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## Product datasheet for SC119361

## Caspase-6 (CASP6) (NM_001226) Human Untagged Clone

## Product data:

Product Type:
Product Name:

## Tag:

Symbol:
Synonyms:
Mammalian Cell
Selection:
Vector:
E. coli Selection:

Fully Sequenced ORF:

Expression Plasmids
Caspase-6 (CASP6) (NM_001226) Human Untagged Clone
Tag Free
Caspase-6
MCH2
None
pCMV6-XL5
Ampicillin ( $100 \mathrm{ug} / \mathrm{mL}$ )
>OriGene ORF within SC119361 sequence for NM_001226 edited (data generated by NextGen Sequencing)
ATGAGCTCGGCCTCGGGGCTCCGCAGGGGGCACCCGGCAGGTGGGGAAGAAAACATGACA GAAACAGATGCCTTCTATAAAAGAGAAATGTTTGATCCGGCAGAAAAGTACAAAATGGAC CACAGGAGGAGAGGAATTGCTTTAATCTTCAATCATGAGAGGTTCTTTTGGCACTTAACA CTGCCAGAAAGGCGGGGCACCTGCGCAGATAGAGACAATCTTACCCGCAGGTTTTCAGAT CTAGGATTTGAAGTGAAATGCTTTAATGATCTTAAAGCAGAAGAACTACTGCTCAAAATT CATGAGGTGTCAACTGTTAGCCACGCAGATGCCGATTGCTTTGTGTGTGTCTTCCTGAGC CATGGCGAAGGCAATCACATTTATGCATATGATGCTAAAATCGAAATTCAGACATTAACT GGCTTGTTCAAAGGAGACAAGTGTCACAGCCTGGTTGGAAAACCCAAGATATTTATCATT CAGGCATGTCGGGGAAACCAGCACGATGTGCCAGTCATTCCTTTGGATGTAGTAGATAAT CAGACAGAGAAGTTGGACACCAACATAACTGAGGTGGATGCAGCCTCCGTTTACACGCTG CCTGCTGGAGCTGACTTCCTCATGTGTTACTCTGTTGCAGAAGGATATTATTCTCACCGG GAAACTGTGAACGGCTCATGGTACATTCAAGATTTGTGTGAGATGTTGGGAAAATATGGC TCCTCCTTAGAGTTCACAGAACTCCTCACACTGGTGAACAGGAAAGTTTCTCAGCGCCGA GTGGACTTTTGCAAAGACCCAAGTGCAATTGGAAAGAAGCAGGTTCCCTGTTTTGCCTCA ATGCTAACTAAAAAGCTGCATTTCTTTCCAAAATCTAATTAA

