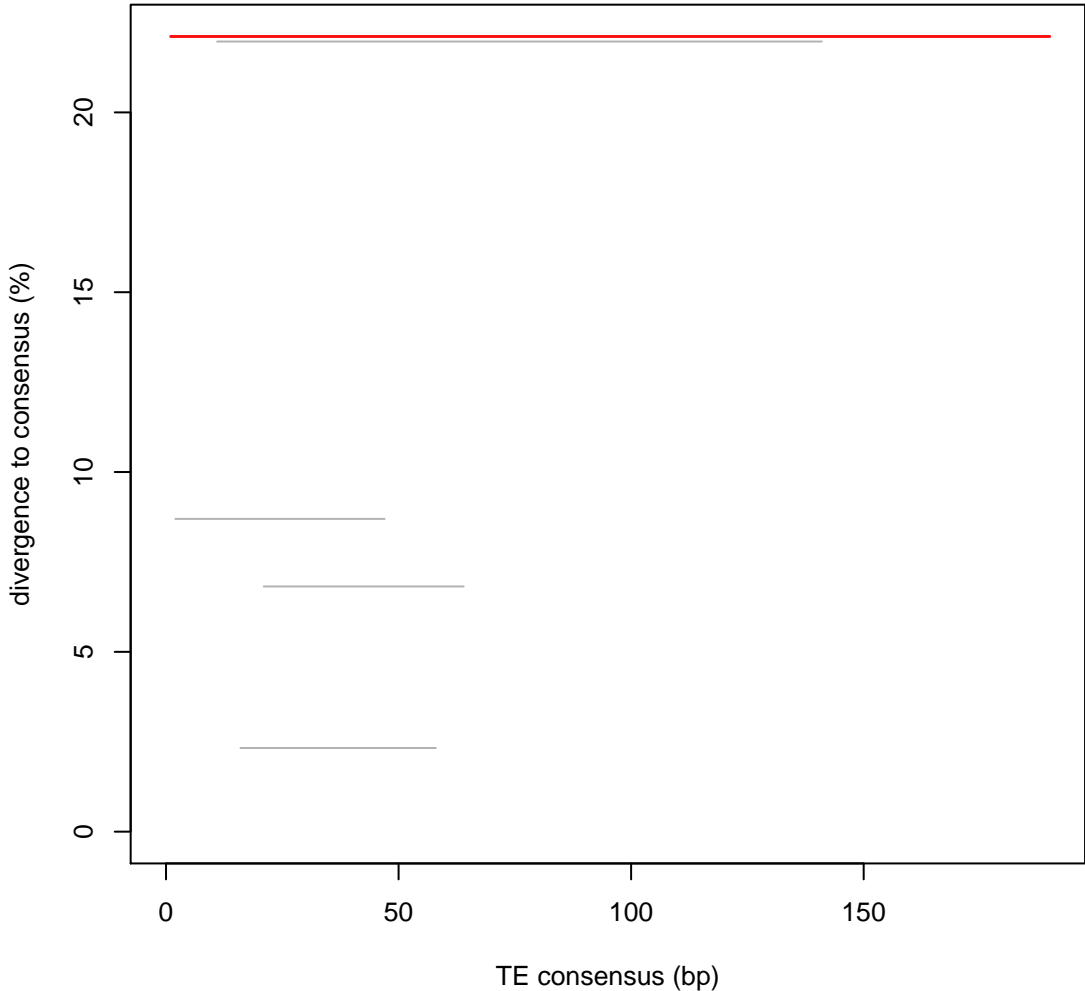
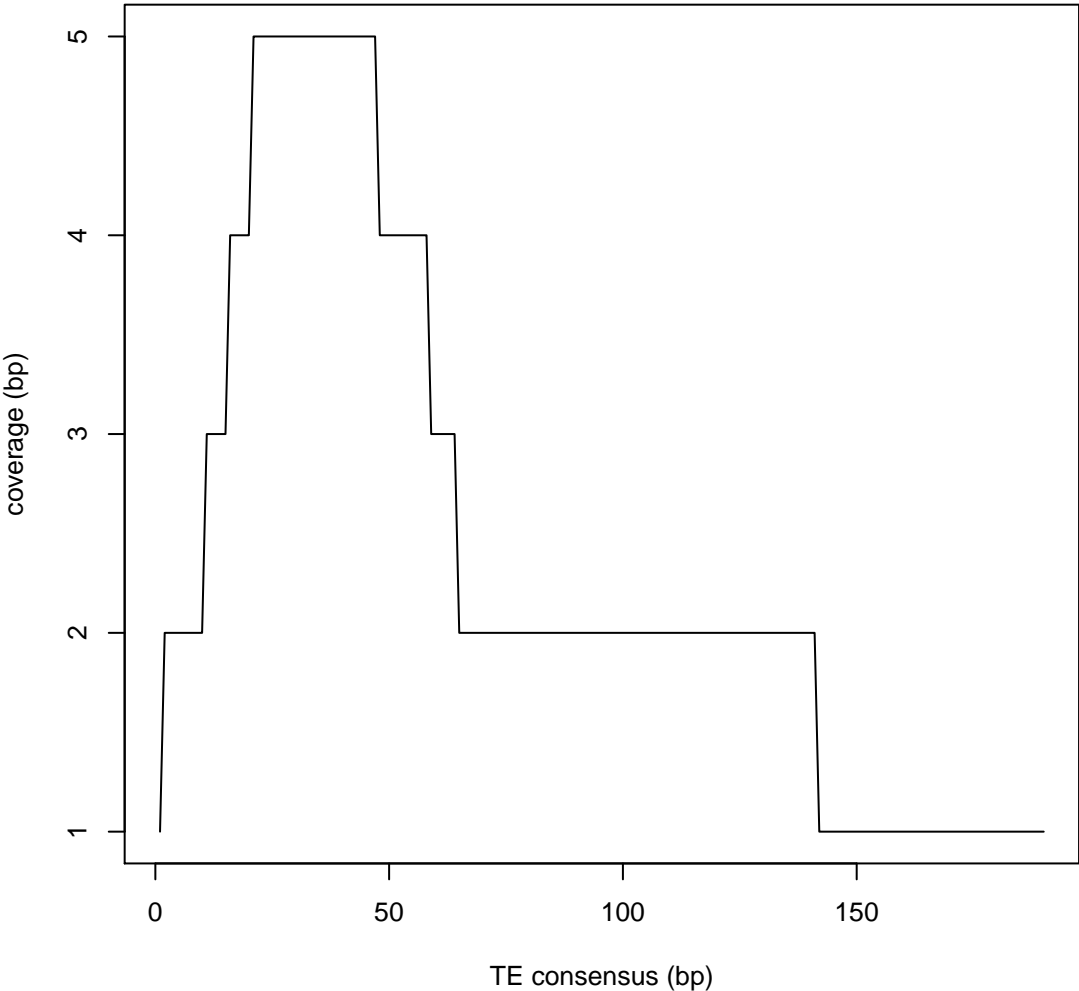


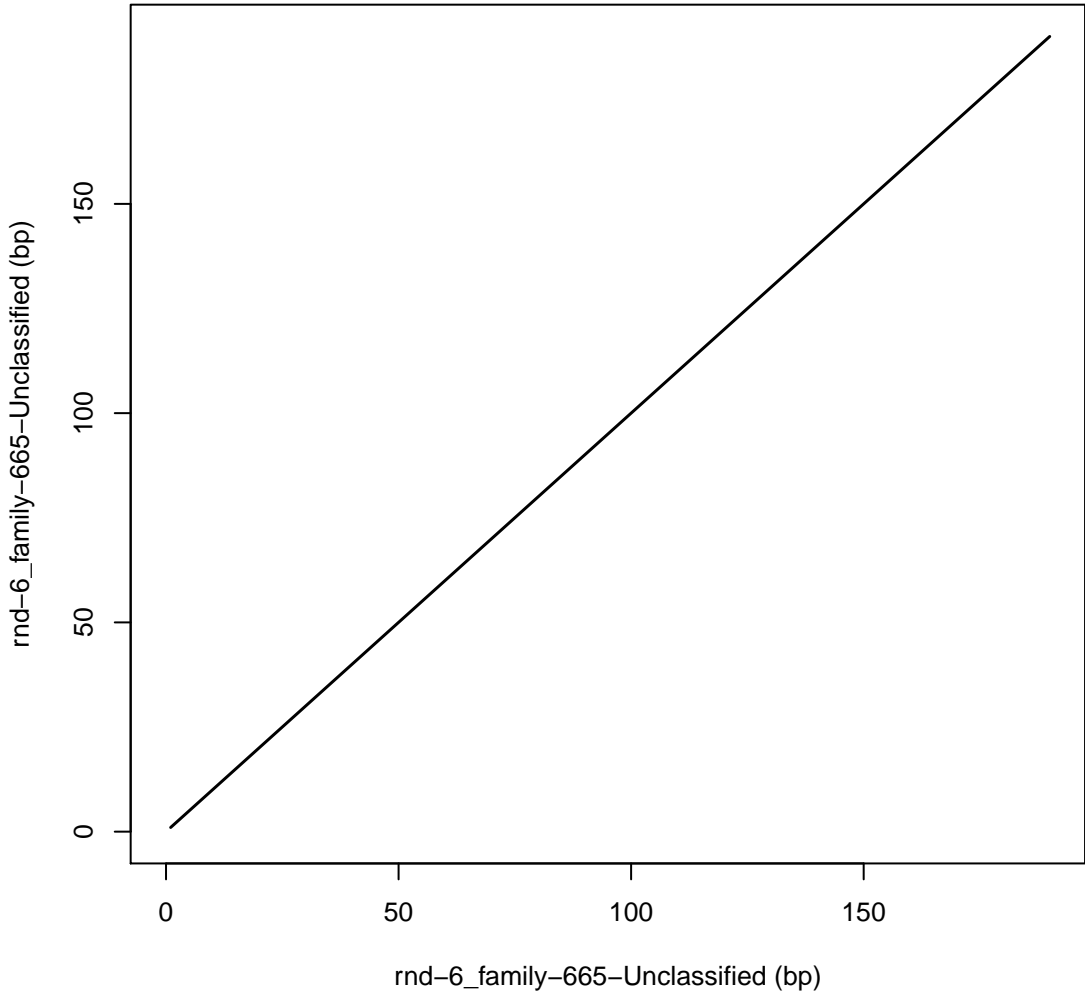
TE: rnd-6_family-665-Unclassified
consensus size: 190bp; fragments: 5; full length: 1 (>=171bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

