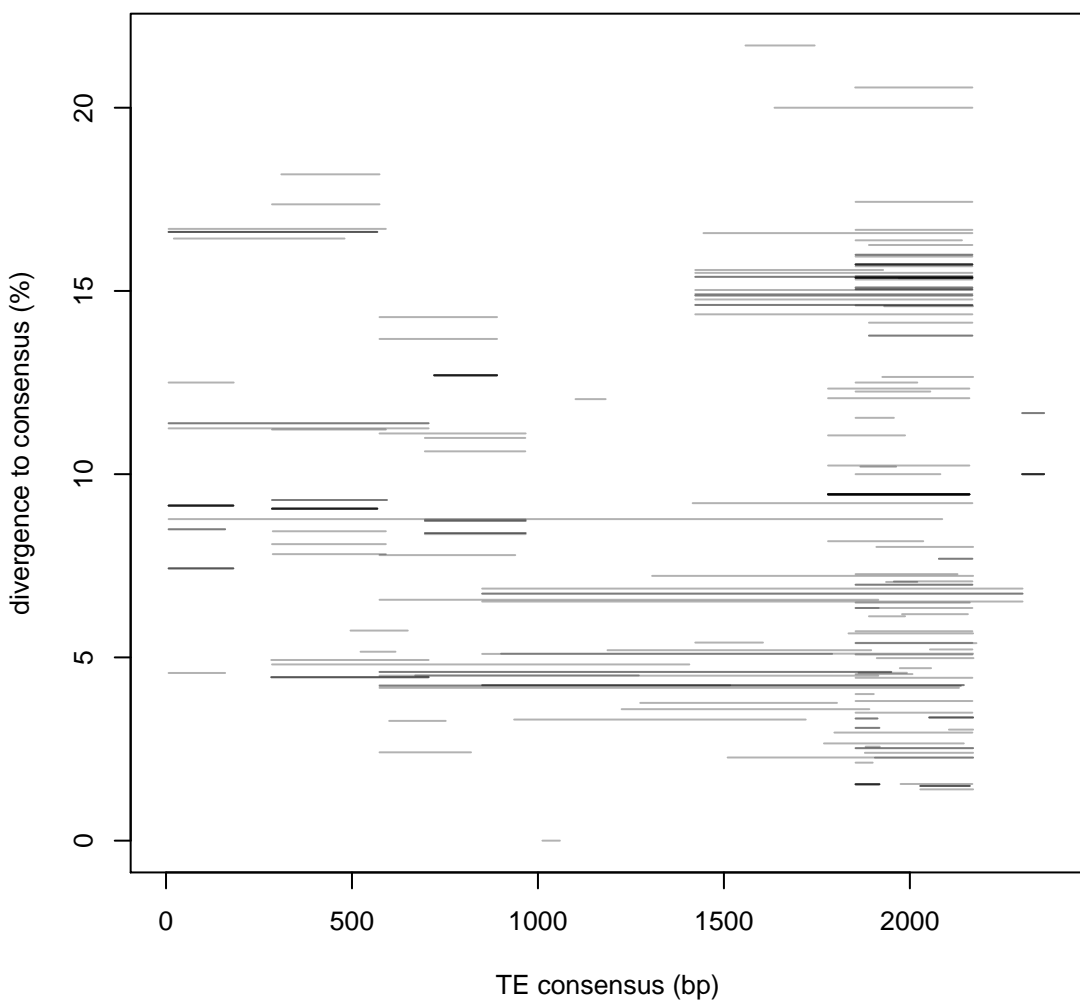
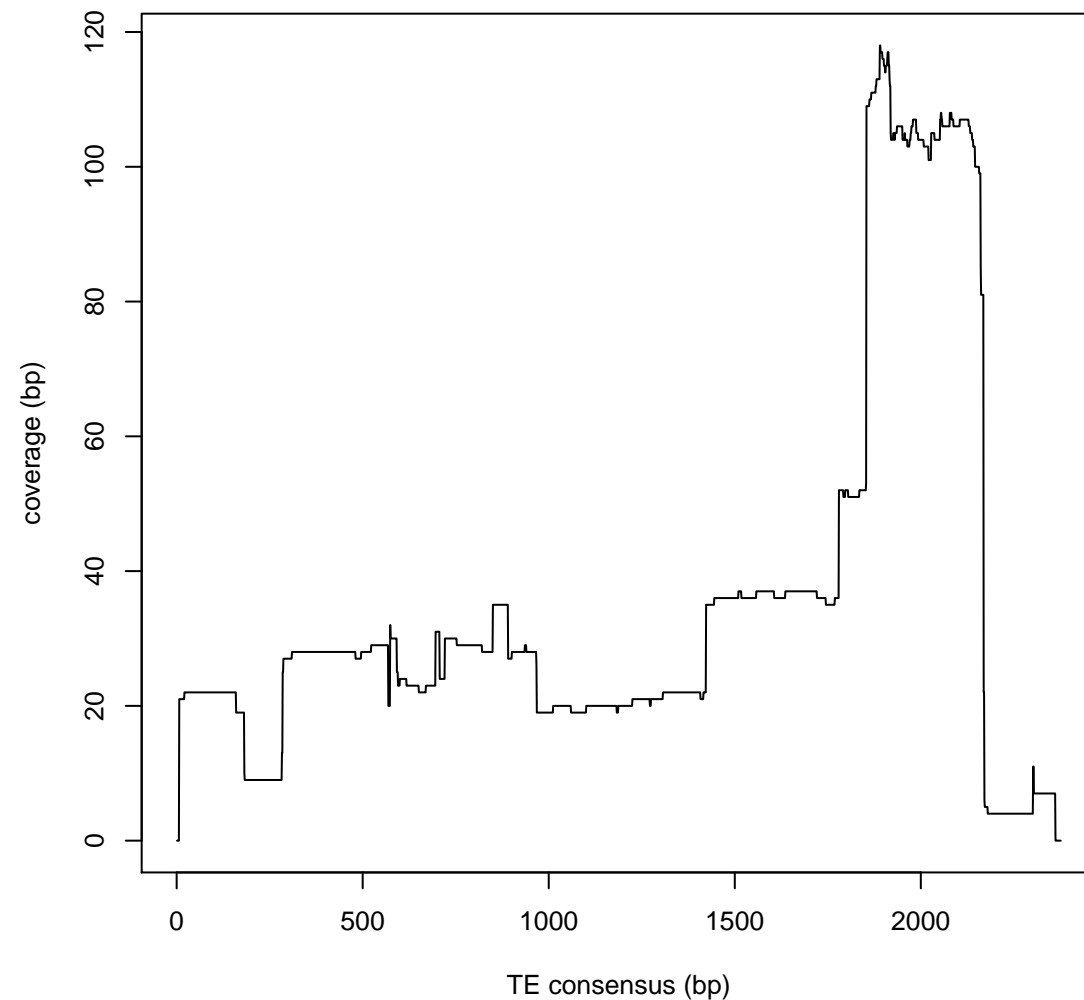


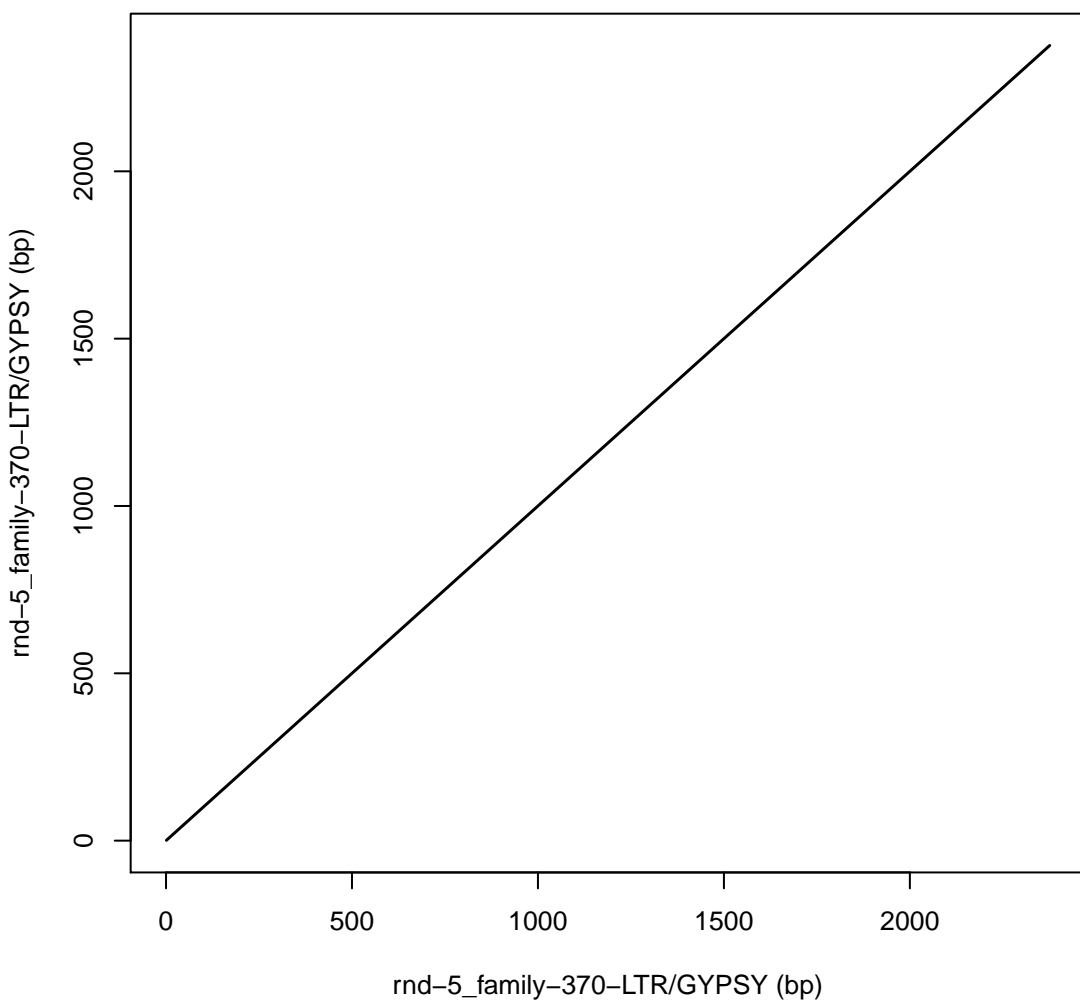
TE: rnd-5_family-370-LTR/GYPSY
consensus size: 2376bp; fragments: 218; full length: 0 (>=2138.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

