abcam

Product datasheet

Recombinant Human Rb protein ab83205

2 Images

Description

Product name Recombinant Human Rb protein

Purity > 95 % SDS-PAGE.

Purified by affinity and FPLC chromatography.

Expression system Baculovirus infected insect cells

Accession NM_000321

Protein length Full length protein

Animal free No

Nature Recombinant

Species Human

Sequence MPPKTPRKTAATAAAAAEPPAPPPPPPEEDPEQDSG

PEDLPLVRLEFE

ETEEPDFTALCQKLKIPDHVRERAWLTWEKVSSVDGVLG

GYIQKKKELWG

ICIFIAAVDLDEMSFTFTELQKNIEISVHKFFNLLKEIDTSTKV

DNAMSR

LLKKYDVLFALFSKLERTCELIYLTQPSSSISTEINSALVLKV

SWITFLL

AKGEVLQMEDDLVISFQLMLCVLDYFIKLSPPMLLKEPYKT

AVIPINGSP

RTPRRGQNRSARIAKQLENDTRIIEVLCKEHECNIDEVKNV

YFKNFIPFM

NSLGLVTSNGLPEVENLSKRYEEIYLKNKDLDARLFLDHD

KTLQTDSIDS

FETQRTPRKSNLDEEVNVIPPHTPVRTVMNTIQQLMMILNS

ASDQPSENL

ISYFNNCTVNPKESILKRVKDIGYIFKEKFAKAVGQGCVEIG

SQRYKLGV

RLYYRVMESMLKSEEERLSIQNFSKLLNDNIFHMSLLACAL

EVVMATYSR

STSQNLDSGTDLSFPWILNVLNLKAFDFYKVIESFIKAEGN

LTREMIKHL

ERCEHRIMESLAWLSDSPLFDLIKQSKDREGPTDHLESAC

PLNLPLQNNH

TAADMYLSPVRSPKKKGSTTRVNSTANAETQATSAFQTQ

1

KPLKSTSLSLF

YKKVYRLAYLRLNTLCERLLSEHPELEHIWTLFQHTLQNEY

ELMRDRHL

DQIMMCSMYGICKVKNIDLKFKIIVTAYKDLPHAVQETFKRV

LIKEEEYD

SIIVFYNSVFMQRLKTNILQYASTRPPTLSPIPHIPRSPYKFP

SSPLRIP

GGNIYISPLKSPYKISEGLPTPTKMTPRSRILVSIGESFGTSE

KFQKINQ

MVCNSDRVLKRSAEGSNPPKPLKKLRFDIEGSDEADGSK

HLPGESKFQQK

LAEMTSTRTRMQKQKMNDSMDTSNKEEK

Predicted molecular weight 108 kDa

Amino acids 1 to 928

Tags His tag N-Terminus

Specifications

Our Abpromise guarantee covers the use of ab83205 in the following tested applications.

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

Applications SDS-PAGE

Gel Supershift Assays

Western blot

EMSA

Form Liquid

Preparation and Storage

Stability and Storage Shipped on dry ice. Upon delivery aliquot and store at -80°C. Avoid freeze / thaw cycles.

pH: 7.9

Constituents: 0.0154% DTT, 0.316% Tris HCI, 0.00584% EDTA, 20% Glycerol (glycerin,

glycerine)

General Info

Function

Key regulator of entry into cell division that acts as a tumor suppressor. Promotes G0-G1 transition when phosphorylated by CDK3/cyclin-C. Acts as a transcription repressor of E2F1 target genes. The underphosphorylated, active form of RB1 interacts with E2F1 and represses its transcription activity, leading to cell cycle arrest. Directly involved in heterochromatin formation by maintaining overall chromatin structure and, in particular, that of constitutive heterochromatin by stabilizing histone methylation. Recruits and targets histone methyltransferases SUV39H1, KMT5B and KMT5C, leading to epigenetic transcriptional repression. Controls histone H4 'Lys-20' trimethylation. Inhibits the intrinsic kinase activity of TAF1. Mediates transcriptional repression by SMARCA4/BRG1 by recruiting a histone deacetylase (HDAC) complex to the c-FOS promoter. In resting neurons, transcription of the c-FOS promoter is inhibited by BRG1-dependent recruitment of a phospho-RB1-HDAC1 repressor complex. Upon calcium influx, RB1 is

dephosphorylated by calcineurin, which leads to release of the repressor complex (By similarity). In case of viral infections, interactions with SV40 large T antigen, HPV E7 protein or adenovirus E1A protein induce the disassembly of RB1-E2F1 complex thereby disrupting RB1's activity.

Tissue specificity

Expressed in the retina.

Involvement in disease

Childhood cancer retinoblastoma

Bladder cancer

Osteogenic sarcoma

Sequence similarities

Belongs to the retinoblastoma protein (RB) family.

Domain

The Pocket domain binds to the threonine-phosphorylated domain C, thereby preventing interaction with heterodimeric E2F/DP transcription factor complexes.

Post-translational modifications

Phosphorylated by CDK6 and CDK4, and subsequently by CDK2 at Ser-567 in G1, thereby releasing E2F1 which is then able to activate cell growth. Dephosphorylated at the late M phase. SV40 large T antigen, HPV E7 and adenovirus E1A bind to the underphosphorylated, active form of pRb. Phosphorylation at Thr-821 and Thr-826 promotes interaction between the C-terminal domain C and the Pocket domain, and thereby inhibits interactions with heterodimeric E2F/DP transcription factor complexes. Dephosphorylated at Ser-795 by calcineruin upon calcium stimulation. CDK3/cyclin-C-mediated phosphorylation at Ser-807 and Ser-811 is required for G0-G1 transition. Phosphorylated by CDK1 and CDK2 upon TGFB1-mediated apoptosis.

N-terminus is methylated by METTL11A/NTM1 (By similarity). Monomethylation at Lys-810 by SMYD2 enhances phosphorylation at Ser-807 and Ser-811, and promotes cell cycle progression. Monomethylation at Lys-860 by SMYD2 promotes interaction with L3MBTL1.

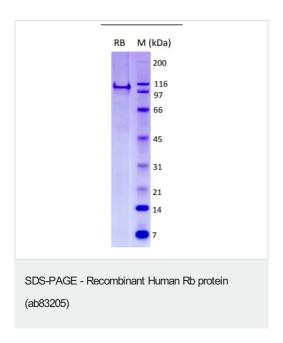
Acetylation at Lys-873 and Lys-874 regulates subcellular localization, at least during keratinocytes

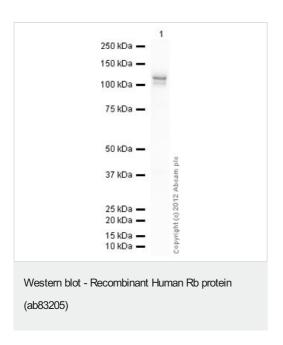
differentiation.

Cellular localization

Nucleus.

Images





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