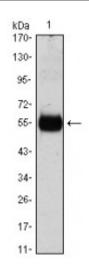


Anti-p27 antibody



Description Mouse monoclonal to p27.

Model STJ98298

Host Mouse

Reactivity Human

Applications ELISA, WB

Immunogen Purified recombinant fragment of human p27 expressed in E. Coli.

Gene ID <u>1027</u>

Gene Symbol CDKN1B

Dilution range WB 1:500-1:2000ELISA 1:10000

Specificity p27 Monoclonal Antibody detects endogenous levels of p27 protein.

Tissue Specificity Expressed in all tissues tested. Highest levels in skeletal muscle, lowest in

liver and kidney.

Purification Affinity purification

Clone ID 3D8

Note For Research Use Only (RUO).

Protein Name Cyclin-dependent kinase inhibitor 1B Cyclin-dependent kinase inhibitor p27

p27Kip1

Clonality Monoclonal

Conjugation Unconjugated

IgG1 **Isotype**

Formulation Ascitic fluid containing 0.03% sodium azide.

Store at -20°C, and avoid repeat freeze-thaw cycles. **Storage Instruction**

HGNC:1785OMIM:600778 **Database Links**

Cyclin-dependent kinase inhibitor 1B Cyclin-dependent kinase inhibitor p27 **Alternative Names**

p27Kip1

Function Important regulator of cell cycle progression. Involved in G1 arrest. Potent

> inhibitor of cyclin E- and cyclin A-CDK2 complexes. Forms a complex with cyclin type D-CDK4 complexes and is involved in the assembly, stability, and modulation of CCND1-CDK4 complex activation. Acts either as an inhibitor

or an activator of cyclin type D-CDK4 complexes depending on its

phosphorylation state and/or stoichometry.

Sequence and Domain Family A peptide sequence containing only AA 28-79 retains substantial Kip1 cyclin

A/CDK2 inhibitory activity.

Cellular Localization Nucleus. Cytoplasm. Endosome. Nuclear and cytoplasmic in quiescent cells.

AKT- or RSK-mediated phosphorylation on Thr-198, binds 14-3-3,

translocates to the cytoplasm and promotes cell cycle progression. Mitogenactivated UHMK1 phosphorylation on Ser-10 also results in translocation to

the cytoplasm and cell cycle progression. Phosphorylation on Ser-10 facilitates nuclear export. Translocates to the nucleus on phosphorylation of Tyr-88 and Tyr-89. Colocalizes at the endosome with SNX6. this leads to

lysosomal degradation.

Post-translational **Modifications**

Phosphorylated; phosphorylation occurs on serine, threonine and tyrosine residues. Phosphorylation on Ser-10 is the major site of phosphorylation in resting cells, takes place at the G(0)-G(1) phase and leads to protein stability. Phosphorylation on other sites is greatly enhanced by mitogens, growth factors, cMYC and in certain cancer cell lines. The phosphorylated form found in the cytoplasm is inactivate. Phosphorylation on Thr-198 is required for interaction with 14-3-3 proteins. Phosphorylation on Thr-187, by CDK1 and CDK2 leads to protein ubiquitination and proteasomal degradation. Tyrosine phosphorylation promotes this process. Phosphorylation by PKB/AKT1 can be suppressed by LY294002, an inhibitor of the catalytic subunit of PI3K. Phosphorylation on Tyr-88 and Tyr-89 has no effect on binding CDK2, but is required for binding CDK4. Dephosphorylated on tyrosine residues by G-CSF. Ubiquitinated; in the cytoplasm by the KPC complex (composed of RNF123/KPC1 and UBAC1/KPC2) and, in the nucleus, by SCF(SKP2). The latter requires prior phosphorylation on Thr-187. Ubiquitinated; by a TRIM21-containing SCF(SKP2)-like complex; leads to its degradation. Subject to degradation in the lysosome. Interaction with SNX6 promotes lysosomal degradation.