

# Methods and Applications in Fluorescence



## PAPER

### Novel benzanthrone probes for membrane and protein studies

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**Keywords:** benzanthrone dyes, lipid membranes, fibrillar protein aggregates, polarity, viscosity

#### Abstract

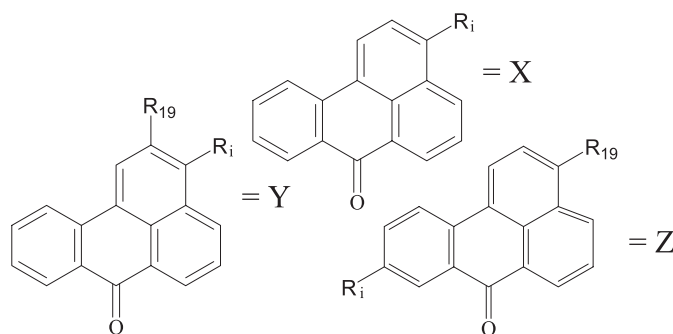
The applicability of a series of novel benzanthrone dyes to monitoring the changes in physicochemical properties of lipid bilayer and to differentiating between the native and aggregated protein states has been evaluated. Based on the quantitative parameters of the dye-membrane and dye-protein binding derived from the fluorimetric titration data, the most prospective membrane probes and amyloid tracers have been selected from the group of examined compounds. Analysis of the red edge excitation shifts of the membrane- and amyloid-bound dyes provided information on the properties of benzanthrone binding sites within the lipid and protein matrixes. To understand how amyloid specificity of benzantrones correlates with their structure, quantitative structure activity relationship (QSAR) analysis was performed involving a range of quantum chemical molecular descriptors. A statistically significant model was obtained for predicting the sensitivity of novel benzanthrone dyes to amyloid fibrils.

## 1. Introduction

Benzanthrone dyes, a well known class of fluorescent probes that emit in the spectral region from yellow-green to red-purple (depending on the dye structure), have found diverse applications in a variety of fields as the components of liquid-crystal systems for electro-optic displays of the 'guest-host' type [1, 2], disperse dyes for textiles [3] and polymers [4, 5], laser dyes [6], etc. Likewise, these compounds are of great interest for biomedical research and diagnostics due to their favorable spectral characteristics, such as large extinction coefficient, marked Stokes shift; negligible fluorescence in an aqueous phase, ability to intramolecular charge transfer (ICT) within chromophoric system, occurring between electron-donating groups in C-3 position and electron-accepting carbonyl group, etc [7-9]. The ability of benzanthrone fluorophores to form ICT state gives rise to significant increase of the dye dipole moment after excitation, thereby inducing the reorientation of solvent dipoles around the excited-state dipole. This property renders these dyes highly suitable for studying the membrane-

related processes, especially those coupled with the changes in polarity and viscosity of lipid bilayer [10, 11]. In the early 1970s Dobretsov and Vladimirov have demonstrated that one representative of benzantrones 3-methoxybenzanthrone (MBA) can be effectively used to monitor the membrane structural changes caused by the shifts in cholesterol (Chol) level, temperature, pH, etc [12]. Subsequently, Yang *et al* have employed this probe as DNA intercalator [13]. Later it has been shown that other benzanthrone derivatives possess marked affinity for cellular (lymphocyte) membranes, responding to the alterations in immune status of a human organism at different pathologies [14, 15].

Our previous investigations revealed that benzanthrone dyes are capable of distinguishing between the oligomeric and fibrillar aggregates of lysozyme [16, 17]. Specifically, cumulative data from the binding and red edge excitation shift studies of several novel benzantrones allowed us to define the parameters reflecting the dye preference to either pre-fibrillar or amyloid protein aggregates [17]. Moreover, it has been shown that benzanthrone dyes are sensitive to the distinctions in fibril morphology [18]. In our recent works we also



**Figure 1.** Structure of the examined benzanthrone dyes. The dye names and the side groups are given in table 1.

evaluated the lipid-associating ability of benzantrones and found that their spectral responses correlate with the changes in bilayer hydration [19–21].

In the present work we extended our previous investigation of benzanthrone dyes and focused our efforts on evaluating the possibility of using newly synthesized fluorophores for probing the structural features of lipid bilayers and aggregating proteins. More specifically, our goal was several-fold: (i) to assess benzanthrone sensitivity to the changes in physicochemical characteristics of the model membranes composed of zwitterionic lipid phosphatidylcholine (PC) and its mixtures with Chol and anionic phospholipids cardiolipin (CL) and phosphatidylglycerol (PG) in different molar ratios; (ii) to assess the applicability of benzantrones as amyloid markers; (iii) to characterize the properties of the dye binding sites within lipid or protein matrix through analyzing the red edge excitation shifts (REES); (iv) to ascertain the factors determining the lipid- and protein-associating abilities of benzantrones.

## 2. Materials and methods

### 2.1. Materials

Chicken egg white lysozyme (HEWL) and Chol were purchased from Sigma (St. Louis, MO, USA). Bovine heart CL, 1-palmitoyl-2-oleoyl-*sn*-glycero-3-phosphocholine (PC) and 1-palmitoyl-2-oleoyl-*sn*-glycero-3-phospho-*rac*-glycerol (PG) were from Avanti Polar Lipids (Alabaster, AL). Benzanthrone dyes (figure 1 and table 1) were synthesized at the Faculty of Natural Sciences and Mathematics of Daugavpils University, Latvia as described in detail elsewhere [7, 22]. All other chemicals were of analytical grade and used without further purification.

### 2.2. Preparation of lipid vesicles

Lipid vesicles composed of PC and PC mixtures with CL, PG or Chol were prepared using the extrusion technique. A thin lipid film was first formed of the lipid mixtures in chloroform by removing the solvent under a stream of nitrogen. The dry lipid residues were subsequently hydrated with 20 mM HEPES, 0.1 mM EDTA, pH 7.4 at room temperature to yield lipid

concentration 1 mM. Thereafter, lipid suspension was extruded through a 100 nm pore size polycarbonate filter (Millipore, Bedford, USA). In this way, 10 types of lipid vesicles containing PC and 5.3, 11.1 or 25 mol% CL, 10, 20 or 40 mol% PG and 10, 20 or 30 mol% Chol, with the content of phosphate being identical for all liposome preparations. Hereafter, liposomes containing 10, 20 or 40 mol% PG are referred to as PG10, PG20 or PG40, respectively, while liposomes bearing 5.3, 11.1 or 25 mol% CL are denoted as CL5, CL11 or CL25, respectively. Accordingly, liposomes with Chol content 10, 20 or 30 mol% are marked as Chol10, Chol20 and Chol30, respectively.

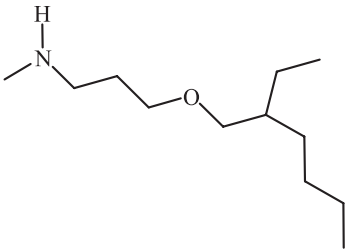
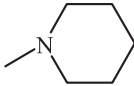
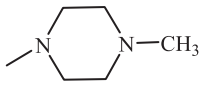
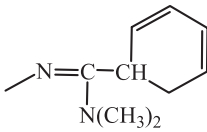
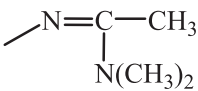
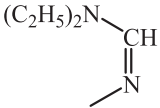
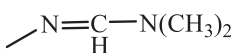
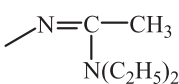
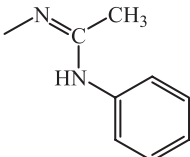
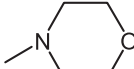
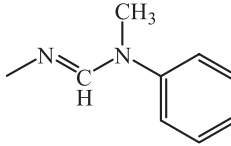
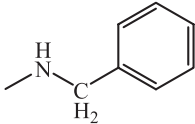
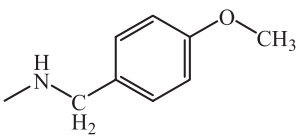
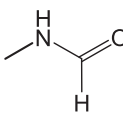
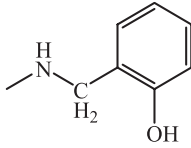
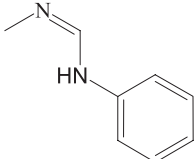
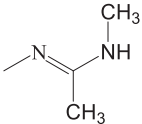
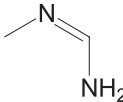
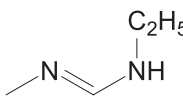
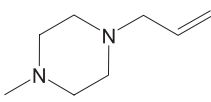
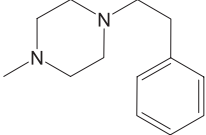

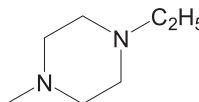
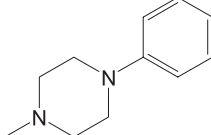
### 2.3. Preparation of lysozyme fibrils

The reaction of lysozyme fibrillization was conducted at 60 °C by continuous shaking of lysozyme solution (20 mg ml<sup>-1</sup>) in 10 mM glycine buffer (pH 1.6) during 19 d [23, 24]. The obtained aggregates, below referred to as  $F_{acs}$ , were used for the screening of the most prospective amyloid tracers of a benzanthrone series. For the investigation of the dye sensitivity to the distinctions in fibril morphology we used other two types of lysozyme fibrils: (i) aggregates ( $F_{ac}$ ), grown by the protein (10 mg ml<sup>-1</sup>) incubation without shaking in 10 mM glycine buffer (pH 2.2) at 60 °C during 6 d [18]; and fibrils ( $F_{eth}$ ) prepared by dissolving the protein (3 mg ml<sup>-1</sup>) in deionized water with subsequent slow addition of ethanol to a final concentration 80%, followed by a constant agitation of a sample at 25 °C during 30 d [18].

### 2.4. Steady-state fluorescence measurements

Steady-state fluorescence spectra were recorded with LS-55 (Perkin-Elmer Ltd, Beaconsfield, UK) or Varian Cary Eclipse (Varian Instruments, Walnut Creek, CA) spectrofluorimeters equipped with a magnetically stirred, thermostated cuvette holder. Fluorescence measurements were performed at 20 °C using 10 mm path-length quartz cuvettes. The excitation and emission slit widths were set at 10 nm. Excitation wavelengths and emission maxima of the examined dyes are given in table 2. Quantum yields of the examined dyes were calculated using rhodamine 101 as a standard according to the formula:

Table 1. Chemical structures of novel benzanthrone dyes.

Dye	Structure	Dye	Structure	Dye	Structure		
A4	X-R1	A6	X-R2	A8	X-R3		
AM1	X-R4	AM2	X-R5	AM15	X-R6		
AM3	X-R7	AM4	X-R8	AM12	X-R9		
ABM	X-R10	AM18	X-R11	IBH	X-R12		
IAH	X-R13	FA	X-R14	ISH	X-R15		
AM17	X-R16	AM20	X-R17	AM19	X-R18		
AM21	X-R19	P11	X-R20	P14	X-R21		
AM2-23	Y-R5	AM4-23	Y-R8	AM15-23	Y-R6		
AM18	Y-R11	3AM39	Z-R7	P9	X-R23		
P10	X-R24						
R1		R2		R3		R4	
R5		R6		R7		R8	
R9		R10		R11		R12	
R13		R14		R15		R16	
R17		R18		R19		R20	
R21		R22		R23		R24	

**Table 2.** The basic photophysical characteristics, partition coefficients and quantum yields of benzanthrone dyes in different systems.

