

Poster Abstracts (continued)

Blue Light Effects, Photochemistry, Luciferase

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| I01 | Class II DNA photolyase from <i>Arabidopsis thaliana</i> contains FAD as cofactor | Oliver Kleiner, Jens Butenandt, Thomas Carell and Alfred Batschauer |
| I02 | Occurrence of P-flavin binding protein in <i>Vibrio fischerii</i> and properties of the protein | Sabu Kasai |
| I03 | Modelling the intermediate IV of the luciferase reaction: characterisation of the complex of 5-decylFMN-4a-OH with <i>Vibrio harveyi</i> luciferase | Benfang Lei, Qizhu Ding and Shiao-Chun Tu |
| I04 | In DNA photolyase from <i>Anacystis nidulans</i> primary electron transfer from tryptophan to FADH⁺ is followed by oxidation of a tyrosine residue | Corinne Aubert, Paul Mathis, André P. M. Eker and Klaus Brettel |
| I05 | DCRY: A <i>Drosophila</i> photoreceptor protein implicated in light entrainment of circadian rhythm | Tomoko Ishiwaka, Akira Matsumoto, Teiichi Tanimura, Shin Togashi, Ryu Ueda and Takeshi Todo |
| I06 | Isolation and characterization of DNA photolyase/cytochrome family in Zebra fish | Yuri Kobayashi, Tomoko Ishikawa and Takeshi Todo |

Flavoproteins, 3D Structures

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| J01 | Crystal structures of „unactivated” p-hydroxybenzoate hydroxylase | Michel H. M. Eppink, Willem J. H. van Berkel, Alex Tepliakov and Herman A. Schreuder |
| J02 | Crystallographic studies of a complex between the ferredoxin-NADP⁺-reductase from the cyanobacterium <i>Anabaena</i> PCC7119 and its functional partner | Renaud Morales, Galina Kachalova, Marie-Hélène Charon and Michael Frey |
| J03 | X-Ray structure of a monomeric α subunit of the sulfite reductase from <i>E. coli</i> | A. Gruez, D. Pignol, J. L. Ferrer, M. Zeghouf, J. Coves, M. Fontecave and J.-C. Fontecilla-Camps |
| J04 | NMR-studies on FMN-binding protein from <i>Desulfovibrio vulgaris</i> (Miyazaki F) | Masaya Kitamura, Hideo Inoue, Edwards Liepinsh and Gottfried Otting |
| J05 | Crystal structures of the tetraheme flavoprotein enzyme fumarate reductase from <i>Shewanella putrefaciens</i> strain MR1 | David Leys, Alexandre Tsapin, Terrance E. Meyer, Michael A. Cusanovich, Yves Guisez Jozef J. Van Beeumen |
| J06 | Crystallographic analysis of C44S NADH oxidase from the human pathogen <i>Streptococcus pyogenes</i> | T. Conn Mallett, Hiroaki Sakai, Derek Parsonage, Al Claiborne and Tomitake Tsukihara |
| J07 | A model of the structure of 6-hydroxy-D-nicotine oxidase | William S. McIntyre, Igor Efimov, Louise M. Cunane and Scott Mathews |

J08	Crystal structure of the flavin reductase Fre from <i>Escherichia coli</i>	Margareta Ingelman, S. Ramaswamy, Vincent Nivière, Marc Fontecave and Hans Eklund
J09	Towards the structure of a soluble fumarate reductase	Sara L. Pealing, Paul Taylor, Graeme A. Reid, Stephen K. Chapman and Malcolm D. Walkinshaw
J10	Characterization of two <i>E. coli</i> methylenetetrahydrofolate reductase (MTHFR) mutants, Asp120Asn and Glu28Gln	Elizabeth E. Trimmer, Brian D. Guenther, Martha L. Ludwig and Rowena Matthews
J11	Characterization of glutathione amide reductase from the purple bacterium, <i>Chromatium gracile</i>	Bjorn Vergauwen, Vesna Kostanjevecki, Yves Guisnez and Robert Bartsch
J12	Domain structure and kinetic analysis of streptococcal L-α-glycerophosphate oxidase	Véronique Charrier, James Luba, Derek Parsonage and Al Claiborne
J13	Refined 2.1 Å structure of the cysteine-sulfenic acid redox center in NADH peroxidase	Joanne I. Yeh, T. Conn Mallett, Edward J. Crane III, Derek Parsonage, Wim G. J. Hol and Al Claiborne
J14	Crystal structure of polyamine oxidase from <i>Zea mays</i> L. at 1.9 Å resolution	Claudia Binda, Alessandro Coda, Riccardo Angelini, Rodolfo Federico, Paolo Ascenzi and Andrea Mattevi
J15	A new functional model for the <i>Escherichia coli</i> sulfite reductase: the $\alpha 1\beta 1$ complex	Jaques Coves, Mahel Zeghouf and Marc Fontecave

