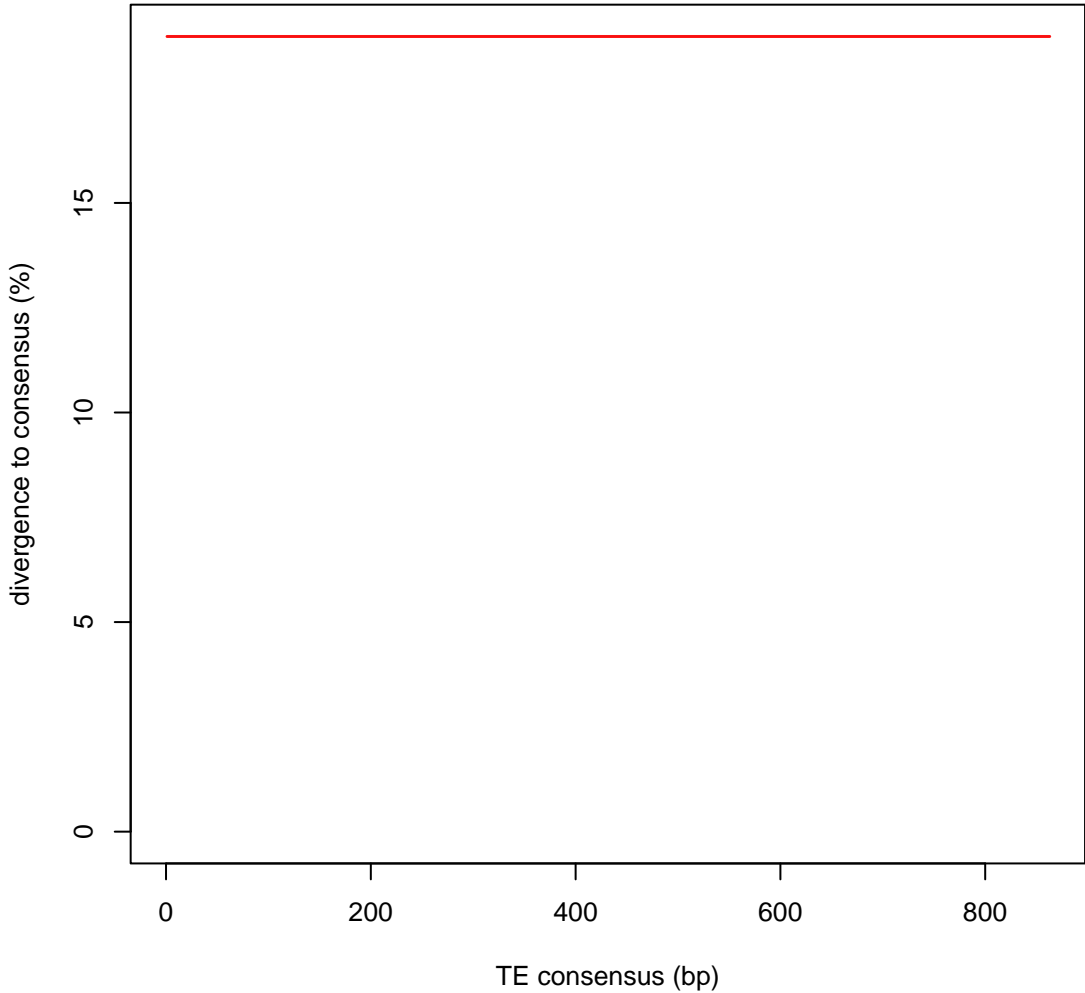
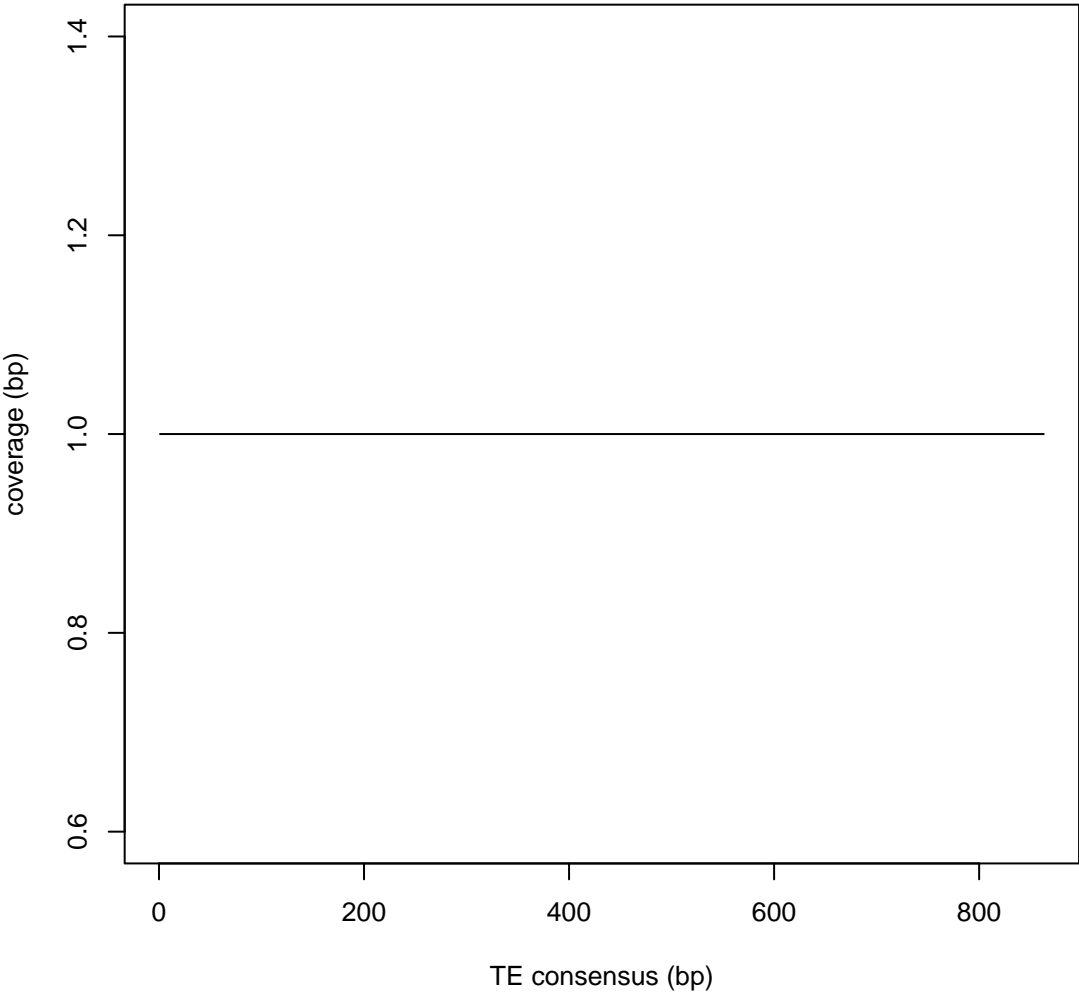


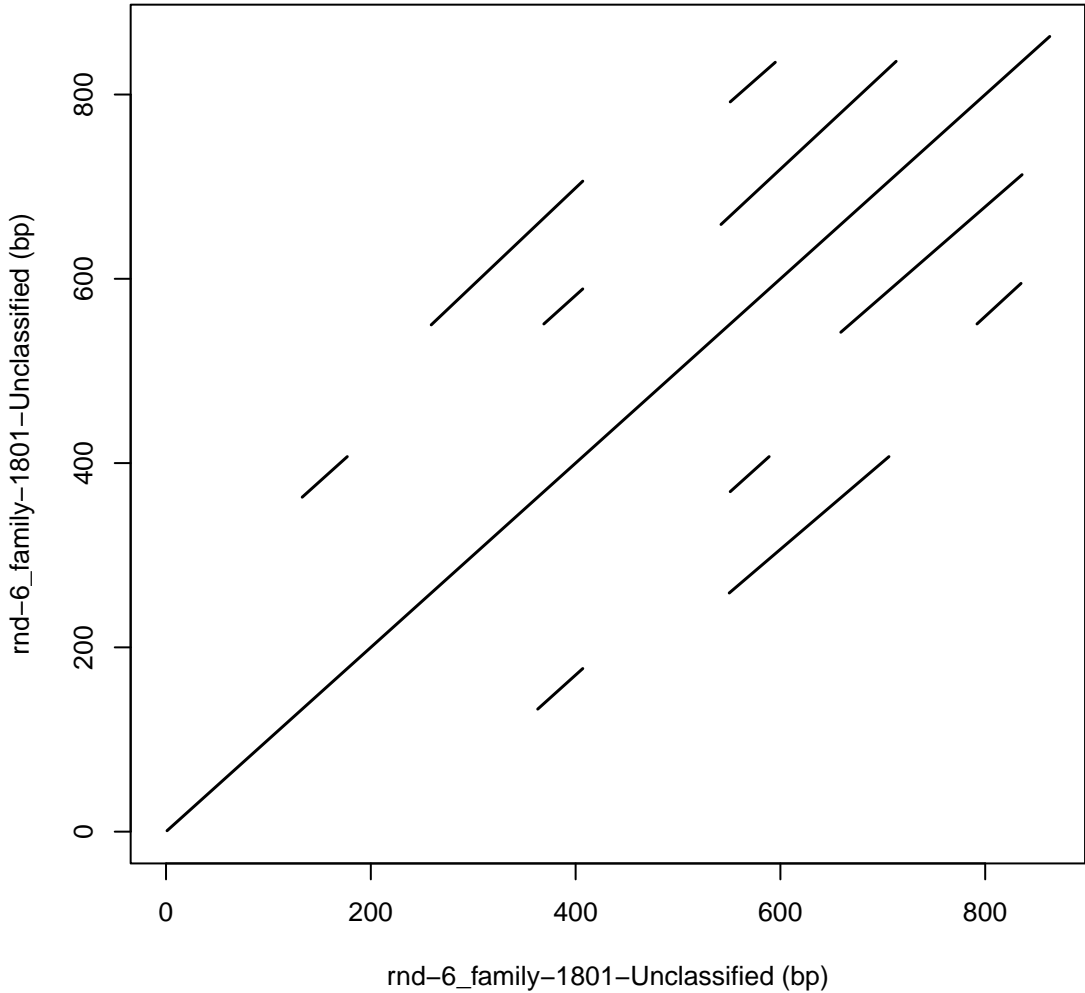
TE: rnd-6_family-1801-Unclassified
consensus size: 863bp; fragments: 1; full length: 1 (≥ 776.7 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

