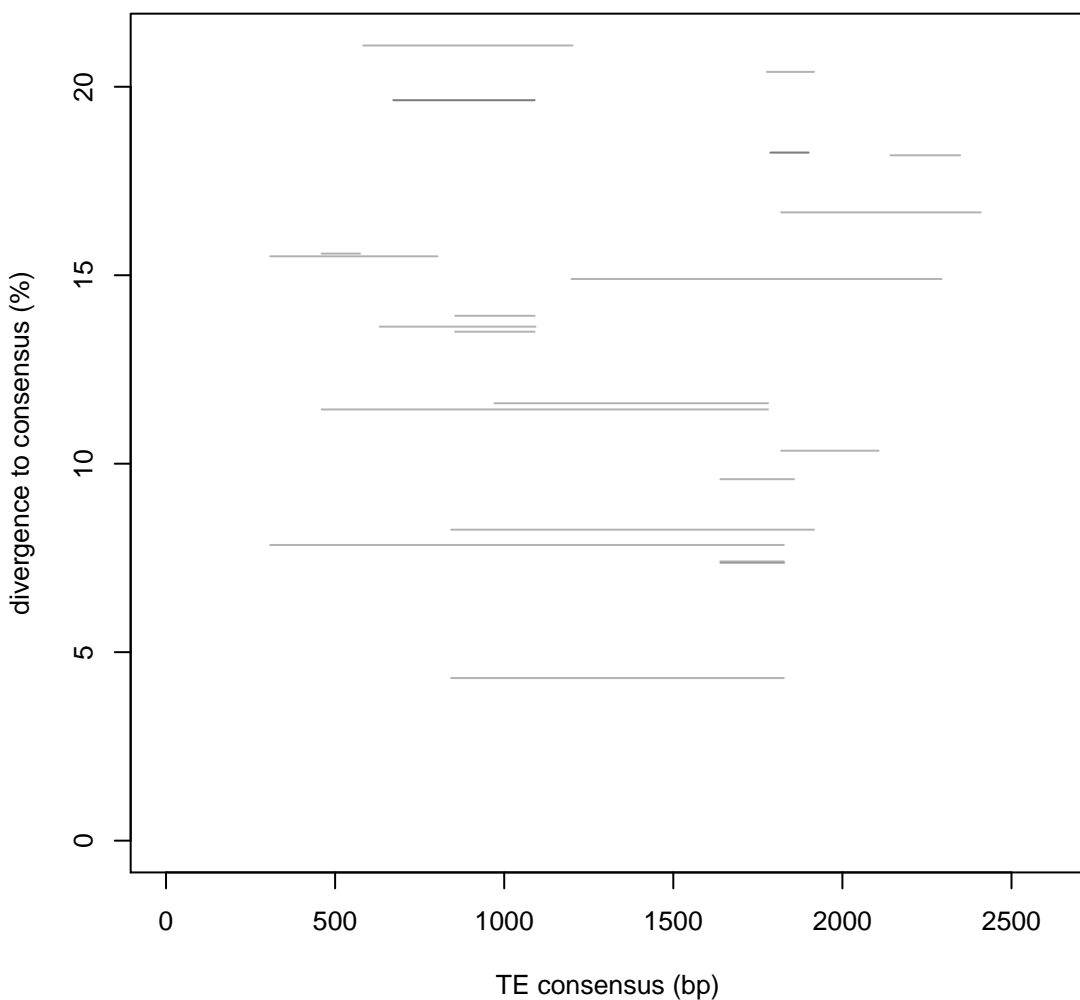
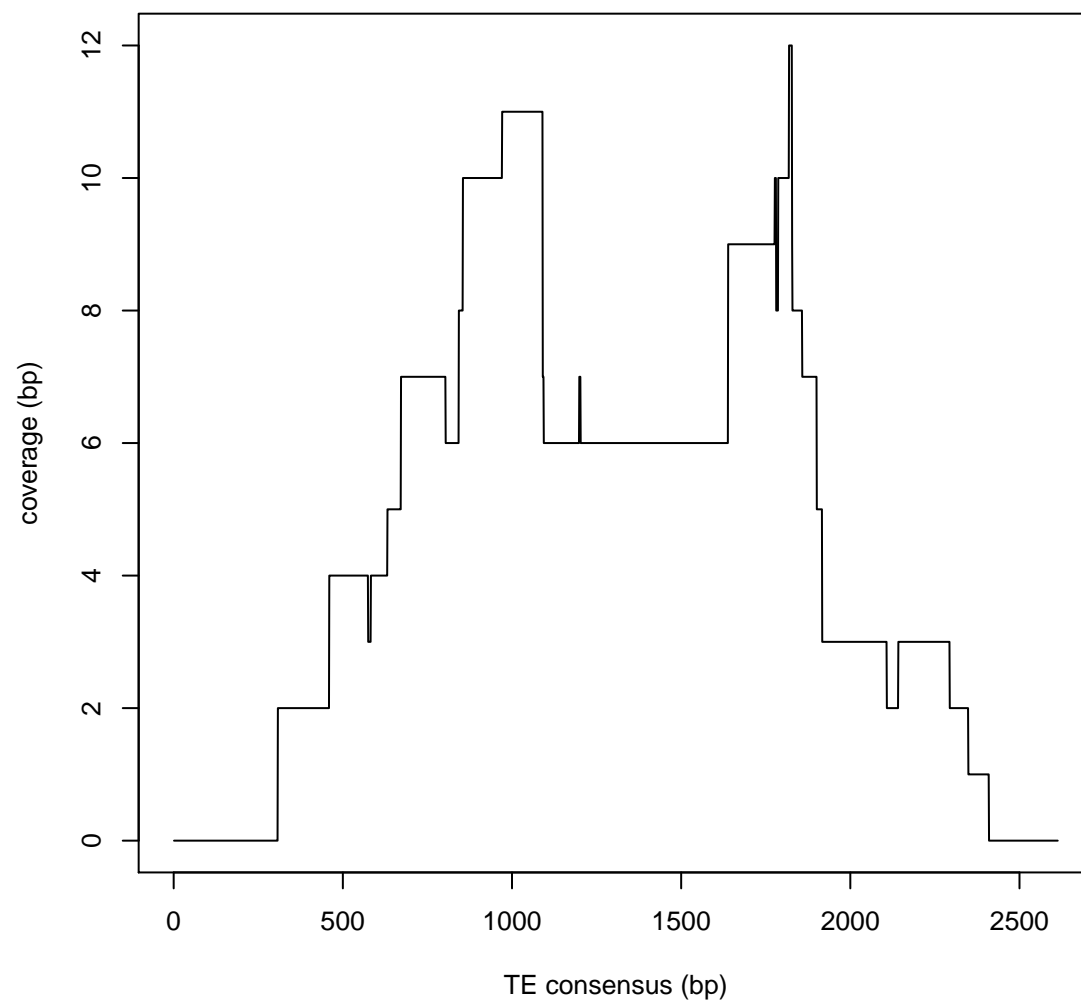


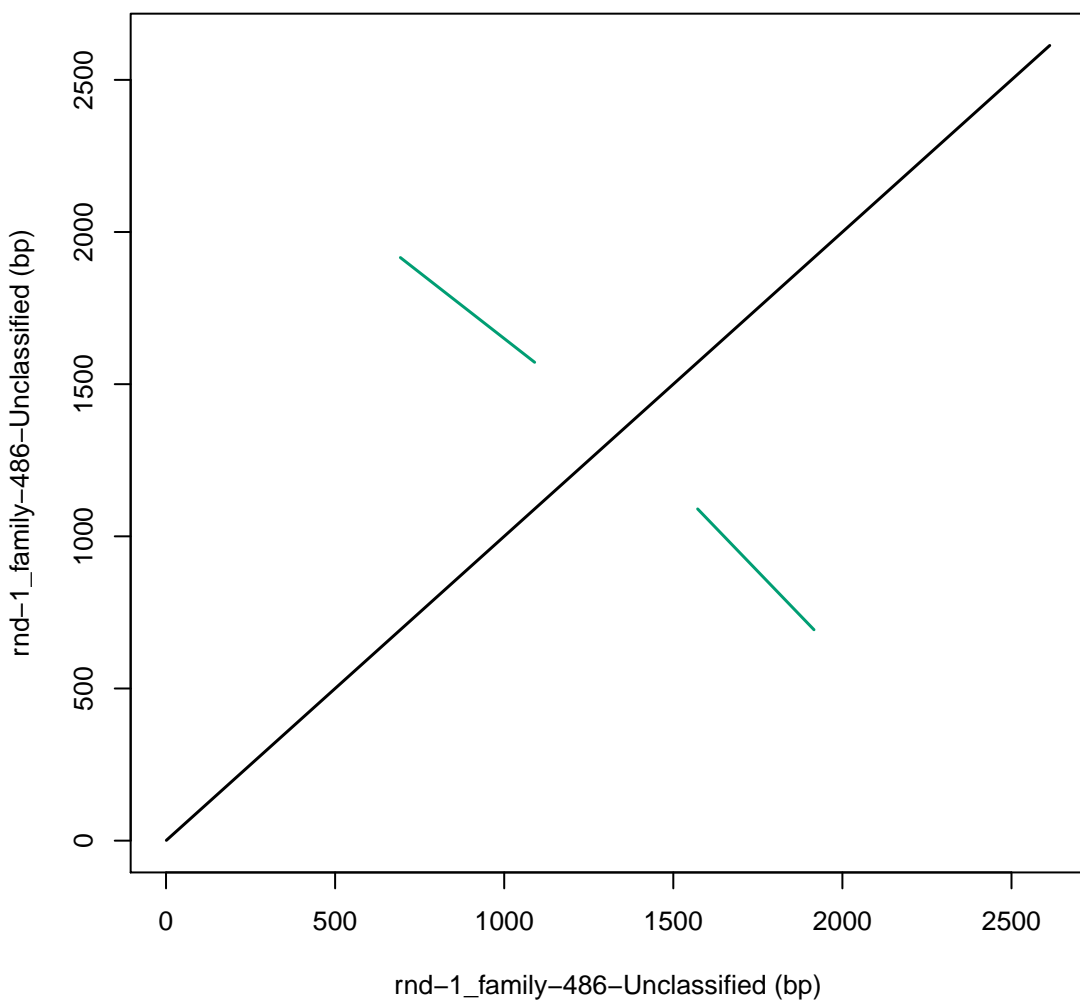
TE: rnd-1_family-486-Unclassified
consensus size: 2613bp; fragments: 23; full length: 0 (≥ 2351.7 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

