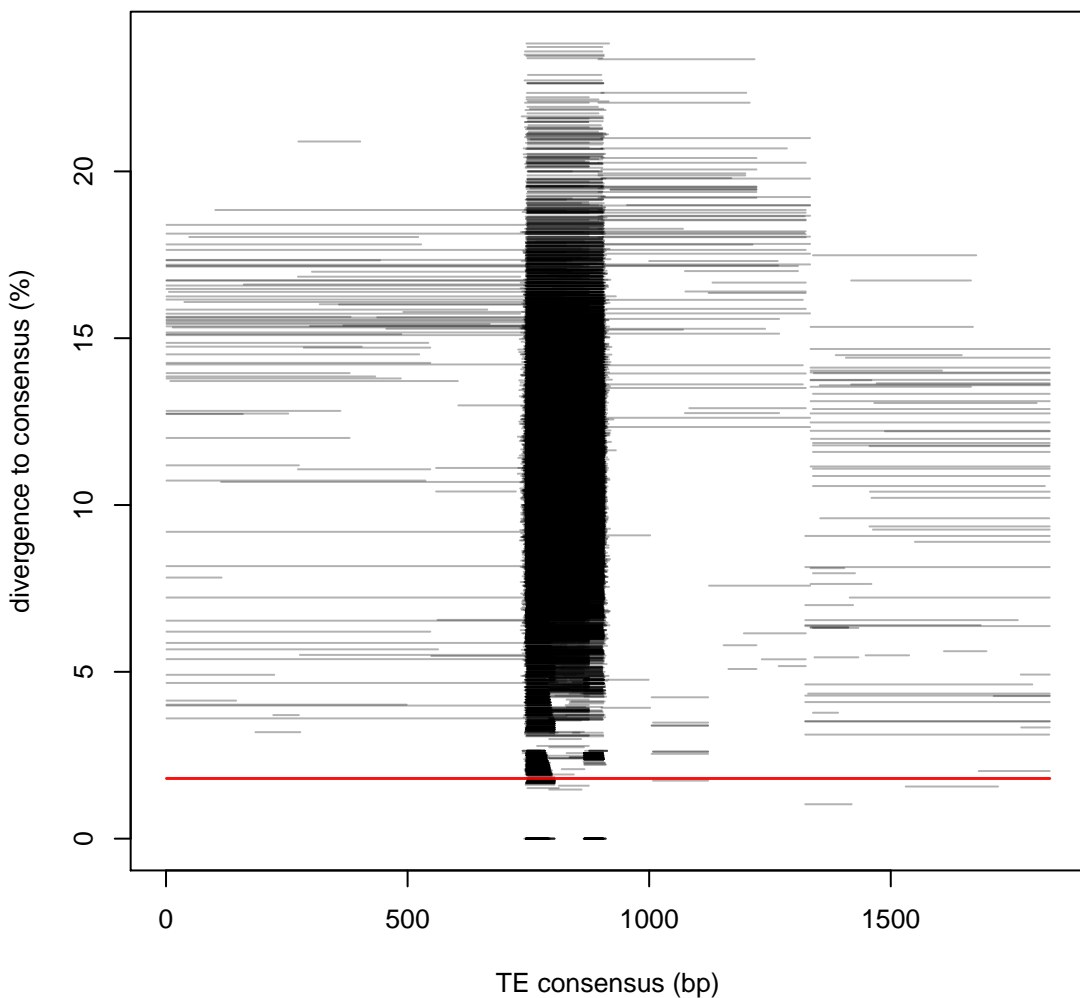
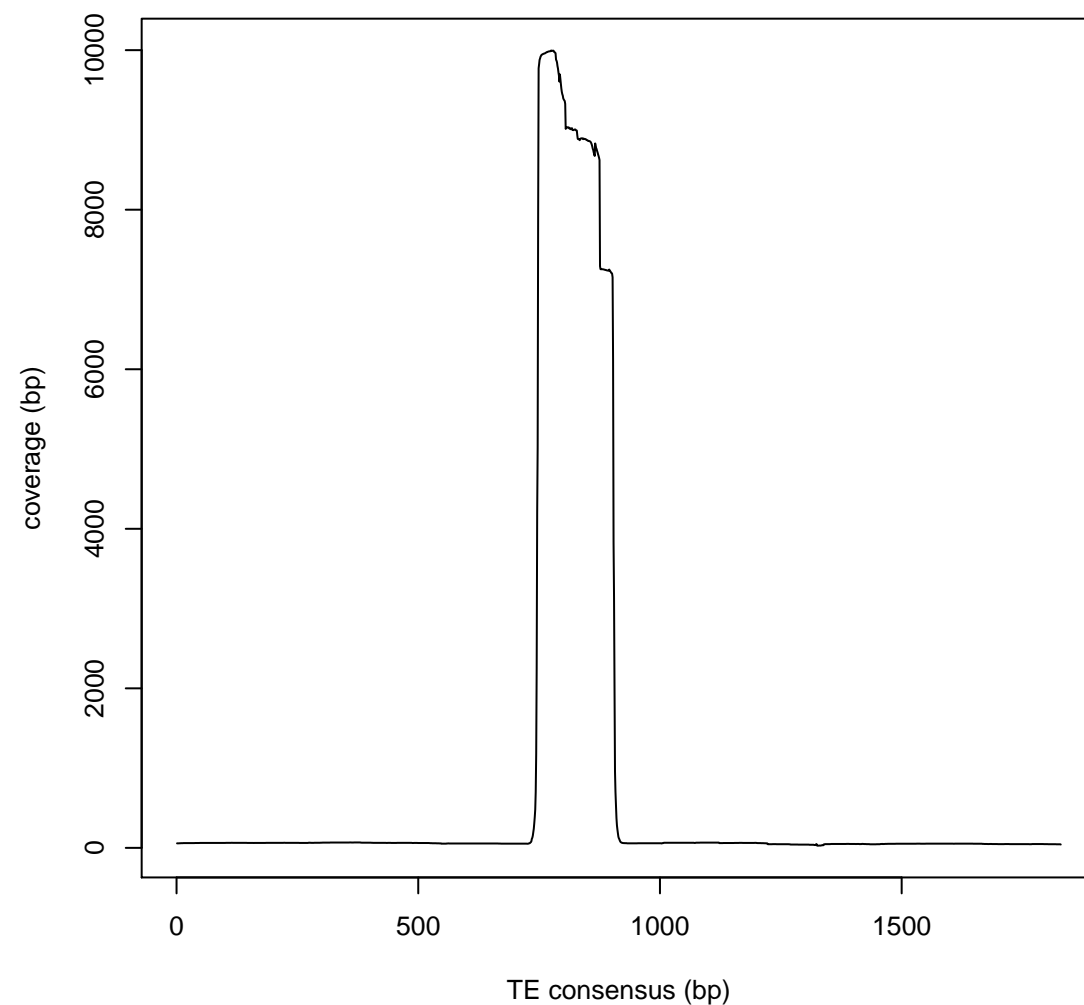


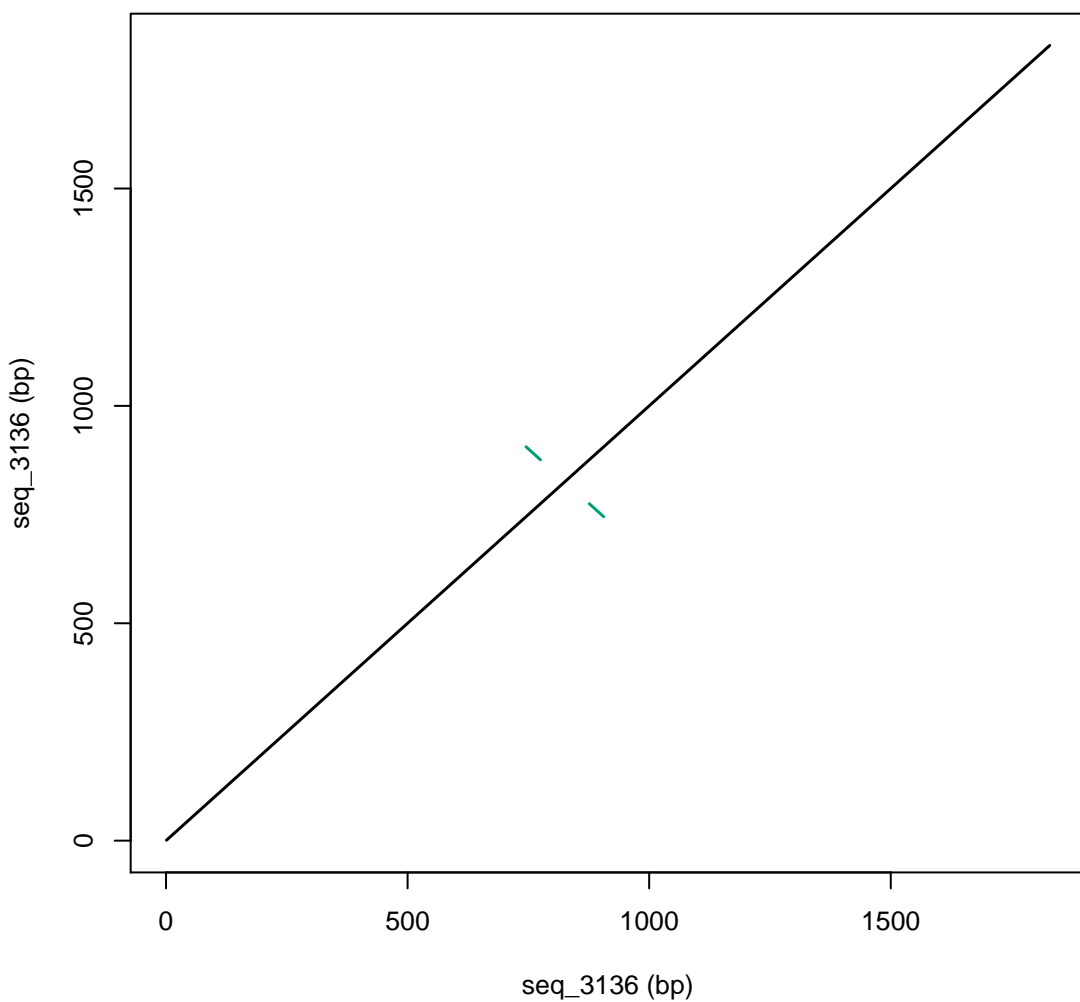
TE: seq\_3136  
consensus size: 1829bp; fragments: 10999; full length: 1 (>=1646.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

