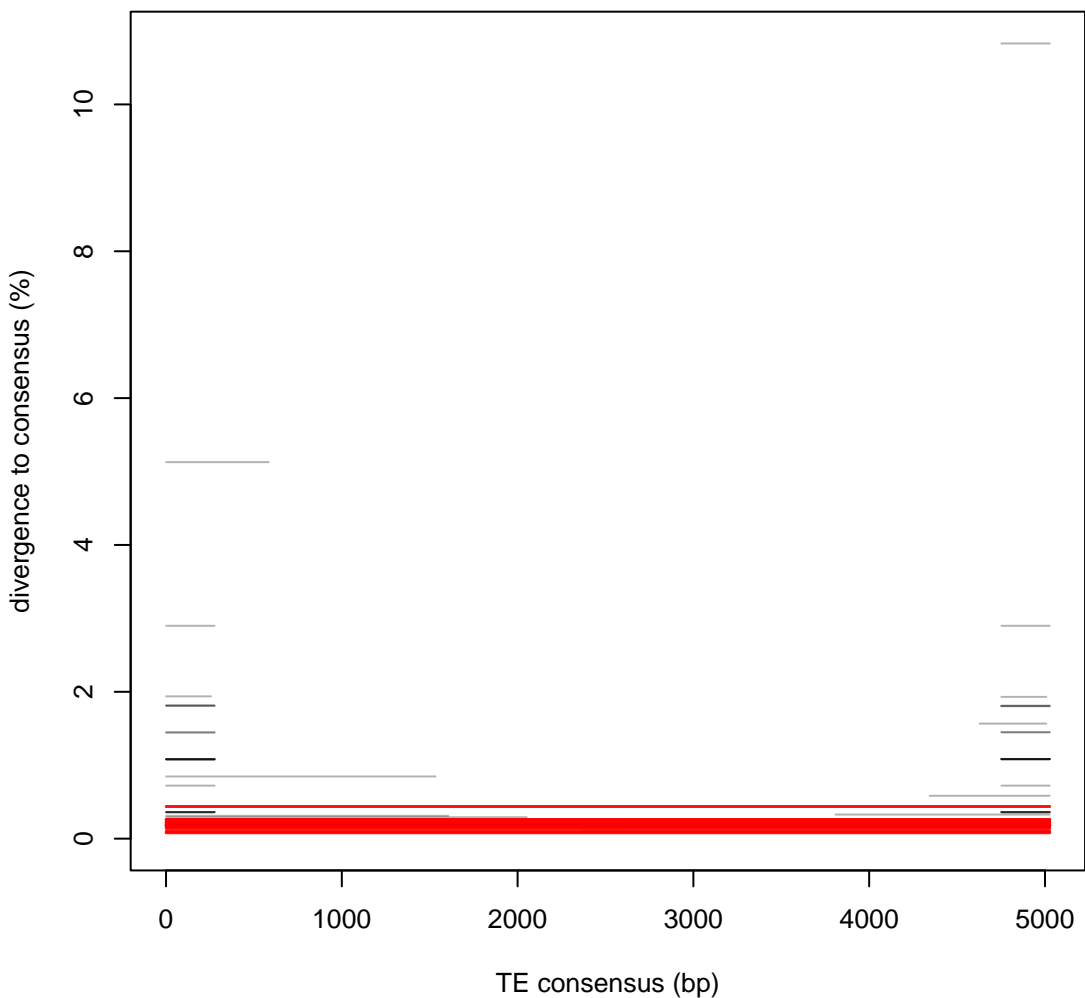
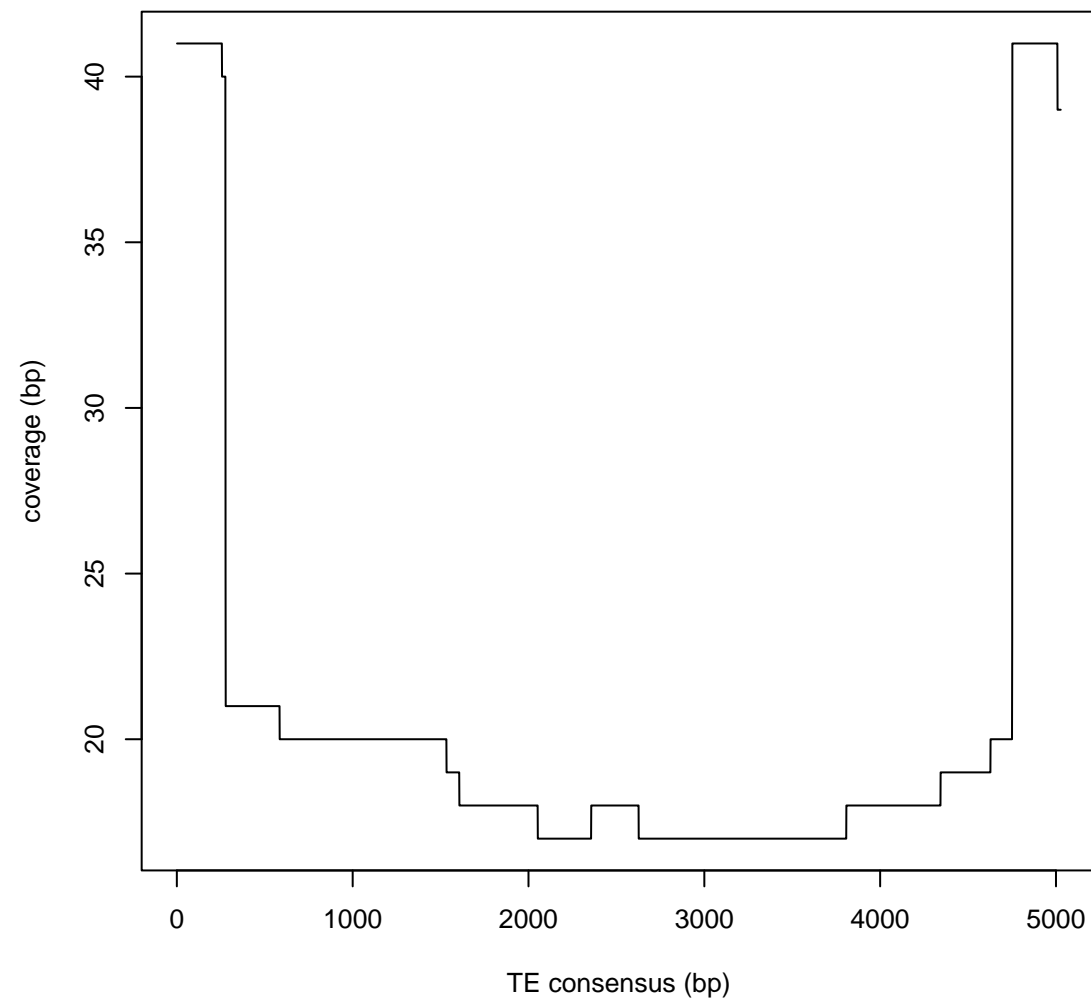


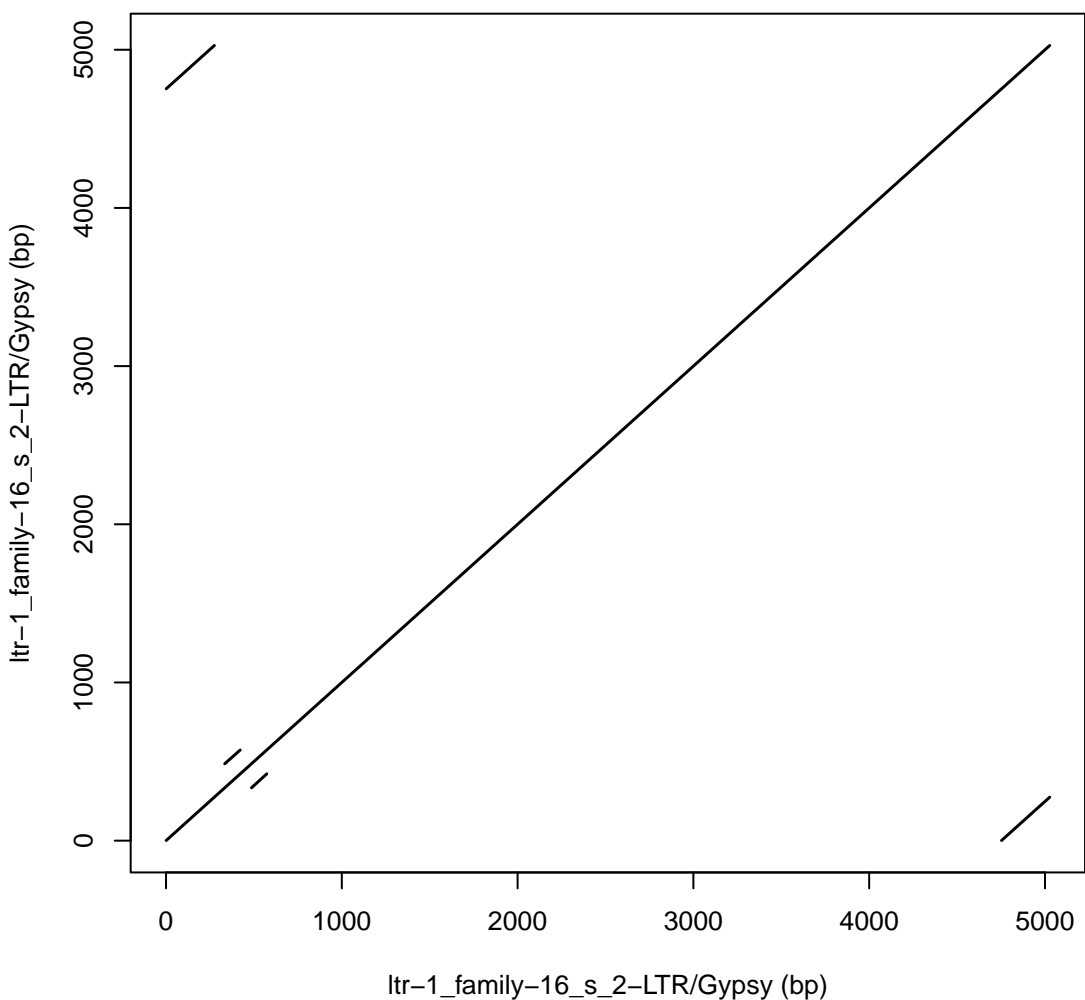
TE: ltr-1_family-16_s_2-LTR/Gypsy
consensus size: 5027bp; fragments: 66; full length: 16 (>=4524.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

