Bub3 promotes Cdc20-dependent activation of the APC/C in S. cerevisiae

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The spindle checkpoint ensures accurate chromosome segregation by sending a signal from an unattached kinetochore to inhibit anaphase onset. Numerous studies have described the role of Bub3 in checkpoint activation, but less is known about its functions apart from the spindle checkpoint. In this paper, we demonstrate that Bub3 has an unexpected role promoting metaphase progression in budding yeast. Loss of Bub3 resulted in a metaphase delay that was not a consequence of aneuploidy or the activation of a checkpoint. Instead, bub3Δ cells had impaired binding of the anaphase-promoting complex/cyclosome (APC/C) with its activator Cdc20, and the delay could be rescued by Cdc20 overexpression. Kinetochore localization of Bub3 was required for normal mitotic progression, and Bub3 and Cdc20 colocalized at the kinetochore. Although Bub1 binds Bub3 at the kinetochore, bub1Δ cells did not have compromised APC/C and Cdc20 binding. The results demonstrate that Bub3 has a previously unknown function at the kinetochore in activating APC/C-Cdc20 for normal mitotic progression.

Introduction

Anaphase onset is regulated by the anaphase-promoting complex/cyclosome (APC/C), a ubiquitin ligase that when bound to the activator Cdc20 targets substrates for ubiquitination and subsequent degradation (London and Biggins, 2014b). If chromosomes are not attached to spindle microtubules in metaphase, the spindle checkpoint signals to inhibit APC/C substrate ubiquitination. Spindle checkpoint activation depends on the kinetochore localization of the highly conserved checkpoint proteins Bub1, Bub3, Mad1, and Mad2 (London and Biggins, 2014b). Bub3 localizes to kinetochore protein Knl1/Spc105 and brings Bub1 (London et al., 2012; Shepperd et al., 2012; Yamagishi et al., 2012; Primorac et al., 2013). Recent studies in yeast and Caenorhabditis elegans show that Mad1 binds Bub1 and activates Mad2 (London and Biggins, 2014a; Moyle et al., 2014). Ultimately, active Mad2 binds Cdc20 and forms a mitotic checkpoint complex (MCC) with Bub3 and BubR1/Mad3 (Sudakin et al., 2001). The MCC binds the APC/C to block the interaction of Cdc20 with its substrates but still allow Cdc20 autoubiquitination, which promotes Cdc20 turnover during the spindle checkpoint arrest (Pan and Chen, 2004; Braunstein et al., 2007; King et al., 2007; Nilsson et al., 2008; Ge et al., 2009; Herzog et al., 2009; Foe et al., 2011; Mansfeld et al., 2011; Chao et al., 2012).

In addition to their roles in spindle checkpoint signaling, Bub1 and Bub3 facilitate chromosome biorientation, the process in which sister chromatid kinetochores attach to microtubules from opposite spindle poles (van der Horst and Lens, 2014). The Bub1-dependent phosphorylation of histone H2A allows the recruitment of shugoshin to the centromere (Kawashima et al., 2010). Shugoshin (Sgo1 in budding yeast) recruits other proteins needed for kinetochore biorientation, such as the chromosome passenger complex (CPC; van der Horst and Lens, 2014). In budding yeast, the spindle checkpoint proteins are not essential; however, bub1Δ and bub3Δ cells have an increased rate of chromosome missegregation and grow more slowly than wild-type cells (Hoyt et al., 1991; Li and Murray, 1991; Warren et al., 2002). Whether the growth defect in bub1Δ and bub3Δ cells is a consequence of the failure to localize the CPC, or caused by the loss of an additional role of Bub1 and Bub3, is unclear.

Here, we find that in the absence of Bub1 or Bub3, the duration of metaphase is longer than in wild-type cells. This result was unexpected because Bub1 and Bub3 are required for spindle checkpoint signaling, which delays cells in metaphase;
therefore, the expectation would be that loss of Bub1 or Bub3 would not cause a delay. We find that in \( bub3 \Delta \) cells, but not in \( bub1 \Delta \) cells, binding of the APC/C and Cdc20 is defective, and the metaphase delay can be rescued by Cdc20 overexpression. Bub3 kinetochore localization is required for the normal timing of anaphase onset and for normal binding of APC/C and Cdc20, suggesting that Bub3 enhances the binding of APC/C and Cdc20 through its interactions with Cdc20 at the kinetochore. Our observations reveal that Bub3 has a previously undiscovered role, independent of the spindle checkpoint, in activating the APC/C in metaphase.

