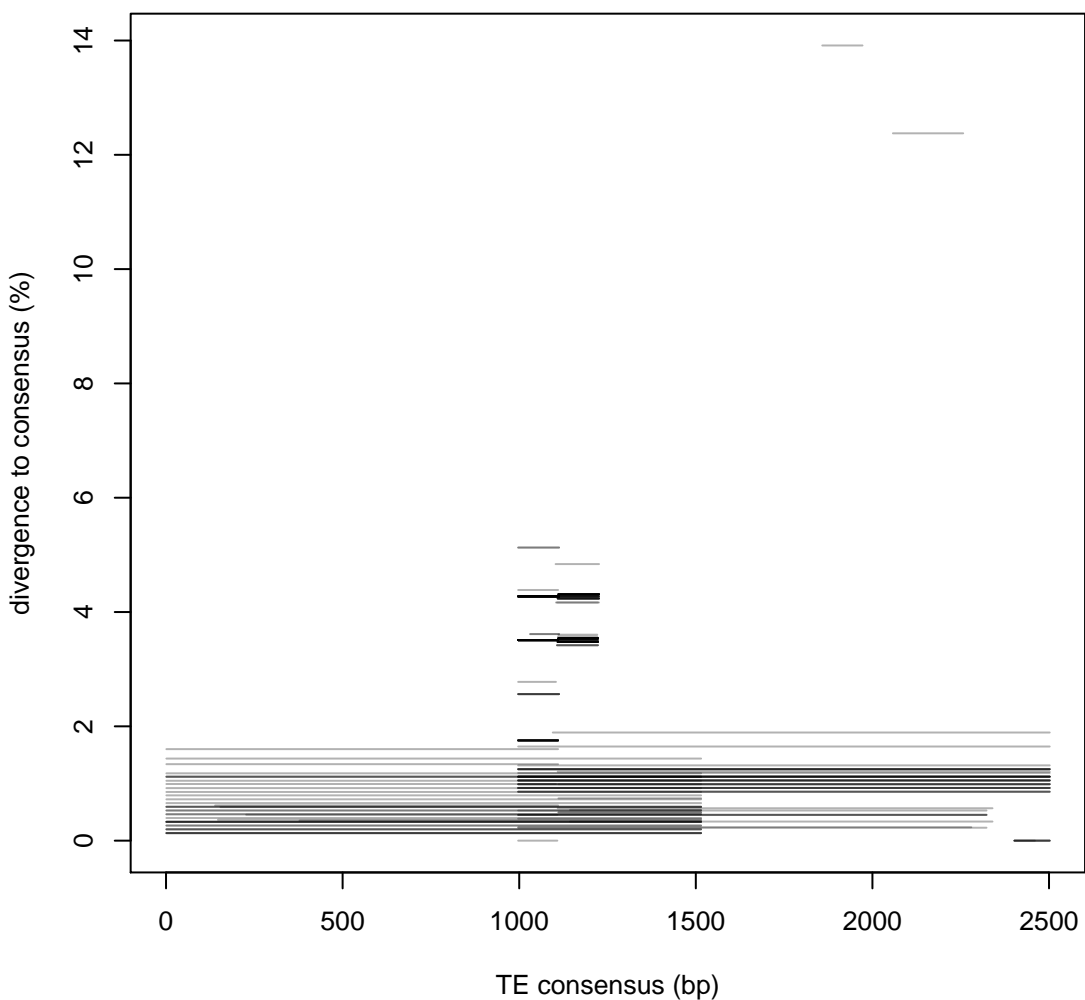
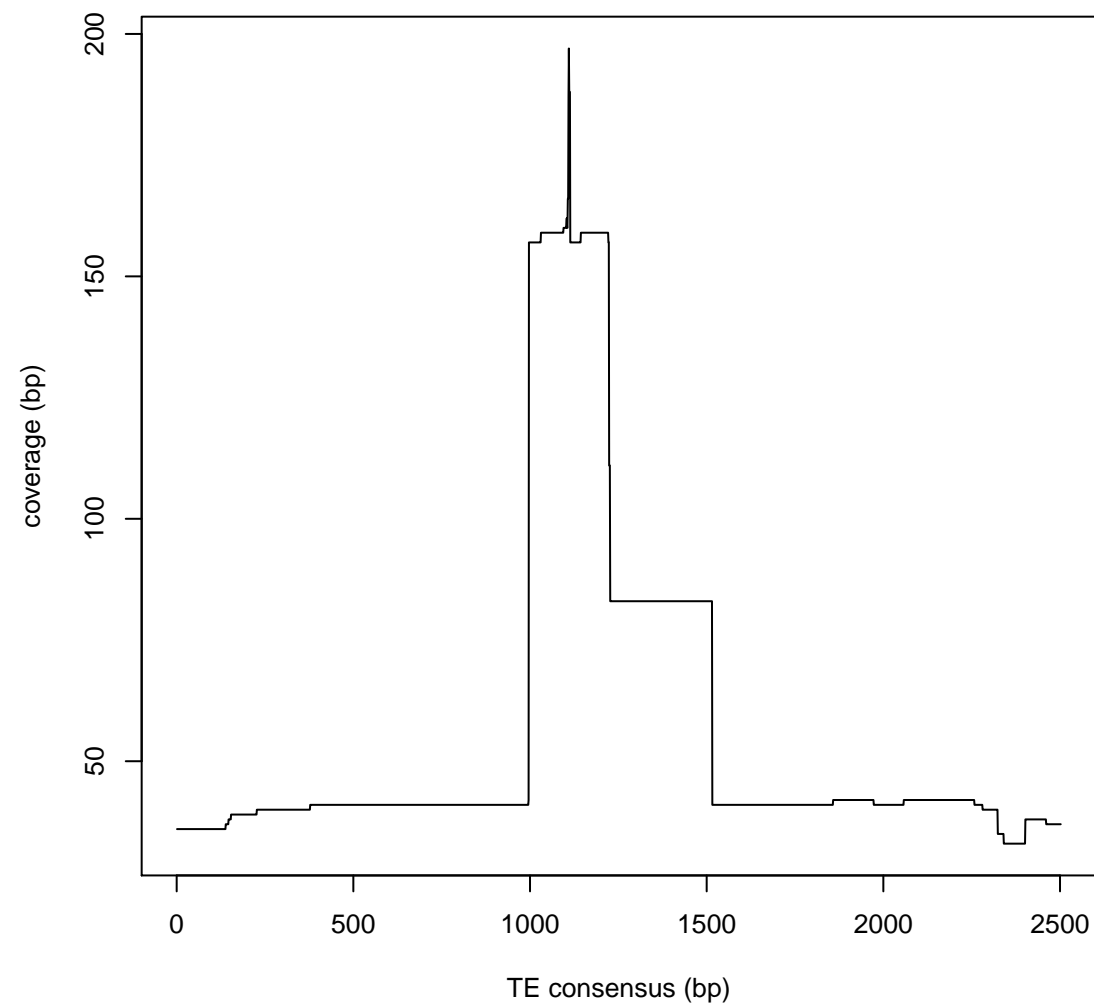


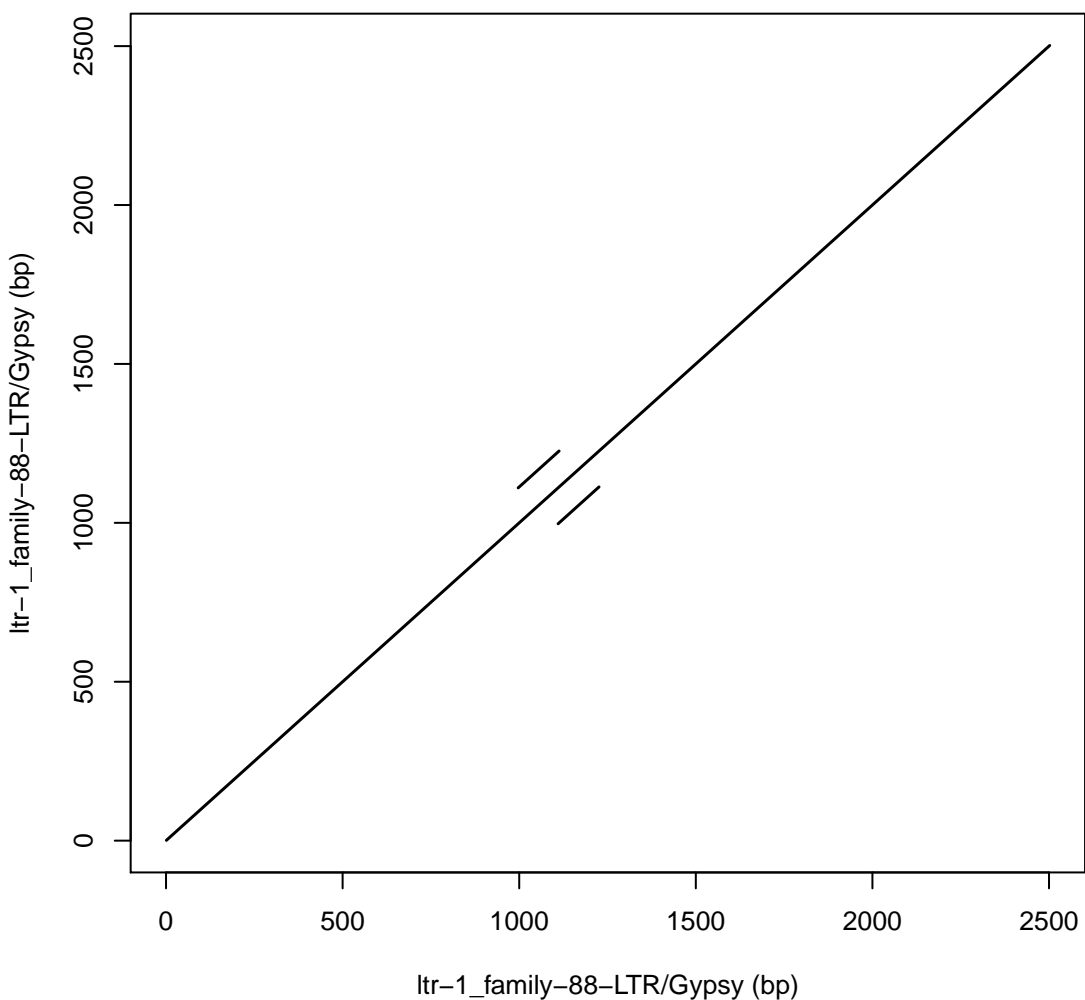
TE: ltr-1_family-88-LTR/Gypsy
consensus size: 2502bp; fragments: 251; full length: 0 (>=2251.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

