

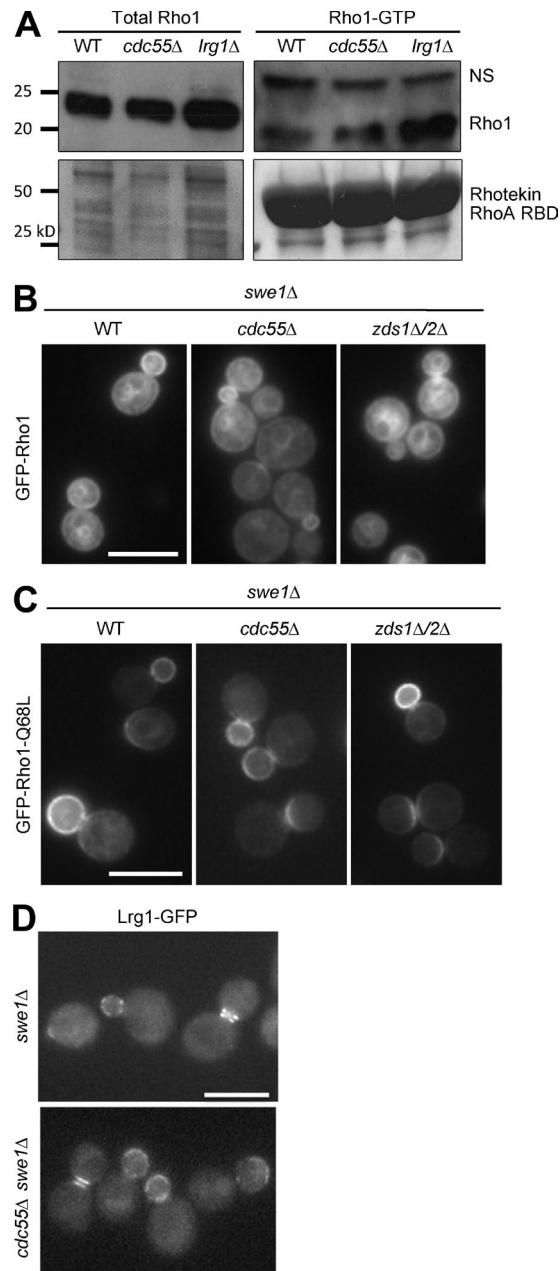
Jonasson et al., <http://www.jcb.org/cgi/content/full/jcb.201508119/DC1>

Figure S1. Activity and localization of Rho1 was not significantly affected in PP2A mutants, associated with Fig. 3. (A) Rho1 activity was does not appear to be significantly affected by loss of *CDC55*. The total cellular amount of active Rho1 (Rho1-GTP) was analyzed in the indicated strains by a previously established pull-down assay for active RhoA/Rho1 using GST-rhotekin RhoA binding domain (RBD) beads [Kimura et al., 2000; Kono et al., 2008; Yoshida et al., 2009]. Rho1 was detected by Western blot analysis using a custom-made rabbit anti-Rho1 antibody (Y1486). Ponceau staining is shown as a loading control for GST-rhotekin RBD. (B and C) Rho1 is localized to the bud cortex in the absence of PP2A subunits. GFP-Rho1 (B) and GFP-Rho1Q68L (C) were expressed under the *RHO1* promoter from the centromeric plasmids, and images were taken in living cells at RT. Note that these experiments were performed in a *swe1Δ* background. Bars, 5 μm. (D) Bud tip and bud neck localization of Lrg1 was not detectably affected by deletion of *CDC55*. In *swe1Δ* strains, Lrg1-GFP was expressed at the endogenous locus, and live cells were imaged at RT. WT, wild type.

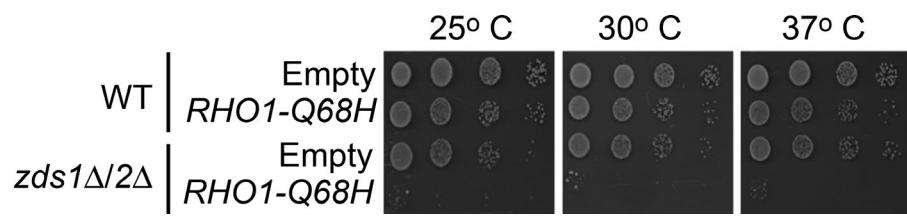


Figure S2. ***zds1Δ zds2Δ* cells are sensitive to expression of active Rho1, associated with Fig. 4.** Serial dilutions of yeast strains expressing *RHO1-Q68H* (active Rho1) under the control of the *RHO1* promoter were spotted on SC-URA plates and grown at different temperatures for 3 d before the plates were imaged. WT, wild type.