**Results and discussion**

**Anaphase onset is delayed in cells lacking kinetochore-localized Bub1 or Bub3**

We used time-lapse microscopy to measure the duration of mitosis in wild-type, \( bub1 \Delta \), and \( bub3 \Delta \) cells. The cells expressed Tub1-GFP to monitor spindle assembly and Spc42-mCherry to monitor the separation of spindle pole bodies (SPBs), the yeast equivalent of the centromere (Fig. 1, A–C). In wild-type cells, the duration from SPB separation to anaphase spindle elongation was 25 ± 1 min (mean ± SEM; Fig. 1, A and D). The duration increased to 44 ± 5 min in \( bub1 \Delta \) cells and 42 ± 3 min in \( bub3 \Delta \) cells, a statistically significant difference from wild-type cells (P < 0.0001, Mann–Whitney test; Fig. 1 D). The \( bub1 \Delta \) and \( bub3 \Delta \) cells also showed more cell-to-cell variation in cell cycle length (Fig. S1).

Bub1 and Bub3 have known functions at the kinetochore. However, in *Drosophila melanogaster* and human cells certain checkpoint proteins have kinetochore-independent roles in regulating cell cycle progression (Meraldi et al., 2004; Lopes et al., 2005). Therefore, we tested whether Bub1 and Bub3 kinetochore localization is required for normal mitotic progression. We performed time-lapse microscopy on *spc105-6A* and *bub3R127A/R239A*, two mutants that disrupt the interaction between Bub3 and kinetochore protein Spc105, preventing Bub3 kinetochore localization (London et al., 2012; Primorac et al., 2013). Bub3 does not require Bub1 for kinetochore localization, but Bub1 does require Bub3; therefore, *spc105-6A* and *bub3R127A/R239A* mutants also fail to recruit Bub1 to the kinetochore (Gillet et al., 2004; Primorac et al., 2013). The duration from SPB separation to anaphase onset was delayed in both mutants when compared with wild type, at 43 ± 4 min in *bub1 Δ* and 42 ± 4 min in *bub3 Δ* cells (Fig. 1 D). These results suggest that Bub1 and Bub3 kinetochore localization is required for normal mitotic progression.

At the kinetochore, Bub1 and Bub3 have known roles in signaling the spindle checkpoint and in recruiting Sgo1 (Biggs, 2013). Because Mad2 and Mad3 are required for checkpoint signaling (Li and Murray, 1991), we tested their role in mitotic progression. The *mad2 Δ* and *mad3 Δ* cells progress through mitosis with similar timing as wild-type cells (Fig. 1 D). To test the role of Sgo1 recruitment in the normal timing of anaphase onset, we analyzed a mutant version of Bub1 with the kinase domain deleted, \( Bub1- \Delta K \). The kinase activity of Bub1 is not needed for spindle checkpoint signaling but is needed for Sgo1 localization (Warren et al., 2002; Fernius and Hardwick, 2007; Kawashima et al., 2010). Bub1-\( \Delta K \) localizes to the kinetochore but does not phosphorylate histone H2A for Sgo1 localization (Fernius and Hardwick, 2007). We find that \( bub1- \Delta K \) cells have a delay in anaphase onset (Fig. 1 D). Furthermore, *sgo1 Δ* cells also have a delay in anaphase onset (Fig. 1 D). The results indicate that Bub1 kinase activity and Sgo1 localization are also required for the normal timing of anaphase onset.

