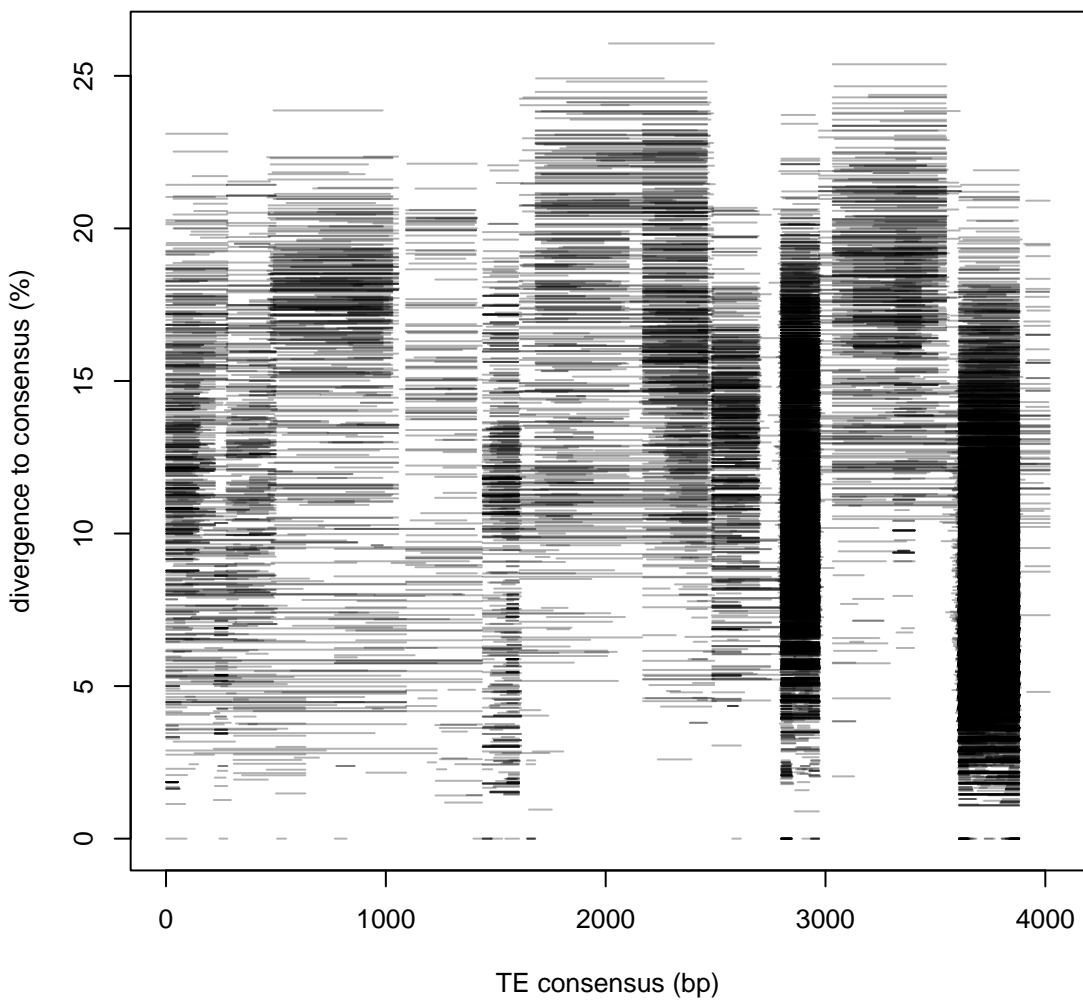
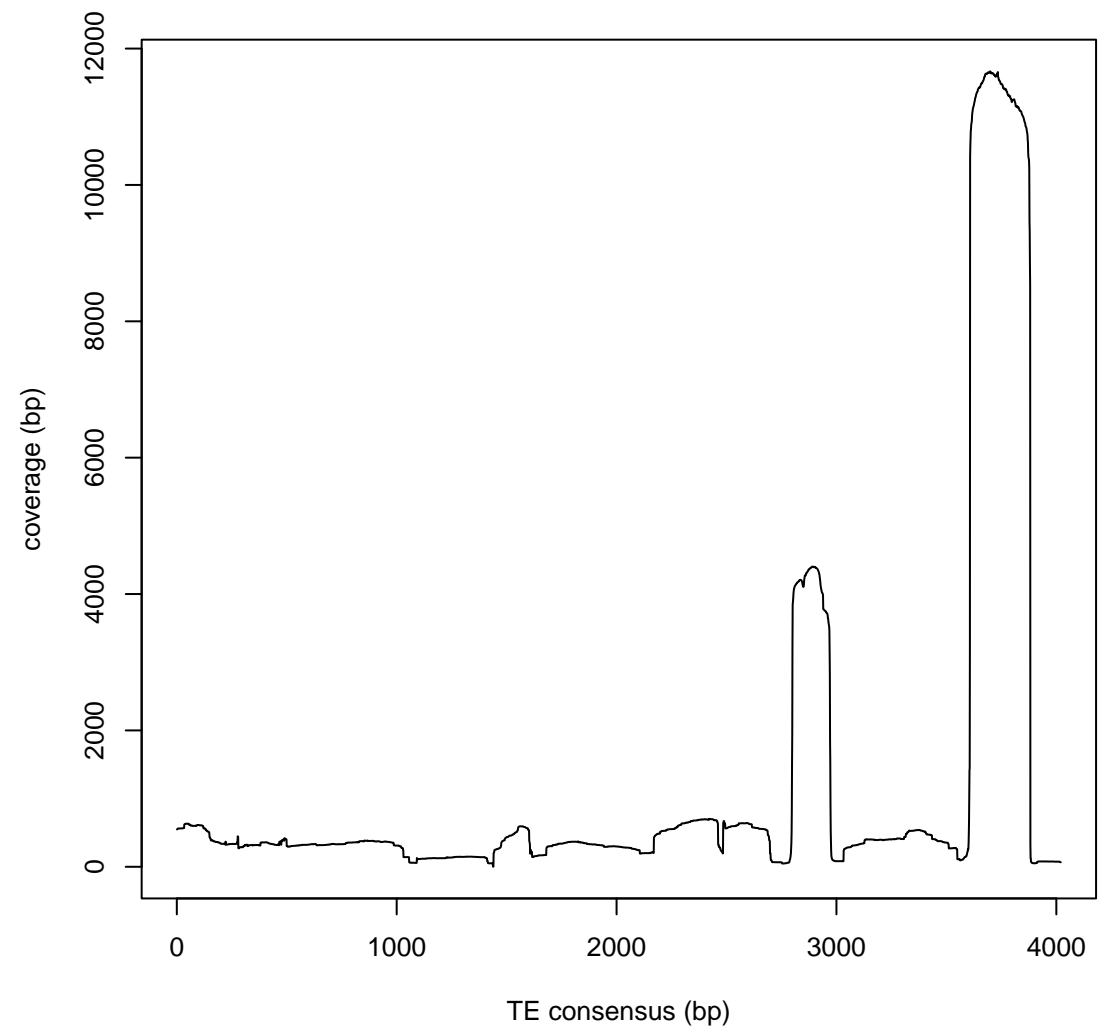


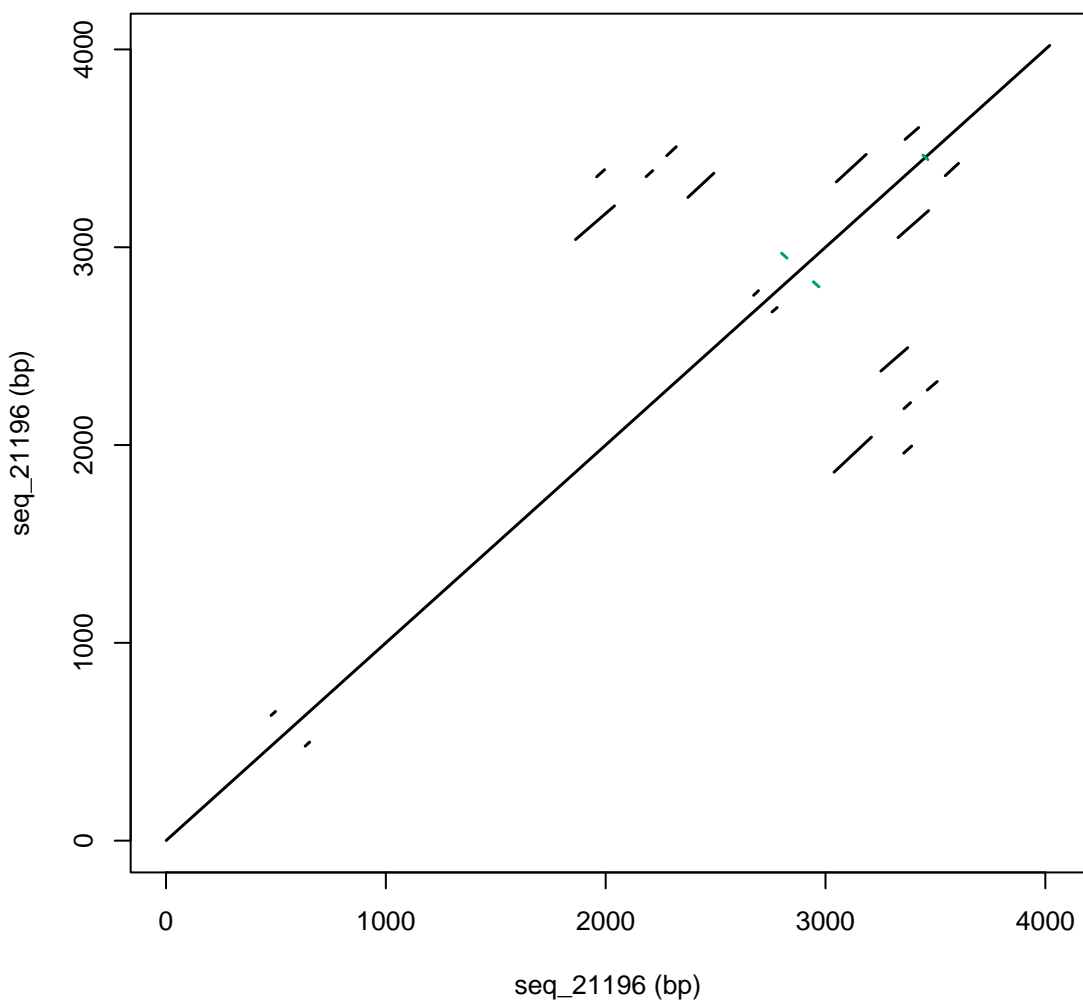
TE: seq_21196
consensus size: 4020bp; fragments: 21860; full length: 0 (>=3618bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

