Role of Budding Yeast Cdc5 in Cell Division: Spatial and Temporal Regulation of M Phase Progression by Polo-Like Kinase

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Members of the polo subfamily of protein kinases have been identified in various eucaryotic organisms from budding yeast to mammals and appear to play pivotal roles in cell division and proliferation. Polo-like kinases (Plks) include mammalian Plk1, Snk, and Fnk/Prk, Xenopus laevis Plx1, Drosophila melanogaster polo, Schizosaccharomyces pombe Plo1, and Saccharomyces cerevisiae Cdc5 [1]. Budding yeast S. cerevisiae has become a excellent model system to study the function of the Plk because the organism contains only one Plk gene CDC5 whereas mammals express at least three different Plks; Plk1, Snk, and Fnk. Among these, Plk1 and Cdc5 functions during M phase but Snk and Fnk seems to act at early stage of cell division prior to M phase. In addition to a high degree of sequence similarity in the kinase domain, polo kinases contain a strikingly conserved motif termed "polo-box" in the noncatalytic C-terminal domain. It is apparent from genetic and biochemical analyses that polo kinases regulate diverse cellular events at various stages of M phase, including centrosome maturation, bipolar spindle formation, activation of Cdc2 through Cdc25C phosphatase at at the G₂/M transition, DNA damage checkpoint adaptation, activation of the anaphase-promoting complex, and cytokinesis in various eucaryotic systems [1]. The roles of these kinases are likely to be conserved among evolutionarily distant eucaryotic cells. It was demonstrated that murine Plk is a functional homolog of S. cerevisiae Cdc5 and that the polo box is required for the ability of Plk to functionally complement the cdc5-1 defect by targeting the catalytic activity of the enzyme to specific subcellular location [2]. Without impairing kinase activity, conservative mutations in the polo box abolish the ability of Cdc5 to functionally complement the defect associated with a cdc5-1 temperature-sensitive mutation, to localize to the spindle pole body (SPB) and cytokinetic neck filaments, and to induce elongated cells with ectopic septin ring structures. Consistent with the polo box-dependent subcellular localization, the C-terminal domain of Cdc5, but not its polo box mutant, is sufficient for subcellular localization, and its overexpression appears to inhibit cytokinesis [7]. These data provide evidence that the polo box is required to direct Cdc5 to both SPB and mother-bud neck through binding to components of specific subcellular locations. To provide new insights into the function of budding yeast polo kinase Cdc5p, novel temperature-sensitive cdc5 mutants were generated by mutagenizing the C-terminal domain. At a semipermissive temperature, the cdc5-3 mutant exhibited a synergistic bud elongation, an indicative of G2/M arrest, and growth defect with loss of HSL1, a component important for

normal G2/M transition. Loss of SWE1, which inactivates the budding yeast Cdk1 homolog Cdc28p by inhibitory phosphorylation, suppressed the cdc5-3 hsl1 Δ defect, suggesting that Cdc5p functions at a point upstream of Swelp. Thus, Cdc5p contributes to the activation of the Swelp-dependent Cdc28p/Clb pathway at G2/M transition [4,5]. Intact polo-box of Cdc5 is essential to localization at mother-bud neck containing septin filament. Overexpression of the C-terminal domain of Cdc5 $(cdc5\Delta N)$, but not the corresponding polo-box mutant, resulted in connected cells. These cells shared cytoplasm with incomplete septa, and possessed aberrant septin ring structures. Moreover, depletion of Cdc5 function leads to an arrest in cytokinesis. Provision of additional copies of endogenous CDC5 remedied this phenotype, suggesting a dominant-negative inhibition of cytokinesis [6]. The polo-box-dependent interactions between Cdc5 and septins (Cdc11 and Cdc12) suggest that direct interactions between cdc5ΔN and septins regulate cytokinetic pathways after completion of chromosome separation. Bbp1p, isolated as a polo-box interacting protein by a yeast two-hybrid screen, localizes to the periphery of the central plaque of the SPB and plays an important role in SPB duplication. Similarly, Cdc5p localized to the cytoplasmic periphery of the SPB. In vitro binding studies showed that Cdc5p interacted with the N-terminal domain of Bbp1p (Bbp1p Δ C). In addition, Bbp1p was required for proper localization of Cdc5p to the SPB. The C-terminal coiled-coil domain of Bbp1p (Bbp1p(243-385)), which is crucial for both the homodimerization and the SPB localization, could target the localization-defective Cdc5p Δ C to the SPB. Consistent with this observation, expression of CDC5 Δ C-BBP1(243-385) under CDC5 promoter control partially complemented the cdc5Δ defect. These data suggest that BbplpΔC interacts with the polo-box domain of Cdc5p, and this interaction is critical for the subcellular localization and mitotic functions of Cdc5p [3]. Thus, Bbp1 and septins provide a spatial compartment of Cdc5 functions by mediating the polo-box-dependent localization of Cdc5 at both SPB and cytokinetic neck filament where Cdc5 timely coordinates various stages of M phase including G2/M transition, mitotic exit, and cytokinesis.

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