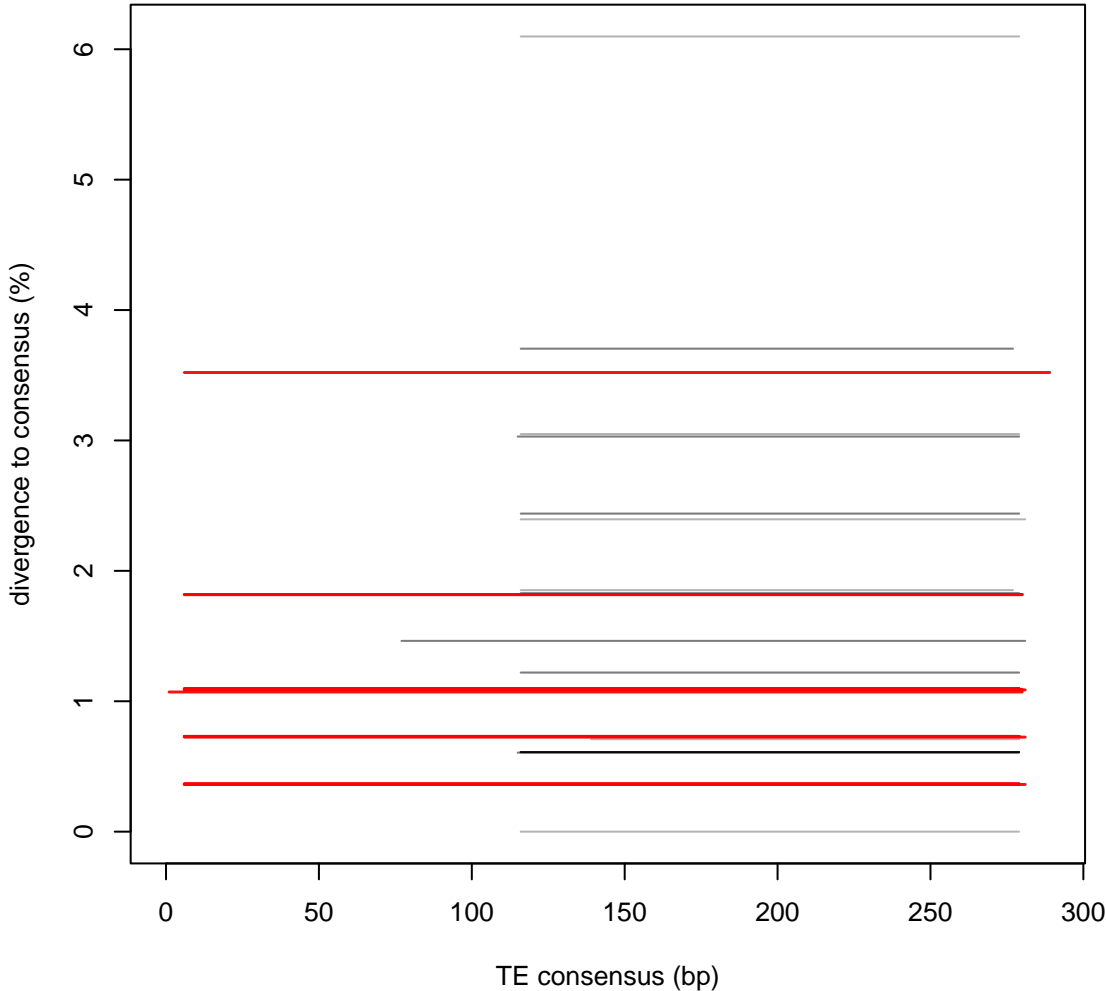
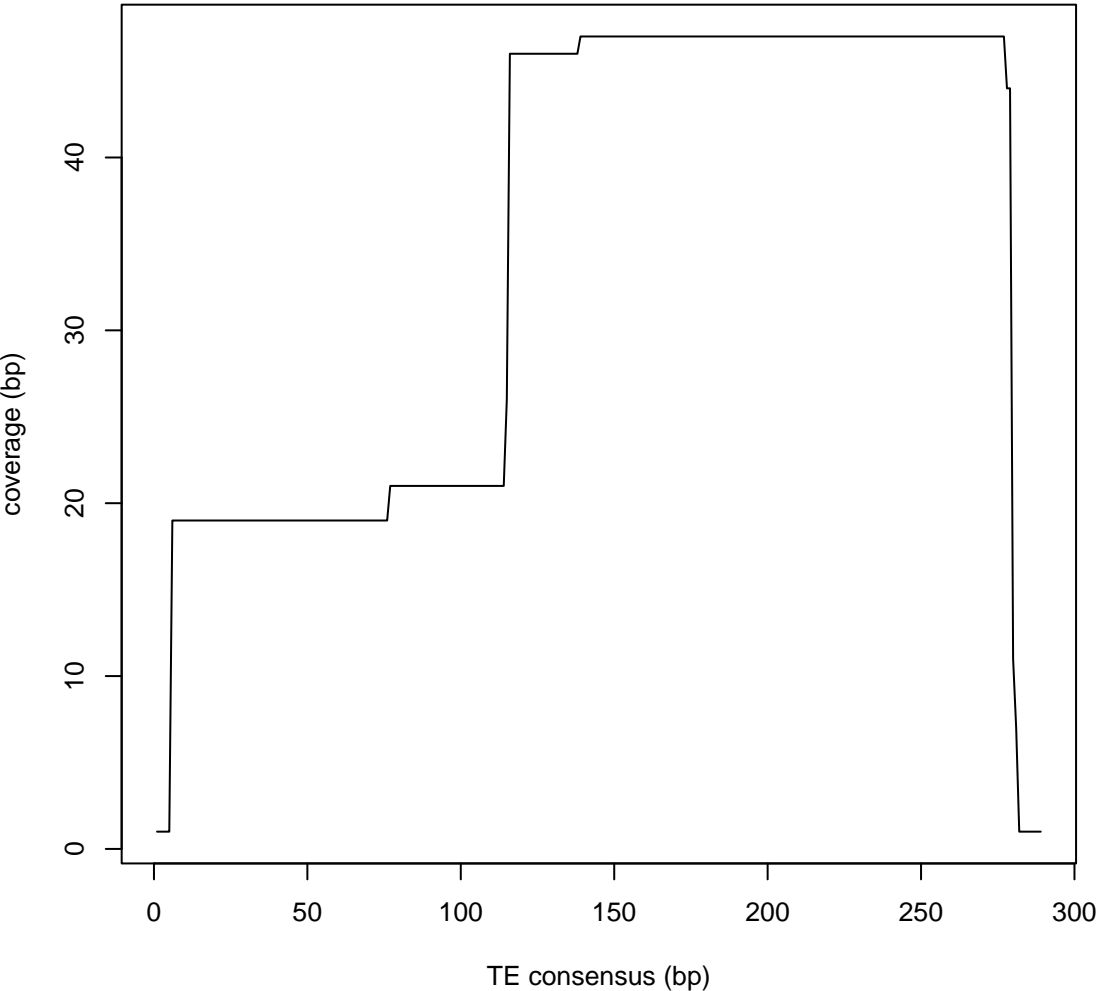


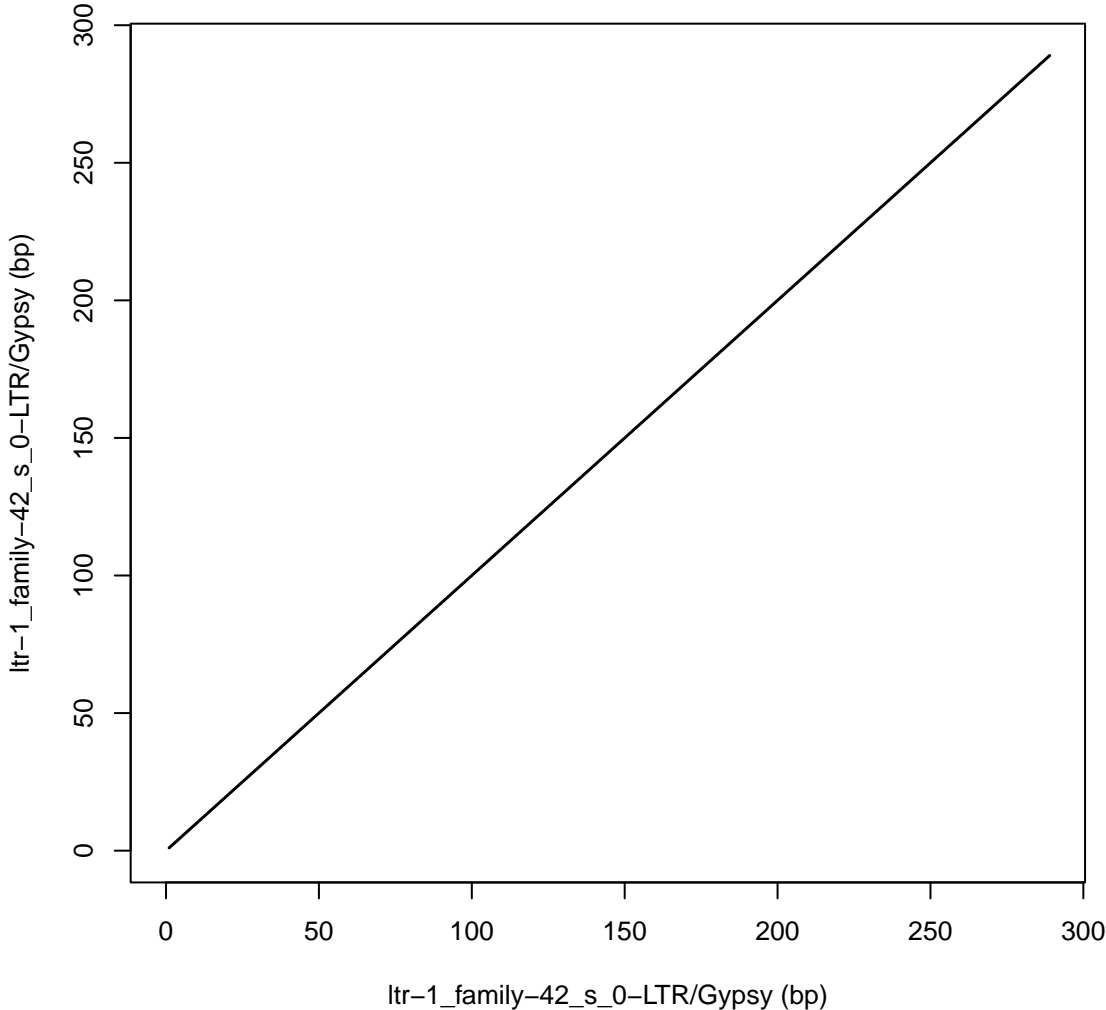
TE: ltr-1_family-42_s_0-LTR/Gypsy
consensus size: 289bp; fragments: 47; full length: 19 (>=260.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

