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1	Tuning Color Variation In Grape Anthocyanins At The Molecular Scale
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34 ABSTRACT

35 Anthocyanins are the main grape pigments. Due to their aromatic cyclic arrangements, they 36 are able to absorb the radiation in the low energy range of the visible spectrum. In the fruit of 37 Vitis vinifera L., the five main anthocyanidins (cyanidin, peonidin, delphinidin, petunidin and 38 malvidin) are present as 3-O-glucosides, as well as their acetyl, p-coumaroyl and caffeoyl 39 ester forms. Despite the huge number of experimental studies dedicated to the anthocyanin 40 profile analysis of grapes and wines, the complete theoretical elucidation of the optical 41 properties of grape anthocyanins is missing. The present work carried out this task through 42 quantum chemistry calculations based on time-dependent density functional theory (TD-43 DFT), compared to experimental spectra. The differences in visible absorption spectra 44 between the most common grape anthocyanins were rationalized according to B-ring substitution, glucosylation and esterification. A particular attention was given to the intra-45 46 molecular copigmentation effect, demonstrating the existence of an intra-molecular charge 47 transfer excited state for the *p*-coumaroyl and caffeoyl ester forms.

- 48
- 49 **KEYWORDS**: UV/Vis absorption, intra-molecular copigmentation, optical properties,
- 50 pigment profile, TD-DFT, grape anthocyanins, wine
- 51

52 **INTRODUCTION**

53 Anthocyanins are a class of π -conjugated compounds belonging to polyphenols. They are 54 glycosides of the so-called anthocyanidins, characterized by a flavylium (2-phenyl-55 benzylpyrilium) skeleton (Fig. 1). The many possible chemical substitutions (with e.g., OH 56 and OMe groups) allow a large variety of compounds. They are pigments responsible for the 57 shiny orange, pink, red, violet and blue colors in roots, stems, flowers and fruit of many plants 58 (e.g., orchids, grapes) (Castañeda-Ovando, Pacheco-Hernández, Páez-Hernández, Rodríguez 59 & Galán-Vidal, 2009). Such a color variety is mainly driven by the capacity of these 60 compounds to absorb green light (at around 520 nm), which has generally been attributed to the resonant structure of the flavylium cation (Allen, 1998; Castañeda-Ovando, et al., 2009). 61 62 This was confirmed by quantum chemistry calculations, showing a highly delocalized π -63 conjugated system, spread over the entire molecule (Di Meo, Sancho Garcia, Dangles & 64 Trouillas, 2012). This extended π -conjugation allows light absorption in the visible range rather than the violet to ultraviolet regions as it is described and theoretically rationalized for 65 66 most of the other natural polyphenols (Anouar, Gierschner, Duroux & Trouillas, 2012).

67 A thorough understanding of optical properties and pigmentation variation has become crucial 68 in various domains including food and wine chemistry. We believe that in a near future 69 fundamental rationalization of grape pigments should become mandatory to supporting the 70 development of fast and non-invasive phenotyping strategies (Rustioni, Basilico, Fiori, Leoni, 71 Maghradze & Failla, 2013). In grapes and wines, five anthocyanidin moieties are present in 72 their glucosylated form, namely peonidin, cyanidin, malvidin, petunidin and delphinidin (Fig. 73 1a). The most common anthocyanin in *Vitis vinifera* L. (*i.e.*, the common wine grape specie) 74 is malvidin-3-O-glucoside (Kennedy, Saucier & Glories, 2006). There also exist anthocyanins 75 having their glucose moiety attached to a carboxylic acid (acetic, *p*-coumaric and caffeic acids 76 in Vitis vinifera L.) by an ester bond, giving rise to a group of acylated pigments (Fig. 1b)

77 (Allen, 1998). In the latter structures, intra-molecular copigmentation effects have been
78 suggested (Dangles, Saito & Brouillard, 1993).

