

**Anti-Aurora B Antibody**  
Catalog # ABO11114**Specification****Anti-Aurora B Antibody - Product Information**

Application	WB
Primary Accession	<a href="#">O70126</a>
Host	Rabbit
Reactivity	Mouse, Rat
Clonality	Polyclonal
Format	Lyophilized

**Description**

Rabbit IgG polyclonal antibody for Aurora kinase B(AURKB) detection. Tested with WB in Mouse;Rat.

**Reconstitution**

Add 0.2ml of distilled water will yield a concentration of 500ug/ml.

**Anti-Aurora B Antibody - Additional Information**

Gene ID 20877

**Other Names**

Aurora kinase B, 2.7.11.1, Aurora 1, Aurora- and IPL1-like midbody-associated protein 1, Aurora/IPL1-related kinase 2, ARK-2, Aurora-related kinase 2, STK-1, Serine/threonine-protein kinase 12, Serine/threonine-protein kinase 5, Serine/threonine-protein kinase aurora-B, Aurkb, Aik2, Aim1, Airk2, Ark2, Stk1, Stk12, Stk5

**Calculated MW**

39384 MW KDa

**Application Details**

Western blot, 0.1-0.5 µg/ml, Rat, Mouse<br>

**Subcellular Localization**

Nucleus . Chromosome . Chromosome, centromere . Cytoplasm, cytoskeleton, spindle . Midbody . Localizes on chromosome arms and inner centromeres from prophase through metaphase and then transferring to the spindle midzone and midbody from anaphase through cytokinesis. Colocalized with gamma tubulin in the mid-body (By similarity). Proper localization of the active, Thr-237-phosphorylated form during metaphase may be dependent upon interaction with SPDYC. Colocalized with SIRT2 during cytokinesis with the midbody (By similarity). .

**Tissue Specificity**

Expressed in testis, intestine and spleen. All of them are tissues that contain a large number of proliferating cells. Expressed during S phase, in a cell-cycle-dependent fashion.

**Protein Name**

Aurora kinase B

## Contents

Each vial contains 5mg BSA, 0.9mg NaCl, 0.2mg Na<sub>2</sub>HPO<sub>4</sub>, 0.05mg Thimerosal, 0.05mg NaN<sub>3</sub>.

## Immunogen

A synthetic peptide corresponding to a sequence at the N-terminus of mouse Aurora B(72-87aa QNKQPFTIDNFEIGRP), different from the related rat sequence by two amino acids.

## Purification

Immunogen affinity purified.

## Cross Reactivity

No cross reactivity with other proteins

## Storage

**At -20°C for one year. After reconstitution, at 4°C for one month. It can also be aliquotted and stored frozen at -20°C for a longer time. Avoid repeated freezing and thawing.**

## Sequence Similarities

Belongs to the protein kinase superfamily. Ser/Thr protein kinase family. Aurora subfamily.

## Anti-Aurora B Antibody - Protein Information

### Name Aurkb

### Function

Serine/threonine-protein kinase component of the chromosomal passenger complex (CPC), a complex that acts as a key regulator of mitosis (By similarity). The CPC complex has essential functions at the centromere in ensuring correct chromosome alignment and segregation and is required for chromatin-induced microtubule stabilization and spindle assembly (By similarity). Involved in the bipolar attachment of spindle microtubules to kinetochores and is a key regulator for the onset of cytokinesis during mitosis (By similarity). Required for central/midzone spindle assembly and cleavage furrow formation (By similarity). Key component of the cytokinesis checkpoint, a process required to delay abscission to prevent both premature resolution of intercellular chromosome bridges and accumulation of DNA damage: phosphorylates CHMP4C, leading to retain abscission-competent VPS4 (VPS4A and/or VPS4B) at the midbody ring until abscission checkpoint signaling is terminated at late cytokinesis (By similarity). AURKB phosphorylates the CPC complex subunits BIRC5/survivin, CDCA8/borealin and INCENP (By similarity). Phosphorylation of INCENP leads to increased AURKB activity (By similarity). Other known AURKB substrates involved in centromeric functions and mitosis are CENPA, DES/desmin, GPAF, KIF2C, NSUN2, RACGAP1, SEPTIN1, VIM/vimentin, HASPIN, and histone H3 (By similarity). A positive feedback loop involving HASPIN and AURKB contributes to localization of CPC to centromeres (By similarity). Phosphorylation of VIM controls vimentin filament segregation in cytokinetic process, whereas histone H3 is phosphorylated at 'Ser-10' and 'Ser-28' during mitosis (H3S10ph and H3S28ph, respectively) (PubMed:<a href="http://www.uniprot.org/citations/11784863" target="\_blank">11784863</a>). AURKB is also required for kinetochore localization of BUB1 and SGO1 (By similarity). Phosphorylation of p53/TP53 negatively regulates its transcriptional activity (By similarity). Key regulator of active promoters in resting B- and T-lymphocytes: acts by mediating phosphorylation of H3S28ph at active promoters in resting B- cells, inhibiting RNF2/RING1B-mediated ubiquitination of histone H2A and enhancing binding and activity of the USP16 deubiquitinase at transcribed genes (PubMed:<a href="http://www.uniprot.org/citations/24034696" target="\_blank">24034696</a>). Acts as an inhibitor of CGAS during mitosis: catalyzes phosphorylation of the N-terminus of CGAS during the G2-M transition, blocking CGAS liquid phase separation and activation, and thereby preventing CGAS-induced autoimmunity (By similarity). Phosphorylates KRT5 during anaphase and

