

CORRECTED VERSION

(19) World Intellectual Property
Organization
International Bureau



(10) International Publication Number
WO 2014/020624 A9

(43) International Publication Date
6 February 2014 (06.02.2014)

(51) International Patent Classification:

A01N 57/10 (2006.01) A01P 5/00 (2006.01)
A01N 63/00 (2006.01) A01P 7/00 (2006.01)
A01N 65/00 (2009.01) A01P 13/00 (2006.01)
A01P 1/00 (2006.01) A01N 57/16 (2006.01)
A01P 3/00 (2006.01)

HN, HR, HU, ID, IL, IN, IS, JP, KE, KG, KN, KP, KR, KZ, LA, LC, LK, LR, LS, LT, LU, LY, MA, MD, ME, MG, MK, MN, MW, MX, MY, MZ, NA, NG, NI, NO, NZ, OM, PA, PE, PG, PH, PL, PT, QA, RO, RS, RU, RW, SC, SD, SE, SG, SK, SL, SM, ST, SV, SY, TH, TJ, TM, TN, TR, TT, TZ, UA, UG, US, UZ, VC, VN, ZA, ZM, ZW.

(21) International Application Number:

PCT/IT20 13/000 193

(84) Designated States (unless otherwise indicated, for every kind of regional protection available):

ARIPO (BW, GH, GM, KE, LR, LS, MW, MZ, NA, RW, SD, SL, SZ, TZ, UG, ZM, ZW), Eurasian (AM, AZ, BY, KG, KZ, RU, TJ, TM), European (AL, AT, BE, BG, CH, CY, CZ, DE, DK, EE, ES, FI, FR, GB, GR, HR, HU, IE, IS, IT, LT, LU, LV, MC, MK, MT, NL, NO, PL, PT, RO, RS, SE, SI, SK, SM, TR), OAPI (BF, BJ, CF, CG, CI, CM, GA, GN, GQ, GW, KM, ML, MR, NE, SN, TD, TG).

(22) International Filing Date:

10 July 2013 (10.07.2013)

(25) Filing Language:

English

(26) Publication Language:

English

(30) Priority Data:

NA2012A000046 2 August 2012 (02.08.2012) IT

Published:

- with international search report (Art. 21(3))
- with information concerning authorization of rectification of an obvious mistake under Rule 91.3 (b) (Rule 48.2(i))

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(88) Date of publication of the international search report: 8 May 2014

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(48) Date of publication of this corrected version: 27 November 2014

(81) Designated States (unless otherwise indicated, for every kind of national protection available): AE, AG, AL, AM, AO, AT, AU, AZ, BA, BB, BG, BH, BN, BR, BW, BY, BZ, CA, CH, CL, CN, CO, CR, CU, CZ, DE, DK, DM, DO, DZ, EC, EE, EG, ES, FI, GB, GD, GE, GH, GM, GT,

(15) Information about Correction: see Notice of 27 November 2014



WO 2014/020624 ^ 9

(54) Title: COMPOSITION COMPRISING NUCLEIC ACIDS OF PARASITIC, PATHOGENIC OR WEED BIOLOGICAL SYSTEMS FOR INHIBITING AND/OR CONTROLLING THE GROWTH OF SAID SYSTEMS

(57) Abstract: The present invention concerns a DNA fragment mixture for the prevention or for the treatment against at least one pathogenic, parasitic or weed species of the plants or the environment, wherein said total DNA consist of random fragments of the total DNA of said at least one pathogenic, parasitic or weed species, and/or at least one phylogenetically similar species, against which said prevention and treatment are directed. Further, the invention concerns a process and relative system allowing the improvement of the production/growth of microorganisms at high yield in bioreactors or photobioreactors, or of plants in different culture systems, characterized in that the nucleic acids of the same organisms produced by said process are removed from culture medium and said culture medium, deprived of said nucleic acid, is used again in said process.

COMPOSITION COMPRISING NUCLEIC ACIDS OF PARASITIC,
PATHOGENIC OR INFESTING BIOLOGICAL SYSTEMS FOR
INHIBITING AND/OR CONTROLLING THE GROWTH OF SAID
SYSTEMS AND PROCESS FOR THE PREPARATION THEREOF

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The present invention concerns nucleic acid compositions of parasitic, pathogenic and infesting biological systems for inhibiting and/or controlling said systems and process for preparation thereof. Further the invention concerns a process and related system allowing the improvement of the production/growth of microorganisms at high yield in bioreactors or photobioreactors, or plants in various culture systems, by separation of nucleic acids produced by said organisms.

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Description

Back-ground of the art

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Since many years, the research for new products inhibiting biological systems demands an enormous scientific, economic and industrial effort and includes the detection of active principles from natural sources or by synthetic way and successive biological and pharmacological tests, both in vitro and in vivo, in addition to field and clinical application tests (Morgan et al. 2011. Health Policy, 100:4-17).

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The drug use (pesticides, herbicides, antibiotics) for the inhibition of biological systems concerns various fields of application, among which there are mainly agriculture and medicine and it is finalized to the control, inhibition or elimination of harmful organisms, whether pathogenic, parasitic, or infesting, and also treatment against the formation of microbial biofilms.

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In spite of the wideness and diversification of such application range, the use of drugs displays shared problems as (i) the specificity of action, (ii) the possible toxicity for human and other species, (iii) the environmental contamination and (iv) the onset of resistances in the populations of the target organisms.

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Generally the drugs (pesticides, herbicides, antibiotics) consist of small and middle size organic molecules and display a large diversification both

in terms of chemical components and action mechanisms. Such drugs have been detected generally based on methods of "random screening", that is estimating the activity and the use potentiality of chemical compounds obtained by synthesis in random way starting from various basic structures resulting from naturally occurring backbones. Accordingly

5 in the course of the last 50 years, a very large number of products, among which more recently also polynucleotides, were marketed.

One of the main problems related to the currently active principles used in agriculture and medicine is not sufficient action specificity thereof for the parasitic, pathogenic and/or infesting organisms and consequently side effects and toxicity on the host organism. For example are known the cases of Paraquat associated to the onset of the Parkinson's disease and products proved to be lethal for human embryos. In addition to the problems of acute toxicity, the chronic exposure to some products displays

10 a recognized risk for cancerogenicity, for which the enforced norms demand for ecotoxicological monitoring and use modalities associated to environmental dynamics and persistence of the various products.

At last, it is not to be ignored the problem of the environmental contamination caused by drugs (pesticides, herbicides, antibiotics). The recent applications of regulations in environmental field have drastically reduced the authorizations to the use of phytodrugs and pesticides and many products will have to be abandoned in the near future for the entering into force of more restrictive regulations. For example, methyl bromide, widely used as fumigant for the control of pathogens in the soil,

15 among which nematodes, has been prohibited due to the inclusion thereof among compounds considered responsible for damage to the ozone layer. Taking into consideration these problems, the majority of more modern active principles were selected for a fast decomposition thereof in the soil or crops. However, this possible advantage in terms of environmental contamination can limit the effectiveness over the time of the used active principles.

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Another shared fundamental problem for currently used products, both in agricultural field and therapeutic applications in medical field, is the

appearance of pharmacological resistance after a period from the starting of the treatment.

For example, with reference to agriculture, in USA, in the early sixties, 2,4 D has been the first herbicide, used on wide scale, for which the onset of resistance in populations of *Dacus carota* was reported (Switzer, (1957), Proc. North Eastern Weed Cont. Conf., 11:315; Whitehead and Switzer, (1963), Can. J. Plant Sci., 43:255). Successively, starting from early 70 years numerous cases of infesting species resistant to triazines (Ryan, (1970), Weed Sci., 18:614; LeBaron and Gressel, Eds., Herbicide Resistance in Plants, Wiley, New York, 1982) were reported. The records then has grown exponentially in the following decades together the diffusion of the new products for chemical and biological fight and intensive use thereof (Powles and Shaner, 2001. Herbicide resistance and world grains, CRC press, New York).

Various onset mechanisms of infesting resistance to herbicide treatment are known. For example, various mechanisms reported in literature are: resistance as a result of selection of multiple copies of target genes (Gaines et al. 2010, Proc. Natl. Acad. Sci. 107(3): 1029-1034); mutation of the target genes (Wakelin et al. 2006. Weed Res. (Oxford), 46(5): 432-440); vacuolar sequestration (Ge et al. 2010. Pest Management Sci. 66:576); expression of enzymes that metabolize the herbicides (Hidayat et al. 1997. Pest. Biochem. Physiol. 57(2): 137-146).

Currently in the agricultural field, the appearance of resistance in over 200 infesting species was detected, selected above all as a result of the use of active principles characterized by the same mechanism of action. In Italy, in Po plain, cases of resistance to Atrazines in populations of infesting herbs are known, like *Solanum nigrum*, *Chenopodium album* and *Amaranthus cruentus*, or, in other regions, those with resistance to enzymatic inhibitors in different other infesting species (*Lolium* spp, *Phalaris paradoxa*, *Papaver rhoeas*, *Sinapis arvensis*, *Echinochloa crus-galliums*, *Sorghum halepense*, *Cyperus difformis*).

Also in medical field, the development of resistance to drugs makes useless the use thereof, constituting one of greatest problems related to

antibiotic therapies (Clatworthy et al. 2007. Nat. Chem. Biol. 3(9): 541 - 548). Although enormous human and economic resources in order to discover new molecules and to study the resistance mechanisms were used, the appearance of the resistance to drugs proved to be faster than the discovery of new drugs.

The problem of the drug-resistance became primary world-wise in the field of health as it results in clinical complications for the antibiotic therapy (increase of the duration of diseases, increase of complications, possibility of epidemics) and demands for additional costs in the treatment of the infections from antibiotic-resistant bacteria (by means of use of further drugs, thus lengthening of the hospitalization). A case of antibiotic-resistance, more and more present and dangerous for the public health and representing one of the main causes of hospital infections, is meticilline resistance of the staphylococci and above all of various strains of *Staphylococcus aureus*. These bacteria, by means of the mechanism of reduced affinity for the target, are able to express a modified protein that does not interact with beta-lactam antibiotics. The European Union has faced this problem presenting a "Communitarian Strategy against the Antimicrobial Resistance" wherein a prudent use of antimicrobial agents in medicine is recommended in order to contain the development of resistances.

The possibility to inhibit the expression of specific genes resulted in great interest since the first publication of Zamecnik and Stephenson paper wherein it is shown that the use of compositions based on antisense oligonucleotides (ASO) could inhibit the viral protein translation (Zamecnik and Stephenson (1978) Proc. Natl. Acad. Sci. 75:285-288). Following this discovery, the most recent experimentations on techniques of fight to parasitic, pathogenic and infesting agents further investigated about the use of various RNA and/or DNA small size molecules (in the order of various tens of bases) selectively binding to targets displaying complementary sequences like mRNA, microRNA, or mitochondrial RNA, thus inhibiting the translation thereof and therefore carrying out a control activity for the targeted species.

These modern methodologies found applications in the fight against parasitic species (e.g. WO 2005/049841 A 1), infesting plants (e.g. WO 201 1/1 1257 A 1) and antimicrobial and oncologic therapy.

5 The currently used approaches are multiple, among which the modification of RNA in order to control the level of specific RNA messenger thus limiting the expression of the gene encoding the same. Such methodology is characterized by the administration by means of suitable constructs of a RNA or DNA sequence aimed to a specific target, for example a gene or parts thereof. The problems associated to the use of such therapeutic
10 approaches include the necessity of characterization for a specific sequence as therapy target. There is also the difficulty to obtain an effective minimal concentration of the construct in the target organ. The recent history of the antisense technology, in addition, reveals that while the identification of ASOs binding specific RNAs is now relatively simple, formulation thereof in compositions displaying an effective application
15 potentiality for the inhibition of the expression of specific genes being still problematic. Various recent studies addressed aforesaid problem, proposing more stable and effective structures and compositions (e.g. WO 201 1/031 520 A 1). As it is well-known this type of techniques displays elevated selectivity in terms of specificity of the action site. Up to now cases of the onset of resistance to the use of ASOs in herbicide and pesticide treatment are not reported as their application has not still reached the stage of commercialization and consequent dissemination of the products. However, taken into consideration the particular action
20 specificity of the ASOs, it is realistic to presume that, as a result of extensive use thereof and consequent selective pressure on the infesting, parasitic and/or pathogenic populations, the application of said ASOs will result in the onset of resistances in the treated populations thus making ineffective the treatments, in the same way as it occurred in the past for other drug types characterized by elevated specificity of action.
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In the field of the bulk cultures in bioreactors and photobioreactors, therefore in confined controlled systems, one of not yet solved major problems in the bulk culture of microorganisms is the onset of saturation

phenomena of the growth curve for said organisms (stationary stage), with consequent impossibility to exceed certain concentrations for volume unit. A typical growth curve of such microorganisms (bacteria, photosynthetic bacteria, yeasts, microalgae and blue-green algae) is characterized by a starting latency stage, a successive exponential stage and a steady-state stage with saturation of the growth curve, followed sometimes also from a cellular death stage.

The impossibility to exceed the threshold of the maximum concentration for volume unit that a microorganism can reach during the culture on large scale in reactors and systems of bulk culture, remarkably limits the potentialities of exploitation of such organisms not allowing ulterior increments of the production beyond the stationary stage.

Various motivations are proposed as cause of the decrease of the cell growth rate of the microorganisms in saturation stage, all mainly related to the limitation of nutritive elements in medium, or auto-limitation of the culture penetrating radiation in the case of photosynthetic organisms (self-shading phenomenon). Various experimental studies demonstrated that the phenomena of the growth curve saturation occur as well in the absence of apparent limiting factors and this phenomenon is associated with accumulation of inhibiting substances in culture medium.

In the light of above exposed it is therefore apparent the need for new methods for the inhibition of harmful organisms or the increase of yield in the production processes of microorganisms overcoming the disadvantages of the prior art methods.

The title invention results from an important observation occurred during ecological studies on the cycle of the organic substance in the soil. In these studies it was observed that the DNA, released by the decomposition of soil accumulated organic material, proves to be inhibiting for the plant species producing said organic material, meanwhile said DNA does not inhibit other species for which said DNA can on the contrary represent a nutritive resource.

