Overexpression of alcohol oxidase in Pichia pastoris

M J. de Hoop¹, J. Cregg², I. Keizer-Gunnink³, Klaas Sjollema³, M Veenhuis³ and G. Ab¹

¹Laboratory of Biochemistry, University of Groningen, Nijenborgh 16, 9747 AG Groningen, The Netherlands, ²Dept of Chemical and Biological Sciences, Oregon Graduate Institute of Science and Technology, 19600 Von Neumann Dr, Beaveiton OR 97006-1999, USA and ³Laboratory for Electron Microscopy, University of Groningen, Kerklaan 30, 9751 NN Groningen, The Netherlands

Received 8 July 1991

The protein import capacity of peroxisomes in methylotrophic yeasts was studied using *Pichia pastoris* containing one or two extra copies of the gene encoding the peroxisomal protein alcohol oxidase. The alcohol oxidase overproduced in this strain was only partially imported and assembled into the active, octameric form of the protein. The excess remained in the cytosol as protein aggregates composed of monomers. These results are discussed in view of the possible application of peroxisomes as storage compartments for heterologous proteins.

Alcohol oxidase, Methylotrophic yeast, Peroxisome

I. INTRODUCTION

Peroxisomes are single-membrane bounded organelles containing a matrix of proteins, generally including catalase and one or more hydrogen peroxide-producing oxidases They are versatile organelles that can vary in number, size and protein content [1,2] In yeast, prolifeiation of peroxisomes is dependent on growth conditions; some of the enzymes involved in the metabolism of particular C- and N-sources in the medium accumulate in the organelle Therefore, the composition of peioxisomes can be easily manipulated in yeast [3-6]. Stiong proliferation of peroxisomes occurs for example when methylotrophic yeasts are shifted from glucose to methanol as the sole source of carbon and energy Under certain conditions, these peroxisomes can occupy up to 80% of the cytosolic volume [7] These peroxisomes have a crystalline matrix composed of alcohol oxidase octamers, which is the active form of this protein [8,9] Also present in the peroxisomes, although not as abundant as alcohol oxidase, are 2 other key enzymes of methanol metabolism, namely catalase and dihydroxyacetone synthase [10]

The import capacity of peroxisomes present in methanol-cultured yeast may offer the possibility to store heterologously expressed proteins, equipped with the appropriate targetting signals, into peroxisomes [11]. Peroxisomal localization could have the advantage that the proteins are kept separate from proteolytic activities in the cytosol This system may have a potential value for storage of heterologously expressed proteins which

Abbreviation Hp-AOX, the alcohol oxidase gene from Hansenula polymorpha

Correspondence address G Ab, Laboratory of Biochemistry, University of Groningen, Nijenborgh 16, 9747 AG Groningen. The Netherlands Fax (31) (50) 634165 are susceptible to proteolysis. In a first attempt to analyze the import capacity of peroxisomes present in methanol-cultured cells, we overexpressed alcohol oxidase and analyzed to what extent the enzyme was imported.

2 MATERIALS AND METHODS

21 Plasmid construction, yeast transformation and genetic analysis A 2 5-kb fragment containing the coding sequence of the alcohol oxidase gene from Hansenula polymorpha (Hp-AOX) [12], flanked by 10 bp 5' and 250 bp 3' untranslated sequences, was inserted into the BamHI site of vector pAOBam This is an expression vector derived from pAO804 [13] by insertion of a BamHI linker (AATTGGATCC) into the unique EcoRI site In the final construct, the Hp-AOX gene was located between the promoter and terminator of the AOXI gene of Pichua pastoris

A transformation system of the related methylotrophic yeast *Pichia* pastoi is [14] was used for overexpression of alcohol oxidase Transformation of *Pichia pastoris* GS115 (*his4*) was performed as described by Cregg et al [14], and was based on complementation of histidine auxotrophy To force homologous integration into the defective *his4* gene, the plasmid DNA was linearizedwith *Stu*I, which cuts approximately in the middle of the *HiS4* gene present in pAOBam (Fig 1A) Transformants were screened for proper integration of the plasmid by Southern blotting The blot was screened with random-primed [α -³²PldCTP-labelled [15] plasmid pYM4, which is pBR322 containing the *HIS4* gene of *P pastoris* [14]

