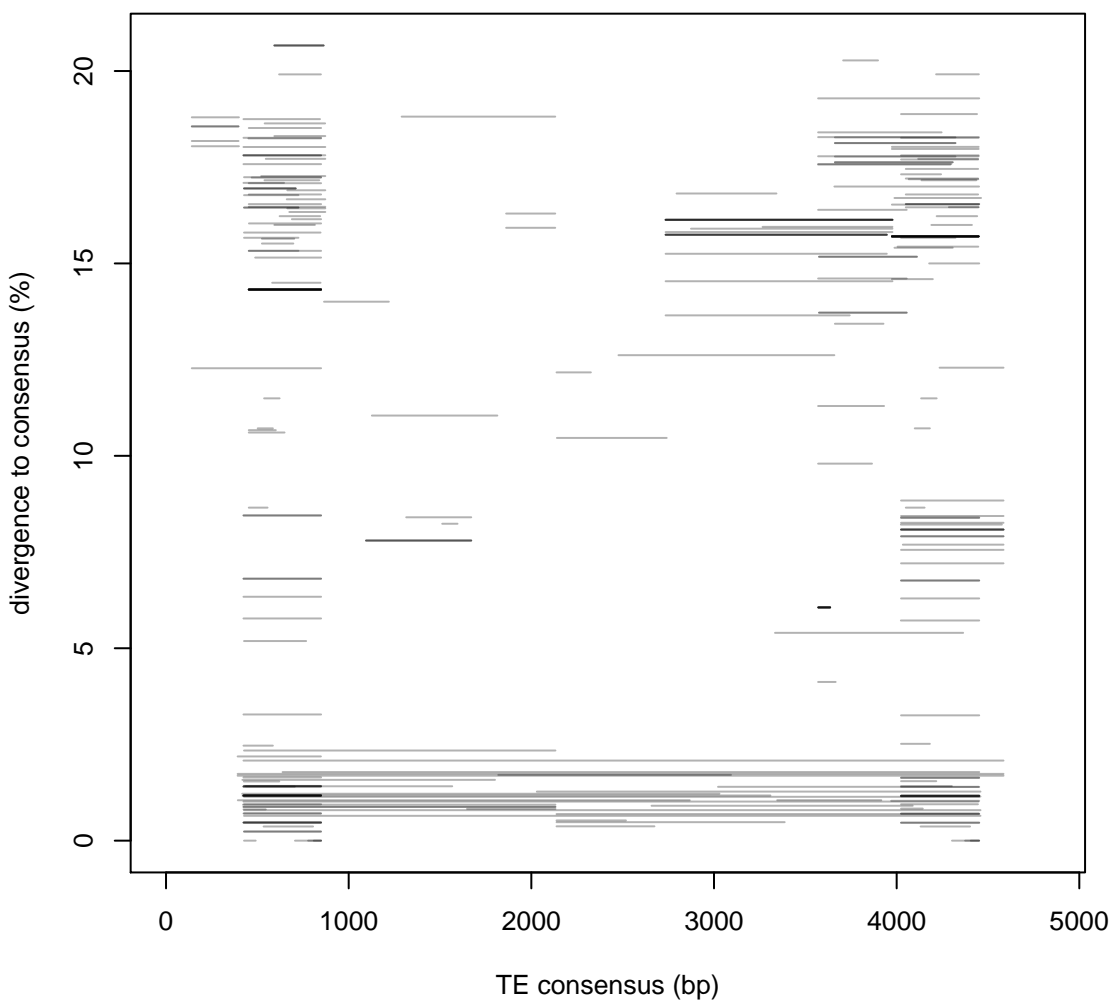
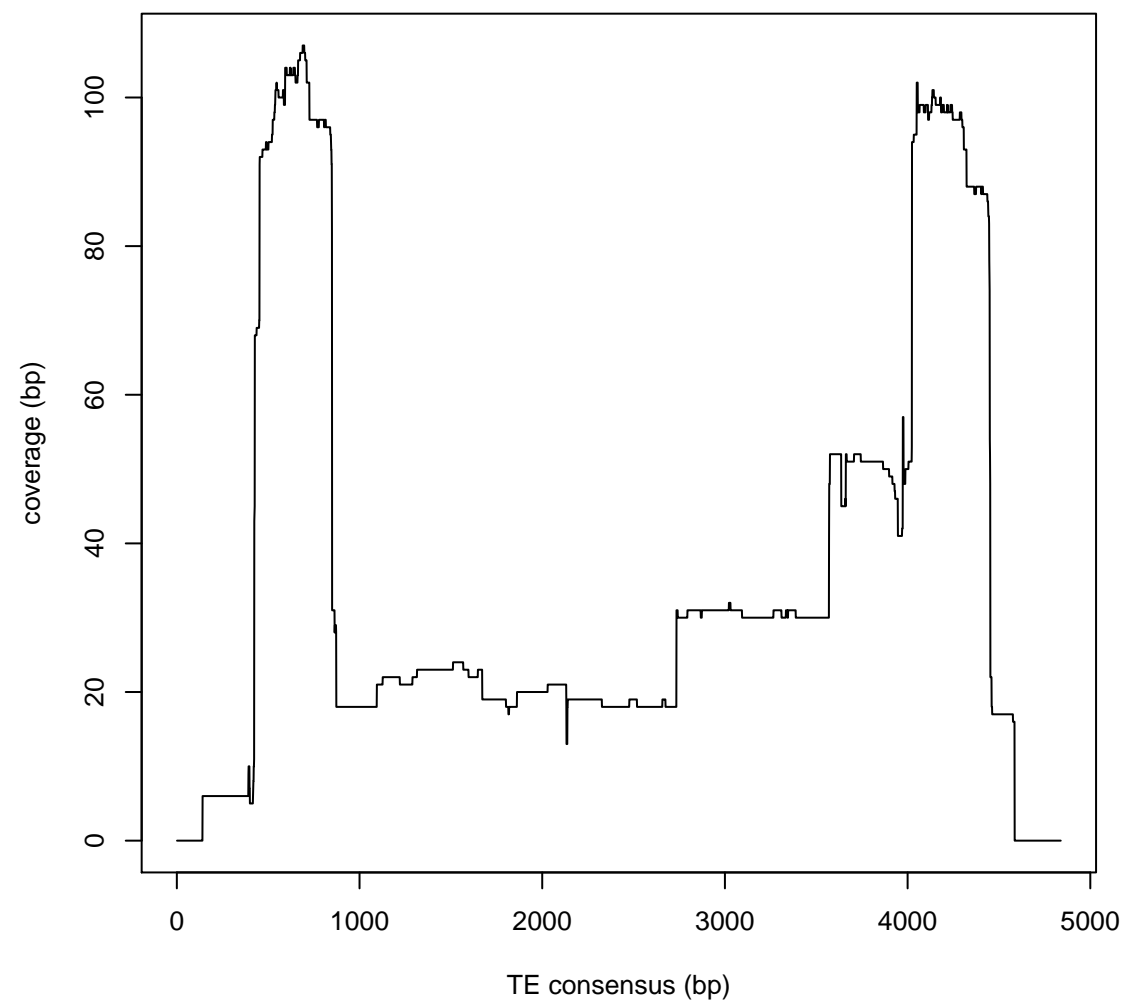


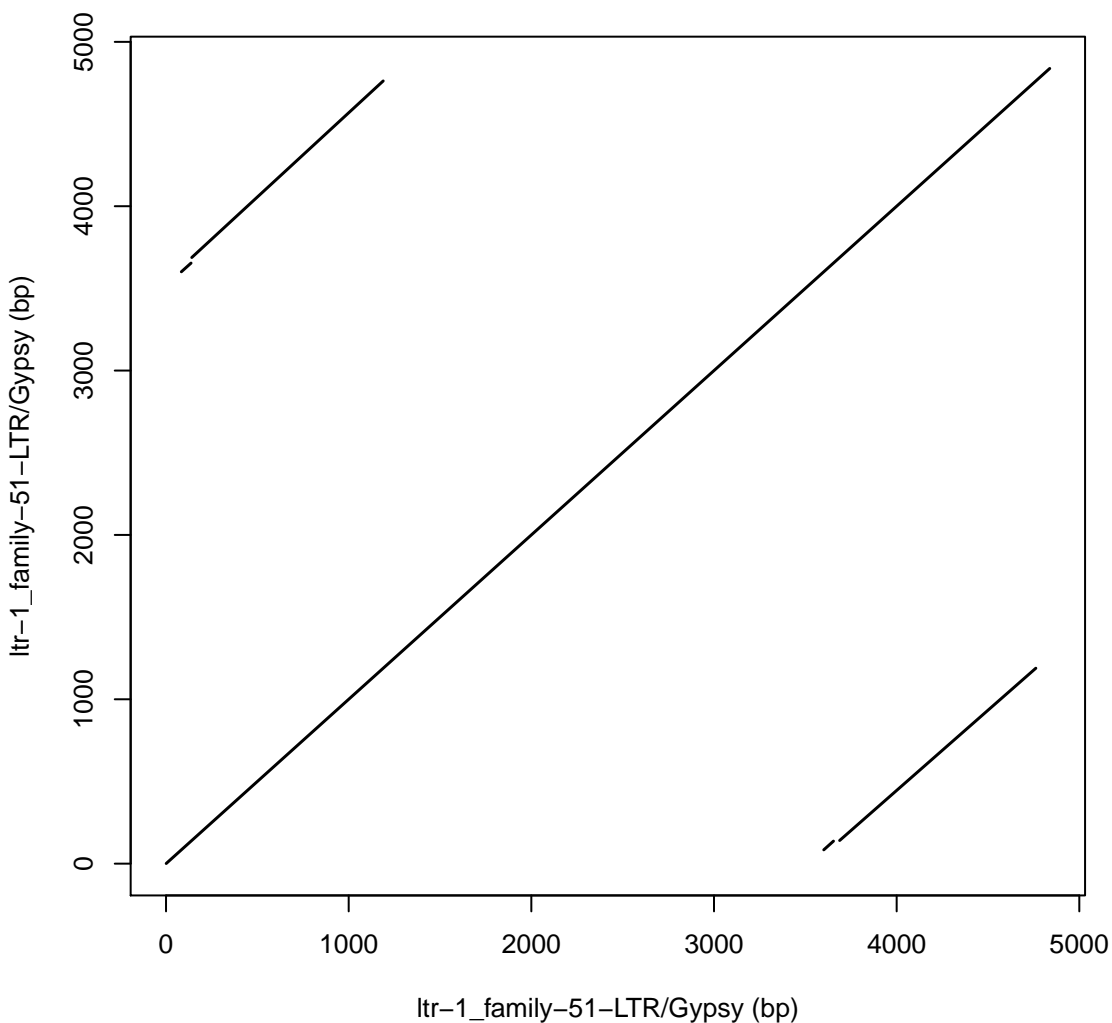
TE: ltr-1_family-51-LTR/Gypsy
consensus size: 4838bp; fragments: 281; full length: 0 (>=4354.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

