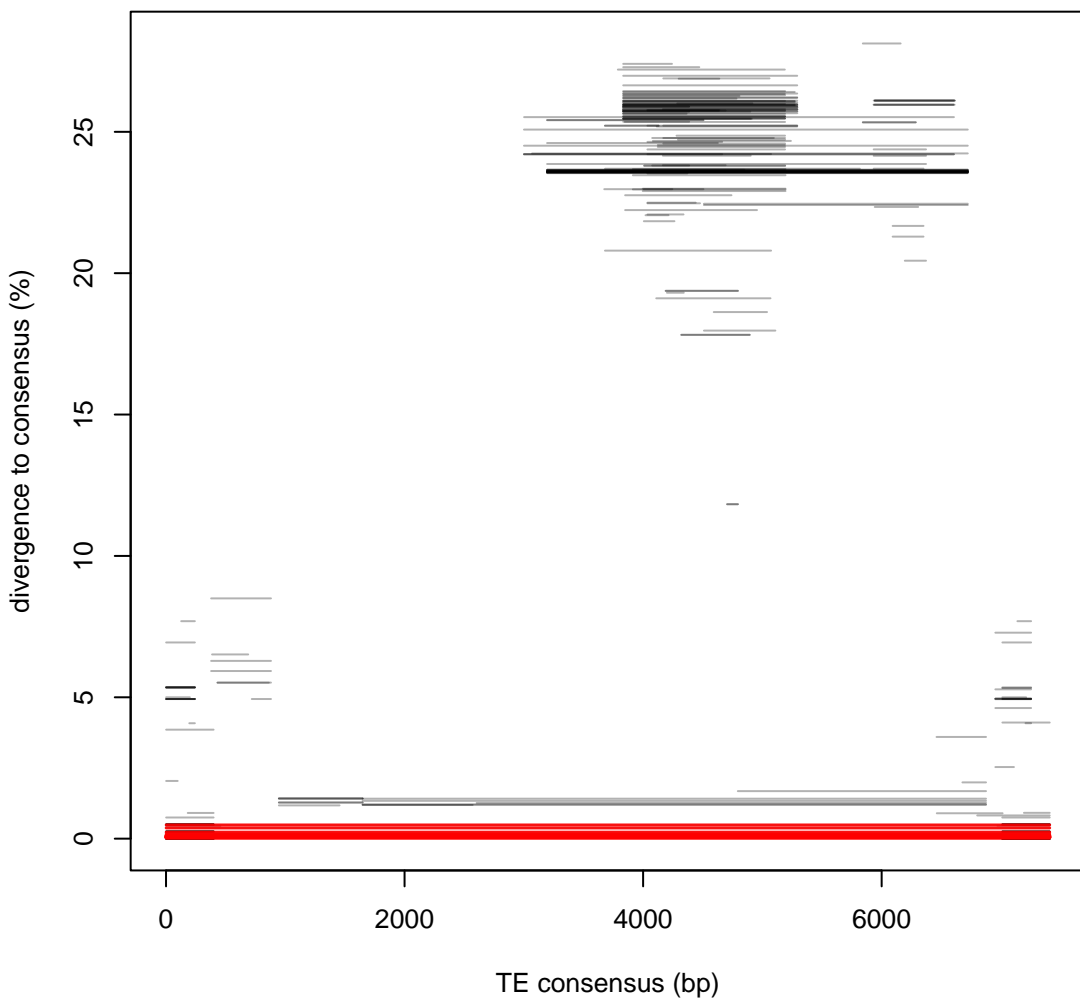
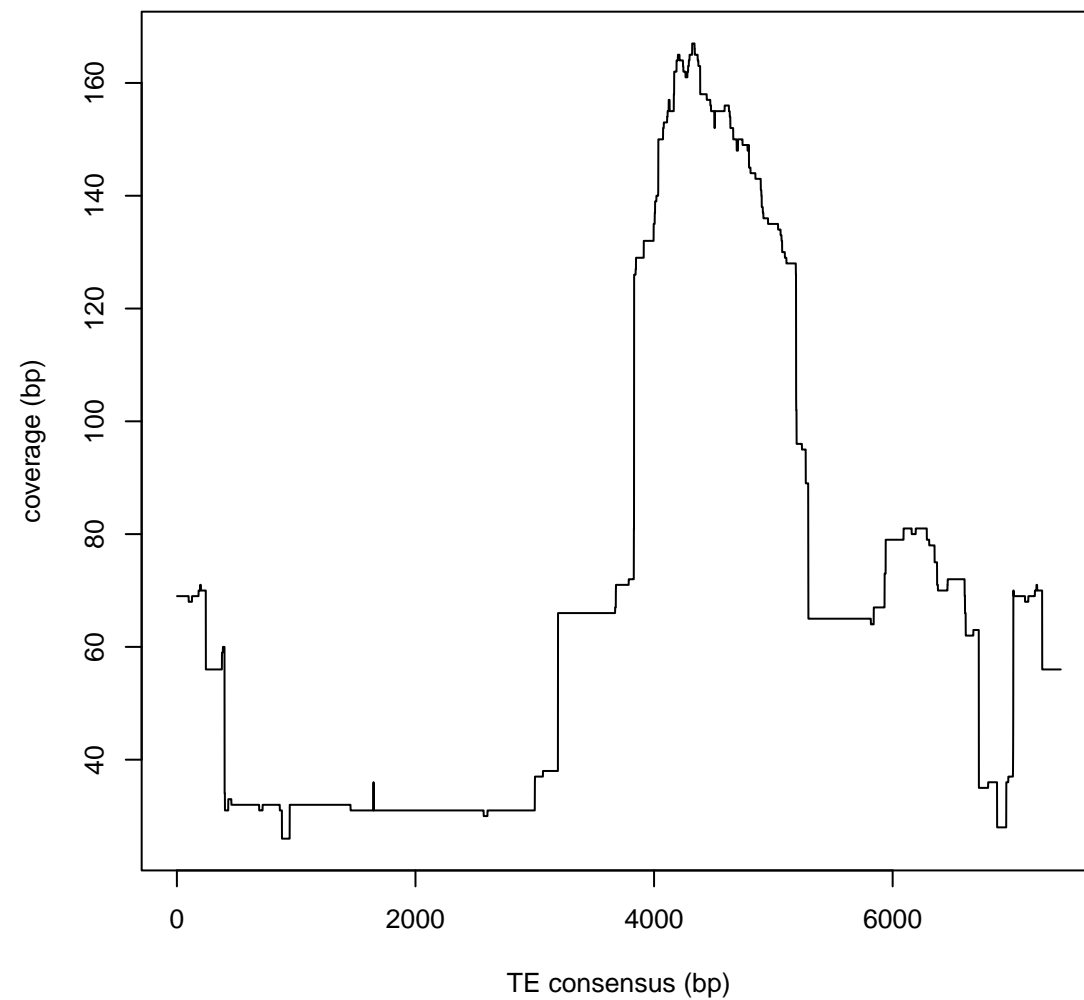


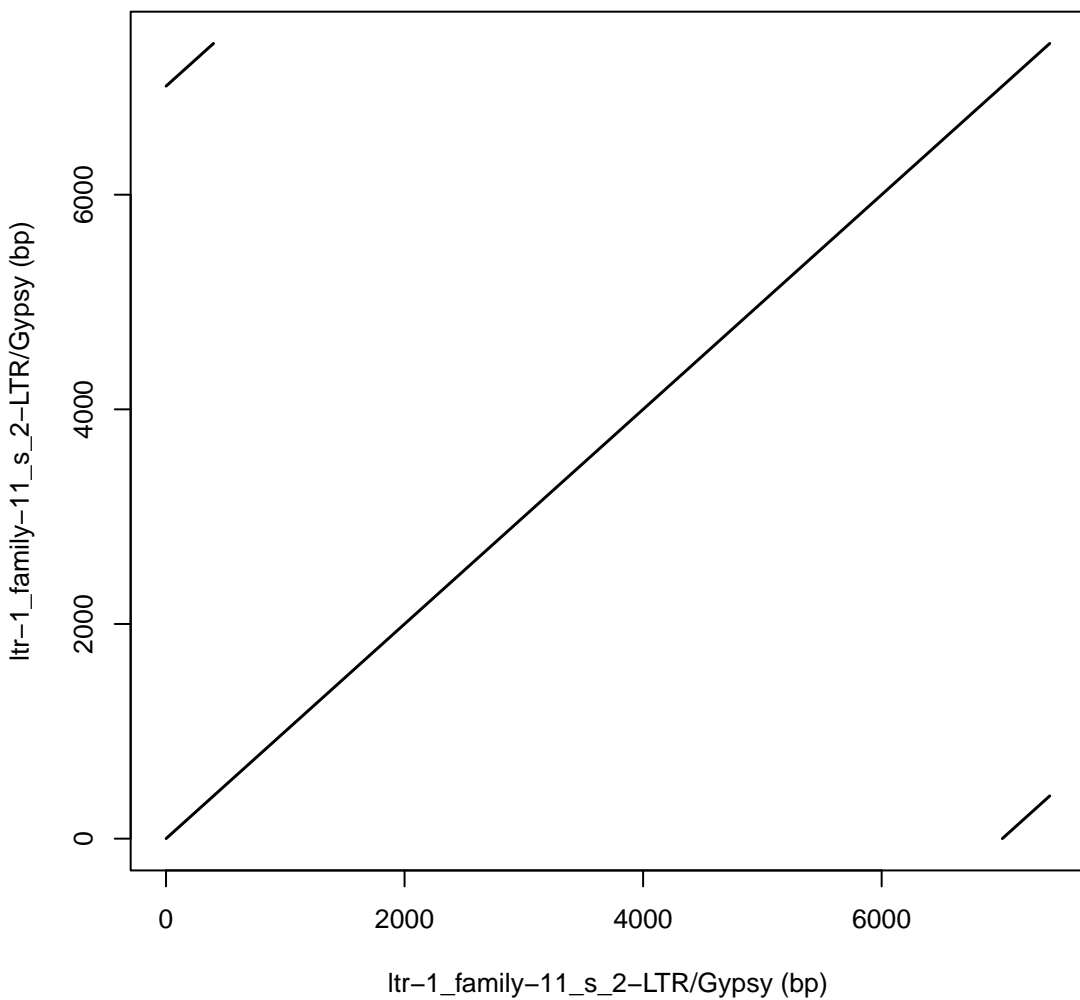
TE: ltr-1_family-11_s_2-LTR/Gypsy
consensus size: 7409bp; fragments: 307; full length: 26 (>=6668.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

