FIFTH INTERNATIONAL SEMINAR ON

ENERGY TRANSFER IN CONDENSED MATTER

Structure, Conformation and Function of Molecular Systems

Largely Related to Photosynthesis

Proceedings

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CONTENTS

OPENING ADDRESS K. Vacek	. 1
Invited Lectures	
STRUCTURE-FUNCTION RELATIONSHIP IN PHOTOSYNTHETIC SYSTEMS IN VIVO	
REGULATION OF ENERGY TRANSFER BY MEMBRANE CONFORMATIONAL	
CHANGES INDUCED BY PROTEIN PHOSPHORYLATION J. Barber FLUORESCENCE KINETICS OF C-PHYCOCYANIN	. 5
FROM Mastigocladus laminosus DEPENDING ON ITS STATE	
OF AGGREGATION H. Scheer and S. Schneider	. 15
PHOTOINACTIVATION OF PHOTOSYSTEM II: ITS INHERENT INSTABILITY	
OR REGULATION OF ENERGY TRANSFER? I. Setlik	. 23
NONLINEAR ABSORPTION OF CHLOROPHYLL in vivo	
P. Hoffmann and D. Leupold	. 32
SPECTROSCOPY AND STRUCTURE-FUNCTION RELATIONSHIP IN PHOTOSYNTHETIC SYSTEMS	
ORIENTATION AND ASSIGNEMENT OF THE CHROMOPHORES IN REACTION	
CENTERS OF Rhodopseudomonas viridis AND Rhodopseudomonas	
sphaeroides J. Breton	40
PHOTOACOUSTIC AND FLUORESCENCE SPECTRA OF ORIENTED	
CHLOROPLASTS L. Lorrain, D. Frackowiak, R. M. Leblanc	42
ENERGY TRANSFER BETWEEN DIMERIC CHLOROPHYLL a AND ALL TRANS	
β-CAROTENE in vitro AS RESOLVED BY FLUORESCENCE PHOTOQUENCHING	
N. E. Binnie, L. V. Haley and J. A. Koningstein	44
PIGMENT STATES AND ENVIRONMENTS IN PHOTOSYNTHETIC STRUCTURES	F 2
FROM THEIR RESONANCE RAMAN SCATTERING M. Lutz and B. Robert	52
FLUORESCENCE DETECTED MAGNETIC RESONANCE STUDIES ON PHOTOSYNTHETIC BACTERIA A. Angerhofer, J. U. von Schütz and H. C. Wolf	59
BACIERIA A. Angernojer, J. J. John Schulz und H. C. Wolj	33
ENERGY TRANSFER IN PRIMARY PHOTOSYNTHETIC PROCESSES - THEORETICAL APPROACHES	
ENERGY TRANSFER IN PRIMARY PHOTOSYNTHETIC PROCESSES - SOME	
RECENT RESULTS AND APPROACHES L. Skála	67
MODELLING OF ENERGY TRANSFER IN PHOTOSYNTHETIC	
SYSTEMS V. Čápek, V. Szöcs and L. Nedbal	76
THEORETICAL ASPECTS OF STRUCTURE-FUNCTION RELATIONSHIP	
IN PRIMARY PHOTOSYNTHETIC PROCESSES F C Patron	Ω1

