# Probing the Dynamics and Functions of Aurora B Kinase in Living Cells during Mitosis and Cytokinesis $^{\boxed{V}}$

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Aurora B is a protein kinase and a chromosomal passenger protein that undergoes dynamic redistribution during mitosis. We have probed the mechanism that regulates its localization with cells expressing green fluorescent protein (GFP)-tagged wild-type or mutant aurora B. Aurora B was found at centromeres at prophase and persisted until ~0.5 min after anaphase onset, when it redistributed to the spindle midzone and became concentrated at the equator along midzone microtubules. Depolymerization of microtubules inhibited the dissociation of aurora B from centromeres at early anaphase and caused the dispersion of aurora B from the spindle midzone at late anaphase; however, centromeric association during prometaphase was unaffected. Inhibition of CDK1 deactivation similarly caused aurora B to remain associated with centromeres during anaphase. In contrast, inhibition of the kinase activity of aurora B appeared to have no effect on its interactions with centromeres or initial relocation onto midzone microtubules. Instead, kinase-inactive aurora B caused abnormal mitosis and deactivation of the spindle checkpoint. In addition, midzone microtubule bundles became destabilized and aurora B dispersed from the equator. Our results suggest that microtubules, CDK1, and the kinase activity each play a distinct role in the dynamics and functions of aurora B in dividing cells.

### **INTRODUCTION**

Aurora/Ipl1p-related proteins are a family of well-conserved serine-threonine kinases involved in various stages of mitosis (Giet and Prigent, 1999). The founding members are Ipl1p from *Saccharomyces cerevisiae* (Francisco and Chan, 1994) and aurora from *Drosophila melanogaster* (Glover *et al.*, 1995). Aurora A functions in centrosome separation and spindle bipolarity (Glover *et al.*, 1995; Roghi *et al.*, 1998; Giet and Prigent, 2000), whereas aurora B appears to function in both early and late mitotic events (Bischoff *et al.*, 1998; Schumacher *et al.*, 1998; Terada *et al.*, 1998). Ablation of aurora B activity by RNA interference (Schumacher *et al.*, 1998; Kaitna *et al.*, 2000; Adams *et al.*, 2001; Giet and Glover, 2001) or by deactivating a temperature-sensitive (ts) mutant (Severson *et al.*, 2000) leads to a wide range of defects in prometaphase chromosomal congregation, anaphase chromosomal segre-

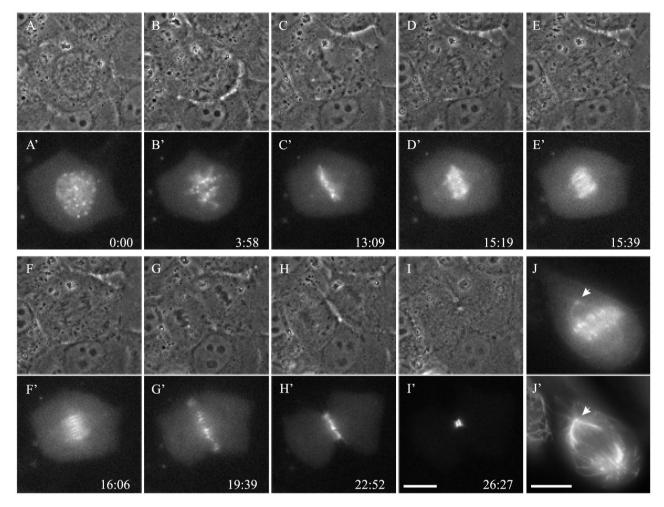
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☑ Online version of this article contains video material for some figures. Online version available at www.molbiolcell.org. Abbreviations used: GFP, green fluorescent protein; INCENP, inner centromere protein; NRK, normal rat kidney.

gation, and cytokinesis. Aurora B also plays a role in chromosomal condensation by phosphorylating the H3 histone (Hsu *et al.*, 2000; Adams *et al.*, 2001; Giet and Glover, 2001).

