

Catalogue #MCA-40B7: Monoclonal Antibody 40B7 to Yeast Plasma Membrane ATPase Pma1p

The Immunogen: Pma1p is an abundant multidomain protein of yeast which is localized in the plasma membrane. It functions as a major regulator of cytoplasmic pH, by pumping proteins out of the cell. It is and part of the P2 subgroup of cation-transporting ATPases. Since Pma1p is a major plasma membrane protein, antibodies which bind to it such as MCA-40B7 are useful makers of yeast plasma membranes.

Diagram of Domain Structure: Generated from sequence of yeast Pma1p sequence [CAA96708](#) (with the SMART program from [EMBL](#) in Heidelberg). From N-terminus (left in diagram) to C-terminus the domains are as follows. Following two regions of low complexity (purple) is PFAM: Cation_ATPase_N which refers to the [Cation ATPase N-terminal domain](#). PFAM: E1_E2 Atpase refers to the [E1 E2 ATPase domain](#). PFAM Hydrolase refers to the [Hydrolase domain](#). Finally there are 6 predicted transmembrane segments (Blue). Scale is number of amino acids.



The Antibody: Mouse monoclonal antibody, in the form of sterile-filtered cell culture fluid from an Integra CL-350 biochamber plus sodium azide. The solution contains 10% bovine serum, and about 0.5 mg/ml antibody. The IgG isotype is not known.

Suggestions for use: For western blots of yeast protein samples (cell lysates), try MCA-40B7 diluted 1/10,000, followed by chemiluminescent detection (ECL). For non-ECL western detection methods, try MCA-40B7 diluted 1/1,000 to 1/5,000. For immunofluorescence on yeast cells, try MCA-40B7 diluted 1/500 to 1/5,000. MCA-40B7 reacts with Pma1p based on the size of the band detected on western blots and the immunofluorescence localization pattern. MCA-40B7 also detects Pma1p-containing fusion proteins on western blots. Antibody preparation contains 10mM sodium azide preservative (Link to <http://www.encorbio.com/MSDS/azide.htm> for Material Safety Data Sheet).. Avoid repeated freezing and thawing, store at 4°C or -20°C.

References:

1. Smardon AM *et al.* The RAVE complex is an isoform-specific V-ATPase assembly factor in yeast. *Mol Biol Cell* 25:356-67 (2014). **ICC/IF; *Saccharomyces cerevisiae***. [PubMed: 24307682](#)
2. Epstein T *et al.* Separation of metabolic supply and demand: aerobic glycolysis as a normal physiological response to fluctuating energetic demands in the membrane. *Cancer Metab* 2:7 (2014). **WB**. [PubMed: 24982758](#)
3. Digne SA & Kozminski KG Secretory Vesicles Deliver Cdc42p to Sites of Polarized Growth in *S. cerevisiae*. *PLoS One* 9:e99494 (2014). **WB; *Saccharomyces cerevisiae***. [PubMed: 24945395](#)
4. Thayer NH *et al.* Identification of long-lived proteins retained in cells undergoing repeated asymmetric divisions. *Proc Natl Acad Sci U S A* 111:14019-26 (2014). [PubMed: 25228775](#)
5. Henderson KA *et al.* Mother-daughter asymmetry of pH underlies aging and rejuvenation in yeast. *Elife* 3:e03504 (2014). [PubMed: 25190112](#)

6. Wade F *et al.* Relationship between homo-oligomerization of a mammalian olfactory receptor and its activation state demonstrated by bioluminescence resonance energy transfer. *J Biol Chem* 286:15252-9 (2011). **WB**. [PubMed: 21454689](#)
7. Gulshan K *et al.* Compartment-specific synthesis of phosphatidylethanolamine is required for normal heavy metal resistance. *Mol Biol Cell* 21:443-55 (2010). **WB; Saccharomyces cerevisiae**. [PubMed: 20016005](#)
8. Soper JH *et al.* {alpha}-Synuclein-induced Aggregation of Cytoplasmic Vesicles in *Saccharomyces cerevisiae*. *Mol Biol Cell* 19:1093-103 (2008). **WB; Saccharomyces cerevisiae**. [PubMed: 18172022](#)
9. Martínez-Muñoz GA & Kane P Vacuolar and Plasma Membrane Proton Pumps Collaborate to Achieve Cytosolic pH Homeostasis in Yeast. *J Biol Chem* 283:20309-19 (2008). **ICC/IF; Saccharomyces cerevisiae**. [PubMed: 18502746](#)
10. Clark SW & Rose MD Arp10p is a pointed-end-associated component of yeast dynein. *Mol Biol Cell* 17:738-48 (2006). [PubMed: 16291862](#)
11. Zhang J *et al.* Characterization of the transport mechanism and permeant binding profile of the uridine permease Fui1p of *Saccharomyces cerevisiae*. *J Biol Chem* 281:28210-21 (2006). **WB; Saccharomyces cerevisiae**. [PubMed: 16854981](#)
12. Shibagaki N & Grossman AR The role of the STAS domain in the function and biogenesis of a sulfate transporter as probed by random mutagenesis. *J Biol Chem* 281:22964-73 (2006). [PubMed: 16754669](#)

Limitations: This product is for research use only and is not approved for use in humans or in clinical diagnosis.

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