Dye	Ethanol				PC			CL5		CL10		Chol30	
	$\lambda_A$ , nm	$\lambda_F$ , nm	$\lg \varepsilon$	$Q_0$	$Q$	$K_p \times 10^4$	$Q$	$K_p \times 10^4$	$Q$	$K_p \times 10^4$	$Q$	$K_p \times 10^4$	
AM1	470	649	4.27	0.005	0.06	$2.7 \pm 0.2$	0.06	$3.0 \pm 0.3$	0.05	$2.8 \pm 0.3$	0.07	$3.1 \pm 0.1$	
AM2*	470	656	4.2	0.003	0.07	$0.4 \pm 0.1$	0.05	$0.5 \pm 0.1$	0.07	$0.7 \pm 0.05$	0.06	$0.5 \pm 0.04$	
AM3*	469	636	4.07	0.013	0.44	$0.8 \pm 0.2$	0.41	$0.8 \pm 0.04$	0.45	$0.78 \pm 0.01$	0.46	$0.84 \pm 0.07$	
AM4*	472	657	4.19	0.002	0.38	$1.3 \pm 0.2$	0.42	$1.4 \pm 0.15$	0.41	$1.1 \pm 0.17$	0.43	$0.97 \pm 0.12$	
ABM	444	644	4.33	0.008	0.27	$6.4 \pm 0.7$	0.32	$8.2 \pm 0.4$	0.5	$10.3 \pm 0.9$	0.36	$6.9 \pm 0.3$	
A6	455	654	4.01	0.018	0.44	$11 \pm 1.5$	0.52	$9.2 \pm 0.9$	0.6	$8.4 \pm 0.7$	0.49	$8.7 \pm 0.5$	
A8*	444	637	3.93	0.015	0.6	$0.9 \pm 0.07$	0.5	$3.4 \pm 0.05$	0.48	$6.2 \pm 1.6$	0.49	$1.7 \pm 0.27$	
A4	530	651	4.19	0.0008	0.1	$1.0 \pm 0.04$	0.09	$1.1 \pm 0.1$	0.09	$0.9 \pm 0.1$	0.23	$0.9 \pm 0.01$	
AM12*	469	628	3.85	0.009	0.6	$18 \pm 0.8$	0.5	$8.7 \pm 0.3$	0.3	$5.8 \pm 0.6$	0.24	$15 \pm 1.5$	
AM15*	475	632	4.15	0.005	0.6	$6.1 \pm 0.2$	0.47	$2.7 \pm 0.2$	0.36	$1.6 \pm 0.05$	0.3	$4.9 \pm 0.4$	
AM18*	465	618	3.53	0.004	0.7	$12 \pm 0.8$	0.4	$9.7 \pm 0.6$	0.2	$7.7 \pm 0.5$	0.5	$7.8 \pm 0.5$	
IAH*	524	644	3.6	0.002	0.06	$4.8 \pm 0.5$	0.05	$7.6 \pm 0.5$	0.04	$5.4 \pm 0.1$	0.04	$6.8 \pm 0.8$	
IBH*	531	642	3.8	0.001	0.13	$6.0 \pm 0.4$	0.06	$11 \pm 0.6$	0.03	$11 \pm 0.4$	0.02	$9.8 \pm 0.7$	
ISH*	529	644	2.82	0.001	0.09	$6.1 \pm 0.5$	0.08	$8.8 \pm 0.4$	0.07	$7.0 \pm 0.4$	0.1	$6.2 \pm 0.2$	
3AM39	474	645	4.03	0.009	0.4	$2.5 \pm 0.1$	0.3	$5.5 \pm 0.2$	0.4	$9.0 \pm 0.4$	0.6	$3.4 \pm 0.1$	
AM2-23	451	638	4.02	0.004	0.5	$5.1 \pm 0.2$	0.4	$7.6 \pm 0.2$	0.3	$9.8 \pm 0.7$	0.4	$8.8 \pm 0.4$	
AM4-23	462	642	4.06	0.007	0.3	$4.1 \pm 0.9$	0.27	$10 \pm 1.7$	0.23	$11.4 \pm 0.5$	0.25	$4.4 \pm 0.7$	
AM15-23	448	633	4.02	0.004	0.2	$6.7 \pm 1.3$	0.4	$8.3 \pm 0.2$	0.3	$9.2 \pm 0.9$	0.5	$10 \pm 1.4$	
AM18-23	448	624	3.72	0.019	0.18	$8.8 \pm 1.3$	0.2	$9.2 \pm 1.8$	0.22	$9.3 \pm 1.7$	0.3	$8.9 \pm 0.3$	
AM20	443	640	4.02	0.007	0.3	$4.1 \pm 0.1$	0.4	$9.8 \pm 0.3$	0.42	$10.1 \pm 0.5$	0.5	$7.1 \pm 0.9$	
AM19	421	549	4.39	0.12	0.18	$13.3 \pm 0.7$	0.12	$14 \pm 0.2$	0.15	$12.8 \pm 1.6$	0.18	$13 \pm 1.2$	
AM21	446	554	3.99	0.03	0.23	$1.9 \pm 0.3$	0.32	$7.7 \pm 0.7$	0.18	$8.1 \pm 1.2$	0.22	$1.5 \pm 0.6$	
P9	446	647	4.01	0.019	0.4	$2.3 \pm 0.1$	0.33	$4.1 \pm 0.2$	0.3	$8.1 \pm 0.7$	0.4	$15 \pm 1.2$	
P14	452	639	3.61	0.022	0.1	$15 \pm 2.6$	0.3	$14.7 \pm 0.9$	0.3	$15.4 \pm 2.1$	0.12	$14.9 \pm 0.6$	
FA	420	554	3.54	0.05	0.08	$9.1 \pm 1.5$	0.08	$8.8 \pm 0.9$	0.09	$9.0 \pm 1.3$	0.07	$9.2 \pm 0.7$	
P11	452	572	4	0.0013						Not determined			
P10	446	589	4	0.007						Not determined			
AM17	464	673	4.15	0.008						Not determined			

Note: Parameters given for dyes marked with asterisk have been reported in our previous works [\*].

$$Q = \frac{Q_s(1-10^{-A_s})S_p}{(1-10^{-A_p})S_s} \cdot \frac{n_p^2}{n_s^2} \quad (1)$$

where  $Q_s$  is the standard quantum yield,  $S_p$  and  $S_s$  are the areas under the fluorescence spectra of the dye and standard, respectively,  $A_p$  and  $A_s$  are absorbances of the dye and standard at a certain excitation wavelength,  $n_p$  and  $n_s$  are refractive indexes of the measured dye solution and standard, respectively.

## 2.5. Time-resolved fluorescence measurements

Time-resolved fluorescence intensity decays of benzanthrone dyes were measured using a Photon Technology International spectrofluorometer equipped by GL-302 Dye Laser (Canada). This machine uses GL-3300 Nitrogen Laser that delivers a crisp pulse at 337 nm. The solution of LC 4230 (POPOP) laser dye (Kodak, USA) in a mixture of toluene with ethanol was used to excite benzanthrone dyes at 421 nm. Instrument response functions were measured at the excitation wavelength using the colloidal silica (Ludox) as a scatterer. To get smooth curves the final decay profile

of each dye was obtained as the average of 6 repeats. The resulted data were fitted by biexponential function using Felix-GX software:

$$I(t) = \sum_{i=1}^n \alpha_i \exp(-t/\tau_i) \quad (2)$$

where  $\tau_i$  and  $\alpha_i$  are decay times and pre-exponential coefficients,  $n$  is the number of fluorescence lifetime components. The goodness of each fit was characterized by the value of reduced chi-square, weighted residuals and autocorrelation function of the weighted residuals. A fit was considered good when  $\chi^2$  fell within the range of 0.8–1.2, and the plots of weighted residuals and autocorrelation function were randomly distributed around zero [11].

The average fluorescence lifetime ( $\langle\tau\rangle$ ) of the dyes in the presence of liposomes was calculated using the following equation [11]:

$$\langle\tau\rangle = \frac{\sum_i \alpha_i \tau_i^2}{\sum_i \alpha_i \tau_i} \quad (3)$$

## 2.6. Partitioning model

The binding of benzanthrone dyes to model membranes has been analyzed in terms of partition model [25]. The total concentration of the dye distributing between aqueous and lipid phases ( $Z_{\text{tot}}$ ) can be represented as:

$$Z_{\text{tot}} = Z_F + Z_L \quad (4)$$

where subscripts  $F$  and  $L$  denote free and lipid-bound dye, respectively. The coefficient of dye partitioning between the two phases ( $K_P$ ) is defined as:

$$K_P = \frac{Z_L V_W}{Z_F V_L} \quad (5)$$

where  $V_W, V_L$  are the volumes of the aqueous and lipid phases, respectively. Given that under the employed experimental conditions the volume of lipid phase is much less than the total volume of the system  $V_t$ , we assume that  $V_W \approx V_t = 1 \text{ dm}^3$ . It is easy to show that

$$Z_F = \frac{Z_{\text{tot}} V_W}{V_W + K_P V_L} = \frac{Z_{\text{tot}}}{1 + K_P V_L} \quad (6)$$

The dye fluorescence intensity measured at a certain lipid concentration can be written as:

$$I = a_f Z_F + a_L Z_L = Z_F \left( a_f + a_L \frac{K_P V_L}{V_W} \right) \quad (7)$$

$$= Z_F (a_f + a_L K_P V_L)$$

where  $a_f, a_L$  represent molar fluorescence of the dye free in solution and in a lipid environment, respectively. From the equations (6) and (7) one obtains:

$$I = \frac{Z_{\text{tot}} V_W (a_f + a_L K_P V_L)}{V_W + K_P V_L} \quad (8)$$

The volume of lipid phase can be determined from:

$$V_L = N_A C_L \sum v_i f_i \quad (9)$$

where  $C_L$  is the molar lipid concentration,  $f_i$  is mole fraction of the  $i$ th bilayer constituent,  $v_i$  is its molecular volume taken as  $1.58 \text{ nm}^3, 3 \text{ nm}^3, 1.6 \text{ nm}^3$  and  $0.74 \text{ nm}^3$  for PC, CL, PG and Chol, respectively [26]. The relationship between  $K_P$  and fluorescence intensity increase ( $\Delta I$ ) upon the dye transfer from water to lipid phase can be written as:

$$\Delta I = I_L - I_W = \frac{K_P V_L (I_{\text{max}} - I_W)}{1 + K_P V_L} \quad (10)$$

where  $I_L$  is the dye fluorescence intensity measured at a certain lipid concentration  $C_L$ ,  $I_W$  is the dye fluorescence intensity in a buffer,  $I_{\text{max}}$  is the limit fluorescence in a lipid environment.

## 2.7. Binding model

Quantitative characteristics of the dye-lysozyme binding were determined in terms of the Langmuir adsorption model by analyzing protein-induced changes in the probe fluorescence intensity at the wavelengths, corresponding to emission maxima for each dye, as described previously [17, 18]. Briefly, approximation of the experimental dependencies

of  $\Delta I$  (fluorescence intensity increase) on  $C_P$  (total protein concentration), and  $\Delta I$  on  $Z$  (total probe concentration), the latter being performed only for  $F_{\text{acs}}$ , by equation (11) allowed us to determine the dye-protein binding parameters—association constant ( $K_a$ ), binding stoichiometry ( $n$ ) and molar fluorescence ( $\alpha$ ), characterizing the difference between molar fluorescence of the bound and free dye:

$$\Delta I = 0.5\alpha \left[ Z + nC_P + 1/K_a - \sqrt{(Z + nC_P + 1/K_a)^2 - 4nC_P Z} \right] \quad (11)$$

