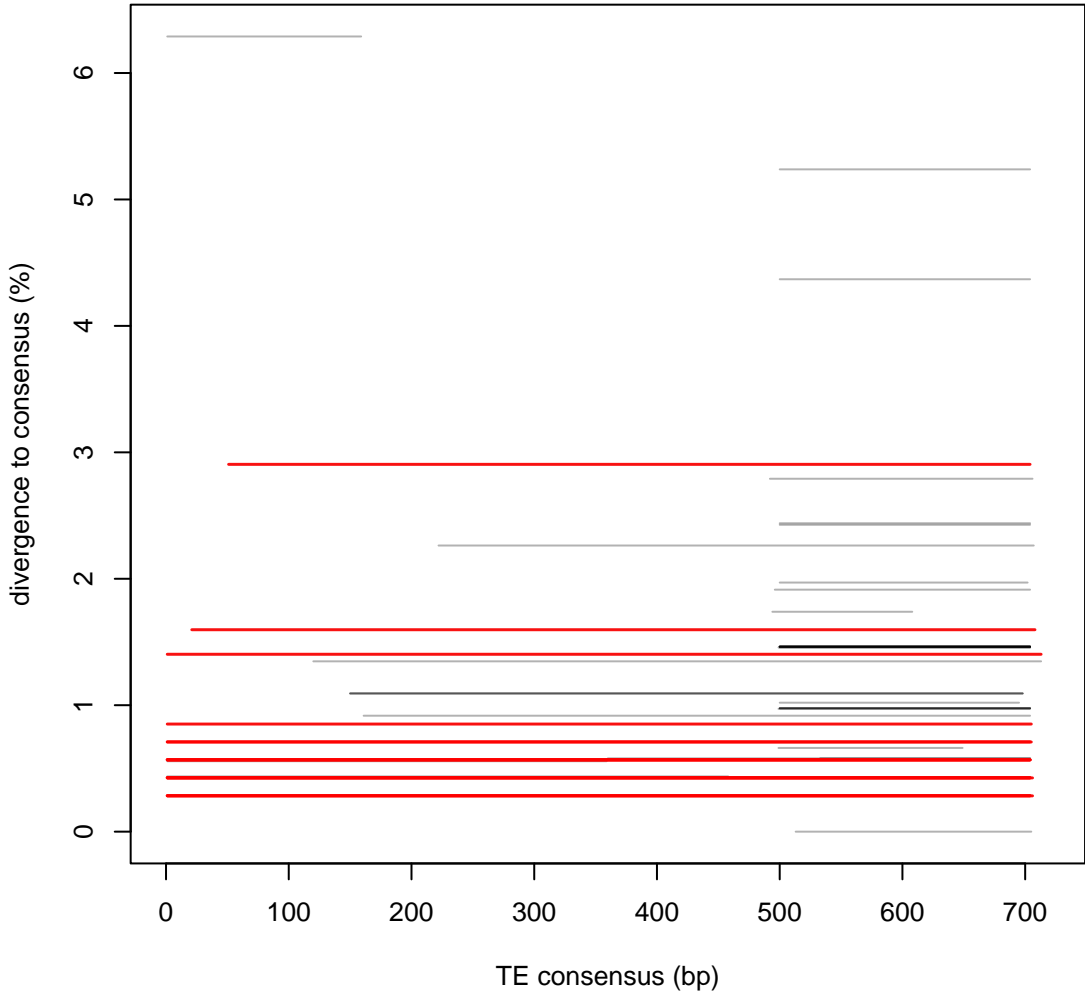
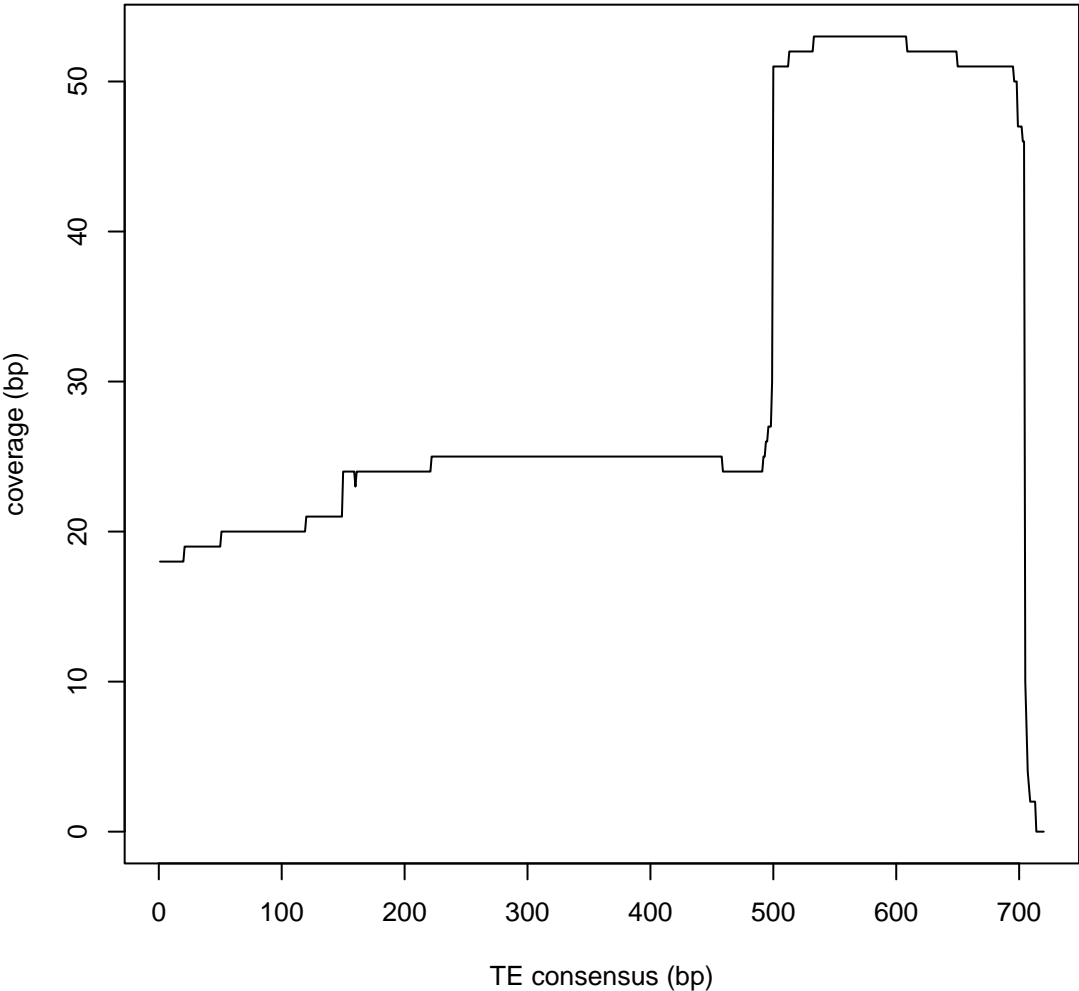


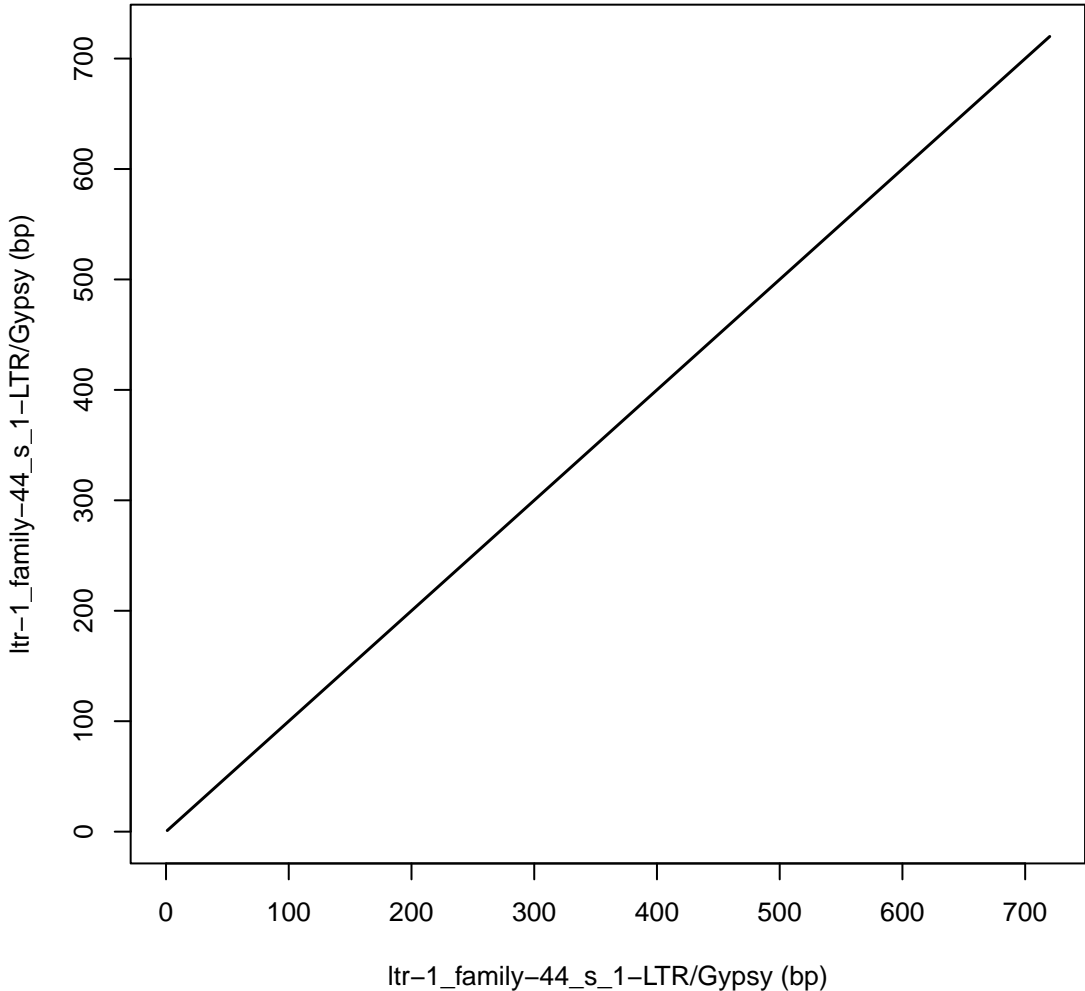
TE: ltr-1_family-44_s_1-LTR/Gypsy
consensus size: 720bp; fragments: 56; full length: 17 (>=648bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

