1	Computational Calculation of Dissolved Organic Matter
2	Absorption Spectra
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22	Synopsis

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The absorption of dissolved organic matter was investigated using computational models. The results indicate that it was possible to approximately recreate DOM spectra by summing individual 24

25 chromophores spectra.



30 ABSTRACT

31 The absorption spectrum of dissolved organic matter (DOM) is a topic of interest to environmental 32 scientists and engineers, as it can be used to assess both the concentration and physicochemical 33 properties of DOM. In this study, the UV-Vis spectra for DOM model compounds were calculated 34 using time-dependent density functional theory (TD-DFT). Summing these individual spectra, it 35 was possible to recreate the observed exponential shape of the DOM absorption spectra. 36 Additionally, by predicting the effects of sodium borohydride reduction on the model compounds, 37 then calculating the UV-Vis absorbance spectra of the reduced compounds, it was also possible to 38 correctly predict the effects of borohydride reduction on DOM absorbance spectra with a larger 39 relative decrease in absorbance at longer wavelengths. The contribution of charge-transfer (CT) 40 interactions to DOM absorption were also evaluated and the calculations showed that intra-41 molecular CT interactions could take place, while inter-molecular CT interactions are proposed to 42 be less likely to contribute.

43 INTRODUCTION

44 The absorption spectrum of dissolved organic matter (DOM) is one of its most studied 45 properties. In the early stages of DOM research, absorption spectra were used as a proxy for the concentration of DOM in surface waters.¹ Over the past decades, the use of the absorption spectra 46 47 has evolved from a quantitative surrogate for concentration, to a qualitative descriptor for 48 physicochemical reactivity and biogeochemical origin. For example, the specific UV absorption 49 at 254 nm (SUVA254) has become one of the most widely used optical parameters to describe DOM 50 physicochemical reactivity and character, in settings from assessment of DOM reactivity during drinking water treatment all the way to biogeochemical source.²⁻⁵ In addition, a wide range of 51 52 metrics derived from the absorption spectra have been developed and implemented by the 53 environmental science and engineering community, like the spectral slope and E2/E3 ratio. The 54 spectral slope describing the approximate exponential decline in chromophoric DOM (CDOM) 55 absorption with increasing wavelength is widely used for tracing changes in the chemical composition of CDOM.^{6; 7} E2/E3 is the ratio of absorbance at 250 nm to at 365 nm and often used 56 as an indicator for humification and molecular weight of humic substances.⁸⁻¹⁰ The lower values 57 58 of E2/E3 ratio may be indicative of the presence of structures with higher molecular weight, 59 aromaticity, and humification degree.

Not all molecules in DOM interact with light. CDOM is defined as the light-absorbing fraction of DOM ([DOM]>[CDOM]) and is the main absorber of light in environmental systems.¹¹⁻
¹⁴ Thus, employing absorbance as a surrogate for the physicochemical properties of DOM (e.g., molecular weight) implies the sometimes arbitrary assumption that the behavior of the chromophores represents the behavior of either the most important components for a specific application (e.g., relationship between $SUVA_{254}$ and formation of disinfection byproducts), or directly correlates with how the rest of the mixture behaves.

67 The exponential shape of DOM UV-Vis absorption spectra is widely observed regardless 68 of source and usually is featureless, but some shoulders or peaks can be observed for example for organic matter from atmospheric sources,¹⁵ from algae provenance,¹⁶ or through Gaussian 69 decomposition of DOM spectra.¹⁷ Two alternative theories have been proposed to explain DOM 70 71 absorbance spectrum. The first one postulates that the absorbance is the sum of the individual 72 DOM chromophores absorbance. The second model postulates that the absorbance is the sum of 73 two terms: 1) the sum of the absorbance of the individual chromophores and 2) a term due to 74 charge-transfer (CT) interactions between electron donor (mostly phenols) and acceptor (mostly quinones and ketones) moieties of DOM, that contributes to the absorption at wavelengths \geq 75 76 350nm. The effects of sodium borohydride reduction on DOM UV-Visa absorption spectra were 77 often used to probe into the CT model. The reduction transforms quinones and ketones functional groups into phenols, decreasing so the amount of CT interactions.¹⁸⁻²¹ We refer the reader to recent 78 79 publications, which describe alternative and complementary mechanistic interpretations for the shape of the UV-Vis spectra of DOM.²¹⁻²⁷ 80

The aforementioned CT interactions were assumed in the literature to be *intra* molecular interactions due to DOM absorption spectra being concentration independent, but CT interactions can in principle be due to both *intra* and *inter* molecular interactions. *Inter* molecular CT interactions were the scope of a previous work that indicated formation of π -stacking interactions in Suwannee river fulvic acid (SRFA) models, with subsequent effects on the calculated absorbance spectra.²⁸ These results offer computational confirmation of CT interactions within DOM, suggesting that they contribute partially to its optical properties. However, it was shown