**The delay in cell cycle progression in \( bub3 \Delta \) cells is not caused by activation of the G2/M, DNA damage, or spindle position checkpoints**

The increased duration of mitosis in \( bub1 \Delta \) and \( bub3 \Delta \) cells could be a result of the activation of a checkpoint, such as the G2/M, DNA damage, and spindle position checkpoints. We deleted genes required for each checkpoint in combination with \( bub3 \Delta \) to determine whether the mutation rescues the delay in anaphase onset. In \( bub3 \Delta \) *chk1 Δ*, *bub3 Δ swe1 Δ*, and *bub3 Δ bub2 Δ* cells, anaphase onset was delayed in comparison to wild-type cells, similar to the delay in *bub3 Δ* cells (Fig. 1 E). The *bub3 Δ mad2 Δ* cells also have a delay in anaphase onset. The results suggest that the loss of Bub3 does not delay anaphase onset by activating the G2/M, DNA damage, spindle position, or spindle checkpoints.

**Metaphase is delayed in cells lacking Bub1 or Bub3**

The budding yeast spindle forms at the end of S phase (Biggs, 2013). Therefore, a cell cycle delay from SPB separation to anaphase onset could be a result of G2 or metaphase delay. To determine whether \( bub1 \Delta \) and \( bub3 \Delta \) cells were delayed in metaphase, we monitored cells released from a metaphase arrest and measured the duration to anaphase onset. Cells are reversibly arrested in metaphase by placing the APC/C activator Cdc20 under the control of the methionine-repressible MET3 promoter (*PMET3CDC20*; Uhlmann et al., 2000). The *PMET3CDC20, bub1 Δ PMET3CDC20, bub3 Δ PMET3CDC20*, and *bub1-ΔK PMET3CDC20* cells arrested in metaphase in medium containing methionine were released from the arrest by washing out the medium and adding medium lacking methionine. The *PMET3CDC20* cells enter anaphase in 22 ± 1 min. The *bub3 Δ PMET3CDC20, bub1 Δ PMET3CDC20*, and *bub1-ΔK PMET3CDC20* cells have a statistically significant delay in anaphase onset compared with wild-type cells and enter anaphase after 36 ± 4, 33 ± 3, and 31 ± 3 min, respectively (Fig. 1 F).

Furthermore, we measured the duration that kinetochores were biooriented, or attached to opposite spindle poles in \( bub1 \Delta \) and \( bub3 \Delta \) cells. To assess bioorientation, we monitored the separation of a tagged kinetochore protein Mtw1-GFP. Biooriented sister chromatid kinetochores are pulled ~0.8 µm apart (Joglekar et al., 2008). In wild-type cells, kinetochores are biooriented for 23 ± 1 min. In \( bub1 \Delta \) and \( bub3 \Delta \) cells, the duration that kinetochores are biooriented is increased to 42 ± 3 min and 44 ± 3 min, respectively (Fig. 1 G). The duration is also increased in *bub1-ΔK, sgo1 Δ*, and *bub3R127A/R239A* cells. In summary,
3 of 9

Figure 1. Anaphase onset is delayed in the absence of kinetochore-localized Bub1 or Bub3. (A–C) Time-lapse images of wild-type, bub1Δ, and bub3Δ cells with Spc42-mCherry and Tub1-GFP. Bars, 2 μm. (D–G) Plots of the mean time (in minutes) for each genotype. At least 50 cells were counted per genotype. Error bars are the SEM. Asterisks indicate a statistically significant difference compared with wild-type cells (*, \( P < 0.0001 \), Mann-Whitney test). (D and E) Time from SPB separation to anaphase onset. (F) Time from metaphase to anaphase onset in strains with CDC20 under the repressible MET3 promoter. (G) Time from Mtw1-GFP foci separation to anaphase onset.

the results suggest that bub1Δ, bub3Δ, bub1ΔK, sgo1Δ, and bub3R127A/R239A cells are delayed in metaphase with bioriented sister chromatid kinetochores.

