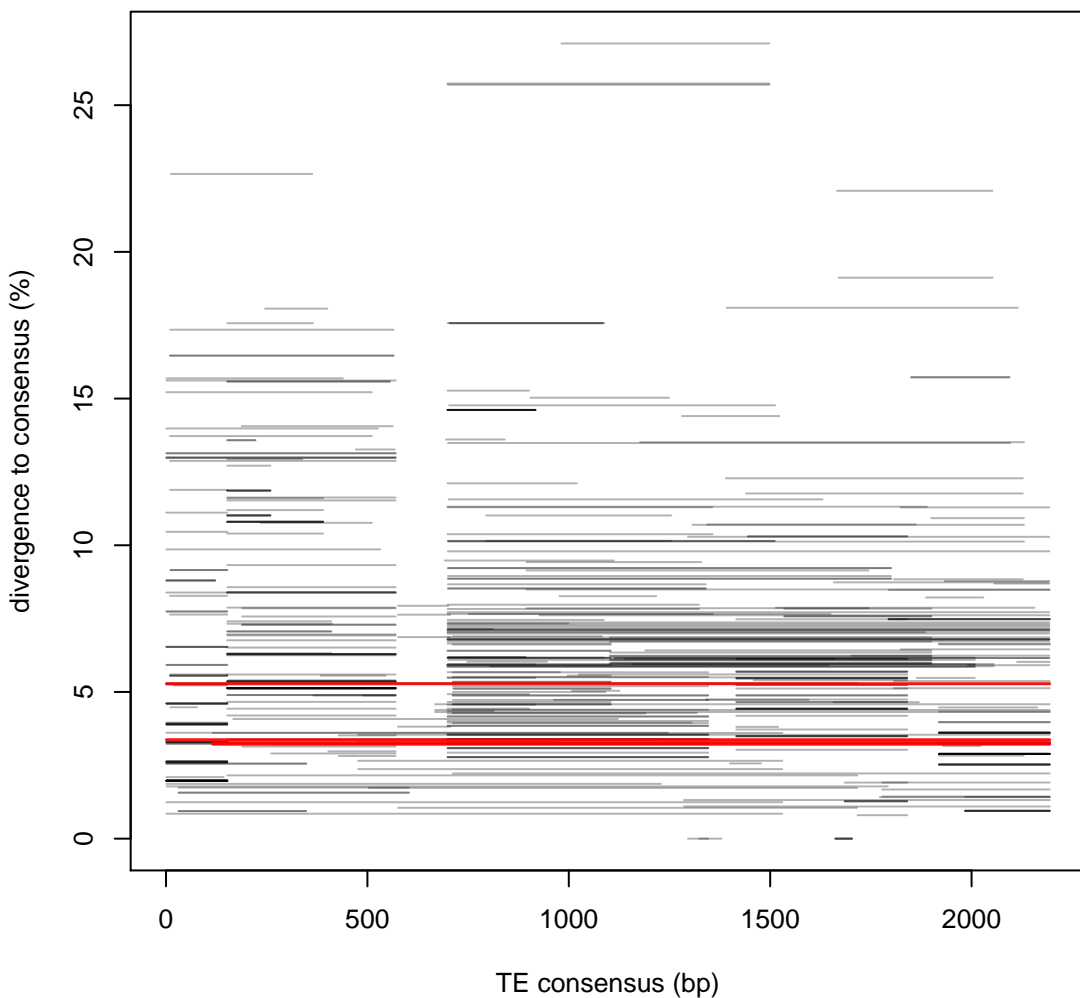
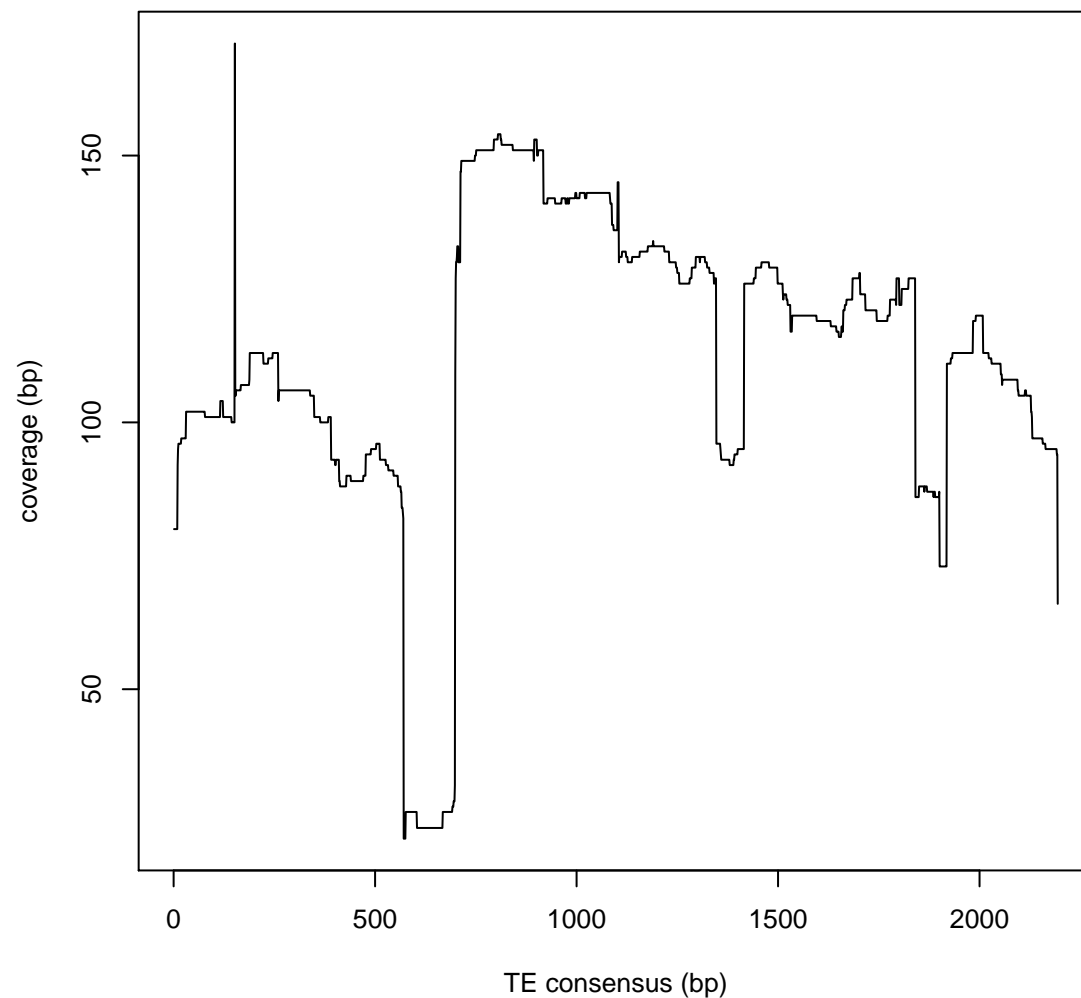


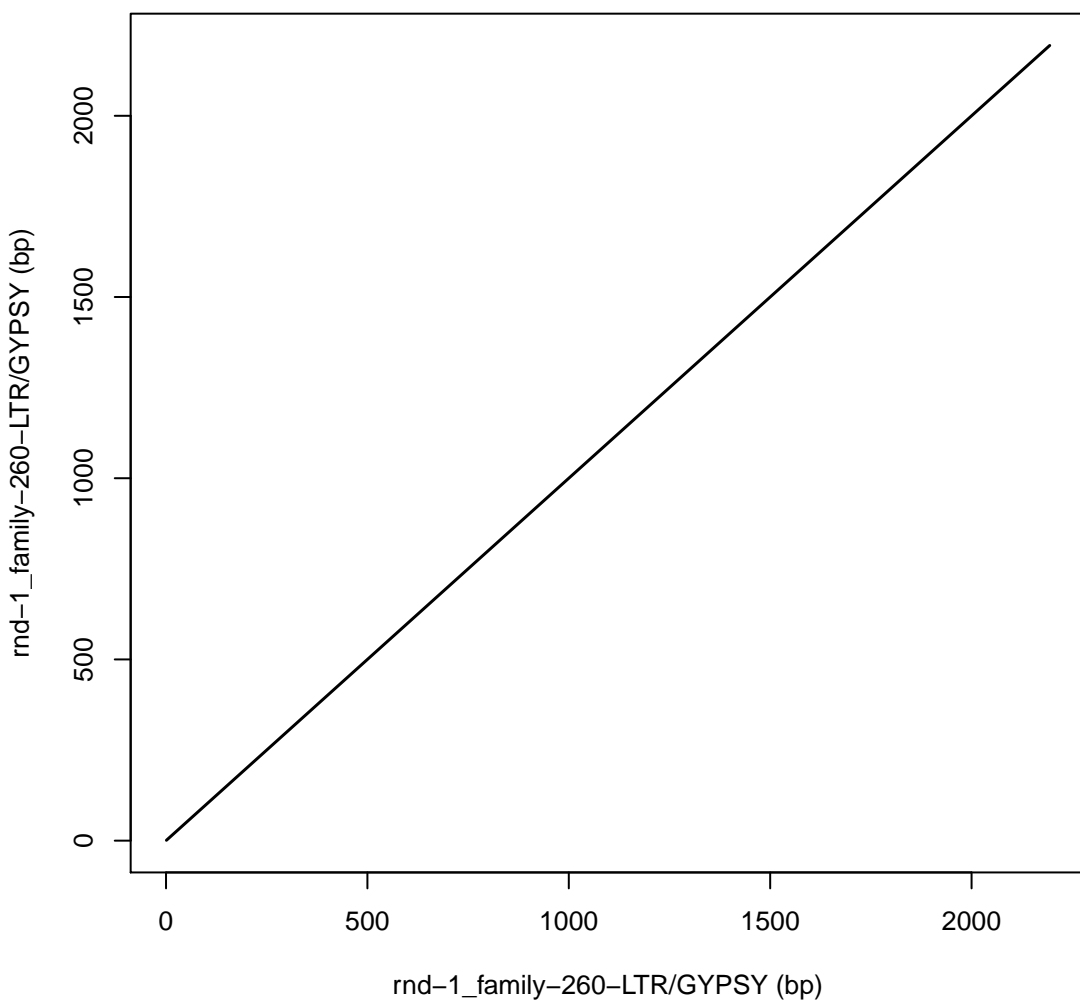
TE: rnd-1_family-260-LTR/GYPSY
consensus size: 2194bp; fragments: 500; full length: 4 (≥ 1974.6 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

