

**Table S1. Yeast Strains used in this study**

Strain	Genotype	Origin
BY4741	<i>MATa his3Δ1 leu2Δ0 met25Δ0 ura3Δ0</i>	<i>Saccharomyces cerevisiae</i> (ATCC® 201388™)
BY4742	<i>MATα his3Δ1 leu2Δ0 lys2Δ0 ura3Δ0</i>	<i>Saccharomyces cerevisiae</i> (ATCC® 201389™)
MMY0266	<i>MATα cdc10(D182N)-Vc::KanMX his3Δ1 leu2Δ0 met25Δ0 ura3Δ0</i>	(Weems and McMurray 2017)
YO802	<i>MATα his3 leu2 lys2 trp1 ura3 CDC12-Vc::KanMX</i>	(Oh et al. 2013)
YO685	<i>MATα his3 leu2 lys2 trp1 ura3 CDC3-Vc::HIS3MX</i>	(Oh et al. 2013)
YO1057	<i>MATα his3 leu2 lys2 trp1 ura3 CDC10-Vc::TRP1</i>	(Oh et al. 2013)
YO671	<i>MATα his3 leu2 lys2 trp1 ura3 CDC11-Vc::HIS3MX</i>	(Oh et al. 2013)
YO619	<i>MATα his3 leu2 lys2 trp1 ura3 SHS1-Vc::HIS3MX</i>	(Oh et al. 2013)
M-1622	<i>MATα his3 leu2 lys2 trp1 ura3 cdc3(G365R)</i>	(Nagaraj et al. 2008)
RBy33	<i>MATa his3Δ200 leu2Δ1 lys2Δ202 trp1Δ63 1cUAS53::URA3</i>	(Brachmann et al. 1996)
RBy159	<i>MATα his3Δ200 leu2Δ1 lys2Δ202 trp1Δ63 1cUAS53::URA3</i>	(Brachmann et al. 1996)
yIG397	<i>MATa ade2-1 can1-100 his3-11,15 leu2-3,112 trp1-1 ura3-1 URA3 3XRGC::pCYC1::ADE2</i>	(Ishioka et al. 1993)
yJM3164	<i>MATa his3Δ200 HTB2-mNeonGreen::HIS3 leu2Δ1 lys2-801 trp1Δ63 ura3-52</i>	Gift from Jeff Moore
yJM1838	<i>MATα his3Δ200 leu2Δ1 lys2-801 trp1Δ63 ura3-52</i>	Gift from Jeff Moore
JTY4204	<i>MATa gal2 leu2 pep4-3 prb1-1122 prc1-451 trp1Δ63 ura3</i>	(Jones 1977)
TH5	<i>MATa leu2-3,112 trp1 tup1 ura3-52 dfr1::URA3</i>	(Huang et al. 1992)
Y0134	<i>MATa his3Δ1 leu2Δ0 met25Δ0 trp1Δ::KanMX ura3Δ0</i>	<i>Saccharomyces</i> Genome Deletion Project (Winzeler et al. 1999; Giaever et al. 2002)
H00398	<i>MATa his3Δ1 HSP82-VN::KIURA3 leu2Δ0 met25Δ0 ura3Δ0</i>	<i>S. cerevisiae</i> VN-Fusion Library (Sung et al. 2013)
H00399	<i>MATa his3Δ1 HSC82-VN::KIURA3 leu2Δ0 met25Δ0 ura3Δ0</i>	<i>S. cerevisiae</i> VN-Fusion Library (Sung et al. 2013)

Strain	Genotype	Origin
H00384	<i>MATa his3Δ1 HSP104-V<sub>N</sub>::KIURA3 leu2Δ0 met25Δ0 ura3Δ0</i>	<i>S. cerevisiae</i> V <sub>N</sub> -Fusion Library (Sung <i>et al.</i> 2013)
H00387	<i>MATa his3Δ1 leu2Δ0 met25Δ0 SSA1-V<sub>N</sub>::KIURA3 ura3Δ0</i>	<i>S. cerevisiae</i> V <sub>N</sub> -Fusion Library (Sung <i>et al.</i> 2013)
