

## Product datasheet for MR216969

### Upf1 (NM\_001122829) Mouse Tagged ORF Clone

#### Product data:

Product Type:	Expression Plasmids
Product Name:	Upf1 (NM_001122829) Mouse Tagged ORF Clone
Tag:	Myc-DDK
Symbol:	Upf1
Synonyms:	B430202H16Rik; NORF1; PNORF-1; Rent1; Upflp
Mammalian Cell Selection:	Neomycin
Vector:	pCMV6-Entry (PS100001)
E. coli Selection:	Kanamycin (25 ug/mL)
ORF Nucleotide Sequence:	>MR216969 representing NM_001122829 Red=Cloning site Blue=ORF Green=Tags(s)

TTTTGTAATACGACTCACTATAGGGCGGCCGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC  
GCC**CGATCGCC**

ATGAGTGTGGAGGCGTACGGCCCCAGCTCGAAACACTCACCTTCTTGGACTGAGGAGGCCGAGCTGC  
TCGGCGCCGACCCAGGGCTCCGAGTTCGAATTCACCGACTCACCTTCCCAGCCAGACGCAGACGCC  
CCCCGGCGGCCCGGGCGGGAGGCCCGGGCGAGCGGGCGCAGGCGCGGCCGCGCCAGCTCGAC  
GCACAAGTTGGACCAGAGGGCATCTTGCAAATGGGGCTGTGGATGACAGTGTGGCAAGACCAGCCAGC  
TGCTAGCTGAGCTGAACCTCGAGGAAGATGAAGAGGACACATACTACACTAAGGACTCCCAGTCCACGC  
CTGCAGTACTGTGGAATCCATGATCCTGCCTGCGTGGTTTACTGTAATACCAGCAAGAAGTGGTTCTGC  
AATGGCCGAGGAAATACTTCTGGCAGCCACATTGTGAATCACCTCGTGAGGGCAAATGCAAGGAAGTGA  
CGCTGCACAAGGACGGGCTCTGGGCGAGACCGTGTGGAGTGCTACAAGTGTGGCTGCCCAACGTCTT  
CCTGCTGGGCTTCATCCCTGCGAAGGCCGACTCTGTGGTGGTGTGTTGTGCAGGCAGCCCTGTGCCAGC  
CAGAGCAGCCTGAAGGACATCAACTGGGACAGCTCACAGTGGCAGCCCTAATCCAGGACCGGTGCTTTC  
TGTCATGGCTGGTCAAGATTCCTCTGAGCAGGAGCAGCTGCGAGCACGGCAGATCACGGCACAGCAGAT  
CAACAAGCTGGAAGAGCTCTGGAAGGAAAATCCTTCAGCCACTCTGGAGGACCTGGAGAAGCCAGGCGTA  
GACGAGGAGCCACAGCAGTGTCTCTGCGTTACGAGGATGCTTACCAGTACCAGAACATCTCGGGCCAC  
TGGTCAAGCTGGAGGCTGACTATGACAAGAAGTTGAAGGAGTACAGACTCAAGATAACATCACGGTCAG  
GTGGGACCTGGGCTTAACAAGAAGAGAATCGCCTTCTCACTTTGCCAAGACTGACTCTGGTAATGAG  
GATTTAGTCATAATTTGGTTAAGAGACATGCGGCTCATGCAGGGTATGAGATCTGTCTGCGGTACAAA  
GGGATCTGGCGCCCTGTGGAAGGGATTGGCCACGTCATCAAGGTTCTGATAATTATGGTATGAGAT  
TGCTATTGAGCTCCGCAGCAGCGTGGGTGCCCTGTGGAAGTACCCACAACCTCCAAGTGGATTTGTG  
TGGAAGTCAACCTCTTTGATAGGATGCAGAGTCACTGAAGACCTTCGCTGTGGACGAGACCTCTGTG  
CAGGGTATATTTACCACAAGCTGCTGGCCACGAGGTGGAGGATGTGGTATCAAGTCCAGCTGCCAAA  
GCGCTTACAGCTCAGGGCTCCCTGACCTCAACCACTCTCAGGTGTATGCTGTGAAGACCGTGTGCAG



