

# GENETICS AND BIOGENESIS OF CHLOROPLASTS AND MITOCHONDRIA

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## Contents

Preface	v
<b>ROLE OF CHLOROPLAST AND NUCLEAR GENES IN PRODUCTION OF CHLOROPLAST PROTEINS</b>	
Role of chloroplast and nuclear DNA genes during evolution of fraction I protein K. Chen, S. Johal and S.G. Wildman	3
Polypeptide chains of the large and small subunits of fraction I protein J.C. Gray, S.D. Kung and S.G. Wildman	13
Studies on the primary structure of the small subunit of ribulose-1,5-diphosphate carboxylase C. Poulsen, S. Strøbaek and B.G. Haslett	17
Ontogeny, insertion, and activation of two thylakoid peptides required for photosystem II activity in the nuclear temperature sensitive T4 mutant of <i>Chlamydomonas reinhardtii</i> F. Kretzer, I. Ohad and P. Bennoun	25
Methods for the detection and characterization of photosynthetic mutants in <i>Chlamydomonas reinhardtii</i> P. Bennoun and N.H. Chua	33
Biochemical studies on a plastid ribosome-deficient mutant of <i>Hordeum vulgare</i> T. Börner, B. Schumann and R. Hagemann	41
Sites of synthesis of chloroplast membrane proteins in <i>Vicia faba</i> W. Hachtel	49
Synthesis of chlorophyll-free thylakoids in <i>Chlorella</i> after clindamycin-treatment and in a temperature sensitive mutant of <i>Chlorella</i> G. Gallig	53
Genes affecting erythromycin resistance and sensitivity of <i>Chlamydomonas reinhardtii</i> chloroplast ribosomes L. Bogorad, J.N. Davidson and M.R. Hanson	61
Genetic control of chloroplast ribosome biogenesis in <i>Chlamydomonas</i> N.W. Gillham, J.E. Boynton, E.H. Harris, S.B. Fox and P.L. Bolen	69
<b>CONTROL OF FORMATION AND ASSEMBLY OF CHLOROPLAST CONSTITUENTS</b>	
Cellular origins of plastid membrane polypeptides in <i>Euglena</i> S. Bingham and J.A. Schiff	79
Synthesis of the major thylakoid polypeptides during greening of <i>Chlamydomonas reinhardtii</i> Y-1 J.K. Hooper	87
Relationship between chloroplastic metabolism and cytoplasmic translation G. Ledoigt and M. Lefort-Tran	95
Temperature-sensitivity of chloroplast ribosome formation in higher plants J. Feierabend	99
Temperature control of chloroplast development R.M. Smillie	103
Chlorophyll synthesis and the development of photosynthetic activity W.T. Griffiths, N.L. Morgan and R.E. Mapleston	111

Genetic regulation of chlorophyll synthesis analyzed with double mutants in barley A. Kahn, N. Avivi-Bleiser and D. von Wettstein	119
MITOCHONDRIAL ATPase COMPLEX	
Mutation in <i>Saccharomyces cerevisiae</i> mitochondrial F <sub>1</sub> leading to aurovertin resistance E. Agsteribbe, M. Douglas, E. Ebner, T.Y. Koh and G. Schatz	135
Mitochondrially encoded oligomycin-resistant mutants of <i>S. cerevisiae</i> : Structural integration of ATPase and phenotype M. Somlo and J. Cosson	143
Investigation of the oligomycin binding protein in yeast mitochondrial ATPase R.S. Criddle, C. Arulanandan, T. Edwards, R. Johnston, S. Scharf and R. Enns	151
Polypeptides encoded by mitochondrial genes in ascomycetes H. Küntzel, M.A. Marahiel, D.E. Leister and P. Nelson	159
Cytoplasmic synthesis of the dicyclohexylcarbodiimide-binding protein in <i>Neurospora crassa</i> W. Sebald, Th. Graf and G. Wild	167
Biochemical genetics of oxidative phosphorylation an approach to the reaction mechanism D.E. Griffiths	175
MITOCHONDRIAL RESPIRATORY COMPLEXES	
Analysis of the polypeptide chains of cytochrome oxidase from beef heart mitochondria C. Buse and G. Steffens	189
Partial sequence of a chloroform-methanol soluble polypeptide from <i>Neurospora</i> mitochondrial membranes W. Machleidt, R. Michel, W. Neupert and E. Wachter	195
Identification of an assembly intermediate of cytochrome oxidase in <i>Neurospora crassa</i> S. Werner and M. Neuner-Wild	199
The assembly of cytochrome <i>c</i> oxidase from <i>Saccharomyces cerevisiae</i> R.O. Poyton and E. McKemie	207
Structure and biosynthesis of cytochrome <i>c</i> oxidase F. Cabral, J. Saltzgaber, W. Birchmeier, D. Deters, T. Frey, C. Kohler and G. Schatz	215
Coordination of mitochondrial and cytoplasmic protein synthesis in <i>Neurospora crassa</i> W. Neupert and A. v. Rücker	231
The use of double mutant strains containing both heat- and cold-sensitive mutations in studies of mitochondrial biogenesis T. Mason, P. Boerner and C. Biron	239
Cold sensitivity of mitochondrial biogenesis in a nuclear mutant of <i>Neurospora crassa</i> R. Kientsch and S. Werner	247
Isolation and characterization of a cytochrome oxidase deficient mutant of <i>Neurospora crassa</i> K. Bruckmooser and S. Werner	253