H00388	<i>MATa his3Δ1 leu2Δ0 met25Δ0 SSA2-V<sub>N</sub>::KIURA3 ura3Δ0</i>	<i>S. cerevisiae</i> V <sub>N</sub> -Fusion Library (Sung <i>et al.</i> 2013)
H00403	<i>MATa his3Δ1 leu2Δ0 met25Δ0 SSA3-V<sub>N</sub>::KIURA3 ura3Δ0</i>	<i>S. cerevisiae</i> V <sub>N</sub> -Fusion Library (Sung <i>et al.</i> 2013)
H00404	<i>MATa his3Δ1 leu2Δ0 met25Δ0 SSA4-V<sub>N</sub>::KIURA3 ura3Δ0</i>	<i>S. cerevisiae</i> V <sub>N</sub> -Fusion Library (Sung <i>et al.</i> 2013)
H00395	<i>MATa his3Δ1 leu2Δ0 met25Δ0 SSB1-V<sub>N</sub>::KIURA3 ura3Δ0</i>	<i>S. cerevisiae</i> V <sub>N</sub> -Fusion Library (Sung <i>et al.</i> 2013)
H00394	<i>MATa his3Δ1 leu2Δ0 met25Δ0 SSB2-V<sub>N</sub>::KIURA3 ura3Δ0</i>	<i>S. cerevisiae</i> V <sub>N</sub> -Fusion Library (Sung <i>et al.</i> 2013)
H00383	<i>MATa CCT7-V<sub>N</sub>::KIURA3 his3Δ1 leu2Δ0 met25Δ0 ura3Δ0</i>	<i>S. cerevisiae</i> V <sub>N</sub> -Fusion Library (Sung <i>et al.</i> 2013)
H00526	<i>CCT7-V<sub>N</sub>::URA3 CDC3-V<sub>C</sub>::HIS3MX</i>	Spore clone from YO685 x H00383 cross
H04802	<i>MATa GIM3-V<sub>N</sub>::KIURA3 his3Δ1 leu2Δ0 met25Δ0 ura3Δ0</i>	<i>S. cerevisiae</i> V <sub>N</sub> -Fusion Library (Sung <i>et al.</i> 2013)
H00385	<i>MATa his3Δ1 leu2Δ0 met25Δ0 ura3Δ0 YDJ1-V<sub>N</sub>::KIURA3</i>	<i>S. cerevisiae</i> V <sub>N</sub> -Fusion Library (Sung <i>et al.</i> 2013)
H00396	<i>MATa his3Δ1 leu2Δ0 met25Δ0 ura3Δ0 ZUO1-V<sub>N</sub>::KIURA3</i>	<i>S. cerevisiae</i> V <sub>N</sub> -Fusion Library (Sung <i>et al.</i> 2013)
H06680	<i>MATa his3Δ1 leu2Δ0 met25Δ0 SIS1-V<sub>N</sub>::KIURA3 ura3Δ0</i>	<i>S. cerevisiae</i> V <sub>N</sub> -Fusion Library (Sung <i>et al.</i> 2013)
H06674	<i>MATa APJ1-V<sub>N</sub>::KIURA3 his3Δ1 leu2Δ0 met25Δ0 ura3Δ0</i>	<i>S. cerevisiae</i> V <sub>N</sub> -Fusion Library (Sung <i>et al.</i> 2013)
H05926	<i>MATa DJP1-V<sub>N</sub>::KIURA3 his3Δ1 leu2Δ0 met25Δ0 ura3Δ0</i>	<i>S. cerevisiae</i> V <sub>N</sub> -Fusion Library (Sung <i>et al.</i> 2013)
H06675	<i>MATa his3Δ1 leu2Δ0 met25Δ0 ura3Δ0 JJJ1-V<sub>N</sub>::KIURA3</i>	<i>S. cerevisiae</i> V <sub>N</sub> -Fusion Library (Sung <i>et al.</i> 2013)
H06685	<i>MATa his3Δ1 leu2Δ0 met25Δ0 ura3Δ0 JJJ2-V<sub>N</sub>::KIURA3</i>	<i>S. cerevisiae</i> V <sub>N</sub> -Fusion Library (Sung <i>et al.</i> 2013)