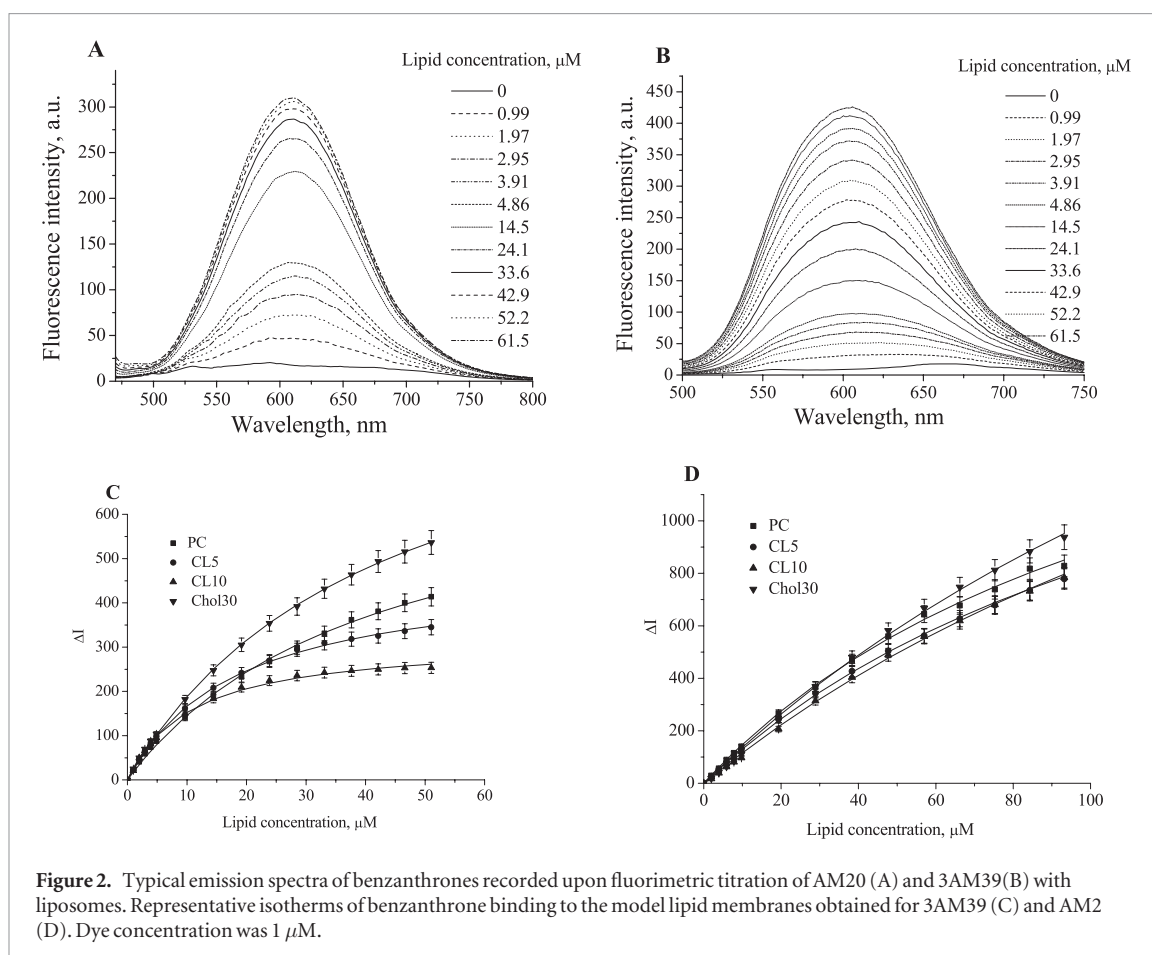
## 2.8. Quantum-chemical calculations

The energy of the highest occupied molecular orbital ( $E_{\text{HOMO}}$ ), the energy of the lowest unoccupied molecular orbital ( $E_{\text{LUMO}}$ ), the solvent-accessible area ( $CAr$ ), the cosmo volume (molecular volume) ( $CV$ ), the molecular length ( $L$ ), height ( $H$ ) and width ( $W$ ), polarizability of the molecule at the electric field strength  $0 \text{ eV}$  ( $P$ ), the total charge on carbon  $\sum Q(C)$  and nitrogen atoms  $\sum Q(N)$  was performed by the semiempirical PM6 method using MOPAC 2012 Version 10.006 W-free academic license [27]. Molecular descriptors  $\mu_g, \mu_e$  (the dipole moment of the ground and excited states, respectively);  $E_g, E_e$  (the energy of the ground and excited states, respectively);  $f$  (oscillator strength);  $\Delta E_e$  (the energy of electronic transition  $S_0 \rightarrow S_1^{\text{abs}}$  from the ground state to the lowest singlet excited state, formed immediately after photon absorption);  $\varphi$  (dihedral angle between the donor and acceptor);  $q_g, q_e$  (the total charge on the donor of the ground and excited states, respectively) was performed with the 6-31 G(d,p) basis set, using density functional theory (DFT) and B3LYP functional [28] with Win-Gamess (version May 1, 2013 R1), offering reasonable estimates of the ground state properties [29]. Semiempirical AM1 method with added polarization (1) and diffuse (1) functions on heavy atoms, and a polarization function on hydrogen atoms was employed for the ground state geometry optimization of the benzanthrone derivatives. Virtual Computational Chemistry Laboratory ([www.vcclab.org](http://www.vcclab.org)) was used for the calculation of lipophilicity of the examined compounds ( $C \log P$ ).

## 3. Results and discussion

### 3.1. Photophysical properties of benzanthrone derivatives

The dyes under study (figure 1 and table 1) are asymmetrical amidino- and amino- benzanthrone derivatives from yellow–green to red–purple colour. Presented in table 2 are the basic photophysical characteristics of the examined benzantrones in ethanol: absorption maximum ( $\lambda_A$ ); extinction coefficient ( $\epsilon$ ), emission maximum ( $\lambda_F$ ). The examined benzanthrone dyes are characterized by the broad absorption and fluorescence spectra with absorption maxima in ethanol within a range from 420 to 530 nm



and emission maxima from 554 to 657 nm depending on the dye structure. These dyes were found to be nearly non-emissive in buffer solution. The corresponding quantum yields of benzantrone dyes in buffer ( $Q_0$ ) are presented in table 2. As was demonstrated by Kirilova and coauthors, benzantrone dyes displayed strong fluorescent solvatochromism due to the donor–acceptor nature of their structures [7]. Specifically, the emission maxima of the hydrophobic probes in ethanol showed  $\sim 30$ – $70$  nm shifts to the longer wavelengths as compared to those in benzene. Furthermore, a comprehensive analysis of the dye spectral properties in a range of solvents of varying polarity allowed to detect the increased Stoke’s shifts and decreased quantum yields with the growth of the environmental polarity [22]. Notably, the  $\lambda_A$  values of ISH, IAH, IBH and A4 are red-shifted compared to other dyes, the property which is most likely associated with higher electron-donating ability of amino group compared to that of amidino group [2, 4, 22].

### 3.2. Association of benzantrone dyes with model lipid membranes

At the first step of the study it was of interest to ascertain how the dye lipid-associating ability depends on the membrane physical properties being varied through introducing anionic phospholipid (CL) or sterol (Chol) into PC bilayer. For this purpose emission spectra of 25 benzantrone dyes were recorded in buffer solution and liposomal suspensions. Typical fluorescence

spectra measured at increasing lipid concentration are presented in figures 2(A) and (B). Fluorescence intensity and quantum yield were found to increase upon the dye transfer from aqueous to lipid phase as a result of reduced polarity of the dye lipid surroundings and restricted fluorophore mobility within the lipid bilayer. To derive the dye partition coefficients in different lipid systems, experimental profiles  $\Delta I(C_L)$  (figure 2) were approximated by equation (10). The resulting binding curves were hyperbolic in shape for all dyes under study (figure 2(C)) except of AM2, AM3, AM4 and IAH for which linear dependencies  $\Delta I(C_L)$  were observed (figure 2(D)). This means that benzantrones AM2, AM3, AM4 and IAH possess considerably lower lipid-associating ability compared to other dyes. It is important to point out that the quantum yields of ISH, IAH, IBH in the lipid vesicles are substantially lower compared to other dyes complicating their use in membrane studies. As seen from table 2, benzantrone derivatives AM1, AM3, AM4, A4, AM19, P14 and FA appeared to be insensitive to the changes in membrane physical properties since inclusion of Chol or CL into PC bilayer didn’t affect the partition coefficients of these dyes. However, as the partition ability of a series of benzantrones depends on lipid bilayer composition, further it was interesting to investigate the lipid binding of the most promising fluorescent probes with a broader variation of the membrane content. To this end, the model lipid membranes were formed from PC and its mixtures with 5.3, 11.1 or 25 mol% CL, 10, 20 or

40 mol% PG and 10, 20 or 30 mol of Chol. The choice of the above components of the examined lipid systems was dictated by at least three reasons [30, 31]:

- (i) CL content in the inner mitochondrial membrane does not exceed 10 mol%, but it can increase considerably till 25–30 mol% in the contact zones between the inner and outer layers of the membrane;
- (ii) to assess the role of headgroup chemical nature and surface charge on the lipid-associating ability of benzanthrone dyes we employed two anionic lipids CL and PG;
- (iii) since Chol exerts stabilizing influence on lipid bilayer structure, it was interesting to ascertain whether this agent can modulate spectral behavior of the examined fluorophores.

The parameters of benzanthrone partitioning into lipid bilayers of different composition are presented in table 3. It turned out that inclusion of anionic lipids and Chol has an ambiguous effect on partition ability of benzanthrone dyes since addition of CL, PG and Chol to PC bilayer gives rise to the decrease of partition coefficients compared to the neat PC membrane for fluorophores of the first group (A6, AM12, AM15 and AM18), whereas benzanthrone dyes of the second group (ABM, 3AM39, AM20, AM15-23, AM4-23, AM2-23 and P9) display the opposite behavior. Since benzanthrone under study are uncharged molecules, the observed tendencies in their behavior cannot be explained by electrostatic dye-lipid interactions and most probably result from the modulating effect of anionic lipids and Chol on the structural and dynamical properties of PC bilayer. Moreover, taking into account the fundamental differences in spectral behavior of A6, AM12, AM15, AM18, on one hand, and ABM, A8, 3-AM39, AM20, AM15-23, AM4-23, AM2-23, P9, on the other hand, it seemed reasonable to scrutinize bilayer location of these two groups of fluorophores.

Therefore, at the next step of the study red edge excitation shift (REES) approach was used to determine the properties of benzanthrone binding sites. The red edge effect is known to originate from the differences in fluorophore-solvent interactions in the ground and excited states, brought about by the changes in the dye dipole moment upon excitation and the rate of solvent reorientation around the excited state fluorophore [32]. In the steady-state fluorescence measurements REES manifests itself as a shift of emission maximum towards higher wavelengths with increasing excitation wavelength. It appeared that REES does not occur for A6, AM12, AM15 and AM18. Recent studies revealed that according to their properties along the normal to bilayer plane, lipid membranes can be divided into three regions: (i) aqueous phase with fast solvent relaxation; (ii) anisotropic polar region where solvent relaxation is restricted and slow; (iii) isotropic hydrophobic part characterized by fast solvent relaxation [33, 34]. Since REES is a characteristic of hydrophilic membrane part,

the fact that AM12, AM18, AM15 and A6 did not show any shift of fluorescence maximum with the change in excitation wavelength indicates that fluorophores of the first group are localized in the nonpolar region of the lipid bilayer.

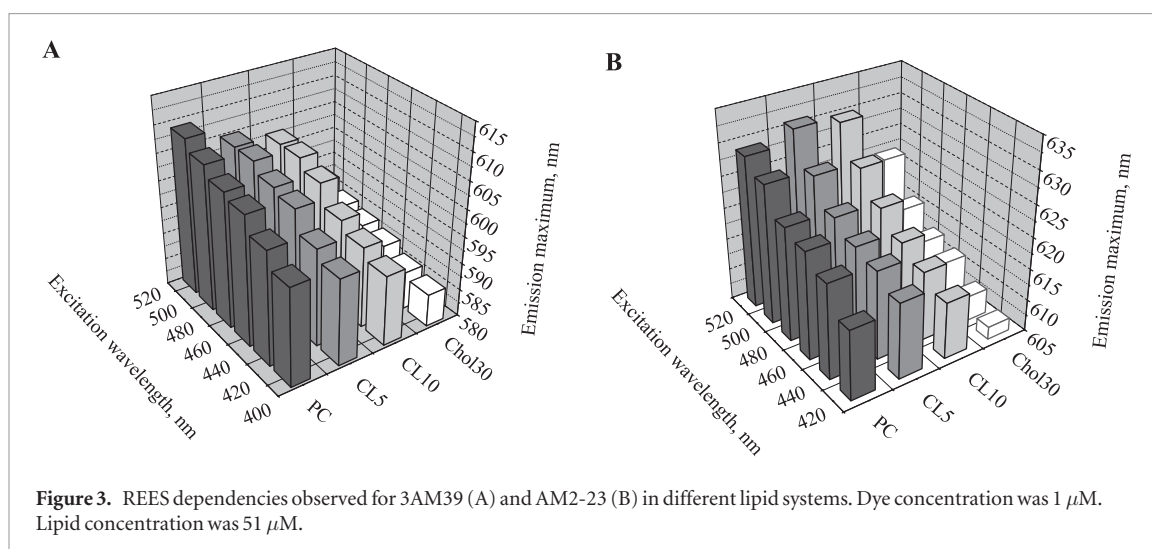
In the meantime, as can be seen in figure 3, benzanthrone of the second group exhibited REES, with the magnitude of emission shift being dependent on the probe structure. The most pronounced REES was observed for AM4-23 та AM20, while the smallest one was revealed for ABM. The REES values were found to rise in the order ABM (6 nm) → 3-AM39 (11 nm) → A8 (12 nm) → AM2-23 (13 nm) → P9 (14 nm) → AM15-23 (14 nm) → AM20 (20 nm) → AM4-23 (24 nm), indicating the increase of emission shift from ABM to AM4-23. The obtained REES values allowed us to range the dyes of the second group in accordance with depth of their bilayer location. As was proposed by Chattopadhyay *et al* (so-called ‘dipstick’ rule), the REES magnitude varies in direct correlation with the depth of probe penetration into membrane interior—the less the REES, the deeper probe locates [32]. Accordingly, ABM is apparently localized in the interfacial region, most likely near the lipid carbonyl groups, while the other dyes prefer the binding sites which are closer to the membrane surface.

