Modern Multidisciplinary Applied Microbiology

Exploiting Microbes and their Interactions



Modern N plinary Applied Microbiology

Exploiting Microbes and Their Interactions

Edited by Antonio Mendez-Vilas





WILEY-VCH Verlag GmbH & Co. KGaA

The Editor

Dr. Antonio Mendez-Vilas Formatex Zurbaran 1 Planta 2a, Local 1 06001 Badajoz Spain

All books published by Wiley-VCH are carefully produced. Nevertheless, authors, editors, and publisher do not warrant the information contained in these books, including this book, to be free of errors. Readers are advised to keep in mind that statements, data, illustrations, procedural details or other items may inadvertently be inaccurate.

Library of Congress Card No.: applied for

British Library Cataloguing-in-Publication Data

A catalogue record for this book is available from the British Library.

Bibliographic information published by

Die Deutsche Bibliothek

Die Deutsche Bibliothek lists this publication in the Deutsche Nationalbibliografie; detailed bibliographic data is available in the Internet at http://dnb.ddb.de>.

© 2006 WILEY-VCH Verlag GmbH & Co. KGaA, Weinheim

All rights reserved (including those of translation into other languages). No part of this book may be reproduced in any form – by photoprinting, microfilm, or any other means – nor transmitted or translated into a machine language without written permission from the publishers. Registered names, trademarks, etc. used in this book, even when not specifically marked as such, are not to be considered unprotected by law.

Printing Strauss GmbH, Mörlenbach

Binding Litges & Dopf Buchbinderei GmbH, Heppenheim

Cover Adam Design, Weinheim

Printed in the Federal Republic of Germany Printed on acid-free paper

ISBN-13: 978-3-527-31611-3 **ISBN-10:** 3-527-31611-6

Preface

This book contains selected papers related to contributions that were presented in short during the 1st International Conference on Environmental, Industrial and Applied Microbiology (BioMicroWorld-2005), held on March 15–18th, 2005 in Badajoz (Spain). http://www.formatex.org/biomicroworld2005, and it is intended to give an overview on the current state-of-the-art of the field.

Focus of the Conference. While Microbiology is about the study of microorganisms (bacteria, viruses, algae, fungi, and protozoa) and of related topics such as microbes interactions, the immune response and molecular genetics, Applied Microbiology is quite interdisciplinary, overlapping aspects of several other academic branches, some of them traditionally near, such as cell biology, molecular and cell biophysics, physiology, parasitology, biochemistry, genetics, medicine, pharmacology, and medical technology, and other not so near as physics, physical (bio)chemistry, materials science, nanotechnology, computer science, information technology, instrumentation, but collaboration with which is resulting in extraordinary advances in this post-genomic world. Thus, cross-disciplinary cooperation in Microbiology has made possible that microbiologists can not only study traditional microorganisms, but also aspects of molecular genetics, biosensors, cancer, aging, immunodeficiency diseases, animal and plant cell cultures, and microscopy, among others. Modern microbiology includes a broad variety of scholarly approaches which lead to a better understanding of all living things at the micrometer-scale/cellular and nanometerscale/molecular level, and which produce beneficial applications in medicine, agriculture, industry, and ecology.

In this context, the Conference called for papers reporting interdisciplinary resear-chers, relating Microbiology with other Sciences as Physico/Chemistry, Environ-mental Science, Genetics, Pharmacology, Nanoscience, Microscopy/Imaging Science, etc. In other words, we are specially (but not exclusively) interested in reports applying the techniques, the training, and the culture of Microbiology to research areas usually associated with other scientific and engineering disciplines. Over 750 participants from over 60 countries attended the Conference, 15% of which participated with a grant from the conference organization. Over 1100 works were presented during the different oral and posters sessions. Good examples of modern interdisciplinary applied microbiology were the works represented by the three Plenary Speakers:

David C. White, Director of the Center for Biomarker Analysis, University of Tennessee, USA

Lecture: Biomarkers to define Interactions in the Environment and Health

Alexander Steinbüchel, Institut für Molekulare Mikrobiologie und Biotechnologie, Münster, GERMANY