Table S1. Yeast strains used in this study

Strain	Genotype	Source
PY3295	MA _T a his3Δ1 leu2Δ0 ura3Δ0 met15Δ0	Open Biosystems
PY3296	MA _T a his3Δ1 leu2Δ0 ura3Δ0 lys2Δ0	Open Biosystems
SY1327	MA _T a rom1Δ::KanMX6 rom2Δ::KanMX6 tus1Δ::KanMX6 + <CEN-ROM2-URA3>	Yoshida et al., 2009
SY1854	MA _T a zds1::HIS3MX6::pGAL1-3xHA-ZDS1-GFP::KanMX6	This study
VR334	MA _T a cdc55::CDC55-13xMyc::KanMX6	This study
VR371	MA _T a cdc55::CDC55-13xMyc::KanMX6 zds1::KanMX6 zds2::HIS3MX6	This study
YSY4	MA _T a ade2 his3 leu2 trp1 ura3 rho1Δ::HIS3 ade3::RHO1::LEU2 (YOC764)	Saka et al., 2001
YSY6	MA _T a ade2 his3 leu2 lys2 trp1 ura3 rho1Δ::LYS2 ade3::rho1-2::HIS3 (YOC729)	Saka et al., 2001
YSY74	MA _T a cdc55Δ::KanMX6	Rossio and Yoshida, 2011
SY1763	MA _T a zds1Δ::KanMX6 zds2Δ::HIS3MX6	Rossio and Yoshida, 2011
VR368	MA _T a pph21Δ::KanMX6	Open Biosystems
VR406	MA _T a pph22Δ::KanMX6	Open Biosystems
VR408	MA _T a pph21Δ::KanMX6 pph22Δ::KanMX6	This study
VR403	MA _T a sit4Δ::KanMX6	Open Biosystems
SY1529	MA _T a lrg1Δ::KanMX6	Yoshida et al., 2009
SY1390	MA _T a swe1Δ::LEU2	Rossio and Yoshida, 2011
YSY71	MA _T a swe1Δ::LEU2 cdc55Δ::KanMX6	Rossio and Yoshida, 2011
YSY80	MA _T a swe1Δ::LEU2 zds1Δ::KanMX6 zds2Δ::HIS3MX6	Rossio and Yoshida, 2011
SY1630	MA _T a zds1::ZDS1-GFP::HIS3MX6	Rossio and Yoshida, 2011
VR63	MA _T a zds1::ZDS1-GFP::HIS3MX6 zds2Δ::HIS3MX6	Rossio and Yoshida, 2011
SY1856	MA _T a zds1::Zds1ΔC400-GFP::HIS3MX6	Rossio and Yoshida, 2011
VR58	MA _T a zds1::Zds1ΔC400-GFP::HIS3MX6 zds2Δ::HIS3MX6	Rossio and Yoshida, 2011
SY1857	MA _T a zds1::Zds1ΔC800-GFP::HIS3MX6	Rossio and Yoshida, 2011
VR60	MA _T a zds1::Zds1ΔC800-GFP::HIS3MX6 zds2Δ::HIS3MX6	Rossio and Yoshida, 2011
SY1779	MA _T a cdc55::CDC55-GFP::HIS3MX6	Rossio and Yoshida, 2011
SY1811	MA _T a cdc55::Cdc55-GFP-NES::KanMX6	Rossio and Yoshida, 2011
SY1808	MA _T a cdc55::Cdc55-GFP-NLS::KanMX6	Rossio and Yoshida, 2011
SY1793	MA _T a cdc55::CDC55-GFP::HIS3MX6 zds1::KanMX6 zds2Δ::HIS3MX6	Rossio and Yoshida, 2011
VR21	MA _T a cdc55::CDC55-GFP-NES::KanMX6 zds1::KanMX6 zds2Δ::HIS3MX6	Rossio and Yoshida, 2011
YPH499	MA _T a ade2 his3 leu2 lys2 trp1 ura3	Sikorski and Hieter, 1989
YOC2581	MA _T a ade2 his3 leu2 lys2 trp1 ura3 zds1Δ::cgHIS3	Sekiya-Kawasaki et al., 2002
SY1762	MA _T a ade2 his3 leu2 lys2 trp1 ura3 zds1Δ::cgHIS3 zds2Δ::KanMX6	This study
YSY123	MA _T a mpk1Δ::KanMX6	This study
VR160	MA _T a mpk1Δ::KanMX6 cdc55Δ::KanMX6	This study
VR194	MA _T a mpk1Δ::KanMX6 zds1Δ::KanMX6 zds2Δ::HIS3MX6	This study
YSY83	MA _T a swe1Δ::LEU2 lrg1::LRG1-3xHA::HIS3MX6	This study
YSY85	MA _T a swe1Δ::LEU2 cdc55Δ::KanMX6 lrg1::LRG1-3xHA::HIS3MX6	This study
YEJ56	MA _T a lrg1::pGAL1-3xHA-LRG1::HIS3MX6	This study
YEJ96	MA _T a cdc55Δ::KanMX6 lrg1::pGAL1-3xHA-LRG1::HIS3MX6	This study
YEJ968	MA _T a swe1Δ::LEU2 cdc55Δ::HIS3MX6	This study
YEJ972	MA _T a swe1Δ::LEU2 cdc55Δ::HIS3MX6 lrg1Δ::KanMX6	This study
YEJ900	MA _T a swe1Δ::LEU2 sac7::SAC7-3xHA::HIS3MX6	This study
YEJ904	MA _T a swe1Δ::LEU2 cdc55Δ::KanMX6 sac7::SAC7-3xHA::HIS3MX6	This study
YEJ163	MA _T a his3Δ200 leu2Δ1 ura3-3Δ cim3-1, ts (CMY763)	Ghislain et al., 1993
YEJ815	MA _T a CIM3 sac7::SAC7-3xHA::HIS3MX6	This study
YEJ803	MA _T a CIM3 cdc55Δ::KanMX6 sac7::SAC7-3xHA::HIS3MX6	This study
YEJ811	MA _T a cim3-1, ts sac7::SAC7-3xHA::HIS3MX6	This study
YEJ807	MA _T a cim3-1, ts cdc55Δ::KanMX6 sac7::SAC7-3xHA::HIS3MX6	This study
YSY186	MA _T a ade2 his3 leu2 lys2 trp1 ura3 fks1Δ::HIS3 fks2Δ::LYS2 ade3::fks1-1154::TRP (YOC1087)	Sekiya-Kawasaki et al., 2002
YSY124	MA _T a leu2-3, 112 ura3-52 trp1-1 his4 can1Δ pck1ΔLEU2 <YCp50-pkc1-2, ts>	Levin and Bartlett-Heubusch, 1992
SY1895	MA _T a swe1Δ::LEU2 lrg1::LRG1-GFP::HIS3MX6	This study
SY1897	MA _T a swe1Δ::LEU2 cdc55Δ::KanMX6 lrg1::LRG1-GFP::HIS3MX6	This study
SY1806	MA _T a sac7Δ::HIS3MX6	Yoshida et al., 2009