79 Despite all the available experimental studies focusing on the grape and wine anthocyanin 80 profiles (Dallas & Laureano, 1994; Ferrandino, Guidoni & Mannini, 2007; Fischer, Löchner 81 & Wolz, 2006; García-Puente Rivas, Alcalde-Eon, Santos-Buelga, Rivas-Gonzalo & 82 Escribano-Bailón, 2006; Mattivi, Guzzon, Vrhovsek, Stefanini & Velasco, 2006; Roggero, 83 Coen & Ragonnet, 1986; Rustioni, Rossoni, Calatroni & Failla, 2011; Rustioni, Rossoni, 84 Cola, Mariani & Failla, 2011; Rustioni, Rossoni, Failla & Scienza, 2013), a profound 85 understanding of the optical variability of these compounds is still missing. Quantum 86 chemistry calculation is an adapted tool for investigation of these properties. In particular time 87 dependent density functional theory (TD-DFT) enables to accurately characterize excited 88 states and evaluate the subsequent absorption properties in the visible range. More than 89 providing accurate absorption wavelengths, TD-DFT calculations provide a complete 90 description of the excited states and thus allow full assignment of electronic transitions from the π to π^* molecular orbitals (MO) responsible for each UV/Vis absorption band. In this 91 92 way, the method supports the tuning of pigmentation in various industrial applications e.g., 93 pigment painting and solar cells. Moreover the intra-molecular interactions suggested from 94 experience (Dangles, et al., 1993) can now be accurately evaluated by quantum calculations; 95 these interactions are known to be responsible for copigmentation, but the process was only 96 recently fully understood using dispersion-corrected DFT (DFT-D) calculations (Di Meo, et 97 al., 2012).

The present work aims at rationalizing the differences in the absorption spectra between a series of grape anthocyanins, which has not been performed yet for these compounds. The series studied here is constituted of five glucosylated (compounds **1-5** in Figure 1a) and three acylated anthocyanins (compounds **6-8** in Figure 1b), which were investigated due to their

- 102 high occurrence in grape. This comprehensive study allows establishing a thorough structure-
- 103 property relationship for grape anthocyanins. In addition and as a new insight, the
- 104 copigmentation effects are carefully rationalized for the acylated anthocyanins.
- 105

106 MATERIAL AND METHODS

107 Experimental data

108 The grape anthocyanin spectra were collected from our database of HPLC chromatograms 109 obtained in 2008, within the framework of the study on competitive copigmentation 110 interactions (Rustioni, Bedgood Jr, Failla, Prenzler & Robards, 2012). Anthocyanins were 111 extracted from the skin of Shiraz and Sangiovese berries. The primary pigment solutions were 112 prepared after the careful removal of skin from 30 berries, and suspended in a 60 mL modelwine-buffer solution (12% ethanol and 2.5 g L^{-1} tartaric acid adjusted at pH 3.3 with aqueous 113 114 NaOH). The pigment extracts were obtained by filtration after this mixture was shacked for 6 115 hours. The extracts were characterized on a Varian Prostar 240I HPLC (Mulgrave, Vic, Australia) using a Phenomenex Gemini 5 µm C18 110A 250 x 4.6 mm column (Lane Cove, 116 NSW, Australia), with a Phenomenex SecurityGuard column, operated at 25°C. For the 117 118 HPLC separation the mobile phase was water: formic acid: acetonitrile (87:10:3 v/v/v, eluent A; 40:10:50 v/v/v, eluent B), and the following gradient sequence was used: from 10 to 25% 119 120 B (10 min), from 25 to 31% B (5 min), from 31 to 40% B (5 min), from 40 to 50% B (10 121 min), from 50 to 100% B (10 min) from 100 to 10% B (5 min), as previously described by 122 Kammerer et al. (Kammerer, Claus, Carle & Schieber, 2004). The flow rate was 0.8 mL min⁻ ¹. The eluent was monitored by a Varian Prostar 335 photodiode array detector (Mulgrave, 123 124 Vic, Australia). Peak identification was achieved using both the retention time and the LCMS 125 analysis as previously described (Rustioni, et al., 2012).

126 Malvidin aglycone, and caffeic and p-coumaric acids (SIGMA-ALDRICH) were used 127 without further purification. To compare their spectra with those obtained in HPLC, three solutions (10 mg L^{-1} for malvidin, 5.7 mg L^{-1} for caffeic acid and 3.2 mg L^{-1} for *p*-coumaric 128 129 acid) were prepared using the same solvent as for the HPLC elution (water:formic 130 acid:acetonitrile 65:10:25 v/v/v). A Jasco 7800 UV-Vis spectrophotometer was used to record 131 absorption spectra from 200 to 600 nm.

132 Considering the high percentage of formic acid providing pH ca. 1.6, the flavylium cation 133 form of the anthocyanins is unambiguously predominant. For this reason and for the sake of 134 comparison with experimental data, all the theoretical calculations were performed 135 considering this cationic form. ~

136

137 **Theoretical methodology**

138 Over the past decade, the density functional theory (DFT) has been validated and extensively 139 used to evaluate conformational, electronic and optical properties of natural polyphenols (Anouar, Gierschner, Duroux & Trouillas, 2012; Trouillas, Marsal, Siri, Lazzaroni & Duroux, 140 141 2006). TD-DFT turned out to be an accurate theoretical method to describe excited states and 142 consequently UV/Visible properties at an acceptable computational time for medium-size 143 polyphenols, namely containing up to 200 atoms (Anouar, et al., 2012; Nave, et al., 2012).

