

## Cleaved-PARP-1 (G215) Polyclonal Antibody

Catalog No: YC0073

**Reactivity:** Human; Mouse; Rat

**Applications:** WB;ELISA

Target: PARP

Fields: >>Base excision repair;>>NF-kappa B signaling

pathway;>>Apoptosis;>>Necroptosis;>>Diabetic cardiomyopathy

Gene Name: PARP1

**Protein Name:** Poly [ADP-ribose] polymerase 1

P09874

P11103

Human Gene Id: 142

**Human Swiss Prot** 

No:

**Mouse Swiss Prot** 

No:

**Immunogen:** The antiserum was produced against synthesized peptide derived from human

PARP. AA range:196-245

**Specificity:** Cleaved-PARP-1 (G215) Polyclonal Antibody detects endogenous levels of

fragment of activated PARP-1 protein resulting from cleavage adjacent to G215.

**Formulation :** Liquid in PBS containing 50% glycerol, 0.5% BSA and 0.02% sodium azide.

Source: Polyclonal, Rabbit, IgG

**Dilution:** WB 1:500 - 1:2000. ELISA: 1:5000. Not yet tested in other applications.

**Purification:** The antibody was affinity-purified from rabbit antiserum by affinity-

chromatography using epitope-specific immunogen.

**Concentration:** 1 mg/ml

1/4



**Storage Stability:** -15°C to -25°C/1 year(Do not lower than -25°C)

Observed Band: 89kD

**Cell Pathway:** Base excision repair;

**Background:** This gene encodes a chromatin-associated enzyme, poly(ADP-

ribosyl)transferase, which modifies various nuclear proteins by poly(ADP-ribosyl)ation. The modification is dependent on DNA and is involved in the regulation of various important cellular processes such as differentiation,

proliferation, and tumor transformation and also in the regulation of the molecular events involved in the recovery of cell from DNA damage. In addition, this enzyme

may be the site of mutation in Fanconi anemia, and may participate in the pathophysiology of type I diabetes. [provided by RefSeq, Jul 2008],

**Function :** catalytic activity:NAD(+) + (ADP-D-ribosyl)(n)-acceptor = nicotinamide + (ADP-

D-ribosyl)(n+1)-acceptor.,function:Involved in the base excision repair (BER) pathway, by catalyzing the poly(ADP-ribosyl)ation of a limited number of acceptor

proteins involved in chromatin architecture and in DNA metabolism. This modification follows DNA damages and appears as an obligatory step in a

detection/signaling pathway leading to the reparation of DNA strand

breaks.,miscellaneous:The ADP-D-ribosyl group of NAD(+) is transferred to an acceptor carboxyl group on a histone or the enzyme itself, and further ADP-ribosyl groups are transferred to the 2'-position of the terminal adenosine moiety, building up a polymer with an average chain length of 20-30 units.,PTM:Phosphorylated

by PRKDC. Phosphorylated upon DNA damage, probably by ATM or ATR.,PTM:Poly-ADP-ribosylated by PARP2.,similarity:Contains 1 BRCT

Subcellular Location:

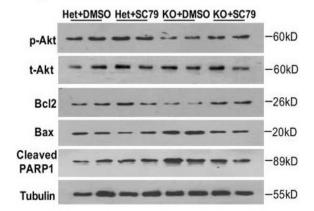
Nucleus . Nucleus, nucleolus . Chromosome . Localizes to sites of DNA damage.

**Expression:** 

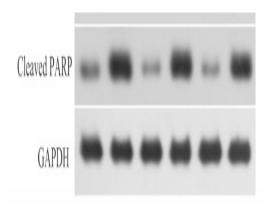
Brain, Colon carcinoma, Fibroblast, Lung, Ovarian carcinoma, Skin,

## **Products Images**

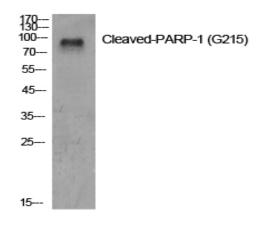




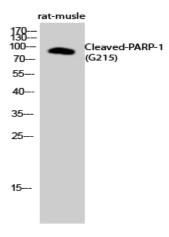
Wang, Bin, et al. "Loss of Tctn3 causes neuronal apoptosis and neural tube defects in mice." Cell death & disease 9.5 (2018): 520



Mao, Dongwei, et al. "RNAi-mediated knockdown of the CLN3 gene inhibits proliferation and promotes apoptosis in drug-resistant ovarian cancer cells." Molecular medicine reports12.5 (2015): 6635-6641.



Western Blot analysis of various cells using Cleaved-PARP-1 (G215) Polyclonal Antibody diluted at 1:500



Western Blot analysis of rat-musle cells using Cleaved-PARP-1 (G215) Polyclonal Antibody diluted at 1:500