PHOTOSYNTHETIC MODEL SYSTEMS

SOME ASPECTS OF CHLOROPHYLL - PROTEIN INTERACTIONS AS STUDIED
BY PORPHYRINE - PEPTIDE MODEL SYSTEMS
P. Pančoška, P. Maloň and K. Bláha 92
SYNTHETIC MODELLING APPROACHES TO PHOTOSYNTHETIC
REACTION CENTER STRUCTURE AND FUNCTION M. R. Wasielewski,
M. P. Niemczyk, W. A. Svec and E. B. Pewitt 101
ELECTRON AND ENERGY TRANSFER IN REAL AND ARTIFICIAL
PHOTOSYNTHETIC SYSTEMS S. G. Boxer, A. Kuki,
D. J. Lockhart, T. R. Middendorf and R. Moog 108
MODEL FOR P680: ONE-ELECTRON WATER OXIDATION BY HYDRATED
CHLOROPHYLL a DIMER RADICAL CATION D. J. Diestler,
M. S. Showell, O. Riveros and F. K. Fong
PHOTOPHYSICAL PROPERTIES OF METALLOPORPHYRINS
R. L. Brookfield, H. Ellu? and A. Harriman 124
NEW SPECTROSCOPIC TECHNIQUES APPLIED TO THE PHOTOSYNTHETIC SYSTEMS
SITE SELECTION SPECTROSCOPY OF MODEL PHOTOSYNTHETIC
SYSTEMS J. Hála
FINE-STRUCTURED SPECTRA OF CHLOROPHYLL UNDER THE CONDITIONS
OF FAST ENERGY TRANSFER R. Avarmaa and K. Rebane
DYNAMIC PROPERTIES OF TRIPLET EXCITONS IN MOLECULAR
CRYSTALS STUDIED BY ELECTRON SPIN ECHO AND PULSED
LASER TECHNIQUES J. Schmidt
MAGNETIC INTERACTIONS IN BACTERIAL PHOTOSYNTHETIC REACTION
CENTERS A. J. Hoff
OPTICAL SPECTROSCOPY AND MAGNETIC RESONANCE
OF PHOTOSYNTHETIC PARTICLES AND MODEL SYSTEMS
T. J. Schaafsma, G. F. W. Searle, C. Dijkema,
L. Nedbal, A. van Hoek, P. Gast and F. G. H. van Wijk 163
To a Constitution
Poster Sections
SYSTEMS IN VIVO
ORGANIZATION OF CHLOROPHYLL IN CHLOROSOMES OF GREEN
SULFUR BACTERIA J. M. Olson, P. D. Gerola, J. Mikkelsen,
J. Knudsen and P. Højrup 17.
ORIENTATION OF PHEOPHYTIN AND P680, THE PRIMARY ELECTRON
ACCEPTOR AND DONOR IN PHOTOSYSTEM II I. Ganago, V. Klimov,
A. Ganago, V. Shuvalov, Yu. Erokhin

