

# 1 Rosé Wine Fining Using Polyvinylpyrrolidone: Colorimetry, 2 Targeted Polyphenomics, and Molecular Dynamics Simulations

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## 10 **S** Supporting Information

11 **ABSTRACT:** Polyvinylpyrrolidone (PVPP) is a fining agent polymer used in winemaking to adjust rosé wine color and to  
 12 prevent organoleptic degradations by reducing polyphenol content. The impact of this polymer on color parameters and  
 13 polyphenols of rosé wines was investigated, and the binding specificity of polyphenols toward PVPP was determined. Color  
 14 measured by colorimetry decreased after treatment, thus confirming the adsorption of anthocyanins and other pigments.  
 15 Phenolic composition was determined before and after fining by targeted polyphenomics (UPLC-electrospray ionization source  
 16 (ESI)-mass spectrometry (MS)/MS). MS analysis showed adsorption differences among polyphenol families. Flavonols (42%)  
 17 and flavanols (64%) were the most affected. Anthocyanins were not strongly adsorbed on average (12%), but a specific  
 18 adsorption of coumaroylated anthocyanins was observed (37%). Intermolecular interactions were also studied using molecular  
 19 dynamics simulations. Relative adsorptions of flavanols were correlated with the calculated interaction energies. The specific  
 20 affinity of coumaroylated anthocyanins toward PVPP was also well explained by the molecular modeling.

21 **KEYWORDS:** rosé wine, PVPP, phenolic compounds, CIELAB, interaction energy, molecular dynamics simulations

## 22 ■ INTRODUCTION

23 Polyphenols are essential molecules found in rosé wines. They  
 24 can be divided into seven families: benzoic acids, hydroxycin-  
 25 namic acids, stilbenes, flavonols, flavan-3-ols, dihydroflavonols,  
 26 and anthocyanins. Their quantities vary depending on different  
 27 factors such as grape variety, geographic origin, and wine-  
 28 making process. In wine, they are responsible for quality and  
 29 sensorial characteristics such as taste and color. More  
 30 specifically, anthocyanins, which are the red grape pigments,  
 31 play an important role in wine color.<sup>1</sup> Polyphenols in rosé  
 32 wines in the presence of SO<sub>2</sub> can also enhance the antioxidant  
 33 effect of these wines by a synergistic effect.<sup>2</sup> However, an excess  
 34 of polyphenols may induce defaults like browning problems  
 35 due to oxidation of polyphenols and especially flavanols.<sup>3–5</sup>  
 36 During storage, the stability of the pink color of rosé wines may  
 37 be an issue as more orange pigments may form because of  
 38 reactions of phenolic acids, flavanols, and anthocyanins, like  
 39 xanthylum derivatives<sup>6,7</sup> or pyranoanthocyanins, one of the  
 40 most important classes of anthocyanins derivatives.<sup>8–10</sup> Some  
 41 thiol aroma compounds may also be trapped by quinones  
 42 during the oxidative process involved in rosé wine aging.<sup>11</sup> One  
 43 way to limit these problems is to reduce the polyphenol  
 44 quantities in the wine using fining agents such as  
 45 polyvinylpyrrolidone (PVPP), a cross-linked synthetic  
 46 polymer of polyvinylpyrrolidone (PVP) (Figure 1) known to  
 47 have polyphenol binding affinities. During fining, PVPP adsorbs  
 48 some polyphenols thus reducing their amounts in alcoholic  
 49 beverages like beer and wine.<sup>12–14</sup> This adsorption of

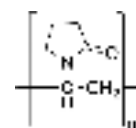


Figure 1. Chemical structure of PVP, (C<sub>6</sub>H<sub>9</sub>NO)<sub>n</sub>.

