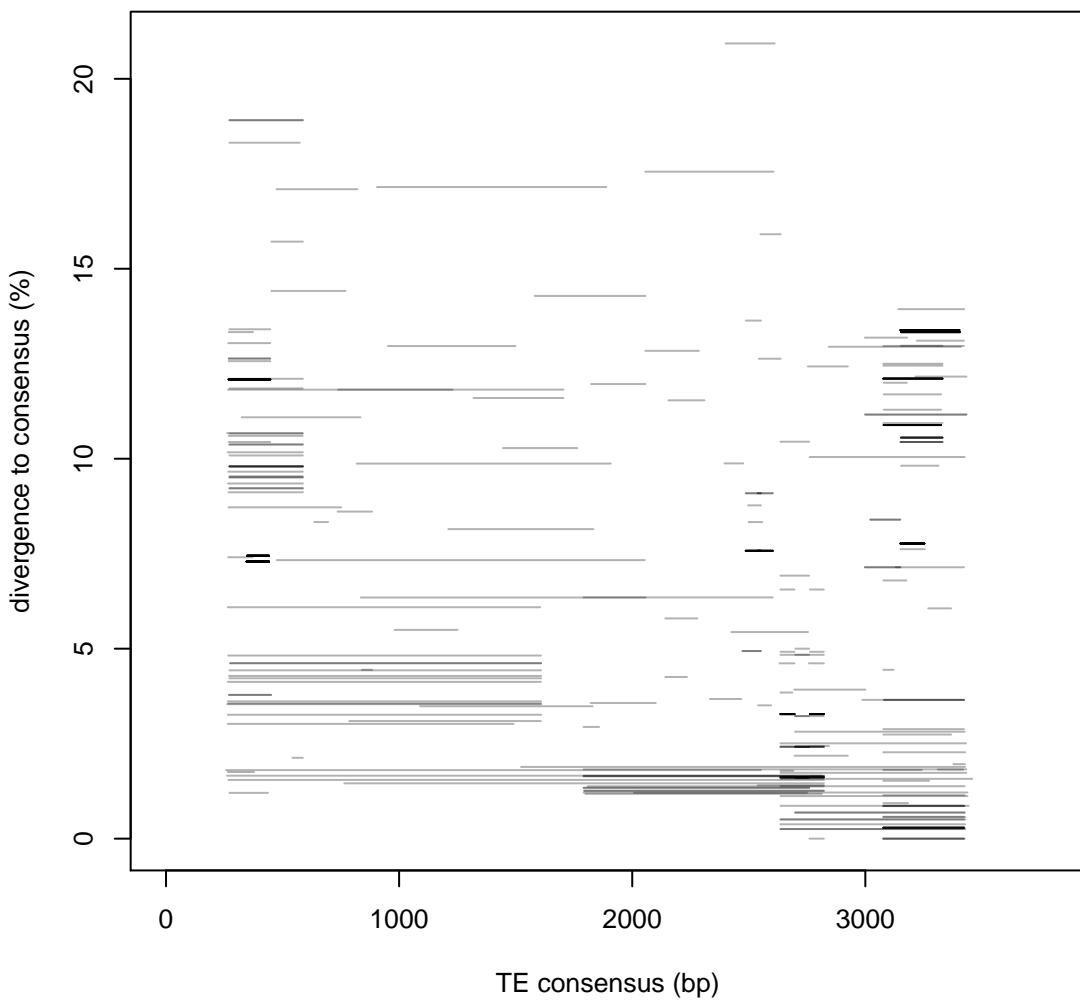
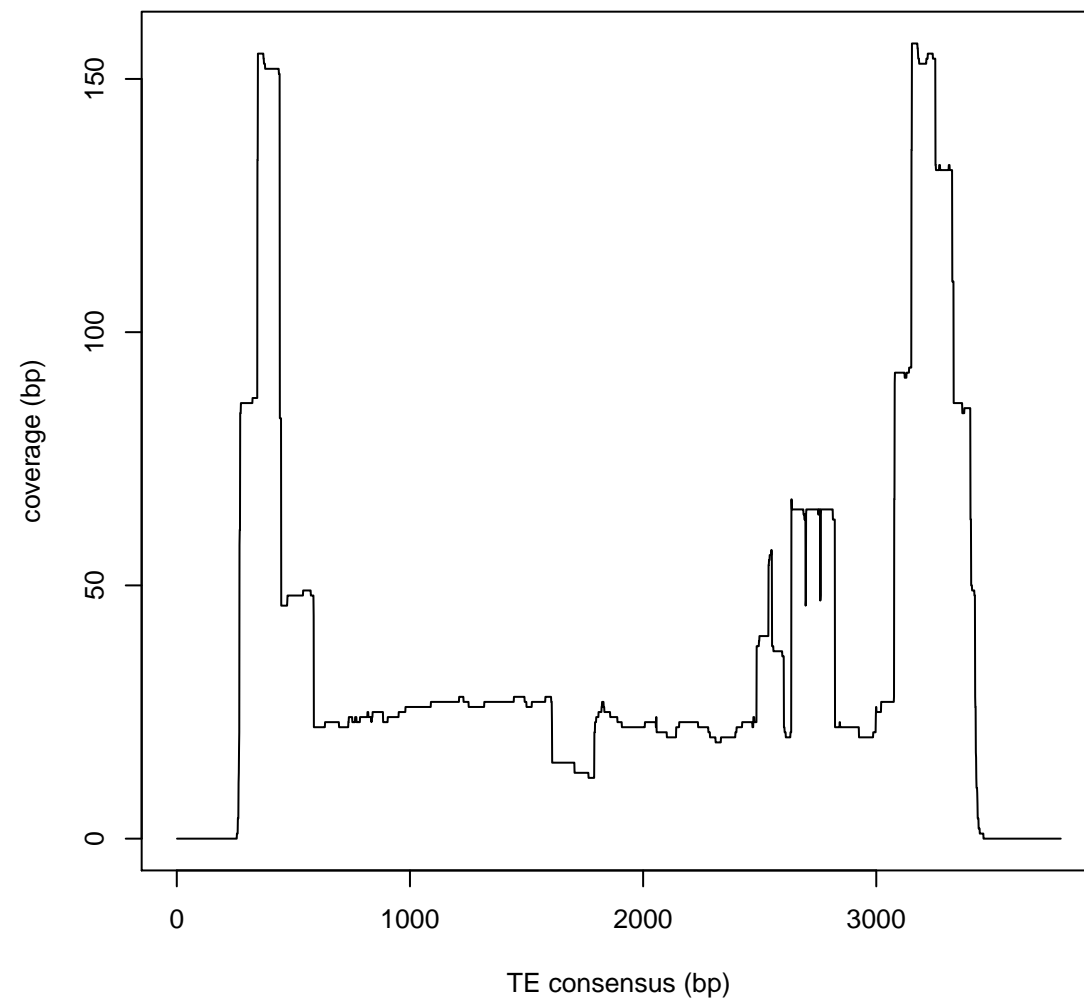


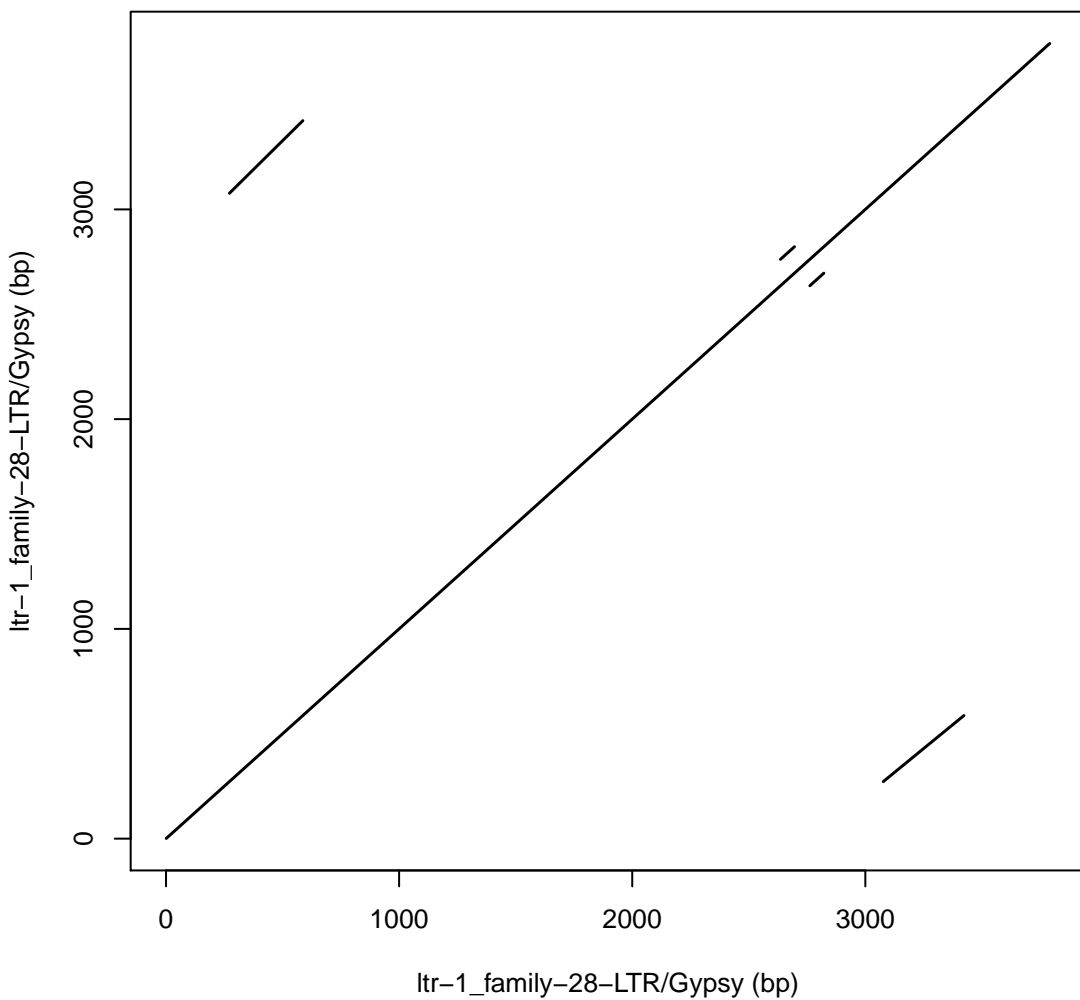
TE: ltr-1_family-28-LTR/Gypsy
consensus size: 3791bp; fragments: 476; full length: 0 (>=3411.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

