

SPECIFICATION

METHOD FOR PRODUCING OSMOTOLERANT PLANTS

FIELD OF THE INVENTION

5 This invention relates to a method for producing plants with novel properties, more specifically, a method for producing salt-tolerant and/or osmotolerant plants which are highly resistant to environmental stresses.

10 DESCRIPTION OF THE PRIOR ART

 One effective way to prevent global warming is greening of uncultivated soils such as desert or salt-accumulated soil. As a means therefor, the development of plants which are resistant to environmental stresses in
15 combination with an engineering solution such as irrigation plays an important role in controlling encroachment of desert, promoting greening and preventing global warming.

 Salt accumulation causes the following damages: (1)
20 accumulated salt lowers the water potential in soil to prevent plants from absorbing water; (2) the salt absorbed (penetrated) into plants disturbs their metabolism; (3) salt inhibits the absorption of other ions necessary for viability (Sato, F., Plant Cell Engineering, Supplement,
25 "Environmental Problems and Phytobiotechnology", pp. 33-39, 1994). Especially, the inhibition of water absorption causes plants to lose turgor pressure and close stoma. Thus, photosynthesis is deteriorated and growth is

seriously inhibited.

Plants have evolved various mechanisms to adapt themselves to such environments. In a simple adaptation model, plant cells keep an osmotic difference between the inside and outside of the cells in some way, and restore turgor by water absorption. For example, halobacteria, which are not plants however, keep an osmotic balance between the inside and outside by accumulating salt in the cells. In this case, however, it is difficult to adapt them to environmental (osmotic) changes, because intracellular metabolic enzymes per se need to be salt-tolerant.

Therefore, a better adaptation mechanism is the synthesis of a specific compound called "compatible solute" for keeping an intracellular osmosis depending on extrinsic osmotic changes as many salt-tolerant plants do so.

As the compatible solute, bipolar compounds such as glycinebetaine or proline and polyols such as pinitol, sorbitol or mannitol are known. These compounds are characterized by low molecular weight, high water-solubility, low metabolizability, non-influence on metabolism, etc., and are suitable for osmoregulation.

Among others, glycinebetaine (hereinafter referred to as betaine) is noted as a compatible solute found in plants and bacteria which are adaptable to salt-accumulated and/or water-deficient environment. Betaine is thought as a compatible solute found in higher plants

such as Chenopodeaceae, Gramineae, Solanaceae, as well as cyanobacteria, *Escherichia coli*, etc. (for example, see Rhodes, D. and Hanson, A.D., *Annu. Rev. Plant Physiol. Plant Mol. Biol.* 44:357-3584, 1993). Betaine is an
5 osmoprotective substance which keeps an osmotic balance with environments (Robinson, S.P. and Jones, G.P., *Aust. J. Plant Physiol.* 13:659-668, 1986) and prevents the dissociation of soluble enzymes due to high salt concentration (Gabbay-Azaria et al., *Arch. Biochem.*
10 *Biophys.* 264:333-339, 1988). In addition, betaine can protect photosystem II complex against high salt concentration by stabilization of neighboring proteins and manganese cluster within photosynthetic oxygen-evolving complex (for example, see Murata et al., *FEBS Lett.*
15 296:187-189, 1992).

In *Escherichia coli* and spinach (*Spinacia oleracea*), betaine is biosynthesized from choline via two steps of oxidation as shown in Fig. 1. *E. coli* contains two dehydrogenases; one is a membrane-bound oxygen-dependent
20 choline dehydrogenase which oxidizes choline to betainealdehyde (Landfald, B. and Strom, A., *J. Bacteriol.* 165:849-855, 1986), and the other is a soluble NAD-dependent betainealdehyde dehydrogenase which oxidizes betainealdehyde to betaine (Falkenberg, P. and Strom, A.R.,
25 *Biochim. Biophys. Acta.* 1034:253-259, 1990). In higher plants, it has been demonstrated that betaine is synthesized in the chloroplasts via a similar pathway to *E. coli*. In spinach (*Spinacia oleracea*), the first step of

oxidation is catalysed by a ferredoxin-dependent choline monooxygenase (Brouquisse, R. et al., Plant Physiol. 90:322-329, 1989) and an NAD-dependent betainealdehyde dehydrogenase which catalyzes the second step of oxidation (Weretilnyk, E.A. et al., Planta. 178:342-352, 1989) has already been isolated. These plants were found to increase the activities of the both enzymes and thereby the amount of betaine under salt stress (for example, see Hanson, A.D. et al., Proc. Natl. Acad. Sci. U.S.A. 82:3678-3682, 1985).

Alternatively, choline oxidase from the gram-positive soil bacterium *Arthrobacter globiformis* is able to oxidize choline to betaine in one-step oxidation reaction (Ikuta, S. et al., J. Biochem. 82:1741-1749, 1977).

Attempts have been made to confer salt tolerance by integrating such two genes as found in *E. coli* and higher plants or choline oxidase gene into a plant to allow it constantly express these genes (for example, see Nomura M. et al., Plant Physiol. 107:703-708, 1995). However, no success has been achieved so far in obtaining a salt-tolerant and/or osmotolerant plant by integrating such genes into a plant, especially a higher plant expressing them stably.

Choline oxidase is commercially available, but its amino acid sequence has not been determined. Therefore, it would be highly desirable to determine a genetic sequence encoding choline oxidase which can efficiently

convert choline into betaine and to integrate it into a plant to allow it to stably express said sequence, whereby producing a plant which is tolerant to environmental (osmotic) changes such as high salt concentration.

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DISCLOSURE OF THE INVENTION

After profound study to solve the above problems, the present inventors succeeded in isolating a novel gene encoding choline oxidase (The Japanese Society of Plant
10 Physiologist, Annual meeting of 1994, the 34th Symposium held March 28-30, 1994) and integrating it into cyanobacteria, brassicaceous and gramineous plants to obtain salt-tolerant and/or osmotolerant plants.

Accordingly, this invention provides a method for
15 producing salt-tolerant and/or osmotolerant plants, which comprises the step of transforming a plant with a recombinant vector carrying a gene encoding choline oxidase, as well as the salt-tolerant and/or osmotolerant plants obtained by said method.

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BRIEF DESCRIPTION OF THE DRAWINGS

FIG. 1 is a schematic representation showing the oxidation process from choline to betaine.

FIG. 2A is a schematic representation showing the
25 constructs used for transformation of *Synechococcus* PCC7942. PAM refers to *Synechococcus* PCC7942 transformed with pAM1044, and PAMCOD refers to *Synechococcus* PCC7942 transformed with pAM1044 carrying the *codA* gene. Dashed

arrows indicate the primers used for PCR. Triangles represent the *conII* promoter. Arrows indicate the orientation of the genes.

FIG. 2B is an SDS-PAGE representation (photograph of electrophoresis) showing the complete replacement of chromosomes by the spectinomycin-resistant gene and *codA* gene in DNA of *Synechococcus* PCC7942. Lane a: λ -HindIII/ ϕ x174-HaeIII fragment; lane b: the wild-type strain of *Synechococcus* PCC7942; lane c: the strain PAM; lanes d and e: the strain PAMCOD (lanes b, c and d show the results of PCR with primers 1 and 2, and lane e shows the results of PCR with primers 1 and 3).

FIG. 3 is a Western blot analysis representation (photograph of electrophoresis) showing the expression of choline oxidase in the *Synechococcus* strains PAM and PAMCOD. Lane a: protein extracts from the strain PAMCOD; lane b: protein extracts from the strain PAM; lane c: purified choline oxidase.

FIG. 4 shows the effect of NaCl on growth. Growth of the *Synechococcus* strains PAM (○) and PAMCOD (●) in the presence of 0.4M NaCl is shown. For comparison, growth of the *Synechococcus* strains PAM (△) and PAMCOD (▲) cultivated on the medium free from NaCl is also shown.

FIG. 5 shows the effect of sorbitol on growth. Growth of the *Synechococcus* strains PAM (○) and PAMCOD (●) in the presence of 0.8M sorbitol is shown. For comparison, growth of the *Synechococcus* strains PAM (△) and PAMCOD (▲) cultivated on the medium free from

sorbitol is also shown.

FIG. 6 shows the effect of NaCl on chlorophyll content. Chlorophyll contents of the *Synechococcus* strains PAM (○) and PAMCOD (●) in the presence of 0.4M NaCl are shown.

FIG. 7 shows the effect of sorbitol on chlorophyll content. Chlorophyll contents of the *Synechococcus* strains PAM (○) and PAMCOD (●) in the presence of 0.8M sorbitol are shown.

FIG. 8 shows the effect of NaCl on photosynthetic activity. Oxygen evolution levels from the *Synechococcus* strains PAM (○) and PAMCOD (●) in the presence of 0.4M NaCl are shown.

FIG. 9 is a schematic representation showing the restriction enzyme map of the *codA* gene.

FIG. 10 is a schematic representation showing the structure of the binary vector plasmid pGAH/*codA* used for transformation of *Arabidopsis*.

FIG. 11 shows Western blot analysis (photograph of electrophoresis) of choline oxidase in soluble fractions of the wild-type and transformant plants of *Arabidopsis*. Lane 1: choline oxidase from a commercial product of *Arthrobacter globiformis* (Sigma Chemical Co., St. Louis, MO, USA); lane 2: soluble fractions of the wild-type plant; lane 3: soluble fractions of a transformant plant.

FIG. 12 shows the effect of NaCl on the growth of *Arabidopsis*. The wild-type (A) and transformant C1-0 (B) plants in the presence of 60mM NaCl are shown (photographs

showing the morphology of the organisms).

FIG. 13 shows the effect of sorbitol on the growth of *Arabidopsis*. The wild-type (W) and transformant (T) plants in the presence of 100, 200 and 400mM sorbitol are shown (photographs showing the morphology of the organisms).

FIG. 14 shows the influence of salt stress on photosynthesis system II in leaves of the wild-type and transformant plants (*Arabidopsis*). ○: the wild-type plant incubated under light conditions; ●: a transformant plant incubated under light conditions; △: the wild-type plant incubated under dark conditions; ▲: a transformant plant incubated under dark conditions. Each data represents the average of triplicate runs with standard deviation of $\pm 5\%$.

FIG. 15 shows the structures of the two chimeric *codA* genes used for transformation of rice, i.e. 35SINTP*codA* and 35SIN*codA*.

FIG. 16 shows NMR charts representing betaine accumulation in rice plants of the wild-type strain, a transformant (A) not expressing the *codA* gene, and a transformant (B) expressing the *codA* gene. In the figure, GB and Ch represent peaks corresponding to betaine and choline, respectively.

FIG. 17 shows the influence of salt stress on photosynthetic system II activity of the wild-type and transformant rice plants.

THE BEST EMBODIMENTS OF THE INVENTION

The gene encoding choline oxidase used in this invention is a gene which encodes a protein capable of converting choline into betaine in a one-step reaction and
5 which may be derived from gram-positive soil bacteria of the genus *Arthrobacter*. For example, it may be preferably derived from *Arthrobacter globiformis* and *Arthrobacter pascens*, especially *Arthrobacter globiformis*.

