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The role of MAPK and SCF in the destruction of Med13 in cyclin C mediated cell death. D.C. Stieg¹, S.D. Willis¹, J. Scuorzo², M. Song², V. Ganesan¹, R. Strich¹, K.F. Cooper¹; ¹Molecular Biology, Rowan University Graduate School of Biomedical Sciences, Stratford, NJ, ²Medicine, Rowan University School of Osteopathic Medicine, Stratford, NJ

In response to stress, the yeast ¹ and mammalian ² cyclin C translocate from the nucleus to the cytoplasm, where it associates with the GTPase Drp1/Dnm1 to drive mitochondrial fragmentation and apoptosis. Therefore, the decision to release cyclin C represents a key life or death decision. In unstressed cells, the cyclin C-Cdk8 kinase regulates transcription by associating with the Mediator of RNA polymerase II. We previously reported that that the Mediator component Med13 anchors cyclin C in the nucleus³. Loss of Med13 function leads to constitutive cytoplasmic localization of cyclin C, resulting in fragmented mitochondria, hypersensitivity to stress and mitochondrial dysfunction due to

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loss of mtDNA. Recently we showed that this molecular switch operates in a two step process. First, efficient cyclin C nuclear release requires its ROS-induced phosphorylation by the MAP kinase Slt2⁴ in a carboxyl terminal region of cyclin C, which includes a putative Med13 interaction site. The second step involves ROS-induced Med13 destruction by the SCF^{Grr1} ubiquitin ligase. Med13 associates with Grr1 in two-hybrid assays, and SCF mediated degradation of Med13 requires active cyclin C-Cdk8⁵. However, phosphorylation of Med13 by cyclin C-Cdk8 does not trigger Med13 destruction. This suggests a model in which this kinase primes the Med13 degron for SCF^{Grr1} and either Slt2, or an as yet unidentified kinase, triggers its destruction. Taken together, these results are consistent with a model in which cyclin C phosphorylation by Slt2 permits its disassociation from Med13, and that Med13 destruction allows full cyclin C release and prevents re-accumulation of the cyclin in the nucleus.

¹Dev. Cell. (2014), 28:161; ²Mol. Biol. Cell. (2015), 26:1030; ³Mol. Biol. Cell (2104) 25:2807; ⁴Mol. Biol. Cell. (2014) 25:1396; ⁵MB0C. (2017) submitted

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