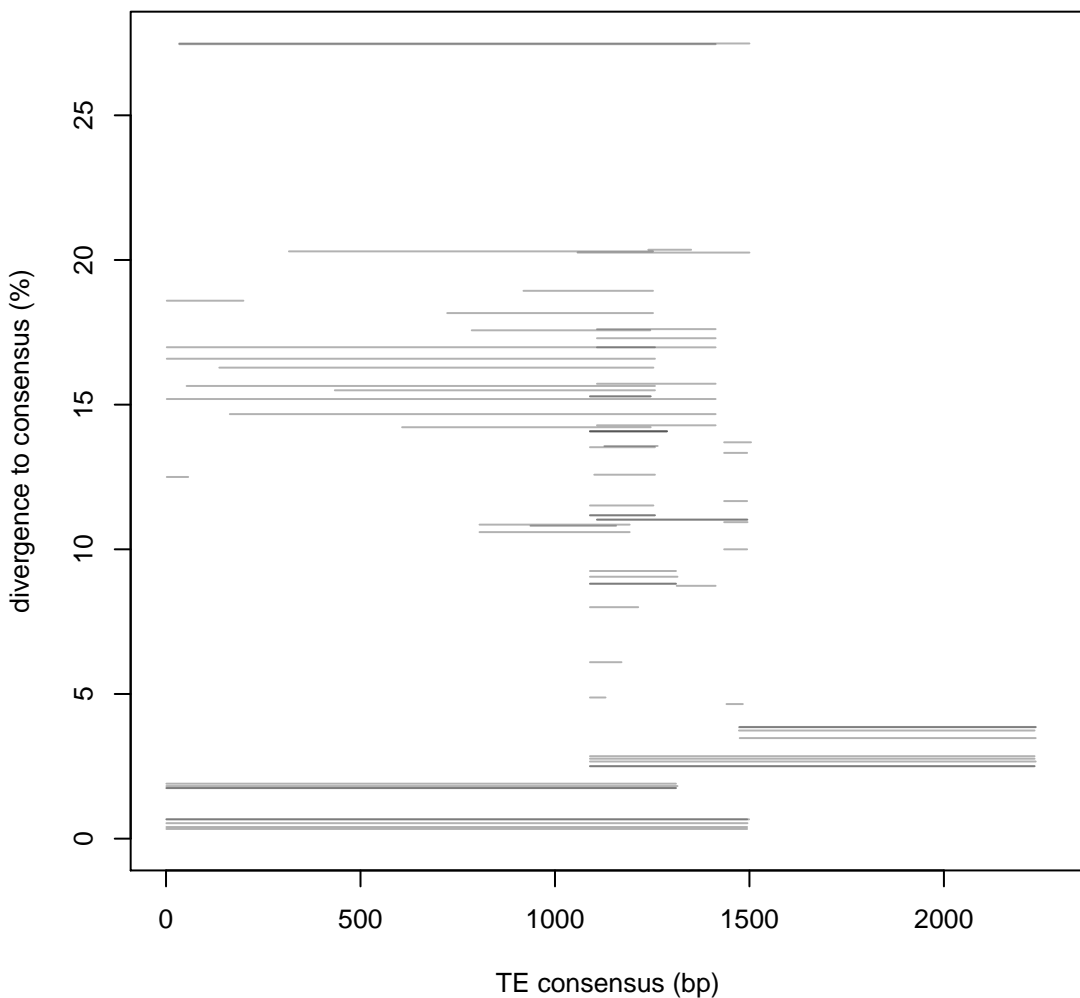
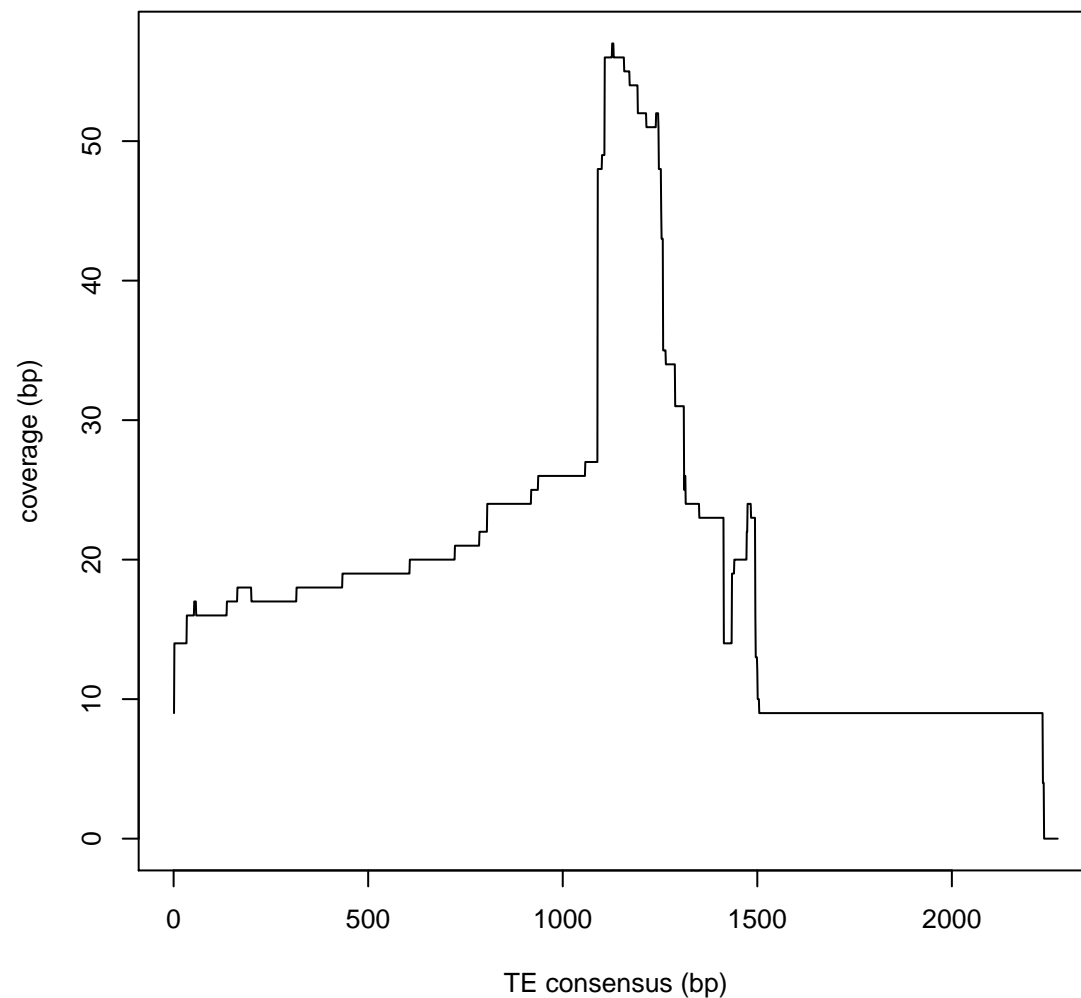


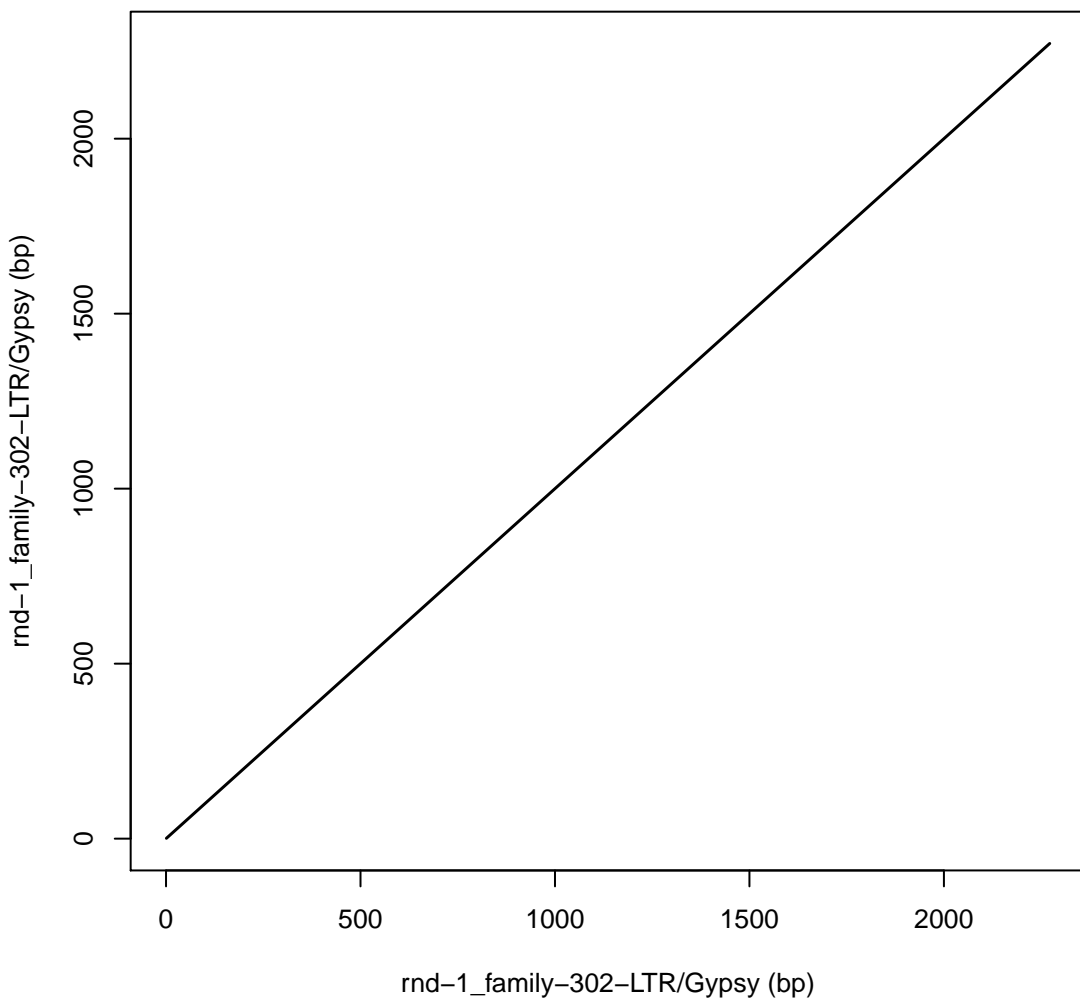
TE: rnd-1_family-302-LTR/Gypsy
consensus size: 2272bp; fragments: 71; full length: 0 (≥ 2044.8 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