The present inventors successfully cloned the *codA*
10 gene encoding choline oxidase from *Arthrobacter globiformis* and determined its nucleotide sequence. The *codA* gene contains an open reading frame of 1641bp, which encodes 547 amino acids. The nucleotide sequence and amino acid sequence of the *codA* gene are shown as SEQ ID
15 NO. 1 in SEQUENCE LIST.

Plants can be transformed with such a choline oxidase-coding gene integrated into appropriate vectors. Then, the gene can be expressed in the plants by introducing into these vectors an appropriate promoter or
20 a sequence responsible for the expression of character.

Any gene having a nucleotide sequence resulting from addition, deletion or substitution of the nucleotide sequence encoding the amino acid of SEQ ID NO. 1 in SEQUENCE LIST or a part thereof may be used as the gene
25 according to this invention so far as it encodes a polypeptide showing choline oxidase activity.

According to the method of this invention, salt tolerance and/or osmotolerance can be conferred to a large

variety of plants ranging from cyanobacteria to higher plants. Cyanobacteria are widely used as model organisms of higher plants because they have basically the same photosynthetic mechanism as higher plants and they are easily transformed to give the results in a short time. Some transformant-type cyanobacteria readily incorporate foreign DNA into their cells to cause efficient recombination. Such cyanobacteria include *Synechococcus* PCC7942, *Synechococcus* PCC6301 (ATCC 27144) and *Synechocystis* PCC6803 (ATCC 27184) (Protein, Nucleic Acid, Enzyme, Vol.35, No.14, pp.2542-2551, 1991; Crit. Rev. Microbiol. Vol.13, No.1, pp.111-132, 1985).

Higher plants include dicotyledons and monocotyledons. In Examples described below, highly salt-tolerant and/or osmotolerant plants could be obtained from a brassicaceous plant as a dicotyledon, but it is not limitative and other families and genera of dicotyledons may be used. The method of this invention may also be applicable to monocotyledons. It was found that a monocotyledonous plant rice, which originally lacks betaine-synthesizing ability, gained this ability, and therefore salt tolerance after transformation according to the method of this invention.

The vectors into which the choline oxidase-coding gene is integrated and the procedures for transformation and selection of the transformant plant materials can be appropriately chosen dependent on the nature of the plant to be transformed.

For example, plasmids such as pUC303 can be used for cyanobacteria. Then, the transformants having desired properties can be selected by the antibiotics resistant genes inserted into these plasmids. This invention
5 succeeded in obtaining the plants which stably show salt tolerance and/or osmotolerance by transforming the cyanobacterium *Synechococcus* PCC7942 with the *codA* gene encoding choline oxidase from *Arthrobacter globiformis*.

When the *Synechococcus* PCC7942 transformed with the
10 *codA* gene was cultivated on a medium supplemented with choline chloride, the *Synechococcus* was found to take up exogeneously supplied choline and convert it into betaine. In view of the report that choline transport is induced by salt stress in several salt-tolerant bacteria, resulting
15 in accumulation of higher level of betaine (Kaenjak, A. et al., J. Bacteriol. 175:2400-2406, 1993), the influence of salt stress on betaine accumulation was examined by treating the transformant *Synechococcus* produced by this invention with NaCl at various concentrations. However,
20 the influence of NaCl concentration on betaine accumulation was not significantly observed, suggesting that the transporter for choline uptake in *Synechococcus* is not specifically induced by salt stress.

It has also been reported that betaine not only
25 acts as an osmoprotectant but also plays an essential role in protection of photosynthetic mechanism in photoautotrophic organisms (Murata, N. et al., FEBS Lett. 296:187-189, 1992). The *Synechococcus* transformed

according to this invention was cultivated in the presence of high concentration NaCl or sorbitol to examine growth, chlorophyll content and photosynthetic activity. As a result, the transformant *Synechococcus* grew well in the presence of either high concentration salt or sorbitol and also showed the similar results for chlorophyll content and photosynthetic activity, as compared with the control non-transformant in which growth, chlorophyll content and photosynthetic activity were all inhibited. These results mean that excellent salt tolerance and osmotolerance were conferred to the *Synechococcus* transformed with the gene encoding choline oxidase according to the method of this invention.

Dicotyledons may be transformed by gene introduction technique using protoplasts or a part of tissue. In case of the gene introduction using tissue pieces, the Ti plasmid from *Agrobacterium* may be used. Tissue pieces of a callused plant may be infected with *Agrobacterium* bearing protoplasts into which the choline oxidase-coding gene has been integrated, selected by resistance to an antibiotic such as kanamycin, and then differentiated in shoots to give a transformant plant.

In this invention, a salt-tolerant and/or osmotolerant plant was obtained by transforming the brassicaceous plant *Arabidopsis thaliana* with the choline oxidase-coding gene as follows.

A binary vector plasmid pGAH-codA carrying the codA gene was prepared and integrated into *Agrobacterium*

tumefaciens EHA101 bearing the Ti plasmid. Hypocotyl
calli of Arabidopsis were infected with the resultant
Agrobacterium EHA101 (pGAH/codA) incorporating the codA
gene, then shoots were formed and selected by kanamycin
5 and hygromycin resistance to induce roots and to form
seeds. The plants obtained from said heterozygous T2
seeds were self-fertilized to give homozygous T3
individuals, which were sown to form transformant plants.
These transformant plants showed that choline oxidase had
10 been transported to the chloroplasts. The transformant
plants grew well even in the presence of high
concentration sodium chloride or sorbitol.

The monocotyledonous plant rice (*Oryza sativa* L. cv.
Nippon bare) can be transformed with two chimeric codA
15 genes prepared on the plasmid pUC119, which are localised
on cytosol or plastide after translation under
transcriptional control of the cauliflower mosaic virus 35S
promoter. Both of the chimeric genes include a rice-
derived intron in the 5' non-translated sequence in order
20 to enhance the expression.

The transformant rice can be produced by the
following procedure. Namely, the transformant plant can
be obtained by introducing said chimeric codA genes into
suspension culture cells from scutellum calli of rice
25 seeds together with the selection marker hygromycin-
resistant gene by a particle gun device, then selecting
the transformed calluses based on the antibiotics
resistance, and redifferentiating them into a plant.

Although the wild-type rice lacks betaine-synthesizing ability, the rice transformed by the method of this invention gained betaine-synthesizing ability. The transformant rice expressing the *codA* gene grew
5 equally to the non-transformed plant without showing any apparent abnormality under the both of geponic and hydroponic conditions. This may conclude that hydrogen peroxide formed as a by-product of betaine synthesis was efficiently detoxified in the cells.

10 Moreover, salt tolerance tests of the transformants cultivated on water with NaCl showed that the inhibition of photosynthetic activity in the transformants was slower than observed in the wild type. This is the first case in which rice has gained betaine-synthesizing ability through
15 a genetic engineering procedure.

These results show that various plants transformed with recombinant vectors carrying the gene encoding choline oxidase have an excellent salt tolerance.

According to this invention, salt-tolerant and/or
20 osmotolerant transformant plants which are highly resistant to environmental stresses can be obtained. The range of plants to which can be conferred salt tolerance and/or osmotolerane by the method of this invention is very wide, from cyanobacteria to higher plants.

25 Especially, this invention is the first case in which salt-tolerant and/or osmotolerant transformant plants were obtained from monocotyledons including most of main crops and expected to be applied in a very wide range.

The following examples further explain this invention in detail, but are not intended to limit the scope of this invention.

EXAMPLES

5 Example 1: Transformation of the cyanobacterium

Synechococcus PCC7942 with the *codA* gene

(1) Cloning of the *codA* gene

The choline oxidase gene was isolated from *Arthrobacter globiformis* by the method described in the
10 Abstracts of Oral Reports in the abstracts, the 34th annual meeting of the Japanese Society of Plant Physiologists, 1994. In brief, 1) choline oxidase is fragmented with cyanogen bromide, 2) the N-terminal amino acid sequences of appropriate fragments are determined, 3)
15 an appropriate part is selected from said amino acid partial sequences to synthesize oligonucleotides corresponding thereto, 4) a partial sequences of the choline oxidase gene is amplified by PCR (Polymerase Chain Reaction) using these oligonucleotides as primers, 5) the
20 amplified partial sequence of the choline oxidase gene is used as a probe to screen the genomic DNA library of *Arthrobacter globiformis*.

Thus obtained positive clones were subcloned into the plasmid pBluescript (SK⁺) (Stratagene) to isolate
25 positive clones, which were subjected to Southern blot analysis. A 3.6kbp *Xba*I-*Xho*I fragment which hybridized to said probe was subcloned into pBluescript and mapped with restriction enzymes. The nucleotide sequence of the

region spanning from the first SalI-site to XhoI-site (about 2.5kbp) was determined.

The results showed that the choline oxidase gene contains an open reading frame of 1641bp which encodes a polypeptide of 547 amino acid residues. The amino acid sequence and the nucleotide sequence of the choline oxidase-coding gene are shown as SEQ ID NO. 1 in SEQUENCE LIST.

10 (2) Transformation of Synechococcus PCC7942 with the *codA* gene

The plasmid pBluescript carrying the *codA* gene was digested with BstEII (position -40 from the initiation of translation) and SmaI (downstream of the stop codon) restriction enzymes. The BstE II-cohesive end was filled in by Klenow fragment (Takara, Tokyo, Japan). The blunt-ended fragment containing the coding region of the *codA* gene and a putative ribosome binding site was inserted into the SmaI site of the plasmid pAM1044. The correct orientation of the gene, which is supposed to be expressed under control of the *conII* promoter of pAM1044, was confirmed by restriction analysis. The *conII* promoter is the consensus sequence of promoters of *E. coli*, which contains the base sequences TTGGACA (-35) and TATAAT (-10).

25 The plasmid pAM1044 and the plasmid containing the *codA* gene were used for transformation of Synechococcus PCC7942 by the method of Elhai et al. The resultant transformant was designated as the strain PAMCOD.

Synechococcus PCC7942 transformed with pAM1044 alone was used as a control and designated as the strain PAM.

Selection of the transformants was done on the BG11 agar plate containing spectinomycin at 30µg/ml. After
5 several inoculations of a single colony to fresh BG11 plates containing spectinomycin, the complete insertion of the spectinomycin-resistant gene and the *codA* gene into all the copies of the chromosomes was confirmed by PCR (Polymerase Chain Reaction) using the primers indicated in
10 Fig. 2A. The complete insertion of the spectinomycin-resistant gene and the *codA* gene into the chromosomes of *Synechococcus* was confirmed by PCR using a combination of primers 1 and 2.

15 Example 2: Confirmation of the gene inserted into transformants

The DNAs from the wild-type strain, the strain PAM and the strain PAMCOD of *Synechococcus* PCC7942 were used as templates for PCR, and the amplified products were
20 analyzed by SDS-PAGE. The results are shown in Fig. 2B.