**APC/C substrates accumulate in bub1Δ and bub3Δ cells**

We monitored the timing of APC/C substrate degradation. Cells were arrested in G1 with α factor and released from the arrest, and then the protein was isolated every 20 min for one cell cycle, adding α-factor back to the cells to ensure that they do not go through another cell cycle. The normal timing of degradation of three APC/C substrates Pds1-9Myc, Clb2, and Clb5 was delayed in bub1Δ and bub3Δ cells compared with wild-type cells (Fig. 2, A–H). In bub1Δ and bub3Δ cells, the APC/C substrates were present at times when they should be degraded, such as in G1 (0, 120, and 140 min), as scored by the presence of unbudded cells without a spindle and with Sic1-3HA, a Cdk inhibitor that is present in G1 (Fig. 2, A–H). These results demonstrate that APC/C substrates are not degraded with normal timing in bub1Δ and bub3Δ cells.
We dissected the tetrads from the BUB3/bub3::LEU2 heterozygote on an agar pad containing medium lacking leucine such that only the bub3::LEU2 spores will germinate (Fig. 3 A). We performed the same analysis with wild-type cells with an integrated vector containing LEU2. In both wild-type and bub3Δ cells, the first cell cycle upon germination was slower at all stages than the next cell cycles, so we averaged the second, third, and fourth cell cycles, counting ≥50 cell divisions for each genotype. Anaphase onset was indeed significantly delayed in bub3Δ cells at 33 ± 1 min compared with 24 ± 1 min in wild-type cells (P < 0.0001, Mann–Whitney test; Fig. 3 B). Similar analysis from the spores of the Bub3/bub3R127A/R239A heterozygote showed that the bub3R127A/R239A cells also have a delay in anaphase onset similar to bub3Δ cells (Fig. 3 B). The mean time of metaphase in the newly germinated bub3Δ and bub3R127A/R239A cells is somewhat decreased when compared with the corresponding log phase–grown cultures, and there is less cell-to-cell variation, suggesting that some cells in the log phase cultures were likely aneuploid (Figs. 3 B and S2). Nonetheless, the results demonstrate that bub3Δ and bub3R127A/R239A cells have a metaphase delay that is not a consequence of aneuploidy. Unfortunately, we could not perform the same analysis for bub1Δ because very few spores of the BUB1/bub1Δ heterozygote germinated, suggesting that two copies of BUB1 are required for...
normal chromosome segregation in meiosis. We conclude that kinetochore-localized Bub3 promotes the normal duration of metaphase.

The binding of Cdc20 to the APC/C is impaired in the absence of Bub3

Because the metaphase activation of the APC/C requires the association of APC/C with Cdc20 (Visintin et al., 1997; Lorca et al., 1998), we used coimmunoprecipitation (co-IP) to examine the interaction between the APC/C and Cdc20 in wild-type, bub1Δ, bub3Δ, sgo1Δ, and bub1-ΔK cells transitioning into anaphase. Cells were arrested in G1 with α-factor, released from the arrest, and then collected at the transition into anaphase. The cells express CDC23-3HA, a tagged component of the APC/C, allowing the IP of the APC/C with anti-HA antibodies. Antibodies against Cdc20 detected the amount of Cdc20 in the IP. In the no tag control, Cdc23 and Cdc20 were not detected in the IP, showing that the IP was specific to tagged Cdc23 (Fig. 4 A).

Surprisingly, there was a severe reduction of Cdc20 associated with the APC/C in bub3Δ cells compared with wild-type cells (Fig. 4, A and B). Cdc20 levels are not reduced in the input, suggesting that the diminished level of Cdc20 coimmunoprecipitated in bub3Δ cells cannot be explained by reduced Cdc20 protein levels (Fig. 4, A and B). Furthermore, similar levels of Cdc23 were immunoprecipitated from each strain (Fig. 4 C). There was a 93% reduction in the ratio of Cdc20 in the IP to Cdc20 in the input (Cdc20input/Cdc20input) in bub3Δ compared with wild type (Fig. 4 B). In contrast to bub3Δ cells, the bub1Δ, sgo1Δ, and bub1-ΔK cells do not have reduced levels of Cdc20 bound to APC/C component Cdc23 (Fig. 4, A and B). These results suggest that the binding of APC/C and Cdc20 is impaired in the absence of Bub3 but not Bub1 or Sgo1.

We also assessed whether kinetochore localization of Bub3 was needed for normal binding of the APC/C with Cdc20. In the bub3R127A/R239A mutant that does not localize to the kinetochore, the amount of Cdc20 immunoprecipitated with APC/C component Cdc23 is also severely reduced when compared with wild type (Fig. 4 A and B). These results suggest that kinetochore-localized Bub3 is required for the normal binding of the APC/C and Cdc20.

