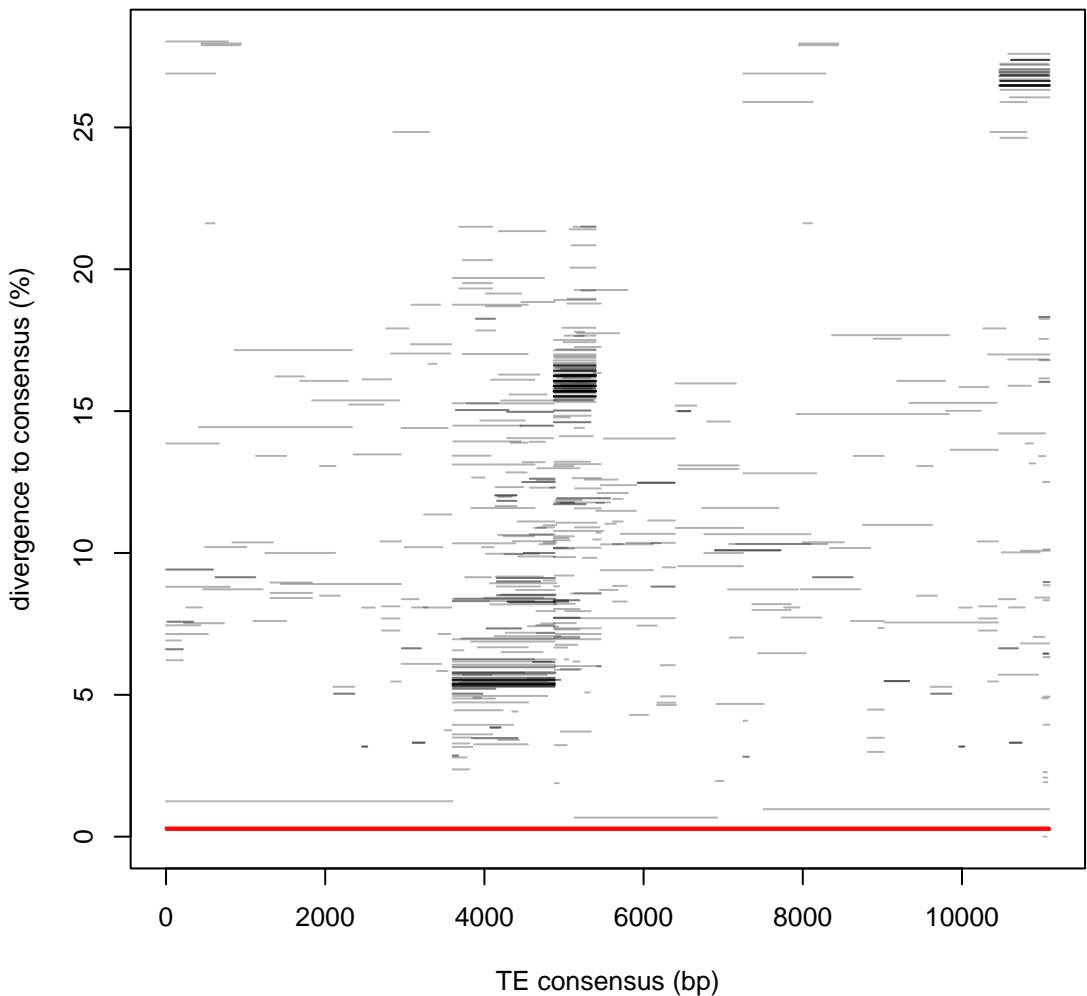
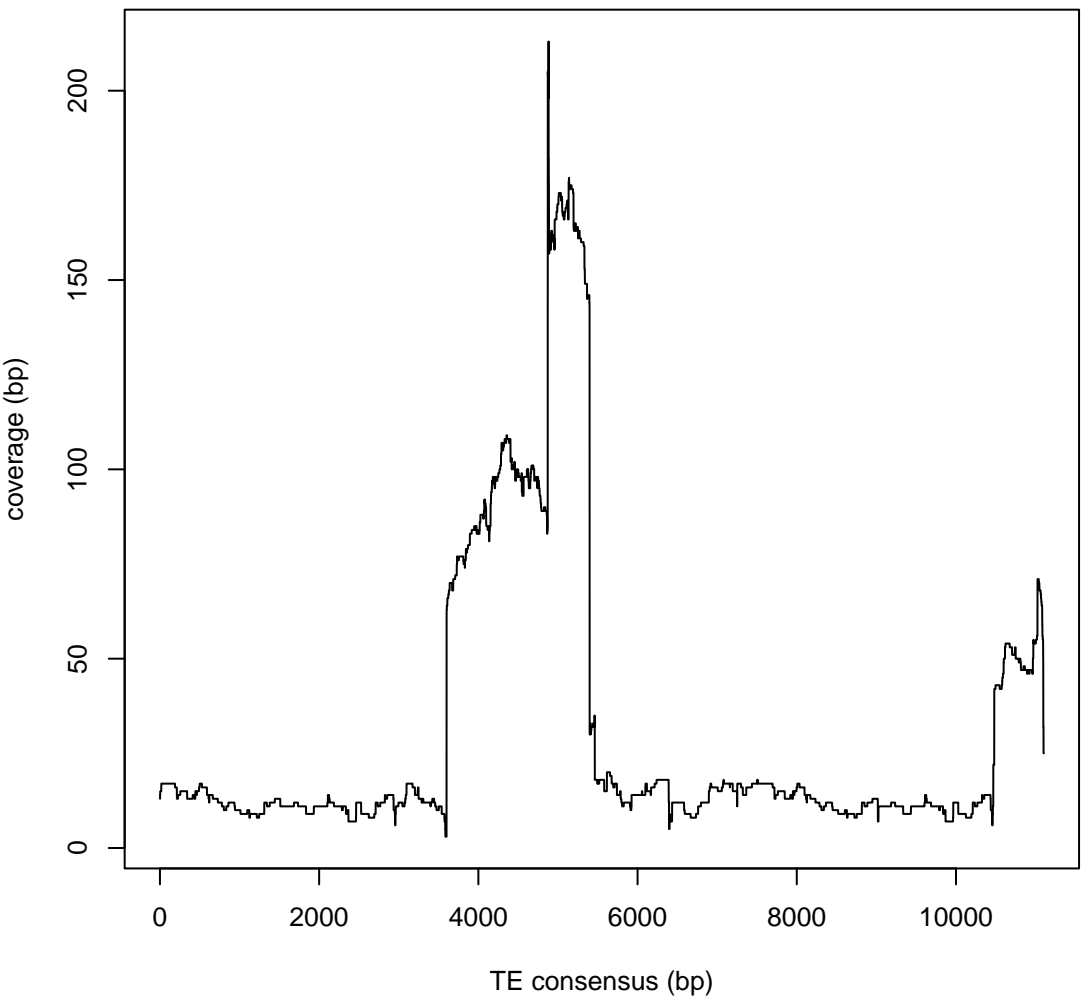


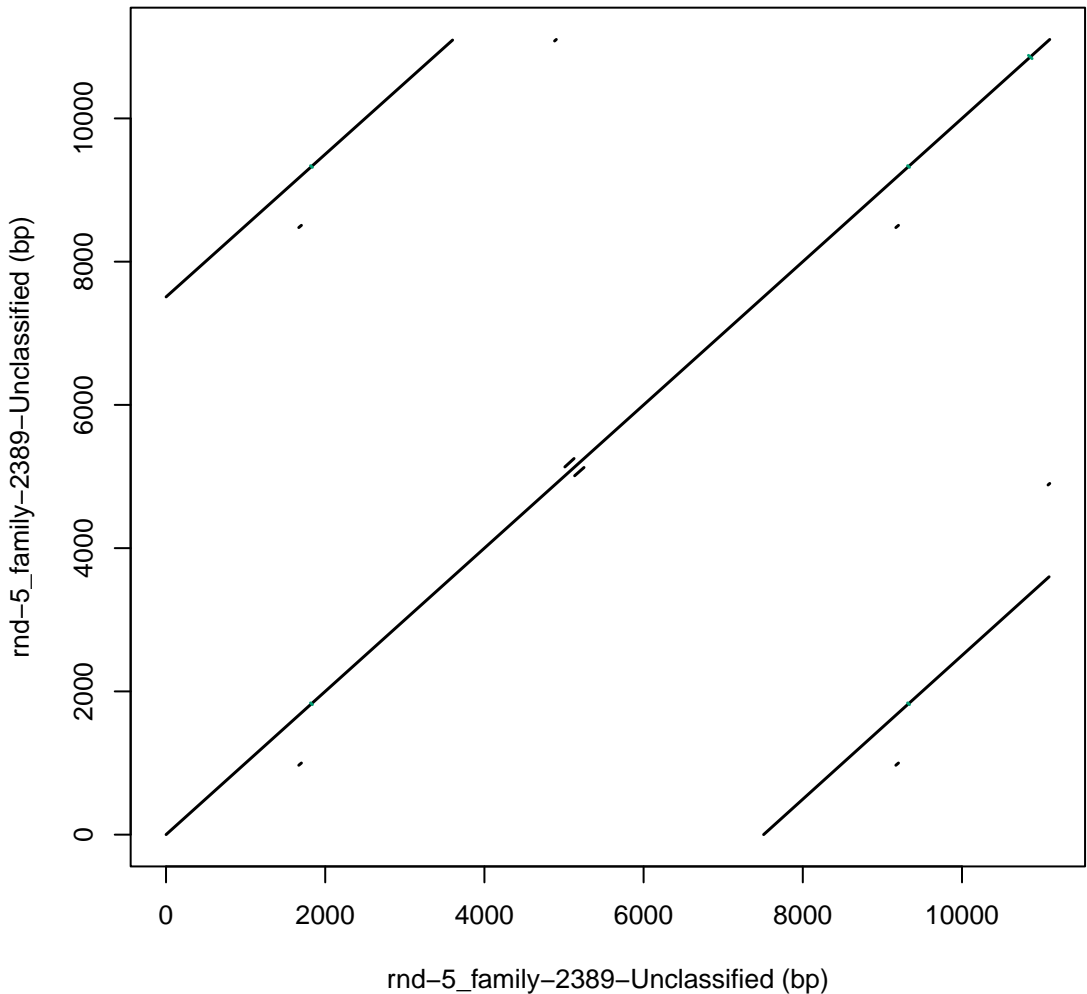
TE: rnd-5\_family-2389-Unclassified  
consensus size: 11102bp; fragments: 644; full length: 2 (>=9991.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

