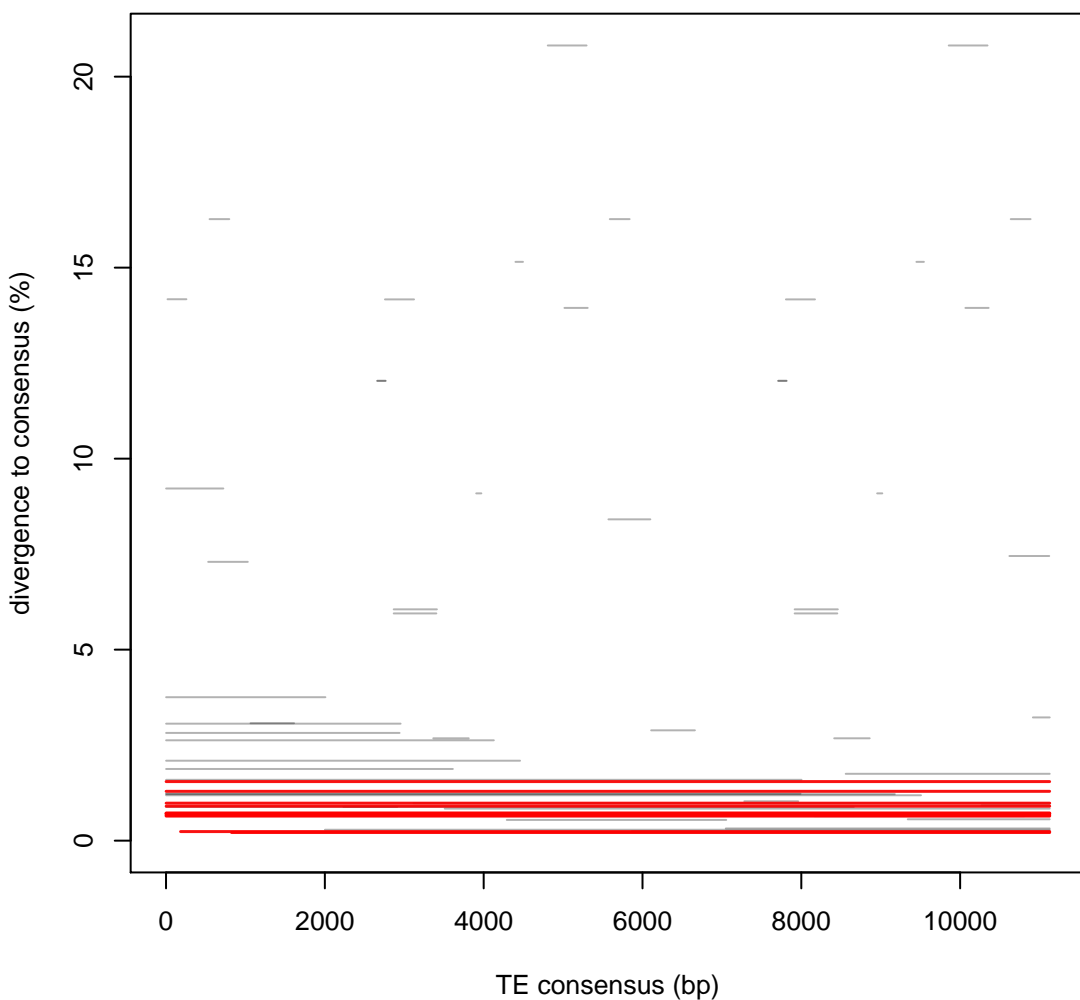
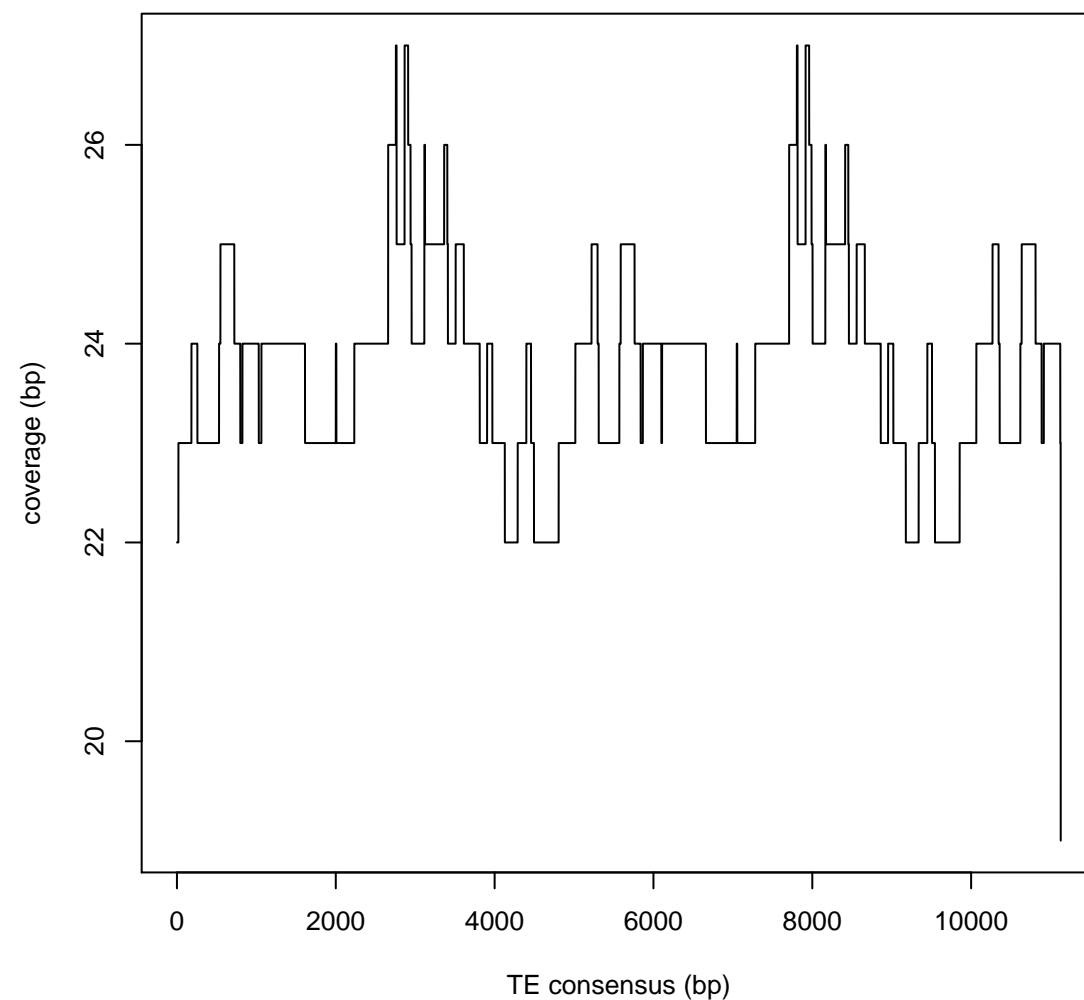


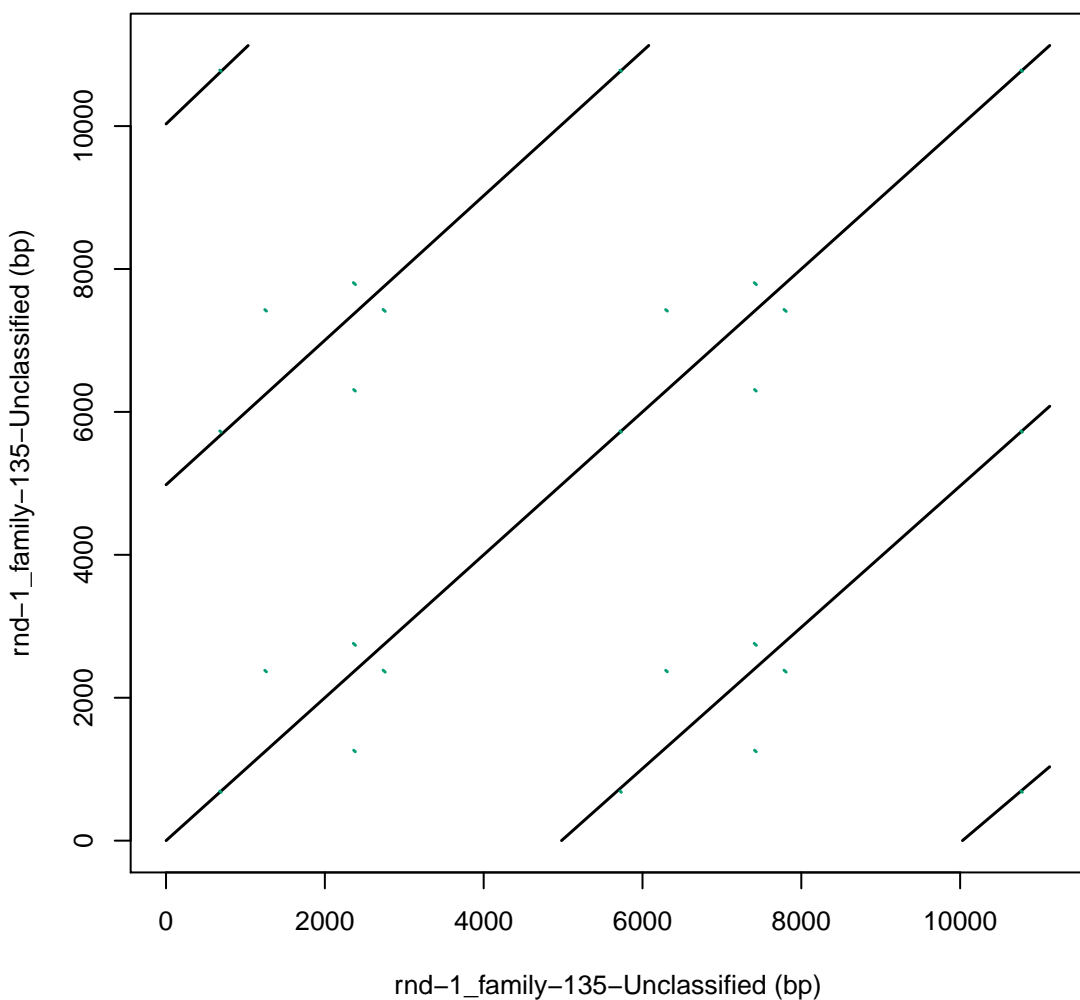
TE: rnd-1\_family-135-Unclassified  
consensus size: 11128bp; fragments: 69; full length: 12 ( $\geq 10015.2\text{bp}$ )



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