telophase (PubMed:<a href="http://www.uniprot.org/citations/29518391" target="\_blank">29518391</a>). Phosphorylates ATXN10 which promotes phosphorylation of ATXN10 by PLK1 and may play a role in the regulation of cytokinesis and stimulating the proteasomal degradation of ATXN10 (By similarity).

#### Cellular Location

Nucleus {ECO:0000250|UniProtKB:Q96GD4}. Chromosome {ECO:0000250|UniProtKB:Q96GD4}. Chromosome, centromere {ECO:0000250|UniProtKB:Q96GD4}. Chromosome, centromere, kinetochore {ECO:0000250|UniProtKB:Q96GD4}. Cytoplasm, cytoskeleton, spindle {ECO:0000250|UniProtKB:Q96GD4}. Midbody {ECO:0000250|UniProtKB:Q96GD4} Note=Localizes on chromosome arms and inner centromeres from prophase through metaphase and then transferring to the spindle midzone and midbody from anaphase through cytokinesis. Colocalized with gamma tubulin in the midbody. Proper localization of the active, Thr-237- phosphorylated form during metaphase may be dependent upon interaction with SPDYC. Colocalized with SIRT2 during cytokinesis with the midbody Localization (and probably targeting of the CPC) to the inner centromere occurs predominantly in regions with overlapping mitosis- specific histone phosphorylations H3pT3 and H2ApT12 {ECO:0000250|UniProtKB:Q96GD4}

#### Tissue Location

Expressed in testis, intestine and spleen. All of them are tissues that contain a large number of proliferating cells Expressed during S phase, in a cell-cycle-dependent fashion

### Anti-Aurora B Antibody - Protocols

Provided below are standard protocols that you may find useful for product applications.

- [Western Blot](#)
- [Blocking Peptides](#)
- [Dot Blot](#)
- [Immunohistochemistry](#)
- [Immunofluorescence](#)
- [Immunoprecipitation](#)
- [Flow Cytometry](#)
- [Cell Culture](#)

### Anti-Aurora B Antibody - Images



Anti-Aurora B antibody, ABO11114, Western blotting Lane 1: Rat Liver Tissue Lysate Lane 2: 22RV Cell Lysate Lane 3: HELA Cell Lysate Lane 4: SW620 Cell Lysate

### **Anti-Aurora B Antibody - Background**

AURKB (aurora kinase B) also called Aik2, AIM-1, ARK2, AurB, STK12, IPL1, PPP1R48 or STK5, localizes to microtubules near kinetochores, specifically to the specialized microtubules called K-fibers. Cell cycle and Northern blot analyses showed that STK12 is expressed in the S phase and persistently thereafter. Western blot analysis indicated that STK12 is localized in the midbodies during anaphase. Northern blot analysis detected strong expression of a 1.5-kb STK12 transcript in thymus, with weaker expression in small intestine, testis, colon, spleen, and brain. The AURKB gene is mapped on 17p13.1. Examination of the role of both kinases in the phosphorylation of CENPA revealed that the reaction is mediated sequentially by AURKA and AURKB in early mitosis. EB1 overexpression enhanced AURKB kinase activity, and knockdown of EB1 with small interfering RNA impaired AURKB activity. EB1 protected AURKB from dephosphorylation/inactivation by protein phosphatase-2A by blocking binding of PP2A to AURKB.