144 Hybrid functionals succeeded at describing most of polyphenol properties; the B3P86 145 functional has appeared particularly adapted to assess both their thermodynamic (Trouillas, et 146 al., 2006) and UV/Visible absorption properties (Anouar, et al., 2012). Using the Pople-type 147 triple- ζ basis set, TD-B3P86/6-311+G(d,p) allowed to establish reliable structure property 148 (UV/Visible absorption) relationships for large series of polyphenols, at a reasonable computational time (Anouar, et al., 2012; Millot, Di Meo, Tomasi, Boustie & Trouillas, 149 150 2012).

B3P86 was thus extrapolated for the present work to evaluate both conformation and UV/Visible absorption properties of peonidin, cyanidin, malvidin, petunidin and delphinidin and their glucoside derivatives (Fig. 1).

Concerning the acylated derivatives, cofacial (non-covalent) intra-molecular interactions 154 155 were examined between the anthocyanidin and the phenolic acyl (i.e., p-coumarovl and 156 caffeoyl) moieties. It is well described that classical hybrid functionals fail at describing these non-covalent interactions, which are mainly dispersive effects (e.g., π -stacking, v- π and long-157 158 distance H-bonds). The empirical dispersion-corrected DFT (DFT-D) is a successful approach 159 to evaluate these non-covalent complexation, also allowing to circumvent the use of high-160 costing post-HF methods (Grimme, 2006). Our recent parameterization of the B3P86-D2 161 functional provided very reliable results to describe non-covalent interaction in polyphenols 162 and copigmentation (anthocyanidin:flavonol) complexes with respect to both post-HF methods and experiments (Di Meo, et al., 2012). Therefore in the present work, the ground 163 164 state geometries of the acylated derivatives were corrected according to this empirical dispersion correction. The cc-pVDZ basis set was used, which has appeared sufficient to 165 reach accuracy, with respect to high-level quantum calculations and experimental data, in the 166 description of anthocyanidin: flavonol complexation.¹ All DFT-D calculations were achieved 167 168 within the resolution of identity (RI) approximation, dramatically decreasing the 169 computational time with a negligible error (Neese, Wennmohs, Hansen & Becker, 2009). 170 When working with DFT-D(B3P86-D2), the geometries were obtained with the ORCA 171 package (Neese, 2012). In the other case i.e., when working with B3P86 (no dispersion 172 correction), the geometries were calculated with GAUSSIAN09 (Frisch, et al., 2009). All TD-173 DFT calculations were achieved with GAUSSIAN09.

¹ It must be stressed that dispersion-corrected B3P86-D2 functional has been assessed and validated within double-z basis set precision.

174 Solvent effects were taken into account using implicit solvent models, in which the solute 175 is embedded in a shape-adapted cavity surrounded by a dielectric continuum. Even if they do 176 not explicitly account for inter-molecular interaction between solute and solvent, implicit 177 solvent models have allowed reaching accuracy to evaluate various properties of polyphenols, 178 including optical properties (Anouar, et al., 2012; Millot, et al., 2012). The qualitative 179 description of the influence of solvent polarity is particularly relevant, which perfectly 180 reproduce structure-property relationships. Methanol ($\varepsilon = 32.61$) was used to simulate the 181 polarity of the experimental conditions used to measure the UV/Visible spectra. The 182 conductor-like screening model (COSMO) (Sinnecker, Rajendran, Klamt, Diedenhofen & 183 Neese, 2006) and the integral-equation formalism polarizable continuum model (IEFPCM) 184 (Cossi, Scalmani, Rega & Barone, 2002) were used with ORCA and GAUSSIAN09, 185 respectively.

186

187 **RESULTS AND DISCUSSION**

188 UV/Visible absorption properties of grape 3-O-glucosylated anthocyanins

The experimental spectra measured for the five 3-*O*-glucosylated anthocyanidins (compounds **1-5** namely cyanidin 3-*O*-glucoside, peonidin 3-*O*-glucoside, delphinidin 3-*O*glucoside, petunidin 3-*O*-glucoside and malvidin 3-*O*-glucoside – also called oenin – as seen in Fig. 1), synthesized in grape berries, consist of three main peaks experimentally located at around i) 520 nm (Band I), ii) 440 nm (Band II) and iii) 330 nm (Band III) (Fig. 2a). The disubstituted B-ring compounds (*i.e.*, cyanidin 3-*O*-glucoside and peonidin 3-*O*-glucosides) have an additional absorption band located at around 375 nm (Band II_b).