A PCR performed on the DNA from the wild-type strain revealed an amplified product of about 400bp (Fig. 2B, lane b). When the DNA from the strain PAM was used as a template, a band of about 2.4kb appeared, representing
25 the insertion of pAM1044 into the chromosomes. The absence of the band of about 400bp, which was observed in the wild-type strain, confirms the complete replacement of the native chromosomes by the mutated chromosomes in the

strain PAM.

When the DNA from PAMCOD was used as a template, the band corresponding to the wild-type chromosomes was not observed (Fig. 2B, lane c). However, the expected
5 band of about 4.1kb was not amplified, either, likely due to a large size of the insert and a high GC content in the codA sequence. Therefore, primer 3 corresponding to the coding region of the codA gene (Fig. 2A) was used in combination with primer 1. The expected band of about
10 2.6kb was amplified (Fig. 2B, lane d), indicating the presence of the codA gene in the chromosomes of the strain PAMCOD.

Example 3: Expression of the codA gene in the 15 Synechococcus strain PAMCOD

The expression of the codA gene in the strain PAMCOD obtained in Example 1 was examined by Western blot analysis using polyclonal antiserum raised against the purified choline oxidase. The results are shown in Fig. 3.
20 Signals were detected in the protein extracts obtained from the strain PAMCOD (lane a) and purified choline oxidase (lane c) at the position of 60kDa. Such a signal was not detected in the protein extracts obtained from the strain PAM (lane b). This result confirms that the codA
25 gene was expressed under control of the conII promoter in Synechococcus PCC7942.

Example 4: Analysis of betaine concentration in cells

The transformed cells were grown in one liter of BG11 medium supplemented with 5mM choline chloride. Salt stress was given by adding various concentrations of NaCl. The harvested cells were treated with 1M H₂SO₄ at 25°C for 20 hours and betaine was recovered from the mixture by means of the periodate precipitation technique (Wall, J.S. et al., *Analyt. Chem.* 32:870-874, 1960). The betaine periodate was dissolved in 1ml of methanol-d₄ (Wako Pure Chemical Industries, Osaka, Japan) containing 2mM 2-methyl-2-propanol (Wako Pure Chemical Industries) as an internal standard. This solution was transferred to an NMR tube and ¹H NMR spectra were measured with a Bruker AMX 360 Wb. Betaine was quantified by comparing the integrated peaks with a standard curve.

The concentration of betaine in the cells of the strain PAMCOD was determined on the basis of the cell volume estimated from the electron micrograph of negatively stained cells. The cytoplasm of a single cell had a cylindrical shape of 2.14 μm in length and 0.82 μm in diameter and the cell volume was estimated to be approximately 1.13 μm³.

The following Table 1 shows the changes in betaine concentration in the cells with the increase of NaCl concentration in the medium. Any trace of betaine could not be detected in the strain PAM lacking the *codA* gene. The betaine concentration in the cells of the strain PAMCOD ranged from 60 to 90mM. Betaine accumulation in the cells of the strain PAMCOD was not significantly affected

growth test in the presence of sorbitol are shown in Fig.

5. The growth of the strain PAM was inhibited by high salt and osmotic stresses, while the strain PAMCOD could grow under these conditions.

5 The results of the chlorophyll content test in the presence of NaCl are shown in Fig. 6 and the results of the chlorophyll content test in the presence of sorbitol are shown in Fig. 7. The results of this test were similar to those of the cell growth test. The chlorophyll
10 content in the strain PAM treated with a high concentration salt gradually decreased and the chlorophyll content in the strain PAM treated with 0.8M sorbitol rapidly decreased, but on the contrary, the strain PAMCOD continued to grow even under salt or sorbitol treatment.

15 The results of the photosynthetic activity test in the presence of NaCl are shown in Fig. 8. Photosynthetic activity of the strain PAM was strongly inhibited by salt stress. In contrast, photosynthetic activity of the cells
20 of the strain PAMCOD was temporarily inhibited at the early stage of salt treatment, but afterward recovered and continuously increased. The similar results were obtained in the photosynthetic activity test in the presence of sorbitol. Interestingly, the temporary decrease of
25 photosynthetic activity was not observed when the cells of the strain PAMCOD were treated with 0.8M sorbitol.

Example 6: Preparation of a binary vector plasmid carrying the *codA* gene

The *rbcS* (ribulose 1,5-bisphosphate carboxylase small subunit) transit signal *XbaI*-*NdeI* fragment (about 200bp) from tobacco (*Nicotiana glauca*) was amplified by PCR using 5'CTGTCTAGATGTAATTAACAATGGCT3' and
5 5'CCACATATGCATGCATTGCACTCT3' as primers to introduce the *XbaI* and *NdeI* sites.

Then, the N-terminal-*BamHI* fragment (about 100bp) of the *codA* gene was amplified by PCR using 5'AACCATATGCACATCGACAACATC3' and 5'GCTCCATCCAGCGGTCCAGC3'
10 as primers to introduce the *NdeI* site. The *BamHI*-*SmaI* fragment (about 1.6kbp) of the *codA* gene was prepared by restriction enzymes. Further, the *SmaI*-C-terminal fragment (about 80bp) of the *codA* gene was amplified by PCR using 5'GAAACAGTCCTGCTTCCACAC3' and
15 5'GCGAGCTCTGCCTACACCGCCAT3' as primers to introduce the *SacI* site.

The *GUS* (β -glucuronidase) gene in the binary vector plasmid pBI221 was replaced by these fragments.

The restriction enzyme map of the *codA* gene is
20 shown in Fig. 9.

The *HindIII*-*EcoRI* fragment containing the 35S promoter and the NOS (nopal synthase) terminator of cauliflower mosaic virus was introduced into the binary vector plasmid pGAH to prepare the plasmid pGAH/*codA* (Fig.
25 10). This plasmid contains kanamycin- and hygromycin-resistant genes.

Example 7: Introduction of the binary vector plasmid into

Agrobacterium

The *Agrobacterium tumefaciens* EHA 101 bearing the Ti plasmid was mixed with the binary vector plasmid pGAH/codA obtained in Example 6, then freeze-d and thawed,
5 and screened on the LB plate containing tetracycline and kanamycin. The resultant *Agrobacterium* in which the codA gene has been integrated was designated as EHA101 (pGAH/codA).

10 Example 8: Transformation of Arabidopsis

The *Arabidopsis thaliana* strain WS was germinated to prepare a hypocotyl segment. This hypocotyl was callused on B5 medium (ICN Biochemicals) (pH 5.7) containing 0.05mg/l of kinetin (Wako Pure Chemical
15 Industries) and 0.5mg/l of 2,4-D (Wako Pure Chemical Industries) to form hypocotyl calli.

Then, the calli were infected with the codA-containing *Agrobacterium* EHA101(pGAH/codA) prepared in Example 7 and cocultivated. After detoxification of
20 *Agrobacterium* by B5 medium containing 250mg/l of vancomycin, 500mg/l of carbenicillin and 200mg/l of Claforan, the cultures were transferred to a differentiation medium (B5 medium containing 25mg/l of kanamycin and 15mg/l of hygromycin) to form shoots. Then,
25 kanamycin- and hygromycin-resistant shoots were selected to induce roots and to form seeds. The resultant T2 seeds are heterozygous transformed in only one of the chromosomes.

Then, the plants obtained from the T2 seeds were self-fertilized and selected by kanamycin and hygromycin to give homozygous T3 seeds.

The plants of the wild-type and transformant strains were grown in a medium (pH 5.2) containing 0.1% HYPONEX (Hyponex Corporation, Marysville, OH, USA) at 22°C for 30 days on water or soil consisting of vermiculite and perlite with illumination of $75\mu\text{mol.m}^{-2}.\text{s}^{-1}$ for 16 hours in a day and in a dark room for the remaining 8 hours unless otherwise indicated, and then used for experiments.

Example 9: Immunological study of the expressed choline oxidase

An antibody was raised against choline oxidase according to the method described by the present inventors in literature (Deshniumn, P. et al., Plant Mol. Biol. 29:897-907, 1995).

Leaf from each 20-day old plant of the wild-type and transformant strains of *Arabidopsis thaliana* was ground in a microcentrifuge at 0°C and the homogenates were centrifuged at 10,000 x g for 10 minutes to prepare soluble fractions. The soluble protein of the supernatant was separated by SDS-PAGE and transferred to a nylon membrane (Immobilon PVDF; Millipore, Bedford, MA, USA). The membrane was incubated with the antibody against choline oxidase and detected with a system consisting of biotinylated secondary antibody, avidin and biotinylated horse radish peroxidase (ABC Kit; Vectastain, Burlingane,

CA, USA).

The results of Western blot analysis are shown in Fig. 11. The presence of an immune responsive protein of 64kDa corresponding to choline oxidase was identified. A
5 small amount of protein of 70kDa corresponding to the precursor of choline oxidase and the rbcS transit peptide were also observed. These results show that the codA gene was correctly integrated and expressed in the chromosomes and that the expressed precursor was processed into a
10 mature protein.

Then, localization of the expressed choline oxidase in the plant was detected with the antibody against choline oxidase by a method described in literature (Mustardy, L. et al., Plant Physiol. 94:334-340, 1990). A
15 small piece of young leaf from the plant was fixed with 1% glutaraldehyde in 0.1M sodium phosphate buffer (pH 7.2) for one hour. After rinsed with the same buffer, the sample was dehydrated with ethanol and placed in Lowicryl K4M resin (TAAB Laboratories Equipment Ltd., Berkshire,
20 U.K.). Immuno-gold labeling was conducted by a method described in literature (Mustardy et al., supra).

As a result, the expressed choline oxidase was found to be localized in stroma of the chloroplasts, indicating that choline oxidase had been transported to
25 the chloroplasts.

Example 10: Determination of betaine and chlorophyll contents in transformant plants

Betaine content in leaf of the plants was calculated by measuring NMR spectra of the quaternary ammonium compound (Wall, J. et al., *Analyt. Chem.* 32:870-874, 1960). 5g of leaf of the wild-type strain and
5 transformant plants were powdered in liquid nitrogen by a ceramic motor. This powder was suspended in 25 ml of 1.0M H_2SO_4 and incubated at 25°C for 2 hours. After insoluble matters were removed, the supernatant was recovered by centrifugation at 1000 x g for 10 minutes. The
10 supernatant was combined with 10ml of KI-I₂ solution and incubated at 0°C for 2 hours. Peridodide-adducts of betaine and choline were recovered by centrifugation at 1000xg for 30 minutes and dissolved into 0.5ml of CD₃OH (Wako Pure Chemical Industries) containing 0.5mM 2-methyl-
15 2-propanol (Wako Pure Chemical Industries) as an internal standard to measure ¹H NMR spectra. Two main peaks corresponding to betaine and choline were observed, and the integrated betaine peaks were used for determination of the concentration.