POLARIZED ABSORPTION, FLUORESCENCE AND FLUORESCENCE LIFETIME	
OF PHYCOBILISOMES IN PVA FILMS D. Frackowiak, L. G. Erokhina,	
A. Balter, J. Szurkowski and B. Szych	180
PHOTOCHEMICAL ACTIVITES OF CHLOROPLASTS AFTER DETERGENT	
TREATMENT E. Apostolova, D. Kafalieva	183
LIGHT-INDUCED CONFORMATIONAL CHANGES IN CHLOROPLAST	
MEMBRANE PROTEINS E. Apostolova, D. Kafalieva	185
ANALYSIS OF THE POLYPEPTIDE COMPOSITION OF VARIOUS	
PHOTOSYNTETIC MEMBRANES BY TWO-DIMENSIONAL GEL-ELECTROPHORESIS	
J. Masojídek, M. Droppa and G. Horváth	187
PECULIARITIES OF THE ENERGY TRANSFER FROM THE CAROTENOIDS	
AND CHLOROPYHLL b TO THE PHOTOSYSTEMS II AND I	
J. Voigt and U. Stahl	190
EVIDENCE OF NONLINEAR COHERENT EFFECTS CONNECTED WITH THE	
ENERGY TRANSFER IN PHOTOSYNTHETIC MEMBRANES	
G. Kehrberg and J. Voigt	192
LIFETIME AND YIELD OF VARIABLE FLUORESCENCE AND THE ENERGY	
TRANSFER BETWEEN THE LHC AND THE PHOTOSYSTEMS M. Senoner	194
SOME ASPECTS OF TEMPERATURE DEPENDENCES OF in vivo CHLOROPHYLL a	
FLUORESCENCE STUDIED IN DIFFERENT MEDIA J. Nauš, L. Dvořák,	
M. Vychodilová, Z. Kupka	196
STUDY OF THE SLOW CHLOROPHYLL a FLUORESCENCE CHANGES IN INTACT	
ISOLATED CHLOROPLASTS INDUCED BY MEDIATOR OF CYCLIC PHOTO-	
PHOSPHORYLATION J. Fejzo, M. Plesničar, Lj. Kolar-Anić	198
STRUCTURAL AND P700 ⁺ HEAT INDUCED CHANGES OF PEA CHLOROPLAST	
MEMBRANES: FLUORESCENT AND ESR STUDIES A. G. Ivanov,	
M. Velitchkova and D. Kafalieva	200
CHANGES IN RELATIVE OPTICAL CROSS-SECTION OF THE TWO PHOTOSYSTEMS	
DURING THE LIFE CYCLE OF chlorella P. Butko	202
ULTRAFAST FLUORESCENCE INDUCTION PHASE AND PRIMARY REACTIONS	
IN PHOTOSYSTEM II OF CHLOROPLASTS J. Deprez, A. Dobek,	
N. E. Geacintov, G. Paillotin and J. Breton	204
EXCITATION ENERGY TRANSFORMATIONS IN BACTERIAL PHOTOSYNTHESIS	
STUDIES BY PICOSECOND FLUORESCENCE OF INTACT CELLS AND	
ISOLATED REACTION CENTERS A. Freiberg, V. Godik,	
S. Kharchenko, K. Timpmann	211
TRIPLET FORMATION IN A PHOTOSYNTHETIC THREE-SPIN SYSTEM:	
THE REACTION CENTER FROM Rhodopseudomonas viridis	
F. G. H. van Wijk, P. Gast and T. J. Schaafsma	213
THE SUBUNIT STRUCTURE AND THE AMINO ACID COMPOSITION	
OF MAIZE PHOSPHOENOLPYRUVATE CARBOXYLASE	
M Stihonová and S Lehlová	215

THEORETICAL APPROACHES

SIMPLE THEORETICAL MODEL OF EXCITATION ENERGY TRANSFER	
IN PRIMARY STAGE OF PHOTOSYNTHESIS V. Kapsa,	
O. Bílek, L. Skála and P. Pančoška	217
MAGNETIC FIELD EFFECTS ON THE CHARGE RECOMBINATION REACTION	
YIELDS IN PHOTOSYNTHETIC REACTION CENTERS	
A. A. Demidenko, E. G. Petrov	219
ELECTRON TRANSFER IN THE INITIAL PHOTOSYNTHESIS STAGES:	
THE ROLE OF CONFORMATION STATES	
V. V. Gorbach, E. G. Petrov	221
THEORETICAL INVESTIGATION OF LASER INDUCED PRIMARY	
PROCESSES OF PHOTOSYNTHESIS 5. Kudžmauskas,	
G. Trinkunas and L. Valkunas 2	223
DENSITY MATRIX FORMULATION OF UNIMOLECULAR PROTON TRANSFER	
REACTION P. Bañacký and Ľ. Vodná	225
PHOTOSYNTHETIC MODEL SYSTEMS	
TIME-DECAY FLUORESCENCE STUDIES OF CHLOROPHYLL a	
IN DIFFERENT SOLVENTS R. Vladkova, D. Kafalieva,	227
V. Kolev and V. Spasov	221
G. D. Olovianishnikova, N. A. Sadevnikova	220
EXCITON DELOCALIZATION ON THE EXCITED SINGLET STATE	. 2
OF CHLOROPHYLL a MONOHYDRATE DIMER AND FLUORESCENCE	
LIFETIMES OF TRANSLATIONALLY EQUIVALENT AGGREGATES	
F. K. Fong, M. S. Showell and A. J. Alfano	221
MONOHYDRATE DIMER MODEL AND METAMERIC TRANSITION IN P700	: J I
PHOTOOXIDATION F. K. Fong and M. S. Showell	? ? ?
DIHYDRATE CHLOROPHYLL DIMER AS REACTION CENTER	
FOR WATER PHOTOLYSIS F. K. Fong, M. S. Showell,	
J. L. You and A. J. Gotch	235
TRANSIENT ABSORPTION OF CHLOROPHYLL a - WATER COMPLEXES	
P. Malý, R. Danielius and R. Gadonas	237
In vitro AGGREGATES OF BCHL c MODELS FOR THE CHLOROSOME	
PROTEIN COMPLEX OF GREEN SULFUR BACTERIA	
G. H. van Brakel and J. M. Olson	239
AGGREGATIONS STATES OF CAROTENOIDS IN BINARY SOLVENTS	
AND IN LYOTROPICS V. D. Kolev	243
MOLECULAR ARRANGEMENTS OF PROTOCHLOROPHYLL FORMS	
B. Böddi and F. Láng	245