The overexpression of Cdc20 rescues the metaphase delay of bub3Δ cells but not bub1Δ, bub1-ΔK, or sgo1Δ cells

Considering that APC/C and Cdc20 binding is impaired in bub3Δ cells, we tested whether increasing Cdc20 levels could suppress the metaphase delay. We overexpressed CDC20 in wild-type, bub3Δ, bub1Δ, bub1-ΔK, and sgo1Δ cells and measured the duration from SPB separation to anaphase onset. Cells with an integrated vector with CDC20 under the galactose-inducible GAL1 promoter (PgalCDC20) were grown in galactose medium to overexpress CDC20. Cells grow slower in galactose medium than glucose medium, so we compared cells with PgalCDC20 to cells with an integrated empty vector, both growing in galactose medium. In wild-type cells, the duration from SPB separation to anaphase onset is 37 ± 2 min in cells overexpressing CDC20, which is similar to the 39 ± 2 min in cells not overexpressing CDC20 (Fig. 5 A). In contrast, the duration from SPB separation to anaphase onset is shorter in bub3Δ cells overexpressing CDC20 at 41 ± 2 min compared with bub3Δ cells not overexpressing CDC20 at 56 ± 3 min (Fig. 5 A). In the bub3Δ cells overexpressing CDC20, there was an increase in the number of dead cells when compared with the bub3Δ cells not overexpressing CDC20, suggesting that the metaphase delay...
may allow increased cell survival. We conclude that overexpression of Cdc20 rescues the metaphase delay in bub3Δ cells.

In contrast to bub3Δ cells, the duration of metaphase is not shortened in bub1Δ, bub1ΔK, or sgo1Δ cells overexpressing CDC20 when compared with the respective genotype that does not overexpress CDC20 (Fig. 5 A). These results and the results from the co-IP experiments suggest that although bub1Δ and bub3Δ cells both have a metaphase delay, the delay may be caused by different mechanisms. In the absence of Bub1, a failure to localize Sgo1 and the CPC to the kinetochore may lead to a metaphase delay. In support of this model, a previous study showed that the growth defect of sgo1Δ cells can be rescued by a mutant of Sli15, sli15(ΔNT), which bypasses the requirement for CPC clustering between sister kinetochores (Campbell and Desai, 2013). However, in the absence of Bub3, the metaphase delay can be rescued by CDC20 overexpression and is therefore likely caused by the impaired interaction between the APC/C and Cdc20.

**Bub3 and Cdc20 colocalize to the kinetochore**

Because Bub3 kinetochore localization is required for the proper timing of anaphase onset and for full APC/C-Cdc20 binding, we examined the localization of Bub3 and Cdc20 at the kinetochore. Although Cdc20 localizes to kinetochores in many organisms, the localization has not been demonstrated in budding yeast (London and Biggins, 2014b). In intact budding yeast cells, kinetochores are clustered near the SPB, and individual kinetochores cannot be resolved. To examine individual kinetochores, we used mitotic chromosome spreads from protoplasts of cells expressing kinetochore protein Ndc10-6HA, Bub3-EGFP, and 12Myc-Cdc20. Immunofluorescence was performed with anti-HA, anti-GFP, and anti-Myc antibodies. As an example of a protein known to localize to the kinetochore, we show the colocalization of Bub3 and Ndc10 in mitotic chromosome spreads (Fig. 5 B). We find that Cdc20 and Bub3 colocalize, and Cdc20 and Ndc10 colocalize (Fig. 5, C and D). In contrast to Bub3, Cdc20 does not localize to all kinetochores at a single time point, suggesting that the localization of Cdc20 to a kinetochore may be transient. These results demonstrate that Bub3 colocalizes with Cdc20 at the kinetochore.

**Conclusion**

This study revealed an unexpected role for Bub3 in activating the APC/C for the normal progression of metaphase. In the absence of Bub3, metaphase is delayed, and binding of APC/C and Cdc20 is impaired. Intriguingly, Bub3 must localize to the kinetochore to prevent the metaphase delay, and Bub3 and Cdc20 colocalize at the kinetochore. Our results suggest a new model for the coordination of the metaphase to anaphase transition with kinetochore-localized Bub3 promoting Cdc20 and APC/C binding.

