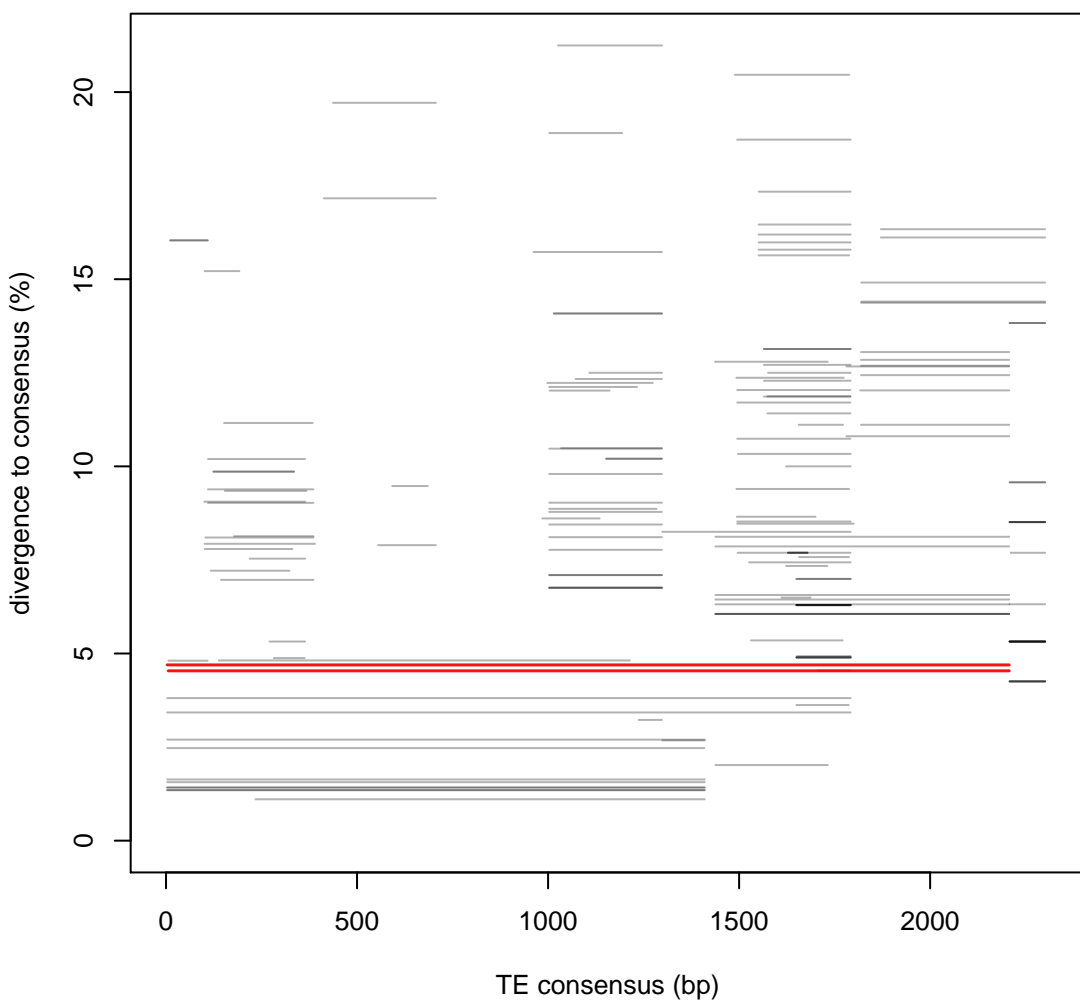
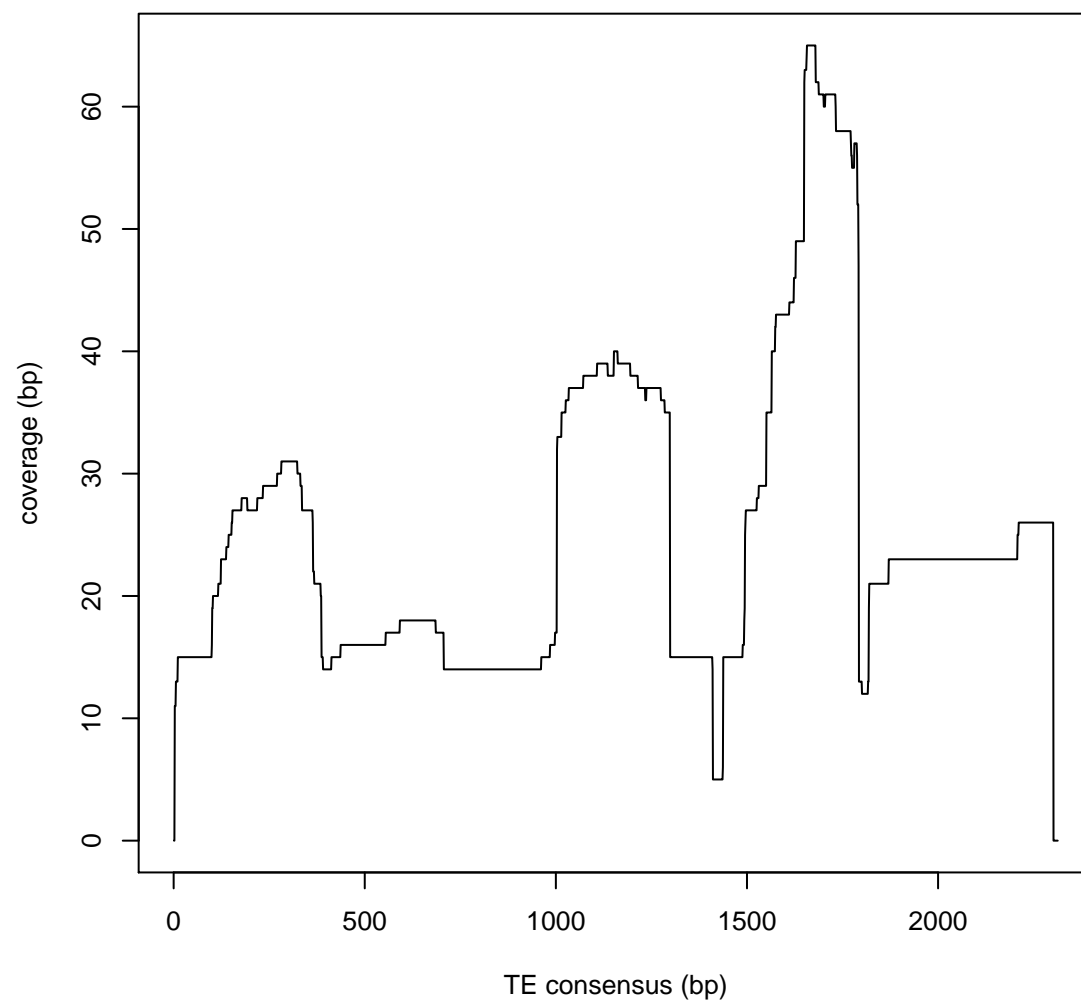


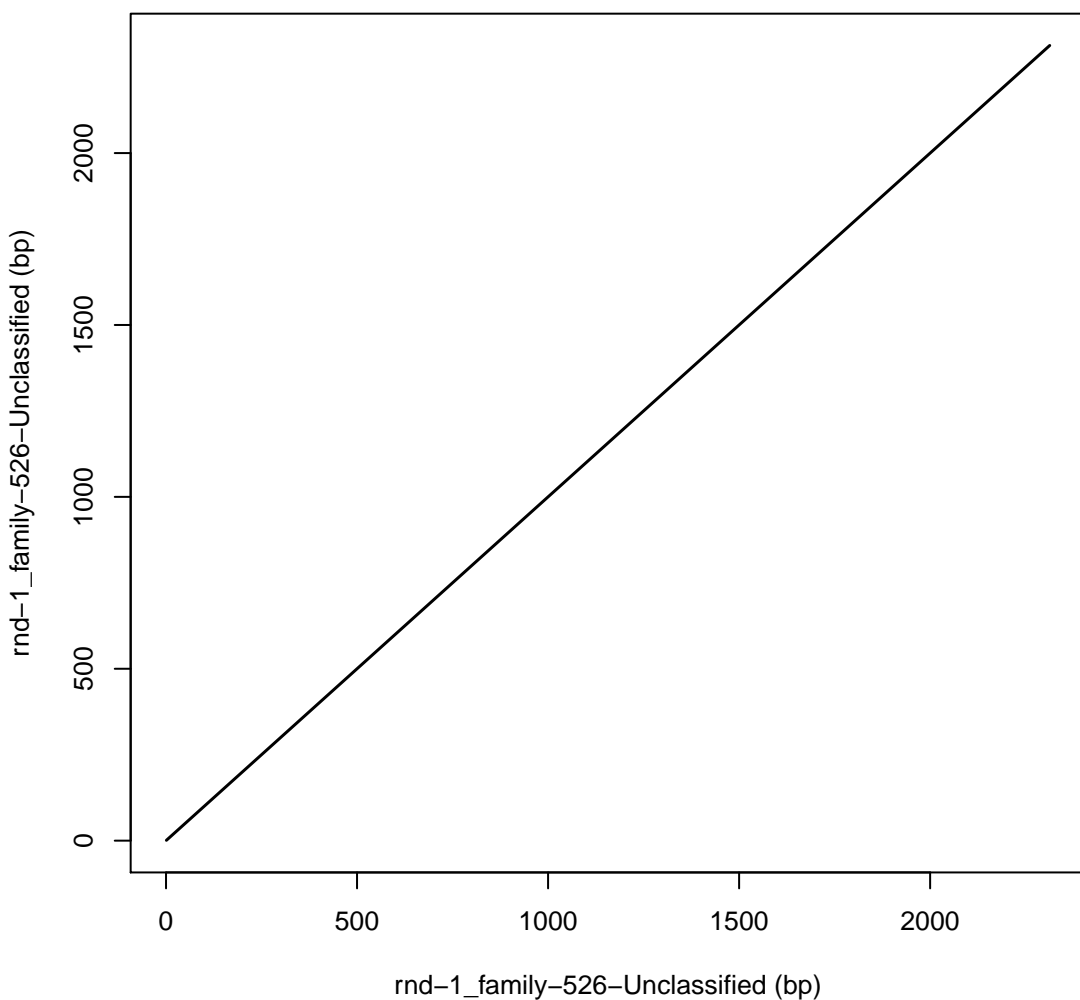
TE: rnd-1\_family-526-Unclassified  
consensus size: 2313bp; fragments: 164; full length: 2 ( $\geq 2081.7$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

