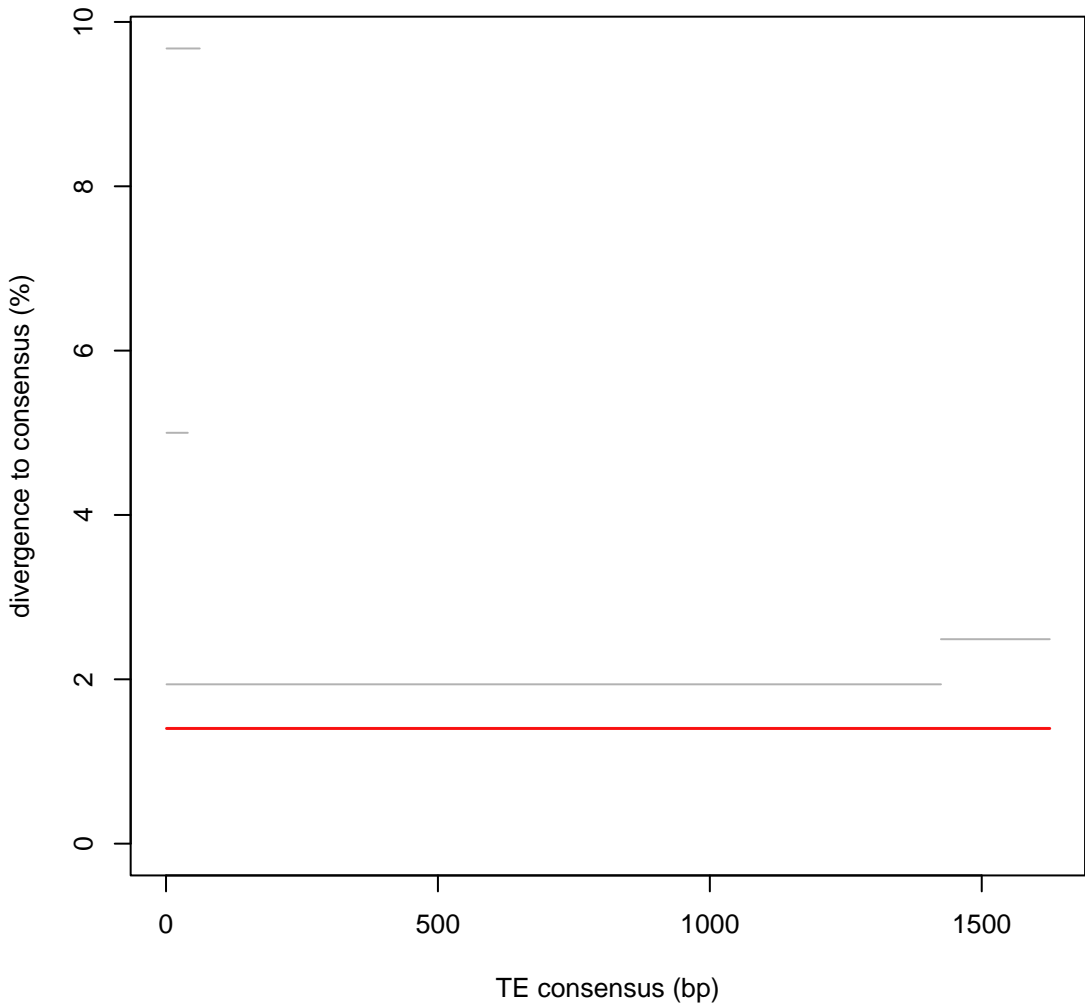
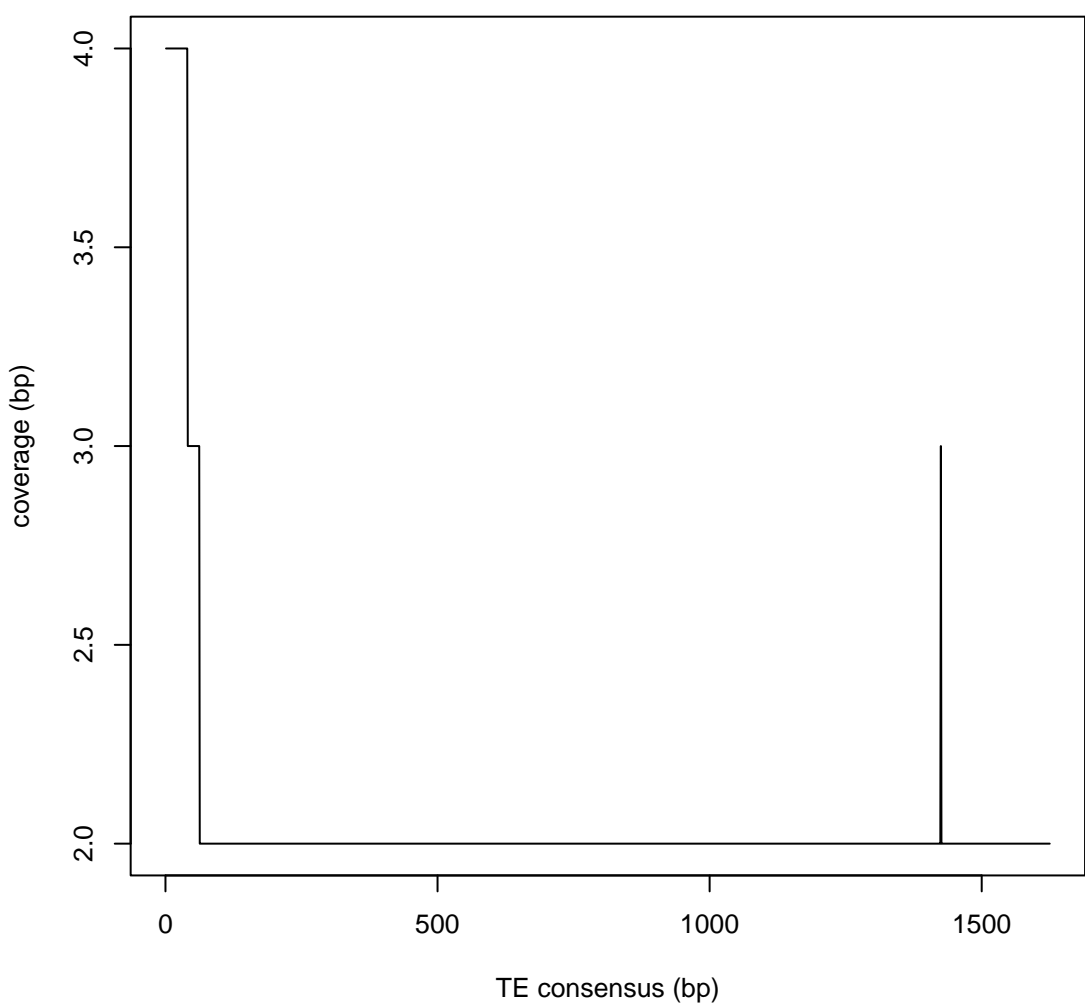


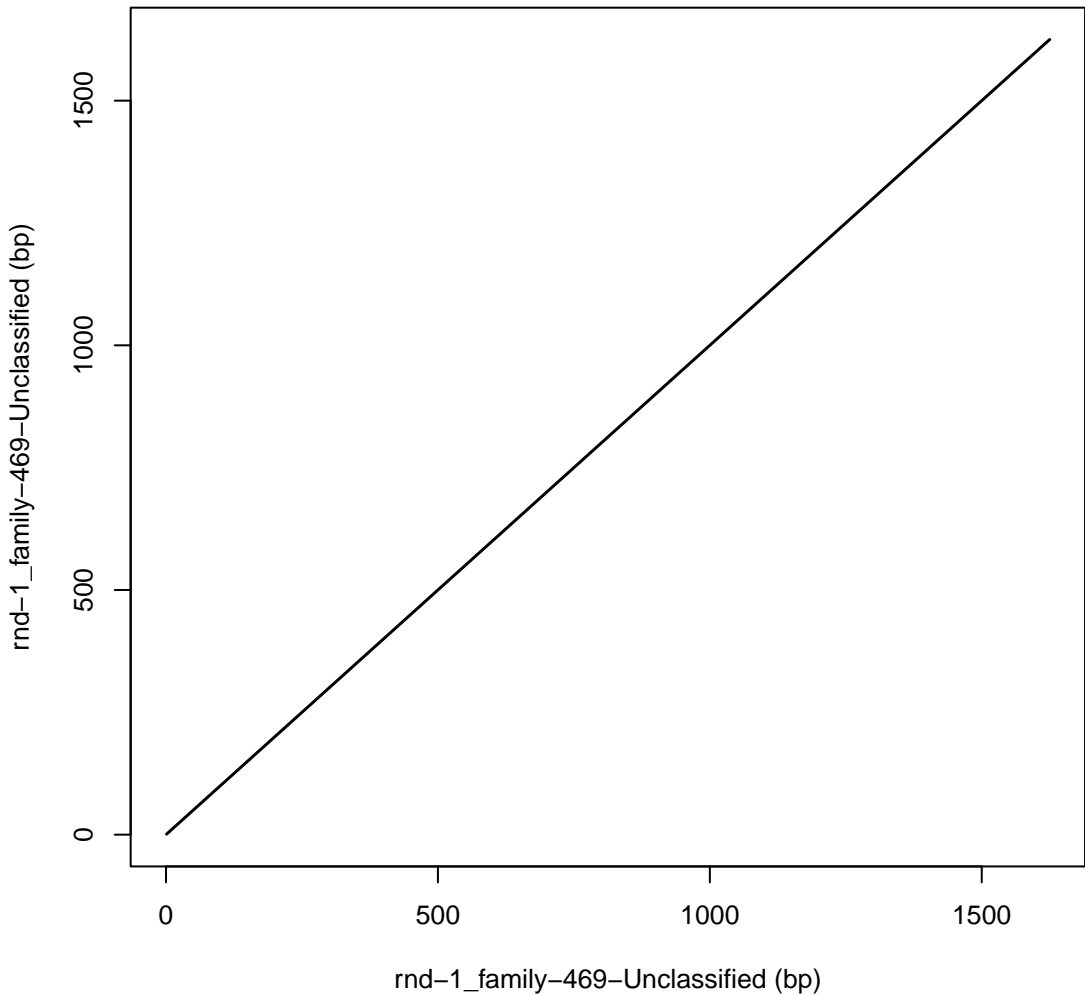
TE: rnd-1_family-469-Unclassified
consensus size: 1625bp; fragments: 5; full length: 1 (>=1462.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

