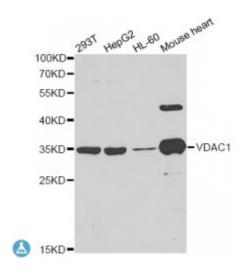


Anti-VDAC1 Antibody



Description This gene encodes a voltage-dependent anion channel protein that is a

major component of the outer mitochondrial membrane. The encoded protein facilitates the exchange of metabolites and ions across the outer mitochondrial membrane and may regulate mitochondrial functions. This protein also forms channels in the plasma membrane and may be involved in transmembrane electron transport. Alternate splicing results in multiple transcript variants. Multiple pseudogenes of this gene are found on

chromosomes 1, 2 3, 6, 9, 12, X and Y.

Model STJ115597

Host Rabbit

Reactivity Human, Mouse

Applications IF, IHC, WB

Immunogen Recombinant fusion protein containing a sequence corresponding to amino

acids 1-283 of human VDAC1 (NP_003365.1).

Gene ID <u>7416</u>

Gene Symbol VDAC1

Dilution range WB 1:500 - 1:2000

IHC 1:50 - 1:200 IF 1:50 - 1:200

Tissue Specificity Heart, liver and skeletal muscle

Purification Affinity purification

Note For Research Use Only (RUO).

Protein Name Voltage-dependent anion-selective channel protein 1 VDAC-1 hVDAC1

Outer mitochondrial membrane protein porin 1 Plasmalemmal porin Porin

31HL Porin 31HM

Molecular Weight 30.773 kDa

Clonality Polyclonal

Conjugation Unconjugated

Isotype IgG

Formulation PBS with 0.02% sodium azide, 50% glycerol, pH7.3.

Storage Instruction Store at -20C. Avoid freeze / thaw cycles.

Database Links HGNC:12669OMIM:604492Reactome:R-HSA-1268020

Alternative Names Voltage-dependent anion-selective channel protein 1 VDAC-1 hVDAC1

Outer mitochondrial membrane protein porin 1 Plasmalemmal porin Porin

31HL Porin 31HM

Function Forms a channel through the mitochondrial outer membrane and also the

plasma membrane, The channel at the outer mitochondrial membrane allows

diffusion of small hydrophilic molecules

Cellular Localization Mitochondrion outer membrane

Post-translational Phosphorylation at Ser-193 by NEK1 promotes the open conformational state

Modifications preventing excessive mitochondrial membrane permeability and subsequent

apoptotic cell death after injury, Phosphorylation by the AKT-GSK3B axis stabilizes the protein probably by preventing ubiquitin-mediated proteasomal

degradation,

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