Lecture: Unspecific Microbial Enzymes for Template-independent Biosynthesis of Polyoxoesters, Polythioesters and Polyamides

Timo Lövgren, Department of Biochemistry and Food Chemistry/Biotechnology, University of Turku, FINDLAND

Lecture: Novel time-resolved Fluorescence based Immunoassays and Real-time PCR assays in Microbiological Applications

I am are very grateful to all members of the Organizing Committee for the hard work done in the Conference preparation (which began over a year before the conference) and for the good job that made the Conference so successful that there are already several candidates to host next edition of BioMicroWorld. We would also like to thank the members of the International Scientific Advisory Committee, as well as the reviewers, for their advice, which has certainly helped to improve the quality, accuracy and relevance of this conference Program and publications.

Finally, we would like to thank *BIOMEDAL S.L.-Advances for the Postgenomic Era* (<u>http://www.biomedal.es</u>) for sponsoring the Conference.

Hoping that this edition will stimulate further meetings focusing on the interdisciplinary nature of current relevant applied research, we hope that readers will find the content fruitful and interesting.

Antonio Mendez-Vilas Editor

FORMATEX, Badajoz, Spain June, 2006

List of Contents

Preface v

Environmental Microbiology, Marine Microbiology, Water/Aqu Geomicrobiology	atic Microbiology
A new potential indicator of virological contamination of surfaces	3

A novel mechanism for bacterial acid resistance: a carbon dioxide-dependent system 8

An individual based model to study the main groups of microbes active in composting process 14

Biotechnological approach for treatment of textile wastewater – A case study 19

Catalytic performance of lignin peroxidase hosted in AOT reverse micellar medium 28 Characterization of β -lactam-resistant genes from a metagenomic library of cold-seep sediments in deep-sea 33

Cobalt-induced stimulation and inhibition of cytochromes synthesis and extracellular nitrite release in *Paracoccus denitrificans* 42

Communication of microbial risk to workers in a wastewater treatment plant 46

De novo protein production under long term starvation in Flexibacter chinensis 50

Development of a biotechnology tool using New Zealand white-rot fungi to degrade pentachlorophenol in soil $-\,A$ summary $\,$ 55

Ecological interactions among protozoan parasites and their avian hosts: an approach 60

Effects of hydrodynamics on biofilm formation 64

Effect of organic management on soil chemical and biochemical properties of a Xerofluvent of the Guadalquivir River Valley (SW Spain) 73

Environmental virological monitoring for the epidemiological surveillance and risk assessment 78

Heavy metal effects on Extremophiles and on enzyme biosynthesis in a new *Bacillus* strain from Mount Rittmann, Antarctica 83

Heavy metal toxicity in *Rhizobium leguminosarum* biovar viciae isolated from soils neighbouring metalomechanics industries 89

Isolation and screening of potential fungi for decolourization of distillery wastewaters 95

Microbial corrosion in zinc surface layer of galvanized steel by biofilms of sulfate reducer bacteria 103

Microbial diversity and metabolic pathway analyses of a mesophilic acetate-degrading methanogenic community in a chemostat cultivation 109

Microbiota associated with Posidonia oceanica in Western Mediterranean sea 114

Novel methodologies for the detection and classification of cultured and uncultured microorganisms from cultural heritage samples 120

Perylene toxicity in the estuarine environment of Ria de Aveiro (Portugal) 125

Phenotypic and genotypic characterization of an outbreak of *Pseudomonas aeruginosa* from wild birds 130

Single and multi-metal removal by an environmental mixed bacterial isolate 136

Study of *Helicobacter pylori* viability in aquatic microcosms by epifluorescence stain and *in situ* hybridization 142

Survival of *Ralstonia solanacearum* biovar 2 in river water: influence of water microbiota 147

The control of biological corrosion in cooling water system of a power plant 153

The effects of various antibiotics on marine bacteria 159

The influence of temperature and type of illumination on the biochemical composition and EPA production of *Nannochloropsis gaditana* 167

Two-phases olive mill solid waste treatment in an anaerobic continuous stirred tank reactor at diluted influent concentrations: evaluation and implicated microorganisms 172