Immunolocalization studies demonstrated that aurora B localizes at chromosomal centromeres during prometaphase, and subsequently relocates to midzone microtubules and midbodies during anaphase and telophase (Schumacher et al., 1998; Adams et al., 2001; Giet and Glover, 2001). This distribution pattern is similar to that of several chromosome passenger proteins, including inner centromere protein (INCENP; Cooke et al., 1987; Mackay et al., 1998), TD60 (Andreassen et al., 1991; Martineau-Thuillier et al., 1998), and survivin (Skoufias et al., 2000; Uren et al., 2000). Furthermore, as for aurora B, INCENP has been implicated in cytokinesis based on gene ablation and mutation experiments (Eckley et al., 1997; Mackay et al., 1998; Cutts et al., 1999; Adams et al., 2000, 2001; Kaitna *et al.*, 2000). The similar behavior of these proteins raises the possibility that they form a complex that is transported by microtubule-based motors to the equator during cytokinesis. This notion is supported by the identification of a complex from Xenopus egg extracts and from HeLa cells extracts that contains both INCENP and aurora B (Adams et al., 2000; Kaitna et al., 2000).

From these studies, it is clear that the dynamics of aurora B and associated proteins plays a critical role in their functions.



**Figure 1.** Dynamics of aurora B in living mitotic NRK cells transfected with aurora B-GFP. Time-lapse phase-contrast (A–I) and corresponding fluorescence (A′–I′) images show the redistribution of aurora B during mitosis and cytokinesis. Time elapsed since prophase is shown in minutes:seconds. Aurora B was localized at centromeres at prophase (A and A′), and persisted through prometaphase (B and B′), metaphase (C and C′), and early anaphase (D and D′). It then started to relocate to midzone microtubules (E and E′), initially spanning the space between separating chromosomes (E and E′ and F and F′) but later condensing into short segments on the equatorial plane (G and G′ and H and H′). These segments then coalesced laterally to form the midbody (I and I′). (J and J′) Separate cell fixed at early anaphase and stained with anti-β-tubulin. Note that aurora B was localized at the spindle pole (arrows). Bars, 10 μm.

Changes in the localization of aurora B may serve as an effective mechanism for defining its substrates in a temporally and spatially dependent manner. Although there is still no direct evidence, it is possible that CDK1 serves as a master switch for regulating the localization of these proteins. An attractive possibility is that signals of CDK1 may be relayed through aurora B to other components of the chromosomal passenger complex and the associated motor molecule MKLP1/ZEN-4 (Terada *et al.*, 1998; Severson *et al.*, 2000). Phosphorylation of these proteins by aurora B may in turn effect the subsequent localization of aurora B at the spindle midzone.

So far, the localization of aurora B and other chromosomal passenger proteins has relied on immunofluorescence of fixed cells. Due to the highly dynamic nature of these proteins, detailed understanding of their function and regulation may benefit greatly from the direct observations and manipulations of living cells. In this study we have transfected rat aurora B as

a green fluorescent protein (GFP) fusion protein into normal rat kidney (NRK) cells. To test the role of its kinase activity, we have also created a GFP fusion protein of kinase-inactive aurora B. We were able to follow the striking relocation of aurora B in living cells, and to test the effects of several factors that may affect aurora B distribution. Our results suggest that both microtubules and CDK1 play a major role in the relocation of aurora B from centromeres to the central spindle during anaphase. In contrast, the kinase activity of aurora B is required for its stable localization once aurora B is relocated along midzone microtubules.

### **MATERIALS AND METHODS**

### Cell Culture, Microscopy, and Image Processing

Normal rat kidney epithelial cells (NRK-52E; American Type Culture Collection, Rockville, MD) were cultured in Kaighn's modified

1100 Molecular Biology of the Cell

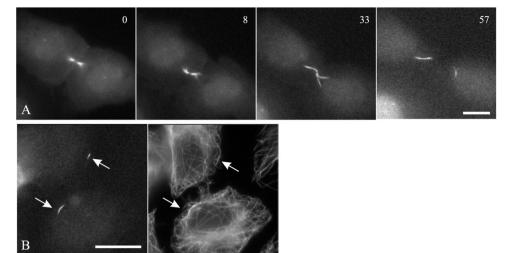


Figure 2. Dissociation of aurora B from the midbody after cytokinesis. Time-lapse images show the dissociation of aurora B-GFP from the midbody as one or more segments that migrate toward the cell center (A). Time elapsed in minutes is given at the upper right corner. Fixation and staining for  $\beta$ -tubulin in a separate cell showed that the condensed segments of aurora B (B, left), already well separated from the region occupied by the midbody, were localized along interphase microtubules (B, right). Bars, 10 μm.