2.2 Preparation of cell extracts, measurement of activity and octamerization

A culture of transformed cells, grown on Difco's Yeast Nitrogen Base without amino acids and with 0 5% glycerol to an OD_{600} of 2-3, was 10-fold diluted with medium containing 1% methanol as the sole carbon source After 16 h, cells were harvested (OD_{600} =2-2 5) and extracts were prepared [8] Alcohol oxidase activity was assayed as described [16] Monomeric and octameric alcohol oxidase were separated by sucrose gradient centrifugation [8] and visualised by Western blotting [17] The antibodies used were either a polyclonal antibody raised against denatured alcohol oxidase of *Hansenula polymorpha* or a monoclonal antibody (OAO11) The polyclonal antibody recognized both alcohol oxidase of *P pastoris* and *H polymorpha*, while the monoclonal antibody was specific for alcohol oxidase from *H polymorpha* (M de Hoop, unpublished results)

23 Immuno-electronmicroscopy

The intracellular localization of alcohol oxidase and dihydro vacetone synthase was determined by immunogoid labelling on thin sections of Lowicryl-embedded cells [10]

3 RESULTS

31. Multi-copy integrations in Pichia pastoris GS115

To generate a strain that overproduces alcohol oxidase when cultured on methanol, P pastoris GS115 was equipped with additional copies of the alcohol oxidase gene from the related yeast H polymorpha The alcohol oxidase gene was placed between the promoter and the terminator of the P pastoris alcohol oxidase gene The recombinant plasmid was targeted into the defective his4 gene of the host strain GS115. To favour homologous recombination, the plasmid was linearized in the HIS4 gene prior to transformation (Fig. 1A). DNA from histidine prototrophic transformants was digested with Bg/II and analyzed by Southern blotting using pYM4 as a probe. This plasmid contained the HIS4 gene and the *E* coli gene encoding β -lactamase [14] Host DNA yielded one band of 3 kb representing the defective his4 gene (Fig 1A and Fig. 2, lane 1) Singlecopy integrations in the his4 locus resulted in the replacement of the 3-kb band by a number of new bands (Fig 1B and Fig. 2, lane 2). Knowing the positions of the BgIII sites, we could identify a band of 2 4 kb as the fragment containing the β -lactamase gene. The bands of 4.0 and 4.5 kb hybridize because they contain HIS4 sequences plus different parts of the plasmid An additional band of 55 kb and an intensified 2.4 kb band indicated a double-copy integration of the plasmid in the defective his4 gene (Fig. 2, lane 3)



Fig 1 Gene targeting of the *H* polymorpha alcohol oxidase gene into the hist locus of *P* pastoris Panel (Λ) depicts the transformation vector (pAOBam) containing the *Hp-AOX* gene (arrow) between the promoter (P) and terminator (T) of the *P* pastoris AOXI gene. Homologous recombination of the plasmid, linearized in the *HIS* sequence, with the his locus (dashed) of the *P* pastoris genome (stippled) is indicated Panel (B) shows the gene configuration after integration in the hist locus. The positions (arrows) and lenght of *Bg/II* fragments (B) are indicated



Fig 2 Southern analysis of *Pichia pastoris* GS115 cells containing no (1), one (2) or two copies (3) of the Hp-AOX gene integrated in the *his4* gene. The *Bg*/II fragments hybridizing with plasmid pYM4 are shown

