

# Product datasheet for SC331709

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OriGene Technologies, Inc.

## RAGE (AGER) (NM 001206932) Human Untagged Clone

### **Product data:**

**Product Type: Expression Plasmids** 

**Product Name:** RAGE (AGER) (NM 001206932) Human Untagged Clone

Tag: Tag Free RAGE Symbol:

Synonyms: RAGE; SCARJ1; sRAGE Vector: pCMV6-Entry (PS100001)

>SC331709 representing NM\_001206932. **Fully Sequenced ORF:** 

Blue=Insert sequence Red=Cloning site Green=Tag(s)

ATGGCAGCCGGAACAGCAGTTGGAGCCTGGGTGCTGGTCCTCAGTCTGTGGGGGGGCAGTAGTAGGTGCT CAAAACATCACAGCCCGGATTGGCGAGCCACTGGTGCTGAAGTGTAAGGGGGCCCCCAAGAAACCACCC CAGCGGCTGGAATGGAAACTGGGAGGAGGCCCCTGGGACAGTGTGGCTCGTGTCCTTCCCAACGGCTCC CTCTTCCTTCCGGCTGTCGGGATCCAGGATGAGGGGATTTTCCGGTGCCAGGCAATGAACAGGAATGGA AAGGAGACCAAGTCCAACTACCGAGTCCGTGTCTACCAGATTCCTGGGAAGCCAGAAATTGTAGATTCT GCCTCTGAACTCACGGCTGGTGTTCCCAATAAGGTGGGGACATGTGTCAGAGGGAAGCTACCCTGCA GGGACTCTTAGCTGGCACTTGGATGGGAAGCCCCTGGTGCCTAATGAGAAGGGAGTATCTGTGAAGGAA CAGACCAGGAGACACCCTGAGACAGGGCTCTTCACACTGCAGTCGGAGCTAATGGTGACCCCAGCCCGG GGAGGAGATCCCCGTCCCACCTTCTCCTGTAGCTTCAGCCCAGGCCTTCCCCGACACCCGGGCCTTGCGC ACAGCCCCATCCAGCCCGTGTCTGGGAGCCTGTGCCTCTGGAGGAGGTCCAATTGGTGGTGGAGCCA GAAGGTGGAGCAGTAGCTCCTGGTGGAACCGTAACCCTGACCTGTGAAGTCCCTGCCCAGCCCTCTCCT CAAATCCACTGGATGAAGGATGGTGTGCCCCTTGCCCCTTCCCCCAGCCCTGTGCTGATCCTCCCTGAG ATAGGGCCTCAGGACCAGGGAACCTACAGCTGTGTGGCCACCCATTCCAGCCACGGGCCCCAGGAAAGC CGTGCTGTCAGCATCAGCATCATCGAACCAGGCGAGGAGGGGCCAACTGCAGGCTCTGTGGGAGGATCA GGGCTGGGAACTCTAGCCCTGGCCCTGGGGATCCTGGGAGGCCTGGGGACAGCCGCCCTGCTCATTGGG 

**Restriction Sites:** Sgfl-Mlul

ACCN: NM 001206932

Insert Size: 1173 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).



## RAGE (AGER) (NM\_001206932) Human Untagged Clone - SC331709

**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:** 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.

**RefSeq:** NM 001206932.1

RefSeq Size:1466 bpRefSeq ORF:1173 bp

Locus ID: 177

UniProt ID: Q15109
Cytogenetics: 6p21.32

**Protein Families:** Druggable Genome, Secreted Protein, Transmembrane

MW: 41.2 kDa

**Gene Summary:** The advanced glycosylation end product (AGE) receptor encoded by this gene is a member of

the immunoglobulin superfamily of cell surface receptors. It is a multiligand receptor, and besides AGE, interacts with other molecules implicated in homeostasis, development, and inflammation, and certain diseases, such as diabetes and Alzheimer's disease. Many alternatively spliced transcript variants encoding different isoforms, as well as non-protein-

coding variants, have been described for this gene (PMID:18089847). [provided by RefSeq, May 2011]

Transcript Variant: This variant (3, also known as RAGE\_v4) uses alternate in-frame acceptor and donor splice sites at two internal coding exons compared to variant 1. This results in a shorter isoform (2) missing two internal protein segments compared to isoform 1. Sequence Note: This Refseq, containing two in-frame translation initiation codons (at nt 8-10 and nt 101-103), is annotated with a CDS starting from the downstream AUG (dAUG) because the AGE receptor encoded by this gene is a known type 1 transmembrane protein requiring signal peptide for its function, and a signal peptide of 22 aa is predicted for the dAUG initiated

protein. Translation initiation from the upstream AUG (uAUG) will add an extra 31 aa to the N-

terminus, and no signal peptide is predicted for the uAUG initiated protein.