20 Chlorophyll content in leaf was measured by the following procedure. Leaf (1g) was powdered in liquid nitrogen by a ceramic motor. The powder was suspended in 10ml of acetone:water (4:1, v/v). After incubation for 30 minutes, insoluble matters were removed and the
25 supernatant was subjected to spectrophotometry (Arnon, D.I. *Plant Physiol.* 24:1-15, 1949).

As a result, the both of betaine and choline were observed in the transformant plant, while only choline was

The seeds of the wild-type strain and the transformant plants were sterilized and inoculated on semi-solid media containing 100, 200 and 400mM sorbitol. The cultures were placed in an incubator maintained at 22°C with illumination of 75 $\mu\text{mol}/\text{m}^2/\text{second}$ for 16 hours in a day and in a dark room for the remaining 8 hours, and regularly observed. The state on the day 15 is shown in Fig. 13 (W on the left represents the wild-type strain and T on the right represents the transformant plants). At a sorbitol concentration of 100mM, some wild-type strains showed no germination or its leaves whitened even if they germinated. However, the transformant plants continued to grow while keeping green. At a concentration of 200mM, growth of the both strains was inhibited, but the inhibition degree of the transformant plants was lower than that of the wild-type strain, and especially, the growth of roots of the transformant plants was remarkably better than that of the wild-type strain. At a concentration of 400mM, the wild-type strain showed no germination and the transformant plants showed no germination, either, or scarcely grew even if they germinated.

Example 12: Photosynthetic activity of transformant plants under salt stress

The influence of salt stress on photosynthetic system II activity of mature leaves was measured by monitoring fluorescence of chlorophyll.

The wild-type strain and the transformant strain

plants grown on a control medium were transferred to HYPONEX medium containing 400mM sodium chloride and incubated under the light or dark conditions described above. After a determined time, leaves were taken from
5 the plants and the efficiency of photosynthetic system II was measured as a ratio of variable chlorophyll fluorescence to maximum chlorophyll fluorescence (Fv/Fm) by using a pulse intensity-modulated fluorometer (PAM-2000; Walts, Effeltrich, Germany) (Annu. Rev. Plant
10 Physiol. Plant Mol. Biol. 42:313-349, 1991).

The results are shown in Fig. 14. After incubation for 2 days under light conditions, the wild-type strain almost lost photosynthetic system II activity, while the transformant plants maintained 50% of the original level
15 of photosynthetic system II activity. Under dark conditions, inactivation due to salt stress was much more moderate, but photosynthetic system II activity of the transformant plants was more tolerant than that of the wild-type strain. When the plants were transferred to the
20 medium containing 200mM sodium chloride, the decrease of photosynthetic system II activity was much more moderate than observed on the medium containing 400mM sodium chloride, but the transformant plants were again more tolerant to salt stress than the wild-type strain.

25

Example 13: Preparation of the chimeric *codA* gene used for transformation of rice

Two chimeric *codA* genes (designated as 35SIN*codA*

and 35SINTPcodA, respectively) which are localized on cytosol and plastid, respectively after translation of the choline oxidase gene (codA) from *Arthrobacter globiformis* under transcriptional control of the cauliflower mosaic virus 35S promoter were prepared on the plasmid pUC119 by the procedure described in Example 6 (see Fig. 15). Considering that the presence of an intron is required for high expression of a gene in rice (for example, see Tanaka, A. et al., *Nucleic Acids Res.* 18:6767-6770, 1990), an intron in the 5' non-translated sequence of the superoxide dismutase gene of rice (SodCc2: Sakamoto, A. et al., *FEBS Lett.* 358:62-66, 1995) was introduced into the both chimeric genes. Further, a DNA sequence derived from the rbcS transit peptide (Coruzzi, G. et al., *EMBO J* 3:1671-1679, 1984) from pea was added to 35SINTPcodA, in order to transfer the codA protein to the chloroplasts.

Example 14: Transformation of rice

Each of the two chimeric codA genes prepared in Example 13 was introduced into suspension culture cells from scutellum calli of rice seeds together with the selection marker hygromycin-resistant gene by a particle gun device. The transformed calli were selected based on the antibiotic resistance and redifferentiated into a plant. Polymerase Chain Reaction (PCR) was conducted on the transformed calli or transformed/redifferentiated individuals showing hygromycine resistance, to assess integration and transcription of the codA gene into the

nuclear genome by Northern blot technique and select 80 to 100 or more transformants for each *codA* gene.

Example 15: Analysis of expression of the *codA* gene in
5 transformant rice

The transformants obtained in Example 14 were screened by Western blot technique to obtain the transformant rice (the present generation) expressing the *codA* gene at the protein level, ultimately including 6
10 individuals carrying the plastid-localized gene and 10 individuals carrying the cytosol-localized gene.

Rice lacks intrinsic choline oxidase activity, but the soluble fractions prepared from leaves or roots of the transformants showed choline oxidase activity. Contrary
15 to expectation, all the individuals of the plastid-type transformants were found to express a lower amount of choline oxidase protein than the cytosol-type, despite the same expression promoter used.

When the expression of the *codA* gene was further
20 examined by Northern blot technique, any significant difference was not found in the amount of the both genes expressed at the transcription level. When processing of the intron was examined by reverse transcriptive PCR, a plurality of splicing variants containing different 3'-
25 acceptor sites which may not bring about normal translation into protein were detected from the mRNA transcribed from the plastid-type gene. This suggested that the low level protein expression by the plants

transformed with the plastid-type gene might be due to abnormal processing of the mRNA precursor. This phenomenon seems to be related to the fact that the sequence encoding the transit peptide used for plastid-targeting of choline oxidase was derived from a dicotyledon (pea rbcS gene). Therefore, it may be readily expected that the expression of the codA in rice chloroplasts would be more efficient and the resultant transformant rice would be more tolerant to salt stress if the sequence encoding the transit peptide was derived from a monocotyledon such as rice rbcS.

Example 16: Betaine biosynthesis in transformant rice

The betaine accumulating in the transformant tissue expressing choline oxidase was detected by proton NMR. Fig. 16 shows the results of the NMR of the wild-type strain, a transformant which does not express the codA gene (Fig. 16A) and a transformant which expresses the codA gene (Fig. 16B).

The transformant which expresses choline oxidase biosynthesized betaine and the amount of betaine accumulation showed a positive correlation with the amount of choline oxidase detected by Western blot technique. Betaine accumulation in leaves was higher than detected in roots and amounted to 4 μ mol/g fresh leaf in individuals highly expressing the codA gene. This is the first case in which rice gained the ability to synthesize betaine through a genetic engineering procedure.

Example 17: Evaluation of salt tolerance of transformant
rice

The transformant rice expressing the codA gene grew
5 well equally to the non-transformant (wild-type) under the
both of geponic and hydroponic conditions without showing
any apparent abnormality. This indicates that hydrogen
peroxide formed as a by-product of betaine biosynthesis
was efficiently detoxified in the cells.

10 Then, the transformants which could be found to
express the codA gene at the level of protein, enzyme
activity and betaine production were grown under the
hydroponic conditions containing sodium chloride. The
influence of Na salt on photosynthetic activity was
15 assessed by chlorophyll fluorescence analysis and compared
with the results of the non-transformant (wild-type).
When the transformant rice and non-transformant rice were
placed in aqueous HYPONEX solutions containing 100mM and
10mM sodium chloride and the fluorescence of chlorophyll
20 was determined with time, the transformant was found to
retard the inhibition of photosynthetic activity (Fig. 17).
Thus, the transformant was found to be more tolerant in
salt environment.

[SEQUENCE LISTING]

SEQ ID NO: 1
Sequence Length: 2400
Sequence Type: nucleic acid
5 Strandedness: double
Topology: linear
Molecule Type: DNA
Sequence Feature
Feature Key: mat peptide
10 Position: 361..2002

Sequence Description:

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TGGACCCAGG ACGCCGGCAA GACGCAGCGC GTGGCCGGCC GGCTGCGACA CGGCACCGTC 180
TGGATCAACG ACTTCCACCC CTACCTCCCA CAGACCGAGT GGGGCGGCCTT CGGCCAGTCC 240
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TACCAGAACA CCAGCCCGCA GGTACCGGC TGGTTCTCTG ACCACGGCAA GGAGAACTAG 360
20
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Met His Ile Asp Asn Ile Glu Asn Leu Ser Asp Arg Glu Phe Asp Tyr
1 5 10 15
ATC GTC GTC GGC GGC GGG TCC GCC GGG GCC GCC GTC GCC GCC CGG CTG 456
25 Ile Val Val Gly Gly Gly Ser Ala Gly Ala Ala Val Ala Ala Arg Leu
20 25 30
AGC GAG GAT CCC GCA GTG AGC GTG GCG CTG GTG GAG GCC GGC CCG GAT 504
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	35	40	45	
	GAC CGC GGC GTG CCC GAG GTG CTG CAG CTG GAC CGC TGG ATG GAG CTG	552		
	Asp Arg Gly Val Pro Glu Val Leu Gln Leu Asp Arg Trp Met Glu Leu			
	50	55	60	
5	CTG GAA TCG GGC TAC GAC TGG GAC TAC CCG ATC GAG CCG CAG GAG AAC	600		
	Leu Glu Ser Gly Tyr Asp Trp Asp Tyr Pro Ile Glu Pro Gln Glu Asn			
	65	70	75	80
	GGC AAC TCC TTC ATG CGC CAT GCC CGT GCC AAG GTC ATG GGC GGC TGC	648		
	Gly Asn Ser Phe Met Arg His Ala Arg Ala Lys Val Met Gly Gly Cys			
10	85	90	95	
	TCC AGC CAC AAC TCC TGC ATC GCC TTC TGG GCC CCG CGC GAG GAC CTG	696		
	Ser Ser His Asn Ser Cys Ile Ala Phe Trp Ala Pro Arg Glu Asp Leu			
	100	105	110	
	GAC GAG TGG GAG GCC AAG TAC GGC GCC ACC GGC TGG AAC GCC GAG GCG	744		
15	Asp Glu Trp Glu Ala Lys Tyr Gly Ala Thr Gly Trp Asn Ala Glu Ala			
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	GCC TGG CCG CTG TAC AAG CGG CTG GAA ACC AAC GAG GAC GCG GGC CCG	792		
	Ala Trp Pro Leu Tyr Lys Arg Leu Glu Thr Asn Glu Asp Ala Gly Pro			
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	CCC CCG AAG GAC CCG ACC GGC GTC GCG CTC CTG GAC GGC TGC GAG CAG	888		
	Pro Pro Lys Asp Pro Thr Gly Val Ala Leu Leu Asp Ala Cys Glu Gln			
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5 Ser Ser Ser Val Ser Tyr Ile His Pro Ile Val Glu Gln Glu Asn Phe
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225 230 235 240
10 AGG CGC TGC ACC GGC GTC GAC ATC GTG GAC TCC GCC TTC GGC CGC ACC 1128
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 465 470 475 480
 ATG TCC CCG CTC GAC CCC GAG CTG CGG GTC AAG GGC GTC ACC GGT CTG 1848
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CLAIMES

1. A method for producing a salt-tolerant and/or osmotolerant plant, which comprises the step of transforming a plant with a recombinant vector carrying a gene encoding choline oxidase.
2. A method according to Claim 1 wherein the gene encoding choline oxidase is derived from the soil bacteria *Arthrobacter*.
3. A method according to Claim 1 or 2 wherein the gene encoding choline oxidase is a base sequence encoding the amino acid sequence of SEQ ID NO. 1 in SEQUENCE LIST or a nucleotide sequence encoding a modified amino acid sequence of SEQ ID NO. 1 resulting from addition and/or deletion of one or more amino acid sequences and/or substitution by other amino acids provided that the nucleotide sequence encodes a protein having choline oxidase activity.
4. A method according to any one of Claims 1 to 3 wherein the salt-tolerant and/or osmotolerant plant is a cyanobacterium.
5. A method according to any one of Claims 1 to 3 wherein the salt-tolerant and/or osmotolerant plant is a higher plant.
6. A method according to Claim 5 wherein the higher plant is a dicotyledon.
7. A method according to Claim 6 wherein the dicotyledon is a brassicaceous plant.
8. A method according to Claim 5 wherein the higher plant is a monocotyledon.



