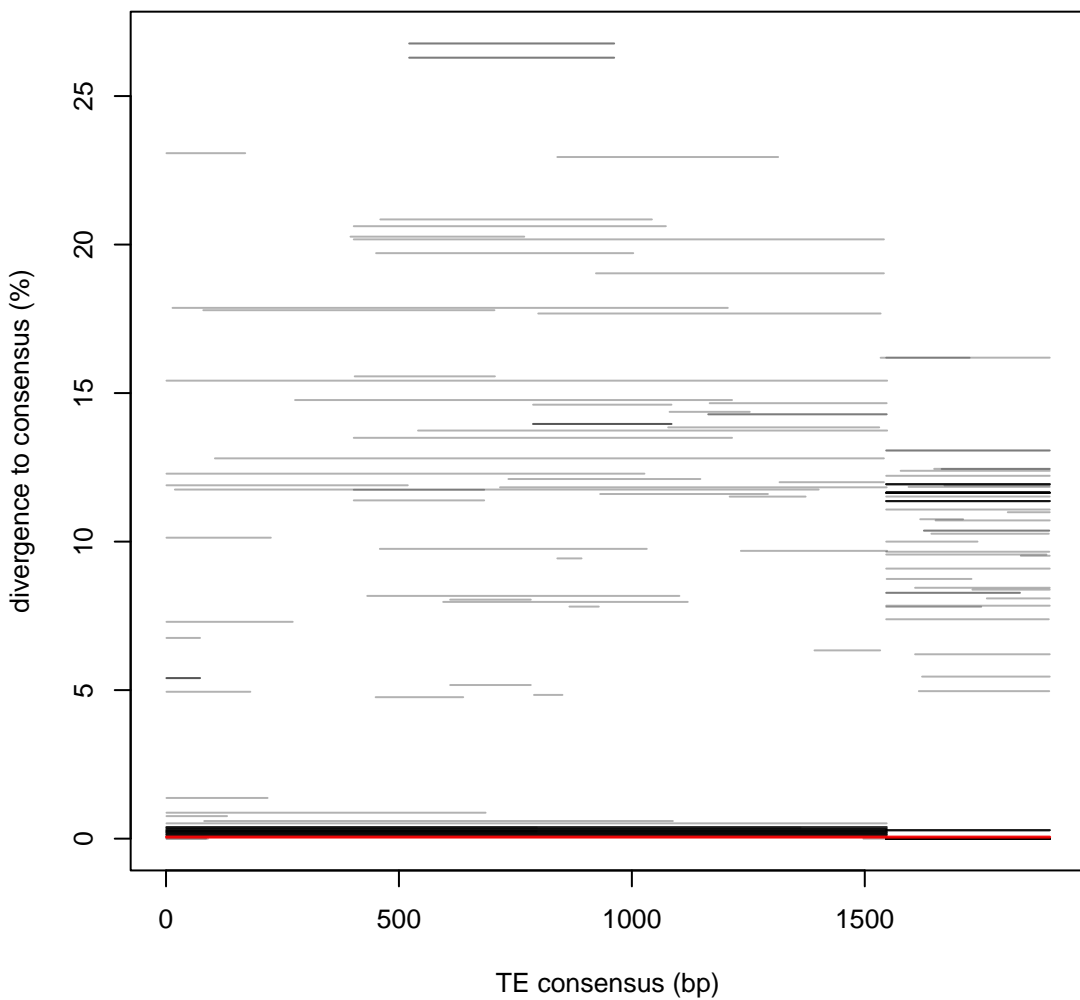
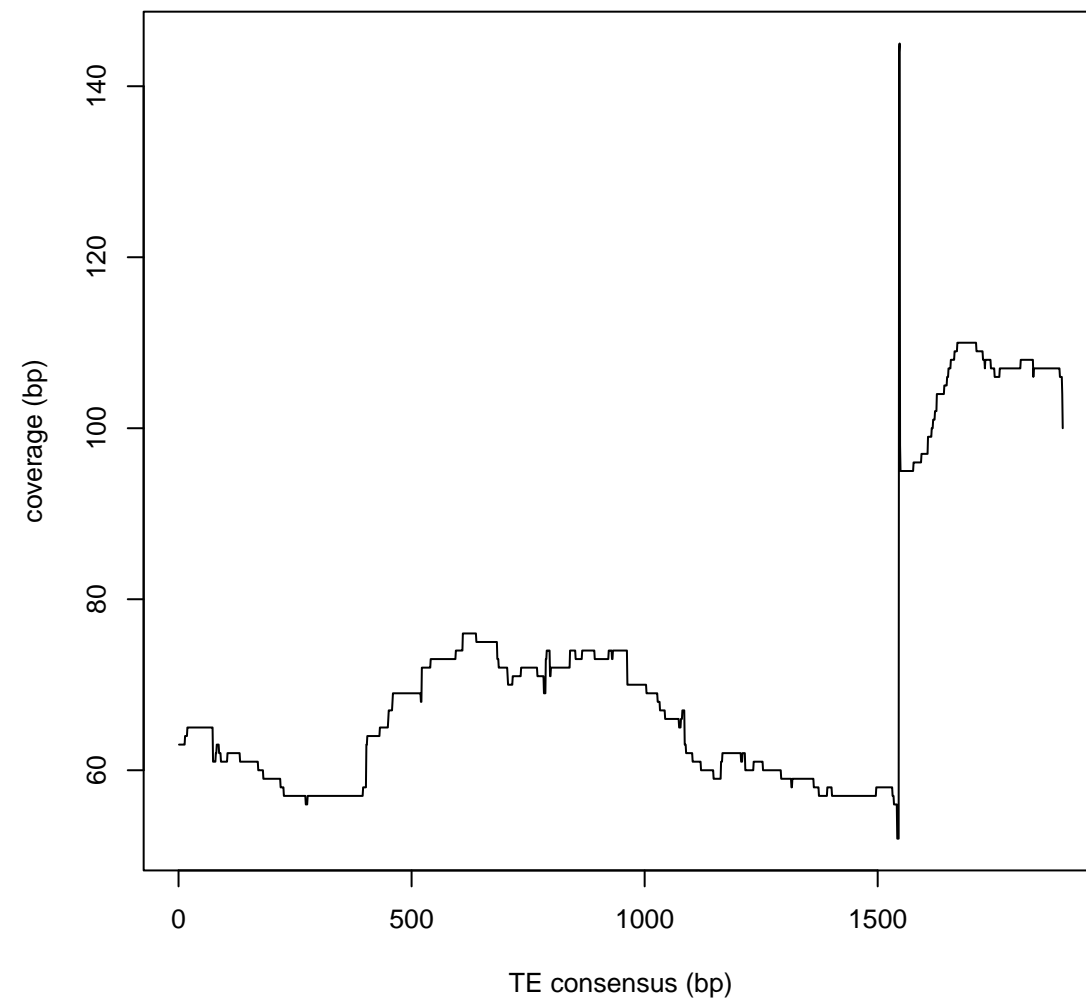


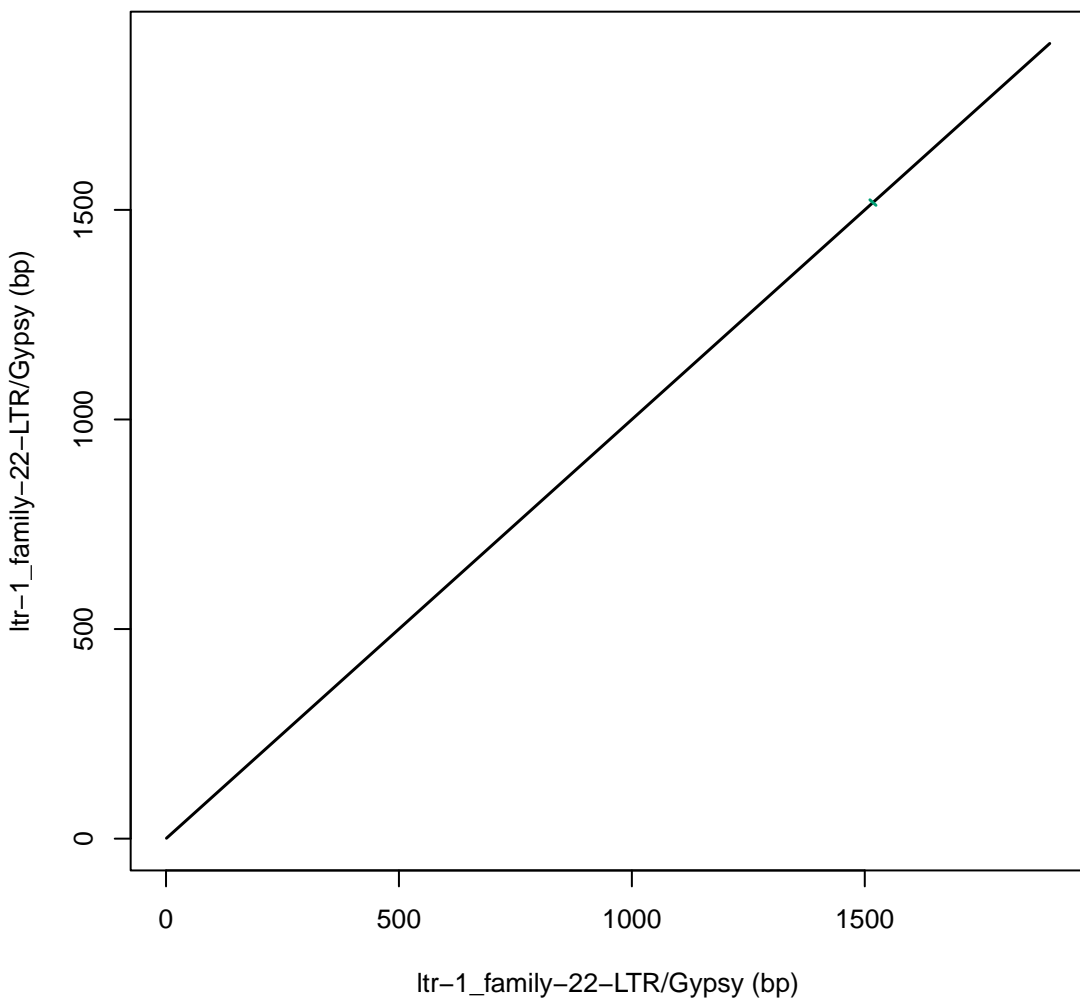
TE: ltr-1_family-22-LTR/Gypsy
consensus size: 1897bp; fragments: 225; full length: 1 (>=1707.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

