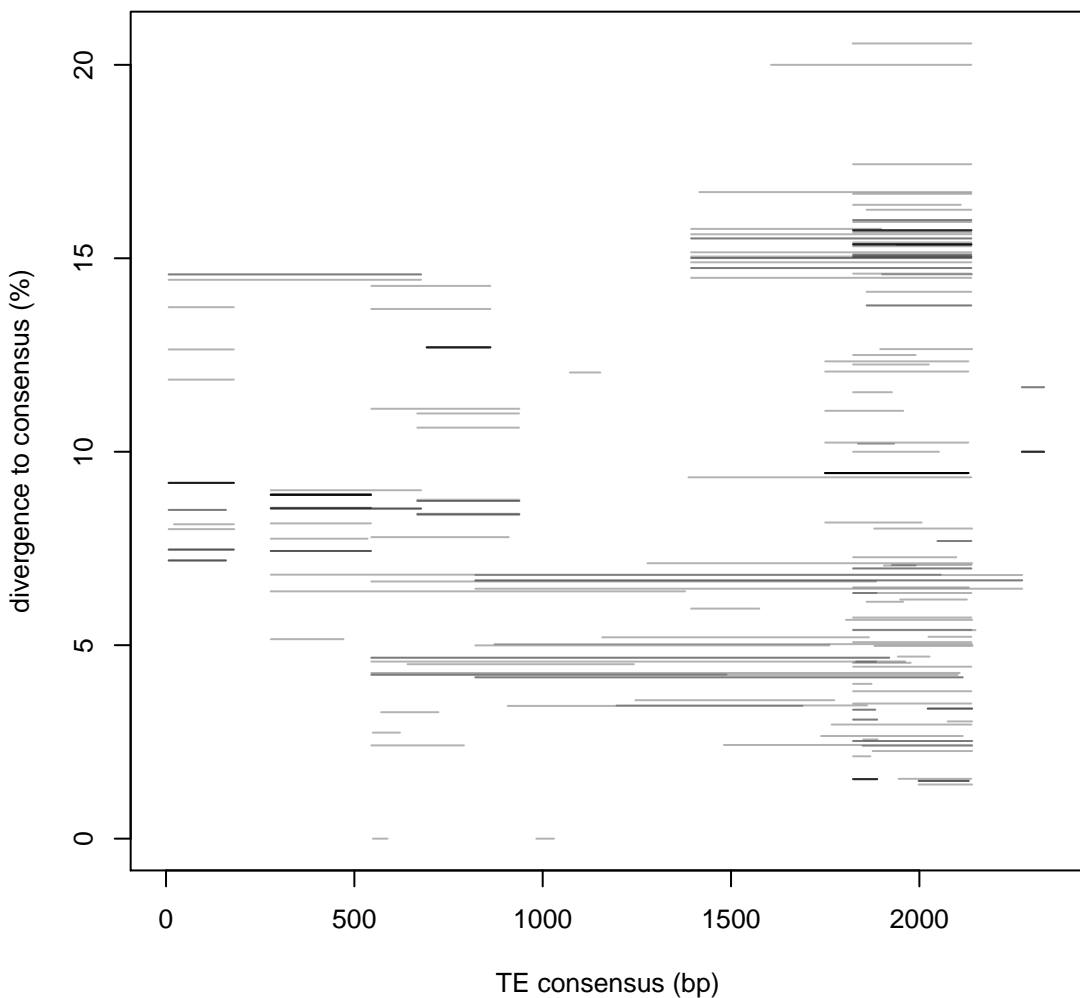
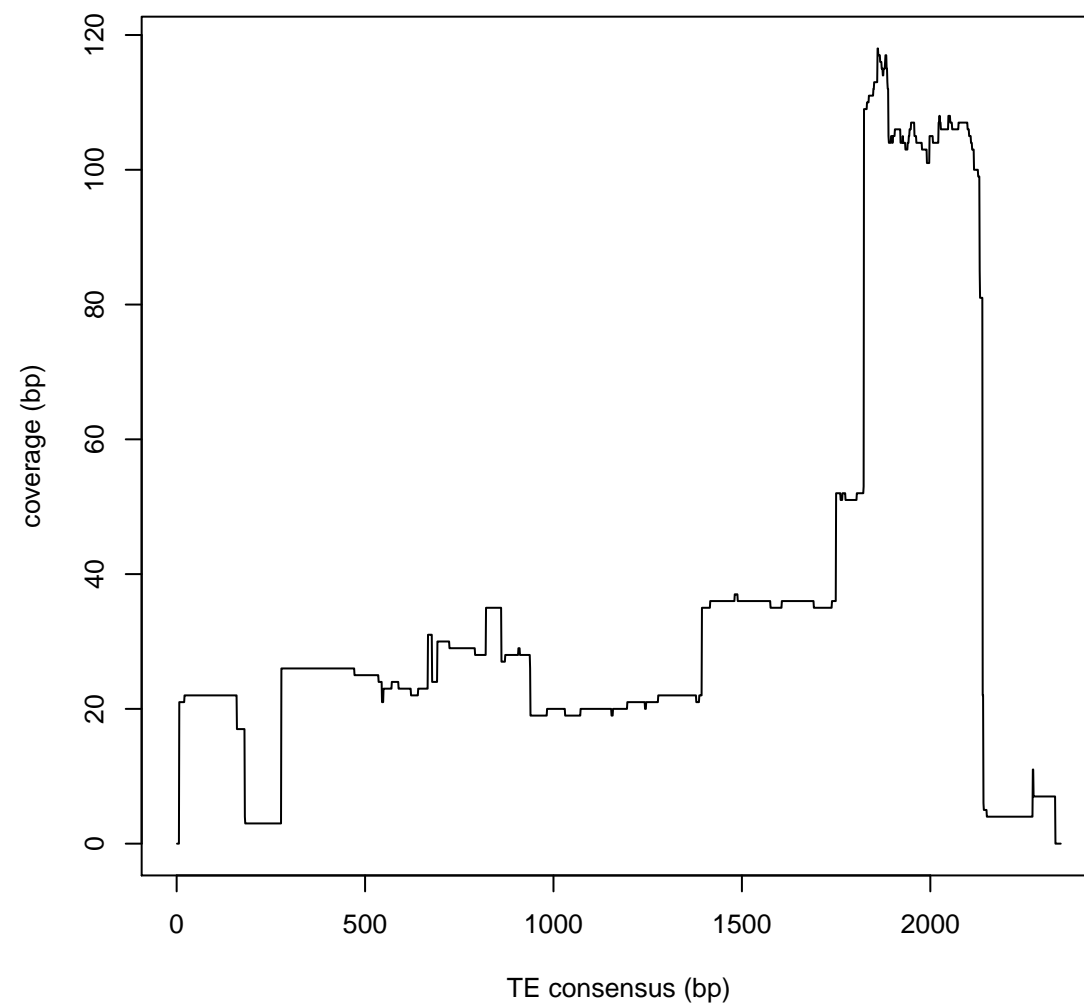


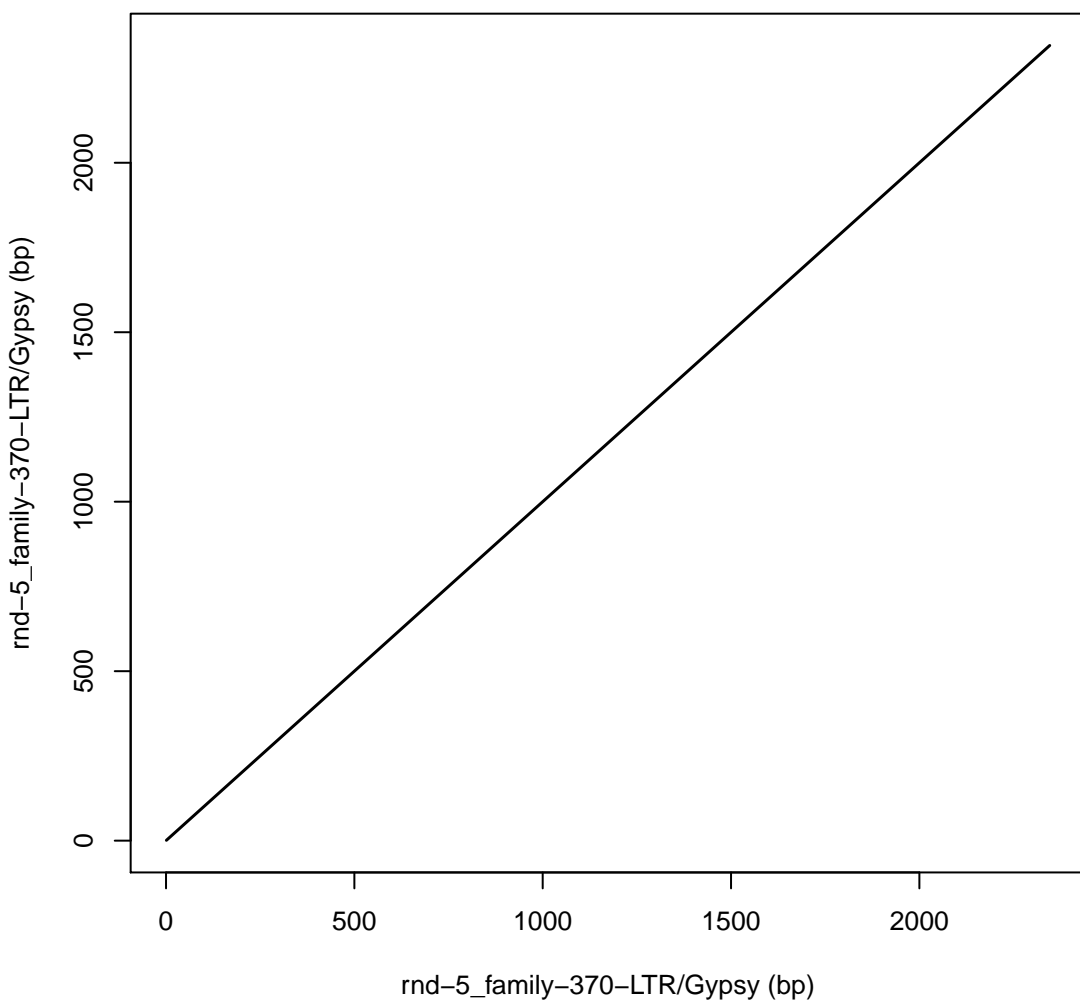
TE: rnd-5_family-370-LTR/Gypsy
consensus size: 2346bp; fragments: 221; full length: 0 (>=2111.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

