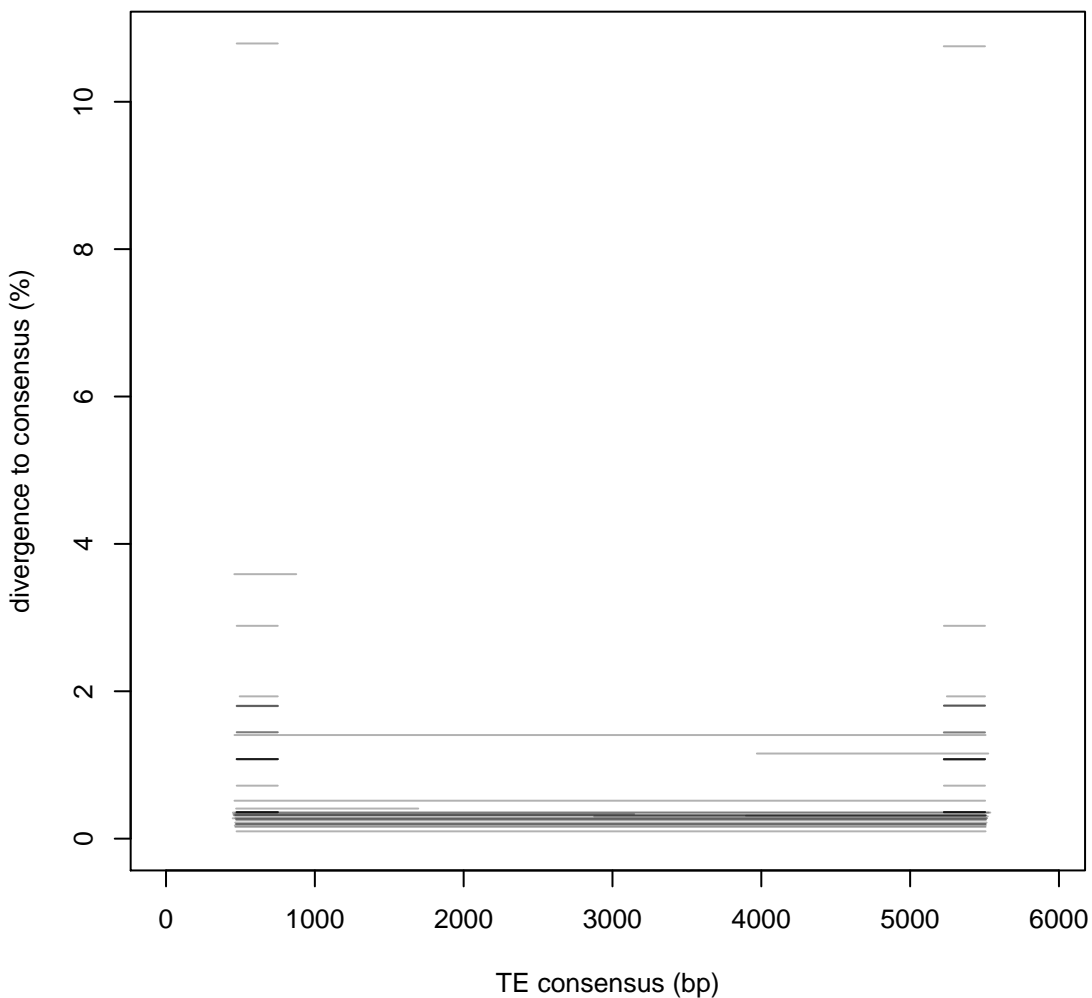
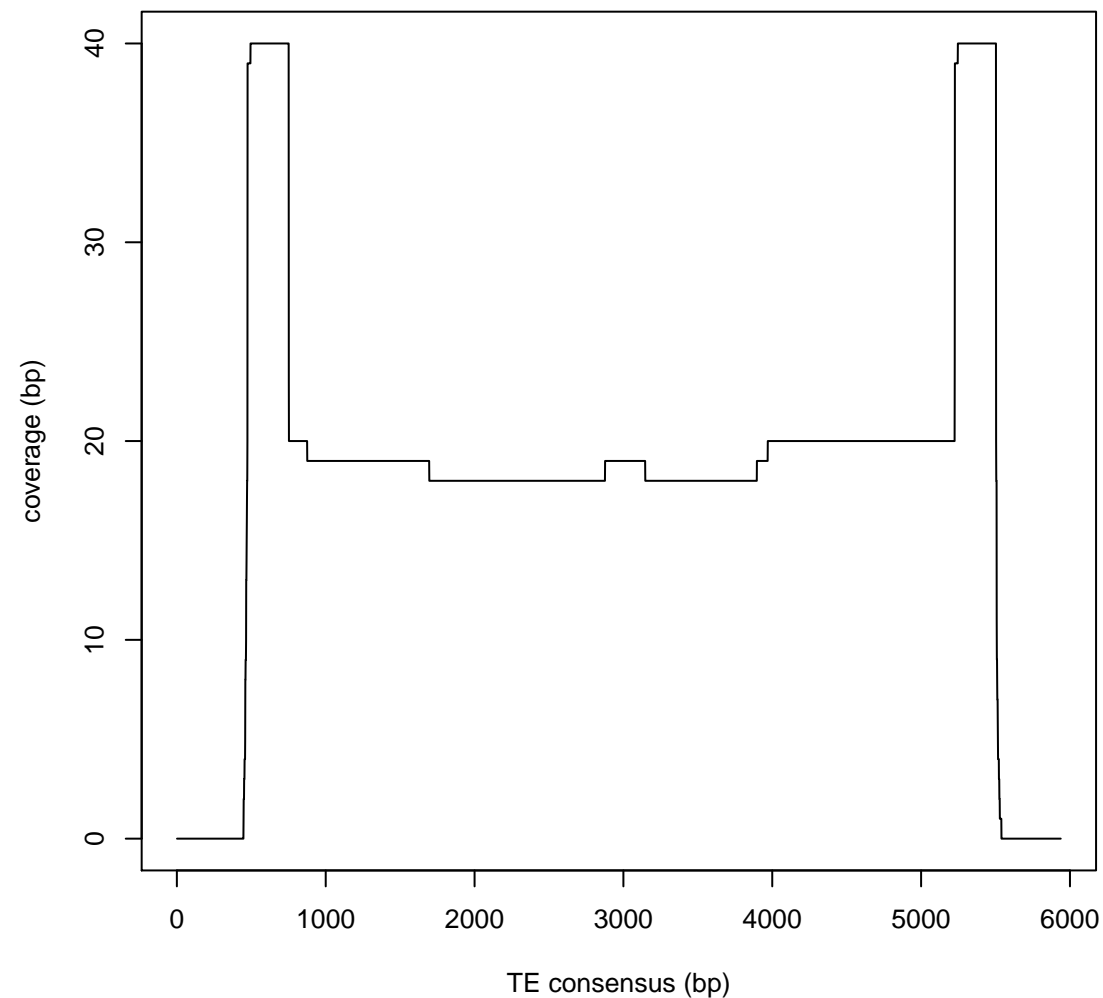


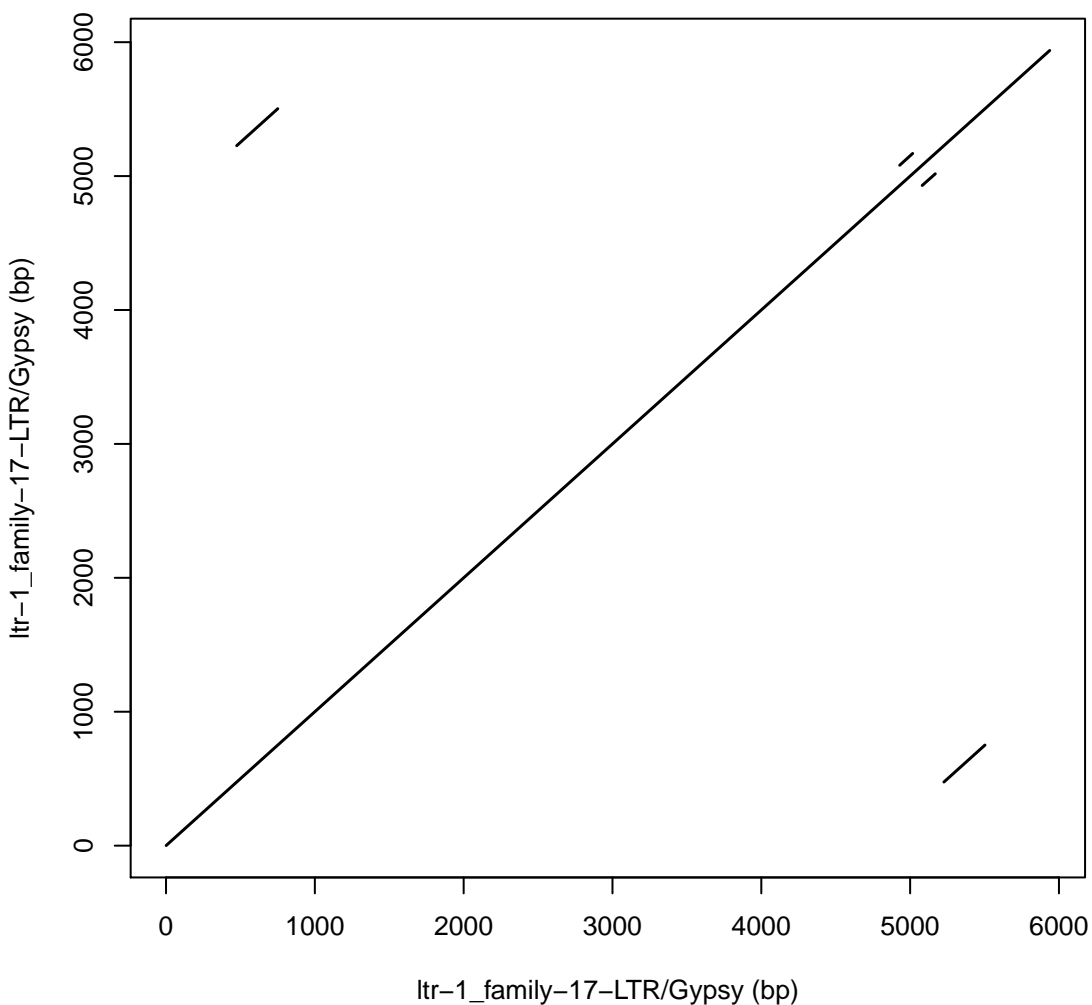
TE: ltr-1_family-17-LTR/Gypsy
consensus size: 5938bp; fragments: 63; full length: 0 (>=5344.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

