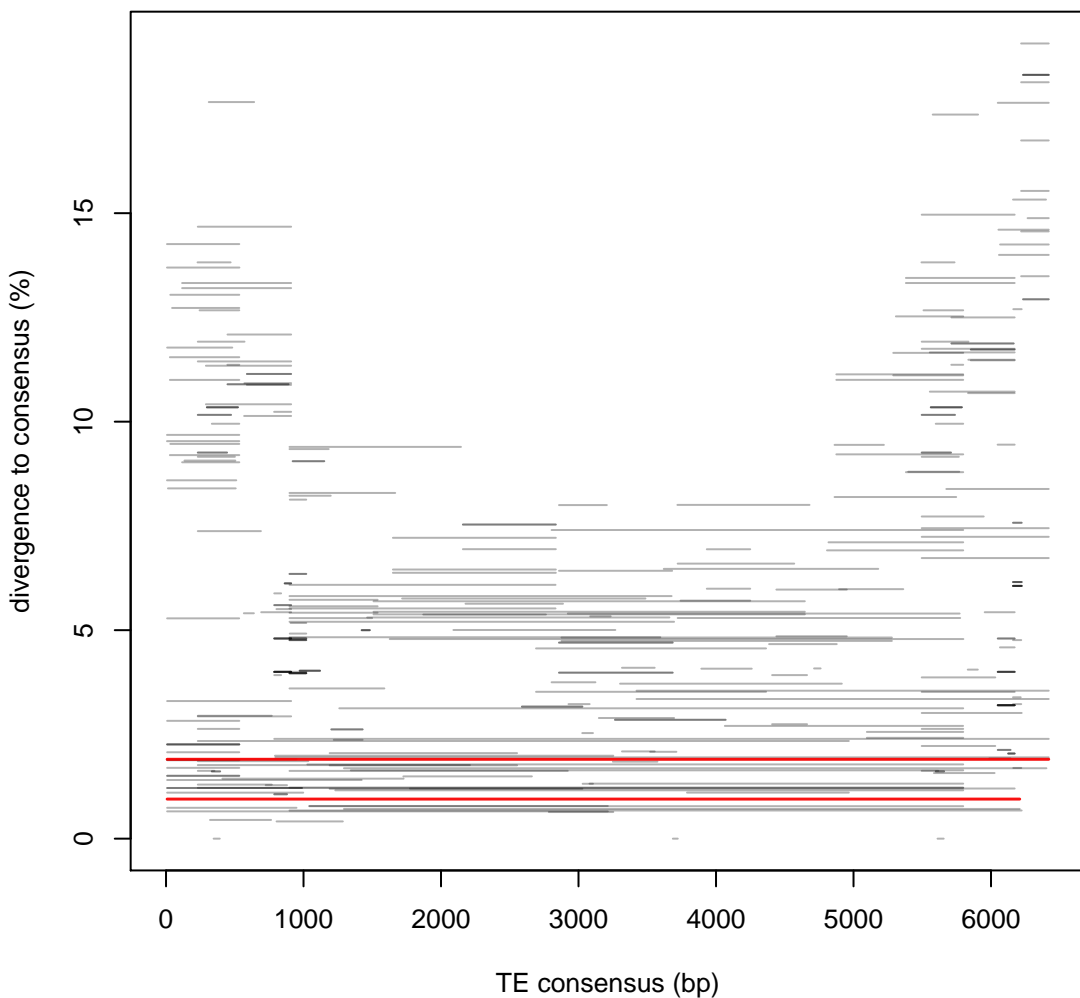
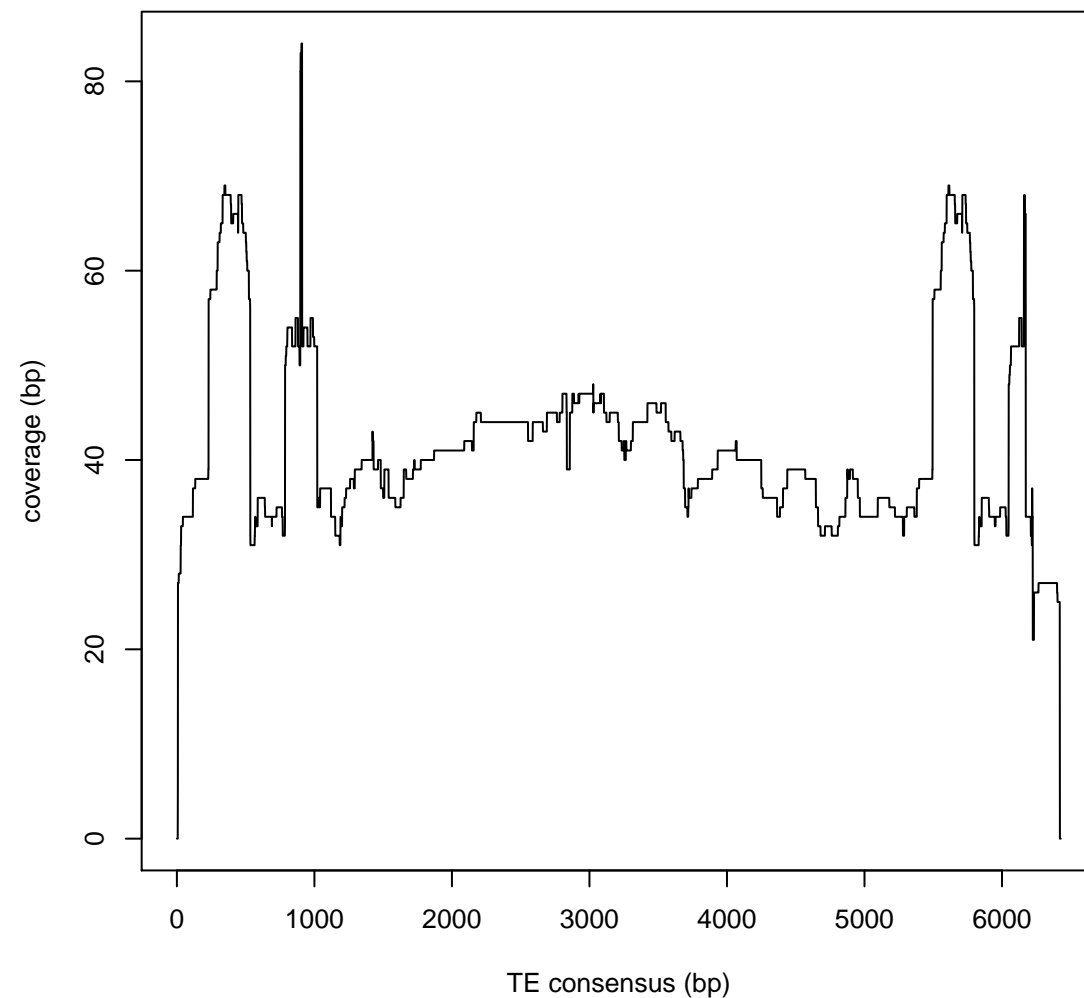


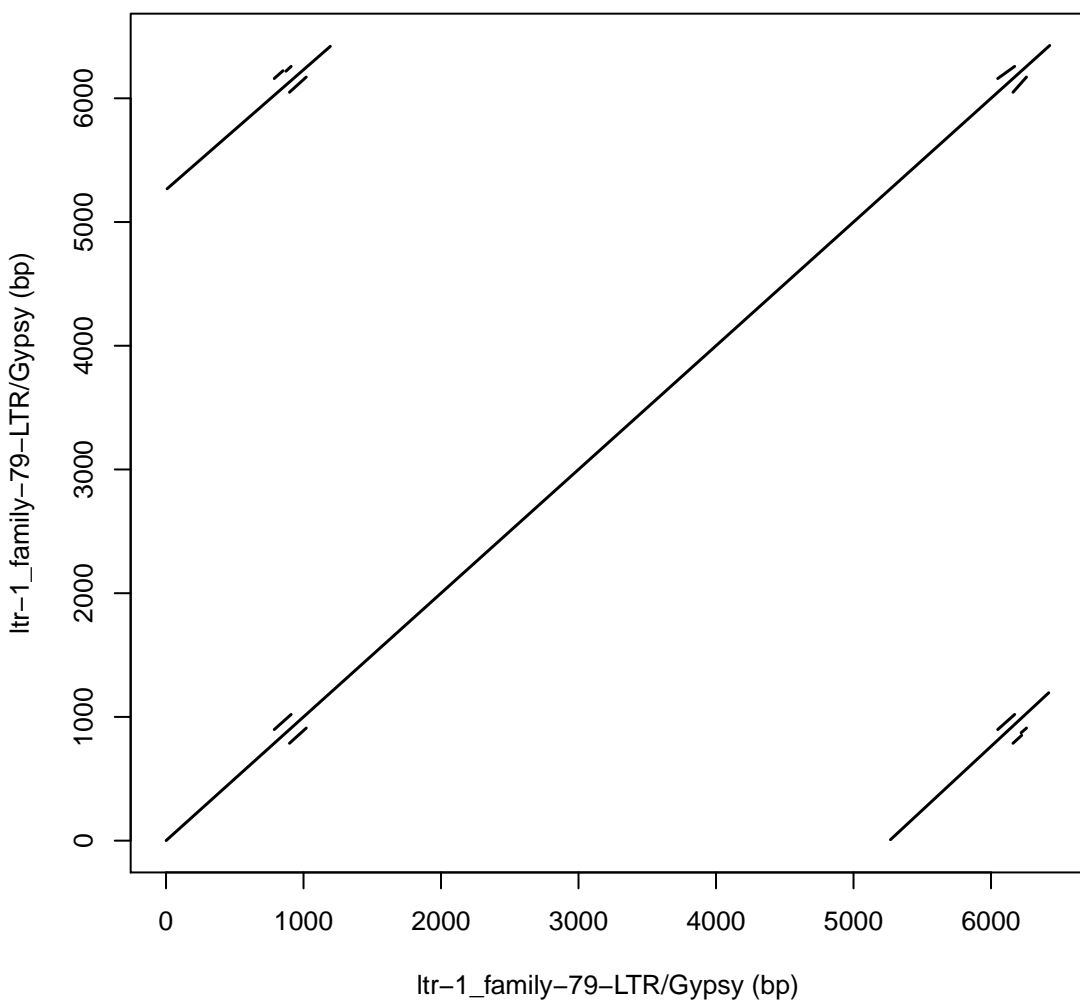
TE: ltr-1_family-79-LTR/Gypsy
consensus size: 6427bp; fragments: 340; full length: 2 (>=5784.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

