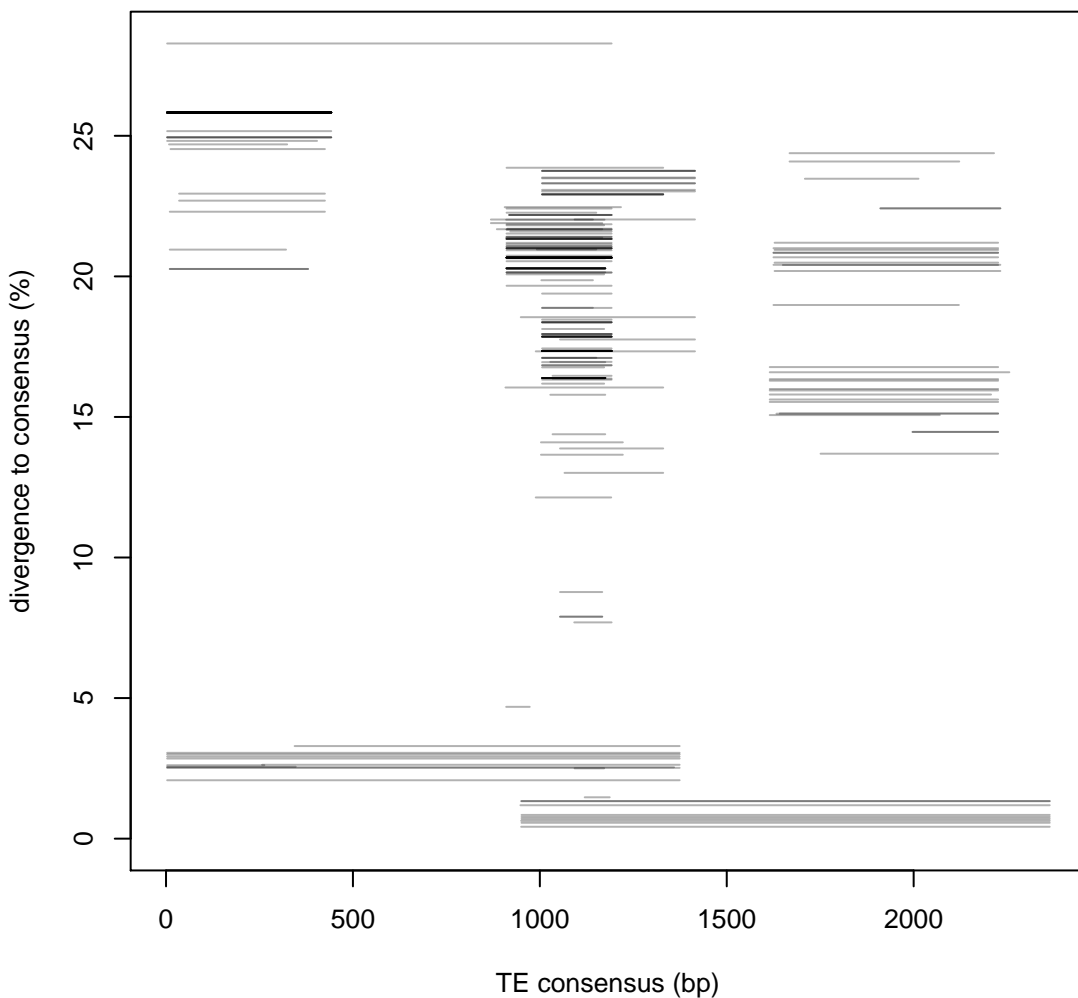
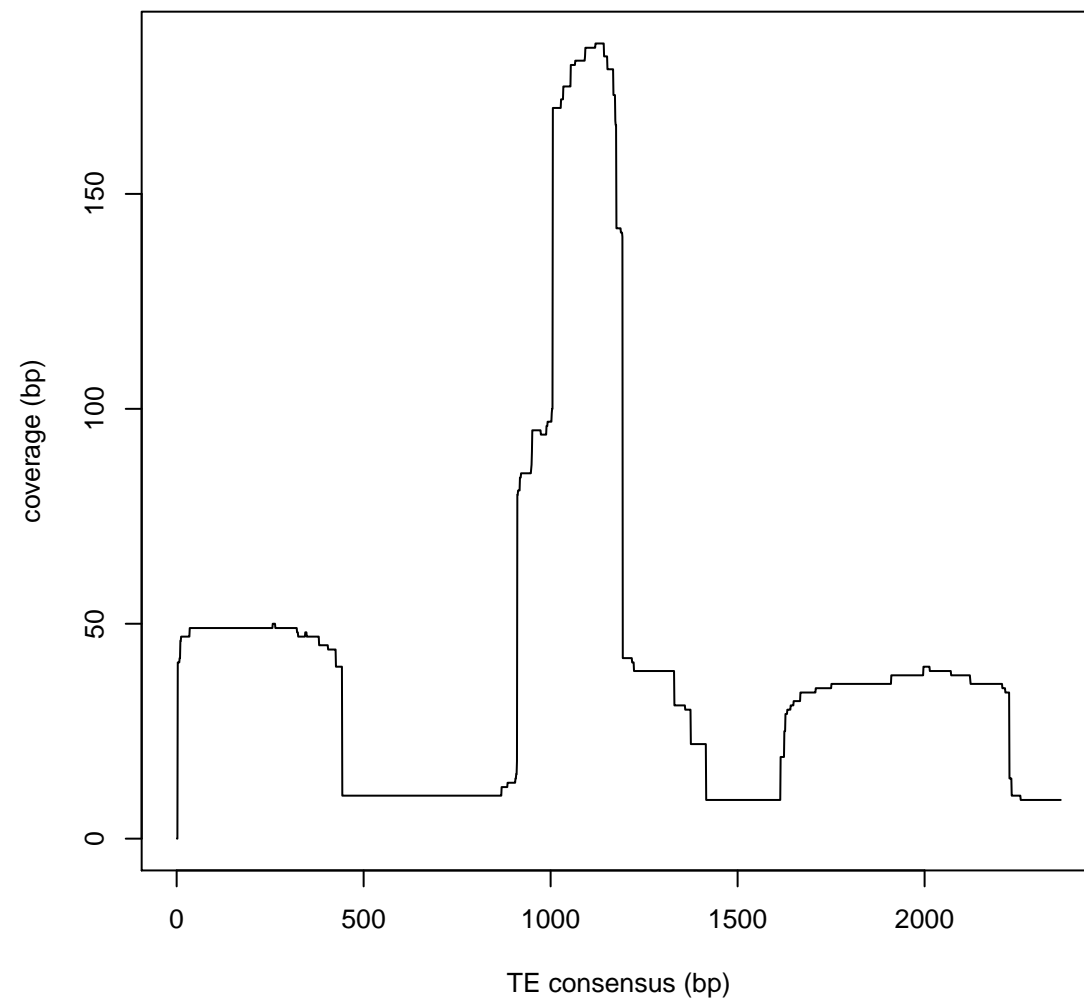


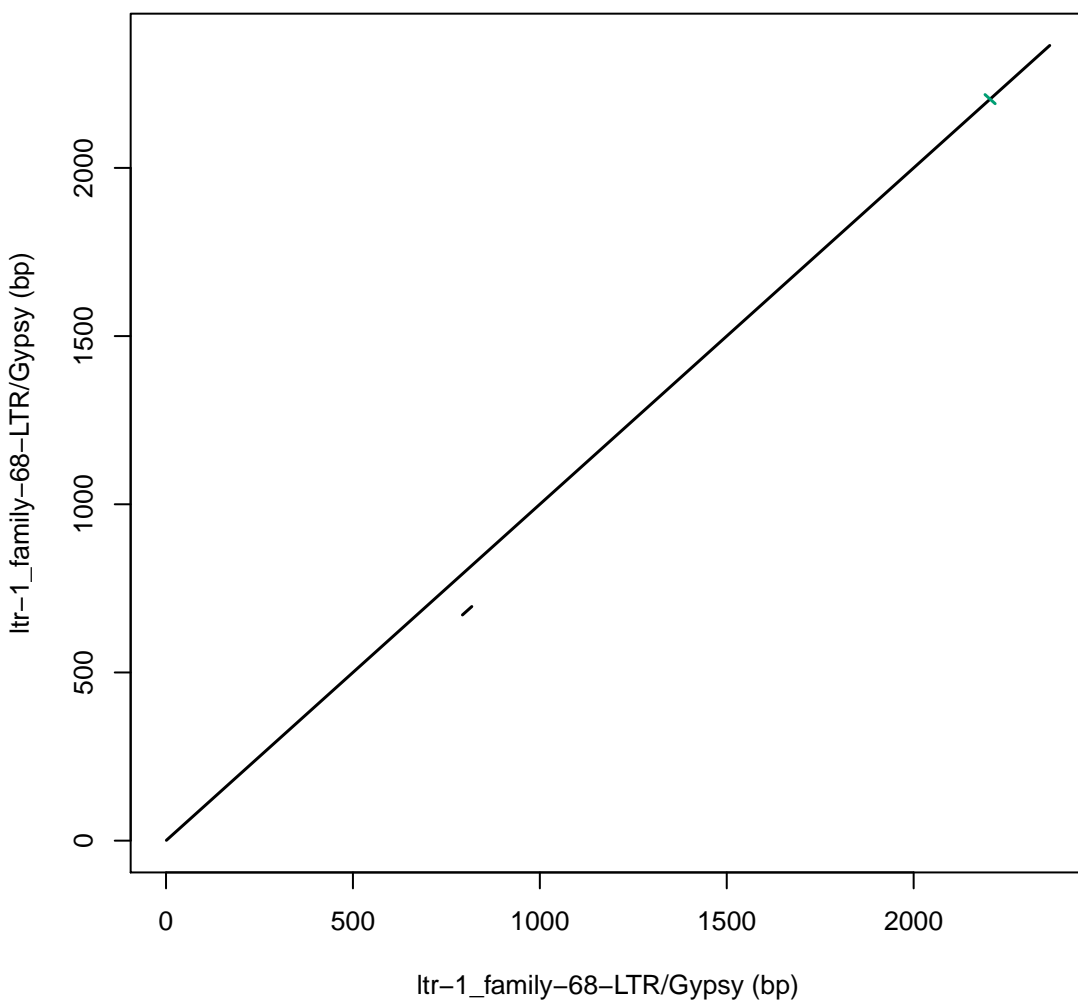
TE: ltr-1_family-68-LTR/Gypsy
consensus size: 2364bp; fragments: 258; full length: 0 (>=2127.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

