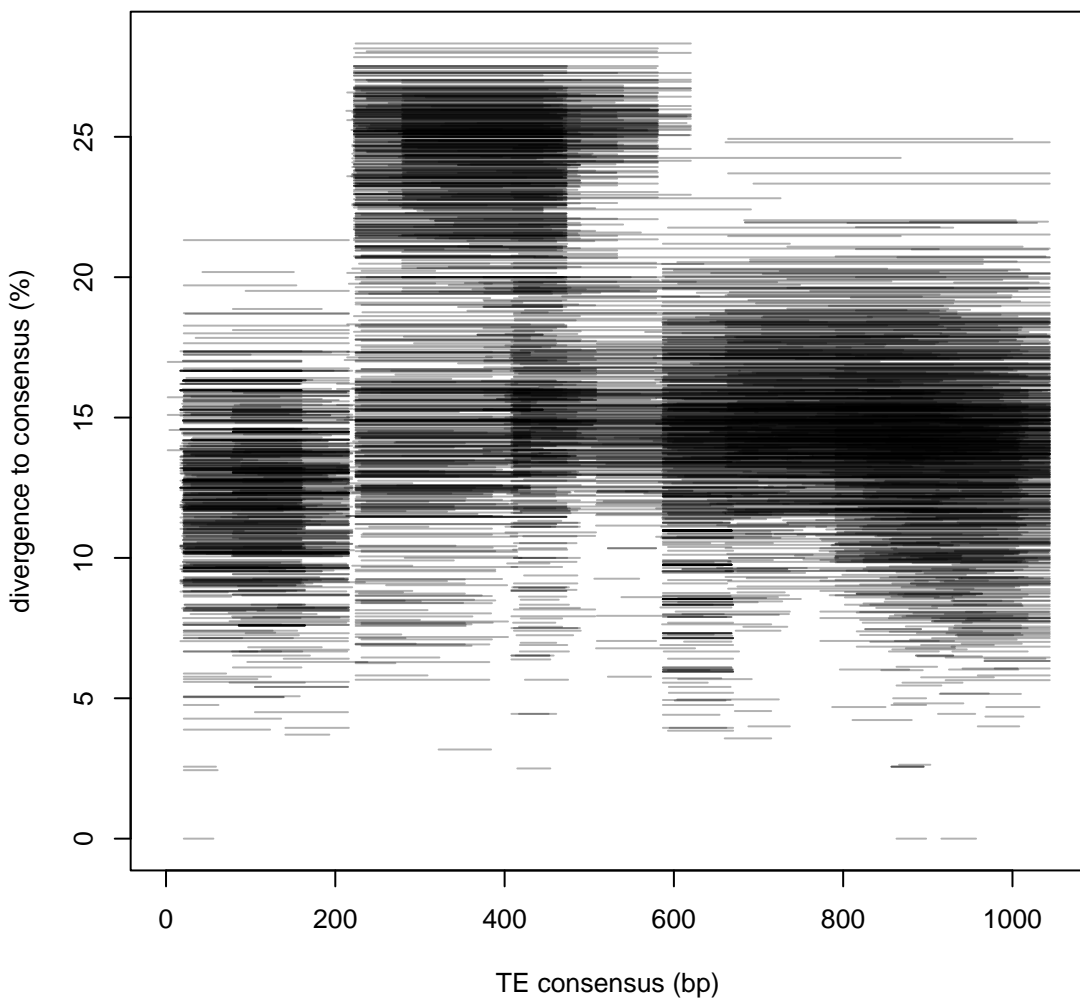
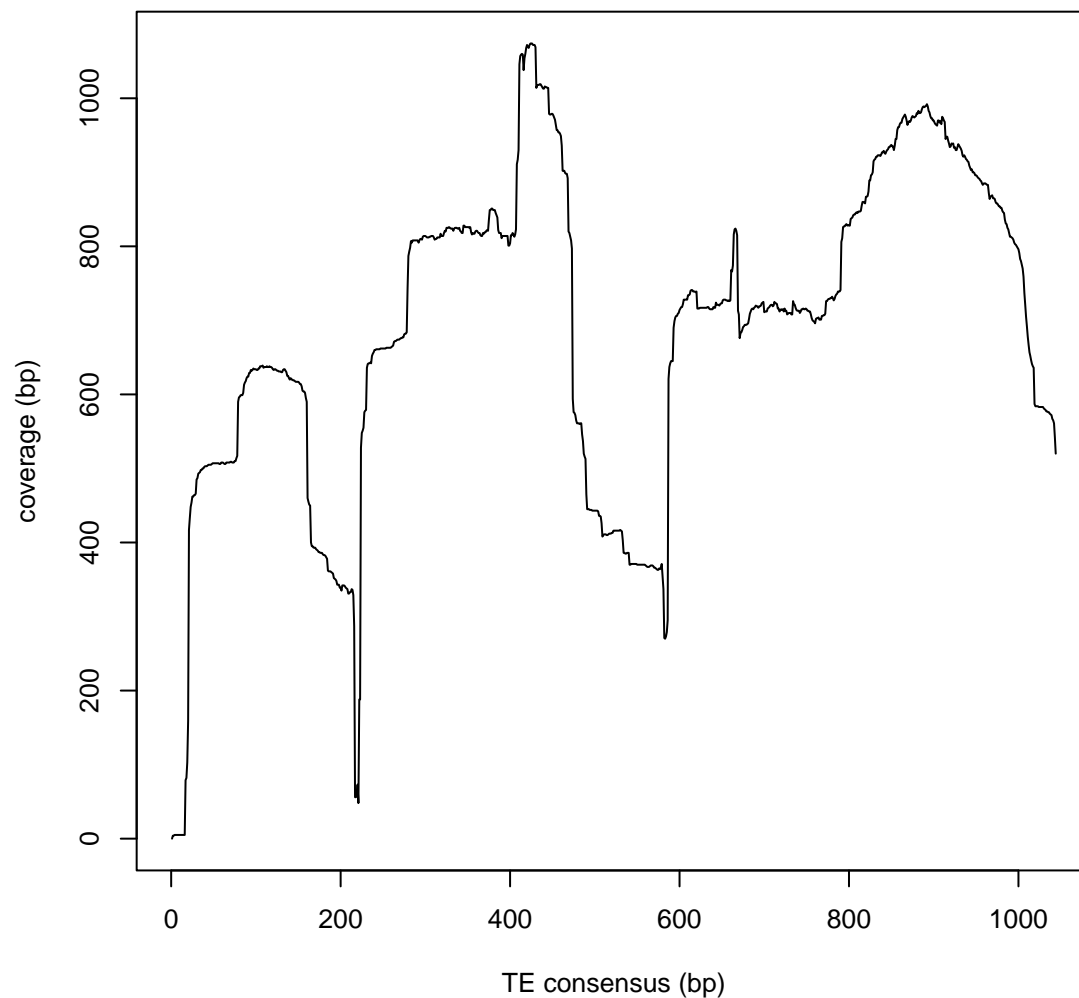


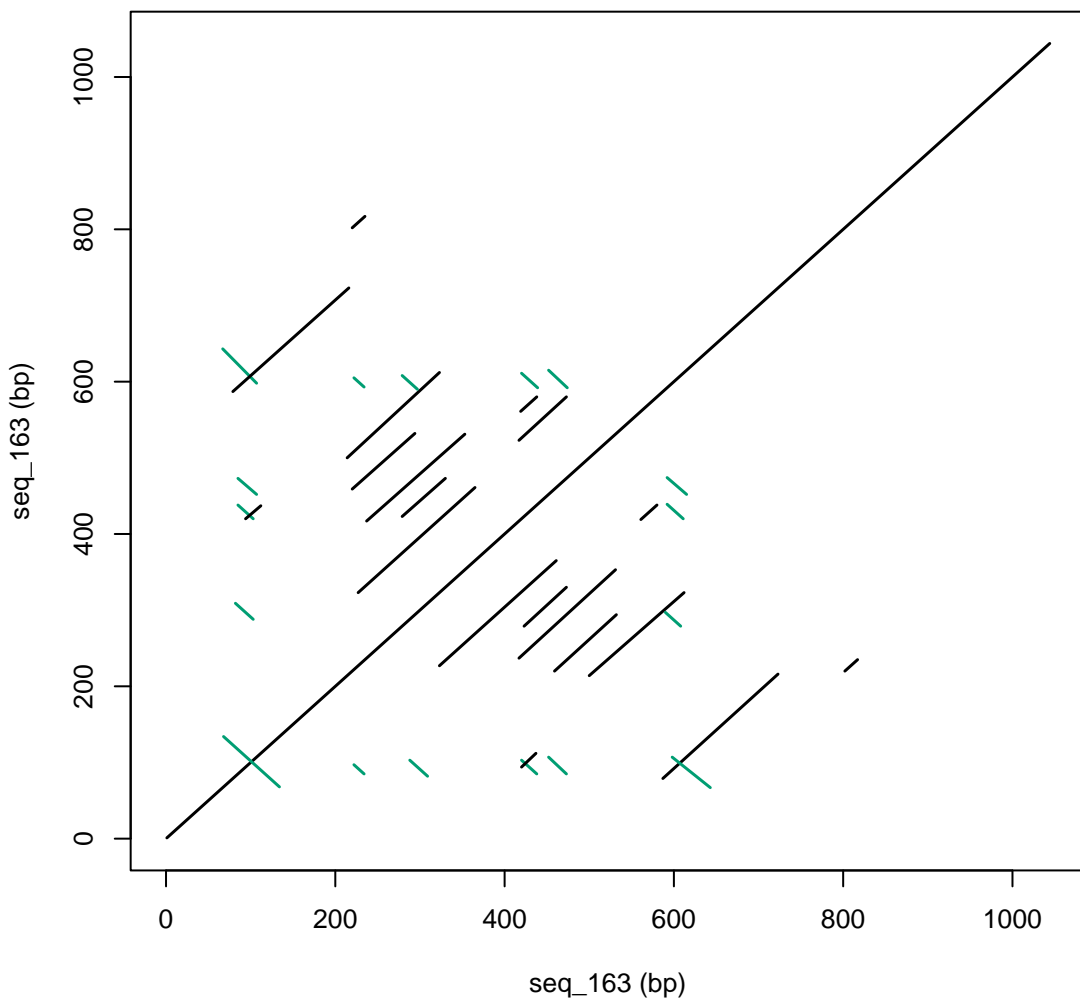
TE: seq_163
consensus size: 1044bp; fragments: 3179; full length: 0 (>=939.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