88 that the formation of π -stacking clusters occurred in water, but that switching the solvent to non-89 polar ones resulted in disruption of the π -stacking structures as well as the whole molecular assembles.²⁸ This result further highlights the dynamic nature of these assemblies. Consequently, 90 91 in non-polar solvents the formation of CT complexes via π -staking clusters is insignificant and 92 cannot contribute to absorption maximum ($\lambda_{a,max}$) of UV-spectra. As DOM spectra are mostly solvent independent,²⁴ this raises the question of whether these inter molecular CT interactions 93 94 contribute significantly to DOM absorbance. *Intra* molecular CT interactions were postulated to 95 exist based on spectrofluorometric observations and also on the effects of sodium borohydride reduction on DOM absorbance and fluorescence properties.^{18; 23} These observations rely on bulk 96 97 DOM properties and with limited insight on what is happening at the molecular level. The 98 possibility of calculating the UV-Vis spectra of molecules allows to test for the possibility of *intra* 99 molecular CT interactions and on which conditions such interactions should take place.

100 Recently it was shown that time - dependent density functional theory (TD-DFT) method 101 reproduces very accurately optical susceptibilities for a wide range of organic compounds that could be found in DOM.^{29; 30} TD-DFT is a first-principles method without any empirical 102 103 spectroscopic parameters. The calculation of transition energies from time-dependent density functional linear response theory has been described in details by Neese.³¹ The linear response 104 105 approach directly yields the transition energy rather than the total energies of the ground- and 106 excited states. Thus, the excited states themselves are never explicitly calculated. Rather, their 107 energies are deduced from the poles of a frequency dependent ground state property. Thus, for the 108 ground state all properties can be calculated from the analytic derivatives of the total energy. The 109 same procedure applies to the excited states as well. The total energy of the excited state is simply

the sum of the ground state energy and the transition energy predicted by the TD-DFT procedure.
The derivatives of this total energy then define all excited state properties.

112 The main objective of this study was to use computational chemistry tools to conduct a 113 detailed evaluation of the UV-Vis spectra of DOM. Specifically, TD-DFT was used to calculate 114 the UV-Vis spectra for a selection of chromophores chemically similar to those chromophores that 115 could be expected in DOM. Calculated spectra were summed together in a superposition-type 116 approach for comparison to DOM absorbance spectra. Additionally, we evaluated whether certain 117 concentration conditions could give rise to inter molecular CT interactions being concentration 118 independent. Finally, TD-DFT was used to calculate the spectra of a selection of compounds that 119 should exhibit intra molecular CT interactions to evaluate in which conditions these CT 120 interactions should occur.

121 MATERIALS AND METHODS

122 Generation of DOM Models

123 In this work, five models (numbered 1 to 5) representing Suwannee River fulvic acid 124 (SRFA) were used. Each model contains a number of molecules, each assigned to a letter (e.g., for 125 model #1, molecules a to m, see Table S1). SRFA was chosen as representative of river headwater 126 allochthonous DOM from the International Humic Substances Society, and because it is in all 127 likelihood, the most studied DOM. The models were created on the basis of SRFA chemical characteristics³² using principles proposed by Vialykh et al.^{33; 34} The algorithm of fragment 128 selection and screening Criteria of each model were described in Vialykh et al.²⁸ Specific chemical 129 130 species were selected for use in this study. These structures are shown in Table S1 of the 131 Supporting Information (SI). The number of each fragment in the models was equal one. Table S2 132 includes the elemental composition for each model, as well as the lowest, and highest molecular 133 weight for the components in the models. Lastly, the mean molecular weight for the models is 134 included. The number average molecular weight for SRFA has been estimated at 724 ± 68 Da based on vapor pressure osmometry,³⁵ 610-760 Da based on hydroxyl radical reactivity,³⁶ 968 Da 135 based on size-exclusion chromatography data³⁷ while the latest high-resolution MS data showed 136 that the range of molecular masses of aquatic humic substances is 290-600 m/z.³⁸ Though we 137 138 attempted to build the models that would reflect the latest data on average molecular weight of 139 molecules present in SRFA, the computational restrictions (time and cost) force us to generate 140 models with the mean molecular weight on the lower end of these estimates (~300-400Da). Larger 141 chromophores can potentially move the absorption band more into the visible as DOM size 142 fractionation experiments indicate that the larger fraction absorbs relatively more visible light than the smaller fractions.³⁹ 143

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145 Absorbance Spectra Calculations

First, the geometry optimization of each molecule was performed by DFT calculations. The properties of a many-electron system can be determined by using functionals, i.e., functions of another function. In the case of DFT, these are functionals of the spatially dependent electron density.