polyphenols by PVPP involves H-bonding between the proton 50  
 donor from the polyphenol and the carbonyl group from PVPP, 51  
 together with polar  $\pi$ -bond overlap (delocalized electrons) and 52  
 hydrophobic reactions.<sup>15,16</sup> In rosé wine production, PVPP is 53  
 regularly used to reduce color and phenolics. This fining 54  
 treatment can be done at the grape must, fermentation, or 55  
 finished wine stages. In this paper, we chose to focus on the 56  
 PVPP treatment at the finished wine stage. The aim of this 57  
 work was to investigate the effect of PVPP on rosé wine color 58  
 adjustment and the selective polyphenol adsorption phenom- 59  
 enon induced by the treatment in a laboratory-standardized 60  
 protocol. To achieve this, we used a targeted polyphenomics 61  
 methodology, which is a metabolomics approach focused on 62  
 polyphenols, recently developed to measure up to 152 phenolic 63  
 compounds in rosé wines by using liquid chromatography 64  
 (LC)-mass spectrometry (MS)/MS in multiple reaction 65

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66 monitoring (MRM) mode.<sup>17</sup> In addition, interaction energy  
67 calculations (at semiempirical quantum mechanical level) and  
68 molecular dynamics simulations including dynamic docking  
69 were carried out to provide deeper insights into the behavior of  
70 the PVPP polymer in a simulated rosé wine solution (ethanol/  
71 water). These studies allowed a better understanding of the  
72 interactions that govern the PVPP-polyphenols affinity during  
73 PVPP treatments on wines.

## 74 ■ MATERIALS AND METHODS

75 **Chemicals.** All chemicals were of analytical reagent grade.  
76 Methanol and formic acid were purchased from Sigma-Aldrich  
77 (Saint-Quentin Fallavier, France). Deionized water was obtained  
78 from a Milli-Q Advantage A10 purification system from Millipore  
79 (Fontenay sous Bois, France). PVPP VINICLAR was obtained from  
80 Laffort (Bordeaux, France).

81 **Wines and Sample Preparation.** Commercial rosé wines from  
82 the same vintage (2015) and from different regions of France were  
83 selected ( $n = 6$ ): Gascogne (1), Languedoc (3), Provence (1),  
84 Roussillon (1). A standardized laboratory protocol was developed to  
85 allow comparison of a PVPP treatment on rosé wines: a 100 g/L stock  
86 solution of PVPP VINICLAR was prepared. After an hour of rest time,  
87 15 mL of wine was supplemented with 120  $\mu$ L of the PVPP stock  
88 solution (final concentration of PVPP = 80 g/hL = maximum legal  
89 use). Samples were mixed for 30 s with a vortex and then were left to  
90 stand for 1 h at constant room temperature (20 °C) and then were  
91 centrifuged at 10 016g (8500 rpm) for 5 min. Initial tests were  
92 performed to determine the minimal contact time needed to reach the  
93 equilibrium (Appendix 1 of the SI). The supernatant was submitted to  
94 spectrophotometric analysis, then was aliquoted in 1.5 mL Eppendorfs,  
95 and was stored at -80 °C until further analyses. For mass  
96 spectrometry analyses, samples were brought back to ambient  
97 temperature, were filtered through a 0.2  $\mu$ m regenerated cellulose  
98 membrane syringe filter (Phenex, Phenomenex, Le Pecq, France), and  
99 then were injected with no further sample preparation. Controls went  
100 through the same steps but without PVPP addition.

101 **Spectrophotometric L\*a\*b\* Measurements.** Color analyses  
102 were performed on a spectrophotometer CM-3600d from KONICA  
103 MINOLTA with a 1.0 cm length glass cell, between 360 and 740 nm  
104 with 10 nm pitch, and were piloted with the SpectraMagic NX  
105 software. The CIELAB coordinates L\*, a\*, b\*, h, and C\* were  
106 obtained using the D65 illuminant and a 10° observer. The CIELAB is  
107 a color space defined in 1976.<sup>18</sup> In this three-dimension system, the L\*  
108 axis indicates that the lightness has a value that extends from 0 (black)  
109 to 100 (white); the a\* and b\* axes represent the chromaticity.  
110 Coordinate a\* has positive values for red colors and negative values for  
111 green colors. Coordinate b\* has positive values for yellow colors and  
112 negative values for blue colors. L\*, a\*, and b\* form a rectangular  
113 coordinate, but any point in this color space can also be defined by the  
114 cylindrical coordinates L\*, C\*, and h. C\* and h, respectively, represent  
115 chroma and hue angle and are, respectively, calculated as  $\sqrt{[(a^*)^2 +$   
116  $(b^*)^2]}$  and  $[\arctan(b^*/a^*)]$ .<sup>19,20</sup> The difference of colors  $\Delta E$  between  
117 two samples may be calculated as  $\sqrt{[(L_1^* - L_2^*)^2 + (a_1^* - a_2^*)^2 +$   
118  $(b_1^* - b_2^*)^2]}$ ; if this value is superior to 1, a color difference can be  
119 perceived by the human eye, and the bigger the  $\Delta E$  value, the easier it  
120 is to notice the color difference.<sup>21</sup>

