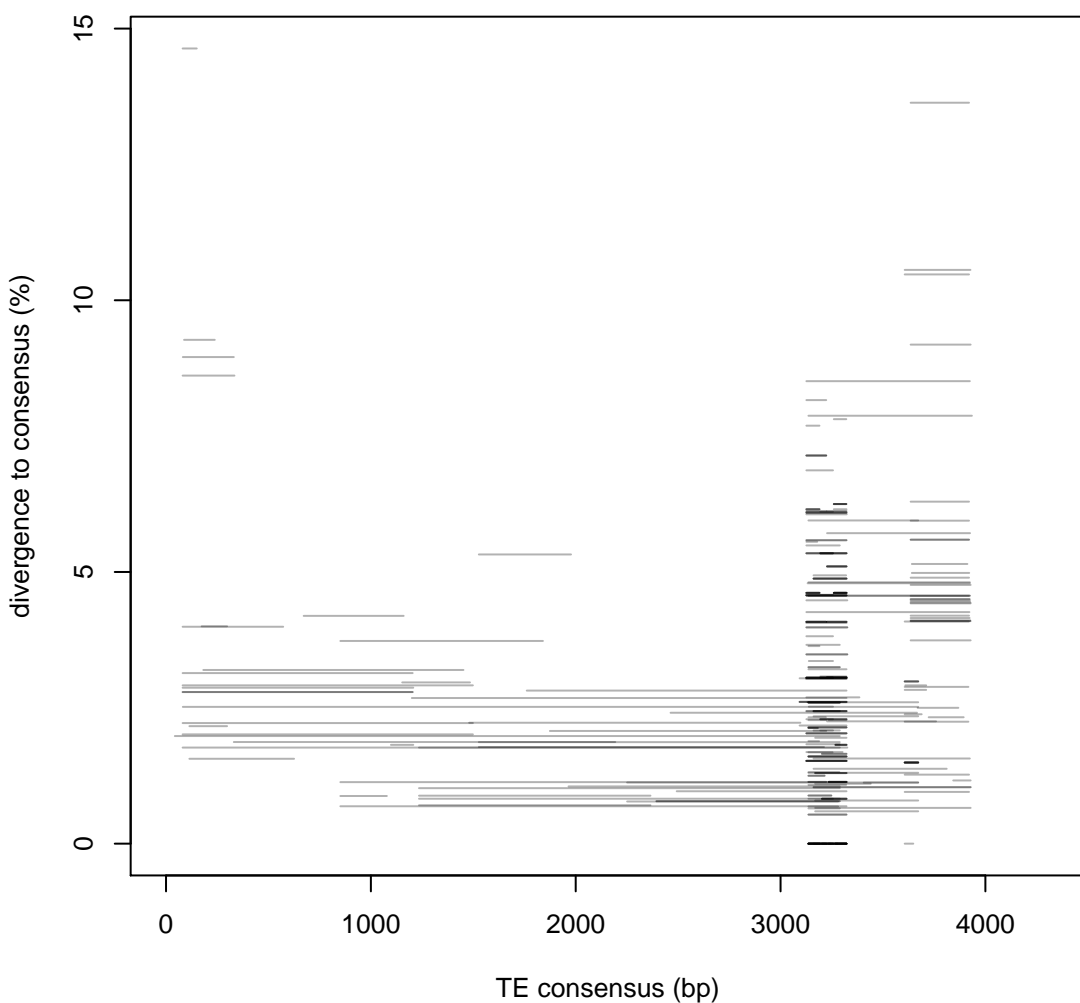
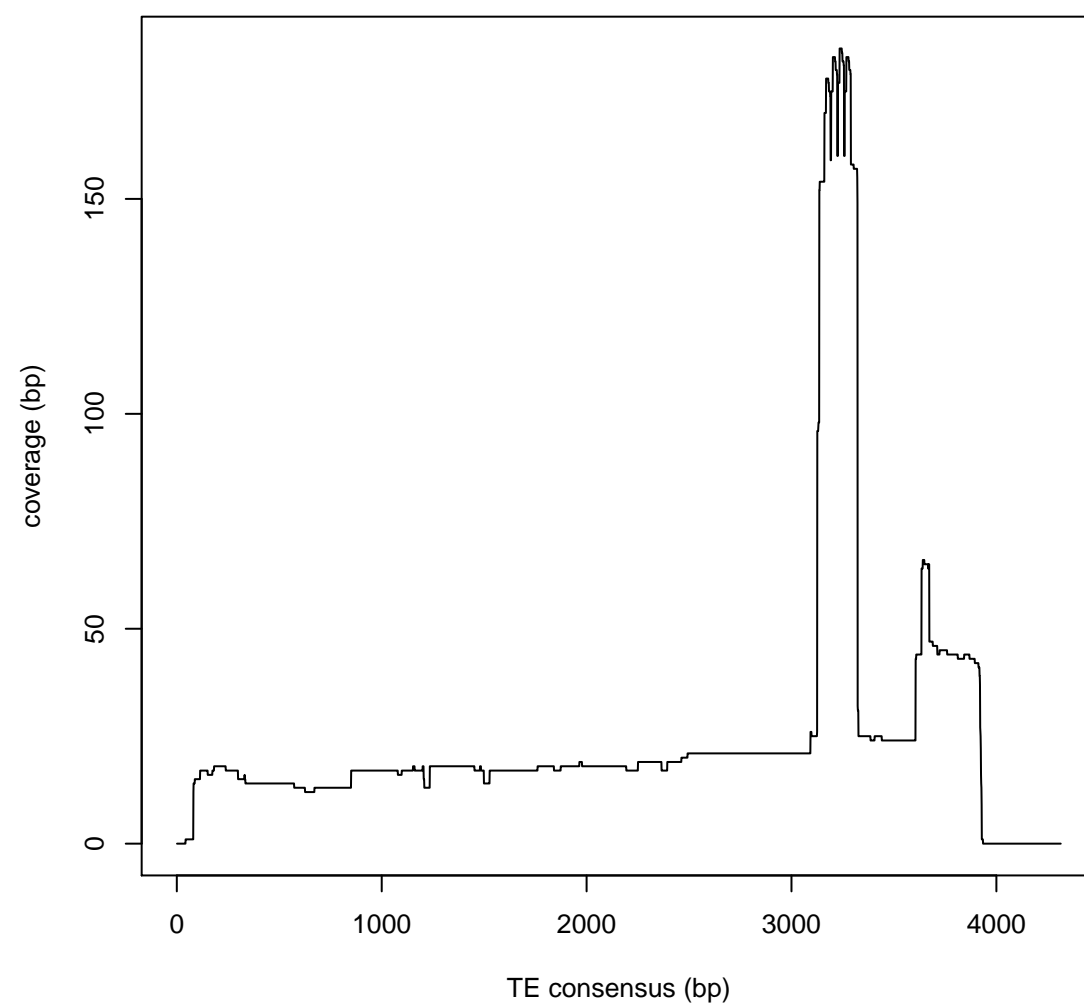


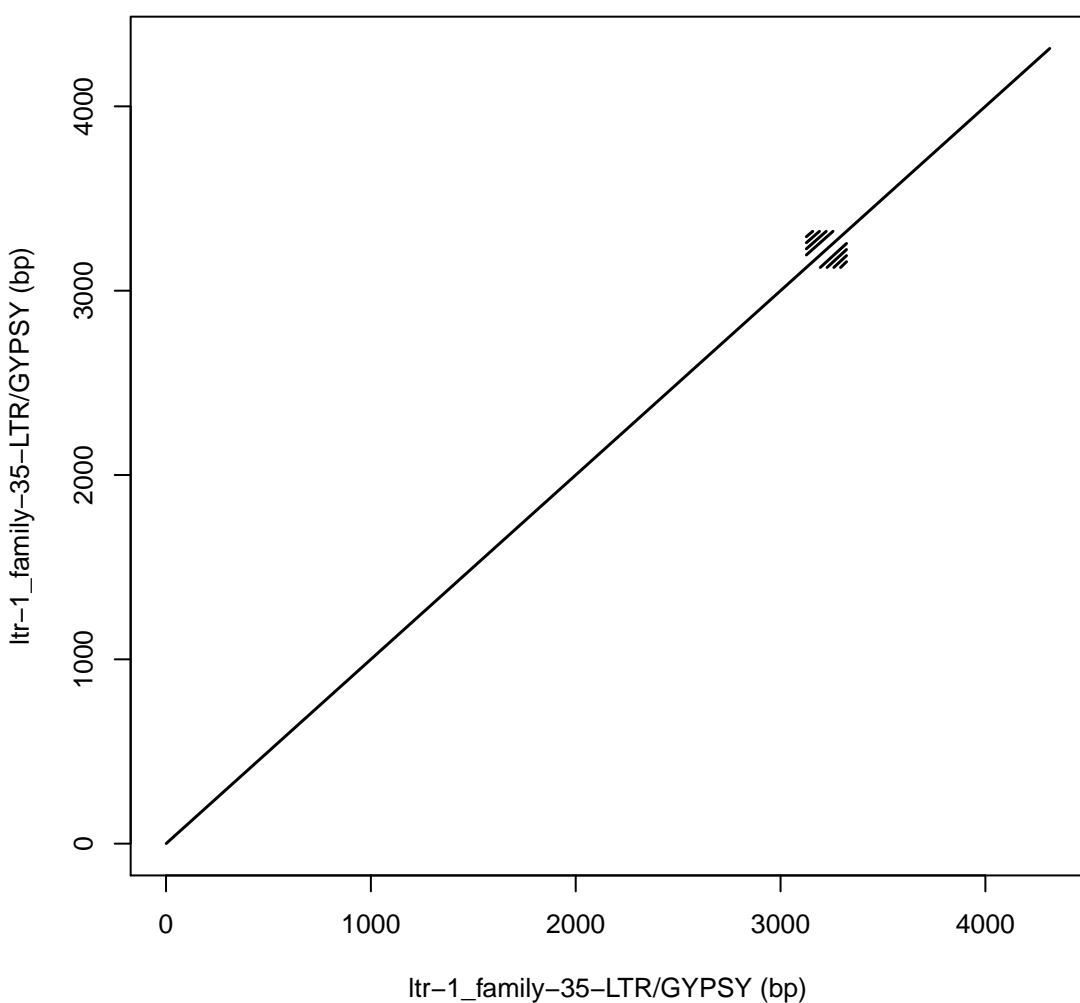
TE: ltr-1_family-35-LTR/GYPSY
consensus size: 4314bp; fragments: 322; full length: 0 (>=3882.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