Further TD-DFT calculations were performed to obtain UV spectra of selected molecules. TD-DFT is a quantum mechanical theory that allows to investigate the properties and dynamics of many-body systems in the presence of time-dependent potentials, such as electric or magnetic fields. The effect of such fields on molecules can be studied with TDDFT to extract features like excitation energies, frequency-dependent response properties, and absorption spectra.

The ORCA 4.0.1 package⁴⁰ was used to carry out the geometry optimizations and spectral 155 156 calculations for individual molecules. Calculations were performed for both fully protonated 157 molecules, that correspond to pH 1-2, and with deprotonated carboxylic groups, which corresponds 158 to pH~5. The geometry optimizations were performed by using the B3LYP functional with def2-159 TZVP basis set in water solvent with SMD solvation model. Tight self-consistent field 160 convergence criteria were set without symmetry constraints. The RIJCOSX approximation was 161 used with auxiliary basis def2/J to accelerate the calculations without notable loss of accuracy. The 162 TD-DFT and sTD-DFT calculations were performed by using B3LYP or BP86 functional on the 163 DFT-optimized structures with def2-TZVP or aug-cc-pVDZ basis sets. The RIJCOSX 164 approximation was employed to speed up the calculations. SMD solvation model was used to 165 conduct the calculations in water solvent. The calculation of the charge transfer in the excited states has been performed by the TheoDORE program package.^{41;42} To plot the absorption spectra 166

167 for individual molecules, Avogadro program⁴³ was used with Gaussian line shape and peak width 168 20. The increase in peak width above 20 resulted in smoothing the curve and disappearing peaks, 169 e.g., peak at 260 nm for flavone (Figure S1). In order to plot the absorption spectra for a mixture 170 of several molecules, the Octave⁴⁴ script was used to calculate Gaussian line shape of absorption 171 spectra with peak width 20. The calculated spectra of coumarin and flavone was compared to their 172 measured spectra to verify the accuracy of the calculation (Figure S1).

UV-Vis spectra for the different models were calculated based on the superposition model, *i.e.*, we assumed that there were no interaction between individual molecules that would lead to additional absorption bands, and the overall spectrum of the model was calculated as a sum of spectra of individual molecules. Additionally, a global model containing all molecules was also calculated (see Table S1 for the structures presents in each model). For different pH the spectra were calculated separately.

Many molecules were used to create a computational model. Some of these molecules are difficult to purchase in pure form or not available commercially, making the comparison of the calculated spectra with spectra of real mixture of molecules not possible.

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Chemicals and Solutions

183 SRFA (catalogue number 1S101F) was obtained from the International Humic Substance 184 Society (IHSS, St-Paul, MN, USA). Coumarin and flavone (purity min. 99%) and HPLC grade 185 methanol were obtained from VWR. All solutions were made using ultrapure water (resistivity 186 18.2 M Ω cm) obtained from a Sartorius Arium pro dispenser or equivalent. The SRFA solution 187 was made according to the methods presented in Leresche et al.⁴⁵ UV-Vis absorbance spectra were 188 measured on a Cary 100 Bio UV-visible spectrophotometer using a 1 cm path length quartz 189 cuvette. 190

191 **Borohydride reduction**

To simulate the effects of borohydride on our model compounds, we applied the following transformation rules: acetophenone to 1-phenylethanol, benzophenone to the corresponding alcohol, esters to the corresponding cleaved alcohol, anthraquinone to anthracene, unconjugated ketones and aldehydes to the alcohol, and conjugated ketones and aldehydes to a mix of conjugated and unconjugated alcohols. Please see Table S1 for the structures of the reduced compounds.

198 **RESULTS & DISCUSSION**

199 Influence of Functional Groups and Chemical Structures on UV-Vis Spectra

200 It is known that the presence of different functional groups or molecular fragments affects UV-Vis spectra,⁴⁶ and because DOM contains a range of functional groups, this can extend the 201 202 absorption of chromophores into the visible region. Light absorbed in the UV-Vis range induces 203 an electron shift to one of the empty molecular orbitals. For the wavelength range above 220 nm, 204 most absorption occurs from extensive conjugation that decreases the energy gap between the 205 highest occupied molecular orbital (HOMO) and the empty molecular orbitals. For example, 206 benzene has a $\lambda_{a,max}$ at ~254 nm, whereas for naphthalene the value is 275 nm, and 356 nm for 207 anthracene (the sequence of linear poly-aromatic hydrocarbons has a shift in wavelength max of 208 \approx 75 nm per ring). In the case of phenol, $\lambda_{a,max}$ is 270 nm, and 262 nm for methylbenzene. As we 209 consider the UV-Vis spectrum of DOM, and the fact that it extends into the visible, it is expected 210 that single ring components will dominate the absorption below 300 nm while in the visible multi-211 ring chromophores are the main light absorbers. This can be observed in the present data set (Table 212 S4 and Figure S2) were most molecules absorbing at wavelengths longer than 400 nm have multi-213 ring chromophores while molecules having their $\lambda_{a,max}$ in the 250-300 nm range are mostly single-214 ring compounds.

