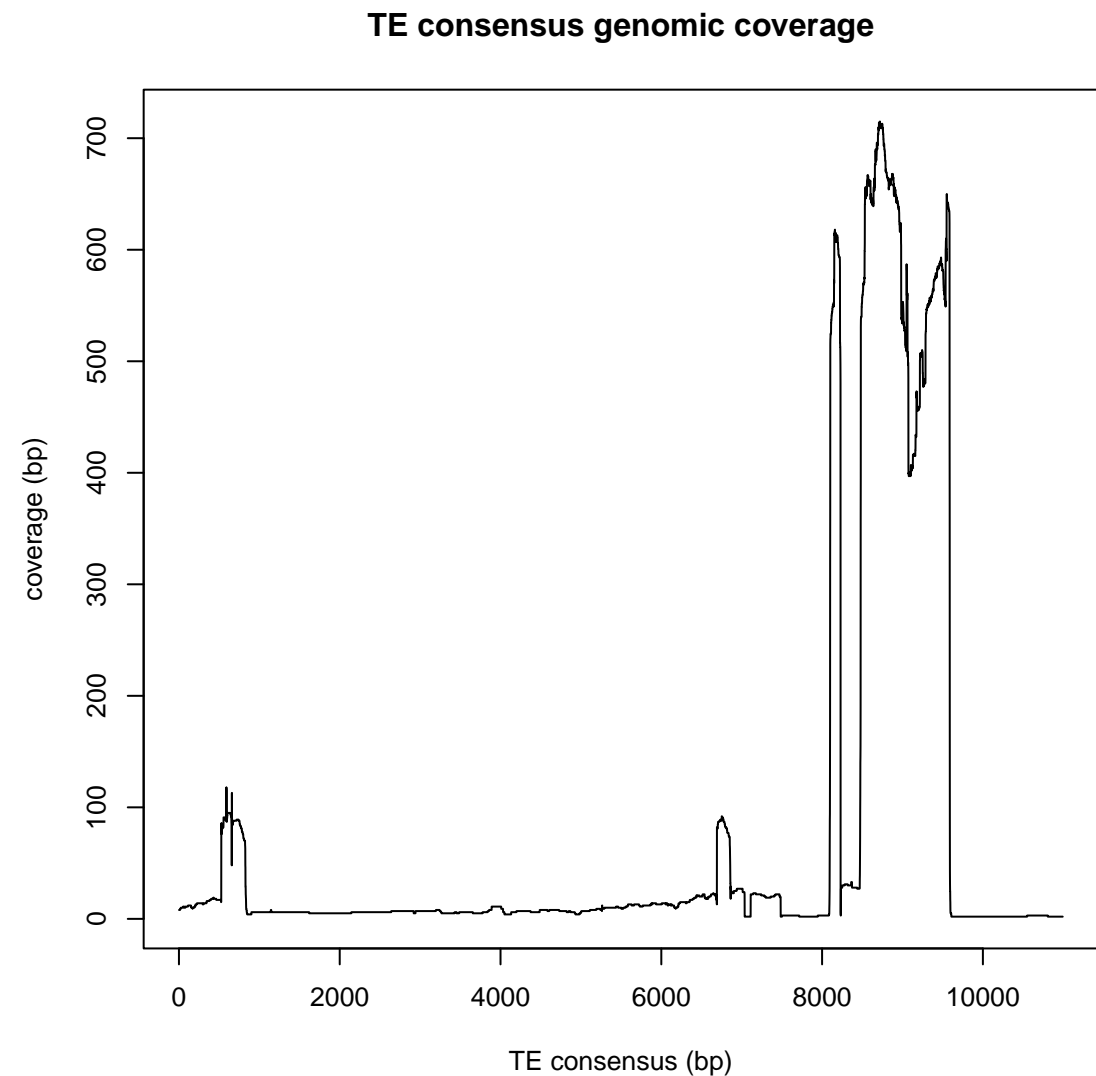
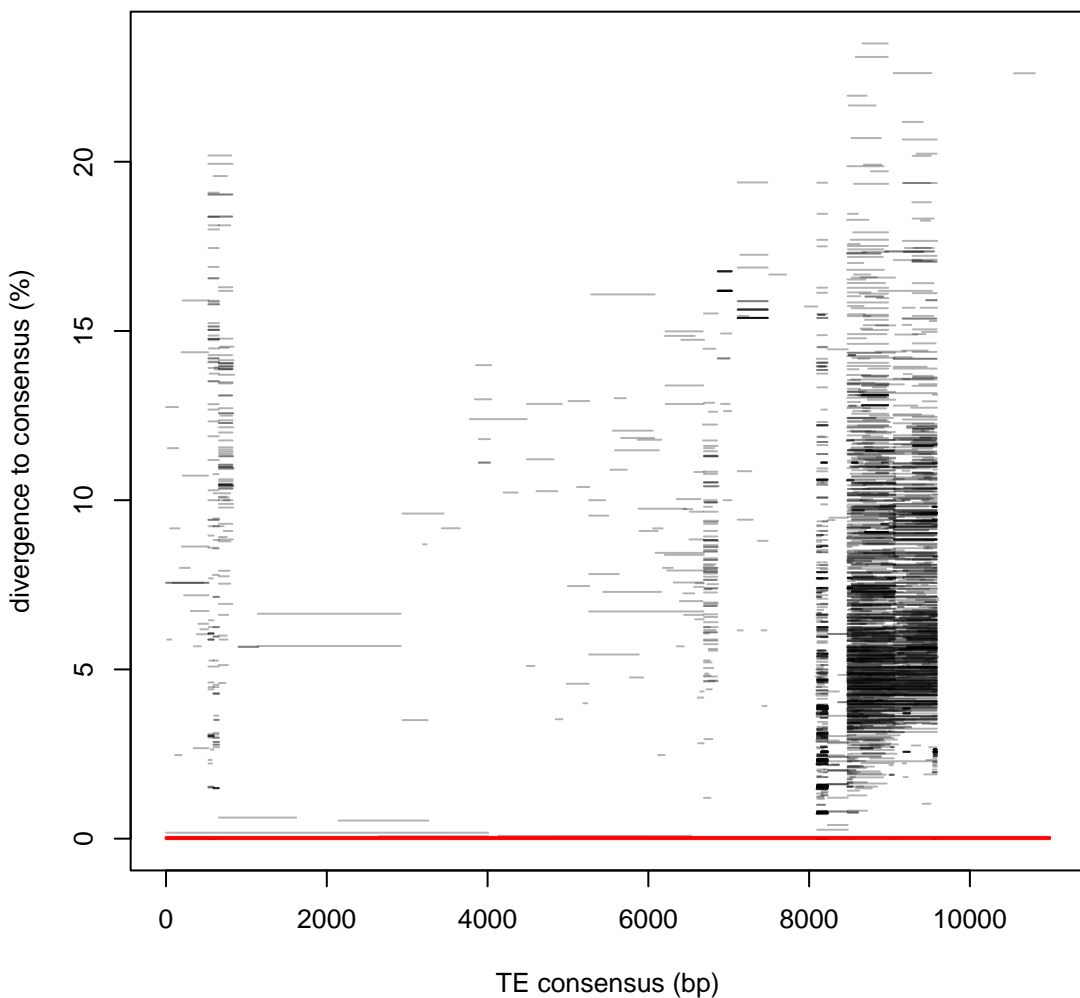
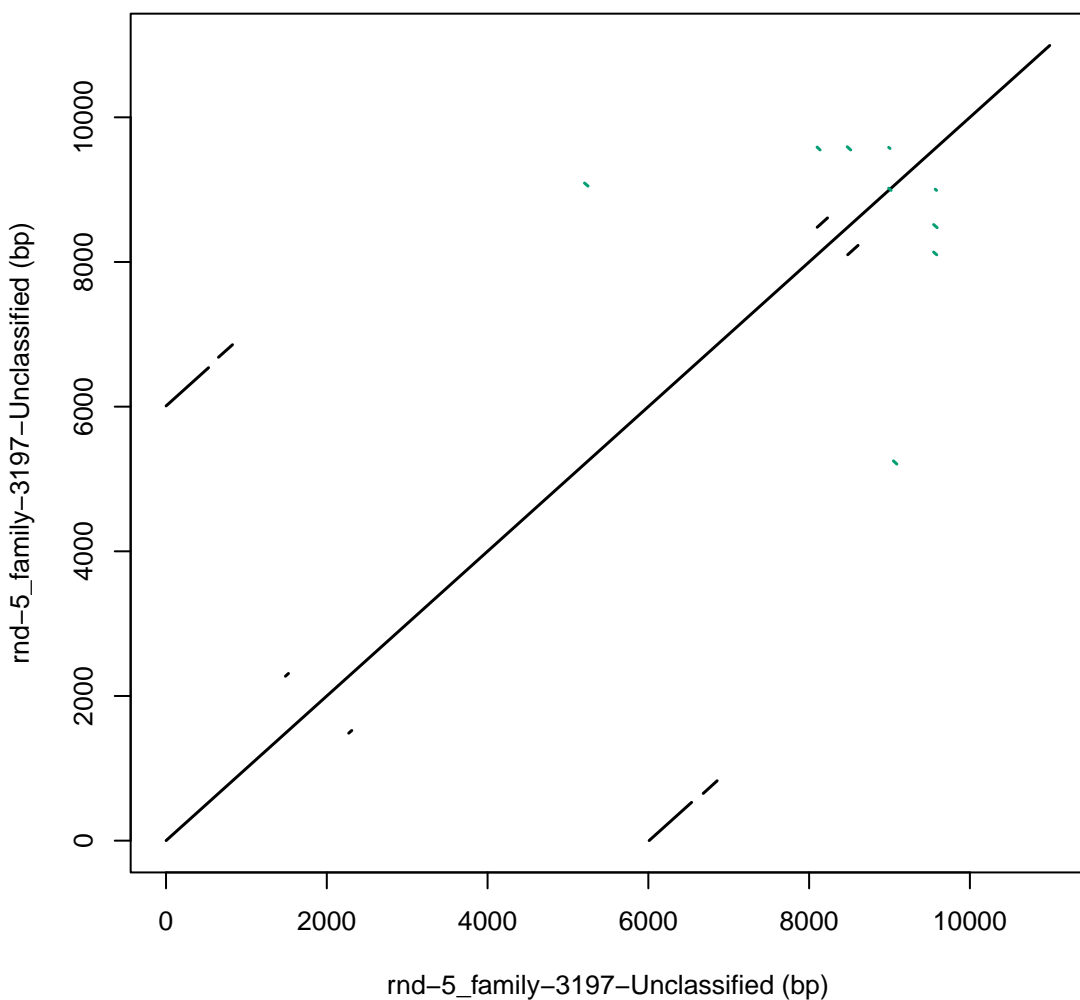


TE: rnd-5\_family-3197-Unclassified  
consensus size: 10993bp; fragments: 2858; full length: 2 (>=9893.7bp)



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

