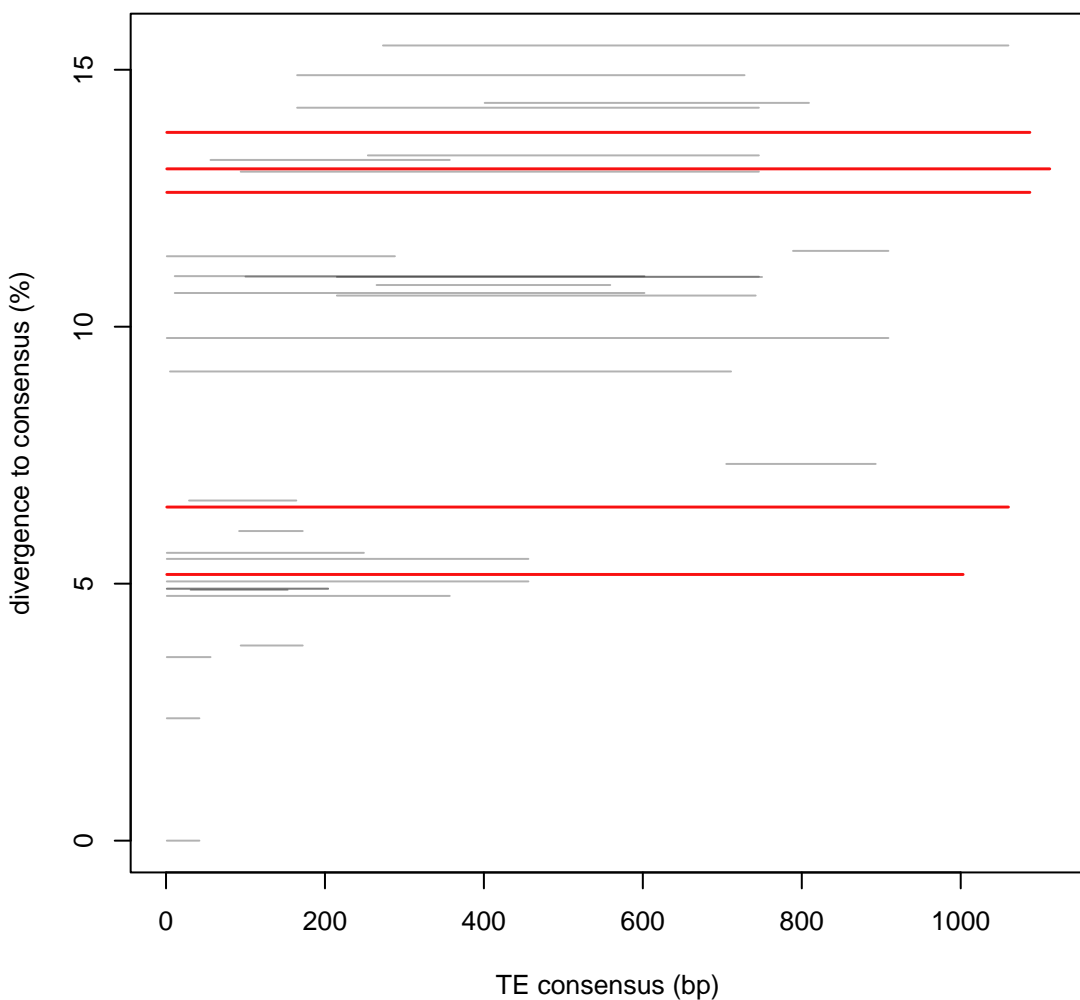
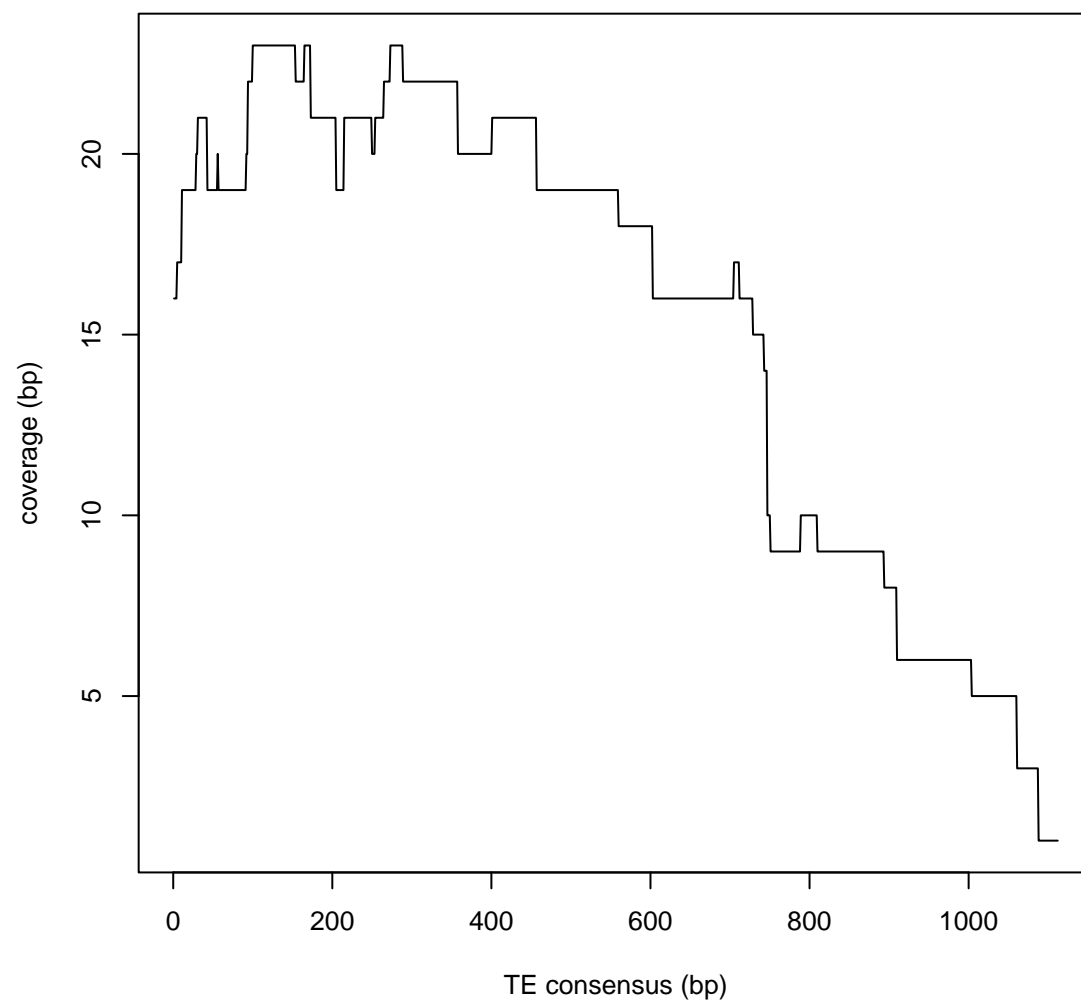


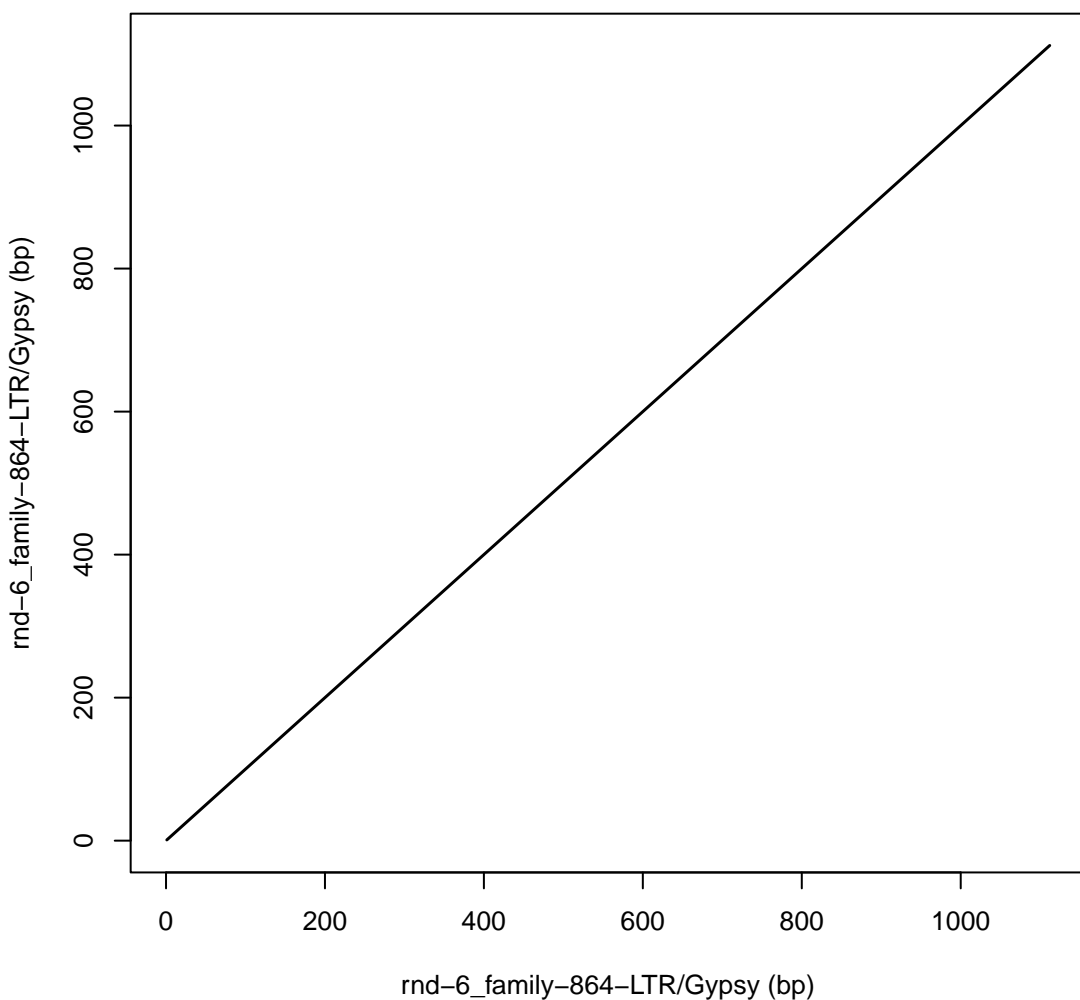
TE: rnd-6_family-864-LTR/Gypsy
consensus size: 1112bp; fragments: 36; full length: 5 (≥ 1000.8 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

