

Control of the integrated stress response by a stochastic molecular switch

Stefan M. Kallenberger^{1,2,3}, Philipp Klein⁴, Hanna Roth⁴, Karsten Roth⁴, Thi Bach Nga Ly-Hartig^{5,6}, Vera Magg⁴, Janez Aleš⁷, Soheil Rastgou Talemi⁸, Yu Qiang⁹, Steffen Wolf⁷, Olga Oleksiuk⁴, Roma Kurilov¹⁰, Barbara Di Ventura¹¹, Ralf Bartenschlager^{4,12}, Roland Eils^{1,2}, Karl Rohr⁹, Fred A. Hamprecht⁷, Thomas Höfer⁸, Oliver T. Fackler¹³, Georg Stoecklin^{5,6}, Alessia Ruggieri⁴

¹ Health Data Science Unit, Heidelberg University Hospital and BioQuant, Heidelberg, Germany

² Digital Health Center, Berlin Institute of Health (BIH) and Charité, Berlin, Germany.

³ Medical Oncology, National Center for Tumor Diseases, Heidelberg University, Heidelberg, Germany

⁴ Department of Infectious Diseases, Molecular Virology, Center for Integrative Infectious Diseases Research, Heidelberg University, Heidelberg, Germany

⁵ Division of Biochemistry, Mannheim Institute for Innate Immunoscience (MI3), Medical Faculty Mannheim, Heidelberg University, Mannheim, Germany

⁶ Center for Molecular Biology of Heidelberg University (ZMBH), DKFZ-ZMBH Alliance, Heidelberg, Germany

⁷ HCI/IWR, Heidelberg University, Heidelberg, Germany

⁸ Division of Theoretical Systems Biology, German Cancer Research Center (DKFZ), Heidelberg, Germany

⁹ Biomedical Computer Vision Group, BioQuant, IPMB, Heidelberg University, Heidelberg, Germany

¹⁰ Applied Bioinformatics Division, German Cancer Research Center (DKFZ), Heidelberg, Germany

¹¹ Faculty of Biology, Institute of Biology II, University of Freiburg, Freiburg, Germany; Centers for Biological Signalling Studies BIOSS and CIBSS, University of Freiburg, Freiburg, Germany

¹² Division Virus-Associated Carcinogenesis, German Cancer Research Center (DKFZ), Heidelberg, Germany

¹³ Department of Infectious Diseases, Integrative Virology, Center for Integrative Infectious Diseases Research, Heidelberg University, Heidelberg, Germany

Stress granules (SGs) are formed in the cytosol as an acute response to environmental cues, such as virus infections or exposition to chemotherapeutic drugs, that activate the integrated stress response (ISR), a central signaling pathway controlling protein synthesis. Using chronic virus infection as stress model, we previously uncovered a unique temporal control of the ISR resulting in recurrent phases of SG assembly and disassembly. Here, we elucidated the molecular network generating this fluctuating stress response, by integrating quantitative experiments with mathematical modeling, and found that the ISR operates as a stochastic switch. Key elements controlling this switch are the cooperative activation of the stress-sensing kinase PKR, the ultrasensitive response of SG formation to the phosphorylation of the translation initiation factor eIF2 α , and negative feedback via GADD34, a stress-induced subunit of protein phosphatase 1. This negative feedback loop was responsible for adaptation to cellular stress as well as the dynamics of stress-induced apoptosis. We identified GADD34 mRNA levels as the molecular memory of the ISR that plays a central role in cell adaptation to acute and chronic stress.