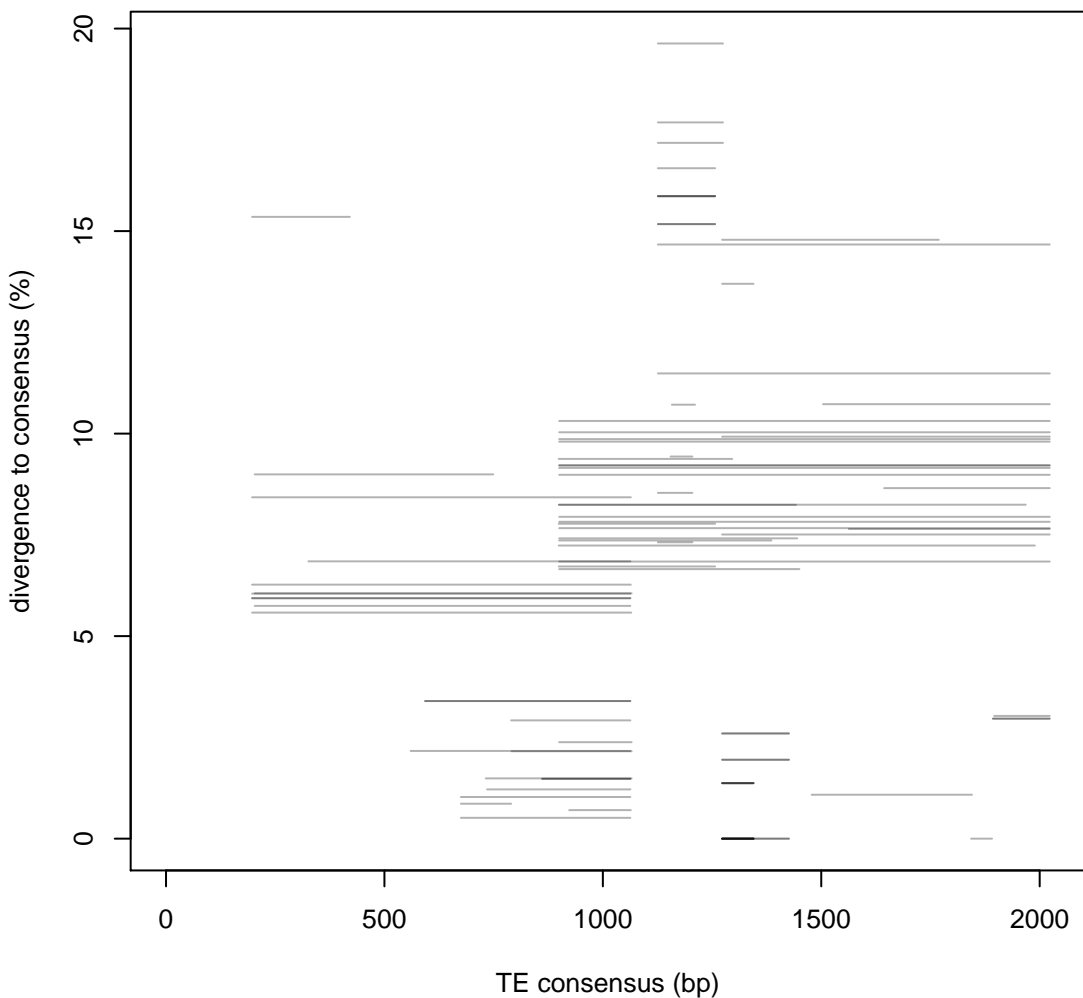
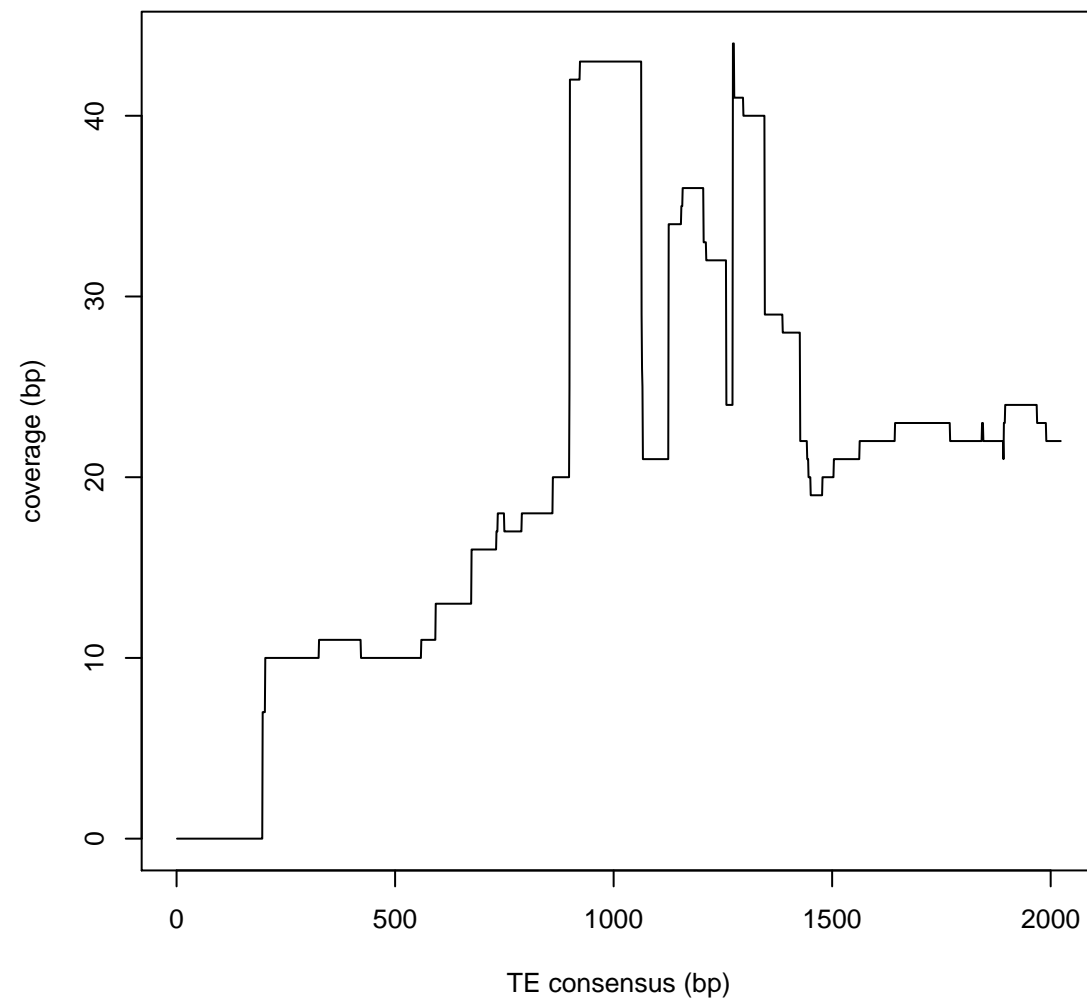


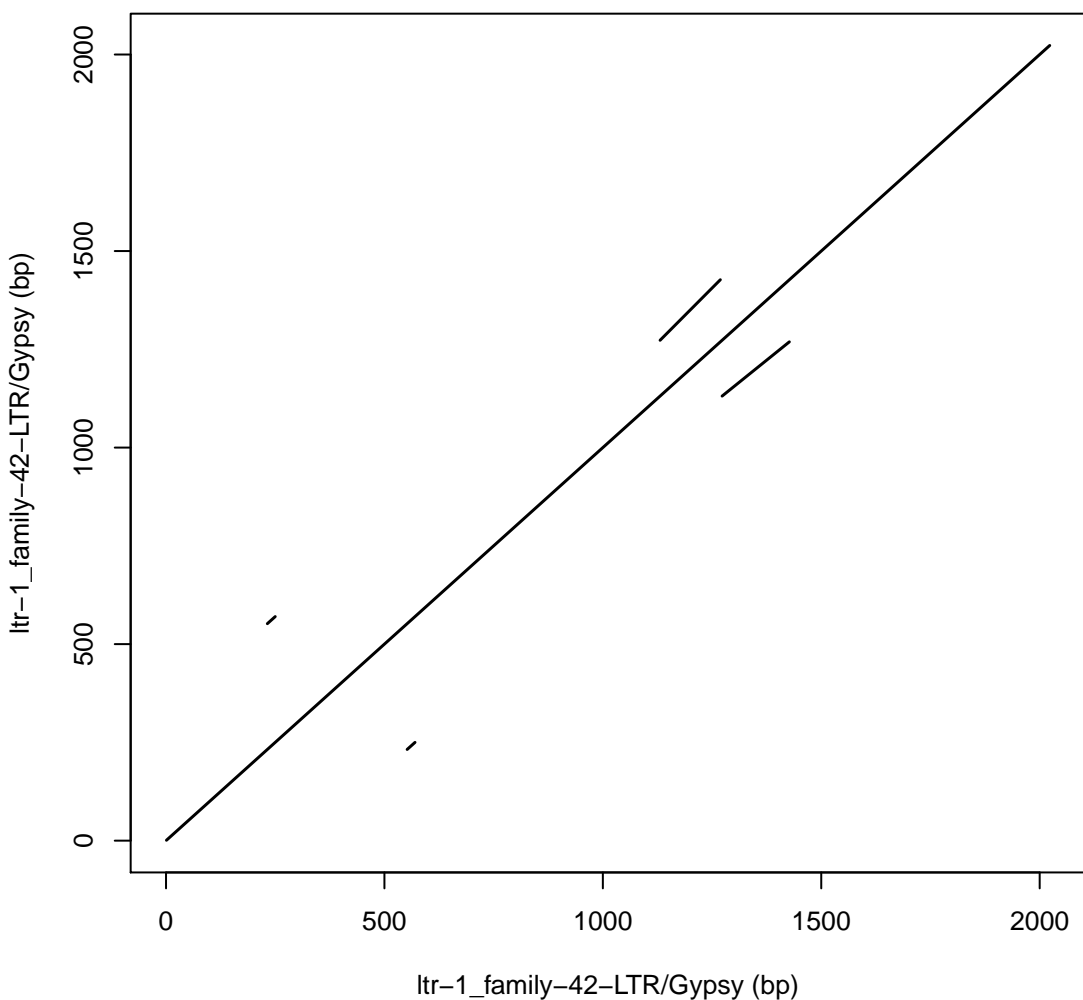
TE: ltr-1_family-42-LTR/Gypsy
consensus size: 2023bp; fragments: 89; full length: 0 (>=1820.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