Subunit structure and arrangement of mitochondrial cytochrome <i>b</i> H. Weiss and B. Ziganke	259
The <i>bc</i> <sub>1</sub> -complex from beef heart prepared by adsorption chromatography in Triton X-100 G. von Jagow, W.D. Engel, P. Riccio and H. Schägger	267
Complex III of yeast: Subunit composition and biosynthesis M.B. Katan and G.S.P. Groot	273
Purification and biogenesis of cytochrome <i>b</i> in bakers' yeast L.-F. Lin and D.S. Beattie	281
Biosynthesis of cytochrome <i>c</i> in the honey bee, <i>Apis mellifera</i> M. Osanai and H. Rembold	289
CHLOROPLAST DNA: GENES AND MOLECULES	
The circular diploid model of chloroplast DNA in <i>Chlamydomonas</i> R. Sager	295
On the search for a molecular mechanism of cytoplasmic inheritance: Past controversy, present progress and future outlook K.S. Chiang	305
Transmission, segregation and recombination of chloroplast genes in <i>Chlamydomonas</i> J.E. Boynton, N.W. Gillham, E.H. Harris, C.L. Tingle, K. Van Winkle- Swift and G.M.W. Adams	313
A uniparental mutant of <i>Chlamydomonas reinhardtii</i> with a variant thylakoid membrane polypeptide N.-H. Chua	323
Plastid distribution and plastid competition in higher plants and the induction of plastom mutations by nitroso-urea-compounds R. Hagemann	331
Structural and functional analysis of <i>Euglena gracilis</i> chloroplast DNA E. Stutz, E.J. Crouse, L. Graf, B. Jenni and H. Kopecka	339
Restriction endonuclease map of chloroplast DNA from <i>Euglena gracilis</i> P.W. Gray and R.B. Hallick	347
The location of rRNA genes on the restriction endonuclease map of the <i>Spinacia oleracea</i> chloroplast DNA R.G. Herrmann, H.-J. Bohnert, A. Driesel and G. Hobom	351
Analysis of the coding capacity of EcoRI restriction fragments of spinach chloroplast DNA P.R. Whitfeld, B.A. Atchison, W. Bottomley and C.J. Leaver	361
Physical and transcriptional mapping of <i>Zea mays</i> chloroplast DNA J.R. Bedbrook and L. Bogorad	369
Studies with chloroplast DNA-plasmid hybrids from <i>Chlamydomonas reinhardtii</i> J.-D. Rochaix	375
Replication of circular chloroplast DNA K.K. Tewari, R.D. Kolodner and W. Dobkin	379
Studies of the growth and replication of spinach chloroplasts and of the location and segregation of their DNA J.V. Possingham and R.J. Rose	387

## MITOCHONDRIAL DNA: GENES AND MOLECULES

Mechanisms and rules for transmission, recombination and segregation of mitochondrial genes in <i>Saccharomyces cerevisiae</i> B. Dujon and P.P. Slonimski	393
Confirmations and exceptions to the phage analogy model: Input bias, bud position effects, zygote heterogeneity, and uniparental inheritance P.S. Perlman, C.W. Birky, Jr., C.A. Demko and R.L. Strausberg	405
On homozygotization of mitochondrial mutations in <i>Saccharomyces cerevisiae</i> A. Putrament, R. Polakowska, H. Baranowska and A. Ejchart	415
Genetic determination of mitochondrial cytochrome <i>b</i> A. Tzagoloff, F. Foury and A. Akai	419
The isolation and simultaneous physical mapping of mitochondrial mutations affecting respiratory complexes J. Rytka, K.J. English, R.M. Hall, A.W. Linnane and H.B. Lukins	427
Genetic analysis of mitochondrial polymorphic proteins in yeast M.G. Douglas, R.L. Strausberg, P.S. Perlman and R.A. Butow	435
Regulation of cytochrome oxidase formation by mutations in a mitochondrial gene for cytochrome <i>b</i> P. Pajot, M.L. Wambier-Kluppel, Z. Kotylak and P.P. Slonimski	443
Mitochondrial genes determining cytochrome <i>b</i> (complex III) and cytochrome oxidase function G.S. Cobon, D.J. Groot Obbink, R.M. Hall, R. Maxwell, M. Murphy, J. Rytka and A.W. Linnane	453
Antimycin- and funiculosin-resistant mutants in <i>Saccharomyces cerevisiae</i> : New markers on the mitochondrial DNA B. Lang, G. Burger, W. Bandlow, F. Kaudewitz and R.J. Schweyen	461
Two mitochondrial antimycin A resistance loci in <i>Saccharomyces cerevisiae</i> E. Pratje and G. Michaelis	467
Mitochondrial inheritance of mucidin resistance in yeast J. Šubík	473
Behaviour of <i>Saccharomyces cerevisiae</i> mutant resistant to Janus Green A. Kruszewska and B. Szczesniak	479
Mitochondrial mutations conferring heat or cold sensitivity in <i>Saccharomyces cerevisiae</i> W.E. Lancashire	481
High spontaneous petite frequency strains of <i>Saccharomyces cerevisiae</i> generated in complementation tests G.D. Clark-Walker, K.M. Oakley, C.R. McArthur and G.L.G. Miklos	491
Extrachromosomal inheritance in a petite - negative yeast - <i>Schizosaccharomyces pombe</i> K. Wolf, G. Seitz, G. Lückemann, B. Lang, G. Burger, W. Bandlow and F. Kaudewitz	497
The mitochondrial genome of yeast: Organization and recombination G. Bernardi	503
The variability of the mitochondrial genome of <i>Saccharomyces</i> strains J.P.M. Sanders, C. Heyting and P. Borst	511
Restriction endonuclease mapping and analysis of grande and mutant yeast mitochondrial DNA R. Morimoto, A. Lewin, S. Merten and M. Rabinowitz	519
The control of mitochondrial DNA synthesis in yeast petite mutants P. Borst, C. Heyting and J.P.M. Sanders	525