[View online »](#)

AGACCACTCAGCCTCATCCAGGGCCCTCCAGGCACAGGCAAGACTGTGACATCAGCCACTATTGTCTACC  
ACCTTGCTCGGCAGGGCAATGGGCCTGTACTGGTTTGTGCTCCAAGTAACATCGCTGTGGACCAGCTCAC  
AGAGAAGATCCACCAGACAGGACTGAAGGTCGTACGCCTCTGTGCCAAGAGCCGTGAGGCCATTGACTCC  
CCAGTGTCTTCTGGCTTTGCACAACCAGATCAGGAACATGGACAGCATGCCTGAGCTGCAGAAGCTGC  
AGCAGCTAAAGGATGAGACAGGCGAGCTGTCATCTGCAGATGAGAAGCGGTACCGGGCGCTTAAGCGCAC  
AGCTGAGAGAGAACTTCTCATGAATGCAGATGTCATATGCTGCACATGTGTGGTGTGGTGACCCGAGG  
CTGGCCAAGATGCAGTTCGGTTCCATCCTCATTGACGAGAGACCCAGGCCACTGAGCCTGAGTGCATGG  
TGCTGTAGTCCCTGGGGCCAAGCAGCTAATCCTCGTTCGGTGACCACTGCCAGCTGGGCCAGTGGTGAT  
GTGCAAGAAGGCAGCCAAGGCCGACTGTCACAATCGCTCTTCGAGCGCTTGGTGGTGTGGGCATCCGG  
CCCATCCGCCTGCAGGTGCAATACCGCATGCACCCTGCACTCAGCGCCTTCCGTCCAACATCTTCTACG  
AGGGCTCATTGCAGAATGGCGTCACTGCAGCGGATCGTGTCAAAAAGGCTTTGACTTCCAGTGGCCACA  
ACCTGACAAGCCTATGTTCTTCTACGTGACGCAGGGCCAGGAGGATTGCCAGCTCTGGCACATCTAC  
CTCAACAGGACGGAGGCCAATGTGGAGAAGATAACTACGAAGCTGTTGAAGGCAGGTGCAAAGCCTG  
ACCAGATCGGCATCATCACCCCTACGAGGGCCAGCGCTTACTTGGTGCAGTACATGCAGTTCAGCGG  
CTCCCTGCACAAAAGCTCTACCAGGAAGTGAAATTGCCAGTGTGGACGCCTTCCAGGGCCGGGAGAAG  
GACTTCATCATTCTGTCTGCGTGCAGCCTAACATCAGGGCATTGGTTCCATAACGACCCCGGC  
GTCTGAATGTGGCTCTCACCAGAGCAAGATATGGCGTGATCATTGTGGTAAACCAAAGGCCCTGTGAA  
GCAGCCCTGTGGAATCACCTGTGAGCTACTACAAGGAACAGAAGGCGTAGTGAAGGGCCGCTCAAC  
AACCTACGTGAGAGCCTCATGCAGTTCAGCAAGCCTCGCAAACCTGTCAACACTGTCAACCCGGGTGCC  
GCTTCATGACTACTGCCATGTACGATGCCCGTGAAGCCATCATCCCCGGGTCTGTCTATGACCGCAGCAG  
CCAGGGCCGGCCCTCGAACATGTACTTCCAGACCCATGACCAGATTAGTATGATCAGCGCAGGCCCCAGC  
CACGTGGCTGCCATGAACATCCCTATTCCTTCAACTTGGTTCATGCCTCCCATGCCGCCACTGGCTACT  
TCGGACAGGCCAACGGCCGGCAGCTGGTGGGGCACCCAAAAACCAAGACTGGCCGTGGGGCCGCCA  
GAAGAACCCTTTGGGCTTCTGGGCCAGCCAGACCACCTTCCAACAGCCAGGCCAGCCAGGACGTG  
GCCTCCAGCCCTTTTACAGGGTGCCCTCACACAGGTTACGTGTCCATGAGCCAGCCCTCTCAGATGA  
GCCAGCCTGGCCTCTCCAGCCAGAAGTGTCCAGGACAGCTACCTCGGTGATGAGTTTAAATCACAGAT  
TGACGTGGCACTCTACAAGACTCCACATACCAGGGAGAGCGGGCATACCAGCACGGCGGGTACCAGGG  
CTGTCCCAGTAC

