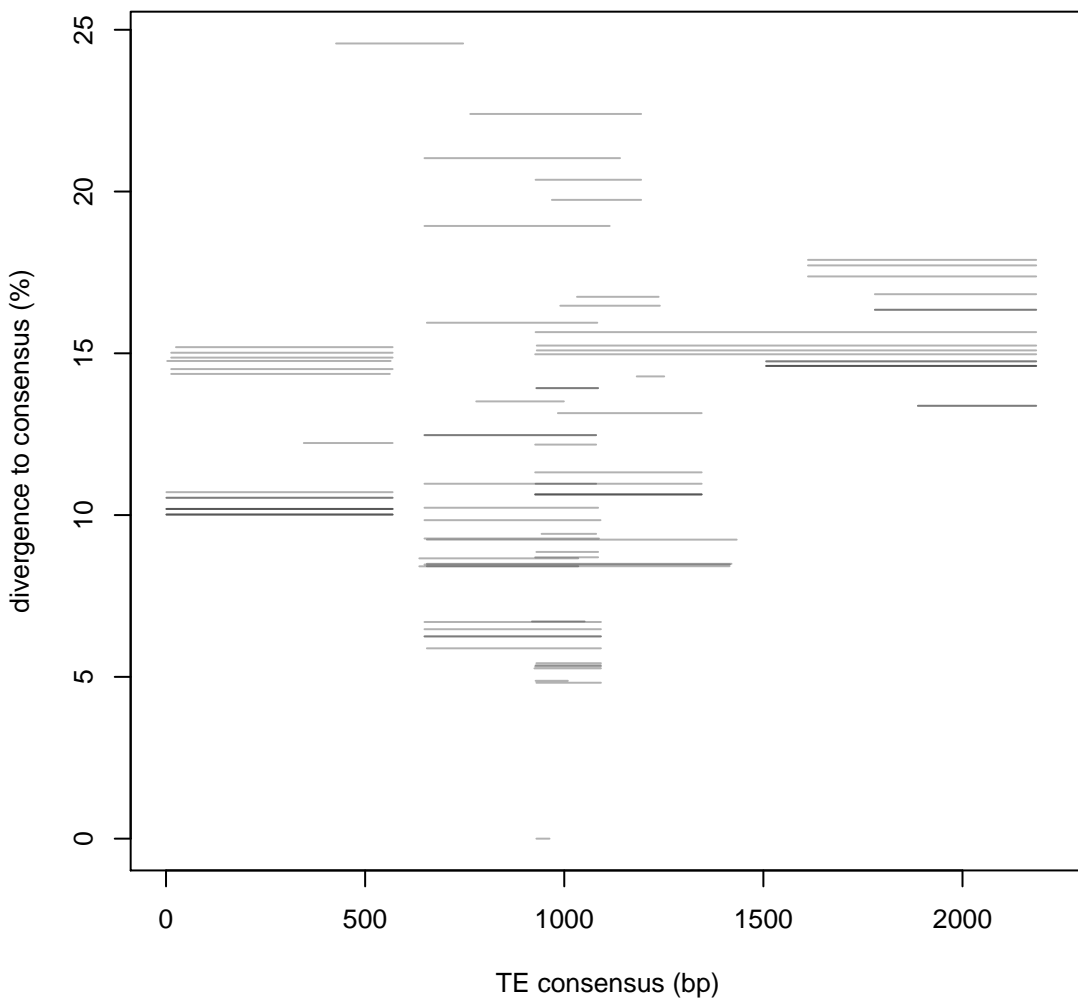
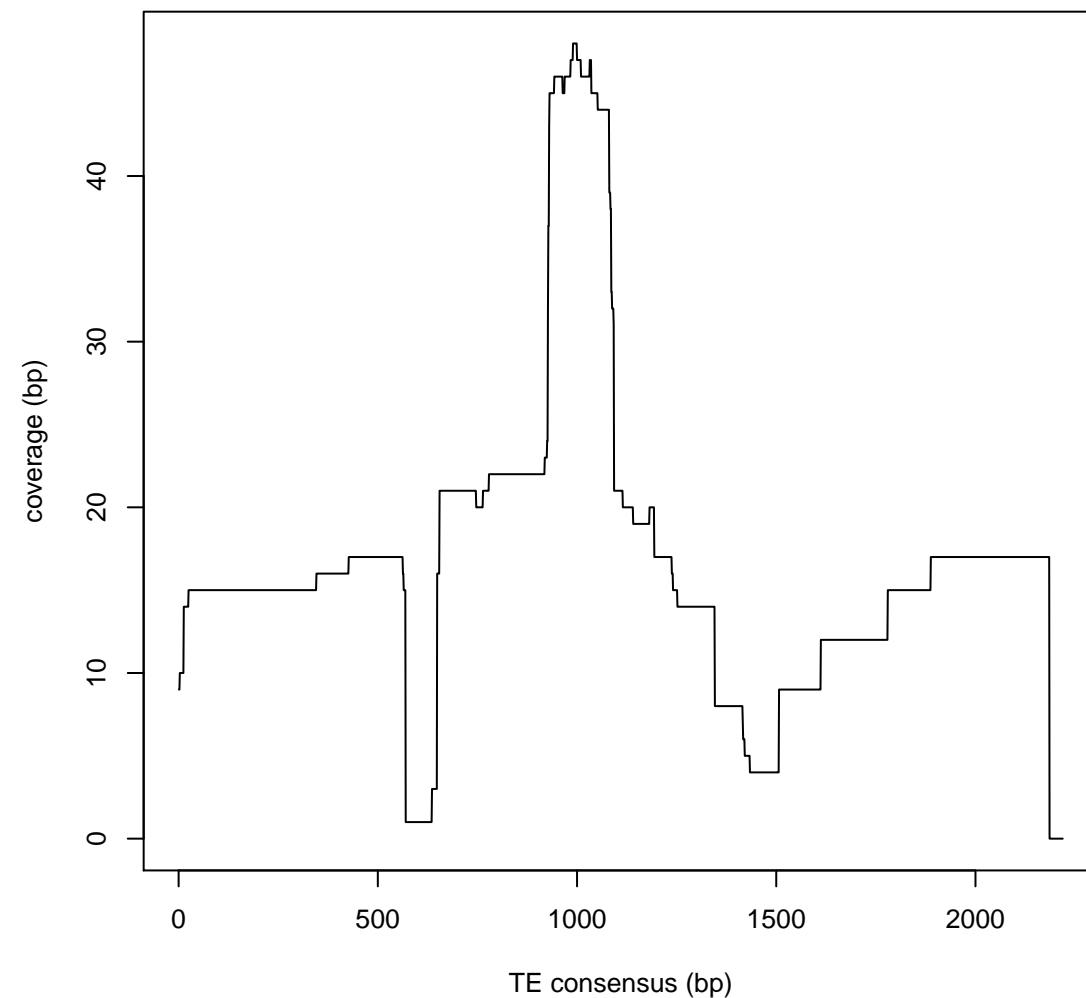


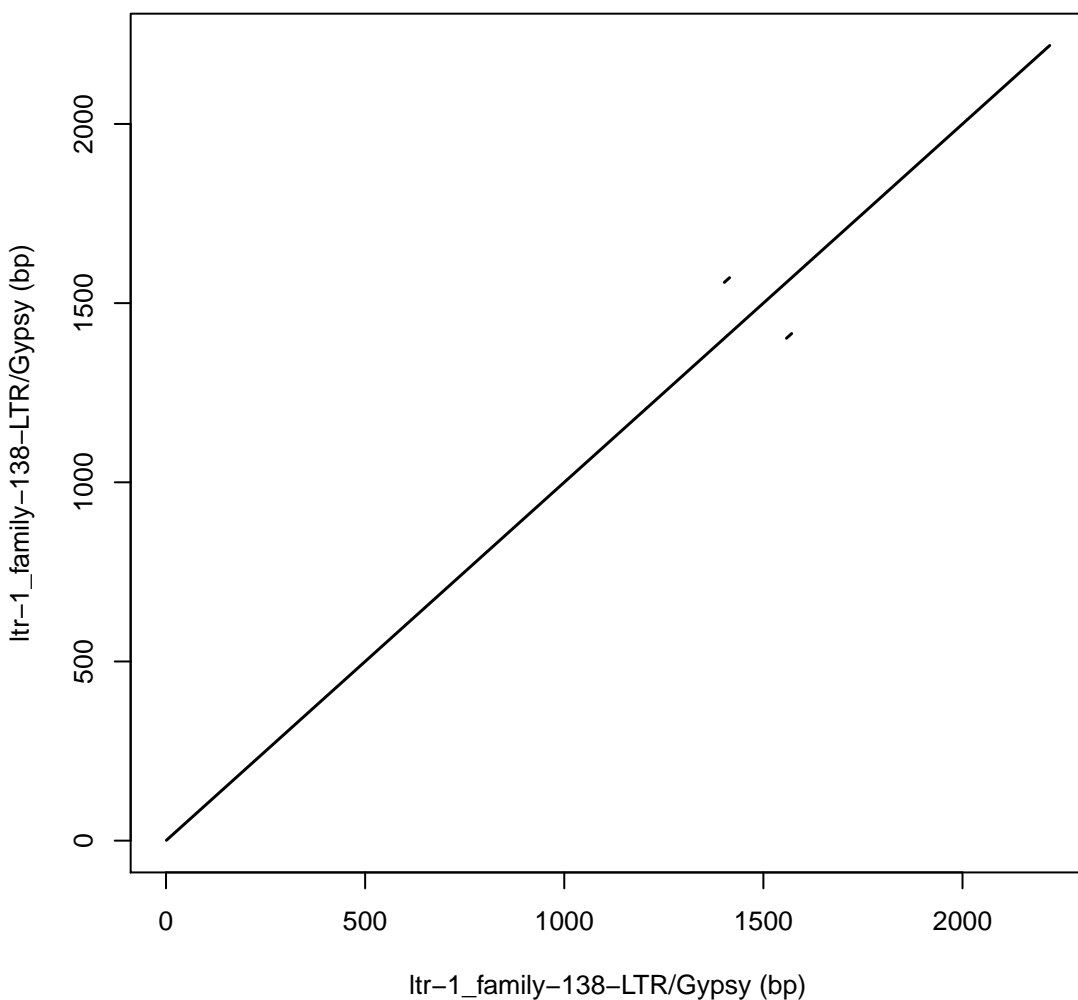
TE: ltr-1_family-138-LTR/Gypsy
consensus size: 2219bp; fragments: 81; full length: 0 (≥ 1997.1 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

