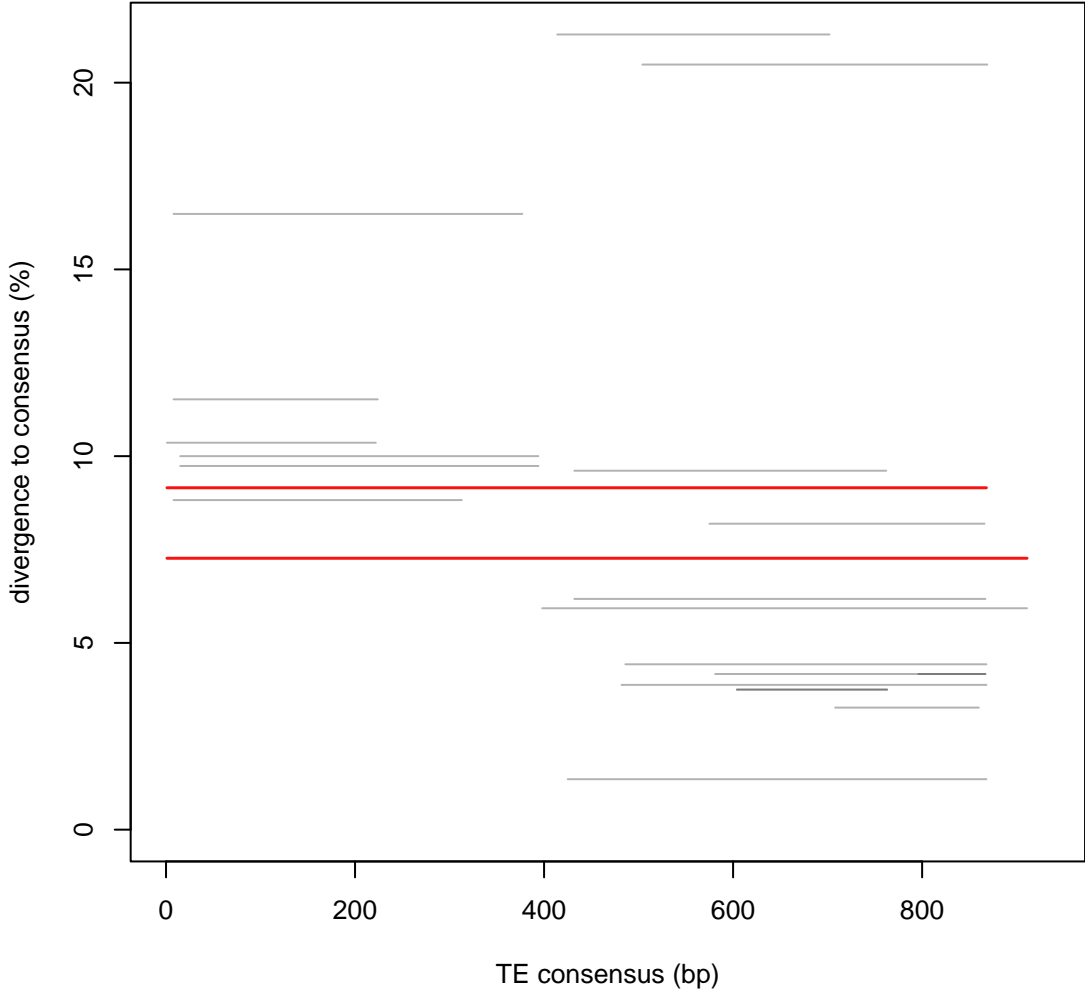
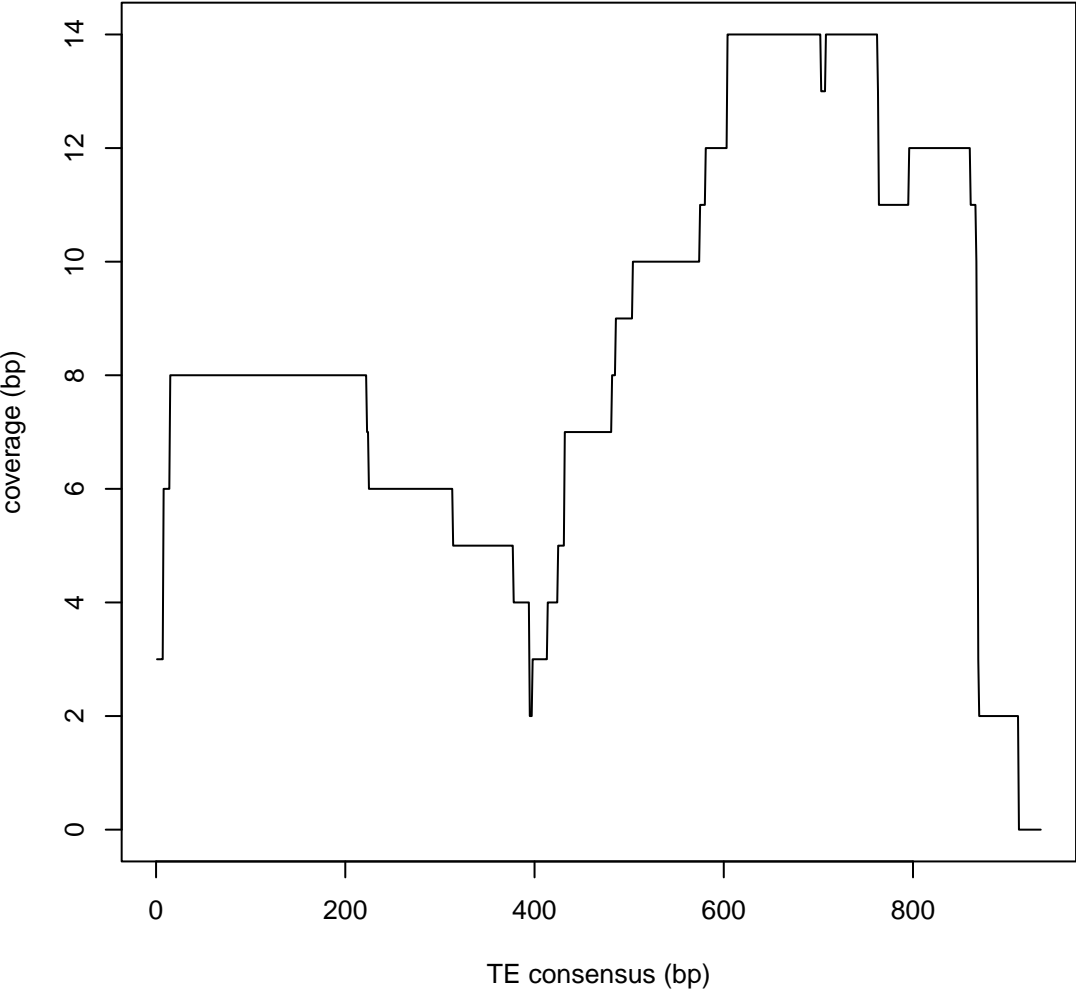


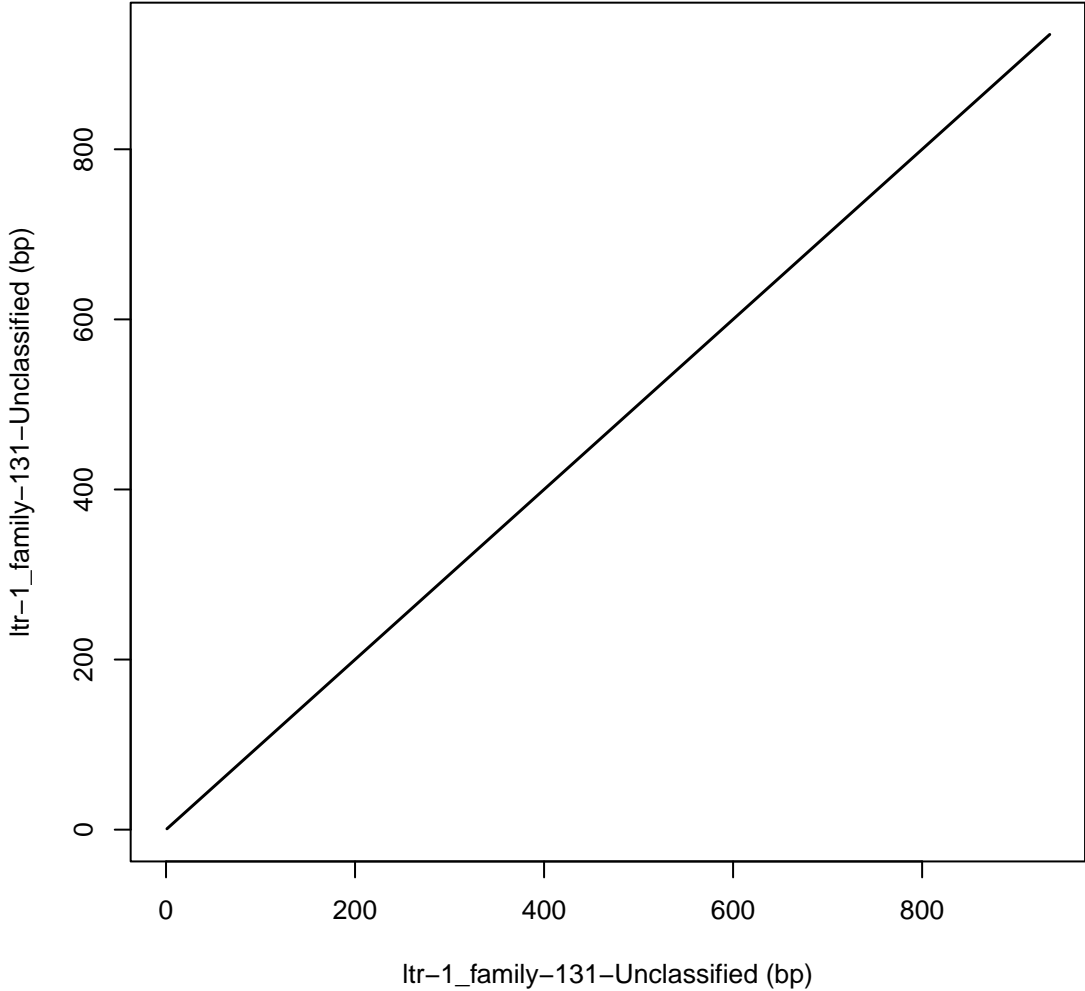
TE: ltr-1\_family-131-Unclassified  
consensus size: 935bp; fragments: 22; full length: 2 (>=841.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

