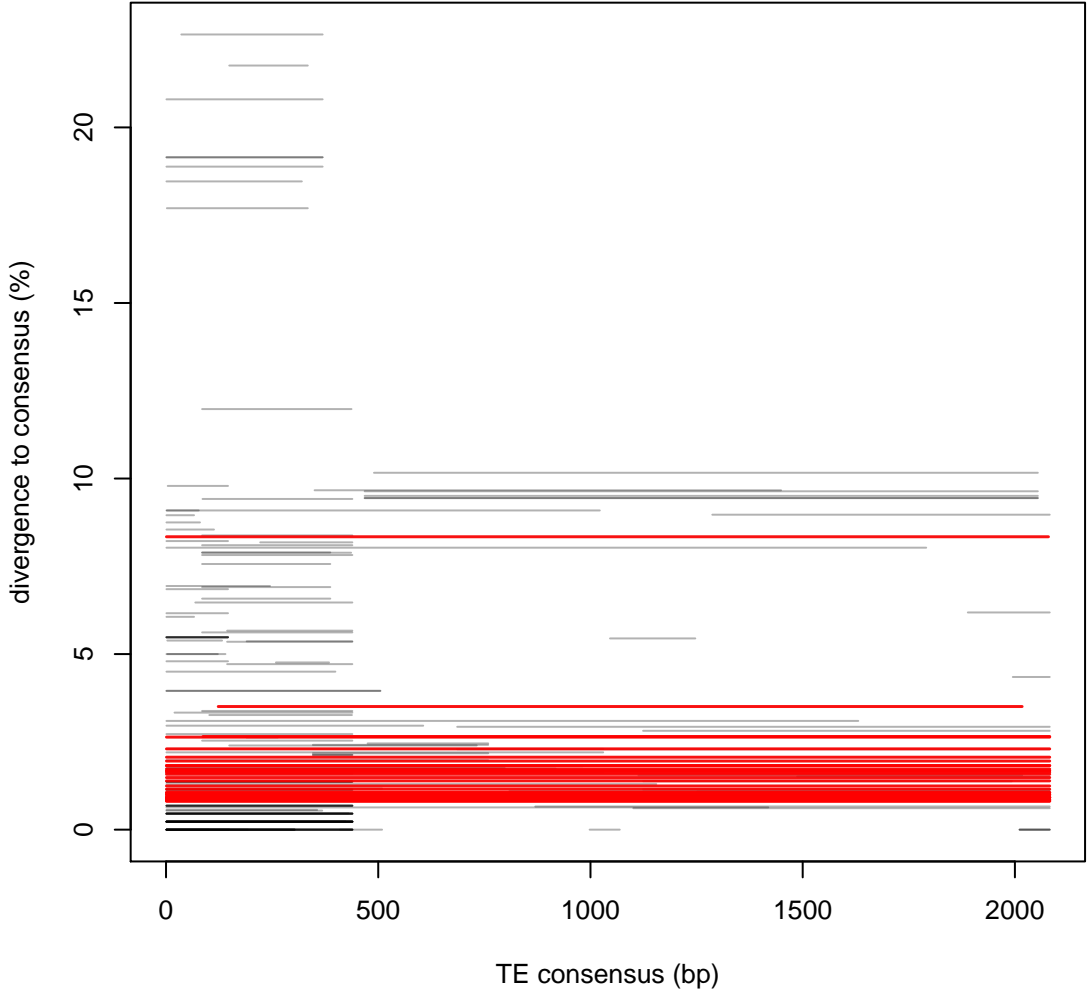
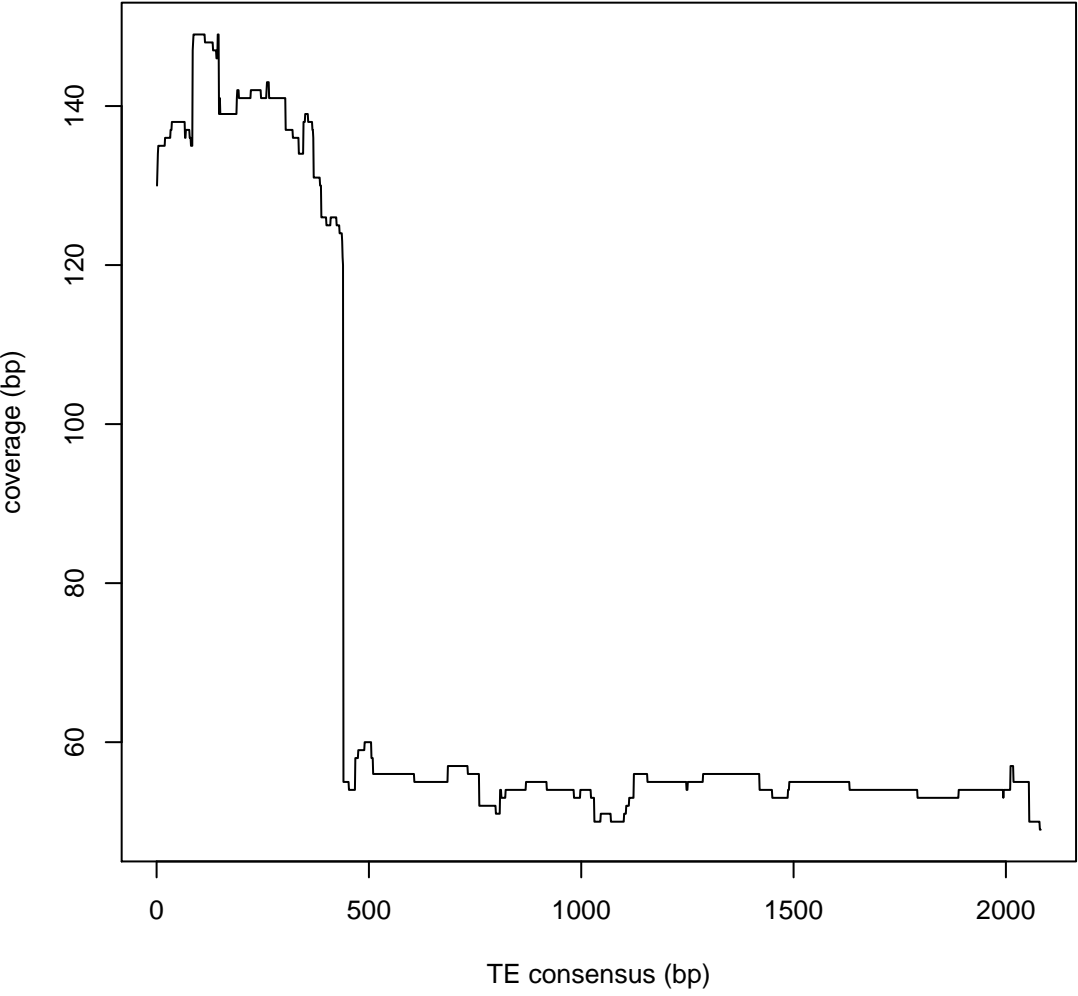


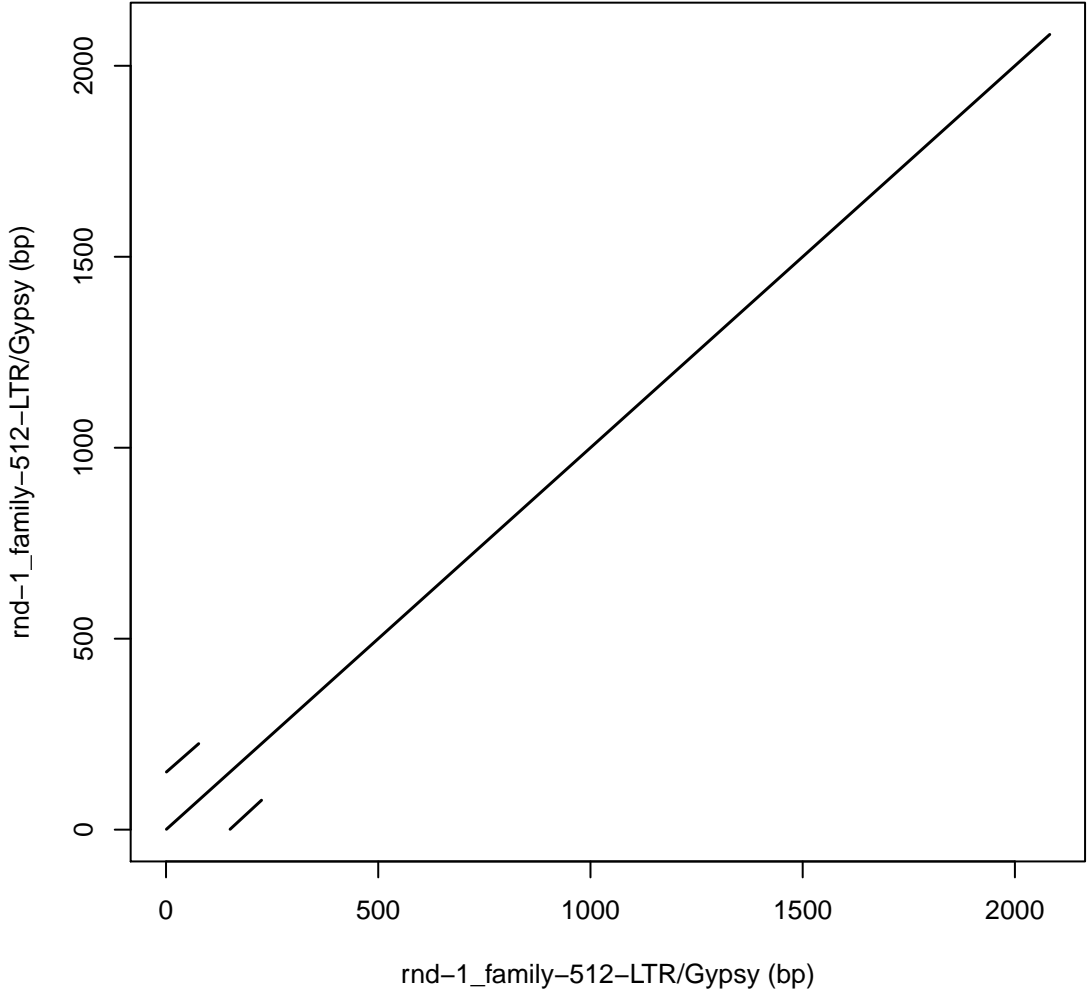
TE: rnd-1_family-512-LTR/Gypsy
consensus size: 2082bp; fragments: 202; full length: 32 (>=1873.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

