

A Reliable Research Partner in Life Science and Medicine

HUS1 Polyclonal Antibody

catalog number: E-AB-64873

Note: Centrifuge before opening to ensure complete recovery of vial contents.

Description

Reactivity Human; Mouse; Rat

Immunogen Recombinant fusion protein of human HUS1 (NP 004498.1).

Host Rabbit
Isotype IgG

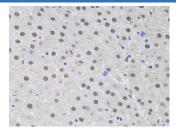
Purification Affinity purification

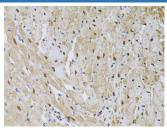
Buffer Phosphate buffered solution, pH 7.4, containing 0.05% stabilizer and 50% glycerol.

Applications Recommended Dilution

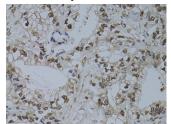
IHC 1:50-1:100

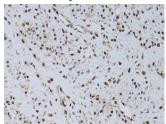
Data





Immunohistochemistry of paraffin-embedded Rat liver using Immunohistochemistry of paraffin-embedded Rat heart using HUS1 Polyclonal Antibody at dilution of 1:100 (40x lens). HUS1 Polyclonal Antibody at dilution of 1:100 (40x lens).





Immunohistochemistry of paraffin-embedded Human gastric Immunohistochemistry of paraffin-embedded Human uterine cancer using HUS1 Polyclonal Antibody at dilution of 1:100 cancer using HUS1 Polyclonal Antibody at dilution of 1:100 (40x lens).

Preparation & Storage

Storage Store at -20°C Valid for 12 months. Avoid freeze / thaw cycles.

Shipping The product is shipped with ice pack, upon receipt, store it immediately at the

temperature recommended.

Background

For Research Use Only

Fax: 1-832-243-6017

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The protein encoded by this gene is a component of an evolutionarily conserved, genotoxin-activated checkpoint complex that is involved in the cell cycle arrest in response to DNA damage. This protein forms a heterotrimeric complex with checkpoint proteins RAD9 and RAD1. In response to DNA damage, the trimeric complex interacts with another protein complex consisting of checkpoint protein RAD17 and four small subunits of the replication factor C (RFC), which loads the combined complex onto the chromatin. The DNA damage induced chromatin binding has been shown to depend on the activation of the checkpoint kinase ATM, and is thought to be an early checkpoint signaling event. Alternative splicing results in multiple transcript variants.

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