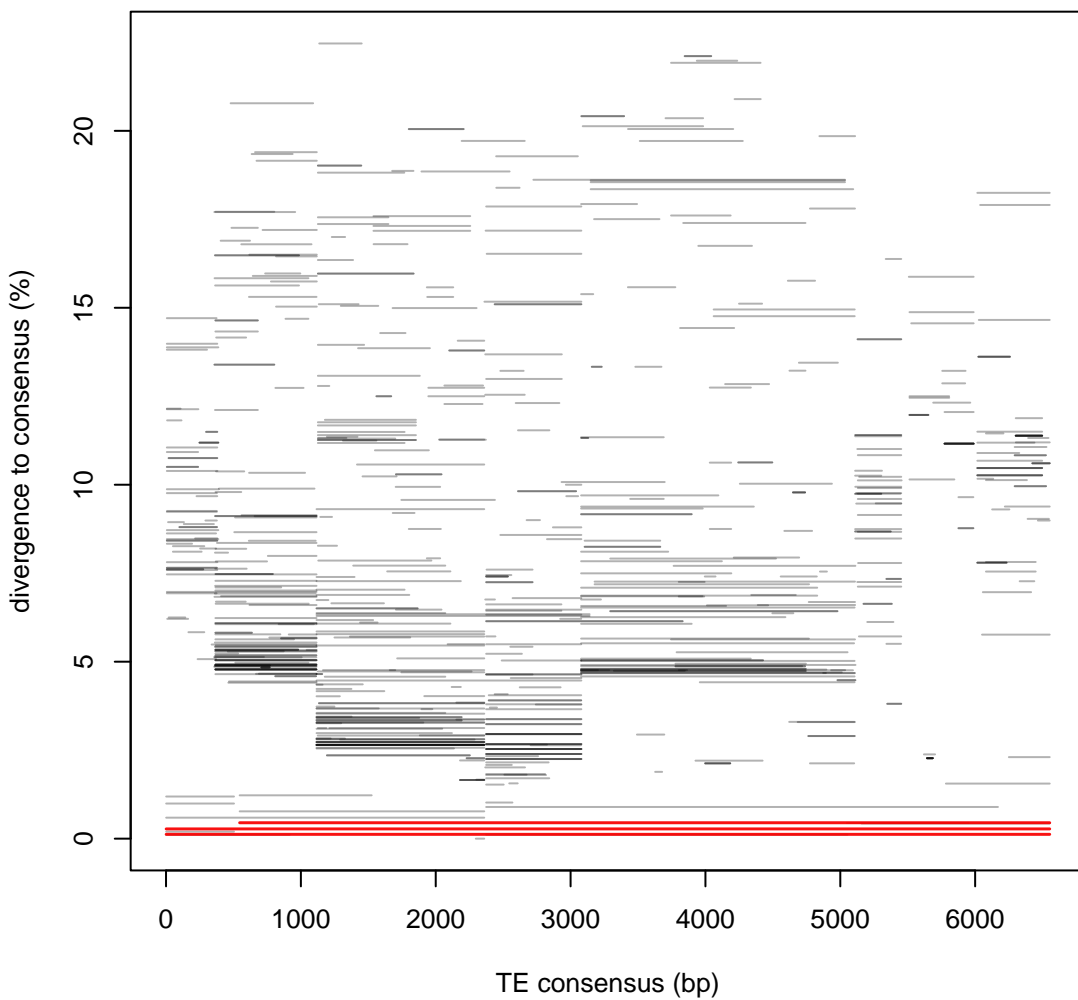
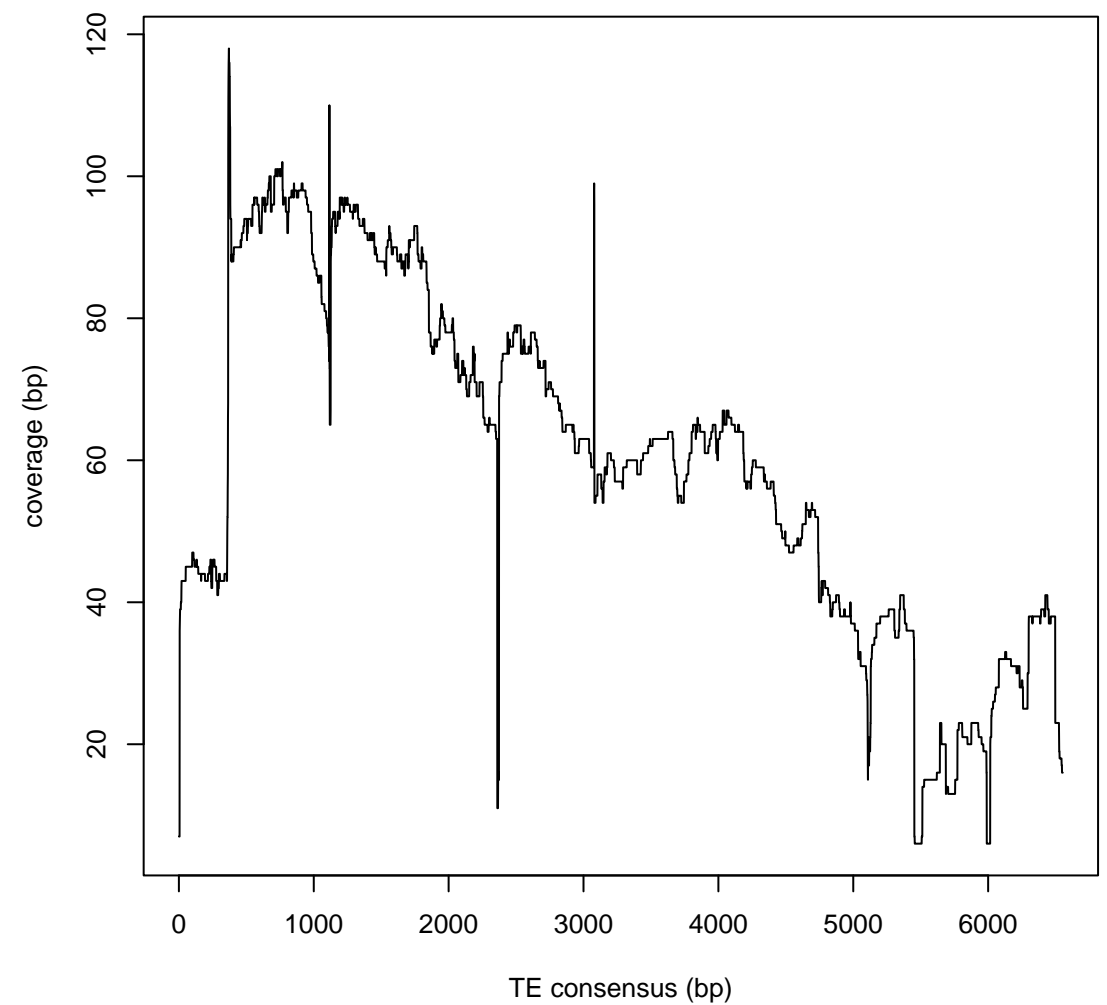


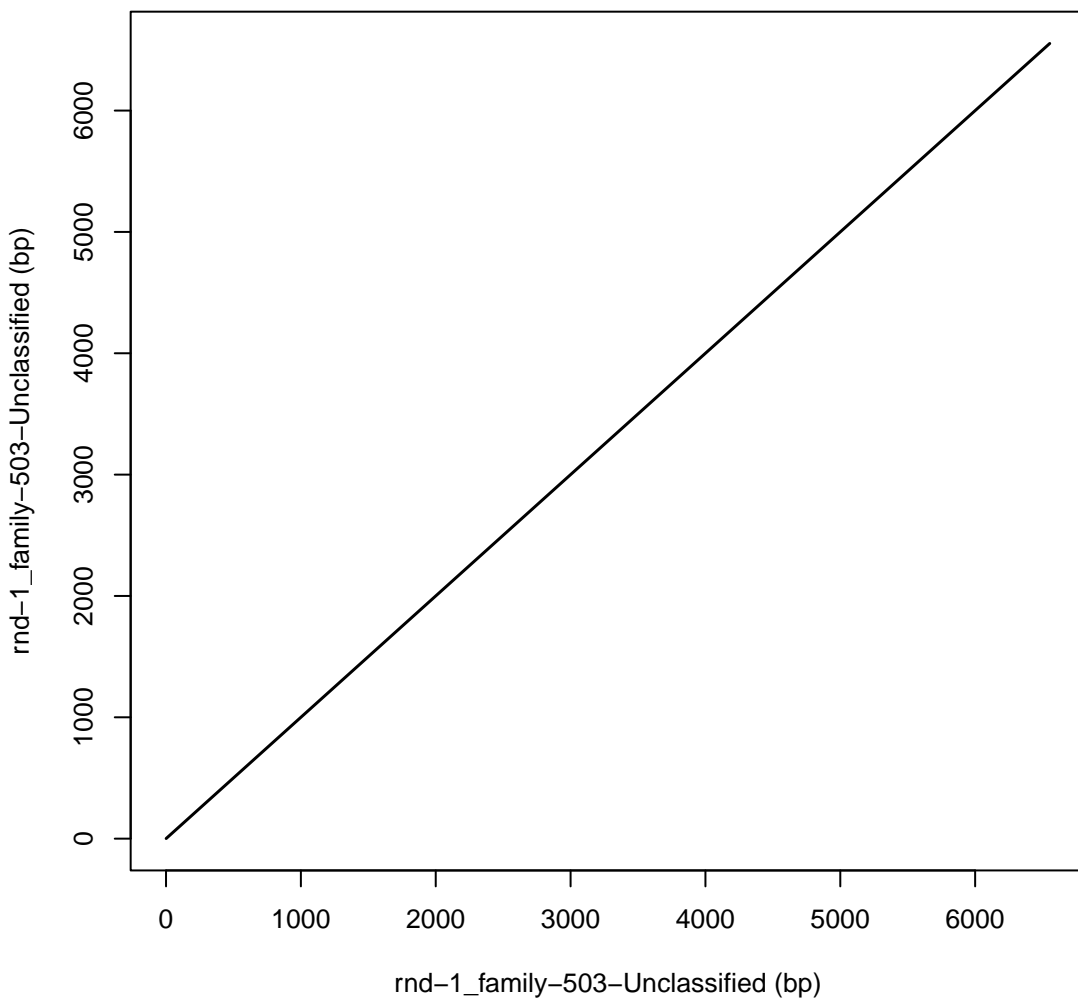
TE: rnd-1\_family-503-Unclassified  
consensus size: 6553bp; fragments: 687; full length: 3 ( $\geq 5897.7$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

