

## Product datasheet for **SC331243**

### eIF3s8 (EIF3C) (NM\_001199142) Human Untagged Clone

#### Product data:

|               |                                                    |
|---------------|----------------------------------------------------|
| Product Type: | Expression Plasmids                                |
| Product Name: | eIF3s8 (EIF3C) (NM_001199142) Human Untagged Clone |
| Tag:          | Tag Free                                           |
| Symbol:       | eIF3s8                                             |
| Synonyms:     | eIF3-p110; EIF3CL; EIF3S8                          |
| Vector:       | pCMV6-Entry (PS100001)                             |



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Fully Sequenced ORF: >SC331243 representing NM\_001199142.  
 Blue=Insert sequence Red=Cloning site Green=Tag(s)

ATGTCGCGGTTTTTACCACCGGTTCCGACAGCGAGTCCGAGTCGTCCTTGTCGGGGGAGGAGCTCGTC  
 ACCAAACCTGTCGGAGGCAACTATGGCAAACAGCCATTGTTGCTGAGCGAGGATGAAGAAGATACCAAG  
 AGAGTTGTCGCGAGTCCCAAGGACAAGAGTTTGAGGAGCTGACCAACCTTATCCGGACCATCCGTAAT  
 GCCAATAAGATTGCTGATGTCACCAAGTCCCTGGAAGAGTTTGAGCTCCTGGGAAAAGCATATGGGAAG  
 GCCAAAAGCATTGTGGACAAAGAAGGTGTCCCGGTTCTATATCCGCATTCTGGCTGACCTAGAGGAC  
 TATCTTAATGAGCTTTGGGAAGATAAGGAAGGGAAGAAGAAGATGAACAAGAACAATGCCAAGGCTCTG  
 AGCACCTTGCGTCAGAAGATCCGAAAATACAACCGTGATTTGAGTCCCATATCACAAGCTACAAGCAG  
 AACCCCGAGCAGTCTGCGGATGAAGATGCTGAGAAAAATGAGGAGGATTCAGAAGGCTCTTCAGATGAG  
 GATGAGGATGAGGACGGAGTCACTGCTGCAACTTTCTGAAGAAGAAATCAGAAGCTCCTTCTGGGGAG  
 AGTCGCAAGTTCCCTAAAAAGATGGATGATGAAGATGAGGACTCAGAAGATTCCGAAGATGATGAAGAC  
 TGGGACACAGGTTCCACATCTTCCGATTCCGACTCAGAGGAGGAAGAAGGGAAACAAACCCGCTGGCC  
 TCAAGATTTCTTAAAAAGCACCACACAGATGAGGACAAGAAGGCAGCCGAGAAGAAACGGGAGGAC  
 AAAGCTAAGAAGAAGCAGCAGGAAATCCAAGCGCTGGATGAGGAGGAGGAGACAATGAAGCGGGG  
 GAGTGGGAAAGGGTCCGGGGCGGAGTGCCTGGTTAAGGAGAAGCCAAAATGTTTGCCAAGGGAAT  
 GAGATCACCCATGCTGTTGTTATCAAGAACTGAATGAGATCCTACAGGCACGAGGCAAGAAGGGAAT  
 GATCGTGTGCCAGATTGAGCTGCTGCAACTGCTGGTTCAGATTGCAGCGGAAAACAACCTGGGAGAG  
 GCGTCAATTGTCAAGATCAAGTTCAATATCATCGCTCTCTATGACTACAACCCCAACCTGGCAACC  
 TACATGAAGCCAGAGATGTTGGGGAAGTGCCTGGACTGCATCAATGAGCTGATGGATATCCTGTTTGA  
 AATCCCAACATTTTTGTTGGAGAGAATATTCTGGAAGAGAGTGAAGACCTGCACAACGCTGACCAACCA  
 CTGCGGTCCGTTGCTGATCCTAACTCTGGTGGAAACGAATGGATGAAGAATTTACCAAAAATAATGCAA  
 AATACTGACCTCACTCCCAAGAGTACGTGGAGCACTTGAAGGATGAGGCCAGGTGTGTGCCATCATC  
 GAGCGTGTGACGCGCTACCTGGAGGAGAAGGGCACTACCGAGGAGGTCTGCCGCATCTACCTGCTGCGC  
 ATCCTGCACACCTACTACAAGTTTGATTACAAGGCCCATCAGCGACAGCTGACCCCGCTGAGGGCTCC  
 TCAAAGTCTGAGCAAGACCAGGCAGAAAATGAGGGCGAGGACTCGGCTGTGTTGATGGAGAGACTGTGC  
 AAGTACATCTACGCCAAGGACCGCACAGACCGGATCCGCACATGTCCATCCTCTGCCACATCTACCAC  
 CATGCTCTGCACTCGCGTGGTACCAGGCCCGCACCTCATGCTCATGAGCCACTGCAGGACAACATT  
 CAGCATGCAGACCCGCGAGTGCAGATCCTTTACAACCGCACCATGGTGCAGCTGGGCATCTGTGCCCTT  
 CGCAAGGCTGACCAAGGACGCACACAACGCCCTGCTGGACATCCAGTGCAGTGGCCGAGCCAAGGAG  
 CTCTGGGCCAGGGCCTGCTGCTGCGCAGCCTGCAGGAGCGCAACCAGGAGCAGGAGAAGGTGGAGCGG  
 CGCGTCAAGTCCCTTCCACCTGCACATCAACCTGGAGCTGCTGGAGTGTGTCTACCTGGTGTCTGCC  
 ATGCTCCTGGAGATCCCTACATGGCCGCCATGAGAGCGATGCCCGCCGACGCATGATCAGCAAGCAG  
 TTCCACCACAGCTGCGCGTGGGCGAGCGACAGCCCTGCTGGTCCCCCTGAGTCCATGCGGGAACAT  
 GTGGTTCGCTGCCTCAAGGCCATGAAGATGGGTGACTGGAAGACCTGTACAGTTTTATCATCAATGAG  
 AAGATGAATGGGAAAGTGTGGGACCTTTTCCCCAGGCTGACAAAGTCCGCACCATGCTGGTTAGGAAG  
 ATCCAGGAAGAGTCACTGAGGACCTACCTTTACCTACAGCAGTGTCTATGACTCCATCAGCATGGAG  
 ACGCTGTCAGACATGTTTGAGCTGGATCTGCCACTGTGCACTCCATCATCAGCAAAATGATCATTAAAT  
 GAGGAGCTGATGGCTCCCTGGACCAGCCAACACAGACAGTGGTATGCACCGCACTGAGCCCACTGCC  
 CAGCAGAACCTGGCTCTGCACTGGCCGAGAAGCTGGGACGCTGGTGGAGAACAACGAACGGGTGTTT  
 GACCACAAGCAGGGCACCTACGGGGGCTACTTCCGAGACCAGAAGGACGGCTACCGCAAAAACGAGGGG  
 TACATGCGCCGCGGTGGCTACCGCCAGCAGCAGTCTCAGACGGCCTACTGA

