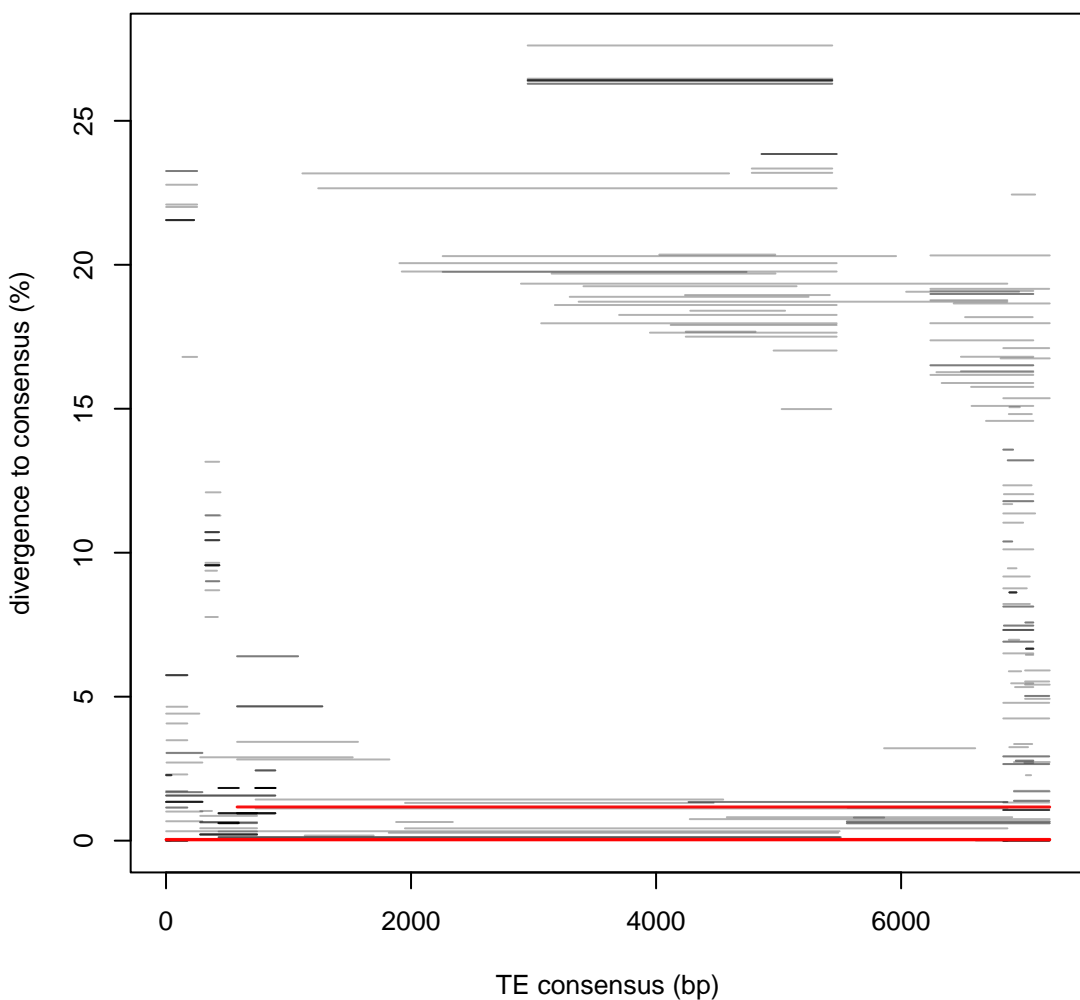
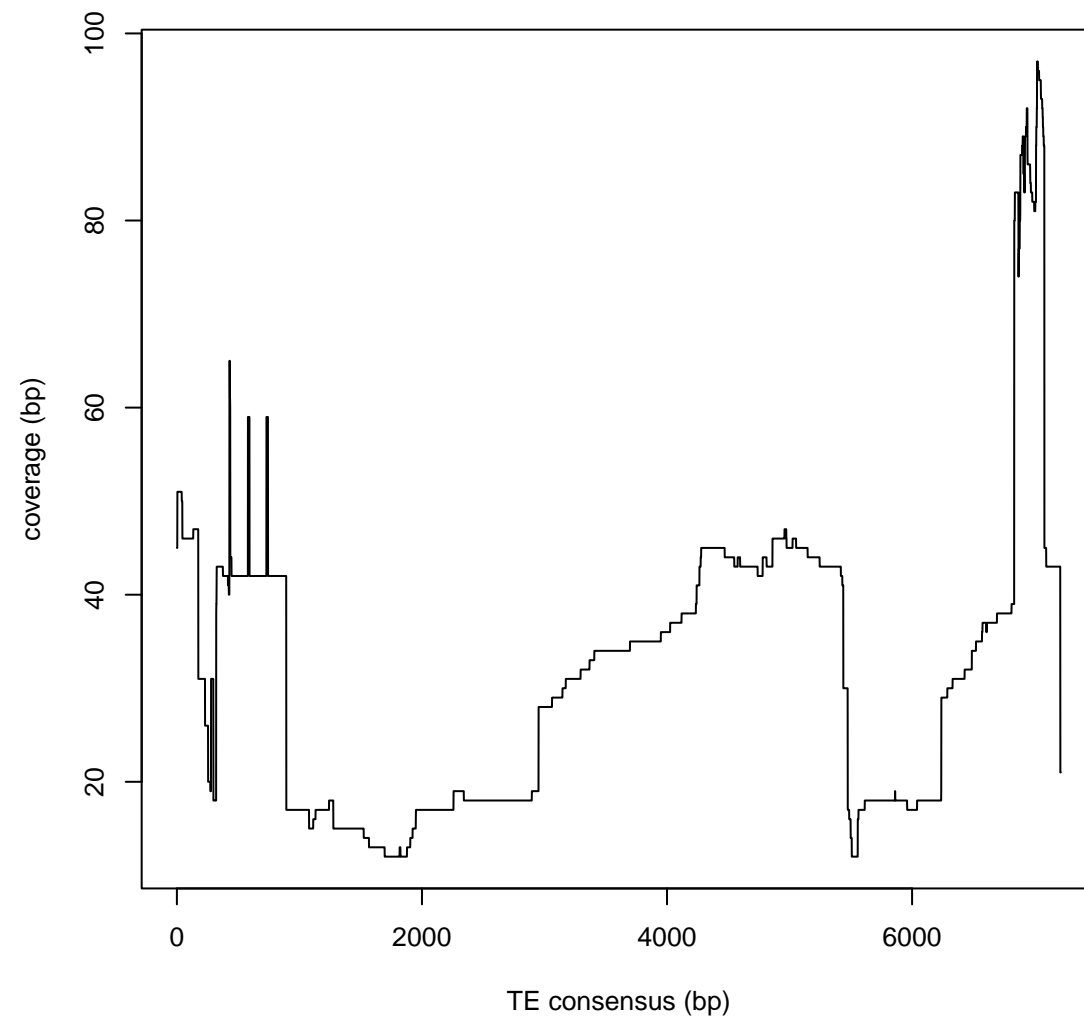


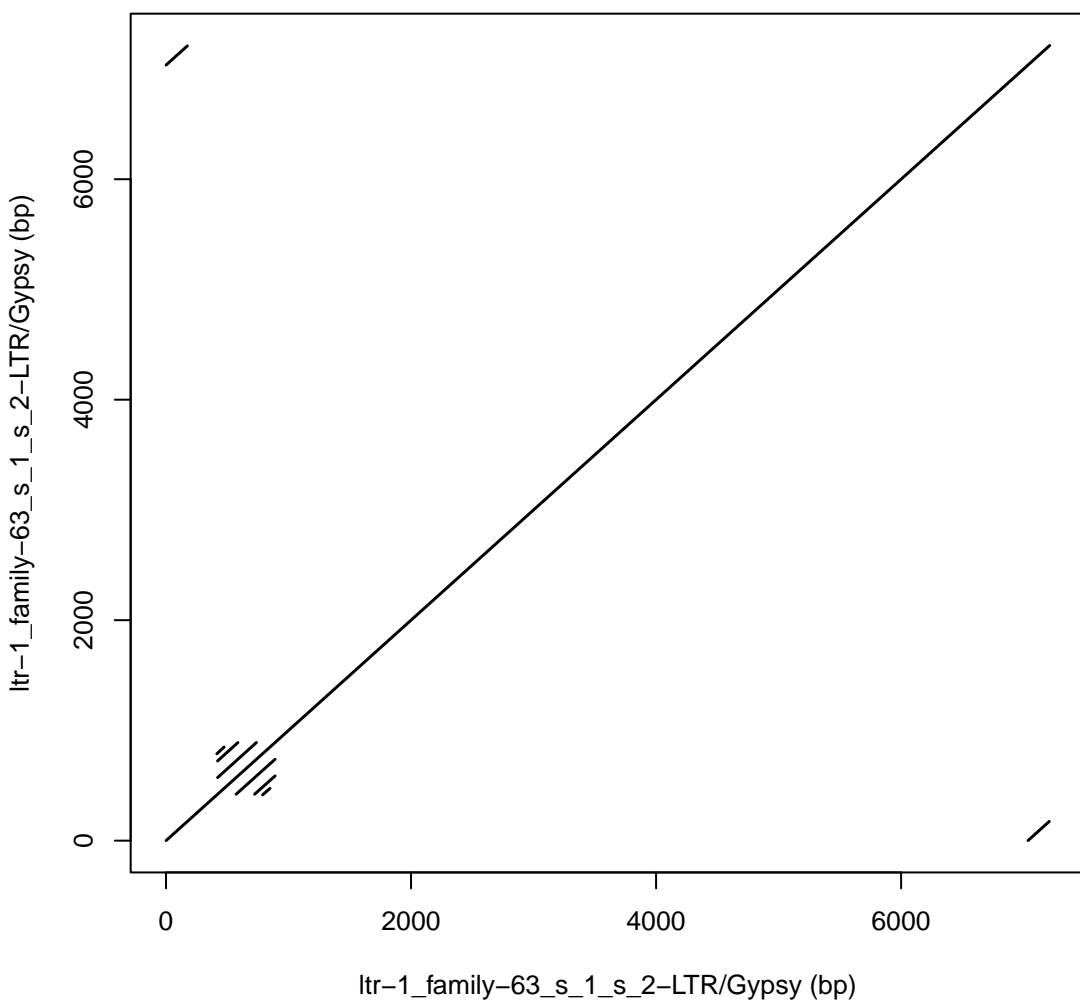
TE: ltr-1_family-63_s_1_s_2-LTR/Gypsy
consensus size: 7213bp; fragments: 308; full length: 3 (>=6491.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