9. A method according to Claim 8 wherein the monocotyledon is a gramineous plant.

10. A salt-tolerant and/or osmotolerant plant produced by the method according to Claim 1, or a progeny thereof having the same properties.

METHOD FOR PRODUCING OSMOTOLERANT PLANTS

ABSTRACT

A method for producing salt-tolerant and/or osmotolerant plants, which comprises the step of
5 transforming a plant with a recombinant vector carrying a gene encoding choline oxidase, as well as the salt-tolerant and/or osmotolerant plants produced by said method or a progeny thereof having the same properties.

Fig. 1

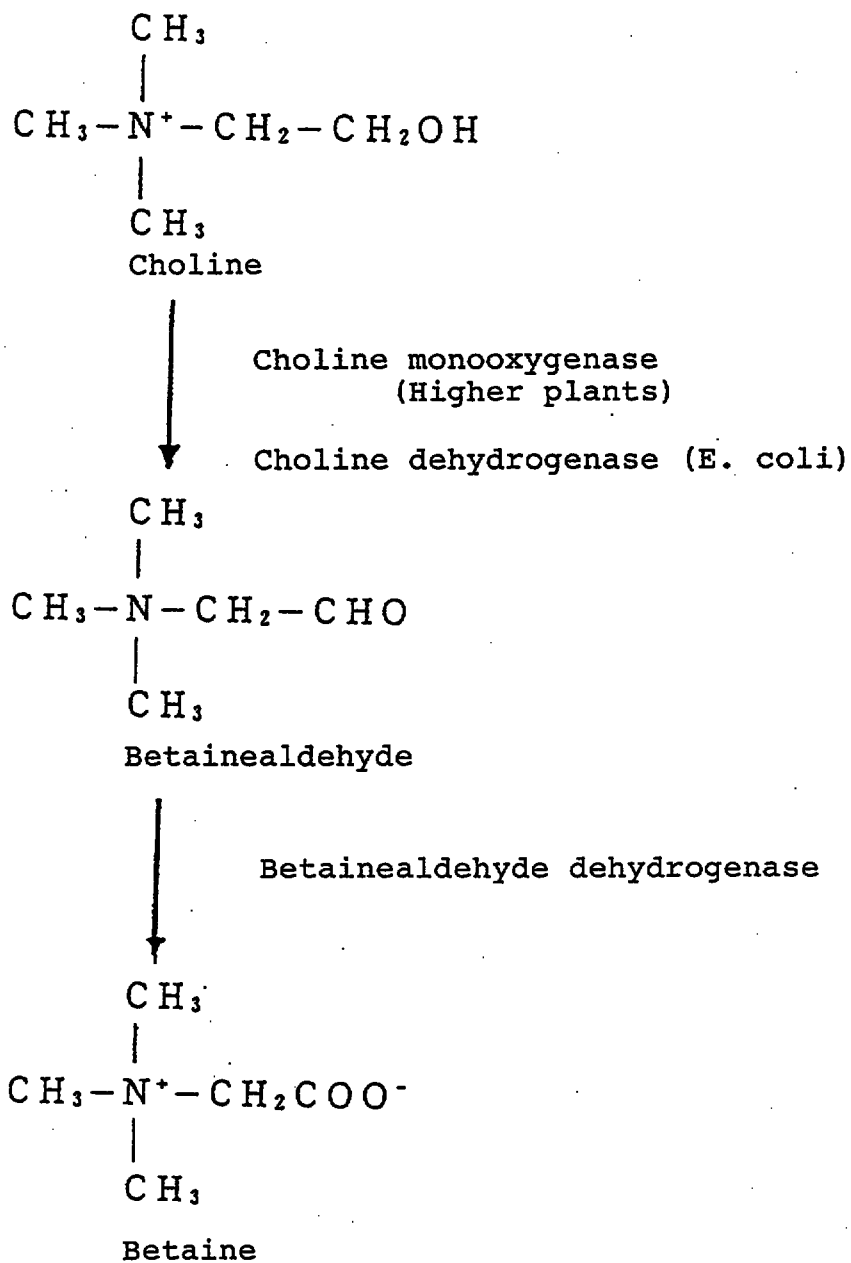
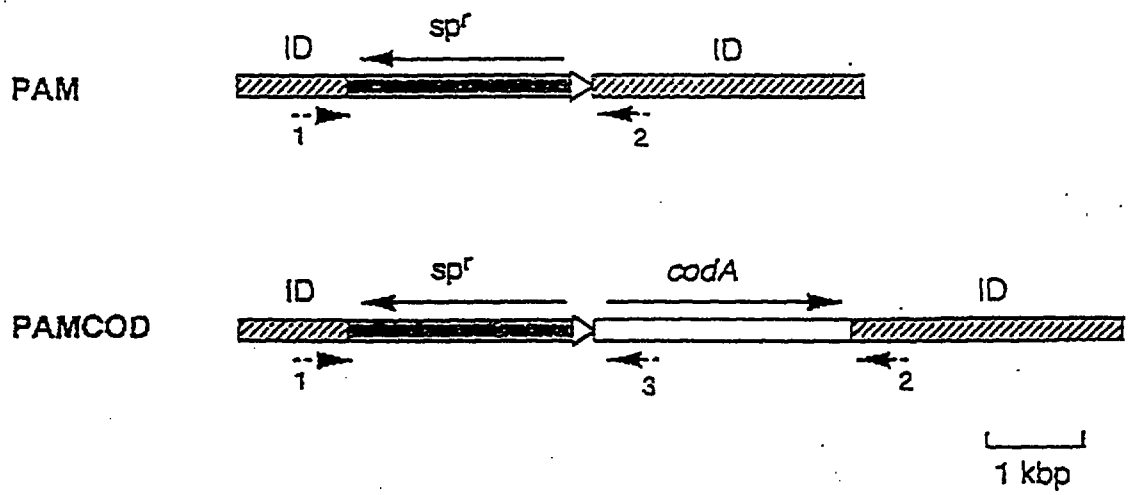


