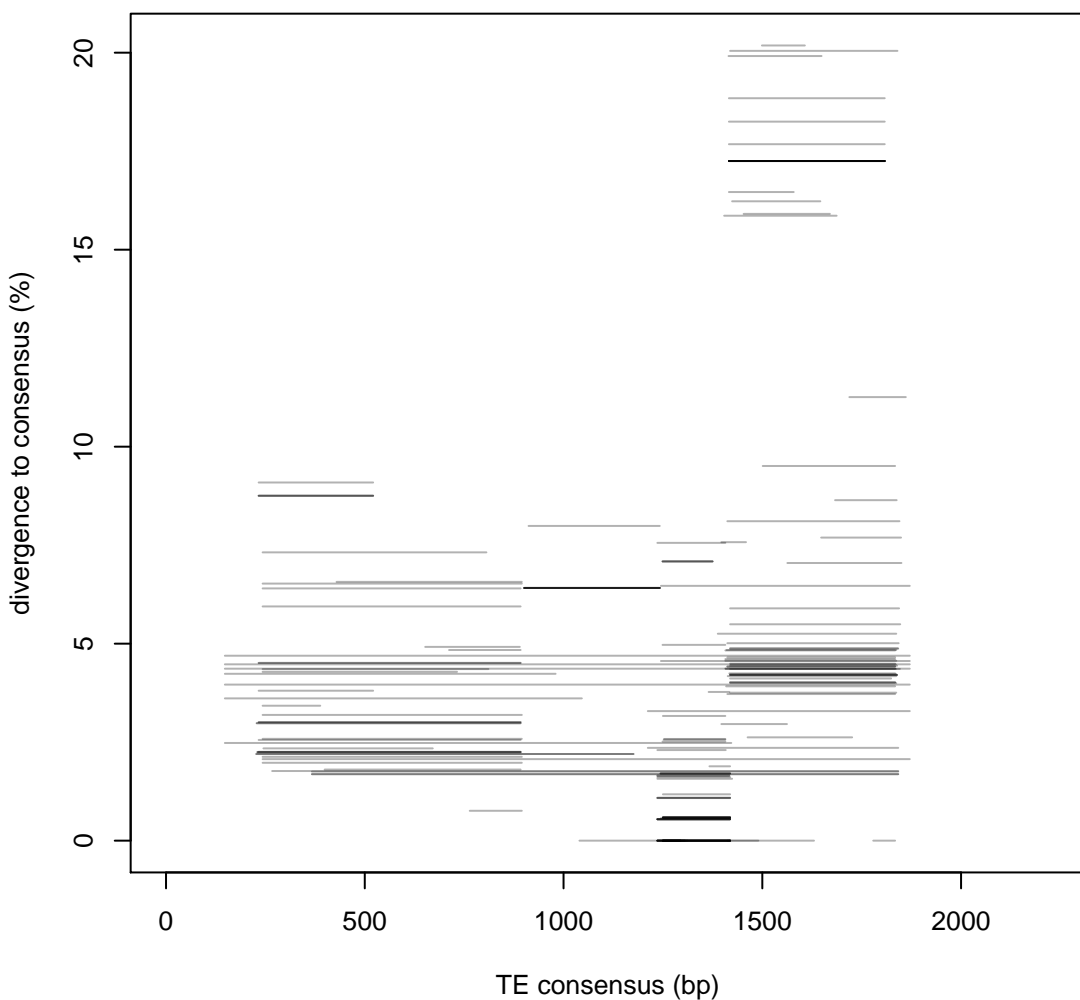
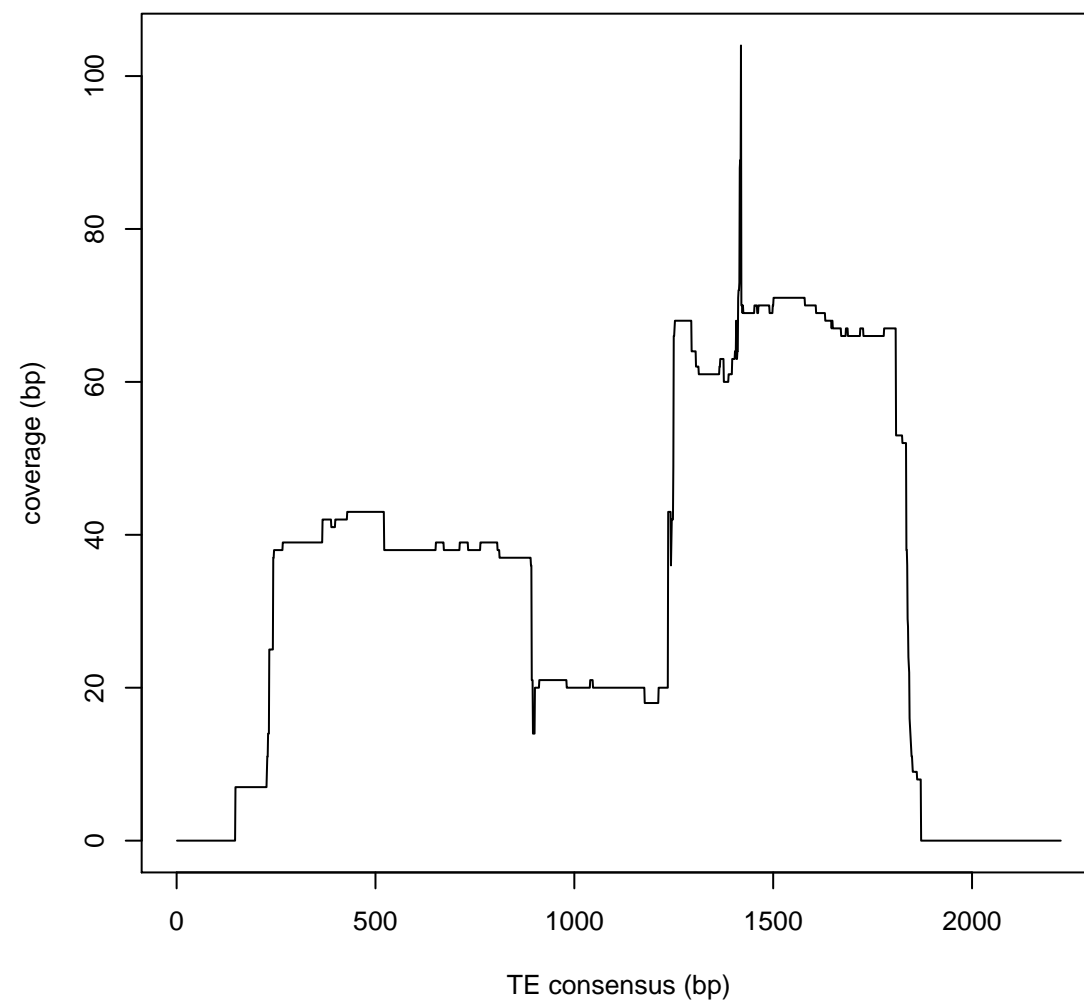


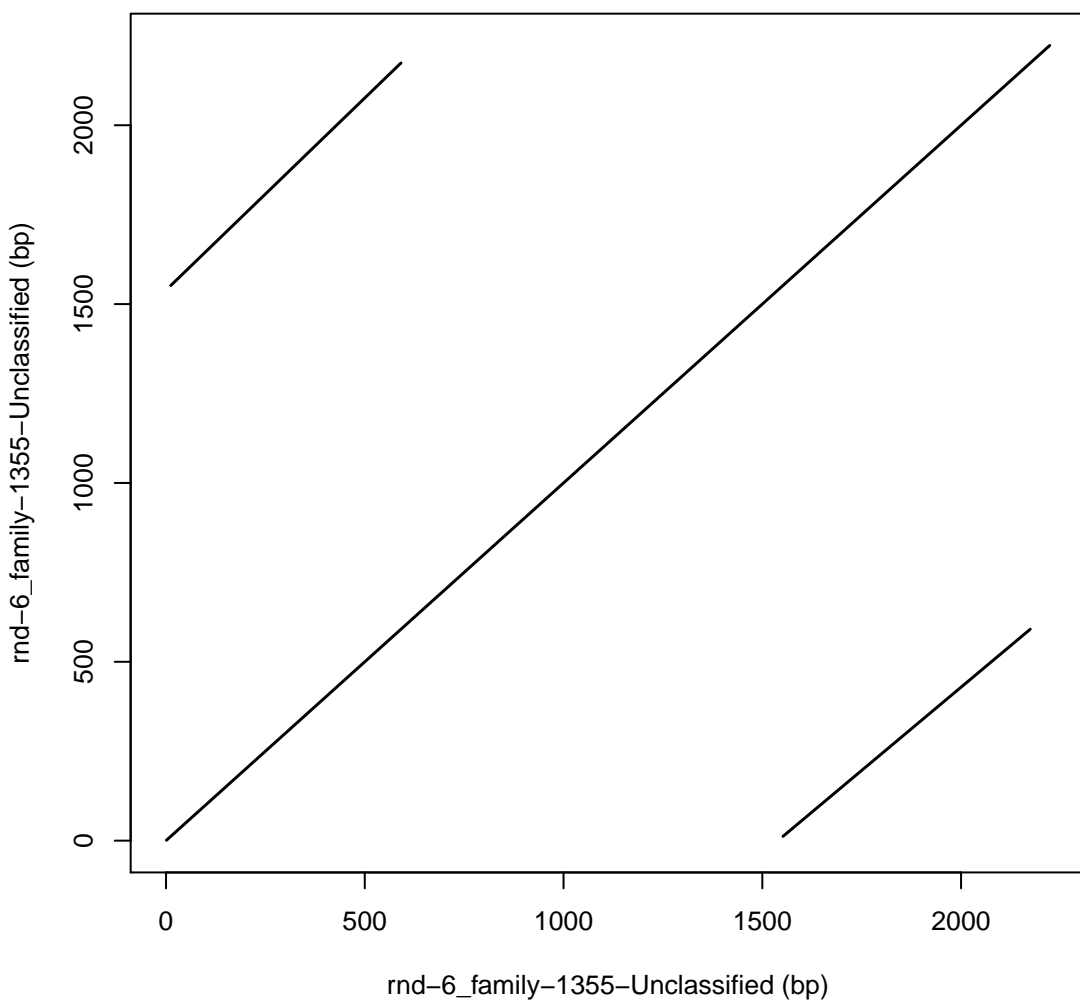
TE: rnd-6_family-1355-Unclassified
consensus size: 2223bp; fragments: 179; full length: 0 (>=2000.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