196

197 *Molecular orbital rationalization of Band I* - The highest absorption wavelength λ_{max} 198 corresponding to Band I of compounds **1-5** are 516, 517, 524, 526 and 527 nm, respectively

199 (Table 1). As already described for anthocyanidins, this wavelength is slightly underestimated 200 by TD-DFT calculations when using standard hybrid functionals (Anouar, et al., 2012; Chai 201 & Head-Gordon, 2008). This band corresponds to a transition from the ground state (S_0) to the 202 first excited state (S_1) and is mainly assigned to the HOMO (highest occupied molecular orbital) to LUMO (lowest unoccupied molecular orbital) electronic transition.² In order to 203 204 simplify notations, the HOMO \rightarrow LUMO electronic transition is quoted H \rightarrow L throughout the 205 text. Both molecular orbitals (MO) are delocalized over the whole anthocyanidin moiety (Fig. 206 3a&b). This allows an efficient overlap between the two molecular orbitals involved in the 207 transition, which provides relatively high oscillator strengths (and therefore high absorption 208 intensities) for all five compounds (f around 0.60 as seen in Table 1). The compounds having a tri-substituted B-ring (compounds 3-5), exhibit a (red)-209 210 bathochromic shift compared to the di-substituted (1 and 2) derivatives (Fig. 2a). This is 211 attributed to the (+M) mesomeric effect that slightly but significantly extends π -conjugation 212 in the B-ring, which subsequently lowered the H-L gap (ΔE_{gap} , Fig. 3a&b). When adding one OH group (i.e., compounds 1 vs. 3 and 2 vs. 4), both H and L are stabilized but to a lower 213 214 extent for H, thus decreasing the gap (e.g., ΔE_{gap} decreases from 2.91 to 2.86 eV, from 1 to 3, 215 see Fig. 3a&b). Such an effect was similarly observed for aglycone flavonoids. (Anouar, et al., 2012) When replacing an OH group by an OMe group, both frontier MOs are destabilized, but 216 217 the H orbital is more impacted by this destabilization, resulting in the decrease of the H-L gap (e.g., ΔE_{gap} is 2.86 and 2.75 eV for 3 and 5, respectively, see S1). 218 219 The theoretical excited states of the aglycone counterpart of compounds 1-5 were also

evaluated for the sake of comparison. From this comparison, it appears that the sugar moiety

 $^{^2}$ In the present work, the orbital analysis is based on the 3D distribution of only the main two MO involved in the main electronic transition. The full analysis taking the contributions of all electronic transitions into account was also achieved (see Supplement Information). However, this latter (thorough) analysis does not influence the conclusions (except concerning one point, which is mentioned in the next footnote); thus for the sake of simplicity, the simpler analysis is kept in the main body of the text.

creates new MOs lying below H-4 (see Supplementary Information), which do not directly influence the absorption spectra *e.g.*, by the formation of new bands. However, the sugar addition induces a slight increase of the H-L gap, due to both the stabilization of H and destabilization of L. This is in agreement with experimental data, as it results in a significant hypsochromic shift of Band I in the glucosides' spectra (*e.g.*, λ_{max} equal 527 and 536 nm for compound **5** and its aglycone counterpart, respectively as seen in Fig. 2b). This does not influence the global spectra, which are similar in shape with both forms.

228 Molecular orbital rationalization of Band II - Band II appears in the experimental spectra 229 as a shoulder of the maximum absorption band, which is located at around 440 nm (Fig. 2a). 230 It corresponds to the second excited state (S₂) and is mainly assigned to the H-1 \rightarrow L and H-231 $2 \rightarrow L$ for di-substituted (*i.e.*, compounds 1 and 2) and tri-substituted (*i.e.*, compounds 3-5) 232 compounds, respectively (Table 1). L being delocalized on the A and C rings, this shoulder 233 appears attributed to this moiety. A thorough analysis of the MO spatial distribution 234 highlights interconversion between both H-1 and H-2 for the di- and tri-substituted 235 compounds; that is, the spatial distribution of H-1 in the di-substituted compound is similar to 236 that of H-2 in the tri-substituted compound (Fig. 3a&b). These MOs are delocalized over the 237 entire anthocyanidin moiety. Moreover, they exhibit the same energy (ca. -7.75 eV for all 238 compounds, as seen in Supplementary Information). For the tri- substituted compounds, H-1 239 is delocalized over the B-ring only. This MO slightly contributes to other excited states, 240 including S_1 (data not shown).