F12 (F12K) medium supplemented with 10% fetal bovine serum (JRH Biosciences, Lenexa, KS), 50 U/ml penicillin, and 50  $\mu$ g/ml streptomycin, on glass chamber dishes as previously described (Mckenna and Wang, 1989). The cells were maintained at 37°C in a stage incubator built on top of an Axiovert S100TV inverted microscope (Carl Zeiss, Thornwood, NY) and viewed with a 100×, numerical aperture 1.30 Fluar lens. All images were acquired with a cooled charge-coupled device camera (ST133 controller and CCD57 chip; Roper Scientific, Treton, NJ) and processed with custom software for background subtraction.

### Plasmid Construction and Transfection

cDNA fragments encoding aurora B and its kinase-inactive mutant were amplified by polymerase chain reaction with FLAG-AIM-1 (WT) and FLAG-AIM-1 (K-R) as templates (Terada et al., 1998), respectively, and subcloned into pEGFP-N1 vector (CLONTECH, Palo Alto, CA). The enhanced green fluorescent protein (EGFP) sequence is located at the 3' end of the aurora B sequence. For transient transfection, NRK cells were plated at a density of  $5 \times 10^4$ cells/ml on a coverslip chamber dish and incubated for 18-24 h. Immediately before transfection, the cells were rinsed once in serum-free F12K or Opti-MEM I medium (Invitrogen, Carlsbad, CA). The cells were transfected with the DNA construct (2  $\mu$ g) by using LipofectAMINE according to manufacturer's instructions (Invitrogen). After 4 h of incubation, the medium containing DNA-LipofectAMINE was replaced with the F12K medium containing 10% FBS, and the cells was cultured for an additional 14-16 h. To maintain the expression of aurora B-GFP, NRK cells on 60-mm culture dishes were transfected with 3  $\mu g$  of aurora B-GFP as described above and cultured in the presence of 800 µg/ml G418 (Invitrogen).

### In Vitro Transcription of Cyclin B $\Delta 90$ and Preparation of Rhodamine-labeled Tubulin

Messenger RNA for the nondegradable mutant of cyclin B (Δ90 cyclin B) was synthesized with the mMESSAGE mMACHINE kit (Ambion, Austin, TX) and prepared for microinjection as described previously with minor modifications (Wheatley *et al.*, 1997): transcribed mRNA was separated from unincorporated nucleotides by a Spin Column 10 (Sigma Chemical, St. Louis, MO) and the yield was estimated by UV absorbance. Tubulin was prepared and labeled with 5- (and 6-) carboxytetramethylrhodamine (Molecular Probes, Eugene, OR) as described previously (Wheatley *et al.*, 1997, 1998).

### Microinjection, Drug Treatment, and Immunofluorescence

Microinjection was performed by low-speed continuous flow with custom-drawn glass needles and a custom-designed pressure control system (Wang, 1992). Nocodazole (Sigma Chemical) was stored at  $-20^{\circ}\text{C}$  as  $10^{3}\times\text{stocks}$  in dimethyl sulfoxide and diluted into prewarmed medium before application to cells.

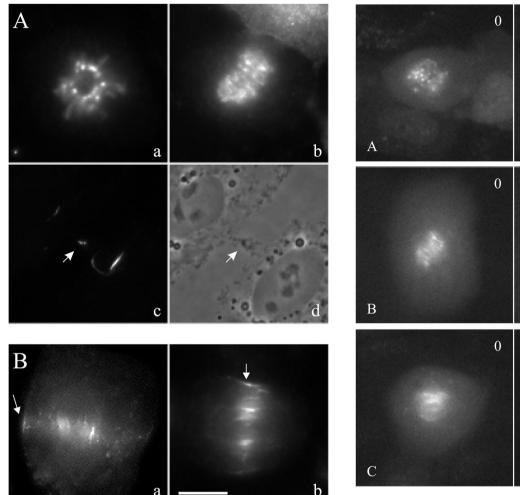
For microtubule and aurora B immunofluorescence, cells were rinsed with warm cytoskeletal buffer and fixed with 4% paraformal-dehyde (EM Scientific, Gibbstown, NJ) in warm cytoskeleton buffer for 10 min (Wheatley and Wang, 1996). They were then rinsed thoroughly in the cytoskeletal buffer and permeablized by incubation with 0.5% Triton X-100 for 5 min. Fixed cells were rinsed with the cytoskeletal buffer, blocked for 10 min with 1% bovine serum albumin (BSA) (Roche Applied Sciences, Indianapolis, IN) in PBS, and then incubated with anti-β-tubulin monoclonal antibodies (Amersham Biosciences, Piscataway, NJ) at a dilution of 1:10 in PBS with 1% BSA or anti-AIM-1 monoclonal antibodies (Murata-Hori *et al.*, 2000) at a dilution of 1:50 for 45 min at 37°C. After washing with PBS/BSA thoroughly, the cells were incubated with Alexa 546-conjugated goat anti-mouse antibodies (Molecular Probes) at a dilution of 1:100 for 30 min at 37°C.

