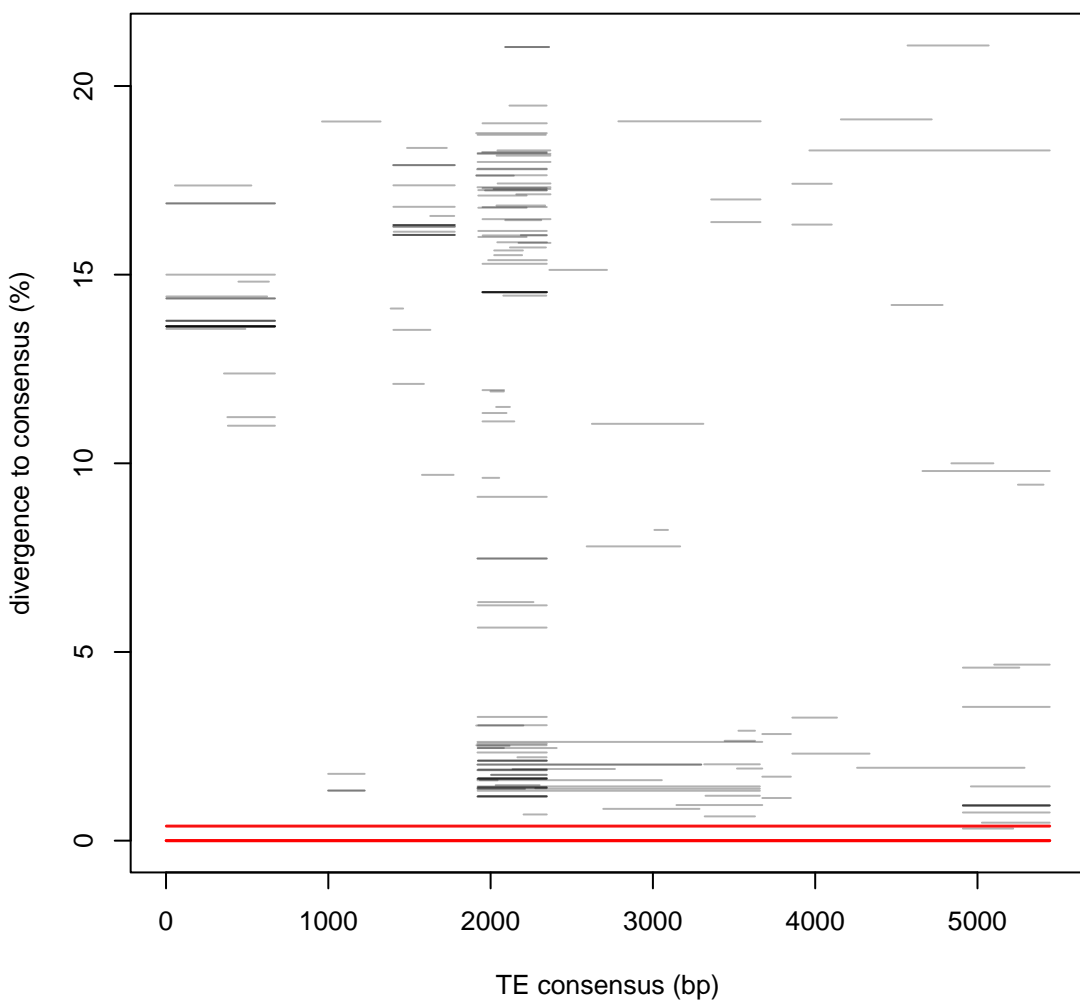
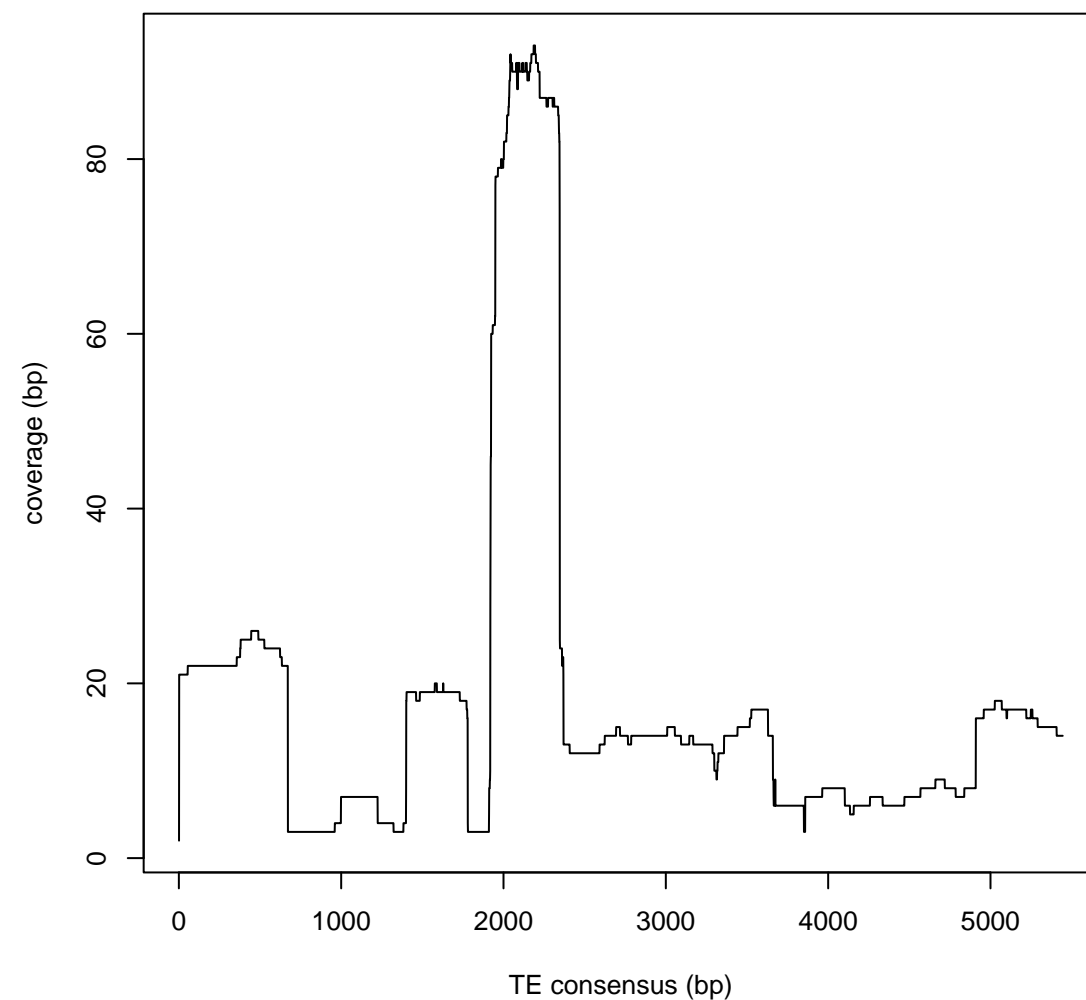


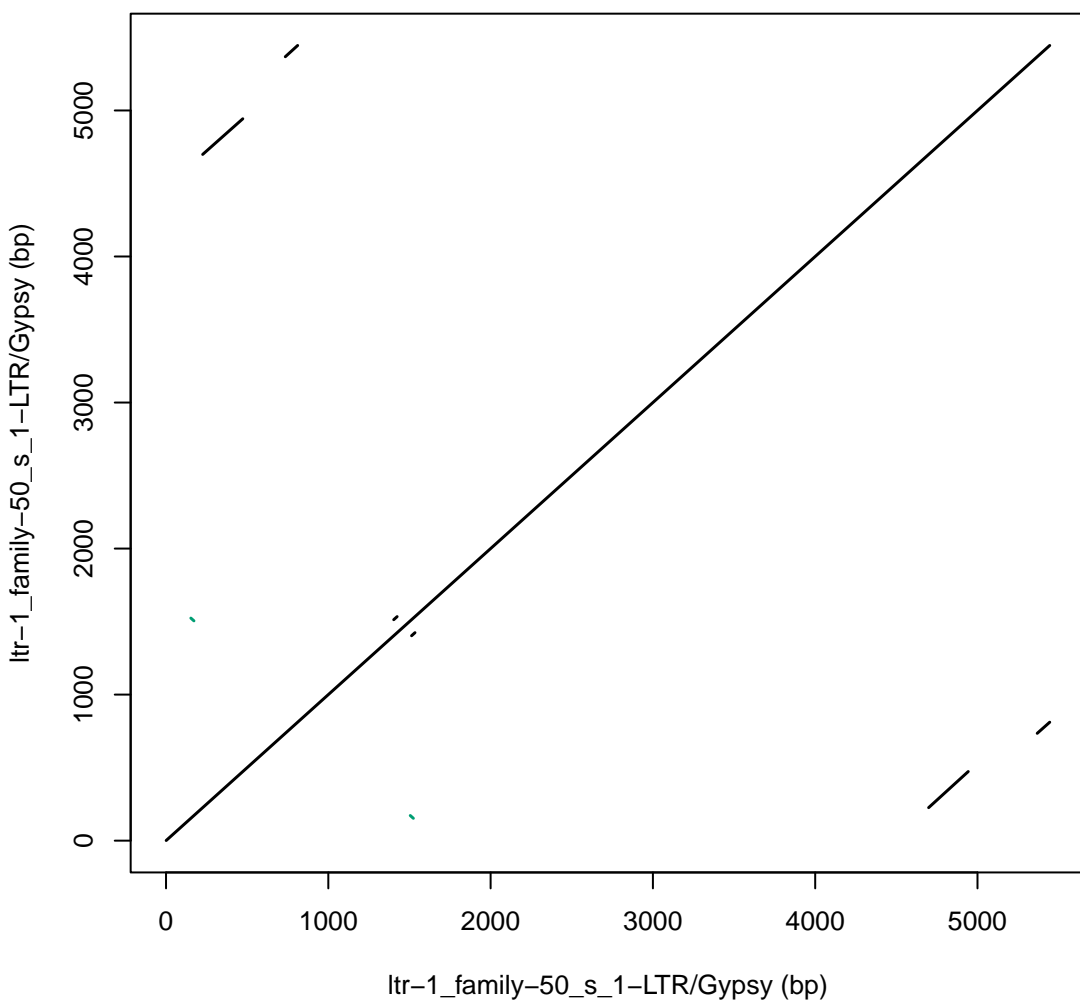
TE: ltr-1_family-50_s_1-LTR/Gypsy
consensus size: 5445bp; fragments: 194; full length: 3 (>=4900.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

