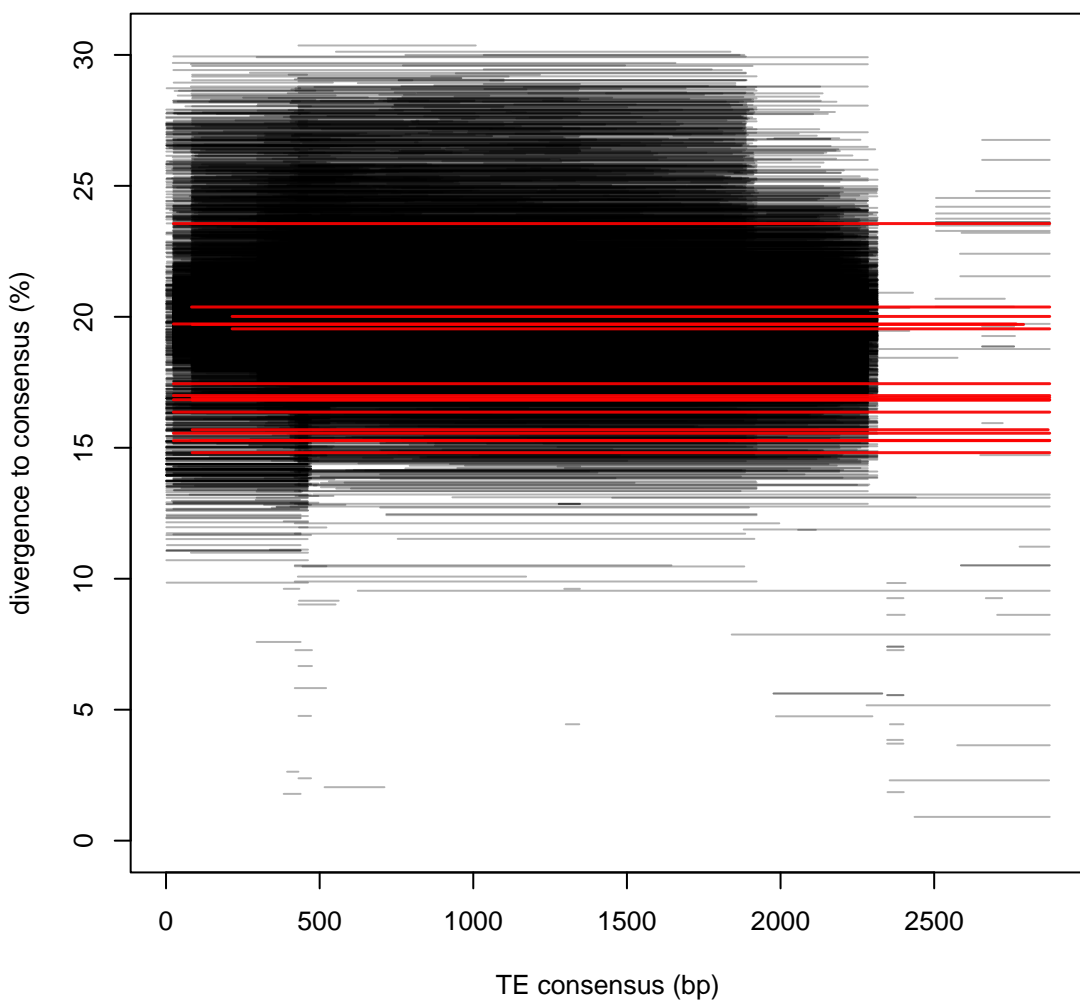
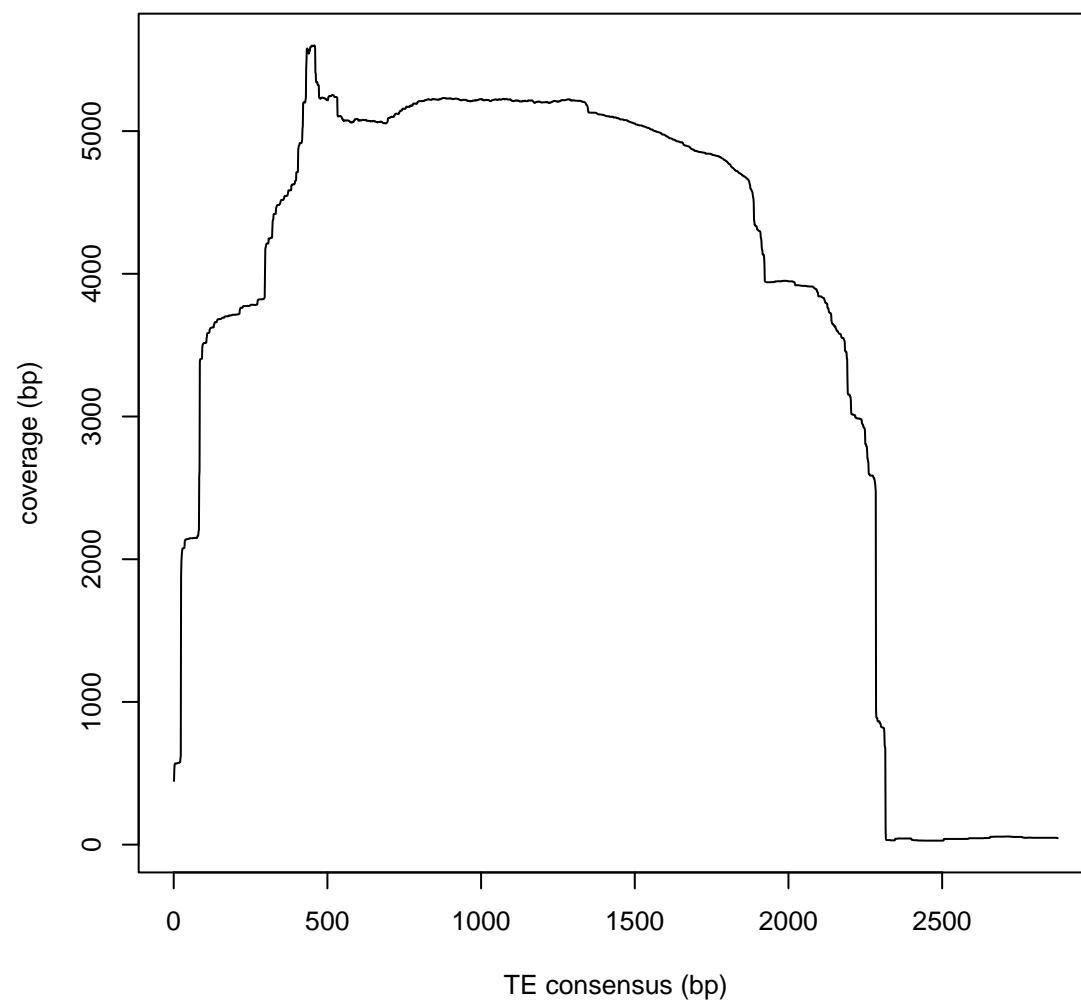


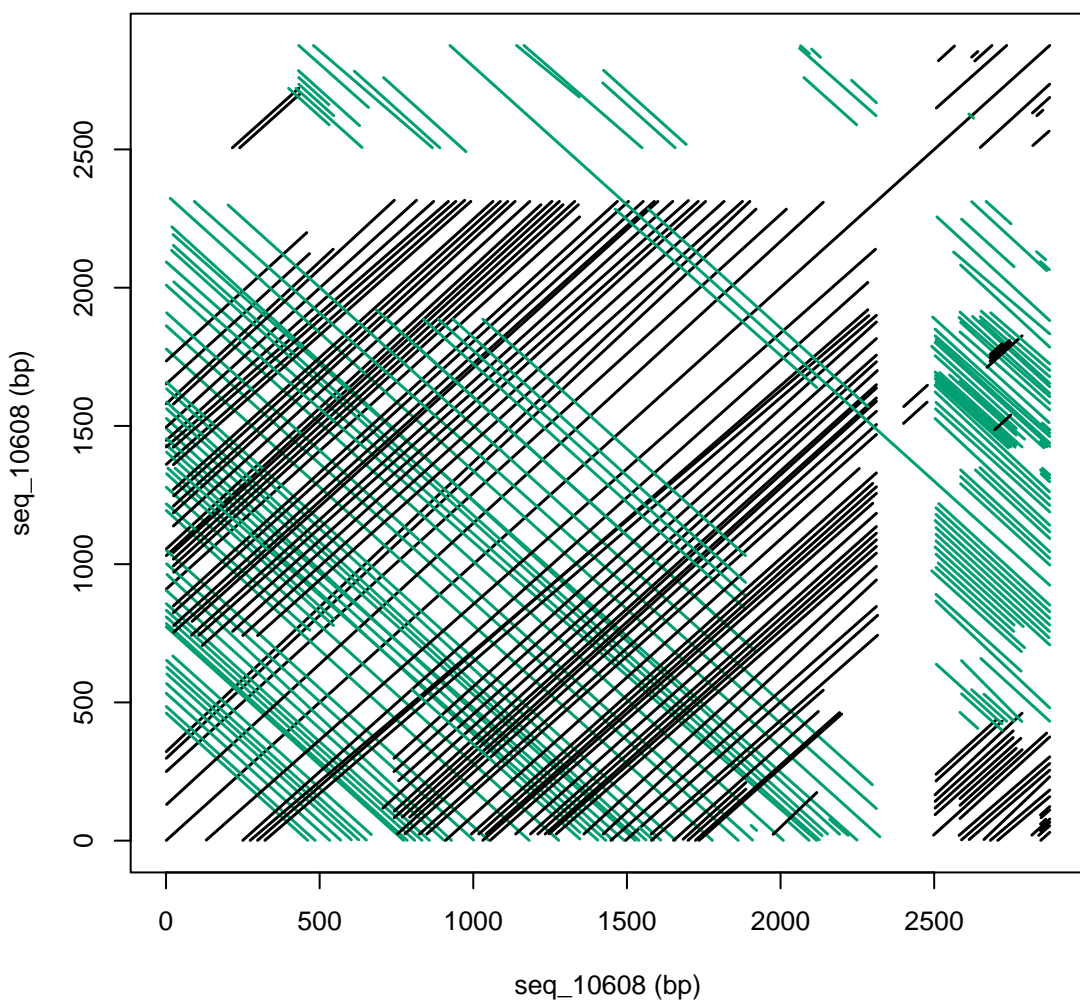
TE: seq\_10608  
consensus size: 2876bp; fragments: 6356; full length: 17 (>=2588.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

