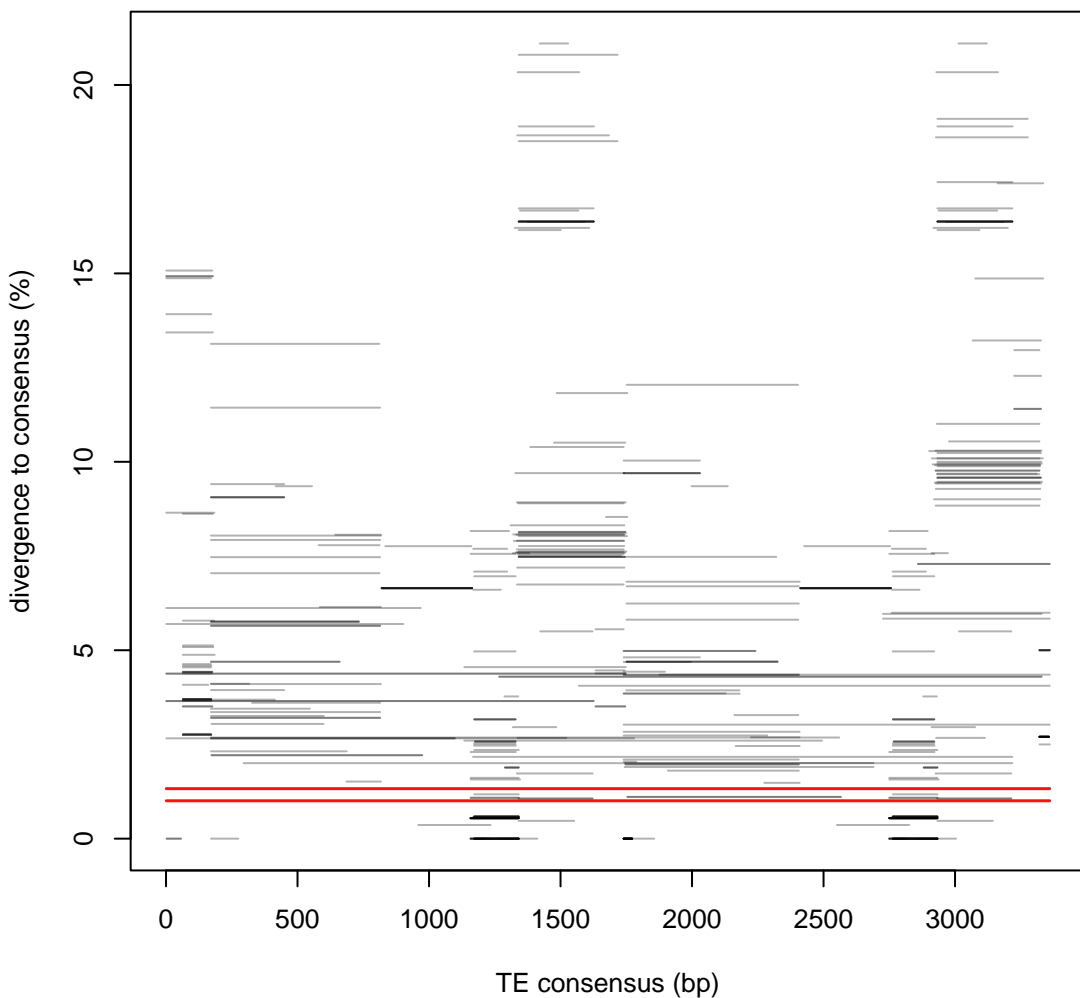
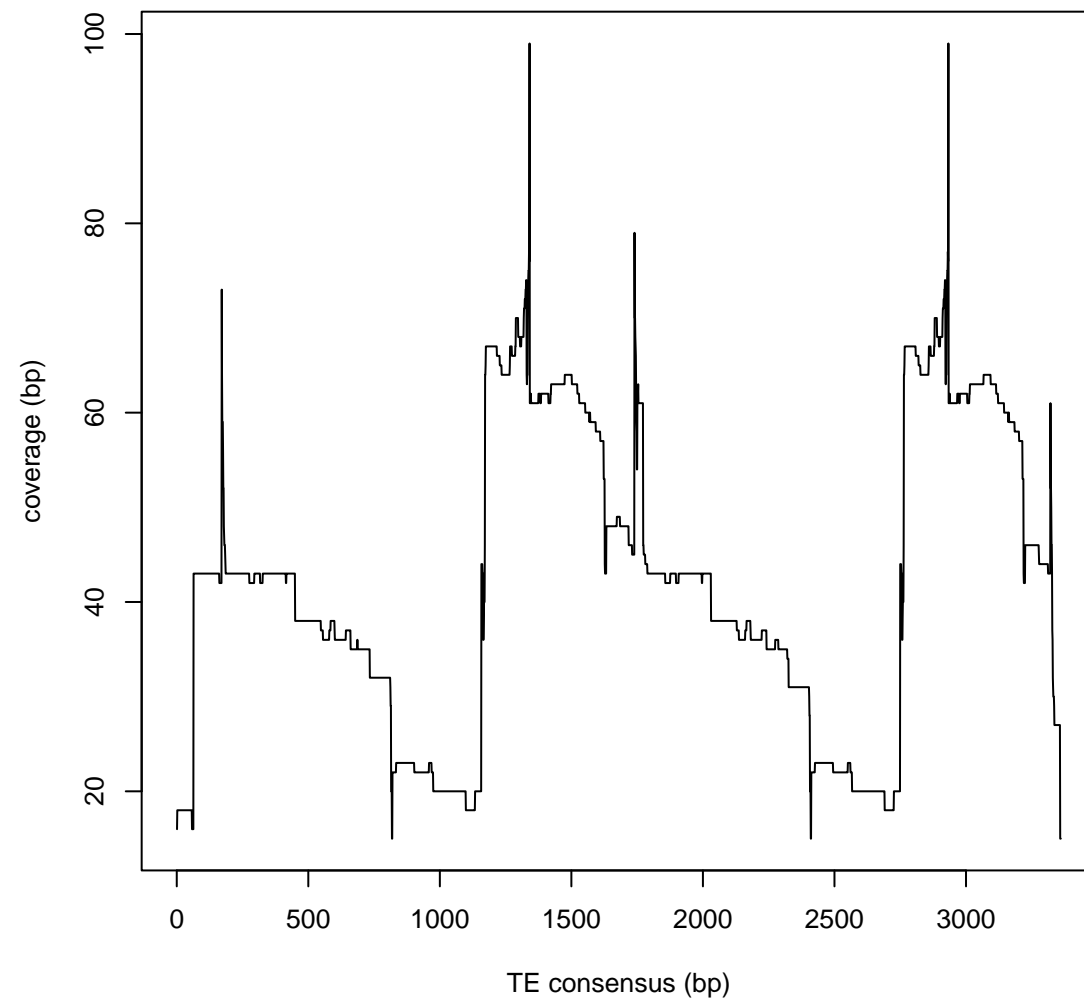


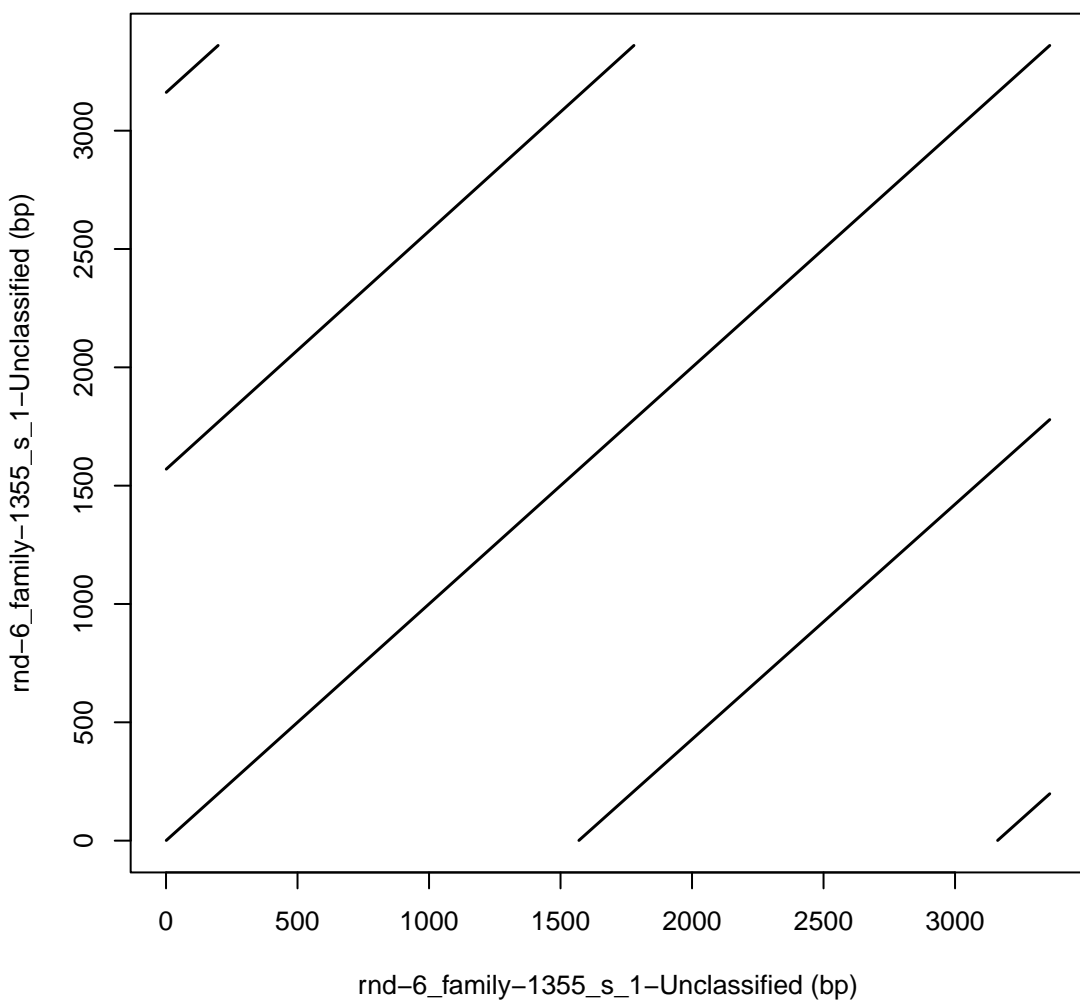
TE: rnd-6_family-1355_s_1-Unclassified
consensus size: 3360bp; fragments: 426; full length: 2 (>=3024bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