SITE SELECTION SPECTRA OF TETRAPHENYLPORPHYRINAMINOACID
SYSTEMS J. Hála, P. Douša, M. Ambrož, I. Pelant
and K. Vacek
INTERACTION BETWEEN CHLOROPHYLL a AND CHLOROPHYLL b IN
NEMATIC LIQUID CRYSTALS J. Szurkowski and D. Frackowiak 251
ENERGY TRANSFER BETWEEN CHLOROPHYLLS a AND b ON PLASTICIZED
POLYETHYLENE PARTICLES G. R. Seely and V. Senthilathipan 253
PHOTOACOUSTIC AND OPTICAL SPECTRA OF CHLOROPHYLL a IN POLY-
ISOBUTYLENE MATRIX M. Kaplanová and M. Dienstbier 255
PHOTOELECTRIC PHENOMENA IN THE CHLOROPHYLL-CONTAINING
MEMBRANE FILTERS Yu. M. Stolovitsky, S. I. Kadoshnikov,
E. Vavřinec, K. Vacek
PHOTOELECTROCHEMICAL AND SPECTRAL PROPERTIES OF CHLOROPHYLL
IN ARTIFICIAL MEMBRANES AND LIPOSOMES S. I. Kadoshnikov,
Yu. M. Stolovitsky
ELECTROCHEMICAL REGULATION OF ELECTRON TRANSFER PREDICTED
BY A MODEL OF CHLOROPHYLL-PROTEIN COMPLEXES? E. Balint,
E. Tombácz, Z. Várkonyi, E. Szalai, J. Hevesi 261
RELATED SYSTEMS
DEVIATIONS IN FÖRSTER-LIKE ENERGY TRANSFER BETWEEN DYE
MOLECULES IN SOLUTION - EVIDENCE FOR AN
INHOMOGENEOUS SPATIAL DISTRIBUTION OF MOLECULES
M. Kaschke, K. Vogler
THERMOSTIMULATED LUMINESCENCE OF CdF ₂ : Eu CRYSTALS A. A. Nagornyi, J. Pospíšil
THE CHARGE CARRIER TRANSPORT STUDY IN LAYERS OF
POLY(N-VINYLCARBAZOLE) AT LOW TEMPERATURE
USING THERMOSTIMULATED LUMINESCENCE
I. Chudáček, J. Pospíšil, I. A. Tale
INDEX OF AUTHORS 271

FLUORESCENCE KINETICS OF C-PHYCOCYANIN FROM MASTIGOCLAUDUS LAMINOSUS DEPENDING ON ITS STATE OF AGGREGATION

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Photosynthetic organisms cover most of their energy needs with sunlight. They have consequently developed a variety of adaption mechanisms to compete efficiently for it. In higher plants, a dominant mechanism is the growth towards the light. Aquatic and microorganisms adapt commonly by chromatic adaptation of the photosynthetic antenna. The Chlorophylls <u>a</u> and <u>b</u> are rather inefficient in collecting green light, and several additional pigment systems have evolved to fill this hole in the action spectrum.