32 Overexpression of alcohol oxidase

Providing *P* pastoris with additional copies of the Hp-AOX gene resulted in higher alcohol oxidase expression levels upon growth on methanol Whereas the increase in activity with the first additional copy was 73%, that with a second copy was considerably smaller (Table I) To examine the localization of the alcohol oxidase, we performed immunogold labeling on Lowicryl-embedded cell material Fig 3A shows the localization of endogenous alcohol oxidase in P pastoris GS115 In these methanol-grown cells, the protein was exclusively located in the peroxisomes; no alcohol oxidase could be found outside the peroxisomes This is in contrast with GS115 cells containing copies of the AOX gene of H polymorpha. In these cells, alcohol oxidase-immunoreactive material was also found outside the peroxisomes, often in irregular, electron-dense structures not surrounded by a membrane (Fig. 3B,C and D)

Extracts of wild-type GS115 and GS115 containing a double-copy integration of the Hp-AOX gene, cultured on methanol, were analyzed by sucrose gradient centrifugation In GS115, only octameric alcohol oxidase could be detected (Fig 4A) The strain, equipped

Table 1

Alcohol oxidase activities in *Pichia pastoris* GS115 containing additional copies of *Hp-AOX*

Yeast strain	Hp AOX copies	AO activity*
GS115	0	085±010
GS115 sc**	1	147±012
Gs115 dc**	2	1.89 ± 0.06

*Activity in µmol H₂O₂ produced per mg protein per min at 37°C, **se means single copy, de means double copy of the *Hp-AOX* gene



Fig 3 Immuno-electronmicloscopy of *Pichia pastoris* GS115 (A. 15 400×) and a transformant containing two copies of the *Hp-AOX* gene (B 31 000 ×, C, 16 800×, and D, 12 600×) The localization of alcohol oxidase on thin sections of methanol-induced transformants is marked by 10-nm gold particles. Organelles are indicated as follows, m = mitochondrion, n = nucleus, p = peroxisome and <math>v = vacuolea = cytosolic aggregate of alcohol oxidase

with two copies of the Hp-AOX gene, contained octameric alcohol oxidase as well as alcohol oxidase-immunoreactive material sedimenting at a rate roughly similar to that of bovine serum albumin run in a parallel



Fig 4 Obigometrization of alcohol oxidase in *Puchia pastoris* GS115 equipped with two copies of the *Hp-AOX* gene. Western blots of sucrose gradient fractions (lane 1 bottom, lane 6 top) of cell lysates are shown A. Western blot of a cell lysate ($200\,\mu$ g) of non-transformed *Puchia pastoris* GS115 probed with polyclonal antibody against alcohol oxidase. B. Western blot containing cell lysate ($100\,\mu$ g) of transformant GS115 overexpressing alcohol oxidase probed with polyclonal antibody against alcohol oxidase. C. identical to (B) but probed with monoclonal antibodies which preferentially recognized alcohol oxid ise from *Hansenula polymorpha*. 'o' indicates position of octameric molecules, while 'm' represents position of monomeric alcohol oxidase. Only the relevant parts of the blots are shown.



Fig 5 Western blots of methanol-induced cell lysates from *Pichia* pastoris (lane 1, 25 μ g) or *Hansenula polymorpha* (lane 2, 25 μ g) probed with polyclonal antibodies made against denatured alcohol oxidase from *Hansenula polymorpha* (A) or the monoclonal antibody OAO11 which preferentially recognized *Hansenula polymorpha* alcohol oxidase (B)

gradient (data not shown) suggesting that it represented monomeric alcohol oxidase (Fig. 4B) The amount of the monomeric alcohol oxidase was estimated to be 20-30% of the total alcohol oxidase. Discrimination of the import system against heterologous alcohol oxidase did not occui This can be concluded from a Western blot analysis using a monoclonal antibody which is specific for alcohol oxidase from Hansenula and does not react with alcohol oxidase from Pichia (Fig. 5). Clearly the alcohol oxidase of Hansenula assembled into octamers (Fig 4C) as did the endogenous alcohol oxidase from P. pastoris (Fig. 4A). Moreover, we have found earlier that peroxisomes of P pastoris are able to recognize and accumulate alcohol oxidase from H polymorpha When the Hp-AOX gene was expressed in a Pichia strain in which both endogenous alcohol oxidase genes were disrupted, all alcohol oxidase was localized exclusively inside peroxisomes and was in the active octamenc form (M de Hoop, unpublished results). Fig 4C shows that the alcohol oxidase of *Hansenula* was able to assemble into octamers in P pastoris Since octamers are normally present inside peroxisomes only [8,16], we conclude that the octameric alcohol oxidasc from Hansenula is localized inside the organelle The overexpression of alcohol oxidase had no effect on import of other matrix proteins, e.g. dihydroxyacetone synthase After immunogold-labelling with antibodies against dihydroxyacctone synthase, gold particles were exclusively confined to the peroxisomal matrix (data not shown)