H00400	<i>MATa his3Δ1 leu2Δ0 met25Δ0 ura3Δ0 SSE2-V<sub>N</sub>::KIURA3</i>	<i>S. cerevisiae</i> V <sub>N</sub> -Fusion Library (Sung <i>et al.</i> 2013)
H06681	<i>MATa his3Δ1 leu2Δ0 met25Δ0 ura3Δ0 SSZ1-V<sub>N</sub>::KIURA3</i>	<i>S. cerevisiae</i> V <sub>N</sub> -Fusion Library (Sung <i>et al.</i> 2013)
H06682	<i>MATa his3Δ1 leu2Δ0 met25Δ0 ura3Δ0 SWA2-V<sub>N</sub>::KIURA3</i>	<i>S. cerevisiae</i> V <sub>N</sub> -Fusion Library (Sung <i>et al.</i> 2013)
H06683	<i>MATa his3Δ1 leu2Δ0 met25Δ0 ura3Δ0 SNO4-V<sub>N</sub>::KIURA3</i>	<i>S. cerevisiae</i> V <sub>N</sub> -Fusion Library (Sung <i>et al.</i> 2013)

Strain	Genotype	Origin
H06684	<i>MATa his3Δ1 leu2Δ0 met25Δ0 ura3Δ0 HSP12-V<sub>N</sub>::KIURA3</i>	<i>S. cerevisiae</i> V <sub>N</sub> -Fusion Library (Sung <i>et al.</i> 2013)
H04024	<i>MATa his3Δ1 leu2Δ0 met25Δ0 ura3Δ0 HSP26-V<sub>N</sub>::KIURA3</i>	<i>S. cerevisiae</i> V <sub>N</sub> -Fusion Library (Sung <i>et al.</i> 2013)
H00397	<i>MATa his3Δ1 HSP31-V<sub>N</sub>::KIURA3 leu2Δ0 met25Δ0 ura3Δ0</i>	<i>S. cerevisiae</i> V <sub>N</sub> -Fusion Library (Sung <i>et al.</i> 2013)
H00397	<i>MATa his3Δ1 HSP32-V<sub>N</sub>::KIURA3 leu2Δ0 met25Δ0 ura3Δ0</i>	<i>S. cerevisiae</i> V <sub>N</sub> -Fusion Library (Sung <i>et al.</i> 2013)
H06677	<i>MATa his3Δ1 HSP33-V<sub>N</sub>::KIURA3 leu2Δ0 met25Δ0 ura3Δ0</i>	<i>S. cerevisiae</i> V <sub>N</sub> -Fusion Library (Sung <i>et al.</i> 2013)
H06676	<i>MATa his3Δ1 HSP42-V<sub>N</sub>::KIURA3 leu2Δ0 met25Δ0 ura3Δ0</i>	<i>S. cerevisiae</i> V <sub>N</sub> -Fusion Library (Sung <i>et al.</i> 2013)
H00401	<i>MATa his3Δ1 leu2Δ0 met25Δ0 PAM18-V<sub>N</sub>::KIURA3 ura3Δ0</i>	<i>S. cerevisiae</i> V <sub>N</sub> -Fusion Library (Sung <i>et al.</i> 2013)
H00379	<i>MATa CCT2-V<sub>N</sub>::KIURA3 his3Δ1 leu2Δ0 met25Δ0 ura3Δ0</i>	<i>S. cerevisiae</i> V <sub>N</sub> -Fusion Library (Sung <i>et al.</i> 2013)
H00380	<i>MATa CCT3-V<sub>N</sub>::KIURA3 his3Δ1 leu2Δ0 met25Δ0 ura3Δ0</i>	<i>S. cerevisiae</i> V <sub>N</sub> -Fusion Library (Sung <i>et al.</i> 2013)
H00380	<i>MATa CCT4-V<sub>N</sub>::KIURA3 his3Δ1 leu2Δ0 met25Δ0 ura3Δ0</i>	<i>S. cerevisiae</i> V <sub>N</sub> -Fusion Library (Sung <i>et al.</i> 2013)
