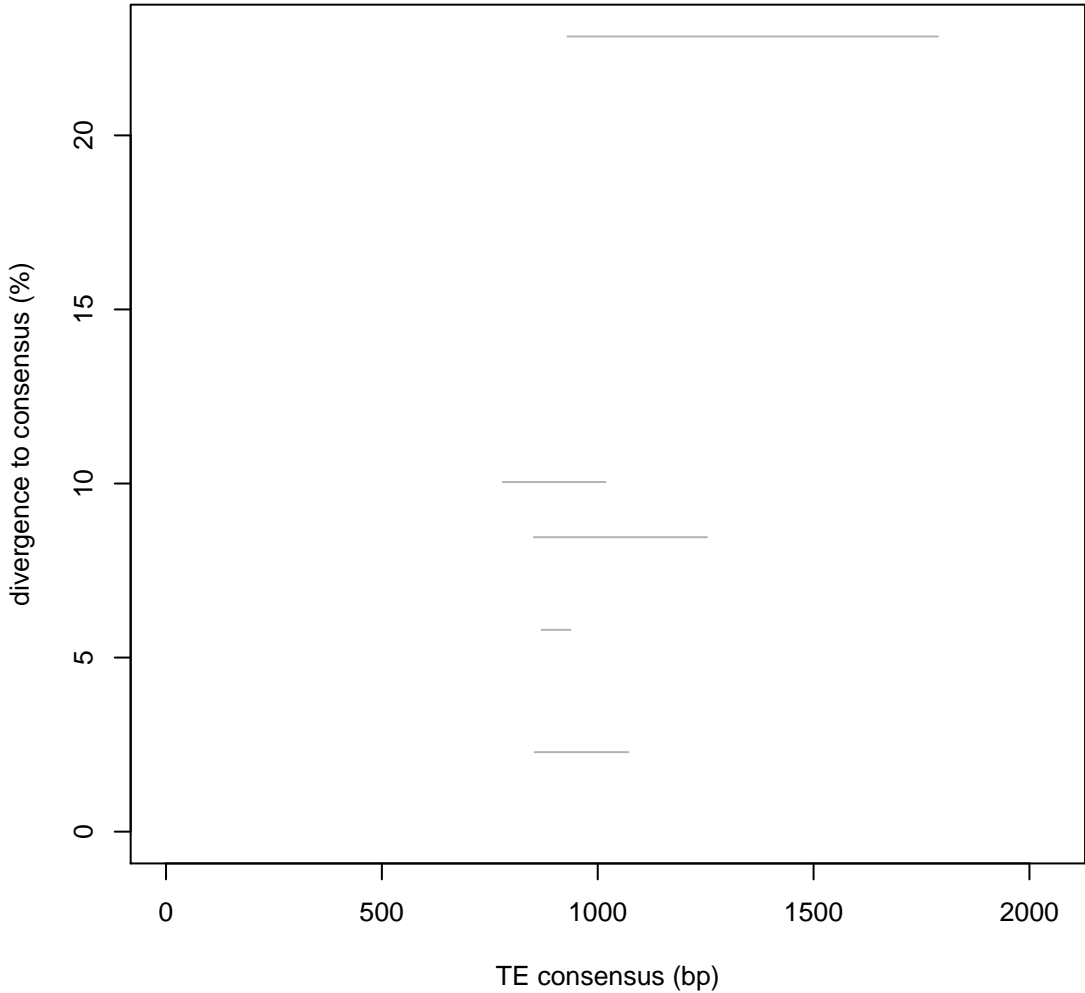
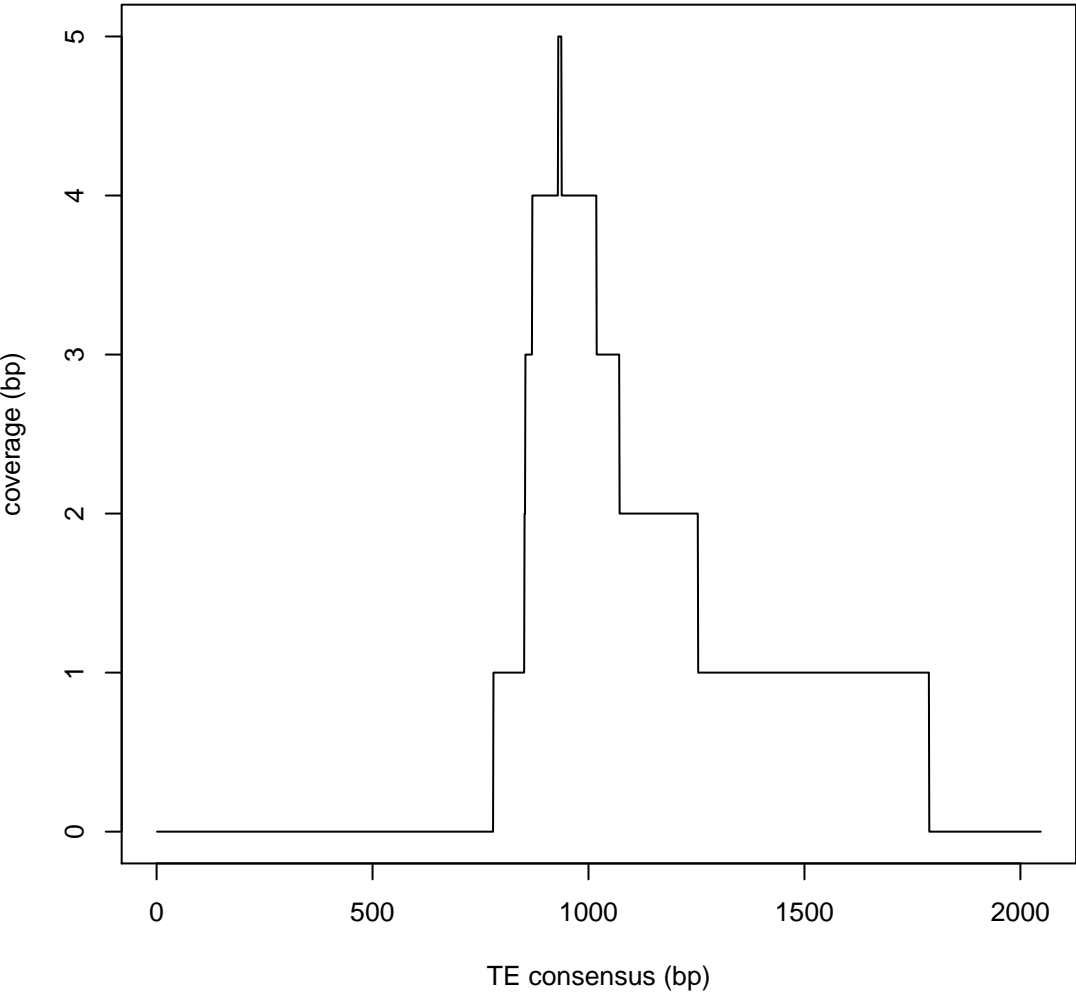


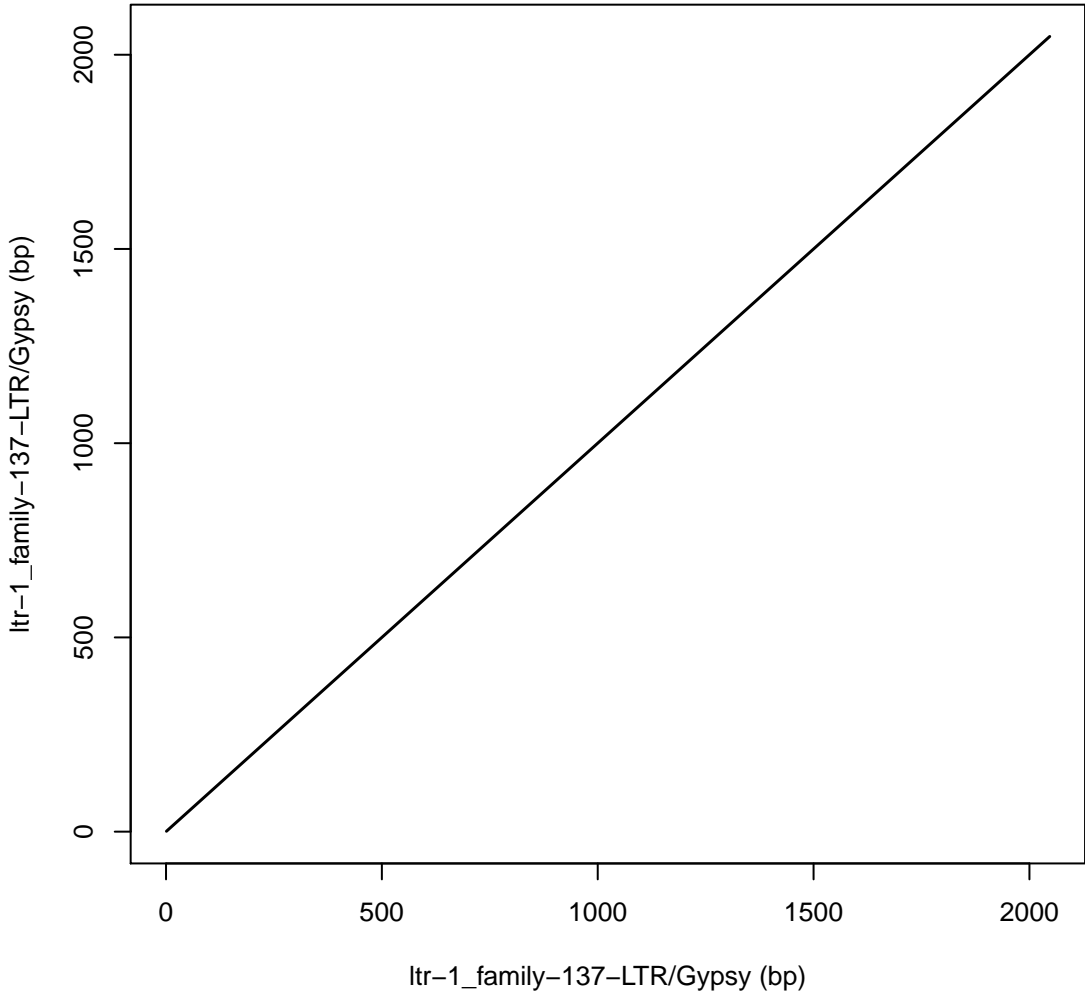
TE: ltr-1_family-137-LTR/Gypsy
consensus size: 2047bp; fragments: 5; full length: 0 (>=1842.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