ACGCGTACGCGGCCGCTCGAGCAGAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCCTGGATT  
ACAAGGATGACGACGATAAGGTTTAA

Protein Sequence: >MR216969 representing NM\_001122829  
 Red=Cloning site Green=Tags(s)

```
MSVEAYGPSSQTLTFLDTEEAELLGADTQGSEFEFTDFTLPSQTQTPPGGPGGAGGPGGAGAGGAAGQLD
AQVGPEGILQNGAVDDSVAKTSQLLAELNFEDEEDTYTKDLPVHACSYCGIHDPACVVYCNNTSKKWFC
NGRGNTSGSHIVNHLVRAKCKEVLHKDGPLGETVLECYNGCRNVFLLGFIPAKADSVVLLCRQPCAS
QSSLKDINWSSQWQPLIQDRCFLSWLVKIPSEQEQLRARQITAQQINKLEELWKENPSATLEDLEKPGV
DEEPQHVLLRYEDAYQYQNI FGPLVKLEADYDKLLESQTQDNITVRWDLGLNKKRIAFFTLPKTDSGNE
DLVIIWLRDMRLMQGDEICLRYKGD LAPLWKGIGHVIKVPDNYGDEIAIELRSSVGAPVEVTHNFQVDFV
WKSTSFDRMQSALKTFVAVDETSVSGYIYHKLLGHEVEDVVIKQQLPKRFTAQGLPDLNHSQVYAVKTVLQ
RPLSLIQPPGTGKTVTSATIVYHLARQNGPVLVCAPSNIAVDQLTEKIHQTGLKVVRLCAKSREAI DS
PVSFLALHNQIRNMDMPQLKQLKDETGELSSADEKRYRALKRTAERELLMNADVICCTCVGAGDPR
LAKMQFRSILIDESTQATEPECMVPVVLGAKQLILVGDHCQLGPVVMCKKAAKAGLSQSLFERLVVLGIR
PIRLQVQYRMHPALSAFPSNIFYEGSLQNGVTAADRVKKGDFQWPQDPKPMFFVYTQGGEEIASSGTSY
LNRTEAANVEKITTLLKAGAKPDQIGIITPYEGQRSYLVOYMQFSGSLHTKLYQEVEIASVDAFQGREK
DFIILSCVRANEHQIGIFLNDPRLNVALTRARYGVII VGNPKALSKQPLWNHLLSYYKEQKALVEGPLN
NLRESLMQFSKPRKLVNTVNPGARFMTTAMYDAREAIIPGSVYDRSSQGRPSNMYFQTHDQISMISAGPS
HYAAMNIPIPFNLVMPMPPPGYFGQANGPAAGRGT PKTKTGRGGRQKNRFGLPGPSQTTLPNSQASQDV
ASQPF SQGAL TQGYVMSQPSQMSQPGLSQPELSQDSYLGDEFKSIDVALSQDSTYQGERAYQHGGVTG
LSQY
```

TRTRPLEQKLISEEDLAANDILDYKDDDDKV

Chromatograms: [https://cdn.origene.com/chromatograms/mm9101\\_f09.zip](https://cdn.origene.com/chromatograms/mm9101_f09.zip)

