

Supplementary material

Yeast strains

Strains were grown in rich medium or synthetic medium as described (Guthrie and Fink, 1991).

Table S1. Yeast strains

Strains	Genotype	Source
EY492	<i>MATα ura3-1 leu2-3,112 trp1-1 his3-11 ade2-1 can1-100 Gal⁺</i>	R. Rothstein
EY699	<i>ura3-1 leu2-3,112 trp1-1 his3-11 ade2-1 can1-100 Gal⁺</i>	R. Rothstein
EY700	EY699 <i>fus3-6::LEU2</i>	E. Elion
EY957	EY699 <i>sst1</i>	E. Elion
EY1775	EY957 <i>ste5::TRP3</i>	E. Elion
EYL300	<i>MATα leu2-3 ura3-52</i>	R. Li
EYL302	<i>MATα leu2-3 ura3-52 cdc42-1</i>	R. Li
EYL427	<i>bni1::KAN ura3-1 leu2-3,112 trp1-1 his3-11 ade2-100 ade3</i>	D. Pellman
EYL428	<i>bni1::KAN ura3-1 leu2-3,112 trp1-1 his3-11 ade2-100 ade3 MATα</i>	D. Pellman
EYL917	EYL427 <i>sst1</i>	B.N. Lee
EYL1096	<i>sst1::LEU2 FUS3GFP::HIS3 ura3-1 leu3-3,112 trp1-1 his3-11 ade2-1 can1-100</i>	A. Levchenko
EYL1098	<i>myo1::KAN ura3-1 leu2-3,112 trp1-1 his3-11 ade2-1 can1-100 Gal⁺</i>	R. Li
EYL1730	<i>bnr1::KAN ura3-1 leu2-3,112 trp1-1 his3-11 ade2-100</i>	D. Pellman
EYL1735	<i>tpm1-2::LEU2 tpm2::HIS3 his3Δ-200 leu2-3,112 lys2-801 trp1-1 ura3-52</i>	A. Bretscher
EYL1736	<i>tpm2::HIS3 his3Δ-200 leu2-3,112 lys2-801 trp1-1 ura3-52</i>	A. Bretscher
EYL1748	<i>bni1::BNII#1::HIS3 bnr1::KAN ura3-1 leu2-3,112 trp1-1 his3-11 ade2-100</i>	D. Pellman
EYL1765	<i>his3Δ1 leu2Δ0 met15Δ0 ura3Δ0</i>	Res Gene
EYL1802	<i>MATα MYO2::HIS3 his3-Δ200 ura3-52 leu2-3,112 lys2-801 ade2-101 Gal⁺</i>	D. Pellman
EYL1807	<i>MATα myo2-66::HIS3 his3-Δ200 ura3-52 leu2-3,112 lys2-801 ade2-101 Gal⁺</i>	D. Pellman
QMY27	<i>sst1::LEU2 bni1::kanR FUS3GFP::HIS3 ura3-1 leu3-3,112 trp1-1 his3-11 ade2-1 can1-100</i>	This study
QMY458	EYL1802 switched to <i>MATα</i>	This study
QMY459	EYL1807 switched to <i>MATα</i>	This study

