

Product datasheet for SC210508

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

Dynamin 2 (DNM2) (NM_001005361) Human 3' UTR Clone

Product data:

Product Type: 3' UTR Clones

Product Name: Dynamin 2 (DNM2) (NM_001005361) Human 3' UTR Clone

Vector: pMirTarget (PS100062)

Symbol: DNM2

Synonyms: CMT2M; CMTDI1; CMTDIB; DI-CMTB; DYN2; DYNII; LCCS5

ACCN: NM_001005361

Insert Size: 886 bp

The sequence shown below is from the reference sequence of NM_001005361. The complete

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

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

AGTGCCTGCACTCTGTATTCTATTAATAAACTAAAATAAAGGGAAGACGCTGCTGGTG

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).





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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.

RefSeq: <u>NM 001005361.3</u>

Summary: Dynamins represent one of the subfamilies of GTP-binding proteins. These proteins share

considerable sequence similarity over the N-terminal portion of the molecule, which contains the GTPase domain. Dynamins are associated with microtubules. They have been implicated in cell processes such as endocytosis and cell motility, and in alterations of the membrane that accompany certain activities such as bone resorption by osteoclasts. Dynamins bind many proteins that bind actin and other cytoskeletal proteins. Dynamins can also self-assemble, a process that stimulates GTPase activity. Five alternatively spliced transcripts encoding different proteins have been described. Additional alternatively spliced transcripts may exist, but their full-length nature has not been determined. [provided by RefSeq, Jun

2010]

Locus ID: 1785

MW: 30.3