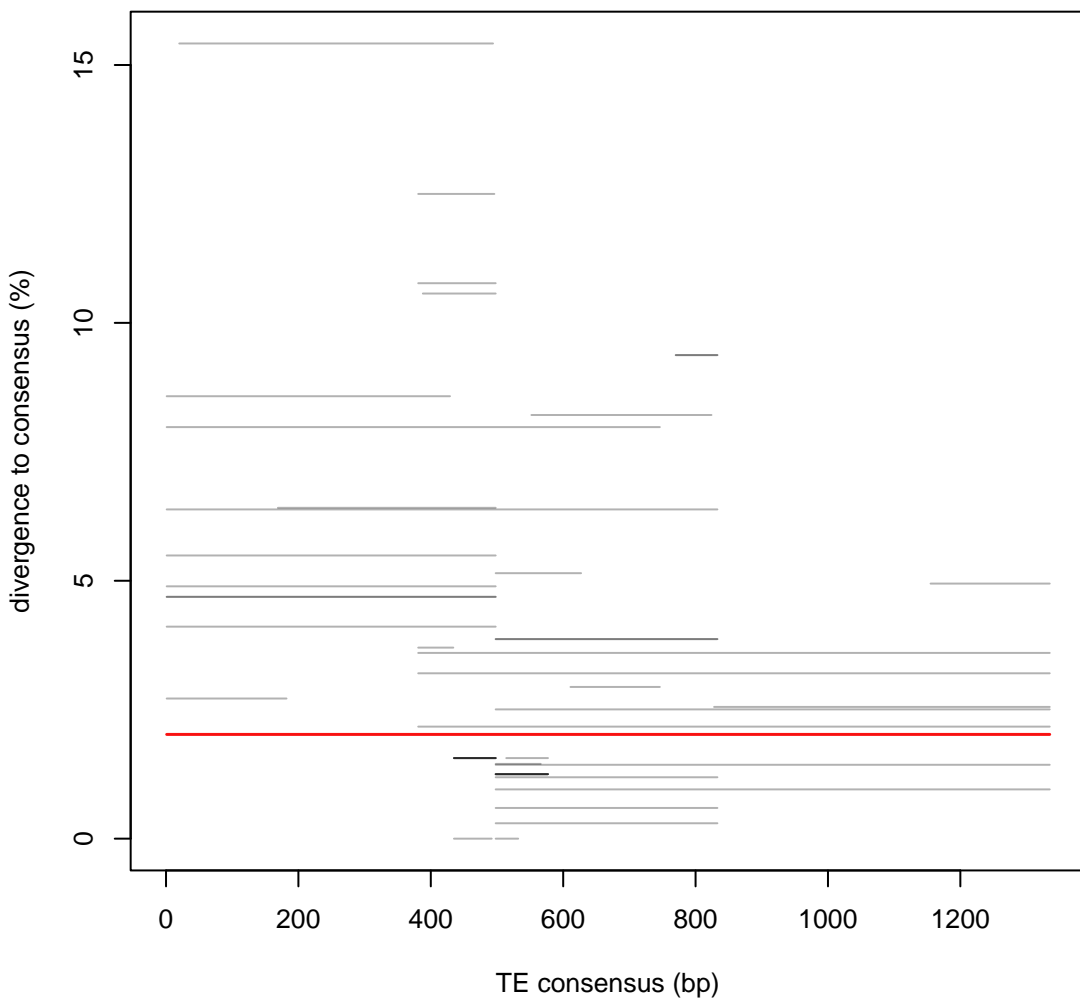
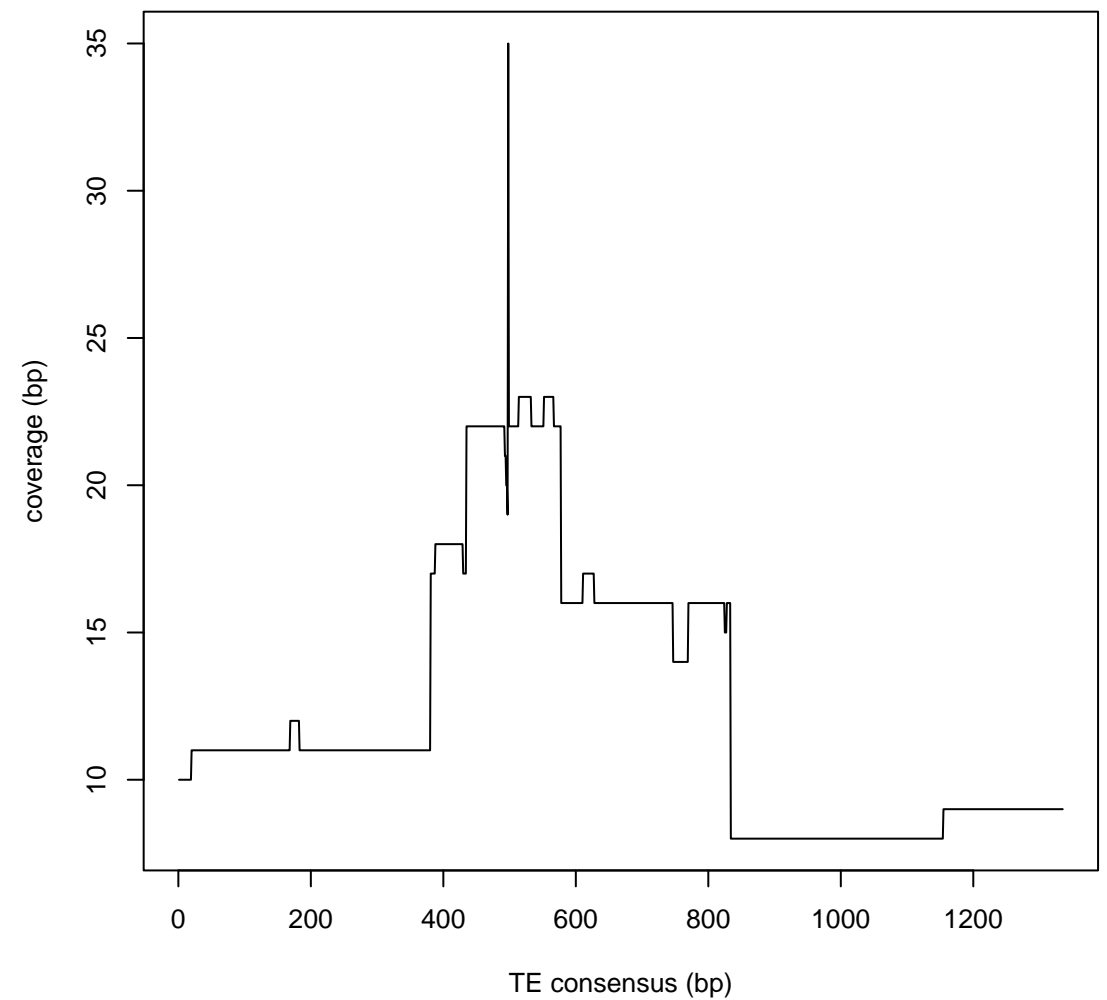


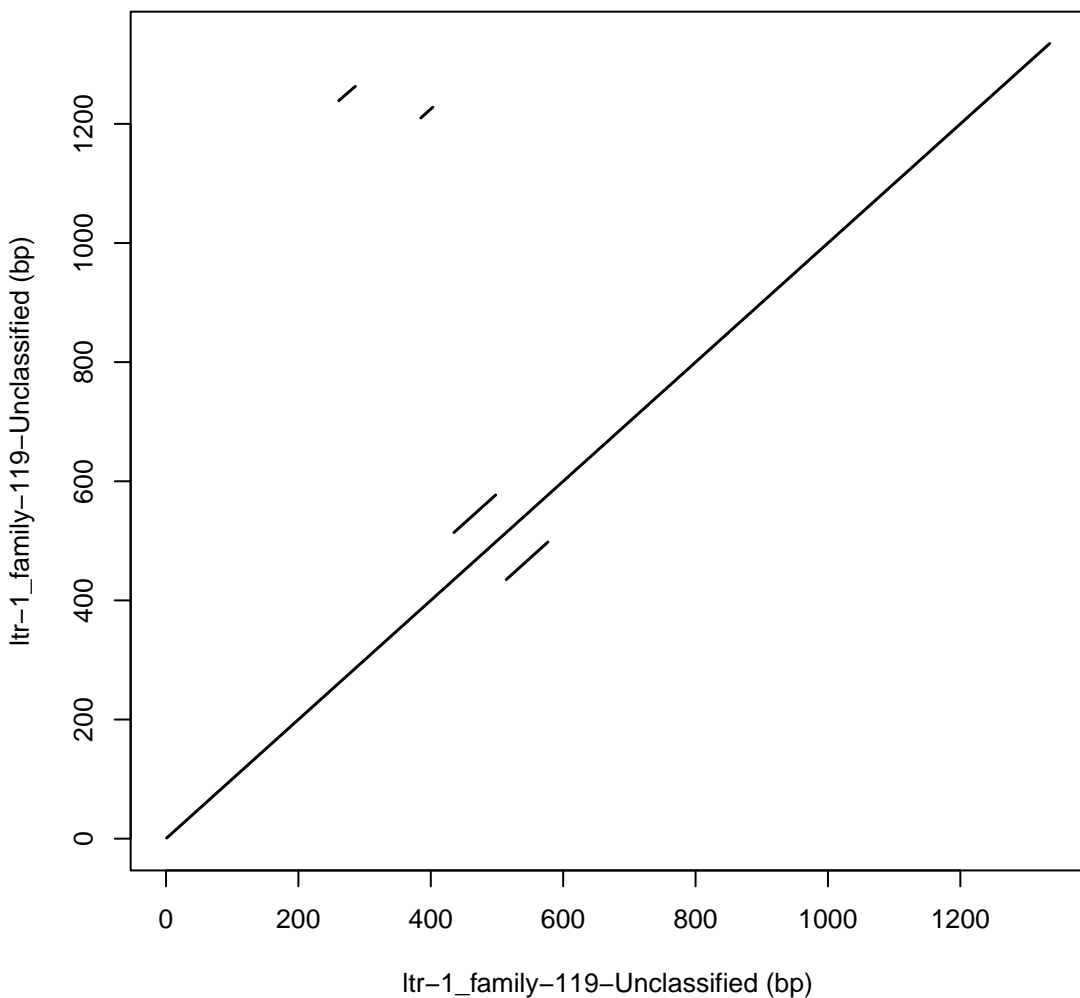
TE: ltr-1\_family-119-Unclassified  
consensus size: 1335bp; fragments: 48; full length: 1 ( $\geq 1201.5$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