Mad2 staining was performed with a method modified from that of Waters *et al.* (1998). The cells were rinsed with PHEM buffer (60 mM PIPES, 25 mM HEPES, 10 mM EGTA, and 4 mM MgSO<sub>4</sub>, pH 6.9), lysed for 1 min 0.1% Triton X-100 in PHEM buffer, and then fixed with 4% formaldehyde in PHEM buffer for 20 min. Fixed cells were rinsed with PBS with 0.05% Tween 20, blocked for 30 min with 1% BSA in PBS, and incubated with anti-Mad2 rabbit polyclonal antibodies (gift of Dr. E.D. Salmon, University of North Carolina, Chapel Hill, NC) at a dilution of 1:50 in PBS with 2.5% BSA for 45 min at 37°C. After washing with PBS with 2.5% BSA, cells were incubated with Alexa 546-conjugated goat anti-rabbit antibodies (Molecular Probes) at a dilution of 1:100 for 30 min at 37°C. Finally, cells were rinsed with PBS and mounted in an antibleaching medium (Wheatley and Wang, 1996).

### **RESULTS**

# Aurora B-GFP as a Probe of Aurora B Dynamics in Dividing NRK Cells

NRK cells were transfected with GFP-tagged aurora B and observed with fluorescence optics during mitosis and cyto-



**Figure 3.** Distribution of endogenous aurora B and aurora B-GFP during cell division. (A) Immunofluorescence localization of aurora B at prometaphase (a), anaphase onset (b), and after cytokinesis (c and d; arrows point to the midbody). The distribution is similar to that seen with aurora B-GFP in Figures 1 and 2. (B) Localization of aurora B during anaphase. Both aurora B-GFP (a) and endogenous aurora B (b) can be found on the equatorial cortex (arrows) and along midzone microtubules. Bar,  $10~\mu m$ .

kinesis (Figure 1). Aurora B-GFP became localized as early as prophase (before nuclear envelope breakdown) at chromosomal centromeres (Figure 1A). Some aurora B-GFP was also found at the spindle pole (Figure 1J), as confirmed by counterstaining for β-tubulin (Figure 1J'). During anaphase, aurora B-GFP dissociated from centromeres and redistributed to midzone microtubules. However, the relocation occurred slightly after anaphase onset, because there was a period of 20–30 s when most aurora B-GFP was still associated with centromeres of separated chromosomes (Figure 1, D–F). Relocated aurora B initially spanned the region between separated chromosomes (Figure 1, E–F), but subsequently condensed into short segments on the equatorial plane that collapsed laterally during cytokinesis to form the midbody (Figure 1, G–I).

**Figure 4.** Effects of nocodazole on the dynamics of aurora B-GFP. NRK cells stably expressing aurora B-GFP were treated with  $10-20~\mu M$  nocodazole at prometaphase (A), early anaphase (B) or midanaphase (C). Images of aurora B-GFP were recorded immediately before (left) and after nocodazole treatment (right). Time elapsed since the addition of nocodazole is given in minutes at the upper right corner. Treatment with nocodazole had no effect on centromeric association of aurora B at prometaphase (A); however, it inhibited the dissociation of aurora B from centromeres at early anaphase (B), and caused the dispersion of aurora B that was localized along midzone microtubules during late anaphase (C). Bar,  $10~\mu m$ .

10

The distribution of aurora B-GFP was followed beyond the completion of cytokinesis, to determine the fate of aurora B-GFP concentrated at the midbody. The condensed structure extended into elongated segments and migrated toward the cell center (Figure 2A). Counterstaining for  $\beta$ -tubulin indicated that the movement took place along microtubules of the interphase network (Figure 2B). These aurora B-containing segments subsequently dispersed beyond detection, before they reached a defined destination.