H00391	<i>MATa GIM4-V<sub>N</sub>::KIURA3 his3Δ1 leu2Δ0 met25Δ0 ura3Δ0</i>	<i>S. cerevisiae</i> V <sub>N</sub> -Fusion Library (Sung <i>et al.</i> 2013)
H00390	<i>MATa GIM5-V<sub>N</sub>::KIURA3 his3Δ1 leu2Δ0 met25Δ0 ura3Δ0</i>	<i>S. cerevisiae</i> V <sub>N</sub> -Fusion Library (Sung <i>et al.</i> 2013)
H00393	<i>MATa PFD1-V<sub>N</sub>::KIURA3 his3Δ1 leu2Δ0 met25Δ0 ura3Δ0</i>	<i>S. cerevisiae</i> V <sub>N</sub> -Fusion Library (Sung <i>et al.</i> 2013)
H00389	<i>MATa his3Δ1 leu2Δ0 met25Δ0 ura3Δ0 YKE2-V<sub>N</sub>::KIURA3</i>	<i>S. cerevisiae</i> V <sub>N</sub> -Fusion Library (Sung <i>et al.</i> 2013)
H00405	<i>MATa his3Δ1 leu2Δ0 LHS1-V<sub>N</sub>::KIURA3 met25Δ0 ura3Δ0</i>	<i>S. cerevisiae</i> V <sub>N</sub> -Fusion Library (Sung <i>et al.</i> 2013)
H03367	<i>MATa his3Δ1 JID1-V<sub>N</sub>::KIURA3 leu2Δ0 met25Δ0 ura3Δ0</i>	<i>S. cerevisiae</i> V <sub>N</sub> -Fusion Library (Sung <i>et al.</i> 2013)
H03970	<i>MATa his3Δ1 HLJ1-V<sub>N</sub>::KIURA3 leu2Δ0 met25Δ0 ura3Δ0</i>	<i>S. cerevisiae</i> V <sub>N</sub> -Fusion Library (Sung <i>et al.</i> 2013)
H05688	<i>MATa his3Δ1 SEC63-V<sub>N</sub>::KIURA3 leu2Δ0 met25Δ0 ura3Δ0</i>	<i>S. cerevisiae</i> V <sub>N</sub> -Fusion Library (Sung <i>et al.</i> 2013)
H01527	<i>MATa ECM10-V<sub>N</sub>::KIURA3 his3Δ1 leu2Δ0 met25Δ0 ura3Δ0</i>	<i>S. cerevisiae</i> V <sub>N</sub> -Fusion Library (Sung <i>et al.</i> 2013)
H01444	<i>MATa his3Δ1 leu2Δ0 met25Δ0 SSQ1-V<sub>N</sub>::KIURA3 ura3Δ0</i>	<i>S. cerevisiae</i> V <sub>N</sub> -Fusion Library (Sung <i>et al.</i> 2013)
H04280	<i>MATa his3Δ1 JAC1-V<sub>N</sub>::KIURA3 leu2Δ0 met25Δ0 ura3Δ0</i>	<i>S. cerevisiae</i> V <sub>N</sub> -Fusion Library (Sung <i>et al.</i> 2013)

Strain	Genotype	Origin
H05828	<i>MATa his3Δ1 leu2Δ0 MDJ1-V<sub>N</sub>::KIURA3 met25Δ0 ura3Δ0</i>	<i>S. cerevisiae</i> V <sub>N</sub> -Fusion Library (Sung <i>et al.</i> 2013)
H01128	<i>MATa his3Δ1 HSP78-V<sub>N</sub>::KIURA3 leu2Δ0 met25Δ0 ura3Δ0</i>	<i>S. cerevisiae</i> V <sub>N</sub> -Fusion Library (Sung <i>et al.</i> 2013)
H01972	<i>MATa his3Δ1 leu2Δ0 MCX1-V<sub>N</sub>::KIURA3 met25Δ0 ura3Δ0</i>	<i>S. cerevisiae</i> V <sub>N</sub> -Fusion Library (Sung <i>et al.</i> 2013)
H01842	<i>MATa his3Δ1 HSP60-V<sub>N</sub>::KIURA3 leu2Δ0 met25Δ0 ura3Δ0</i>	<i>S. cerevisiae</i> V <sub>N</sub> -Fusion Library (Sung <i>et al.</i> 2013)

