Regulators of mitochondrial translation in Saccharomyces cerevisiae

Siep, M.

Publication date
2001

Citation for published version (APA):

General rights
It is not permitted to download or to forward/distribute the text or part of it without the consent of the author(s) and/or copyright holder(s), other than for strictly personal, individual use, unless the work is under an open content license (like Creative Commons).

Disclaimer/Complaints regulations
If you believe that digital publication of certain material infringes any of your rights or (privacy) interests, please let the Library know, stating your reasons. In case of a legitimate complaint, the Library will make the material inaccessible and/or remove it from the website. Please Ask the Library: https://uba.uva.nl/en/contact, or a letter to: Library of the University of Amsterdam, Secretariat, Singel 425, 1012 WP Amsterdam, The Netherlands. You will be contacted as soon as possible.
References


Rasmussen, S.W. A 37.5 kb region of yeast chromosome X includes the *SME1, MEF2, GSH1* and *CSD3* genes, a *TCP-1*-related gene, an open reading frame similar to the *DAL80* gene, and a tRNA(Arg). *Yeast* **11**, 873-83 (1995).


Rijnberg, M. Analysis of the *COX1* translational activator MSS51. in *Student report, University of Amsterdam*. (1999).


Wallis, M.G., Groudinsky, O., Slonimski, P.P. & Dujardin, G. The NAM1 protein (NAM1p), which is selectively required for cox1, cytb and atp6 transcript processing/stabilisation, is located in the yeast mitochondrial matrix. Eur J Biochem 222, 27-32 (1994).


Wienholds, E. Functional analysis of MSS51, a possible translational activator of COX1 mRNA. in Student report, University of Amsterdam. (, 1998).