A segment of mitochondrial DNA carrying oligomycin resistance K. Wakabayashi	535
Gene identification by coupled transcription-translation of yeast mitochondrial DNA A.F.M. Moorman and L.A. Grivell	539
Mitochondrial mutations that affect mitochondrial transfer ribonucleic acid in <i>Saccharomyces cerevisiae</i> G. Faye, M. Bolotin-Fukuhara and H. Fukuhara	547
Structure and genetics of the 2 $\mu$ m circular DNA in yeast M. Guerineau, C. Grandchamp and P.P. Slonimski	557
Electron microscopical analysis of native and cloned 2- $\mu$ m DNA from <i>Saccharomyces cerevisiae</i> C.P. Hollenberg and H.-D. Royer	565
The study of the genetic function of <i>Paramecium</i> mitochondrial DNA using species hybrids A. Tait, J.K.C. Knowles, J.C. Hardy and H. Lipps	569
Organization and expression of the mitochondrial genome in HeLa cells G. Attardi, M. Albring, F. Amalric, R. Gelfand, J. Griffith, D. Lynch C. Merkel, W. Murphy and D. Ojala	573
Functional organization and evolution of animal mitochondrial DNA W.B. Upholt and I.B. Dawid	587
Physical map and replication of rat mitochondrial DNA K. Koike, M. Kobayashi, S. Tanaka and H. Mizusawa	593
Measurement of the relative rate of mitochondrial DNA synthesis under experimentally varied conditions D. Bogenhagen and D.A. Clayton	597
Use of antibiotic inhibitors in studies of replication and repair of animal mitochondrial deoxyribonucleic acid G.G. Gause, Jr., V.S. Mikhailov, S.I. Tomarev and R.D. Zinovieva	605
Hormonal control of mitochondrial DNA replication in maturing oocytes M. Barat, C. Dufresne, H. Pinon, M. Tourte and J.-C. Mounolou	613
TRANSCRIPTION AND TRANSLATION APPARATUS OF CHLOROPLASTS	
<i>In vitro</i> transcription and translation of chloroplast DNA of <i>C. reinhardi</i> S.J. Surzycki, J.A. Surzycki and R. Lutz	621
Localization of the gene coding for the large subunit of ribulose bisphosphate carboxylase on the chloroplast genome of <i>Chlamydomonas</i> <i>reinhardi</i> S. Howell, P. Heizmann and S. Gelvin	625
Characterization of the RNA compounds synthesized by isolated chloroplasts H.J. Bohnert, A.J. Driesel and R.G. Herrmann	629
Incorporation of <sup>32</sup> P-orthophosphate into nucleoside 5'-triphosphates and RNA by isolated pea chloroplasts J. Bennett and Y. Milewska	637
Phylogenetic origin of chloroplast 16S ribosomal RNA D.E. Buetow, M.S. Kissil and L. Zablen	641
A sequence analysis of low-molecular-weight rRNA from chloroplasts of flowering plants T.A. Dyer and C.M. Bowman	645
Chloroplast ribosomal proteins of <i>Euglena gracilis</i> . Immunological studies G. Freyssinet, F. Morlé and V. Nigon	653

