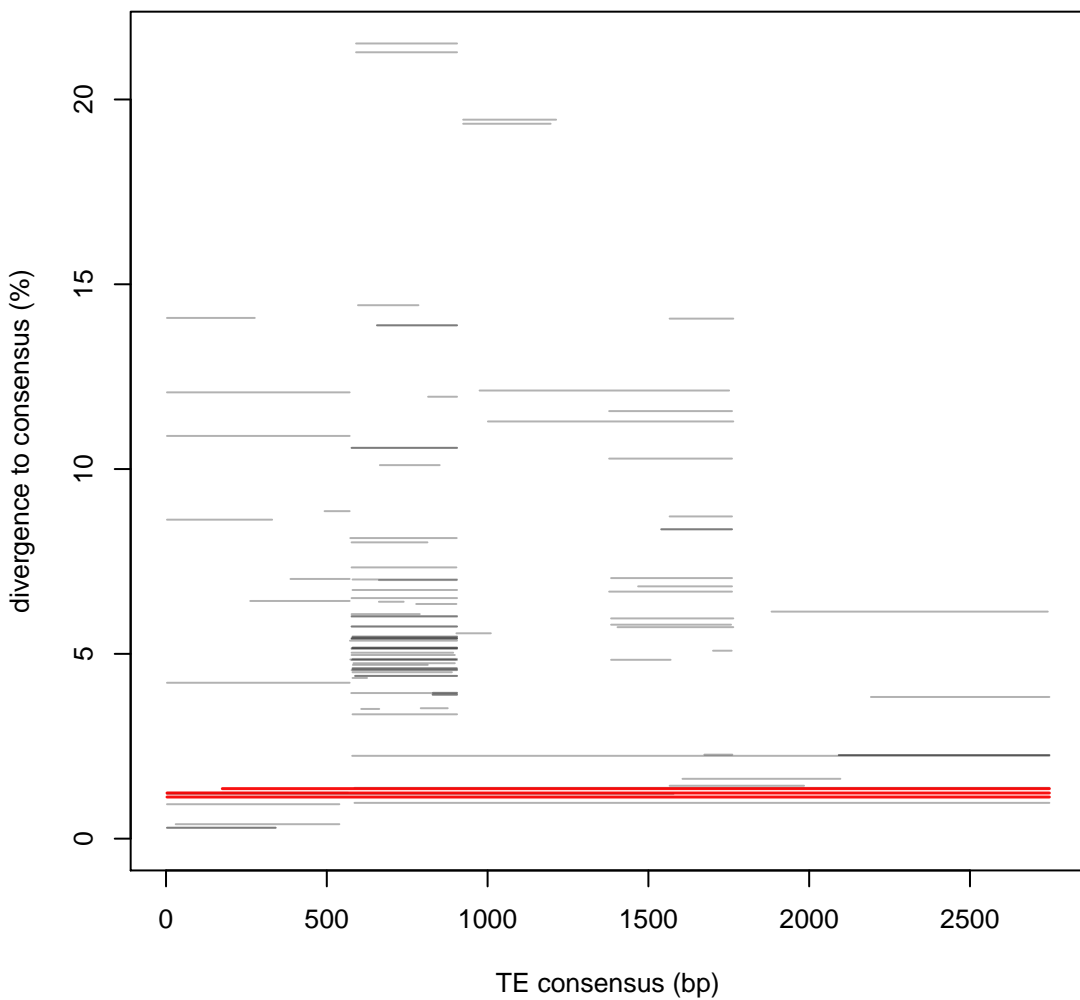
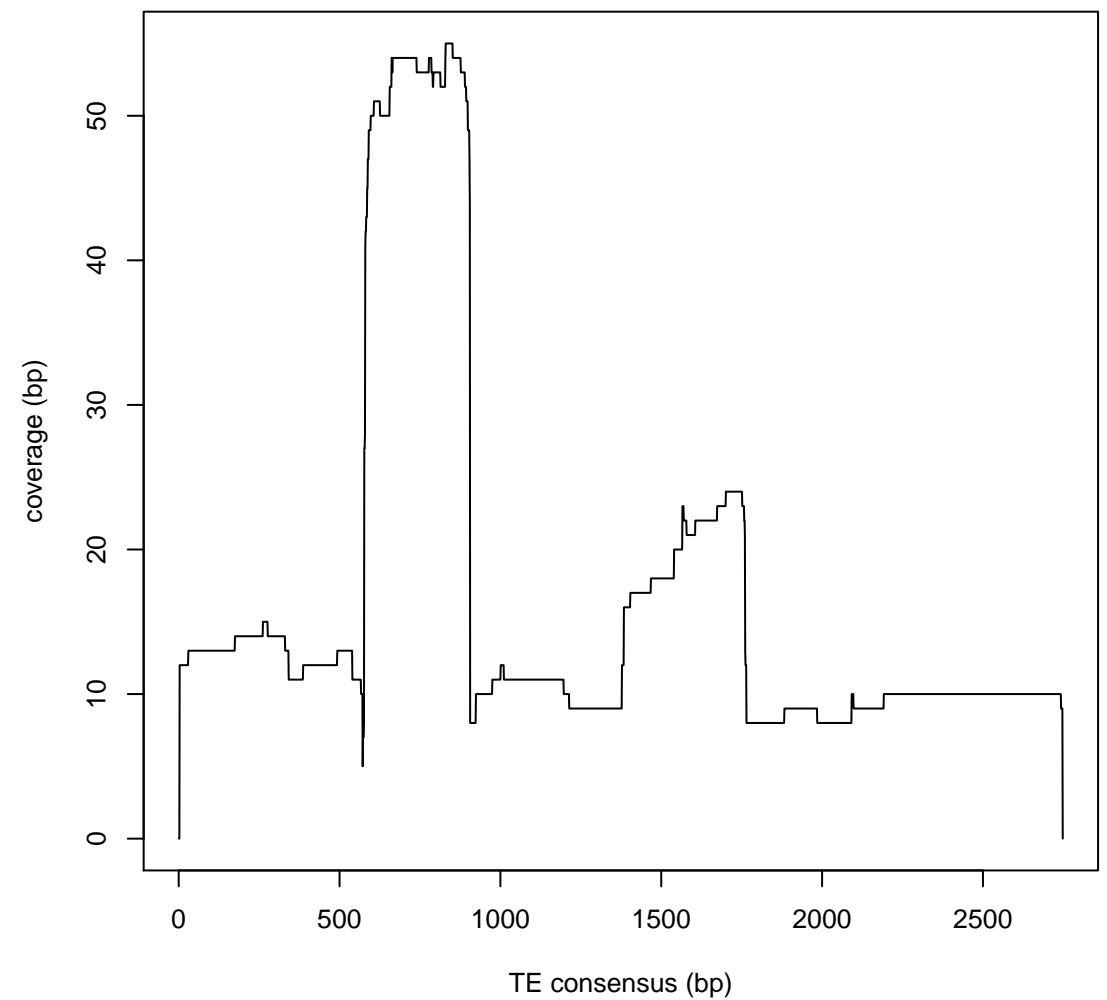


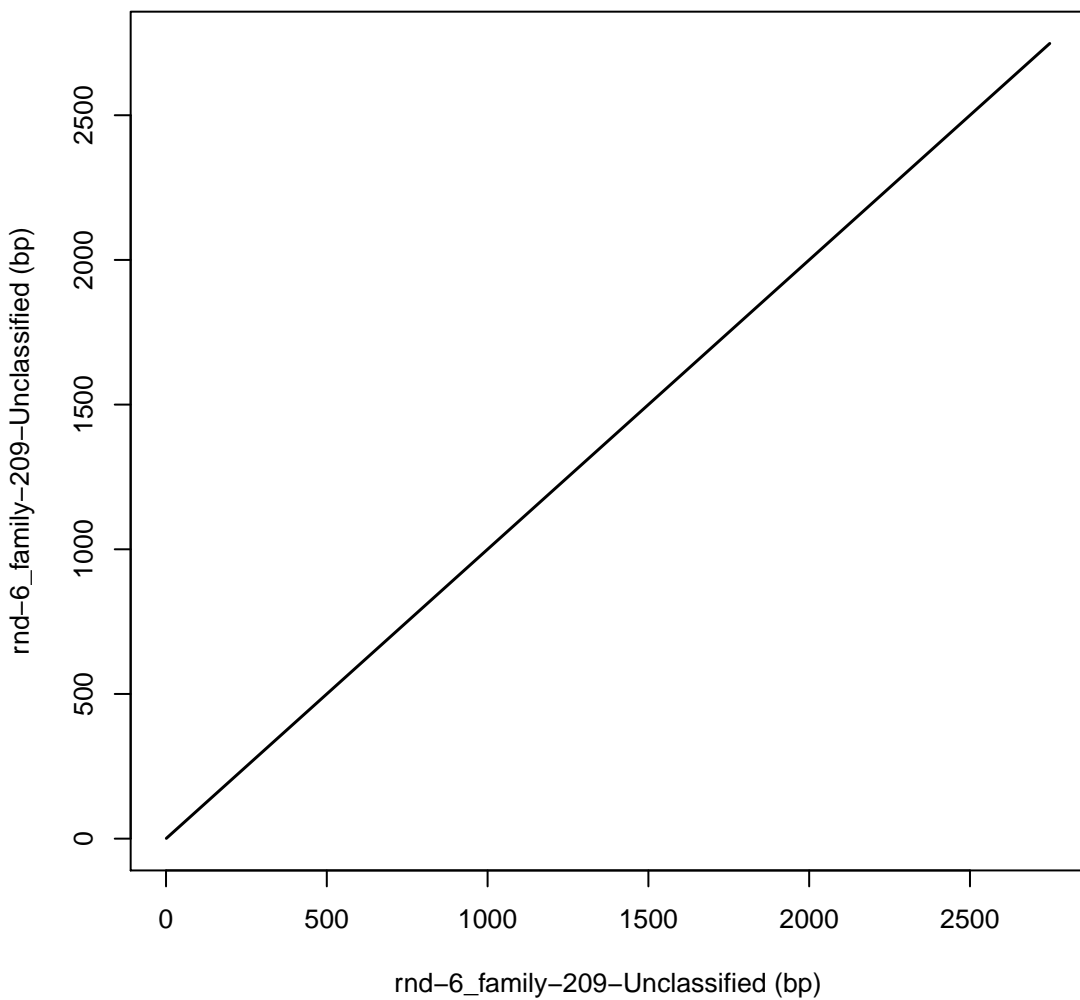
TE: rnd-6_family-209-Unclassified
consensus size: 2748bp; fragments: 101; full length: 3 (≥ 2473.2 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

