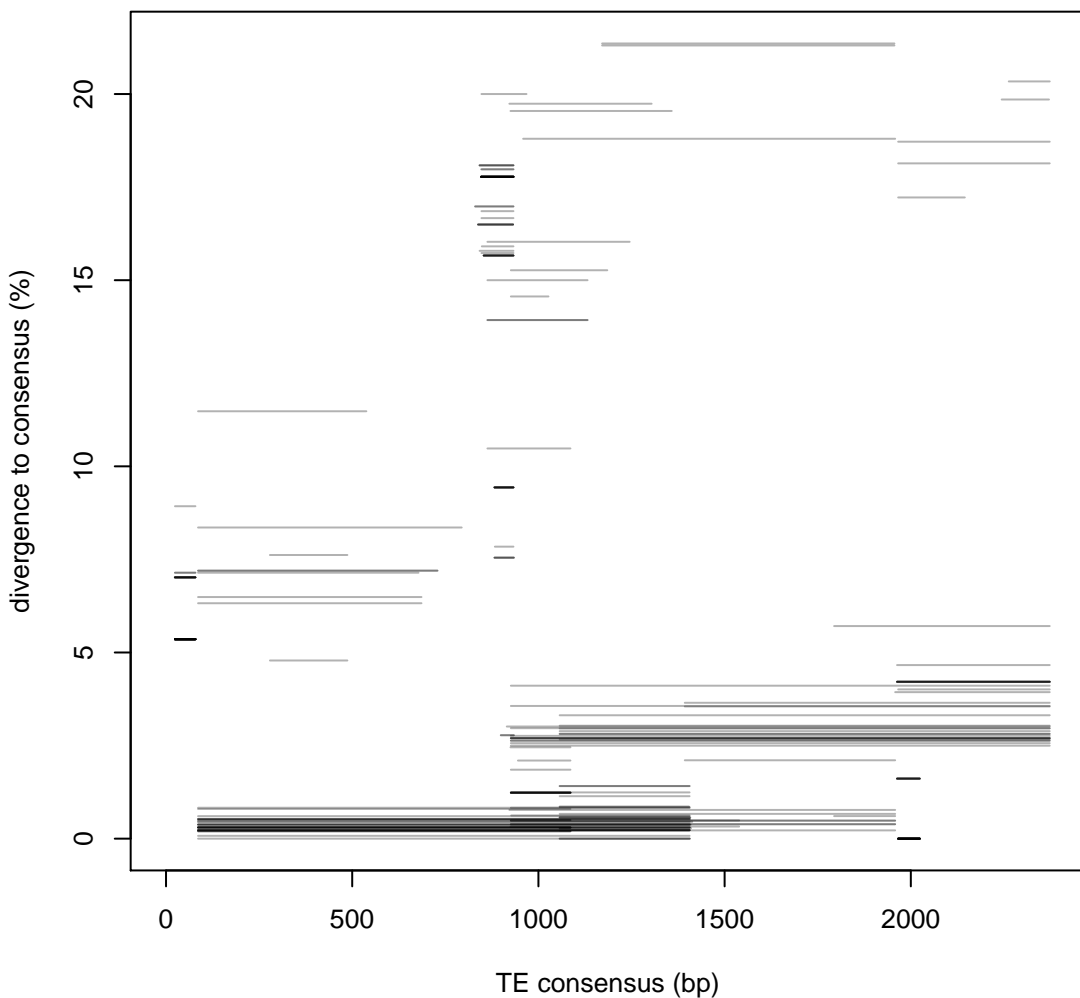
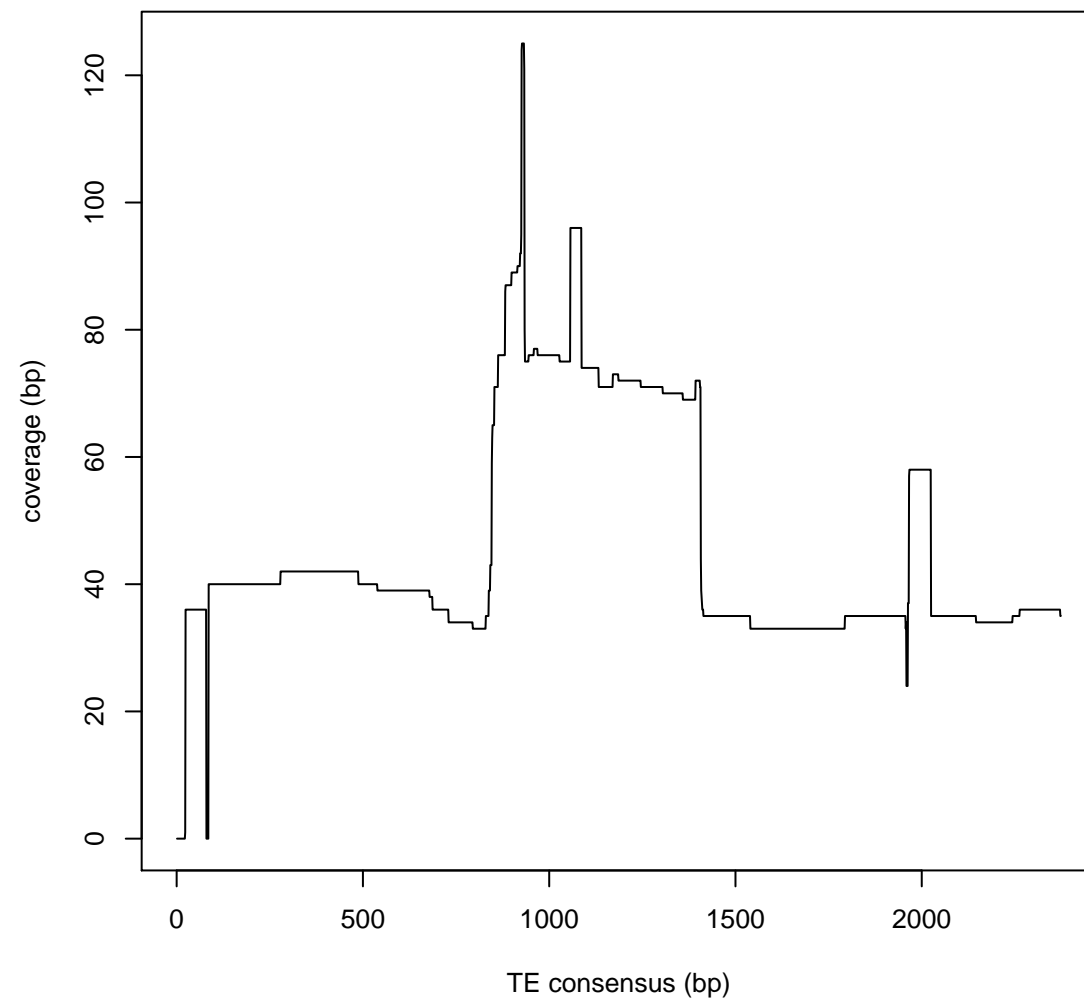


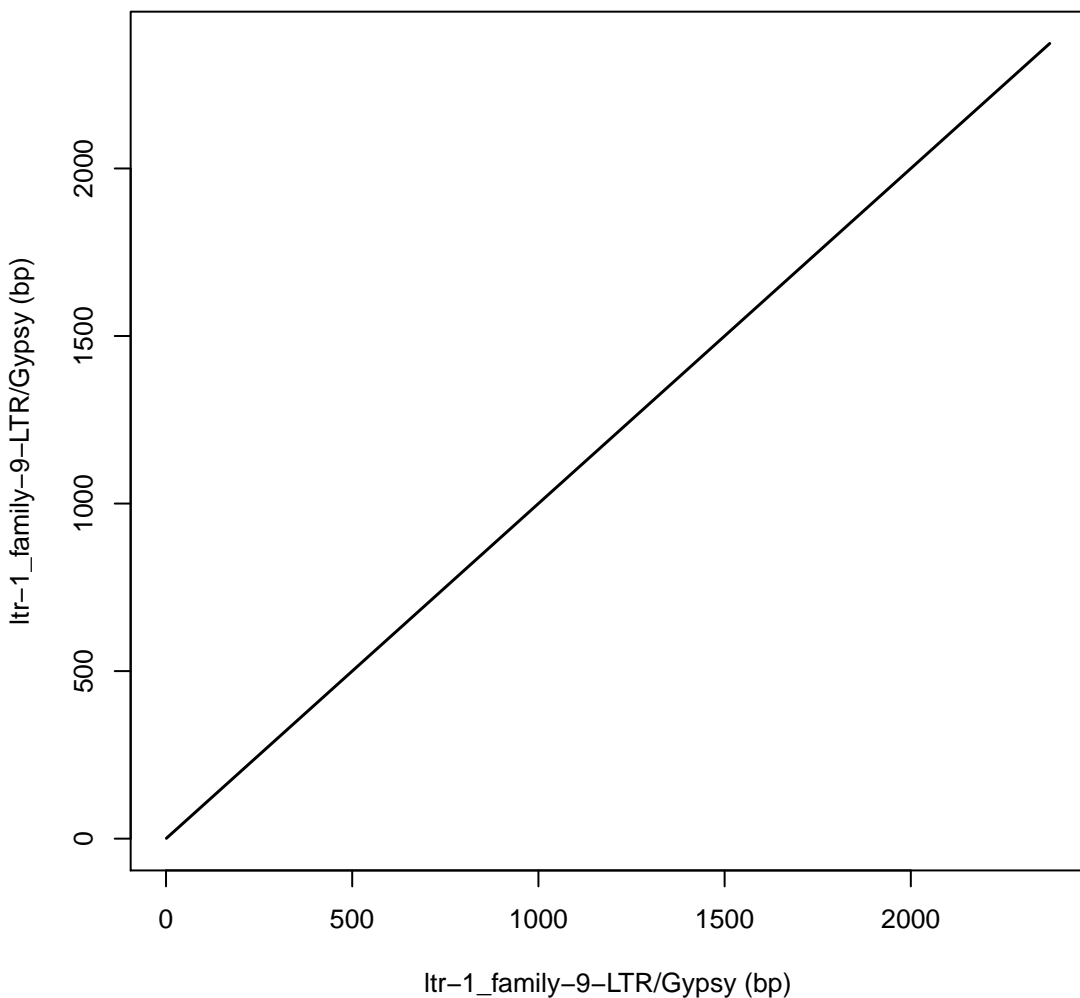
TE: ltr-1_family-9-LTR/Gypsy
consensus size: 2373bp; fragments: 237; full length: 0 (>=2135.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

