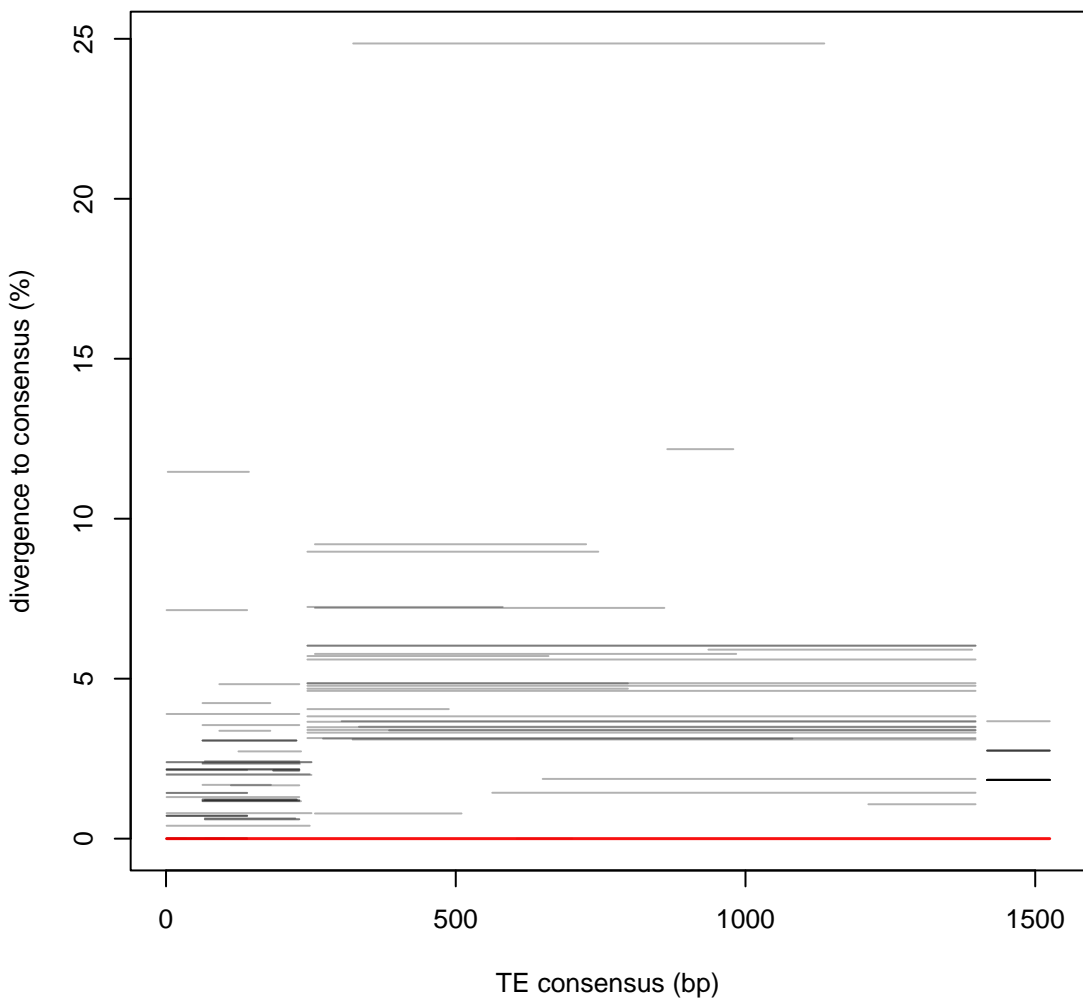
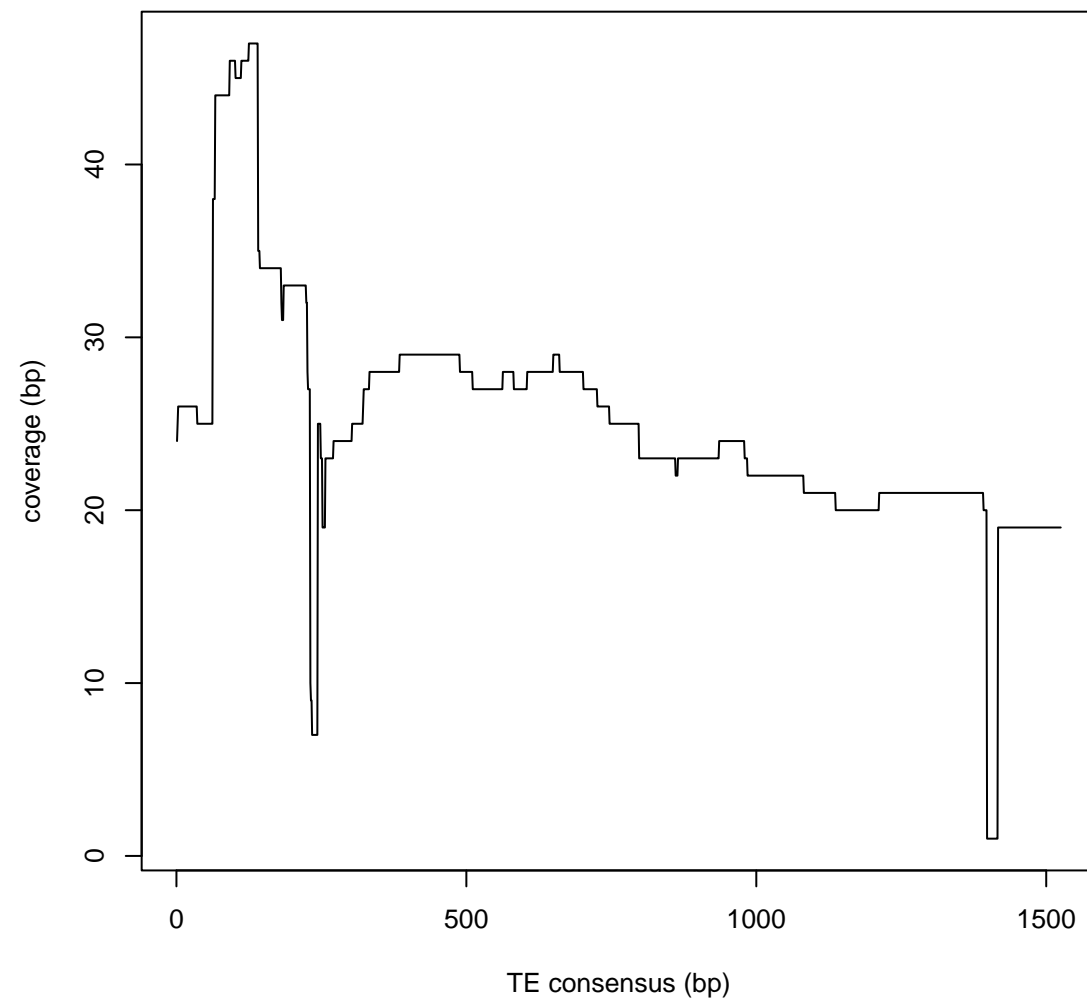


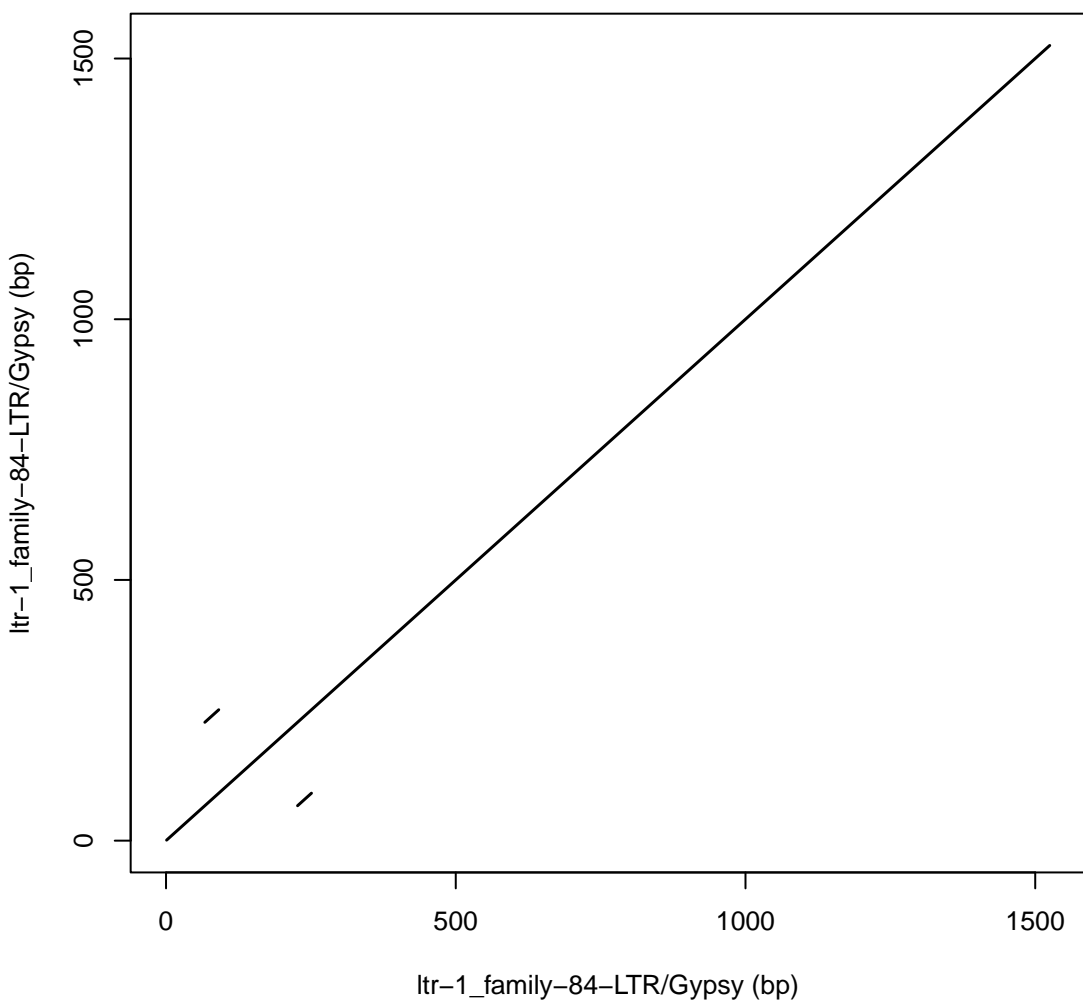
TE: ltr-1_family-84-LTR/Gypsy
consensus size: 1525bp; fragments: 103; full length: 1 (>=1372.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