Monooxygenases

K01	Is charge transfer complex formation essential for reduction of <i>p</i>-hydroxybenzoate hydroxylase?	Barrie Entsch, Mariliz Ortiz-Maldonado and David P. Ballou
K02	Catalytic mechanism of 2-hydroxybiphenyl 3-monooxygenase	Winfried A. Suske, Hans-Peter Kohler and Willem J. H. van Berkel
K03	Purification and some properties of acetophenone monooxygenase	Mariëlle J. H. Moonen, Ivonne M. C. M. Rietjens and Willem J. H. van Berkel
K04	Effect of Substrate and Flavin Activation on the Hydroxylation Reaction of <i>p</i>-Hydroxybenzoate Hydroxylase: Studies of Site Directed Mutants Substituted with 8-Cl-FAD	Mariliz Ortiz-Maldonado, David P. Ballou and Vincent Massey
K05	Heterologous Expression and Kinetic Characterization of Human Squalene Monooxygenase	Brian P. Laden and Todd D. Porter
K06	Novel two-component phenol hydroxylase from a thermophilic <i>Bacillus</i> strain	Ulrike Kirchner, Willem J. H. van Berkel and Rudolf Müller
K07	Development of biocatalyst for acetophenone monooxygenase catalyzed reactions: kinetic and mechanistic characterization of the enzyme and improvement of the biocatalyst	N. M. Kamerbeek and D. B. Janssen

K08	Coenzyme recognition by flavoprotein aromatic hydroxylases	Willem J. H. van Berkel, Michel Eppink and Herman Schreuder
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Flavins and Oxygen, Various

L01	Kinetics, mechanism and regulation of elementary steps of catalysis of pyruvate oxidase from <i>Lactobacillus plantarum</i>	Kai Tittmann, Ralph Golbik, Sandro Ghisla and Gerhard Hübner
L02	Studies on the peroxide- reducing system of <i>Thermus aquaticus</i>.	Catriona Logan and Stephen G. Mayhew
L03	Overproduction of <i>Lactococcus lactis</i> NADH oxidase and its application to metabolic engineering	Derek Parsonage, Michiel Kleerebezem, Iris I. van Swam and Al Claiborne
L04	Converting a dehydrogenase into an oxidase	Graeme A. Reid, Stephen K. Chapman, Lars Østergaard and Martin L. Goble
L05	What protein features make p-hydroxybenzoate hydroxylase react rapidly with oxygen?	Willem van Berkel, Michel Eppink, Herman Schreuder, Mariliz Ortiz-Maldonado, David Ballou and Barrie Entsch
L06	The intermediates involved in the catalytic reaction of cyclohexanone monooxygenase	Dawei Sheng, David P. Ballou and Vincent Massey

Amine Dehydrogenation

M01	Substrate inhibition in trimethylamine dehydrogenase	Peter Roberts, Jaswir Basran, Emma K. Wilson, Russ Hille and Nigel S. Scrutton
M02	Bacterial amine oxidation through horizontal gene transfer from an eukaryotic source	Susann Schenk and Karl Decker
M03	A novel heterotrimeric flavoprotein involved in bacterial nicotine catabolism	Susann Schenk, André Hoelz and Karl Decker
M04	High level expression and characterization of recombinant human liver MAO B	Paige Newton-Vinson and Dale E. Edmondson
M05	A mechanism for monoamine oxidase involving a redox-active disulfide	Rona R. Ramsay
M06	Involvement of mitochondrial matrix in the holoenzyme formation of dimethylglycine dehydrogenase	Carmen Brizio, Annegret Otto, Ernesto Quagliariello, Roderich Brandsch, Salvatore Passarella and Maria Barile
M07	Hydrogen tunnelling in amine dehydrogenases from methylotrophic bacteria	Jaswir Basran, Peter Roberts, Michael J. Sutcliffe and Nigel S. Scrutton
M08	Investigating the mechanism of C-H bond breakage in heterotetrameric sarcosine oxidase from <i>Arthrobacter</i> sp. 1-IN	Richard J. Harris, Rolandas Meskys, Michael Sutcliffe and Nigel S. Scrutton