Restriction Sites: Sgfl-Mlul  
 ACCN: NM\_001199142  
 Insert Size: 2742 bp

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|-------------------------------|---------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|
| <b>OTI Disclaimer:</b>        | 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).                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    |
| <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_001199142.1</a>                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    |
| <b>RefSeq Size:</b>           | 3265 bp                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           |
| <b>RefSeq ORF:</b>            | 2742 bp                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           |
| <b>Locus ID:</b>              | 8663                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              |
| <b>UniProt ID:</b>            | <a href="#">Q99613</a>                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            |
| <b>Cytogenetics:</b>          | 16p11.2                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           |
| <b>Protein Families:</b>      | Druggable Genome                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  |
| <b>MW:</b>                    | 105.3 kDa                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         |
| <b>Gene Summary:</b>          | <p>Component of the eukaryotic translation initiation factor 3 (eIF-3) complex, which is required for several steps in the initiation of protein synthesis (PubMed:17581632, PubMed:25849773, PubMed:27462815). The eIF-3 complex associates with the 40S ribosome and facilitates the recruitment of eIF-1, eIF-1A, eIF-2:GTP:methionyl-tRNA<sub>i</sub> and eIF-5 to form the 43S pre-initiation complex (43S PIC). The eIF-3 complex stimulates mRNA recruitment to the 43S PIC and scanning of the mRNA for AUG recognition. The eIF-3 complex is also required for disassembly and recycling of post-termination ribosomal complexes and subsequently prevents premature joining of the 40S and 60S ribosomal subunits prior to initiation (PubMed:17581632). The eIF-3 complex specifically targets and initiates translation of a subset of mRNAs involved in cell proliferation, including cell cycling, differentiation and apoptosis, and uses different modes of RNA stem-loop binding to exert either translational activation or repression (PubMed:25849773).[UniProtKB/Swiss-Prot Function]</p> <p>Transcript Variant: This variant (3) differs in the 5' UTR compared to variant 1. Variants 1-4 encode the same isoform (a).</p> |