## Literature cited

- Brachmann R. K., M. Vidal, and J. D. Boeke, 1996 Dominant-negative p53 mutations selected in yeast hit cancer hot spots. *Proc Natl Acad Sci U S A* 93: 4091–4095. <https://doi.org/10.1073/pnas.93.9.4091>
- Giaever G., A. M. Chu, L. Ni, C. Connelly, L. Riles, *et al.*, 2002 Functional profiling of the *Saccharomyces cerevisiae* genome. *Nature* 418: 387–391. <https://doi.org/10.1038/nature00935>
- Huang T., B. J. Barclay, T. I. Kalman, R. C. von Borstel, and P. J. Hastings, 1992 The phenotype of a dihydrofolate reductase mutant of *Saccharomyces cerevisiae*. *Gene* 121: 167–171. [https://doi.org/10.1016/0378-1119\(92\)90177-q](https://doi.org/10.1016/0378-1119(92)90177-q)
- Ishioka C., T. Frebourg, Y. X. Yan, M. Vidal, S. H. Friend, *et al.*, 1993 Screening patients for heterozygous p53 mutations using a functional assay in yeast. *Nat Genet* 5: 124–129. <https://doi.org/10.1038/ng1093-124>
- Jones E. W., 1977 Proteinase mutants of *Saccharomyces cerevisiae*. *Genetics* 85: 23–33.
- Nagaraj S., A. Rajendran, C. E. Jackson, and M. S. Longtine, 2008 Role of nucleotide binding in septin-septin interactions and septin localization in *Saccharomyces cerevisiae*. *Mol. Cell. Biol* 28: 5120–5137. <https://doi.org/10.1128/MCB.00786-08>
- Oh Y., J. Schreiter, R. Nishihama, C. Wloka, and E. Bi, 2013 Targeting and functional mechanisms of the cytokinesis-related F-BAR protein Hof1 during the cell cycle. *Mol. Biol. Cell* 24: 1305–1320. <https://doi.org/10.1091/mbc.E12-11-0804>
- Sung M.-K., G. Lim, D.-G. Yi, Y. J. Chang, E. B. Yang, *et al.*, 2013 Genome-wide bimolecular fluorescence complementation analysis of SUMO interactome in yeast. *Genome Res* 23: 736–746. <https://doi.org/10.1101/gr.148346.112>

- Weems A. D., and M. A. McMurray, 2017 The step-wise pathway of septin hetero-octamer assembly in budding yeast. *Elife* 6. <https://doi.org/10.7554/eLife.23689>
- Winzler E. A., D. D. Shoemaker, A. Astromoff, H. Liang, K. Anderson, *et al.*, 1999 Functional characterization of the *S. cerevisiae* genome by gene deletion and parallel analysis. *Science* 285: 901–906.  
<https://doi.org/10.1126/science.285.5429.901>