Unless otherwise indicated, all yeast strains used in this study were isogenic or congenic to BY4741 named PY 3295 or BY4742 named PY 3296.

Table S2. Plasmids used in this study

Plasmid	Gene	Marker and vector type	Source
PB2308	ROM2	URA3, CEN	Yoshida et al., 2009
pGEX4T-1	GST	Amp ^R	NA
pSY371	GST-Rho1-Q68L	Amp ^R	This study
pSY372	ZDS1-GFP	NA	This study
pSY373	CDC55-GFP	NA	This study
pSY374	ZDS1	URA3, leu2d, 2 μ (gTOW6000)	Makanae et al., 2013
pSY65	GFP-RHO1	URA3, CEN	Yoshida et al., 2009
pSY63	GFP-Rho1-Q68L	URA3, CEN	Yoshida et al., 2009
PB1561	RHO1-Q68H	URA3, CEN	Yoshida et al., 2009
PB1544	PKC1-R398P	URA3, CEN	BYP5216 (YGRC)
VR11	CDC55	URA3, 2 μ	This study
pSY375	pGAL ₁ -6xHis-SAC7	URA3, 2 μ	This study
PB1354	pGAL ₁ -PKC1-R398P	NA	Kono et al., 2012
pNV7-MKK1P386	pGAL ₁ -MKK1-S386P	URA3, Yep (pAT597)	Watanabe et al., 1995
PB1543	PKC1	URA3, 2 μ	This study

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