

## **Product datasheet for SC303461**

## BAG2 (NM\_004282) Human Untagged Clone

## **Product data:**

**Product Type:** Expression Plasmids

**Product Name:** BAG2 (NM\_004282) Human Untagged Clone

Tag: Tag Free Symbol: BAG2

Synonyms: BAG-2; dJ417I1.2

Mammalian Cell

Selection:

None

Vector: pCMV6-XL5

E. coli Selection: Ampicillin (100 ug/mL)

Fully Sequenced ORF: >OriGene sequence for NM\_004282 edited

TAGCAATATTTTAATTATCTATCTAGAGATTTTTTAGATTG

Restriction Sites: Please inquire ACCN: NM\_004282

**Insert Size:** 900 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).



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## BAG2 (NM\_004282) Human Untagged Clone - SC303461

**OTI Annotation:** The ORF of this clone has been fully sequenced and found to be a perfect match to

NM\_004282.2.

**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 004282.2</u>, <u>NP 004273.1</u>

 RefSeq Size:
 1556 bp

 RefSeq ORF:
 636 bp

 Locus ID:
 9532

 UniProt ID:
 095816

 Cytogenetics:
 6p12.1

**Protein Families:** Druggable Genome

**Gene Summary:** BAG proteins compete with Hip for binding to the Hsc70/Hsp70 ATPase domain and promote

substrate release. All the BAG proteins have an approximately 45-amino acid BAG domain near the C terminus but differ markedly in their N-terminal regions. The predicted BAG2 protein contains 211 amino acids. The BAG domains of BAG1, BAG2, and BAG3 interact specifically with the Hsc70 ATPase domain in vitro and in mammalian cells. All 3 proteins bind

with high affinity to the ATPase domain of Hsc70 and inhibit its chaperone activity in a Hiprepressible manner. [provided by RefSeq, Jul 2008]