121 **UPLC-QqQ-MS Parameters.** Polyphenol analyses were performed  
122 with a Waters Acquity UPLC system connected to a triple quadrupole  
123 mass spectrometer equipped with an electrospray ionization source  
124 (ESI) operating in switching positive and negative mode. The UPLC  
125 system included a binary pump, a cooled autosampler maintained at 7  
126 °C and equipped with a 5  $\mu$ L sample loop, a 100  $\mu$ L syringe and a 30  
127  $\mu$ L needle, a thermostated column department, and a DAD detector.  
128 MassLynx software was used for instrument control and data  
129 acquisition, and then TargetLynx software was used for data  
130 processing. Quantitative analyses were performed by UHPLC-QqQ-  
131 MS using the multiple reaction monitoring (MRM) detection mode  
132 under the conditions (high-performance liquid chromatography

(HPLC) elution conditions, MS and MRM parameters, calibration 133  
standards) described in Lambert et al.<sup>17</sup> 134

**Computational Methods. Building Molecular Structures.** The 135  
molecular structures of the 49 more abundant molecules (Appendix 2 136  
of the SI), monomer, and tetramer of PVP were built using the 137  
GaussView program.<sup>22</sup> The structures were built considering an 138  
environment at wine pH (3.5). In the case of anthocyanins, their 139  
flavylium ion and hydrated form have been considered. The 140  
geometries of these molecules were optimized at density functional 141  
theory (DFT) level<sup>23</sup> using the B3LYP method<sup>24,25</sup> with 6-31G(d,p) 142  
as basis set, which has been implemented in Gaussian 03 package 143  
program.<sup>26</sup> 144

**In Silico Calculation of Interaction Energies.** A semiempirical 145  
quantum mechanical strategy complemented with Monte Carlo 146  
conformational sampling<sup>27–29</sup> was used to calculate the interaction 147  
energy of molecule1–molecule2 complexes. In this case, the 148  
molecule1 represents the PVP tetramer, and the molecule2 represents 149  
each one of the 49 targets. 150

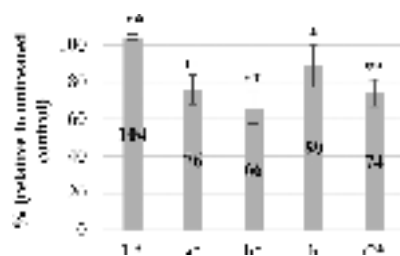
**Molecular Dynamics Simulation (MDS).** The polyvinylpyrrolidone 151  
(PVP) monomer was used to generate 50 PVP chains of 20, 30, and 40 152  
monomers long using LEAP module of AmberTools software.<sup>30</sup> 153  
Subsequently, using PACKMOL software,<sup>31</sup> these 50 chains were 154  
randomly distributed within a virtual sphere of 45 Å radius centered in 155  
the origin 0, 0, 0 (axes X, Y, Z, respectively). The chains were 156  
separated one from each other by a distance of at least 3 Å. These 157  
steps generated a PVP spherical microparticle of 90 Å diameter with 158  
the aim of transforming this PVP microparticle to a PVPP 159  
microparticle (the highly cross-linked version of PVP). The LEAP 160  
module was used to perform the cross-linking procedure.<sup>32</sup> It is based 161  
on a cyclic iteration scheme, and each cycle consists of three steps: (1) 162  
random breaking of a pyrrolidone ring, (2) covalent bonding of the 163  
obtained COOH group to the nearest amino group (only if it is 5 Å 164  
away), and (3) minimization of the system performed with the 165  
steepest descent algorithm and the universal force field (UFF) using 166  
openbabel software.<sup>33</sup> The steps from 1 to 3 are repeated until 60% of 167  
the pyrrolidone rings are broken. The graphical scheme of the 168  
formation of the PVPP microparticle is shown in Appendix 3 of the SI. 169