4 DISCUSSION

Methylotrophic yeasts are an attractive host for the expression of heterologous proteins for several reasons Firstly, methanol is a cheap C-source Secondly, methylotrophic yeasts can be grown to high cell densities (130 g dry cell weight/l) in continuous culture [18]. Thirdly, the promoter of the highly expressed and tightly regulated alcohol oxidase gene can be used to drive synthesis of a foreign protein. Finally, integrating expression vectors yielding stable transformants have been designed [13]. The peroxisomes present in methanol-cultured yeast might add yet another advantage to this expression system, because the organelles may be used for storage of heterologously expressed proteins known to be susceptible to proteolytic cleavage.

The results presented here show that alcohol oxidase expressed from several gene copies in P pastoris is not fully imported and assembled into peroxisomes Obviously, the balance between alcohol oxidase synthesis and its post-translational import is disturbed in these cells, suggesting that saturation of either the import capacity of the peroxisomal space or of the peroxisomal import machinery had occurred Both situations should result in a cytosolic accumulation of all peroxisomal matrix proteins. A similar situation exists in peroxisome-deficient cells, where peroxisomal matrix enzymes are located in the cytosol. In these cells cytosolic crystals composed of active alcohol oxidase are observed [19]. Such alcohol oxidase crystalloids were never observed in the cytosol of strains described in this study and also dihydroxyacetone synthase was still only detectable in the matrix and was not observed in the cytosol. Therefore, we believe that the rate of alcohol oxidase import versus synthesis is limiting, rather than the capacity of peroxisomes for protein storage The fact that peroxisomes can occupy 80% of cell volume in methanollimited continuous cultures [20] demonstrates that peroxisomes should have a larger storage capacity than seen in the multi-copy gene experiments described here Even under the conditions where peroxisomes do not proliferate such as glycerol [16] or glucose [21], single organelles appear to have the ability to internalize the high amounts of alcohol oxidase synthesized in transformants containing multi-copy plasmids Because in our experiments alcohol oxidase is being expressed from multiple copies of the strong alcohol oxidase promoter, it is reasonable to assume that the rate of alcohol oxidase synthesis is exceptionally high in these strains Why would a rapid rate of synthesis cause saturation of the import system and result in aggregation in the cytosol? Too high an expression rate might oversaturate the capacity of cytosolic unfoldases [22] in their task of holding alcohol oxidase monomers in an import-competent conformation. This might result in the formation of alcohol oxidase aggregates as observed in our studies Once aggregated, the alcohol oxidase is probably no longer suitable for import

Concerning the prospects to compartmentalize heterologous proteins in peroxisomes, one has to take into account that peroxisomes have a large storage capacity, but that the efficiency of import may be inversely related to rate of synthesis of peroxisomal proteins Fuithermore, it is also likely that different heterologous proteins are imported at different rates and therefore, it may be necessary to adjust expression rates for optimal compartmentalization of each protein To improve import of heterologous proteins into peroxisomes, either the expression of endogenous peroxisomal proteins has to be reduced, or the import machinery must be further increased by e.g. overexpression of chaperones, like members of the HSP70 family [22,23]

