

## c-Abl rabbit pAb

## Cat No.:ES1819

For research use only

## Overview

Product Name	c-Abl rabbit pAb
Host species	Rabbit
Applications	WB;IHC;IF;ELISA
Species Cross-Reactivity	Human;Mouse;Rat
Recommended dilutions	Western Blot: 1/500 - 1/2000.
	Immunohistochemistry: 1/100 - 1/300.
	Immunofluorescence: 1/200 - 1/1000. ELISA:
	1/20000. Not yet tested in other applications.
Immunogen	The antiserum was produced against synthesized
	peptide derived from human ABL1. AA
	range:861-910
Specificity	c-Abl Polyclonal Antibody detects endogenous levels
	of c-Abl protein.
Formulation	Liquid in PBS containing 50% glycerol, 0.5% BSA and
	0.02% sodium azide.
Storage	Store at -20 $^\circ\!{ m C}$ . Avoid repeated freeze-thaw cycles.
Protein Name	Tyrosine-protein kinase ABL1
Gene Name	ABL1
<b>Cellular localization</b>	Cytoplasm, cytoskeleton. Nucleus. Mitochondrion .
	Shuttles between the nucleus and cytoplasm
	depending on environmental signals. Sequestered
	into the cytoplasm through interaction with 14-3-3
	proteins. Localizes to mitochondria in response to
	oxidative stress (By similarity); [Isoform IB]:
	Nucleus membrane; Lipid-anchor. The myristoylated
	c-ABL protein is reported to be nuclear.
Purification	The antibody was affinity-purified from rabbit
	antiserum by affinity-chromatography using
	epitope-specific immunogen.
Clonality	Polyclonal
Concentration	1 mg/ml
Observed band	125(200kd BCR-ABL complex)
Human Gene ID	25
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Human Swiss-Prot Number Alternative Names

Background

P00519

ABL1; ABL; JTK7; Tyrosine-protein kinase ABL1; Abelson murine leukemia viral oncogene homolog 1; Abelson tyrosine-protein kinase 1; Proto-oncogene c-Abl; p150

This gene is a protooncogene that encodes a protein tyrosine kinase involved in a variety of cellular processes, including cell division, adhesion, differentiation, and response to stress. The activity of the protein is negatively regulated by its SH3 domain, whereby deletion of the region encoding this domain results in an oncogene. The ubiquitously expressed protein has DNA-binding activity that is regulated by CDC2-mediated phosphorylation, suggesting a cell cycle function. This gene has been found fused to a variety of translocation partner genes in various leukemias, most notably the t(9;22) translocation that results in a fusion with the 5' end of the breakpoint cluster region gene (BCR; MIM:151410). Alternative splicing of this gene results in two transcript variants, which contain alternative first exons that are spliced to the remaining common exons. [pr



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