The phycobiliproteins are one such group of antenna pigments. They are used in cyanobacteria, red alga and cryptophytes (1). In the former two, they are highly aggregated together with colorless linker polypeptides, in the phycobilisomes. These are microscopic structures situated at the outer surface of the thylakoid membrane, which transfer their excitation energy efficiently (quantum yield 295%) to the chlorophyllous reaction centers within the membrane. The phycobilisomes show a remarkable structure. They consist of a central core of allophycocyanin (APC) to which rods of phycocyanin (PC) and phycocrythrin (PE) are attached. This morphology matches the energetic ordering of the chromophores required for an energy transfer chain from PE via PC and APC to the chlorophylls and reaction centers.

This report deals with one major pigment of the phycobilisomes, e.g. PC from two cyanobacteria (<u>Spirulina platensis</u> and <u>Mastigocladus</u> <u>laminosus</u>). Three subjects shall be considered: The molecular structure and covalent protein linkages of the chromophores, the <u>in situ</u> structures of the chromophores as determined by chromophore-protein interactions, and the fluorescence kinetics within PC aggegates of increasing complexity.

Chromophore structure

PC contains like the other phycobiliproteins linear tetrapyrrolic chromophores. They are distinct from the mammalian type bile pigments, e.g. biliverdin, by the presence of one hydrogenated ring (generally ring A) and by covalent linkages to the apoprotein. The thioether linkage shown in fig.1 is well established. Additional linkages have been discussed (see references in 1) which are less stable under the conditions hitherto used for structure elucidation, e.g. chromic acid degradation and proteolysis.

We have recently investigated Ehrlich's diazo-reaction (2) as the key reaction for a milder degradation method. In a three-step sequence (3). PC is first denatured, then reduced with sodium borohydride, and the resulting "phycorubin" finally treated with aromatic diazonium salts. The chromophores are thereby split into halves at the central methylene bridge. Depending on the presence or absence of a covalent linkage, the resulting two azo-dipyrromethenones remain either attached to the peptide chain, or become free and can be separated from the former by solvent extraction or gel chromatography. The sequence has two advantages: Information is gained on the C-5 and C-15 methine substituents (which is lost during chromic acid degradation), and the entire reaction sequence can be carried out below 4°C and at neutral pH. A cleavage of labile bonds, or the formation of new bonds is therefore precluded. A quantitative study has now been performed with PC from the two aforementioned species and their subunits. It was found, that there is always only that one dipyrromethene chromophore attached covalently to the protein, which contains the hydrogenated ring and hence the thioether bond. The fully unsaturated dipyrromethenone chromophore corresponding to the other half of the original PC chromophores was free, and it was identical with the respective ring C,D-fragment of the diazo-product of mesobilirubin. The structure of the PC chromophores is thus the one shown in fig.1. Rapoport and Glazer (4) have reported after the completion of this work, that one of the three chromophores in PC from Synechococcus 6301 is bound to the protein and hydrogenated at ring D. We did not find the expected azo-product in the products derived from S. platensis or M. laminosus, but the sensitivity may not have been sufficient for the detection of one out of three chromophores.

Native state of the chromophores

The spectra and most other properties of PC are quite unusual for bile pigments of the structure shown in fig.1. Free chromophores of this type have only moderate absorptivity in the visible spectral range, and their radiative lifetimes are short (20-70psec for bilipeptides depending on binding sites and isolation procedure). Both properties render them poor candidates for antenna pigments from the biophysycists point of view. These and other properties detrimental to photosynthesis, are greatly improved in phycocyanin. The native pigment has an absorptivity which is almost one order of magnitude larger, and it has a high fluorescence quantum yield (radiative lifetimes \$1.5nsec). Since these functionally important properties are reversibly abolished by denaturation of the protein, e.g. with 8M urea, they must be entirely due to non-covalent chromophore-protein interactions.

