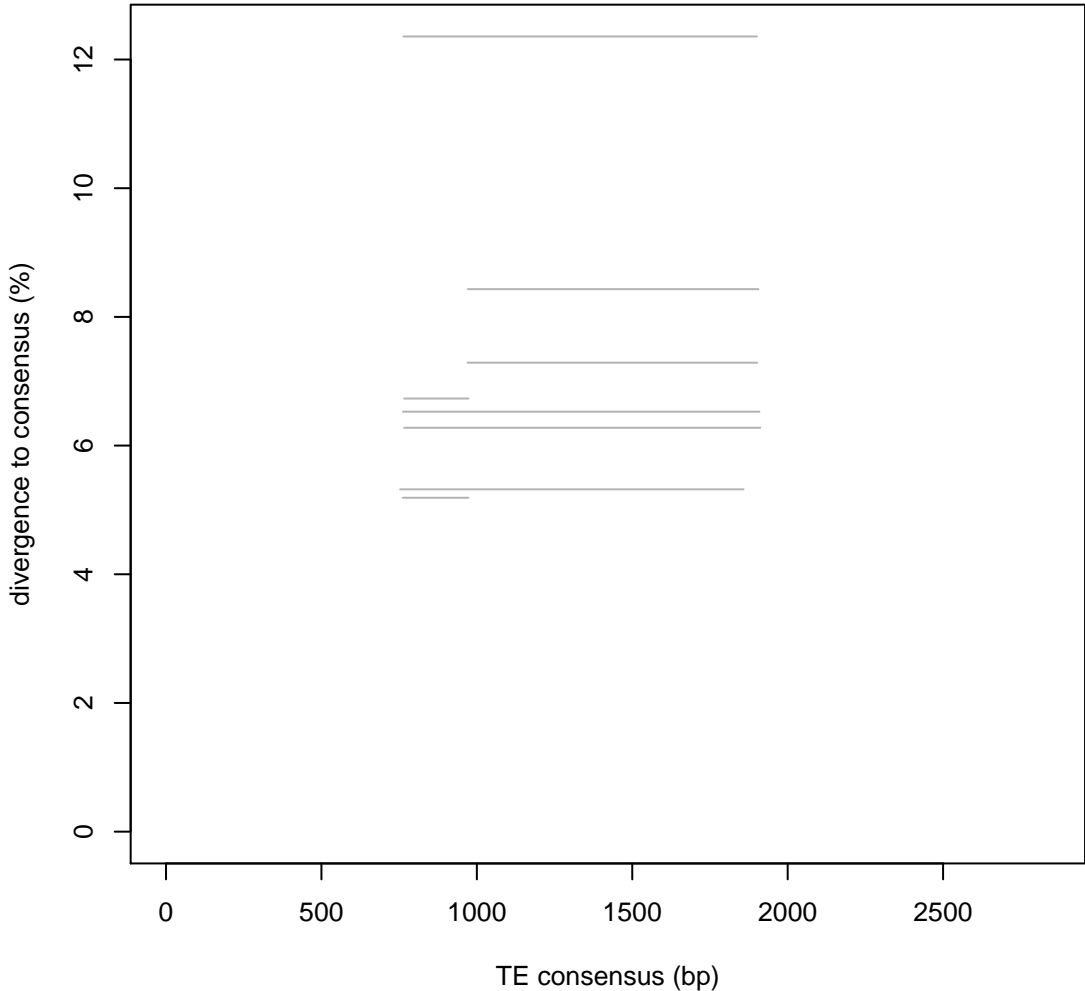
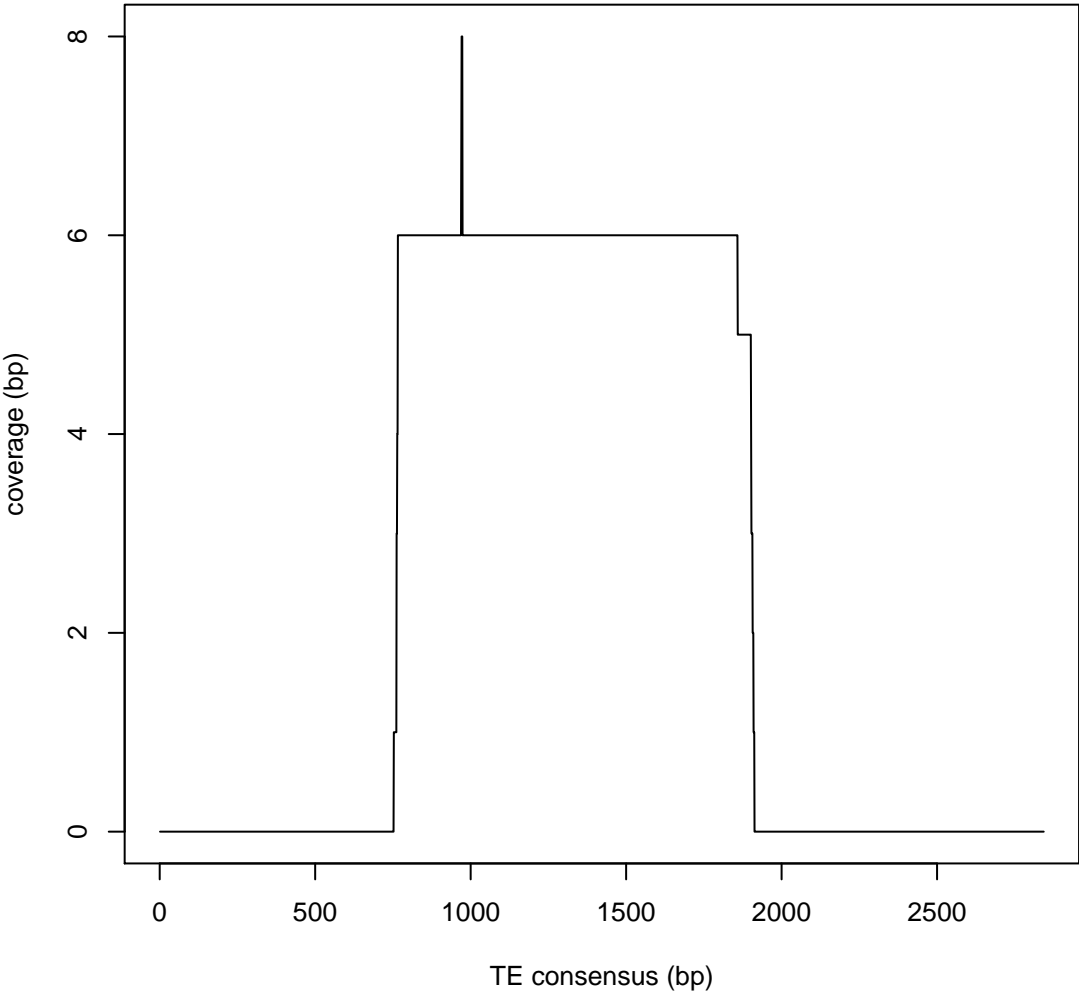


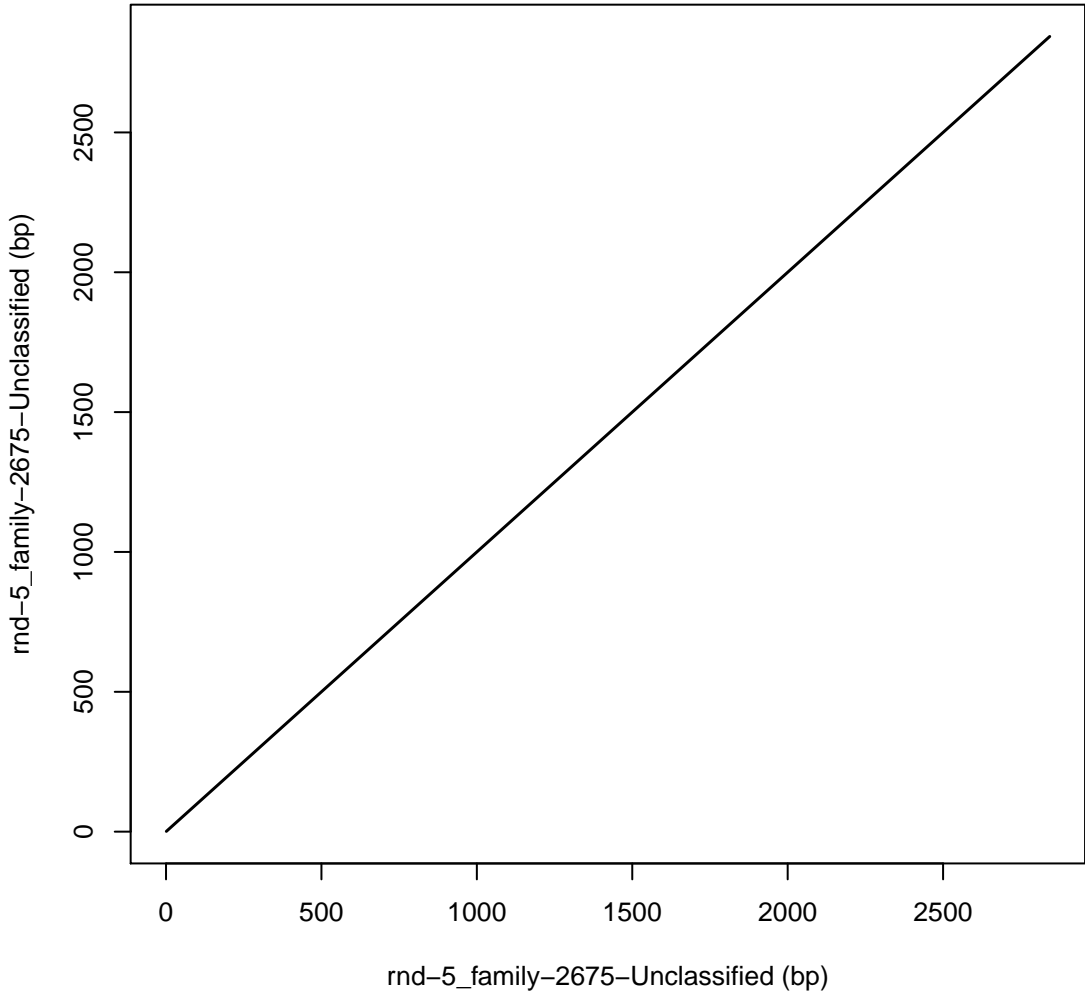
TE: rnd-5_family-2675-Unclassified
consensus size: 2843bp; fragments: 8; full length: 0 (>=2558.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

