

Product datasheet for SC202226

OriGene Technologies, Inc.

9620 Medical Center Drive, Ste 200 Rockville, MD 20850, US Phone: +1-888-267-4436 https://www.origene.com techsupport@origene.com EU: info-de@origene.com CN: techsupport@origene.cn

CDK11A (NM_024011) Human 3' UTR Clone

Product data:

Product Type: 3' UTR Clones

Product Name: CDK11A (NM_024011) Human 3' UTR Clone

Vector: pMirTarget (PS100062)

Symbol: CDK11A

Synonyms: CDC2L2; CDC2L3; CDK11-p46; CDK11-p58; CDK11-p110; p58GTA; PITSLRE

ACCN: NM_024011

Insert Size: 558 bp

Insert Sequence: >SC202226 3'UTR clone of NM_024011

The sequence shown below is from the reference sequence of NM_024011. The complete

sequence of this clone may contain minor differences, such as SNPs.

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

AAGCTC

CGAGATTTCGATTCCACCGCCGCCTTCTATGAAAGG

Restriction Sites: Sgfl-Mlul

OTI Disclaimer: Our molecular clone sequence data has been matched to the sequence identifier above as a

point of reference. Note that the complete sequence of this clone is largely the same as the

reference sequence but may contain minor differences, e.g., single nucleotide

polymorphisms (SNPs).

Components: The cDNA clone is shipped in a 2-D bar-coded Matrix tube as 10 ug dried plasmid DNA. The

package also includes 100 pmols of both the corresponding 5' and 3' vector primers in

separate vials.





CDK11A (NM_024011) Human 3' UTR Clone - SC202226

RefSeq: <u>NM 024011.4</u>

Summary: This gene encodes a member of the serine/threonine protein kinase family. Members of this

kinase family are known to be essential for eukaryotic cell cycle control. Due to a segmental duplication, this gene shares very high sequence identity with a neighboring gene. These two genes are frequently deleted or altered in neuroblastoma. The protein kinase encoded by this gene can be cleaved by caspases and may play a role in cell apoptosis. Alternative splicing

results in multiple transcript variants. [provided by RefSeq, Sep 2015]

Locus ID: 728642 MW: 20.8