

# **Product datasheet for TA326360**

#### OriGene Technologies, Inc.

9620 Medical Center Drive, Ste 200 Rockville, MD 20850, US Phone: +1-888-267-4436 https://www.origene.com techsupport@origene.com EU: info-de@origene.com CN: techsupport@origene.cn

## HSP70-1A (HSPA1A) Mouse Monoclonal Antibody [Clone ID: 3A3]

#### **Product data:**

**Product Type:** Primary Antibodies

Clone Name: 3A3
Applications: WB

Recommended Dilution: ICC: 1:500, WB: 1:5000; IP: 1-2ug

**Reactivity:** Human, Mouse, Rat, Amphibian, Chicken, Fish, Saccharomyces cerevisiae, Fruit fly

Host: Mouse Isotype: IgG1

Clonality: Monoclonal

Immunogen: Human Recombinant Hsp70 overexpressed in E.coli

**Formulation:** PBS pH7.2, 50% glycerol, 0.09% sodium azide

**Concentration:** lot specific

Purification: Protein G Purified

Conjugation: Unconjugated

Storage: Store at -20°C as received.

**Stability:** Stable for 12 months from date of receipt.

Gene Name: heat shock protein family A (Hsp70) member 1A

Database Link: NP 005336

Entrez Gene 24472 RatEntrez Gene 193740 MouseEntrez Gene 3303 Human

P0DMV8





Background:

Hsp70 genes encode abundant heat-inducible 70-kDa hsps (hsp70s). In most eukaryotes hsp70 genes exist as part of a multigene family. They are found in most cellular compartments of eukaryotes including nuclei, mitochondria, chloroplasts, the endoplasmic reticulum and the cytosol, as well as in bacteria. The genes show a high degree of conservation, having at least 50% identity. The N-terminal two thirds of hsp70s are more conserved than the C-terminal third. Hsp70 binds ATP with high affinity and possesses a weak ATPase activity which can be stimulated by binding to unfolded proteins and synthetic peptides. When hsc70 (constitutively expressed) present in mammalian cells was truncated, ATP binding activity was found to reside in an N-terminal fragment of 44 kDa which lacked peptide binding capacity. Polypeptide binding ability therefore resided within the C-terminal half. The structure of this ATP binding domain displays multiple features of nucleotide binding proteins. All hsp70s, regardless of location, bind proteins, particularly unfolded ones. The molecular chaperones of the hsp70 family recognize and bind to nascent polypeptide chains as well as partially folded intermediates of proteins preventing their aggregation and misfolding. The binding of ATP triggers a critical conformational change leading to the release of the bound substrate protein. The universal ability of hsp70s to undergo cycles of binding to and release from hydrophobic stretches of partially unfolded proteins determines their role in a great variety of vital intracellular functions such as protein synthesis, protein folding and oligomerization and protein transport.

**Synonyms:** HEL-S-103; HSP70-1; HSP70-1A; HSP70.1; HSP70I; HSP72; HSPA1

**Note:** Detects several members of the heat shock protein 70kDa gene family including Hsp70,

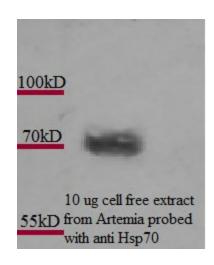
Hsc70, p75 and following heat shock, Hsp72 from yeast, Drosophila, fish, mouse, avian,

amphibian and human samples.

**Protein Pathways:** Antigen processing and presentation, Endocytosis, MAPK signaling pathway, Prion diseases,

Spliceosome

## **Product images:**



10ug cell free extract from Artemia probed with anti-Hsp70