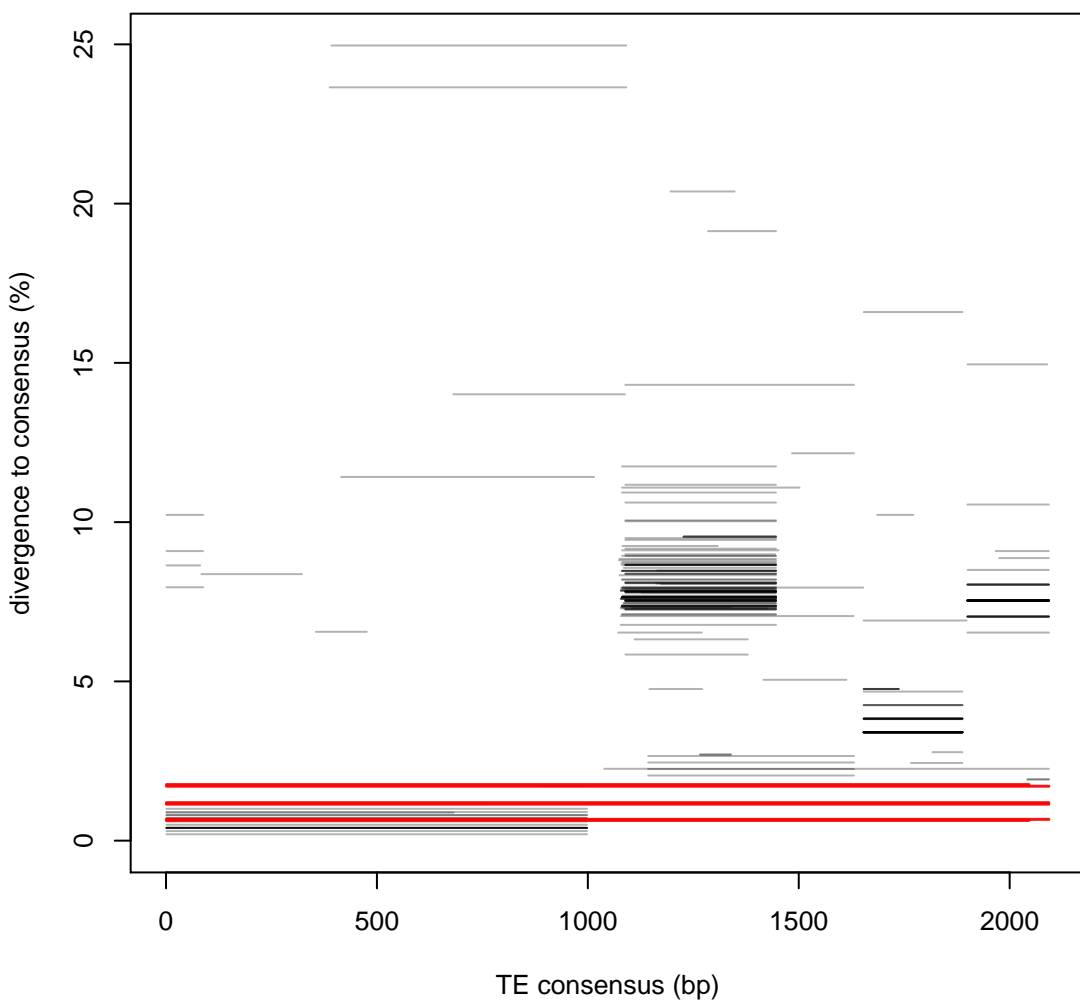
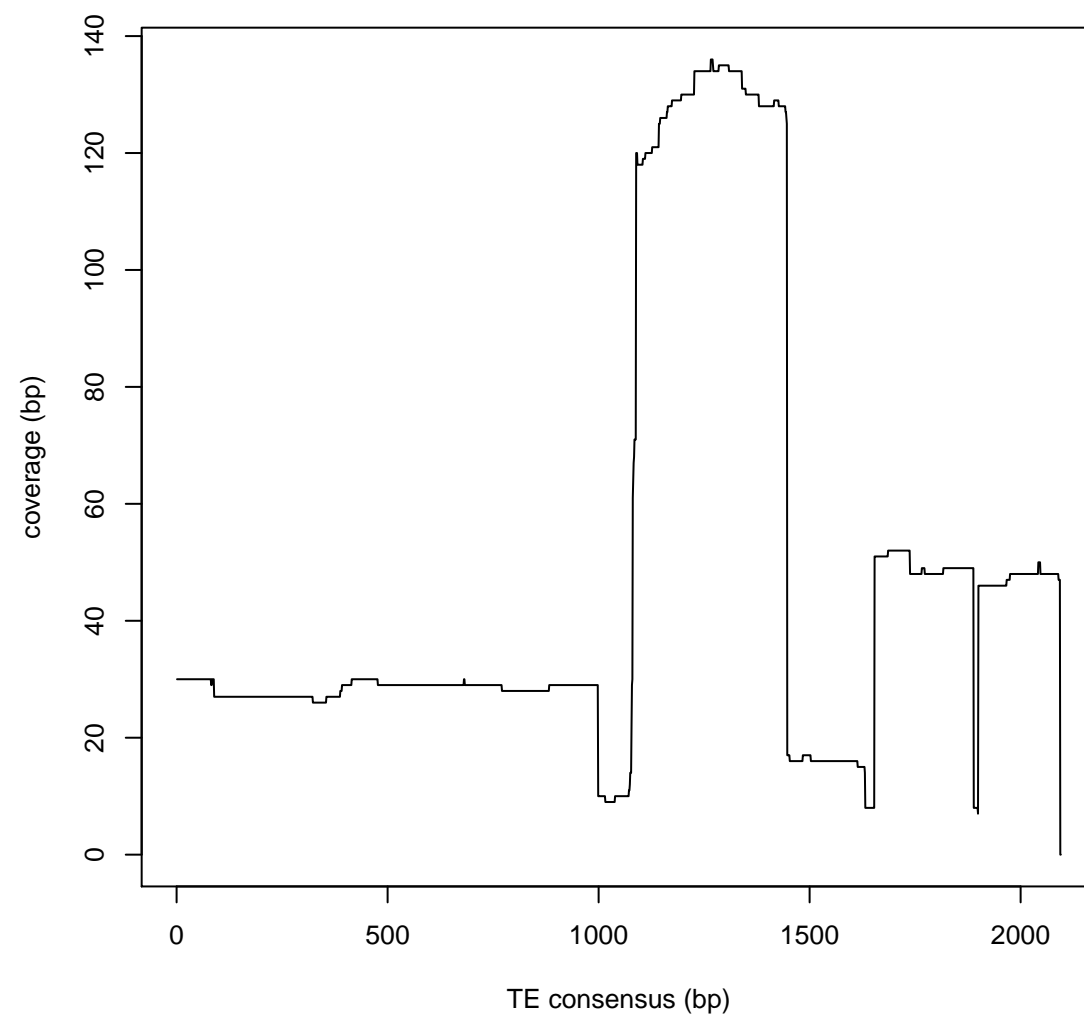


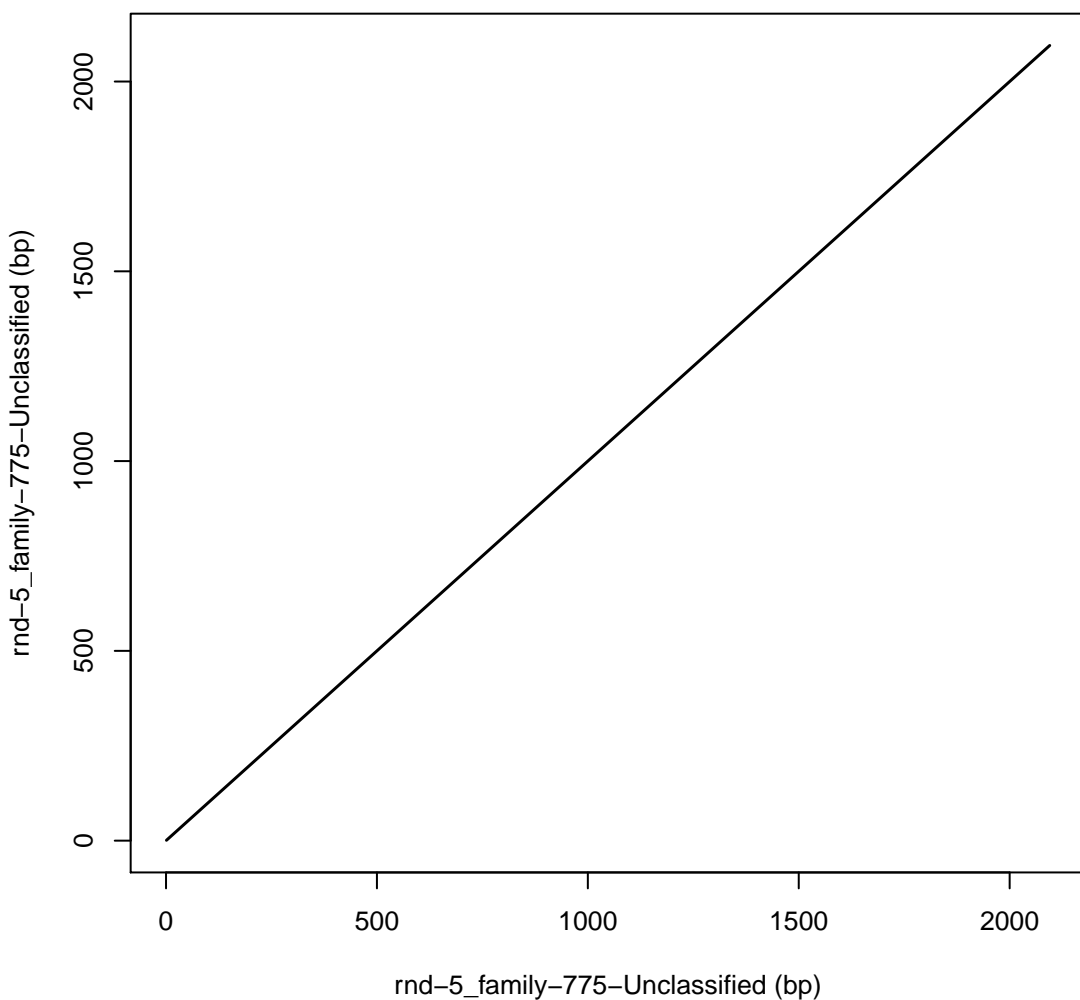
TE: rnd-5\_family-775-Unclassified  
consensus size: 2095bp; fragments: 260; full length: 6 ( $\geq 1885.5$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

