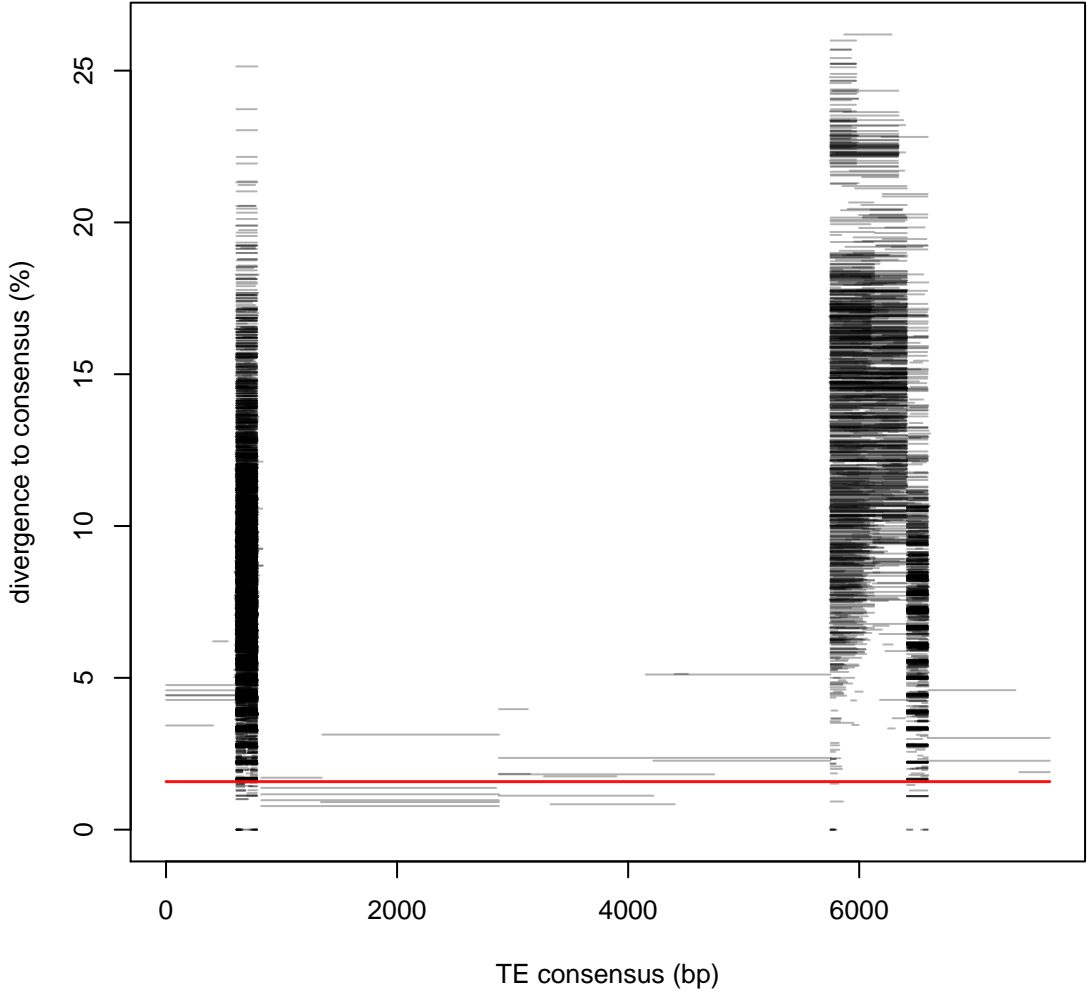
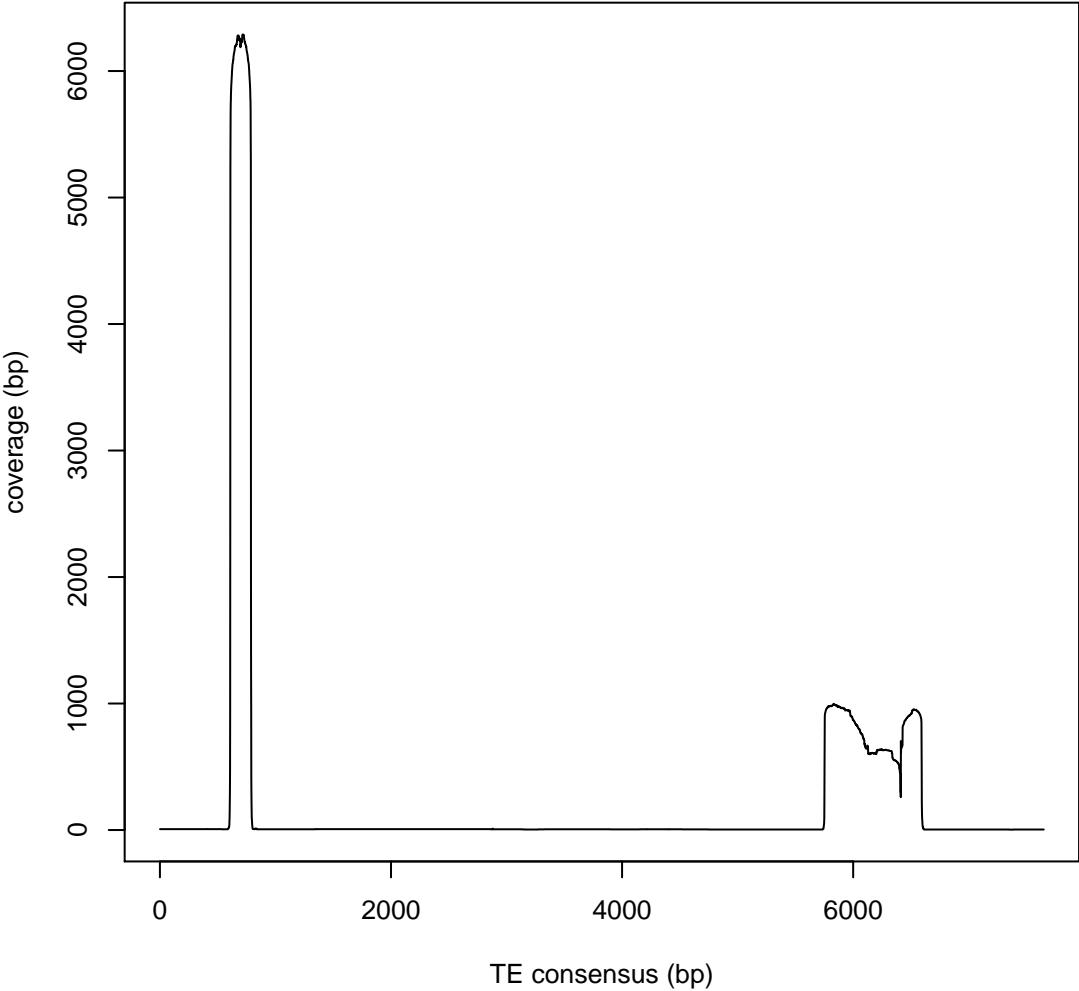


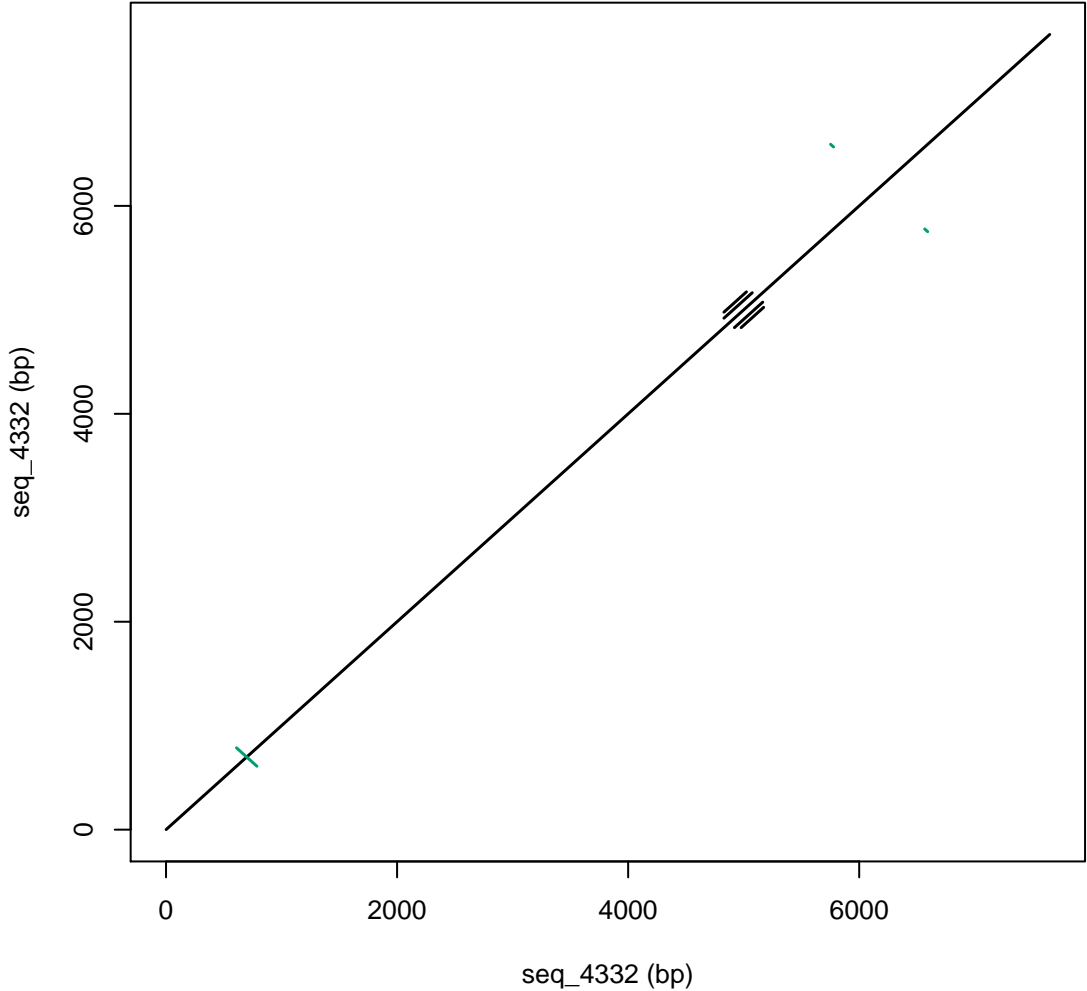
TE: seq\_4332  
consensus size: 7649bp; fragments: 9208; full length: 1 (>=6884.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

