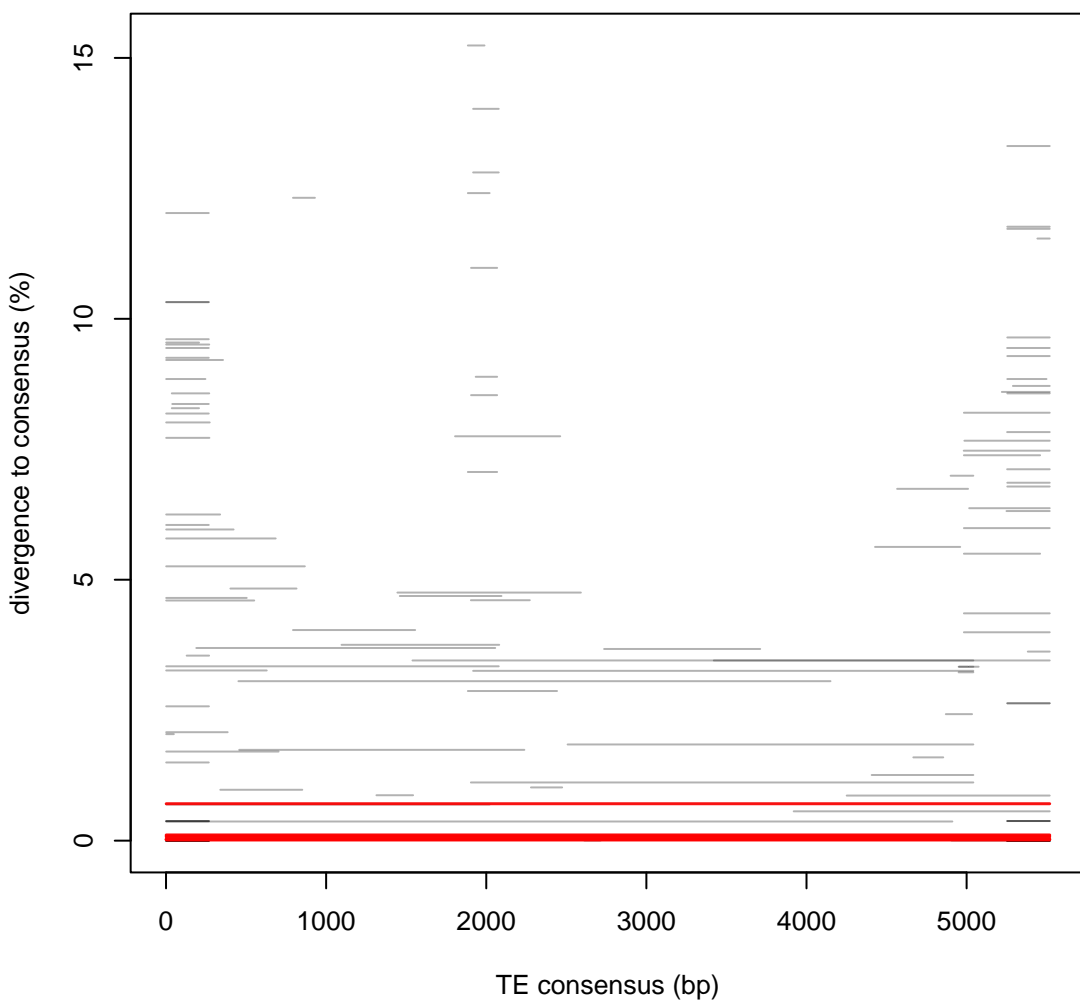
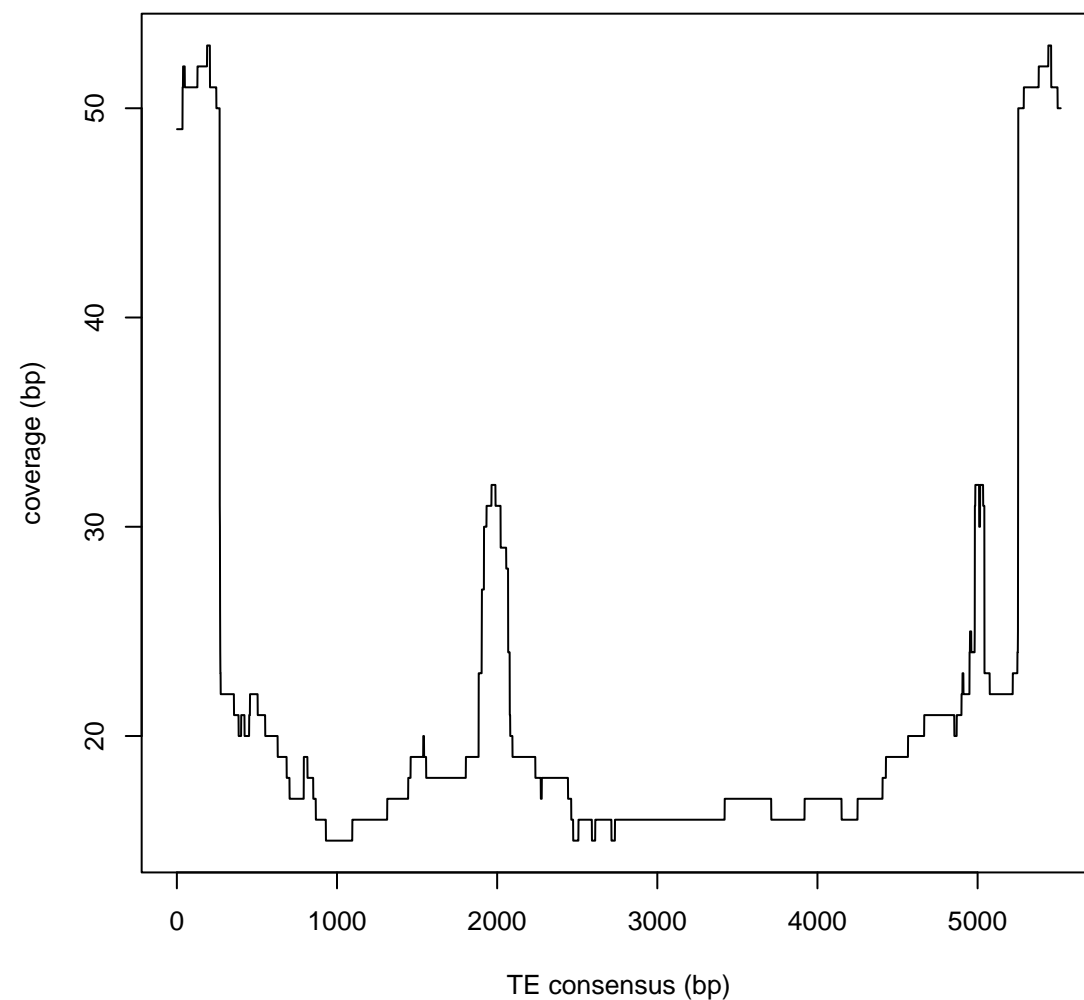


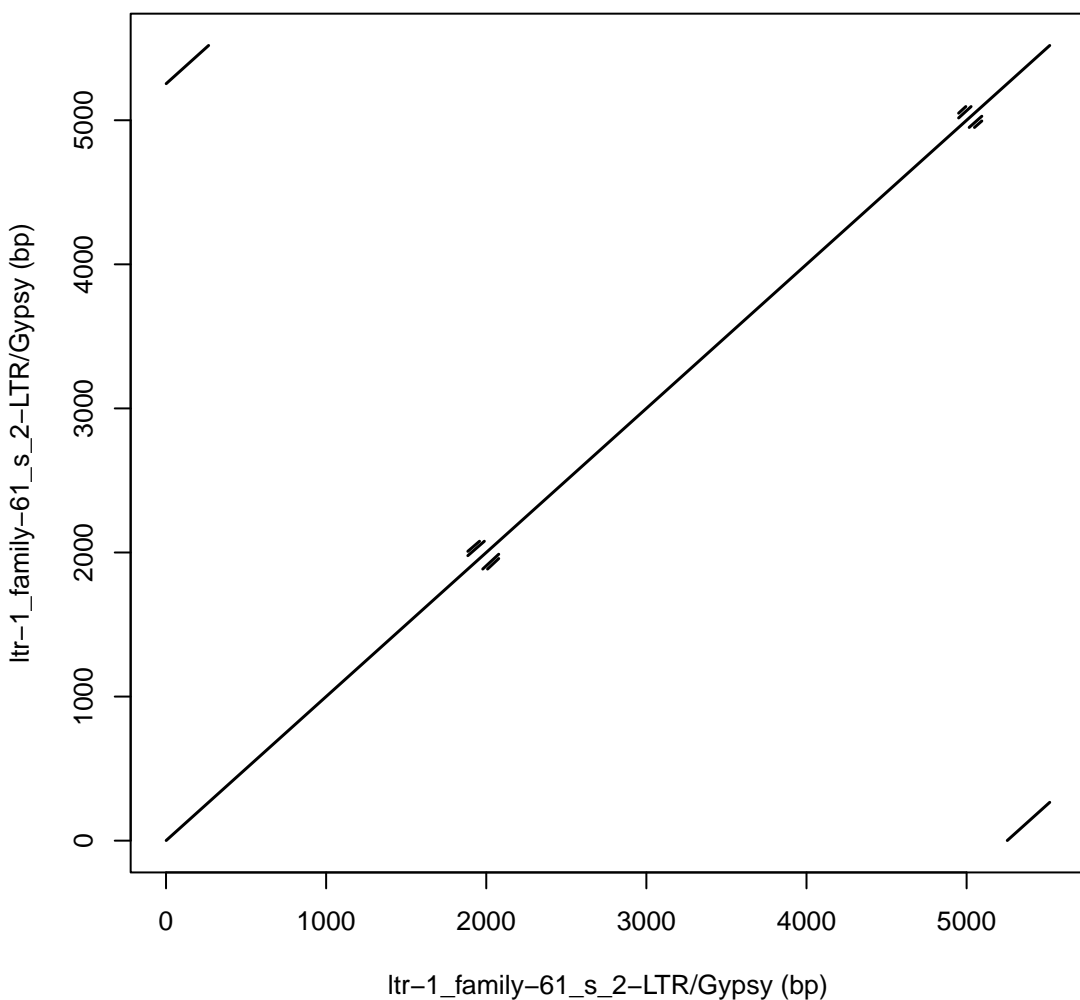
TE: ltr-1_family-61_s_2-LTR/Gypsy
consensus size: 5519bp; fragments: 137; full length: 8 (>=4967.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

