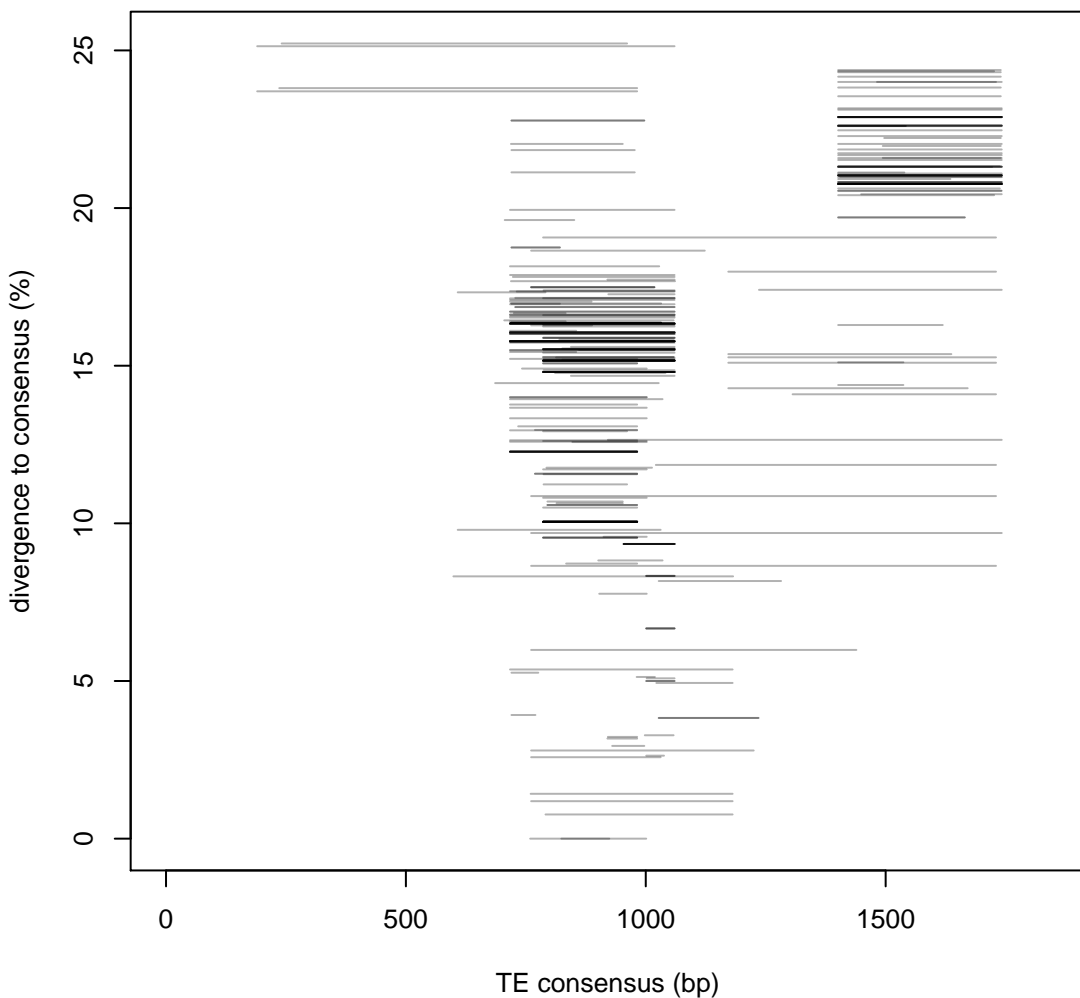
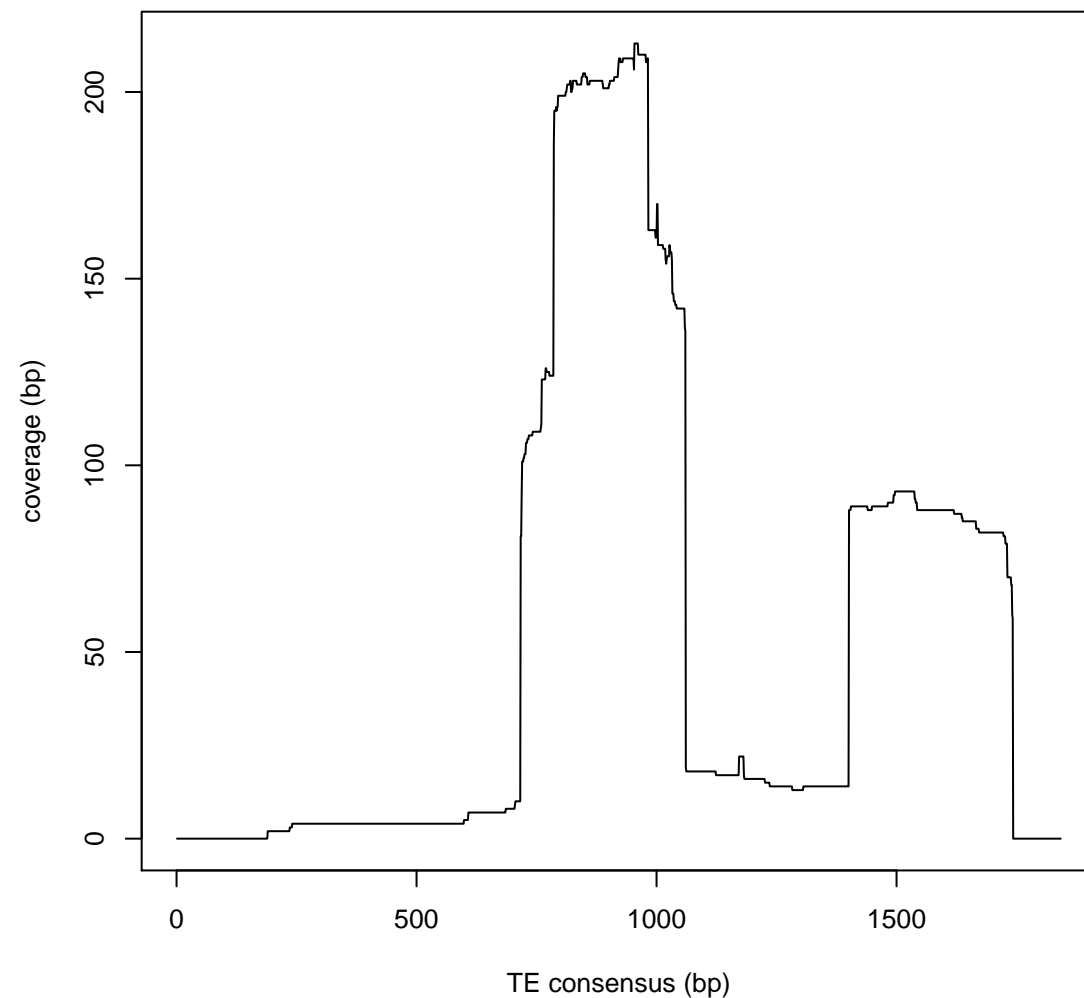


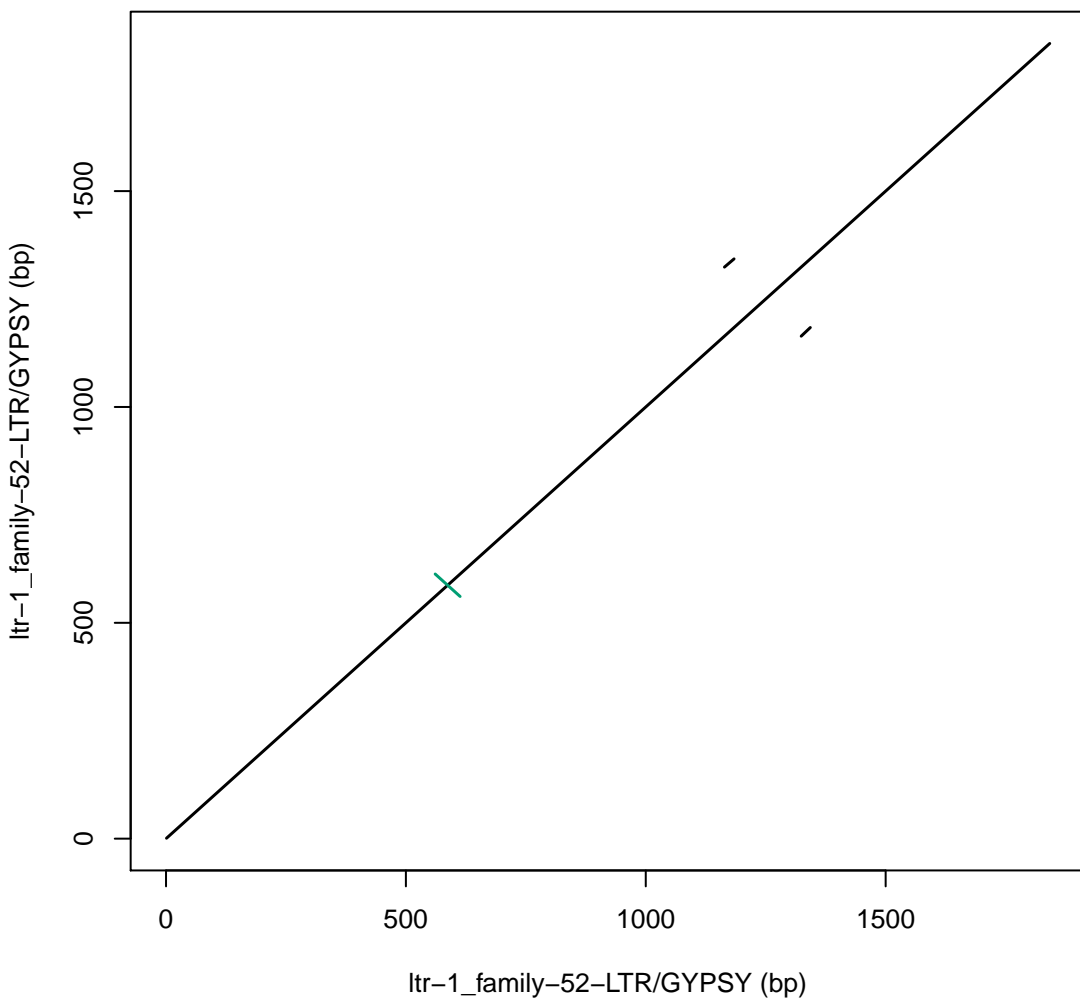
TE: ltr-1_family-52-LTR/GYPSY
consensus size: 1842bp; fragments: 334; full length: 0 (>=1657.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