Flavones are plants pigments and potentially a class of DOM chromophores.⁴⁷ Spectra calculation for a series of flavones molecules substituted with various electron donating or withdrawing groups showed that the substituents shift the $\lambda_{a,max}$ by up to 120 nm (Table S5 and Figure 1). The Hammett σ constants (that relate a functional group to its electron donating or withdrawing character) vs $\lambda_{a,max}$ relationships follows a U-shape non-linear relationship with shorter $\lambda_{a,max}$ values for substituents having σ values close to 0 and longer $\lambda_{a,max}$ values for electron 221 poor and electron rich substituents. This observation is similar to the one made in a study for substituted squaraine dyes⁴⁸ but differs from the results of a study on substituted triphenylamines 222 were a linear relationship was observed between σ and $\lambda_{a,max}$.⁴⁹ The U-shape can be rationalized 223 224 by the absorbance spectra having a relatively important absorption band around 300 nm and for 225 some of the substituents to present additional absorption band at longer wavelengths that shift 226 $\lambda_{a,max}$ (Figure 1). Note that if there are more than one absorption maxima, $\lambda_{a,max}$ corresponds to the 227 longest wavelength maxima. The wavelength of maximum absorption of the 300 nm feature ($\lambda_{\rm M}$) 228 can be seen to follow a linear relationship with the σ substituent (Figure 1) and it can be concluded 229 that the observed U-shape relationship between $\lambda_{a,max}$ and the Hammett σ substituent comes from 230 the appearance of additional absorption bands for some of the compounds.

231 Analysis of UV-Vis spectra by the region of $\lambda_{a,max}$ revealed that molecules containing 232 quinone groups have $\lambda_{a,max}$ above 400 nm (Table S4). For example, 2,3-dicarboxy-1,4-233 benzoquinone has a $\lambda_{a,max}$ at 611 nm in deprotonated form and at 506 nm in protonated form. The 234 presence of three conjugated aromatic rings (with or without functional groups) results in $\lambda_{a,max}$ 235 above 400 nm as well. The presence of ketone groups conjugated with an aromatic ring result in 236 $\lambda_{a,max}$ by the 300-400 nm region. Presence of two or more conjugated double bonds and COOH 237 group also results in $\lambda_{a,max}$ between 300-400 nm. Molecules containing a single aromatic ring with 238 different functional groups (except ketone) tend to have $\lambda_{a,max}$ between 250-300 nm. $\lambda_{a,max}$ in the 239 same range was observed for molecules with two conjugated double bonds.

DOM being a mixture of compounds, its absorbance spectrum reflects the spectra of the individual components. For the aforementioned superposition model, DOM spectrum is simply the sum of the spectra of the individual components and one can model DOM spectrum by averaging



243 the individual components. For the CT model, one must add to the spectra average a term 244 corresponding to CT interactions. Investigations on these CT interactions are presented below.

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Figure 1. Calculated absorbance spectra for the substituted flavones in the A) meta and in the B) para position. Absorption maximum wavelength ($\lambda_{a,max}$) and absorption maximum wavelength for the peaks centered around 300 nm ($\lambda_{\rm M}$) of a series of flavones backbone substituted molecules in the C) meta and in the D) para position. The substituents are presented in Table S4.

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251 Simulated UV-Vis Spectra of the SRFA models

252 The UV-Vis spectra of the models were created by averaging the UV-Vis spectra of the 253 models' molecules (see above). Since the real samples of DOM contain thousands of individual 254 molecules and our models contained 8-21 molecules and 65 for the global model, the quantitative 255 comparison is not feasible. Thus, only qualitative comparison between calculated and experimental 256 UV-spectra was performed. UV-Vis spectra calculated for the models decayed with increasing 257 wavelength. The shape of the spectra was close to exponentially decaying curve as the absorption

258 spectrum of SRFA (Figure 2). Although all models include molecules that absorb light in a region 259 above 400nm, the calculated UV-Vis spectra for models #4 and 5 were the closest to the 260 experimental data, whereas spectra for model #2 had the largest discrepancy with experimental 261 data. The observed variation in the UV-Vis spectra between models might be a result of the mean 262 molecular weight of the models. Thus, for models #4 and 5 these values were 497 and 391Da 263 accordingly, since significant part of the molecules included in the models had large molecular 264 weight (above 400Da) and in the structure of each molecule there were several conjugated groups 265 that absorb light, which produces a broadening of the individual absorption peaks and additionally 266 contributes to increase of absorption at longer wavelengths. The opposite could be said about 267 model #2 that is composed mainly from small molecules with molecular weight below 260Da. 268 Though there were fragments that absorb above 400 nm (pH1) and 600 nm (pH5) the number of 269 fragments that did not absorb light above 250 was significantly higher than in other models, e.g., 270 seven fragments in model#2 vs one fragment in model #4. The spectra for the other molecules in 271 model#2 were presented as a line with one or two individual sharp absorption peaks, rather than a 272 smooth exponentially decaying curve. As a result, a larger discrepancy between UV-spectra 273 calculated for model#2 and experimental data was observed in the range between 250-400 nm.