Therefore, while attempting to interpret the observed increase of partition coefficients for fluorophores of the second group, the influence of anionic lipids and Chol on physicochemical properties of the lipid-water interface should be considered. One of the possible effects of CL, PG and Chol involves the alterations in bilayer hydration. First of all it should be noted that PC, in contrast to CL, doesn’t contain donor groups for hydrogen bonds (CL is capable of forming a network of intra- or intermolecular hydrogen bonds) [35]. Shibata *et al* based on FTIR data, hypothesized that the amount of membrane-bound water increases significantly in the presence of cone-shaped CL molecules (CL enhances the hydration of ester C=O groups) [36]. These findings are in good agreement with the results of molecular dynamic simulations of CL-containing lipid membranes [35]. Likewise, the ability to increase bilayer hydration was also reported for PG and Chol. In particular, several authors demonstrated that hydroxyl group of PG can form hydrogen bonds with lipid molecules as well as with water [37, 38]. Moreover, an additional argument in favor of the increase in the degree of bilayer hydration in the presence of anionic lipids, stems from the analysis of spectral responses of a number of environment-sensitive fluorophores [39, 40]. Specifically, this effect was found by the measurements of the generalized polarization of Laurdan (2-dimethylamino-6-dodecanoylnaphthalene) in micelles composed of anionic gangliosides G1, G2, G3, and model membranes, containing dipalmitoylphosphatidylcholine (DPPC) and its mixtures with CL [41, 42]. Laurdan and Prodan (2-dimethylamino-6-propionynaphthalene) molecules possess donor and

**Table 3.** Parameters of benzanthrone dye partitioning into lipid systems.

Dye	Parameter	PC	CL5	CL10	CL25	PG10	PG20	PG40	Chol10	Chol20	Chol30
ABM	$K_p \times 10^4$	6.4 ± 0.7	8.2 ± 0.4	10.3 ± 0.9	11.5 ± 1.1	6.9 ± 0.9	7.4 ± 0.7	7.7 ± 1.1	7.9 ± 0.2	8.1 ± 0.4	6.9 ± 0.3
	$\Delta I_{max}$	197 ± 3	170 ± 3	188 ± 4	290 ± 15	224 ± 6	230 ± 0.9	244 ± 4	225 ± 5	353 ± 8	412 ± 7
	$K_p \times 10^4$	11 ± 1.5	9.2 ± 0.9	8.4 ± 0.7	7.8 ± 0.5	8.8 ± 0.3	8.2 ± 0.3	7.1 ± 1.1	10.1 ± 1.2	9.0 ± 0.1	8.7 ± 0.5
A6	$\Delta I_{max}$	280 ± 5	319 ± 3	331 ± 8	368 ± 2	313 ± 10	354 ± 31	364 ± 21	392 ± 9	419 ± 9	423 ± 10
	$K_p \times 10^4$	0.9 ± 0.07	3.4 ± 0.5	6.2 ± 1.6	7.8 ± 0.5	4.2 ± 0.1	5.3 ± 0.7	8.1 ± 0.9	5.5 ± 0.5	6.4 ± 0.5	6.5 ± 0.7
	$\Delta I_{max}$	334 ± 19	401 ± 16	514 ± 18	534 ± 20	577 ± 5	680 ± 23	632 ± 15	519 ± 21	532 ± 4	498 ± 5
AM12	$K_p \times 10^4$	18 ± 0.8	8.7 ± 0.3	5.8 ± 0.6	3.1 ± 0.8	8.1 ± 0.6	7.9 ± 0.4	5.6 ± 0.5	17 ± 1.2	14.8 ± 0.8	15 ± 1.5
	$\Delta I_{max}$	316 ± 3	287 ± 8	267 ± 16	311 ± 14	232 ± 34	208 ± 5	197 ± 15	528 ± 8	342 ± 25	480 ± 8
	$K_p \times 10^4$	6.1 ± 0.2	2.7 ± 0.2	1.6 ± 0.5	1.4 ± 0.4	4.1 ± 0.5	3.4 ± 0.5	2.9 ± 0.8	6.2 ± 0.3	5.7 ± 1.4	4.9 ± 0.4
AM15	$\Delta I_{max}$	401 ± 12	373 ± 15	353 ± 58	323 ± 12	321 ± 21	297 ± 16	281 ± 20	385 ± 11	342 ± 48	280 ± 9
	$K_p \times 10^4$	12 ± 0.8	9.7 ± 0.6	7.7 ± 0.5	5.4 ± 0.7	8.4 ± 0.7	6.9 ± 0.5	3.7 ± 0.6	9.6 ± 1.1	8.1 ± 0.7	7.8 ± 0.5
	$\Delta I_{max}$	542 ± 10	417 ± 23	334 ± 7	401 ± 14	365 ± 13	402 ± 5	423 ± 25	487 ± 23	455 ± 31	554 ± 14
3AM39	$K_p \times 10^4$	2.5 ± 0.1	5.5 ± 0.2	9.0 ± 0.4	12.8 ± 1.2	3.6 ± 0.7	5.2 ± 0.3	7.4 ± 0.4	2.7 ± 0.3	2.9 ± 0.2	3.4 ± 0.1
	$\Delta I_{max}$	749 ± 24	471 ± 4	315 ± 4	286 ± 9	694 ± 11	444 ± 20	370 ± 9	821 ± 13	808 ± 65	1099 ± 65
	$K_p \times 10^4$	4.1 ± 0.1	9.8 ± 0.3	10.1 ± 5	12 ± 0.8	9.4 ± 0.6	12.4 ± 0.8	14.4 ± 1.3	7.1 ± 0.9	7.5 ± 0.8	8.2 ± 0.7
AM20	$\Delta I_{max}$	458 ± 4	334 ± 3	303 ± 4	237 ± 4	253 ± 5	190 ± 3	181 ± 4	354 ± 16	314 ± 8	382 ± 10
	$K_p \times 10^4$	6.7 ± 1.3	8.3 ± 0.2	9.2 ± 0.9	9.4 ± 0.5	7.7 ± 1.5	8.4 ± 0.6	9.2 ± 0.5	11.1 ± 0.8	11.9 ± 0.9	10 ± 1.4
	$\Delta I_{max}$	399 ± 28	472 ± 4	373 ± 9	563 ± 25	393 ± 11	385 ± 11	386 ± 9	716 ± 21	824 ± 14	722 ± 25
AM4-23	$K_p \times 10^4$	4.1 ± 0.9	10 ± 1.7	11.4 ± 0.5	6.3 ± 1.2	7.8 ± 0.5	8.5 ± 1.2	8.7 ± 0.8	4.1 ± 0.4	4.3 ± 0.7	4.4 ± 0.7
	$\Delta I_{max}$	409 ± 53	259 ± 17	230 ± 16	310 ± 27	152 ± 6	171 ± 9	227 ± 9	362 ± 15	386 ± 5	424 ± 43
	$K_p \times 10^4$	5.1 ± 0.2	7.6 ± 0.2	9.8 ± 0.7	11.4 ± 0.5	7.5 ± 1.2	8.4 ± 0.1	9.9 ± 2.1	5.4 ± 0.5	8.0 ± 1.6	8.8 ± 0.4
AM2-23	$\Delta I_{max}$	727 ± 12	653 ± 7	524 ± 1	498 ± 7	449 ± 37	410 ± 0.25	445 ± 15	514 ± 14	455 ± 43	411 ± 31
	$K_p \times 10^4$	2.3 ± 0.1	4.1 ± 0.2	8.1 ± 0.7	9.6 ± 0.4	4.4 ± 0.4	6.2 ± 0.7	8.4 ± 0.2	12.6 ± 1	13 ± 0.8	15 ± 1.1
	$\Delta I_{max}$	513 ± 25	448 ± 10	467 ± 5	502 ± 9	303 ± 3	284 ± 7	286 ± 16	238 ± 6	312 ± 11	355 ± 10



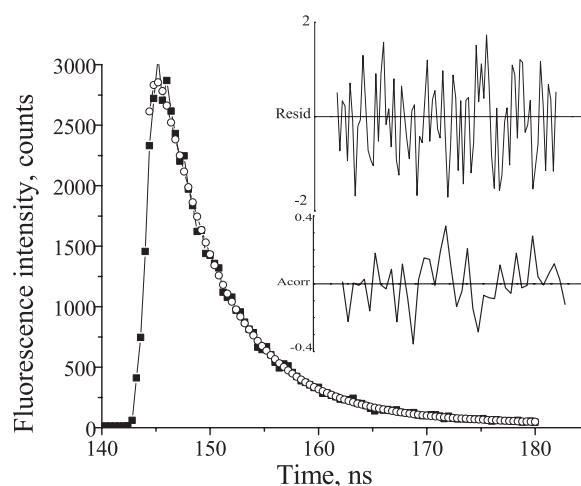


acceptor moieties, high ground and excited state dipole moments, which are also the characteristic features of benzantrones [22, 43]. The two emission bands of the dyes are associated with a pair of non-relaxed (or locally excited) and solvent-relaxed (or twisted internal charge transfer) S1 states, the latter being predominant in polar solvents. Fluorescence emission maxima and lifetimes of Laurdan and Prodan are highly sensitive to the presence of water in the membrane [44, 45]. Specifically, they show long-wavelength spectral shifts  $\sim 50$  nm and the shorter excited state lifetimes on the bilayer transition from the gel to the fluid phase [44, 46]. Furthermore, the novel dye 2-amino-N-hexadecyl-benzamide (Ahba) was also sensitive to the modifications of the bilayer core and headgroup region, as revealed by the steady-state and time-resolved anisotropy measurements [46]. A wealth of evidence indicates that Chol is also capable of inducing the elevation of water content in polar region of lipid bilayer [47]. Specifically it was shown that Chol inclusion into PC bilayer alters lipid packing density allowing a greater number of water molecules to penetrate in the membrane headgroup region [47]. Another support to the idea that ABM, A8, 3-AM39, AM20, AM15-23, AM4-23, AM2-23 and P9 can be employed as reporters of the changes in membrane polarity comes from 2 to 5 nm shifts of fluorescence maxima in the presence of anionic lipids and Chol compared to those observed in neat PC bilayer.

