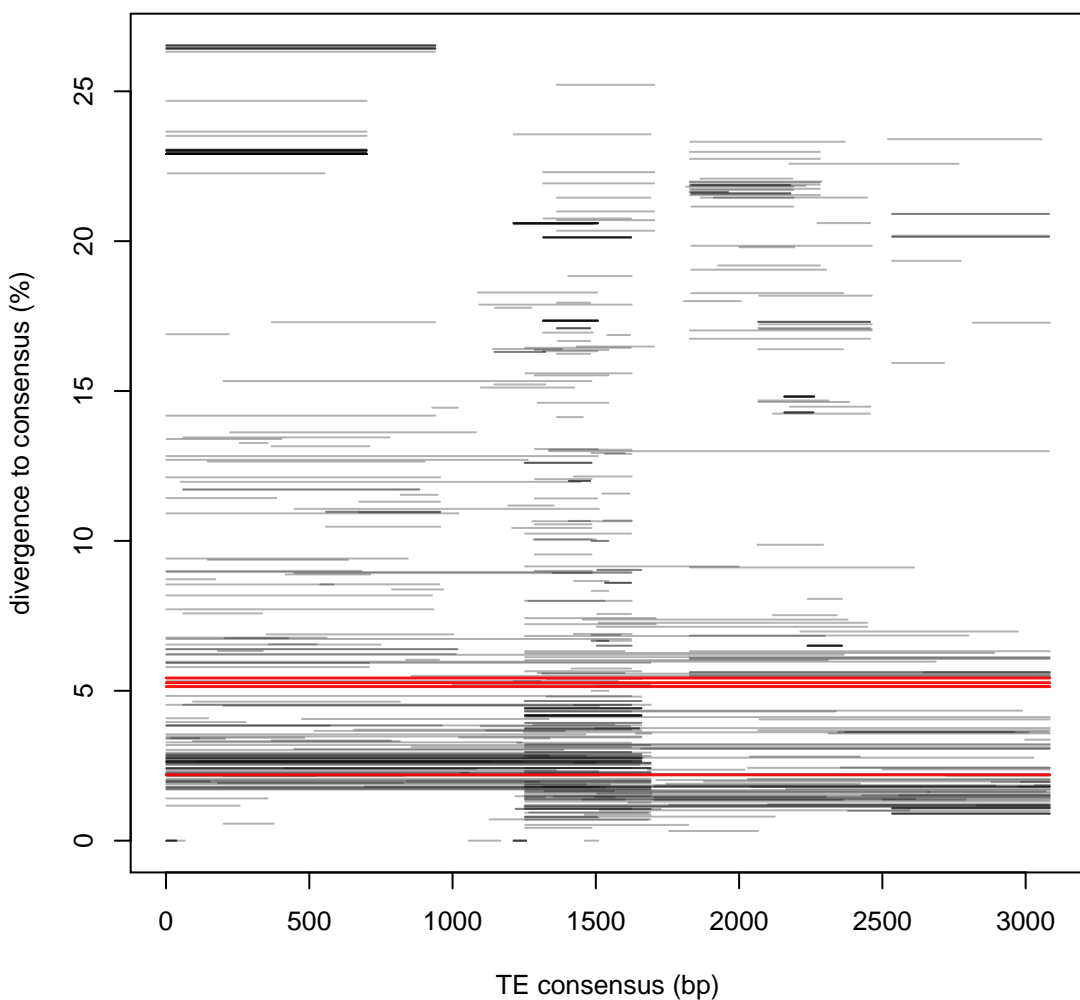
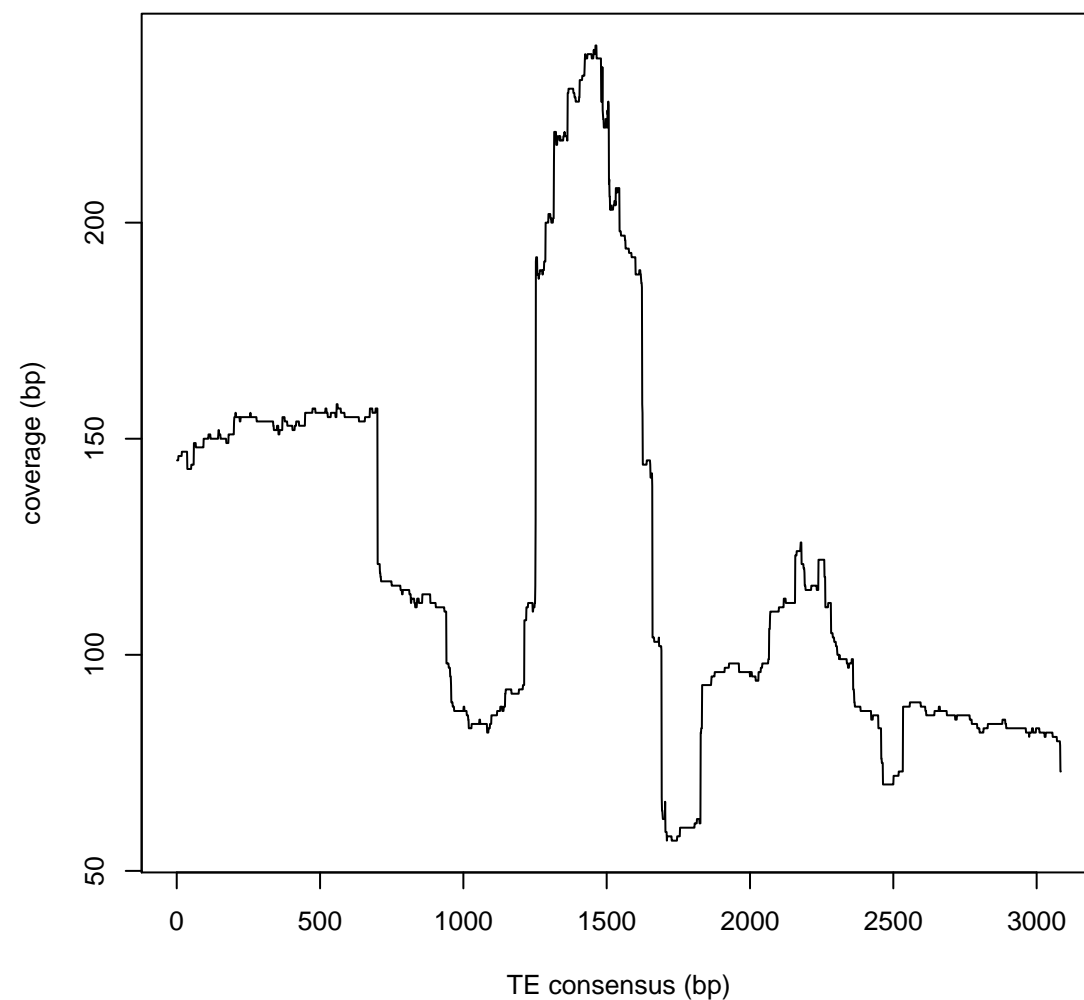


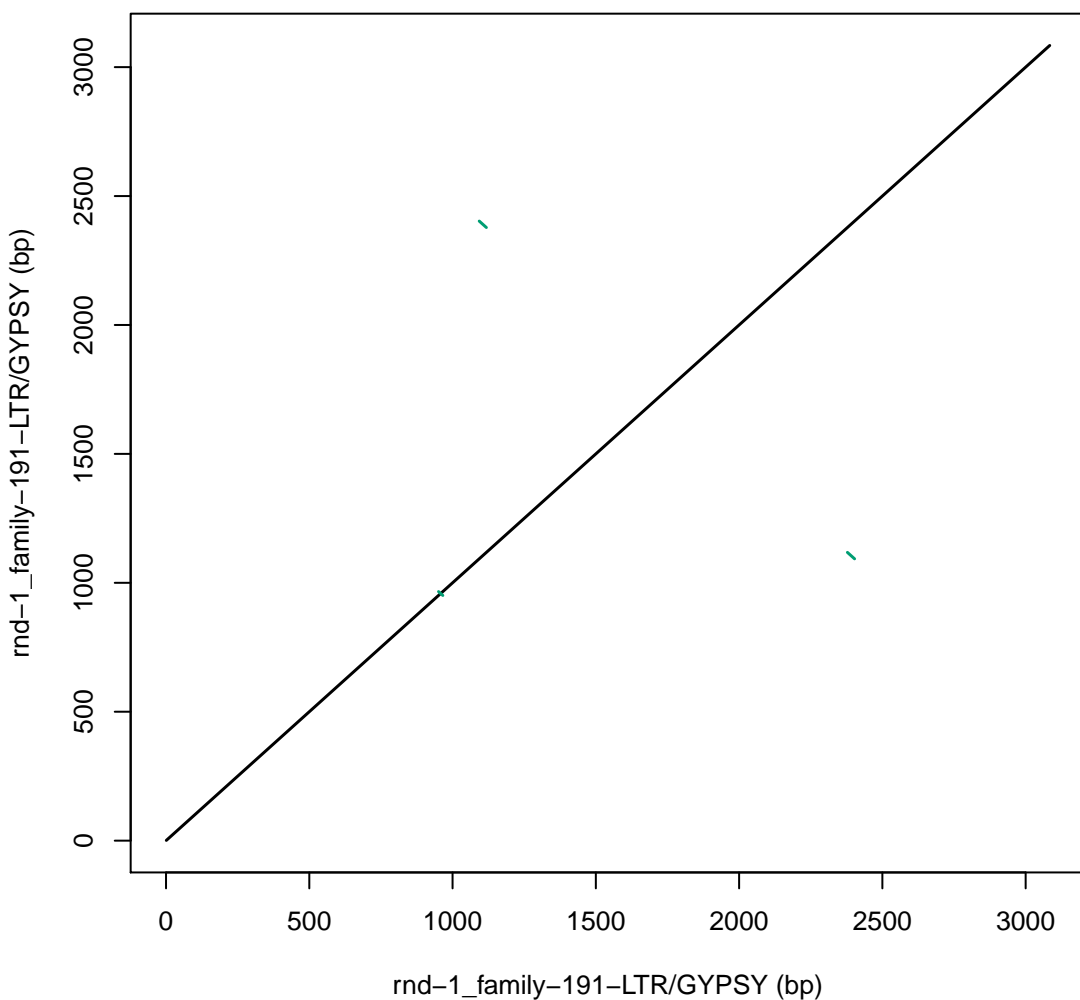
TE: rnd-1_family-191-LTR/GYPSY
consensus size: 3084bp; fragments: 542; full length: 4 (>=2775.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

