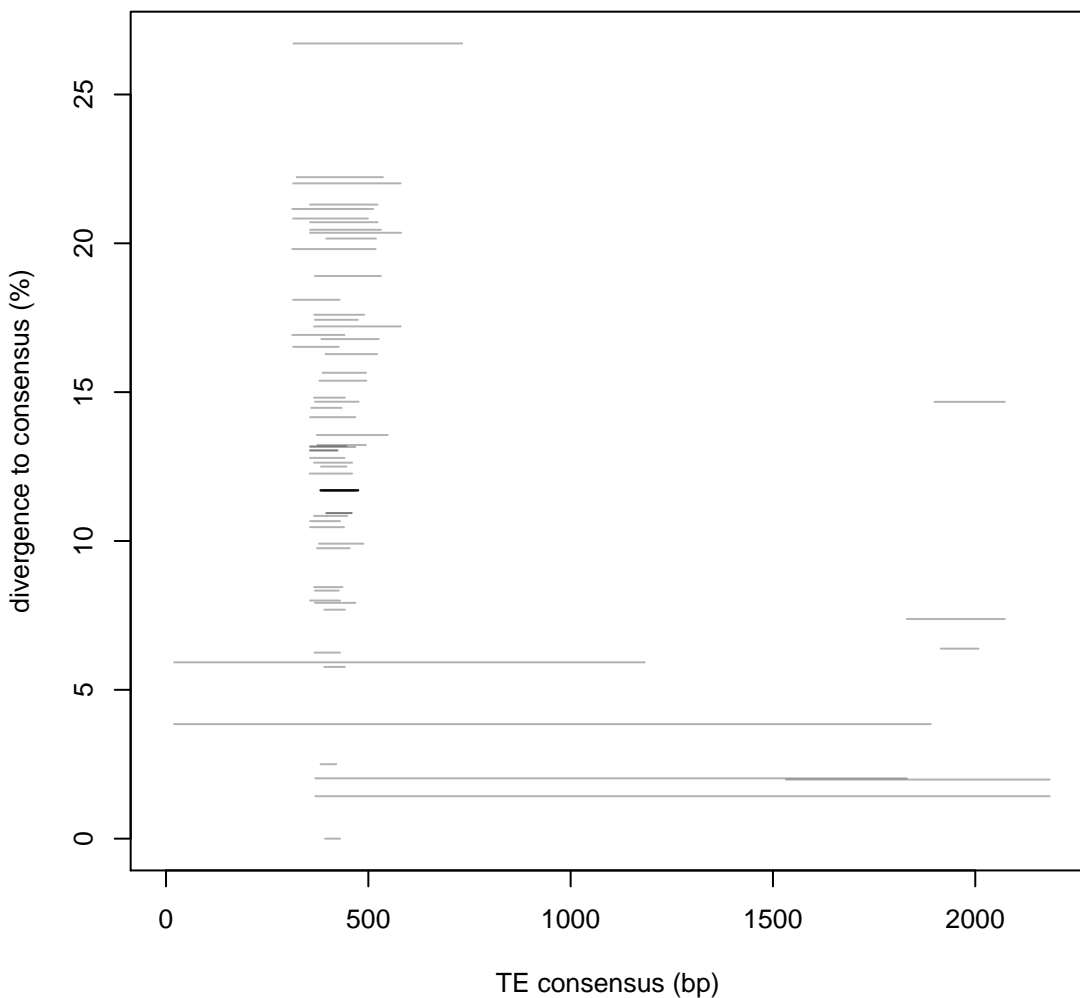
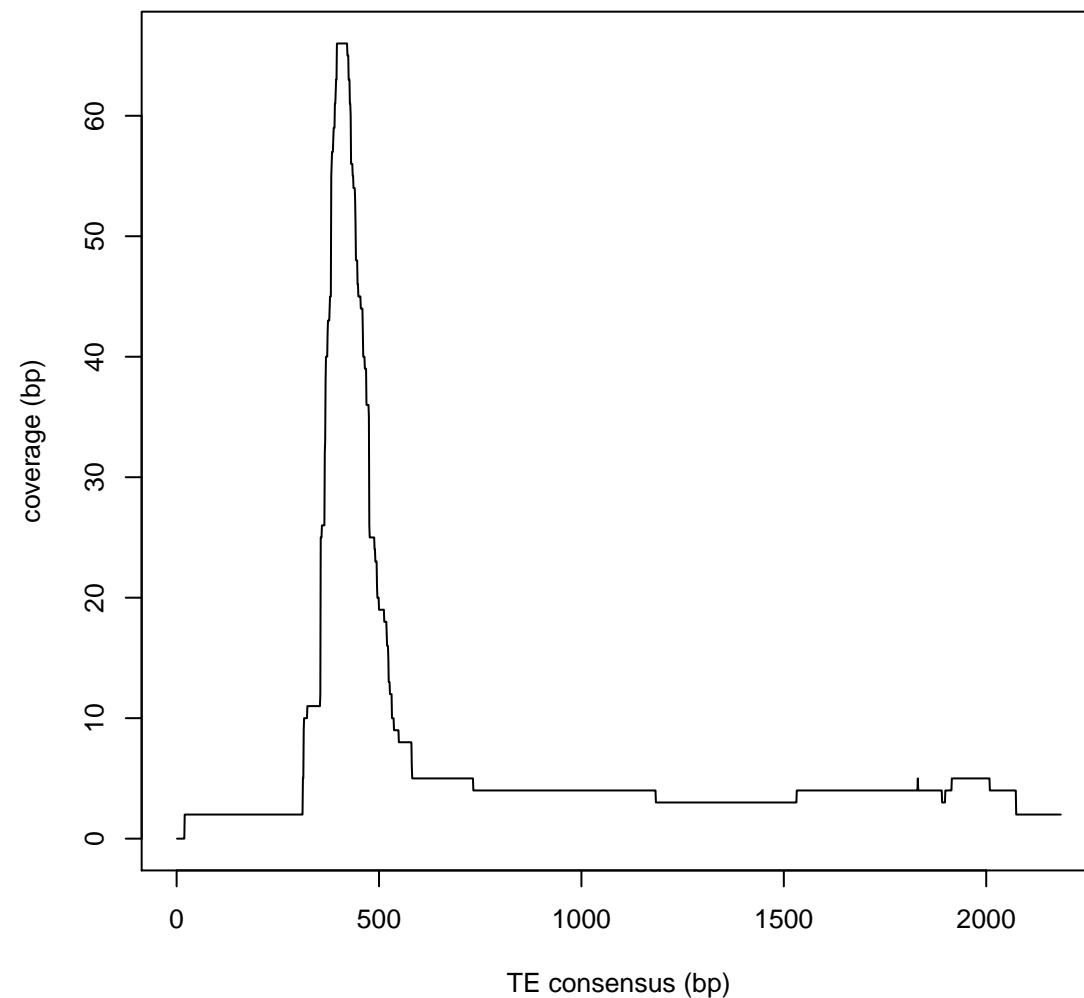


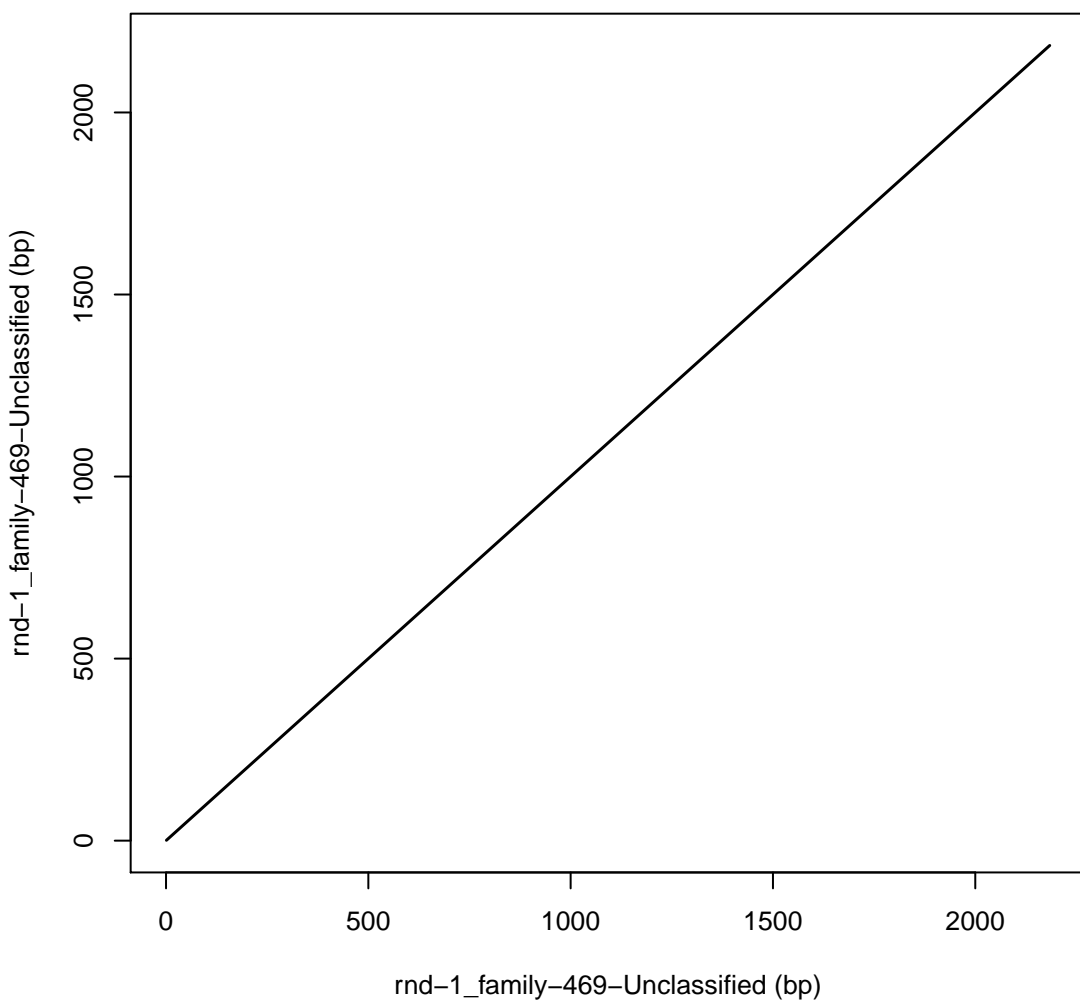
TE: rnd-1\_family-469-Unclassified  
consensus size: 2184bp; fragments: 70; full length: 0 ( $\geq 1965.6$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