In the case of forest ecosystems this inhibiting effect of specific extracellular DNA only on species of plants with homologous DNA,

represents a regulating mechanism of the natural co-existence among various species and favours the biodiversity of plant communities. In fact, when a single species, for any reason, is under conditions suitable to increase the own dominance on other species, in the time, necessarily, will accumulate larger residues of self organic substance, whose decomposition and consequent DNA release will produce an inhibiting effect on the same species thus reducing the competitive ability and therefore the dominance. On the same phenomenon is based the so-called "fatigue of the soil" (either referred as "soil sickness") in agriculture, that is productivity losses of the repeated mono-specific cultures in the time not resulting from nutritive nature problems, but from accumulation of plant residues of the same cultures. The observation of species-specific inhibition due to accumulation of self DNA, during the cycle of the organic substance, explains the aforesaid phenomena of productivity loss and said finding never was reported in the scientific literature.

On the base of these studies, a second important result was reached, consisting of the observation that microbial populations (for example bacteria, microalgae and fungi) produce and secrete, during the growth, self DNA molecules in form of different size fragments which, by accumulation in the population growth substrate, exert a inhibiting effect on the growth of said population.

It is known that a microbial population typically grows initially in exponential way then enters in a so-called stationary stage, followed, possibly, from a death stage. Various growth limiting factors of such microbial populations are known, among which, fundamentally, the nutritive limits and/or accumulation of toxic substances of various nature. In literature it is reported the production and secretion of extracellular DNA from various organisms (Peters and Pretorius, 2011. *Chemical Clinics Acta*. 412:806-811), but up to now it is not reported that an extracellular DNA release during the growth of a microbial population produces an inhibiting effect on the growth of said population, in absolutely analogous way as observed for the plants affected by the accumulation of self DNA in the soil. The observations this invention is based on reveal a general,

biological law according to which the release of self DNA by an organism proves to be regulating the growth of said organism and of the same species population organisms. This result leads to new remarkable application scenarios in agro-pharmaceutical field. Particularly, the control of any harmful species becomes possible through the exposure to self total, random fragmented DNA, such to reproduce what observed in the natural cycles of decomposition of the organic substance or in the extracellular secretion phenomena during the growth of the microbial populations. Thus it is hypothesized a possible application use of compositions whose active principle consists of total and random fragmented DNA of a to be controlled species.

The tests that are illustrated below demonstrate the effectiveness of such method for any treated species. Those reported are easy replicable experiments and do not demand any a priori knowledge of the genome of the to be treated species in terms of sequencing and/or detection of specific target genes. For the preparation of the inhibiting compositions only the extraction of the total DNA from organic material of the target species (for example from leaves of plants, fungus mycelium, microbial biomass) and successive random fragmentation of said extracted DNA are sufficient. The inventive set up procedure simulates the process of natural decomposition of the organic substance in the soil and produces a mixture of variously sized polynucleotides comprising from tens to thousands of bases. Such fragments, representing the total DNA of the to be controlled species, produce highly selective effect on organisms with homologous DNA whereas not act on a single target gene as it occurs, on the contrary, in the case of the various types of antisense oligonucleotides. This aspect potentially constitutes a remarkable advantage because it eliminates the possibility of appearance of resistances in the treated populations, because for no organism it is presumably possible to develop simultaneously resistances to the inhibition of all self functions. It is known, in fact, according to the scientific literature, that the onset speed of resistance phenomena in populations of parasitic, pathogenic and infesting agents is closely correlated to the number of genes responsible

for such resistance. More precisely, lower the number of necessary genes for the resistance onset, greater the speed according to which this resistance will be positively selected inside of the to be controlled population (Prather et al. the 2000. Herbicide resistance: definitions and management strategies. University of California, Division of Agriculture and Natural Resources, Oakland - Brent and Hollomon, 2007. Fungicide resistance in crop pathogens: how can it be managed? Croplife International, Brussel).

The invention concerns therefore the use of compositions for inhibiting parasitic, pathogenic and/or infesting species wherein, like active principle, random fragments of total self DNA, identical or similar to the same genome of the to be inhibited species are used. Further the present invention concerns a composition comprising said random fragments of total DNA.

In addition the inventors of the present invention developed a process and the related system for the production/growth of microorganisms in bioreactors or photobioreactors or plants in systems for hydroponic culture, on substrates outside the soil, or in soil, by means of separation of nucleic acids of said microorganisms or plants.

Below in the present patent application, the term "saturation level" means that stage established in the course of growth of the microorganisms, when the concentration of nucleic acids or fragments thereof in culture medium is such to result in a steady-state stage regarding the course of growth of the microorganisms, with saturation of the growth curve thereof.

In the same way, the term "exhaust medium" means a culture medium wherein the level of concentration of nucleic acids and fragments thereof reached a concentration resulting in a saturation level on the growth course of the microorganisms.

As to the term "reclamation" it means the operation suitable to remove nucleic acids and fragments thereof from exhaust medium.

"Regenerated medium" means medium obtained after the reclamation of exhaust medium.

DETAILED DESCRIPTION OF THE INVENTION

The present invention is based on the experimental evidence that the exposure of an organism to extracts of total self DNA results in an inhibiting effect on the functions of said organism, thus limiting the growth or interfering in fundamental way with physiology thereof at cellular level.

5 In detail, the total DNA of a parasitic, pathogenic and/or infesting organism extracted and administered without no selection of specific fragments or use of specific constructs, can be used as inhibiting product for the same organism or other organisms with identical or similar genome. The inhibiting action is exerted when the parasitic, pathogenic and/or infesting
10 species is contacted, within growth substrate thereof or at systemic level, with randomly fragmented total self DNA. Once absorbed, such DNA produces a limiting effect on the growth and a generalized inhibiting effect on the various functionalities of the organism, otherwise than it occurs in the case of applications of specific nucleic acid sequences aiming to "gene
15 silencing" or "RNA interference" effects. The adsorption and the inhibiting activity depend on the concentration and fragmentation level of the nucleic acids the system is exposed to. A greater effect is pointed out depending on the fragmentation degree of used molecules (for example after sonication or following natural decomposition). Experiments were carried
20 out using various average lengths of DNA fragments and the inhibiting effect proves to be more effective for fragment mixtures with higher length frequency of approximately 200 bases.

Specifically, the production of random fragments of total DNA extracted from various species according to illustrative experimentations was carried
25 out with various approaches in order to demonstrate that the fragmentation technique does not influence the effectiveness of the inhibiting effect. Said fragmentation can be carried out by means of natural degradation (exposure to environmental factors), artificial degradation (by combustion, enzymatic or mechanical way), or by synthesis (methods for
30 random amplification like DOP, that is by PCR with degenerated oligonucleotides).

The authors of the present invention carried out studies on plants, fungi, insects, yeasts, algae and protozoa. The results of such experimentation

unequivocally demonstrated the general subsistence of the described inhibiting effect on any treated species.

In the case of the plants, the presence in aqueous solution of the total DNA of a species, obtained by simple extraction and randomly
5 fragmented, also at low concentration, in contact with the roots of the same species, quickly produces radical necrosis, leaf chlorosis, interruption of the meristem activities and, when in contact with the seeds, suppression of the germination ability. On the contrary, the exposure of the roots and/or seeds of a species to solutions of randomly fragmented
10 total DNA of different species does not produce noticeable inhibitions, also for exposures at high concentration, i.e. showing a highly species-specific effect. It is therefore apparent that the presence of randomly fragmented total DNA in a growth environment can carry out a selective inhibiting function on organisms with identical or similar genome not influencing
15 other species. It is possible to find out this inhibiting effect both when the total DNA is artificially fragmented and inserted in the growth medium of the biological system and when the DNA is produced and secreted by said biological system during the growth thereof.

This therefore makes possible an action targeted only against the
20 parasitic, pathogenic and/or infesting species, by treatments with formulations containing DNA specific for this species, with the enormous advantage that said molecules will display the inhibiting effect only on the target species, but not on the to be protected or other species, also in case of possible repeated exposures.

The same principle observed by the inventors of the present invention can
25 be favourably employed in order to increase the yields of processes for production of microorganisms in bioreactors and photobioreactors, tanks for hydroponic culture and culture systems outside the soil. In fact, the inventors of the present invention, by means of laboratory tests using
30 bioreactors for growth of algae, yeasts and bacteria, observed a supernatant accumulation of variously sized DNA fragments (approximately 50-800 bp) belonging to the same produced microorganisms, during the growth stages at increasing concentrations. It

was observed that this accumulation occurs in the growth stages at increasing concentrations and that results in an inhibiting effect on the growth of said culture when in the bioreactors the steady-state stage and successive death stage are reached (figure 11A and 11B).

5 The experimental tests pointed out that a self-inhibiting species-specific effect exists, as cultures of a given species display to be inhibited only by supernatants of cultures of the same species. Further it was observed that the removal of the DNA from the culture medium is suitable to remove the inhibiting effect allowing the growth stage to be restored and higher cell
10 densities in bioreactors (figure 11 C) and photobioreactors to be reached. Analogously, in tanks for hydroponic cultures of plants, in the presence of decreasing production in absence of complete replacement of the circulating nutritive solution, it was observed an accumulation in the substrate of DNA of the cultured species. Also in this case, the removal of
15 the accumulated DNA from the growth substrate allowed the productivity to be restored.

Taking into consideration these experimental results, therefore, it is possible to develop technological systems for the removal of DNA occurring in cellular cultures (bioreactors and photobioreactors),
20 hydroponic culture tanks and in culture systems outside soil in order the productions to be optimized.

It is therefore a specific object of the present invention the use of a composition comprising or consisting of a mixture of total DNA fragments wherein said total DNA is randomly fragmented for the prevention and the
25 treatment against at least one pathogenic, parasitic or infesting species of the plants or environment, wherein said total DNA is total DNA of said pathogenic, parasitic or infesting species, and/or at least one phylogenetically similar species, against which said prevention and treatment are directed. As above said, the fragment mixture according to
30 the present invention is obtained by fragmentation of the total DNA extracted from said pathogenic, parasitic or infesting species against which said prevention and treatment are directed, without carrying out any selection of fragments, that is using all fragments obtained from the acid

nucleic. Therefore according to the present invention randomly fragmented whole DNA, that is without carrying out any selection of specific and/or specifically sized sequences, is used. Fragments of said total DNA are obtained by means of random fragmentation of the extracted total DNA, or
5 by means of random fragment synthesis starting from the total DNA. The substantial difference of the technique of the present invention compared to the "gene silencing" and "RNA interference" current techniques is that in the current state of the art the specific regions of to be used nucleic acids must be actively selected, whereas in the case of the technique of the
10 present invention nor previous knowledge neither selection for the to be used specific sequences are necessary. Said total DNA can be extracted from said at least one pathogenic, parasitic or infesting species or artificially synthesised. The nucleic acid can be amplified and/or fragmented by means of chemical and/or physical type procedures and
15 the obtained fragments possibly can be further amplified. The production methods of random fragments of total DNA include, for example, chemical, biochemical and molecular methods, sonication techniques, heat treatments and pyrolysis procedures,. For example, following the extraction of total DNA from leaf tissue from *Arabidopsis thaliana*, the
20 exposure of the seeds from the same species to such extract containing DNA in not fragmented form does not display significant inhibiting effects. Differently following a sonication treatments of the extract using a dipping sonicator for three or four cycles lasting three minutes at 80% power produces a mixture of variously sized random fragments with increasing
25 inhibiting activity, in function of reached fragmentation degree, on the germination of *A. thaliana* seeds. The same mixtures of random fragments of total DNA extracted from *A. thaliana* did not display inhibiting effect on other treated species, as *Lycopersicon esculentum* and *Avena sativa* (used as control biotest in this experiment). Analogous results were
30 observed when the extract of total DNA was randomly fragmented with other above cited chemical-physical methods. Finally the same results are obtained exposing seeds of *A. thaliana* to mixtures of random fragments of the total DNA wherein said fragments were obtained by amplification of

total DNA using DOP. The composition can be used as biocide, herbicide, fungicide, insecticide, acaricide, nematocide, antiprotozoic, algaecide, bactericide. The composition can be administered to said at least one pathogenic, parasitic or infesting species by surface contacting, cytotropic administration, systemic administration by means of, for example, injection, ingestion or inhalation, or adsorption. The composition can be formulated in a form, for dry or liquid treatments, selected in the group consisting of dispersion, for example in form of aerosol, suspension, wettable or soluble powders, emulsions in water or other solvents, dispersible granules, suspensions of microcapsules, emulsifiable concentrates, fluid pastes, macro emulsions, oil dispersions, baits. According to a particular embodiment, the nucleic acid random fragments can be engineered in vectors. The composition according to the present invention can comprise further pesticides selected from the group consisting of fungicides, insecticides, nematocides, miticides, arthropocides, bactericidal, algaecides.

The invention concerns further a method for the prevention or treatment against at least one pathogenic, parasitic or infesting species of the plants or environment, said method comprising or consisting of the following steps of: a) extraction of the total DNA of at least one of said pathogenic, parasitic or infesting species against which said prevention and treatment are directed; b) production of random fragments of said total DNA in order to obtain a mixture of DNA random fragments; and c) contacting said mixture of random fragments with said pathogenic, parasitic or infesting species.

A composition comprising or consisting of a mixture of total DNA random fragments for the use in the prevention and treatment against at least one pathogenic, parasitic or infesting species of man or animal, wherein said total DNA is the DNA of said at least one pathogenic, parasitic or infesting species, against which said prevention and treatment are directed, is a further object of the present invention. Said total DNA random fragments can be obtained by degradation of the total DNA by means of chemical and/or physical methods, or by synthesis of random fragments of said total

DNA. The composition according to the present invention can be formulated for dry or liquid treatments, or aerosol dispersion, in form of suspensions, wettable or soluble powders, emulsions in water or other solvents, dispersible granules, suspensions of microcapsules, emulsifiable concentrates, fluid pastes and macro emulsions, oil dispersions, creams, capsules and tablets. The administration routes include contact or surface, systemic and cytotropic treatments, by injection, absorption, ingestion and inhalation, as well as any other administration route useful to the application end. The composition according to the present invention can include mixtures with other chemical compounds acting as tackifying agents, wetting agents, suspending agents, excipients, amending agent and solvents, surface-active agents, as well as other substances useful to the application end. At last, the composition according to the invention can include other pharmaceutical compounds in formulations and mixtures of not antagonistic but possibly synergic media with the activity of the treatments. In such cases the combination of compounds generally will correspond to weight ratios from about 0% to 100%.

