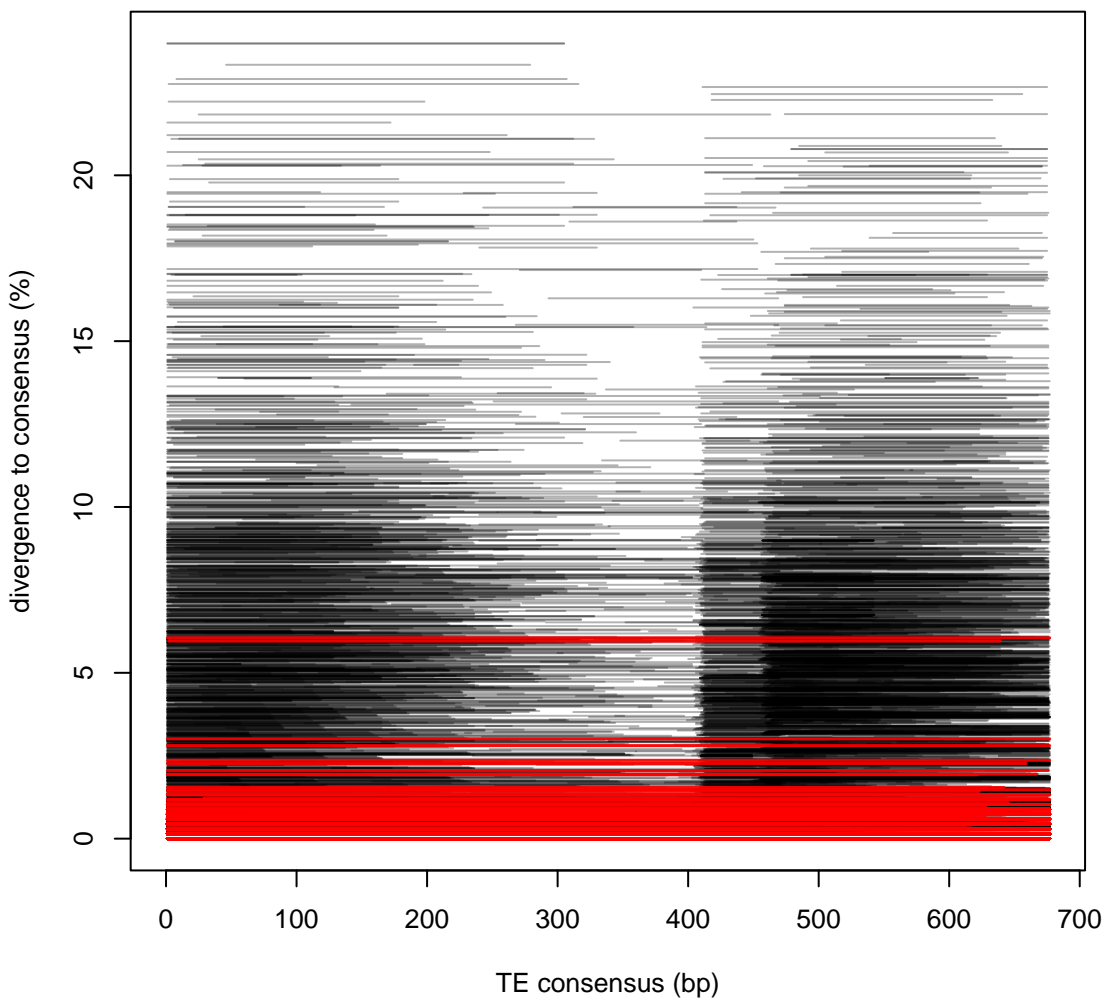
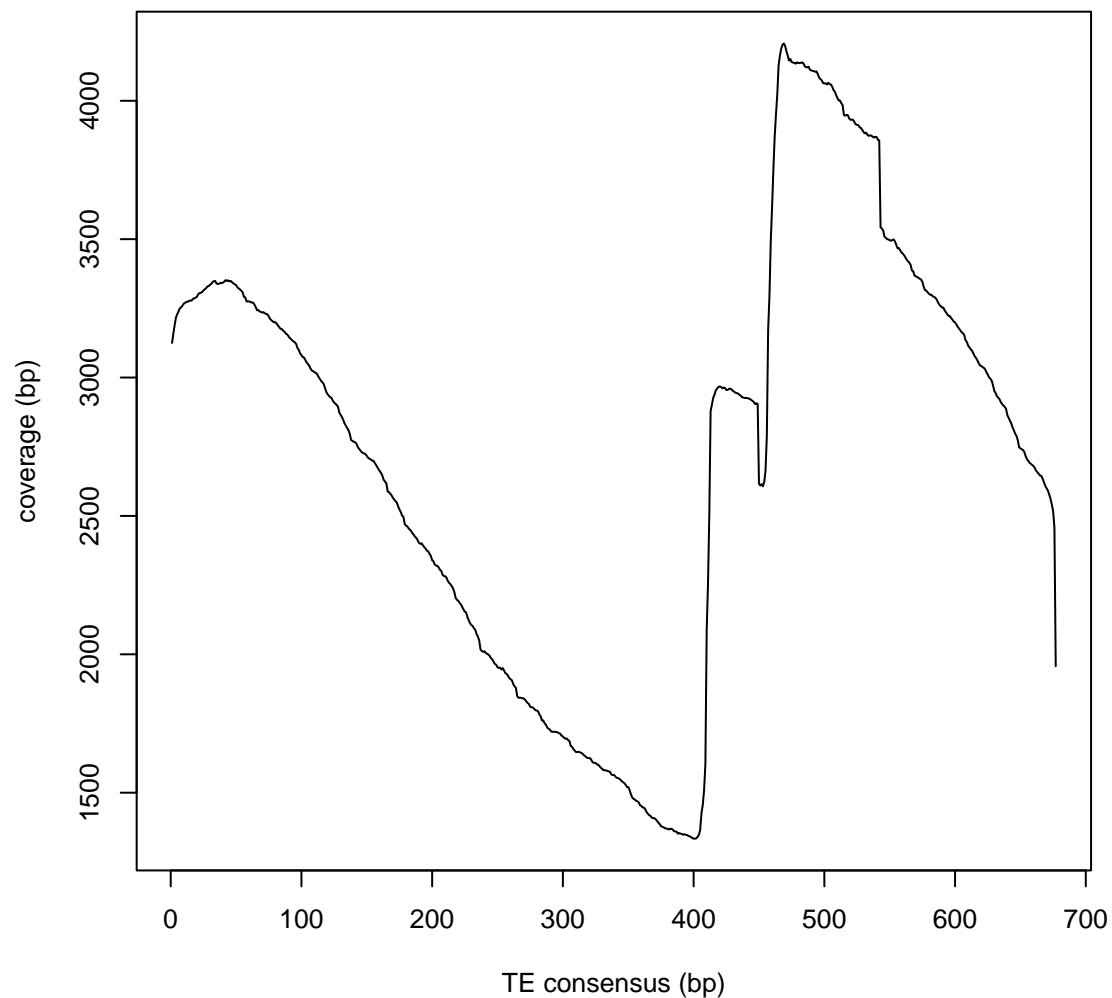


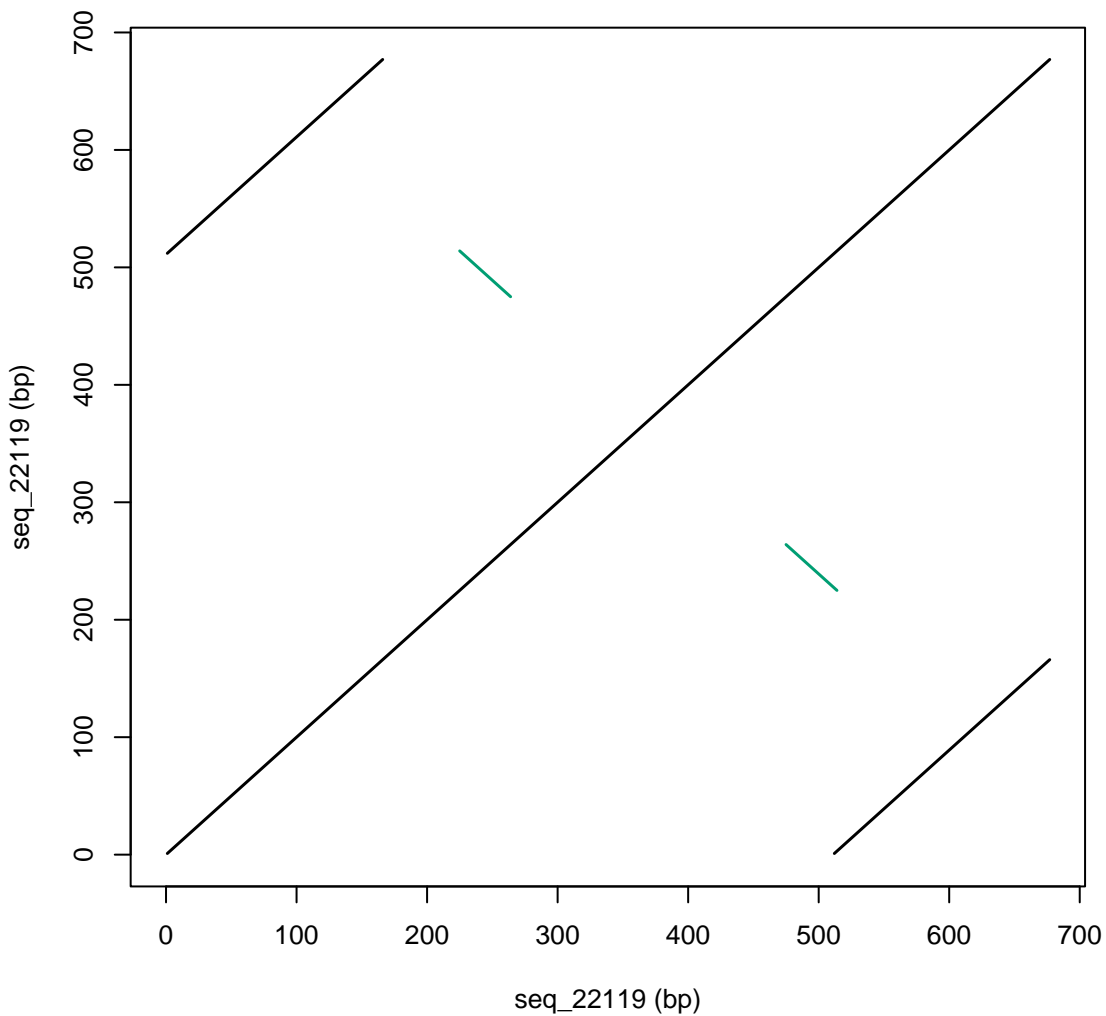
TE: seq_22119
consensus size: 677bp; fragments: 7574; full length: 741 (>=609.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

