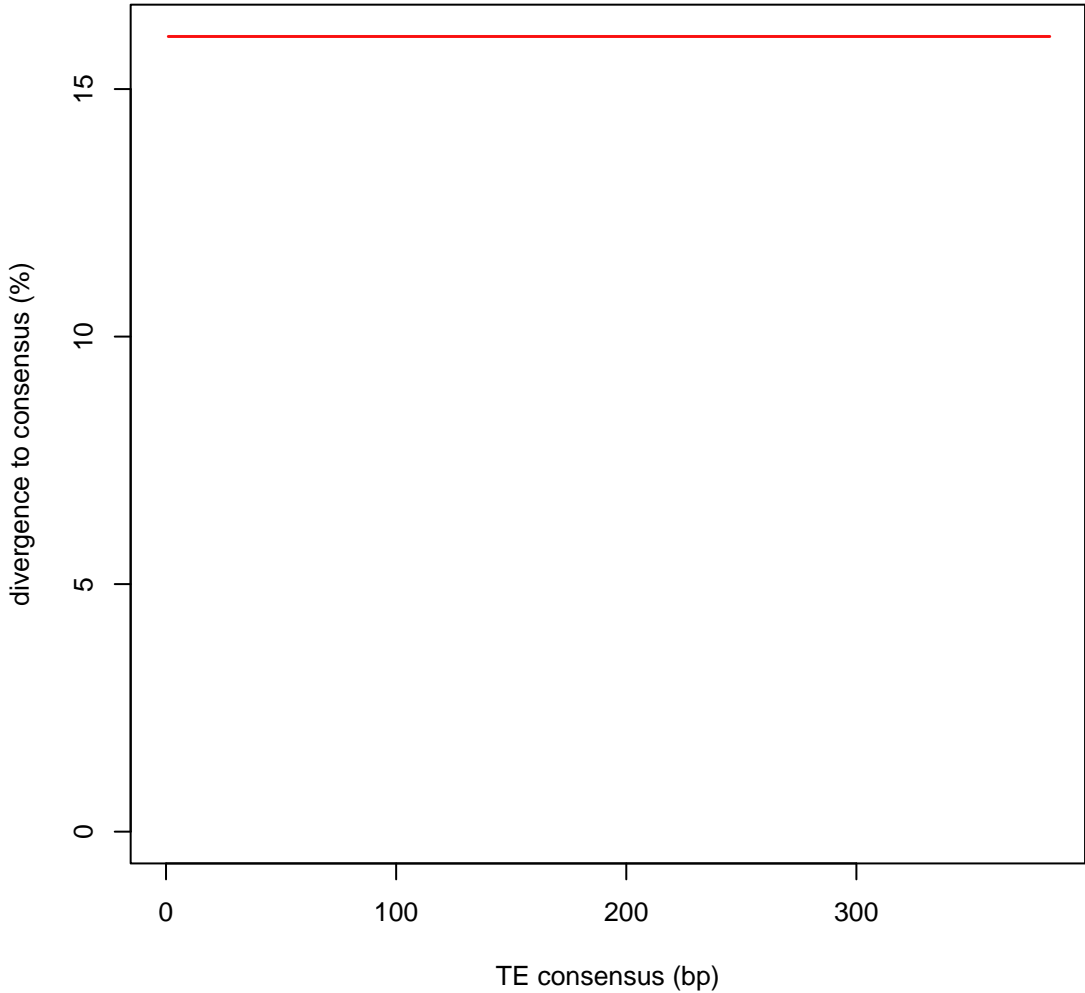
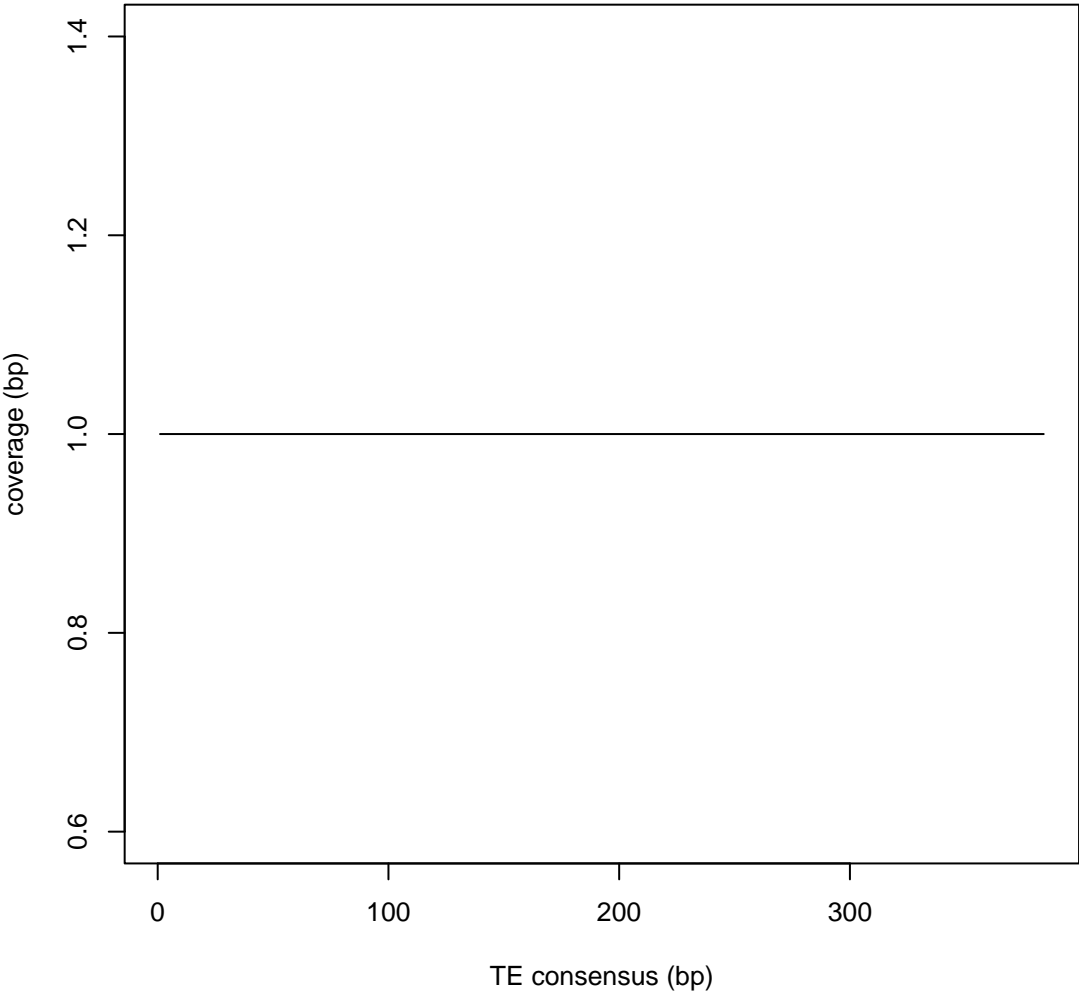


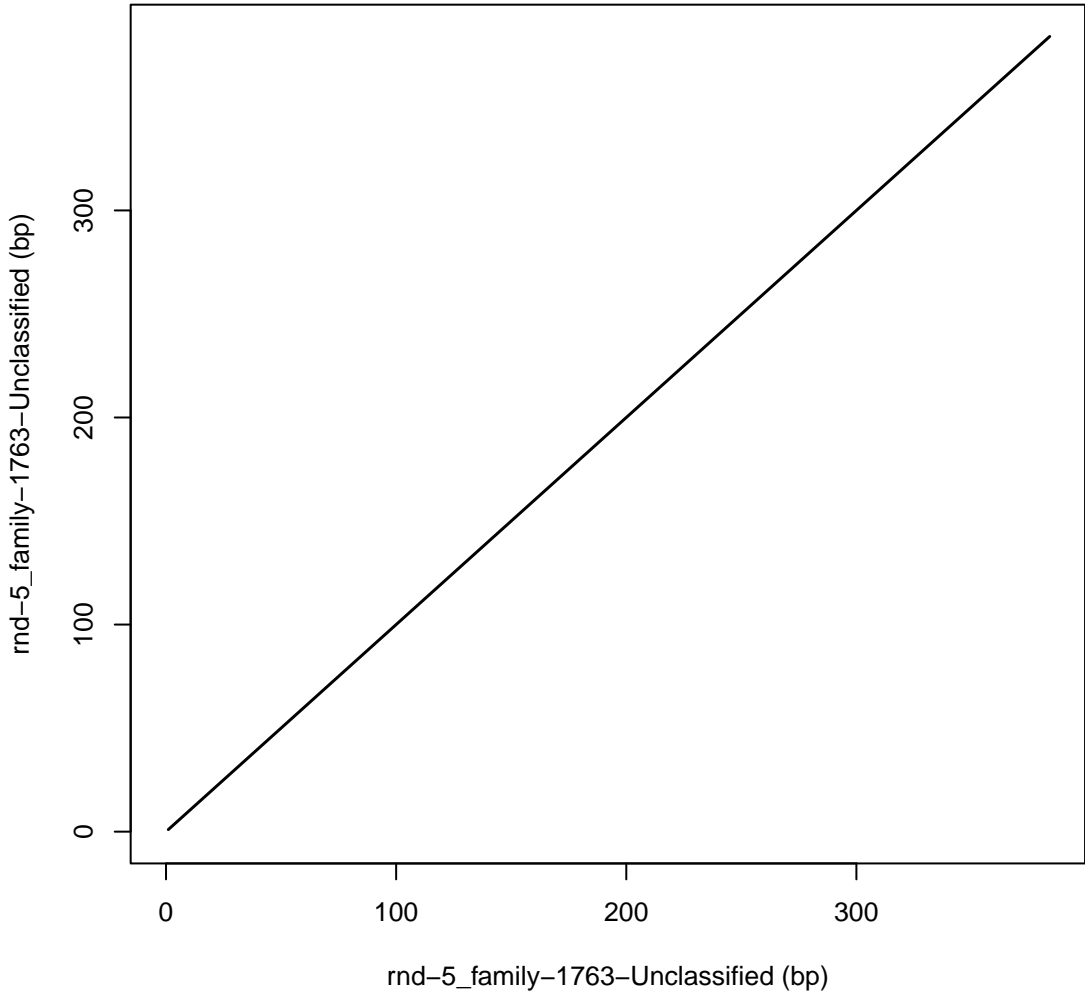
TE: rnd-5\_family-1763-Unclassified  
consensus size: 384bp; fragments: 1; full length: 1 (>=345.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