To explain the decrease of partition coefficients of A6, AM 12, AM 15 and AM18 in the presence of CL, PG and Chol, we should search for the mechanism that may underlie the effect of these lipids on physicochemical characteristics of a hydrophobic region of lipid bilayer. Specifically, liposomes composed of 1-palmitoyl-2-oleoyl-*sn*-glycero-3-phosphocholine (POPC) are in a fluid phase at 20  $^{\circ}\text{C}$  [48]. However, similar to other lipids in a fluid phase, they form liquid ordered state, possessing higher acyl chain order, when more than 20% Chol is introduced into the bilayer [49, 50]. Since in liquid-crystalline lipid phase Chol can produce tighter lateral packing of lipid molecules, the observed  $K_p$  decrease

can be a consequence of Chol condensing effect on PC bilayer [26]. The situation becomes more complicated in the case of CL-containing membranes, where partition coefficient is reduced in value, since the ordering effect of CL on zwitterionic lipid bilayer is still a matter of controversy. Particularly, recent monolayer study revealed that the order of PC hydrocarbon chains increases upon CL inclusion [51]. However, fluorescence anisotropy measurements indicate that inclusion of up to 20 mol% CL in PC bilayer does not change bilayer fluidity [52]. Besides, on the basis of FTIR data Shibata *et al* [36] showed that CL negative charge tends to move the  $\text{N}^+$  end of P-N dipole parallel to the membrane surface, thereby causing the rearrangement of water bridges at the bilayer surface and stabilizing in such a way the intermolecular hydrogen-bonded network including hydration water. Interestingly, it was found that PG also exerts a stabilizing influence on zwitterionic bilayers [53]. The changes in hydration extent is thought to considerably affect molecular organization of a lipid bilayer, since the increase of water content in headgroup region was reported to modify the alignment of choline-phosphate dipole and lateral packing of hydrocarbon chains [54]. Presumably, the above factors prevent partitioning of A6, AM12, AM15 and AM18 in the CL- and PG-containing lipid bilayers, so we can assume that these dyes are sensitive to the changes in membrane packing density.

To better characterise the membrane partitioning of the examined compounds, in the following experiments we performed time-resolved fluorescence measurements for five fluorophores of the second group (ABM, AM4-23, AM20, 3AM39, AM2-23) and A6 which belongs to the first group of the dyes. The representative time-resolved fluorescence intensity decays of benzantrones together with the parameters responsible for goodness of each fit are given in figure 4 and table 4. As can be seen from table 4, the fluorescence intensity decays in lipid membranes are adequately fitted by biexponential function where the longer lifetime component ( $\tau_1$ ) corresponds to the lifetime of the lipid-bound dye (lower environmental



**Figure 4.** Time-resolved fluorescence intensity decay of ABM in PC liposomes (■), together with its exponential fit (○). Concentration of ABM was  $2.4 \mu\text{M}$ . The excitation wavelength was 460 nm and emission was monitored at 600 nm. Lipid concentration was  $33 \mu\text{M}$ . Shown in insets are the weighted residuals and autocorrelation function of weighted residuals.

polarity compared to buffer solution), while the shorter component accounts for the free dye in buffer (polar environment). The data presented in table 4 indicate that inclusion of anionic lipids PG, or CL and Chol into PC membrane leads to significant variations in fluorescence lifetimes of benzantrones ( $\tau_1$  and  $\tau_2$ ), as well as in fractional contributions of each decay time to the overall intensity ( $\alpha_1$  and  $\alpha_2$ ).

Generally, while exploring the fluorescence decay profiles the parameter, such as the average fluorescence lifetime is used (equation (3)) [11]. As can be seen from table 4, inclusion of CL, PG and Chol into PC bilayer gives rise to the increase of  $\langle\tau\rangle$  values compared to the neat PC membranes. To interpret this fact, one should bear in mind that the excited-state dipole moments of benzantrone dyes are rather high compared to the ground-state counterparts, pointing to a substantial redistribution of the  $\pi$ -electron densities in a more polar excited state [11, 20, 22]. As a result, significant change in the dipole moment upon excitation indicates that the excited state is a twisted intramolecular charge-transfer (TICT) in nature. Many fluorescent compounds, especially those with flexible skeletons in the excited state show lifetime increase due to their sensitivity to membrane dynamics [11]. Indeed, as mentioned above, fluorescence lifetimes of e.g. Laurdan and Prodan increased when the bilayer goes from the fluid to the less hydrated gel phase [44, 46]. Since, as it was hypothesized above, A6 is localized in the hydrophobic membrane region, an increase of its  $\langle\tau\rangle$  values is likely a consequence of CL, PG and Chol ability to affect the packing and ordering of lipid bilayers. Obviously, A6 belongs to the group of molecular probes sensitive to rigidity of their local environment. Lipid-associating properties of ABM, AM4-23, AM20, 3AM39 and AM-23 seem to be different compared to A6, as judged from the decrease of average fluorescence lifetime in the presence of anionic lipids and Chol. According to the results of REES measurements, these

dyes are localized in the polar part of lipid bilayer, so the observed decrease of  $\langle\tau\rangle$  values for ABM, AM4-23, AM20, 3AM39 and AM-23 may reflect the increase of microenvironmental polarity in CL, PG and Chol-containing membranes, as was also evidenced by Laurdan and Prodan [44, 46].

At the next step of the study fluorescence anisotropy of A6, ABM, AM4-23, AM20, 3AM39 and AM-23 was measured in the various lipid systems. Fluorescence anisotropy of a membrane-bound probe is determined by the rate of its rotational diffusion. Since diffusive motions depend on free volume of the dye microenvironment [11], anisotropy values reflect the changes in lipid packing density. As illustrated in table 5, only anisotropy of A6 appeared to be sensitive to the changes in membrane composition. The observation that anisotropy values are higher in Chol-containing membranes in comparison with pure PC liposomes confirms the above hypothesis that A6 resides in the hydrophobic membrane region experiencing condensing effect of Chol.

### 3.3. Benzantrone dyes as amyloid markers

In the following, it seems of importance to illustrate that applicability of the novel benzantrones is not limited by the membrane studies. One possible application of these dyes in the protein research field involves detection and structural analysis of particular class of protein aggregates, amyloid fibrils, associated with a number of pathologies. To exemplify, here we present the data on benzantrone interaction with the native lysozyme and  $F_{acs}$  fibrils. To estimate the quantitative parameters of the dye-protein complexation (association constant ( $K_a$ ) and binding stoichiometry ( $n$ )), the experimental dependencies of fluorescence increase on the protein concentration  $\Delta I(C_p)$  (figure 5) were approximated by the Langmuir adsorption model (equation (11)). Likewise, for more accurate evaluation of the dye specificity for a certain

**Table 4.** Fluorescence lifetimes of benzanthrone dyes in different lipid systems.

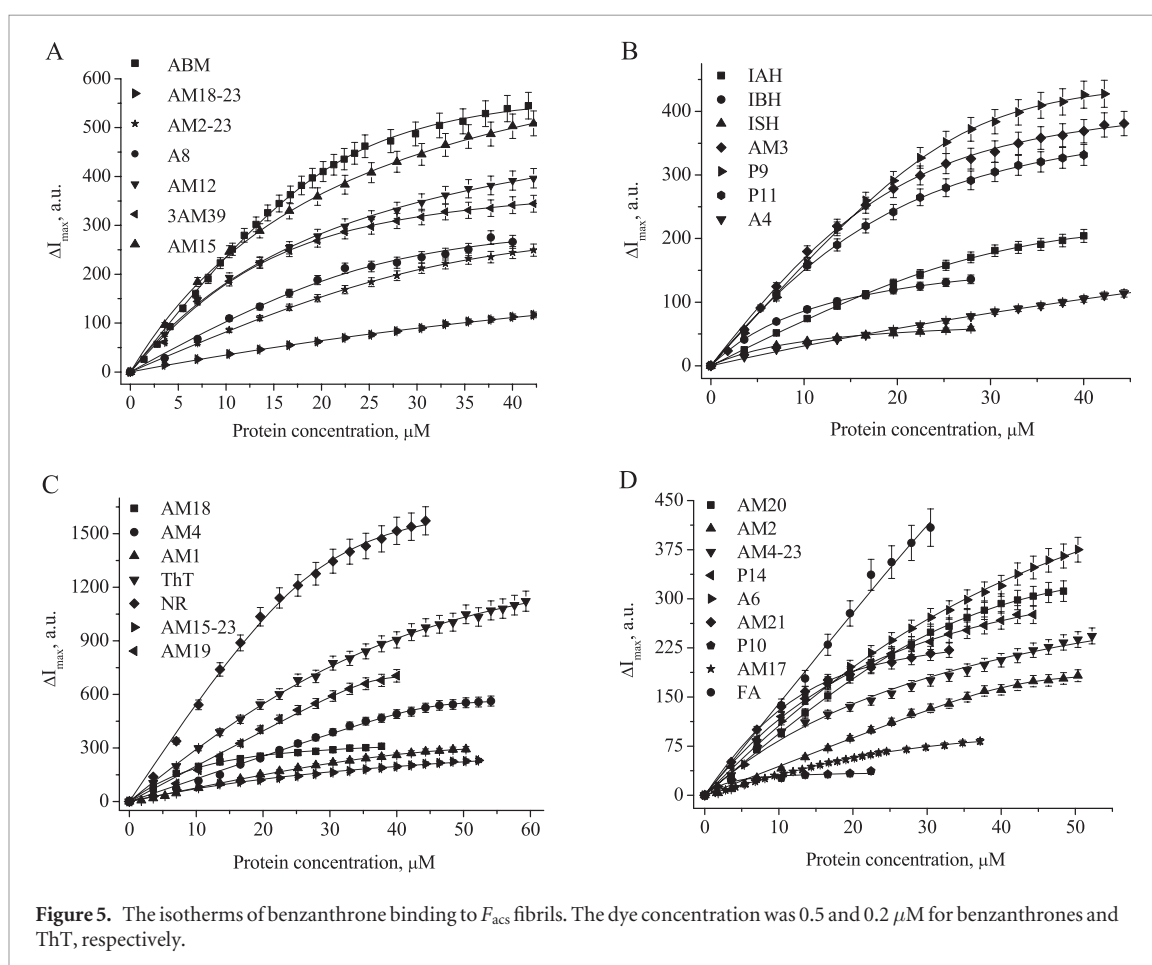
Dye	Parameter	PC	CL5	CL10	CL25	PG10	PG20	PG40	Chol30
A6	$\alpha_1$	0.85	0.82	0.81	0.79	0.77	0.79	0.69	0.83
	$\tau_{1, ns}$	5.46	5.59	5.75	5.81	5.74	5.82	5.99	6.58
	$\alpha_2$	0.15	0.18	0.19	0.21	0.23	0.21	0.31	0.17
	$\tau_{2, ns}$	0.63	0.77	0.89	0.93	1.25	1.31	1.34	1.48
	$\langle \tau \rangle, ns$	5.36	5.45	5.57	5.61	5.46	5.56	5.56	6.35
ABM	$\chi^2$	1.09	1.17	1.18	1.08	1.3	1.1	1.0	1.4
	$\alpha_1$	0.8	0.77	0.79	0.79	0.81	0.77	0.7	0.75
	$\tau_{1, ns}$	6.49	6.37	6.24	6.22	6.33	6.11	5.9	8.07
	$\alpha_2$	0.2	0.23	0.21	0.21	0.23	0.23	0.3	0.25
	$\tau_{2, ns}$	1.98	2.11	1.57	2.12	1.92	1.83	1.95	2.96
AM4-23	$\langle \tau \rangle, ns$	6.17	5.99	5.95	5.88	5.98	5.76	5.41	7.51
	$\chi^2$	0.9	1.0	1.11	0.96	1.04	1.2	1.24	0.93
	$\alpha_1$	0.69	0.71	0.73	0.79	0.76	0.78	0.77	0.65
	$\tau_{1, ns}$	5.4	5.25	5.15	4.9	5.23	4.97	4.89	5.3
	$\alpha_2$	0.31	0.29	0.27	0.21	0.24	0.22	0.23	0.35
AM20	$\tau_{2, ns}$	2.2	1.92	1.99	2.1	2.2	2.14	1.97	2.03
	$\langle \tau \rangle, ns$	4.9	4.81	4.75	4.61	4.87	4.66	4.75	4.74
	$\chi^2$	0.89	0.98	0.93	1.01	0.95	1.11	0.95	1.14
	$\alpha_1$	0.88	0.91	0.79	0.81	0.76	0.82	0.68	0.75
	$\tau_{1, ns}$	5.61	5.37	5.4	5.3	5.8	5.64	5.9	5.29
3AM39	$\alpha_2$	0.12	0.09	0.21	0.19	0.24	0.18	0.32	0.25
	$\tau_{2, ns}$	1.8	1.3	1.6	1.8	2.1	1.89	2.21	2.14
	$\langle \tau \rangle, ns$	5.45	5.27	5.12	5.04	5.42	5.38	5.34	4.91
	$\chi^2$	1.02	0.8	0.83	0.9	1.01	0.94	0.87	1.14
	$\alpha_1$	0.89	0.79	0.8	0.79	0.83	0.84	0.89	0.86
AM2-23	$\tau_{1, ns}$	6.13	6.27	6.22	6.21	6.04	5.95	5.57	5.35
	$\alpha_2$	0.11	0.21	0.2	0.21	0.17	0.16	0.11	0.14
	$\tau_{2, ns}$	1.73	1.93	1.93	2	1.33	1.4	1.1	1.71
	$\langle \tau \rangle, ns$	5.98	5.96	5.91	5.89	5.83	5.75	5.46	5.17
	$\chi^2$	0.9	0.91	0.89	1.0	1.1	1.2	1.2	0.91
AM2-23	$\alpha_1$	0.72	0.71	0.71	0.72	0.71	0.8	0.72	0.75
	$\tau_{1, ns}$	7.1	6.4	6.1	6.7	6.2	6.11	5.98	5.31
	$\alpha_2$	0.28	0.29	0.29	0.28	0.29	0.2	0.24	0.25
	$\tau_{2, ns}$	3.5	2.7	2.6	2.7	1.9	1.1	1.6	1.7
	$\langle \tau \rangle, ns$	6.52	5.86	5.58	6.16	5.74	5.78	5.57	4.95
	$\chi^2$	0.93	0.87	0.98	0.82	0.83	1.2	1.1	1.1