It is also a specific object of the present invention a process for high yield production of microorganisms in bioreactors or photobioreactors, or for culture systems of plants both inside and outside soil, characterized in that the nucleic acids from the same organisms produced by said process are removed from culture medium and the culture medium deprived of said nucleic acid can be used again in said process.

According to the process of the invention, the nucleic acids are removed from the culture medium or growth substrate using separation techniques selected from the group consisting of: external removal unit, preferably consisting of an external container and a system for generation of electric, or magnetic, or electromagnetic fields and at least one duct and at least one displacement means for the extraction of concentrated nucleic acids from the preceding system; integrated removal unit, preferably consisting of a system of generation of electric, or magnetic, or electromagnetic fields and at least one duct and at least one displacement means for the extraction of concentrated nucleic acids from the preceding system, the

whole integrated within the same bioreactor; techniques applicable in situ and outside units, preferably centrifugation techniques; filtration techniques; treatments with DNase and other degrading enzymes for nucleic acids; heat treatments; acidifying treatments

5 Preferably, according to the process of the invention, said nucleic acids are removed from the culture medium by means of at least one external or bioreactor or photobioreactor integrated removal unit or systems for hydroponic culture using techniques selected from the group consisting of application of static electric fields, application of static magnetic fields,
10 application of dynamic magnetic and/or electric fields, centrifugation, filtration, treatment with DNase, heat treatment, acidifying treatment.

A system for production of microorganisms comprising at least one bioreactor or photobioreactor or system for hydroponic culture, at least one unit for nucleic acid removal external to said at least one bioreactor or
15 photobioreactor or system for hydroponic culture is an ulterior aspect of the present invention, said unit for nucleic acid removal being connected to said at least one bioreactor or photobioreactor or system for hydroponic culture by means of at least one first duct and at least one first displacement means for the withdrawal of culture medium containing acid
20 nucleic, and a at least one second duct and at least one second displacement means for the re-introduction in said at least one bioreactor or photobioreactor or system for hydroponic culture the culture medium from which the nucleic acids are removed; said system comprising further at least one third displacement means for the withdrawal of separated
25 nucleic acid. Preferably said duct can be a tube. Said displacement means can be preferably a pump.

The reclamation of culture medium is realizable both continuously using treatments for removal of the DNA from culture medium during the growth stage and carrying out a treatment of exhaust medium when saturation
30 stage of the cellular growth occurs.

The reclamation of culture medium can be carried out "in situ" directly in the bioreactors or photobioreactors using treatments with enzymes or other compounds active for the degradation of the DNA occurring in the

supernatant of the cellular cultures, or using techniques for application of electric, magnetic or electromagnetic fields. The reclamation using these techniques proves to be viable taking advantage of the polarity characteristics of nucleic acid molecules therefore separable by electric fields, or using techniques for magnetization of DNA molecules by binding the same to magnetic molecules and applying magnetic fields for the separation thereof. Alternatively, the reclamation can be carried out using a removal system placed outside the bioreactor (Figure 12 A) or hydroponic culture (Figure 13A) with re-circulation of regenerated medium or a bioreactor (Figure 12 B) or hydroponic culture integrated (Figure 13B) removal system.

The techniques for removal of the DNA from culture medium can be: techniques for application of static electric fields (FIGURE 14); techniques for application of static magnetic fields (FIGURE 15); techniques for application of dynamic magnetic and/or electric fields; centrifugation; filtration techniques; treatment with DNase; heat treatments; acidifying treatment.

In a preferred embodiment of the invention (figure 12 A), said at least one bioreactor (1) can be connected to said at least one external removal unit (4) in such a way that the exhaust culture medium is transferred in at least said a removal unit (4) by means of at least one first duct (2) and at least one first displacement means (3) for the withdrawal of culture medium containing nucleic acid. In such unit (4), the reclamation can be carried out using one of the aforesaid techniques for DNA removal from medium. At this point the separated nucleic acid will be collected by the removal unit (4), preferably by means of at least one third duct (5) and at least one third displacement means (6) for the withdrawal of separated nucleic acid, whereas the regenerated medium will be re-introduced in said at least one bioreactor (1) by means of at least one second duct (7) and at least one second displacement medium (8).

Further the integration of the fertirrigation equipments in open field or protected culture with systems for removal of nucleic acids represents a further application of the present invention. A possible diagram of such

application is reported in Figure 16 wherein a tank with a solution containing nuclease is integrated in a fertirrigation equipment.

The use of total DNA random fragments for the control and inhibition the parasitic, infesting and/or pathogenic species in agriculture and medicine, compared to the existing techniques and products, results in significant advantages.

A first advantage results from the absolute selectivity of the treatments with total DNA random fragments of the parasitic, infesting and/or pathogenic species, because said total DNA random fragments do not display effects on species different than treated ones.

The use of total DNA random fragments of the parasitic, infesting and/or pathogenic species does not induce the appearance of resistance in the same species populations, because action mechanism thereof is different than that of the currently marketed or under experimentation products (including the methods of RNA interference and ASO), as based on a multiple and contemporary interference with the entire genome with the entire transcriptional and protein synthesis system, thus not allowing specific resistance and selection of resistant populations to be set up.

A further advantage results from the absence of toxicity caused from the nucleic acid use, that, being primary metabolites, ubiquitous in nature, do not result in environmental contamination. In fact, the administration of nonspecific nucleic acid fragments displays no toxicity for species other than treated ones.

The present invention affords further a remarkable simplification of the research for new drugs with consequent great economic benefit. In fact, the detection of parasitic, pathogenic and infesting species constitutes the only condition necessary for the preparation of the specific drug for the pharmacological treatment. On the contrary, the current pharmacological research for products based on techniques like RNA interference or use of antisense oligonucleotides, require not only the knowledge of the full genomic sequence of the target organism, but also the detailed knowledge of the gene functionalities in order to detect possible specific sequences to be targeted for inhibiting activity.

On the other hand, the traditional pharmacological research is based on "random screening" studies of synthetic compounds or naturally occurring molecules. Such substances, once detected and tested for their activity, in the majority of the cases are characterized also by a general toxicity for other organisms and humans. Practically, the application of such products often is associated to environmental contamination concerns due to the toxicity thereof.

A further advantage results from the relative easy availability and bulk production of nucleic acid meeting the requirements of title patent. In fact, random fragments of the total DNA of the target organism can be obtained by approaches and methodologies now extremely diffused in biomolecular field. The techniques can be based on extraction of the genetic material from the target organisms, or by means of synthesis.

Further, the stability of DNA fragments can assure the persistence over the time also under unfavourable environmental conditions (e.g. in the soil), resulting therefore, in durable applications resulting in both environmental and economic benefits.

A further advantage results from the fact that the DNA for the proposed uses can be obtained by extraction from naturally occurring material or by synthesis based on synthesis and extraction techniques and amplification as well, now widely diffused and established in molecular biology. Significant but not restrictive examples of possible production techniques include the dissolution of the cellular tissues, inactivation of the cellular nucleases and recovery of nucleic acids from the solution containing biological lysates.

Nucleic acids are further synthesizable starting from template molecules or de novo according to different approaches.

The considered molecules can be amplified, thus producing multiple copies identical or similar to the considered template, for example by cloning or PCR based techniques.

PCR variants are useful in order to produce the active principle to be used assuring random amplification of the nucleic acid fragments from a template sample. For example, an approach like Random Amplified

Polymorphic DNA (RAPD) is very suitable to our end not requiring, otherwise than the classic PCR, knowledge of the starting DNA template sequence.

5 Emulsion PCR represents another example of innovative approach suitable to assure in an extremely quick way the amplification of various DNA fragments starting from a genomic DNA sample atomized so as to obtain 300 - 800 nucleotide long fragments, or from amplicons obtained by other approaches.

10 Further, as to the production processes of microorganisms in bioreactors and photobioreactors and tanks for hydroponic culture the removal of exhaust culture medium and the re-suspension of the cells in "regenerated" medium, at equal cellular concentration for volume unit and under the same contour conditions, allow the restoring of the culture growth stage.

15 In addition to the advantage of the production increase resulting from the removal of the inhibiting factor, a further remarkable advantage results from the possibility of re-use of exhaust medium, after regeneration thereof, for other successive cultures, possibility not viable in the current systems of production.

20 The present invention now will be described, illustrative, by an illustrative but not limitative way, with particular reference to embodying examples and enclosed drawings, wherein:

Figure 1 shows a conceptual view of the object of the invention.

25 Figure 2 shows the inhibition of the germination of seeds of *Lepidium* and *Acanthus* by exposure to self DNA at optimal concentration of 200 ppm.

Figure 3 shows the inhibition of *Quercus ilex*, *Quercus pubescens*, *Hedera elix*, *Ampelodesma mauritanica*, *Festuca drimeja*, *Coronilla emerus*, *Medicago marina*, *Alnus cordata*, *Robinia pseudoacacia*, *Pinus halepensis* plants by exposure to self DNA.

30 Figure 4 shows the inhibition of *Arabidopsis thaliana* plants by exposure to self and *Lycopersicon esculentum* DNA.

Figure 5 shows the inhibiting effect on the spore germination and hyphal growth of *Aspergillus niger* when said fungus is exposed to self or other

fungus species DNAs (*Trichoderma hartianum*).

Figure 6 shows the inhibition of *Sarcophaga carnaria* insect by exposure to self DNA.

5 Figure 7 shows the completed metamorphoses, in percentage compared to not exposed control, in larvae of *Sarcophaga carnaria* dipter exposed for 4 weeks to homologous DNA at three different concentrations, or fungus (*Penicillium chrysogenum*) or bacterium (*Bacillus subtilis*) extracted heterologous DNA.

10 Figure 8 shows the viable cell counts in *Bacillus subtilis* bacterium in cultures exposed for 24 hours to homologous DNA at three different concentrations and in not exposed control.

15 Figure 9 shows the germination of the spores in the *Trichoderma harzianum* fungus, in percentage compared to not exposed control, in cultures exposed to homologous DNA at three different concentrations, or to heterologous DNA extracted from other fungus (*Aspergillus niger*), insect (*Sarcophaga carnaria*) or bacterium (*Bacillus subtilis*).

Figure 10 shows growth dynamics of *Scenedesmus obliquus* microalga in two cultures exposed to homologous DNA at different concentration, and not exposed control.

20 Figure 11 shows the accumulation of extracellular DNA in the liquid substrate of two different bioreactors.

25 Figure 12 shows: (a) the diagram of system for production of microorganisms in bioreactor characterized from an external unit for DNA removal; (b) the diagram of system for production of microorganisms in bioreactor wherein the removal of DNA from culture medium occurs by bioreactor integrated removal unit.

30 Figure 13 shows: (a) a tank for hydroponic culture characterized from the presence of an external removal DNA unit and recirculation of culture medium within the hydroponic system; (b) a tank for hydroponic culture wherein the removal of the DNA from culture medium occurs by means of removal unit integrated to the same tank.

Figure 14 shows a specific diagram of the external removal unit for the system represented in figure 12, wherein the separation of nucleic acid

occurs by application of an electric field.

Figure 15 shows a further diagram of the removal unit for the system showed in figure 12, wherein the separation of nucleic acid occurs by application of a magnetic field.

5 Figure 16 shows an diagram of plant culture wherein the substrate is treated with nuclease by integration with fertirrigation system.

In all the examples of reported experiments, the nucleic acid composition used in the different treatments was prepared according to the procedure outlined in Figure 1. In particular, the total DNA extracted with standard
10 procedures from organic material (leaves, fungus mycelium, microbial biomass) was treated by sonication for at least three cycles lasting three minutes at maximum power with dipping sonicator until to obtain the production of composition of random fragments falling in 50 and 1000 kb size range. The verification of the fragmentation level is carried out by
15 standard procedures using agarose or polyacrylamide gel electrophoresis and staining techniques, Sybr safe type and UV visualization.

EXAMPLE 1: Inhibition of *Acanthus mollis* and *Lepidium sativum* plants by exposure to self DNAs

A first experiment was carried out on *Acanthus mollis* and *Lepidium sativum* plants, the latter species was selected because it is particularly
20 sensitive to toxins. DNAs of acanthus, *Acanthus mollis*, and watercress, *Lepidium sativum*, were obtained by direct extraction from leaves of the two species and stored in distilled H₂O. Successively 10 previously sterilized seeds of *A. mollis* or *L. sativum*, in Petri plates (9 cm diameter)
25 are placed on a sheet of sterile filter paper. The seeds of each species are treated separately with the DNA of the two species at concentrations of 2, 20 and 200 ppm whereas sterile H₂O was added to the control. The germination of the two species and the total root length were quantified after 7 days of incubation at 24 °C by observation and measurement of the
30 roots. Each treatment was repeated thrice.

The treatment of the seeds of the two species with DNA extracted from plants of the two species, applied separately, allowed to estimate the effect of the DNA on the root growth and the optimal activity concentration.

The results of the experimentation, reported in Figure 2, show that both seeds of *Lepidium* and seeds of *Acanthus* are inhibited in the germination by the exposure to self DNA at optimal concentration of 200 ppm. On the contrary, the exposure of seeds to DNA from other species does not show noticeable effects on the seed germination.

EXAMPLE 2: Inhibition of *Quercus ilex*, *Quercus pubescens*, *Hedera elix*, *Ampelodesma mauritanica*, *Festuca drimeja*, *Coronilla emerus*, *Medicago marina*, *Alnus cordata*, *Robinia pseudoacacia*, *Pinus halepensis* plants by exposure to self DNAs

A second experiment concerned the analysis of the germination and root growth of 10 species of natural environment plants. Surface sterilized seeds of *Quercus ilex*, *Quercus pubescens*, *Hedera elix*, *Ampelodesma mauritanica*, *Festuca drimeja*, *Coronilla emerus*, *Medicago marina*, *Alnus cordata*, *Robinia pseudoacacia*, *Pinus halepensis* plants, were separately treated with DNAs of all the species applied at the concentration of 500 ppm. Shortly, in Petri plates (9 cm diameter) are placed 10 seeds of each species on a sheet of sterile filter paper. The different DNAs at concentration of 500 ppm were added to the plates whereas only sterile H₂O was added to the control. The germination of the species seeds and the total root length were quantified after 7 days of incubation at 24 °C by observation and measurement of the roots. Each treatment was repeated thrice. The results, reported in Figure 3 (average of the tests carried out on the above reported different species), shows the inhibition of the germination resulting from the exposure to self DNA and the absence of inhibition in the presence of heterologous DNA.

