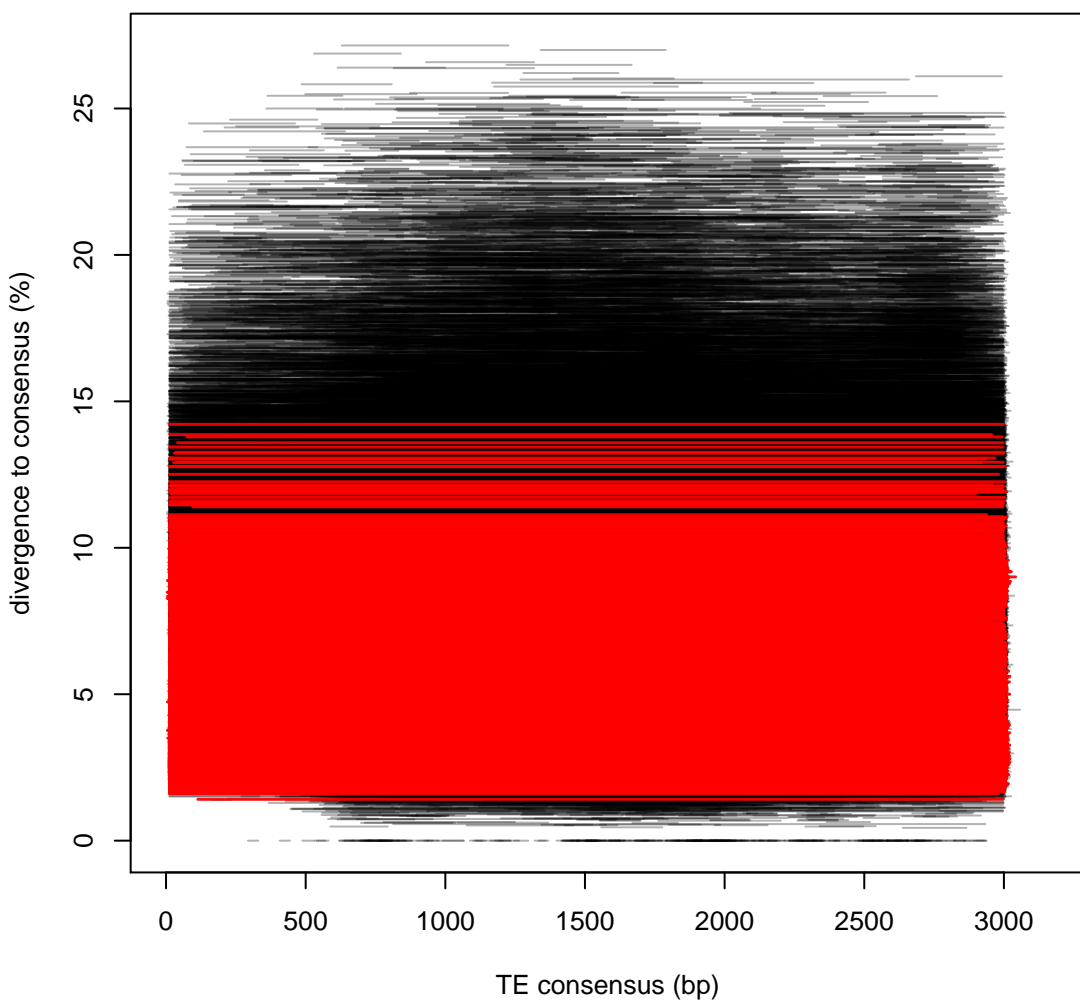
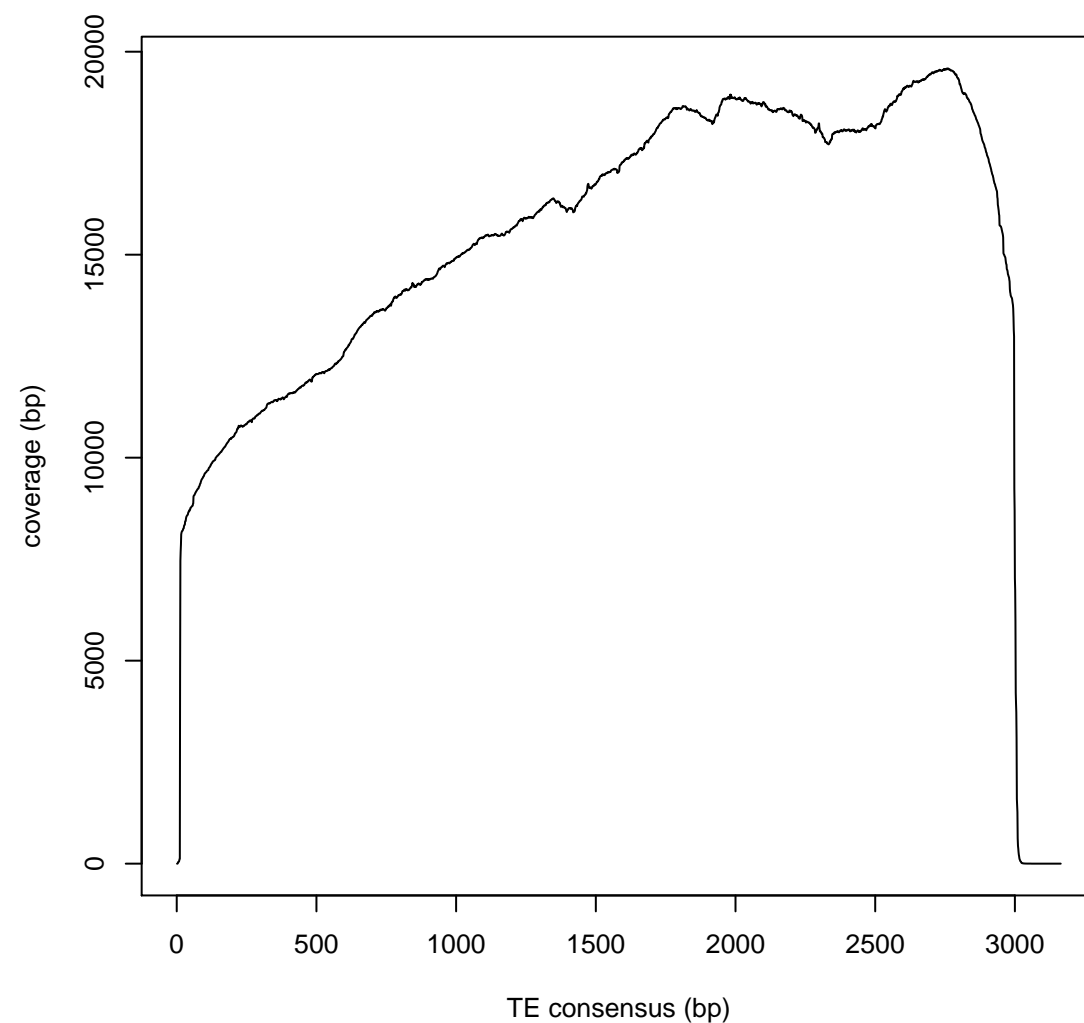


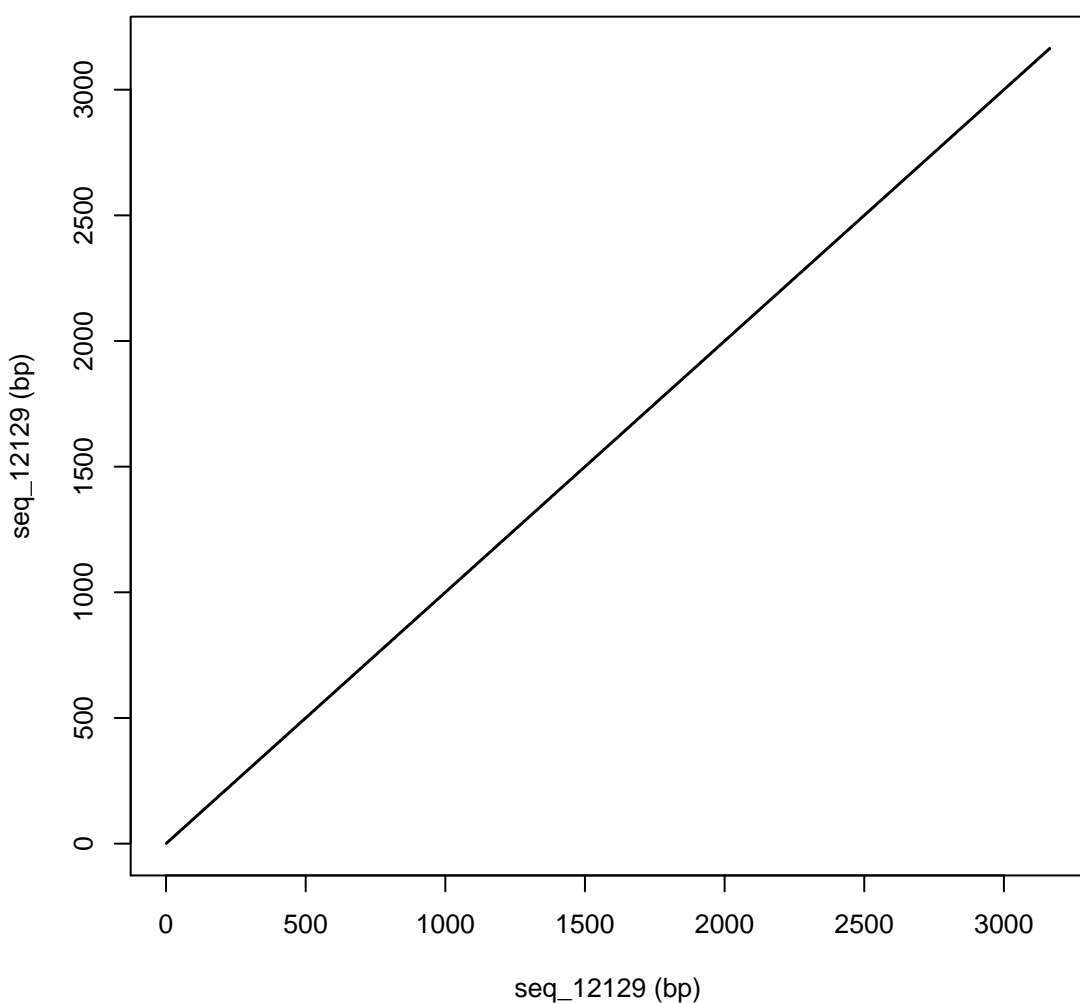
TE: seq_12129
consensus size: 3164bp; fragments: 56112; full length: 2777 (>=2847.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