The role of Bub3 in activating APC/C-Cdc20 seems contradictory to its known role in inhibiting APC/C activity during spindle checkpoint signaling. However, the results are consistent when considering that although substrate ubiquitination is blocked during checkpoint signaling, APC/C-Cdc20 autoubiquitinites (Pan and Chen, 2004; Braunstein et al., 2007; King et al., 2007; Nilsson et al., 2008; Ge et al., 2009; Herzog et al., 2009; Roe et al., 2011; Mansfeld et al., 2011; Chao et al., 2012). In vitro experiments using purified budding yeast checkpoint proteins from cells with an active spindle checkpoint show that Bub3-Mad3 stimulates the binding of APC/C and Cdc20 to promote Cdc20 autoubiquitinitation (Foster and Morgan, 2012). Our in vivo experiments suggest that Bub3 enhances APC/C and Cdc20 binding for normal metaphase progression, independent of spindle checkpoint signaling. We propose that Bub3 normally promotes APC/C and Cdc20 binding; however, when the spindle checkpoint is active, the association of Bub3 with the MCC blocks the interaction...
of the APC/C with its substrates but still allows Cdc20 auto-
ubiquitination. In summary, Bub3’s role in promoting binding
of APC/C and Cdc20 is important for normal cell cycle progres-
sion and spindle checkpoint signaling.

Materials and methods
Budding yeast strains
All strains are W303 derivatives and are described in Table S1. Deletion
and tagged strains were made by PCR amplification and transformation of
marked cassettes (Longtine et al., 1998; Janke et al., 2004). For deletions,
primers with homology overhangs flanking the open reading frame were
used to amplify the cassette. For tagged strains, primers flanking the stop
codon of the target genes were used. The P°CDC20 strain was made
by PCR amplifying the CDC20 ORF and cloning it behind the GAL1,10
promoter in pRS306 and then integrating the plasmid into the URA3 locus.
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used in this study.

Microscope image acquisition
Cells were imaged with an inverted microscope (Ti-E; Nikon) equipped
with a 60x objective (Plan Apochromat NA 1.4 oil), a Lambda 10–3 opti-
cal filter changer and SmartShutter, GFP, and mCherry filters, and a
charge-coupled device camera (CoolSNAP HQ2; Photometrics) at 25°C.
Z stacks of 5–6 sections were acquired in 5–10-min intervals for 12–16 h
with a combination of 12.5% neutral density filter and 6.25% neutral den-
sity filter and exposure times of 50–900 ms. Z stacks were combined into
a single maximum-intensity projection using the maximum-intensity projec-
tion function in NIS-Elements software (Nikon). Images were cropped to
the same size to show individual cells dividing over time with GFP- and
mCherry-tagged proteins.

Time-lapse microscopy
Cells were incubated at 30°C to mid–log phase. Wild-type control cells and
mutant cells were put on two side-by-side agar pads containing synthetic

Figure 5. Anaphase onset timing is restored
by CDC20 overexpression in bub3Δ cells but not bub1Δ, sgo1Δ, or bub1-1K cells, and Bub3
colocalizes with Cdc20 at the kinetochore.
(A) Plot of the mean time (minutes) from SPB
separation to anaphase onset when CDC20 is
overexpressed by an inducible GAL1 promoter.
At least 50 cells were counted per genotype.
Error bars are the SEM. Asterisks indicate statis-
tically significant difference compared with wild
type [WT] + GAL1-CDC20 (*, P < 0.0001,
Mann–Whitney test). (B–D) Immunofluorescence
of mitotic chromosome spreads showing Bub3-
EGFP and Ndc10-6HA (B), Bub3-EGFP and
12Myc-Cdc20 (C), and Ndc10-6HA and 12Myc-
Cdc20 (D). Bars, 1 μm.
complete medium and imaged together for each video at 25°C. Wild-type cells were checked for the same mean cell cycle duration for each experiment to ensure that conditions were the same in each experiment. Images were acquired in 5-min intervals for 12–16 h with exposure times of 50–900 ms according to fluorescence intensity. In the galactose induction experiments, cells were incubated at 30°C to mid-log phase, and galactose was added to the medium for 3 h before the imaging. Time-lapse microscopy was analyzed for cell cycle duration, and data were graphed in Prism (GraphPad Software). The significance was calculated using the Mann–Whitney test.