Use of rRNA gene restriction patterns in *Colibacillosis* by *E. coli* O78, O88 and non typable strains in wild birds 178

Industrial Microbiology – Future Bioindustries

Acrylic acid removal from synthetic wastewater and actual industrial wastewater by pure culture of bacteria isolated from the acrylonitrile-butadiene-styrene resin manufactured wastewater treatment system 189

Biodegradation of hydrolyzed polyacrylamides in aqueous solution 194 Biomethanation of sugar industry wastewater by using down-flow anaerobic filter 200

Biosynthesis of calcium gluconate by Aspergillus niger in shake flask 206

Combined effect of doxorubicin and metal on the yeast Candida utilis 216

Hydrodynamic analysis and inoculum effect on *Rhizopus nigricans* growth for cellulase production in bubbling column bioreactor 221

Inactivation mechanisms of his-tagged D-amino acid oxidase from *Trigonopsis variabilis* 227

Killer activity of yeasts isolated from Spanish dry-cured ham 232

Limiting factors of photo-hydrogen production by *Rhodopseudomonas palustris* WP3-5 236

Mineral-phosphate solubilization activity of iron ore associated microflora 241

Optimisation of two recombinant whole cell systems for the production of optically pure D-amino acids 246

Single cell protein production by *Saccharomyces* sp no12 by utilizing Lignocellulosic waste and its nutritional evaluation 251

Study on the hemoprotein at 503 nm in Salmonella typhimurium 256

The knowledge flow and commercialisation along the continuous chain: education – research – implementation by innovative methods 261

Xanthan gum production using whey for preculture preparation 265

Food Microbiology

An assessment of differential media for the recovery of histamine producing bacteria 271

Aeromonas growth under low temperatures 275

Comparison of two processes for isolation of exopolysaccharide produced by $Lactobacillus\ acidophilus\ 280$

Detection of *Listeria monocytogenes* from fresh and marine water fish using Real Time PCR, PCR and standard ISO methods 286

Direct identification of *Campylobacter* species in poultry samples by multiplex PCR 291

Ecology and biodiversity of microbial populations in spontaneous grape must fermentations 296

Effects of NH₄VO₃ on mitochondrial electron-transport chain and glutathione peroxidase activity of Saccharomyces cerevisiae UE-ME₃ 301

Identification by PCR of trichothecene-producing Fusarium 306

Isolation of *Arcobacter butzleri* from meats and its susceptibility to various acids and humectants 311

PCR assay for identification of *Aspergillus* section *Nigri* using internal transcribed spacer regions 1 and 2 316

Progress in prevention of aflatoxin contamination in food by preharvest application of a yeast strain, *Pichia anomala* WRL-076 322

Quantitative models application to fulfil microbiological criteria in foods 327

Rapid biased evolution of genetically unstable wine yeast hybrids under non-selective conditions 332

Rodamine resistance as marker for monitoring yeasts in wine fermentations 337

Strength of attachment affects survival of *Salmonella* on inoculated cantaloupe treated with sanitizers 342

Sulfometuron methyl resistance as genetic marker for monitoring yeast populations in wine fermentation 351

Use of carbon dioxide to control the microbial spoilage of bullfrog (*Rana catesbeiana*) meat 356

Agriculture, Soil, Forest Microbiology

Agricultural non point sources control against microbiological contaminations of drinking water resources in dairy mountain areas 364

Antagonistic activity of actinomycetes and fungi from horticultural compost against phytopathogen microorganisms 370

Application of arbuscular mycorrhizal fungal in vitro biofertilizers in agro-industries 375

Biological control of flax scorch using *Glomus intraradices* and *Trichoderma atroviride* 380

Characterization of Brenneria sp. from poplar cankers in Spain 385

Environmental purification capacity and control of microbiological contamination of water resources in dairy mountain area 390

Group specific primers for studying Methylobacterium biodiversity on crop plants 397

Microbial biodiversity in soils under a conservation agriculture regime 404

Molecular analysis of microbial communities in northern terrestrial systems 408

MycorID: a molecular database based on ITS-RFLP analysis for identification of ectomycorrhizae at Iberian Peninsula 411

