

BRCA1 rabbit pAb

Cat No.: ES1788

For research use only

Overview

Product Name BRCA1 rabbit pAb

Host species Rabbit

Applications WB;IHC;IF;ELISA Species Cross-Reactivity Human;Rat

Recommended dilutions Western Blot: 1/500 - 1/2000.

Immunohistochemistry: 1/100 - 1/300. Immunofluorescence: 1/200 - 1/1000. ELISA:

1/5000. Not yet tested in other applications.
The antiserum was produced against synthesized

Immunogen The antiserum was produced against synthesized

peptide derived from human BRCA1. AA

range:1391-1440

Specificity BRCA1 Polyclonal Antibody detects endogenous

levels of BRCA1 protein.

Formulation Liquid in PBS containing 50% glycerol, 0.5% BSA and

0.02% sodium azide.

Storage Store at -20° C. Avoid repeated freeze-thaw cycles.

Protein Name Breast cancer type 1 susceptibility protein

Gene Name BRCA1

Cellular localization Nucleus . Chromosome . Cytoplasm . Localizes at

sites of DNA damage at double-strand breaks (DSBs); recruitment to DNA damage sites is mediated by

ABRAXAS1 and the BRCA1-A complex

(PubMed:26778126). Translocated to the cytoplasm during UV-induced apoptosis (PubMed:20160719). .; [Isoform 3]: Cytoplasm.; [Isoform 5]: Cytoplasm.

The antibody was affinity-purified from rabbit antiserum by affinity-chromatography using

epitope-specific immunogen.

Clonality Polyclonal Concentration 1 mg/ml

Observed band

Purification

Human Gene ID 672 Human Swiss-Prot Number P38398



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Alternative Names

Background

BRCA1; RNF53; Breast cancer type 1 susceptibility protein; RING finger protein 53 This gene encodes a nuclear phosphoprotein that plays a role in maintaining genomic stability, and it also acts as a tumor suppressor. The encoded protein combines with other tumor suppressors, DNA damage sensors, and signal transducers to form a large multi-subunit protein complex known as the BRCA1-associated genome surveillance complex (BASC). This gene product associates with RNA polymerase II, and through the C-terminal domain, also interacts with histone deacetylase complexes. This protein thus plays a role in transcription, DNA repair of double-stranded breaks, and recombination. Mutations in this gene are responsible for approximately 40% of inherited breast cancers and more than 80% of inherited breast and ovarian cancers. Alternative splicing plays a role in modulating the subcellular localization and physiological function of this gene. Many alternatively spliced transcript varian