For the di-substituted compounds (1 and 2), there exists a specific absorption band, quoted Band II_b in Table 1, which is absent in the experimental spectra of the tri-substituted compounds (3, 4 and 5). Quantum chemistry calculations fully rationalize this experimentally observed difference. Band II_b is assigned to the H-2 \rightarrow L electronic transition. In the disubstituted compounds, H-2 is slightly delocalized on the A&C-rings (Fig. 3a), allowing

electronic transitions with low oscillator strengths (f = 0.15 and 0.18 for 1 and 2, respectively). In the tri-substituted compounds, due to the MO redistribution described above, H-1 is almost fully delocalized on the B-ring, preventing this electronic transition involving the A&C-rings. Moreover, in this case, the energetic difference between H-1 and L is lowered. Thus for the tri-substituted compounds, Band II_b is weak and red-shifted; it cannot be experimentally observed under classical conditions (i.e., room temperature and classical resolutions).

253

254 Molecular orbital rationalization of Band III - Band III is mainly assigned to the H-4→L 255 electronic transition. In agreement with the experimental data, Band III exhibits relatively low 256 oscillator strengths compare to those of Band I, for all five compounds (f is ranging from 0.02 257 to 0.08, see Table 1). The (blue)hypsochromic shift experimentally observed from the tri- to the di-substituted compounds (e.g., $\lambda = 348$ and 328 nm for 5 and 1, respectively, see Table 258 259 1) is rationalized by the influence of the sugar moiety, which surprisingly, but in full agreement with the experimental data, borrows part of the H-4 orbital, in the di-substituted 260 261 compounds (Fig. 3a&b). In this case the sugar moiety stabilizes H-4, thus increasing the 262 energy difference between H-4 and L, inducing the hypsochromic shift. This is probably only 263 an (energetic) indirect effect, as the electronic excitation does not involve the sugar moiety (Fig. S2). 3 264

265

266 UV/Visible absorption properties of grape acylated anthocyanins

The spectral modulation related to esterification is discussed for the malvidin derivatives, as they are the main grape anthocyanidins.

³ This is clearly observed when analyzing all MO involved in this electronic transition.

270 Conformational analysis - Concerning the three malvidin 3-O-glucoside esters

271 (compounds 6, 7 and 8, see Fig. 1) one, two (7A and 7B) and two (8A and 8B) conformers

272 were obtained, respectively (Fig. 4).

273 For each possible conformer, the Boltzmann distribution percentages D_{Bolt} were estimated

274 (Table 2) from the relative energies $\Delta E_{relative}$ (as calculated between all conformers with

275 B3P86-D2/cc-pVDZ), according to the following formula:

276
$$D_{Bolt} = \exp\left(-\frac{\Delta E_{relative}}{RT}\right)$$

where R is the ideal gas constant and T equals 298 K.

278 The non-covalent intra-molecular interactions between the acyl and anthocyanidin moieties 279 play a crucial role in the stabilization of these conformers. A strong H-bond (distance lower 280 than 1.65 Å) between the 5-OH and the ester keto groups is observed in 6, 7B, 8A and 8B 281 (Fig. 4). A weaker H-bond is observed between the 5-OH and the O atom involved in the 282 esterification (distance of 2.39 Å) for 7A. Other non-covalent interactions are observed in 7A, 7B and 8A (Fig. 4), namely i) one H-bonding between the 7-OH and the 4"-OH groups 283 (distance of 1.80 and 2.27 Å, respectively) for 7A and 7B, which is not present in compound 8 284 285 as 4"-OH is absent in this case, and ii) π -stacking interaction in **7B** and **8A** (distance of 4.20) Å). 286

- 287 The conformation populations of **7** and **8**, show a predominance of **7A** ($D_{Bolt} = 90\%$) and 288 **8A** ($D_{Bolt} = 83\%$) for both compounds, respectively. This clearly suggests that both H-bonding 289 and π -stacking interactions are the driving forces of this intra-molecular folding.
- 290

UV/Visible absorption spectra of 6, 7 and 8: a matter of intra-molecular copigmentation When compared to the glycosilated malvidin (5), the experimental spectra of 6, 7 and 8
clearly show the influence of esterification, mainly in the presence of the phenolic acid moiety

