

Anti-DCPS Antibody (9T672)

Product Details

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| Ig Type: | IgG2b/Kappa |
| Reactivity: | Human (predicted:Mouse) |
| Clone: | 9T672 |
| Purification: | Protein G purified |

Applications

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|---------------------------------|---|
| Verified Activity: | 1. Tissue: Human breast cancer |
| | Section type: Formalin fixed & Paraffin-embedded section |
| | Retrieval method: High temperature and high pressure |
| | Retrieval buffer: Tris/EDTA buffer, pH 9.0 Primary ab dilution: 1:100 |
| | Primary ab incubation condition: 1 hour at room temperature |
| | Secondary ab: SP Kit (Mouse) |
| | Counter stain: Hematoxylin (Blue) |
| | Comment: Color brown is the positive signal for TMAB-05010 |
| | 2. Blocking buffer: 5% NFDm/TBST |
| | Primary ab dilution: 1:1000 |
| | Primary ab incubation condition: 4°C overnight |
| | Secondary ab: Goat Anti-Mouse IgG H&L (HRP) |
| | Lysate: HeLa, HepG2, A549, EL4.IL-2 |
| Protein loading quantity: 20 µg | |
| Exposure time: 60 s | |
| Predicted MW: 40 kDa | |
| Observed MW: 35 kDa | |
| Application: | WB,IHC-P,IHC-Fr,IF |
| Recommended | WB: 1:500-1000; IHC-P: 1:100-500; IHC-Fr: 1:100-500; IF: 1:100-500 |

Properties

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|----------------------|---|
| Stability & Storage: | Store at 2°C-8°C for 1 month. Store at -20°C or -80°C for 12 months. Avoid repeated freeze-thaw cycles. |
| Shipping: | Shipping with blue ice. |

Antigen Details

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| Gene ID: | 28960 |
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Research Background

Eukaryotic cells primarily utilize exoribonucleases and decapping enzymes to degrade their mRNA. DcpS is a scavenger pyrophosphatase that hydrolyzes the residual cap structure following 3' to 5' decay of an mRNA. Following mRNA degradation DcpS releases N-7 methyl guanosine monophosphate and 5'-diphosphate terminated cap or mRNA products. The central histidine within the DcpS HIT motif is critical for decapping activity and defines the HIT motif as a new mRNA decapping domain, making DcpS the first member of the HIT family of proteins with a defined biological function. HIT proteins are homodimeric and contain two conserved 100-amino-acid HIT fold domains with independent active sites that are each sufficient to bind and hydrolyze cognate substrates.

Inhibitor · Natural Compounds · Compound Libraries · Recombinant Proteins

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