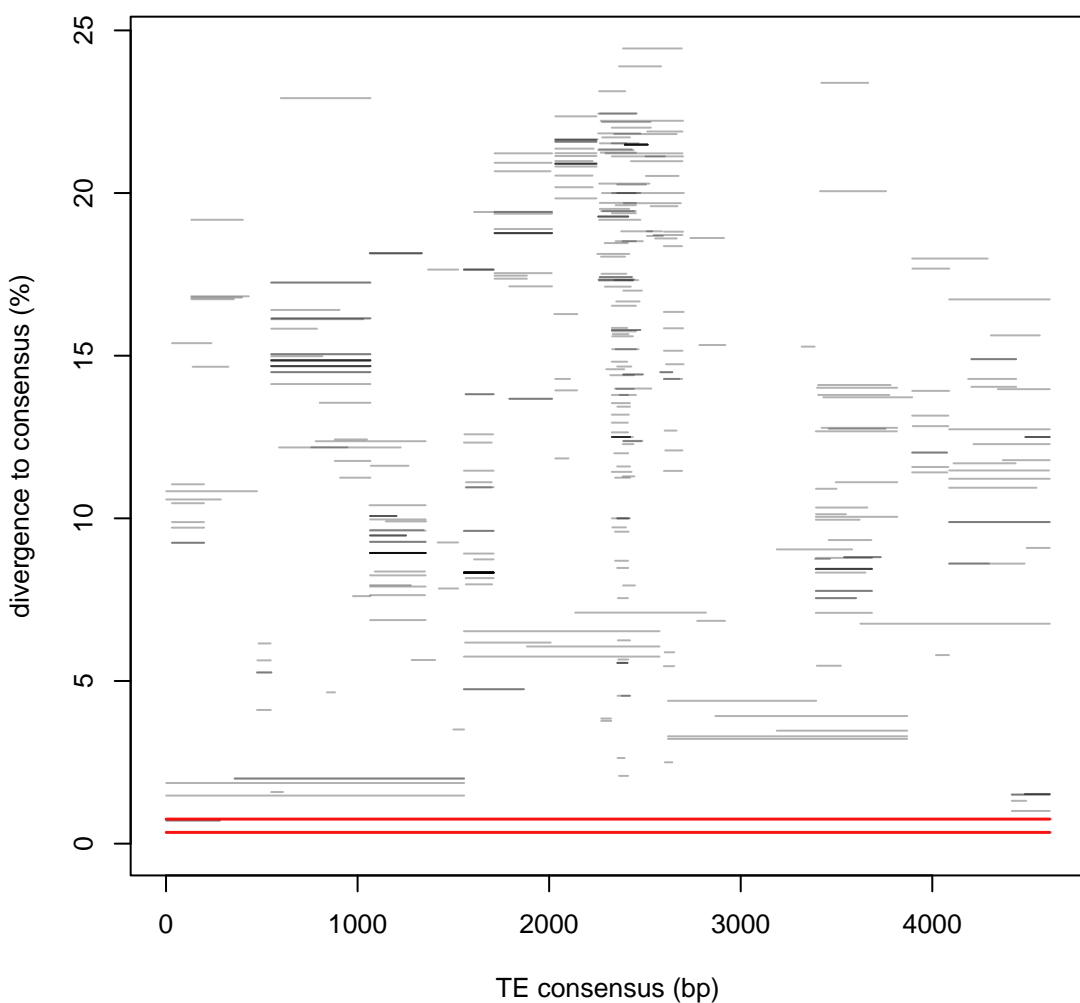
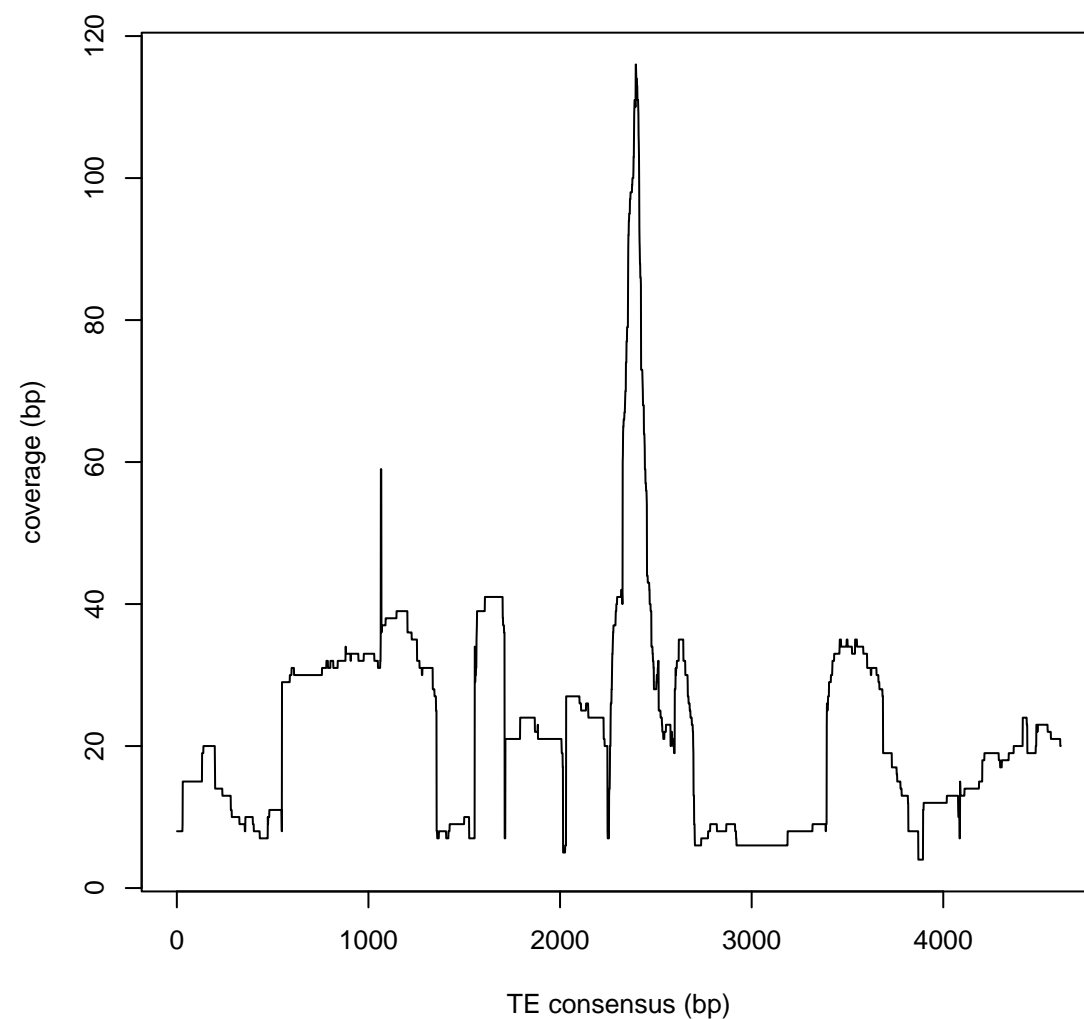


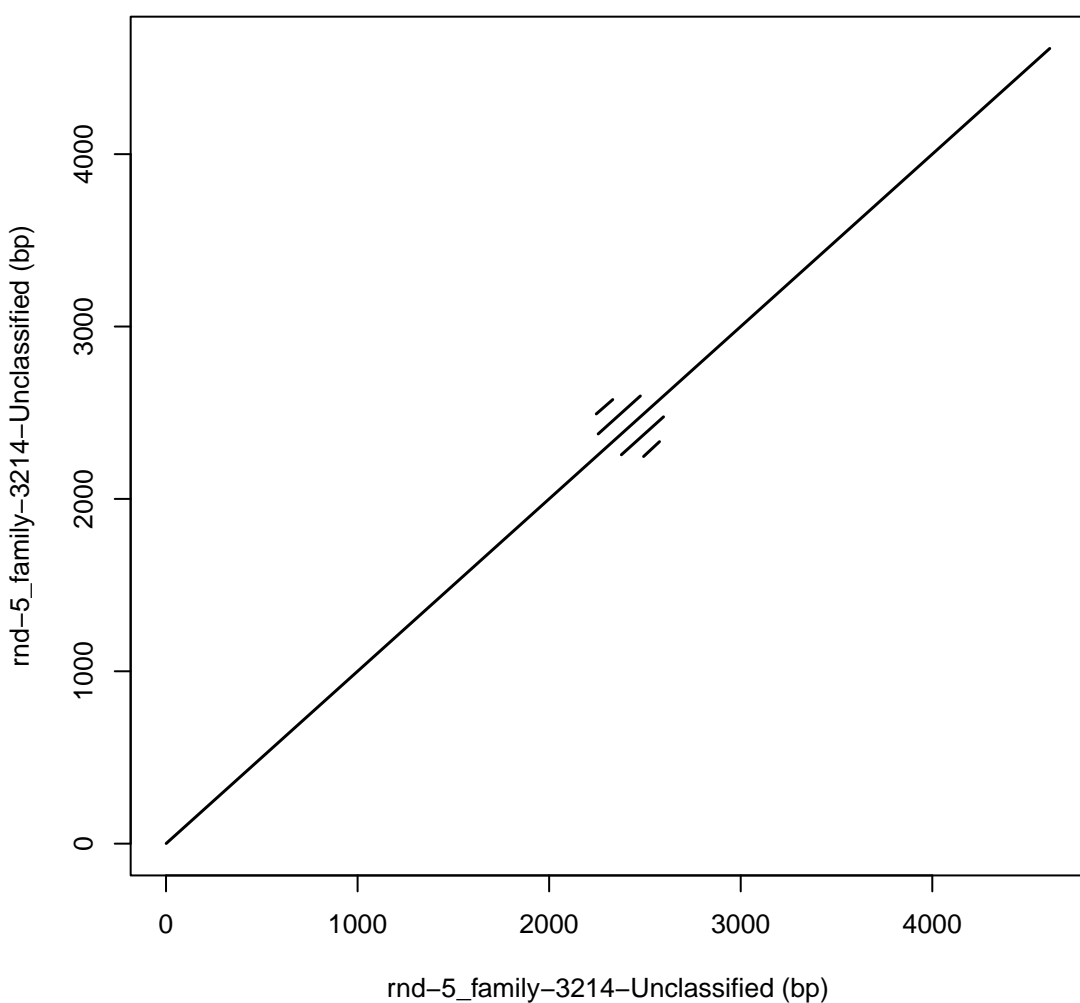
TE: rnd-5\_family-3214-Unclassified  
consensus size: 4613bp; fragments: 395; full length: 2 ( $\geq 4151.7$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