A chloroplast membrane fraction enriched in chloroplast ribosomes M.M. Margulies and J. Weistrop	657
The tRNAs and aminoacyl-tRNA synthetases of <i>Euglena</i> chloroplasts W.E. Barnett, S.D. Schwartzbach and L.I. Hecker	661
tRNAs and aminoacyl-tRNA synthetases in plant organelles J.H. Weil, G. Burkard, P. Guillemaut, G. Jeannin, R. Martin and A. Steinmetz	667
TRANSCRIPTION AND TRANSLATION APPARATUS OF MITOCHONDRIA	
Characterization and translation of yeast mitochondrial RNA F. Hendler, A. Halbreich, S. Jakovcic, J. Patzer, S. Merten and M. Rabinowitz	679
The mitochondrial RNAs of <i>Neurospora crassa</i> : Their function in translation and their relation to the mitochondrial genome A.M. Kroon, P. Terpstra, M. Holtrop, H. de Vries, C. van den Bogert, J. de Jonge and E. Agsteribbe	685
Dual origin of mRNA associated proteins in Ehrlich ascites mitochondria N.G. Avadhani, V.A. Aroskar, F.S. Lewis, G.J. Hansel and M.P. Wolf	697
Mitochondrial transcription in rat liver. Studies on the synthesis of poly(A)-containing RNA C. Saccone, P. Cantatore, G. Pepe, R. Gallerani, C. De Giorgi and C. De Benedetto	701
Properties and purification of poly(A) polymerase from rat liver mitochondria R. Gallerani, C. De Benedetto, C. De Giorgi and C. Saccone	709
The <i>poky</i> mutant of <i>Neurospora crassa</i> A.M. Lambowitz	713
The proteins of <i>Neurospora crassa</i> mitochondrial and cytoplasmic ribosomes H. de Vries and C. van den Bogert	721
Significance of 80-S ribosomes associated with <i>Neurospora crassa</i> mitochondria R. Michel, G. Hallermayer, M.A. Harmey, F. Miller and W. Neupert	725
Comparative studies of ribosomes from mitochondria, chloroplasts and cytoplasm. Morphology and electrophoretic behavior B.J. Stevens, J.-J. Cury, G. Ledoigt and J. André	731
Protein composition of the bovine mitochondrial ribosome T.W. O'Brien, D.E. Matthews and N.D. Denslow	741
Transfer RNAs of yeast mitochondria N.C. Martin and M. Rabinowitz	749
Isoacceptor tRNA species in yeast mitochondria. Methionine and formyl- methionine specific tRNAs coded by mitochondrial DNA R. Martin, J.M. Schneller, A.J.C. Stahl and G. Dirheimer	755
Isoaccepting tRNA <sup>Ser</sup> in mitochondria from <i>Saccharomyces cerevisiae</i> : Mitochondrially coded and cytoplasmic species G. Baldacci, C. Falcone, L. Frontali, G. Macino and C. Palleschi	759
Imported tRNA: Its synthetase as a probably transport protein Y. Suyama and J. Hamada	763
Characterization of rRNA and tRNA from mitochondria of <i>Locusta migratoria</i> H. Feldmann and W. Kleinow	771
Immunological study of yeast mitochondrial phenylalanyl-tRNA synthetase J.M. Schneller, C. Schneller and A.J.C. Stahl	775
Mitochondrial protein synthesis in higher plants C.J. Leaver	779

## GENERAL ASPECTS OF MITOCHONDRIAL BIOGENESIS

Mitochondrial phospholipid synthesis and the phospholipid exchange proteins K.W.A. Wirtz, R.H. Lumb, H.H. Kamp, G.M. Helmkamp, H. van den Bosch and L.L.M. van Deenen	785
Incorporation of mitochondrial membrane proteins into liposomes G.D. Eytan	793
The role of mitochondria-bound 80S ribosomes in mitochondrial biogenesis W.F. Bennett, A. Gutierrez-Hartmann and R.A. Butow	801
Studies on the synthesis of mitochondrial proteins in the cytoplasm and on their transport into the mitochondrion G. Hallermayer and W. Neupert	807
<i>In vitro</i> synthesis and transport into mitochondria of cytoplasmically translated proteins M.A. Harmey, G. Hallermayer and W. Neupert	813
Specific labelling of mitochondrially synthesized proteins in yeast cells in the absence of antibiotics W. Bandlow	819
Integration and disintegration of proteins synthesized in mitochondria H.-D. Hofmann, E. Hundt and B. Kadenbach	827
Synthesis of mitochondrial DNA, -proteins and -phospholipids in the young sea urchin embryo <i>Sphaerechinus granularis</i> H. Bresch	831
Inhibition of cytoplasmic protein synthesis by mitochondrial soluble factors in rat liver and Walker carcinosarcoma N. González-Cadavid, B. Dorta and A. Carmona	835
Mammalian embryos: A model for studying the dependence of growth and differentiation processes on mitochondrial biogenesis and function R. Bass	843
Unmasking of mitochondrial precursors stored in the yolk platelets of <i>Artemia salina</i> dormant gastrulae C.G. Vallejo and R. Marco	847
Screening tests for suppressors of respiratory deficient mutants in <i>Schizosaccharomyces pombe</i> and model for a mitochondrial partial suppression of nuclear pleiotropic strain A. Goffeau, F. Labaille, O. Mohar and A.-M. Colson	851
Respiration deficient mutants with intact mitochondrial genomes: Casting a wider net H.R. Mahler, T. Bilinski, D. Miller, D. Hanson, P.S. Perlman and C.A. Demko	857
Assembly of the cyanide-insensitive respiratory pathway in <i>Neurospora crassa</i> D.L. Edwards, J.H. Chalmers, Jr., H.J. Guzik and J.T. Warden	865
Physiological and genetical analysis of the respiratory chain of <i>Paramecium</i> J. Doussière, A. Adoutte, A. Sainsard, F. Ruiz, J. Beisson and P. Vignais	873
Genetic control of glycerol-3-phosphate dehydrogenase synthesis in <i>Neurospora</i> J.B. Courtright	881
Primary antimitochondrial activity of carcinogens in <i>Saccharomyces cerevisiae</i> V. Egilsson, I.H. Evans and D. Wilkie	885
Author index	893



