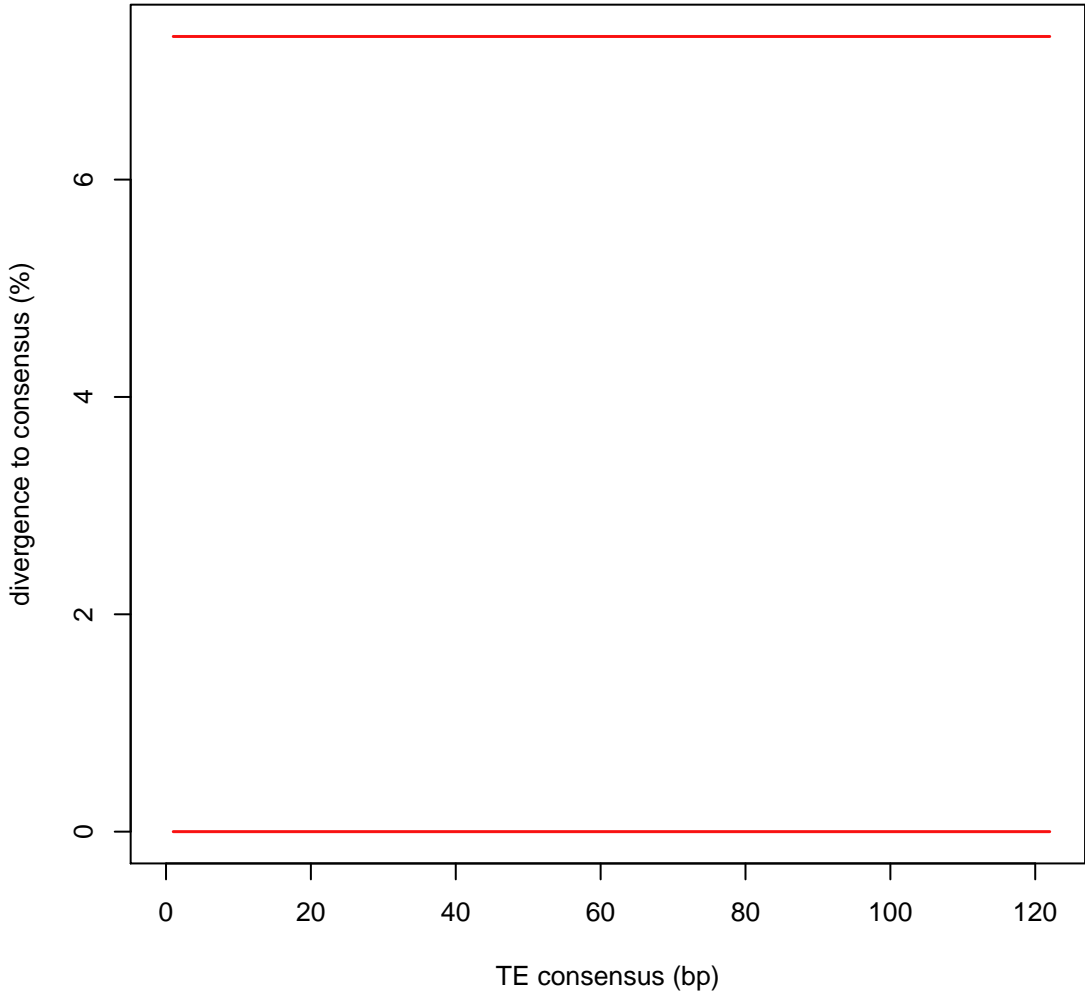
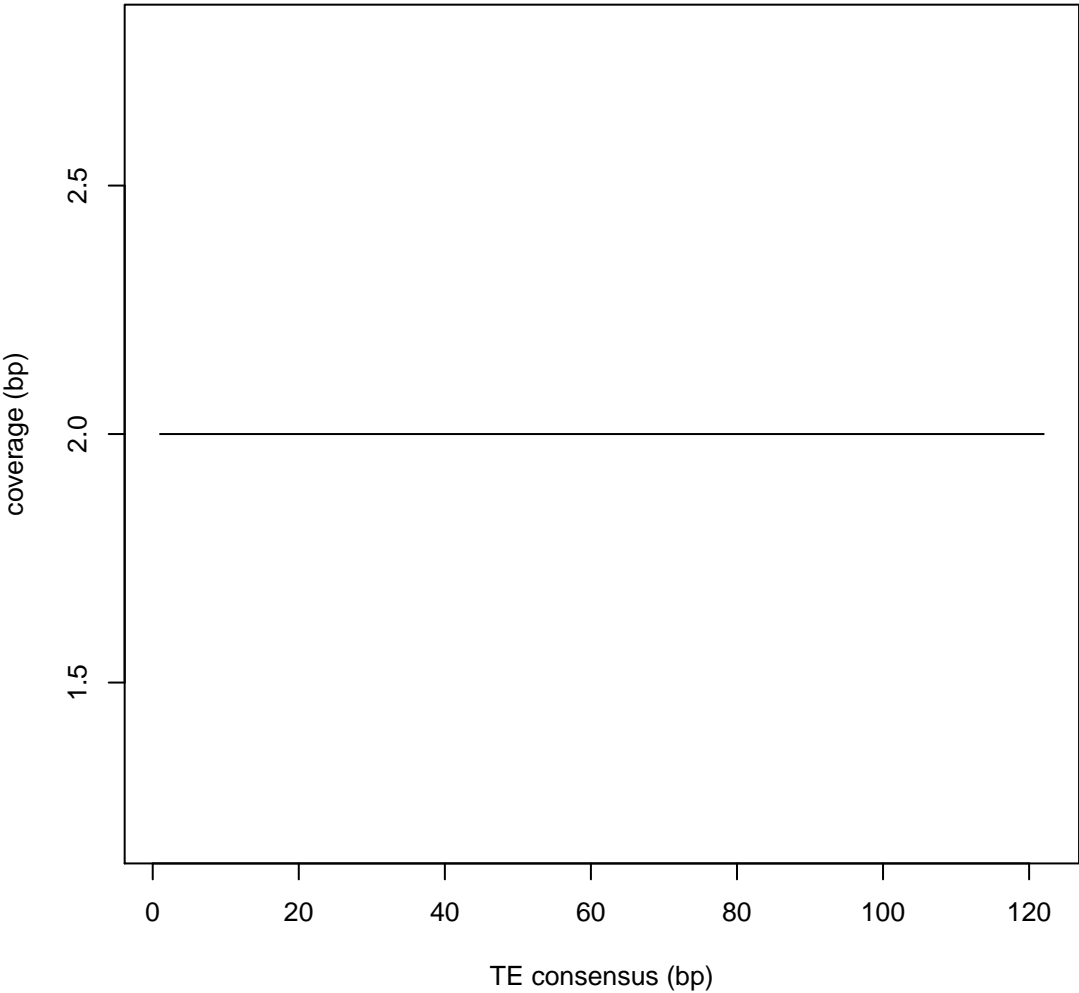


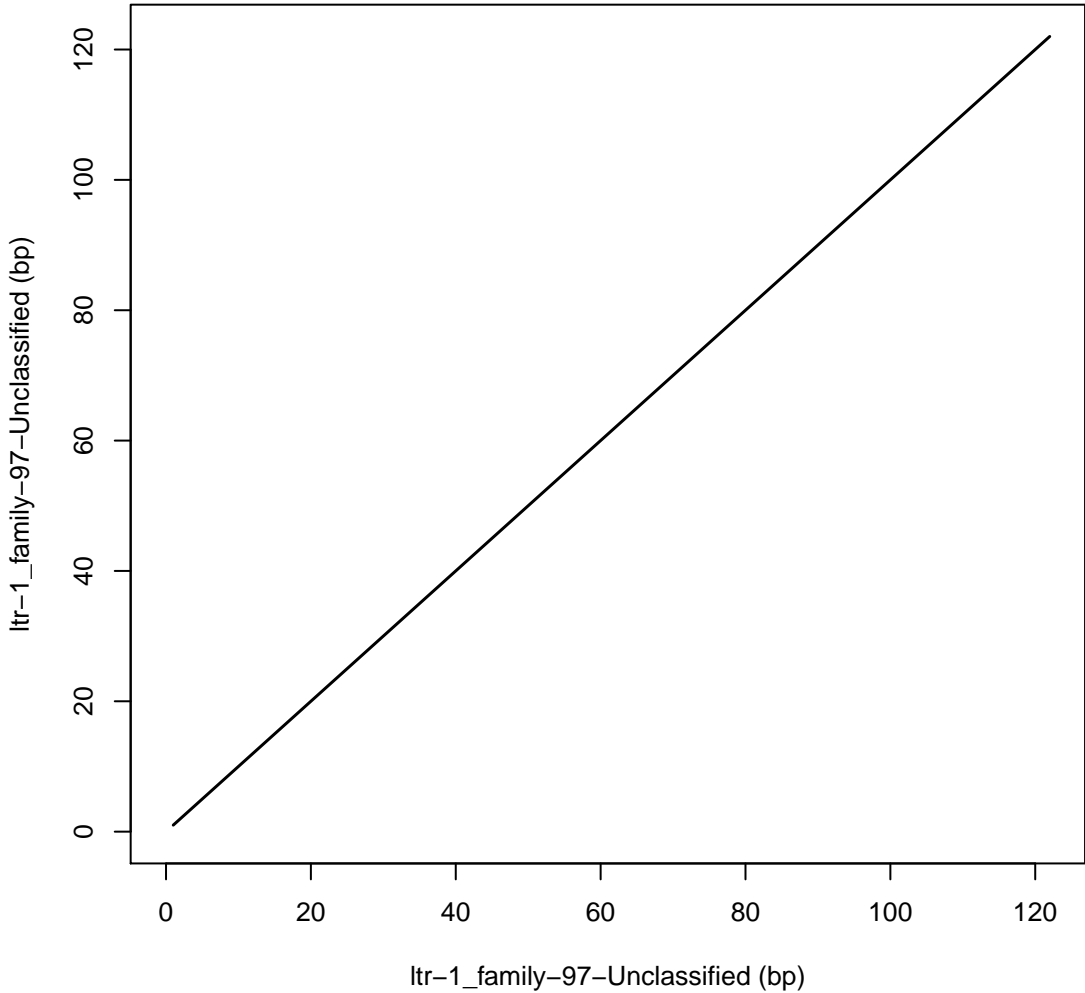
TE: Itr-1\_family-97-Unclassified  
consensus size: 122bp; fragments: 2; full length: 2 (>=109.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

