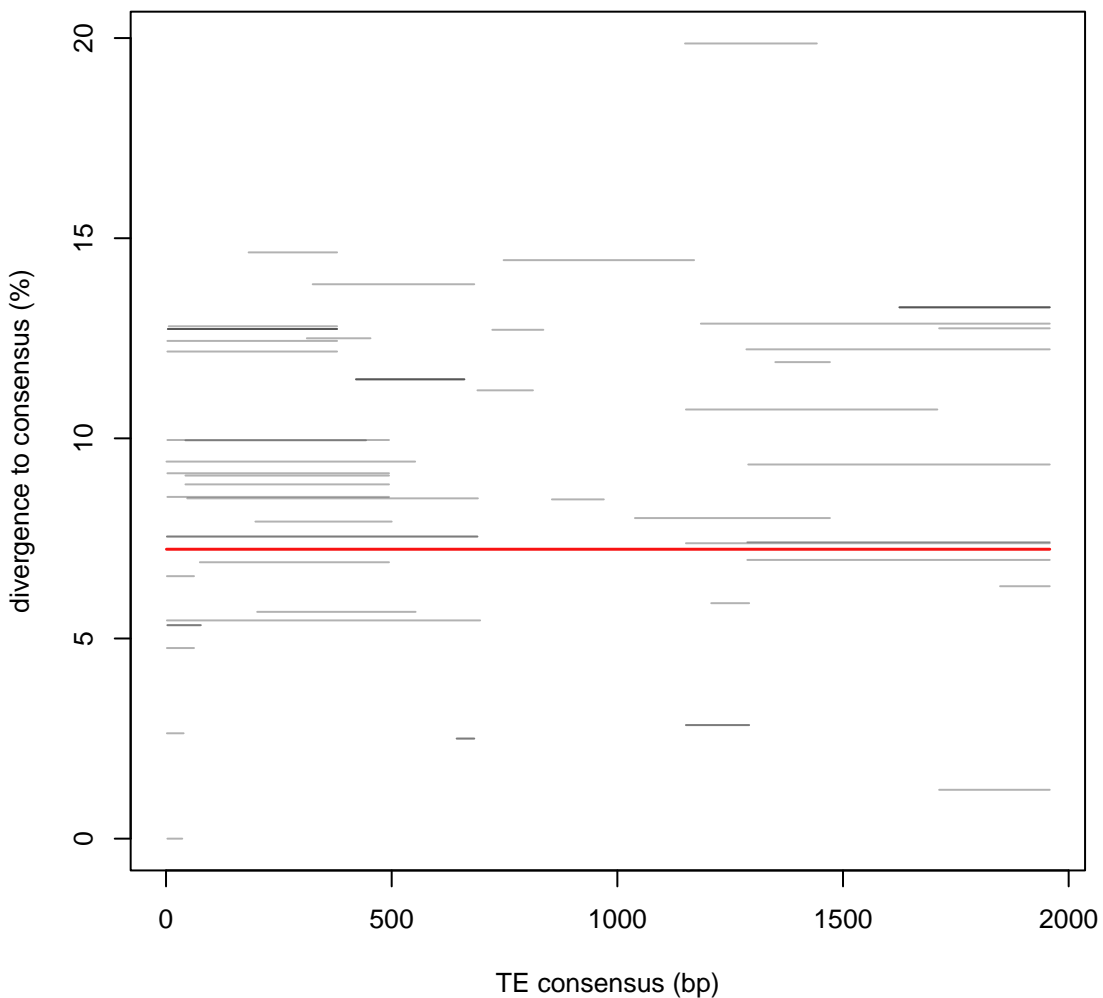
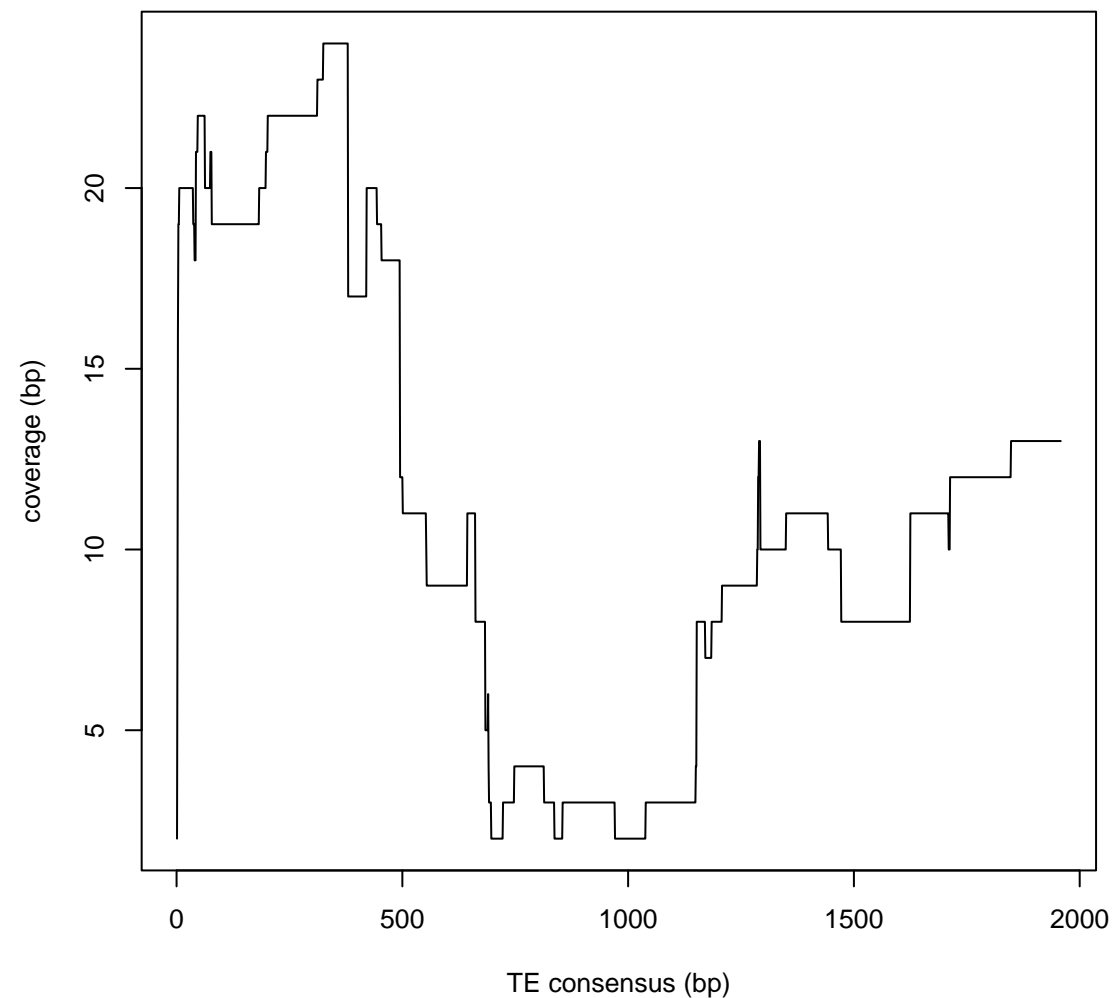


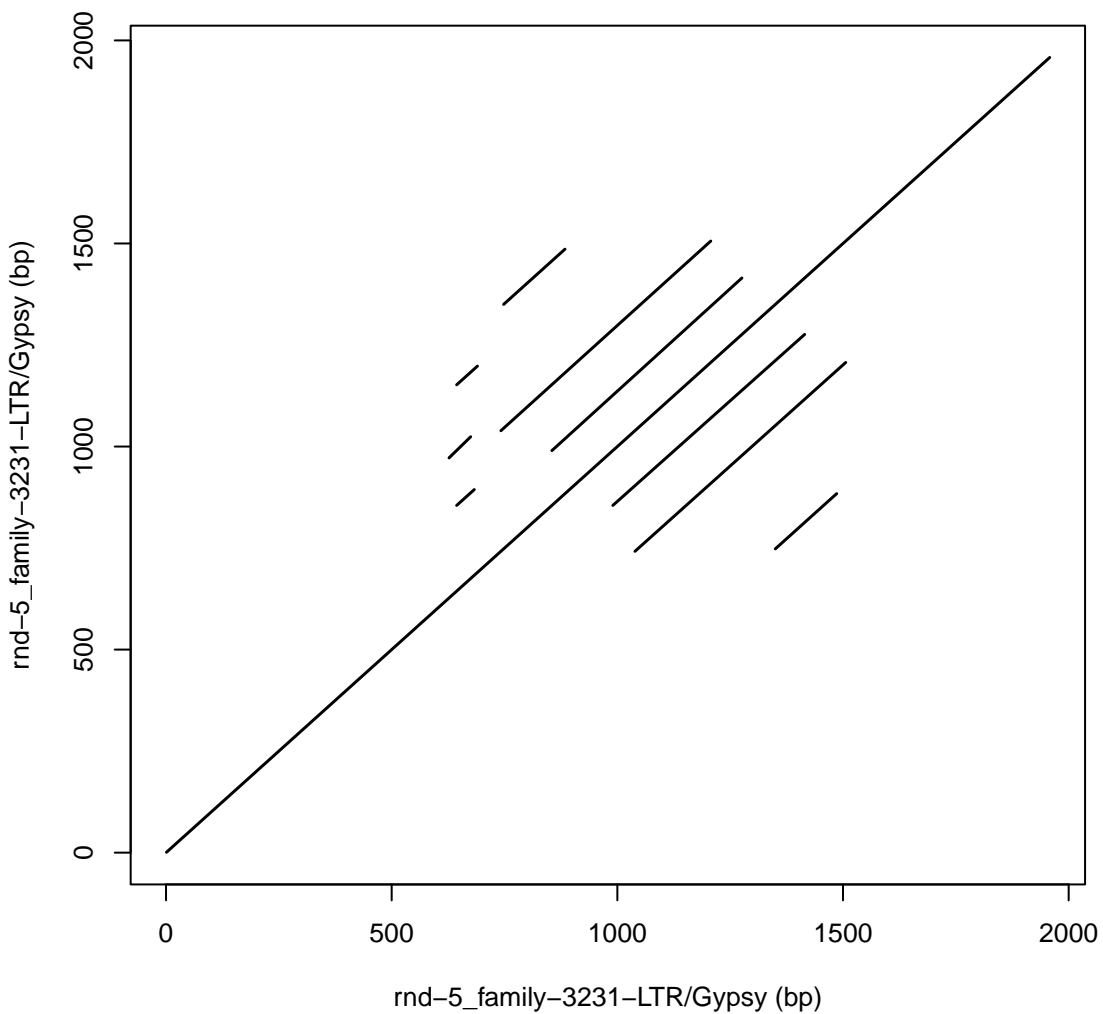
TE: rnd-5_family-3231-LTR/Gypsy
consensus size: 1958bp; fragments: 58; full length: 1 (>=1762.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

