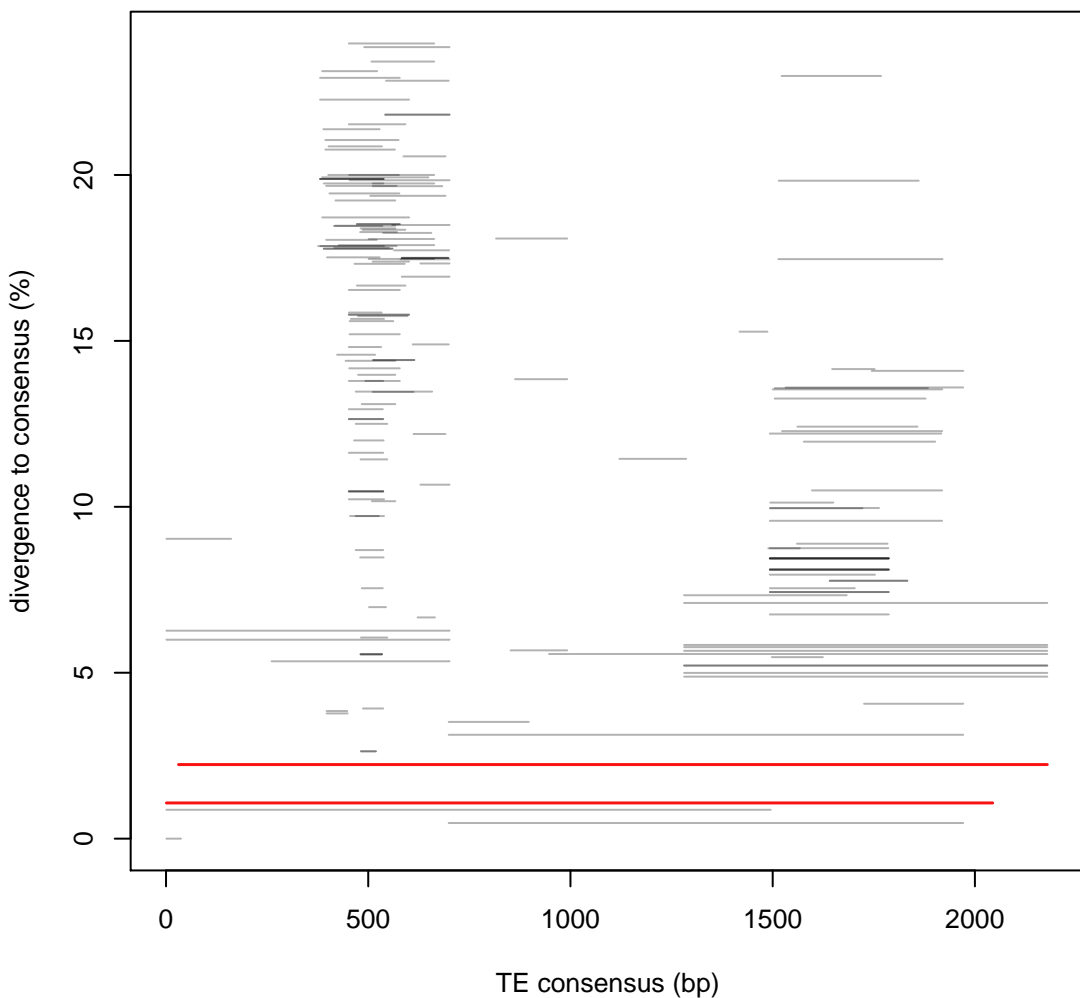
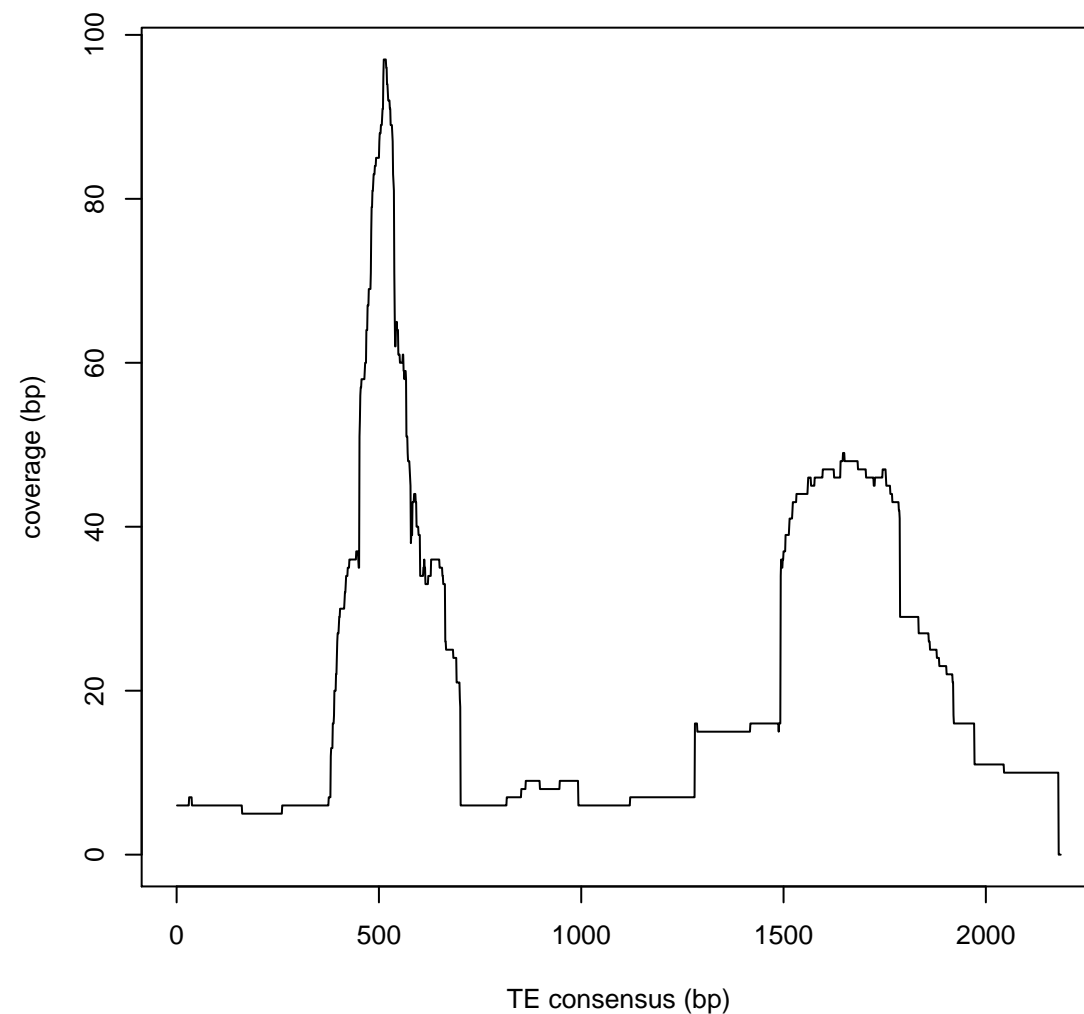


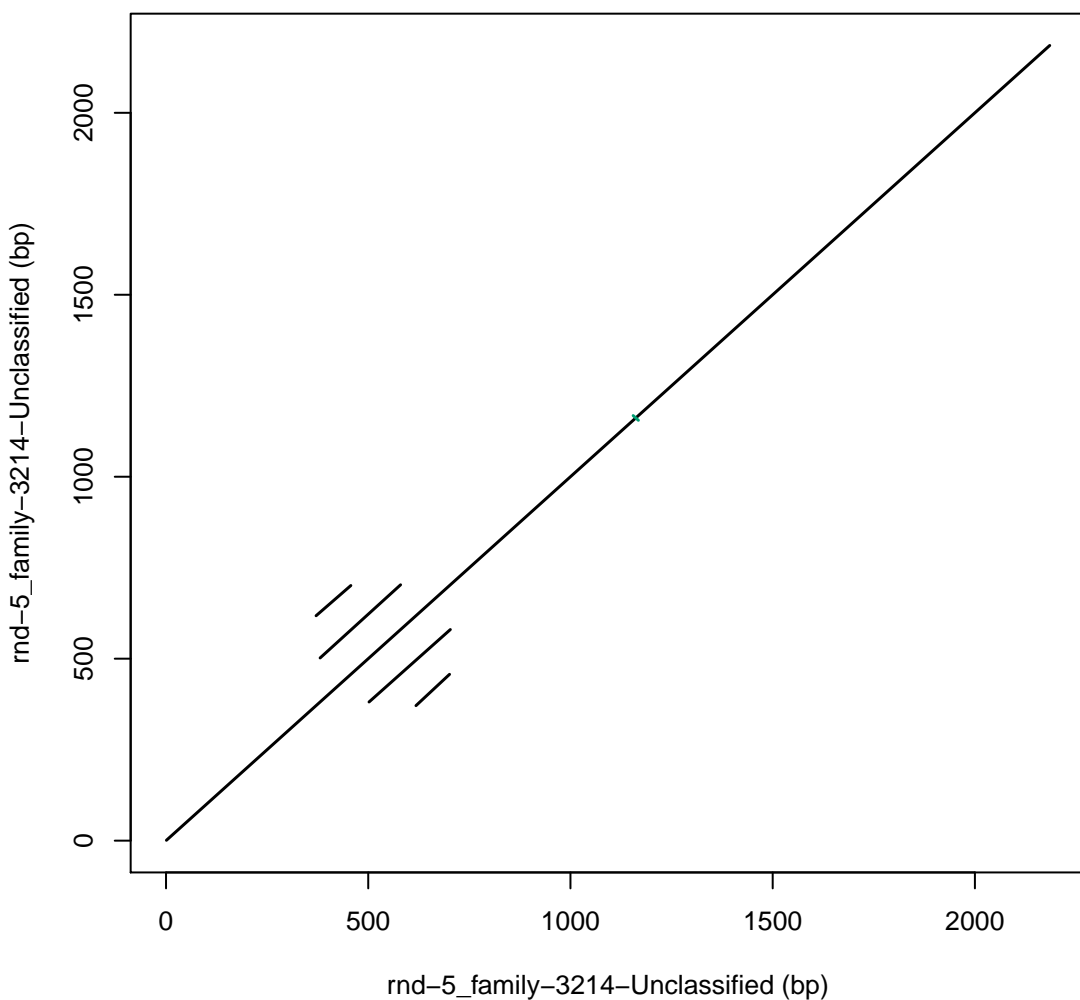
TE: rnd-5\_family-3214-Unclassified  
consensus size: 2185bp; fragments: 175; full length: 2 ( $\geq 1966.5$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