A systematic survey of bile pigments structurally related to I and the effect of various treatments of the chromophores (pH, complexation) on their properties has led to a model (fig.2), in which the chromophores are rigidly fixed in an extended conformation (1). Free bile pigment chromophores are by contrast rather flexible and have predominantly cyclic conformations. The energetically unfavorable conformation of the native PC chromophores, which is essential to their function as light harvesting pigments, has recently been substantiated by the high-resolution x-ray structure of PC from M. laminosus (5).

The picture is further complicated by the fact that PC contains three chromophores at two different subunits. The α -subunit carries one, the β -subunit 2 chromophores. Two of them (α and β 1) are at binding sites with similar primary and tertiary structure (5), the third one is distinct with regard to its binding site. It is also possible to distinguish three distinct chromophores spectroscopically from the absorption and circular dichroism properties of PC subunits. The isolated α -subunit absorbs maximally in the red at 618 nm, and a similar component is present in integral PC (fig.3). The β -subunit absorbs at 606 nm, and two chromophore absorptions (\approx 585 and 615nm) can be resolved by circular dichroism spectroscopy and by fluorescence polarisation (6). The cd-anisotropy of the β -subunit is only 60% of the α -subunit, and its visible maximum shifted by 15 nm to the blue (fig.4). Thus it contains one cd-active, twisted chromophore absorbing at the blue side of the unstructured absorption band, and one cd

Fig.1: Molecular structure of phycocyanin chromophore drawn in the cyclic-helical conformation characteristic for free bile pigments.

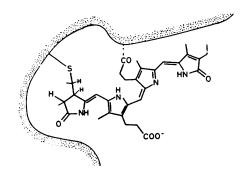


Fig.2: Native conformation of phycocyanin chromophore.
Schematic representation of an extended conformation which is suggested from spectroscopic data.

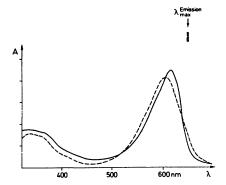


Fig.4: Circular dichroism spectra of subunits of phycocyanin from Mastigocladus laminosus

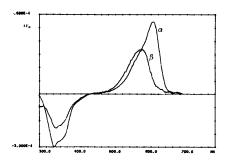


Fig.3: Absorption spectra of subunits of phycocyanin from Mastigocladus laminosus

inactive (planar?) chromophore absorbing at the red side of the absorption maximum. However, a correlation between these two absorption bands of the 8-chromophore and the known binding sites is presently not yet possible.

Fluorescence Kinetics

In order to elucidate the fluorescence Kinetics of PC and its implications on the energy transfer, we have begun to vary systematically the complexity of PC aggregates and then studied their polarized time resolved fluorescence (7-9). Five different aggregation states have been investigated: The α - and β -subunits in their monomeric states, the $(\alpha\beta)$ heterodimer (= monomer), the $(\alpha\beta)$ heterohexamer (= trimer) and entire phycobilisomes. In each case, the pigments were excited with a tunable mode-locked dye laser (photon flux at sample ≤1018photons cm 1 pulse 1, pulse width ≈1psec fwhm, 80 MHz repetition rate). The polarized, time-resolved fluorescences decay curves have been recorded with a synchroscan streak camera. From the decay traces with the emission polarizing filter being oriented parallel (I \rightarrow) or perpendicular (I \rightarrow) to the polarisation filter of the exciting light, the isotropic fluorescence (I + 2I) and the anisotropic fluorescence (expressed as difference function I - I, see ref.9) have been caculated.

All decay curves were originally fit under the assumption of a biexponential decay function. Under idential conditions (\$\lambda\$, exc. = 600nm, \$\lambda\$, em. \$\lambda\$620nm), the isotropic decay rates of the subunits, the monomer and the trimer were similar, but the decay rate is sharply increased in the phycobilisomes due to the presence of an acceptor, e.g. APC. The high rate of transfer to acceptors absorbing at longer wavelengths can be seen directly under conditions conditions of short-wavelength (\$590nm) excitation and long-wavelength detection (\$2620nm) (see below and fig.5). It necessitates the introduction of an additional rise-term of the fluorescence. In the anisotropic decay of the increasingly complex aggregates, there is a pronounced increase in the depolarisation rate between monomer and trimer. A second increase is seen between trimer and phycobilisomes, for which the anisotropic decay was beyond the detection limit of the system.