The distribution of aurora B-GFP was generally consistent with immunolocalization of aurora B in other systems (Schumacher *et al.*, 1998; Adams *et al.*, 2001; Giet and Glover, 2001). To confirm that aurora B-GFP mimics the behavior of endog-

1102 Molecular Biology of the Cell

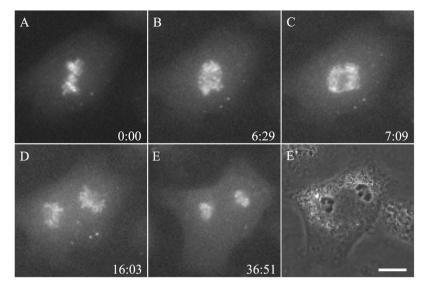


Figure 5. Effect of  $\Delta 90$  cyclin B on the dynamics of aurora B. An NRK cell stably expressing aurora B was microinjected with mRNA for  $\Delta 90$  cyclin B at prometaphase. Neither chromosomal congregation nor anaphase onset appeared affected. However, aurora B remained associated with centromeres 30 min after anaphase onset (E), and cytokinesis was inhibited (E'). The phase image corresponding to E is shown in E'. Time elapsed since metaphase is given in minutes: seconds at the lower right corner. Bar,  $10~\mu m$ .

enous aurora B, we fixed and stained nontransfected cells with antibodies against aurora B (Figure 3). As for aurora B-GFP, endogenous aurora B was localized at centromeres in prometaphase cells and remained associated with centromeres until shortly after anaphase onset (Figure 3A). Its subsequent relocation to midzone microtubules and the equatorial cortex (Figure 3B), and to bundle structures after the completion of cortical ingression, was also similar to what was observed with aurora B-GFP (Figure 3A). However, we were unable to detect the association of endogenous aurora B at spindle poles by immunofluorescence staining, possibly due to the limited amount of aurora B at these structures.

Based on the intensity of immunofluorescence staining, we were able to estimate the relative amount of aurora B in cells expressing aurora B-GFP and in nontransfected cells. Transfection under the present condition was found to increase the total amount of aurora B by an average of approximately twofold, which is likely lower than what was obtained previously when cells were transfected with a higher amount of DNA (Tatsuka *et al.*, 1998).

## Involvement of Microtubules and CDK1 in Dynamics of Aurora B

The ability to image aurora B in living cells allowed us to probe the mechanism that regulates its redistribution during

**Table 1.** Effect of kinase inactive mutant of aurora B on mitosis NRK cells were transiently transfected with aurora B-GFP or aurora B (K-R)-GFP. Cells expressing GFP proteins in prophase or prometaphase were observed with phase optics to assess the behavior of chromosomes.

	Chromosome miscongregation	Chromosome missegregation
aurora B (WT)	0.0% (0/16)	0.0% (0/16)
aurora B (K-R)	46.7% (14/30)	36.7% (11/30)

cell division. We first treated aurora B-GFP-expressing cells with nocodazole to determine whether microtubules are required for its dynamic relocation. No obvious change was observed in the centromeric localization of aurora B-GFP when cells were treated at prophase or prometaphase (100%) for n = 10; Figure 4A), suggesting that microtubules were not involved in maintaining the centromeric localization of aurora B during early mitosis. In contrast, microtubules were required for the dissociation of aurora B-GFP from centromeres and the relocation to the spindle midzone during anaphase, as indicated by the maintenance of aurora B-GFP on separated chromosomes when cells were treated with nocodazole shortly after anaphase onset (100% for n =10; Figure 4B). Application of nocodazole during later stages of anaphase caused aurora B-GFP to disperse from the spindle midzone (100% for n = 10; Figure  $\bar{4}C$ ), suggesting that microtubules were also involved in maintaining its midzone localization. Each of these experiments with nocodazole was repeated 10 times with consistent results.

We speculate that the dissociation of aurora B-GFP from the centromeres may be regulated by cyclin B degradation/ CDK1 deactivation. To address the involvement of CDK1, we microinjected the mRNA for Δ90 cyclin B, a nondegradable active domain of cyclin B, into dividing NRK cells transfected with aurora B-GFP. This treatment was previously shown to inhibit the degradation of cyclin B at anaphase onset, thus maintaining the CDK1 activity at a high level (Glotzer et al., 1991; Wheatley et al., 1997). The injection had no effect on metaphase chromosomal alignment or anaphase chromosomal separation (Figure 5, A–C; Wheatley et al., 1997). However, aurora B-GFP failed to dissociate from centromeres and cytokinesis failed as well (100% for n = 11; Figure 5, C-E). Thus, CDK1 inactivation is required for the redistribution of aurora B from chromosomes to the spindle midzone. To test whether CDK1 inactivation alone is sufficient for the release of aurora B, prometaphase cells were treated with a CDK1 inhibitor, purvalanol A (Chang et al., 1999; Chadebech et al., 2000). The treatment had no effect on the centromeric association (our unpublished data), suggest-

