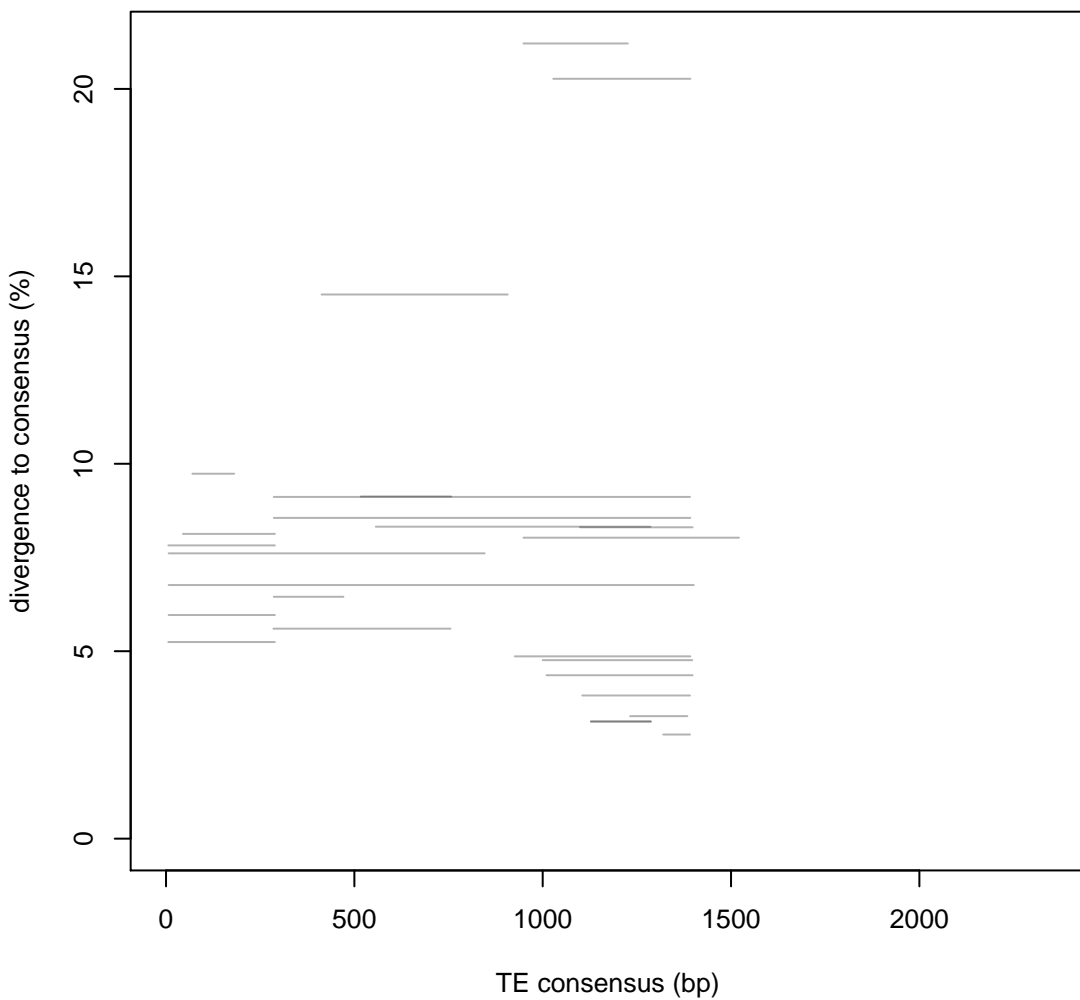
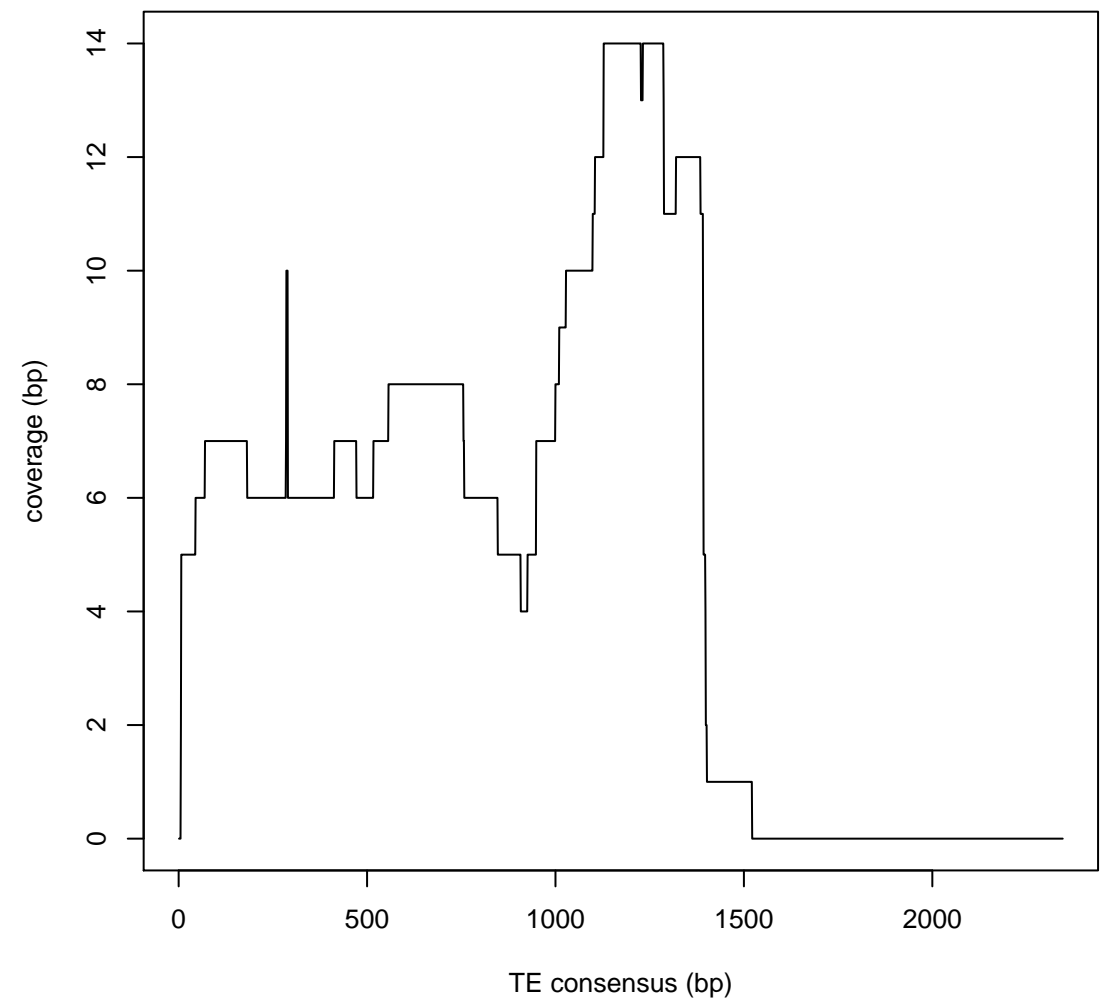


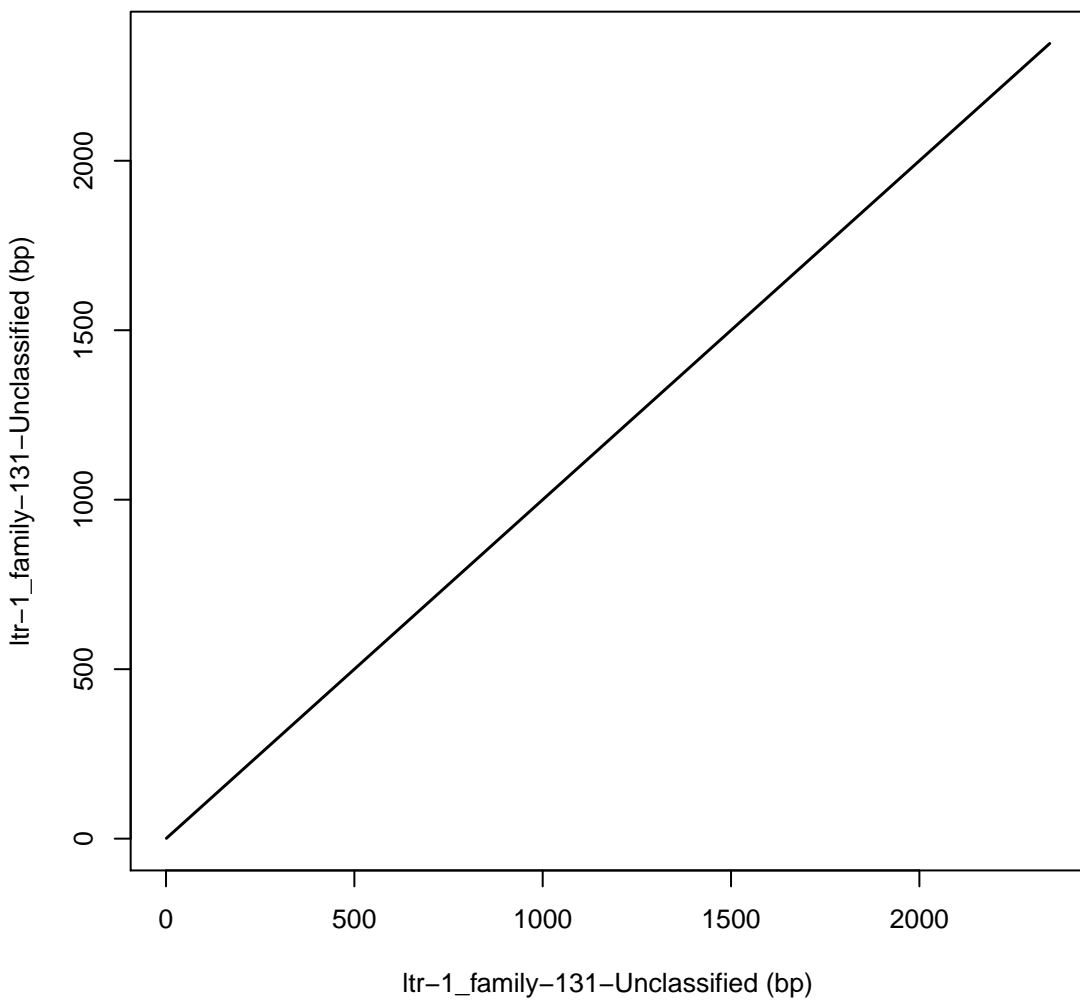
TE: ltr-1\_family-131-Unclassified  
consensus size: 2346bp; fragments: 26; full length: 0 ( $\geq 2111.4$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

