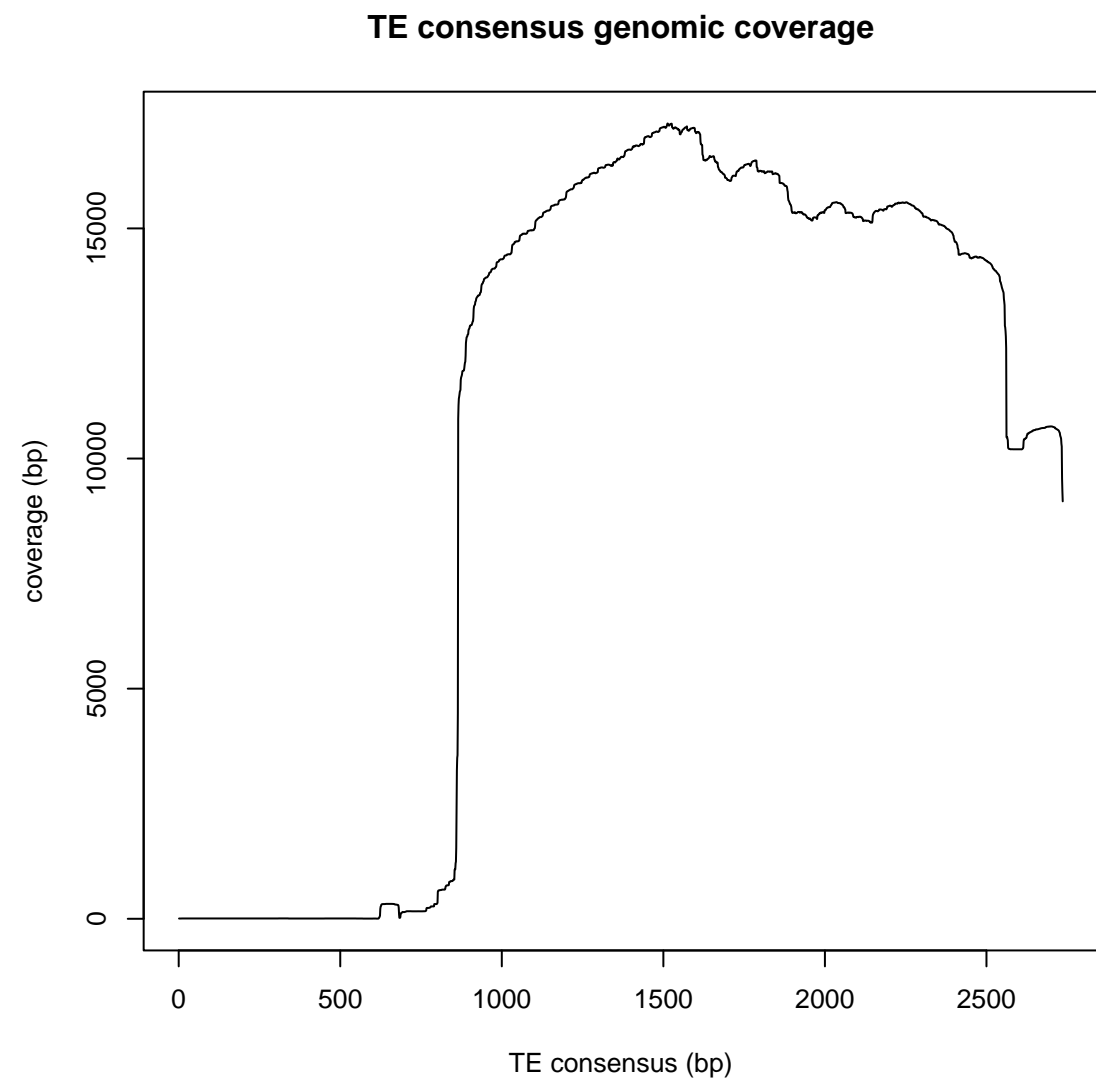
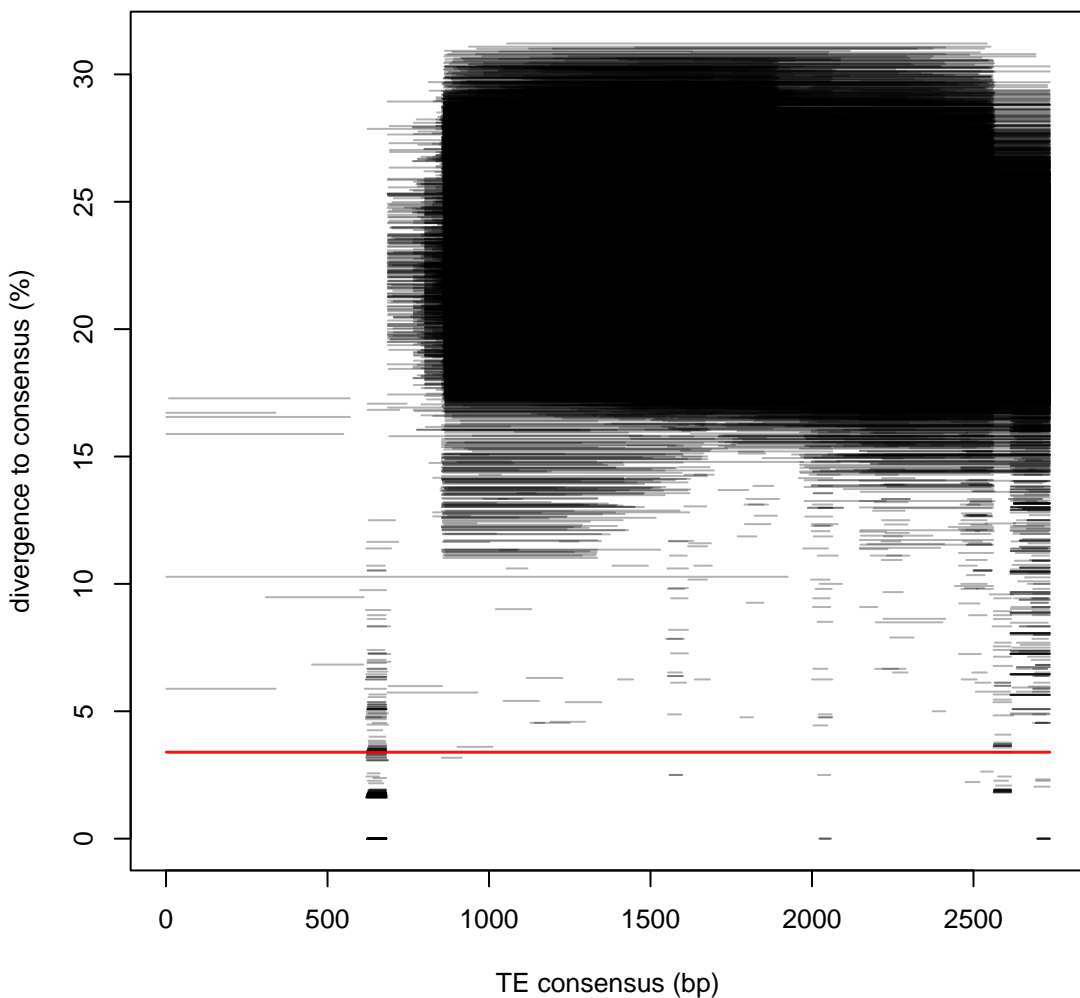
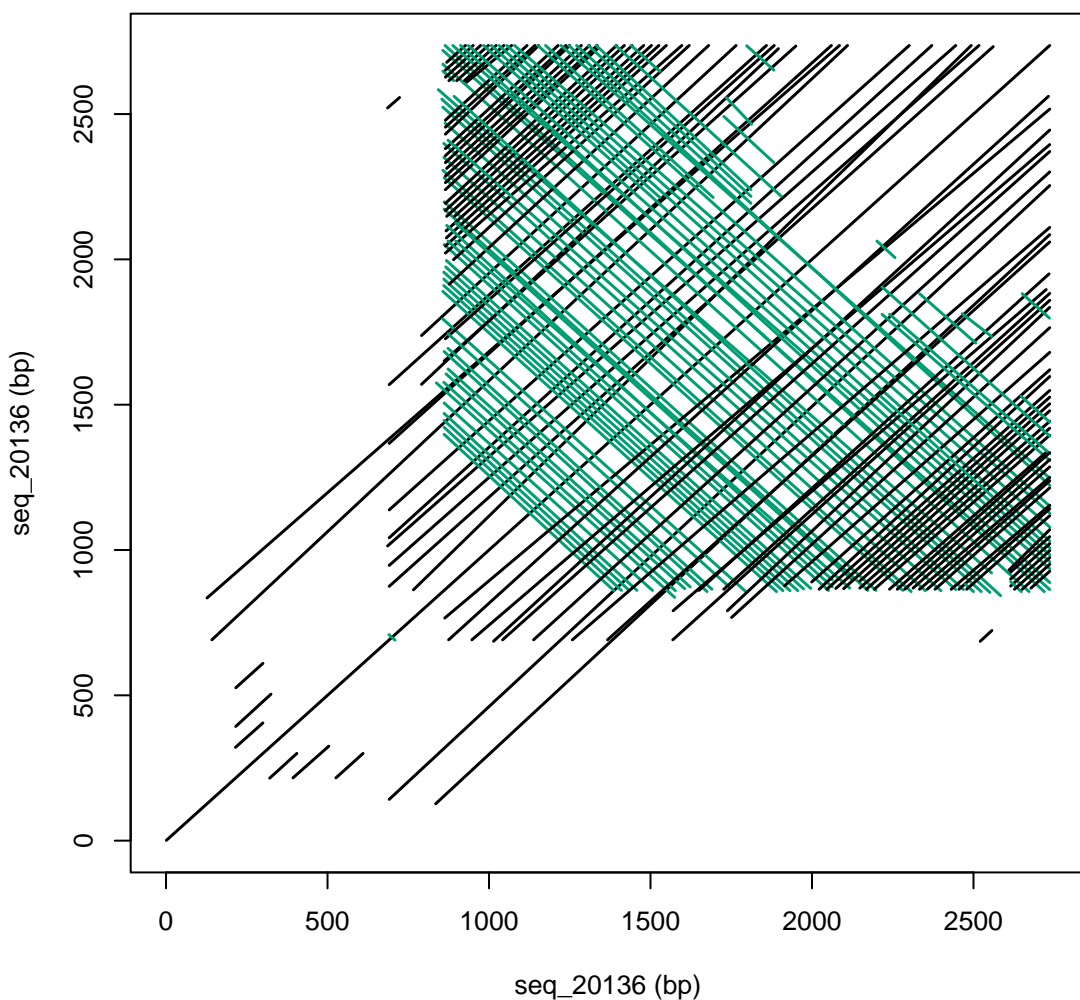


TE: seq_20136
consensus size: 2736bp; fragments: 22060; full length: 1 (>=2462.4bp)



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

