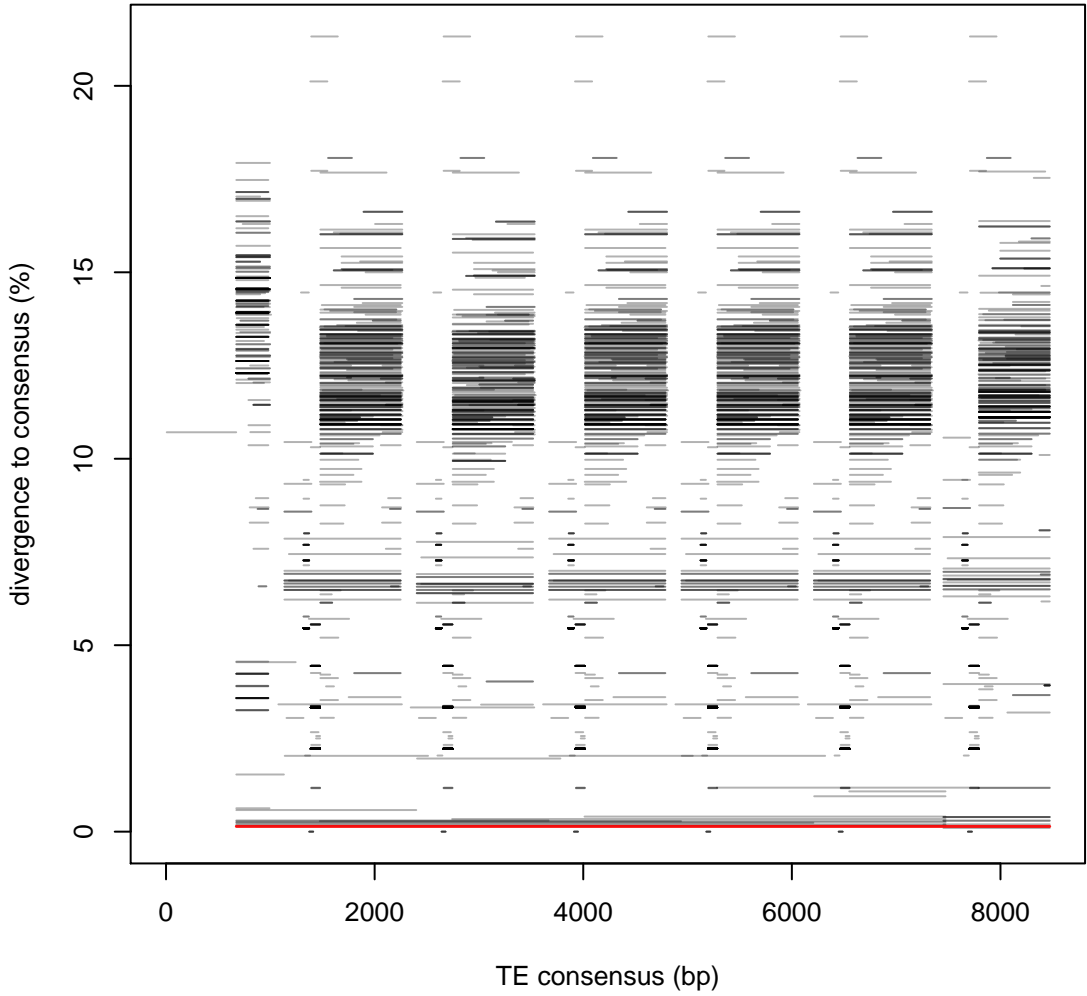
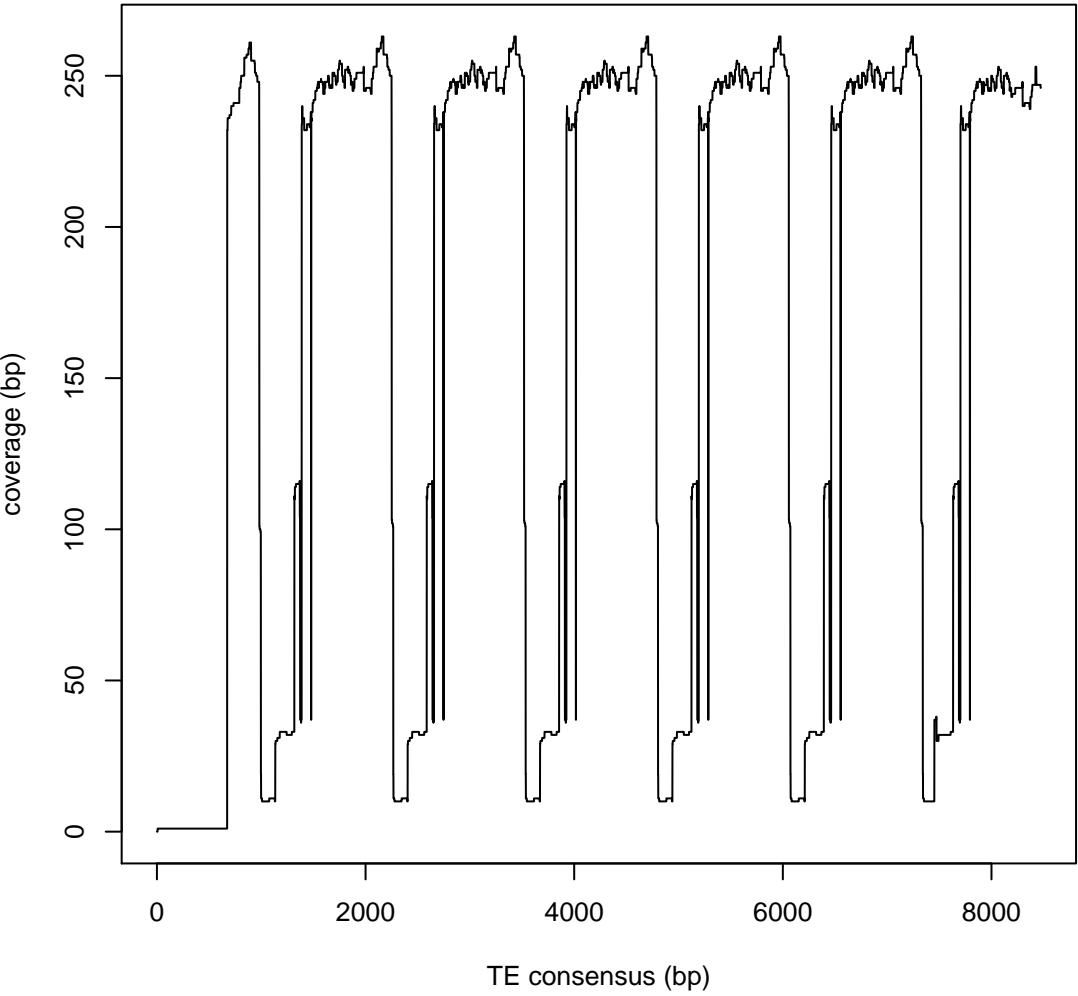


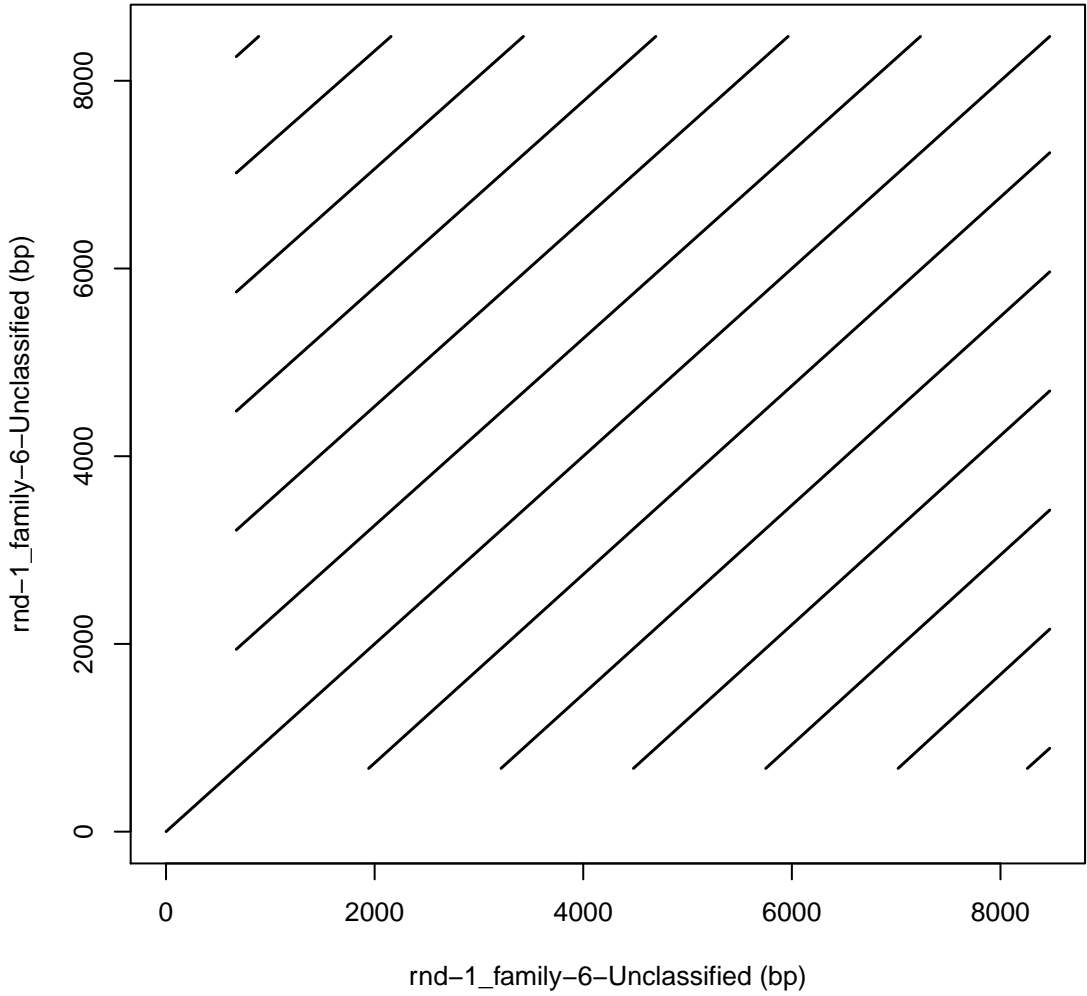
TE: rnd-1_family-6-Unclassified
consensus size: 8472bp; fragments: 3858; full length: 1 (>=7624.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