COORDINATION OF MITOCHONDRIAL AND CYTOPLASMIC PROTEIN  
SYNTHESIS IN NEUROSPORA CRASSA

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INTRODUCTION

The formation of some multi-component mitochondrial enzyme complexes such as cytochrome oxidase requires the cooperation of cytoplasmic and mitochondrial translation. Products from both systems are integrated to form functional enzyme complexes with definite subunit stoichiometry. Control of the coordination of these two distinct systems may be exerted at many levels, so that a general study of the coordination presents many difficulties. Recently, a detailed review on these problems was presented by SCHATZ and MASON (1).

In the present investigation we have limited ourselves to one facet of the interaction between cytoplasmic and mitochondrial translation, viz. the synthesis and membrane integration of mitochondrial translation products under conditions when cytoplasmic translation does not operate. We have linked the general aspects of translation to the synthesis and assembly of specific mitochondrial products, namely the subunits of cytochrome oxidase.

RESULTS

1. Amino acid incorporation into mitochondrial membrane proteins of cells with blocked cytoplasmic translation

Cells were prelabelled with  $^{14}\text{C}$ -leucine to obtain homogeneous labelling of total cellular protein. At 16 hrs growth (log phase) cycloheximide (CHI) was added. After the times indicated in Fig. 1, aliquots were withdrawn, mitochondria prepared and the  $^3\text{H}/^{14}\text{C}$ -ratios of the mitochondrial membrane protein determined. Labeling with  $^3\text{H}$ -leucine was performed at a very low added leucine concn. (0.3 nmol/mg cell protein), a procedure usually employed to effectively label mitochondrial translation products.

When leucine was added 2.5 min after addition of CHI, incorporation proceeds for about 10-15 min and then stops. When the cells were preincubated for 62.5 min, incorporation of  $^3\text{H}$ -labelled leucine still takes place and furthermore continues for a longer time span. The level of incorporation is lower, however, than in the cells exposed to CHI for a shorter time.

This observation leads to suggest that the incorporation of  $^3\text{H}$ -leucine in the presence of CHI is not a true measure of the rate of mitochondrial protein

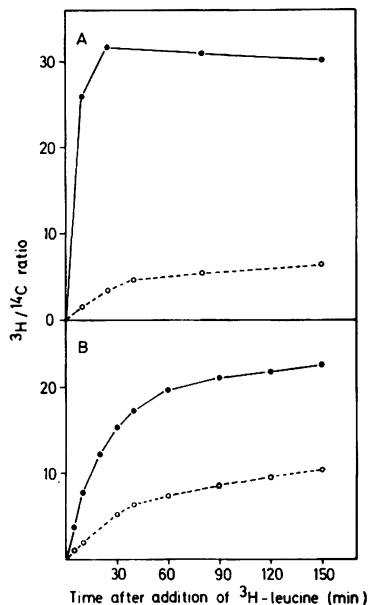


Fig. 1. Time course of  $^3\text{H}$ -leucine incorporation into mitochondria in whole cells as a function of time of incubation in cycloheximide (0.1 mg/ml). Cells were prelabelled with  $^{14}\text{C}$ -leucine. Incorporation was carried out with (A) low and (B) high concn. of added leucine (see text)

—●—●— 2.5 min CHI preincub.;  
-o---o- 62.5 min CHI preincub.

cells to CHI. Leucine concn. increases by a factor of about 10 during the first 90 min. (b) It was found that leucine is rapidly metabolized, a large part of the  $^3\text{H}$ -radioactivity appearing in  $\alpha$ -ketoisocaproic acid.

The effect of preincubation of cells with CHI on the protein synthesis of isolated mitochondria was measured and the results are presented in Fig. 2. The mitochondria from cells kept for 1 hr in the presence of CHI displayed a rate of incorporation which was very similar to that of mitochondria from cells treated for 2.5 min with CHI (or to that of mitochondria from untreated cells). Even after treatment of cells for 4 hrs the rate of incorporation was not reduced by more than about 30%. Similar observations were made by Sebald et al. (3).

It appears therefore that the mitochondrial protein synthesizing system is not essentially linked to the cytoplasmic system as suggested by some investigators (see ref. (1)).

## 2. Translational activity of mitochondrial ribosomes in cycloheximide blocked cells

In view of the unexpected lack of tight coupling of cytoplasmic and

synthesis (see also ref. (2)). This lack of validity may be related to changes in pool size and to metabolism of intracellular leucine.

