

Product datasheet for **SC331347**

BAT3 (BAG6) (NM_001199697) Human Untagged Clone

Product data:

Product Type:	Expression Plasmids
Product Name:	BAT3 (BAG6) (NM_001199697) Human Untagged Clone
Tag:	Tag Free
Symbol:	BAT3
Synonyms:	BAG-6; BAT3; D6S52E; G3
Vector:	pCMV6-Entry (PS100001)



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Fully Sequenced ORF: >SC331347 representing NM_001199697.
 Blue=Insert sequence Red=Cloning site Green=Tag(s)

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ATGGAGCCTAATGATAGTACCAGTACCGCTGTGGAGGAGCCTGACAGCTTGGAGGTGTTGGTGAAGACC
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GCCTTGTGATGATCCTTAG
  
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Restriction Sites: SgfI-MluI
ACCN: NM_001199697
Insert Size: 2712 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:	<ol style="list-style-type: none">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_001199697.1
RefSeq Size:	3121 bp
RefSeq ORF:	2712 bp
Locus ID:	7917
UniProt ID:	P46379
Cytogenetics:	6p21.33
Protein Families:	Druggable Genome, Stem cell - Pluripotency
MW:	96.8 kDa
Gene Summary:	<p>This gene was first characterized as part of a cluster of genes located within the human major histocompatibility complex class III region. This gene encodes a nuclear protein that is cleaved by caspase 3 and is implicated in the control of apoptosis. In addition, the protein forms a complex with E1A binding protein p300 and is required for the acetylation of p53 in response to DNA damage. Multiple transcript variants encoding different isoforms have been found for this gene. [provided by RefSeq, Jul 2008]</p> <p>Transcript Variant: This variant (5) differs in the 5' UTR and lacks three alternate in-frame segments compared to variant 1. The resulting isoform (c) has the same N- and C-termini but is shorter compared to isoform a.</p>