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Supplemental information

Global phosphoproteomics pinpoints uncharted

Gcn2-mediated mechanisms of translational control

Ladislav Dokládal, Michael Stumpe, Benjamin Pillet, Zehan Hu, Guillermo Miguel Garcia Osuna, Dieter Kressler, Jörn Dengjel, and Claudio De Virgilio

Supplemental Information

Supplemental Figures

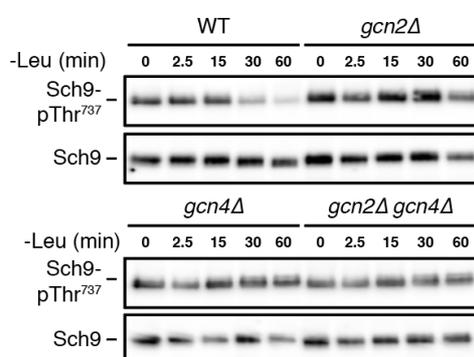


Figure S1. Gcn2 and Gcn4 are Required for Proper TORC1 Inactivation in Leucine-Starved Cells, Related to Figure 1

Wild-type (WT), *gcn2Δ*, *gcn4Δ*, and *gcn2Δ gcn4Δ* cells were grown exponentially (time point 0) and starved for leucine (-Leu) for the indicated time periods. Phosphorylation of the TORC1 target residue Thr⁷³⁷ in Sch9 was monitored by immunoblotting with anti-Sch9-pThr⁷³⁷ and anti-Sch9 antibodies.

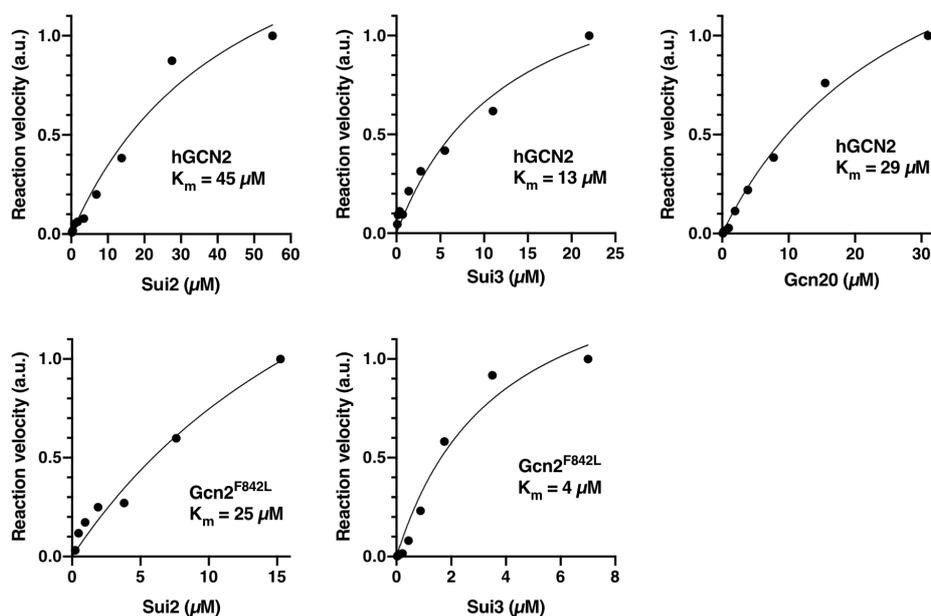


Figure S2. Gcn2 Targets Sui3 and Gcn20 *in Vitro*, Related to Figures 2A and 2B

Kinetic analyses of human GCN2 and yeast Gcn2^{F842L} phosphorylation of yeast Sui2, Sui3, and/or Gcn20. Recombinant substrates were purified from bacteria and examined for phosphorylation by human GCN2 and/or yeast Gcn2^{F842L}. The V_{max} values for the hGCN2-Sui3 and hGCN2-Gcn20 reactions were 67% and 60%, respectively, when compared to the one of the hGCN2-Sui2 reaction (set to 100%). Gcn20 was not subjected to phosphorylation by Gcn2^{F842L} because endogenous Gcn20 co-immunoprecipitating with Gcn2^{F842L} may compete in this assay with the recombinant Gcn20 substrate. Because Gcn2 had very low activity towards Sui2 (control), the kinetic *in vitro* analyses we performed with hyperactive Gcn2^{F842L}. Data are expressed in arbitrary units (a.u.).

