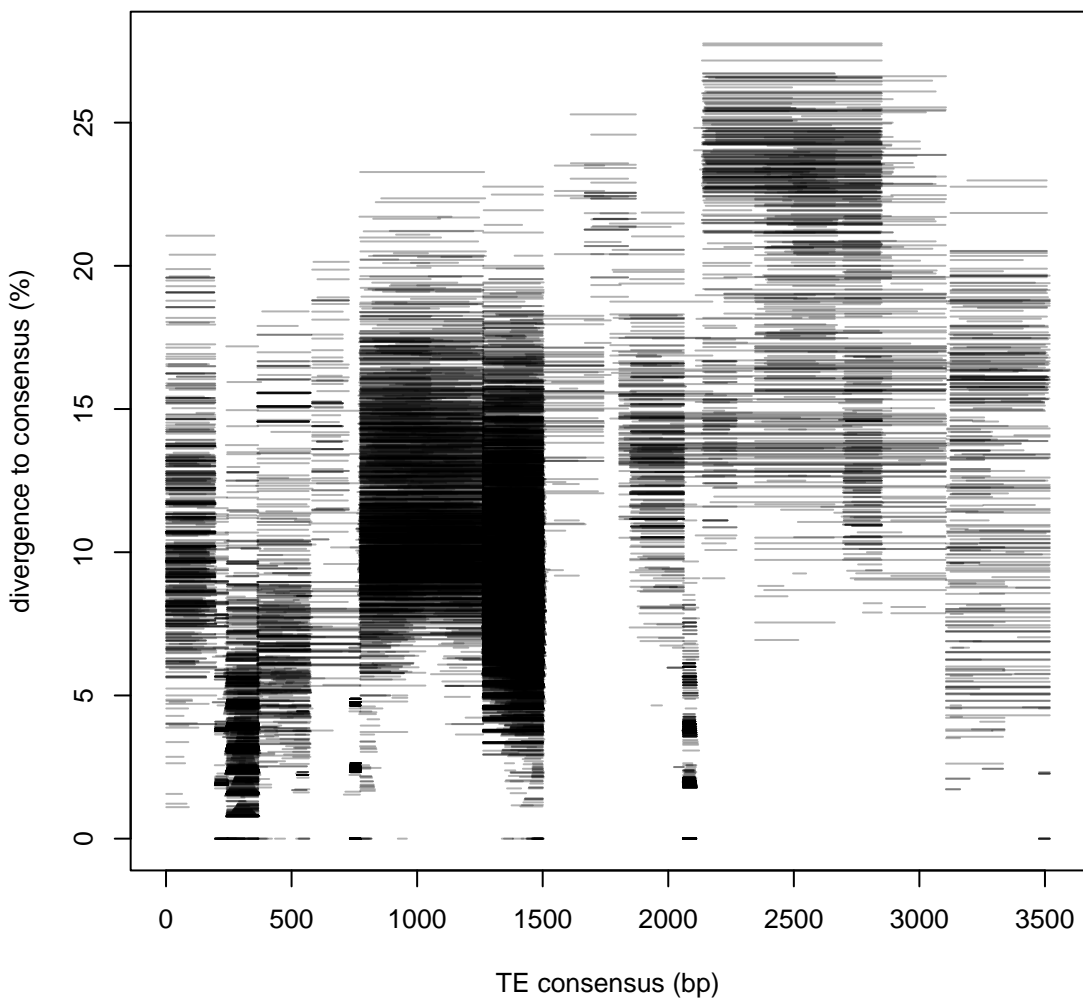
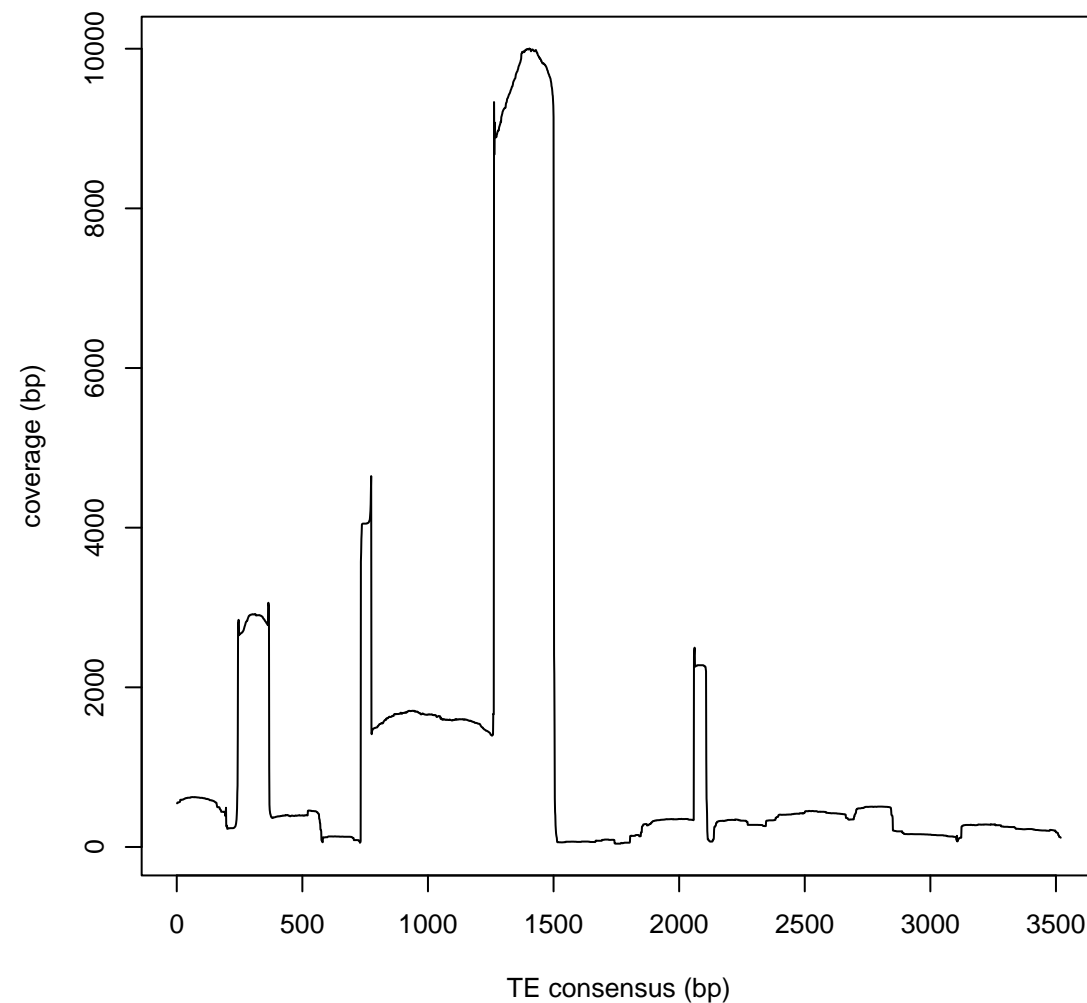


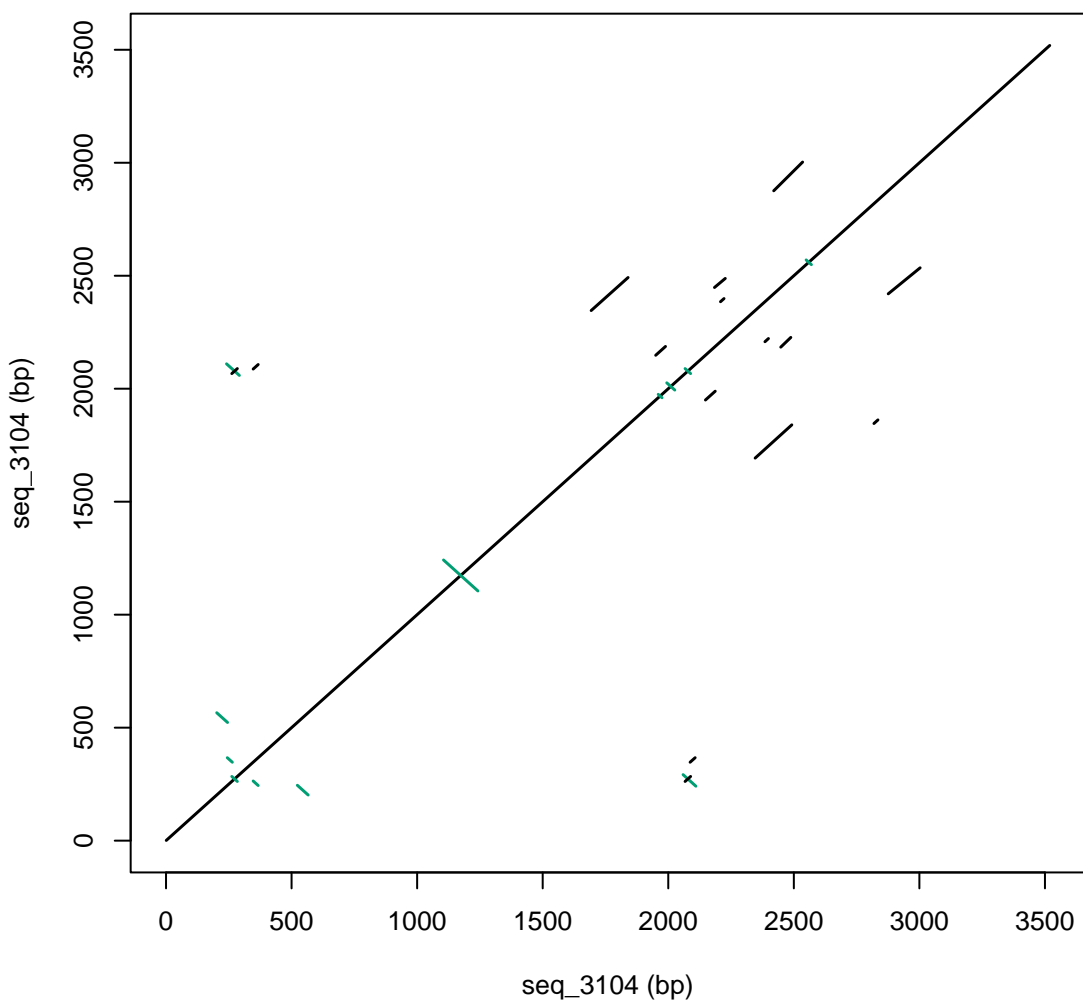
TE: seq_3104
consensus size: 3519bp; fragments: 24586; full length: 0 (>=3167.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

