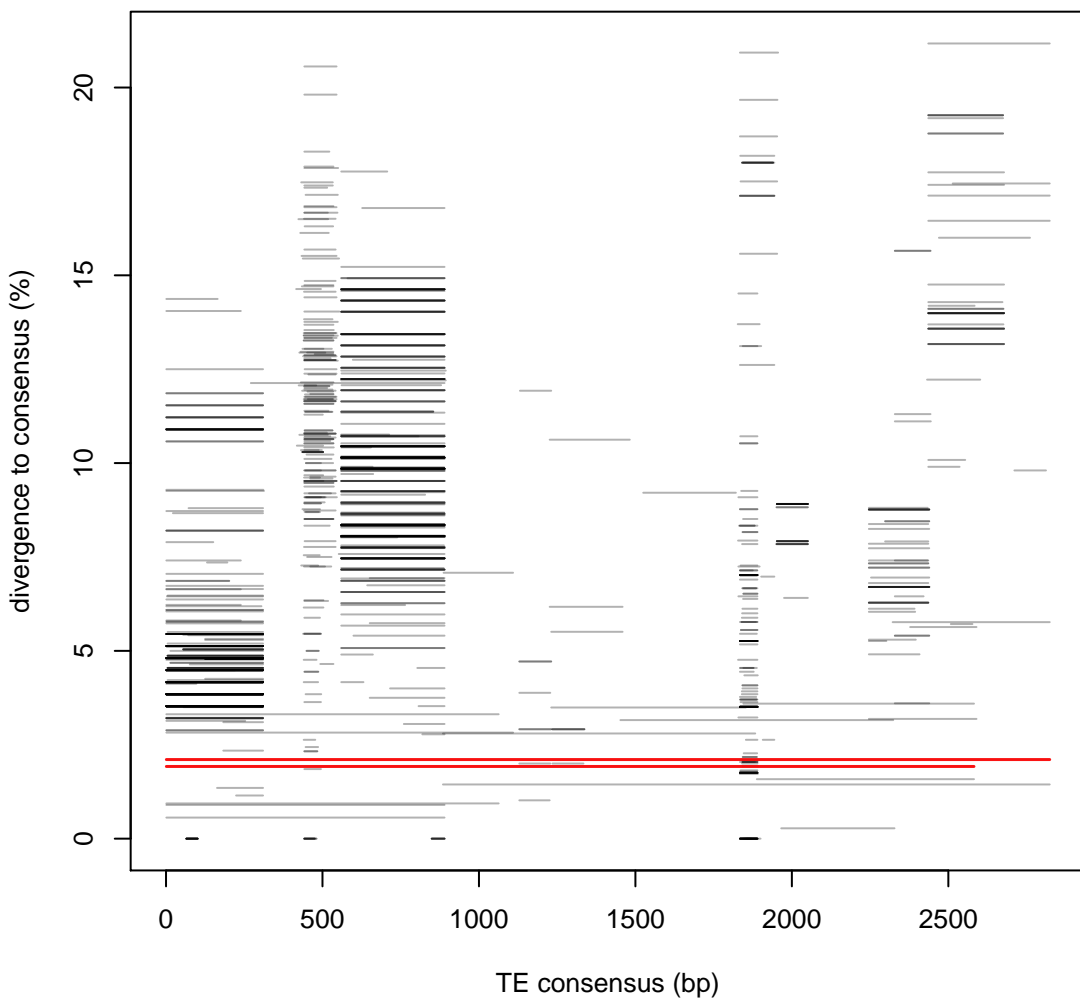
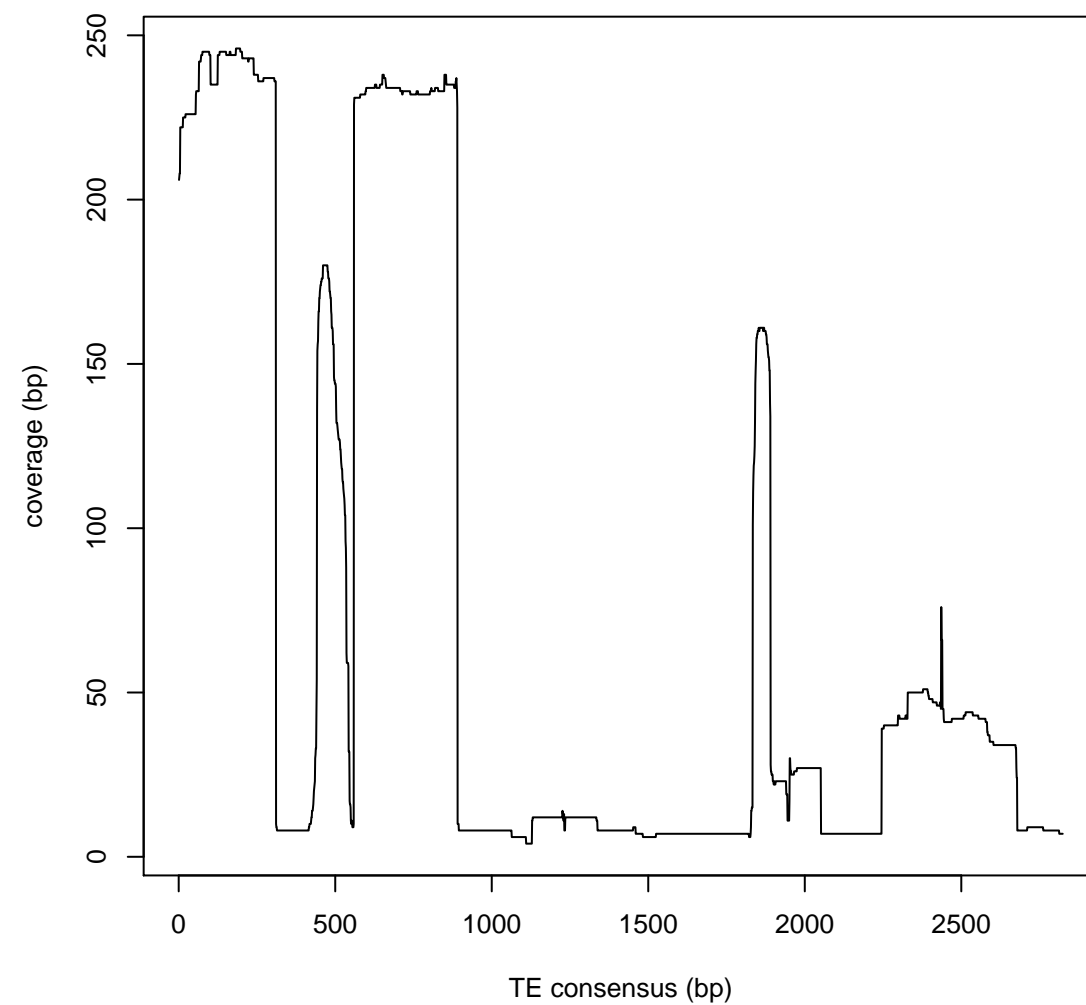


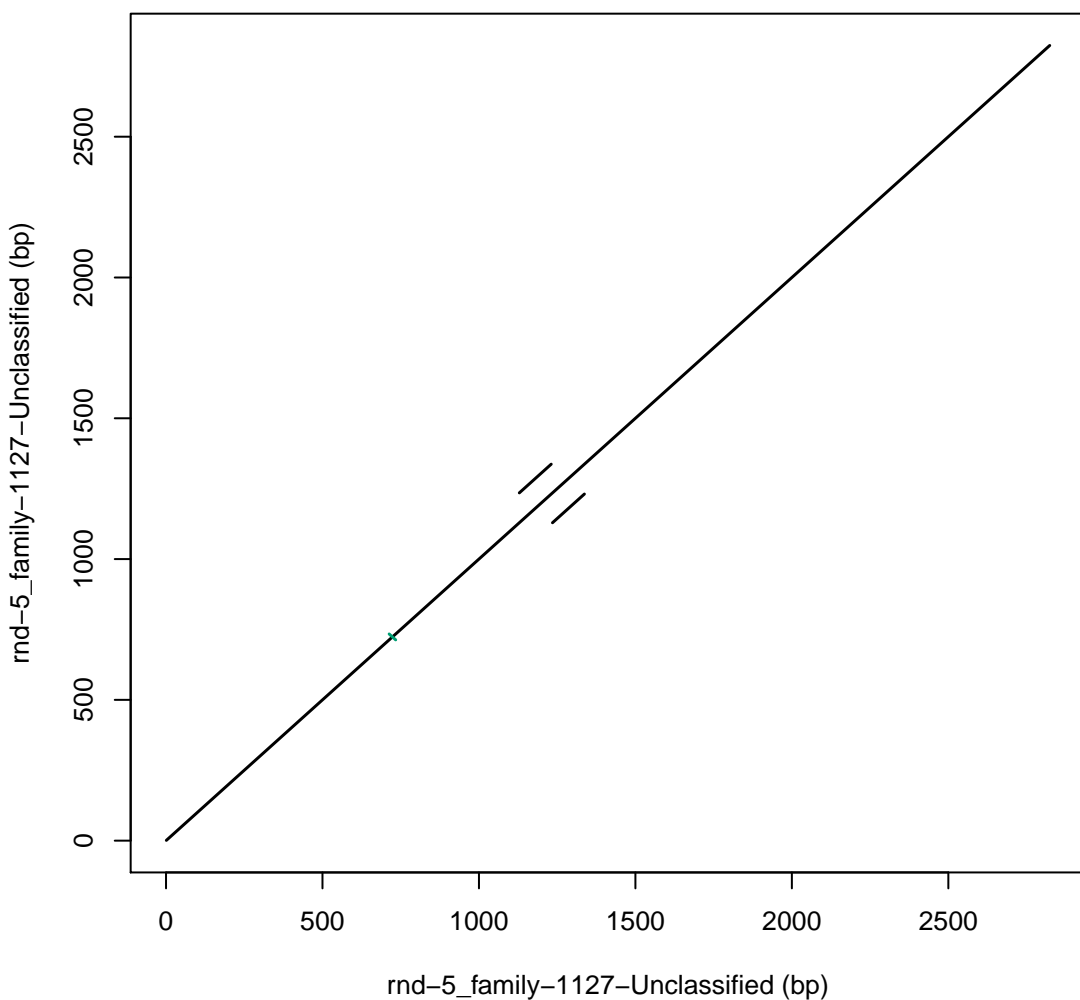
TE: rnd-5\_family-1127-Unclassified  
consensus size: 2824bp; fragments: 961; full length: 2 (>=2541.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