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Figure 2. UV-Vis spectra of Suwanee River Fulvic Acid (SRFA) (dashed line) measured at pH 7,
of the 5 simulated models (continuous lines) and of the global model (that contains all the 5
models).

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It can be observed that the models' absorbances decrease at longer wavelength and that absorption peaks are observable. These peaks are likely due to the absorbance of a single model compound and would likely vanish if many more compounds would be included in the models. Indeed, the global model that includes all the models does not present pronounced peaks (Figure 2). Comparing the model's spectra to the one of SRFA, it is possible to see that the model's absorbance decreases faster than SRFA with increasing wavelength. This discrepancy could be due to the mean molecular weight of the models being lower than the one of SRFA (Note, for computational time reasons, it is not actually possible to calculate the absorbance of molecules having molecular weights above \approx 800 Da). The high molecular weight fractions of SRFA are known to absorb relatively more in the visible than the low molecular weight fractions³⁹ and the calculated models are probably more representative of the low molecular weight fractions of SRFA.

292 Evaluation of Charge-Transfer interactions

293 Inter molecular charge-transfer interactions

294 It should be noted that the absorbance spectra calculated in the present article only include 295 the individual components' contributions and would not integrate any inter molecular CT 296 interactions. The current working model for DOM charge-transfer interactions is that they exhibit structural similarities to eumelanin (a pigment that protects the skin against sunlight ⁵⁰) in terms 297 298 of their three-dimensional configuration. The absorbance of eumelanin-like compounds presents 299 a CT complex absorbance in the visible spectra that also has the particularity to be concentration dependent.⁵¹ By contrast, DOM absorbance spectra are concentration independent. This raises the 300 question of whether conditions exist such that inter molecular interactions do not exhibit a 301 302 concentration dependence. This was evaluated using the following model: the equilibrium 303 equation for the formation of a CT complex between a DOM electron acceptor and donor can be written as $^{A+D} \xrightarrow{CT}$ with A and D an electron acceptor and donor respectively, CT the charge-304 305 transfer complex and K the corresponding equilibrium constant. For CT to be [DOM] independent, 306 [CT] should be directly proportional to [DOM]. Note, we chose a criterion where less than 10% 307 variation is not considered to be a significant change, this choice is arbitrary, but we do estimate 308 that this corresponds to the limit of what could be detected by absorbance measurements of DOM

for wavelengths \geq 400nm. See Text S1 for the derivation of the reaction formula and Table S7 for the presentation of the evaluation results for several [A]₀/[D]₀ ratios and various values of *K*.

311 The evaluation results indicates that for the $[A]_0/[D]_0$ ratios values of 0.01 and 0.1, inter 312 molecular interactions would exhibit a concentration dependence for equilibrium constant below ≈ 1000 (corresponding $\Delta G = -17.1$ kJ mol⁻¹) while above such number the concentration 313 314 dependence would be negligible. For the equimolar $[A]_0/[D]_0$ ratio, *inter* molecular interactions 315 would exhibit a concentration dependence for equilibrium constant below ≈ 10000 (corresponding 316 $\Delta G = -22.8 \text{ kJ mol}^{-1}$). For comparison with these ΔG values, a water hydrogen bond has an energy of -21.6 kJ mol⁻¹ (Note, this value is per mol of water molecules and corresponds to a value of 317 318 -10.8kJ mol⁻¹ of hydrogen bond).⁵² Overall, these calculations indicate that in order to not exhibit 319 concentration dependency, two molecules have to be bonded through either two bonds with 320 strength equivalent to hydrogen bonds or more than two less energetic bonds. As DOM spectrum 321 is concentration independent, this exercise indicates that if there is any inter molecular CT complexes in DOM, the molecules need to interact relatively strongly. Our previous work suggest 322 that the probable main mechanism for intermolecular CT is π -stacking.²⁸ For usual H-bonds, 323 324 namely X-H----Y with nitrogen or oxygen as X and Y, the binding strength is about -17 to -27 kJ mol^{-1.53} At the same time π - π stacking between two single aromatic rings is -10 to -12 kJ mol^{-1,54} 325 326 clearly in usual case H-bond is evidently stronger. However, there are other situations where Hbond may also be quite weak (e.g. C-H----O, whose binding energy is lower than -4 kJ mol⁻¹), 327 328 while π - π stacking may also be quite strong (e.g. cyclo[18]carbon dimer, the binding energy of which is about -38 kJ mol⁻¹).⁵⁵ Additionally, the combination of both π - π stacking and H-bonds 329 330 could prevent π -stacking aggregates from dissociation.

