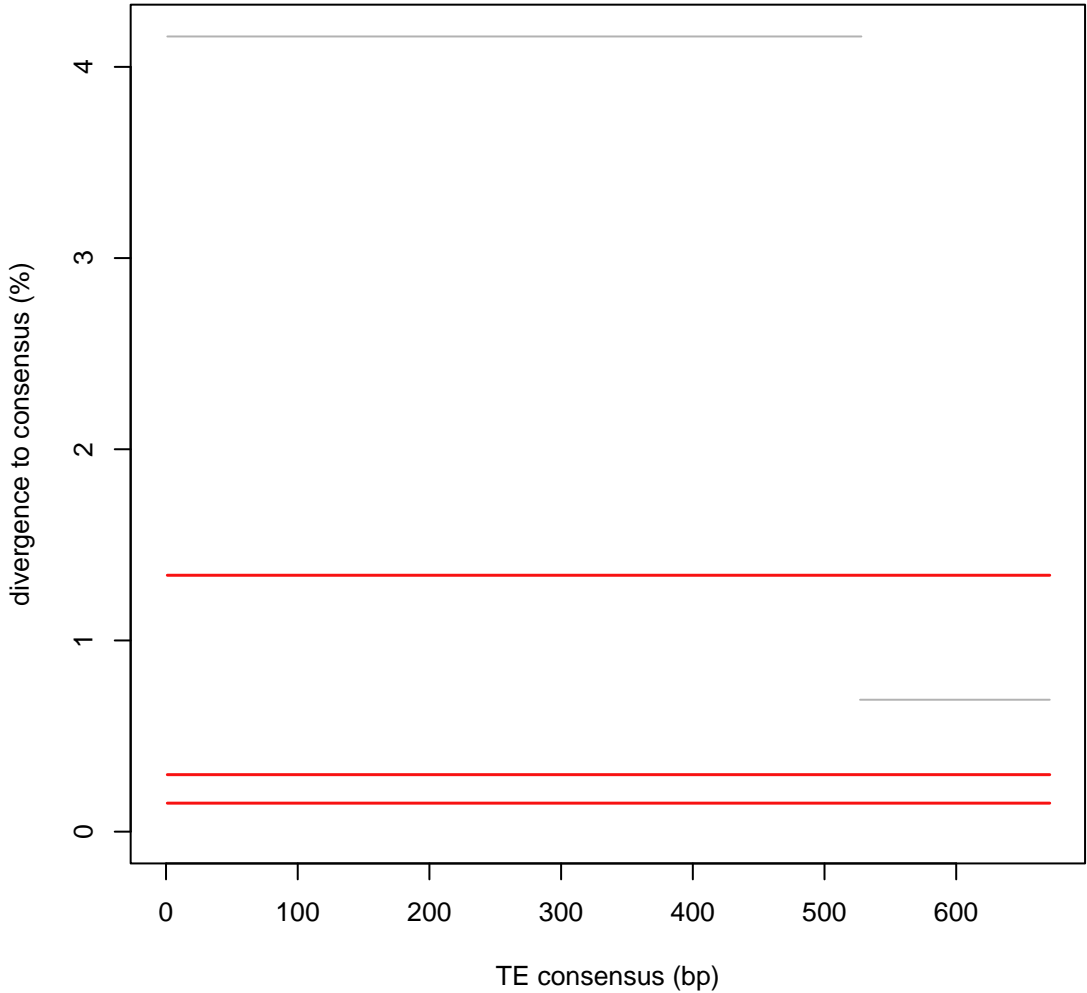
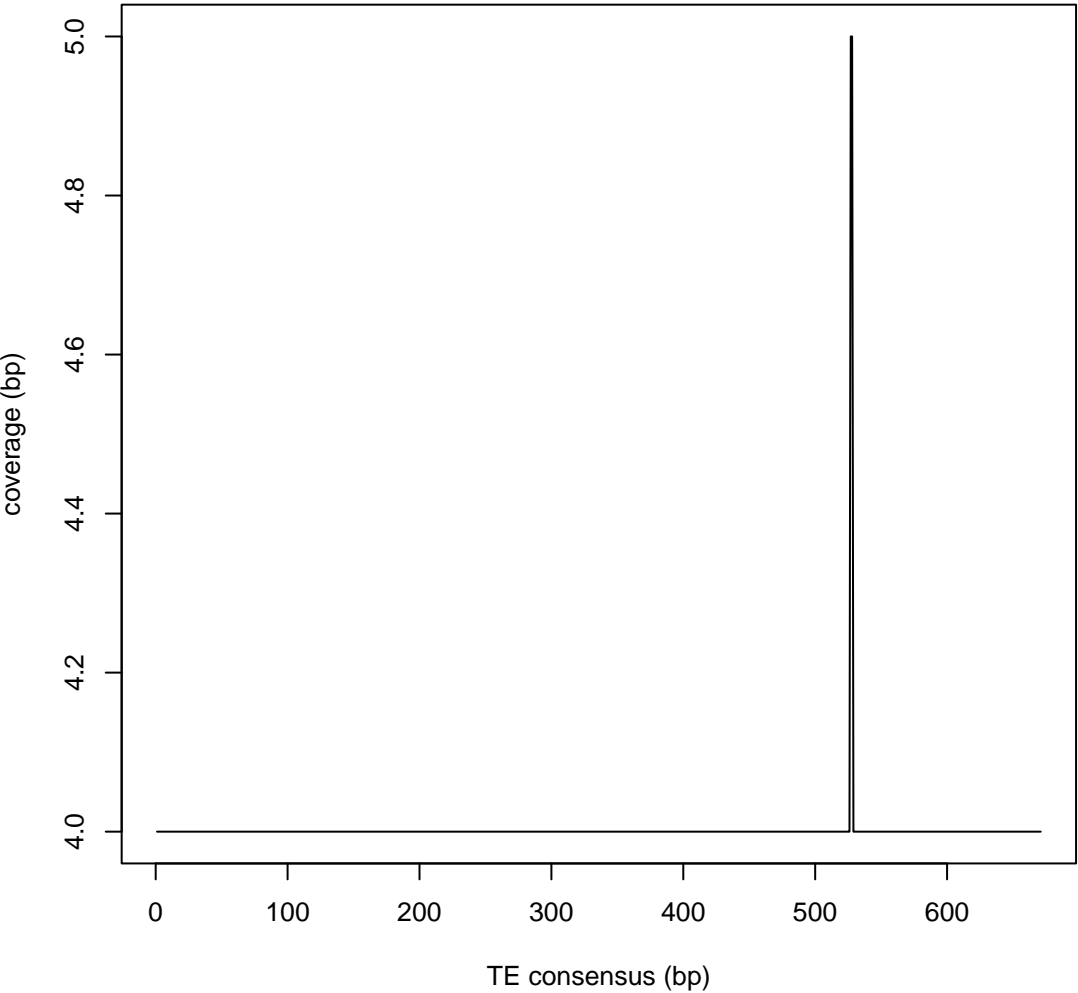


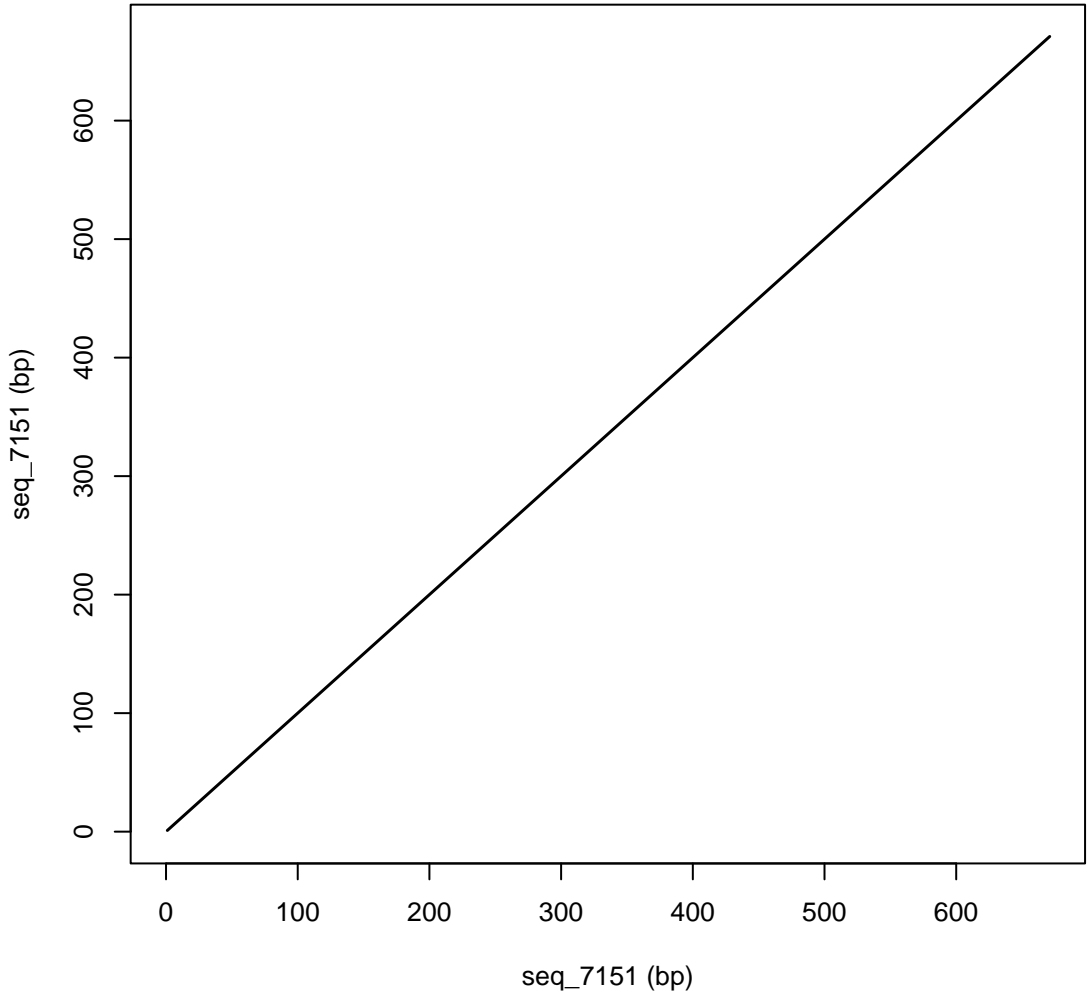
TE: seq_7151
consensus size: 671bp; fragments: 5; full length: 3 (>=603.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