**Spore imaging**

A heterozygous bub3Δ strain (LY1897) was constructed by deleting one copy of BUB3 in a wild-type diploid strain (LY1877). A heterozygous bub3Δ:bub3Δ strain (LY2094) was constructed by mating a bub3Δ:bub3Δ haploid strain (LY1894) with a wild-type haploid strain (LY1930). Cells were sporulated, and 16–20 tetrads were dissected, arranged on –leucine agarose pads, and mounted on a coverglass surrounded by a 1-h interval for 3 h followed by 10-min intervals for 10–12 h to monitor germination and the first four divisions of the spores.

**Mitosis time course protein isolation**

For the mitosis time course, wild-type (LY1959), bub3Δ (LY1990), and bub1Δ (LY1988) cells were incubated in YPD (yeast, peptone, dextrose) at 30°C until saturation, diluted 1:20 into 40 ml of fresh YPD, and incubated at 30°C for 3 h. 5 µM α-factor was added for 2 h at 25°C, and cells were checked for >85% shmoos. α-Factor was washed out using 20 ml of fresh YPD three times at room temperature, and cells were resuspended in 40 ml YPD and incubated at 25°C. After 90 min, α-factor was added back to the culture to prevent the cells from going into the next cell cycle. The cell culture (5 ml) was harvested and snap frozen in liquid nitrogen every 20 min after resuspension, and a sample of cells were fixed using 4% paraformaldehyde to check for spindle morphology.

**APC/C-Cdc20 co-IP**

For the co-IP of Cdc20 and Cdc23-3HA, cells were grown in YPD to saturation overnight at 25°C, diluted into 20 ml of fresh YPD and incubated for 3 h at 25°C. 10 µM α-factor was added for 1.5 h at 25°C. α-Factor was washed out three times using 20 ml YPD, and cells were resuspended in 20 ml YPD at 25°C. Cells were harvested when they were transitioning into anaphase. Cell pellets were divided into two tubes and resuspended in lysis buffer + 1 mM PMSF + protease inhibitor tablet (Roche). Glass beads were added to each tube, and cells were vortexed at 4°C for 7 × 1 min maximum speed, with 1 min on ice in between each vortex. Cell lysates were centrifuged at 13,200 rpm for 5 min. 50–100 µl of cleared lysate was diluted 1:5 using 3×SDS reducing sample buffer and boiled at 95°C for 5 min. 10 µl was loaded for Western blot. 1 µg of mouse α-MyoD (Roche) was mixed with the remaining cleared lysate and incubated on ice for 20 min. The mixture was then preclarified for 2 min at 13,200 rpm, and the supernatant was mixed with 35 µl protein G Dynabeads and incubated on a rotation platform at 4°C for 1 h. The beads were then washed three times with 500 µl Cdc20 bead buffer (200 mM NaCl, 50 mM Tris-Cl, pH 7.4, 50 mM NaF, 5 mM EDTA, 0.1% NP-40, and 1 mM DTT) and two times with 500 µl low salt kinase buffer (10 mM NaCl, 20 mM Hepes-KOH, pH 7.4, and 5 mM MgCl2) the last wash was transferred to a new microcentrifuge tube, and 20 µl of SDS reducing sample buffer was used to resuspend the beads. The beads were then boiled at 95°C for 5 min, and 10 µl supernatant was loaded for Western blotting. Cdc20 and Cdc23-3HA were blotted on two separate blots because of their close sizes.

