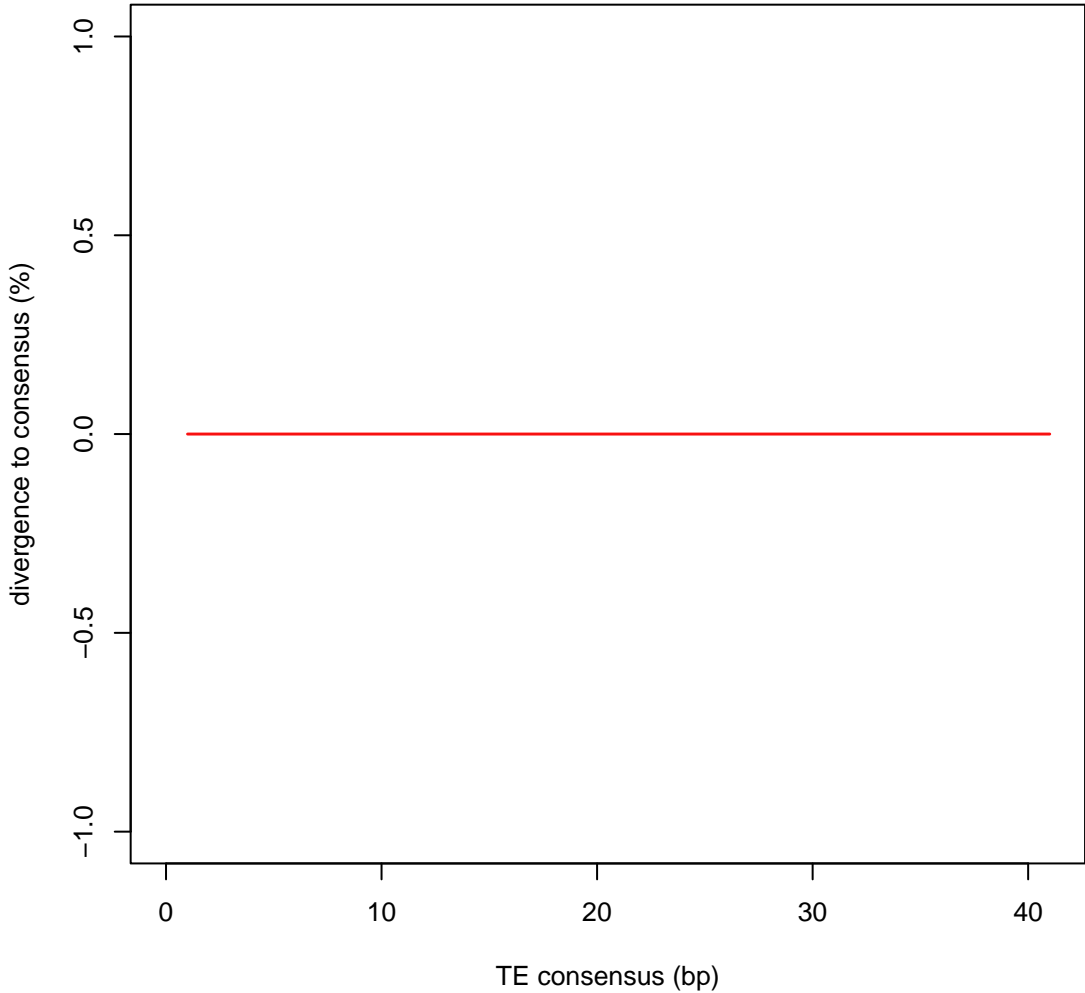
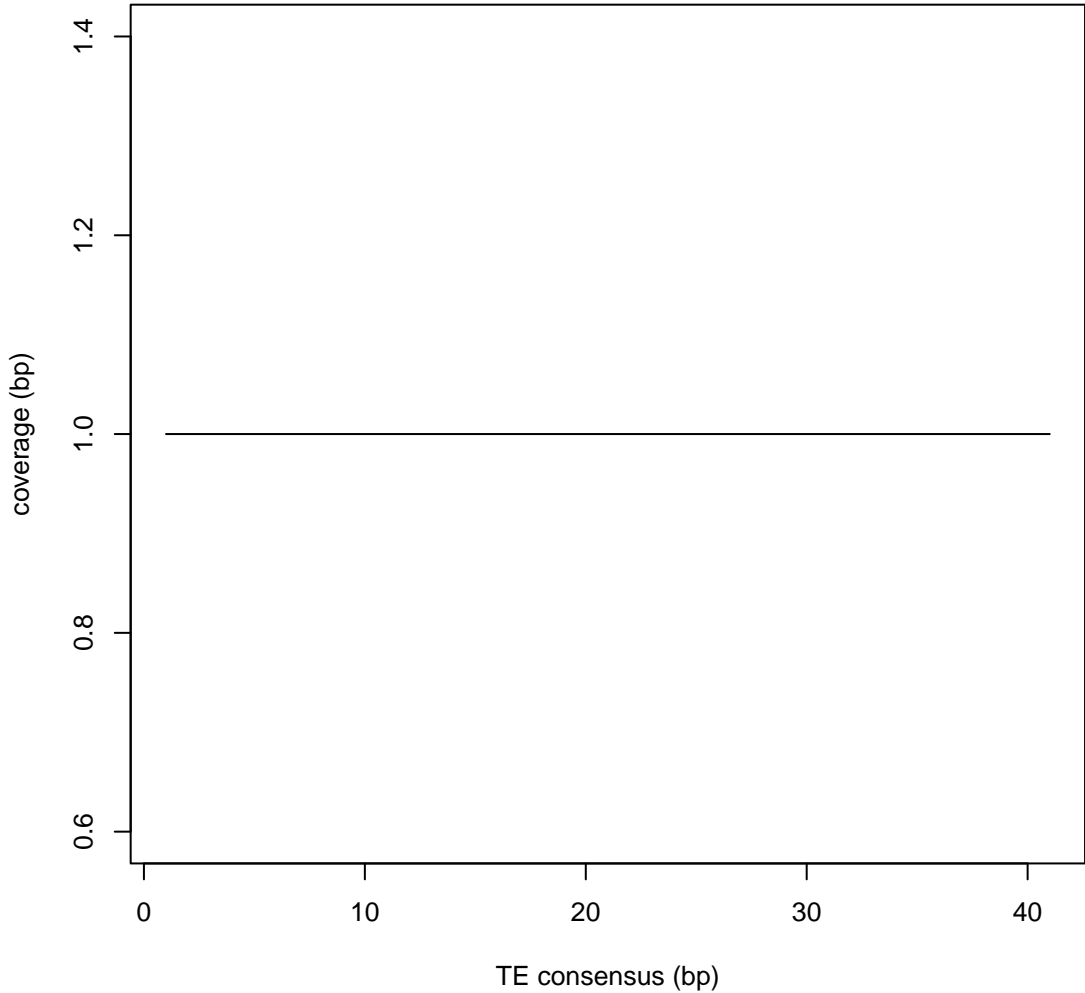


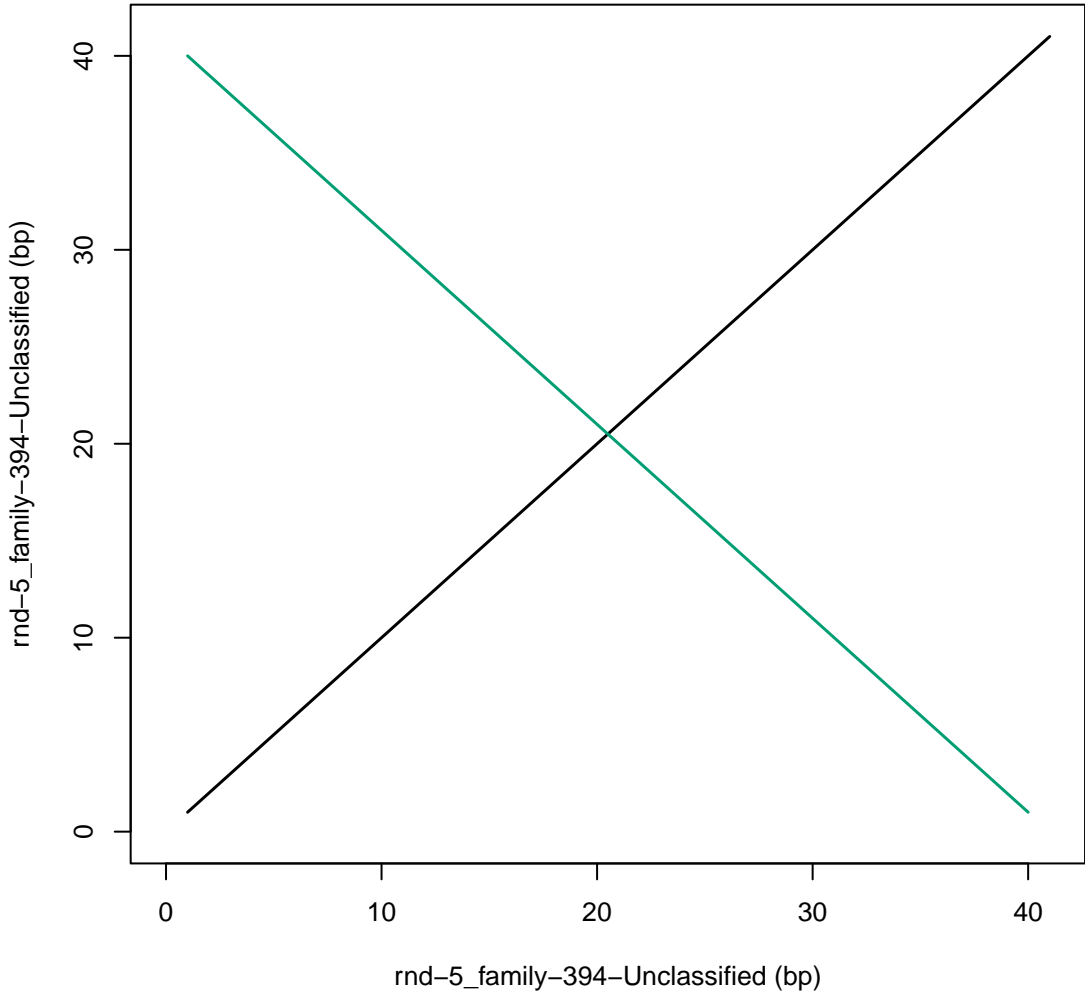
TE: rnd-5_family-394-Unclassified
consensus size: 41bp; fragments: 1; full length: 1 (>=36.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