332 Intra molecular charge-transfer interactions

333 To assess for the possibility of *intra* molecular CT, the spectra of molecules that could 334 undergo intra molecular CT were calculated (Figure 3A). These molecules were constructed using 335 a flavone or a coumarin tetra-methyl substituted backbone. Then each methyl position was 336 alternatively replaced by a $(CH_2)n$ -hydroxybenzene (n=2 or 3) for the coumarin backbone or 337 $(CH_2)n$ - dihydroxybenzene for the flavone backbone. As negative controls, $(CH_2)n$ -338 methylbenzene and (CH₂)n-dimethylbenzene were substituted to the hydroxybenzene and 339 dihydroxybenzene respectively. Additionally, the spectra of molecules known to exhibit unconjugated CT interactionss²² were also calculated (molecules 8, 9a and 9b). In these molecules 340 341 the CT might occur due to π -staking interactions between non-conjugated aromatic rings. 342 Molecules used in biological imaging that possess fluorescence properties assigned to conjugated intra molecular CT complexes^{56; 57} were also calculated (Figure 3B). The donor and acceptor in 343 344 these molecules are connected via conjugated backbone (e.g., alternation of single and double bonds) resulting in CT via delocalized π -orbitals. These molecules possess an electron acceptor 345 346 part (1-benzopyran-4-one) and an electron donor part (benzene ring) that was substituted in the 347 para position with diverse electron rich ($N(CH_3)_2$, OH, OCH₃) functional groups and for controls 348 an electron poor functional group (NO₂) and a H substituent.



Figure 3. Structure and calculated absorbance spectra of the compounds investigated for *intra* molecular charge-transfer (CT) interactions. A) molecules investigated for unconjugated *intra* molecular CT interactions. B) molecules investigated for conjugated *intra* molecular CT interactions. Note, Figure S4 presents additional compounds to Figure 3A substituted on the substitution of the other methyl groups of the coumarin as well as similar compounds based on a flavone backbone. Figure S5 presents additional compounds to Figure 3B having the coumarin backbone hydroxyl group replaced by either a hydrogen atom or a -OCH₃ group.

359 The absorbance spectrum of molecule 8 measured in acetonitrile presents an absorbance band between 500-600 nm that was assigned to a CT complex.⁵⁸ The calculations fail to capture 360 361 this 500-600 nm band (Figure 3A) with the calculated spectra only slightly different from the 362 spectra of the dihydroxybenzene and quinone subunits (Figure S3). The reasons for this 363 discrepancy could be the difference in solvent; water being a protic polar solvent can make 364 hydrogen bonds with the hydroquinone and quinone moieties of molecule 8 that acetonitrile would 365 not. The water hydrogen bonds would compete with the formation of *intra* molecular CT complex, 366 and it is plausible that molecule 8 only exhibits CT properties in an aprotic solvent. An alternative 367 explanation is that the assignment of the CT complex to *intra* molecular was not correct and that 368 the observed 500-600 nm $\lambda_{a,max}$ was due instead to *inter* molecular CT complex. The spectrum 369 measured in dioxane of molecules 9 presents a weak absorption band centered around 500 nm 370 attributed to *intra* molecular CT interactions while for molecule 9b a strong CT absorption band

is visible at 350-600nm.⁵⁹ The calculated spectrum of molecule 9b presents an $\lambda_{a,max}$ at λ =630 nm 371 372 that can be attributed to intra molecular CT interactions (Figure 3A) while for molecule 9 a less intense $\lambda_{a,max}$ can be observed at λ =470 nm that can also be attributed to *intra* molecular CT 373 374 interactions. We also calculated molecules similar to molecule 9b with various carbon atoms (n=2-375 9) between the two rings, the increase in carbon atoms increased the distance between the two 376 rings. The results of the calculation are presented in Figure S3 and indicate that CT is observed 377 only for n= 2 or 3, when the distance between two rings less than 5Å while for n \geq 4 no CT is 378 observed.

379 For the coumarin and flavone backbone molecules grafted with hydroxy or dihydroxy-380 benzene groups, none exhibited non-conjugated CT absorption band (Figure 3A and S4). One 381 possible explanation is that the formation of *intra* molecular CT complexes is competing with the 382 water hydrogen bonds and that the formation of such CT is not favorable in water. As molecule 383 9b presents some CT absorption properties, it could indicate that to present CT properties in water, 384 molecules have either to be constrained in their shape in a way that allows π -stacking formation 385 or that intra molecular CT complexes have to outcompete water hydrogen bonds. Overall, the 386 calculations indicate that intramolecular CT between non-conjugated molecular parts would 387 probably depend on π -stacking, between two or multiple aromatic rings in close proximity.

