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## **Datasheet**

## MME monoclonal antibody, clone HI10a (PerCP)

Catalog Number: MAB13780

Regulatory Status: For research use only (RUO)

Product Description: Mouse monoclonal antibody

raised against human MME.

Clone Name: HI10a

Immunogen: Acute CALLA leukemia blast cells.

Host: Mouse

Theoretical MW (kDa): 100

Reactivity: Human

**Applications:** Flow Cyt

(See our web site product page for detailed applications

information)

Protocols: See our web site at

http://www.abnova.com/support/protocols.asp or product

page for detailed protocols

Form: Liquid

Conjugation: PerCP

Purification: Protein A/G purification

**Purity: >90%** 

Isotype: IgG1, kappa

Recommend Usage: Flow Cytometry (20 uL/10<sup>6</sup> cells)

The optimal working dilution should be determined by

the end user.

Storage Buffer: In PBS, pH 7.4 (protein stabilizer,

0.09% sodium azide).

Storage Instruction: Store in the dark at 4°C. Avoid

prolonged exposure to light.

Entrez GenelD: 4311

Gene Symbol: MME

Gene Alias: CALLA, CD10, DKFZp686O16152,

MGC126681, MGC126707, NEP

Gene Summary: This gene encodes a common acute lymphocytic leukemia antigen that is an important cell surface marker in the diagnosis of human acute lymphocytic leukemia (ALL). This protein is present on leukemic cells of pre-B phenotype, which represent 85% of cases of ALL. This protein is not restricted to leukemic cells, however, and is found on a variety of normal tissues. It is a glycoprotein that is particularly abundant in kidney, where it is present on the brush border of proximal tubules and on glomerular epithelium. The protein is a neutral endopeptidase that cleaves peptides at the amino side of hydrophobic residues and inactivates several peptide hormones including glucagon, enkephalins, substance P, neurotensin, oxytocin, and bradykinin. This gene, which encodes a 100-kD type II transmembrane glycoprotein, exists in a single copy of greater than 45 kb. The 5' untranslated region of this gene is alternatively spliced, resulting in four separate mRNA transcripts. The coding region is not affected by alternative splicing. [provided by RefSeq]