294 (compounds 7 and 8). The three acylated anthocyanins exhibit an experimental bathochromic 295 shift of Band I that is lower for 6 ($\Delta\lambda max$ around 2, 5 and 4 nm for 6, 7 and 8, respectively). 296 The bathochromic shift of Band I is mainly rationalized by π -stacking complexation. In 297 conformations 7B and 8A, the maximum absorption wavelength (related to Band I) and other 298 very close absorption wavelengths are assigned to the $H \rightarrow L$ and $H - 1 \rightarrow L$ electronic transitions 299 (Table 2a). In all conformers, H and L are mainly delocalized on the anthocyanin moiety (Fig. 300 3c-f), and the electron transition presents no charge-transfer character. In the particular case of 301 conformers **7B** and **8A**, H is slightly located on the phenolic ring of the ester moiety (Fig. 302 3d&e). This can be better evaluated by the LCAO (linear combination of atomic orbital) 303 coefficients, which reflect the atomic weight of a given MO. For H of 7B and 8A, these 304 coefficients are 13% and 21% on the acyl moiety, respectively (Table S2). However, a 305 detailed analysis of MO contributions (see Supplementary Information) shows that the 306 esterification affects neither the spatial distribution nor the energy of H and L, showing that 307 the bathochromic shift is not rationalized by the $H \rightarrow L$ contribution. The main contribution to 308 the modification of the excited state (with respect to that of compound 5) comes from H-1, 309 which is highly delocalized over the phenolic ester moiety for all conformers of 7 and 8 (Fig. 310 3c-f). Therefore the H-1 \rightarrow L electronic transition exists only because of the formation of a 311 charge transfer excited state (CT-ES), inducing a global bathochromic shift attributed to 312 esterification. In conformers 7B and 8A, H-1 is also located on the anthocyanin moiety (the LCAO coefficients of H are 57% and 49% on the anthocyanin moiety, for 7B and 8A, 313 314 respectively see Table S2), which increases the corresponding oscillator strength (Table 2).

315 Compound 7 exhibits a slightly lower experimental bathochromic shift than 8 ($\Delta\lambda max = 4$ 316 and 5 nm, for both compounds respectively, see Table 2a). In contrast, the theoretical 317 bathochromic shift is higher with the former compound (7) i.e., 24.7 *vs.* 18.9 nm, for both 318 compounds respectively (Table 2a). This is consistent with the higher electron donor capacity

of the ortho-dihydroxyphenyl (catechol) moiety. Thus the lower experimental bathochromic shift observed for **7** is attributed to a lower population of the π -stacked conformation (**7B**) in solution compared to that of **8** (**8A**). This highlights the importance of i) charge-transfer in the excited state and ii) populations for all the possible copigmentation complexes, as already shown in our previous work (Di Meo, et al., 2012).

324

325 A new band (III_b) at 330 nm - The acylated compounds exhibit a new absorption band, 326 quoted Band III_b, experimentally located at around 330 nm (Fig. 2c), which is somehow a 327 modulation of Band III. This band is actually the sum of the minor anthocyanic absorption 328 Band III (Table 1 and Fig. 2a) and the (red)bathochromic-shifted band mainly attributed to the stand-alone phenolic acid (e.g., $\lambda = 324$ and 336 nm for caffeic acid and the caffeoyl ester of 329 330 malvidin 3-O-glucoside, respectively and $\lambda = 309.6$ and 316 nm for p-coumaric acid and the 331 p-coumaroyl ester of malvidin 3-O-glucoside, respectively, see Table 2b and Fig. 2c). It is 332 theoretically assigned to several contributions predominantly H-1 \rightarrow L+1, H-4/5/6 \rightarrow L and 333 $H \rightarrow L+2$ electronic transitions (Table 2b). The first contribution is purely attributed to the 334 phenol acid moiety, both MO being delocalized on this moiety. These MO are stabilized in 335 the presence of the anthocyanin moiety, H being more stabilized than L, which should induce 336 a hypsochromic shift. The global bathochromic shift is due to the other two contributions. 337 Among these contributions, the H-4 \rightarrow L contribution corresponds to that already observed for 338 compound 5, as the main contribution of Band III.

339

340 CONCLUSION

Grapes and wines represent agricultural product of utmost importance in food industry. The color of the final product is a major descriptor of quality. A deep understanding of the optical properties of the main *Vitis vinifera* L. pigments represents an effective support for the

344 industries to address market requirements. This is mandatory to develop new analytical 345 methods based on the optical properties and to deeply understand fruit potential. As described 346 above, quantum chemistry calculations have allowed elucidating the structure (optical) 347 properties of a series of grape pigments. These calculations provide a molecular picture of the 348 pigments as well as a molecular orbital description of the excited states related to the different 349 absorption bands responsible for color. Subtle effects such as copigmentation can also be 350 explained by using the adapted theoretical methodologies, namely DFT-D. All the theoretical 351 data support the experimental observations. The full MO description of light absorption, in the 352 visible range, enables to tune optical properties (according to the chemical structures of the 353 different grape pigments), and subsequently color variation. Quantum calculation can now be 354 considered as a promising analytical tool for wine industry.

