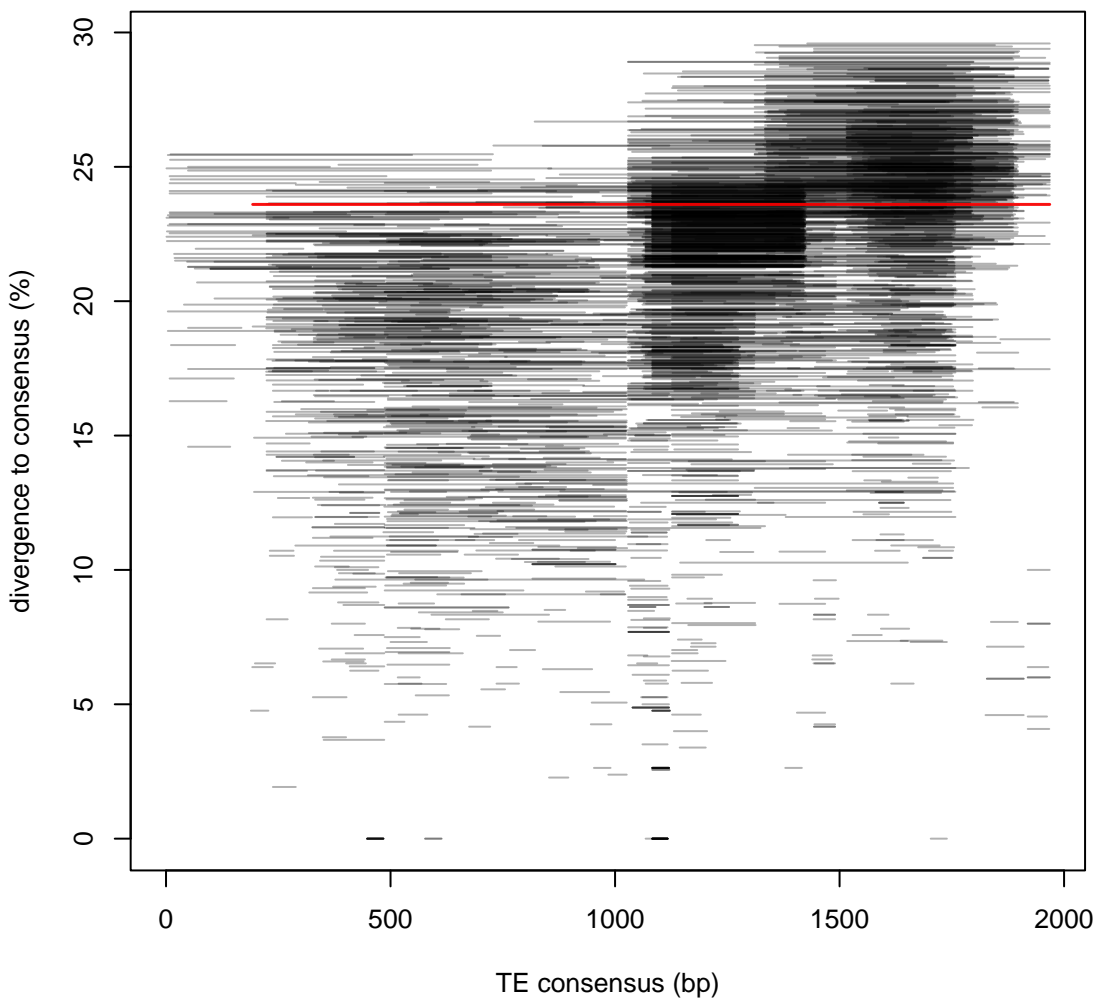
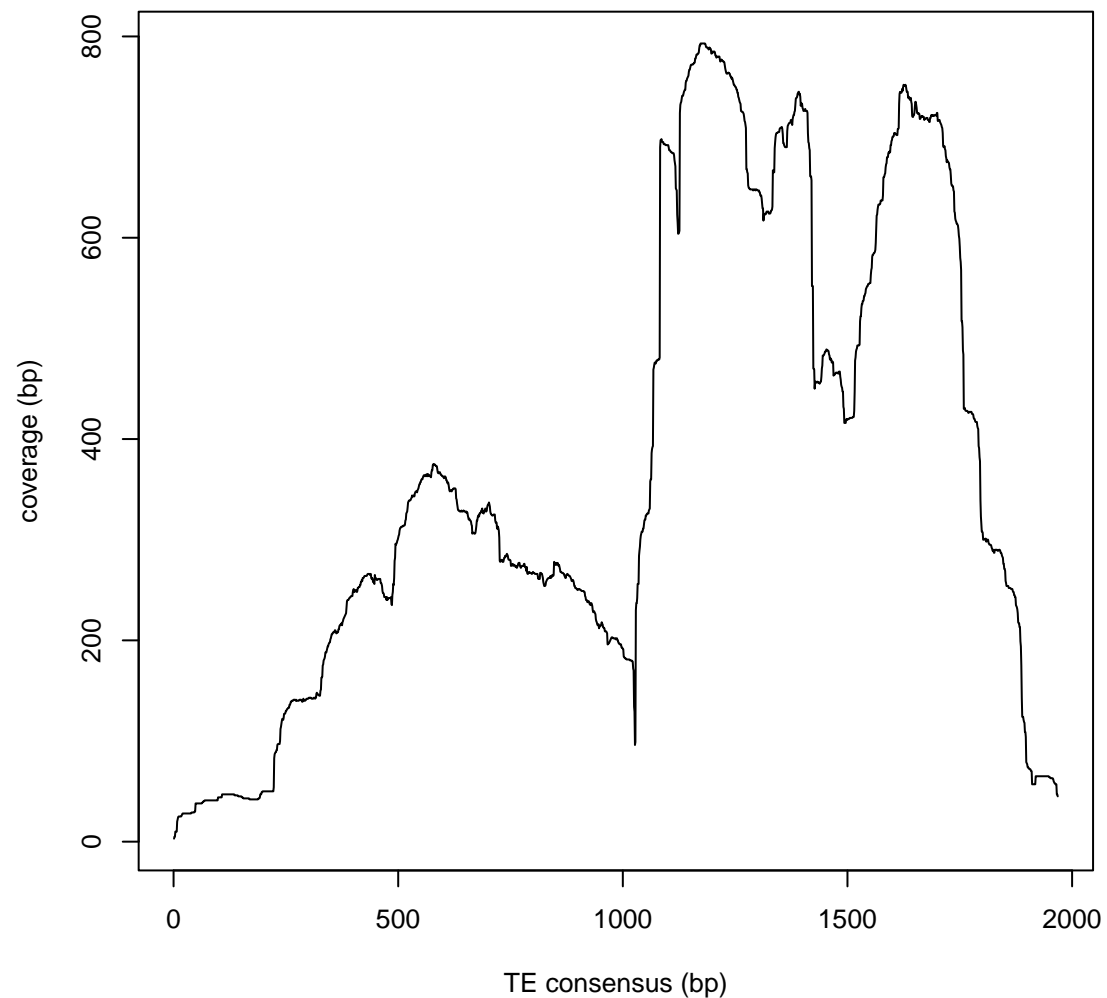


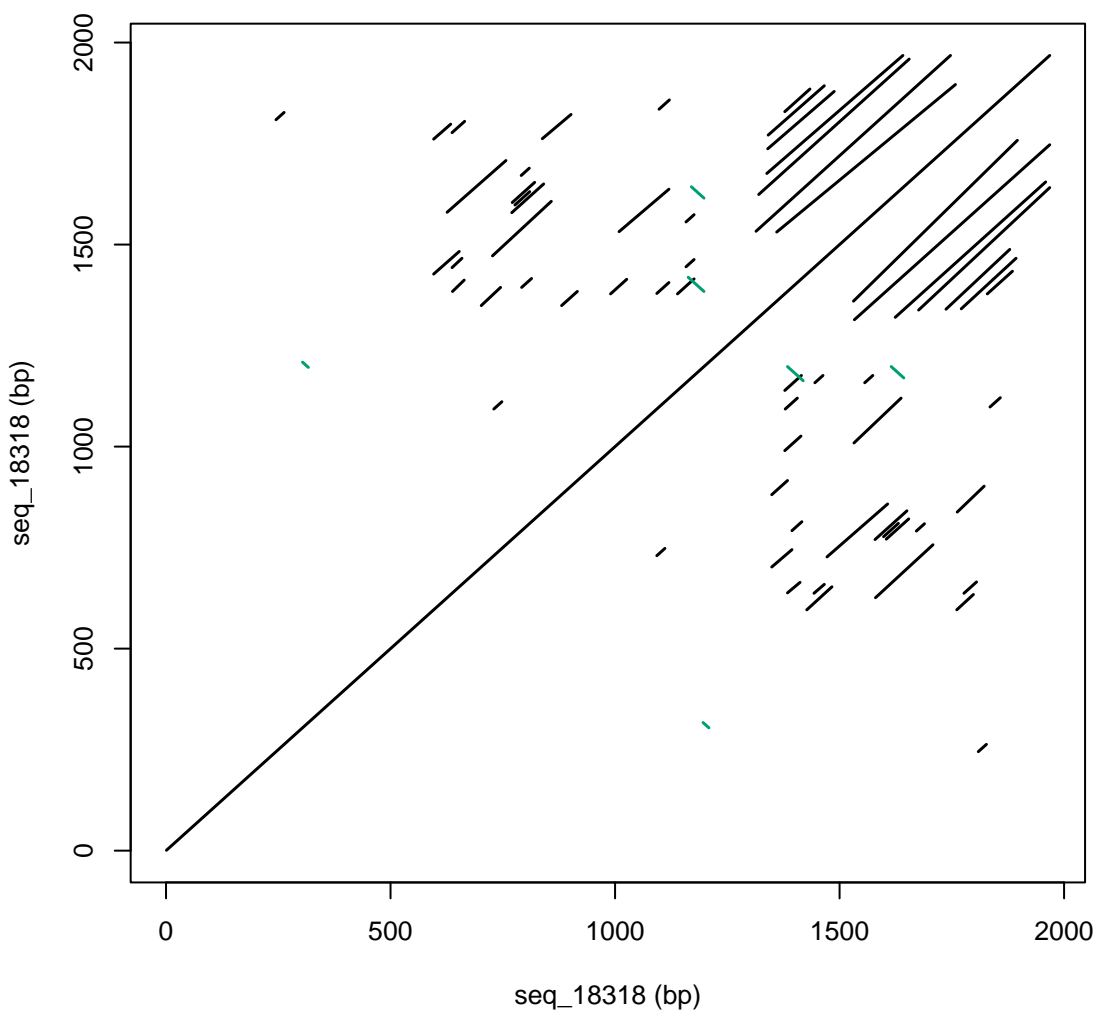
TE: seq_18318
consensus size: 1968bp; fragments: 2254; full length: 1 (>=1771.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

