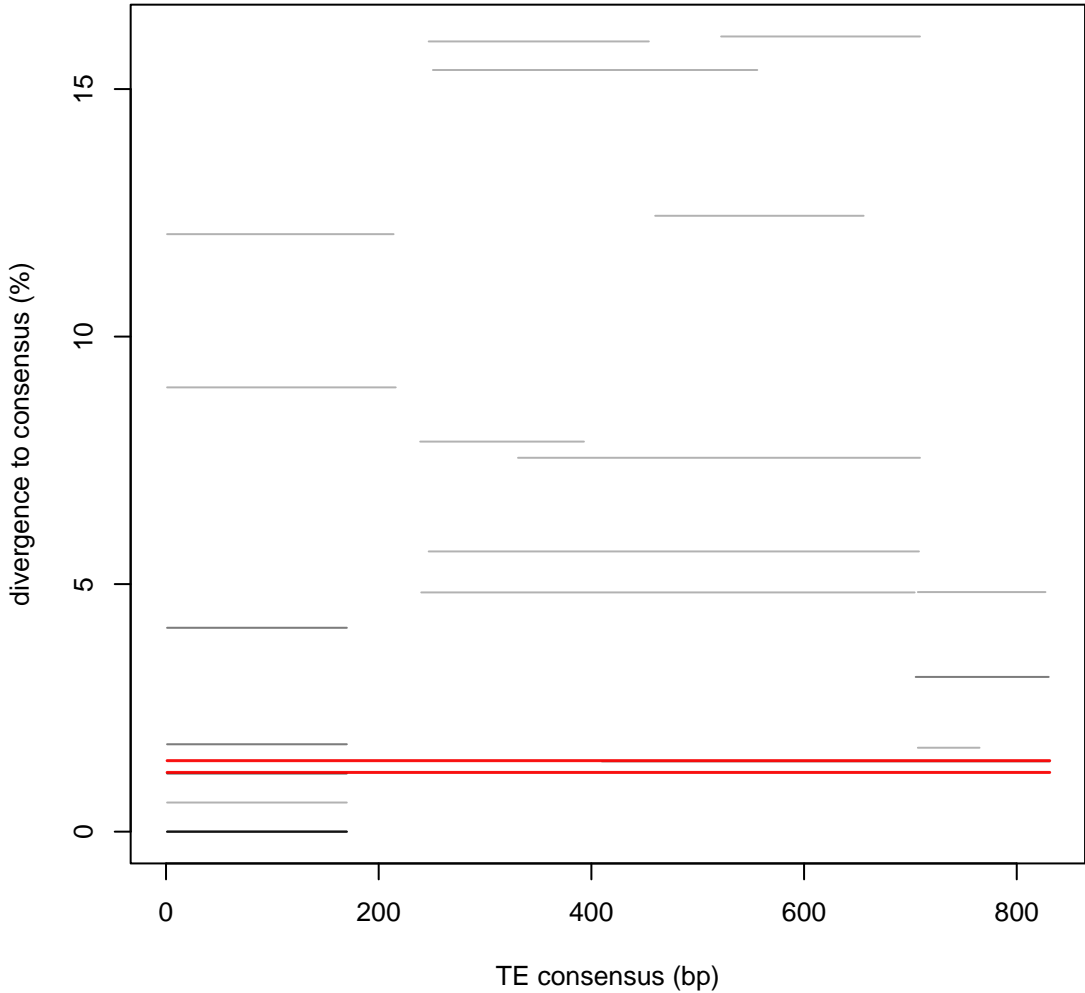
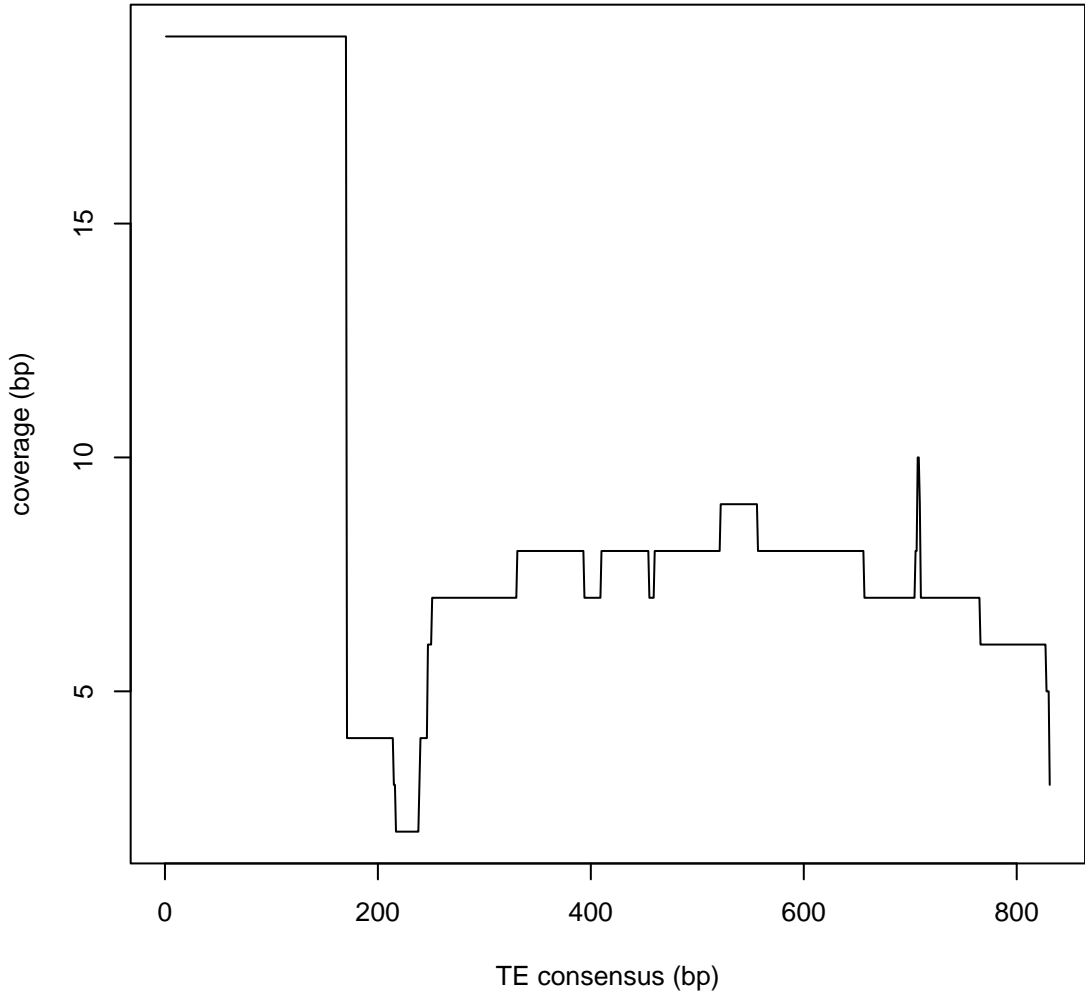


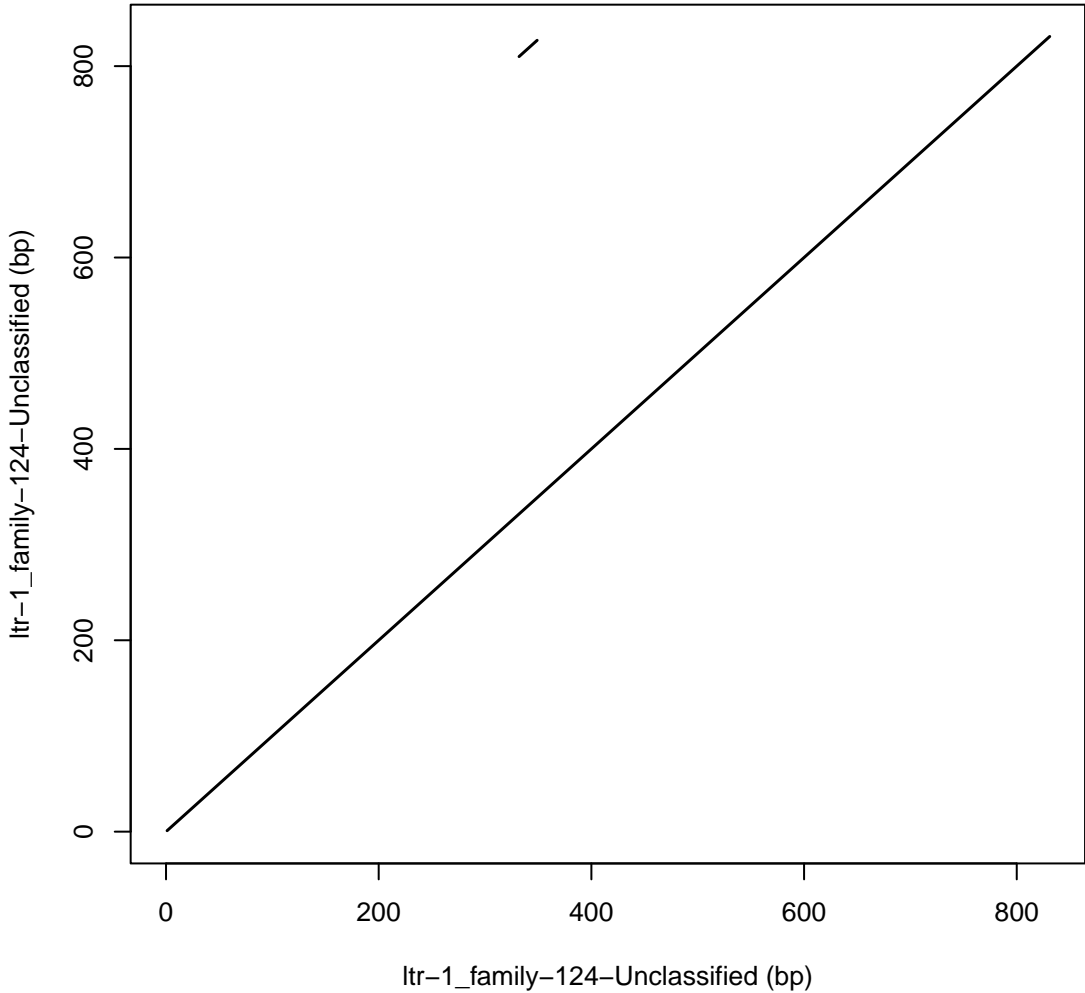
TE: ltr-1\_family-124-Unclassified  
consensus size: 831bp; fragments: 32; full length: 2 ( $\geq 747.9$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

