# **QARS Polyclonal Antibody**

Catalog Number: E-AB-61432



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

### **Description**

Reactivity Human, Mouse

**Immunogen** Recombinant fusion protein of human QARS (NP\_005042.1).

Host Rabbit
Isotype IgG

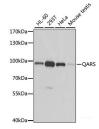
Purification Affinity purification
Conjugation Unconjugated

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

## **Applications** Recommended Dilution

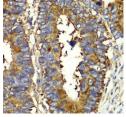
WB 1:500-1:2000 IHC 1:50-1:200 IF 1:50-1:100

## **Data**

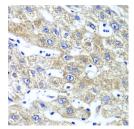


Western blot analysis of extracts of various cell lines using QARS Polyclonal Antibody at dilution of 1:1000.

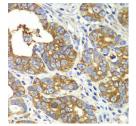
### Observed Mw:88kDa Calculated Mw:86kDa/87kDa



Immunohistochemistry of paraffin-embedded Human colon carcinoma using QARS Polyclonal Antibody at dilution of 1:100 (40x lens).



Immunohistochemistry of paraffin-embedded Human liver damage using QARS Polyclonal Antibody at dilution of 1:100 (40x lens).



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

#### For Research Use Only

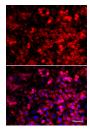
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Immunofluorescence analysis of U2OS cells using QARS Polyclonal Antibody

## **Preparation & Storage**

Storage

Store at -20°C. Avoid freeze / thaw cycles.

### **Background**

Aminoacyl-tRNA synthetases catalyze the aminoacylation of tRNA by their cognate amino acid. Because of their central role in linking amino acids with nucleotide triplets contained in tRNAs, aminoacyl-tRNA synthetases are thought to be among the first proteins that appeared in evolution. In metazoans, 9 aminoacyl-tRNA synthetases specific for glutamine (gln), glutamic acid (glu), and 7 other amino acids are associated within a multienzyme complex. Although present in eukaryotes, glutaminyl-tRNA synthetase (QARS) is absent from many prokaryotes, mitochondria, and chloroplasts, in which Gln-tRNA(Gln) is formed by transamidation of the misacylated Glu-tRNA(Gln). Glutaminyl-tRNA synthetase belongs to the class-I aminoacyl-tRNA synthetase family. Alternative splicing results in multiple transcript variants.

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