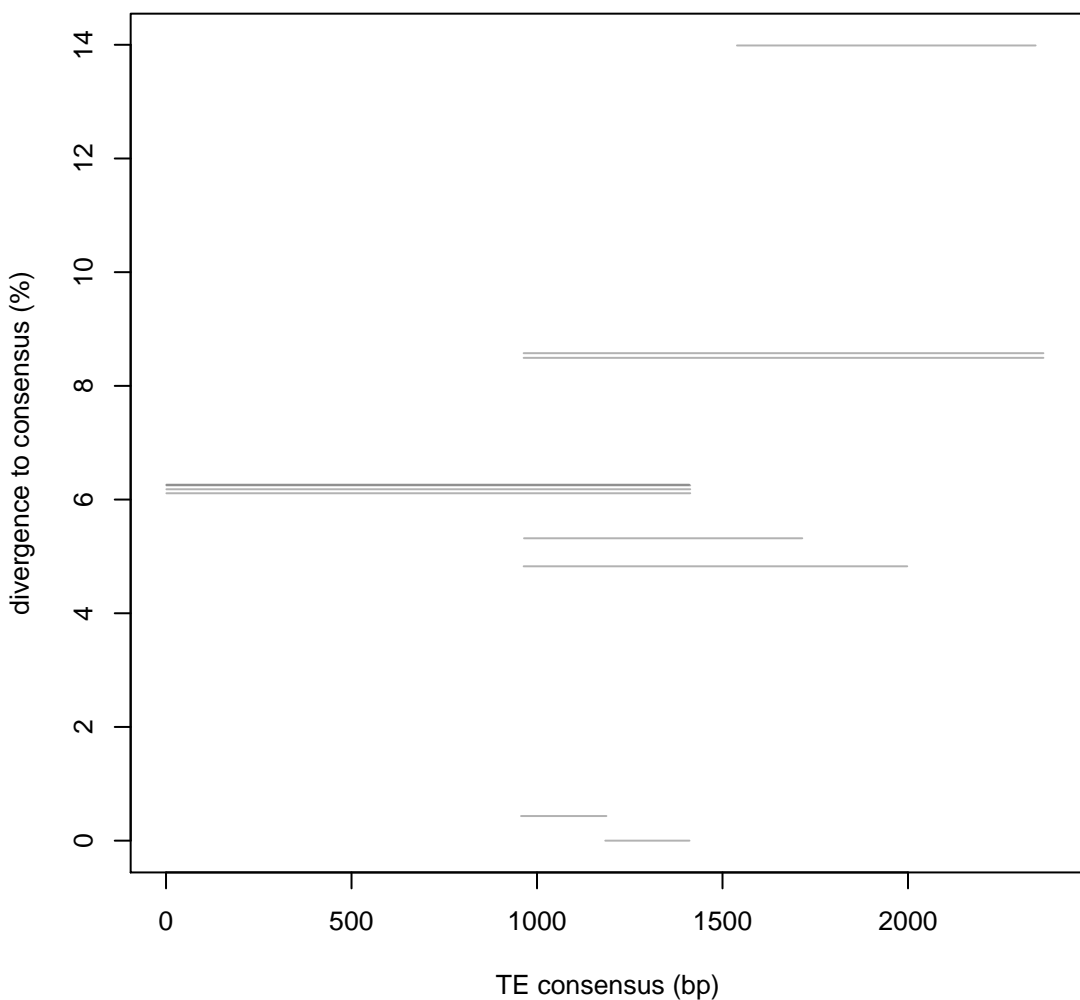
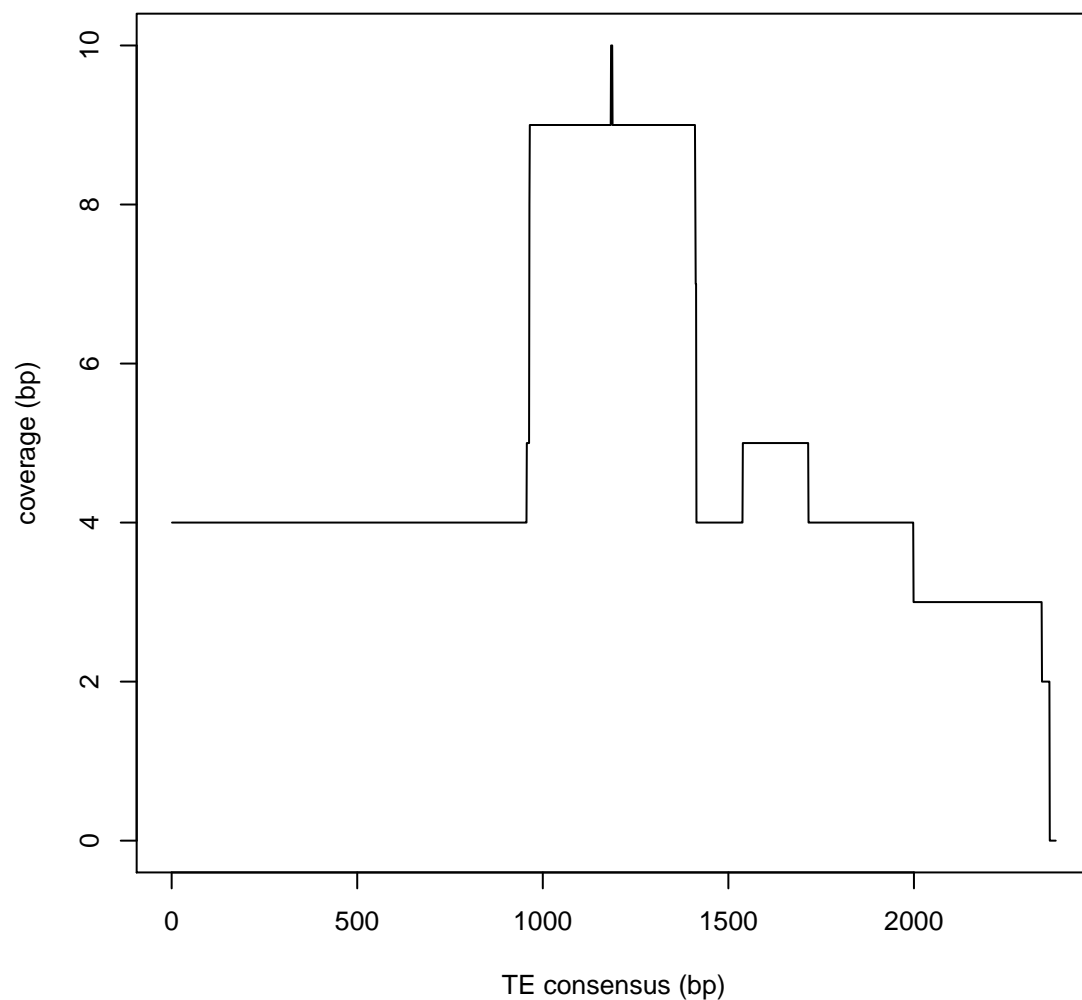


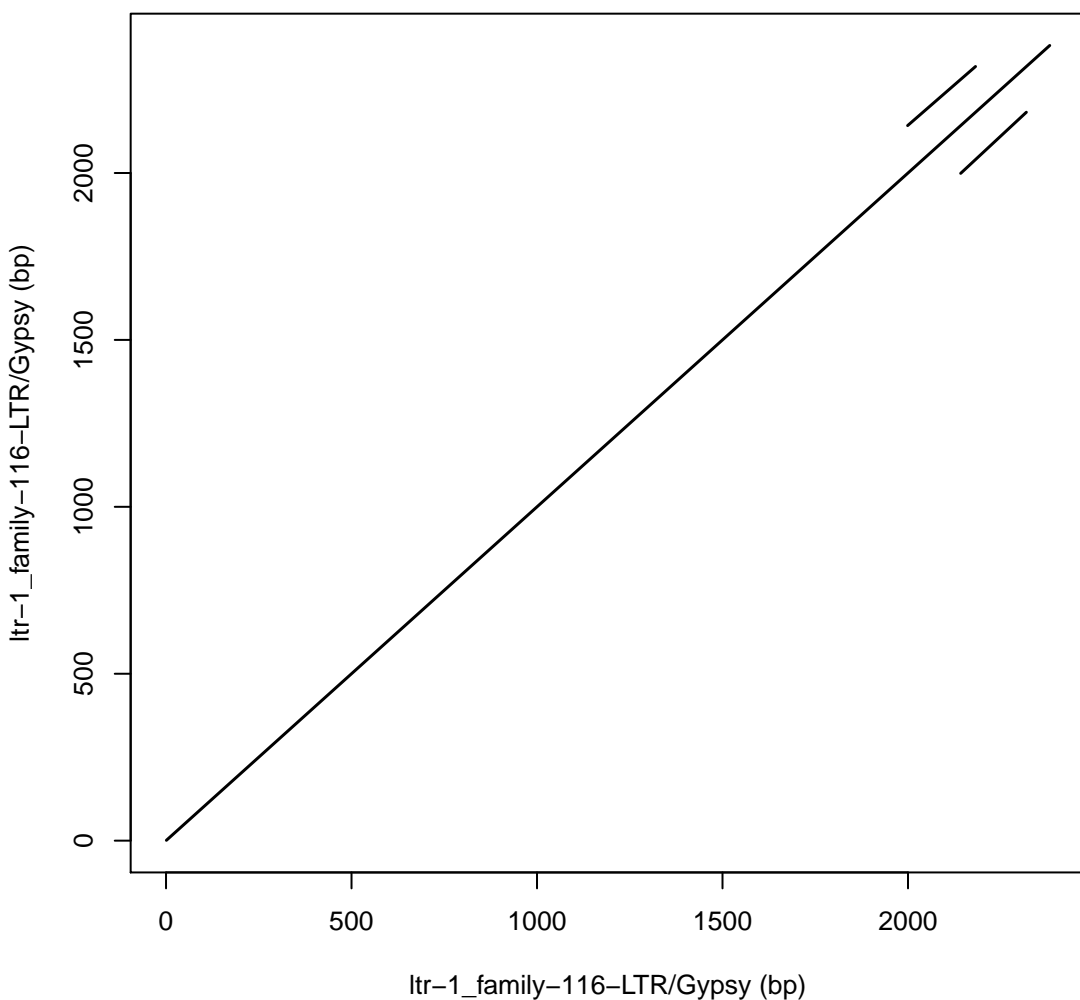
TE: ltr-1_family-116-LTR/Gypsy
consensus size: 2382bp; fragments: 11; full length: 0 (≥ 2143.8 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

