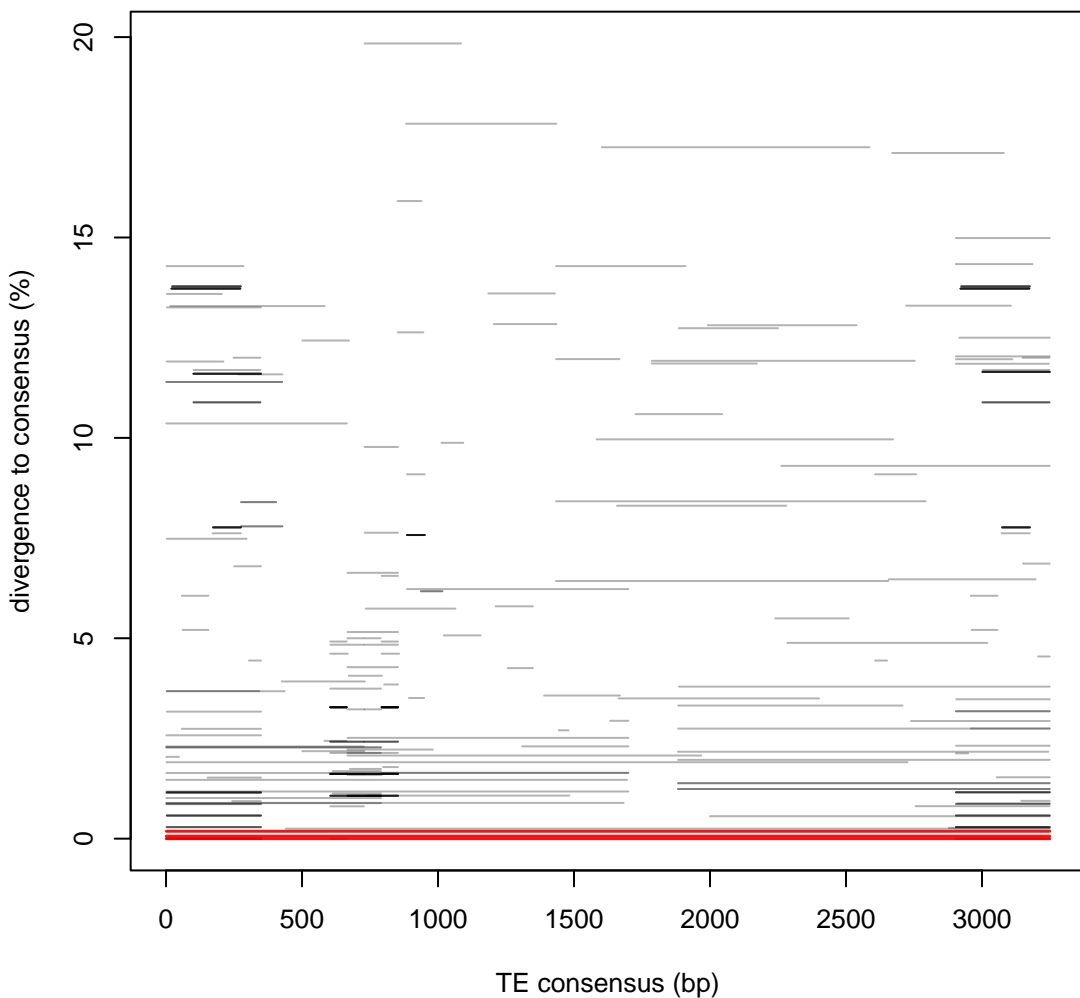
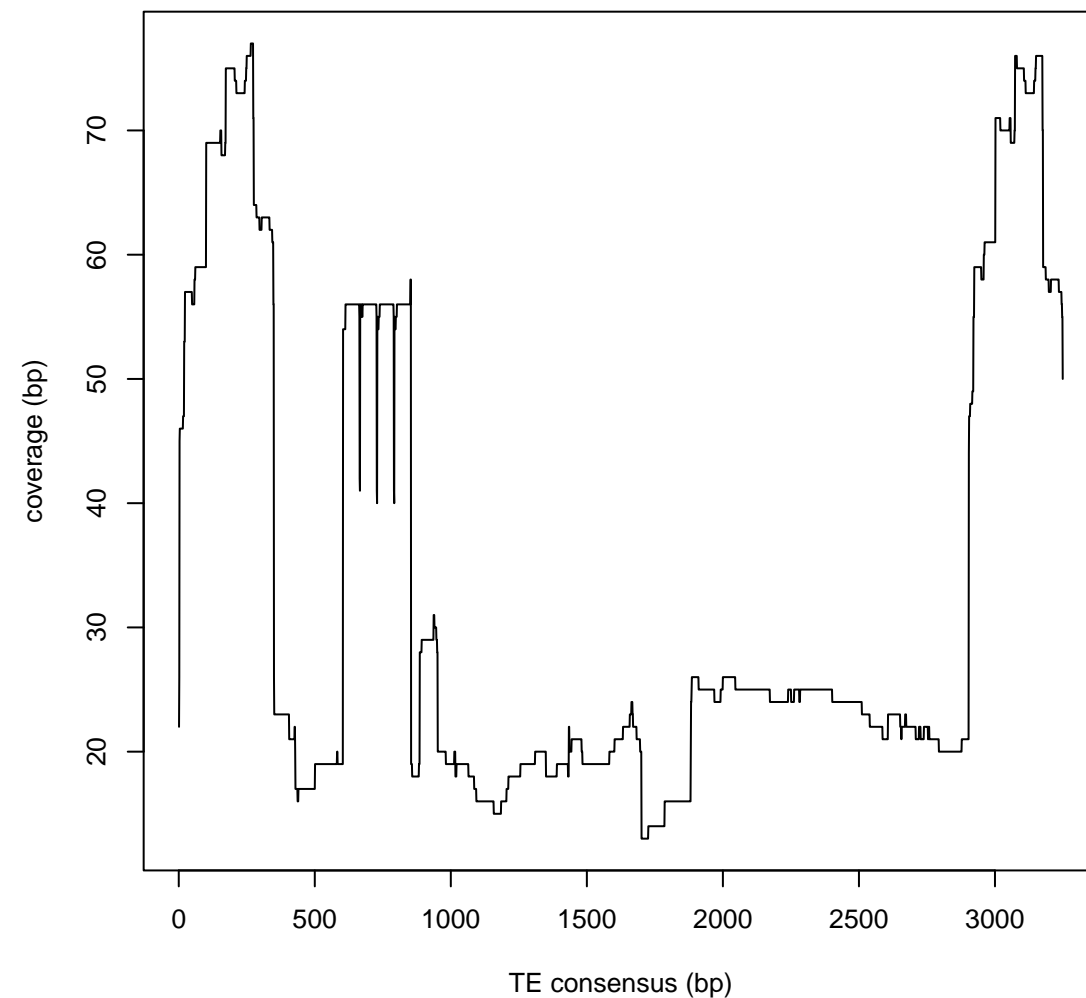


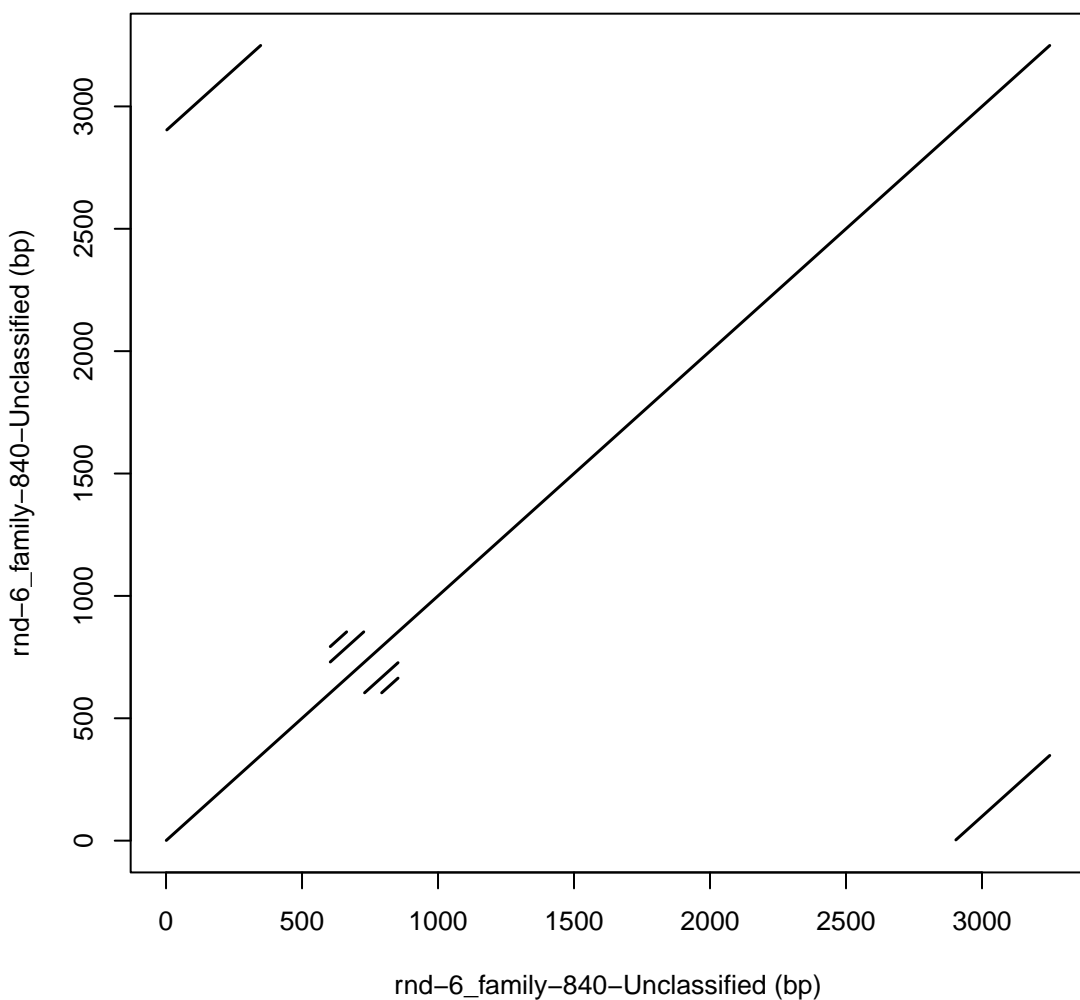
TE: rnd-6\_family-840-Unclassified  
consensus size: 3249bp; fragments: 305; full length: 3 (>=2924.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