For the flavone substituted molecules investigated for conjugated *intra* molecular CT complex, (Figure 3B and S5), $\lambda_{a,max}$ was of ~455 nm for the N(CH₃)₂ group, ~370 nm for the OH and OCH₃ groups and ~510 nm for the control electron withdrawing NO₂ group and ~356 nm for the H group with mean CT contribution to the singlet excited state of 69%, 44%, 50%, 77% and 30% for the N(CH₃)₂, OH, OCH₃, NO₂ and H group respectively. These results indicate that intramolecular CT interactions are affected by the substitution of electron donor or acceptor groups 394 in conjugation with the chromone moiety. The occurrence of an electron donating –OCH₃ or -OH 395 groups in the para-position of the phenyl ring mildly enhances the charge transfer by 4-25% (Table 396 S8) compared to control molecule where R- is hydrogen. The spectral response is qualitatively 397 similar to the hydrogen control molecule ($\lambda_{a,max}$ red shift of up to 44nm). The presence of an 398 electron donating $-N(CH_3)_2$ group causes an increase of CT up to 50% and red shift of $\lambda_{a,max}$ up to 399 134 nm resulting in a new spectral peak in the visible region above 400 nm. Interestingly electron 400 withdrawing -NO₂ group in the same position opens competing charge transfer channels that cause 401 a new spectral peak with $\lambda_{a,max}$ 480-510 nm and higher CT contribution (75-81%). Thus, -OCH₃ 402 or -OH groups didn't induce absorption in the visible range except for the N(CH₃)₂ group while 403 the control molecule (NO₂) possessing two electron accepting parts showed red shifted absorption. 404 This result is similar to what was observed for substituted β -enaminones where the molecule 405 having two electron accepting parts had an absorption spectrum that was red shifted compared to the molecule having an electron accepting and an electron donating part.⁶⁰ 406

407 Effects of sodium borohydride reduction

408 *Sodium borohydride reduction*

Sodium borohydride DOM reduction has been used by many ^{18; 20; 21; 61-70} for investigations on DOM absorption and fluorescence properties. The calculation of the borohydride reduction products of the compounds selected in our models is possible. This allows us to evaluate the effects that borohydride reduction should have on our model's absorbance properties, which are based uniquely on the superposition of individual chromophores. One difficulty doing such exercise is the prediction of the correct reduction products.

415 Borohydride reduction was presented originally in the environmental chemistry literature 416 as mainly reducing carbonyls groups to alcohols within DOM.^{18; 71} This is probably too simplistic 417 since additional side reactions can take place, e.g., for α , β -unsaturated aldehydes and ketones, the 418 reduction of double bonds adjacent to the carbonyl group was seen to take place leading to the formation of a mixture of the saturated and the unsaturated alcohol.⁷² Esters are potentially also 419 reduced under large borohydride excess conditions.⁷³ The experimental conditions proposed for 420 DOM reduction are a 25-fold borohydride mass excess for 2-4 days of reaction,⁷¹ which probably 421 422 allows for reaction of some ester moieties within DOM. Identification of the borohydride reduction products confirms that the reduction is not limited to the conversion of carbonyls groups to 423 424 alcohols. For example, three rings opening products (one unsaturated and two saturated) are obtained upon coumarin's borohydride reduction,⁷⁴ anthrones or anthracenes are obtained upon 425 reduction of substituted anthraquinones,⁷⁵ and dihydro esters are the reduction's products of a 426 series of methyl (p-substituted phenyl)-acrylates.⁷⁶ We adopted the rules presented in the 427 experimental section to calculate the different products of borohydride reduction. The molecules 428 429 before and after reduction are presented in Table S1.

430 *Individual molecules*

431 In most cases, borohydride reduction of individual molecules leads to a blue shift in their 432 UV-Vis spectra (Figure 4), however, a red shift after reduction was observed for several molecules 433 (e.g., molecules c and m in model #1; molecules e and f in model #3 and molecule k in model #5, 434 see Table S1 for the molecules before/after reduction). The blue shift in $\lambda_{a,max}$ varies for different 435 molecules and was between <1 nm to 317 nm at pH 1 and between 8 nm to 364 nm at pH 5, 436 whereas the red shift was between 3.5 nm to 66 nm at pH 1 and between 16 nm and 29 nm at pH 437 5. These results are similar to the one observed upon borohydride reduction of a series of model 438 quinones where a blue shift in $\lambda_{a,max}$ was observed for all but one of the quinones (anthraquinone-2.6-disulfonate) following borohydride reduction.¹⁸ 439



Figure 4. Effects of sodium borohydride reduction on the absorbance spectra. A) structure and calculated absorbance spectra of the parent molecules (11 and 12) and of the borohydride reduced products (11a and 12a). B) Calculated UV-Vis absorbance spectra of the global model before (black) and after (red) borohydride reduction (at pH5) and absorbance ratio (blue, I/I₀). The absorbance ratio was smoothed using a 30 points (\approx 25nm) adjacent averaging algorithm.