EXAMPLE 3: Inhibition of *Arabidopsis thaliana*, *Lycopersicon esculentum*, *Lepidium sativum* and *Lens esculentum* plants by exposure to self DNAs

A third experiment concerned the evaluation, again by germination and root growth tests, of the toxicity on various plant species by nucleic acid extracted from same species. DNAs of *Arabidopsis thaliana*, *Lycopersicon esculentum*, *Lepidium sativum* and *Lens esculentum* were obtained by direct extraction from the respective plants and stored in distilled H₂O. Successively 10 previously sterilized seeds of each species are placed in

Petri plates (9 cm diameter) on a sheet from sterile filter paper. The seeds of each plant are treated separately with the DNAs of the four species at concentrations of 2, 20 and 200 ppm whereas sterile H₂O was added to the control. The experiments were carried out in growth rooms under controlled conditions and complete sterility. The germination of the four species and the total radical length were quantified after 7 days of incubation at 24°C by observation and measurement of the roots. Each treatment was repeated thrice. The four species shown an analogous behaviour, with a remarkable inhibiting effect in the presence of self DNA and the absence of effects in the presence of DNAs of the other species. The inhibiting effect proved to be positively correlated to the DNA concentration. For exemplary purpose *Arabidopsis thaliana* in the presence of self and *Lycopersicon esculentum* DNAs, respectively, are reported. Similar inhibition results in *Arabidopsis thaliana*, when seeds of this species are exposed to DNA of the same species obtained by amplification of fragments of the same DNA using PCR techniques, are observed.

EXAMPLE 4: Inhibition of *Aspergillus niger* and *Trichoderma harzianum* fungi by exposure to self DNAs

A fourth experiment was carried out on the *Aspergillus niger* fungus in order to estimate the effect of self DNA and DNA isolated from another fungus, i.e. *Trichoderma harzianum*, on the cellular growth. Spores of *Aspergillus niger* are obtained by pure cultures in laboratory on agar treated substrate (PDA, potato dextrose agar). The spores were withdrawn under sterility conditions and diluted at concentration of 1×10^6 spores/ml. The experiment of germination was carried out in liquid substrate (PDB 10%) in 96 well ELISA plates. The comparative treatment was carried out with DNA of *Trichoderma harzianum*, used as heterologous, whereas the control was not treated. DNAs extracted from both the species were applied at concentrations of 100 and 1000 ppm. Shortly, in each well, with a total volume of 100 μ I, said two DNAs separately at different concentrations, together with 10 μ I of liquid nutritive substrate (PDB, potato dextrose broth), sterile water and *A. niger* spores were added. The

germination of the spores and the length of the germinative tube were quantified by spectrophotometric readings and optical microscope after 20 hours of incubation at 24°C. The results, reported in Figure 5, show a remarkable inhibiting effect on the germination of the spores and the hyphal growth of *A. niger* only when such fungus was exposed to self DNA.

EXAMPLE 5: Inhibition of *Sarcophaga carnaria* insect by exposure to self DNA.

A fifth experiment was carried out on *Sarcophaga carnaria* insect in order to estimate the effect of self DNA on the life-cycle. Larvae of *Sarcophaga carnaria* dipter were grown in laboratory pure culture at the temperature of 10°C and fed with minced meat. The experiment of DNA toxicity was carried out in square plastic plates (size 12 x 12 cm, height 2 cm). The comparative treatments were carried out with DNAs of *Bacillus subtilis* and *Lepidium sativum*, used as heterologous DNA. As control only minced meat without addition of other treatments was used. Dipter and other two species extracted DNAs were added to the minced meat at concentrations of 2, 20 and 200 ppm, under mixer stirring. Shortly, in each plate DNA was added at the various concentrations, stirred with 1 g of minced meat. The plates were incubated at 10°C in the dark for 21 days. The development, the survival and the time required for the formation of the pupae are monitored every 3 days for the 21 days of incubation. The larvae under control conditions, as well as those treated with heterologous DNA, displayed a regular life-cycle. On the contrary, the exposure to self DNA inhibited the life-cycle causing the death of the larvae proportionally to the treatment concentration. Figures 6 and 7 report the above described results.

EXAMPLE 6: Inhibition of the *Bacillus subtilis* microorganism by exposure to self DNA

In order to demonstrate the possible use of nucleic acid as antibiotics an evaluation of toxicity on *Bacillus subtilis* treated with self DNA at various concentrations was carried out. The experiment was performed using as growth substrate 4 ml of LB (Luria Broth) inoculated with 10 µl of *Bacillus*

subtilis preculture. The treatment consisted in the preparation of the cultures in the presence of *Bacillus subtilis* DNA at final concentrations of 4, 40, and 400 ppm. The cultures were incubated under stirring at 35 °C for 24 h with three repeats of treatment. After 24 hours of incubation, from each test tube 0,5 ml was withdrawn and serially diluted in LB medium, from which 100 microliters of agar treated LB medium in Petri plates were plated. The plates were incubated at 28 °C until the appearance of colonies (CFU - colonies forming units). The results reported in Figure 8 show a remarkable concentration-dependent decrease of CFUs, as response to the treatment.

EXAMPLE 7: Inhibition of *Trichoderma harzianum* fungus by exposure to self DNA

In order to demonstrate the possible use of nucleic acids as fungicide and action specificity thereof an experiment on the germination of the spores in the *Trichoderma harzianum* fungus was carried out. Spores of *Trichoderma harzianum* were obtained by pure laboratory cultures on agar treated substrate (PDA, potato dextrose agar). The spores were withdrawn under sterility conditions and diluted at concentration of 1×10^6 spores/ml. The experiment of germination was carried out in liquid substrate (PDB 10%) in 96 well ELISA plates. The treatment was carried out with homologous or heterologous DNA, that is extracted from the same species of *Trichoderma*, or from a different species of fungus (*Aspergillus niger*), from an insect (*Sarcophaga carnaria*) or from a bacterium (*Bacillus subtilis*). The DNA extracted from the different species was applied at concentrations of 8, 80 and 800 ppm. Shortly, in each well, with a total volume of 100 μ l, DNA separately at different concentrations, together with 10 μ l of liquid nutritive substrate (PDB, potato dextrose broth), sterile water and *Trichoderma* spores were added. The germination of the spores and the length of the germinative tube were quantified by spectrophotometric readings and optical microscope after 20 hours of incubation at 24 °C.

Figure 9 indicates the results of the experiment showing a remarkable concentration-dependent inhibiting effect, on the germination of the *Trichoderma* spores only by DNA of the same fungus species. On the

contrary, the treatment with various species DNAs display stimulating effects on the germination (percentage values compared to not exposed control higher than 100%).

EXAMPLE 8: Inhibition of the *Scenedesmus obliquus* microalga by exposure to self DNA

5 In order to demonstrate the possible DNA use as algicide product a growth test of *Scenedesmus obliquus* green alga under optimal control conditions and self DNA presence was carried out in the culture substrate (CHU#10). The treatments were carried out at two different concentrations (50 and
10 500 ppm) with two repeats. Figure 10 shows the growth dynamics of the alga and shows a remarkable concentration-dependent inhibiting effect of the homologous DNA compared to not exposed control.

EXAMPLE 9: Inhibition of the *Physarium polycephalum* protozoan by exposure to self DNA

15 In order to demonstrate the possible DNA use as antiprotozoic product an experiment was carried out on *Physarium polycephalum* protozoan. As experimental material the culture kit produced from "Carolina Biological Supply" was used. The cultures started on Petri plates with water-agar in order to favour the movement of the organism. As nutriment oat flakes
20 were used. A first bulk culture was carried out on 20 plates and protozoic biomass produced after 15 days was collected and used for DNA extraction using Quiagen kit. The successive experiment consisted of the preparation of three Petri plates filled up with water agar to which two small portions of oats flakes were added, one for control wetted with 5 ml
25 of distilled water and the other with addition of 5 ml of water with DNA of the protozoan at concentration of 200 ppm. The experiment was repeated other two times, with the variation that the added DNA was from bacterium (*Bacillus subtilis*) and insect (*Sarcophaga carnaria*). The results of the experiments shown the absence of growth of *Physarium polycephalum* on
30 the substrate treated with DNA of the same protozoan whereas the organism did not shown growth differences under control conditions or heterologous DNA presence.

EXAMPLE 10: Study of production process of yeasts, bacteria and algae

in bioreactors and photobioreactors

Considering the above reported demonstrations about the inhibiting effect on different species when exposed to self DNAs, check analyses were carried out on the extracellular DNA presence in the growth substrate in bioreactors and photobioreactors with cellular cultures at high density when conditions of growth slowing down, even if under optimal nutritive substrate presence, occur. The study involved the sampling of liquid supernatant of different cultures in bioreactors in exponential growth, slowing down and steady-state stages. The analysis concerned cultures of *Saccharomyces cerevisiae* yeast, *Bacillus subtilis* bacterium and *Phaeodactylum tricornutum* and *Scenedesmus obliquus* microalgae. Samples of the cellular culture supernatants obtained by two cycles of centrifugation at 3000 rpm for 15 minutes were analyzed in order to separate possible cell residues and then subjected to gel electrophoresis after treatment with Syber-Safe and fluorescence evaluation. Figure 11 shows the results of some of these analyses from which it is apparent the accumulation of extracellular DNA in the liquid substrate of the bioreactors. This accumulation is clearly associated to the slowing down of the growth of the different cellular cultures and to the attainment of the steady-state stage (Figure 11A and 11B). Figure 11C shows clearly as the removal of the extracellular DNA from culture medium by chemical-physical procedures and the following introduction of regenerated substrate into the reactor results in the elimination of the inhibiting effect and a consequent restoring of the cellular culture growth.

29
CLAIMS

1. Use of a composition comprising or consisting of a mixture of DNA fragments for the prevention and the treatment against at least one pathogenic, parasitic or infesting species of the plants or environment,
5 wherein said DNA is the total DNA of said at least one pathogenic, parasitic or infesting species, and/or at least one phylogenetically similar species, against which said prevention and treatment are directed.
2. Use according to claim 1, wherein said DNA fragments are obtained by random fragmentation.
- 10 3. Use according to each of preceding claims, wherein said total DNA is extracted from said at least one pathogenic, parasitic or infesting species or artificially synthesized.
4. Use according to each of preceding claims, wherein said total DNA is amplified and/or fragmented by physical chemical procedures and
15 obtained fragments are possibly amplified.
5. Use according to each of preceding claims, wherein said composition is used as biocide, herbicide, fungicide, insecticide, acaricide, nematocide, antiprotozoic, algaecide, bactericide.
6. Use according to each of preceding claims, wherein said
20 composition is administered to said at least one pathogenic, parasitic or infesting species by surface contact, cytotropic administration, systemic administration.
7. Use according to claim 6, wherein the systemic administration is carried out by injection, ingestion or inhalation.
- 25 8. Use according to each of preceding claims, wherein said composition is formulated in a form selected from the group consisting of dispersion, suspension, wettable or soluble powders, emulsions in water or other solvents, dispersible granules, suspensions of microcapsules, emulsifiable concentrates, fluid pastes, macro emulsions, oil dispersions,
30 baits..
9. Use according to each of preceding claims, wherein said total DNA fragments are engineered in vectors.

10. Use according to each of preceding claims, wherein said composition further comprises pesticide compounds selected from the group consisting of fungicides, insecticides, nematocides, miticides, artropocides, bactericidal, algaecides.

5 11. Method for the prevention or the treatment against at least one pathogenic, parasitic or infesting species of the plants or environment, said method comprising or consisting of the following steps of: a) extraction of the total DNA of at least one of said pathogenic, parasitic or infesting species against which said prevention and treatment are
10 directed; b) fragmentation of said total DNA in order to obtain a mixture of DNA random fragments of nucleic acids; and c) contacting said mixture of fragments with said pathogenic, parasitic or infesting species.

12. Composition comprising or consisting of a mixture of DNA fragments for the use in the prevention or treatment against at least one
15 pathogenic, parasitic or infesting species of human or animal, wherein said DNA is the total DNA of said at least one pathogenic, parasitic or infesting species, and/or at least one phylogenetically similar species, against which said prevention and treatment are directed.

13. Composition comprising or consisting of a mixture of DNA fragments according to claim 12 for the use according to claim 12, wherein
20 said DNA fragments are obtained by random fragmentation.

14. Composition comprising or consisting of a mixture of DNA fragments according to any of claims 12-13 for the use according to any of claims 12-13, wherein said pathogenic species is a virus.

25 15. Process for the production of microorganisms in bioreactors or photobioreactors or systems for hydroponic culture characterized in that the nucleic acids of the same microorganisms produced by said process are removed from culture medium and the culture medium deprived of said nucleic acid is used again in said process.

30 16. Process according to claim 15, wherein nucleic acids of the same microorganisms produced by said process are removed from culture medium using separation techniques selected from the group consisting of: external removal unit, preferably consisting of an external container

and a system of generation of electric, or magnetic, or electromagnetic fields and at least one duct and at least one displacement means for the extraction of concentrated nuclei acids from the preceding system; integrated removal unit, preferably consisting of a system of generation of electric, or magnetic, or electromagnetic fields and at least one duct and at least one displacement means for the extraction of concentrated nucleic acids from the preceding system, the whole integrated within the same bioreactor; techniques applicable in situ and outside units, preferably centrifugation techniques; filtration techniques; treatments with DNase and other degrading enzymes for nucleic acids; heat treatments; acidifying treatments.

17. Process according to each of claims 15-16, wherein said nucleic acids are removed from the culture medium by means of at least one external or bioreactor or photobioreactor integrated removal unit or systems for hydroponic culture using techniques selected from the group consisting of application of static electric fields, application of static magnetic fields, application of dynamic magnetic and/or electric fields, centrifugation, filtration, treatment with DNase, heat treatment, acidifying treatment.

18. Process according to each of claims 15-17, wherein said nucleic acids isolated from culture medium are collected and optionally fragmented.

19. Process according to claim 15-18 wherein, when for the separation of nucleic acids an external separating unit is used, the culture medium deprived of said nucleic acids is used again in said process.

20. Process according to each of claims 15-19, wherein said nucleic acids are DNA.

21. Process according to each of claims 15-20, wherein said microorganisms are selected from pathogenic, parasitic or infesting microorganisms.