Nitrification and denitrification associated with N_2O production in a temperate N-fertilized irrigated Uruguayan rice field 416

Nitrifier bacterial activity linked to mineralization of soil organic matter: individual based simulations 421

Nitrogen biological fixation ability by *Rhizobium legominosarum* biovar *phaseoli* on cultivars of *Phaseolus vulgaris* L. 427

Phenotypic and genetic diversity of fluorescent *Pseudomonas* recovered from different host plants 435

Population of Aspergillus flavus on pistachio buds and flowers 440

Removal of microorganisms present in lettuces and soil irrigated with treated wastewaters 446

Soil amendment with sludge generated from metal finishing industries and its impact on metabolic quotient 451

Spatial properties in Individual-based Modelling of microbial systems. Study of the composting process. 461

Ultrastructural and cytochemical aspects of spores germination of *Mucor javanicus* wehmer 466

Bioremediation

Activity of soil microbial communities to monitor the efficiency of a metal phytoremediation process with *Thlaspi caerulescens* 473

Bioprocesses under practical aspects – Multimedia environmental impact assessment of in situ – soil remediation with genetically modified microorganisms 478

Characterization of *Cladosporium oxysporum* and C. *sphaerospermum* using Polyaromatic hydrocarbons (PAHs) as their sole carbon source in tropical coastal seawater 483

Comparison of microbial communities native to three differently polluted ecological niches in the industrial site of Bagnoli (Naples,Italy) 488

Decolorization of Methyl Orange (as a model azo dye) by the newly discovered *Bacillus sp* 494

Degradation of Polyethylene film strips by soil fungal isolates 500

Enhanced biodegradation of polycyclic aromatic hydrocarbons using nonionic surfactants in soil slurries. 506

Enumeration of naphthalene and phenanthrene degrading bacteria as an indicator of hydrocarbon pollution in surface waters of Guayanilla, Puerto Rico: Impact of seasonal variations 511

Formation of Cr(V) and Cr(III) in *Arthrobacter oxydans* exposed to high concentrations of Cr(VI) 516

Metabolism of cyanate and cyanide in the alkalophilic bacterium *Pseudomonas* pseudoalcaligenes CECT5344 521

Microbial ecology of authothermal aerobic digestion (ATAD): diversity, dynamics and activity of bacterial communities involved in treatment of a municipal wastewater. 526

Microbial function after assisted natural remediation of a trace element polluted soil 536

Microbiological characterization of ammonium oxidizing biofilms in rotating biological contactor (RBC) using different support materials 541

Occurrence of two metabolic pathways in benzo[a]pyrene degradation by a Deuteromycete fungus *Fusarium solani* 546

Oleophilic fertilizers and Bioremediation: A new perspective. 551

On Site bioremediation and washing techniques in a cobble beach affected by Prestige oil spill 556

Towards the applicability of rhodococci in monoaromatic compounds bioremediation 561

Transport behaviour of chemotactic bacteria in model aquifers 566

Microbial Biotechnology

Biomass selection and acclimatization for toluene degradation in a biofilter 573

Cloning and overexpression of an alkaline and thermostable α-amylase from a native Iranian Bacillus species (*Bacillus* sp-GSH) isolated from the soil of Rasht 577

Evaluation of performance and kinetics of mesophilic and thermophilic anaerobic digestion for treatment of palm oil mill effluent 582

Expression in *E. coli* of a recombinant form of pulmonary surfactant protein precursor proSP-B $_{\Delta C}$ 587

Influence of grape sanitary quality on yeast killer effect at the winery 592

Inhibitory effects of UV-absorption compounds in hemicellulose hydrolysate on xylitol production by yeast 596

Production and Application of 5-Aminolevulinic Acid from *Rhodobacter capsulatus* SS3 Cultivating in Monosodium Glutamate Effluent 602

Protein profile of Streptomyces clavuligerus strains related to regenerating cells 609

Microfactories – Microbial Production of Chemicals and Pharmaceuticals. Biopolymers

