

Product datasheet for MC226325

Nde1 (NM_001285504) Mouse Untagged Clone

Product data:

Product Type:	Expression Plasmids
Product Name:	Nde1 (NM_001285504) Mouse Untagged Clone
Tag:	Tag Free
Symbol:	Nde1
Synonyms:	2810027M15Rik; AU042936; AW822251; mNudE; Nude
Mammalian Cell Selection:	Neomycin
Vector:	pCMV6-Entry (PS100001)
E. coli Selection:	Kanamycin (25 ug/mL)
Fully Sequenced ORF:	>MC226325 representing NM_001285504 Red=Cloning site Blue=ORF Orange=Stop codon

TTTTGTAATACGACTCACTATAGGGCGCCGGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC
GCC**CGATCGC**

ATGTCCCTGGAAGACTTTGAGCAGCGTTTGAATCAAGCCATTGAAAGAAATGCCTTCCTAGAGAGTGAGC
TGGATGAGAAGGAGAATCTTCTAGAATCTGTGCAGAGGCTGAAGGATGAAGCCCGAGATCTGCGGCAGGA
ATTGGCTGTGCAACAGAAGCAAGACAAGCCCCGGACCCCATGCCAGGCTCAGGGCAAGCCAAGAGGACA
GACATGGCTGTGCAGGCCACAGGCTCTGTACCGTCTACTCCAGTAGCTCACCGAGGACCTAGCTCTGGTT
TGAACACACCAGGAATGTTGAGACGTGGTCTGGACAGCTCCACTAGTGGGACGCCACTCACACCTGCAGC
CCGGATATCAGCCCTAAACATCGTTGGGGACCTGCTTCGAAAGTTGGGGCCCTGGAGTCCAAGCTAGCA
TCATGCAGGAACCTCATGTATGATCAGTCCCAAGCCGGACAAGTGGGCCAGCCTCAGGACGAGGAACCA
AAAACAGAGATGGTGTGACAGAAGGCCAGGCAGTACCAAGTGTGGGCGATAAAGGGTCAGGAAAACGCTT
GGAGTTTGGGAAGCCAGCCTCAGAGCCAGCATCACCGCGCTACCATCCGCCAGGGTGTGGTCAAGCTT
TTGCTT**TAG**

ACGCGTACGCGGCCGCTCGAGCAGAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCCTGGATT
ACAAGGATGACGACGATAAGGTTTAA

Restriction Sites:	SgfI-MluI
ACCN:	NM_001285504
Insert Size:	639 bp



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OTI Disclaimer:	Our molecular clone sequence data has been matched to the reference identifier above as a point of reference. Note that the complete sequence of our molecular clones may differ from the sequence published for this corresponding reference, e.g., by representing an alternative RNA splicing form or single nucleotide polymorphism (SNP).
OTI Annotation:	Clone contains native stop codon, and expresses the complete ORF without any c-terminal tag.
Components:	The ORF clone is ion-exchange column purified and shipped in a 2D barcoded Matrix tube containing 10ug of transfection-ready, dried plasmid DNA (reconstitute with 100 ul of water).
Reconstitution Method:	<ol style="list-style-type: none"> 1. Centrifuge at 5,000xg for 5min. 2. Carefully open the tube and add 100ul of sterile water to dissolve the DNA. 3. Close the tube and incubate for 10 minutes at room temperature. 4. Briefly vortex the tube and then do a quick spin (less than 5000xg) to concentrate the liquid at the bottom. 5. Store the suspended plasmid at -20°C. The DNA is stable for at least one year from date of shipping when stored at -20°C.
Note:	Plasmids are not sterile. For experiments where strict sterility is required, filtration with 0.22um filter is required.
RefSeq:	<u>NM_001285504.1, NP_001272433.1</u>
RefSeq Size:	2145 bp
RefSeq ORF:	639 bp
Locus ID:	67203
UniProt ID:	<u>Q9CZA6</u>
Cytogenetics:	16 A1
Gene Summary:	<p>Required for centrosome duplication and formation and function of the mitotic spindle. Essential for the development of the cerebral cortex. May regulate the production of neurons by controlling the orientation of the mitotic spindle during division of cortical neuronal progenitors of the proliferative ventricular zone of the brain. Orientation of the division plane perpendicular to the layers of the cortex gives rise to two proliferative neuronal progenitors whereas parallel orientation of the division plane yields one proliferative neuronal progenitor and a post-mitotic neuron. A premature shift towards a neuronal fate within the progenitor population may result in an overall reduction in the final number of neurons and an increase in the number of neurons in the deeper layers of the cortex.[UniProtKB/Swiss-Prot Function]</p> <p>Transcript Variant: This variant (d) lacks an exon in the 5' end that results in the use of a downstream AUG, compared to variant a. The encoded isoform (d) has a shorter N-terminal and a distinct C-terminus, compared to isoform a. Sequence Note: This RefSeq record was created from transcript and genomic sequence data to make the sequence consistent with the reference genome assembly. The genomic coordinates used for the transcript record were based on transcript alignments.</p>