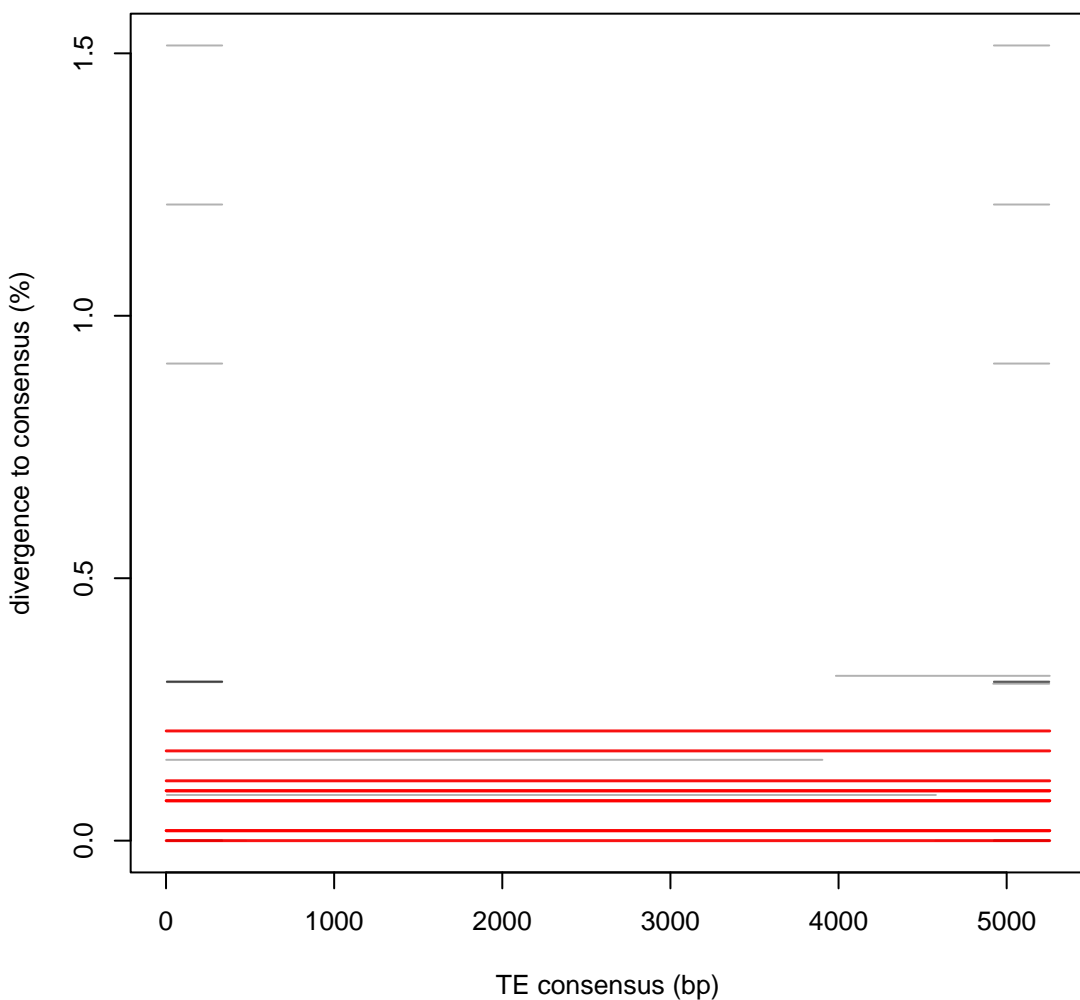
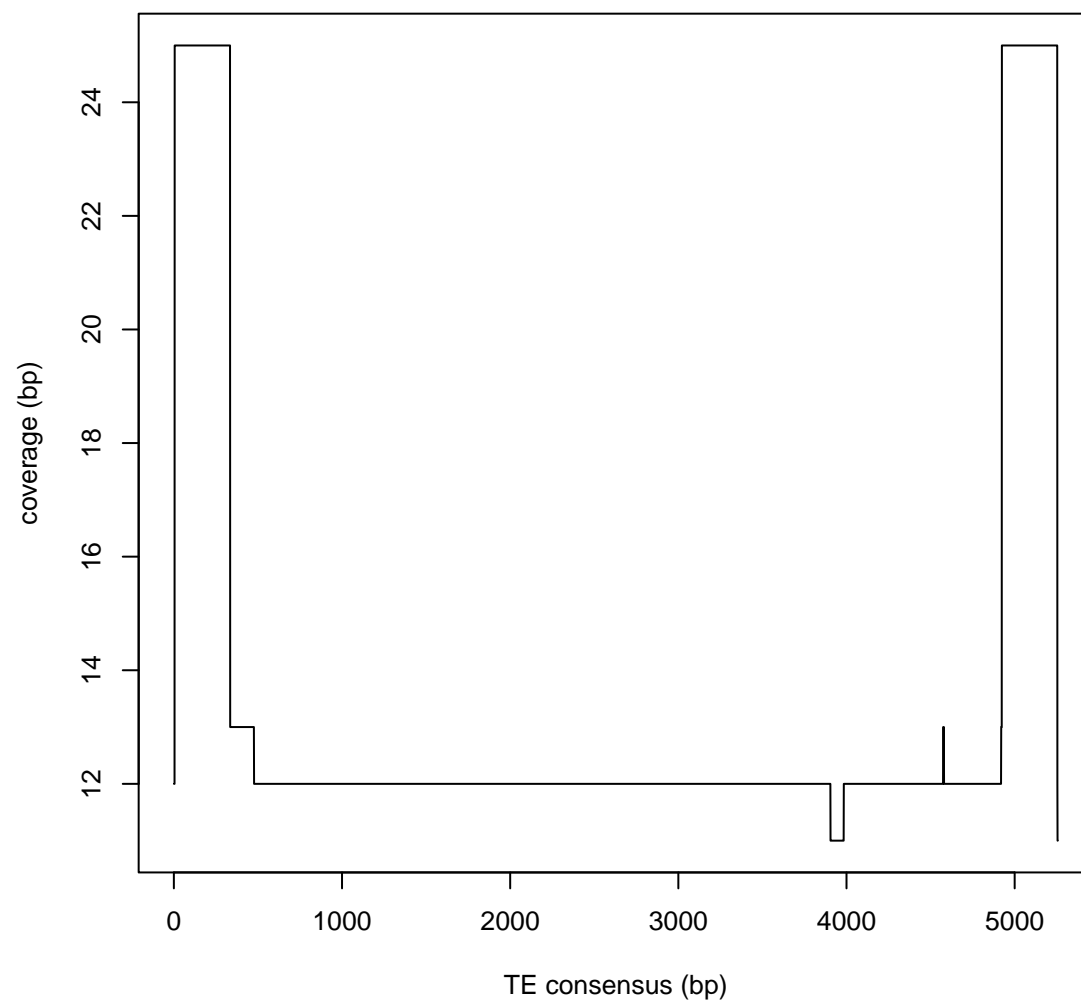


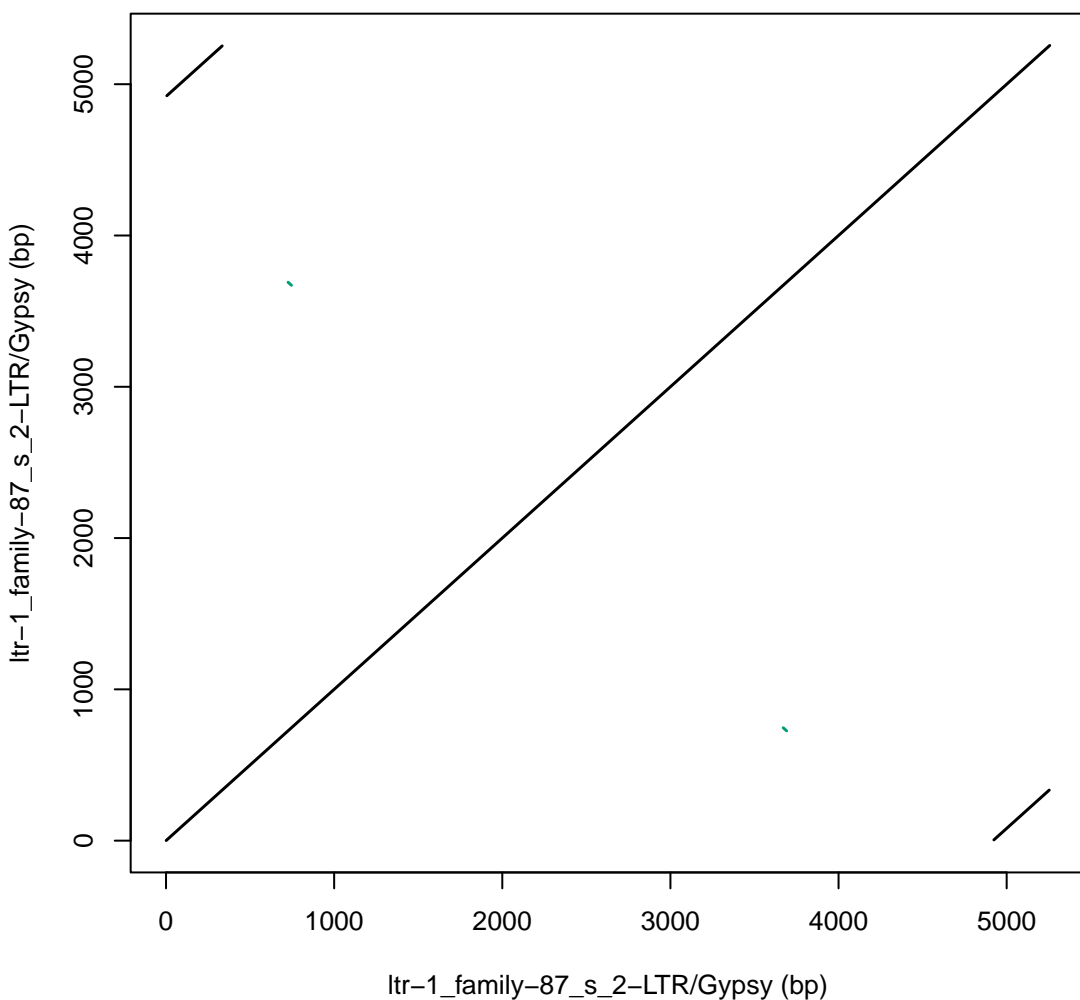
TE: ltr-1_family-87_s_2-LTR/Gypsy
consensus size: 5256bp; fragments: 40; full length: 10 (>=4730.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

