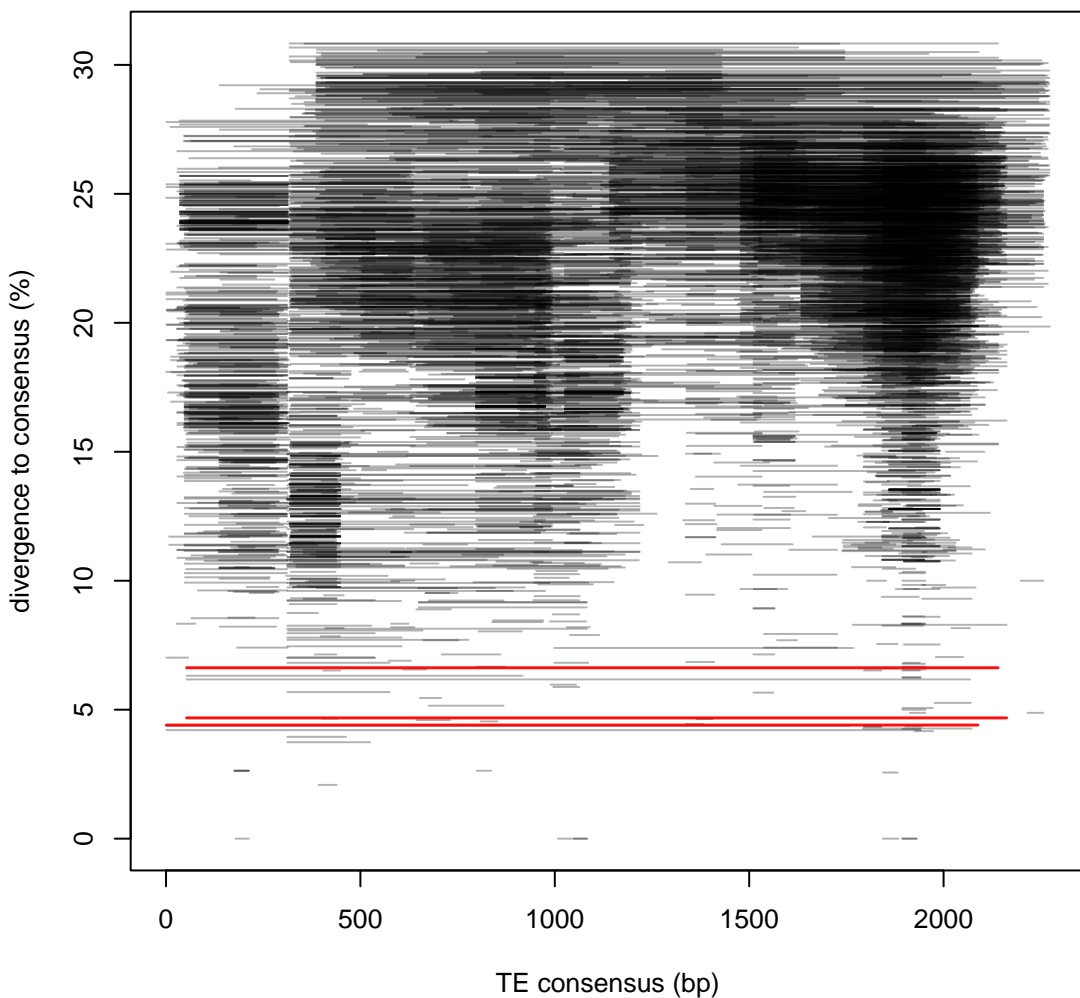
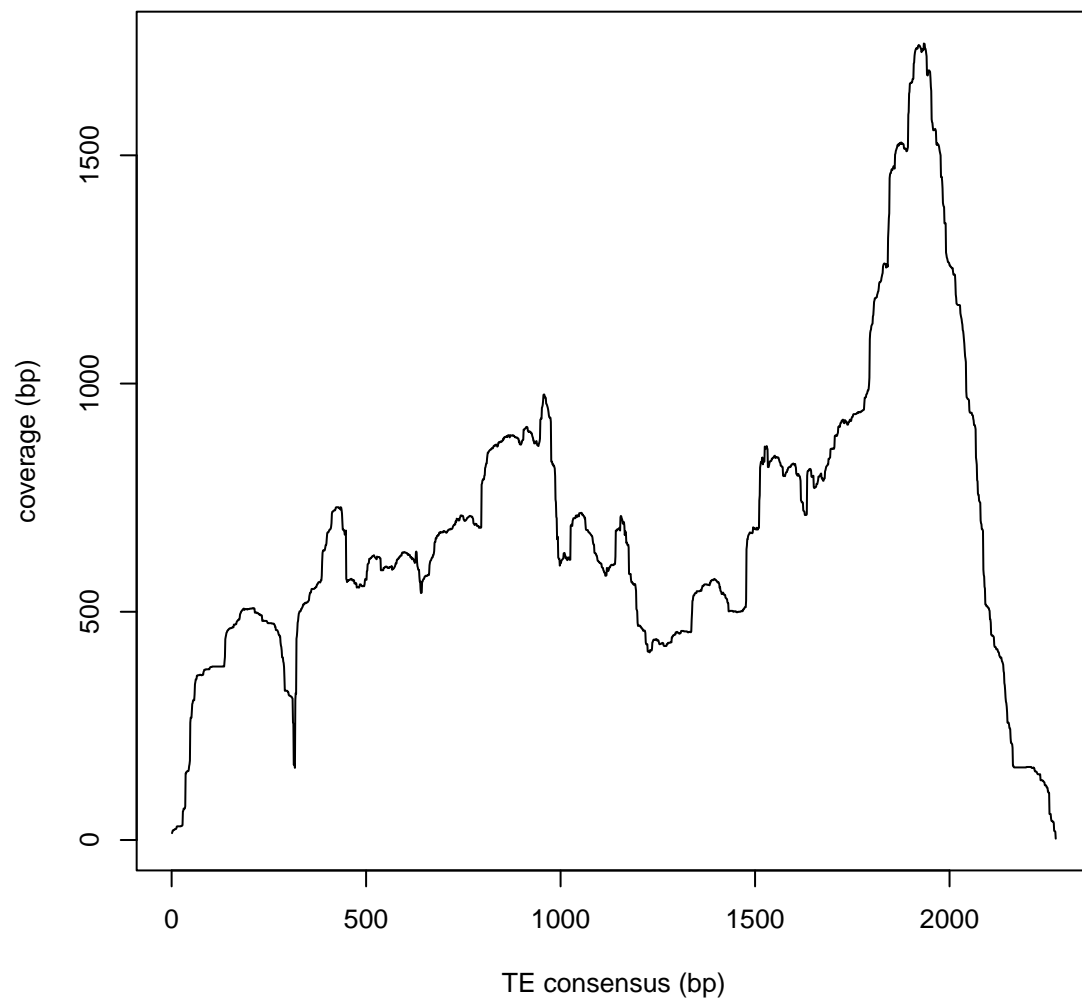


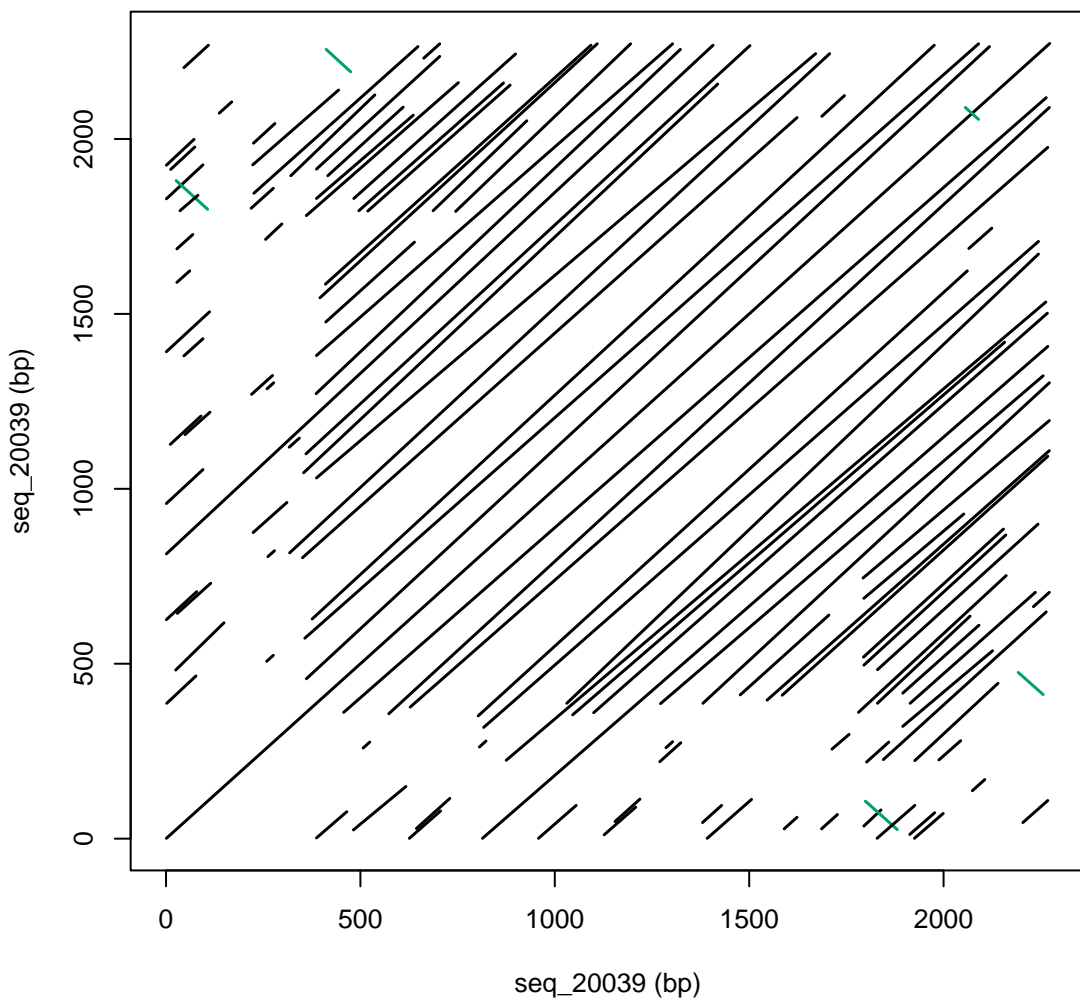
TE: seq_20039
consensus size: 2273bp; fragments: 4370; full length: 3 (≥ 2045.7 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

