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Product datasheet

Recombinant Human KAP1 protein ab131899

1 References 画像数 1

製品の詳細

製品名 Recombinant Human KAP1 protein

発現系Wheat germアクセッション番号Q13263

タンパク質長 Full length protein

Animal free No

由来 Recombinant

生物種 Human

配列 MAASAAAASAAASAASGSPGPGEGSAGGEKRSTAPSAAASA

SASAAASS

PAGGGAEALELLEHCGVCRERLRPEREPRLLPCLHSACSACL

GPAAPAAA

NSSGDGGAAGDGTVVDCPVCKQQCFSKDIVENYFMRDSGSKA

ATDAQDAN

QCCTSCEDNAPATSYCVECSEPLCETCVEAHQRVKYTKDHTV

RSTGPAKS

RDGERTVYCNVHKHEPLVLFCESCDTLTCRDCQLNAHKDHQY

QFLEDAVR

NQRKLLASLVKRLGDKHATLQKSTKEVRSSIRQVSDVQKRVQ

VDVKMAIL

QIMKELNKRGRVLVNDAQKVTEGQQERLERQHWTMTKIQKHQ

EHILRFAS

WALESDNNTALLLSKKLIYFQLHRALKMIVDPVEPHGEMKFQ

WDLNAWTK

SAEAFGKIVAERPGTNSTGPAPMAPPRAPGPLSKQGSGSSQP

MEVQEGYG

FGSGDDPYSSAEPHVSGVKRSRSGEGEVSGLMRKVPRVSLER

LDLDLTAD

SQPPVFKVFPGSTTEDYNLIVIERGAAAAATGQPGTAPAGTP

GAPPLAGM

AIVKEEETEAAIGAPPTATEGPETKPVLMALAEGPGAEGPRL

ASPSGSTS

SGLEVVAPEGTSAPGGGPGTLDDSATICRVCQKPGDLVMCNQ

CEFCFHLD

CHLPALQDVPGEEWSCSLCHVLPDLKEEDGSLSLDGADSTGV

VAKLSPAN

1

QRKCERVLLALFCHEPCRPLHQLATDSTFSLDQPGGTLDLTL

IRARLQEK

LSPPYSSPQEFAQDVGRMFKQFNKLTEDKADVQSIIGLQRFF

ETRMNEAF

GDTKFSAVLVEPPPMSLPGAGLSSQELSGGPGDGP

予測される分子量 115 kDa including tags

領域 1 to 835

タグ GST tag N-Terminus

特性

Our Abpromise quarantee covers the use of ab131899 in the following tested applications.

The application notes include recommended starting dilutions; optimal dilutions/concentrations should be determined by the end user.

アプリケーション Western blot

SDS-PAGE

ELISA

製品の状態 Liquid

前処理および保存

保存方法および安定性 Shipped on dry ice. Upon delivery aliquot and store at -80°C. Avoid freeze / thaw cycles.

pH: 8.00

Constituents: 0.31% Glutathione, 0.79% Tris HCI

関連情報

機能 Nuclear corepressor for KRAB domain-containing zinc finger proteins (KRAB-ZFPs). Mediates

gene silencing by recruiting CHD3, a subunit of the nucleosome remodeling and deacetylation (NuRD) complex, and SETDB1 (which specifically methylates histone H3 at 'Lys-9' (H3K9me)) to the promoter regions of KRAB target genes. Enhances transcriptional repression by coordinating the increase in H3K9me, the decrease in histone H3 'Lys-9 and 'Lys-14' acetylation (H3K9ac and

H3K14ac, respectively) and the disposition of HP1 proteins to silence gene expression.

Recruitment of SETDB1 induces heterochromatinization. May play a role as a coactivator for CEBPB and NR3C1 in the transcriptional activation of ORM1. Also corepressor for ERBB4.

Inhibits E2F1 activity by stimulating E2F1-HDAC1 complex formation and inhibiting E2F1 acetylation. May serve as a partial backup to prevent E2F1-mediated apoptosis in the absence of

RB1. Important regulator of CDKN1A/p21(CIP1). Has E3 SUMO-protein ligase activity toward

itself via its PHD-type zinc finger.

組織特異性 Expressed in all tissues tested including spleen, thymus, prostate, testis, ovary, small intestine,

colon and peripheral blood leukocytes.

パスウェイ Protein modification; protein sumoylation.

配列類似性 Belongs to the TRIM/RBCC family.

Contains 2 B box-type zinc fingers.

Contains 1 bromo domain.
Contains 1 PHD-type zinc finger.

Contains 1 RING-type zinc finger.

ドメイン

The HP1 box is both necessary and sufficient for HP1 binding.

The PHD-type zinc finger enhances CEBPB transcriptional activity. The PHD-type zinc finger, the HP1 box and the bromo domain, function together to assemble the machinery required for repression of KRAB domain-containing proteins. Acts as an intramolecular SUMO E3 ligase for autosumoylation of bromodomain.

The RING-finger-B Box-coiled-coil/tripartite motif (RBCC/TRIM motif) is required for interaction with the KRAB domain of KRAB-zinc finger proteins. Binds four zinc ions per molecule. The RING finger and the N-terminal of the leucine zipper alpha helical coiled-coil region of RBCC are required for oligomerization.

Contains one Pro-Xaa-Val-Xaa-Leu (PxVxL) motif, which is required for interaction with chromoshadow domains. This motif requires additional residues -7, -6, +4 and +5 of the central Val which contact the chromoshadow domain.

翻訳後修飾

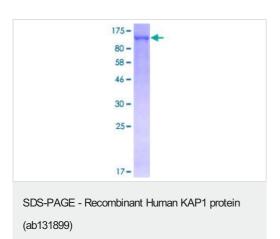
Phosphorylated upon DNA damage, probably by ATM or ATR. ATM-induced phosphorylation on Ser-824 represses sumoylation leading to the de-repression of expression of a subset of genes involved in cell cycle control and apoptosis in response to genotoxic stress. Dephosphorylation by the phosphatases, PPP1CA and PP1CB forms, allows sumoylation and expression of TRIM28 target genes.

Sumoylation/desumoylation events regulate TRIM28-mediated transcriptional repression. Sumoylation is required for interaction with CHD3 and SETDB1 and the corepressor activity. Represses and is repressed by Ser-824 phosphorylation. Enhances the TRIM28 corepressor activity, inhibiting transcriptional activity of a number of genes including GADD45A and CDKN1A/p21. Lys-554, Lys-779 and Lys-804 are the major sites of sumoylation. In response to Dox-induced DNA damage, enhanced phosphorylation on Ser-824 prevents sumoylation and allows de-repression of CDKN1A/p21.

細胞内局在

Nucleus. Associated with centromeric heterochromatin during cell differentiation through CBX1.

画像



12.5% SDS-PAGE analysis of ab131899 stained with Coomassie Blue.

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