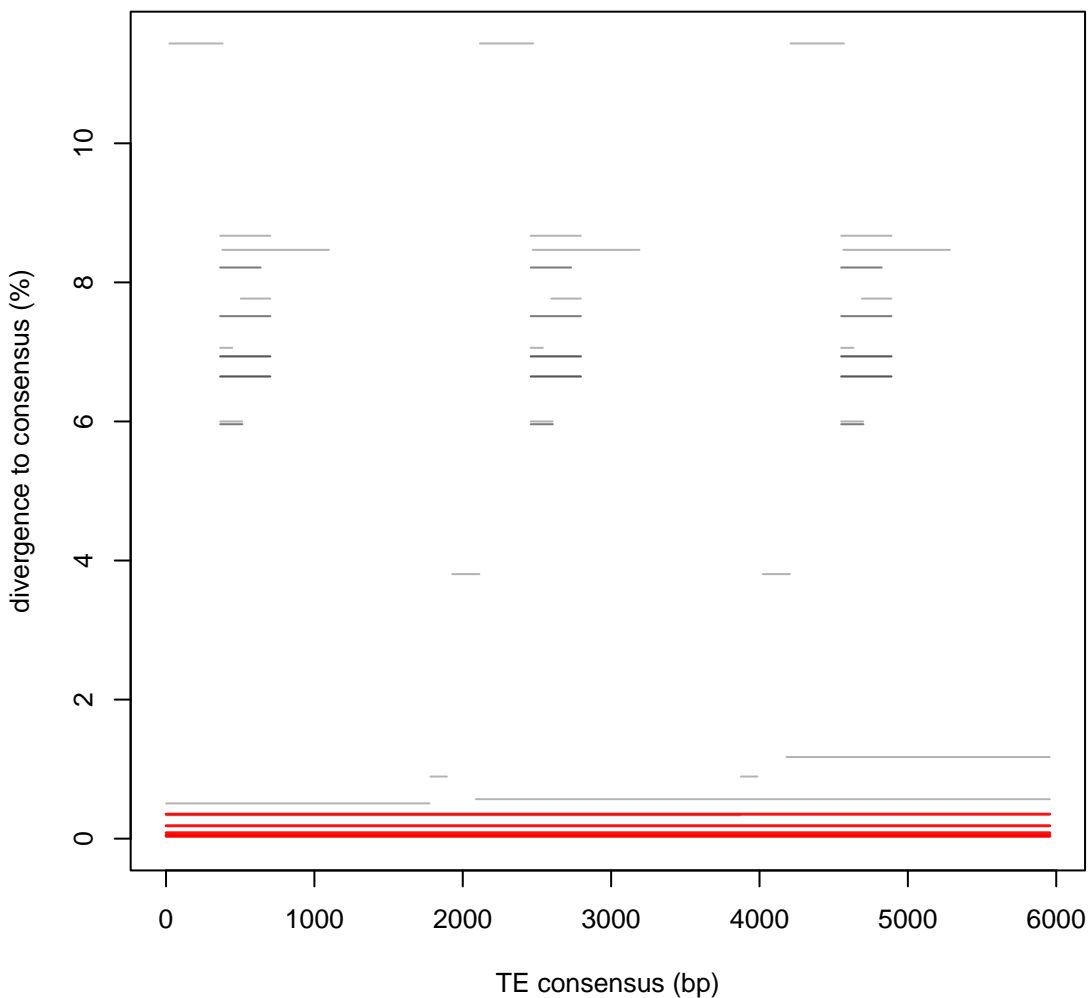
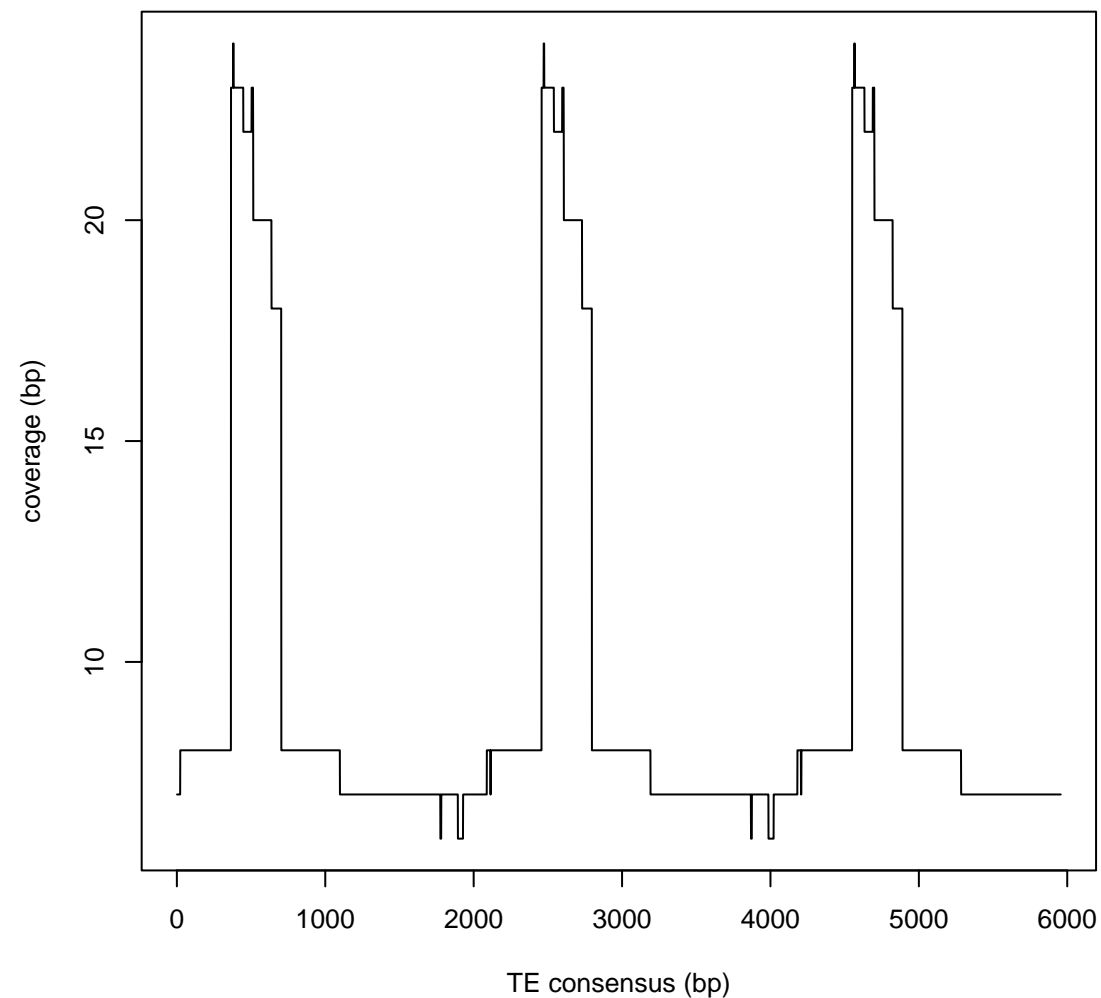


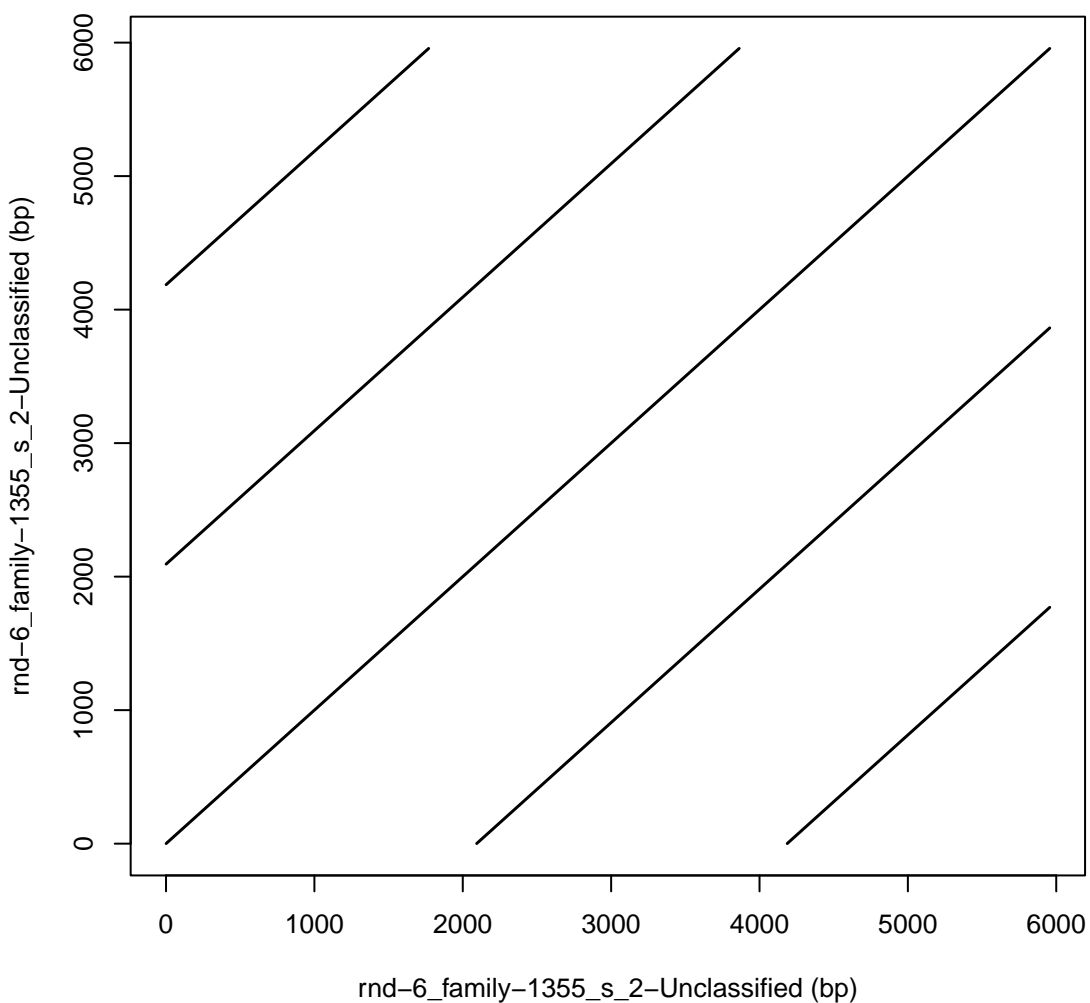
TE: rnd-6_family-1355_s_2-Unclassified
consensus size: 5956bp; fragments: 67; full length: 5 (\geq 5360.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