QMY483	<i>myo3::HIS3 ura3 leu2 trp1 his3 ade2</i>	R. Li
QMY484	<i>myo5::TRP1 ura3 leu2 trp1 his3 ade2</i>	R. Li
QMY485	<i>myo3::HIS myo5::TRP1 ura3 leu2 trp1 his3 ade2</i>	R. Li
QMY550	<i>ura3 leu2 trp1 his3 ade2</i>	R. Li
QMY551	<i>rho1-104 ura3 leu2 trp1 his3 ade2</i>	R. Li
QMY553	<i>rho1::HIS3 rho1-2 LEU2 ura3 leu2 trp1 his3 ade2</i>	R. Li
QMY574	<i>rho3::KAN his3Δ1 leu2Δ0 met15Δ0 ura3Δ0</i>	This study
QMY576	<i>rho5::KAN his3Δ1 leu2Δ0 met15Δ0 ura3Δ0</i>	This study
QMY630	<i>myo2-66::HIS3</i>	This study
QMY631	<i>myo4::KAN</i>	This study
QMY632	<i>myo2-66::HIS3 myo4::KAN</i>	This study
5126	<i>rho4::KAN his3Δ1 leu2Δ0 met15Δ0 ura3Δ0</i>	Res Gene
7230	<i>rho2::KAN his3Δ1 leu2Δ0 met15Δ0 ura3Δ0</i>	Res Gene
22032	<i>MATa rho5::KAN his3Δ1 leu2Δ0 met15Δ0 ura3Δ0</i> <i>MATα rho5::KAN his3Δ1 leu2Δ0 met15Δ0 ura3Δ0</i>	Res Gene
22277	<i>MATa rho3::KAN his3Δ1 leu2Δ0 met15Δ0 ura3Δ0</i> <i>MATα rho3::KAN his3Δ1 leu2Δ0 met15Δ0 ura3Δ0</i>	Res Gene

All strains except EY492, EYL428, 22032 and 22277 are *MATa*.

QMY27 was produced by cross of EYL428 with EYL1096. QMY630-632 were created by crossing EYL1685 with EYL1807 (*she1* is *myo4*). QMY574 and QMY576 were made by sporulation of 22277 and 22032.

Plasmid construction

All *BNI1* plasmids were made from a pRS316-based vector (PB1025, former name p182, see Evangelista et al., 1997) containing the *BNI1* coding sequence with 800 base pairs (bp) upstream and 1600 bp downstream sequence. The Rho-binding domain deletion mutant, *bni1ΔRBD* (QMB16, deletes aa 90-343), was made from two PCR amplified fragments of *BNI1* using the following primers: -788~268 fragment: 5'-ACGCGGATCCTTCACCGCTTTCGCACCTACTT-3' (*Bam*HI site at 5' terminus), 5'-TCCCCGCGGCGATTTTTTATTCAACGGCCT-3' (*Sac*II site at 3' terminus) and 1031~2472 fragment: 5'-CTACCGCGGCGAATCCTGGCAGACAACTT-3' (*Sac*II site at 5' terminus), 5'-TGCATGTCTCGAGTTCATACCTCTGGTGCCTTC-3'. The PCR products were cut with *Bam*HI/*Sac*II and *Sac*II/*Xho*I respectively, then performed a 3-way ligation with *Bam*HI/*Xho*I vector fragment of PB1025 to produce QMB16. The *Bam*HI/*Xho*I fragment of QMB16 containing *bni1ΔRBD* mutation was inserted into EBL334 to produce QMB79 (*bni1ΔRBD-GFP*). The *bni1ΔFH1FH2* mutant (QBM5, deletes

aa 1230-1748) was made by cloning the *BNII* gene into *Bam*HI-*Eag*I sites of pBR322 to make QMB3. A *BNII* fragment from the *Xho*I site to the FH1 domain (nucleotides 3572-4791) was PCR amplified with primers: 5'-GAACTCGAGCCTA AATTCTTCAG-3' and 5'-AGATCCGCGGCAGTAGAGAGATCTTCTGCG-3' (*Sac*II site at 5' terminus). This PCR product was used to replace the *Xho*I-*Sac*II fragment of *BNII* gene in QMB3 to make QMB4. Then the *Bam*HI-*Eag*I fragment of QMB4 was cloned into pRS316 to make QMB5. To make *bni1ΔFH1* (QMB21, deletes aa 1230-1328), fragment from *Xho*I site to FH1 domain (3572-4791) was PCR amplified by the same way as for FH domain deletion, but added a *Sph*I site instead of *Sac*II site to the second primers: 5'-GATGCATGCAGTAGAGAGAT CTTCTGCG-3'. Fragment from the end of FH1 to *Sac*II site (5086-6352) were PCR amplified with 5' primer: 5'-CATGCATGCGCATCGCAAATCAAATCAGCT-3' (*Sph*I site at 5' terminus) and 3' primer: 5'-CTTCCGCGGCTAGATTTTGCCTT-3'. The first and second PCR products were digested with *Xho*I/*Sph*I and *Sph*I/*Sac*II respectively, then used in a 3-way ligation with the *Xho*I/*Sac*II fragment of QMB3 to make QMB19. The *Bam*HI-*Eag*I fragments of QMB19 was cloned into pRS316 to make QMB21. The same strategy was used to make *bni1ΔFH2* (QMB22, deletes aa 1492-1640), except a different 3' primer (5'-TGTAGCATGCCCTCACGCC CTCCAGTCTG-3') for the first PCR, which amplifies fragment from *Xho*I site to FH2 domain (3572-5577), and a different 5' primers (5'-CGATGCATGCTCCATTGAGCAGTTAGTTAA-3') for the second PCR, which amplifies fragment from the end of FH2 domain to *Sac*II site (6021-6352). QMB34 (*bni1ΔCT*, deletes aa 1749-1953) was made by cloning *Bam*HI-*Eag*I fragment of PB1046 (Lee et al., 1999) into pRS315.