Isolation and screening of oleagenous microorganisms for the production of a nutraceutical – single cell oil 615

Manufacturing and characterization of bacterial cellulose tubes using two different fermentation techniques 619

Microbial biotechnology and bioengineering aspects for large-scale xylitol bioproduction from lignocellulosic residues 623

Neural network based software sensors: application to biosurfactant production by *Candida lipolytica* 628

Microbial Physiology, Metabolism and Gene Expression

Conventional and real-time PCR assays for detection and quantification of the expression of fumonisin biosynthetic gene *fum5* in *Fusarium verticillioides* 635

Induction of chromosome replication as a heat stress response in Escherichia coli 640

Medium Effect on Heterologous Protein Leaky Expression in Escherichia coli 645

Modulation of gene expression by 3'-UTR regions in yeast: a promoter-independent alternative 649

Molecular properties of two nitroreductases from *Rhodobacter capsulatus* B10 involved in 2,4-dinitrophenol reduction 654

Nitrate assimilation in *Rhodobacter capsulatus* E1F1: purification and biochemical characterization of nitrite and hydroxylamine reductases 660

Proteases of *Trichoderma* strains from Hungarian winter wheat rhizosphere 664

The yeast external invertase as a reporter to study regulation of *Candida albicans* promoter sequences in *Saccharomyces cerevisiae* 669

Towards understanding the acetic acid resistance in *Gluconacetobacter europaeus* 674

Transcriptome dynamics of ethanologenic yeast in response to 5-hydroxymethylfurfural stress related to biomass conversion to ethanol 679

Medical Microbiology

Detection and characterization of integrons, transposons, plasmids and genomic types in multidrug resistant clinical isolates of *Salmonella enterica* serovar Virchow. 687

Effect of the essential oils of Brazilian species of the genus *Cunila* on the growth and biofilm formation by *Aeromonas*. 692

Human and bovine constitute reservoirs of different sub-populations of *Staphylococcus aureus* in possession of the highly prevalent enterotoxin gene cluster egc_{like}. 697

Improved cytocompatibility of titanium alloy by coating with pure titanium films using sputter-deposition 702

Is intestinal microbiota bound up with changing lifestyle? 708

PCR procedures for detection of emergent multidrug resistant lineages of *Salmonella* serovar Typhimurium. Evolution of a lineage carrying the plasmid pUO-StVR2 713

Analytical Techniques, Imaging Techniques, Microscopy

Biotechnology, microbiology and secondary school 721

Ciliates as potential biosensors for heavy metal pollution 726

Concentration, detection and identification of Infectious Enteroviruses in Sewage 731

Effect of inoculum on the survival of pathogenic agents in different composts 737

Microbial diversity and chimerae from PCR amplified products 744

Screen-printed metal oxides-based enzyme biosensors for food analysis 751

The study of the fermentative growth of *Saccharomyces cerevisiae* S288C using auxoaccelerostat technique 756

Methods in Basic and Applied Microbiology. Microbiology Education

Comparison of different analytical methods for determination of type B trichothecenes in wheat and ochratoxin A in beer 763

DVC-FISH procedure to enumerate specific viable cells of *Lactobacillus delbrueckii* subsp. *bulgaricus* DN-100182 772

Fluorescence imaging and new fluorescence techniques applied to cellular pathopharmacology 779

Impact of *Bifidobacterium animalis* DN-173010 on Human Intestinal Microbiota by Fluorescent in situ Hybridization 784

Scanning electron microscope study of fish and rice flour coextrudates 791
Thin section and freeze-fracture electron microscopy of *Paracoccus denitrificans bacterium* 796

Subject Index 800

Environmental Microbiology, Marine Microbiology, Water/Aquatic Microbiology, Geomicrobiology



A new potential indicator of virological contamination of surfaces

M. Verani¹, B. Casini¹, E. Rovini¹, P. Paone¹, A. Mansi², R. Lombardi², and A. Carducci¹

Keywords: TTVirus, Viral indicator, environmental monitoring

1 Introduction

TT virus is a widespread infectious agent of humans identified in 1998 (1). The studies carried out till now have evidenced a large diffusion of the virus in the world population with a persistent viremia in infected people characterized by a low pathogenic potential. TTV is present in blood, serum, faeces, pharyngeal and nasal swabs and other biological fluids, it has a high environmental resistance and several ways of elimination (2,3). All this characteristics, linked to the technical problems and great limits that present the direct research of pathogenic viruses in environment, suggest the utilization of TT virus as a new possible indicator of presence of haematic and entero-oral transmission viruses (4). TTV could be researched in particular settings like hospitals, clinical laboratories, etc. in witch the presence, at high concentration, of these pathogenic viruses represent a real risk for health-care workers (5).