protein conformation, we employed the following additional quantities: (i)  $I/I_0$ —relative fluorescence intensity increase upon the dye binding to native or fibrillar protein; (ii)  $I/I_{nat}$ —specificity of the dyes to protein fibrillar aggregates; (iii)  $Q$ —quantum yield of the fibril-bound dye. The binding characteristics of benzanthrone dyes associated with native protein and lysozyme fibrils are presented in table 6. The titration of IAH, IBH, ISH, AM21, AM19 with native lysozyme results in saturation of the binding curves only at millimolar protein concentration indicating that association constants are very low for these dyes. Notably, for majority of the dyes elevating concentrations of native lysozyme did not produce any change in the dye emission as evidenced from relatively low quantum yields of fluorophores comparable with their quantum yields in the free state. Such a

behavior may be a consequence of very low affinity of benzanthrone derivatives for protein monomers. It appeared that in the presence of lysozyme fibrillar aggregates A4, IAH and IBH showed the highest fluorescence intensity increase on going from the aqueous phase to protein matrix, while the binding preferences were the best for IAH, ABM, AM4, IBH, AM12, A4, 3AM39, AM15, AM2-23 and AM18-23. Interestingly, the  $K_a$  values for A6, A8, P11, P14, ABM, AM4-23 were 2–60 times higher for native protein in comparison with fibrillar one, whereas AM1, AM4, AM12, AM15, AM18, AM2-23, P9, IAH display the opposite behavior. In addition, fluorescence maxima of A6, AM1, AM4, AM12, AM15, AM18, AM2-23, AM21, P9, A8, P11, P14, IAH, ABM and AM4-23 in monomeric protein ( $\lambda_{em}^{nat}$ ) were shifted to the long-wavelength region relative to their position in fibrils

**Table 5.** Fluorescence anisotropy of benzanthrone dyes in different lipid systems.

Liposomes	ABM	A6	AM423	AM20	3AM39	AM223
PC	0.28	0.21	0.22	0.18	0.17	0.16
CL5	0.28	0.22	0.23	0.18	0.17	0.17
CL10	0.28	0.24	0.22	0.19	0.18	0.16
CL25	0.27	0.23	0.21	0.19	0.17	0.16
PG10	0.28	0.21	0.21	0.18	0.17	0.18
PG20	0.28	0.21	0.22	0.18	0.18	0.17
PG40	0.28	0.22	0.22	0.19	0.18	0.16
Chol10	0.27	0.23	0.21	0.18	0.17	0.17
Chol 20	0.27	0.25	0.22	0.19	0.18	0.17
Chol 30	0.28	0.26	0.22	0.19	0.17	0.16

**Figure 5.** The isotherms of benzanthrone binding to  $F_{acs}$  fibrils. The dye concentration was 0.5 and 0.2  $\mu\text{M}$  for benzanthrone and ThT, respectively.

( $\lambda_{em}^{F_{acs}}$ ), whereas the other dyes were characterized by blue-shifted maxima (table 6). Together with fluorescence maxima position the higher  $K_a$  values observed for A6, A8, P11, P14, ABM, AM4-23 in native lysozyme are suggestive of the sensitivity of these dyes to polarity of their environment. Accordingly, the  $K_a$  values recovered for AM1, AM4, AM12, AM15, AM18, AM2-23, P9, IAH respond to the changes in viscosity of binding sites, since fluorophore surroundings is more rigid in the case of fibrillar protein [55, 56]. To compare the properties of the dyes under study with those of the classical amyloid marker, the binding parameters have been derived also for Thioflavin T (ThT). The results presented in table 6 indicate that benzanthrone dyes and ThT possess relatively high association constant

0.2–1  $\mu\text{M}^{-1}$ , except of P14, AM18, AM21, AM17, AM19 and FA, for which  $K_a$  values range from 5 to 450  $\mu\text{M}^{-1}$ . The values of  $n$  in the presence of amyloid fibrils were approximately equal to 0.05, although the binding stoichiometry for IAH, IBH, ISH, P10 was in the range 0.3–0.5, whereas for AM19, FA, AM4, AM17, ThT this parameter ranges from 0.01 to 0.02. Noteworthy, the classical marker ThT showed higher binding preference to amyloid fibrils in comparison with the examined dyes. However, the association constants of benzanthrone and their quantum yields were in many cases higher, giving the impetus for the use of these dyes in amyloid research. To range the examined dyes according to their specificity to fibrillar protein, three parameters have been used: fluorescence

**Table 6.** Binding characteristics of benzanthrone dyes associated with native lysozyme and  $F_{acs}$  fibrils.

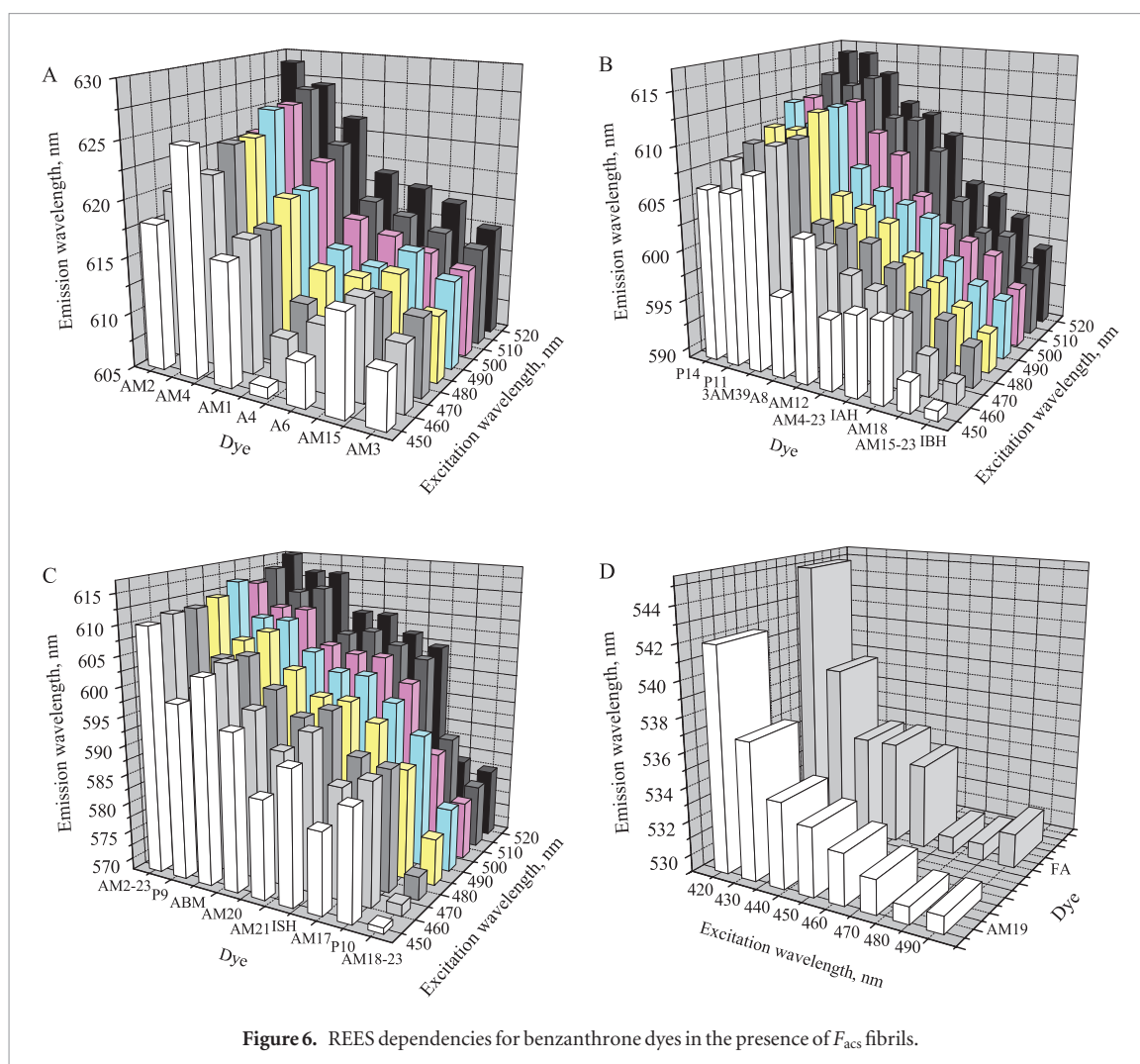
Dye	Native lysozyme					$F_{acs}$					
	$K_a, \mu M^{-1}$	$n$	$I/I_0$	$\lambda_{em}^{nat}, nm$	$Q_n$	$K_a, \mu M^{-1}$	$n$	$I/I_0$	$I/I_{nat}$	$\lambda_{em}^{Facs}, nm$	$Q$
AM1	0.1	0.52	0	609	0.002	0.5	0.05	13	$\infty$	617	0.15
AM2	0.33	0.23	1	655	0.004	0.2	0.12	36	16.4	616	0.16
AM3	—	—	1	653	0	0.25	0.13	14	14	610	0.18
AM4	0.36	0.027	2.2	594	0.009	0.63	0.02	44	17	624	0.33
ABM	4.9	0.54	6.4	564	0.015	0.93	0.06	75	13	603	0.35
A6	3.5	0.11	4.6	600	0.07	0.4	0.07	9	2.3	612	0.23
A8	0.17	0.09	0.8	573	0.024	0.08	0.16	8	10	598	0.47
A4	0.09	0.07	3	621	0.003	0.46	0.06	161	42	614	0.04
AM12	0.33	0.68	1.2	596	0.007	0.62	0.05	29	26	604	0.34
AM15	0.02	6	2.2	609	0.009	1	0.04	41	21	615	0.17
AM18	0.3	0.03	0	575	0.001	5.6	0.12	23	$\infty$	597	0.08
IAH	0.002	0.057	1.6	600	0.061	0.2	0.33	135	132	602	0.2
IBH	0.001	0.1	1	616	0.042	0.33	0.43	98	81	595	0.1
ISH	0.001	0.27	1.1	626	0.034	0.96	0.4	49	37	600	0.03
3AM39	0.13	2	1.4	643	0.013	0.61	0.07	27	14	611	0.23
AM2-23	0.37	0.25	3.8	594	0.015	0.47	0.06	33	9	610	0.22
AM4-23	30	0.06	5	591	0.017	0.43	0.05	11	4	597	0.22
AM15-23	0.87	0.03	1.8	650	0.007	0.87	0.04	28	20	593	0.12
AM18-23	—	—	1	636	0	0.20	0.05	22	22	562	0.17
AM20	0.16	1.3	1.6	606	0.026	0.42	0.04	12	3.4	597	0.36
AM19	0.003	9.9	0.7	593	0.043	414	0.01	2.7	4	530	0.22
AM21	8.5	0.003	1	560	0.037	5	0.04	4	4.3	587	0.19
P9	0.13	0.7	0.9	574	0.075	0.53	0.07	13	8	599	0.34
P14	22	0.1	3	576	0.043	1.6	0.04	8	3.5	605	0.17
FA	50	0.22	1.1	655	0.098	446	0.01	2.6	1.2	531	0.19
P11	9	0.05	3.2	573	0.013	0.14	0.13	13	9	604	0.37
P10	0.024	38	0	661	0.021	0.8	0.5	3	$\infty$	590	0.013
AM17	2.2	0.87	1.5	597	0.005	16	0.02	4	13	579	0.035
ThT	0.23	0.029	2.4	487	0.007	0.51	0.02	99	82	482	0.55