Table S2. Plasmids used in this study

Plasmids	Description	Source	
QMB5	<i>bni1ΔFH1FH2</i>	<i>CEN</i> <i>URA3</i>	This study
QMB16	<i>bni1ΔRBD</i>	<i>CEN</i> <i>URA3</i>	This study
QMB21	<i>bni1ΔFH1</i>	<i>CEN</i> <i>URA3</i>	This study
QMB22	<i>bni1ΔFH2</i>	<i>CEN</i> <i>URA3</i>	This study
QMB34	<i>bni1ΔCT</i>	<i>CEN</i> <i>LEU2</i>	This study
QMB75	<i>MYO2DN</i>	<i>GAL1</i> <i>LEU2</i>	S. Reck-Peterson
QMB79	<i>bni1ΔRBD-GFP</i>	<i>CEN(ADH1)</i> <i>URA3</i>	This study
QMB80	<i>MYO2-HA</i>	<i>CEN</i> <i>HIS3</i>	L. Weisman
EBL95	<i>FUS1-LACZ</i>	<i>2</i> <i>LEU2</i>	Elion et al. (1990)
EBL204	<i>STE5-CTM</i>	<i>CEN(GAL1)</i> <i>LEU2</i>	Pryciak et al. (1998)

EBL284	<i>STE11-4</i>	<i>CEN</i>	<i>HIS3</i>	H. Madhani
EBL332	<i>BNI1-HA4</i>	<i>CEN</i>	<i>URA3</i>	D. Pellman
EBL334	<i>BNI1-GFP</i>	<i>CEN(ADH1)</i>	<i>URA3</i>	D. Pellman
EBL358	<i>STE5-MYC9</i>	<i>CEN</i>	<i>URA3</i>	Mahanty <i>et al.</i> (1999)
EBL365	<i>STE5-MYC9</i>	<i>2</i>	<i>URA3</i>	Mahanty <i>et al.</i> (1999)
EBL367	<i>STE5-GFP</i>	<i>CEN(CUP1)</i>	<i>URA3</i>	Mahanty <i>et al.</i> (1999)
EBL444	<i>TAgNLS^{K128T}-STE5-M9</i>	<i>2</i>	<i>URA3</i>	Mahanty <i>et al.</i> (1999)
EBL453	<i>STE5-MYC9</i>	<i>2</i>	<i>LEU2</i>	Mahanty <i>et al.</i> (1999)
EBL511	<i>GFP-STE20</i>	<i>CEN</i>	<i>URA3</i>	Leberer <i>et al.</i> (1997)
EBL664	<i>GFP-CDC24</i>	<i>CEN(MET)</i>	<i>LEU2</i>	Toenjes <i>et al.</i> (1999)
PYEE1102	<i>FUS3-HA</i>	<i>CEN</i>	<i>HIS3</i>	Elion <i>et al.</i> (1993)

Supplemental References

Elion, E.A., Grisafi, P.L. and Fink, G.R. (1990). FUS3 encodes a *cdc2+*/CDC28-related kinase required for the transition from mitosis into conjugation. *Cell*. **60**, 649-664.

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