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370 **TABLES**

Table 1. Theoretical excitation energies (E, eV), absorption wavelengths (λ , nm), oscillator 371 strengths (f), main electronic transition contribution and experimental absorption wavelength 372 (λ_{exp}, nm) of anthocyanidin 3-O-glucosides. The calculations (excited states + geometries) 373 374 were performed with IEFPCM-B3P86/6-311+G(d,p).

	Band	Compound	Е	λ	f	Electronic transition	λετη
	I	1	2.59	478.0	0.58	$H \rightarrow L (67\%)$	516
	_	2	2.57	482.8	0.59	H→L (68%)	517
		3	2.57	482.8	0.57	H→L (67%)	524
		4	2.56	483.4	0.61	H→L (68%)	526
		5	2.49	497.1	0.59	H→L (68%)	527
	II	1	2.95	419.8	0.07	H-1→L (67%)	~440
		2	2.92	424.8	0.03	H-1→L (68%)	~440
		3	2.95	420.1	0.11	H-2→L (67%)	~440
		4	2.93	423.7	0.06	H-2→L (67%)	~440
		5	2.92	424.0	0.05	H-2→L (68%)	~440
	II _b	1	3.23	384.0	0.15	H-2→L (70%)	373
		2	3.22	385.2	0.18	H-2→L (69%)	375
		3	2.68	463.2	0.03	H-1→L (70%)	-
		4	2.70	456.0	0.01	H-1→L (70%)	-
		5	2.68	461.9	0.07	H-1→L (70%)	-
	III	1	3.88	319.5	0.02	H-4→L (62%)	328
		2	3.88	319.8	0.02	H-4→L (62%)	328
		3	3.73	332.2	0.06	H-4→L (66%)	346
		4	3.72	332.9	0.08	H-4→L (66%)	347
		5	3.69	336.2	0.08	H-4→L (66%)	348
376							

377	Table 2. Theoretical excitation energies (E, eV), absorption wavelengths (λ , nm), oscillator
378	strengths (f), electronic transition contribution and experimental absorption wavelength (λ_{exp} ,
379	nm) of the different optimized conformation of acetyl (6), caffeyl (7) and p-coumaryl (8)
380	esters of malvidin 3-O-glucoside, and related theoretical and experimental bathochromic
381	shifts ($\Delta\lambda th$ and $\Delta\lambda exp$, nm) concerning (a) Band I and (b) Band III. Boltzmann
382	distributions D _{Boltz} are expressed in %. The calculations were performed with IEFPCM-
383	B3P86/6-311+G(d,p)//COSMO-B3P86(D2)/cc-pVDZ.
384	(a)

384

Conformation	D _{Boltz}	Е	λ	f	Electronic transition contribution	$\Delta\lambda th^a$	λ_{exp}	Exp Δλ <i>exp</i>
6	90%	2.51	494.9	0.58	H→L (68%)	-2.2	529	2
7A	90%	2.47	502.2	0.17	H-1→L (57%)	5.0	531	4
					H→L (40%)			
		2.50	496.5	0.38	H→L (54%)	-0.6		
7B	6%	2.38	521.8	0.10	H-1→L (52%)	24.7		
		2.49	498.1	0.37	H-1→L (46%)	1.0		
					H→L (49%)			
8A	83%	2.40	516.0	0.41	H→L (66%)	18.9	532	5
		2.59	479.5	0.03	H-1→L (58%)			
8B	15%	2.50	495.2	0.57	H→L (67%)	-1.9		
	Conformation 6 7A 7B 8A 8B	Conformation DBoltz 6 90% 7A 90% 7B 6% 8A 83% 8B 15%	Conformation D _{Boltz} E 6 90% 2.51 7A 90% 2.47 7B 6% 2.38 2.49 2.49 8A 83% 2.40 2.59 15% 2.50	Conformation D _{Boltz} E λ 6 90% 2.51 494.9 7A 90% 2.47 502.2 7A 2.50 496.5 7B 6% 2.38 521.8 2.49 498.1 8A 83% 2.40 516.0 2.59 479.5 8B 15% 2.50 495.2	Conformation D _{Boltz} E λ f 6 90% 2.51 494.9 0.58 7A 90% 2.47 502.2 0.17 2.50 496.5 0.38 0.38 7B 6% 2.38 521.8 0.10 2.49 498.1 0.37 0.37 8A 83% 2.40 516.0 0.41 2.59 479.5 0.03 0.37	$\begin{array}{c c c c c c c } \mbox{Conformation} & D_{\rm Boltz} & E & \lambda & f & Electronic transition contribution \\ \hline \begin{tabular}{lllllllllllllllllllllllllllllllllll$	$\begin{array}{c c c c c c c } \mbox{Conformation} & D_{\rm Boltz} & E & \lambda & f & Electronic transition contribution & \end{displaysismed lines } \end{displaysismed lines } \\ \hline {\bf 6} & 90\% & 2.51 & 494.9 & 0.58 & H \rightarrow L (68\%) & -2.2 \\ \hline {\bf 7A} & 90\% & 2.47 & 502.2 & 0.17 & H - 1 \rightarrow L (57\%) & 5.0 \\ & & & & & & & & & & & & & & & & & & $	$ \begin{array}{c c c c c c c c c c c c c c c c c c c $