The obtained PVPP microparticle was added in the center of a 170  
solvent box of the following sizes: 150, 150, and 150 Å (axes X, Y, and 171  
Z, respectively). Subsequently, the box was solvated considering a 172  
90:10 mixture of water and ethanol with the aim of simulating the 173  
main components of rosé wine. The amounts of ethanol and water 174  
molecules (TIP3) were obtained on the basis of their corresponding 175  
molecular experimental density (0.789 g cm<sup>-3</sup> for ethanol and 1 g 176  
cm<sup>-3</sup> for water). Subsequently, the 30 polyphenols that had the best 177  
interaction energies (less than or equal to -2.7 kcal mol<sup>-1</sup>, calculated 178  
with semiempirical methods) were added to the inside of the box (8 Å 179  
away from the surface of the PVPP microparticle). Finally, two MDS 180  
were run using the Desmond/Maestro software academic version 4.4<sup>34</sup> 181  
carried out in an NPT ensemble for about 50 ns. The first MDS 182  
considers the anthocyanins in their flavylium ion form, and the second 183  
MDS considers them in their hydrated form. The default relaxation 184  
protocol implemented in Desmond was used. The OPLS force field 185  
was applied to the system. From the results of the MDS, 1000 frames 186  
were extracted, which were analyzed using VMD 1.9.2 software.<sup>35</sup> 187

**Statistical Analyses.** All the experiments were carried out in 188  
triplicate (biological replicates). Statistical analyses, including means, 189  
standard deviations, and analysis of variance (ANOVA), were 190  
performed using Excel (Microsoft, Redmond, WA, USA). 191

## 192 ■ RESULTS AND DISCUSSION

**Effect of PVPP Treatment on Rosé Wine Colors.** As 193  
expected, the CIELAB coordinates of the wines were modified 194  
by the PVPP treatment. The values of the different color 195  
coordinates measured on the control wines and after treatments 196  
are available as supplementary data together with the 197  
corresponding statistical analyses (Appendix 4 of the SI). The 198  
normalized average measures (in percentage compared to 199  
control wines) for all samples are reported in Figure 2. 200



**Figure 2.** Average CIELAB coordinates measured in the six wines after PVPP treatment. Asterisks (\*) and (\*\*) on the tops of the bars indicate a significant difference with the control (Tukey (HSD):  $p < 0.05$  and  $p < 0.01$ , respectively). Results are normalized to the untreated control (100%).

201 Lightness  $L^*$  has increased by only 4% on average, which is a  
202 small change but statistically significant in all the wines. This  
203 limited rise can be explained by the fact that our wines had  
204 relatively high initial  $L^*$  values.

205 The  $a^*$  and  $b^*$  coordinates, respectively, decreased by 24%  
206 and 34% on average for all the wines, which indicated a  
207 statistically significant reduction of the red and yellow color  
208 components, respectively. We can then assume that PVPP  
209 affected color-active polyphenols such as anthocyanins and  
210 their derivatives and flavonols. This color drop can be due to  
211 the reduction of these pigment concentrations but also to the  
212 reduction of the copigmentation effect because of the  
213 adsorption of copigments.<sup>36–38</sup> The yellow color measured by  
214 the  $b^*$  coordinate may also be linked to oxidation phenomena  
215 which are very common in wines. PVPP may have removed  
216 orange pigments resulting from oxidation of compounds such  
217 as flavanols<sup>3–6,39</sup> or from reactions of anthocyanin pig-  
218 ments,<sup>8–10</sup> inducing a reduction of the  $b^*$  value.

219 Logically, the  $C^*$  parameter followed the same trend and  
220 decreased by 26%, reflecting a color intensity loss in the treated  
221 wines.

222 The  $h$  parameter also slightly decreased by 11%, which is the  
223 result of the  $b^*$  component being more affected by PVPP than  
224 the  $a^*$  one.