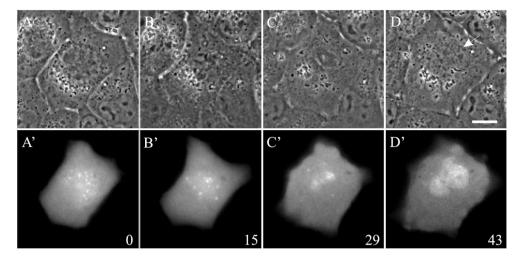


Figure 6. Dynamics of aurora B(K-R)-GFP in a cell that failed to congregate the chromosomes at prometaphase. An NRK cell transiently expressing aurora B(K-R)-GFP was monitored by timelapse phase contrast (A-D) and fluorescence optics (A'-D'). Time elapsed in minutes since nuclear envelope breakdown is shown at the lower right corner. Chromosomes in this cell failed to congregate (B and B'), and started to decondense ~30 min after nuclear envelope breakdown (C and C'). A nuclear envelope then formed around the chromosomes (D, arrow). There was no sign of cytokinesis throughout the period of observation. Bar, 10 μm.

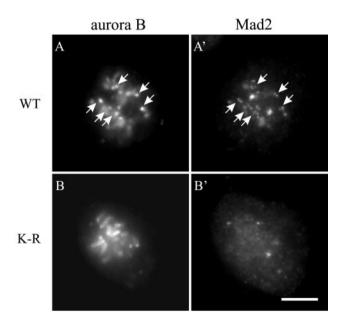
ing that the release of centromeric aurora B requires additional events such as dephosphorylation by a phosphatase, proteolysis, or the assembly of midzone microtubules.

# Involvement of Kinase Activity of Aurora B in Mitosis and Cytokinesis

To address the possibility that the kinase activity of aurora B may regulate its localization, we constructed a GFP-tagged, kinase inactive mutant of aurora B, by using FLAG-AIM-1(K-R) as a template (Terada et al., 1998). NRK cells were transiently transfected with wild-type aurora B-GFP or aurora B(K-R)-GFP. No mitotic defect was observed in cells expressing wild-type aurora B-GFP (Table 1). Time-lapse imaging of aurora B(K-R)-GFP indicated an apparently normal localization at centromeres during prometaphase (Figure 6A'). However, expression of aurora B(K-R)-GFP caused strong dominant negative effects (Table 1). Defects in mitosis were observed in >80% of the cells (n = 30). A large fraction of cells (46.7%) failed to congregate their chromosomes into a metaphase plate (Figure 6, A-C). Interestingly, 66.7% of these cells exited mitosis directly within 1.5 h of nuclear envelope breakdown, by forming a nuclear envelope around chromosomes without going through anaphase (Figure 6D, arrow). Aurora B(K-R)-GFP gradually dispersed from the centromeres as the nuclear envelope reformed. Localization of a spindle checkpoint component, Mad2 (Li and Murray, 1991), indicated that Mad2 disappeared from the kinetochores of these prometaphase-arrested cells (Figure 7, B and B'), despite the absence of a metaphase plate. In control cells, Mad2 dissociates from kinetochores only as chromosomes become congregated into the metaphase plate (Figure 7, A and A'; Chen et al., 1996; Li and Benezra, 1996).

Other cells expressing aurora B(K-R)-GFP (36.7%) managed to enter anaphase, but with defects in chromosomal segregation similar to what was reported after the ablation or deactivation of aurora B (Figure 8B, arrows; Kaitna *et al.*, 2000; Severson *et al.*, 2000; Adams *et al.*, 2001; Giet and Glover, 2001). All aurora B(K-R)-GFP-expressing cells showing mitotic defects also failed cytokinesis. In cells that failed to enter anaphase, there was no detectable initiation of cytokinesis (Figure 6D'). In cells that entered anaphase, cy-