To obviate these effects, leucine was added at concns. sufficiently high to offset these changes. At the concn. added (500 nmol/mg cell protein), leucine uptake still takes place and within 10 min about 75% of the added radioactivity was accumulated in the cells. At this high added leucine concn. incorporation proceeds in both the short and long time CHI preincubation. Incorporation even continues at 150 min after addition of  $^3\text{H}$ -leucine. When the initial rate of the 62.5 min preincubation sample is compared with the rate observed at 60 min with the 2.5 min preincubated sample, it can be seen that the rates are not equal. The longer preincubated sample shows a rate faster by a factor of six than the 2.5 min sample. Therefore, even at this high added leucine concn. the incorporation does not truly reflect the real rate of protein synthesis. This discrepancy between real and apparent rates can be explained by the following observations:

(a) The intracellular concns. of amino acids were found to strongly increase after exposing

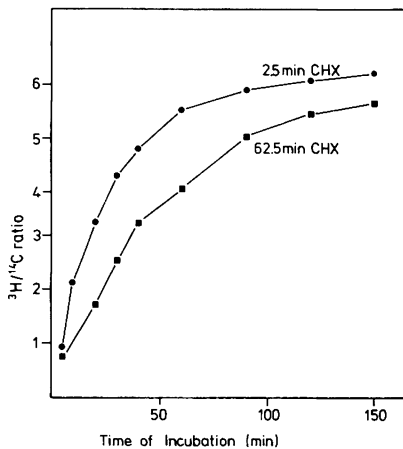


Fig. 2. Time course of  $^3\text{H}$ -leucine incorporation into isolated mitochondria from  $^{14}\text{C}$ -prelabelled cells and the effect of preincubation of cells in cycloheximide on this incorporation. For experimental details see ref.(8).

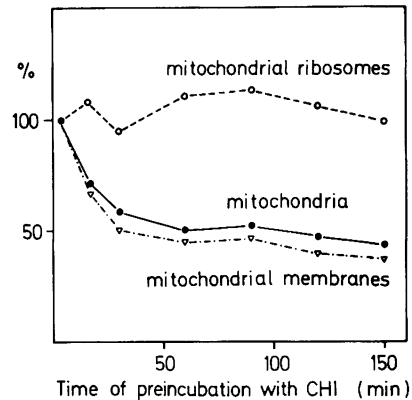


Fig. 3. The effect of cycloheximide preincubation on the *in vivo* incorporation of leucine into different mitochondrial fractions. Whole cells were pulse labelled with  $^3\text{H}$ -leucine for 5 min at the indicated times. All incorporation measurements were corrected for the intracellular leucine pool. The 100% value represents incorporation after 2.5 min preincub.

mitochondrial protein synthesis, the rate of translation on mitochondrial ribosomes was measured by a different approach. Cells were preincubated for different time periods in the presence of CHI and then pulsed with  $^3\text{H}$ -leucine for 5 min. To aliquots of these samples a chase of unlabelled leucine was given for 10 min. Mitochondrial ribosomes were then isolated from the pulsed cells. Mitochondria and mitochondrial membranes were isolated from the chased cells.  $^3\text{H}$ -radioactivity was measured in these fractions and the actual rates of leucine incorporation were determined by correcting for the intracellular leucine concns. at the different times. The result is shown in Fig. 3. The amount of leucine in nascent polypeptide chains is found to be constant at all times of CHI preincubation. The amount of leucine incorporated into mitochondrial membrane protein however decreases during the first 30 min to about 50% compared to the control. (Preincubation for 2.5 min is taken as a control because the addition of CHI is necessary to assess specifically radioactivity in mitochondrial translation products). After 30 min preincubation incorporation stays constant at a level of about 40%.

To measure the velocity of replacement of labelled nascent chains on mitochondrial ribosomes by unlabelled ones, i.e. to determine the rate of elongation, cells were pulse labelled in the presence of CHI and then exposed to a leucine chase of increasing times. The rate of elongation was found to be the same, irrespective of

whether the cells were preincubated with CHI for 2.5 min or for 62.5 min before the pulse. Furthermore, when ribosomes from corresponding mitochondria were analysed by gradient centrifugation, the same distribution of radioactive nascent chains was seen among mono- and polymeric ribosomes, more than 80% of the  $^3\text{H}$ -leucine being associated with polymers.

These observations firmly support the conclusion that mitochondrial translation continues in the absence of cytoplasmic translation. They also suggest that at least part of the polypeptides formed, are integrated into the mitochondrial membrane. They further suggest that part of the translation products are degraded since completed mitochondrial translation products were neither detected in the mitochondrial matrix nor in the postmitochondrial supernatant.

