



## Human BAG3 peptide (DAG-P1401)

This product is for research use only and is not intended for diagnostic use.

### PRODUCT INFORMATION

<b>Antigen Description</b>	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 protein encoded by this gene contains a WW domain in the N-terminal region and a BAG domain in the C-terminal region. 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 Hip-repressible manner. [provided by RefSeq, Jul 2008]
<b>Purity</b>	70 - 90% by HPLC.
<b>Conjugate</b>	Unconjugated
<b>Sequence Similarities</b>	Contains 1 BAG domain.Contains 2 WW domains.
<b>Format</b>	Liquid
<b>Preservative</b>	None
<b>Storage</b>	Shipped at 4°C. Upon delivery aliquot and store at -20°C or -80°C. Avoid repeated freeze / thaw cycles. Information available upon request.

### GENE INFORMATION

<b>Gene Name</b>	<a href="#">BAG3 BCL2-associated athanogene 3 [ Homo sapiens (human) ]</a>
<b>Official Symbol</b>	BAG3
<b>Synonyms</b>	BAG3; BCL2-associated athanogene 3; BIS; MFM6; BAG-3; CAIR-1; BAG family molecular chaperone regulator 3; docking protein CAIR-1; BCL2-binding athanogene 3; bcl-2-binding protein Bis;

<b>Entrez Gene ID</b>	<a href="#">9531</a>
<b>mRNA Refseq</b>	<a href="#">NM_004281.3</a>
<b>Protein Refseq</b>	<a href="#">NP_004272.2</a>
<b>UniProt ID</b>	O95817
<b>Chromosome Location</b>	10q25.2-q26.2
<b>Pathway</b>	Apoptosis Modulation and Signaling, organism-specific biosystem;
<b>Function</b>	chaperone binding; protein binding; protein complex binding;