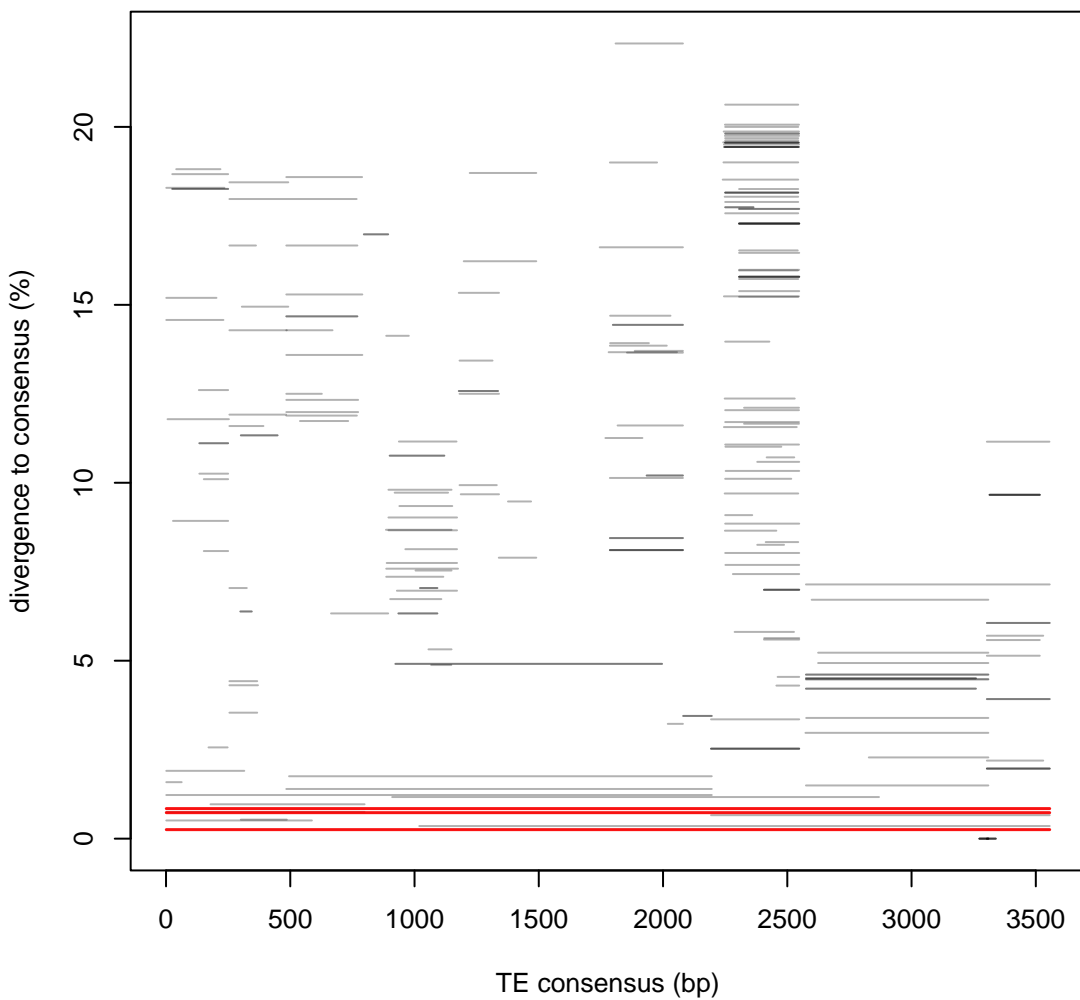
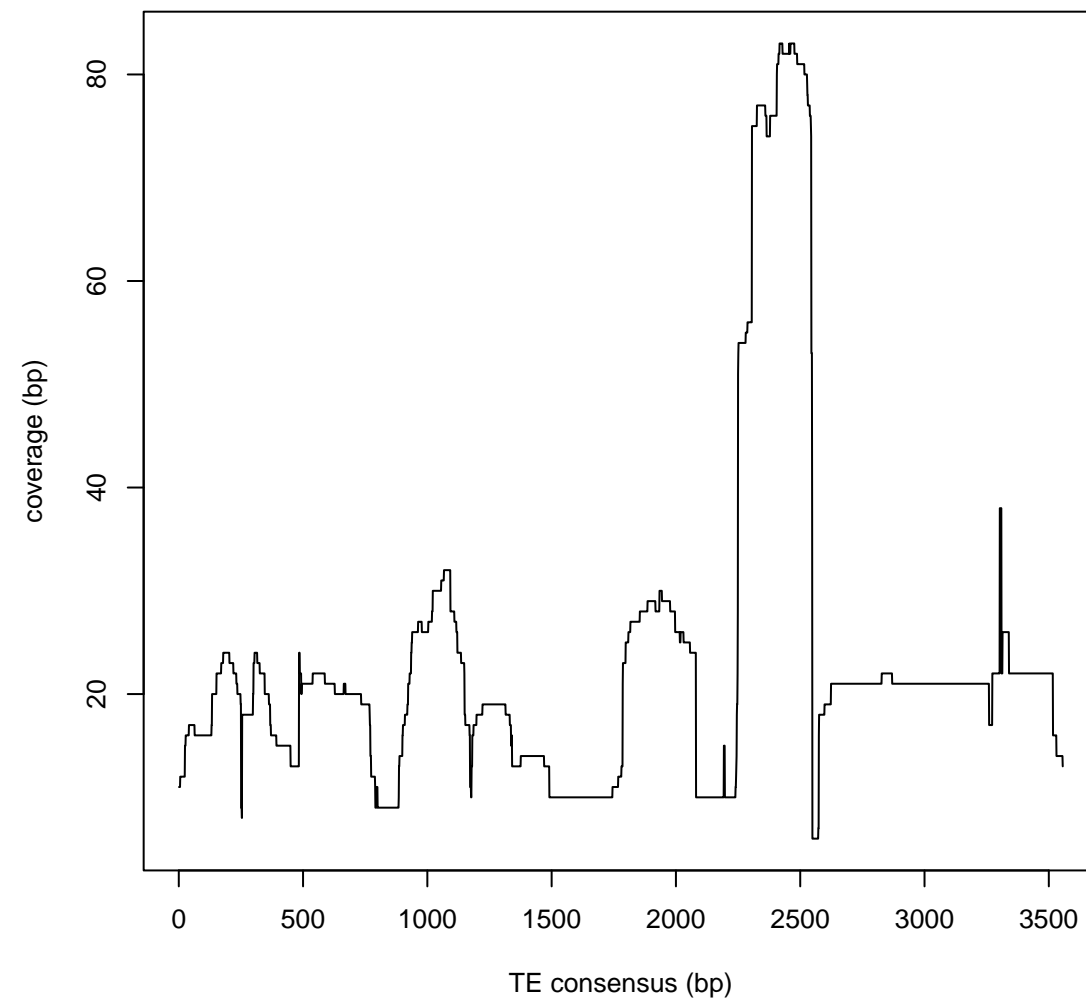


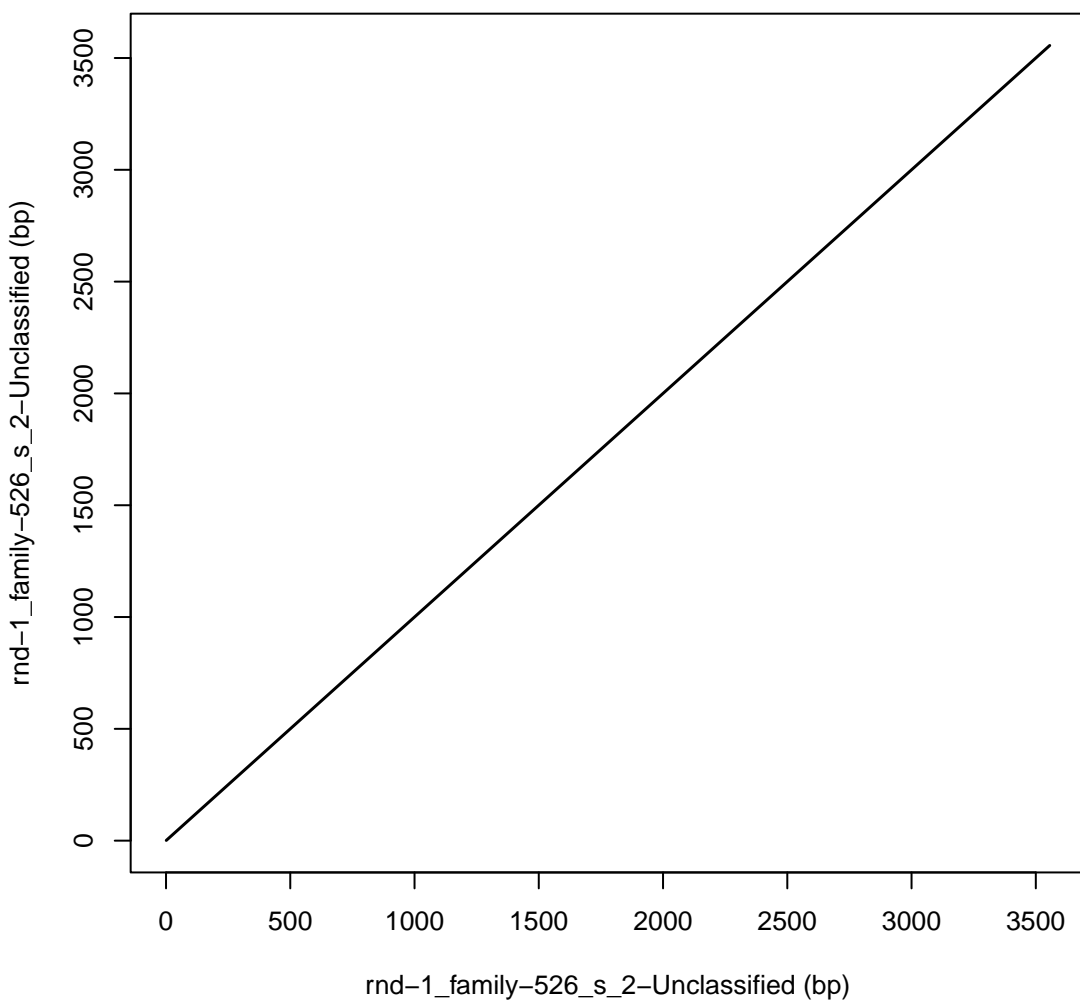
TE: rnd-1_family-526_s_2-Unclassified
consensus size: 3556bp; fragments: 244; full length: 3 (>=3200.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

