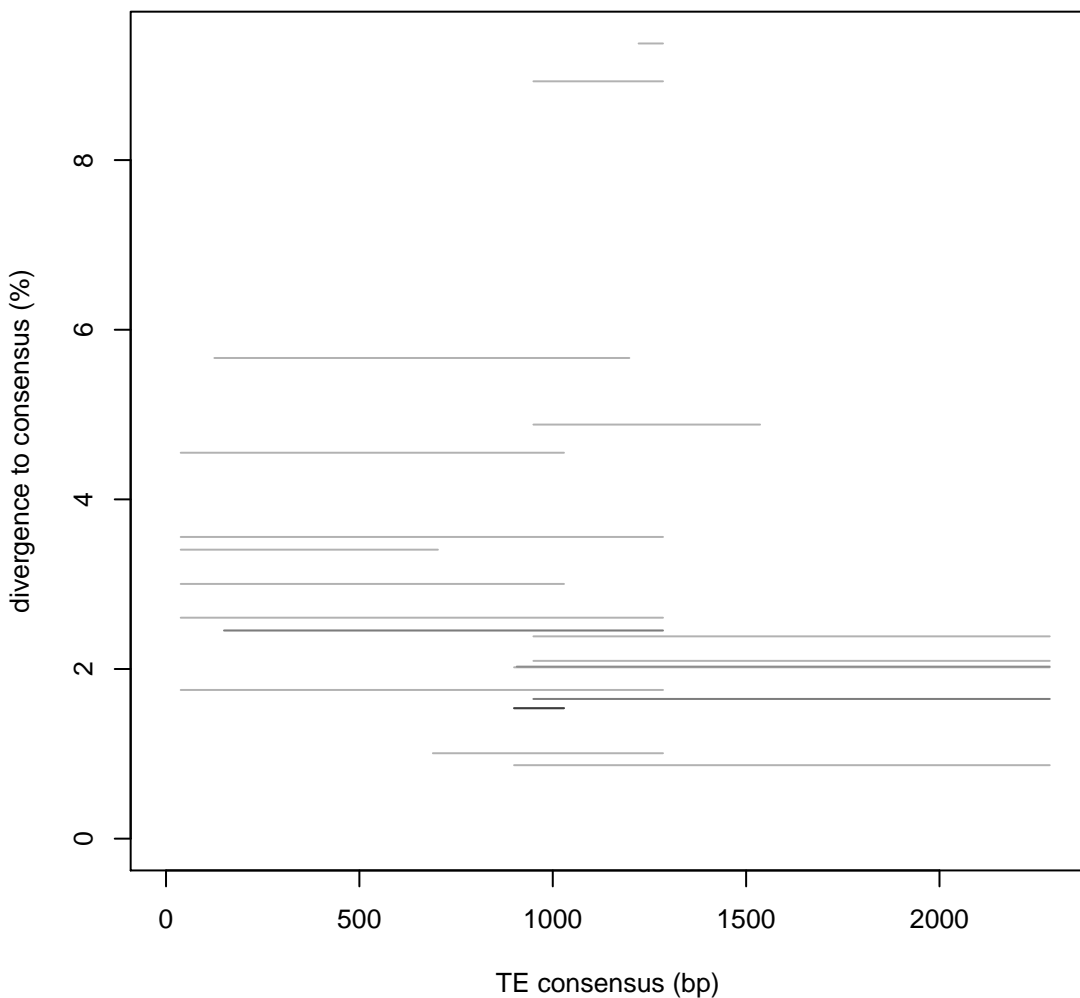
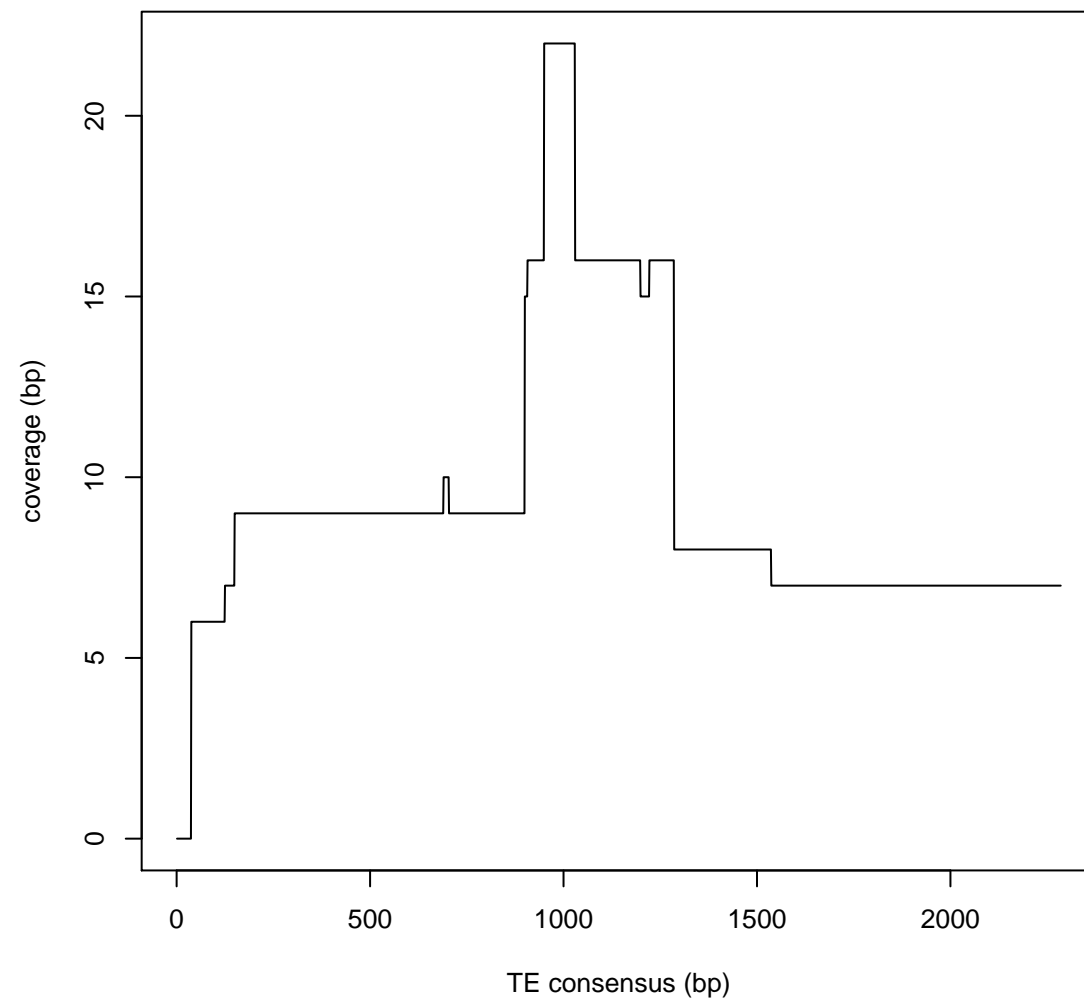


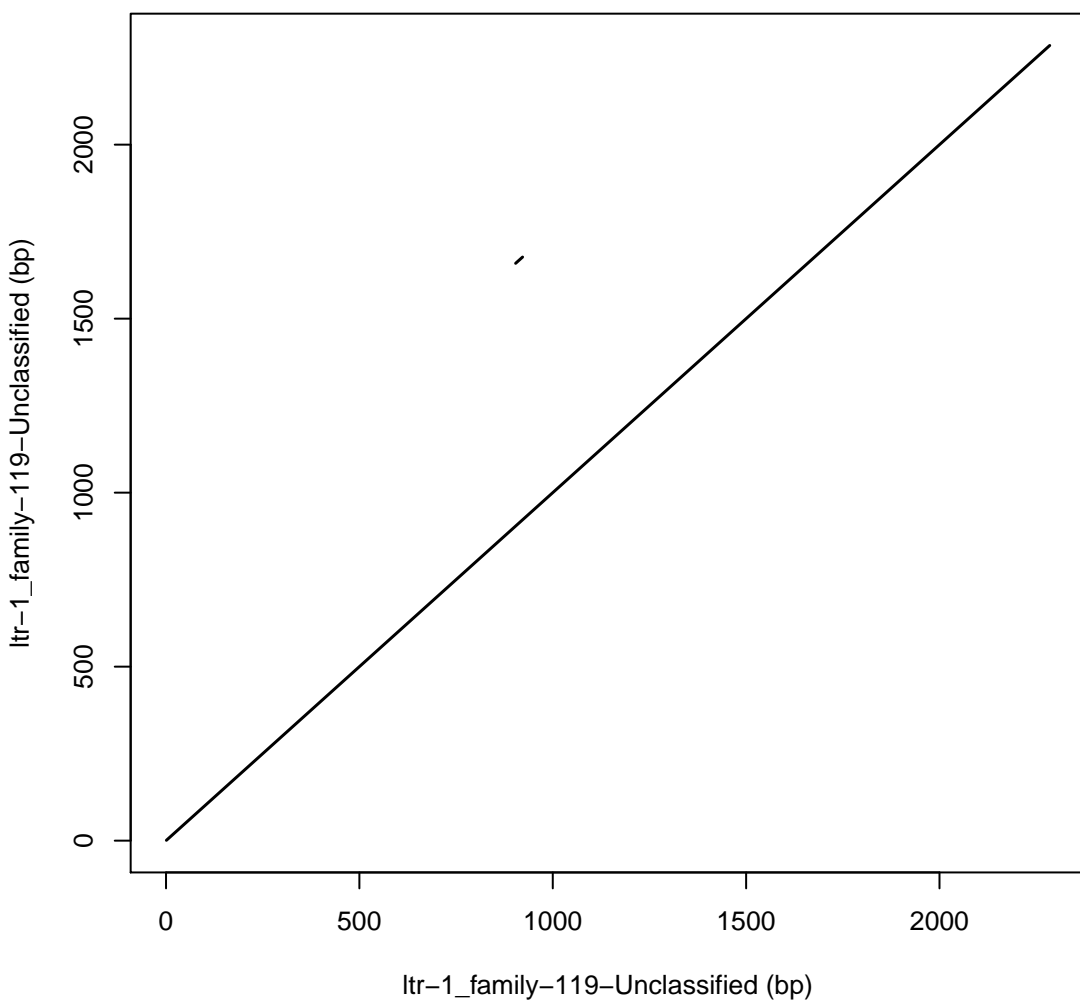
TE: ltr-1_family-119-Unclassified
consensus size: 2285bp; fragments: 24; full length: 0 (>=2056.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