225 In all six wines, the  $\Delta E$  value between the colors of the wines  
226 before and after PVPP fining is superior to 1 (Appendix 4 of the

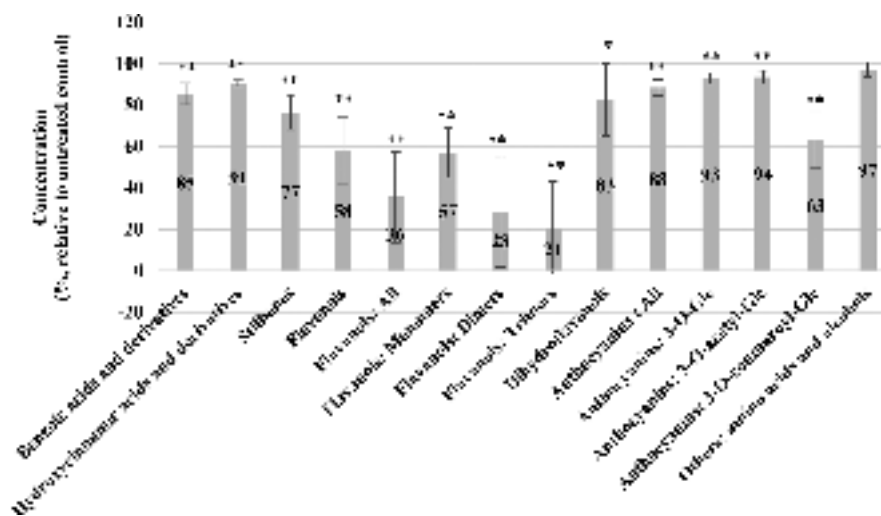
SI) meaning that a standard observer can see a difference in  
227 color. For two of the six wines,  $3, 5 < \Delta E < 5$ , so a clear  
228 difference in color is noticed. For the others,  $\Delta E > 5$ , so the  
229 observer notices two different colors.<sup>21</sup>

**Mass Spectrometry Results.** The polyphenol composition  
231 of the rosé wines was analyzed before and after treatment by  
232 PVPP by LC-MS/MS as previously described in Lambert et  
233 al.<sup>17</sup> The concentrations of the different compounds for all the  
234 wines (before and after treatment) are available as supple-  
235 mentary data (Appendix 5 of the SI).

236 For each molecule family, except for the alcohols and amino  
237 acids, the quantities in the treated wines are statically different  
238 from those measured in the initial wines according to the  
239 ANOVA analysis. However, the absorption capacity of rosé  
240 wine polyphenols to PVPP was very different from one family  
241 to another, confirming the existence of a very selective  
242 adsorption process (Figure 3). Three polyphenol families were  
243 the most affected by the PVPP treatment: flavonols, flavanols,  
244 and some anthocyanins, namely, coumaroylated anthocyanins.

245 Forty-two percent of the initial flavonols were adsorbed by  
246 PVPP. This is in accordance with the lower value of the  
247 CIELAB  $b^*$  value after treatment. Flavonols are pale yellow  
248 pigments,<sup>40</sup> and a positive  $b^*$  value stands for yellowish colors,  
249 so a smaller quantity of these molecules may have contributed  
250 to the reduction of the  $b^*$  coordinate. It is also likely that  
251 orange pigments such as xanthylum derivatives,<sup>6,7</sup> not targeted  
252 in our LC-MRM-MS method, are removed by the treatment.  
253 Furthermore, flavonols are a family of molecules involved in  
254 copigmentation<sup>36</sup> that is responsible for color enhancement in  
255 the wines by increasing the red color of anthocyanins. A  
256 reduction of the concentration of these molecules would limit  
257 the copigmentation effect and induce a reduction of the wine  
258 color, leading to a lower  $a^*$  value.

259 Concerning flavanols, 64% of their total content was  
260 adsorbed by PVPP, which represented the most impacted  
261 family of polyphenols on average for all the wines. Some  
262 important selectivity was observed within this group of  
263 polyphenols as adsorption increased with oligomerization.  
264 Indeed, trimers were slightly more adsorbed than dimers  
265 (79% vs 72%) and much more adsorbed than monomers  
266



**Figure 3.** Average rosé wine polyphenols concentrations measured in the six wines after PVPP treatment. Asterisks (\*) and (\*\*) on the tops of the bars indicate a significant difference with the control (Tukey (HSD):  $p < 0.05$  and  $p < 0.01$ , respectively). Results are normalized to the untreated control (100%).

