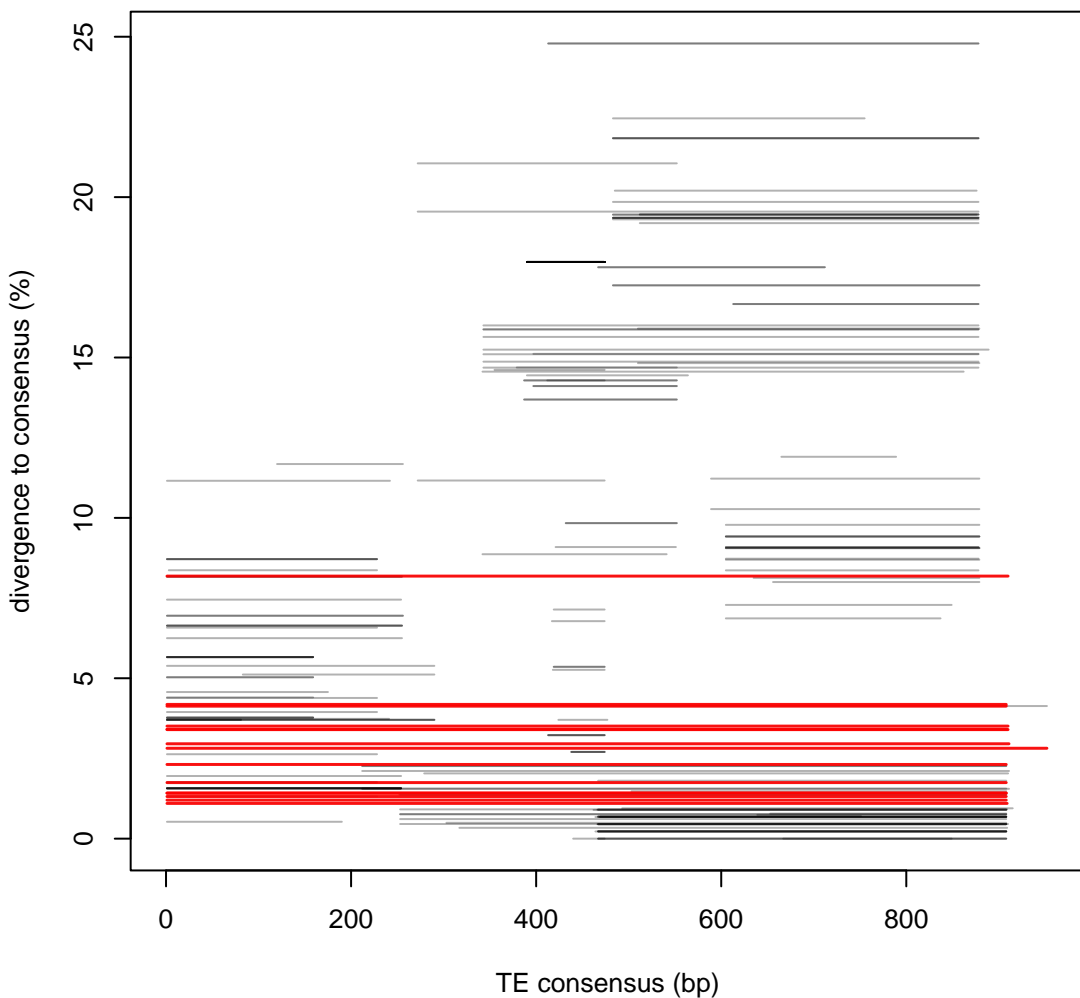
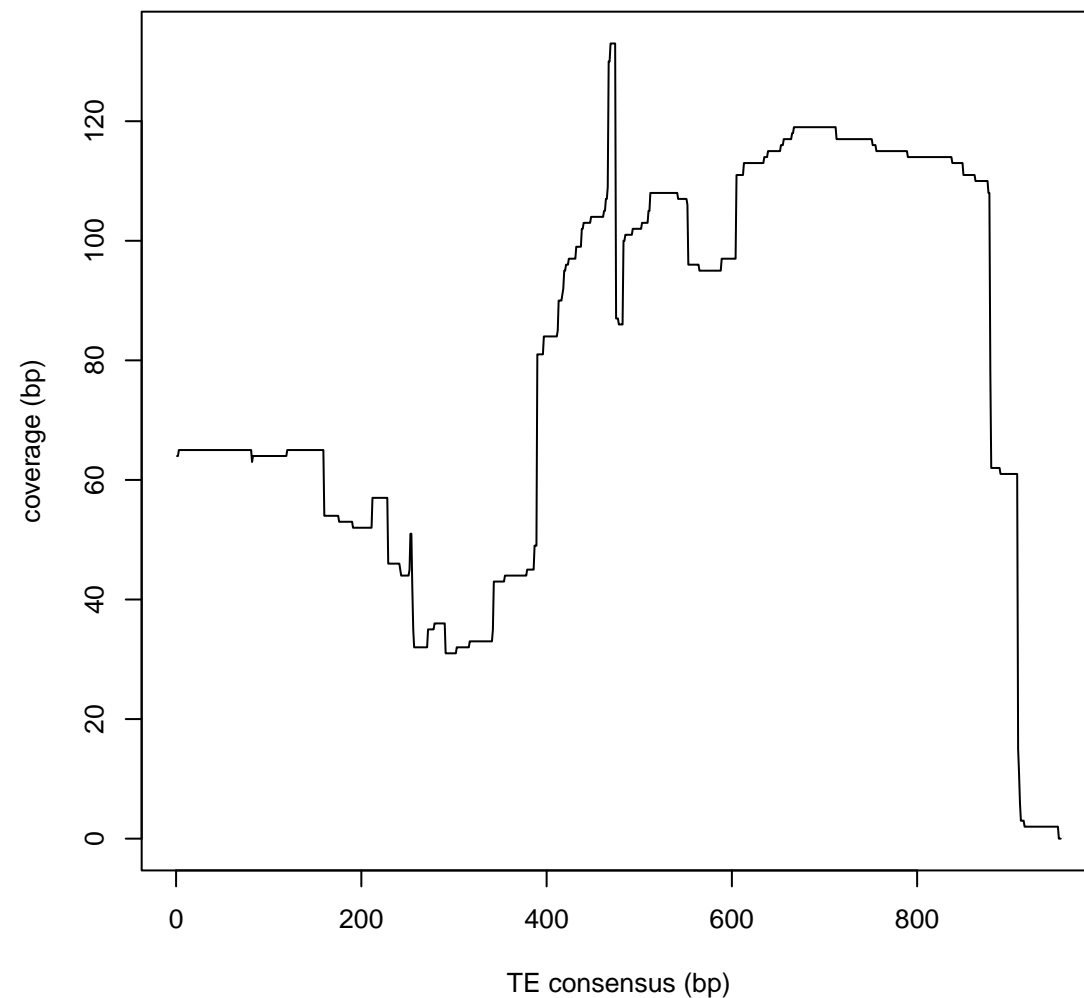


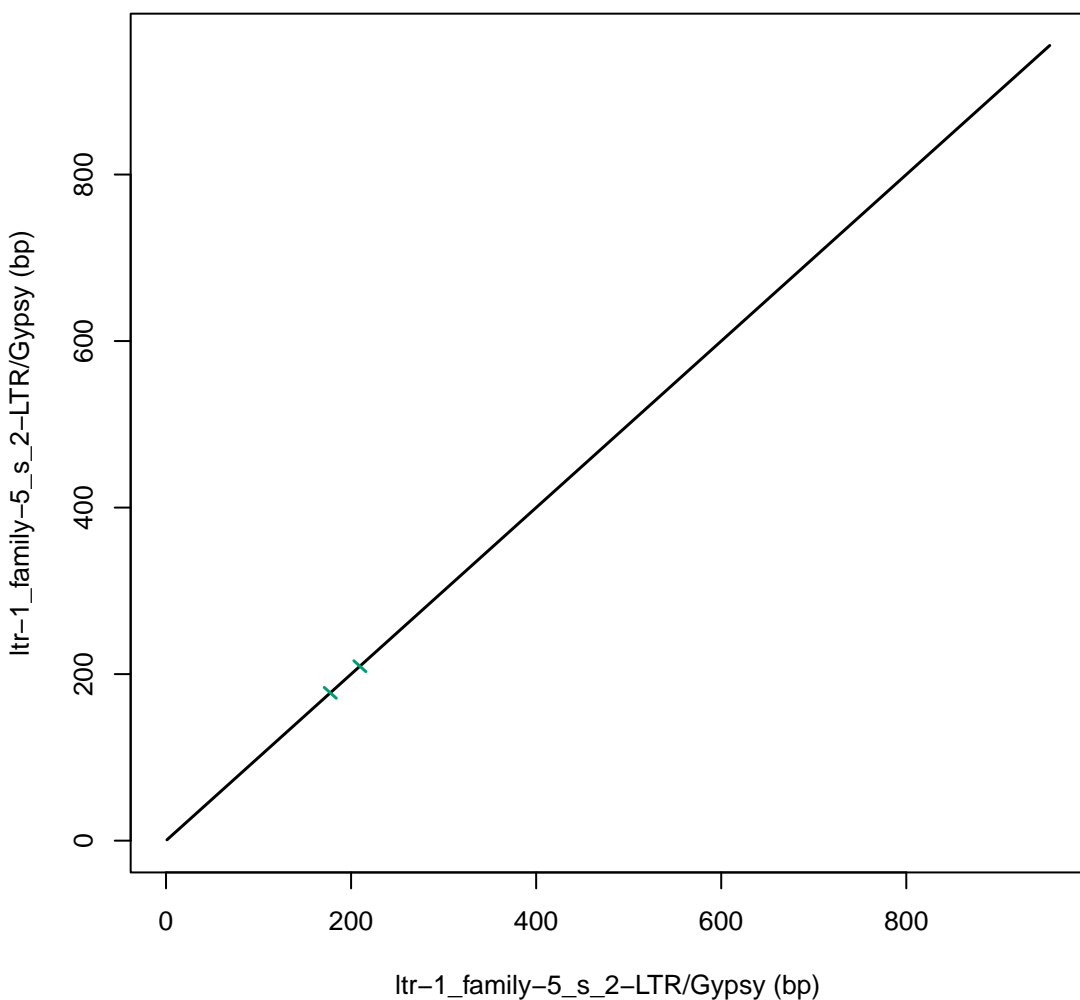
TE: ltr-1_family-5_s_2-LTR/Gypsy
consensus size: 955bp; fragments: 231; full length: 15 (>=859.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

