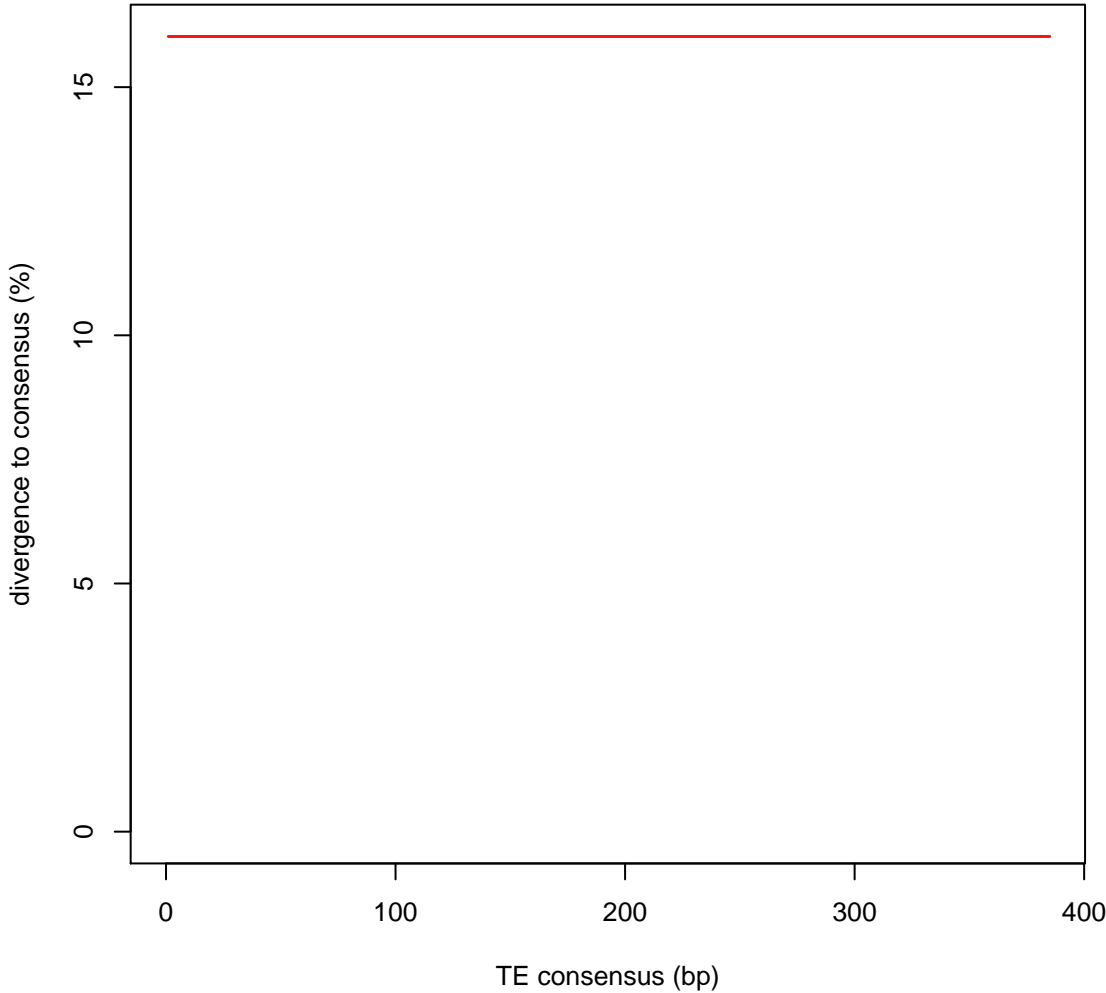
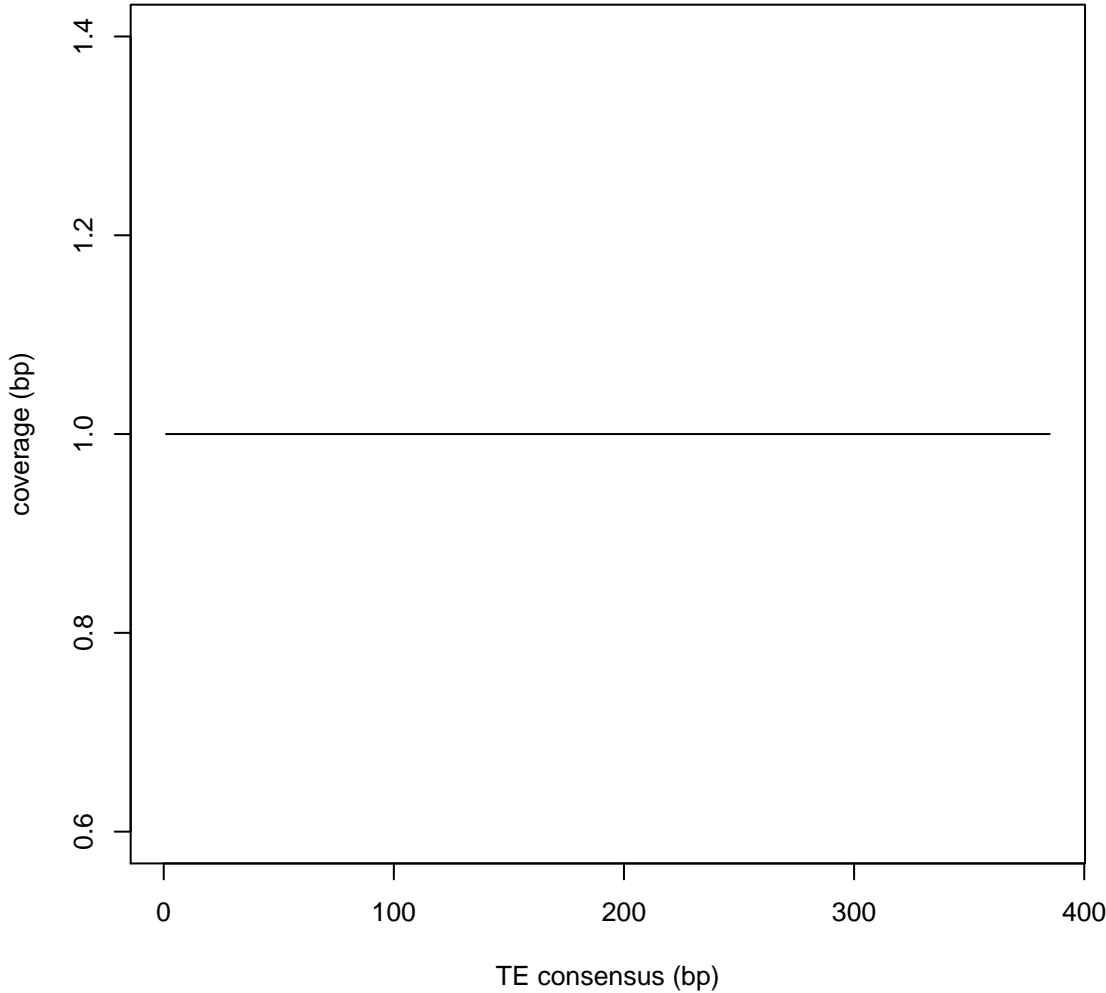


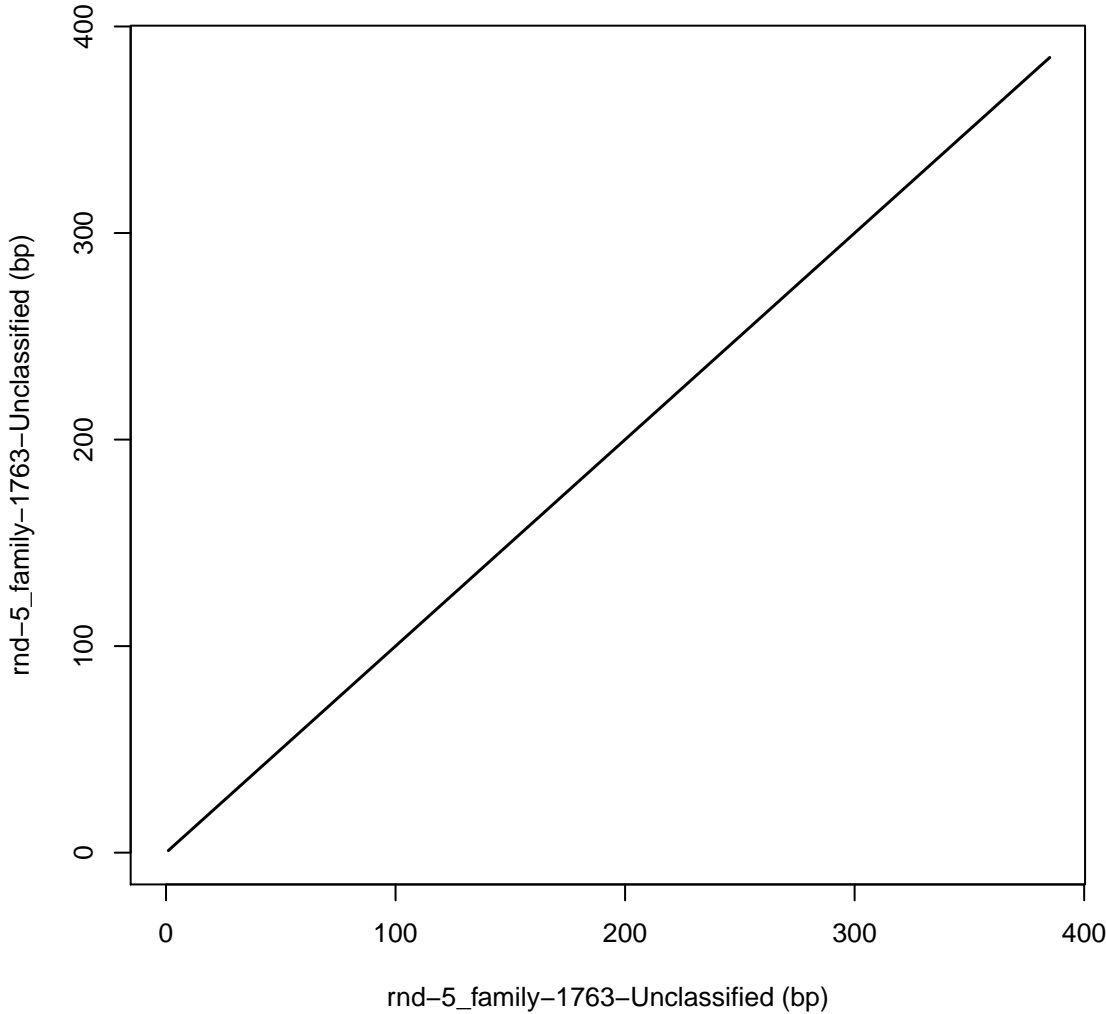
TE: rnd-5_family-1763-Unclassified
consensus size: 385bp; fragments: 1; full length: 1 (>=346.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

