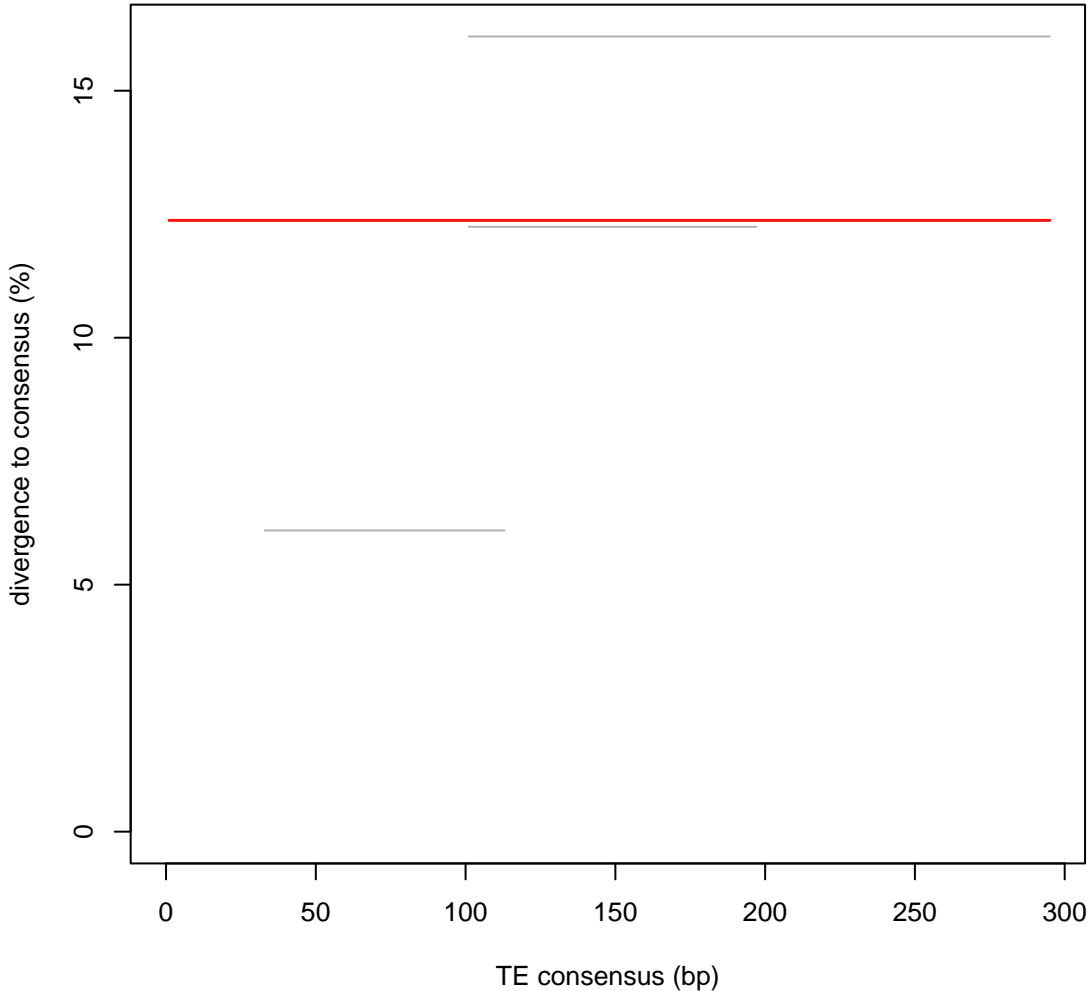
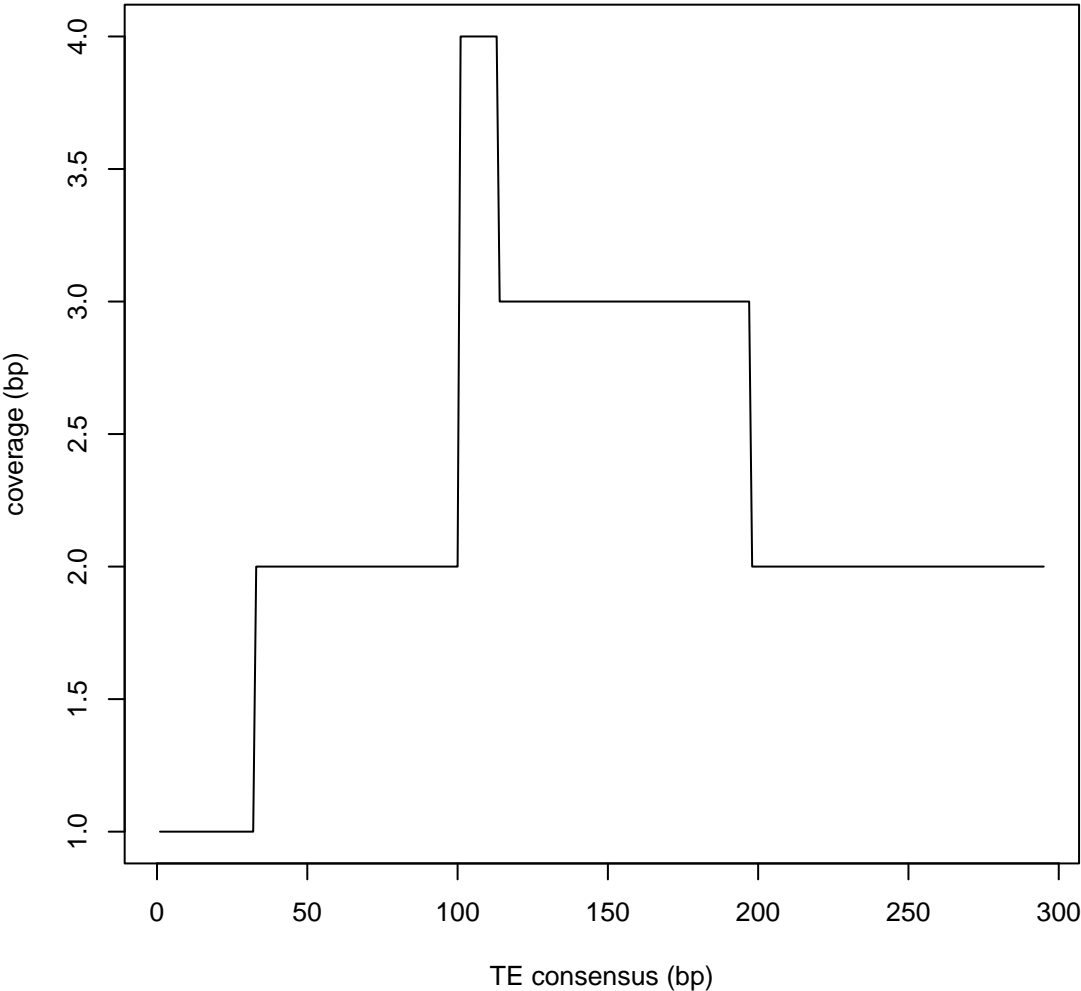


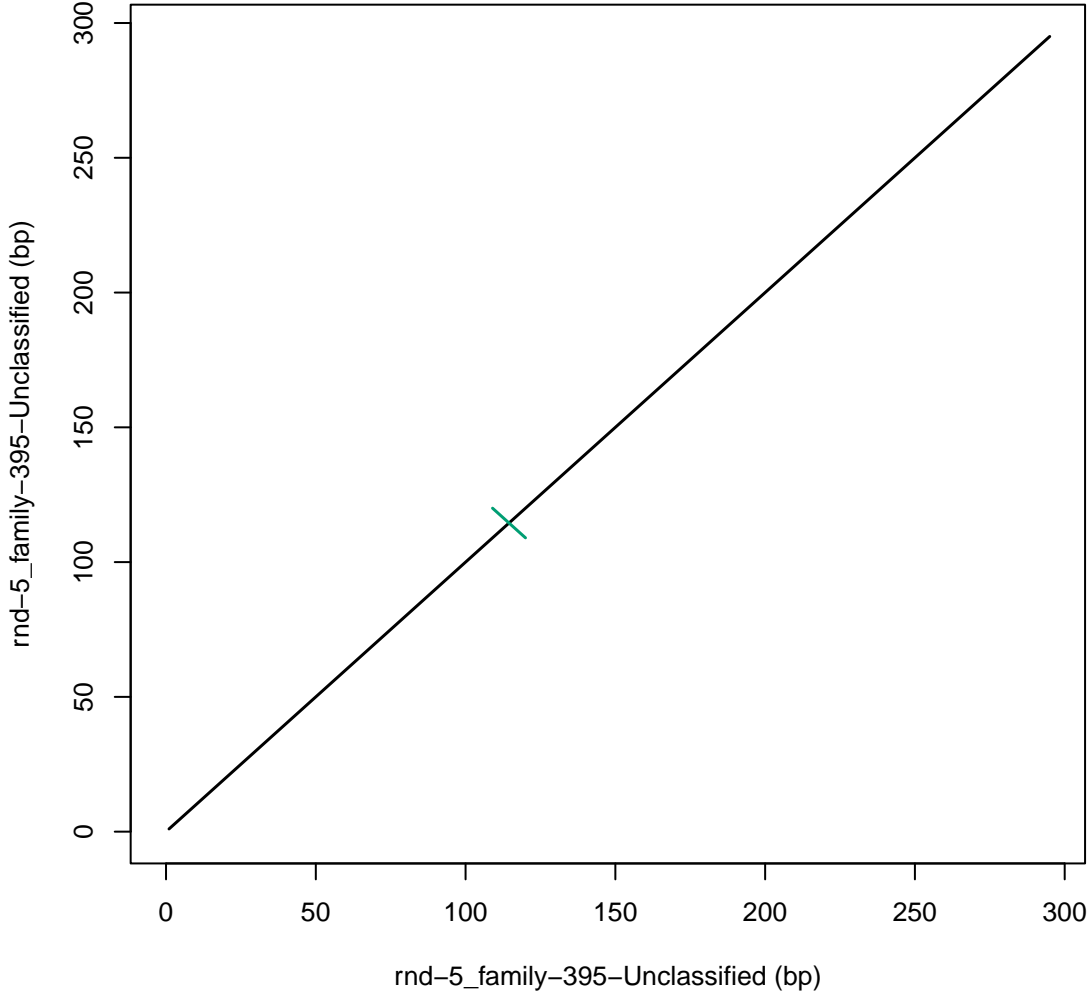
TE: rnd-5_family-395-Unclassified
consensus size: 295bp; fragments: 4; full length: 1 (>=265.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

