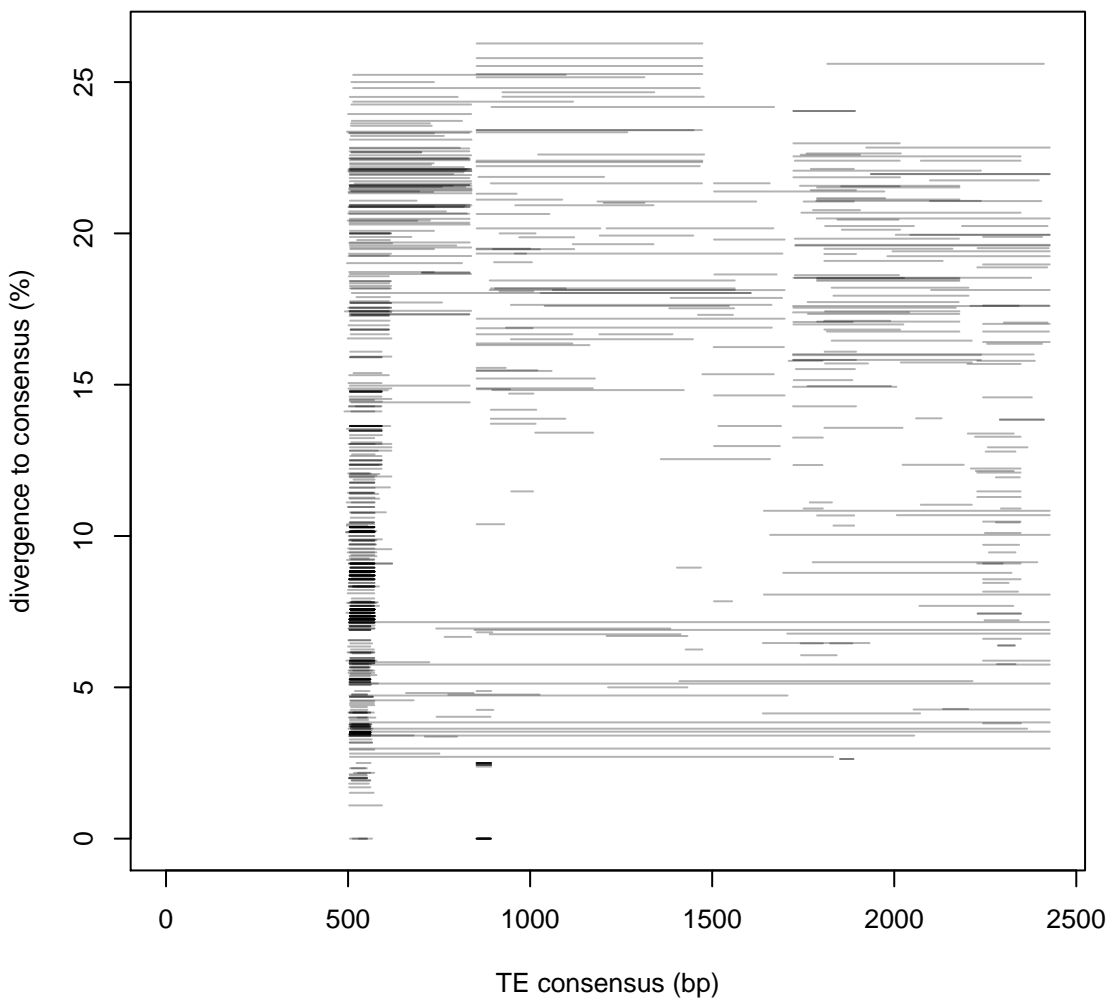
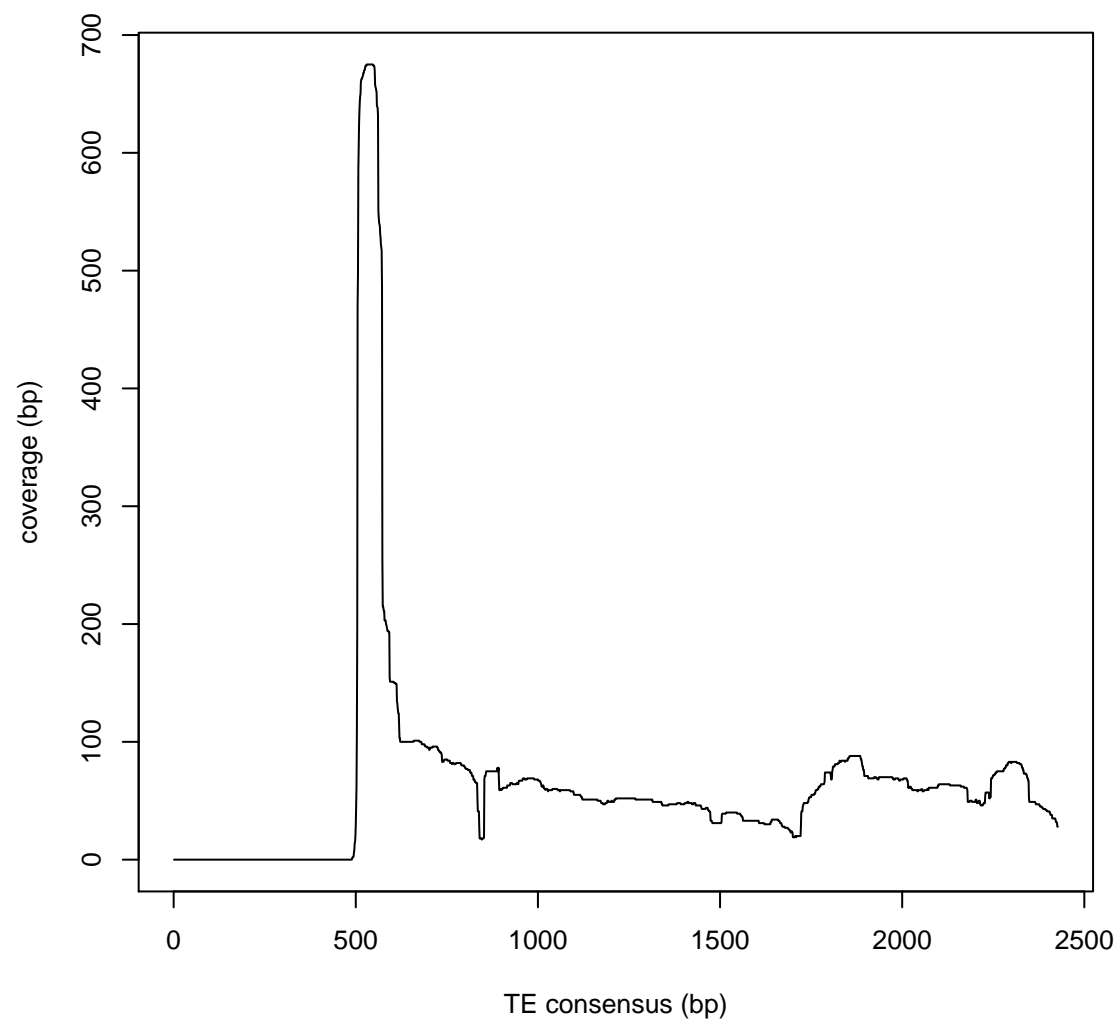


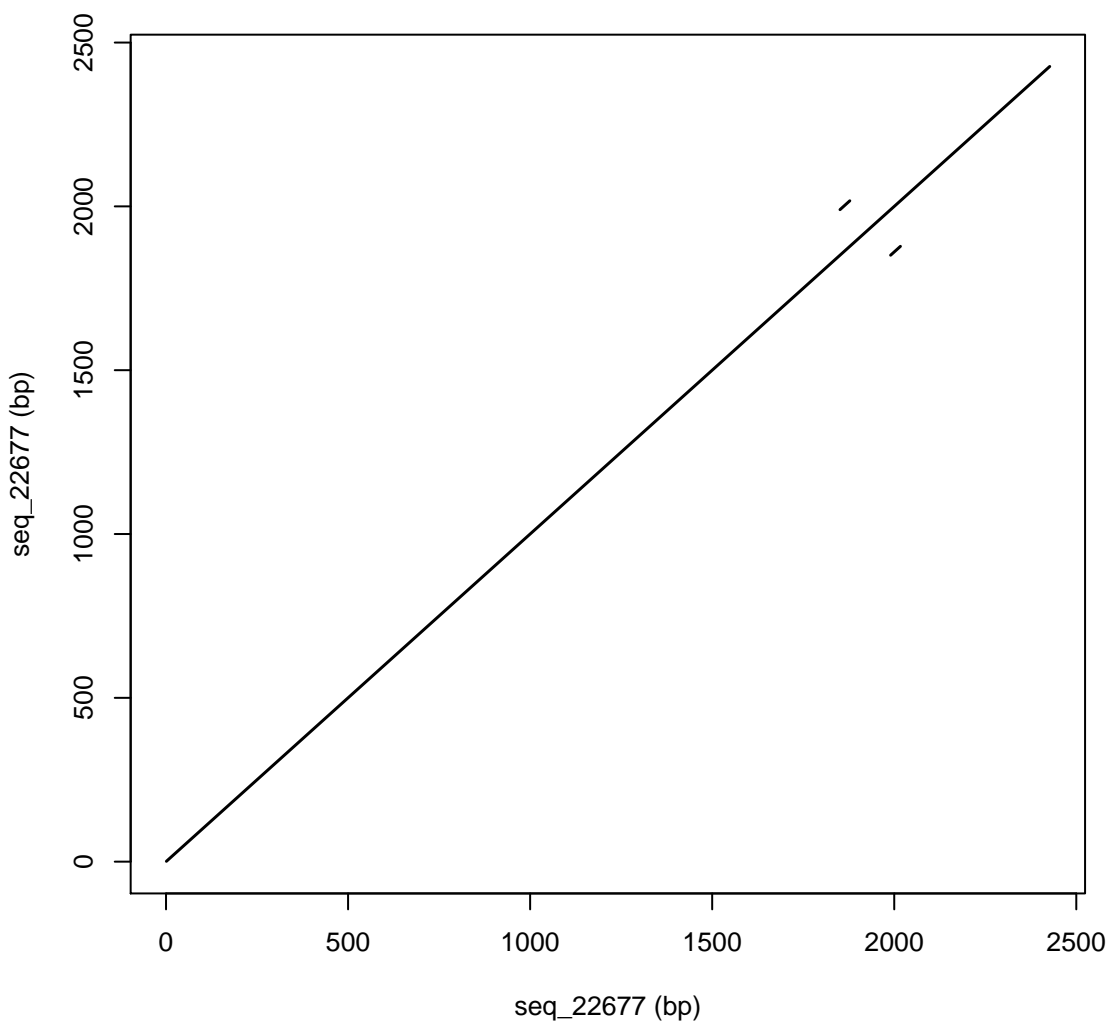
TE: seq_22677
consensus size: 2427bp; fragments: 948; full length: 0 (>=2184.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