Clone variation with respect to NM_001226.3

| ead Nucleotide | >OriGene 5' read for NM_001226 unedited |
| :---: | :---: |
| Sequence: | TGTAATACGACTCACTATAGGGCGGCCGCGAATTCGGCACGAGGGGAGGGCAAGGTGTCT |
|  | GGCTGCGCGTTTGGCTGCAATGAGCTCGGCCTCGGGGCTCCGCAGGGGGCACCCGGCAGG |
|  | TGGGGAAGAAAACATGACAGAAACAGATGCCTTCTATAAAAGAGAAATGTTTGATCCGGC |
|  | AGAAAAGTACAAAATGGACCACAGGAGGAGAGGAATTGCTTTAATCTTCAATCATGAGAG |
|  | GTTCTTTTGGCACTTAACACTGCCAGAAAGGCGGGGCACCTGCGCAGATAGAGACAATCT |
|  | TACCCGCAGGTTTTCAGATCTAGGATTTGAAGTGAAATGCTTTAATGATCTTAAAGCAGA |
|  | AGAACTACTGCTCAAAATTCATGAGGTGTCAACTGTTAGCCACGCAGATGCCGATTGCTT |
|  | TGTGTGTGTCTTCCTGAGCCATGGCGAAGGCAATCACATTTATGCATATGATGCTAAAAT |
|  | CGAAATTCAGACATTAACTGGCTTGTTCAAAGGAGACAAGTGTCACAGCCTGGTTGGAAA |
|  | ACCCAAGATATTTATCATTCAGGCATGTCGGGGAAACCAGCACGATGTGCCAGTCATTCC |
|  | TTTGGATGTAGTAGATAATCAGACAGAGAAGTTGGACACCAACATAACTGAGGTGGATGC |
|  | AGCCTCCGTTTACACGCTGCCTGCTGGAGCTGACTTCCTCATGTNGTACTCTGTTGCAGA |
|  | AGGATATTATTCTCACCGGGAAACTGTGAACGGCTCATGGTACATTCAAGAATTGTGTGA |
|  | GATGTTGGGAAAATATGGNNCTCTCCTTTAAAGTCACAGAACTCCTCACACTGGTGAACA |
|  | NGNANAGTTCTCAGCGCCCGAGTGGACTTTNNGCAAGACCCNAGTGCANNTGGAAGNAAG |
|  | NCAGNTCCCTGGTTTTGCCTCA |
| 3' Read Nucleotide Sequence: | >OriGene 3' read for NM_001226 unedited |
|  | CGGCCGCAATCTAGNATCGAGTTTTTTTTTTTTTTTTTTGAATTTTGAATATTTTATTAA |
|  | GGGATGATTTAATTCCTGGATTTCAAAAACCTTTAAAAACATCAAACAATAAACTTTTAT |
|  | AAAAAAGTGACAAAATACAAGTTAAATCAGCTAGATTTTTGTGTAACCCTGCAGTTTATT |
|  | AAAAAATAATACAAATGCTTATAAATTTTTATTTATCTAATTTCCTGAGGATTTTGTTTC |
|  | TCCATATCAAACCCTTTCACCATGGCCAACATGAACTTTTTTTTTTTTTTTTTAAGGCAG |
|  | TTCTCTGCTAGGCATTAAACTTTAAAACATTTGAATCATTGGACCATAATGCTTCACCCT |
|  | AACGATATTTATATAAAAGGAAGAGAAAGACATTTTCTTTTTTTTGAGACGGAGTTTCAC |
|  | TCGTTGCCCAGGCTGGAGTGCAATGGCGCAATCTCGGCTCACCGCAGCCTCCACCTCCTG |
|  | GGTTCAAGTGATTCTCCTGCCTCAGCCCTCCAAGTAGCTGGGATTGCAGGCATGCGCCGC |
|  | CATGCCTAGCTAAATTTTTTTTTGCATTTTTAGTAGAGACGGGGCTTCTCCATGTTGGTC |
|  | AGGCTGGTCTCGAACTCCCGACCTCAGGTGATCCGCCCACCTTGGACTCCCAAAGTGCTG |
|  | GGATTACAGGTGTGAGTAACCACGCCTGGCTGAGAAAGCCATTTTTAATACAGAGTGTAA |
|  | AATTAGATAGCCCCTATTAATTAATTAGATTTCTGGAAAGAAATGCAGCTCTTTAGTTAG |
|  | CATTGAGGCAAACAGGGAACCTGCTTCTTTCCATTGCACTTGGGTCTTTGCAAAGTCCCT |
|  | TGGGGCTGAAAACTTCCTGTCACCATGTGAGGGTTTGTGACTCTAAGGAGAGCATATTTT |
|  | CCACATTTACAAAATTTGATGTCCATGACCGTACAAGTTCCGGGAGATAATTTTCTGCAC |
|  | AATAA |
| Restriction Sites: | Notl-Notl |
| ACCN: | NM_001226 |
| Insert Size: | 1600 bp |

## OTI Disclaimer:

Components:

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).

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: | 1. Centrifuge at $5,000 \mathrm{xg}$ for 5 min . <br> 2. Carefully open the tube and add 100 ul of sterile water to dissolve the DNA. <br> 3. Close the tube and incubate for 10 minutes at room temperature. <br> 4. Briefly vortex the tube and then do a quick spin (less than 5000 xg ) to concentrate the liquid at the bottom. <br> 5. Store the suspended plasmid at $-20^{\circ} \mathrm{C}$. The DNA is stable for at least one year from date of shipping when stored at $-20^{\circ} \mathrm{C}$. |
| :---: | :---: |
| RefSeq: | NM 001226.3 NP 001217.2 |
| RefSeq Size: | 1661 bp |
| RefSeq ORF: | 882 bp |
| Locus ID: | 839 |
| UniProt ID: | P55212 |
| Cytogenetics: | 4 q 25 |
| Domains: | CASc, ICE_p10, ICE_p20 |
| Protein Families: | Druggable Genome, Protease, Stem cell - Pluripotency |
| Protein Pathways: | Apoptosis |
| Gene Summary: | This gene encodes a member of the cysteine-aspartic acid protease (caspase) family of enzymes. Sequential activation of caspases plays a central role in the execution-phase of cell apoptosis. Caspases exist as inactive proenzymes which undergo proteolytic processing at conserved aspartic acid residues to produce two subunits, large and small, that dimerize to form the active enzyme. This protein is processed by caspases 7,8 and 10 , and is thought to function as a downstream enzyme in the caspase activation cascade. Alternative splicing of this gene results in multiple transcript variants that encode different isoforms. [provided by RefSeq, Oct 2015] <br> Transcript Variant: This variant (alpha) represents the longer transcript and encodes the longer isoform (alpha). 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. |

