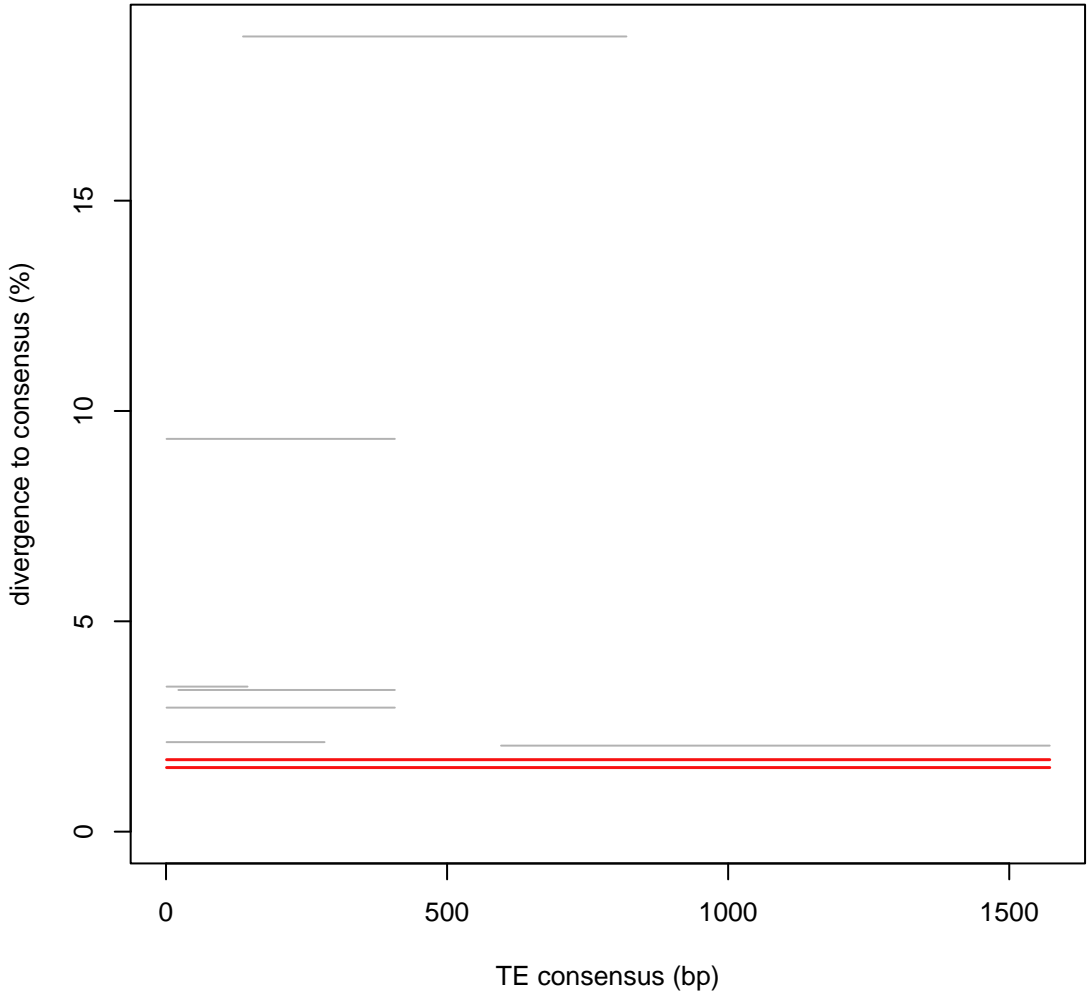
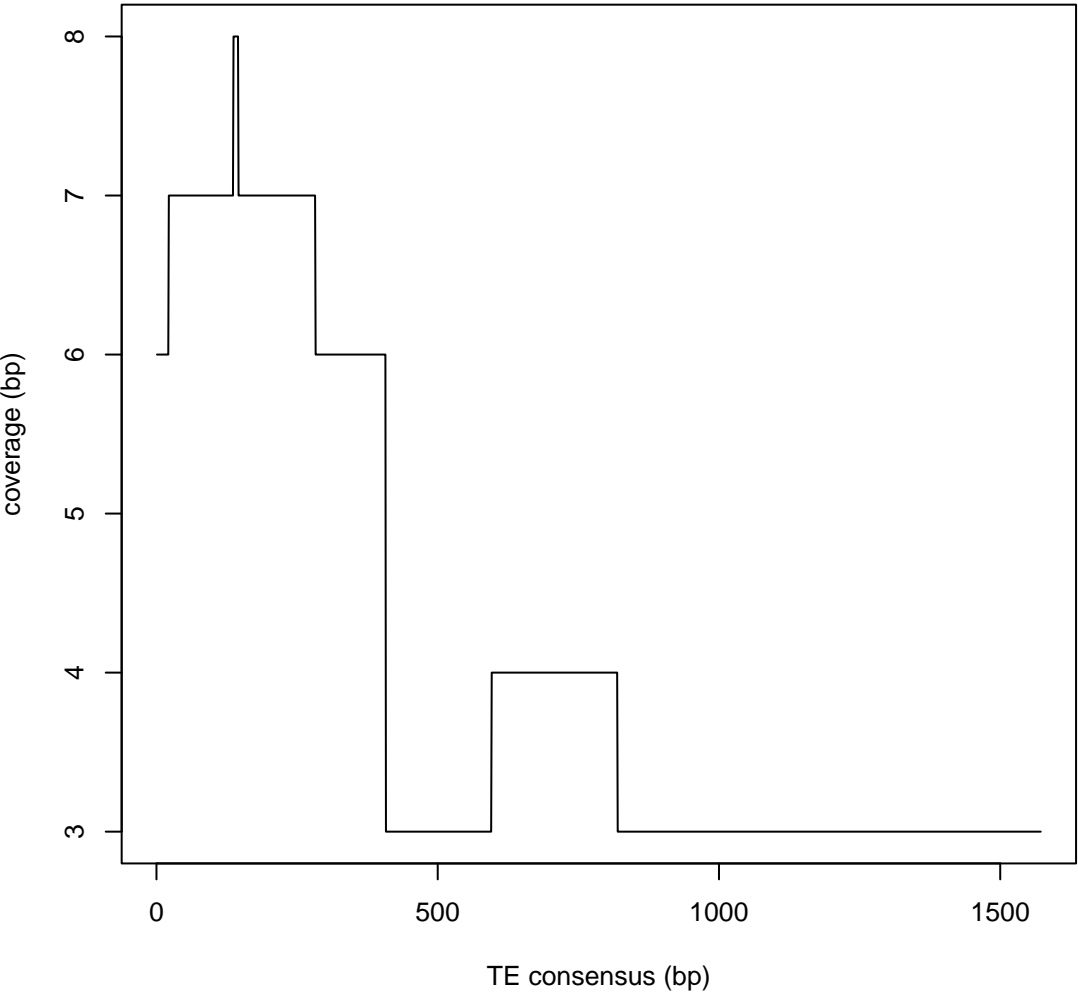


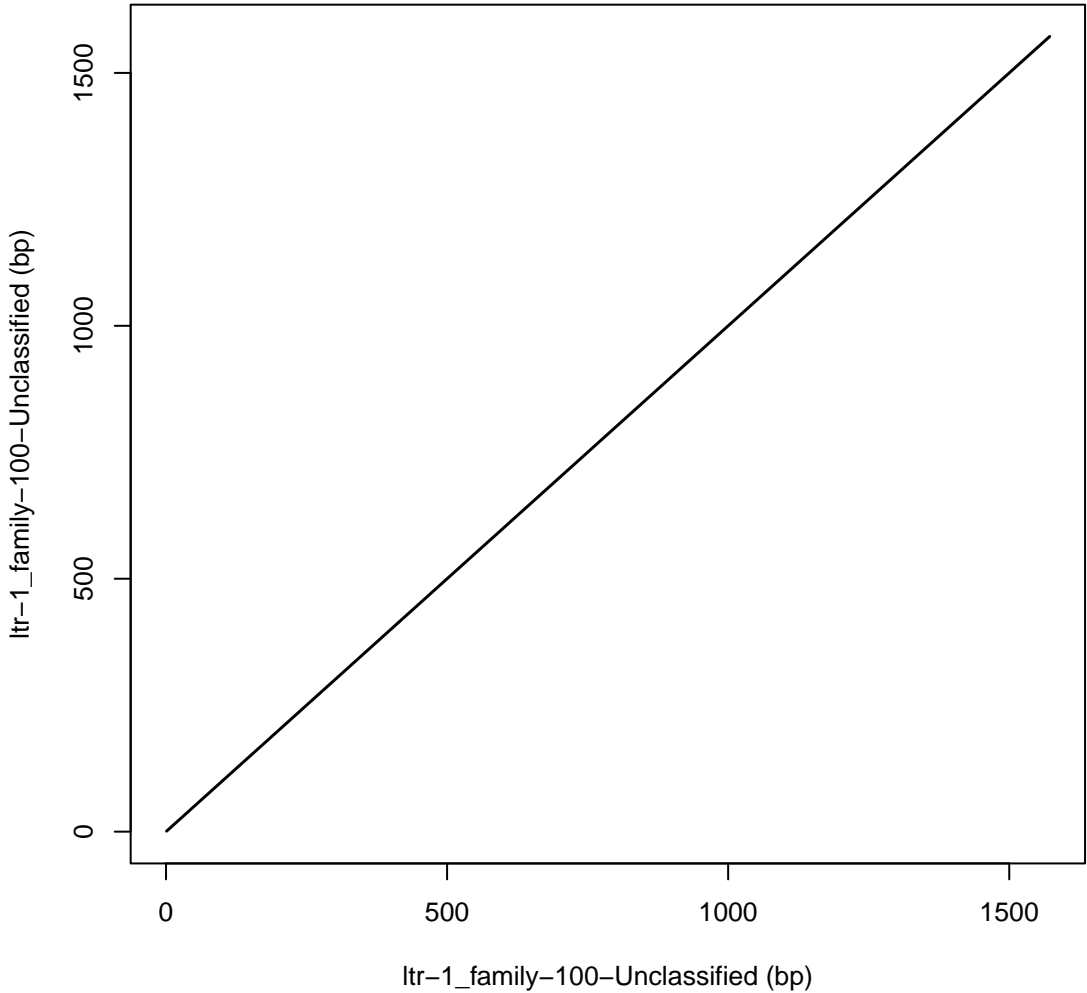
TE: ltr-1_family-100-Unclassified
consensus size: 1572bp; fragments: 9; full length: 2 (≥ 1414.8 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

