

## **Product datasheet for SC330808**

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# hHR23A (RAD23A) (NM\_001270363) Human Untagged Clone

#### **Product data:**

**Product Type:** Expression Plasmids

Product Name: hHR23A (RAD23A) (NM\_001270363) Human Untagged Clone

Tag: Tag Free Symbol: RAD23A

Synonyms: HHR23A; HR23A

**Vector:** pCMV6-Entry (PS100001)

Fully Sequenced ORF: >SC330808 representing NM\_001270363.

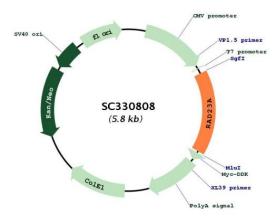
Blue=Insert sequence Red=Cloning site Green=Tag(s)

CTCCTGAGTCAGAACTTTGATGACGAG<mark>TGA</mark>

Restriction Sites: Sgfl-Mlul



#### Plasmid Map:



**ACCN:** NM\_001270363

**Insert Size:** 927 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:** <u>NM 001270363.1</u>

 RefSeq Size:
 1656 bp

 RefSeq ORF:
 927 bp

 Locus ID:
 5886

 UniProt ID:
 P54725

 Cytogenetics:
 19p13.13

**Protein Families:** Druggable Genome

**Protein Pathways:** Nucleotide excision repair

MW: 33.4 kDa

**Gene Summary:** The protein encoded by this gene is one of two human homologs of Saccharomyces

cerevisiae Rad23, a protein involved in nucleotide excision repair. Proteins in this family have a modular domain structure consisting of an ubiquitin-like domain (UbL), ubiquitin-associated domain 1 (UbA1), XPC-binding domain and UbA2. The protein encoded by this gene plays an important role in nucleotide excision repair and also in delivery of polyubiquitinated proteins to the proteasome. Alternative splicing results in multiple transcript variants encoding

multiple isoforms. [provided by RefSeq, Jun 2012]

Transcript Variant: This variant (3) lacks an alternate in-frame exon in the 3' coding region compared to variant 1. It encodes isoform 3 which is shorter than 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.