**Western blotting and quantification**

For the Western blots for the mitosis time course and the co-IP, the membrane was cut according to corresponding size of the proteins and incubated with mouse α-myc (1:500, 9E10, Roche), mouse α-HA (1:1,000, 12CA5, Roche), rabbit α-Cdk2 (1:1,000, 9E10, sc9071, Santa Cruz Biotechnology, Inc.), rabbit α-Cdk28 (1:3,000, PSTAIRE, sc-53, Santa Cruz Biotechnology, Inc.), goat α-Clb5 (1:1,000, y-19, sc-6704, Santa Cruz Biotechnology, Inc.), and goat α-Cdc20 (1:500, yC-20, sc-6731, Santa Cruz Biotechnology, Inc.). PageRuler Prestained Protein Ladder (Thermo Fisher Scientific) was used, and migration of the bands was indicated in each blot. For the mitotic time course experiments, the secondary antibodies used were donkey anti–rabbit HRP (GE Healthcare), sheep anti–mouse HRP (GE Healthcare), and donkey anti–goat HRP (sc-2033, Santa Cruz Biotechnology, Inc.) at 1:5,000. ECL substrates were added to the washed membrane, and exposure was performed using Classic blue sensitive x-ray film on Konica Minolta X-ray film processor. For the co-IP experiments, the secondary antibodies used were IRDye 800CW donkey anti–goat IgG (C41105-01, LCiCOR Biosciences) and IRDye 680RD donkey anti–rabbit IgG (C41009-02, LC-COR Biosciences) at 1:15,000. Blots were processed according to the Li-COR Western blot protocol and imaged using Odyssey CLX (Li-COR Biosciences). All quantifications were processed using Image Studio software (Li-COR Biosciences) and graphed in Prism.

**Immunofluorescence**

Mitotic cells were fixed with 4% paraformaldehyde for 5 min at room temperature. Cells were washed with PBS, and cell wall was digested with 1 mg/ml zymolyase (Zymo Research) buffered with 1 M sorbitol. Harvested protoplasts were washed with 1 M sorbitol, applied on a slide, and hypronot solution (0.01 M sorbitol and 0.1% Triton X-100) was added. The slides were washed with PBS and blocked with 5% bovine serum albumin (Sigma-Aldrich) for 1 h at 25°C. For detection of Bub3-GFP, Ndc10-6HA, and 12Myc-Cdc20, the primary antibodies chicken anti-GFP (1:600, Novus Biologicals), mouse anti-HA (1:500, Santa Cruz Biotechnology, Inc.), and rabbit anti-Myc (1:500, Santa Cruz Biotechnology, Inc.) were applied, respectively, for 15 h at 4°C. Slides were washed twice with PBS, once with PBS containing 0.1% Tween 20, and once with PBS. Secondary antibodies, Alexa Fluor 488–conjugated goat anti–chicken (1:200, Molecular Probes), Alexa Fluor 594–conjugated goat anti–mouse (1:200, Molecular Probes), and Alexa Fluor 394–conjugated goat anti–rabbit (1:200, Molecular Probes) were applied for 1 h at 25°C for imaging. Slides were washed three times with PBS, and DNA was stained with 1 mg/ml DAPI.

**Online supplemental material**

Fig. S1 shows the variability in the duration of anaphase onset of spindle checkpoint mutants using time-lapse microscopy. Fig. S2 shows the variability in the duration of anaphase onset of newly germinated spores. Table S1 shows strains used in this study. Online supplemental material is available at http://www.jcb.org/cgi/content/full/jcb.201412036/D1. Additional data are available in the JCB DataViewer at http://dx.doi.org/10.1083/jcb.201412036.dv.

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**References**


Figure S1. **Variability in metaphase duration shown in a scattered dot plot.** The time from SPB separation to anaphase onset in spindle checkpoint mutants and mutants that fail to localize proteins to the kinetochore. Each dot represents one data point collected from time-lapse microscopy, and 50 data points were collected for each strain (means ± SEM). KD, knockdown; WT, wild type.

Figure S2. **Variability in metaphase duration is reduced in spore imaging.** Scattered dot plot showing the time from SPB separation to anaphase onset in spore imaging time-lapse microscopy of bub3Δ and bub3R127A/R239A. Each dot represents one data point collected from time-lapse microscopy, and ≥50 data points were collected for each strain. WT, wild type.
### Table S1. Strains used in this study

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