Fig. 2

A



B

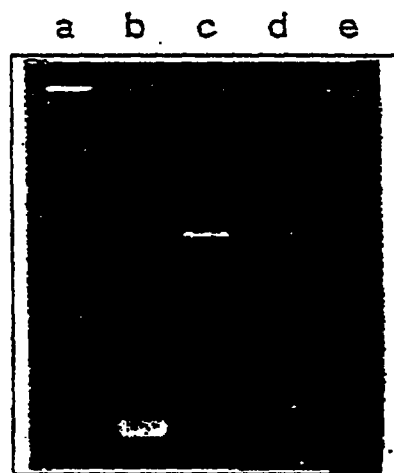


Fig. 3

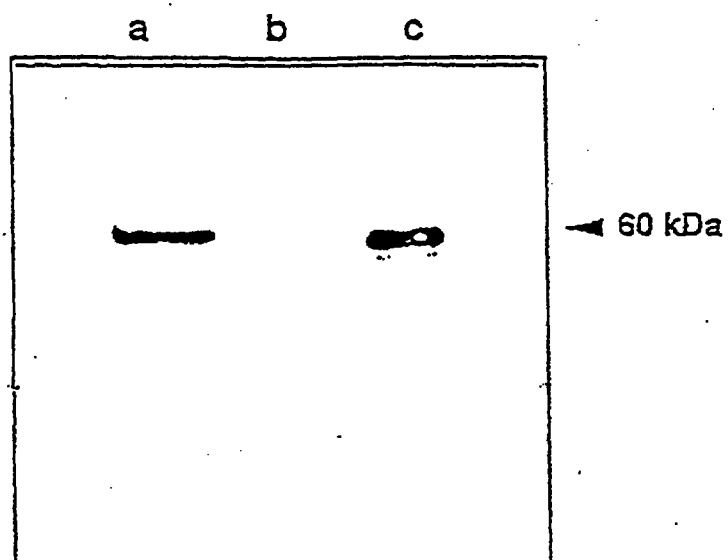


Fig. 4

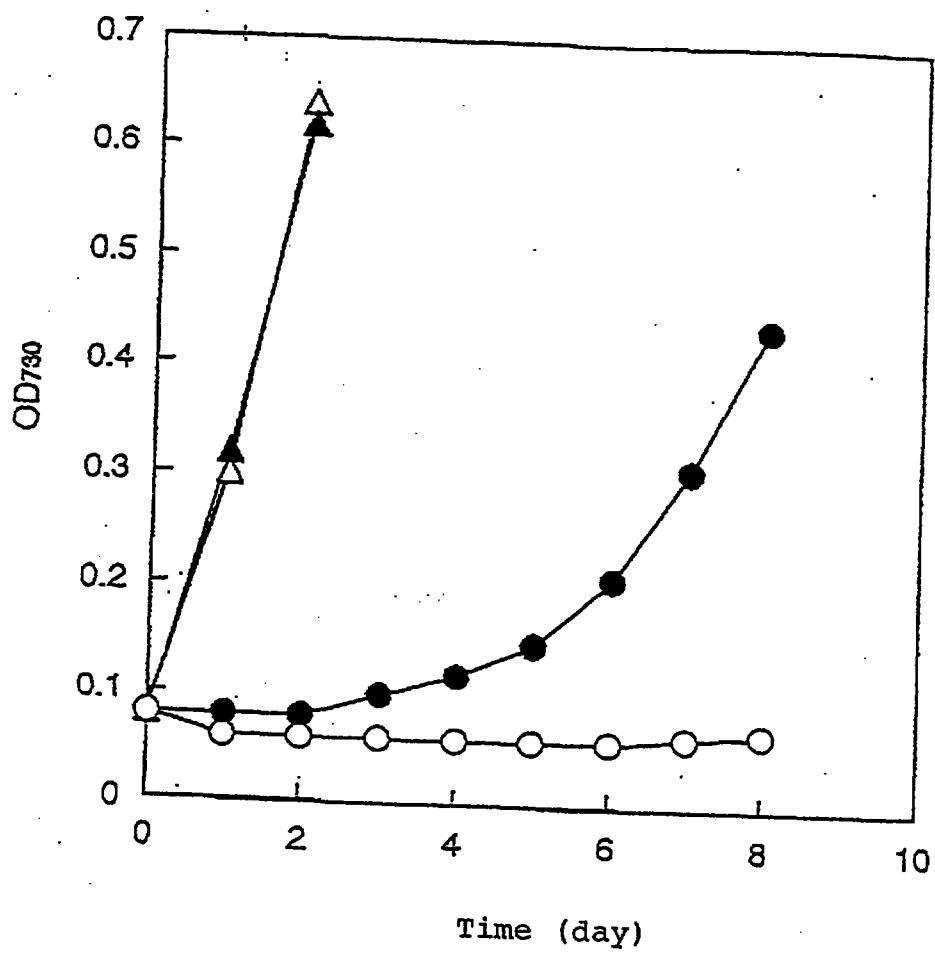


Fig. 5

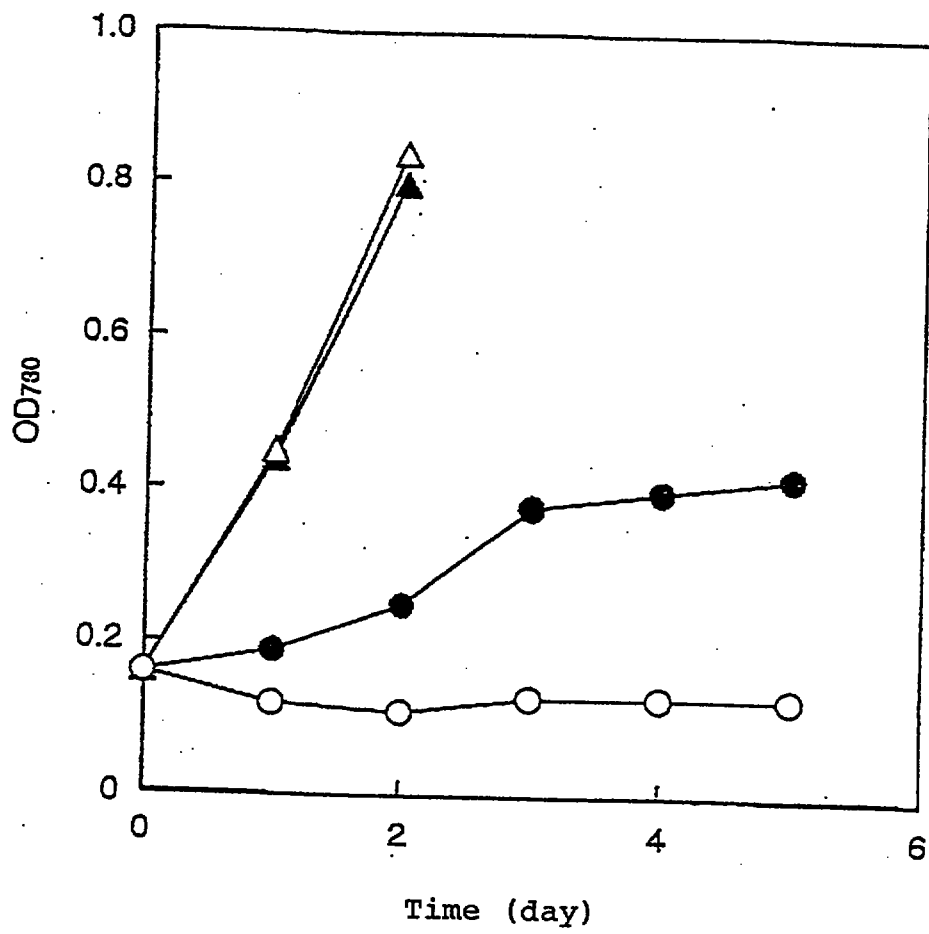


Fig. 6

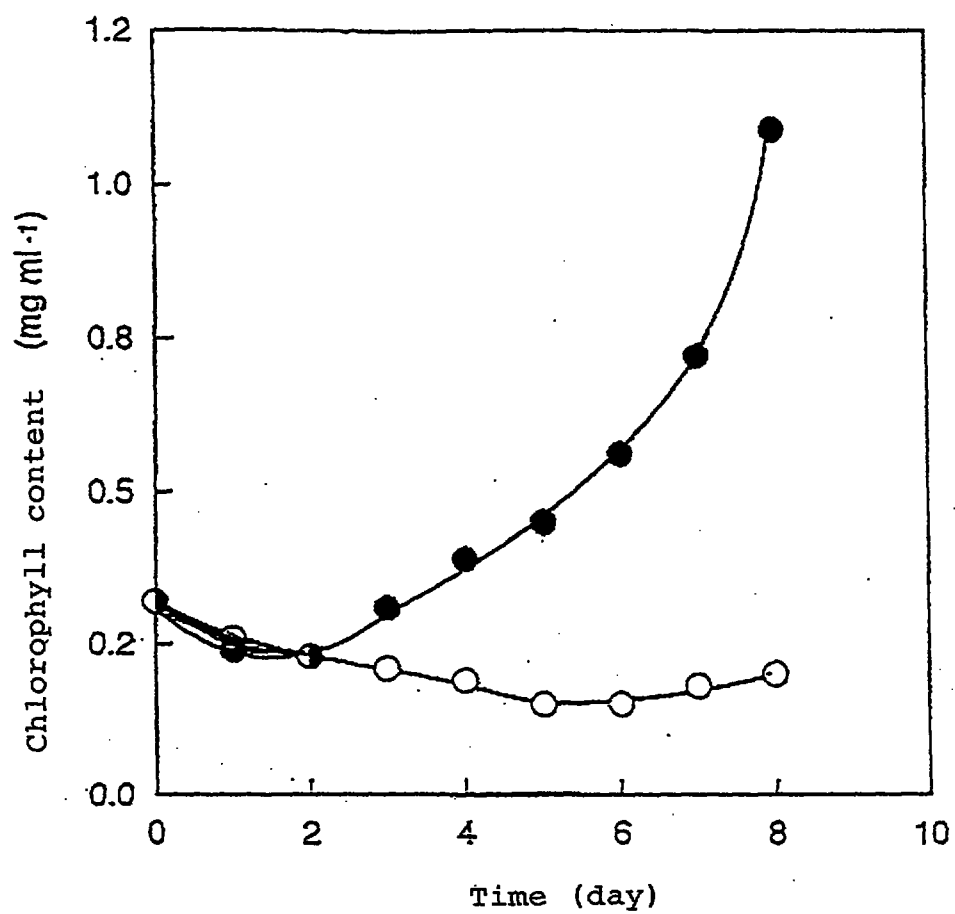


Fig. 7