Acknowledgements We thank Dr M A G Gleeson for his interest and stimulating discussions and are grateful to Prof Dr W Harder for critically reading the manuscript A plasmid containing the Hp-AOXgene was kindly provided by Drs B Distel and A Ledeboer Part of this work was carried out at the Salk Institute Biotechnology and Industrial Associates, Inc (SIBIA), La Jolla, CA, USA in the group of Dr G Thill We would like to thank him for giving M de Hoop this opportunity This was supported by a Wood/Whelan Research Fellowship from the International Union of Biochemistry and the 'Groninger Universiteitsfonds'

REFERENCES

- Bremer, J., Osmundsen, H., Christiansen, R.Z. and Borreback, B (1981) Methods Enzymol 72, 506–520
- [2] Tolbert, N E (1981) Ann Rev Biochem 50, 133-157
- [3] Fukui, S., Tanaka, A., Kawamoto, S., Yashura, S., Teranishi, Y and Osumi, M (1975) J Bacteriol 123, 317–328
- [4] Zwart, K, Veenhuis, M, van Dijken, JP and Harder, W (1980) Arch Microbiol 126, 117–126
- [5] Veenhuis, M, Mateblowski, M, Kunau, WH and Harder, W (1987) Yeast 3, 77–84
- [6] Veenhuis, M and Haider, W (1987) in Peroxisomes in Biology and Medicine (Fahimi, H D and Sies, H eds) pp 436–458, Springer, Berlin-Heidelberg
- [7] Veenhuis, M. van Dijken, J.P., Pilon S.A.F. and Harder, W (1985) Arch Microbiol 143, 153-163
- [8] Goodman J M, Scott, C W, Donahue, P N and Atherton, J P (1984) J Biol Chem 259, 8485–8493
- [9] Veenhuis, M, Harder, W, van Dijken, P. Mayer, F (1981) Mol Cell Biol 1, 949–957
- [10] Douma, A.C., Veenhuis, M., de Koning, W., Evers, M. and Harder, W. (1985) Arch. Microbiol. 143, 237–243
- [11] Gould, S J, Keller, G A, Hosken N, Wilkinson, J and Subramani, S (1989) J Cell Biol 108, 1657–1664
- [12] Ledeboer, A M, Edens, L, Maat, J, Visser, C, Bos, J W, Verrips, C T, Janowicz, Z, Eckart, M, Roggenkamp, R and Hollenberg, C P (1985) Nucleic Acids Res 13, 3063-3082
- [13] Digan, M E, Tschopp, J, Grinna, L, Lair, S V Craig, W S. Velicelebi, G, Siegel, R, Davis, G R and Thill, G (1988) Dev Ind Microbiol 29, 59-65
- [14] Cregg, J M, Barringer, K J, Hessler, A Y and Madden, K R (1985) Mol Cell Biol 5, 3376-3385
- [15] Feinberg, A.P. and Vogelstein, B. (1983) Anal Biochem 132, 6-13
- [16] Roggenkamp, R, Didion, T and Kowallik, KV (1989) Mol Cell Biol 9, 988–994
- [17] Burnette, W N (1981) Anal Biochem 112, 195-203
- [18] Cregg, J M and Madden, K R (1987) in Biological Research on Industrial Yeasts, volume II (Stewart, G G, Russell, I, Klein, R D and Hiebsch, R R, eds) pp 1-18, CRC Press, Boca Raton
- [19] van der Klei, I (1991) Thesis, University of Groningen, The Netherlands
- [20] Veenhuis, M., van Dijken, J.P. and Harder, W. (1983) Adv. Microbiol Physiol. 24, 1-82.
- [21] Distel, B., van der Leij, I., Veenhuis, M. and Tabak, H.F. (1988) J. Cell Biol. 107, 1669–1675
- [22] Deshaies, R J , Koch, B D , Werner-Washburne, M , Craig, E A and Schekman, R (1989) Nature 332, 800-805
- [23] Alvares, K., Carrillo, A., Yuan, P.M., Kawano, H., Morimoto, R.I., Reddy, J.K. (1990) Proc. Natl. Acad. Sci. USA 87, 5293-5297