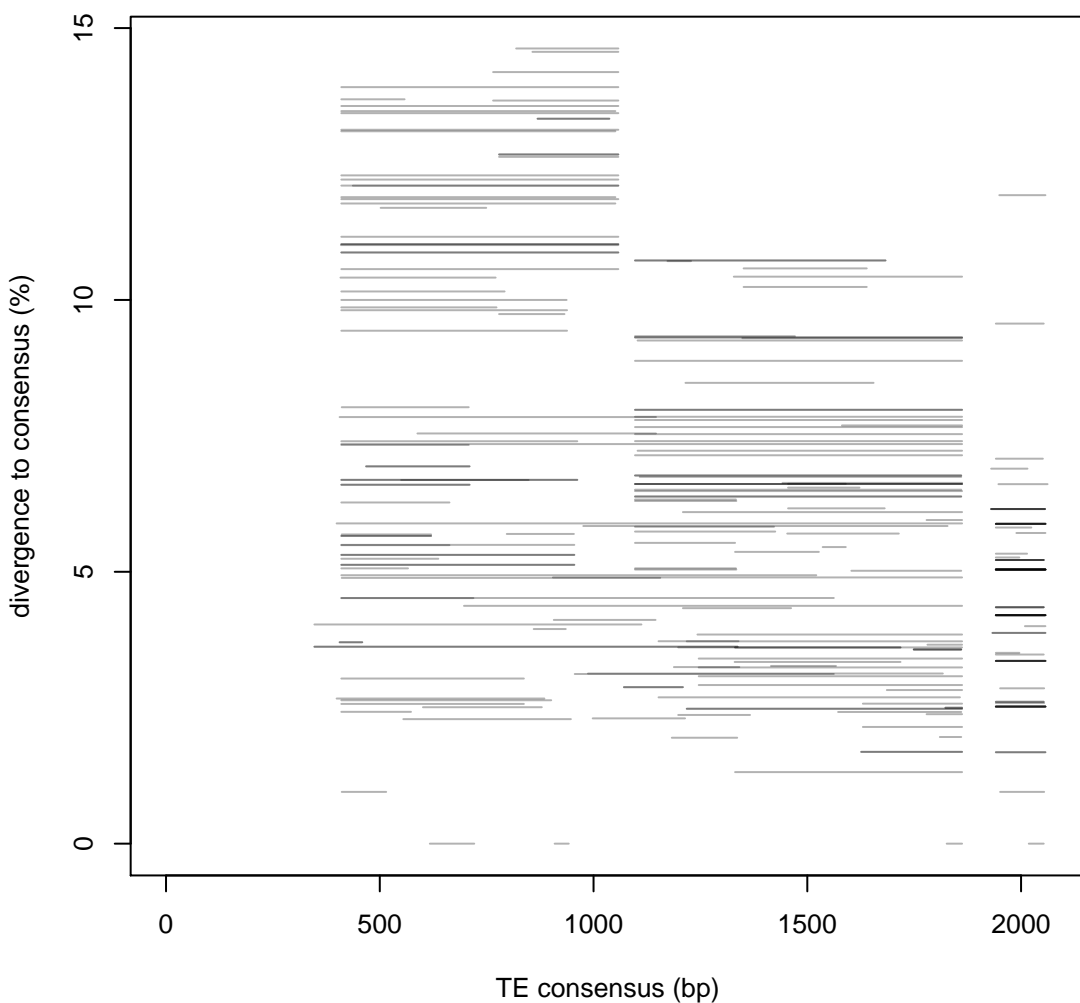
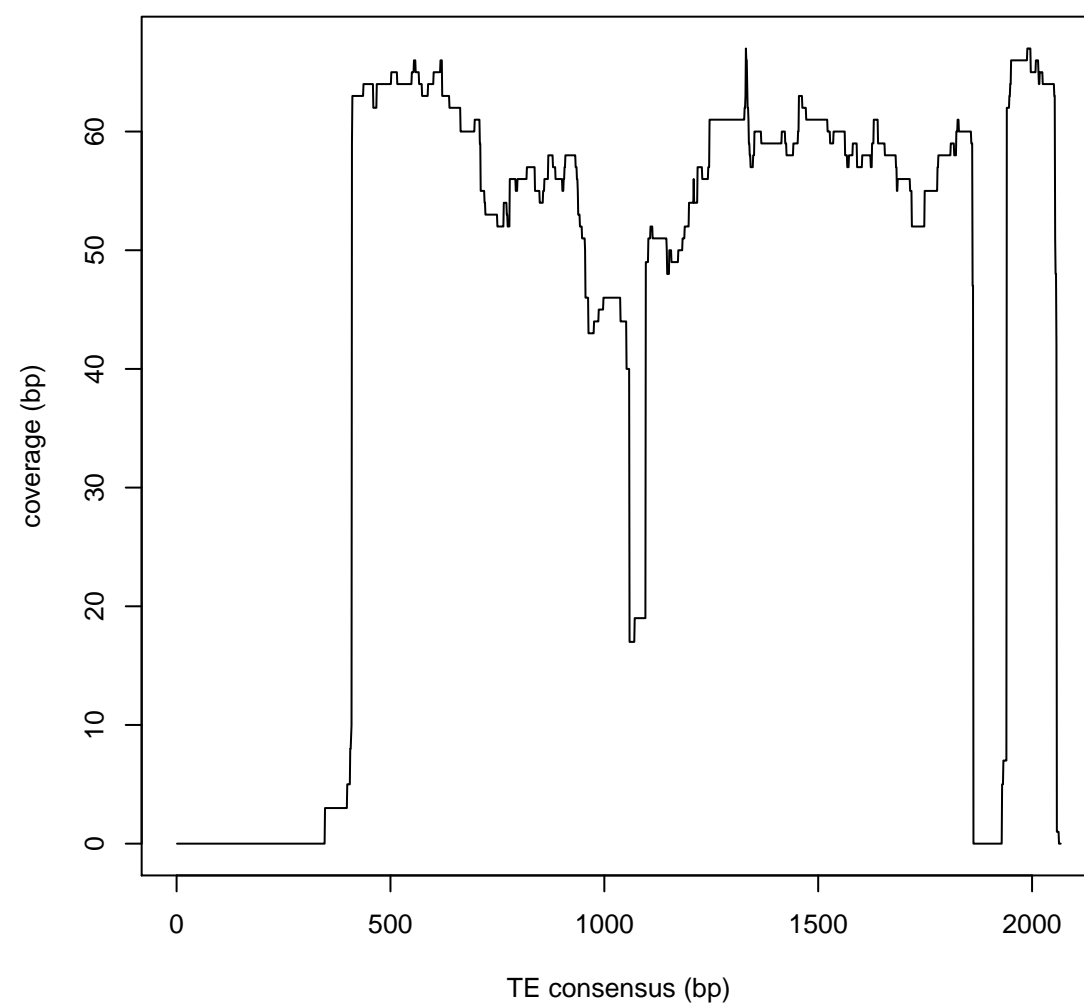


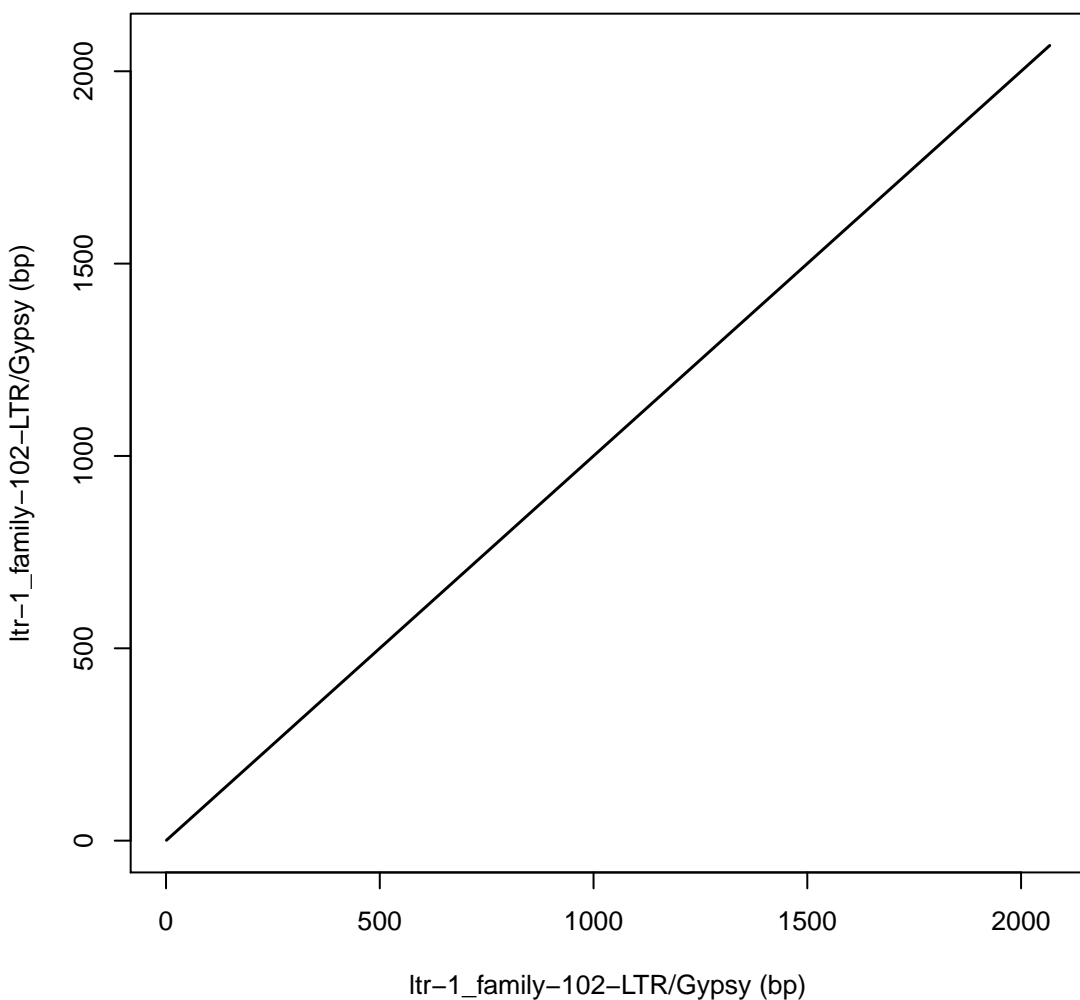
TE: ltr-1_family-102-LTR/Gypsy
consensus size: 2067bp; fragments: 248; full length: 0 (≥ 1860.3 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

