abcam

Product datasheet

Anti-Liver Arginase antibody [ARG1/1125] ab215843

画像数3

製品の概要

製品名 Anti-Liver Arginase antibody [ARG1/1125]

製品の詳細 Mouse monoclonal [ARG1/1125] to Liver Arginase

由来種 Mouse

アプリケーション 適用あり: Protein Array, IHC-P, WB

種交差性 交差種: Human

免疫原 Recombinant fragment within Human Liver Arginase aa 1-150. The exact sequence is proprietary.

87 amino acids long. NCBI Accession No. 440934.

Database link: P05089

ポジティブ・コントロール Human hepatocellular carcinoma tissue. 293T cells. Huma liver lysate. Recombinant Human Liver

Arginase protein fragment.

特記事項

The Life Science industry has been in the grips of a reproducibility crisis for a number of years.

Abcam is leading the way in addressing this with our range of recombinant monoclonal antibodies and knockout edited cell lines for gold-standard validation. Please check that this product meets

your needs before purchasing.

If you have any questions, special requirements or concerns, please send us an inquiry and/or contact our Support team ahead of purchase. Recommended alternatives for this product can be

found below, along with publications, customer reviews and Q&As

製品の特性

製品の状態 Liquid

保存方法 Shipped at 4°C. Store at +4°C short term (1-2 weeks). Upon delivery aliquot. Store at -20°C long

term. Avoid freeze / thaw cycle.

バッファー pH: 7.2

Preservative: 0.05% Sodium azide Constituents: 0.05% BSA, 99% PBS

精製度 Protein A/G purified

ポリモノ モノクローナル **クローン名** ARG1/1125

アイソタイプ lgG3

1

Cytoplasm.

アプリケーション

The Abpromise guarantee <u>Abpromise保証は、</u>次のテスト済みアプリケーションにおけるab215843の使用に適用されます アプリケーションノートには、推奨の開始希釈率がありますが、適切な希釈率につきましてはご検討ください。

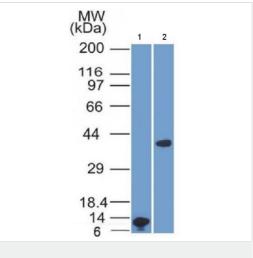
アプリケーション	Abreviews	特記事項
Protein Array		Use at an assay dependent concentration.
IHC-P		Use a concentration of 2 - 4 μ g/ml. Primary incubation for 30 minutes at room temperature. Staining of formalin-fixed tissues requires heating tissue sections in 10mM Tris with 1mM EDTA, pH 9.0, for 45 min at 95°C followed by cooling at room temperature for 20 minutes.
WB		Use a concentration of 1 - 2 μg/ml. Predicted molecular weight: 35 kDa.

ターゲット情報

パスウェイ	Nitrogen metabolism; urea cycle; L-ornithine and urea from L-arginine: step 1/1.
関連疾患	Defects in ARG1 are the cause of argininemia (ARGIN) [MIM:207800]; also known as hyperargininemia. Argininemia is a rare autosomal recessive disorder of the urea cycle. Arginine is elevated in the blood and cerebrospinal fluid, and periodic hyperammonemia occurs. Clinical manifestations include developmental delay, seizures, mental retardation, hypotonia, ataxia, progressive spastic quadriplegia.
配列類似性	Belongs to the arginase family.

画像

細胞内局在



Western blot - Anti-Liver Arginase antibody [ARG1/1125] (ab215843)

All lanes : Anti-Liver Arginase antibody [ARG1/1125] (ab215843) at 1 μg/ml

Lane 1 : recombinant Human Liver Arginase protein fragment

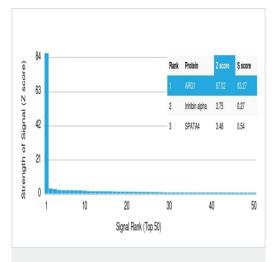
Lane 2: Human liver lysate

Predicted band size: 35 kDa



Immunohistochemistry (Formalin/PFA-fixed paraffinembedded sections) - Anti-Liver Arginase antibody
[ARG1/1125] (ab215843)

Immunohistochemical analysis of formalix-fixed paraffin-embedded Human hepatocellular carcinoma tissue, labeling Liver Arginase using ab215843 at 4 μ g/mL.



Protein Array - Anti-Liver Arginase antibody [ARG1/1125] (ab215843)

ab215843 was tested in protein array against over 19000 different full-length human proteins.

Z- and S- Score: The Z-score represents the strength of a signal that a monoclonal antibody (MAb) (in combination with a fluorescently-tagged anti-lgG secondary antibody) produces when binding to a particular protein on the HuProtTM array. Z-scores are described in units of standard deviations (SD's) above the mean value of all signals generated on that array. If targets on HuProtTM are arranged in descending order of the Z-score, the S-score is the difference (also in units of SD's) between the Z-score. S-score therefore represents the relative target specificity of a MAb to its intended target.

A MAb is specific to its intended target if the MAb has an S-score of at least 2.5. For example, if a MAb binds to protein X with a Z-

score of 43 and to protein Y with a Z-score of 14, then the S-score for the binding of that MAb to protein X is equal to 29.

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