267 (43%). These results are in accordance with the study  
 268 published by Mitchell et al.<sup>41</sup> where the authors showed that  
 269 tendency of the proanthocyanidins to bind to PVPP increased  
 270 with the degree of polymerization ( $(n = 3) > (n = 2) > (n =$   
 271  $1)$ ). As the number of units increases, so does the number of  
 272 hydroxyl groups and aromatic rings. This, respectively, implies  
 273 more hydrogen-bonding sites and hydrophobic interactions,  
 274 inducing a better PVPP affinity.<sup>14,16</sup> The same tendencies were  
 275 also reported for binding of flavanols to different proteins and  
 276 peptides, like salivary proteins, gelatins, casein, and poly-L-  
 277 proline, and precipitation of the resulting complexes that  
 278 increased significantly with the degree of polymerization of  
 279 proanthocyanidins.<sup>42–48</sup>

280 PVPP did not remove an important proportion of  
 281 anthocyanins with a mean adsorption capacity of 12% for all  
 282 forms. However, the coumaroylated anthocyanins (anthocya-  
 283 nin-3-O-coumaroyl-Glc) showed a much higher affinity toward  
 284 PVPP with a 37% average decrease. Although PVPP was  
 285 previously reported to adsorb anthocyanins in wines,<sup>49,50</sup> this is  
 286 the first time to our knowledge that such an affinity is described  
 287 for coumaroylated anthocyanins. This type of acylated  
 288 molecules with an additional phenol ring is less polar than  
 289 other anthocyanins.<sup>51</sup> Thus, a stronger hydrophobic interaction  
 290 might be responsible for this peculiar affinity.

291 The selective affinity of PVPP toward six phenolic  
 292 compounds was already investigated by Durán-Lara et al.<sup>52</sup>  
 293 They showed that PVPP exhibits a higher affinity for quercetin  
 294 (flavonol) and catechin (flavanol: monomer), a moderate  
 295 affinity for epicatechin (flavanol: monomer) and gallic acid  
 296 (benzoic acid), and a lower affinity for 4-methylcatechol  
 297 (alcohol) and caffeic acid (hydroxycinnamic acid). This is in  
 298 accordance with our results, where the affinity order for these  
 299 different polyphenol families is as follows: flavanols (mono-  
 300 mers)  $\approx$  flavonols  $>$  benzoic acids  $>$  hydroxycinnamic acids  $>$   
 301 alcohols.

302 **Computational Results.** Structure based computational  
 303 models were investigated to better understand the different  
 304 affinities observed between PVPP and rosé wine polyphenols  
 305 (results are shown in Table 1).

306 If we consider all components together, there is no  
 307 correlation between the polyphenol adsorption percentage by  
 308 PVPP and the calculated interaction energies. The correspond-  
 309 ing graph is available as supplementary data (Appendix 6 of the  
 310 SI).

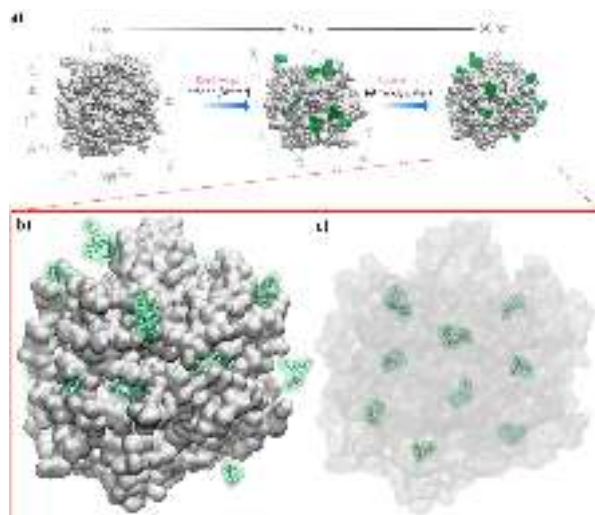
311 However, correlation within the flavanol family was observed  
 312 between the experimental adsorption and the calculated  
 313 interaction energy (Appendix 7 of the SI). For flavanols,  
 314 adsorption of trimers was higher than that of dimers that was  
 315 higher than that of monomers. The same trend can be observed  
 316 in the in-silico calculations: the best interaction energy is  
 317 obtained for the trimers, followed by the dimers and, finally, the  
 318 monomers (Table 1).

