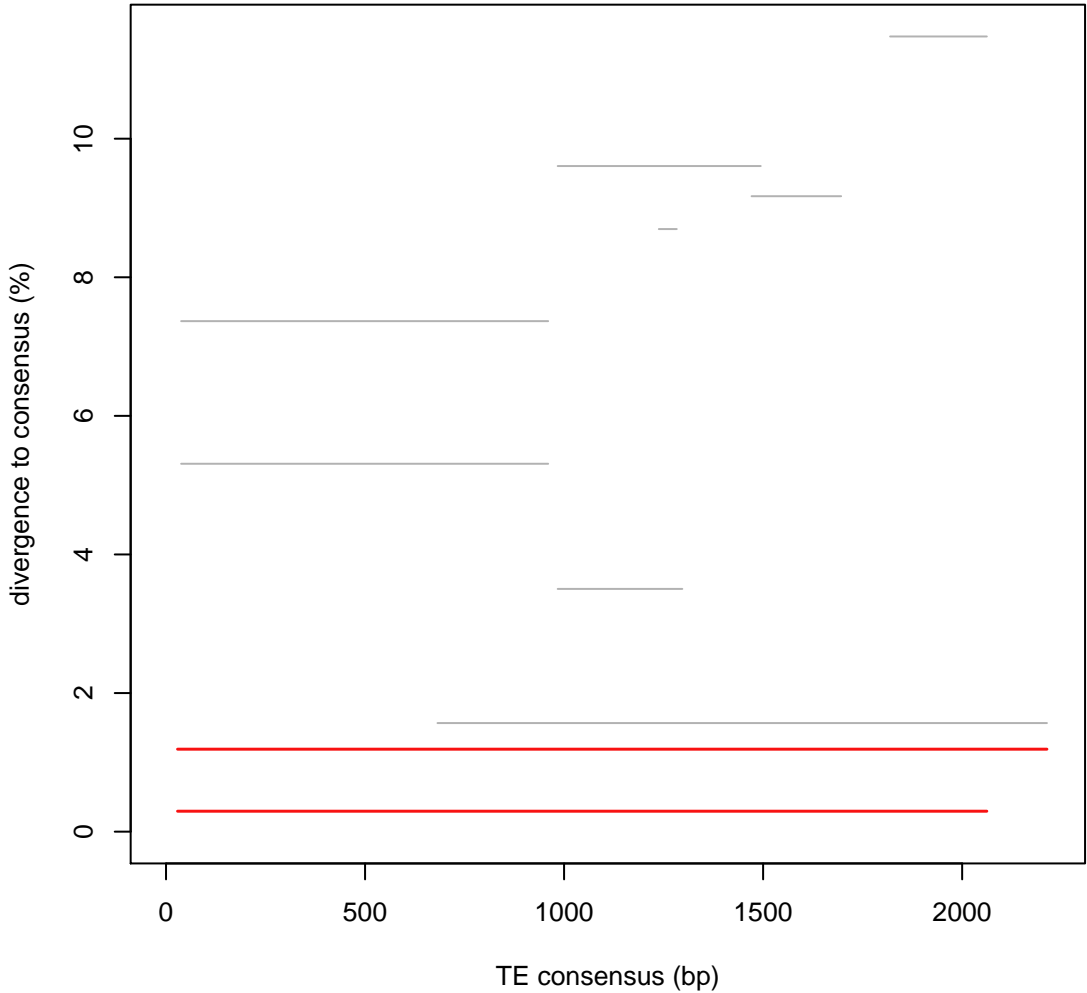
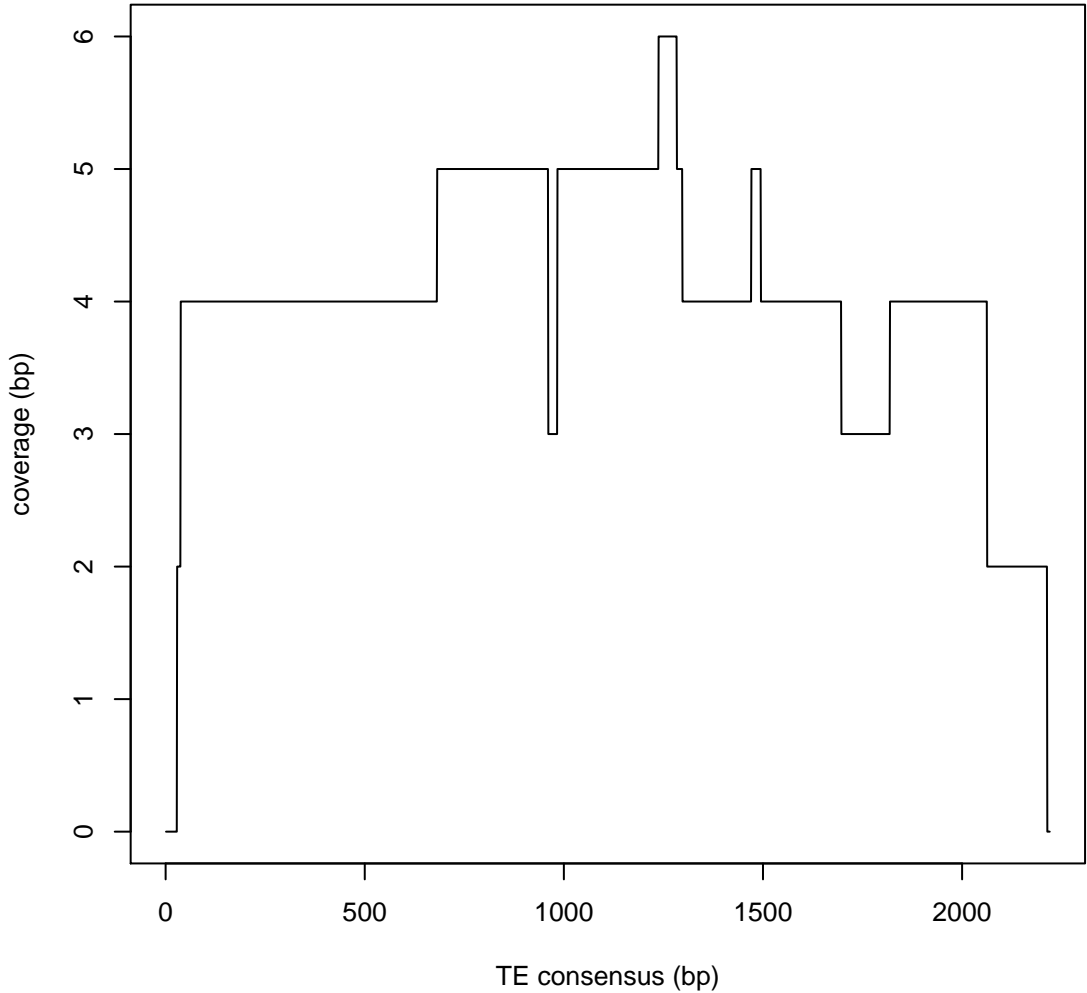


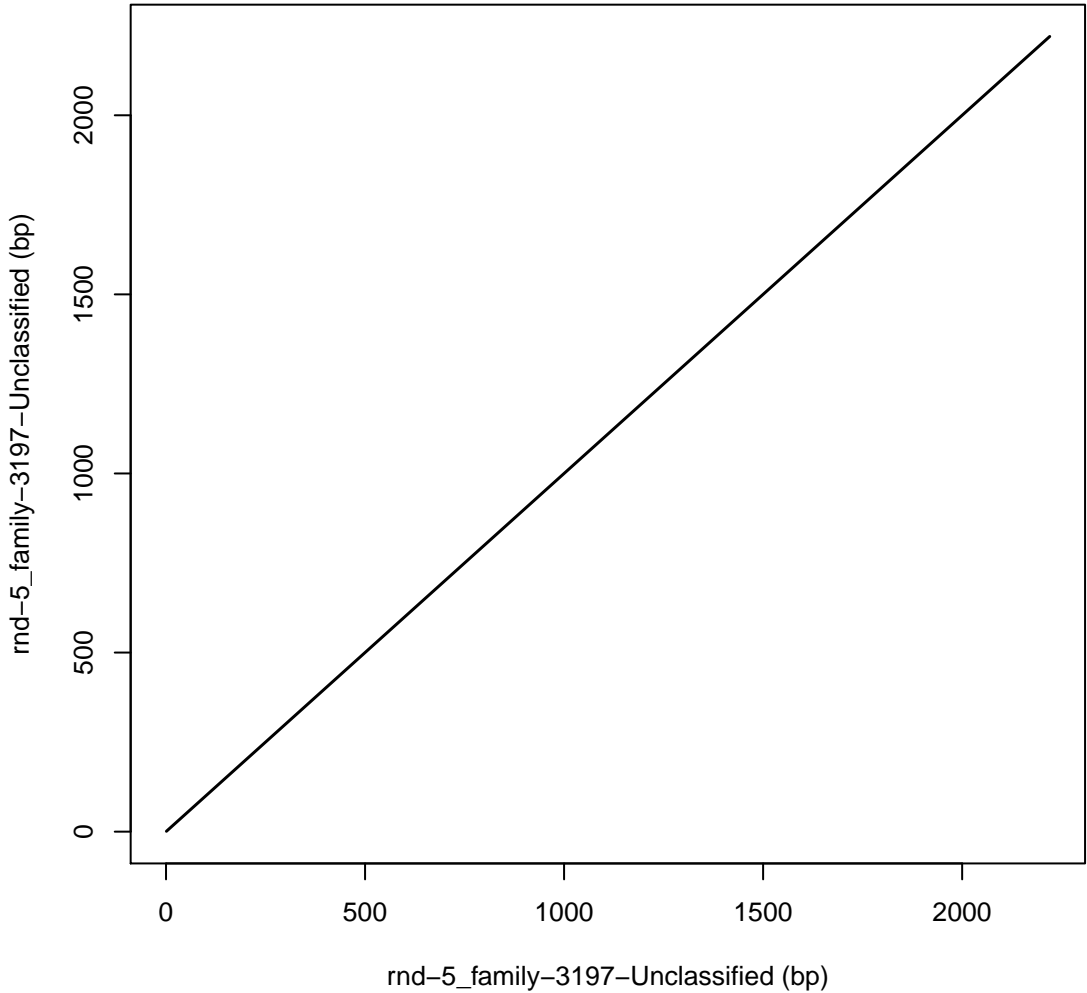
TE: rnd-5_family-3197-Unclassified
consensus size: 2220bp; fragments: 10; full length: 2 (>=1998bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