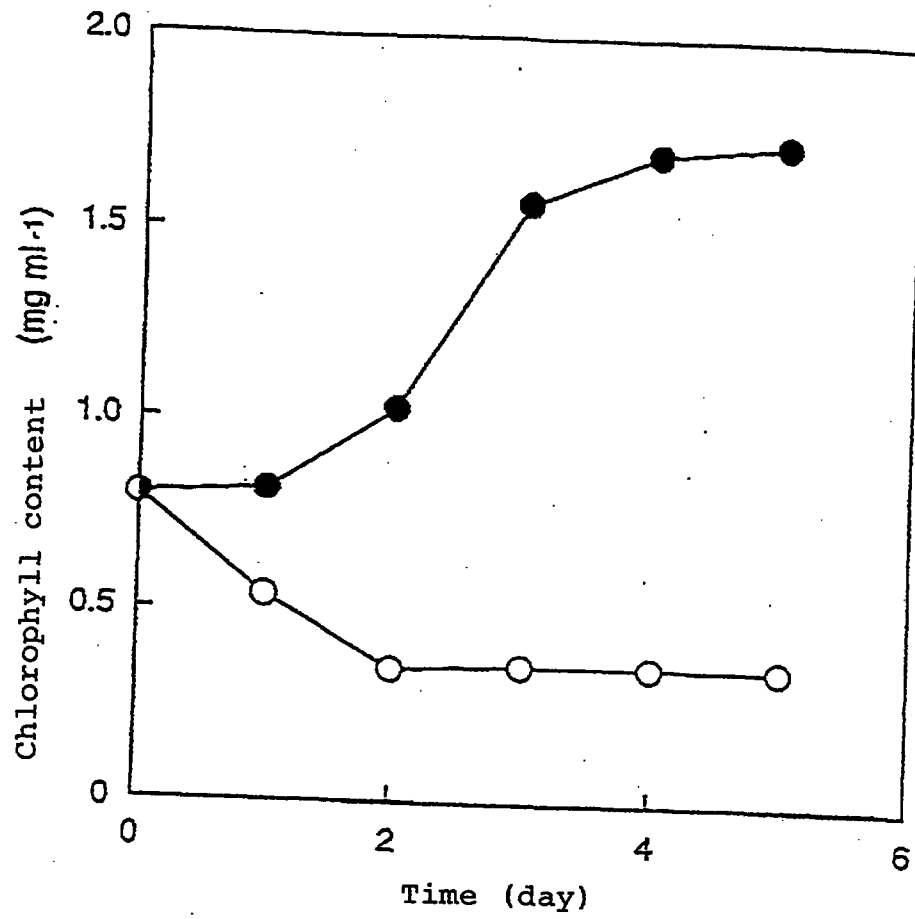


Fig. 8

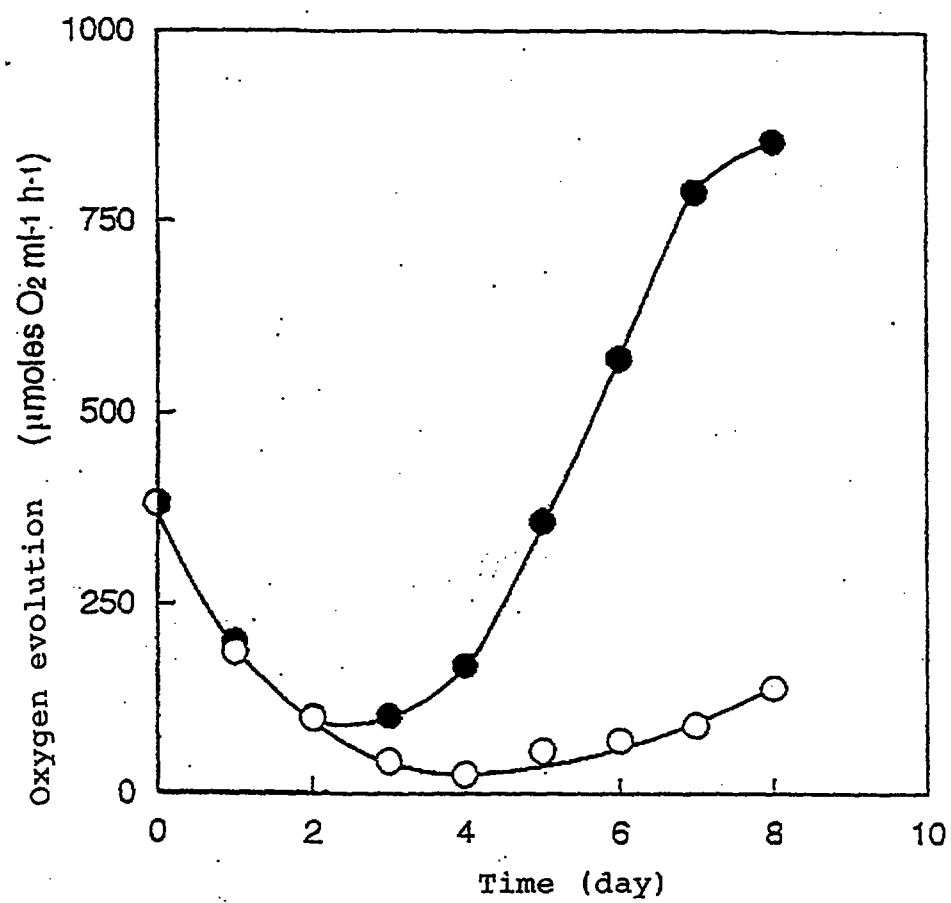


Fig. 9

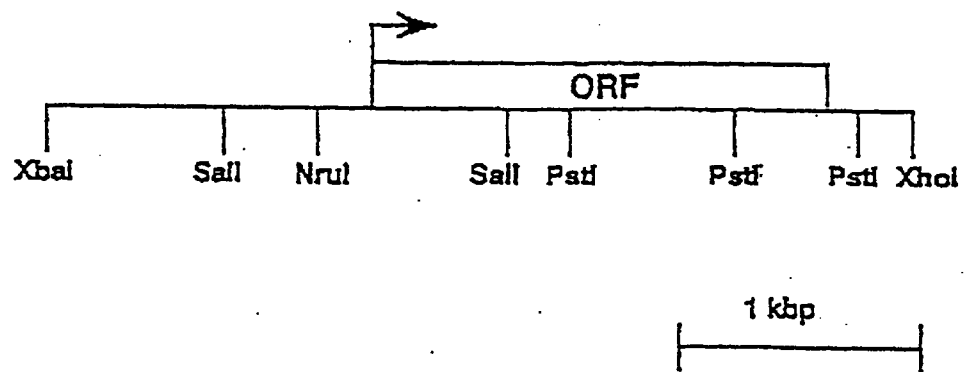


Fig. 10

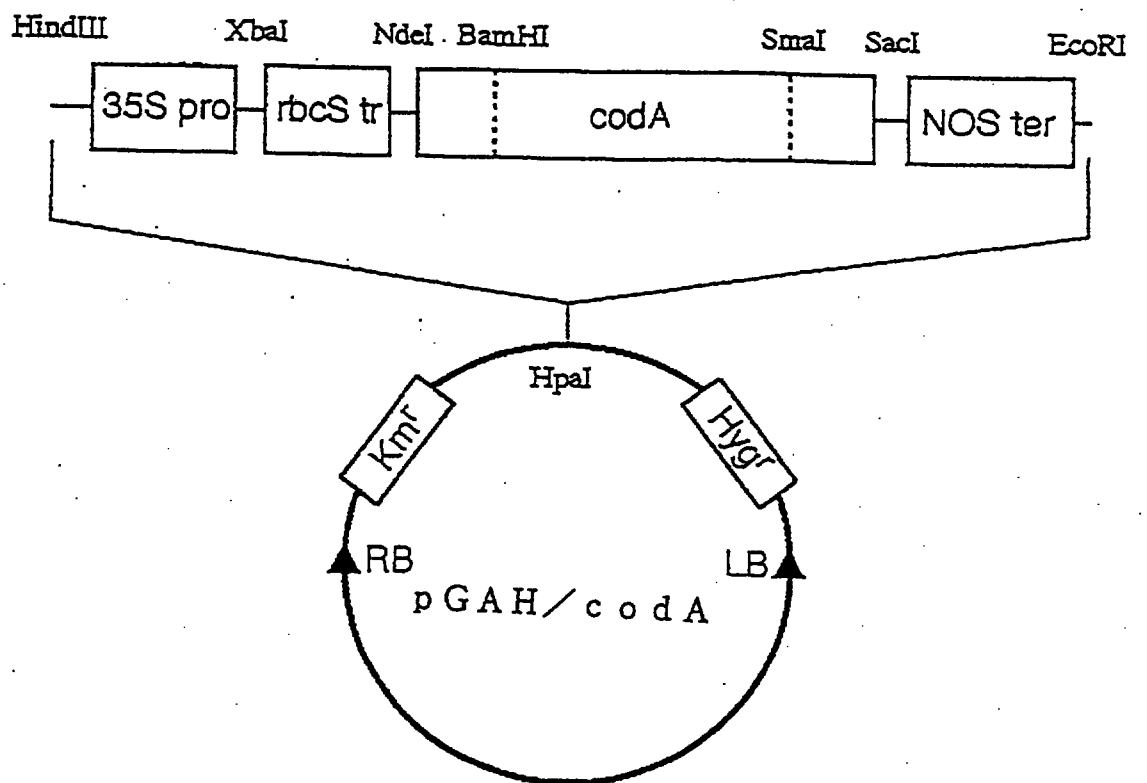


Fig. 11

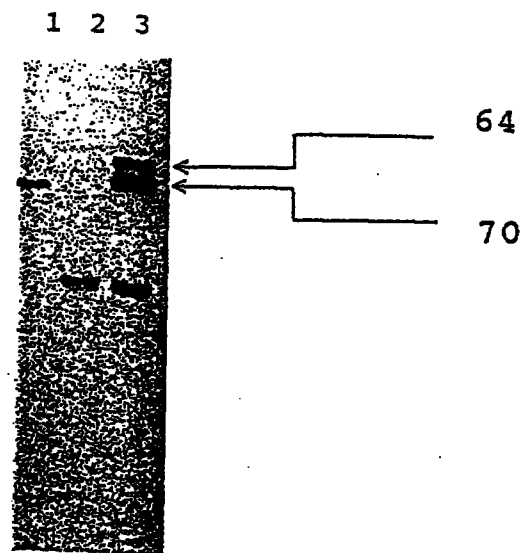
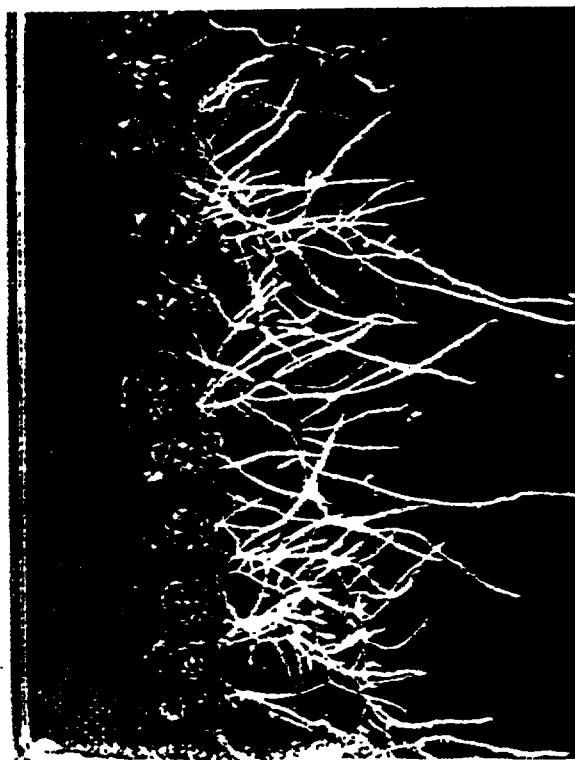