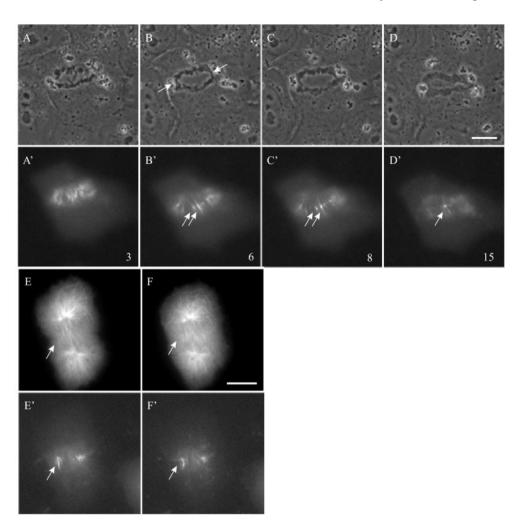
tokinesis failed after an initial ingression. Aurora B(K-R)-GFP was able to redistribute from centromeres to midzone microtubules (Figure 8A'), and to migrate into the equatorial plane to form discrete segments as in control cells (Figure 8, B'-C'). However, in contrast to the wild-type aurora B-GFP that became highly organized on the equatorial plane (Figure 1, F'-H'), aurora B(K-R)-GFP gradually dispersed from the spindle midzone (Figure 8, C'-D'). Microinjection of rhodamine-labeled tubulin into these aurora-B(K-R)-GFP-expressing cells indicated that, although midzone microtu-



**Figure 7.** Bypass of the spindle checkpoint in a prometaphase-arrested cells expressing aurora B(K-R)-GFP. Immunofluorescence indicates little Mad2 staining at kinetochores of a prometaphase-arrested cell transiently expressing aurora B(K-R)-GFP (B and B'), whereas a control prometaphase cell transiently expressing aurora B-GFP shows Mad2 staining on a number of kinetochores (A and A', arrows). Bar,  $10~\mu m$ .

1104 Molecular Biology of the Cell

Figure 8. Dynamics of aurora B(K-R) in a cell that entered anaphase. An NRK cell transiently expressing aurora B(K-R)-GFP was monitored by time-lapse phase (A–D) and fluorescence optics (A'-D'). Time elapsed in minutes after anaphase onset is shown at the lower right corner. Although anaphase onset took place normally, one or more chromosomes failed to segregate (B, arrows). Kinase-inactive aurora B (K-R)-GFP showed initial localization along midzone microtubules (A'), followed by concentration toward the equatorial plane to form segments (B', arrows). However, the intensity of these segments decreased subsequently, reducing them to small aggregates (C' and D', arrows). Cytokinesis was also inhibited (D). Rhodamine-labeled tubulin was microinjected into a cell expressing aurora-B(K-R)-GFP and the images of tubulin (E and F) and aurora-B(K-R) (E' and F') were recorded over a period of 8 min. Arrows indicate the positions in the spindle midzone where aurora-B(K-R)-GFP was concentrated. Midzone microtubules remained disorganized as aurora B dissociated from the equatorial region. Bar, 10  $\mu$ m.



bules were present, they never became organized as in control cells (Figure 8, E and F; Wheatley *et al.*, 1997). These observations suggest that the kinase activity of aurora B was not required for the initial localization to centromeres and the relocation to midzone microtubules, but was required for the organization of midzone microtubules and the stable localization of aurora B along the equator.

#### **DISCUSSION**

### Dynamics of Aurora B during Cell Division

Several significant aspects of aurora B distribution in living mammalian cells were noted in the present study. First, aurora B was localized not only at centromeres but also at spindle poles. The weakness of the polar signal might explain why this feature was not detected in immunofluorescence studies. Although the functional significance is unclear, a similar localization at spindle poles was observed with other kinases involved in cytokinesis such as polo (Arnaud *et al.*, 1998). Second, aurora B-GFP remained associated with separated chromosomes for a brief period at anaphase before relocating to interzonal microtubules, sug-

gesting that this event lies downstream of proteolysis involved in chromosomal separation. Third, the pattern of subsequent distribution, from broad localization between chromosomes to tight segments on the equatorial plane, suggests an active transport mechanism that brings aurora B and associated proteins toward the ends of microtubules. The kinesin-like motor that colocalizes with aurora B in the spindle midzone, known as CHO1/MKLP1, PAV-KLP, or ZEN-4 in different organisms, most likely plays a key role in this process (Nislow *et al.*, 1992; Powers *et al.*, 1998; Adams *et al.*, 1998; Raich *et al.*, 1998).

Taking advantage of the ability to observe aurora B in living cells, we have probed several factors that may affect the dynamics of aurora B. Our results indicated that neither microtubules nor the kinase activity of aurora B is required for the initial localization of aurora B at centromeres during prophase and prometaphase. A similar resistance to nocodazole was reported for the centromeric association of human survivin, a chromosome passenger protein with a similar localization behavior (Skoufias *et al.*, 2000). These results argue against an active transport mechanism that delivers aurora B to the centromeres along kinetochore microtubules.

