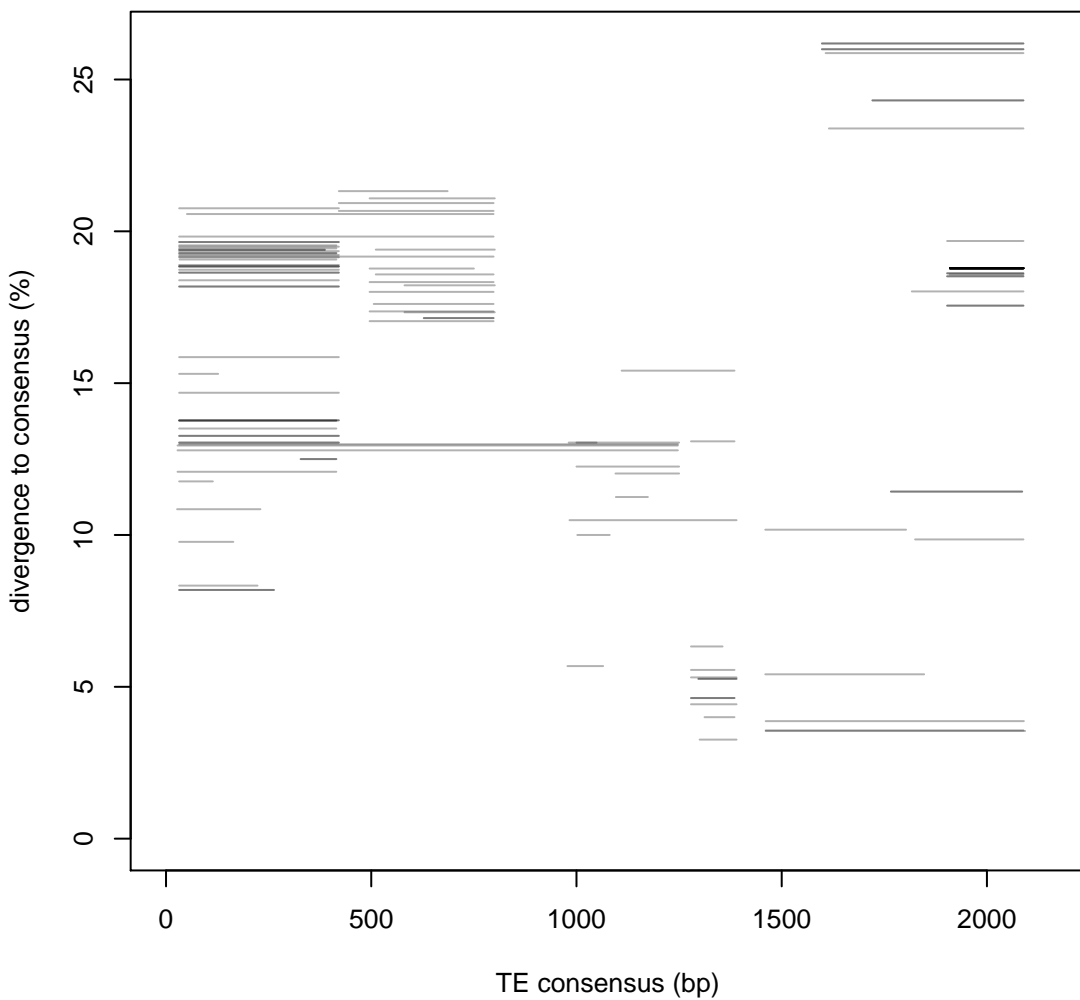
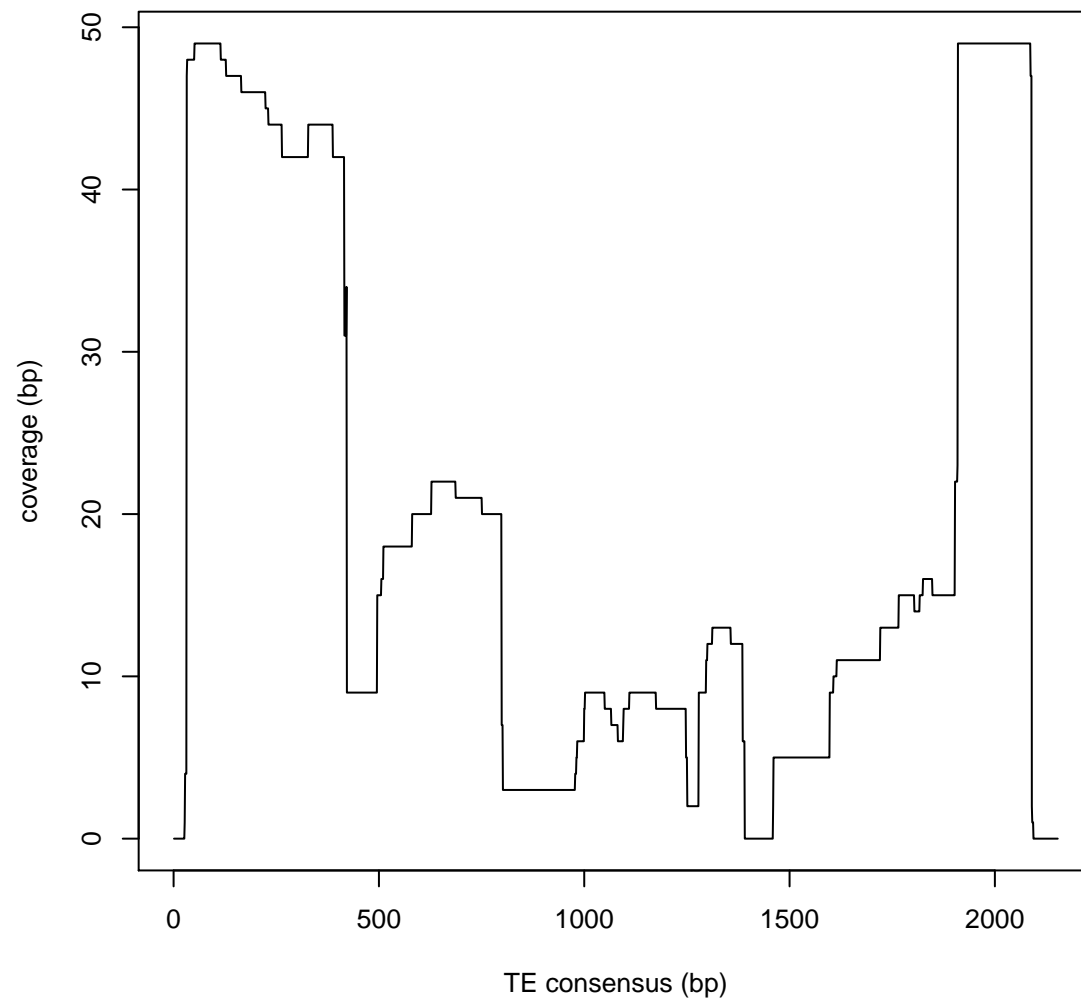


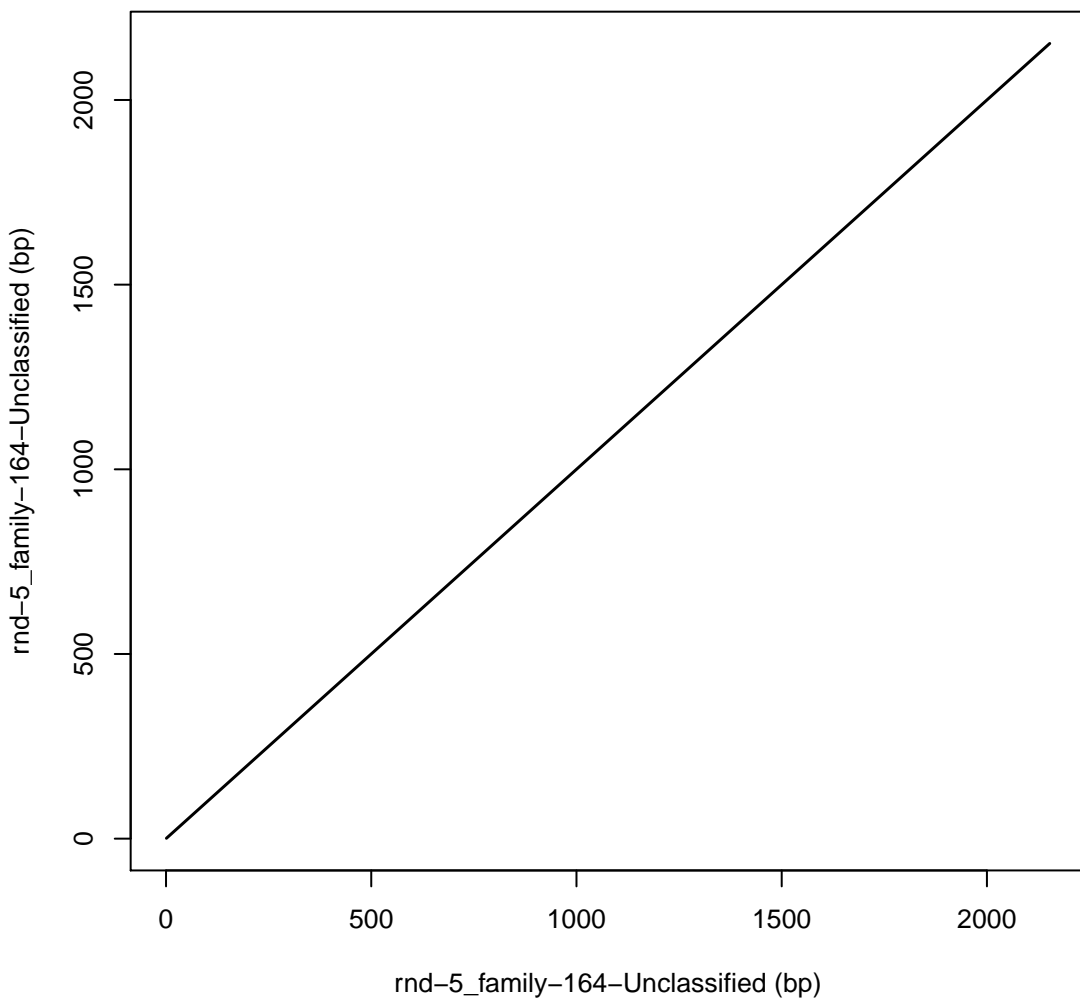
TE: rnd-5_family-164-Unclassified
consensus size: 2153bp; fragments: 138; full length: 0 (>=1937.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