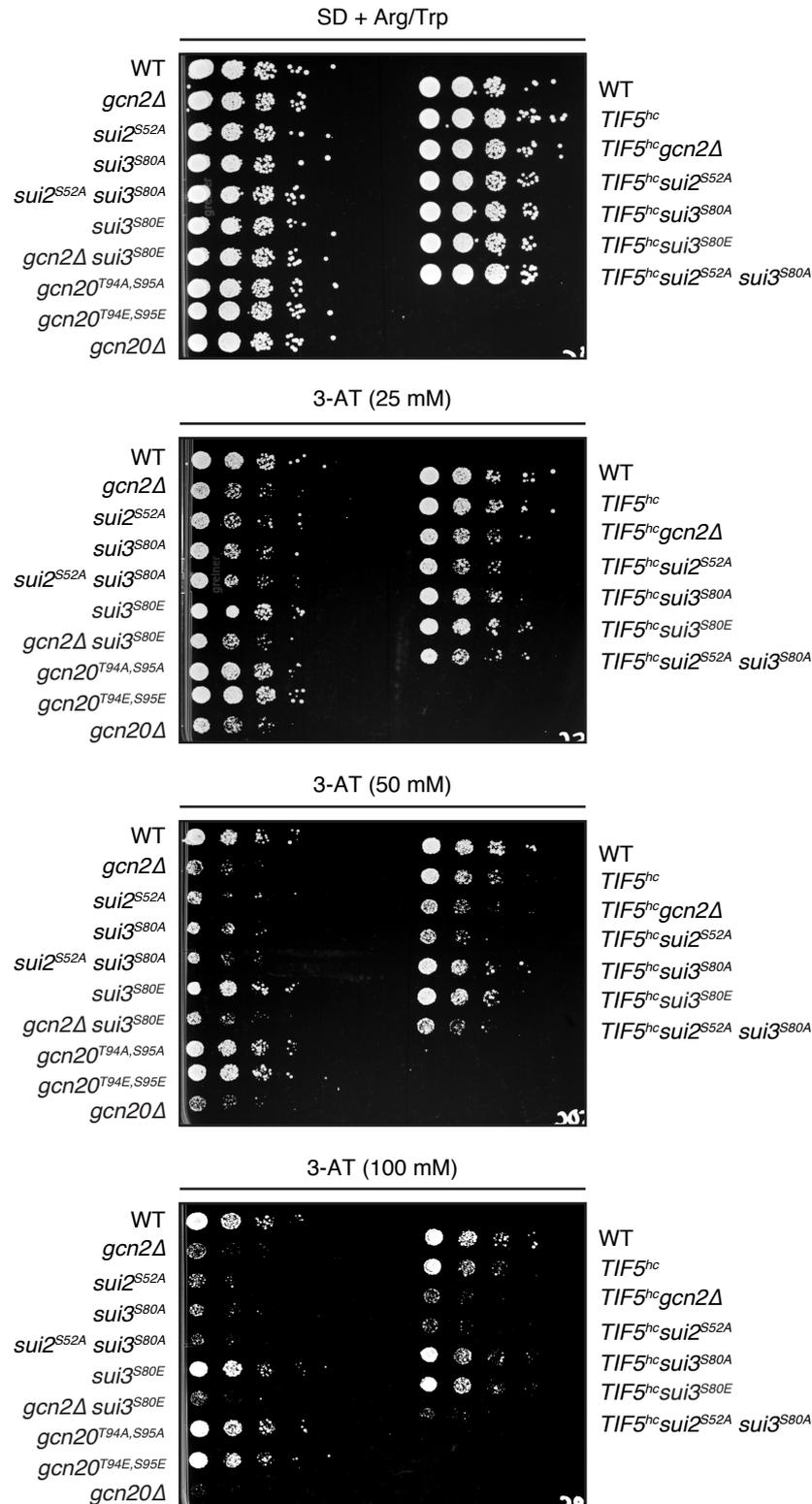


Figure S3. 3-AT Sensitivity of *sui2*, *sui3*, *gcn20*, and *gcn2Δ* Mutants Overexpressing, or Not, *Tif5* from a High-copy Plasmid (*TIF5^{hc}*), Related to Figures 2D and 3E

Exponentially growing strains with the indicated genotypes were spotted (10-fold serial dilutions) and grown for 3 days at 30°C on SD + Arg/Trp or on plates containing the indicated concentrations of 3-AT. The pictures of the control plate (SD + Arg/Trp) and the one with 100 mM 3-AT were used to assemble Fig. 2D and 3E.

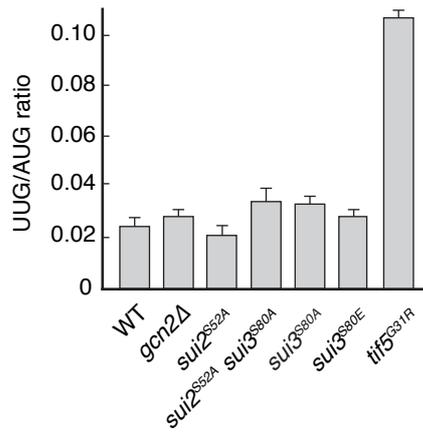


Figure S4. Mutation of Ser⁸⁰ in Sui3 Has Little Impact on Near Cognate Start Codon Recognition on the *HIS4-lacZ* Reporter, Related to Figure 3

Strains with the indicated genotypes expressing either *his4-UUG-LacZ* or *HIS4-AUG-LacZ* reporters from plasmids were assayed for β -galactosidase activities. *tif5^{G31R}* (control) was expressed from a plasmid over the wild-type allele. The respective values were used to calculate the mean UUG/AUG ratio (n = 3; + SD).