Since rotational depolarization of the chromophores fixed to the proteins (molecular weights bettween 16 and several 188 kDa) is

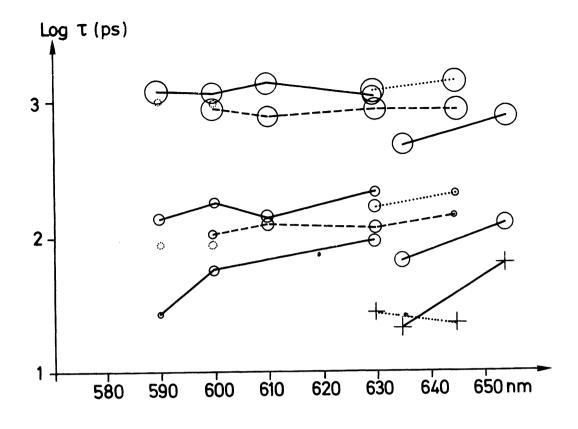


Fig.5: Isotropic fluorescence decay of phycobilisomes from Matiqocladus
Iaminosus. Characteristic emission wavelengths (selected by cut-off filter below 630nm, interference filter at longer wavelengths) are given on the abcissa. Excitation wavelengths were at 580nm (----), 590nm (-----) and 610nm (.....). The circles correspond to decay terms, the crosses to rise terms. The area of the circles is proportional to the intergrated intrensities of the respective decaying components. All experimental data were fit under the assumption of a tri-exponential decay law. If bi-exponentials gave an equally good fit, the data are given as dotted circles (590 and 600nm emission, excitation at 580nm).

negligible on the observation time scale, the depolarisation has been interpreted as being due to energy transfer. The increased depolarisation rate in the trimer is rationalized as a result of an increased number of possible acceptors and by the fact, that the closest chromophore-chromophore distance in the trimer does not occur within a monomeric unit, but rather between β - and α -chromophores located on adjacent monomers (5). The absence of a polarized component in the fluorescence of the phycobilisome is then due to an even increased number of acceptors, i.e. essentially no leakage fluorescence from directly excited chromophores is observed. (No data are hitherto available for inter-chromophore distances between trimeric units in the phycobilisome rods.)

Although most decays can be fit well under the assumption of a bi-exponential decay law, more than two components are sometimes necessary. This is in particular true under conditions were rise-terms are encountered, e.g. if the differences between excitation and detection wavelengths (AL) become substantial (fig. 5). If a series of distinct energy transfer steps is assumed to occur within a given assembly, a set of constant decay times with varying amplitudes and signs is expected for any selected pairs of excitation and emission wavelengths. An analysis allowing for three exponentials in the fluorescence of phycobilisomes does indeed show a nearly constant rate-constant for the long-lived component, which is probably mainly due to APC (fig.5). However, the short term(s) are variable and increase with increasing AD. This indicates, that more than one time constant may be hidden under these terms, whose relative amplitudes change with the probing wavelengths. Experiments with phycobilisomes from a different species have required a fit with four exponentials, whose rate constants are furthermore depudent on the length of the rods (10).

A high complexity of the energy transfer even within isolated phycobiliproteins is indicated by recent results of Holzwarth et al. (11) obtained by single-photon timing techniques. The fluorescence decay data for trimeric aggregates of PC were subjected to a global analysis. It was necessary to use three or, better, even four components, and in particular the fast decaying components were also dependent on the presence or absence of colorless linker polypeptides. This indicates, that even in a relatively simple system, there is already a quite complex energy transfer pattern. It is, however, presently not yet clear which of the energy transfer steps seen in isolated biliproteins are relevant to those observed in integral

- 21 -

phycobilisomes. More work is needed to relate these processes to each other.

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