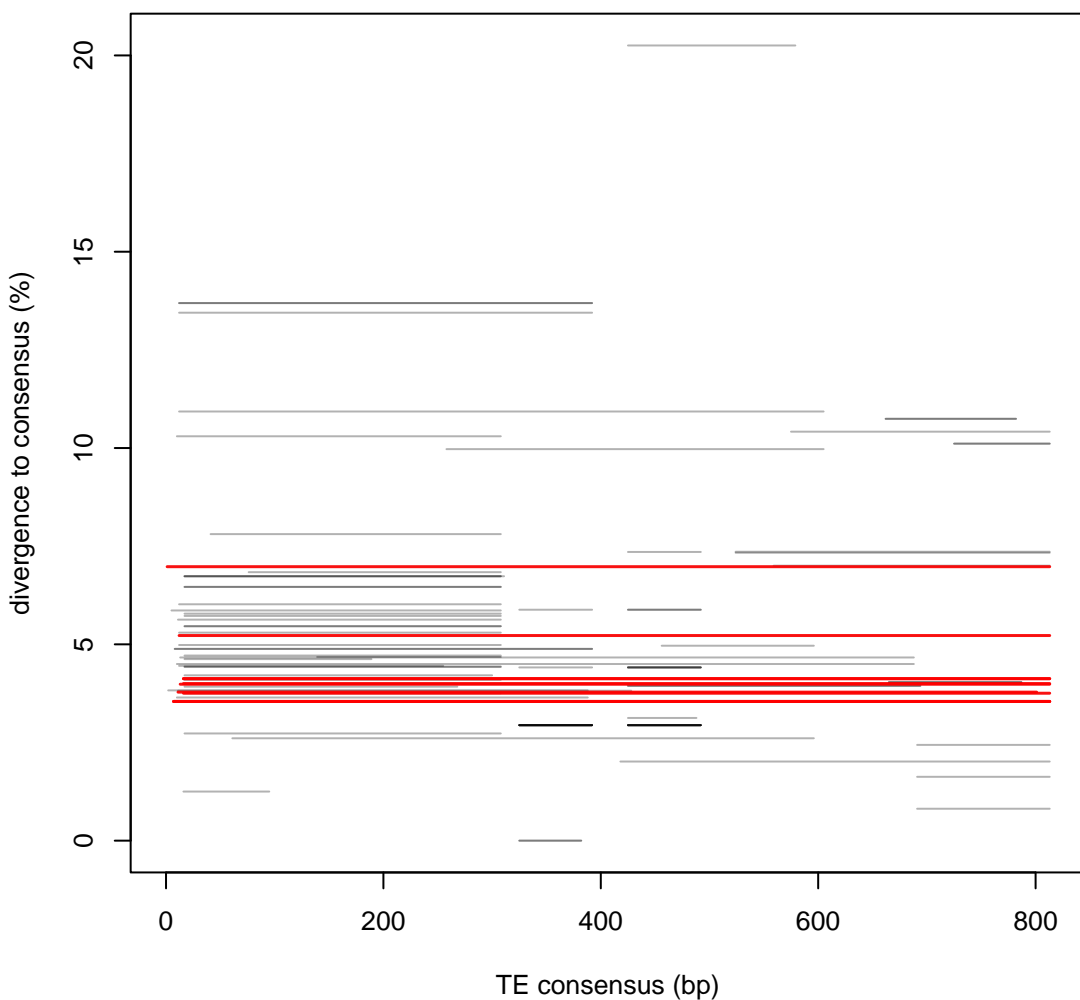
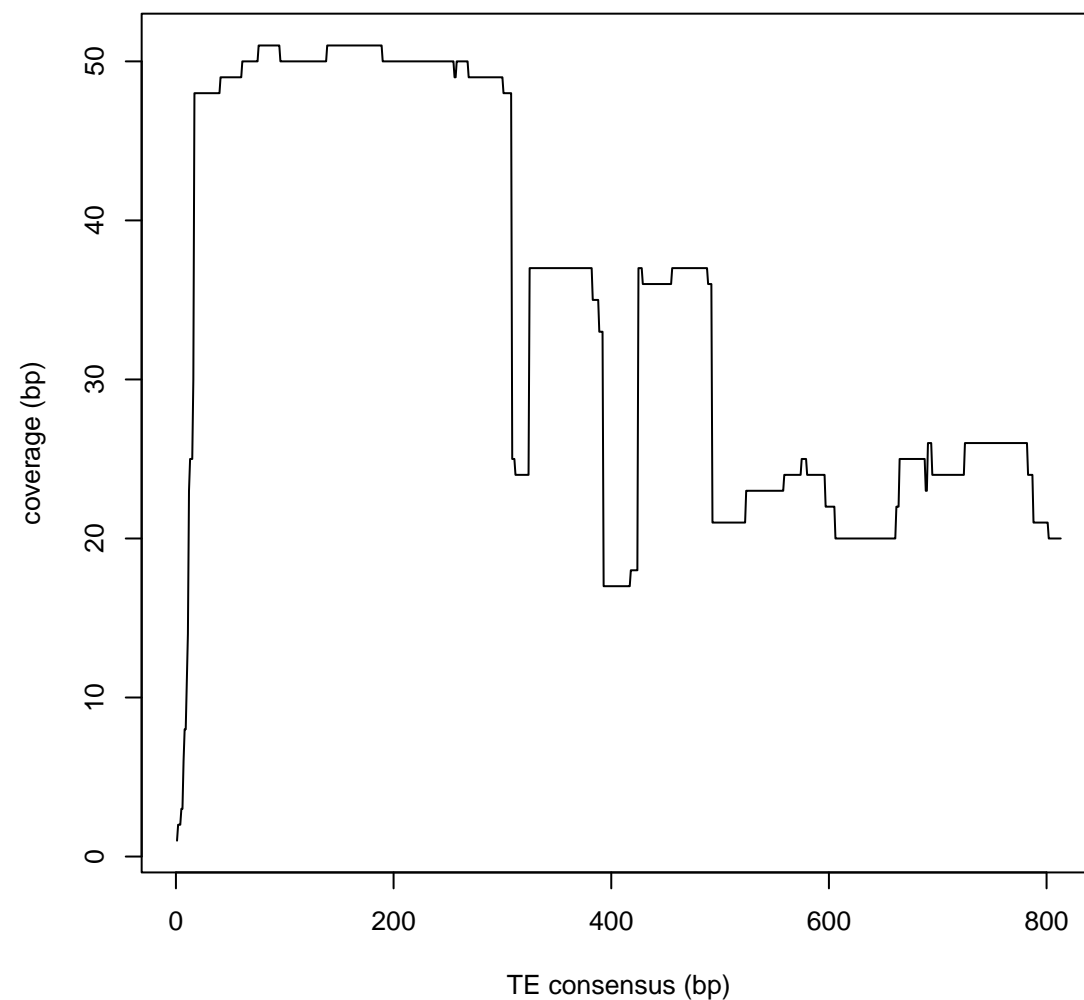


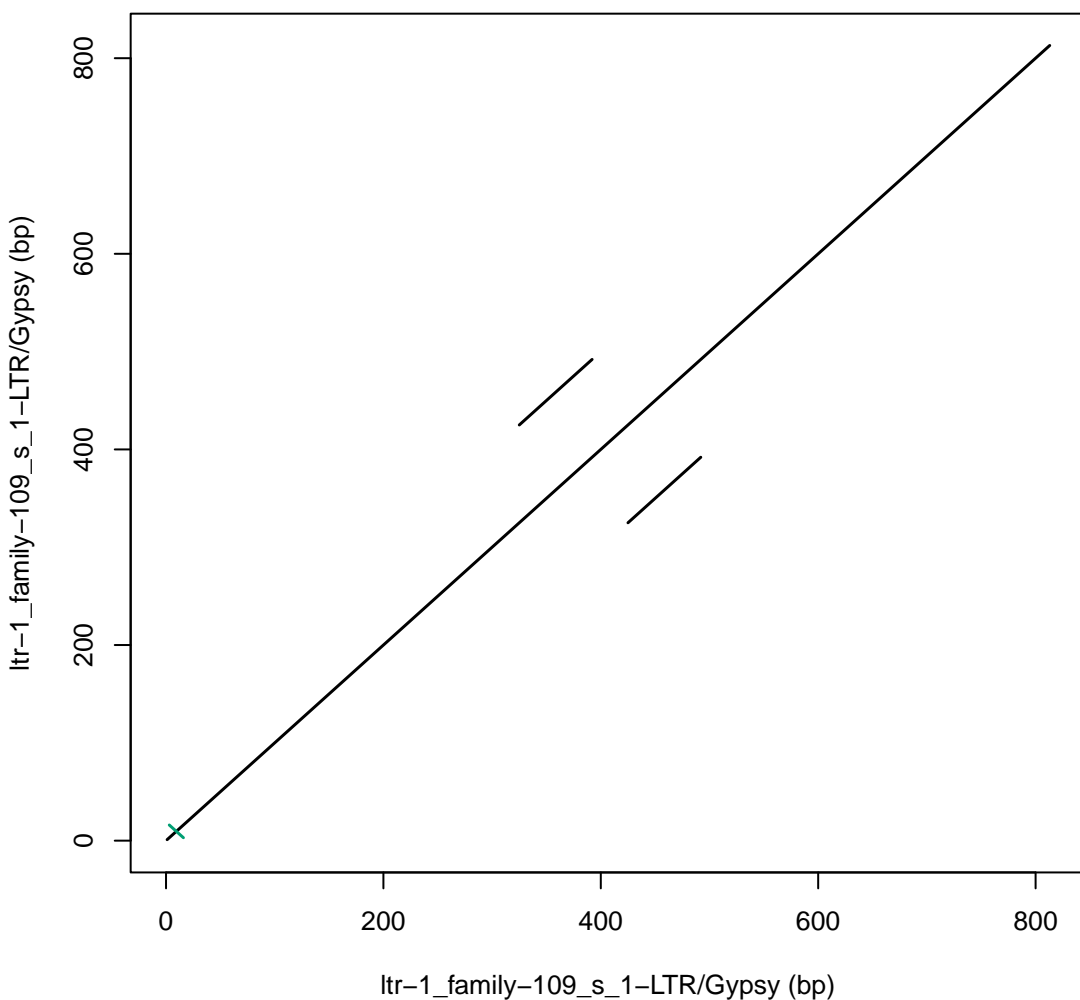
TE: ltr-1_family-109_s_1-LTR/Gypsy
consensus size: 813bp; fragments: 101; full length: 11 (>=731.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

