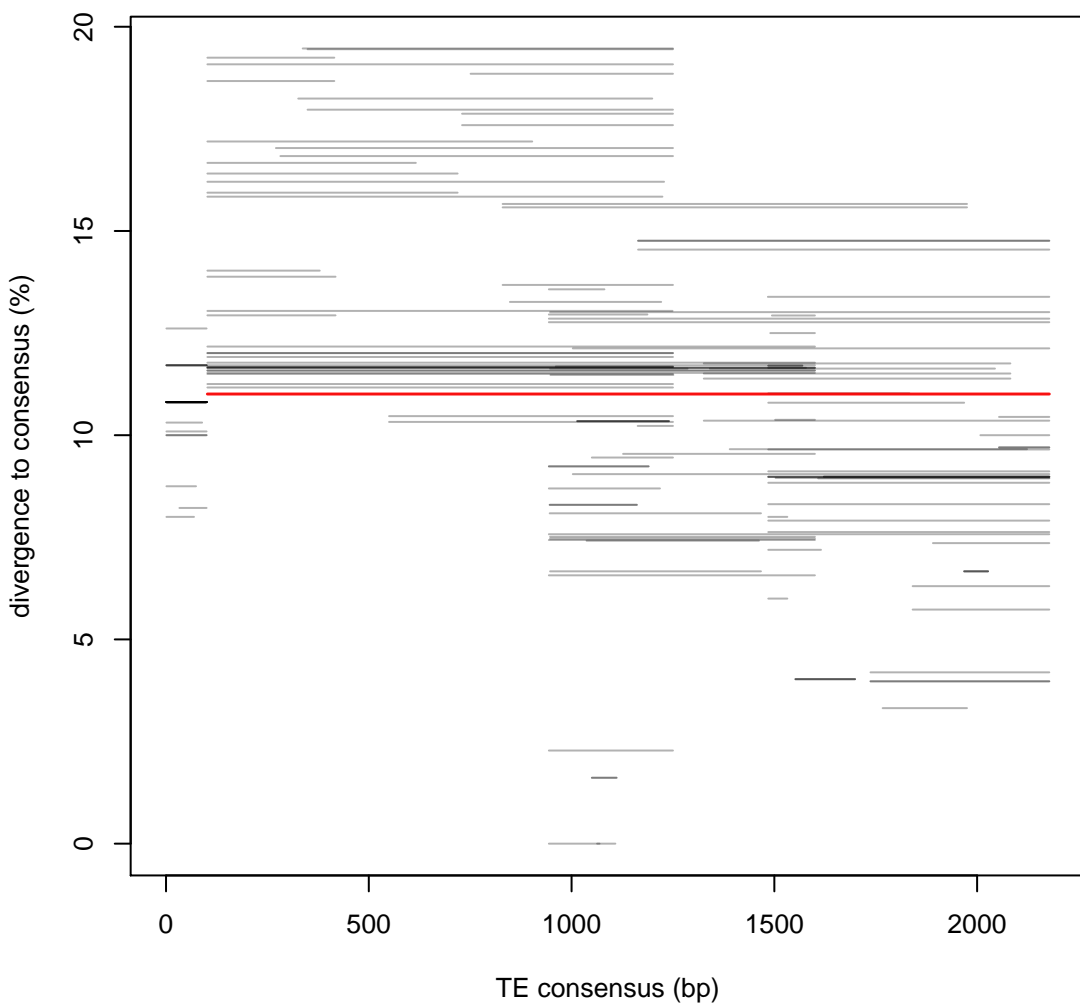
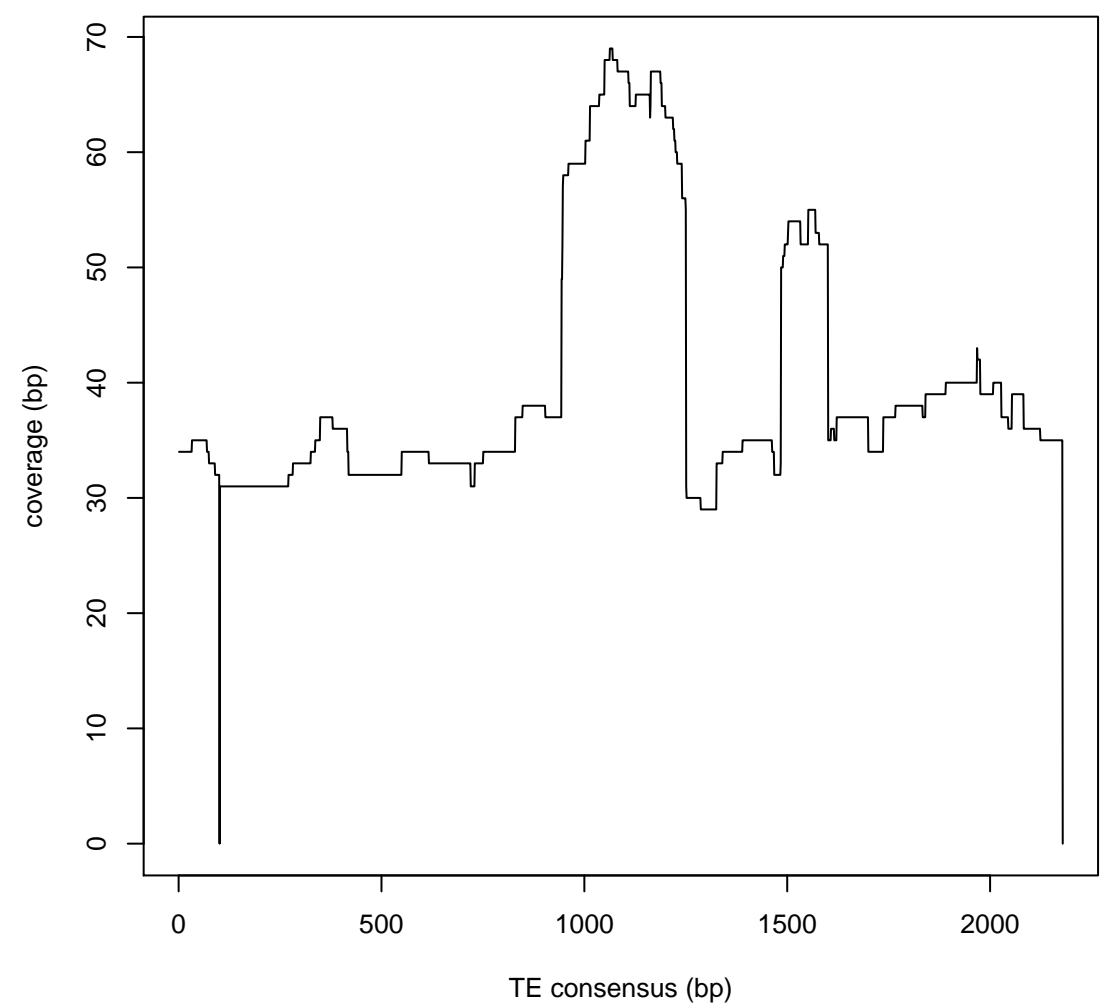


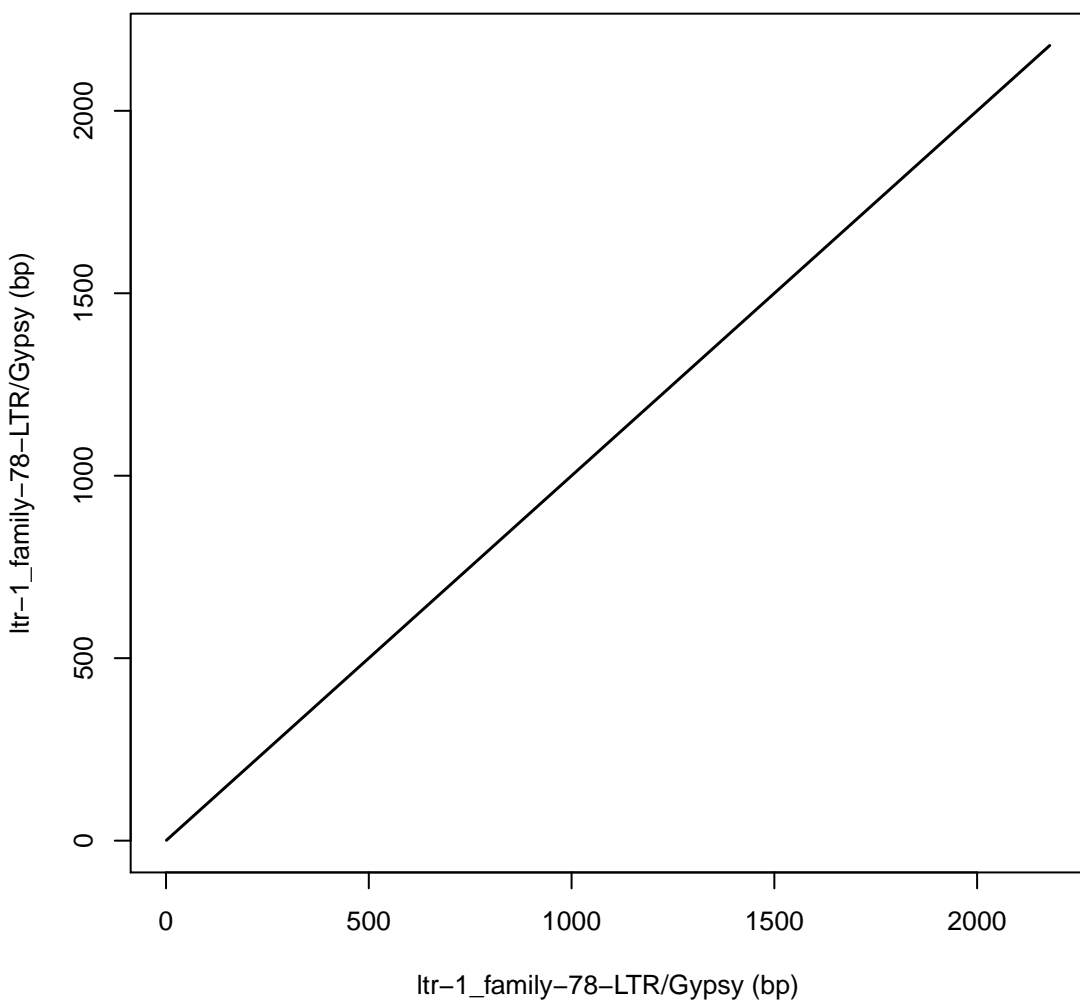
TE: ltr-1_family-78-LTR/Gypsy
consensus size: 2179bp; fragments: 166; full length: 1 (>=1961.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

