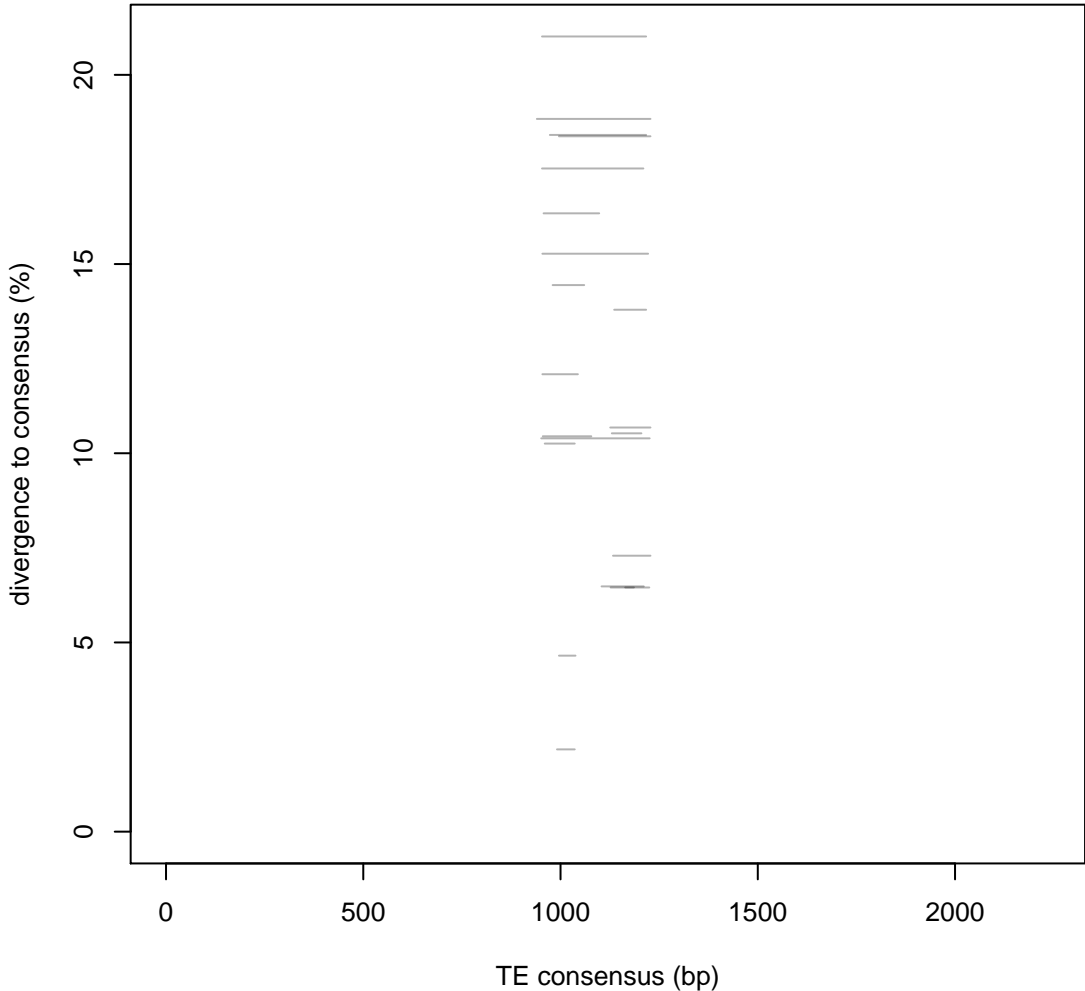
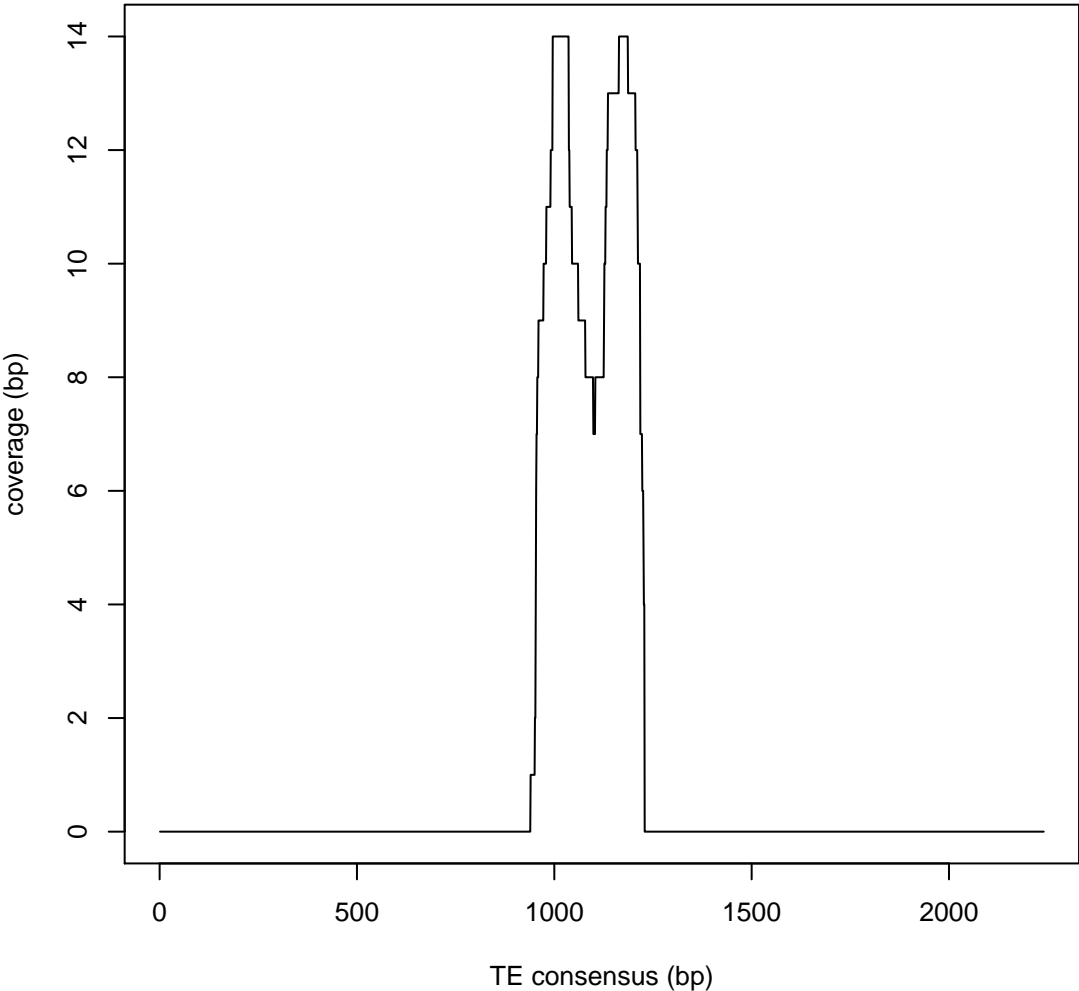


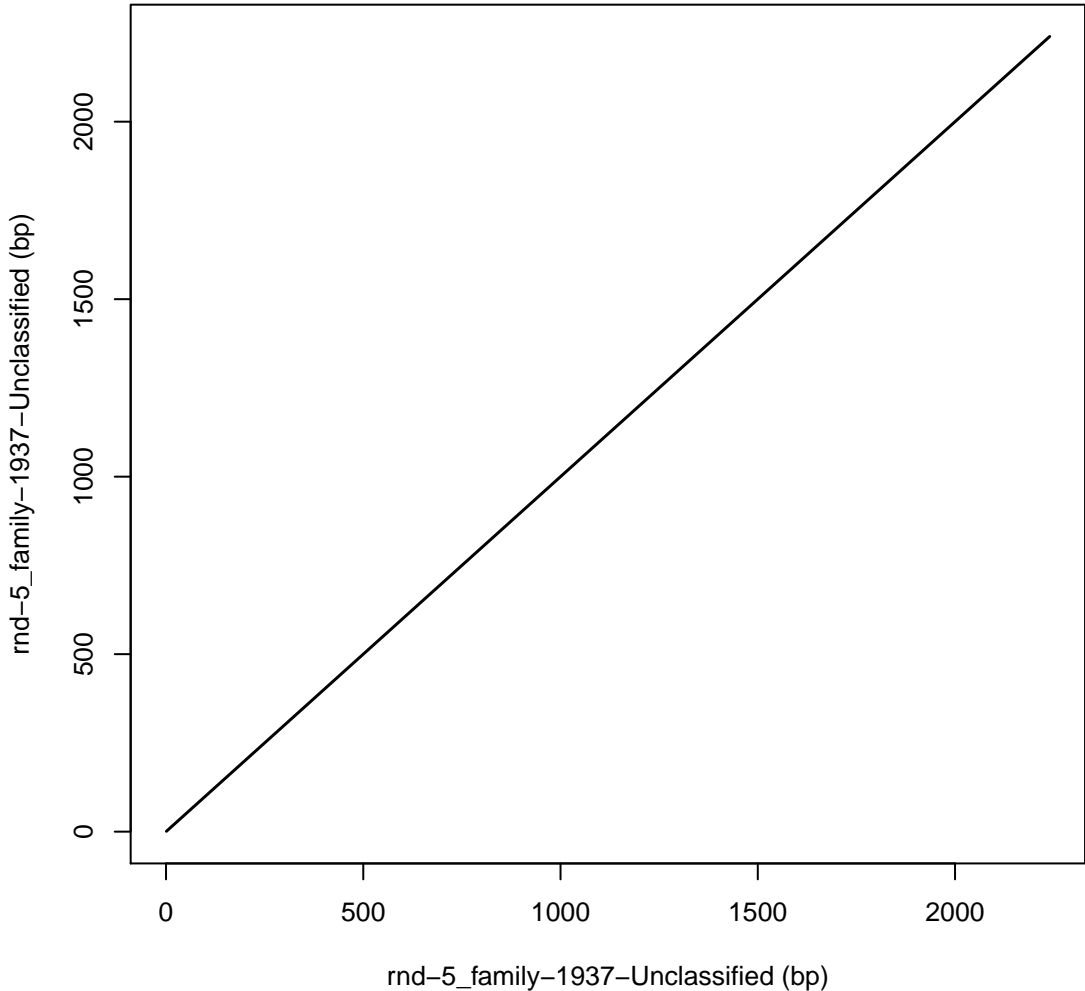
TE: rnd-5\_family-1937-Unclassified  
consensus size: 2240bp; fragments: 21; full length: 0 (>=2016bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

