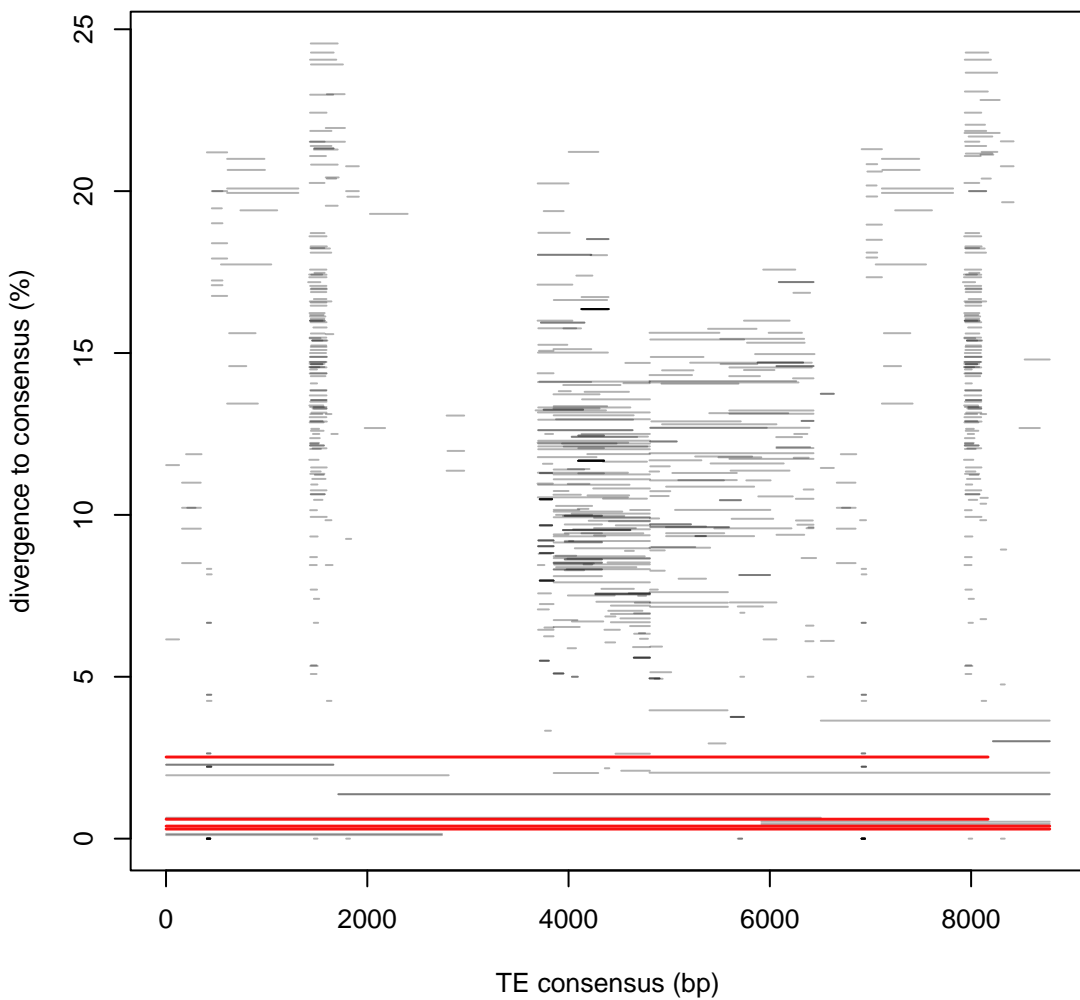
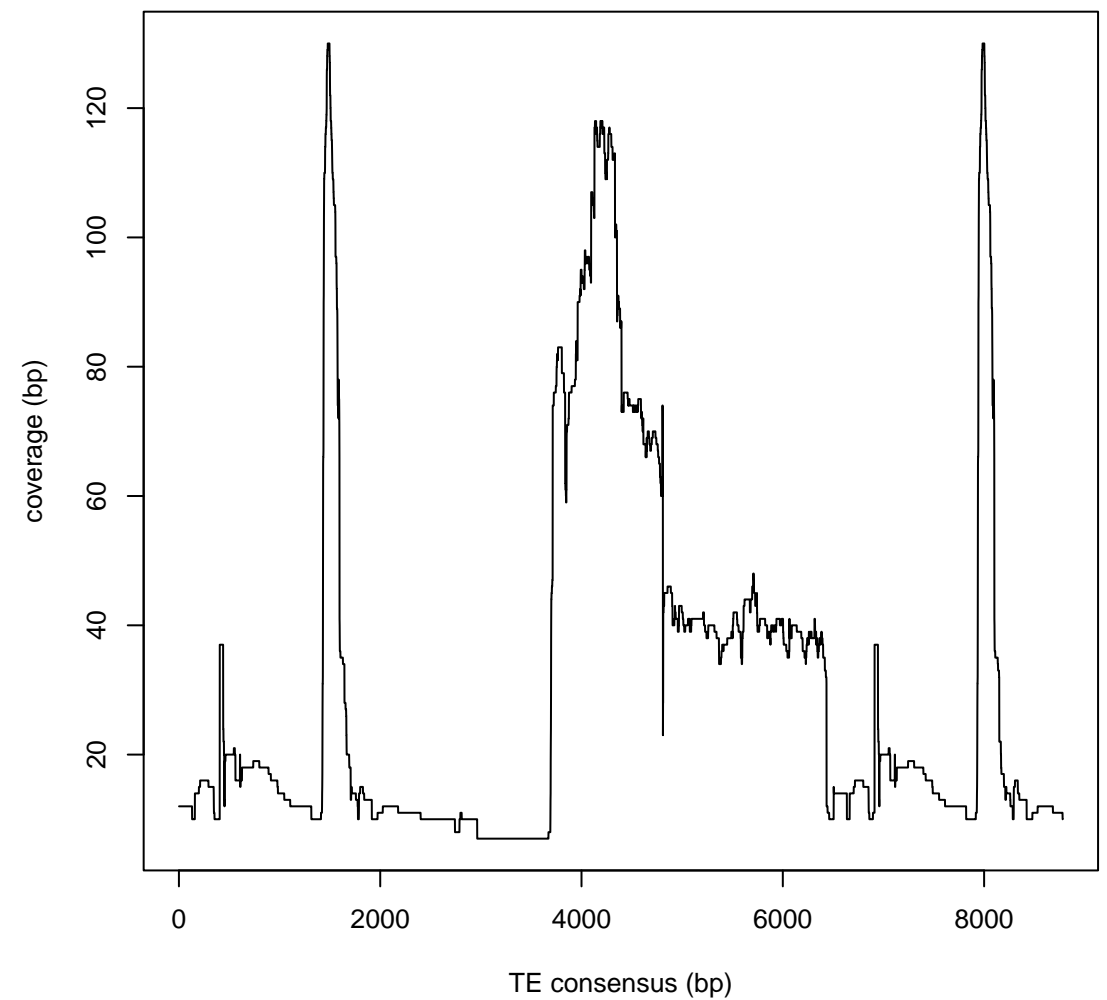


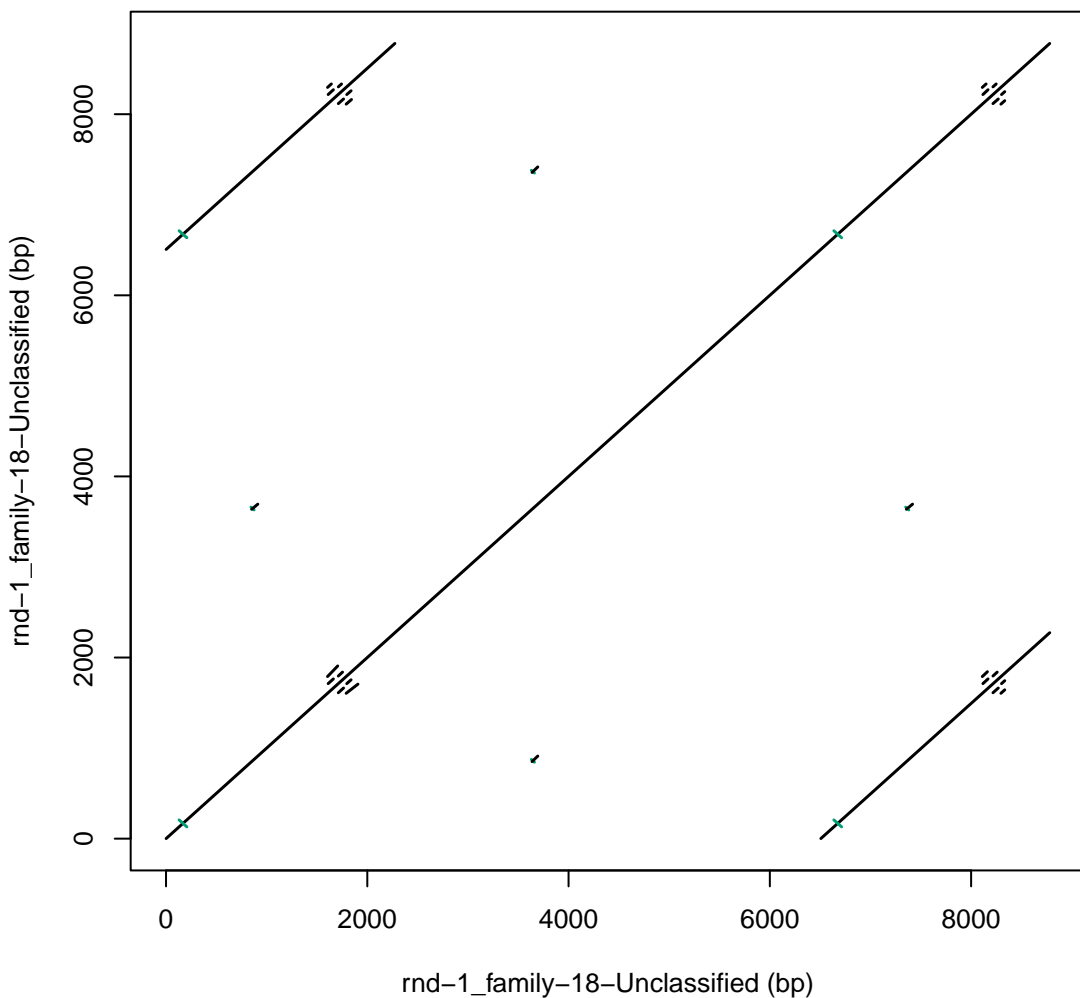
TE: rnd-1\_family-18-Unclassified  
consensus size: 8781bp; fragments: 752; full length: 4 (>=7902.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

