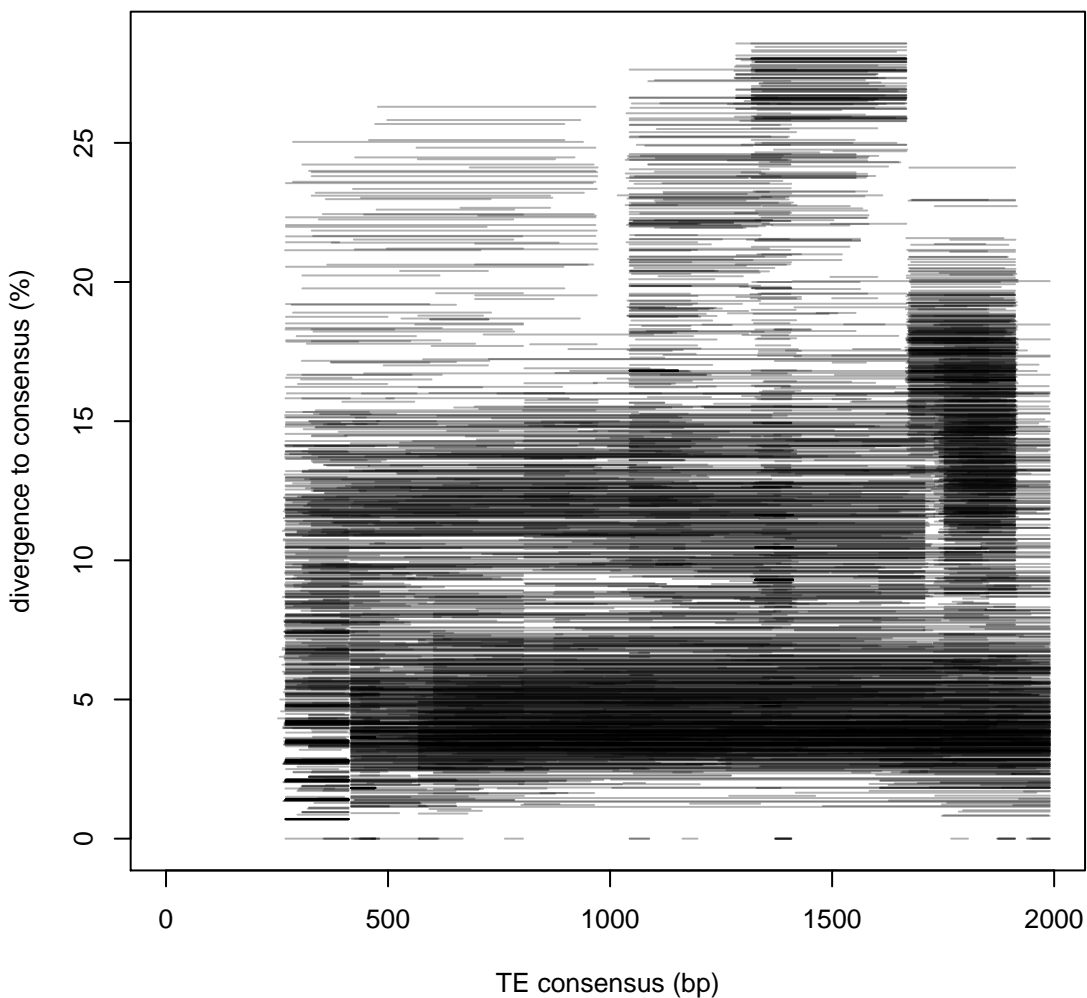
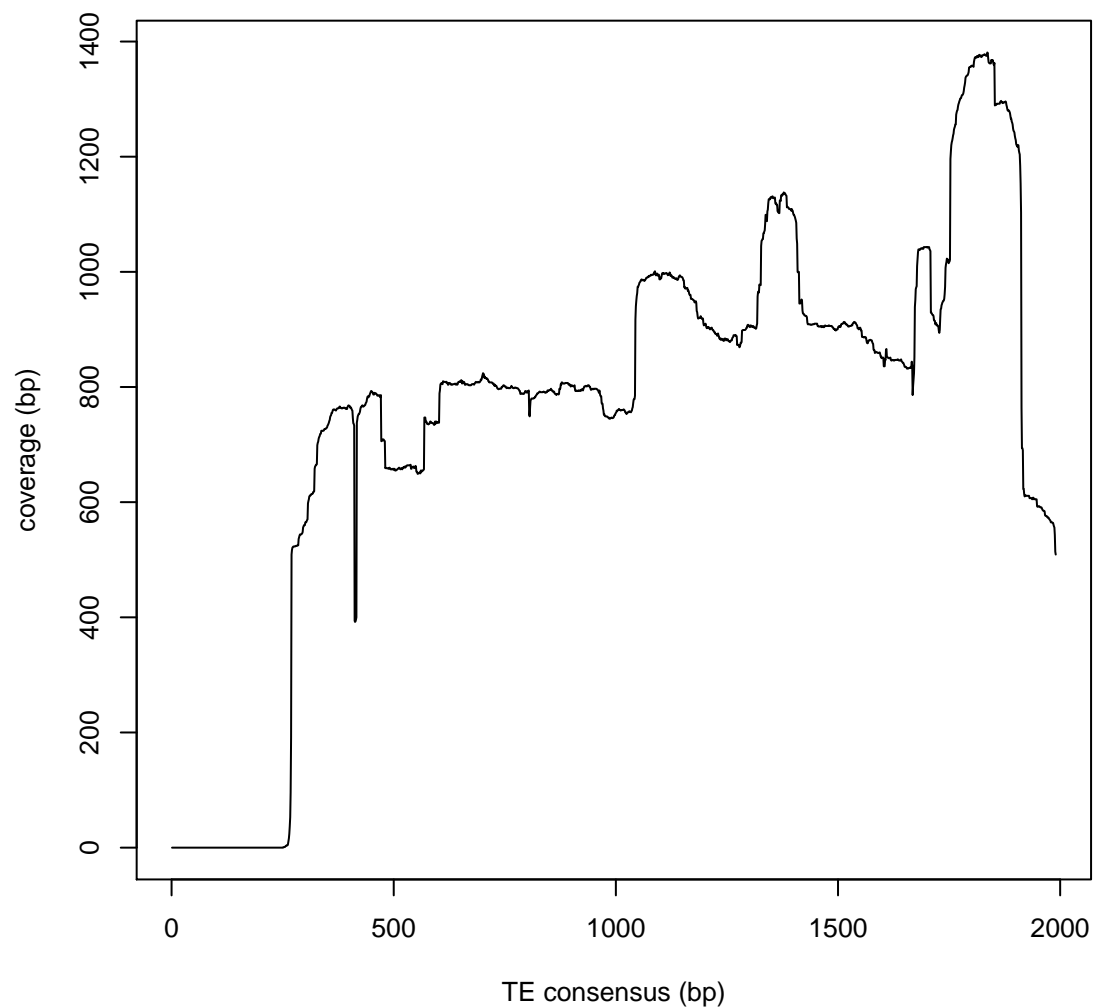


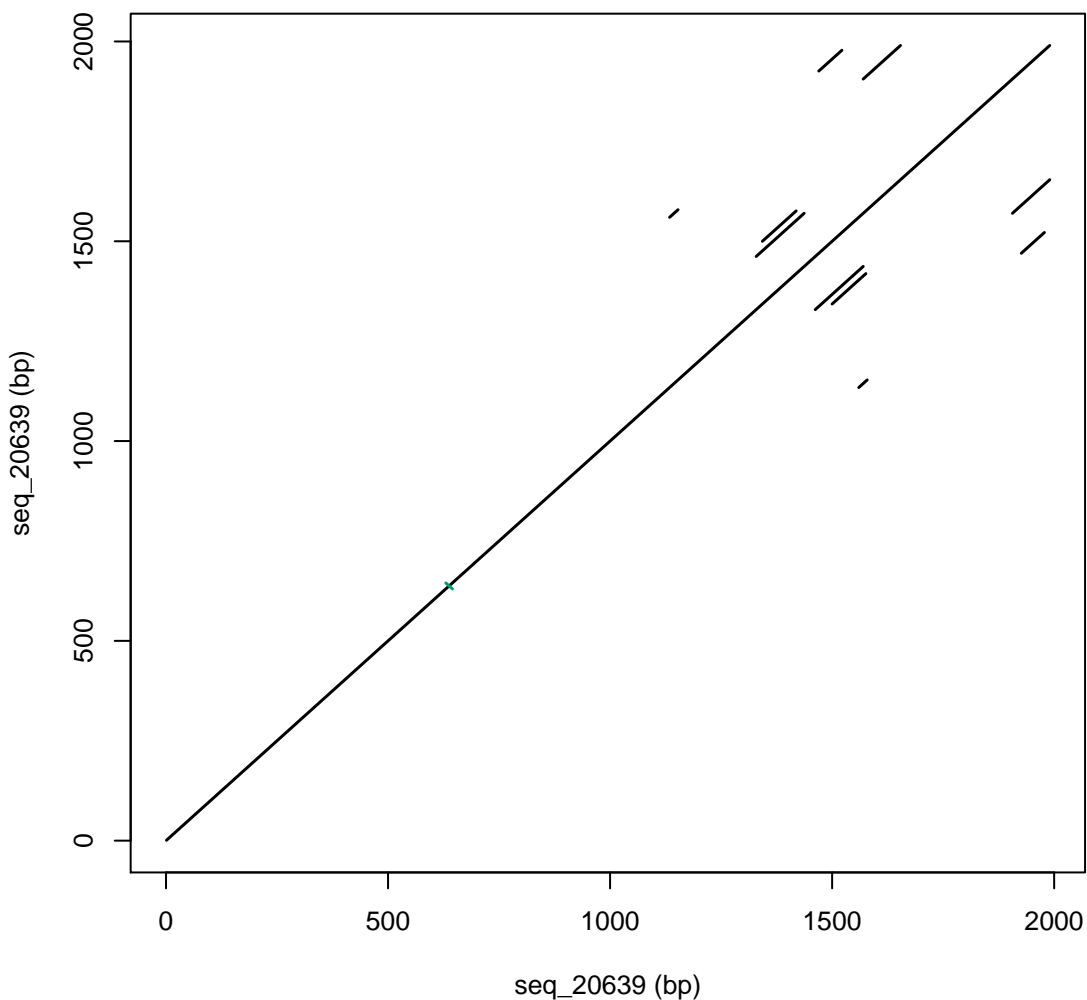
TE: seq_20639
consensus size: 1990bp; fragments: 3624; full length: 0 (>=1791bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