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| M09 | The reductive half-reaction of trimethylamine dehydrogenase with trimethylglutamine | Mei-Huei Jang, Jaswir Basran, Nigel S. Scrutton and Russ Hille |
| M10 | Interaction of FAD analogues with the C406A mutant apoenzyme of human liver monoamine oxidase A | Ravi K. Nandigama and Dale E. Edmondson |

α-Hydroxy Acid Dehydrogenases

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| N01 | X-ray studies of recombinant rat kidney long-chain hydroxy acid oxidase and of the recombinant flavin-binding domain of bakers yeast flavocytochrome b_2 | Louise Cunane, Fraser Welsh, John Barton, Zhi-wei Chen, Ahmed Belmouden, K. H. Diêp Lê, Florence Lederer, Steve Chapman, Graeme Reid and F. Scott Mathews |
| N02 | On the mechanistic value of the dehydrohalogenation reaction of β-halogeno hydroxy acids catalysed by FMN-dependent hydroxy acid-oxidising enzymes: a mutational analysis with flavocytochrome b_2 | Sabrina Bodevin and Florence Lederer |
| N03 | D-lactate dehydrogenase model: mechanism of the oxidation of mandelic acids by functionalized flavin mimics with metal ions | Hideaki Ohshiro, Shin-ichi Kondo and Yumihiko Yano |
| N04 | (S)-mandelate dehydrogenase from <i>Pseudomonas putida</i>: mechanistic roles of the conserved residues H274 and R277 | Isabelle E. Lehoux and Bharati Mitra |
| N05 | On the role of Asp 180 in L-lactate monooxygenase | Stephen A. Sanders, Ute Müh, Charles H. Williams Jr. and Vincent Massey |
| N06 | On the mechanism of glucose oxidase | Ben J. L. Williams and Stephen E. J. Rigby |
| N07 | Reaction mechanism of L-lactate oxidase from <i>Aerococcus viridans</i> | Kazuko Yorita, Toshiyuki Watanabe, Hideo Misaki and Vincent Massey |
| N08 | The substrate specificity of L-mandelate dehydrogenase | Douglas J. Robertson, Graeme A. Reid and Stephen K. Chapman |
| N09 | The membrane-associated (S)-mandelate dehydrogenase from <i>Pseudomonas putida</i>: characterization of a highly active, soluble mutant | Yang Xu and Bharati Mitra |

Acyl-CoA Dehydrogenases

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| O01 | Mechanism-based inactivation of medium-chain Acyl-CoA dehydrogenase by a cytotoxic thioester: bioactivation of 5,6-dichloro-4-thia-5-hexenoyl-CoA | Jennifer F. Baker-Malcom, M. W. Anders and Colin Thorpe |
| O02 | Synthesis and activity of substrate analogs for glutaryl-CoA dehydrogenase | Shannon L. McKinney, Carrie L. Donley, Gregory K. Sewall, Patricia L. Kultgen and Colleen M. Byron |
| O03 | Difference raman studies of hexadienoyl-CoA bound to medium chain acyl CoA dehydrogenase | Alasdair F. Bell, Jiaquan Wu, Kim M. Sabaj, Avery W. Stephens, Marian T. Stankovich and Peter J. Tonge |

O04	Redox potential measurements of short chain acyl-CoA dehydrogenase (SCAD) active site mutants	Jackson D. Pellett, Donald F. Becker, James A. Fuchs and Marian T. Stankovich
O05	Interactions of rat acyl-CoA oxidase with substrate analogs used as active-site probes	Haruhiko Tamaoki, Chiaki Setoyama, Kyosuke Sato, Yasuzo Nishina, Kiyoshi Shiga and Retsu Miura
O06	Biochemical characteristics of recombinant human isovaleryl-CoA dehydrogenase pre-treated with ethylenediaminetetraacetate	Al-Walid A. Mohsen and Jerry Vockley
O07	Probing the mechanism of medium chain acyl-CoA dehydrogenase (MCAD) using spectrally active alternative-substrates and products	Teresa R. Lamm and Marian Stankovich
O08	Substrate chain length specificity of acyl-CoA dehydrogenases: studies on different mutants	Burkhard Küchler, Andy Nandy, Abdel Ghany and Sandro Ghisla
O09	Probing the active site of the medium chain acyl-CoA dehydrogenase: 4-OH-cinnamoyl-CoA as a sensitive probe of polarization and ionization	Irina Rudik, Alasdair Bell, Peter Tonge and Colin Thorpe
O10	Substrate polarization of medium chain acyl-CoA dehydrogenase (MCAD)	Avery Stephens, Kim M. Sabaj, Peter J. Tonge and Marian T. Stankovich