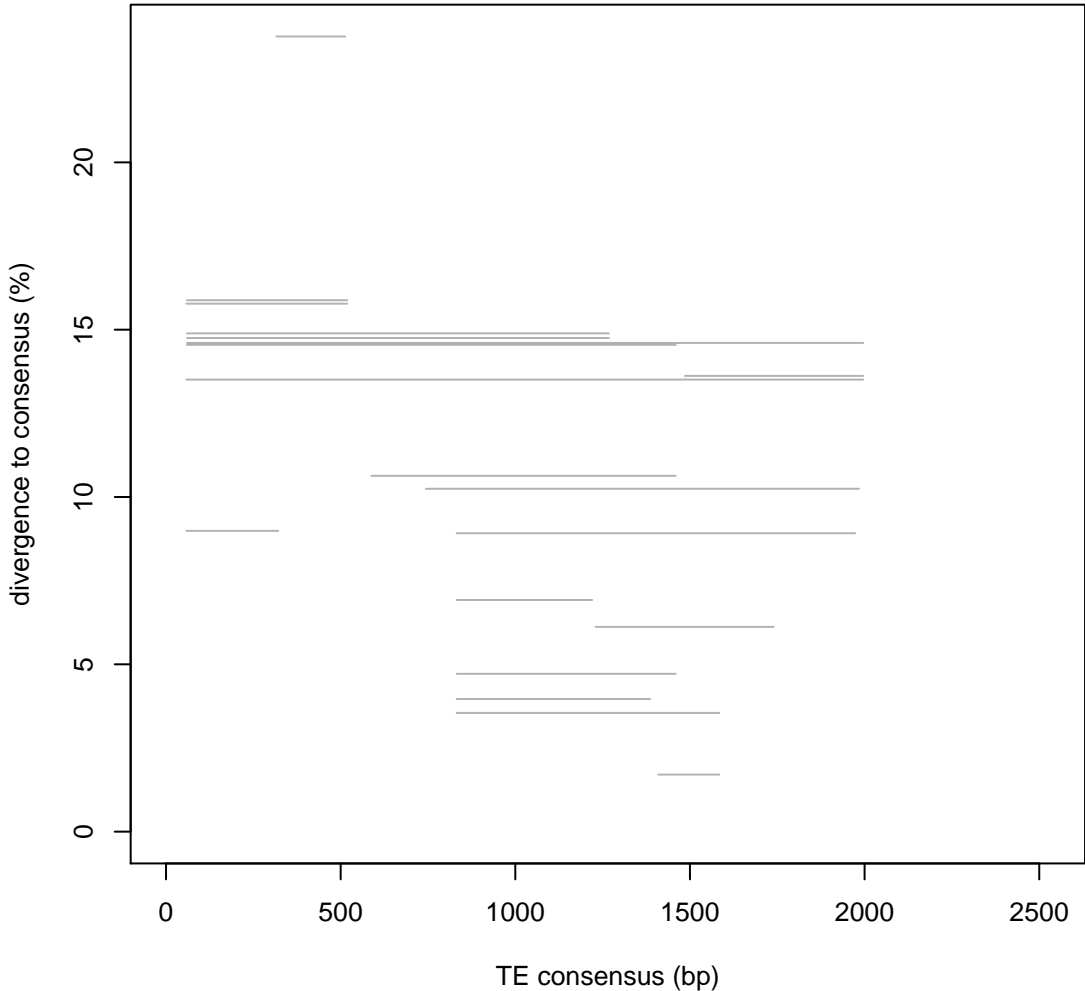
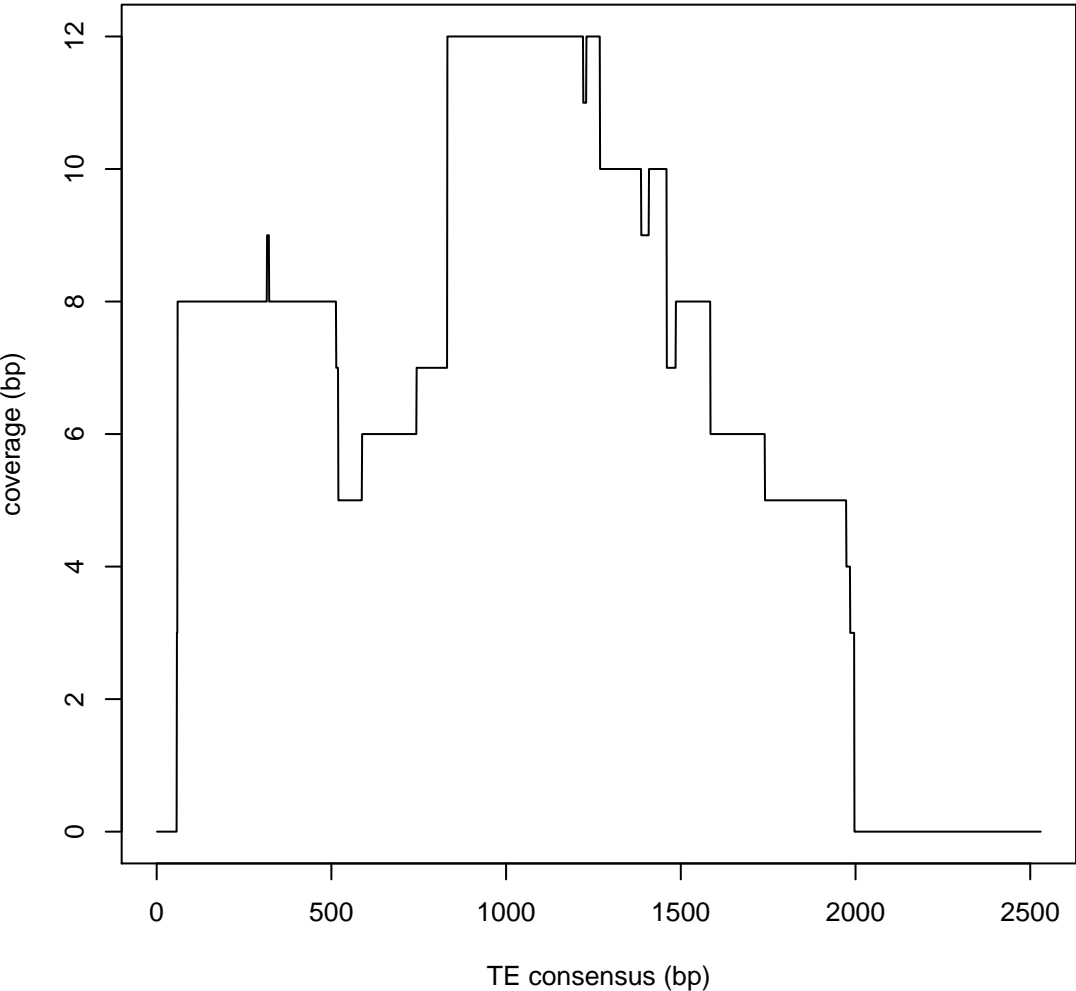


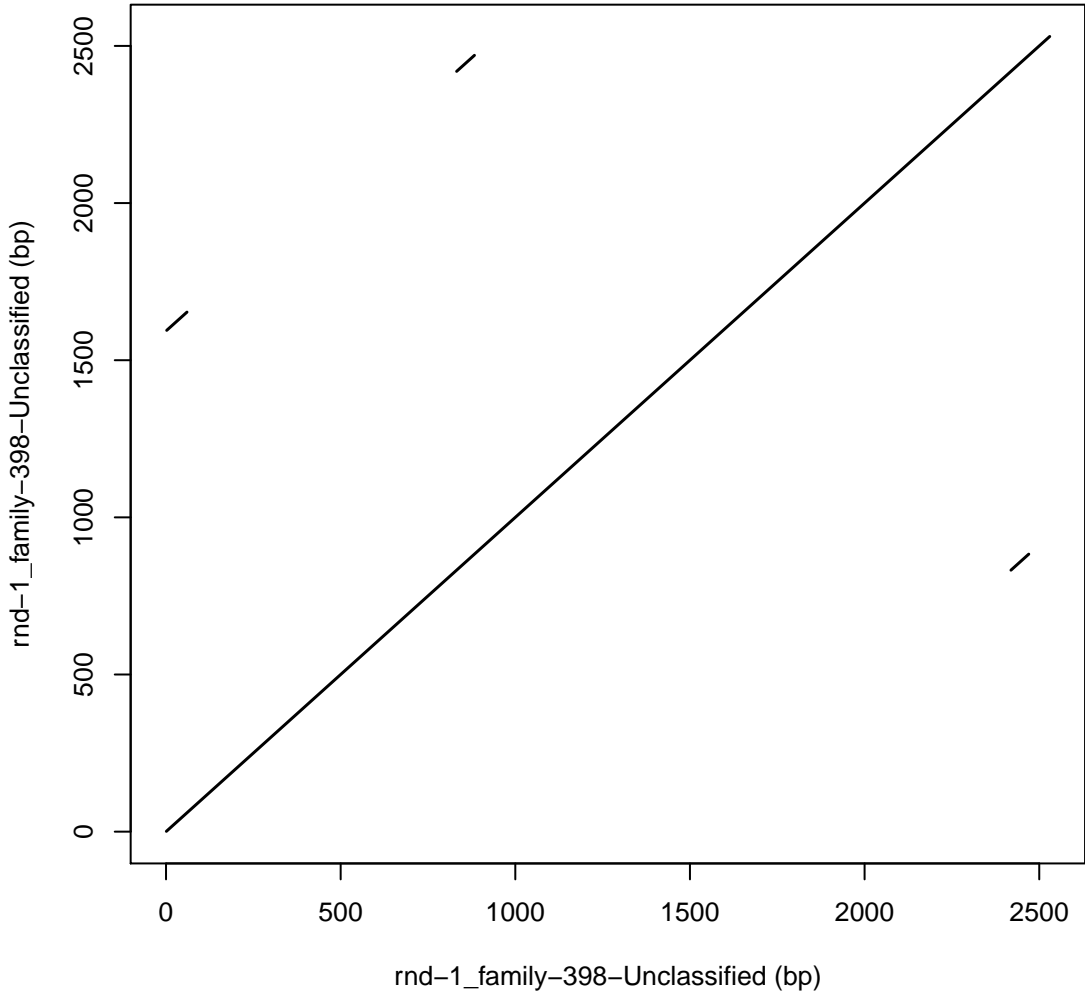
TE: rnd-1_family-398-Unclassified
consensus size: 2530bp; fragments: 19; full length: 0 (>=2277bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