### 3. Translation products formed in mitochondria in cycloheximide blocked cells

Cells homogeneously labelled with  $^{14}\text{C}$ -leucine were exposed to CHI for 2.5 min, then a pulse of  $^3\text{H}$ -leucine was given for 2.5 min, followed by a 60 min chase. This experiment was made in an attempt to catch the two systems as close to equilibrium as possible. Mitochondria were isolated and subjected to SDS gel electrophoresis (Fig. 4).  $^{14}\text{C}$ -labelled total mitochondrial proteins are mainly found at apparent

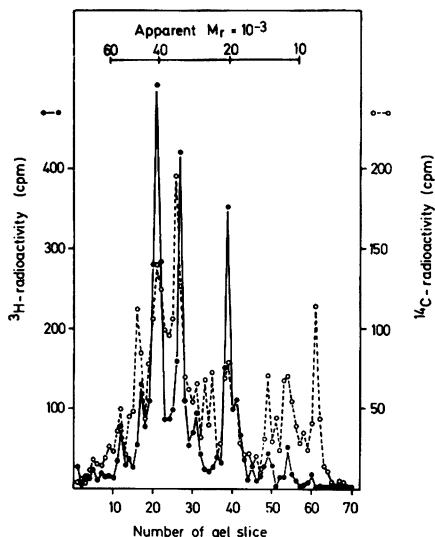


Fig. 4. SDS gel electrophoretic analysis of mitochondrial translation products synthesized in  $^{14}\text{C}$ -prelabelled cells during a short pulse.

Cells were pretreated with cycloheximide for 2.5 min,  $^3\text{H}$ -leucine was then applied for 2.5 min and chased for 60 min with unlabelled leucine.

For experimental details see (8).

$^{14}\text{C}$ -labelled total mitochondrial proteins are mainly found at apparent MWs between 60,000 and 10,000. Well-defined mitochondrial translation products are displayed by the  $^3\text{H}$ -label, which can be attributed to cytochrome oxidase, cytochrome b and OS-ATPase. It should be especially noted that the proportion of mitochondrial translation products with low apparent MWs (10,000 and less) is extremely low.

In a further experiment, cells were preincubated with CHI for 2.5 min, 62.5 min and 122.5 min, respectively, then  $^3\text{H}$ -leucine was incorporated for 45 min and a chase of unlabelled leucine was given for further 10 min. The SDS gel electrophoretic analysis of the isolated mitochondria is presented in Fig. 5.

Clearly, the same products are formed in cells, in which cytoplasmic translation was blocked for 120 min compared to those blocked only for 2.5 min. However, the translation products with apparent MWs of more than 20,000 are present in lower proportion in cells treated with

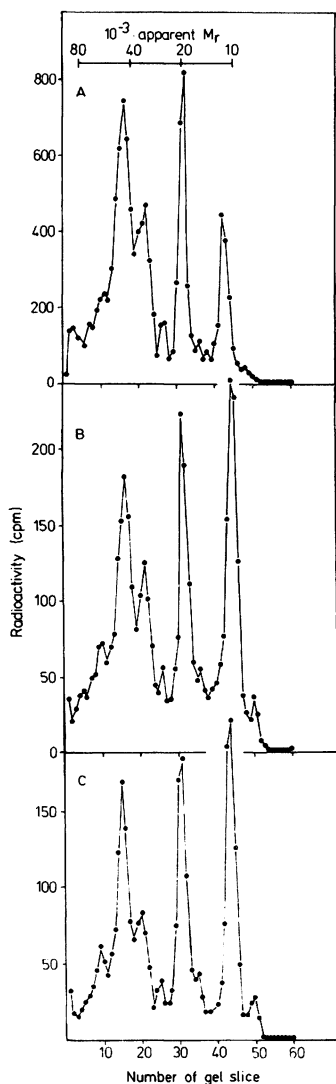


Fig. 5. SDS gel electrophoretic analysis of mitochondrial translation products synthesized after different times of pretreatment of cells with cycloheximide. Cells were incubated with cycloheximide for (A) 2.5 min, (B) 62.5 min and (C) 122.5 min.  $^3\text{H}$ -leucine was added to the cells and after further 45 min incubation a chase of unlabelled leucine was given for 10 min.

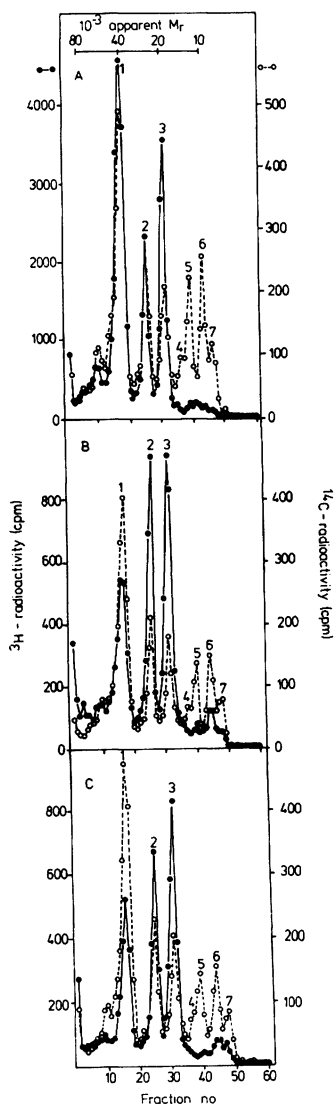


Fig. 6. SDS gel electrophoretic analysis of cytochrome oxidase immunoprecipitated from the total homogenate of cells which were prelabelled with  $^{14}\text{C}$ -leucine. Incubation of cells with cycloheximide and labelling with  $^3\text{H}$ -leucine was carried out as described in legend to Fig. 5. Samples A, B and C correspond to samples A, B and C in Fig. 5.

cycloheximide for longer time. Furthermore, translation products are found in increasing amounts in the low apparent MW region (11,000 and less) after longer preincubation of cells with CHI (cf. Fig. 4).