³⁸⁵ 386

 a the shift is performed with respect to malvidin 3-O-glucoside λ_{max} .

6 <u>(b)</u>										
Ester	Conformation	D _{boltz}	E	λ	f	Electronic transition contribution	Alth	Exp		
Lister	Comornation				J		$\Delta \pi l h$	λ_{exp}	Δλεχρ	
Acetic ester	6	100	3.64	340.8	0.07	H-3→L (65%)	nd	348	nd	
Caffeoyl	7A	90%	3.66	339.1	0.03	H-5→L (48%)	1.1	324	12	
ester						H-6→L (36%)				
						H-1→L+1 (31%)				
						H→L+2 (15%)				
			3.72	333.5	0.64	H-1→L+1 (62%)	-4.5			
	7B	6%	3.51	353.7	0.32	H-1→L+1 (68%)	15.7			
						H-4→L+1 (10%)				
			3.64	340.5	0.03	H-5→L (65%)	2.5			
<i>p</i> -Coumaroyl	8A	83%	3.68	336.6	0.25	H-1→L+1 (55%)	19.6	316	6	
ester						H-5→L (33%)				
			3.74	331.5	0.22	H-5→L (47%)	14.5			
						H-1→L+1 (34%)				
						H→L+2 (15%)				
	8B	15%	3.70	334.8	0.68	H-1→L+1 (64%)	17.7			
						H-4→L+(14%)				
			3.75	331.1	0.13	H-6→L (62%)	14.0			
						H-4→L (20%)				
						H-1→L+1 (19%)				
						H-6→L (15%)				

^{*a*} the shift is performed with respect to the stand-alone acid λ_{max} (i.e., caffeic or p-coumaric acid in solution). 387 Acceleration

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- 481

483 FIGURE CAPTIONS

484

485 Figure 1. Chemical structures of the five glucosylated (a) and three acylated (b) studied

- 486 anthocyanins.
- 487
- 488 **Figure 2.** Experimental spectra of (a) anthocyanidin 3-*O*-glucosides (compounds 1-5); (b)
- 489 malvidin aglycone and malvidin 3-O-glucoside (5); and (c) malvidin 3-O-glucoside (5), oenin

490 esters (compounds 7 and 8), and relative phenolic acids.

491 All the spectra are normalized to appear in at a similar scale: anthocyanic molecules are

492 normalized at a relative absorption value of 100 in the main peak (ca. 530 nm); phenolic acids

493 are normalized according to the corresponding ester (ca. 330 nm).

494

495 Figure 3. Molecular orbital (MO) diagrams of the di-substituted (a) compound 1 and (b) tri-

496 substituted compound **3** and the different conformations of compounds **7** and **8**: (c) **7A** (d) **7B**,

497 (e) **8A** and (f) **8B**. The anthocyanidin moiety is in red, the sugar moiety is in blue and the acyl 498 moiety is in green. The calculations were performed with IEFPCM-B3P86/6-311+G(d,p) and

499 IEFPCM-B3P86/6-311+G(d,p)//COSMO-B3P86(D2)/cc-pVDZ with **1&3** and **7&8**, 500 respectively.

501

Figure 4. Optimized geometries of the acetyl ester (**6**), caffeic ester (**7A**; **7B**) and p-coumaric ester (**8A**; **8B**) of malvidin 3-*O*-glucoside; the anthocyanidin, sugar and the acyl moieties are in red, blue and green, respectively. The most relevant non-covalent intra-molecular interactions between the acyl and anthocyanidin moieties are underlined by broken lines. The calculations were performed with IEFPCM-B3P86/6-311+G(d,p)//COSMO-B3P86(D2)/ccpVDZ.









512 HIGHLIGHTS:

- 513 Grape anthocyanins colors were rationalized using quantum chemistry calculations.
- Accepter 514 Absorption differences related to pigment substitutions were explained.

