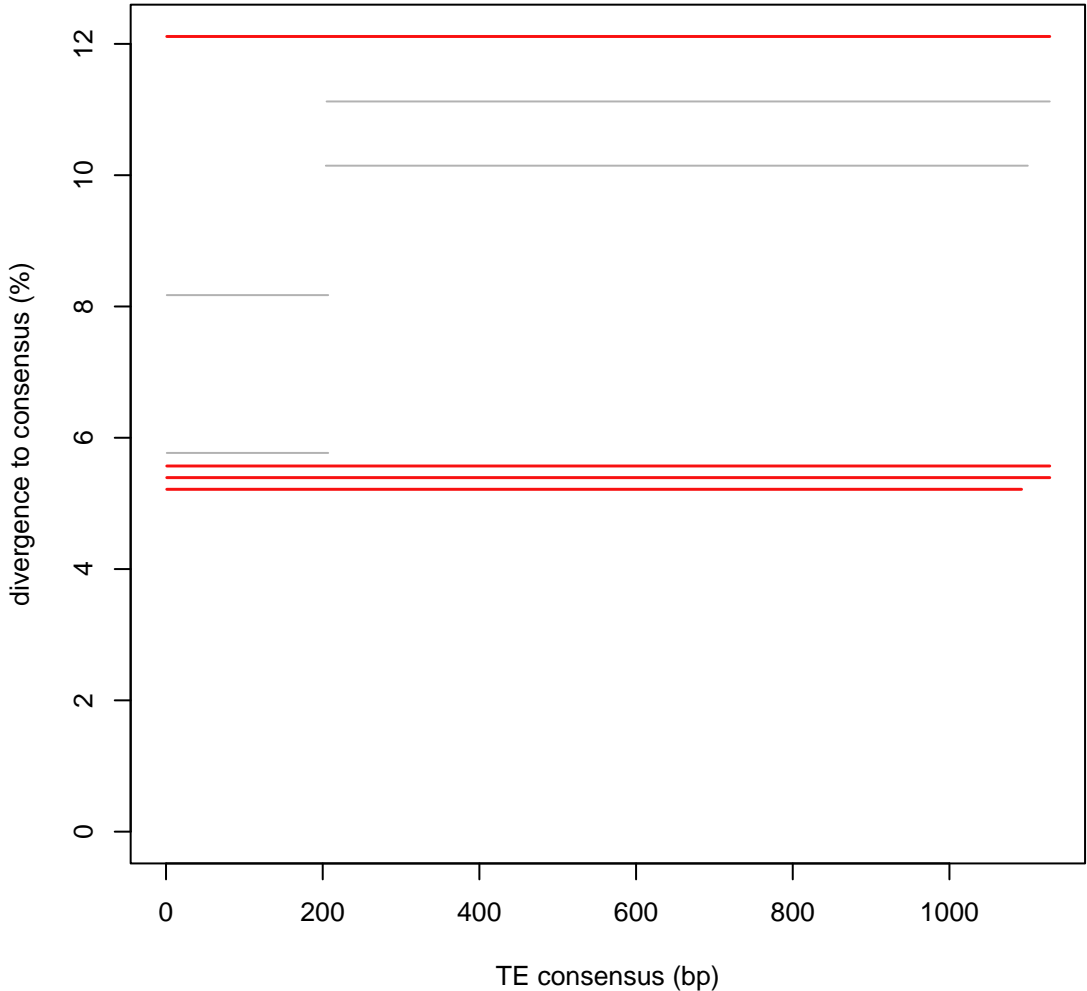
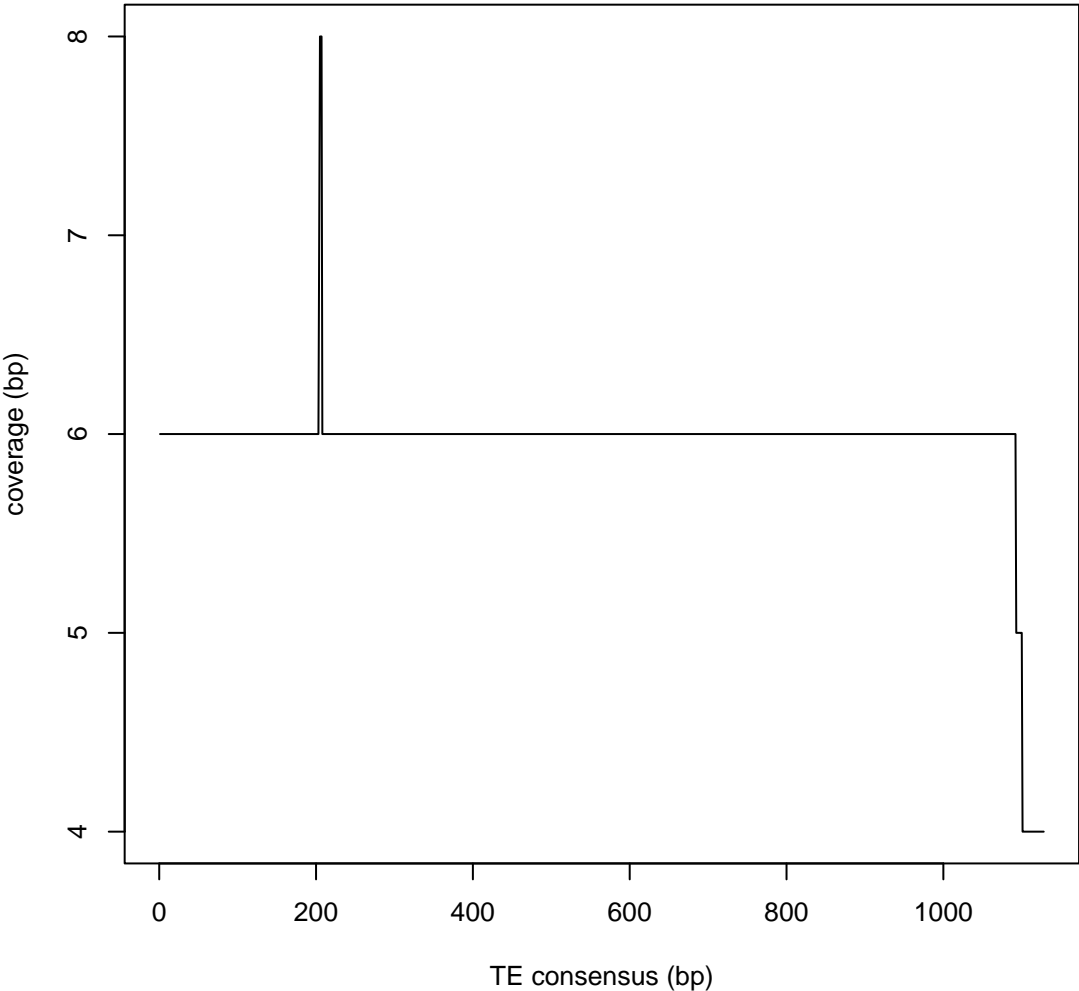


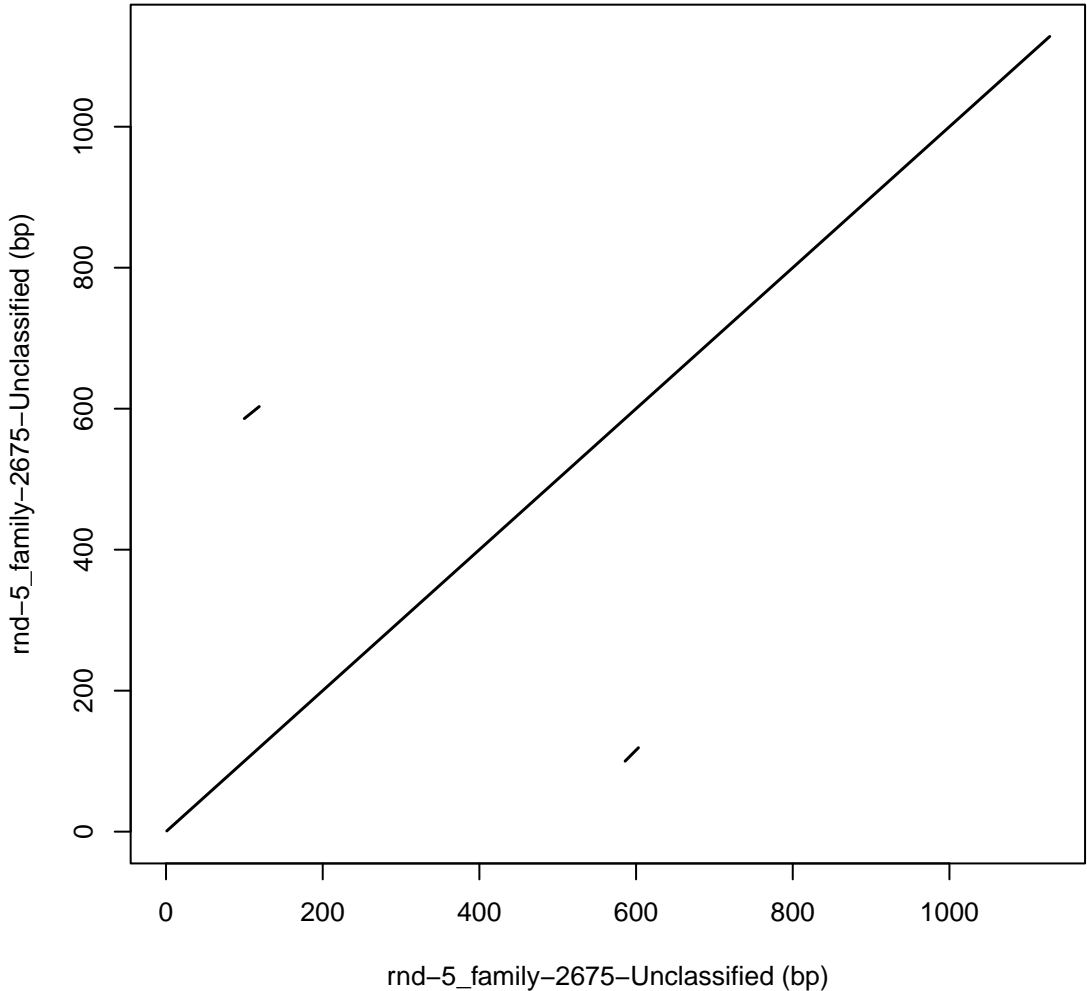
TE: rnd-5_family-2675-Unclassified
consensus size: 1128bp; fragments: 8; full length: 4 (>=1015.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