These observations suggest that original translation products are degraded to low MW components and that part of the degradation products are retained in the membrane. When mitochondrial translation products are formed only during a short pulse period immediately after blocking cytoplasmic translation, this degradation appears to be negligible. It is tempting to suggest that under these conditions cytoplasmically synthesized partner proteins are available in such amounts that the newly formed mitochondrial partners can be assembled to functional complexes and thereby preserved in the mitochondrial membrane. After longer exposure of the cells to CHI, the pools of cytoplasmic partner proteins are exhausted, the mitochondrially translated components are continued to be synthesized but only part of them can be integrated into the membrane. Another part may be degraded probably before being integrated.

#### 4. Synthesis and assembly of cytochrome oxidase components in cells with blocked cytoplasmic translation

We next studied the influence of the availability of cytoplasmically synthesized subunits of cytochrome oxidase on the production and insertion of the mitochondrially synthesized subunits.  $^{14}\text{C}$ -homogeneously labelled cells were exposed to  $^3\text{H}$ -leucine in the presence of CHI as described in Fig. 5. After incubation, the cells were homogenized and extracted with a buffer containing Triton X-100. Cytochrome oxidase was precipitated from these extracts with an antibody against the whole enzyme. SDS gel electrophoretic analysis of the immunoprecipitates is shown in Fig. 6.

The  $^{14}\text{C}$ -radioactivity on the gels displays the seven subunits of the enzyme. After short CHI treatment, the three mitochondrially synthesized subunits (1-3) are found to be labelled with  $^3\text{H}$  in about the same ratio, at which they are present in the  $^{14}\text{C}$ -profile, subunit 3 being somewhat higher. After 62.5 min and 122.5 min preincubation of cells still considerable synthesis of subunits 1-3 occurs. The proportions of the three subunits however change. Subunits 1 and 2 decrease with respect to subunit 3. This is in close agreement with the findings with whole mitochondrial membranes (cf. Fig. 5). In the samples preincubated with CHI for longer times, lower MW material is precipitated which may represent breakdown products of subunits 1-3 which are recognized by the antibody.

These data show that immunoprecipitation of cytochrome oxidase from total cell homogenate appears to capture not only fully assembled enzyme but also precursor polypeptides. However, if immunoprecipitation was carried out with isolated mitochondria, a quite different picture was observed. The labelling pattern in this case resembles that of the chemically isolated complex (4), showing that after short CHI exposure only small amounts of  $^3\text{H}$ -labelled subunits 1 and 2 are found

(see also (5)), whereas after 62.5 min and 122.5 min CHI none of these subunits were detected. This is in agreement with the data of Schwab et al. (4), that the pool of cytoplasmically formed subunits allows assembly of total enzyme only for a few min. Subunit 3 may get into the immunoprecipitate by an exchange process or by specific attachment to a complete enzyme complex.

#### DISCUSSION

The data presented show that mitochondrial translation in *Neurospora* continues even when cytoplasmic translation is shut off for periods as long as 2 hrs, i.e. one half of a doubling period of the cells. We find no evidence for a mechanism which regulates mitochondrial translation in such a way that synthesis of mitochondrial translation products is stopped, when these products cannot any longer become assembled to functional membrane complexes. With respect to the question, what then happens to these products, the experimental data presented would suggest that (a) part of these products are taken up into the membrane and remain there, and (b) another part of the translation products is degraded to lower MW components which partly remain in the mitochondrial membrane. Such a mechanism would require the existence of specific mitochondrial proteinases, since it was observed that pre-existing mitochondrial proteins are not degraded when the cells are kept in the presence of CHI. Based on experiments which were carried out on isolated mitochondria to investigate the formation of cytochrome oxidase components in a reconstituted system, we have arrived at a similar conclusion, viz. that mitochondrial translation products are formed in the absence of cytoplasmic partner proteins but are subject to proteolytic breakdown, probably since they cannot be assembled to functional complexes (6).

Cycloheximide treatment of cells may result in a situation analogous to that in some nuclear mutants. In such mutants defective assembly of cytochrome oxidase or other complexes may result from the inability of cytoplasmic protein synthesis to provide certain partner proteins (for review see (1)). Specific degradation of mitochondrially synthesized partner proteins may occur in these mutants in the same way as in CHI treated cells.

Further studies of the regulatory processes on the level of transcription and translation seem desirable (e.g. (7)), however our limited knowledge of these reactions imposes severe experimental limitations.

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