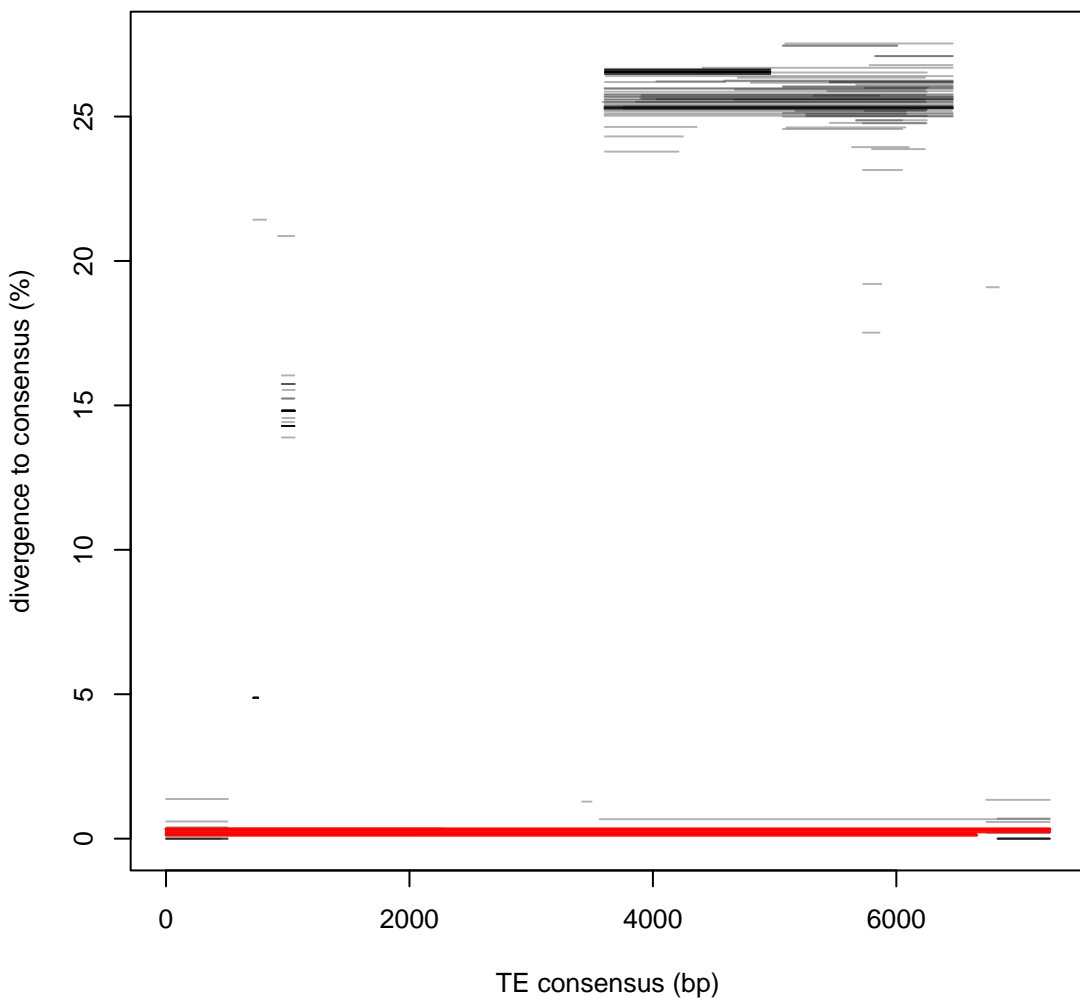
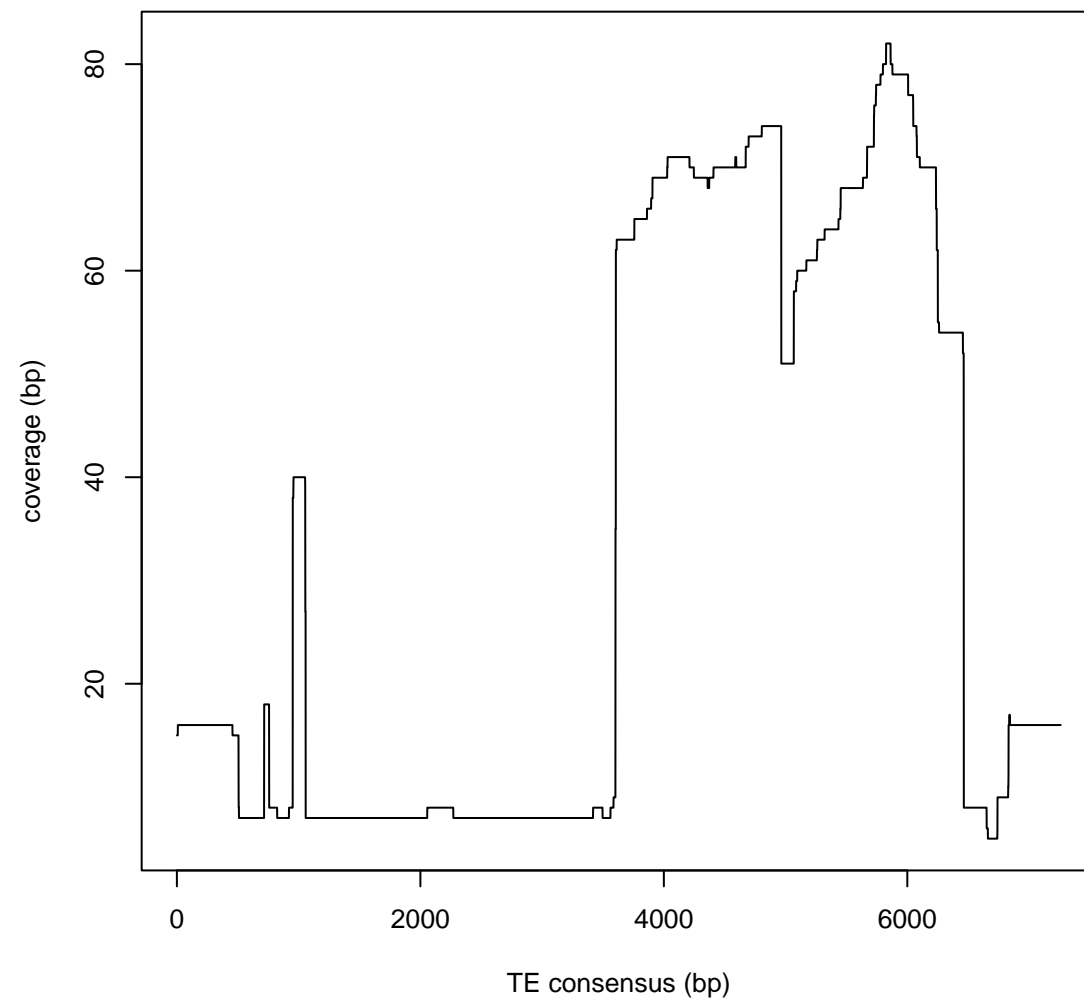


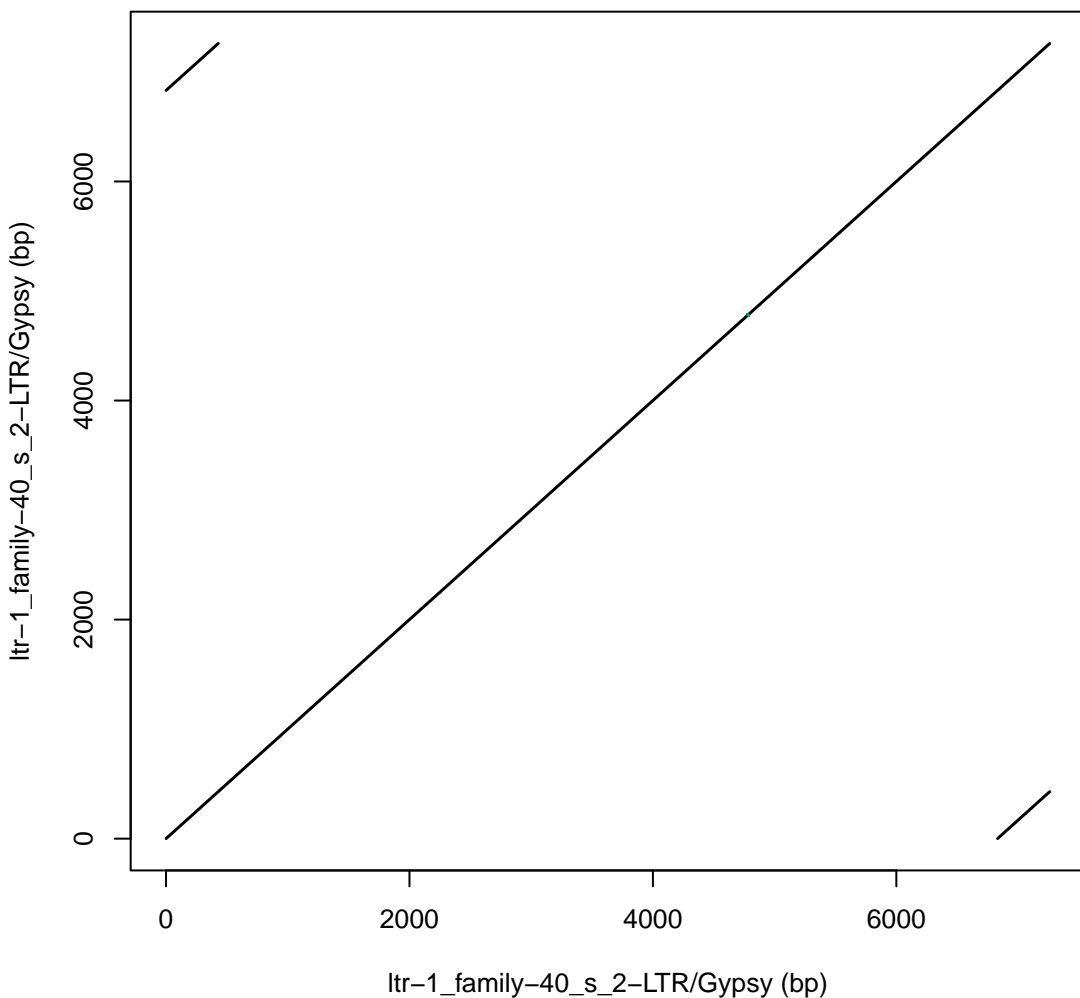
TE: ltr-1_family-40_s_2-LTR/Gypsy
consensus size: 7260bp; fragments: 176; full length: 6 (>=6534bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

