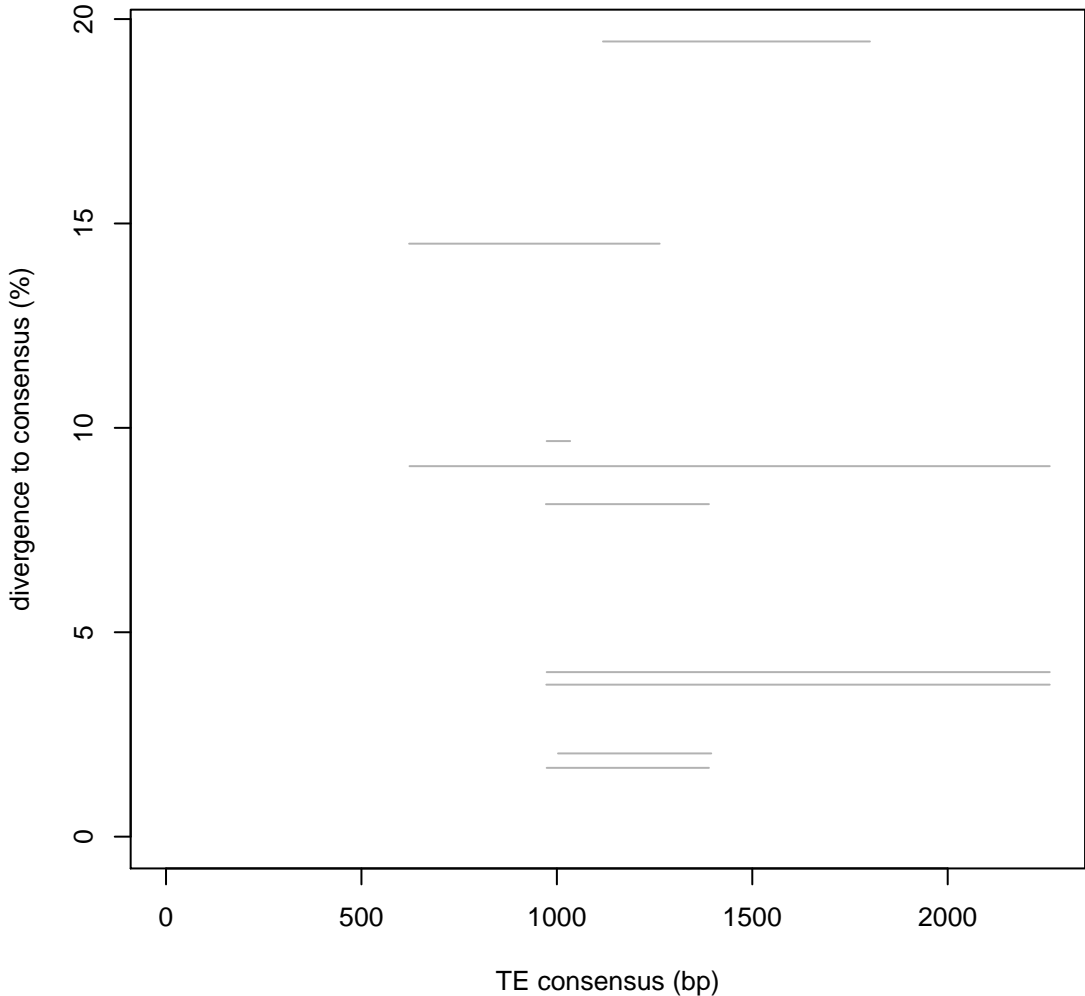
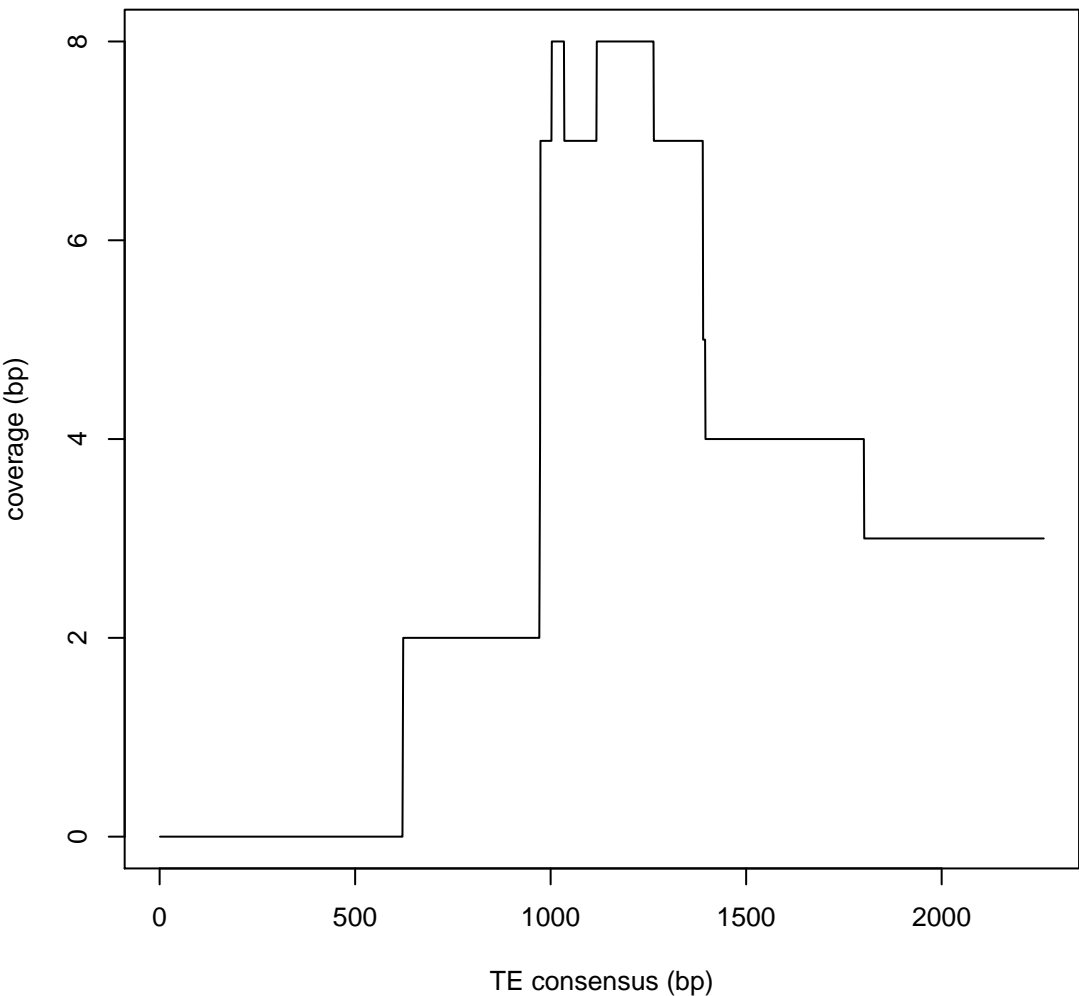


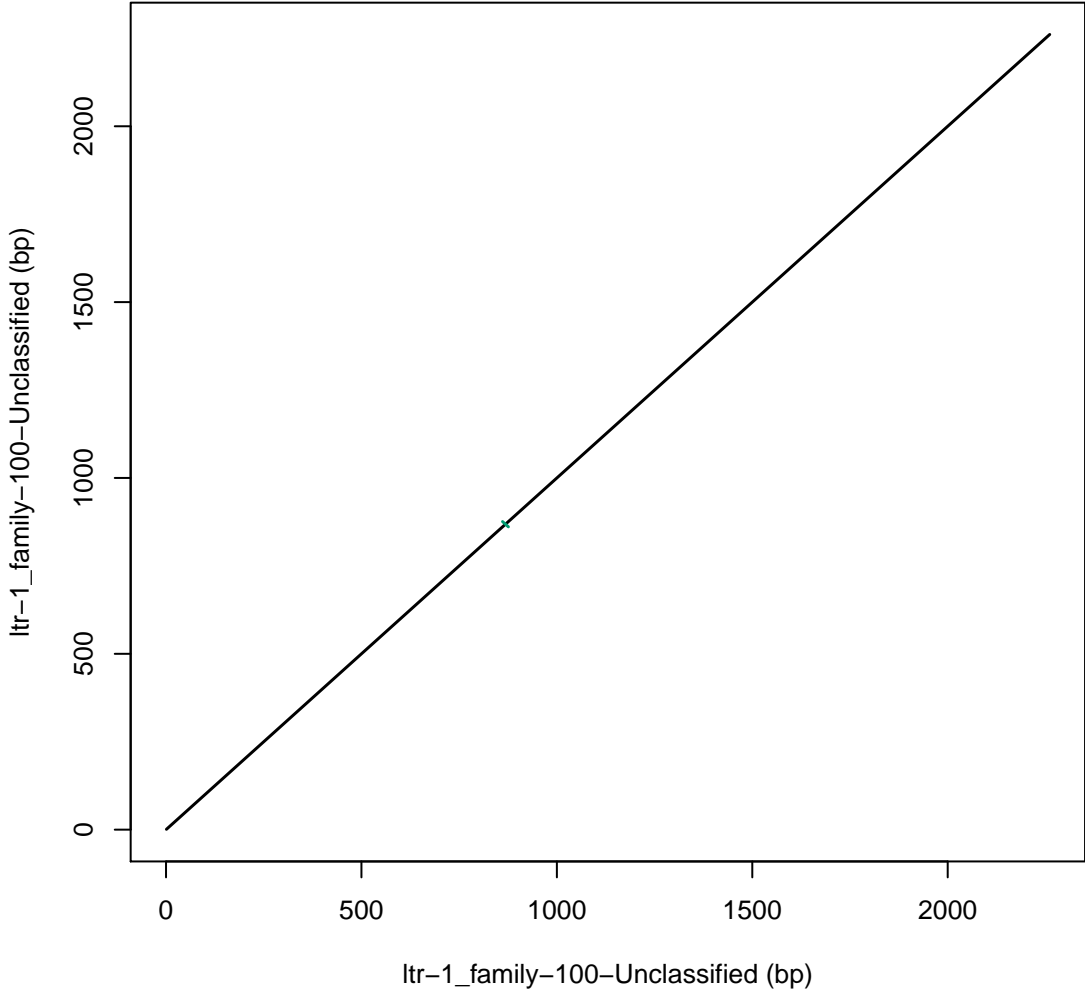
TE: ltr-1_family-100-Unclassified
consensus size: 2261bp; fragments: 9; full length: 0 (>=2034.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