2 Materials and methods

The study was carried out in two stages:

- 2.1 Estimating the sensitivity limits of the technique for TTV detection on surfaces artificially contaminated.
- 2.2 Environmental monitoring utilising the moist sensible technique.

2.1.1 Preparation of artificial samples

The serum from a TTV virus-infected subject with viral title of 46×10^6 copies/ml has been diluted from 10^{-1} to 10^{-4} in bovine fetal serum negative for TTV-DNA presence. 10 microliters of the whole serum and the dilutions, in addition to a negative control were spread on a sterilized stainless steel plate. The plate was then dried for about 10 minutes.

2.1.2 Estimation of sensitivity limits of sample purification and DNA extraction methods

Two analytical protocols have been compared on artificial samples:

Department of Experimental Pathology, Medical Biotechnologies, Infectivology and Epidemiology, University of Pisa, Via S. Zeno 35 56127 Pisa, ITALY
 Institute of Occupational Safety and Prevention, Department of Occupational Hygiene, Rome, ITALY

- a) Two eluents were tested contemporary: beef extract (BE) 3% at pH 9 and bovine serum albumin (BSA) 1% with NaCI 0.85%. For elution trials, the surface was repeatedly wiped with a cotton swab impregnated with 1 ml of eluent. The swab was then dipped in a test tube containing 1 ml of the same eluent, kept in a refrigerator for 2 hours at 4°C during which time it was shaken every 15 minutes, and finally mixed with vortex. The eluates were then recovered and extracted using a commercial kit "QIAamp DNA Mini Kit" (Qiagen). The recovery test was repeated three times.
- b) Application of a commercial kit (DNA IQTM SYSTEM), generally used in forensic field for the detection and purification of DNA present in several biological samples, on the body and on objects for personal use; it was modified in order to achieve the present work using the Lysis Buffer of the kit as eluent to wash the artificially contaminated surface (250 μl) and cutting the swab of 1 cm long to allow its introduction in a spin basket put in a centrifuge test-tube containing 50 μl of the same eluent. Subsequently the swab was treated as indicated by the protocol of the commercial kit. Also in this case the recovery tests were repeated four more times.

2.1.3 Qualitative detection and quantitative determination of the TTV genome by PCR

The qualitative detection of the TTV genome was carried out by "nested PCR", which uses specific primers drawn by the UTR region of the TTV genome (6). For every reaction negative and positive controls were inserted. In order to quantify the TTV genome the TaqMan-Applied Biosystems Prism 7700 (Poster City, California) system has been used, drawing the primers and probe from the UTR region (7).

2.2 Application of the selected method for the environmental monitoring

The most sensible technique is actually used for an environmental monitoring in different sites of an hospital associated with research of an other biological indicator: haemoglobin (Table 1). A total of 74 selecting points were sampled, chosen mainly for the high probability of becoming contaminated, such as work benches, centrifuges, biosafety cabinet, and other instrumentation.

For TTV and haemoglobin detection, cotton swabs soaked in the relative eluent were wiped repeatedly on area of 36 cm squares. For the detection of haemoglobin a kit used for the blood detection hidden in faeces (Kit OC-Hemocard-Alfabiotech®-Wasserman) was modified and applied to the purpose (8).

Table 1 Environmental n	nonitoring
--------------------------------	------------

ENVIRONMENT	ANALYZED SAMPLES
Clinical Lab.	12
Surgery	31
Cardiac Unit Intensive Terapy	16
Rianimation	9
Surgery passage	5
TOT	74