22. System for production of microorganisms comprising at least one bioreactor or photobioreactor (1) or system for hydroponic culture (11), at least one unit for nucleic acid removal (4) external to said at least one

bioreactor or photobioreactor (1) or system for hydroponic culture (11),
said unit for nucleic acid removal (4) being connected to said at least one
bioreactor or photobioreactor (1) or system for hydroponic culture (11) by
at least one first duct (2) and at least one first displacement means (3) for
5 the withdrawal of culture medium containing nucleic acids, and at least
one second duct (7) and at least one second displacement means (8) for
the re-introduction in said at least one bioreactor or photobioreactor (1) or
system for hydroponic culture (11) of culture medium from which nucleic
acids are removed; said system comprising further at least one third duct
10 (5) and at least one third displacement means (6) for the withdrawal of
separated nucleic acid.

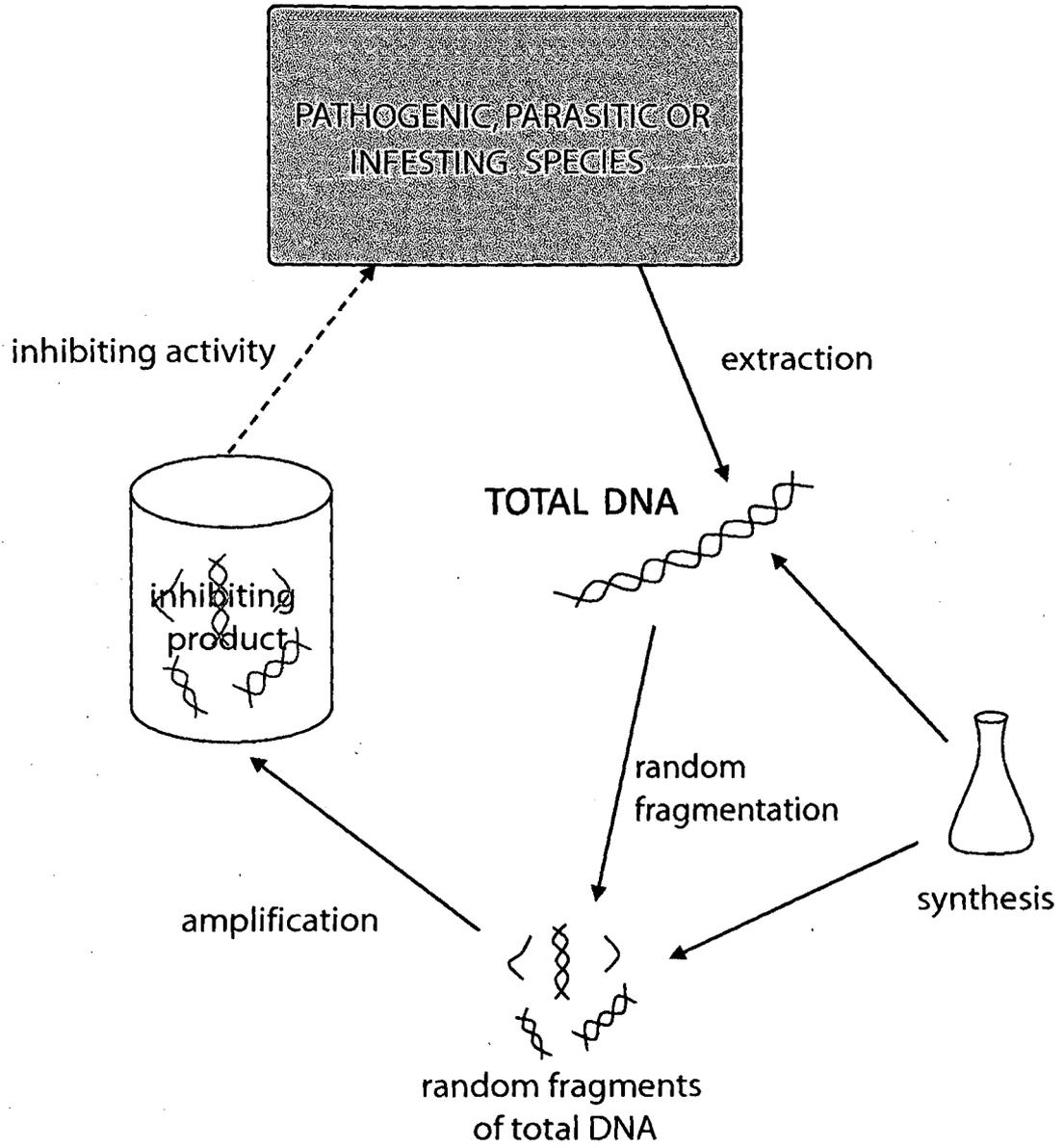


Fig. 1

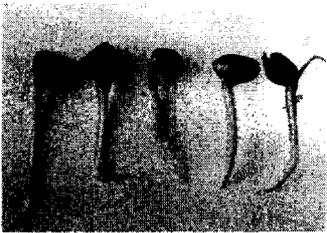
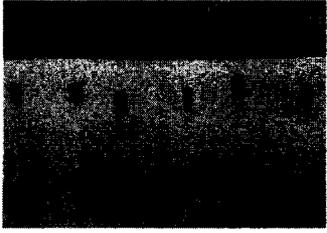
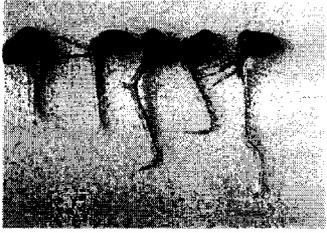
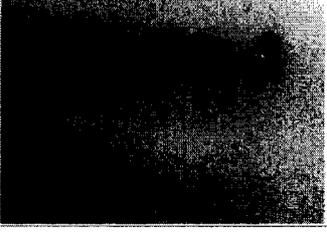
treatment	species	
	<i>Lepidium sativum</i>	<i>Acanthus mollis</i>
control water		
DNA <i>Lepidium</i> (200 ppm)		
DNA <i>Acanthus</i> (200 ppm)		

Fig.2

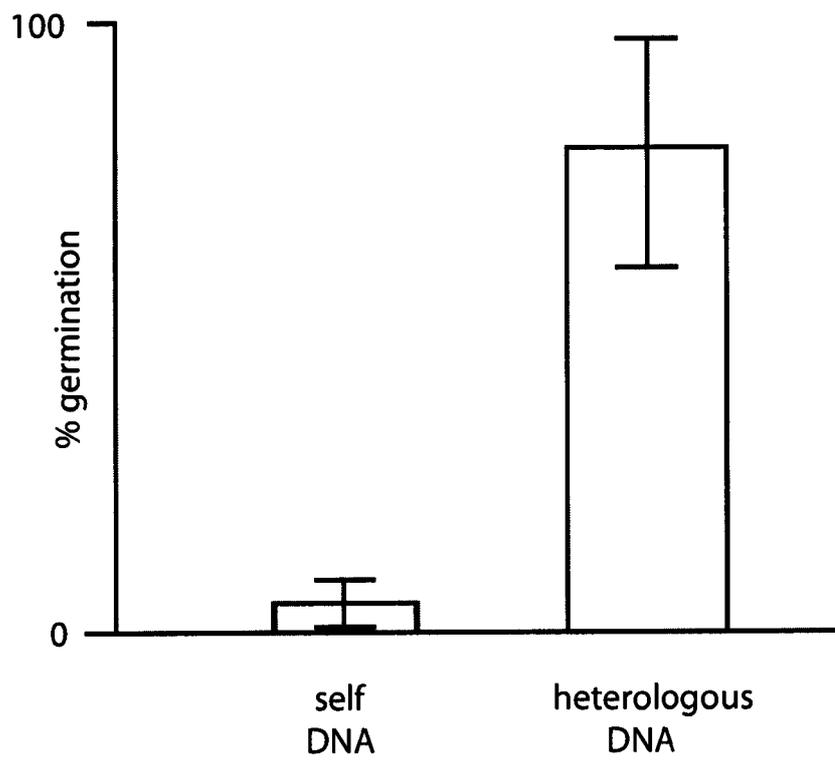


Fig. 3

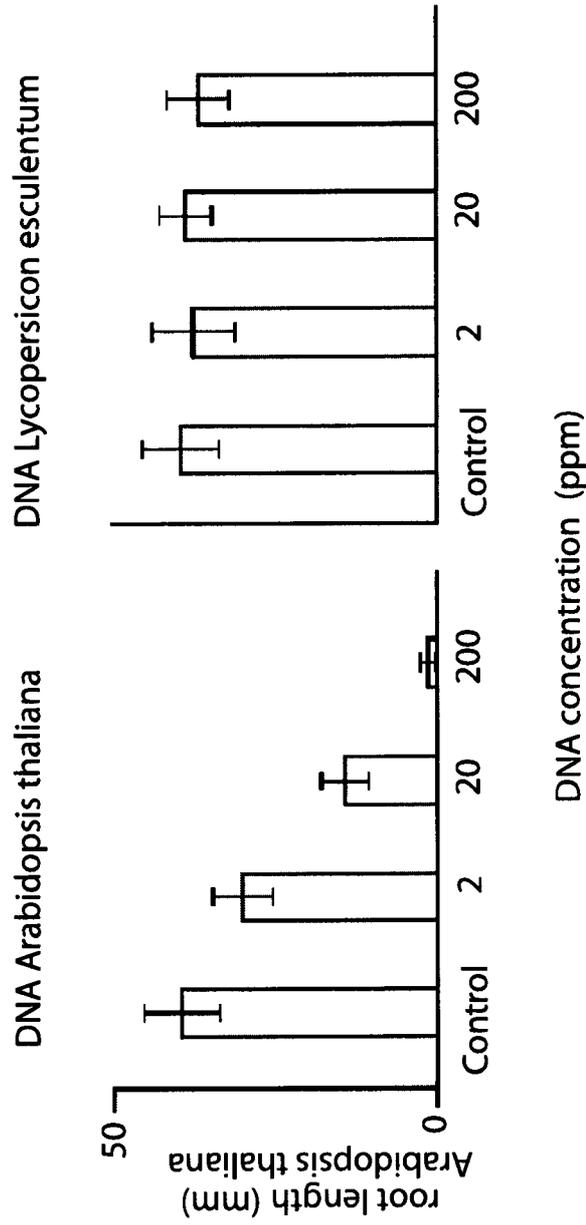
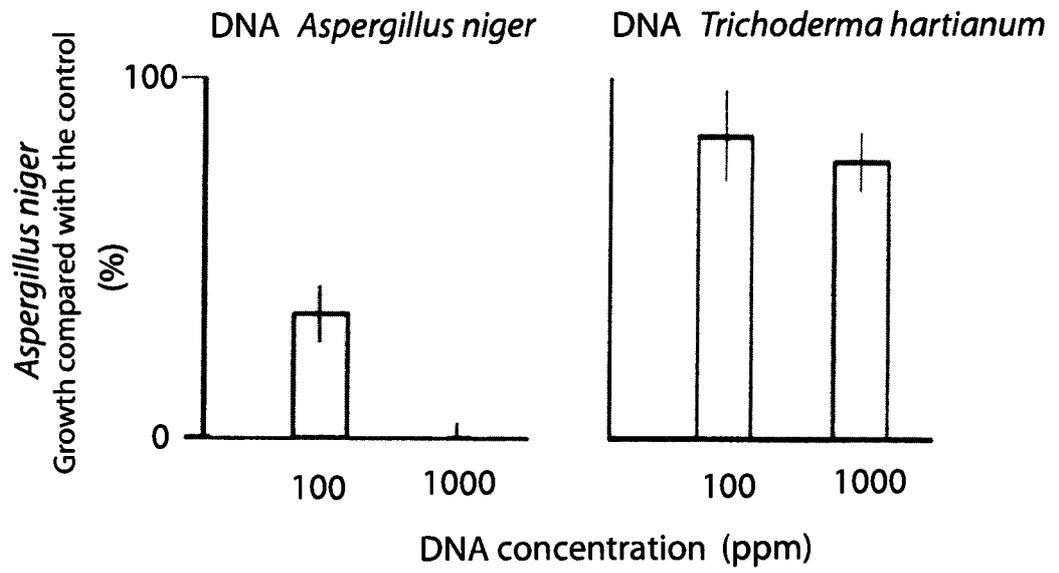


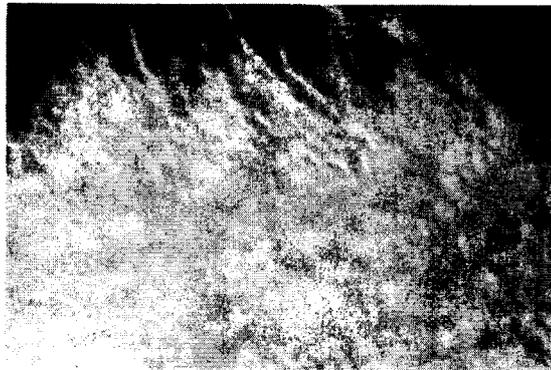
Fig. 4

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Aspergillus niger

Control



DNA *Aspergillus*
(1000 ppm)

