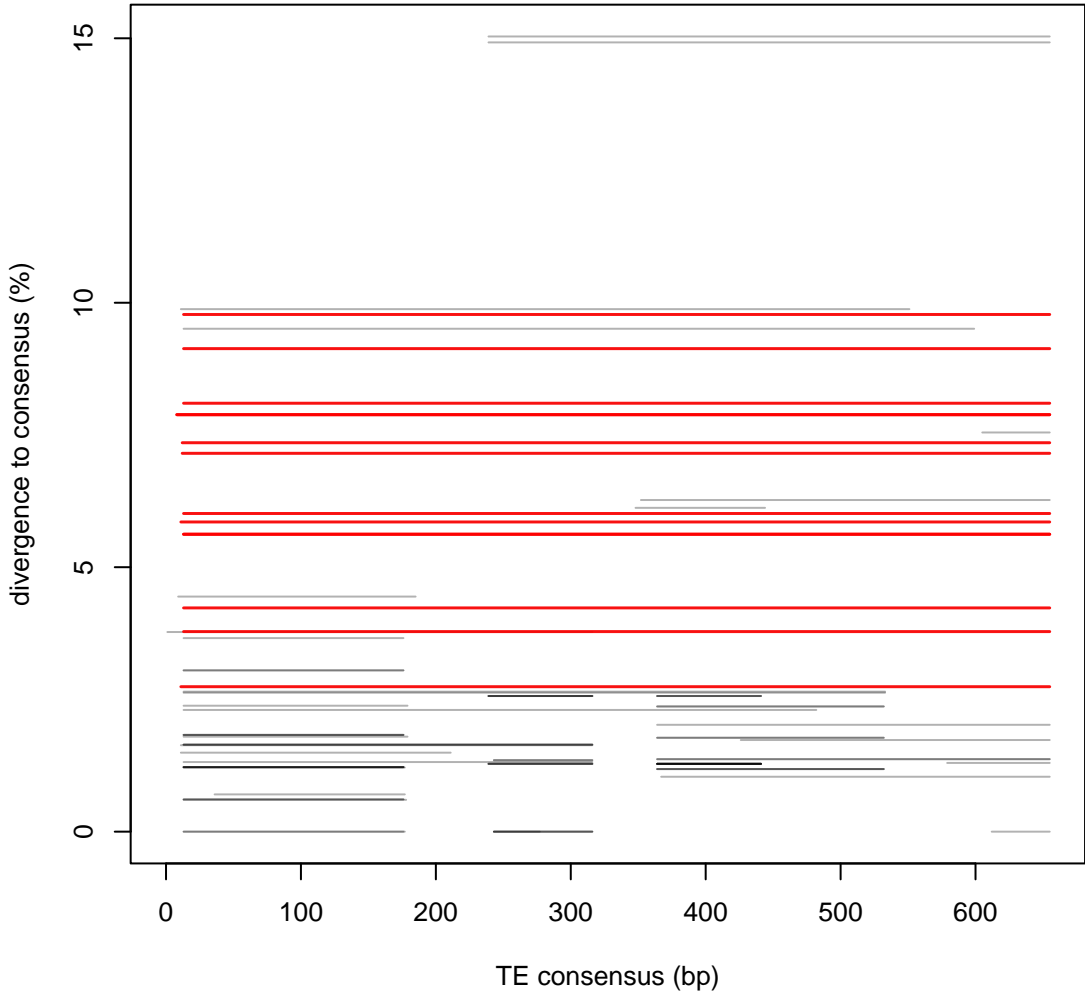
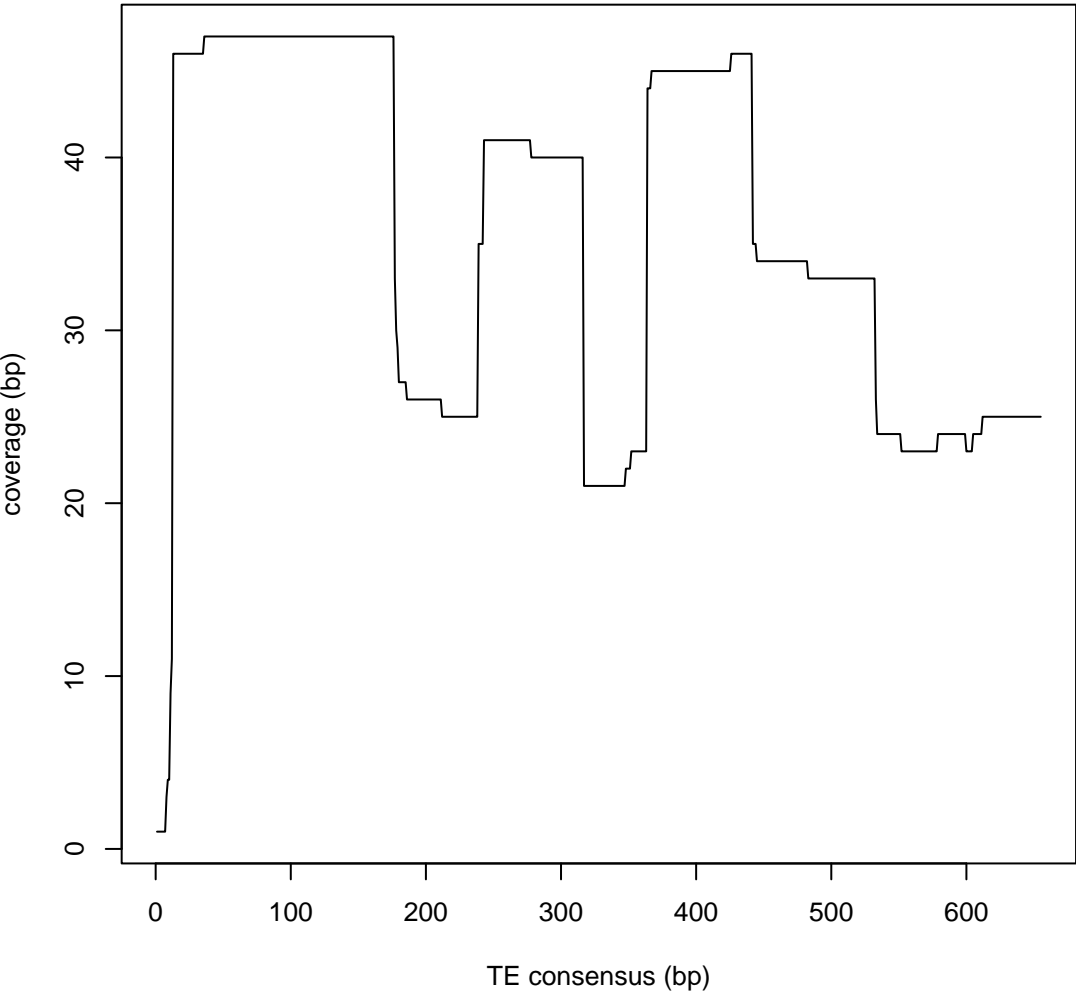


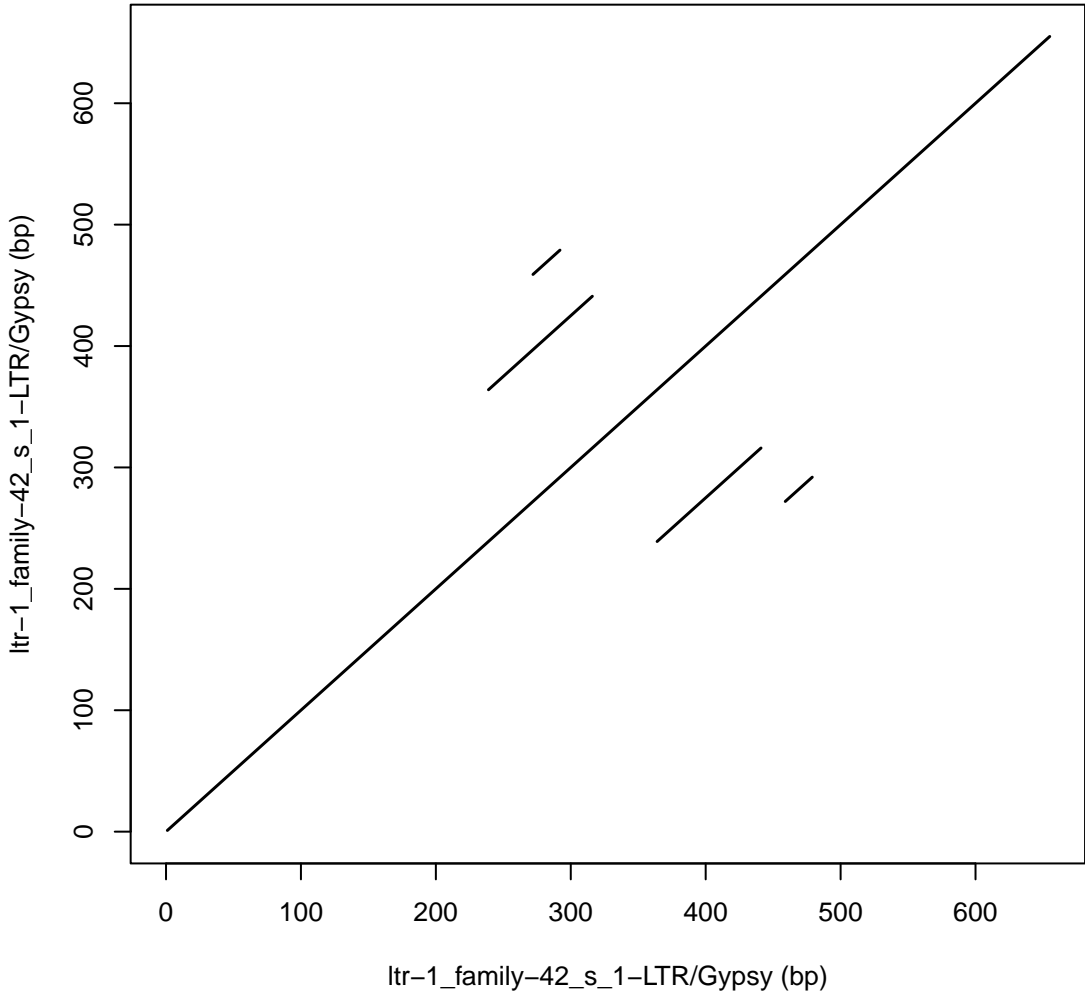
TE: ltr-1_family-42_s_1-LTR/Gypsy
consensus size: 655bp; fragments: 91; full length: 14 (>=589.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