Restriction Sites: SgfI-MluI

Cloning Scheme:



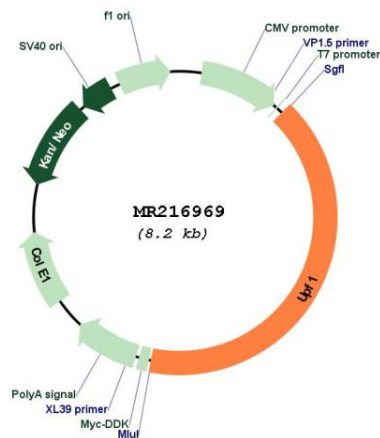
ACCN: NM\_001122829

ORF Size: 3372 bp

<b>OTI Disclaimer:</b>	The molecular sequence of this clone aligns with the gene accession number as a point of reference only. However, individual transcript sequences of the same gene can differ through naturally occurring variations (e.g. polymorphisms), each with its own valid existence. This clone is substantially in agreement with the reference, but a complete review of all prevailing variants is recommended prior to use. <a href="#">More info</a>
<b>OTI Annotation:</b>	This clone was engineered to express the complete ORF with an expression tag. Expression varies depending on the nature of the gene.
<b>Components:</b>	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).
<b>Reconstitution Method:</b>	<ol style="list-style-type: none"><li>1. Centrifuge at 5,000xg for 5min.</li><li>2. Carefully open the tube and add 100ul of sterile water to dissolve the DNA.</li><li>3. Close the tube and incubate for 10 minutes at room temperature.</li><li>4. Briefly vortex the tube and then do a quick spin (less than 5000xg) to concentrate the liquid at the bottom.</li><li>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.</li></ol>
<b>RefSeq:</b>	<a href="#">NM_001122829.2</a> , <a href="#">NP_001116301.1</a>
<b>RefSeq Size:</b>	4657 bp
<b>RefSeq ORF:</b>	3375 bp
<b>Locus ID:</b>	19704
<b>UniProt ID:</b>	<a href="#">Q9EPU0</a>
<b>Cytogenetics:</b>	8 34.15 cM
<b>MW:</b>	124 kDa

**Gene Summary:**

RNA-dependent helicase and ATPase required for nonsense-mediated decay (NMD) of mRNAs containing premature stop codons. Is recruited to mRNAs upon translation termination and undergoes a cycle of phosphorylation and dephosphorylation; its phosphorylation appears to be a key step in NMD. Recruited by release factors to stalled ribosomes together with the SMG1C protein kinase complex to form the transient SURF (SMG1-UPF1-eRF1-eRF3) complex. In EJC-dependent NMD, the SURF complex associates with the exon junction complex (EJC) (located 50-55 or more nucleotides downstream from the termination codon) through UPF2 and allows the formation of an UPF1-UPF2-UPF3 surveillance complex which is believed to activate NMD. Phosphorylated UPF1 is recognized by EST1B/SMG5, SMG6 and SMG7 which are thought to provide a link to the mRNA degradation machinery involving exonucleolytic and endonucleolytic pathways, and to serve as adapters to protein phosphatase 2A (PP2A), thereby triggering UPF1 dephosphorylation. UPF1 can also activate NMD without UPF2 or UPF3, and in the absence of the NMD-enhancing downstream EJC indicative for alternative NMD pathways. Plays a role in replication-dependent histone mRNA degradation at the end of phase S; the function is independent of UPF2. For the recognition of premature termination codons (PTC) and initiation of NMD a competitive interaction between UPF1 and PABPC1 with the ribosome-bound release factors is proposed. The ATPase activity of UPF1 is required for disassembly of mRNPs undergoing NMD (By similarity). Essential for embryonic viability.[UniProtKB/Swiss-Prot Function]

**Product images:**

Circular map for MR216969