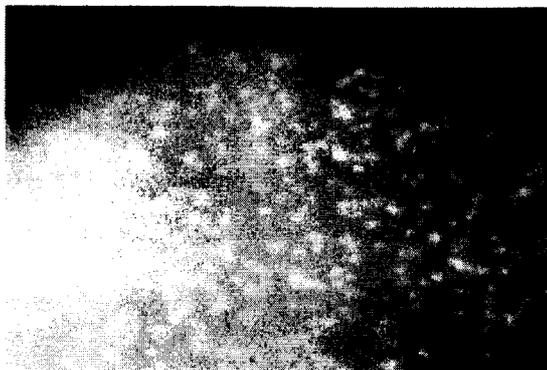


Fig. 5

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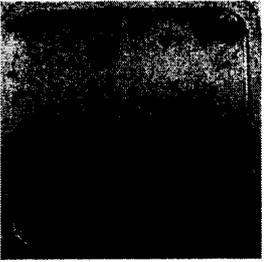
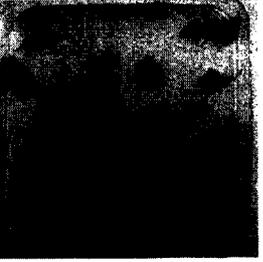
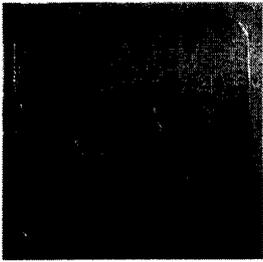
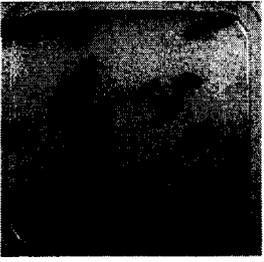
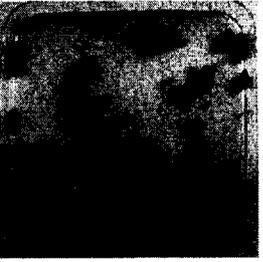
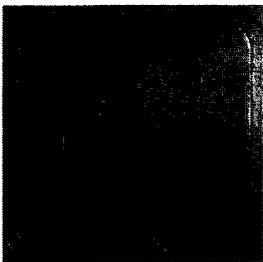
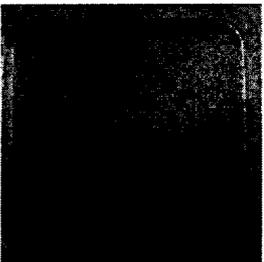
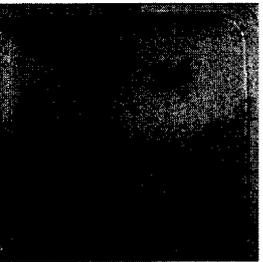
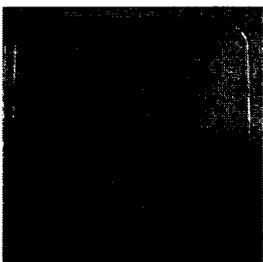
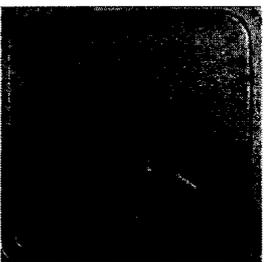
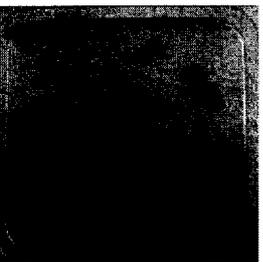
<i>Sarcophaga carnaria</i>	$t = 0$	$t = 10$ days	$t = 20$ days	number of living flies
Control				25
heterologous DNA (200 ppm)				24
self DNA (20 ppm)				2
self DNA (200 ppm)				0

Fig. 6

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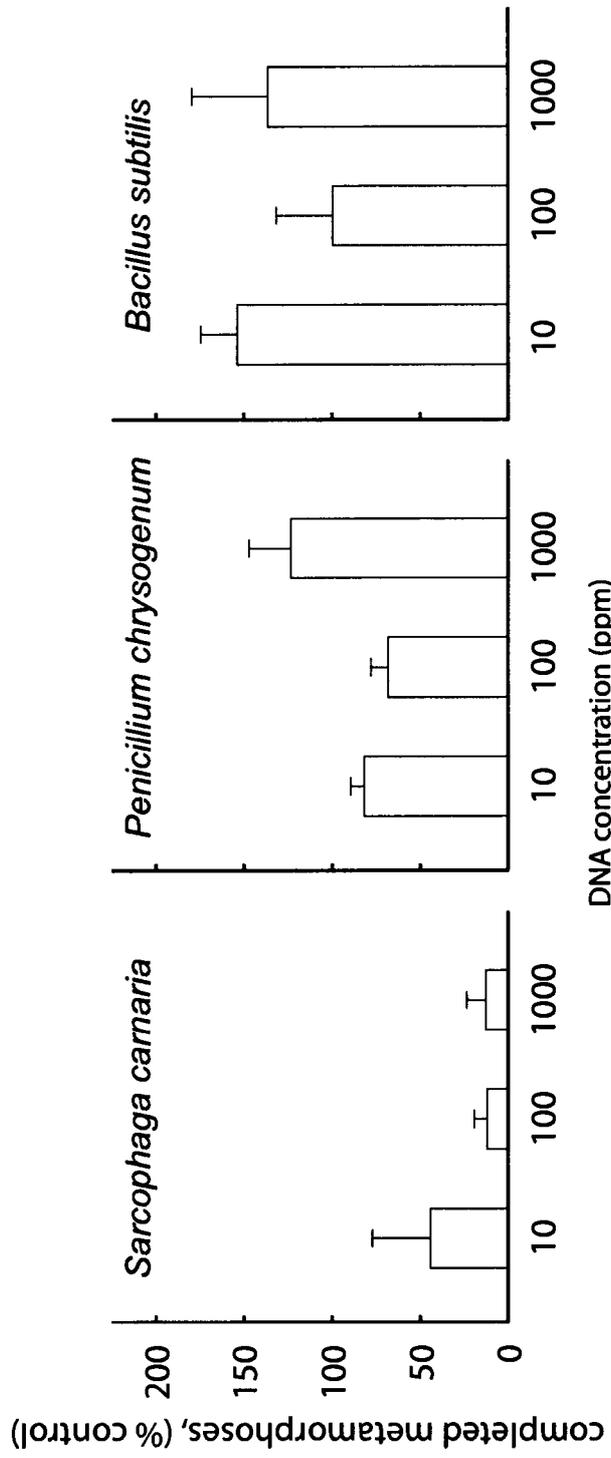


Fig. 7

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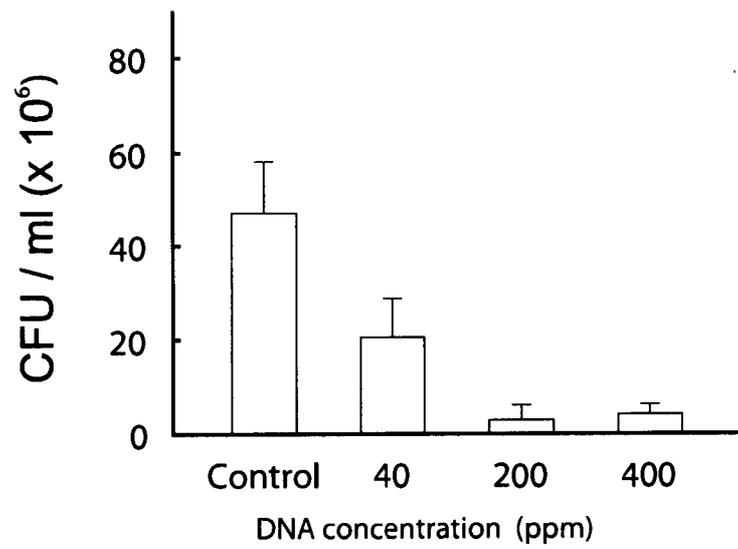


Fig. 8

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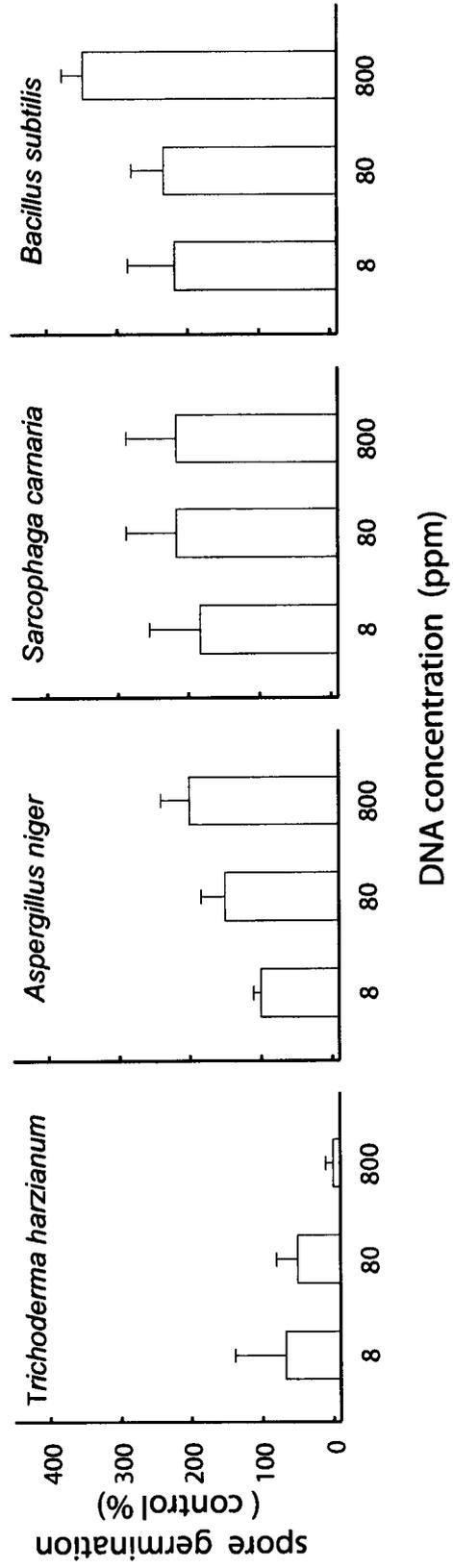


Fig. 9

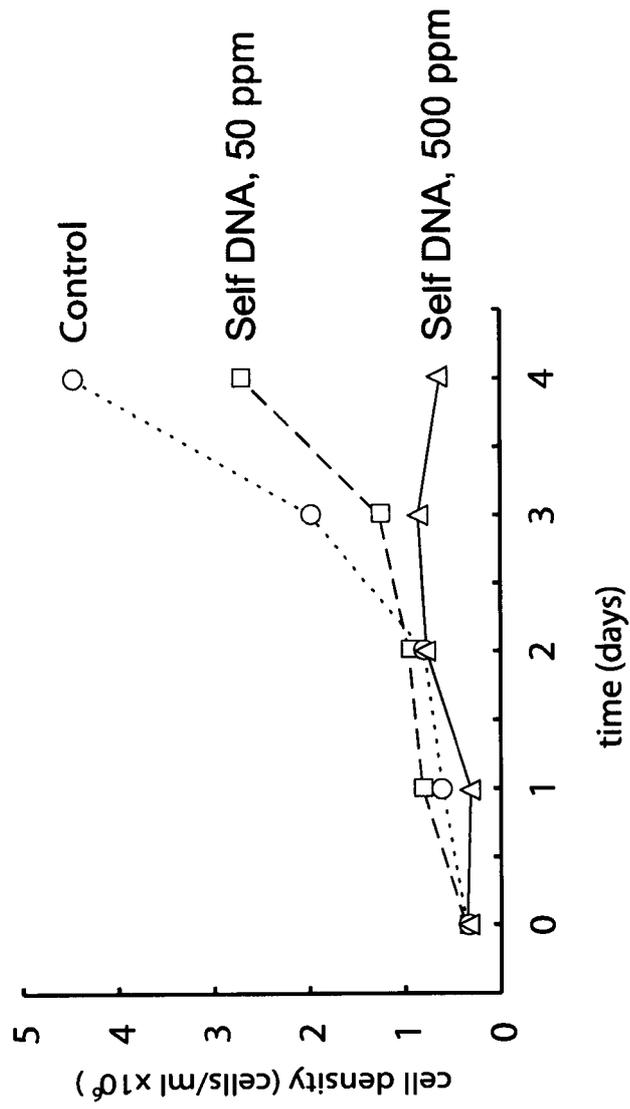
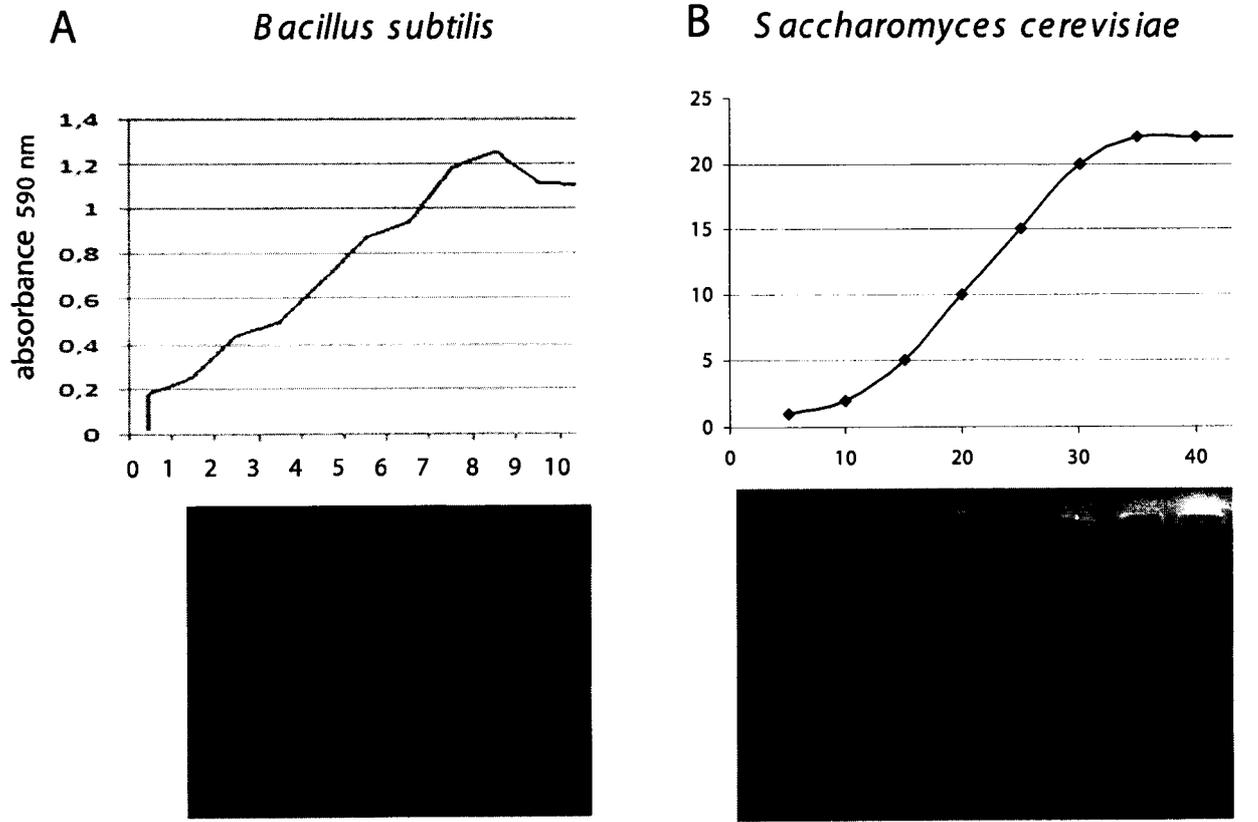


