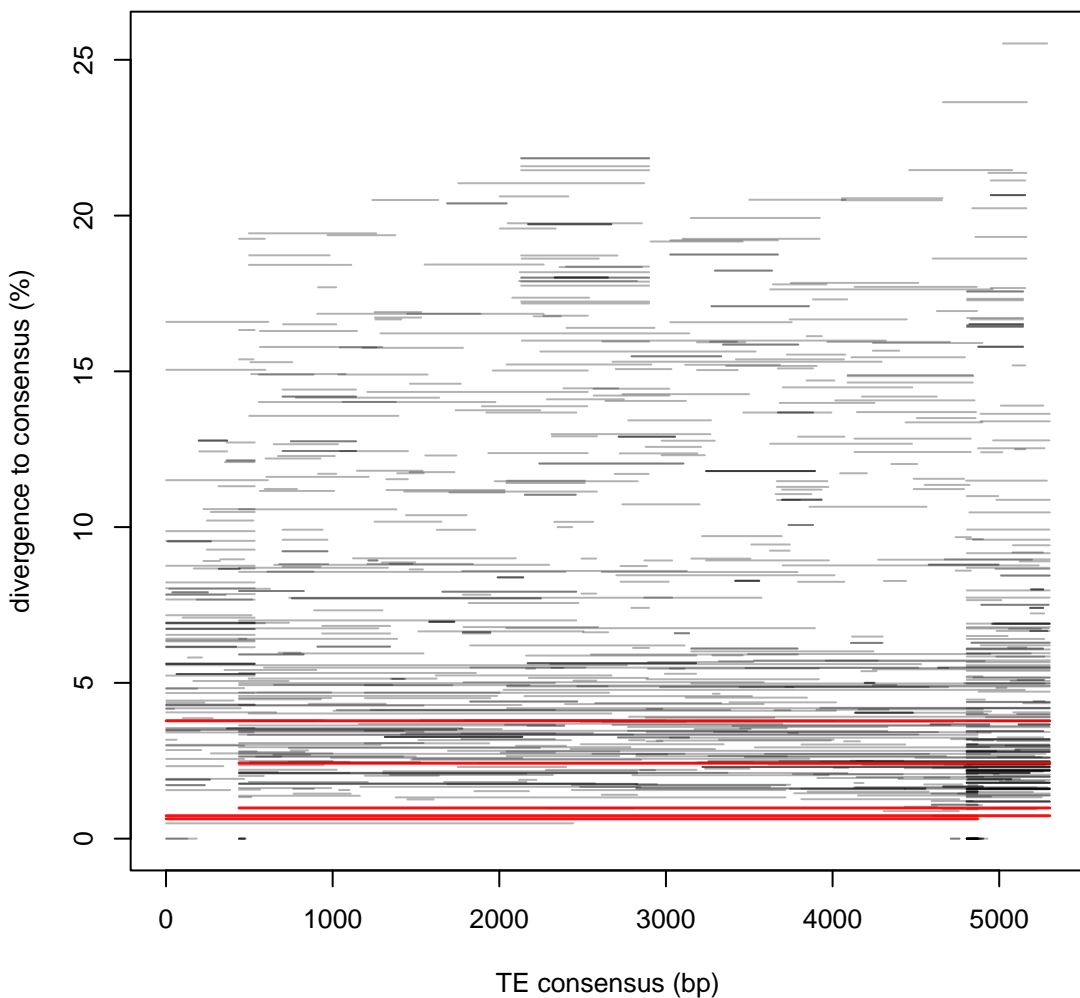
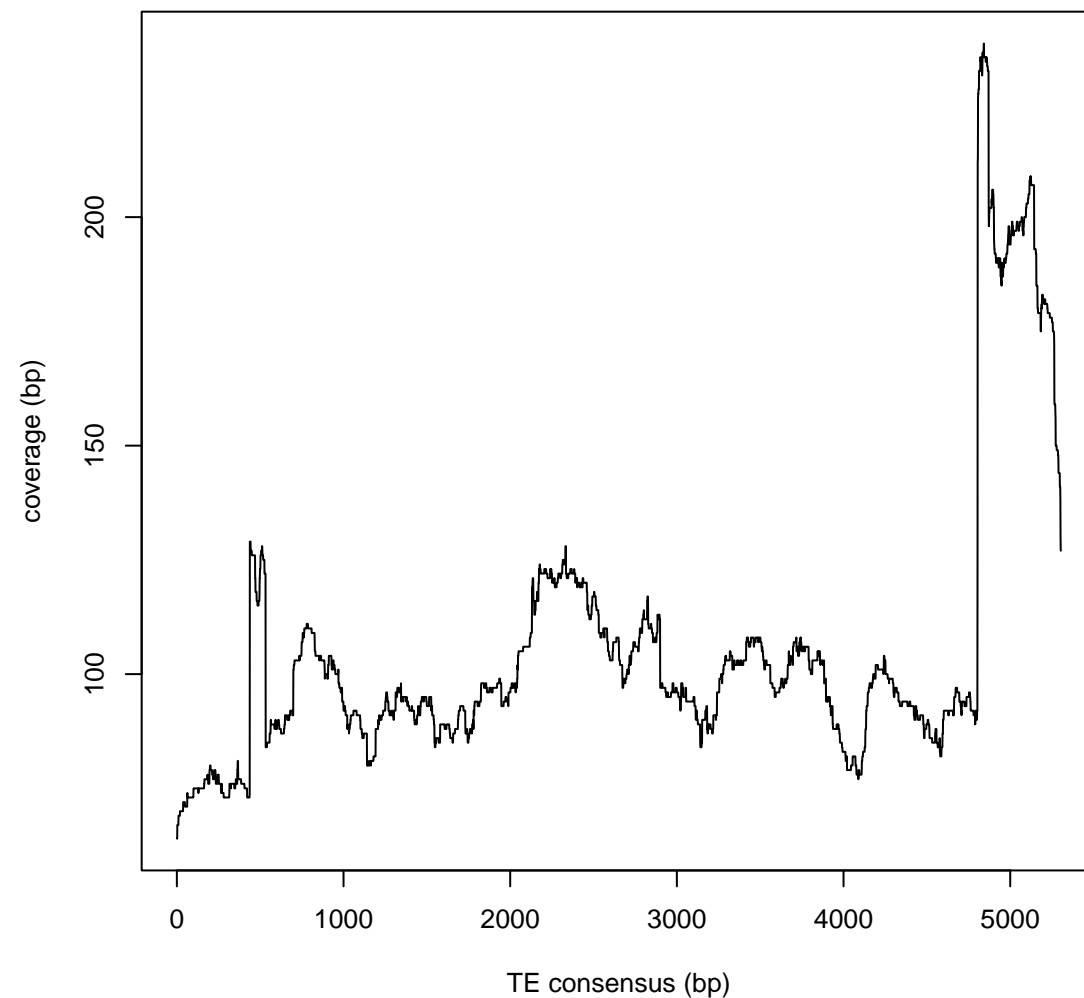


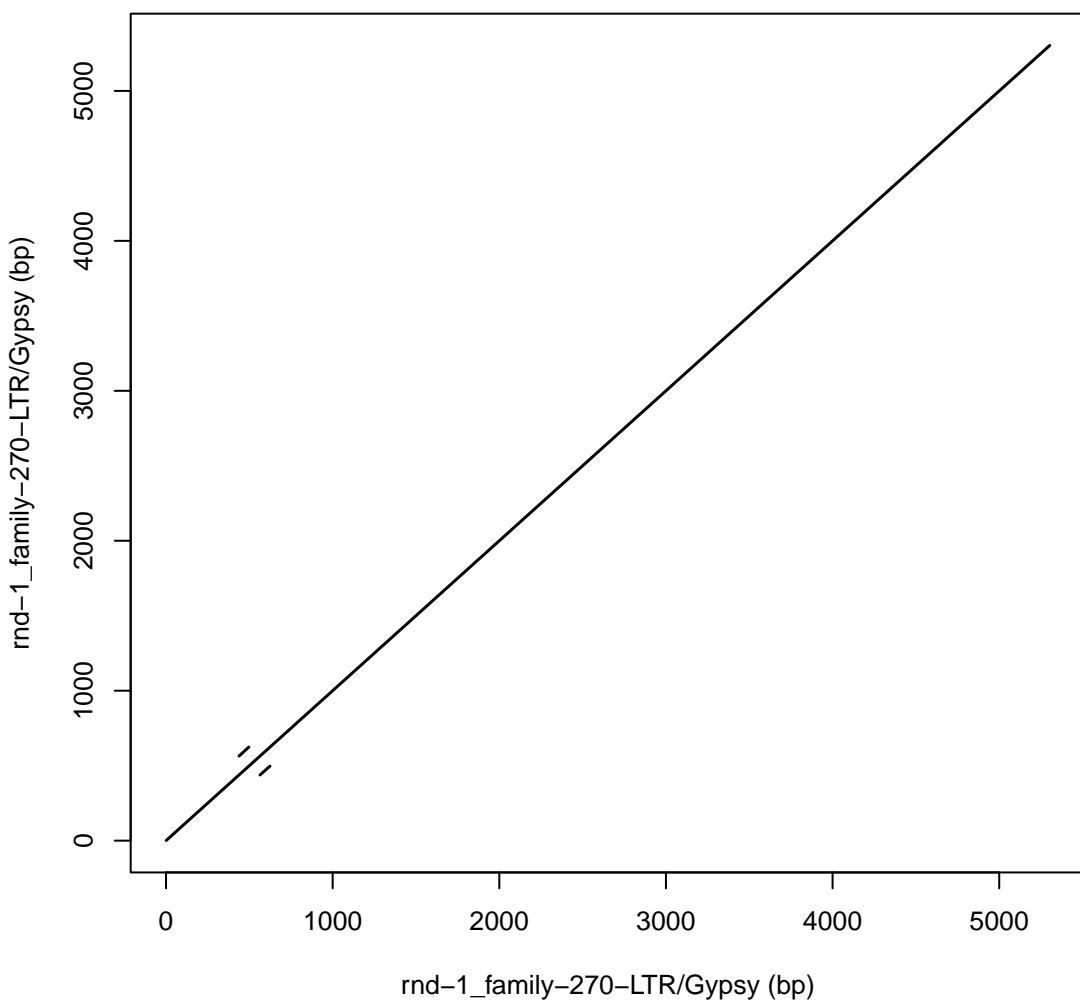
TE: rnd-1_family-270-LTR/Gypsy
consensus size: 5303bp; fragments: 894; full length: 5 (>=4772.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

