

## Product datasheet for **SC326601**

### **EYS (NM\_001142801) Human Untagged Clone**

#### **Product data:**

|                           |  |
|---------------------------|--|
| Product Type:             | Expression Plasmids  |
| Product Name:             | EYS (NM_001142801) Human Untagged Clone  |
| Tag:                      | Tag Free   |
| Symbol:                   | EYS  |
| Synonyms:                 | bA74E24.1; bA166P24.2; bA307F22.3; C6orf178; C6orf179; C6orf180; dj22117.2; dj303F19.1; dj1018A4.2; EGFL10; EGFL11; RP25; SPAM |
| Mammalian Cell Selection: | Neomycin   |
| Vector:                   | pCMV6-Entry (PS100001)   |
| E. coli Selection:        | Kanamycin (25 ug/mL)   |



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**Fully Sequenced ORF:** >SC326601 representing NM\_001142801.  
 Blue=Insert sequence Red=Cloning site Green=Tag(s)

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GCTCGTTTAGTGAACCGTCAGAATTTTGTAAACGACTCACTATAGGGCGGCCGGGAATTCGTGACTG
GATCCGGTACCGAGGAGATCTGCCGCCCGCATCGCC
ATGACTGACAAATCAATCGTCATTCTGAGCCTGATGGTTTTTACAGCTCTTTCATAAATGGAAAAACA
TGTAGACGGCAATTGGTGAAGAATGGCATCCACAACCCTCATCATATGTGGTAAATGGACACTAACA
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TGCCTGAAGATGCAACCTATGTGAACGATCCTGAAGATAATAATTCTTCATGTTGGTCCCACATGAA
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CTATGTTTTCTCAGATGGGCTGGCAACATGTATCTGAAAATACAACCTGATGATCAAGAAAATGAGTGT
CAACATGAAGCTGTTTGTAAAGTGAATAAATAGACCCAGAAGAATCCTCAACCGGTGATACCACAC
CAAATTCAGCAACATATAGAAAAGTTTATACAGCATGACCAGGTGGGATTTATCGTAAGAATATAA
ACGCGTACGCGGCCGCTCGAGCAGAAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCTGGAT
TACAAGGATGACGACGATAAGGTTTAAACGGCCGGC
  
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- Restriction Sites:** SgfI-MluI
- ACCN:** NM\_001142801
- Insert Size:** 1860 bp
- 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:** This TrueClone is provided through our Custom Cloning Process that includes sub-cloning into OriGene's pCMV6 vector and full sequencing to provide a non-variant match to the expected reference without frameshifts, and is delivered as lyophilized plasmid DNA.
- 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).

|                               |   |
|-------------------------------|---|
| <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_001142801.1</a>  |
| <b>RefSeq Size:</b>           | 5462 bp   |
| <b>RefSeq ORF:</b>            | 1860 bp   |
| <b>Locus ID:</b>              | 346007  |
| <b>UniProt ID:</b>            | <a href="#">Q5T1H1</a>  |
| <b>Cytogenetics:</b>          | 6q12  |
| <b>Protein Families:</b>      | Druggable Genome  |
| <b>MW:</b>                    | 69.9 kDa  |
| <b>Gene Summary:</b>          | <p>The product of this gene contains multiple epidermal growth factor (EGF)-like and LamG domains. The protein is expressed in the photoreceptor layer of the retina, and the gene is mutated in autosomal recessive retinitis pigmentosa. Multiple transcript variants encoding different isoforms have been found for this gene. [provided by RefSeq, Dec 2008]</p> <p>Transcript Variant: This variant (2) lacks several 3' exons but contains an alternate 3' terminal exon, and it thus differs in its 3' coding region and 3' UTR, compared to variant 1. The resulting isoform (2) has a substantially shorter and unique C-terminus, compared to isoform 1. 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> |