quantum yield,  $I/I_0$  and  $I/I_{nat}$ . The amyloid specificity was found to decrease in the following row: ThT  $\rightarrow$  IAH  $\rightarrow$  ABM  $\rightarrow$  AM4, IBH  $\rightarrow$  AM12  $\rightarrow$  A4, 3AM39  $\rightarrow$  AM15, AM2-23  $\rightarrow$  AM18-23, AM3, AM18, P9, P11, ISH  $\rightarrow$  AM1  $\rightarrow$  AM15-23  $\rightarrow$  AM20, A8  $\rightarrow$  AM2  $\rightarrow$  AM4-23  $\rightarrow$  P14  $\rightarrow$  A6, AM21  $\rightarrow$  P10  $\rightarrow$  AM19  $\rightarrow$  AM17  $\rightarrow$  FA.

Next, it seems of interest to discuss the nature of putative dye-fibril binding sites. The cavities, channels or grooves, abundant in the structure of amyloid fibrils, can serve as potential binding sites for benzanthrone. Several lines of evidence indicate that classical amyloid tracer ThT predominantly occupies the channels between the side chains, being the common motif in cross- $\beta$  structure and running parallel to fibril axis [57, 58]. Molecular dimensions of benzanthrone dyes ( $\sim 1.1$ – $2$  nm in length,  $\sim 0.7$ – $1$  nm in width and  $\sim 0.3$ – $0.8$  nm in thickness) are close to ThT size (long axis ca. 1.5 nm, short axis 0.6 nm and 0.4 nm in thickness, respectively). Thus, it cannot be excluded that fibril location of benzanthrone derivatives is similar to that of ThT. Therefore, we can suppose that benzanthrone associate with the channels

formed by side chain rows consisting of 4–5 amino acid residues and running in the direction of  $\beta$ -sheet propagation [58, 59].

By analogy with model membranes, to confirm the above assumption concerning the fibril binding sites of benzanthrone, at the next step of the study we focused our efforts on analyzing the effect of solvent dynamics on the spectral properties of the dyes by examining the red edge excitation shift. The REES value was calculated as the difference in emission maxima of the probes excited at 450 nm and 520 nm for all dyes, excepting AM19 and FA for which the limiting excitation wavelengths were 420 and 490 nm. In the case of protein-bound dyes the presence of REES points to the restricted mobility of both the protein polar groups and the protein-bound water molecules on nanosecond scale (i.e. during the excited state lifetime) [60]. As shown in figure 6 all benzanthrone exhibited REES with emission shift in the range 0–21 nm. According to the REES values, the dyes under study can be tentatively divided into two groups: (i) the first group (P10, AM4, AM15, AM3, AM18, IAH, 3AM39, AM2-23, AM12, IBH) displaying low REES (0–7 nm) and (ii) the second



**Figure 6.** REES dependencies for benzanthrone dyes in the presence of  $F_{acs}$  fibrils.

group (the remaining dyes) characterized by the high REES (8–21 nm) with the average magnitude of emission shift being equal to 10 nm. The observed differences in REES values of these groups can be explained by at least two reasons: (i) heterogeneity of the binding sites, and (ii) distinct changes in the dye dipole moment after excitation. To verify the assumption that only the characteristics of binding sites affect the REES values, the dipole moment of the ground and excited states were derived from the quantum-chemical calculations (table 7). It is noteworthy at this point that all dyes belonging to the second group have more polar binding sites compared to the first group: the shifts of their emission maxima in the presence of fibrils relative to those in buffer were in the range 63–13 nm. Nevertheless, there was no correlation between the REES values and the shifts of emission maxima. Thus, one can assume that the REES values of the second group of benzantrones are independent of microenvironmental polarity, being sensitive to the medium viscosity. On the other hand, the magnitude of REES for fluorophores of the first group whose binding sites are less polar, showed an inverse correlation with the shifts of emission maxima in fibril-bound state, suggesting that polarity essentially contribute to spectral and binding characteristics of these dyes. Taking into account that the wet ‘steric

zipper’ interface is abundant of polar residues lined with water molecules, with every polar residue being hydrated [61], it can be supposed that benzantrones of the second group, possessing high REES values, associate with the hydrated binding sites of amyloid fibrils. In contrast, AM4, AM15, AM3, AM18, IAH, 3AM39, AM2-23, AM12 and IBH, belonging to the first group, are most probably interact with the dry ‘steric zipper’ interface. At the dry ‘steric zipper’ interface, enriched with hydrophobic amino acids, the dye-water dipole-dipole interactions are hampered [62], thereby causing the REES weakening or even its elimination.

In our previous work 13 benzanthrone dyes were tested for their sensitivity to the variations in lysozyme fibril morphology [18]. The highest specificity to the  $F_{eth}$  amyloid aggregates was observed for AM2 and AM1, whereas A6 and A8 proved to be most specific to  $F_{ac}$  fibrils. Interestingly, A6 and A8 are representatives of benzantrones associating with wet ‘steric zipper’ interface of  $F_{eth}$ , whereas AM2 presumably interacts with the dry interface, as revealed by examining the red edge excitation shift. Furthermore, all the probes were associated with the wet interface of  $F_{ac}$ . It should be noted that benzantrone dyes possess a relatively high hydrophobicity, which originates from the absence of charged groups and the presence of aromatic benzan-

**Table 7.** Quantum chemical characteristics of benzanthrone dyes (6-31 G (d,p), GAMESS).

Dye	$\mu_g$ , D	$\mu_e$ , D	$E_g$ , Hartree	$f$	$\Delta E_e$ , $\text{cm}^{-1}$	$E_e$ , Hartree	$\varphi$ , degrees	$q_g^{\text{benz}}$	$q_e^{\text{benz}}$
A6	5.36	10.94	-979.2	0.21	22592	-979	24.5	0.14	0.20
AM1	6.44	10.24	-1188	0.264	22487	-1189	94.3	0.1	-0.022
AM2	8.03	12.06	-995.2	0.352	22665	-995	44	0.12	0.003
AM3	6.42	9.77	-956	0.289	22891	-956	-132.3	0.13	0.022
AM4	8.2	12.33	-1074	0.369	22479	-1074	-43.9	0.12	-0.01
AM12	6.99	11.03	-1147.6	0.388	23116	-1147	-45	0.16	0.059
AM15	6.76	10.23	-1035	0.292	22979	-1034	-126.1	0.12	0.012
AM18	5.79	9.55	-1147.6	0.343	22745	-1147	130	0.16	0.066
AM2-23	5.67	9.3	-3568	0.233	22955	-3568	106.1	0.062	-0.034
AM21	8.27	12.03	-956	0.358	23012	-956	33	0.15	0.057
P9	5.26	11.54	-1074	0.215	22084	-1074	23.4	0.14	0.18
AM18-23	4.95	9.18	-3721	0.298	22729	-3721	124.5	0.1	-0.025
A8	5.12	11.41	-1034.5	0.21	22028	-1034	25.1	0.14	0.19
P10	4.31	20.62	-1226	0.141	20156	-1226	26.9	0.23	0.21
P11	4.94	17.51	-1112	0.153	20156	-1112	27.3	0.18	0.23
P14	5.67	10.94	-1305	0.249	22269	-1305	15.4	0.19	0.24
A4	4.97	8.23	-1291.3	0.214	22592	-1291	-124	0.14	0.037
IAH	5.96	10.08	-1169	0.254	23124	-1169	-111.7	0.17	0.050
IBH	5.54	9.40	-1054	0.244	23149	-1054	-113.6	0.16	0.046
ISH	6.27	10.25	-1129	0.236	23366	-1129	-107.9	0.15	0.028
AM19	5.43	8.09	-877.4	0.213	23100	-877	-118	0.31	0.22
FA	3.53	5.12	-897.3	0.173	24149	-897	-130	0.44	0.38
ABM	3.83	9.82	-1015	0.204	22165	-1015	24	0.19	0.25
AM17	3.84	26.1	-1109	0.009	21931	-1109	-113.6	0.16	-0.63
3AM39	5.15	14.66	-3529.1	0.254	21326	-3539	20.5	0.22	-0.05
AM4-23	5.67	9.56	-3647	0.249	22624	-3647	-115.2	0.064	-0.05
AM20	6.76	9.84	-956	0.248	22866	-956	-118	0.11	-0.08
AM15-23	5.57	9.47	-3608	0.259	22955	-3608	-121.9	0.082	-0.02

throne moiety. These facts indicate that fibrils prepared by shaking most probably possess a denser packing of stacking  $\beta$ -sheets, creating prerequisites for fluorophore binding to dry 'steric zipper' interface. In any case, the binding parameters presented in this work confirmed our assumption about benzanthrone sensitivity to amyloid morphology.

Recently, an attempt has been made to understand how the partition ability of benzanthrone dyes depends on their physicochemical properties through analyzing a series of molecular descriptors [20]. The quantitative structure property relationship (QSPR) analysis showed high correlation between experimentally determined partition coefficients and geometrical parameters of the examined dyes, their lipophilicity and dipole moment, pointing to multiplicity of factors contributing to lipid bilayer affinity of these compounds. Therefore, it seemed of interest to develop a QSAR model for predicting the sensitivity of novel benzanthrone dyes to amyloid fibrils. For this purpose, 21 quantum-chemical descriptors for 28 tested dyes were calculated using semiempirical PM6, *ab initio* 6-31 G(d,p) methods. The obtained parameters are presented in tables 7 and 8. As seen in table 7, the changes of the dipole moments of benzantrones after excitation fall in the range 3.3–6.3 D. The highest increases of the dipole moment were observed for P10,

P11, AM17 and 3AM39 (ca ~9.5–22.3 D), while the smallest ones were found for AM19 and FA (2.7 and 1.6 D, respectively). Interestingly, this quantity showed marked positive correlation with the dye sensitivity to amyloid fibrils (see the amyloid specificity row). Indeed, the dipole moment changes determine the sensitivity of the dyes to the polarity of their local environment, thereby affecting their specificity to amyloid fibrils [63]. Comparison of the REES values with the differences  $\mu_e - \mu_g$  for each dye suggests that REES is influenced by the characteristics of benzanthrone binding sites, since the dyes with significant change in the dipole moment upon excitation (A6, P9, A8, P11, P14, ABM, AM17,  $\mu_e - \mu_g \sim 5.3$ –22.3 D) and the dyes with small  $\mu_e - \mu_g$  values (AM1, AM2, AM21, AM18-23, A4, ISH, AM19, FA, AM4-23, AM20, AM15-23,  $\mu_e - \mu_g \sim 1.6$ –4.2 D) exhibit substantial REES. Notably, for amidino benzantrones (IAH, IBH, ISH, FA) the changes in the dipole moment upon excitation are smaller than  $\mu_e - \mu_g$  for amino benzantrones. Interestingly, the values of the dihedral angle between benzanthrone moiety and its substitution in C-3 position were found to fall in the range ( $-132^\circ$ – $130^\circ$ ), with the magnitude of  $\varphi$  change being more pronounced for amidino-benzantrones, pointing to higher conjugation degree between the donor–acceptor system for amino-benzanthrone dyes [64].

