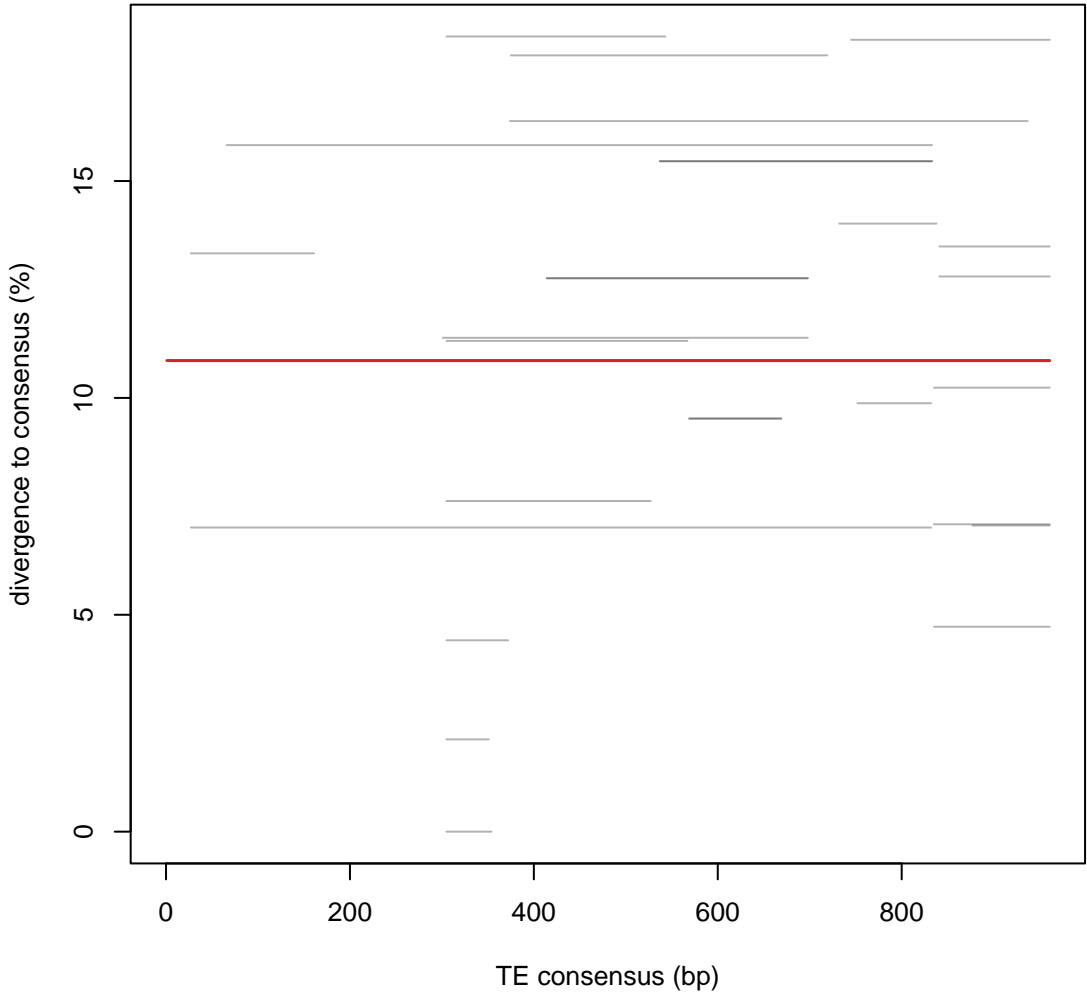
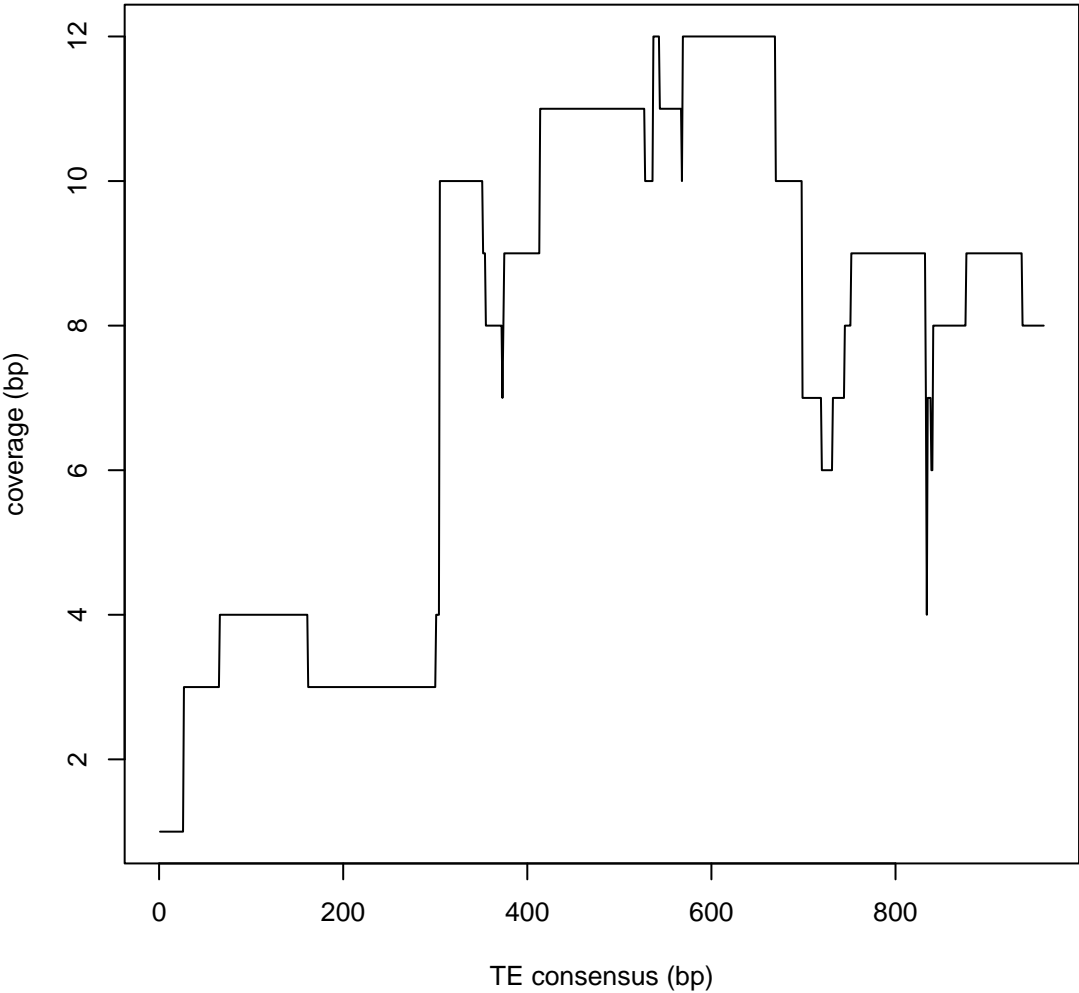


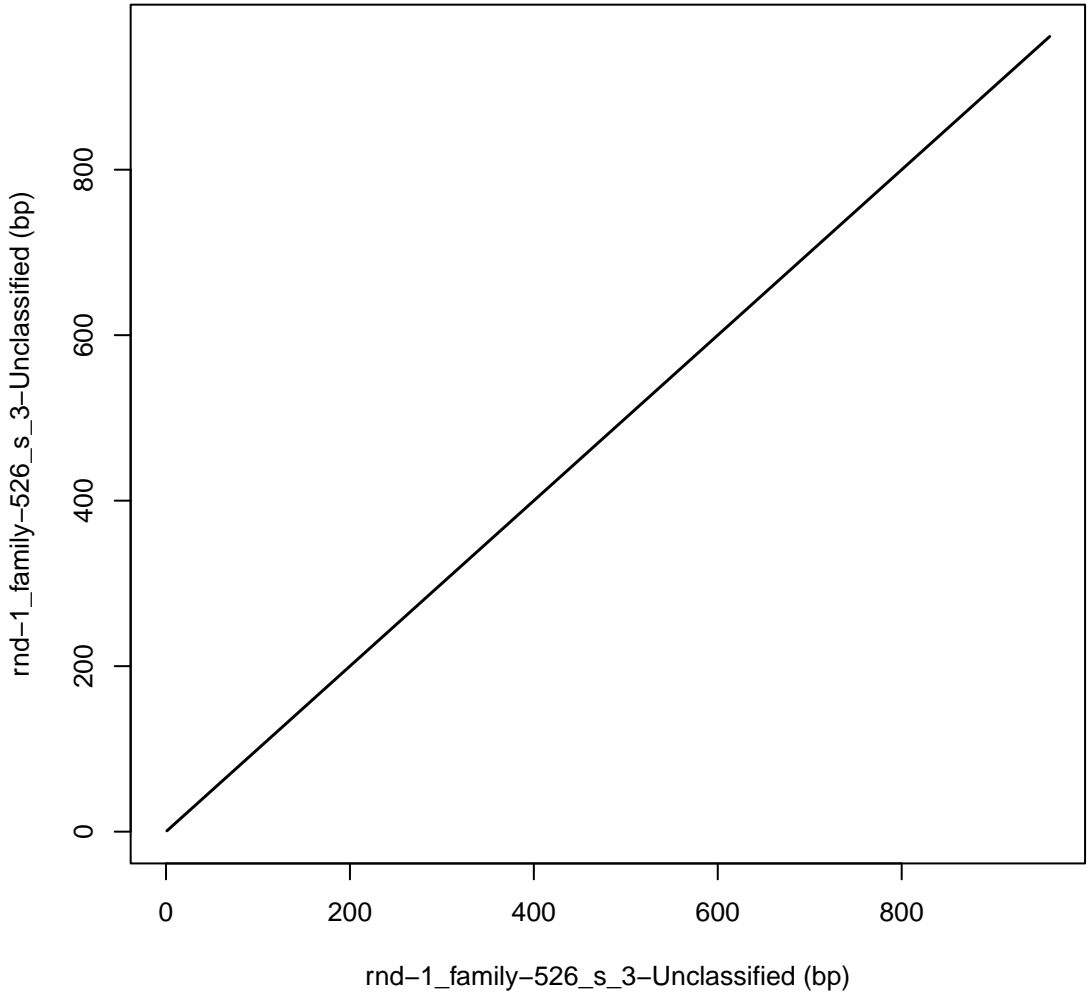
TE: rnd-1_family-526_s_3-Unclassified
consensus size: 961bp; fragments: 28; full length: 1 (>=864.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