Fig. 12

B



A

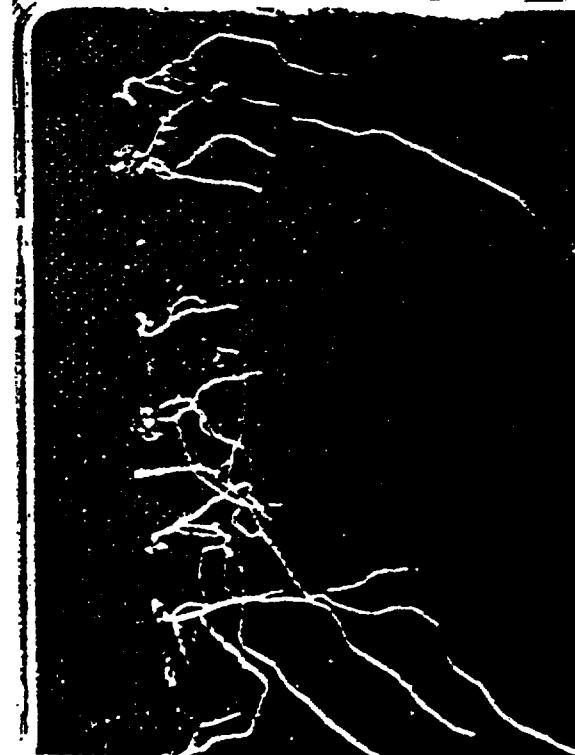


Fig. 13

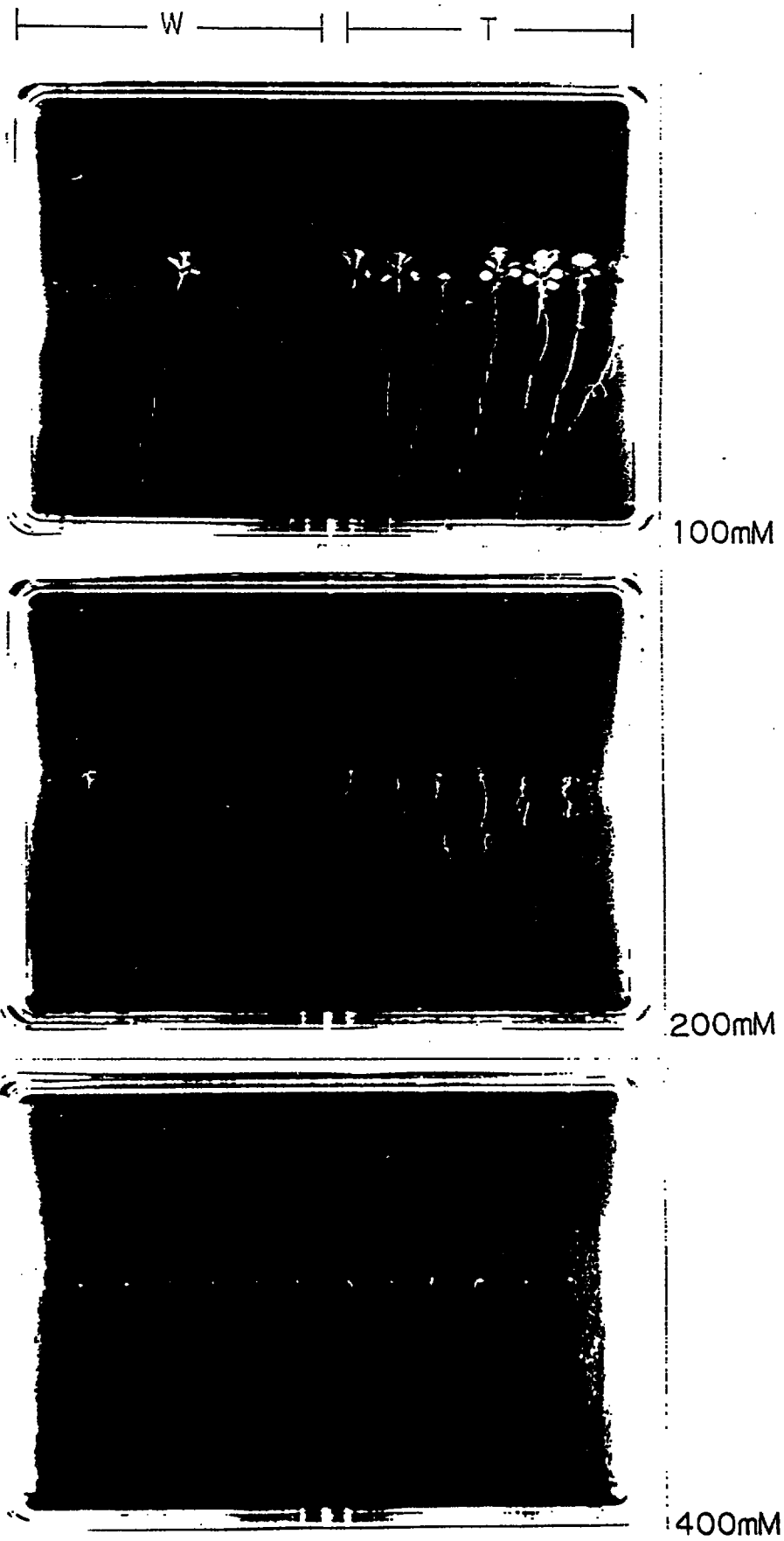


Fig. 14

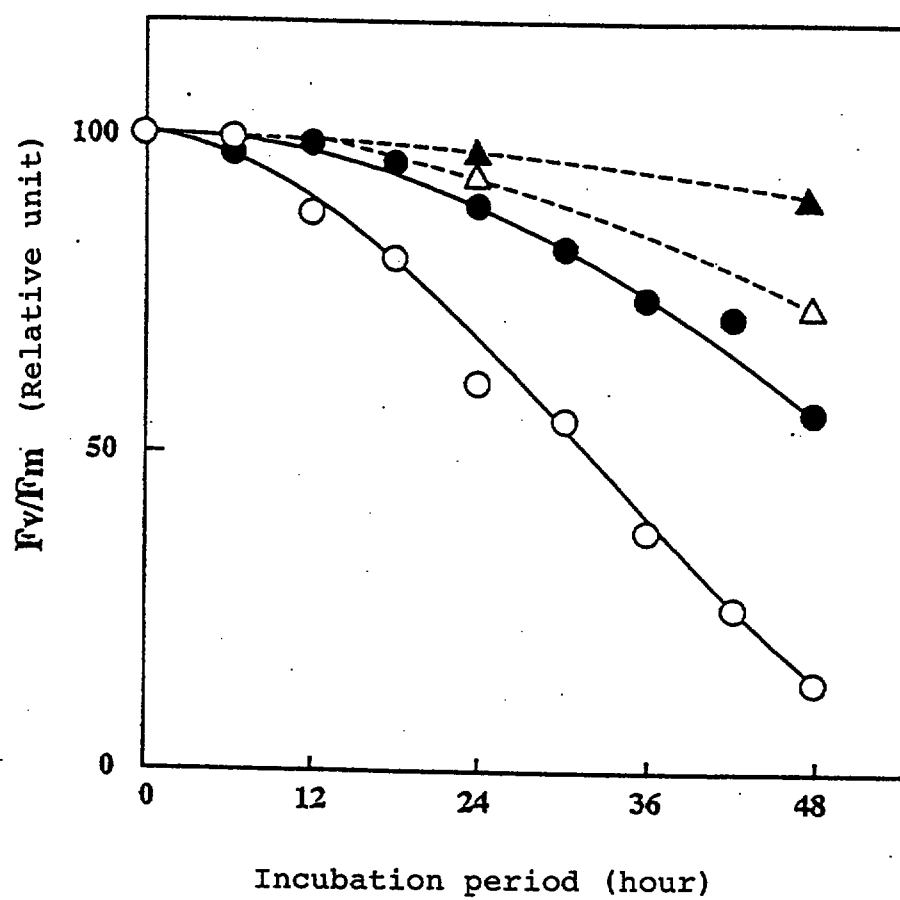


Fig. 15

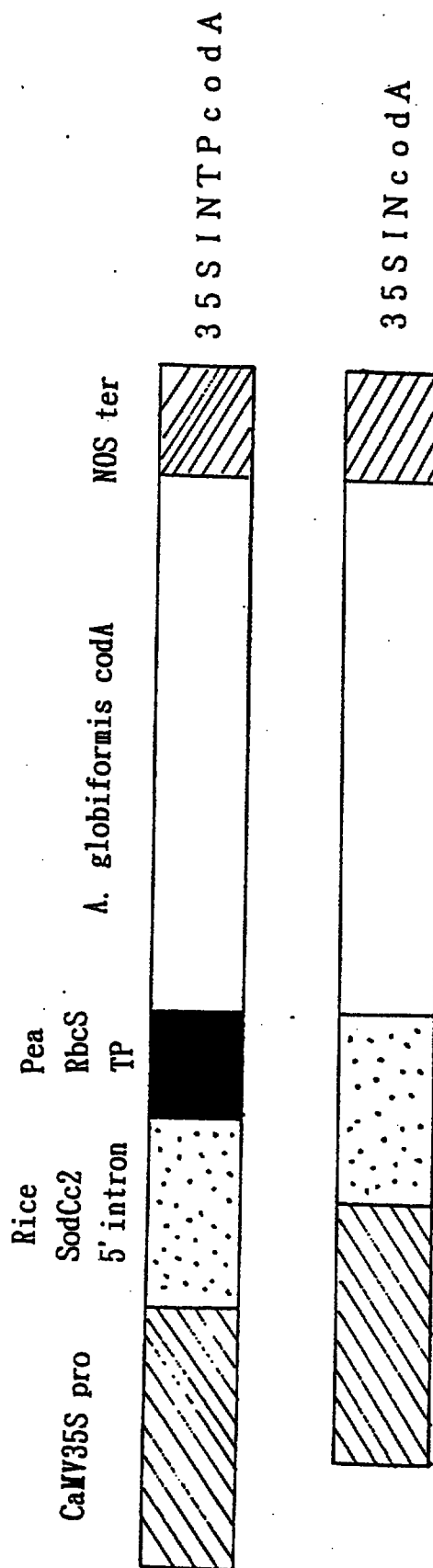


Fig. 16

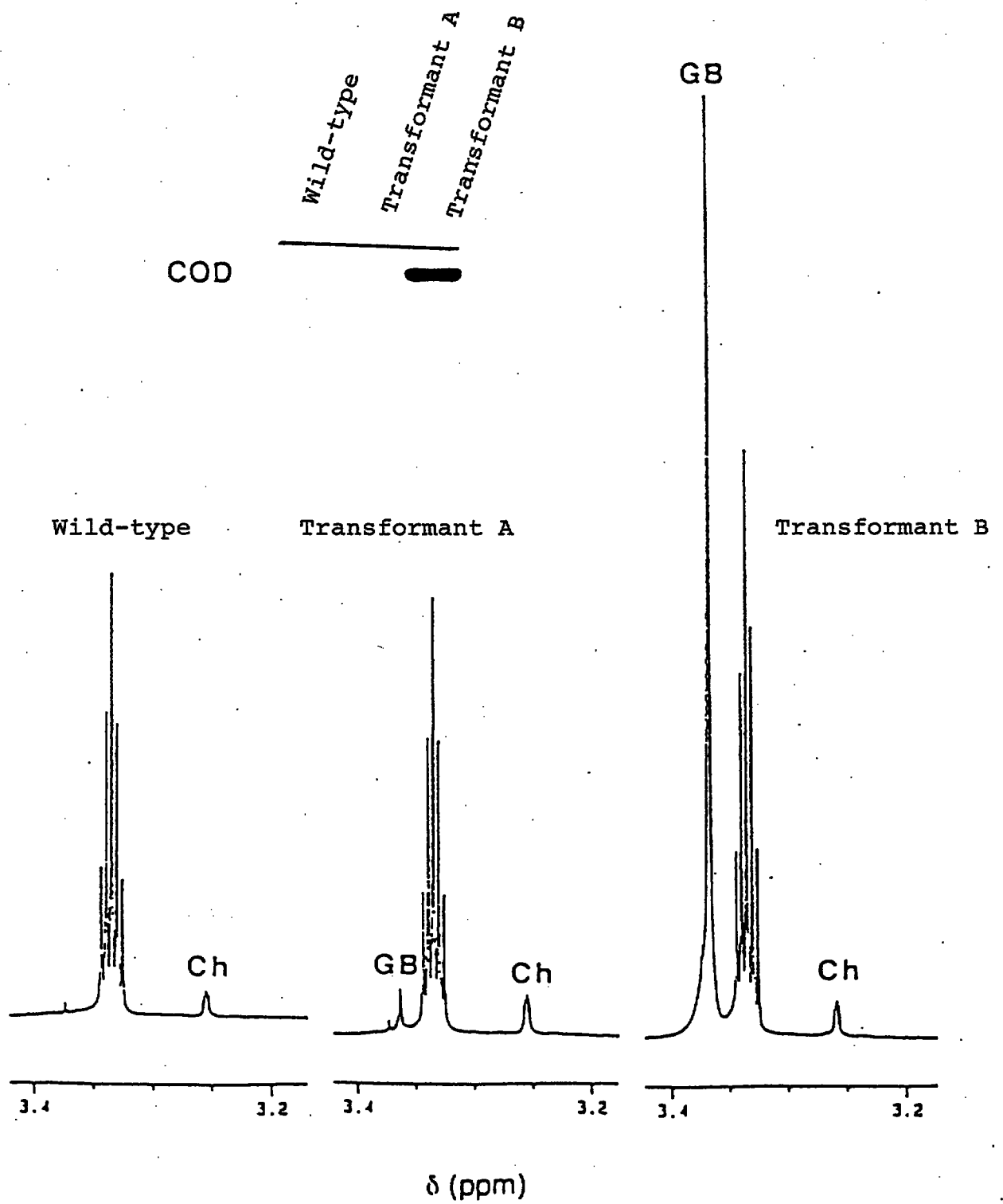


Fig. 17

