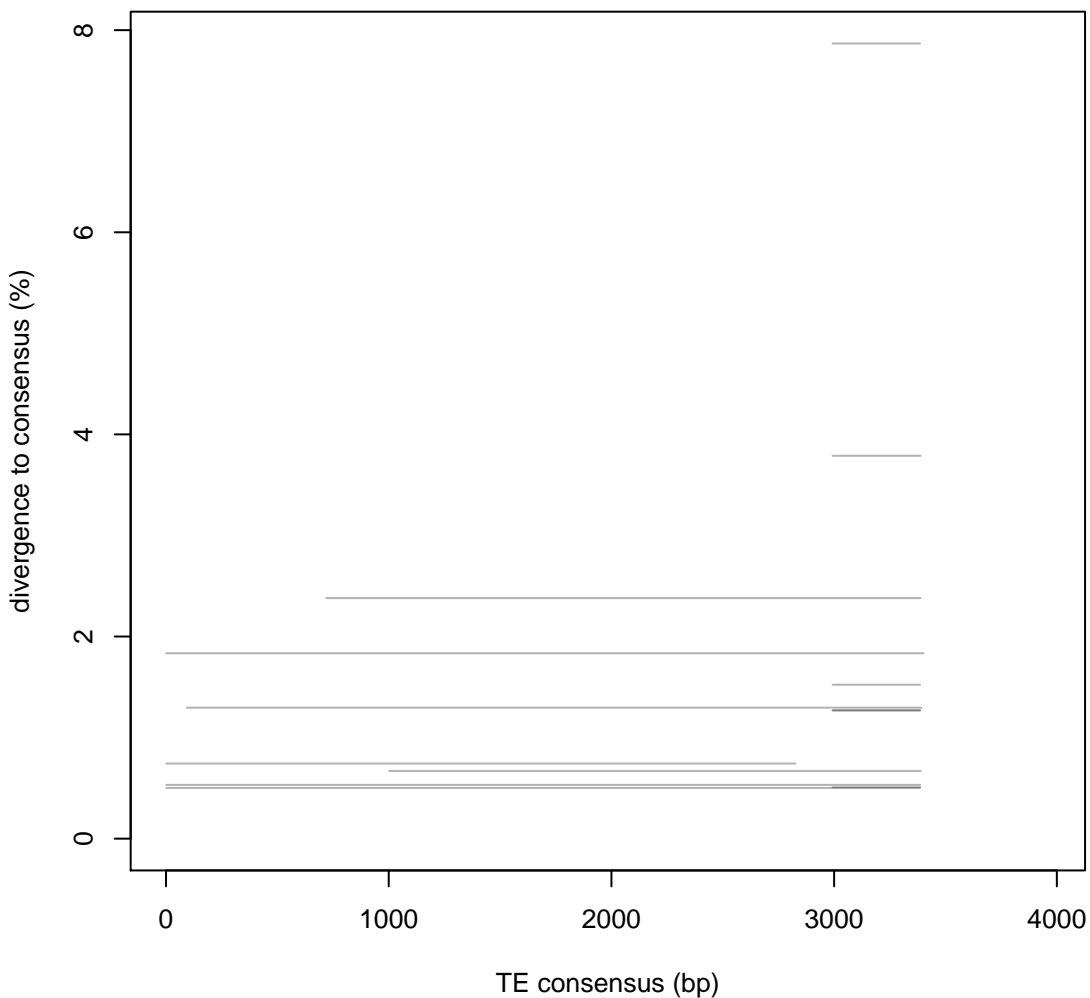
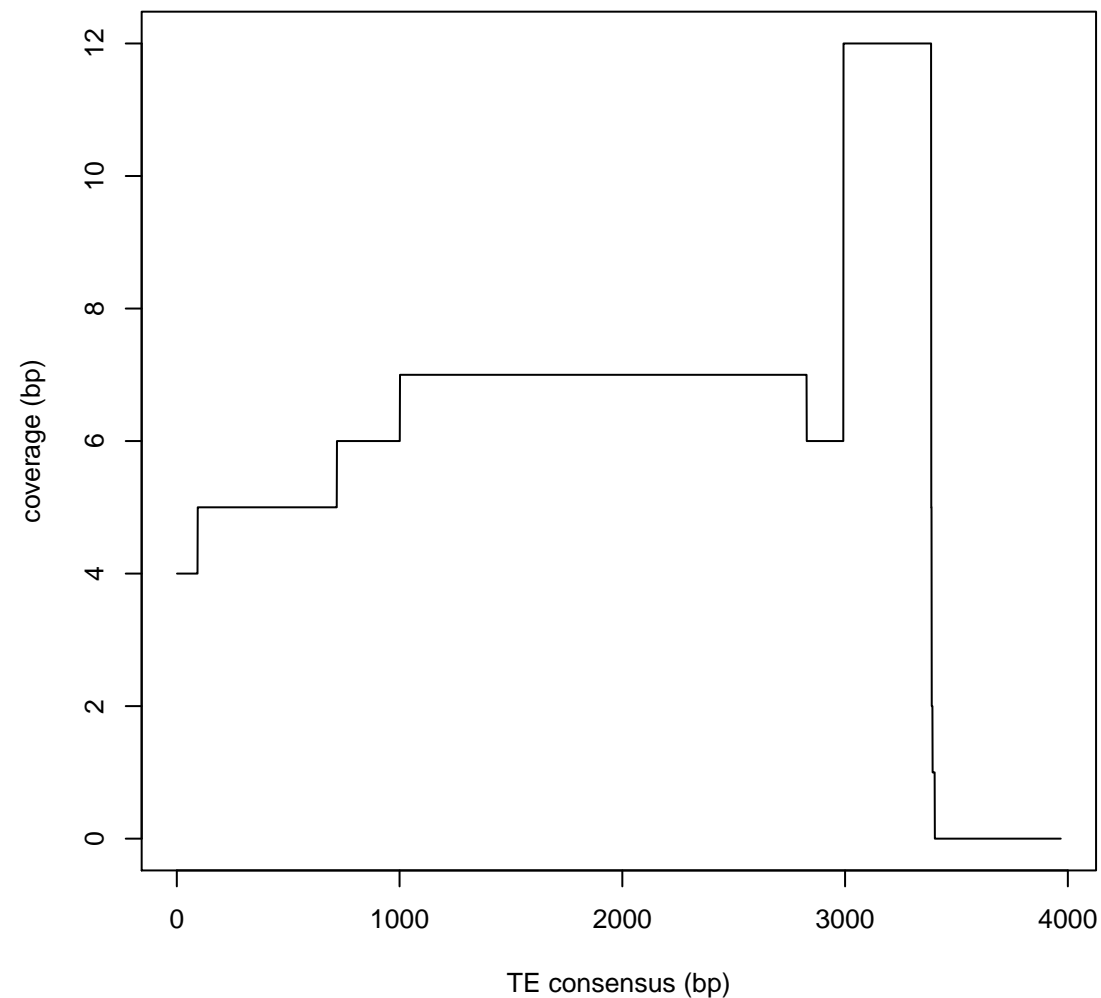


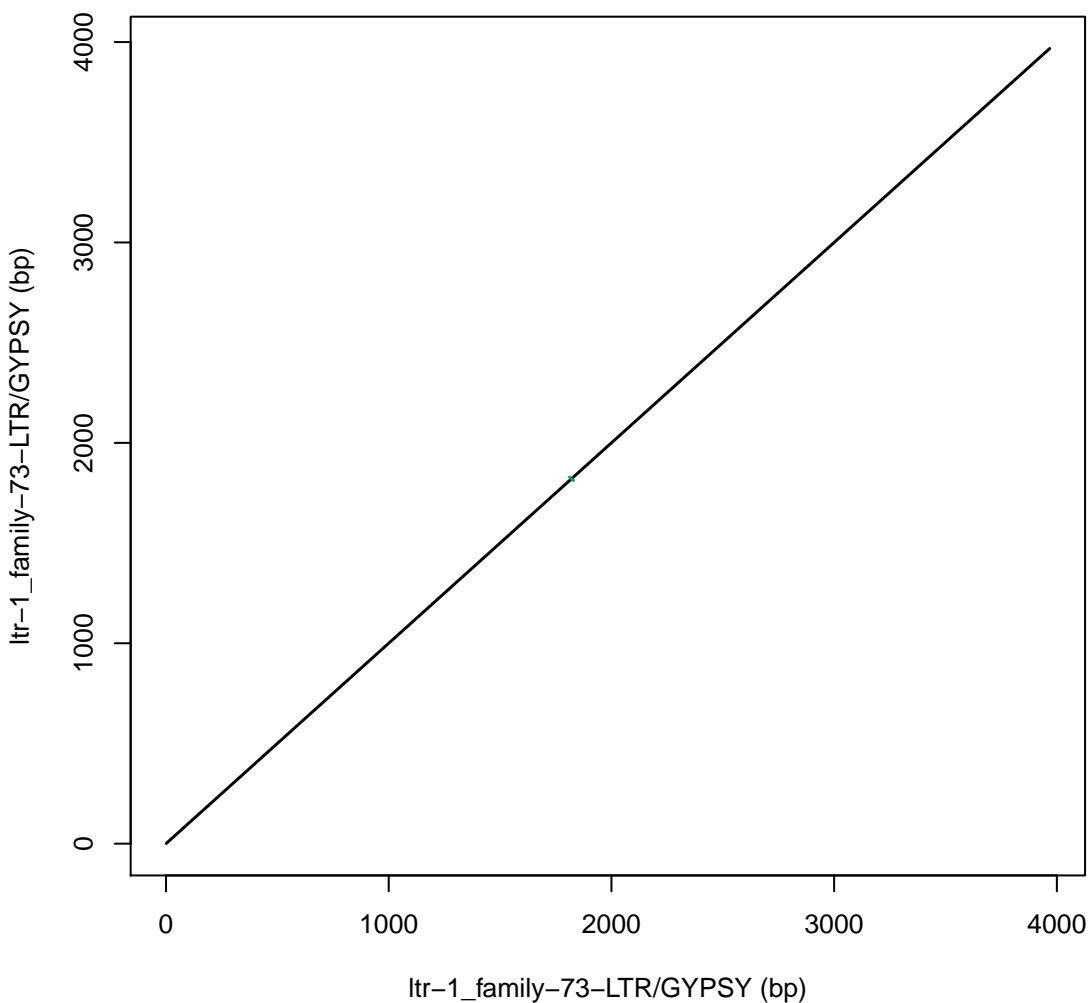
TE: ltr-1_family-73-LTR/GYPSY
consensus size: 3968bp; fragments: 13; full length: 0 (>=3571.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