447 Effects of sodium borohydride on the spectra

For the global model, a blue shift after reduction was observed with a $\lambda_{a,max}$ change of 25 448 nm (pH1) and 123 nm (pH5) (Table S9). At the same time for models #1 and #3 a red shift in $\lambda_{a,max}$ 449 450 was observed at pH1. The largest blue shift in $\lambda_{a,max}$ (237 nm at pH5) was observed in model #2 451 which were composed from small fragments with molecular size between 120-260 Da, whereas 452 for models #4 and #5, composed mostly from fragments with molecular weight above 350 Da, the 453 blue shift was much lower, 24 or 69 nm and 12 or 60 nm at pH 1/pH5, accordingly. Thus, 454 borohydride reduction could significantly decrease the $\lambda_{a,max}$ of small molecules due to destruction 455 of quinone groups, whereas the reduction had lesser effects on the $\lambda_{a,max}$ of molecules with more 456 complex structure and molecular weight above 400 Da.

457 UV-Vis spectrum calculated for the global model (Fig 3B) indicates that borohydride 458 reduction decreased the absorbance, a trend similar to the experimental results.¹⁸ The calculated 459 relative decrease in absorbance (I/I₀) is similar to the measured one with a decrease that is more 460 marked at longer wavelength in the 200-400 nm region. The only area that diverges from this

461 behavior is in the 480-520 nm range. The divergence comes most probably from the very low 462 intensity measured in this wavelength range. Overall, our results indicate that the superposition 463 model could also explain well the effects of borohydride reduction on DOM absorbance spectra 464 not only the CT model as postulated in Ma et al.¹⁸

465 **ENVIRONMENTAL IMPLICATIONS**

466 The work presented here continues the development of computational tools to assess the 467 origins of the optical properties of DOM. Computational approaches offer an advantage when 468 evaluating the superposition model for understanding DOM's optical properties since transitions 469 can be determined computationally for compounds not commercially available. The work 470 presented herein indicates that a superposition of chromophores is largely able to reproduce the 471 absorbance spectral shape and effects of borohydride reduction. These spectral properties have 472 previously been argued to be consistent only with a charge-transfer model, and a recent study 473 highlights that this behavior is also consistent with the behavior of a model charge-transfer 474 complex (quinhydrone).²⁷ Therefore, the absorption properties of DOM and their responses to 475 borohydride reduction appear to be consistent with both a superposition model and charge-transfer 476 model.

One aspect of the relationship between CDOM and DOM that still requires clarification is the overall mass ratio between both, [CDOM]/[DOM]. It is expected that this ratio will be less than unity, as not all components of DOM will absorb. In the case of SFRA, we can use the developed models to ascertain what this ratio would be. Given that DOM is mostly quantified in terms of carbon concentration, we use carbon ratios to ascertain the value for [CDOM]/[DOM]. Table S10 shows our calculated values. The procedure to obtain this value was based on defining a chromophore as a series of consecutive sp² hybridized carbon units, including oxygen atoms

bonded to sp² carbons. Figure S6 shows an example of the delineation used. We did this only for 484 485 chromophores that would absorb at wavelengths above 260nm. As shown in Table S10, the ratio 486 of [CDOM] to [DOM] is roughly 53%, indicating that a little more than half carbon atoms are attached to a chromophore. This would translate into ~53% of the DOC in SFRA being 487 chromophoric. This number can be compared to the ¹³C nuclear magnetic resonance distribution 488 from the IHSS website,⁷⁷ for SRFA (catalogue number 1S101F) the fraction of carbonyl, carboxyl 489 490 and aromatic SRFA carbons is 51%, which is a number close to the 53% calculated here. It should 491 be noted that not all carboxylic groups should be part of a chromophore and that the 51% should 492 be considered as an upper limit maximum.

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501 Associated Content

502 Supporting information

503 The supporting information is available free of charge on the ACS website at DOI:

The supplemental information includes further figures, tables and text sections detailing the models' chemical structures, the effects of sodium borohydride on the structures, the molecules absorption by wavelengths, the effects of substituents on the absorption maximum, the estimation of *inter* molecular CT interactions, additional compounds investigated for *intra* molecular CT interactions and an example of the calculation of the chromophoric part of a structure.

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