**Table 8.** Quantum chemical characteristics of benzanthrone dyes (PM6, MOPAC).

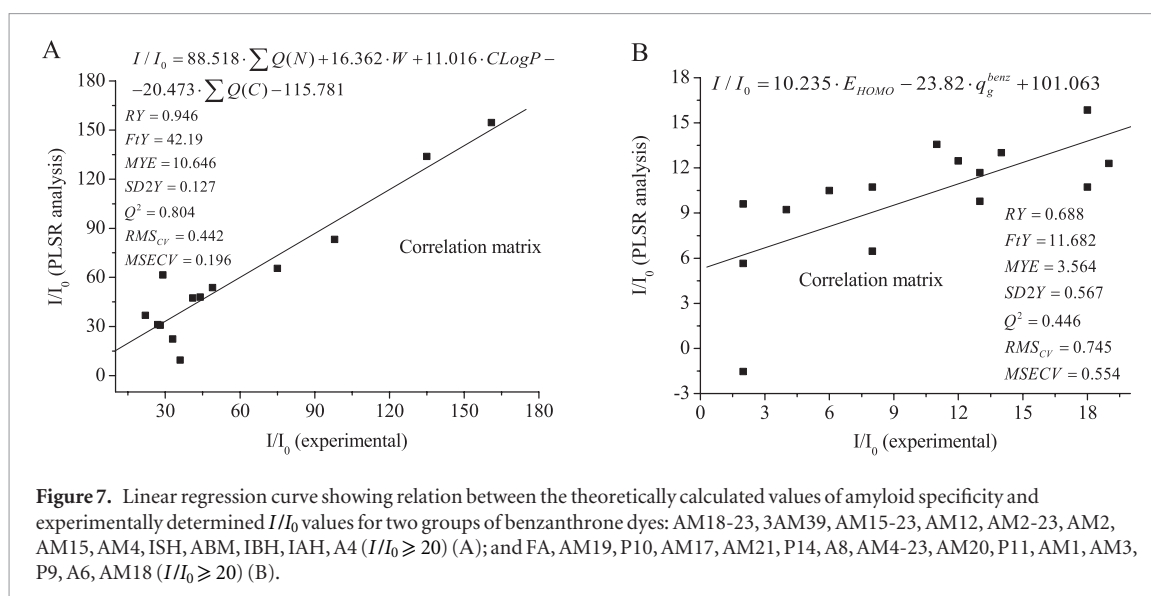
Dye	CAr, Å <sup>2</sup>	CV, Å <sup>3</sup>	E <sub>HOMO</sub> eB	E <sub>HOMO</sub> eB	L, Å	∑Q(N)	∑Q(C)	CLogP	W, Å	H, Å	P, Å <sup>3</sup>
A6	319	368	-8.0	-1.1	12.9	-0.30	-1.99	6.27	7.1	3.5	43.4
AM1	387	460	-8.5	-1.3	13.4	-0.72	-2.16	6.23	8.3	7.2	50.7
AM2	335	380	-8.3	-1.1	13.9	-0.78	-1.61	3.95	7.2	4.6	44.0
AM3	323	358	-8.3	-1.2	13.8	-0.7	-1.30	3.96	7.5	3.6	42.5
AM4	362	416	-8.1	-1.1	14.8	-0.51	-1.61	4.68	7.6	4.0	46.7
AM12	373	426	-8.3	-1.2	15.3	-0.49	-1.68	5.32	7.8	3.8	44.0
AM15	355	403	-8.1	-1.1	13.2	-0.84	-1.72	4.70	9.2	3.7	47.4
AM18	374	427	-8.3	-1.3	15.9	-0.47	-1.59	5.52	7.1	3.4	49.3
AM2-23	347	407	-8.4	-1.4	13.9	-0.74	-1.45	4.80	7.4	4.7	47.1
AM21	324	357	-8.5	-1.2	13.9	-0.95	-1.25	4.15	7.2	2.8	41.2
P9	358	411	-8.5	-1.3	15.8	-0.67	-2.00	4.98	7.4	3.8	45.5
AM18-23	395	454	-8.6	-1.5	16.1	-0.70	-1.41	5.27	7.8	3.8	54.9
A8	340	390	-8.5	-1.3	14.5	-0.63	-1.80	4.49	6.8	3.8	43.6
P10	389	455	-8.4	-1.4	16.7	-0.63	-2.26	6.62	7.6	4.7	52.3
P11	365	425	-8.5	-1.3	14.5	-0.62	-2.20	5.33	8.2	4.8	46.6
P14	430	507	-8.8	-1.4	18.2	-0.68	-2.62	6.47	7.4	4.3	54.8
A4	451	550	-8.4	-1.3	18.7	-0.43	-3.84	7.94	8.7	3.8	54.6
IAH	373	431	-8.5	-1.3	12.1	-0.42	-1.75	5.85	11.4	4.2	47.9
IBH	343	399	-8.5	-1.3	11.3	-0.42	-1.80	5.76	8.3	2.8	44.6
ISH	359	407	-8.3	-1.2	15.7	-0.42	-1.5	5.28	7.2	2.5	46.8
AM19	279	312	-8.6	-1.3	11.4	-0.98	-0.63	3.38	6.9	2.6	36.2
FA	278	308	-9.0	-1.7	11.7	-0.37	-0.51	3.87	6.9	2.2	35.8
ABM	312	359	-8.2	-1.3	11.8	-0.29	-1.36	4.52	7.9	3.4	41.9
AM17	363	411	-8.6	-1.5	13.9	-0.92	-1.55	6.07	7.6	6.3	46.3
3AM39	347	387	-8.4	-1.5	13.9	-0.68	-1.14	4.65	8.1	1.9	47.0
AM4-23	371	445	-8.4	-1.3	13.9	-0.85	-1.93	5.61	7.2	4.7	50.9
AM20	318	357	-8.4	-1.2	13.4	-0.79	-1.32	3.77	6.8	4.4	40.9
AM15-23	365	526	-8.4	-1.4	13.4	-0.75	-1.62	5.33	7.4	5.2	49.7

To correlate the amyloid sensitivity of benzanthrone dyes  $I/I_0$  with a series of calculated molecular descriptors, the partial least squares regression (PLSR) was performed using the resources of [www.vcclab.org/lab/pls/](http://www.vcclab.org/lab/pls/) [65]. Allowing for the foresaid differences between amyloid specificity of benzanthrone derivatives, it was reasonable to perform separate correlation analysis for the dyes highly sensitive to protein aggregates (with  $I/I_0 \geq 20$  (AM18-23, 3AM39, AM15-23, AM12, AM2-23, AM2, AM15, AM4, ISH, ABM, IBH, IAH, A4) and those possessing low amyloid specificity with  $I/I_0 < 20$  (FA, AM19, P10, AM17, AM21, P14, A8, AM4-23, AM20, P11, AM1, AM3, P9, A6, AM18)). To determine the parameters appropriate for PLSR analysis, initially for each group of benzanthrone dyes we performed linear correlation analysis and selected the descriptors, for which correlation coefficients with  $I/I_0$  were higher than 0.6, while the mutual correlation coefficients between the descriptors were below 0.9 [66]. It appeared that for the first group of the probes the highest linear correlation was observed between  $I/I_0$  and  $\sum Q(N)$ ,  $\sum Q(C)$ ,  $CLogP$ ,  $W$  (correlation coefficients are 0.62, -0.74, 0.77, 0.64, respectively); whereas for the second group—with  $E_{HOMO}$ ,  $E_{LUMO}$ ,  $q_g^{benz}$  (correlation coefficients are 0.64, 0.60, -0.61, respectively).

As seen in figure 7(A), the partial least square analysis for the benzanthrone dyes with high sensitivity to amyloid fibrils yields a statistically significant model (Fisher's coefficients  $FtY = 42.19$ ) with high correlation coefficient ( $RY = 0.946$ ) and low mean absolute error ( $MYE = 10.646$ ). In addition, the presented model has very high predictive ability (i.e. cross-validation correlation coefficient  $Q^2 = 0.804$ ), and satisfactory values of mean squared error of cross-validation ( $MSE_{cv} = 0.196$ ) and standard deviation of cross-validation ( $RMS_{cv} = 0.446$ ). The opposite situation was observed for fluorophores with  $I/I_0 < 20$ , since a weak correlation was found. Moreover, predictive ability was relatively low ( $Q^2 = 0.446$ ), but the  $RMS_{cv}$  and  $MSE_{cv}$  values were higher (figure 7(B)). Interestingly, the regression equation obtained for benzanthrone dyes possessing high amyloid specificity ( $I/I_0 \geq 20$ ) is more stable and predictable than the early regression model of sensitivity of ThT derivatives to A $\beta$ 1-40 and insulin fibrils [65, 66].

In the following, let's dwell on the regression model derived for amyloid-specific benzanthrone dyes in more detail (figure 7(A)). General regression equation  $I/I_0$  contains molecular descriptors with a weak mutual correlation. The impact of each descriptor  $I_j$  on the general





value of  $I/I_0$  was about 0.35,  $-0.4$ ,  $0.32$ ,  $0.49$  for  $\sum Q(N)$ ,  $\sum Q(C)$ ,  $CLogP$ ,  $W$ , respectively. Based on the analysis of contribution of each independent descriptor, we can conclude that: (i) the increase in molecular width of benzanthrone dyes  $W$  exerts a positive impact on the probe amyloid specificity to  $F_{acs}$  fibrils; (ii) the marked positive effect on  $I/I_0$  value is inherent to lipophilicity  $CLogP$ ; (iii) the dye sensitivity to amyloid fibrils is proportional to the charge of donor group of benzanthrone molecule ( $\sum Q(N)$ ) and decreases with lowering the total charge on benzanthrone skeleton ( $\sum Q(C)$ ).

A marked positive correlation between amyloid specificity and  $W$  value is indicative of significant role of steric interactions in the benzanthrone complexation with fibrils and their sensitivity to the viscosity of the environment. Indeed, it was previously demonstrated that steric interactions affect the quantum yield of ThT and its derivatives associating with amyloid fibrils [66, 67]. The increase of  $I/I_0$  value with molecular width also point to significance of aromatic interactions in benzanthrone association with lysozyme aggregates. At the same time, the observed positive correlation between amyloid specificity and lipophilicity testifies that benzanthrone binding to amyloid fibrils is controlled also by hydrophobic interactions. It is noteworthy that aromatic and hydrophobic residues significantly contribute to the ThT affinity for amyloid fibrils. Specifically, based on the results of molecular docking, Biancalana and coworkers demonstrated that ThT binds presumably to the fibril grooves formed by tyrosine and leucine [57]. An important role of aromatic interactions in ThT-fibril complexation was also reported by Wolfe *et al* [68].

Taken together, the results of QSAR analysis showed that hydrophobic ( $\sim CLogP$ ) and van-der-Waals/ion-dipole dye protein interactions (electron-donating power of the substituent at the C-3 position ( $\sum Q(N) - \sum Q(C)$ )) are more essential than aromatic and steric interactions ( $W$ ). The obtained model could be effectively employed for preliminary selection of benzanthrone dyes with high amyloid specificity.

## 4. Conclusion

To summarize, the present study has been undertaken to evaluate the applicability of the novel benzanthrone dyes to monitoring the changes in physicochemical properties of lipid membranes, as well as to detection and characterization of the protein fibrillar aggregates. All the examined dyes were found to associate with liposomes and amyloid fibrils with the magnitude of binding being dependent on the dye structure. Based on comprehensive analysis of the spectral characteristics of benzantrones bound to liposomes, we can recommend ABM, 3AM39, AM20, AM15-23, AM4-23, AM2-23 and P9 for examining membrane-related processes coupled with polarity changes. The most prospective amyloid tracers were selected from 28 studied compounds. The results of quantum-chemical calculations and QSAR analysis revealed a positive correlation between amyloid specificity of benzanthrone dyes and a series of descriptors, such as lipophilicity, molecular width of the probe and electron-donating power of the substituent at the C-3 position. A statistically significant model was obtained for predicting the sensitivity of novel benzanthrone dyes to amyloid fibrils.

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