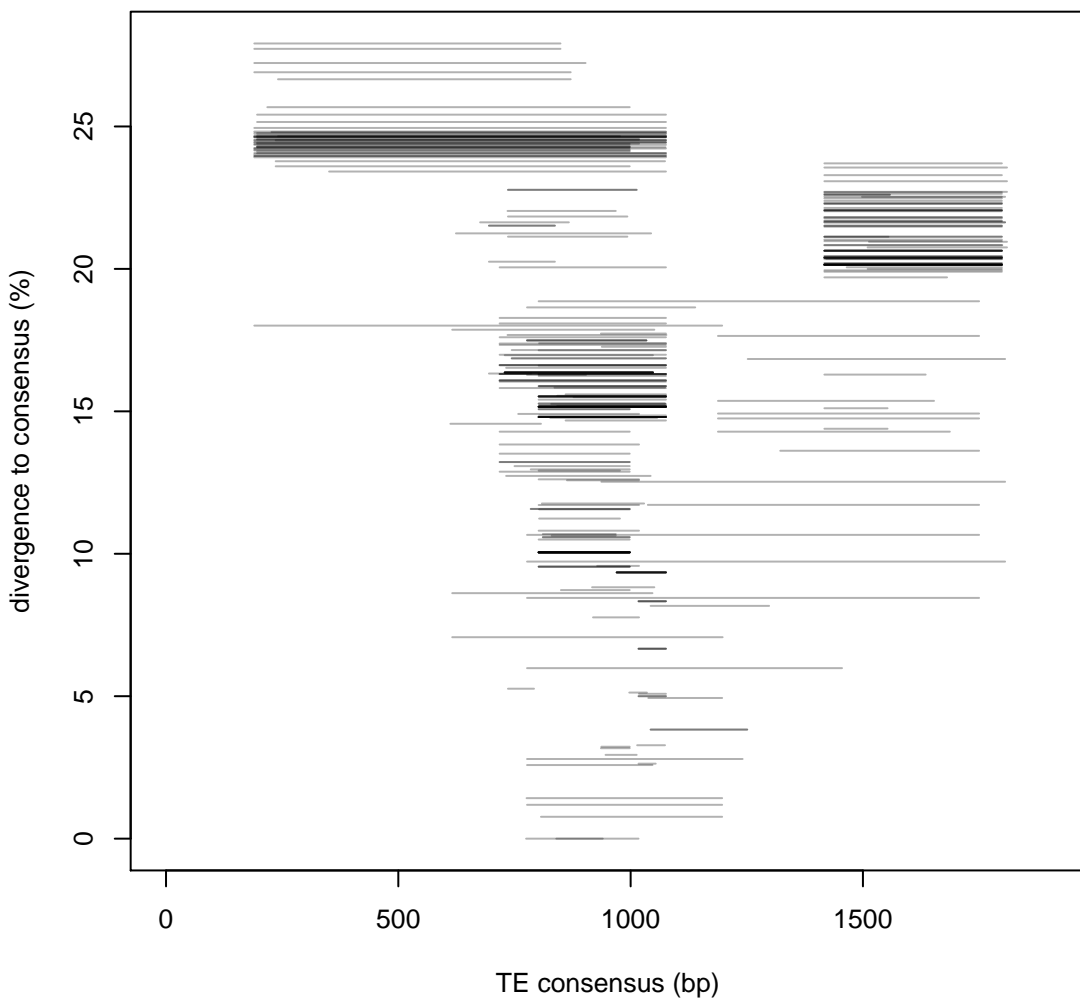
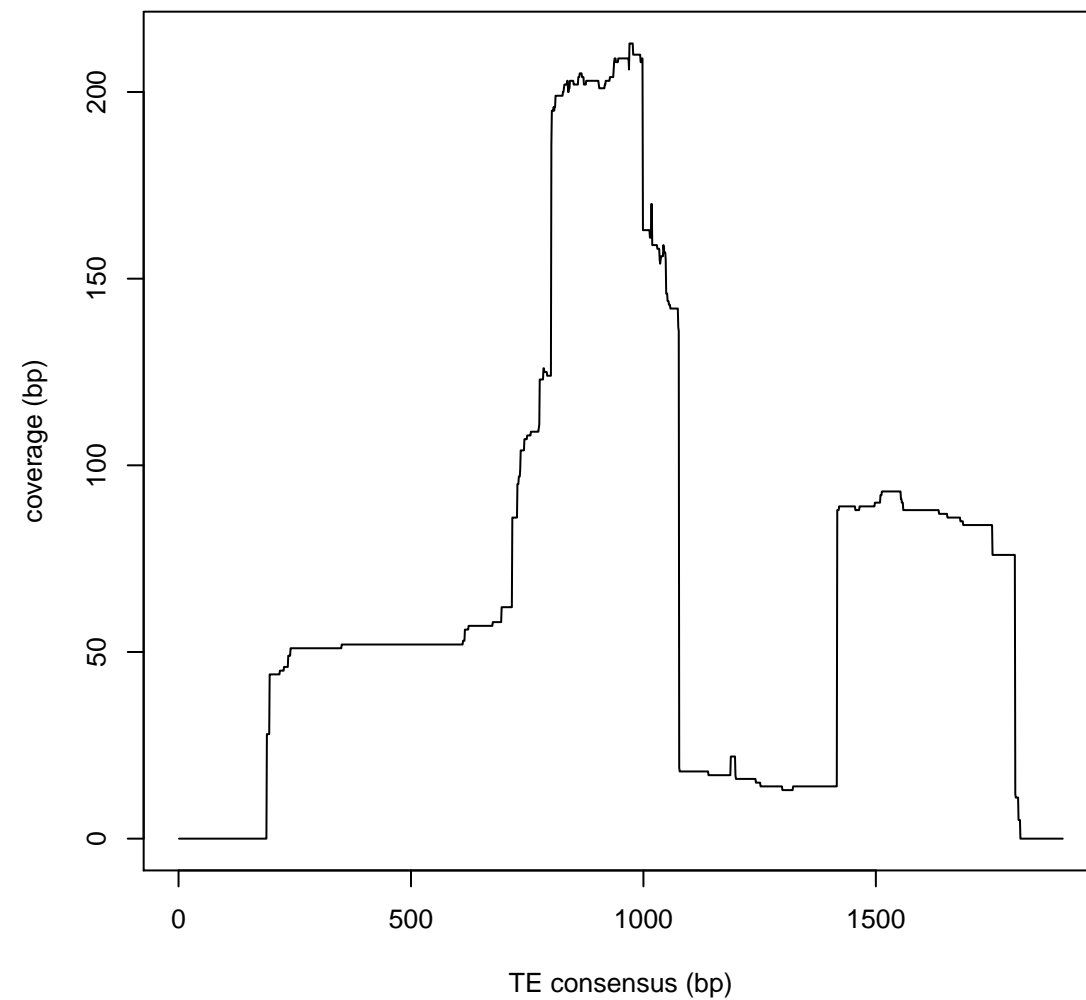


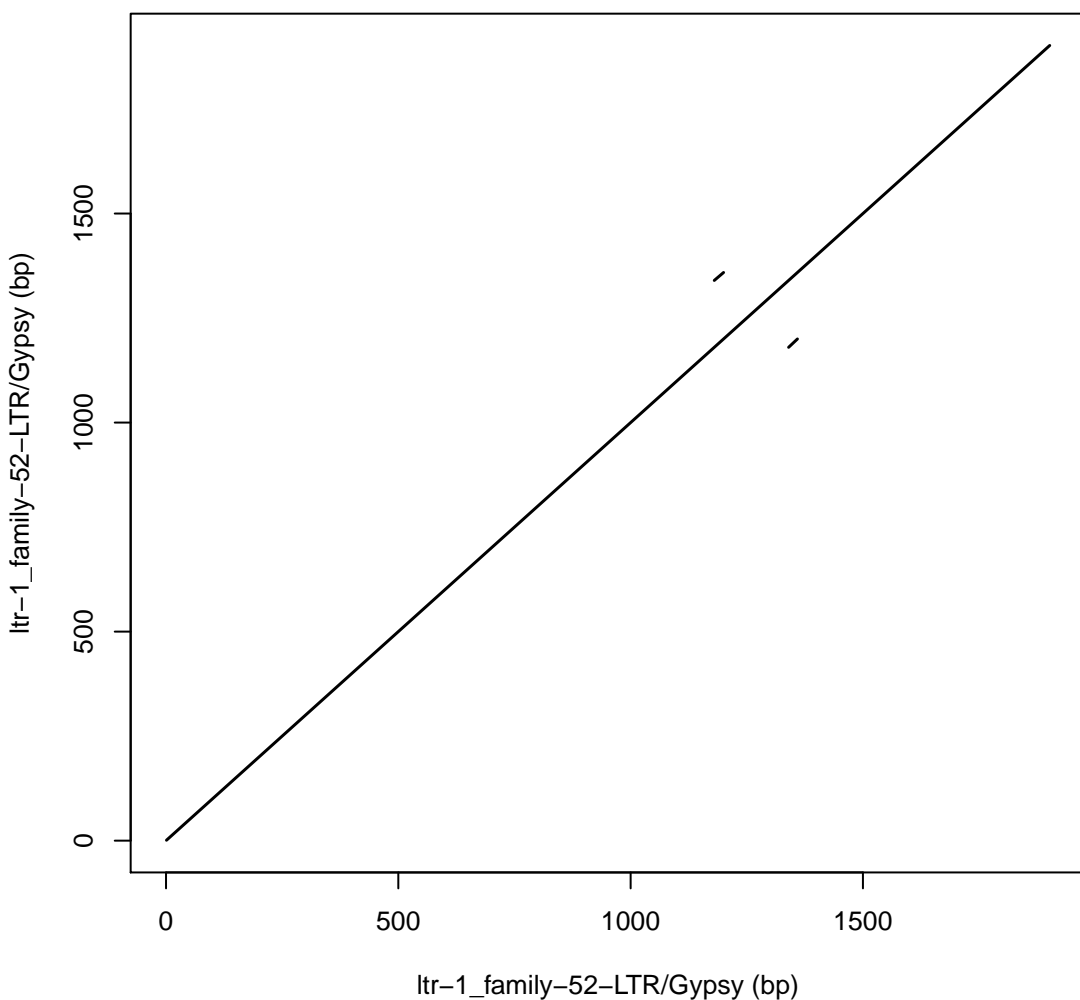
TE: ltr-1_family-52-LTR/Gypsy
consensus size: 1902bp; fragments: 334; full length: 0 (>=1711.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

