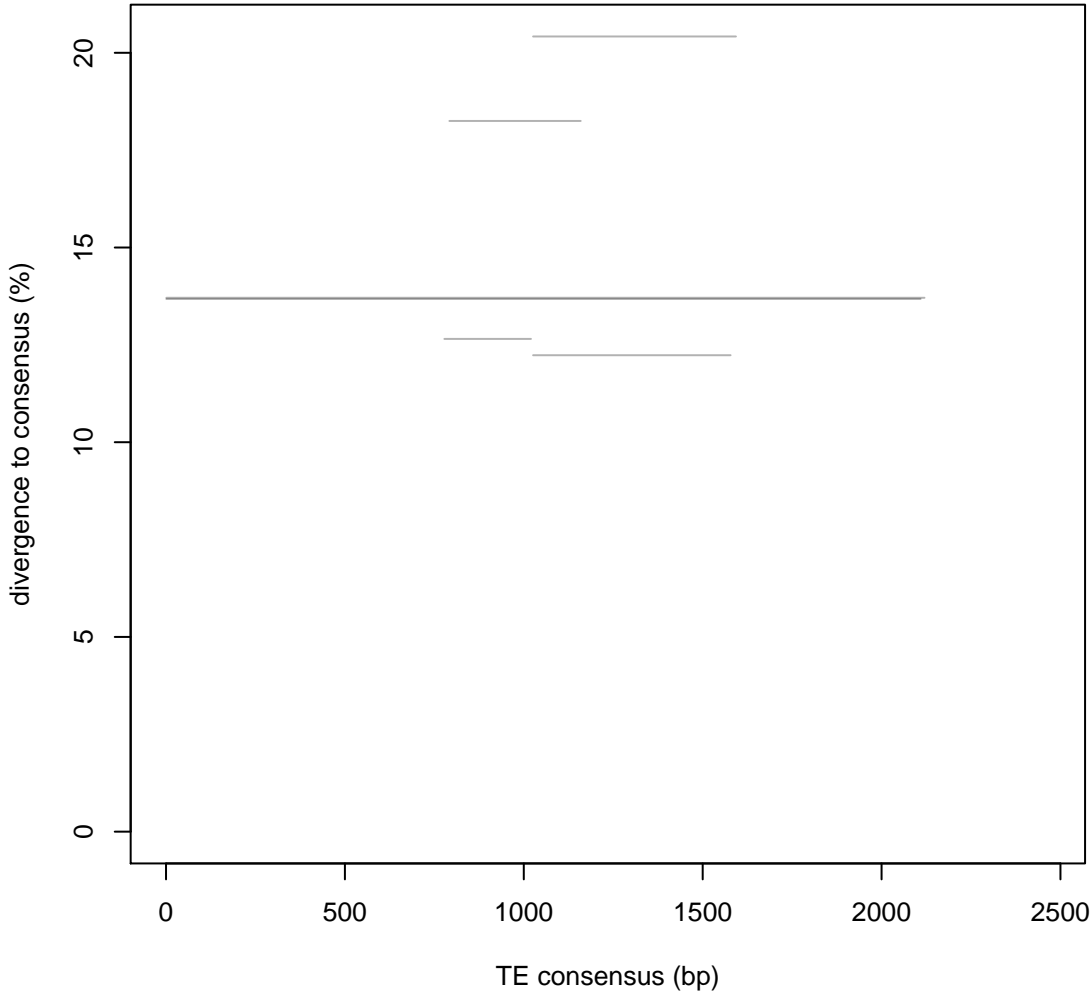
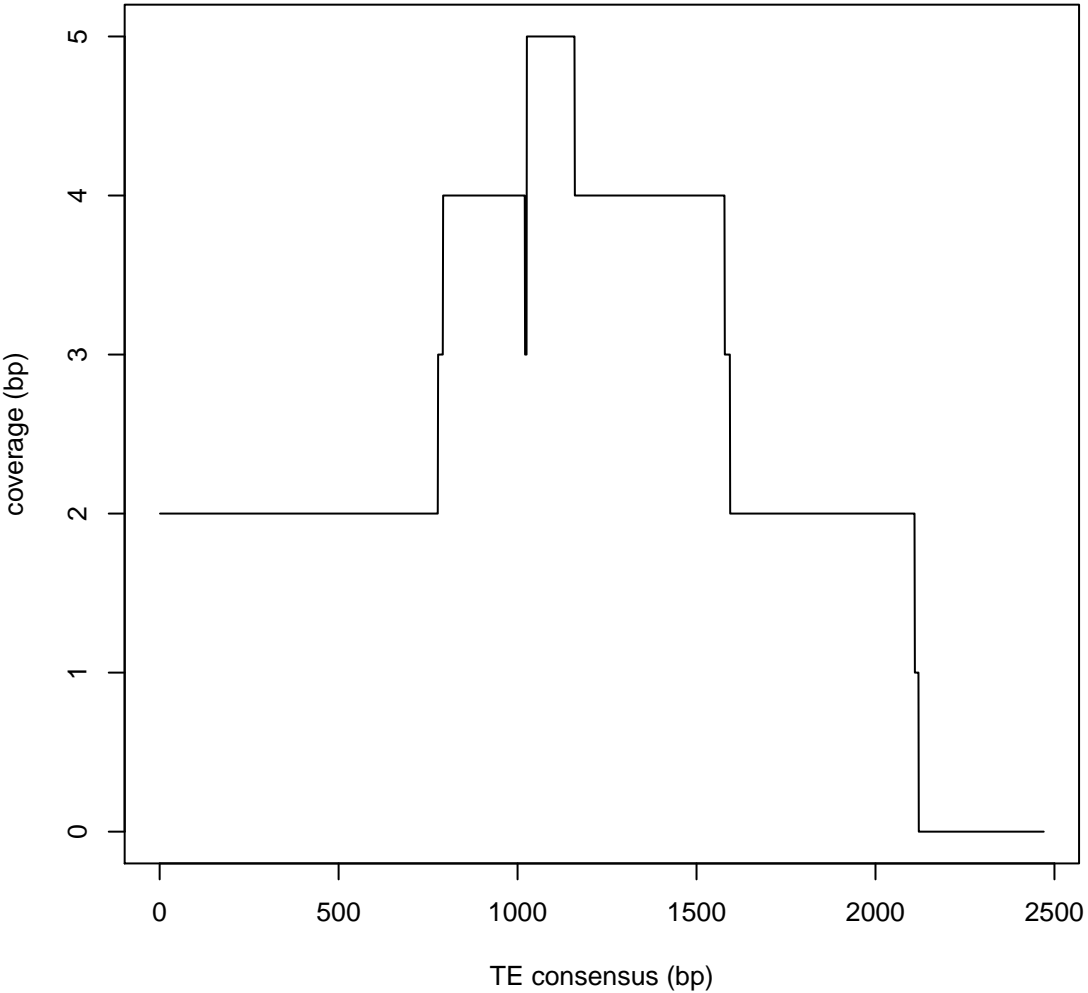


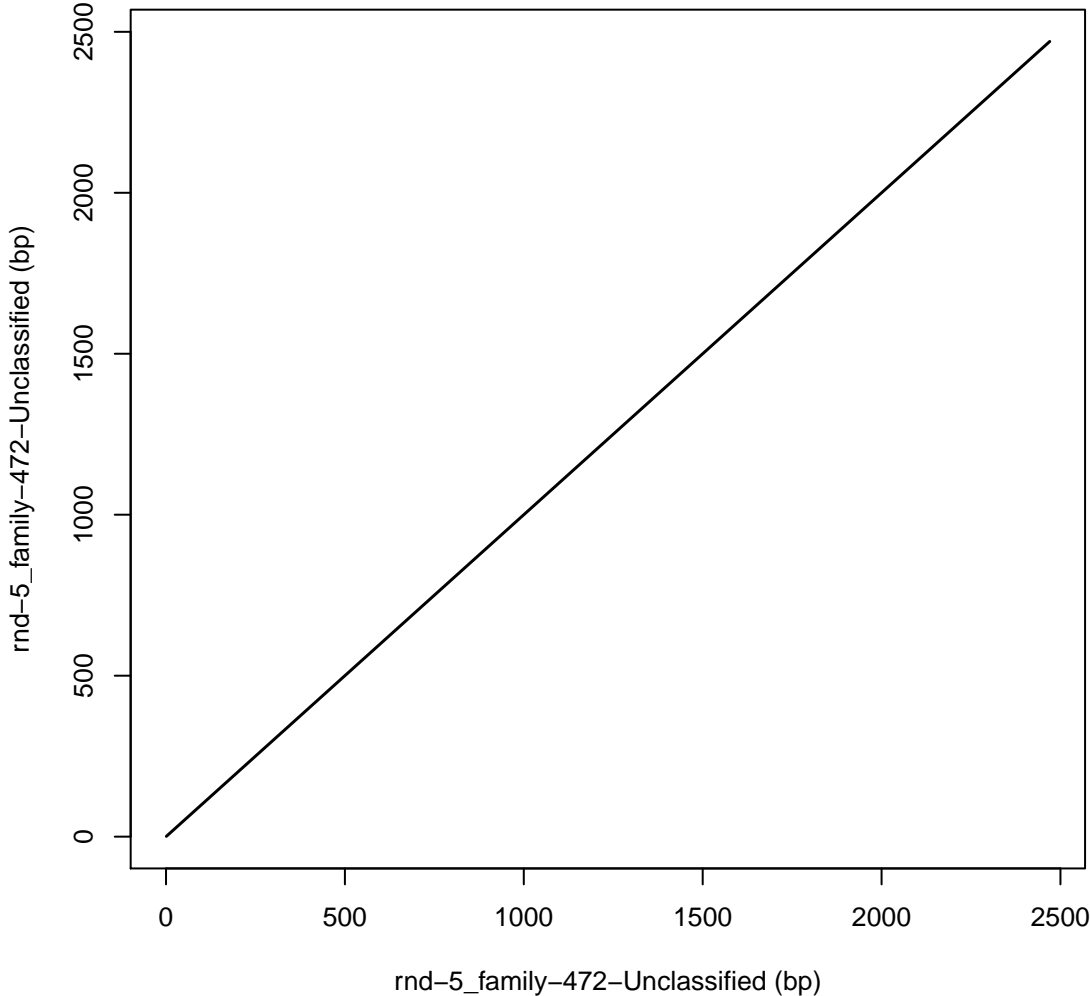
TE: rnd-5_family-472-Unclassified
consensus size: 2470bp; fragments: 6; full length: 0 (>=2223bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