Fig. 10

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Experimental evidences of DNA accumulation in bioreactor supernatant and related inhibiting effect on the growth



fluorescence of supernatant from microbial cultures on 1% agarose gel + SYBR safe

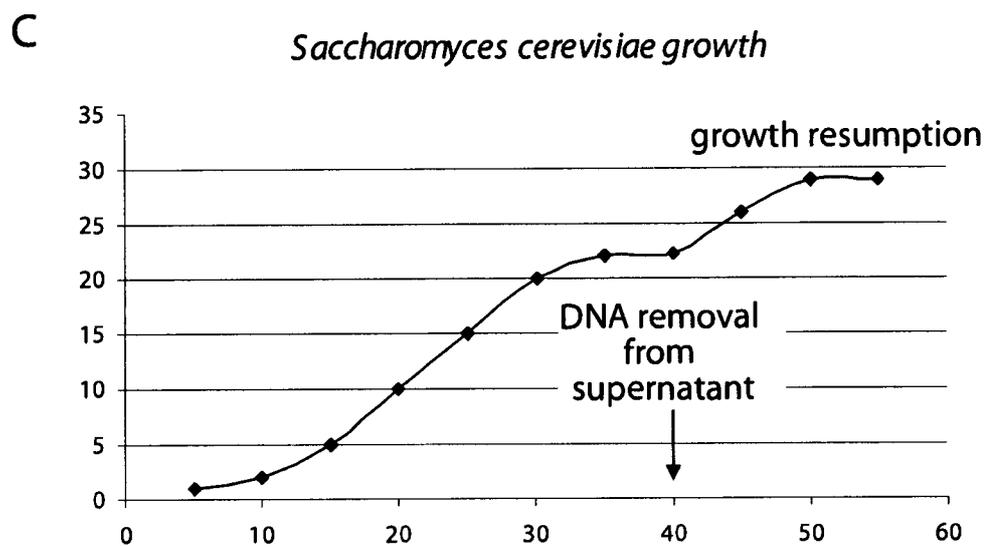


Fig. 11

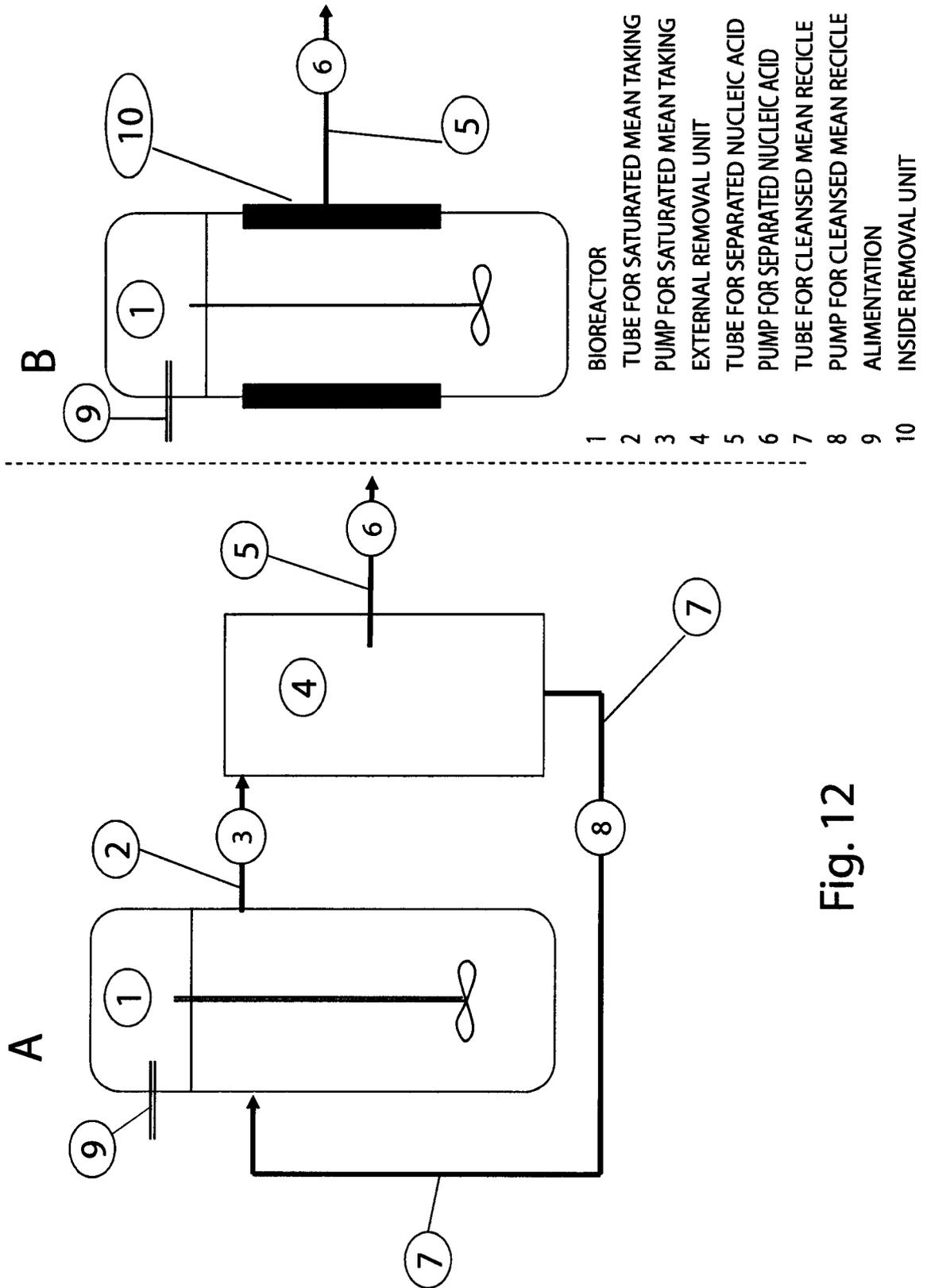


Fig. 12

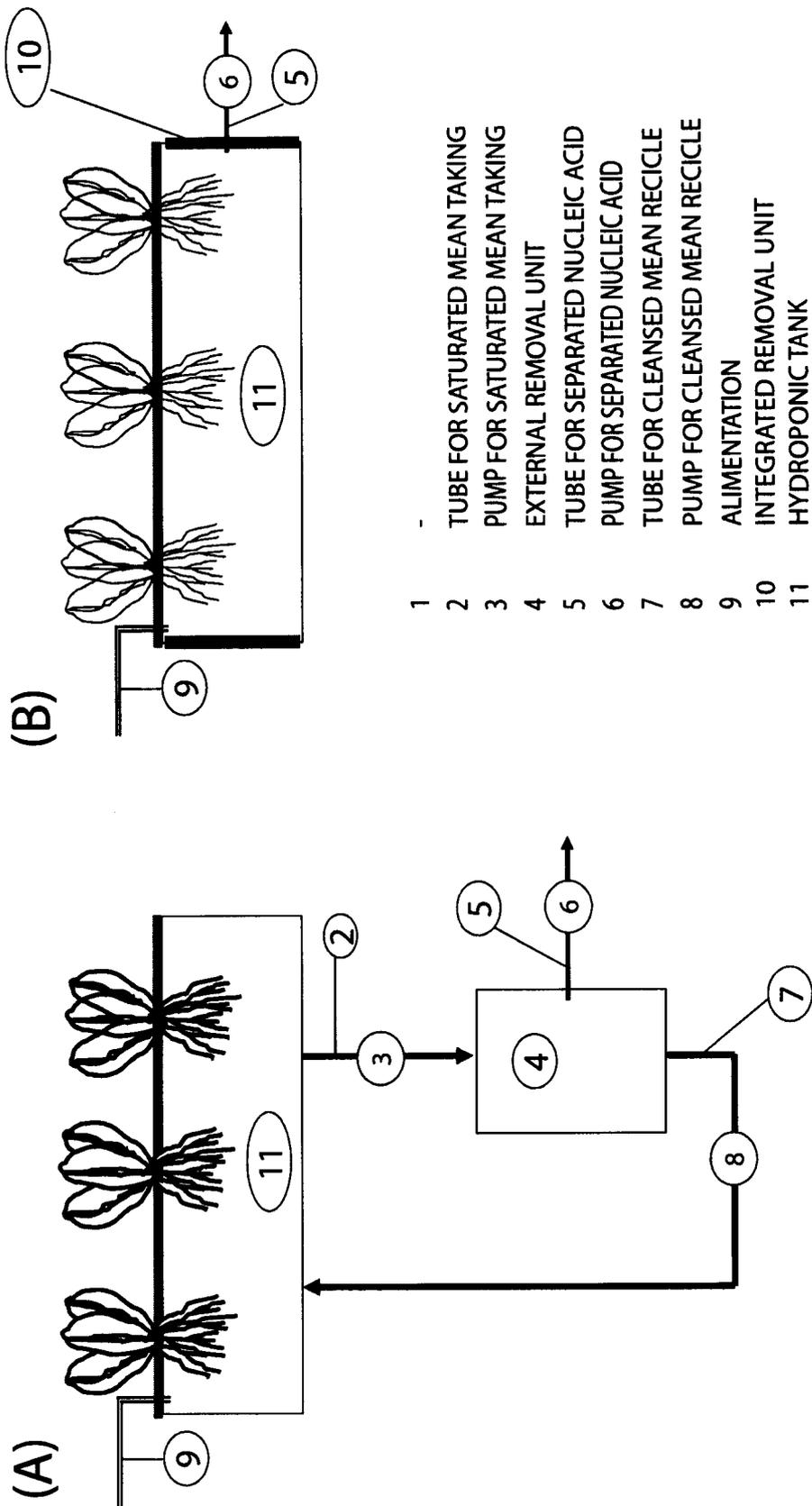
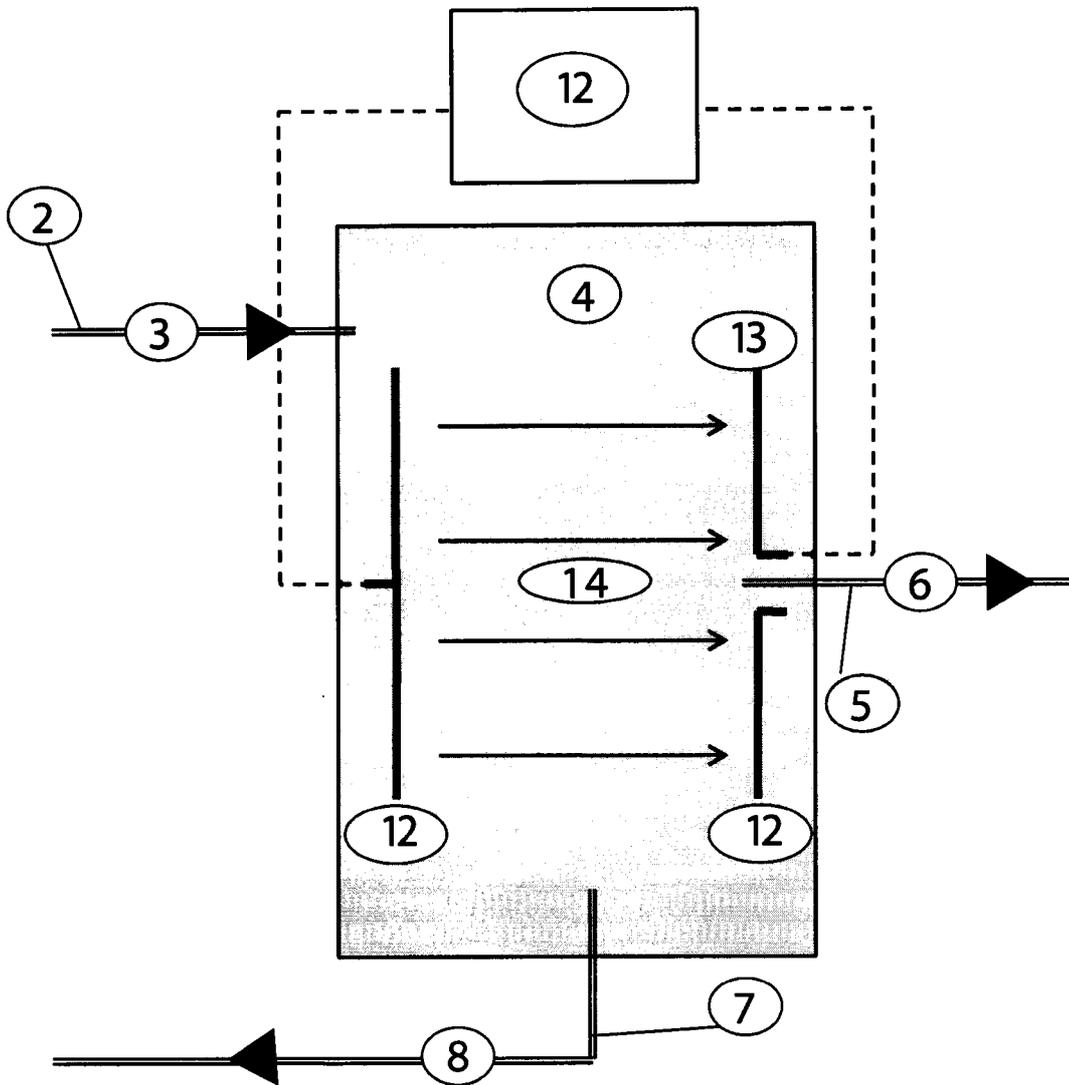


