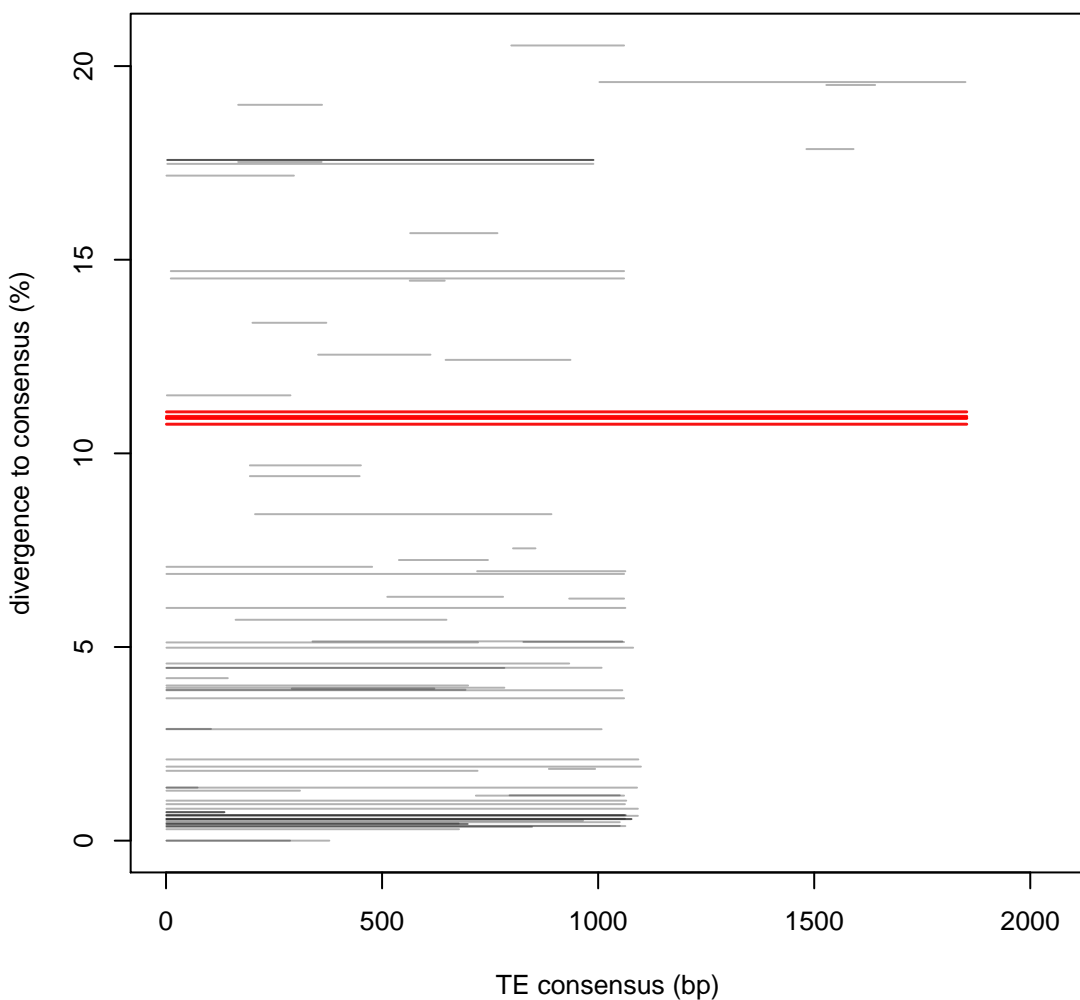
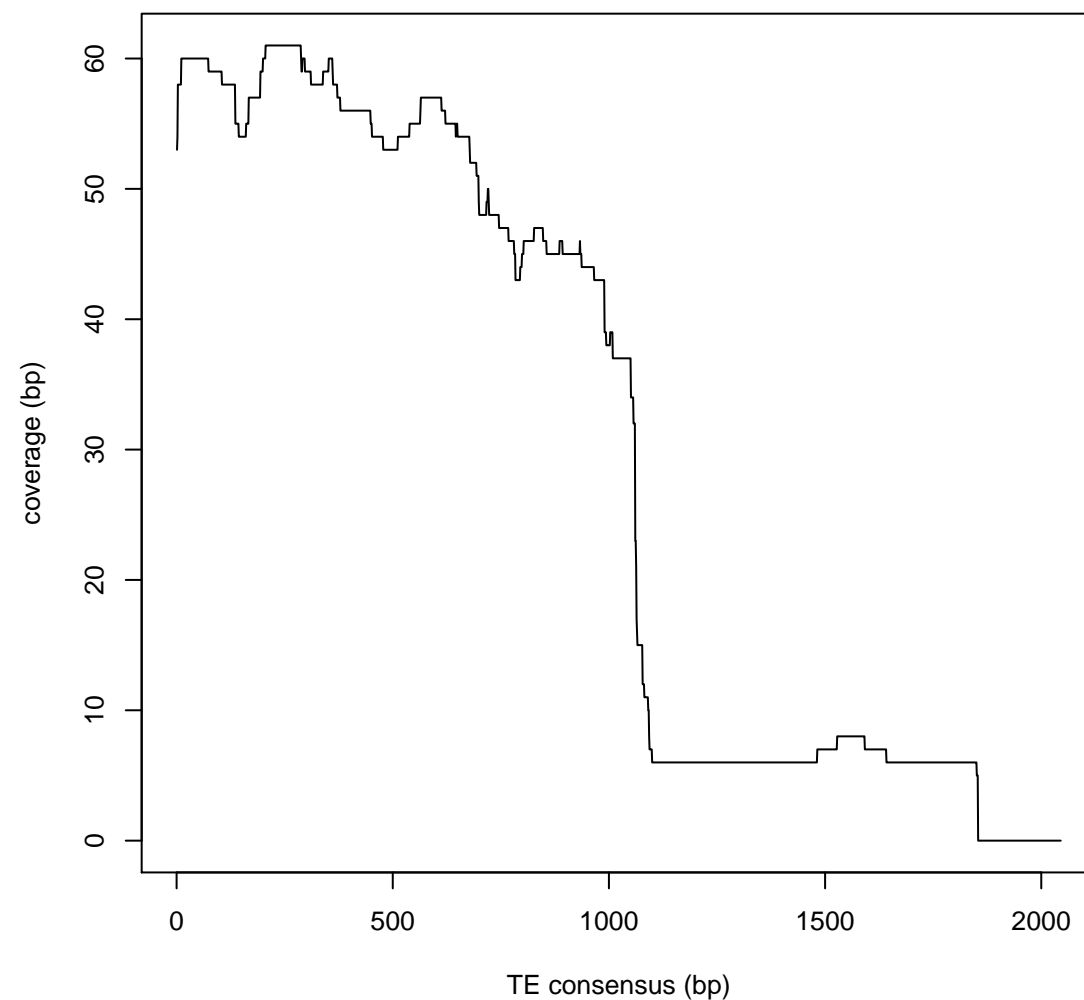


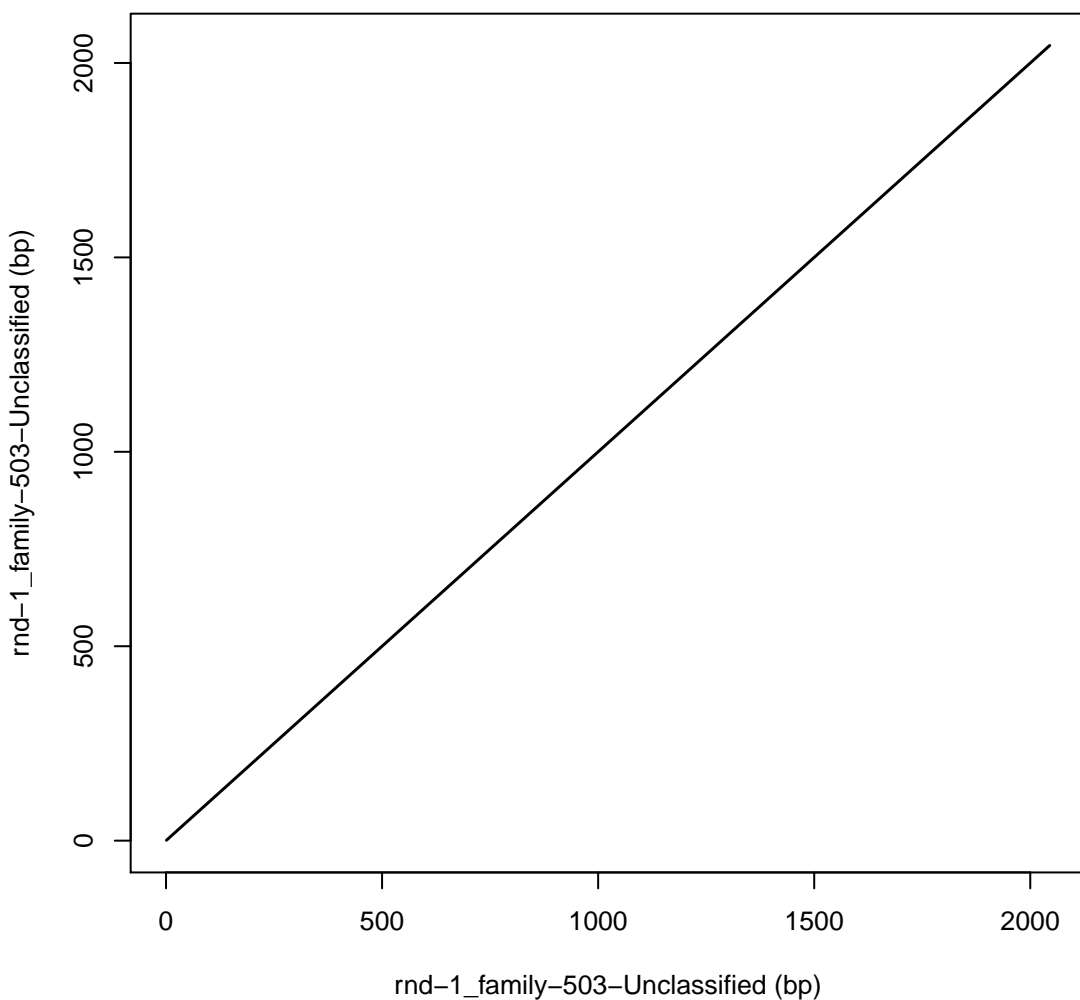
TE: rnd-1_family-503-Unclassified
consensus size: 2045bp; fragments: 86; full length: 5 (≥ 1840.5 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