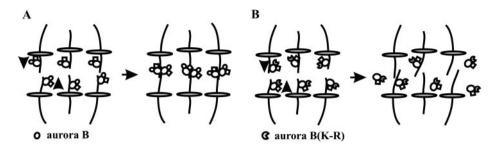


Figure 9. Model showing the possible effects of aurora B on midzone microtubules. Complexes of the chromosomal passenger proteins move toward ends of midzone microtubules (arrows). In cells expressing wild-type aurora B, complexes of chromosomal passenger proteins maintain their microtubule association and promote the formation of midzone microtubule bundles when they are transported to the ends of

microtubules (A). In cells expressing kinase-inactive aurora B, the complex falls off microtubules as they reach the ends, and midzone microtubules remain disorganized (B).

The subsequent relocation of aurora B to interzonal microtubules requires both microtubules and CDK1 deactivation, as indicated by the sensitivity to nocodazole and  $\Delta 90$ cyclin B. However, this process appears to be independent of the kinase activity of aurora B as indicated with cells expressing aurora B(K-R), a kinase inactive mutant of aurora B. Because CDK1 deactivation causes inhibition of the formation of interzonal microtubules (Wheatley et al., 1997), the failure of aurora B relocation may reflect a consequence of the defect in central spindle organization, although it is also possible that the process involves additional steps regulated directly by CDK1. Treatment with nocodazole further suggested that microtubules were required for maintaining the localization of aurora B in the spindle midzone, similar to the behavior of INCENP that may function as an adapter protein for the association of aurora B with interzonal microtubules (Wheatley et al., 2001).

### Functions of Aurora B Kinase in Cell Division

Transient transfection of aurora B(K-R) resulted in chromosome miscongregation and missegregation in NRK cells. The phenotypes are similar to those induced by the inhibition of aurora B expression in *Drosophila* cells (Adams *et al.*, 2001; Giet and Glover, 2001), indicating that the kinase activity of aurora B is crucial to its functions in mitosis. Interestingly, Mad2 was released from the kinetochores despite the failure in chromosomal congregation, suggesting that the checkpoint was short-circuited upstream from the regulation of Mad2-kinetochore association. Because Mad2 release is controlled by the attachment of microtubules at kinetochores (Waters *et al.*, 1998), aurora B may function both to promote interactions of kinetochores with microtubules and/or motors, and to relay such interactions for the regulation of the spindle checkpoint.

The most intriguing observation in the present study is that kinase-inactive aurora B(K-R)-GFP was able to redistribute from the chromosomes to the spindle midzone, but unable to localize stably along midzone microtubules or to organize into an array of concentrated segments on the equatorial plane. This defect was not simply due to the missegregation of chromosomes, because inhibition of topoisomerase II caused a total inhibition of chromosomal segregation while allowing normal cytokinesis in *Caenorhabditis elegans* (Severson *et al.*, 2000), or causing only mislocalized cleavage furrows in mammalian cells (Wheatley *et al.*, 1998). Instead, the results suggest that aurora B(K-R), and associated proteins, fell off the microtubules as they reached the ends. We propose that a substrate of aurora

B is involved in maintaining stable microtubule associations of chromosomal passenger proteins as they are transported along midzone microtubules. The chromosomal passenger proteins in turn promote the organization of midzone microtubule bundles for the localization of cleavage signals (Figure 9). This model explains why cytokinesis fails after initiation in aurora B-ablated cells, and why the depletion of aurora B inhibits the interzonal localization of INCENP (Adams *et al.*, 2001) and the ZEN-4/PAV-KLP motor (Schumacher *et al.*, 1998; Severson *et al.*, 2000; Adams *et al.*, 2001; Giet and Glover, 2001).

To date the only confirmed substrate of aurora B is histone H3, which is known to promote the chromosomal condensation reaction (Hsu *et al.*, 2000; Adams *et al.*, 2001; Giet and Glover, 2001). However, the kinase activity of aurora B was found to reach its peak just after the inactivation of p34<sup>cdc2</sup> (Bischoff *et al.*, 1998), suggesting that the kinase activity is required primarily at prometaphase to anaphase. Therefore, an important task is to identify additional substrates of aurora B that regulate the movement of prometaphase chromosomes and the organization of midzone microtubules.

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#### M. Murata-Hori et al.

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