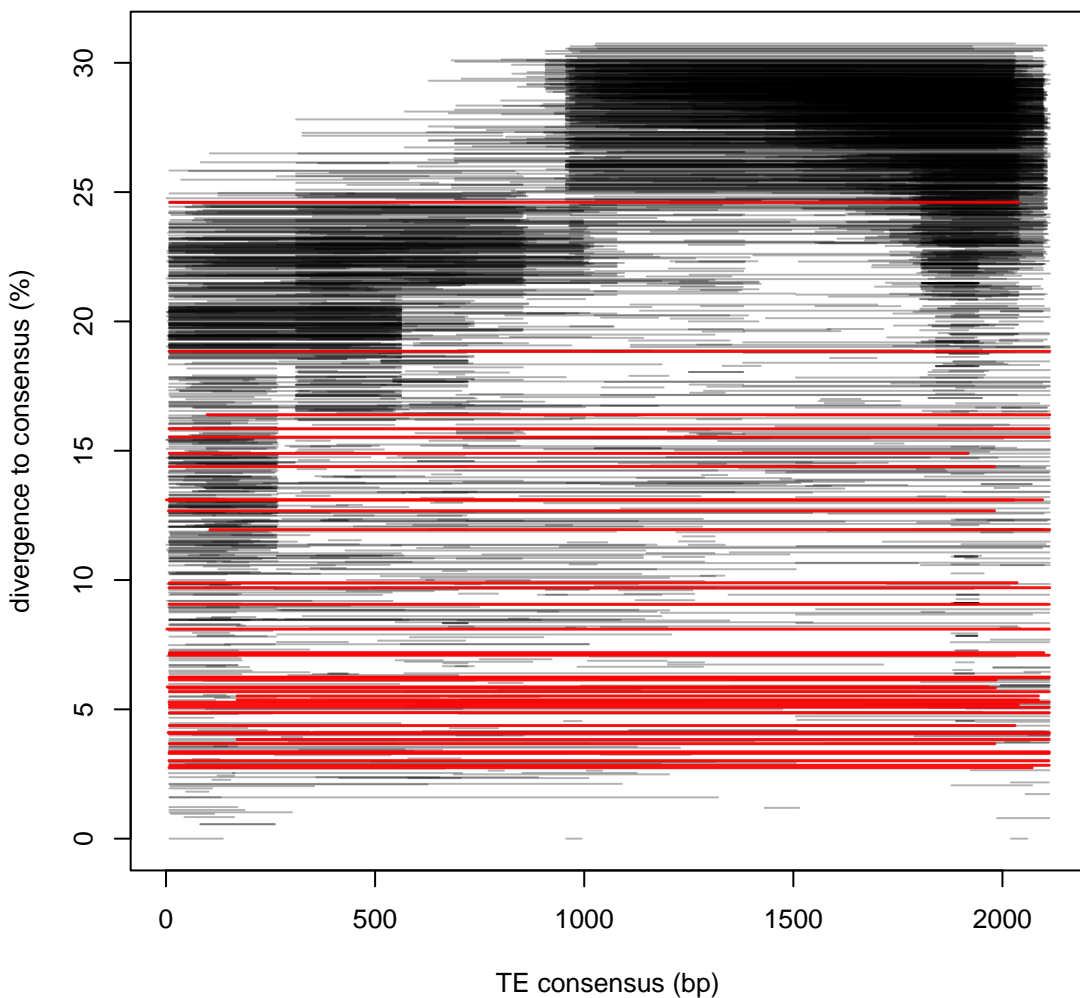
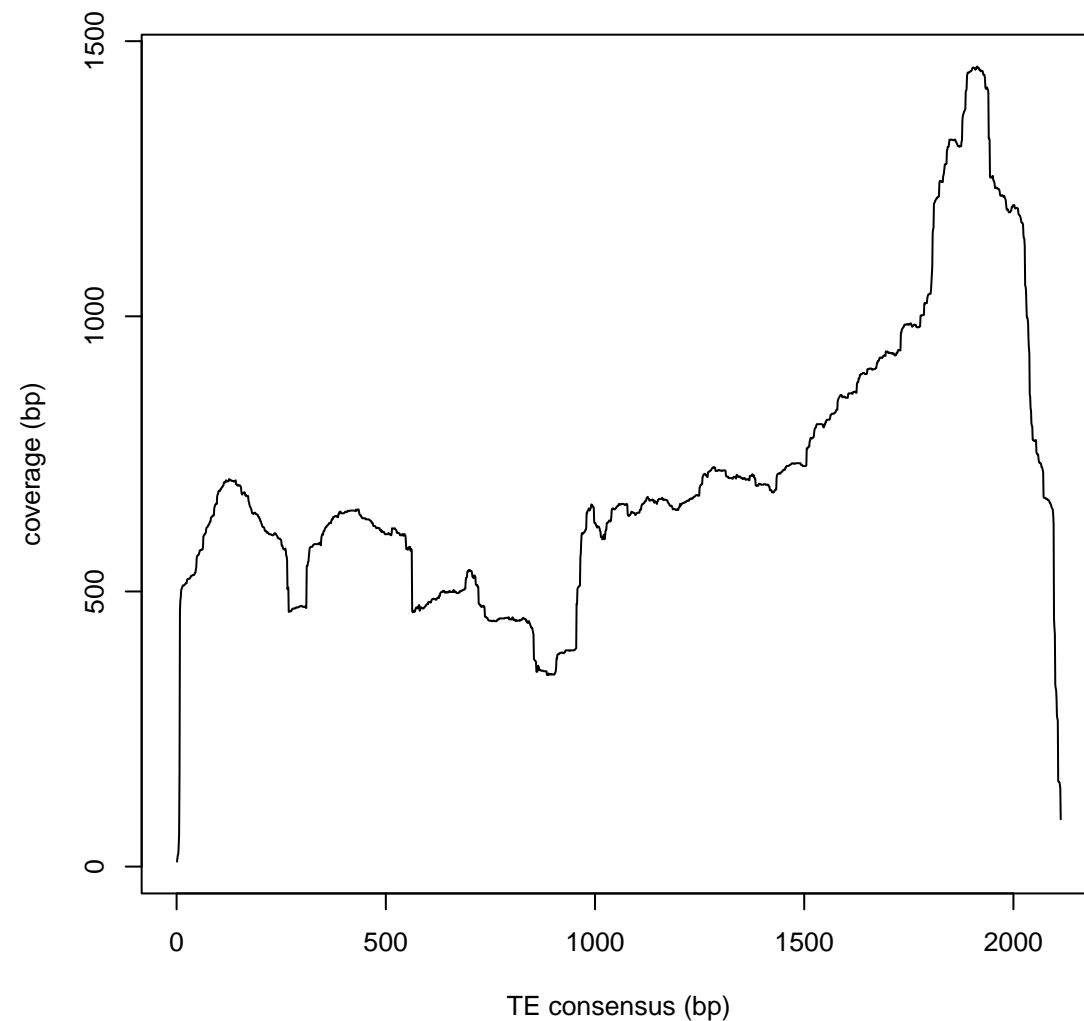


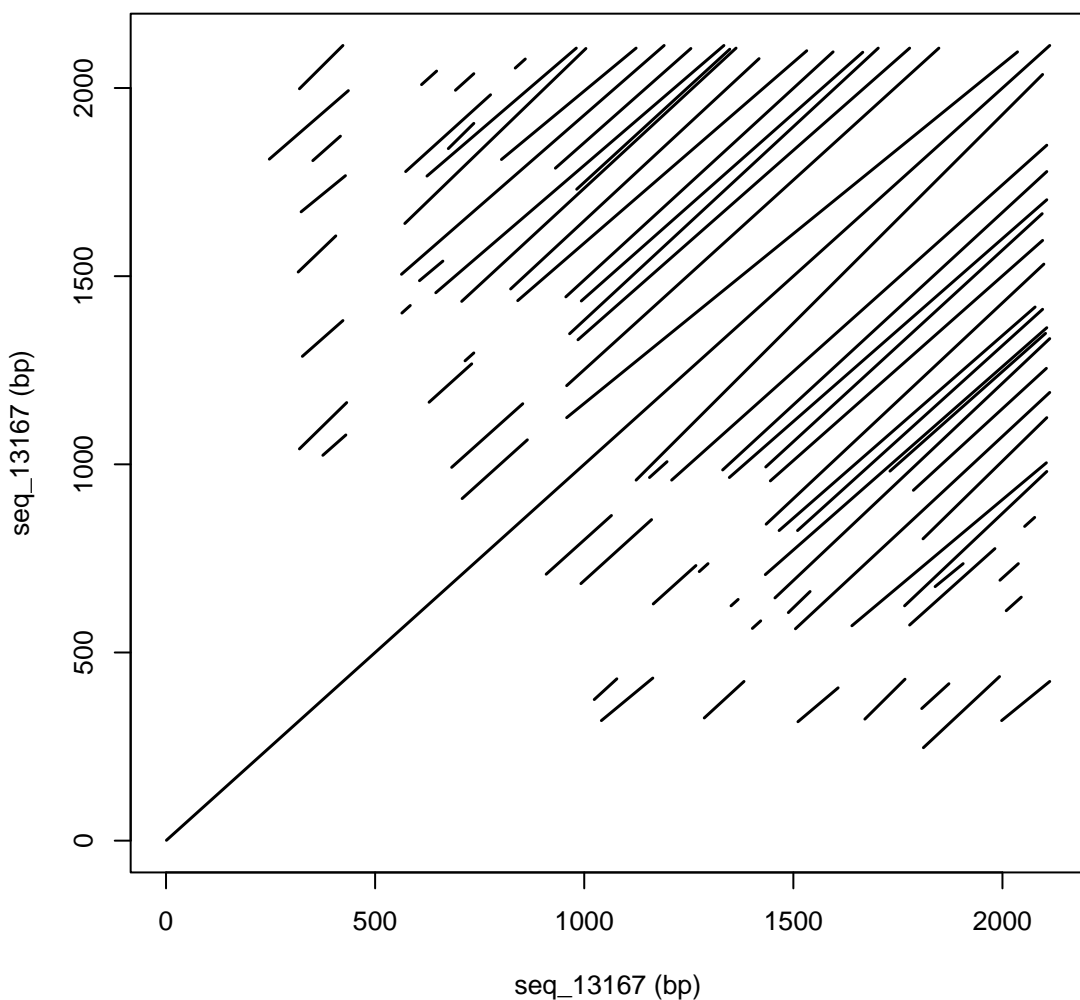
TE: seq_13167
consensus size: 2113bp; fragments: 2780; full length: 40 (>=1901.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