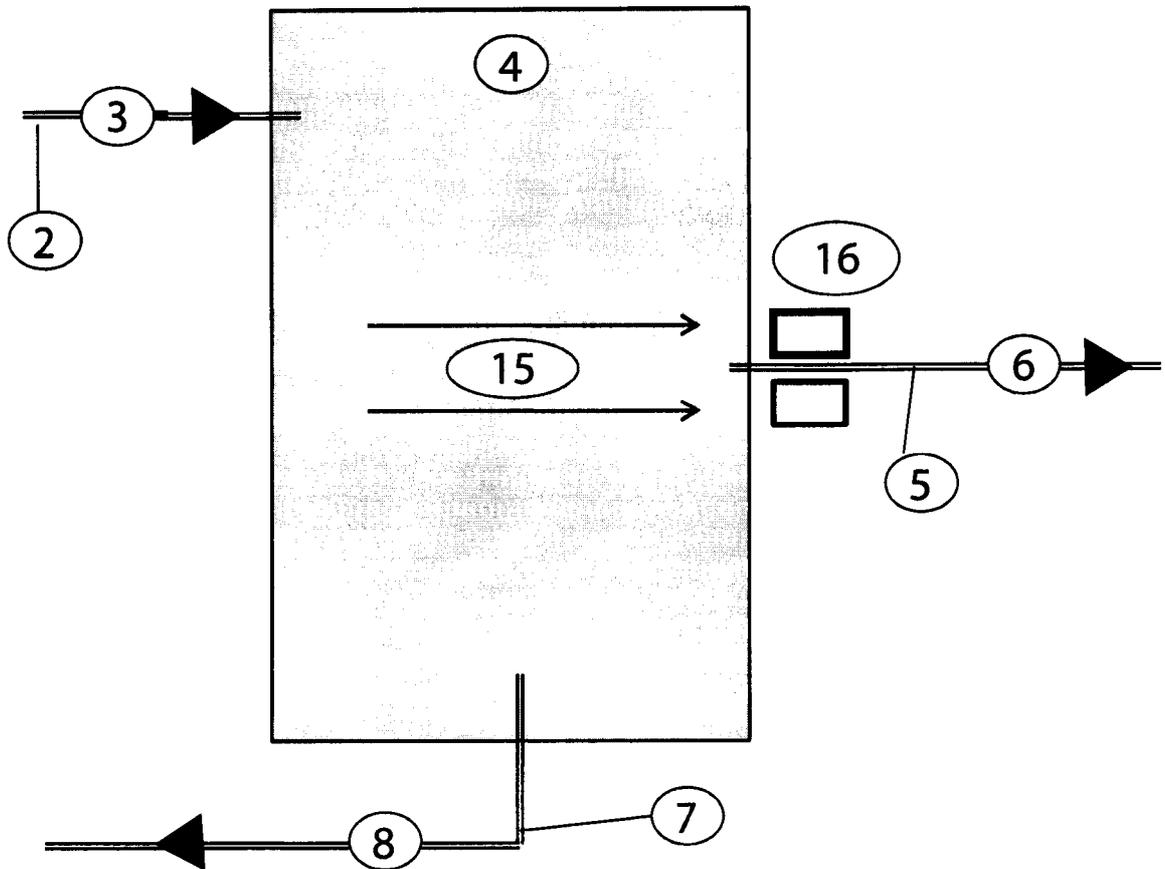
Fig. 13

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- 1 -
- 2 TUBE FOR SATURATED MEAN TAKING
- 3 PUMP FOR SATURATED MEAN TAKING
- 4 EXTERNAL REMOVAL UNIT
- 5 TUBE FOR SEPARATED NUCLEIC ACID
- 6 PUMP FOR SEPARATED NUCLEIC ACID
- 7 TUBE FOR CLEANSED MEAN RECICLE
- 8 PUMP FOR CLEANSED MEAN RECICLE
- 9 -
- 10 -
- 11 -
- 12 CHECK ELECTRONIC OF THE FIELD
- 13 METAL PLATES
- 14 ELECTRIC FIELD

Fig. 14



- 1 -
- 2 TUBE FOR SATURATED MEAN TAKING
- 3 PUMP FOR SATURATED MEAN TAKING
- 4 EXTERNAL REMOVAL UNIT
- 5 TUBE FOR SEPARATED NUCLEIC ACID
- 6 PUMP FOR SEPARATED NUCLEIC ACID
- 7 TUBE FOR CLEANSED MEAN RECICLE
- 8 PUMP FOR CLEANSED MEAN RECICLE
- 9 -
- 10 -
- 11 -
- 12 -
- 13 -
- 14 -
- 15 MAGNETIC FIELD
- 16 PERMANENT MAGNET/ELECTROMAGNET

Fig. 15

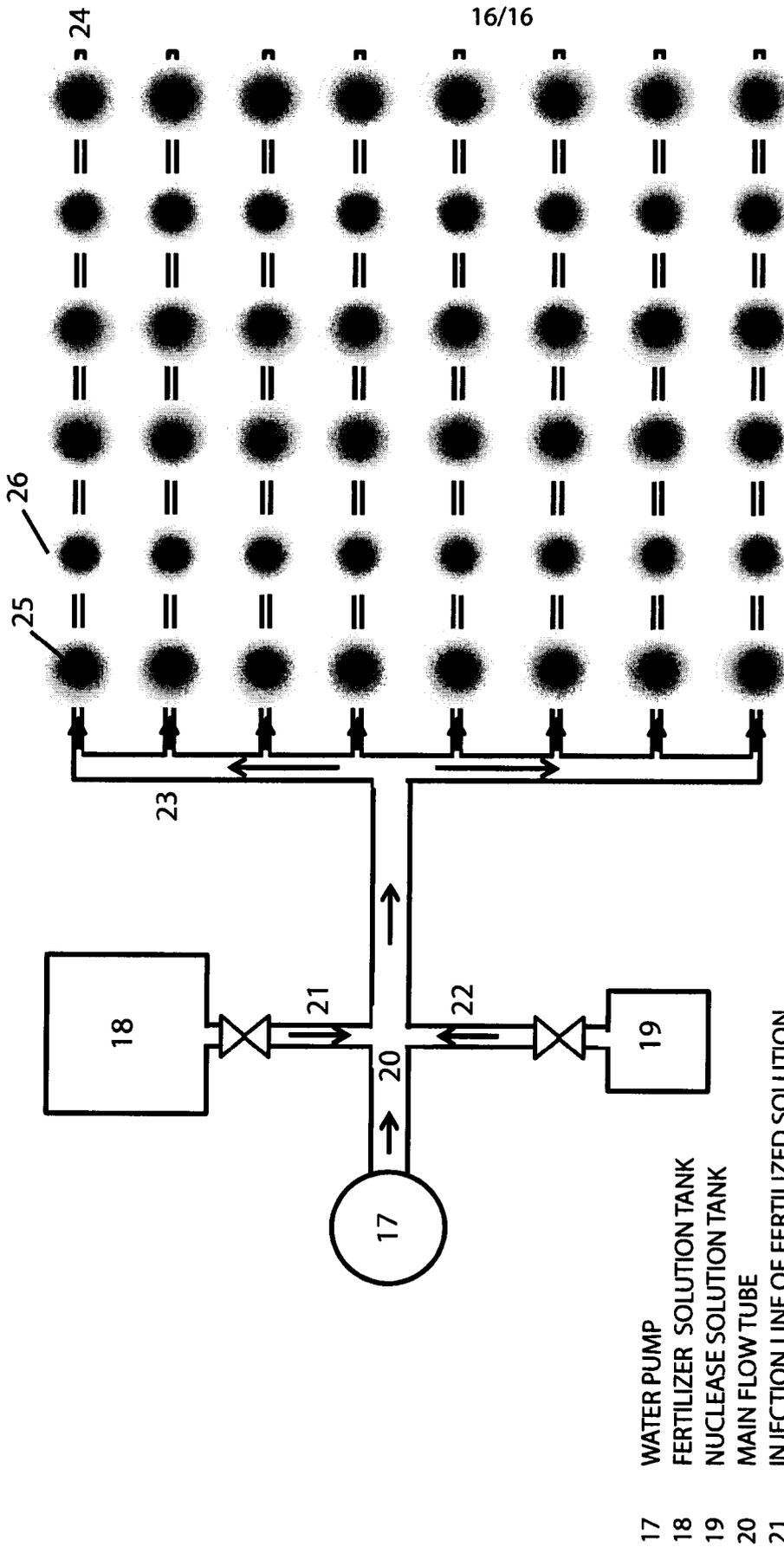


Fig. 16

INTERNATIONAL SEARCH REPORT

International application No PCT/IT2013/000193

A. CLASSIFICATION OF SUBJECT MATTER
 I NV . A91N57/ 10 A01N63/ 00 A01N65/OO A01P1/0 Θ A01 P3/G0
 A01 P5/G0 A01P7/G0 A01 P13/OO A01N57/16

ADD .
 According to International Patent Classification (IPC) or to both national classification and IPC

B. FIELDS SEARCHED
 Minimum documentation searched (classification system followed by classification symbols)
 A01N C07H C12 P

Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched

Electronic data base consulted during the international search (name of data base and, where practicable, search terms used)
 EPO-Internal , CHEM ABS Data , WPI Data

C. DOCUMENTS CONSIDERED TO BE RELEVANT

Category*	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
A	wo 2011/ 112570 AI (MONSANTO TECHNO LOGY LLC [US] ; SAMMONS ROBERT D [US] ; IVASHUTA SERGEY I) 15 September 2011 (2011-09 - 15) paragraph s [0004] , [0006] - [0G09] page 10 , paragraph 50 page 11 , paragraph s 53 , 56-68 - page 17 page 21 , paragraph 74 - page 22 page 63 , paragraph s 104 , 105 page 65 , paragraph 107 - 112 - page 67 page 80 , paragraph 148- 151 - page 81 page 90 , paragraph 174- 176 - page 91 ----- -/--	1 - 14

Further documents are listed in the continuation of Box C. See patent family annex.

* Special categories of cited documents :

"A" document defining the general state of the art which is not considered to be of particular relevance "E" earlier application or patent but published on or after the international filing date "L" document which may throw doubts on priority claim(s) or which is cited to establish the publication date of another citation or other special reason (as specified) "O" document referring to an oral disclosure, use, exhibition or other means "P" document published prior to the international filing date but later than the priority date claimed	"T" later document published after the international filing date or priority date and not in conflict with the application but cited to understand the principle or theory underlying the invention "X" document of particular relevance; the claimed invention cannot be considered novel or cannot be considered to involve an inventive step when the document is taken alone "Y" document of particular relevance; the claimed invention cannot be considered to involve an inventive step when the document is combined with one or more other such documents, such combination being obvious to a person skilled in the art "&" document member of the same patent family
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Date of the actual completion of the international search 29 October 2013	Date of mailing of the international search report 21/03/2014
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Name and mailing address of the ISA/ European Patent Office, P.B. 5818 Patentlaan 2 NL - 2280 HV Rijswijk Tel. (+31-70) 340-2040, Fax: (+31-70) 340-3016	Authorized officer Zanobi ni , Alessandra
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INTERNATIONAL SEARCH REPORT

International application No
PCT/IT2013/000193

C(Continuation). DOCUMENTS CONSIDERED TO BE RELEVANT		
Category	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
X	<p>wo 2005/04984 1 AI (COMMW SCIENT IND RES ORG [AU] ; BAYER BIOSCIENCE NV [BE] ; WATERHOUSE PE) 2 June 2005 (20 05-06-02) page 1, paragraph s 1, 2 page 4, paragraph 6 - page 6, paragraph 8 page 8, paragraph 3 page 10, paragraph 2 - page 11, paragraph 2 page 21, paragraph 3 - page 25, paragraph 1 exampl es 1-3</p> <p align="center">-----</p>	1- 14

INTERNATIONAL SEARCH REPORT

International application No.
PCT/IT2013/00G193

Box No. II Observations where certain claims were found unsearchable (Continuation of item 2 of first sheet)

This international search report has not been established in respect of certain claims under Article 17(2)(a) for the following reasons:

1. Claims Nos.:
because they relate to subject matter not required to be searched by this Authority, namely:

2. Claims Nos.:
because they relate to parts of the international application that do not comply with the prescribed requirements to such an extent that no meaningful international search can be carried out, specifically:

3. Claims Nos.:
because they are dependent claims and are not drafted in accordance with the second and third sentences of Rule 64(a).

Box No. III Observations where unity of invention is lacking (Continuation of item 3 of first sheet)

This International Searching Authority found multiple inventions in this international application, as follows:

see additional sheet

1. As all required additional search fees were timely paid by the applicant, this international search report covers all searchable claims.

2. As all searchable claims could be searched without effort justifying an additional fees, this Authority did not invite payment of additional fees.

3. As only some of the required additional search fees were timely paid by the applicant, this international search report covers only those claims for which fees were paid, specifically claims Nos.:

4. No required additional search fees were timely paid by the applicant. Consequently, this international search report is restricted to the invention first mentioned in the claims; it is covered by claims Nos.:

1-14

Remark on Protest

- The additional search fees were accompanied by the applicant's protest and, where applicable, the payment of a protest fee.
- The additional search fees were accompanied by the applicant's protest but the applicable protest fee was not paid within the time limit specified in the invitation.
- No protest accompanied the payment of additional search fees.

FURTHER INFORMATION CONTINUED FROM PCT/ISA/ 210

This International Searching Authority found multiple (groups of) inventions in this international application, as follows:

1. claims: 1-14

The use of a composition comprising or consisting of a mixture of DNA fragments for the prevention and the treatment against one pathogenic, parasitic or weed species of the plants or environment, wherein said DNA is the total DNA of said at least one pathogenic, parasitic or weed species and/or at least one phylogenetically similar species, against which said prevention and treatment are directed. A method for the prevention or the treatment against at least one pathogenic, parasitic or weed species of the plants or environment, said method comprising or consisting of the following steps of: a) extraction of the total DNA of at least one of said pathogenic, parasitic or weed species and/or at least one phylogenetically similar species, against which said prevention and treatment are directed; b) fragmentation of said total DNA in order to obtain a mixture of DNA random fragments of nucleic acids; and c) contacting said mixture of fragments with said pathogenic, parasitic or weed species. A composition comprising or consisting of a mixture of DNA fragments for the prevention and the treatment against one pathogenic, parasitic or weed species of human or animal, wherein said DNA is the total DNA of said at least one pathogenic, parasitic or weed species and/or at least one phylogenetically similar species, against which said prevention and treatment are directed.

2. claims: 15-22

A process for the production of micro-organisms in bioreactors or photo-bioreactors or systems for hydroponic culture characterized in that the nucleic acids of the same micro-organisms produced by said process are removed from culture medium and the culture medium is used again in said process, wherein said micro-organisms are selected from pathogenic, parasitic or weed micro-organisms; the system used to carry out the process.

INTERNATIONAL SEARCH REPORT

Information on patent Family members

International application No PCT/IT2013/O00193

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