

# **Product datasheet for SC329863**

## RAGE (AGER) (NM 001206954) Human Untagged Clone

### **Product data:**

**Product Type:** Expression Plasmids

**Product Name:** RAGE (AGER) (NM\_001206954) Human Untagged Clone

Tag: Tag Free
Symbol: AGER

Synonyms: RAGE; SCARJ1; sRAGE

Vector: pCMV6-Entry (PS100001)

Fully Sequenced ORF: >SC329863 representing NM\_001206954.

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

GTGCAGAACTGA

Restriction Sites: Sgfl-Mlul



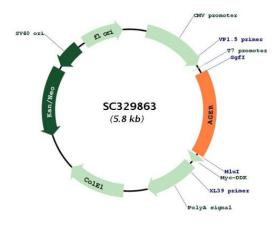
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#### Plasmid Map:



**ACCN:** NM\_001206954

**Insert Size:** 978 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).

**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 001206954.1



#### RAGE (AGER) (NM\_001206954) Human Untagged Clone - SC329863

RefSeq Size:1321 bpRefSeq ORF:978 bpLocus ID:177

UniProt ID: Q15109
Cytogenetics: 6p21.32

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

MW: 35.3 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 (8, also known as RAGE\_v8) lacks two internal coding exons, and uses an alternate donor splice site at another coding exon compared to variant 1. This results in a frame-shift, and a shorter isoform (8) with a distinct C-terminus 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.