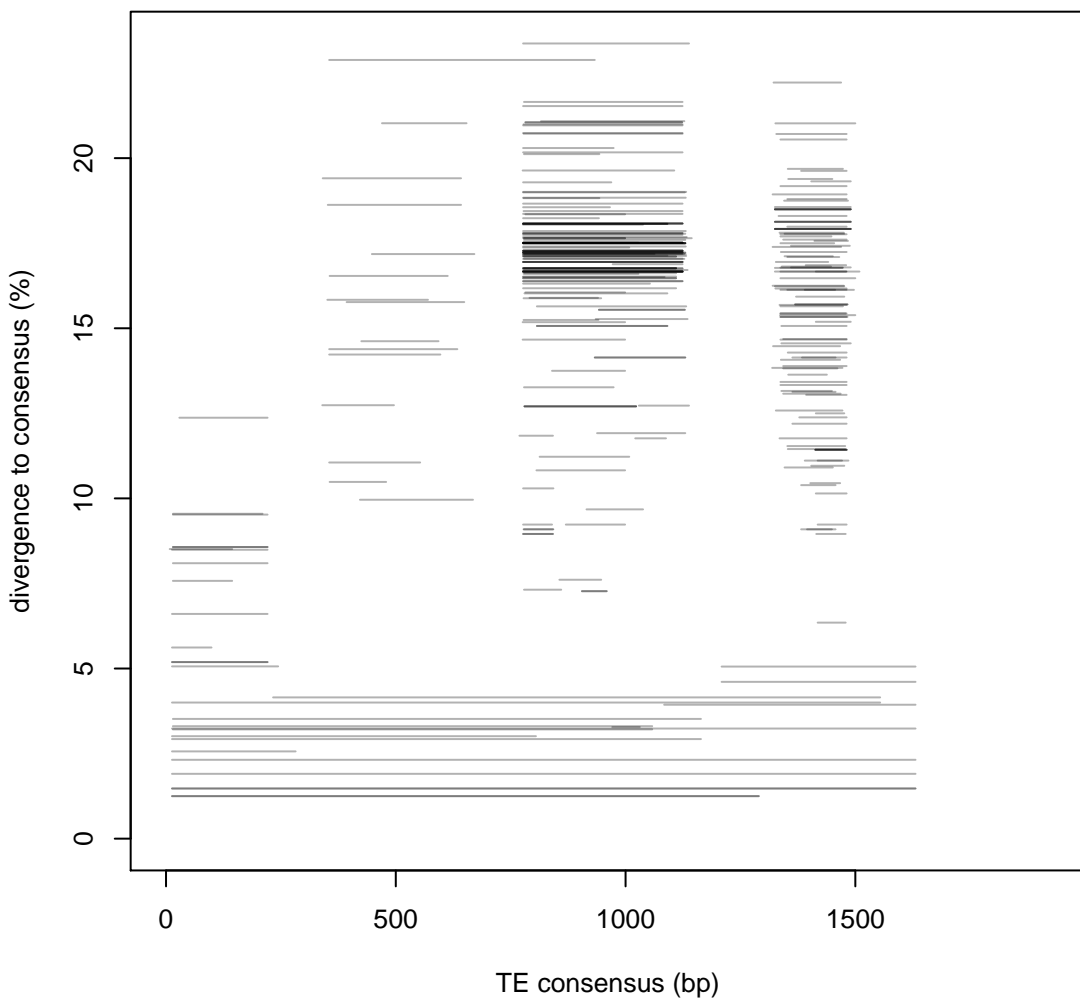
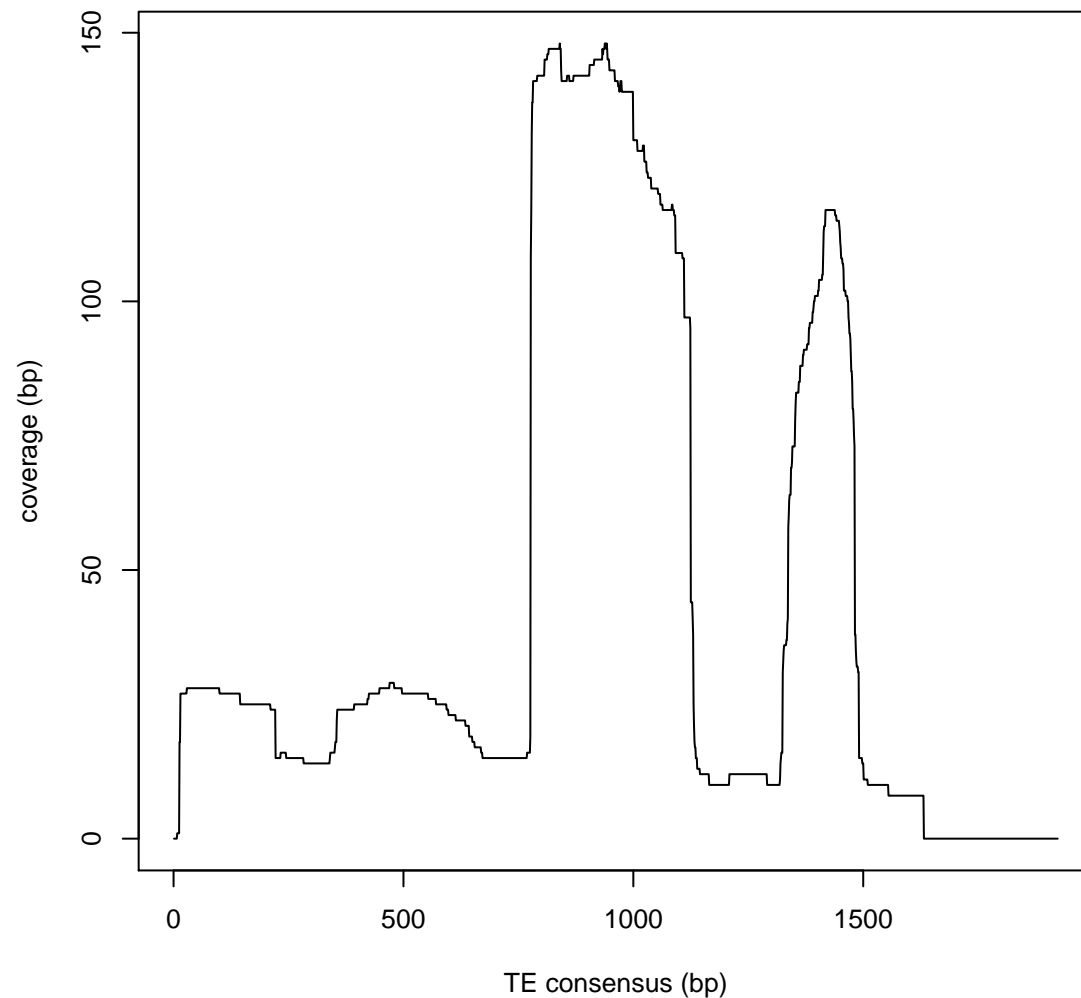


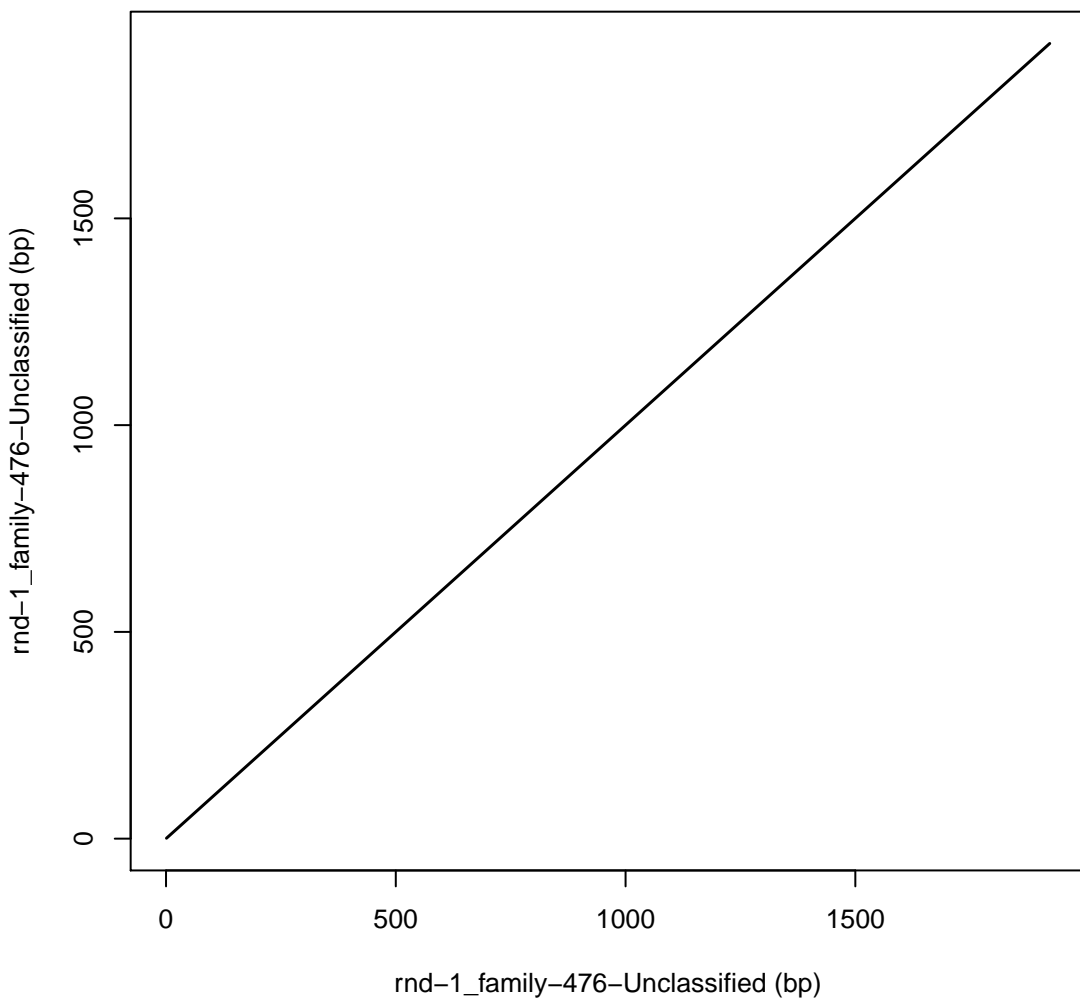
TE: rnd-1_family-476-Unclassified
consensus size: 1923bp; fragments: 303; full length: 0 (≥ 1730.7 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

