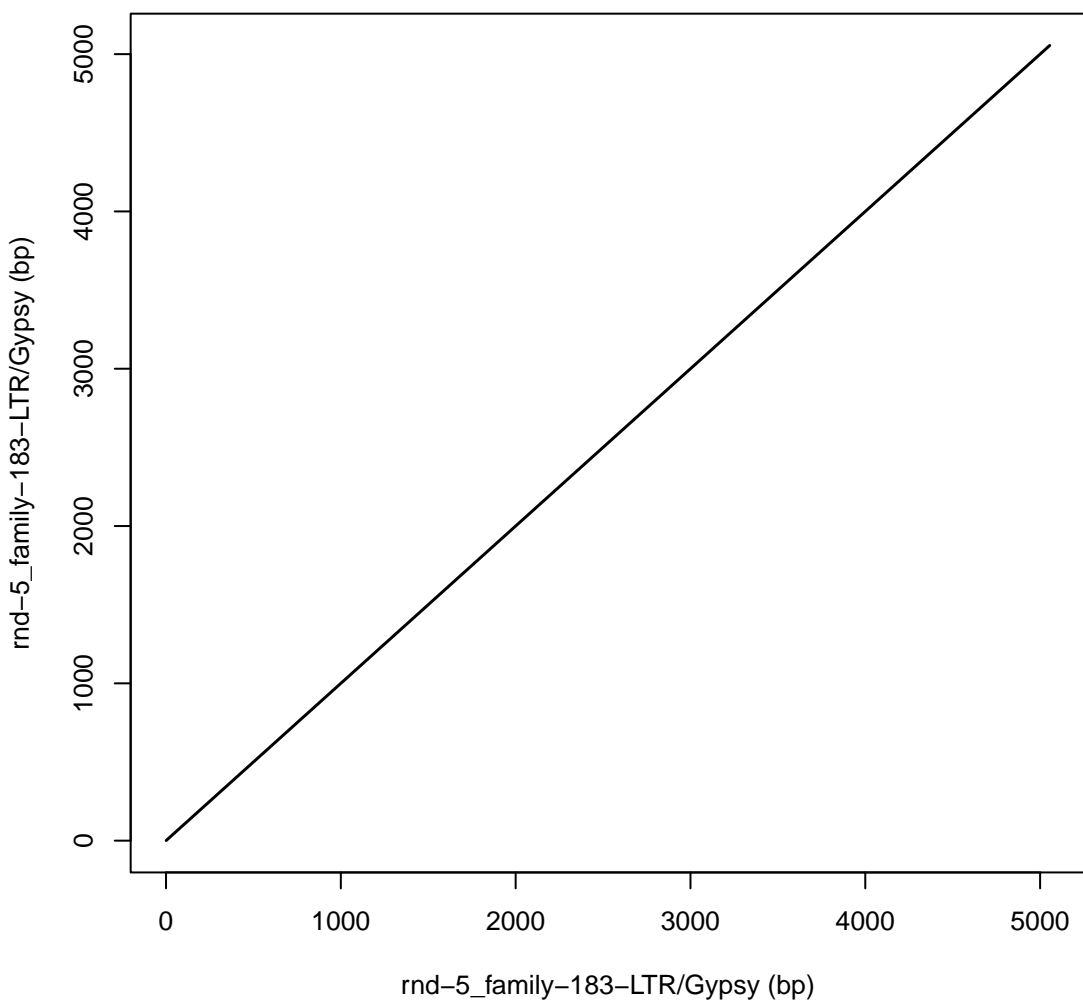
TE: rnd-5_family-183-LTR/Gypsy
consensus size: 5055bp; fragments: 265; full length: 2 (≥ 4549.5 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

