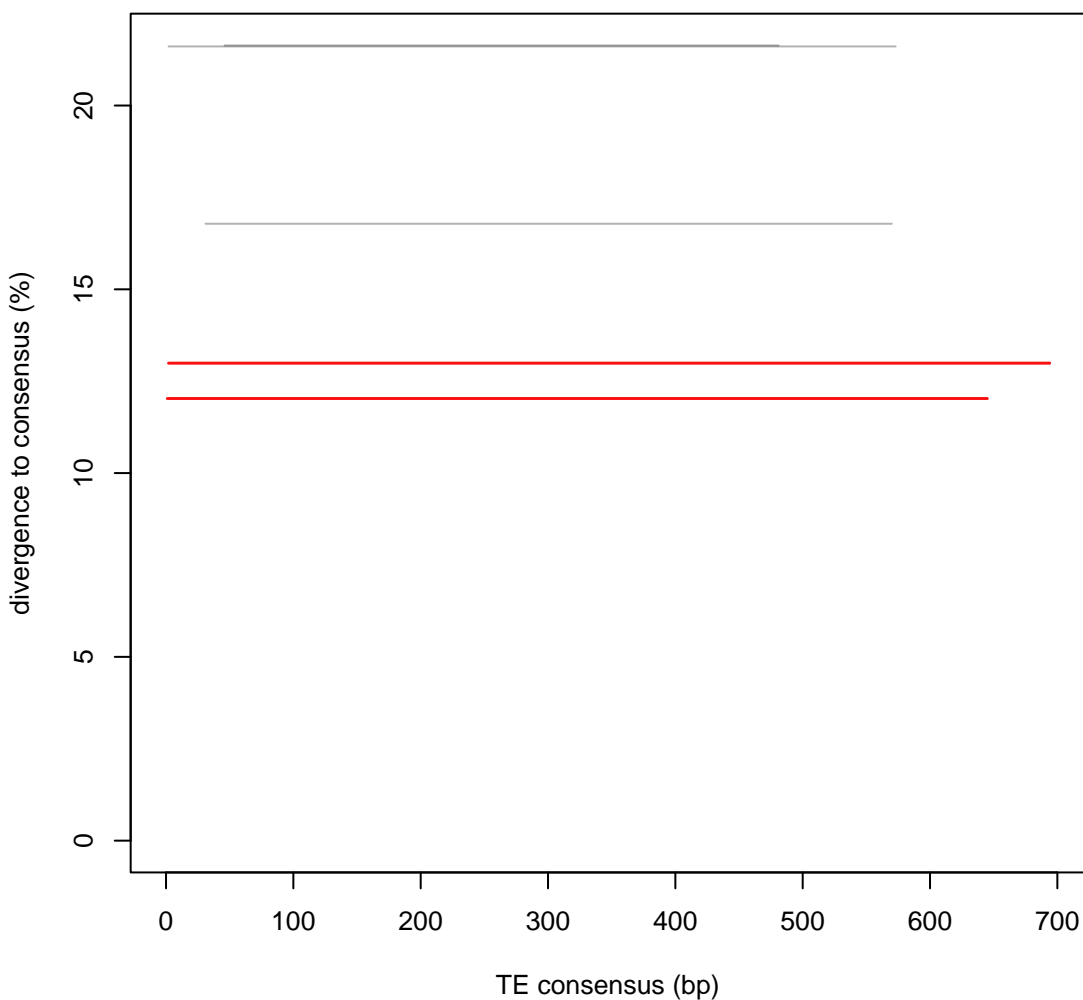
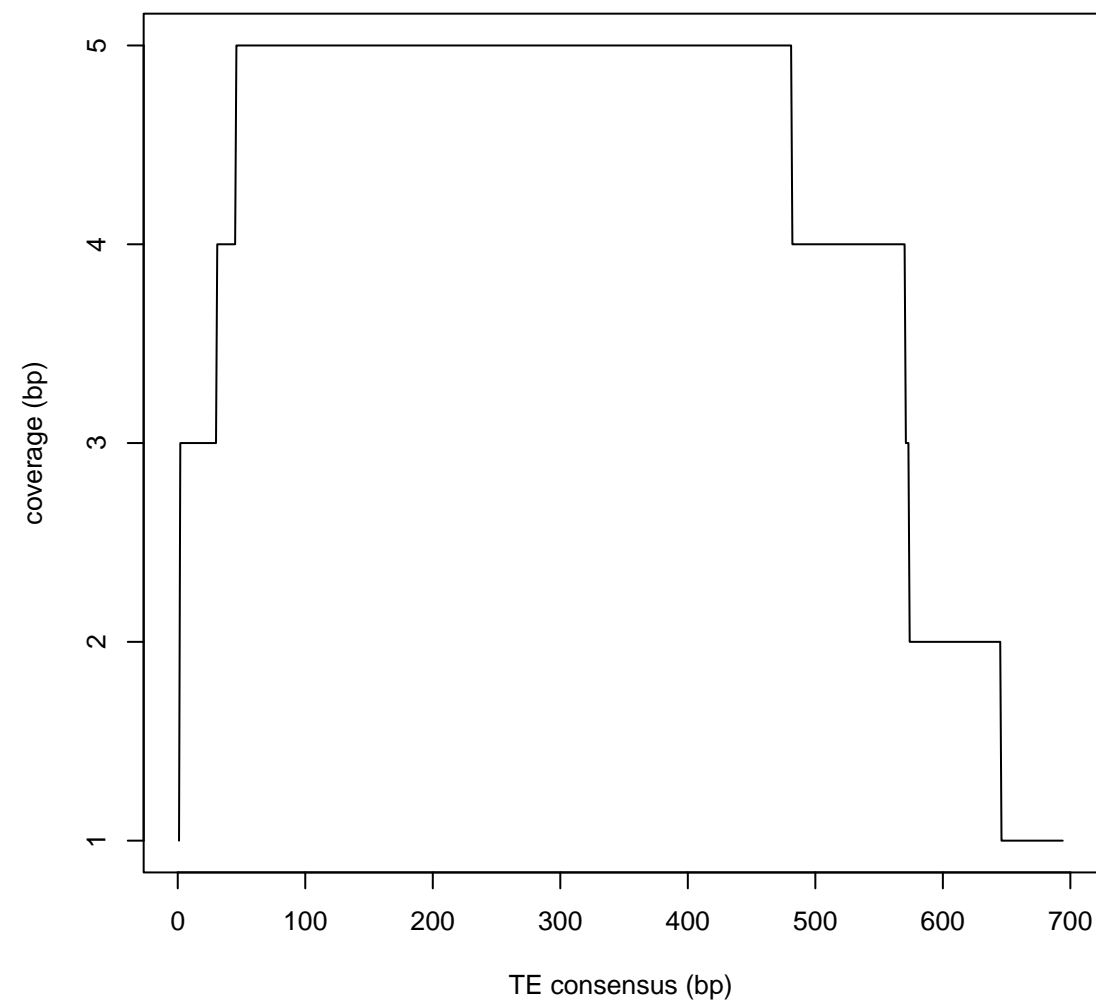


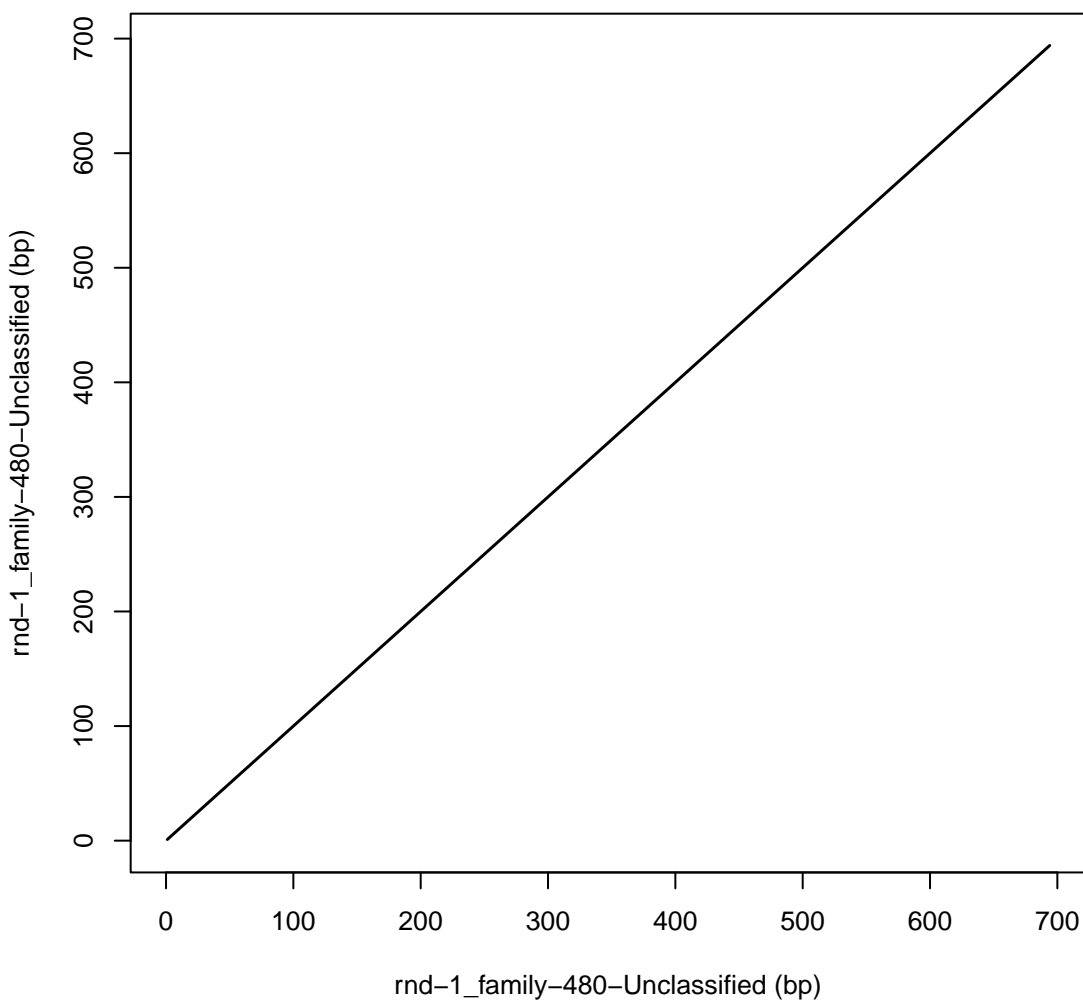
TE: rnd-1_family-480-Unclassified
consensus size: 694bp; fragments: 5; full length: 2 (≥ 624.6 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

