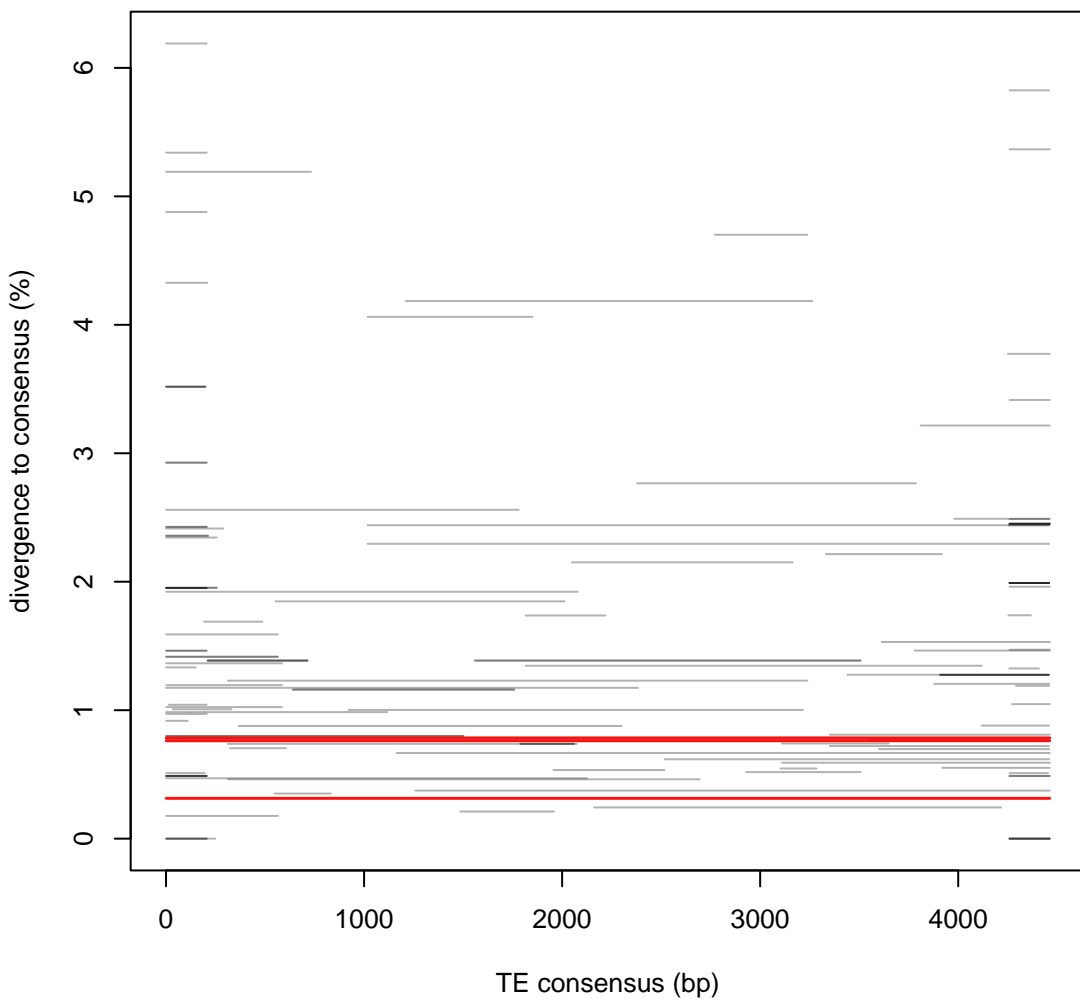
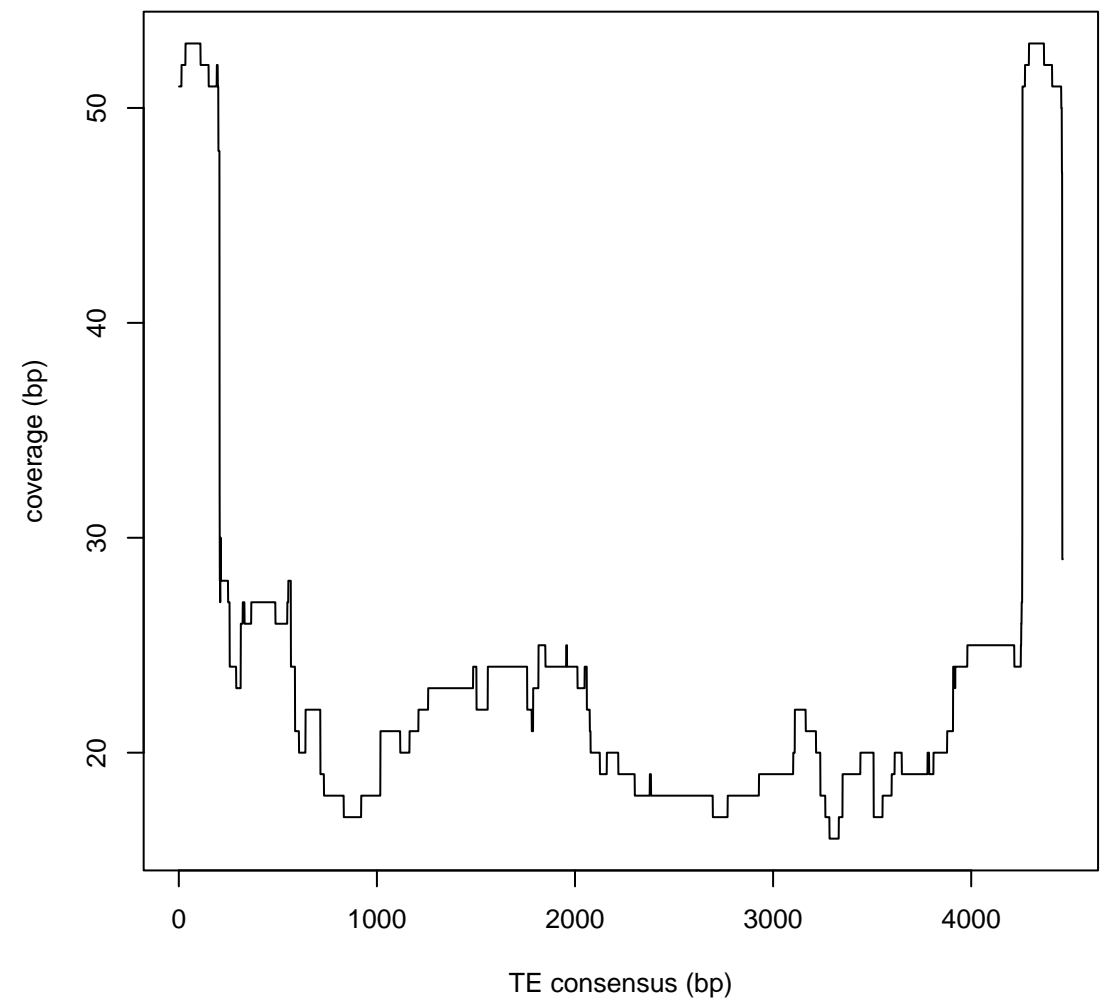


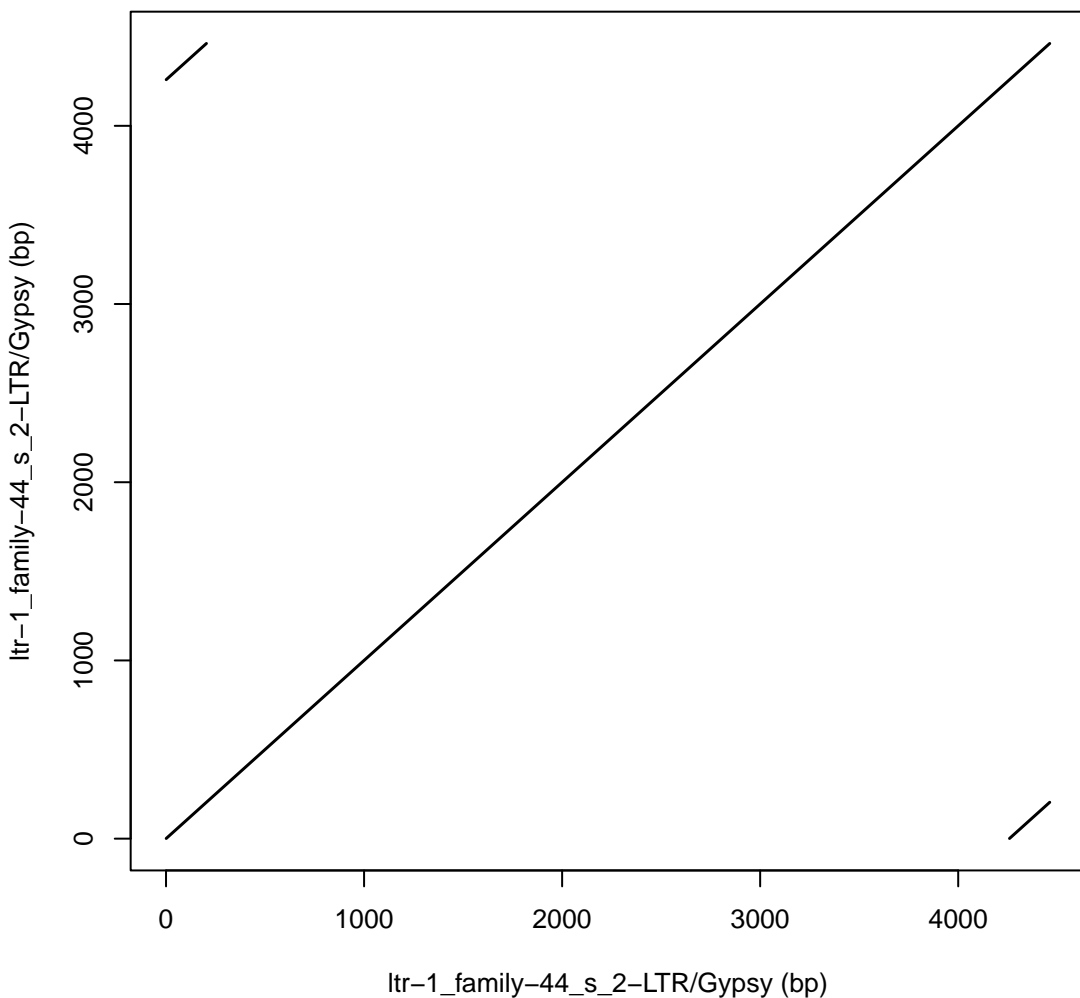
TE: ltr-1_family-44_s_2-LTR/Gypsy
consensus size: 4462bp; fragments: 135; full length: 3 (>=4015.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