319 We can also observe that coumaroylated anthocyanins and  
 320 other anthocyanins have very different behaviors. At wine pH, it  
 321 is possible to find anthocyanins in their cationic form ( $A^+$ ) and  
 322 in greater proportion in their hydrated form (AOH). The latter  
 323 form is due to nucleophilic attack of water on the flavylum ion  
 324 of the anthocyanins. The anthocyanins under their hydrated  
 325 form showed higher interaction energies than their flavylum  
 326 ion form (Table 1). Also, when comparing these values with the  
 327 experimental adsorption, an improvement in the correlation  
 328 was obtained for anthocyanins in their hydrated form, with an  
 329  $r^2$  higher than 0.9 (Appendix 7b of the SI).

**Table 1. Adsorption Percentages and Interaction Energies Calculated at Semi-Empirical Quantum Mechanical Level**

id	name	adsorption percentage	interaction energy kcal mol <sup>-1</sup> (A <sup>+</sup> /AOH)
1	gallic acid	17 ± 6	-2.473
2	protocatechuic acid	10 ± 3	-2.422
3	syringic acid	1 ± 7	-2.572
4	ethyl ester of gallic acid	41 ± 14	-2.473
5	caffeic acid	12 ± 9	-2.548
6	cis-caftaric acid	10 ± 2	-2.633
7	trans-caftaric acid		-2.642
8	ferularic acid	6 ± 8	-2.594
9	ferulic acid	66 ± 26	-2.580
10	p-coumaric acid	3 ± 19	-2.513
11	cis-coutaric acids	12 ± 20	-2.535
12	trans-coutaric acids		-2.601
13	caffeic acid ethyl ester	5 ± 22	-2.542
14	coumaric acid ethyl ester	12 ± 21	-2.515
15	2 S-glutathionyl caftaric acid GRP (grape reaction product)	8 ± 8	-3.380
16	cis-piceid	7 ± 22	-2.973
17	trans-piceid	33 ± 42	-2.965
18	cis-resveratrol	83 ± 15	-2.657
19	trans-resveratrol	20 ± 30	-2.663
20	quercetin glucuronide	44 ± 16	-2.955
21	myricetin glucuronide	19 ± 12	-3.194
22	(+)-catechin	45 ± 14	-2.772
23	dimer cat B1 (procyanidin B1)	72 ± 27	-3.223
24	dimer cat B2 (procyanidin B2)	66 ± 30	-3.107
25	dimer cat B3 (procyanidin B3)	81 ± 23	-3.187
26	(-)-epicatechin	38 ± 12	-2.822
27	trimer cat1 (procyanidin C1)	74 ± 35	-3.518
28	trimer cat2 (procyanidin C2)	80 ± 21	-3.392
29	astilbin	17 ± 18	-3.004
30	delphinidin 3-O-Glc	15 ± 6	(-3.201/-3.423)
31	malvidin 3-O-Glc	6 ± 3	(-3.201/-3.312)
32	peonidin 3-O-Glc	8 ± 2	(-3.213/-3.465)
33	petunidin 3-O-Glc	12 ± 4	(-3.221/-3.488)
34	cyanidin 3-O-acetyl-Glc	12 ± 4	(-3.290/-3.480)
35	delphinidin 3-O-acetyl-Glc	15 ± 7	(-3.210/-3.494)
36	malvidin 3-O-acetyl-Glc	6 ± 3	(-3.303/-3.412)
37	peonidin 3-O-acetyl-Glc	6 ± 3	(-3.178/-3.443)
38	petunidin 3-O-acetyl-Glc	11 ± 5	(-3.225/-3.426)
39	cyanidin 3-O-coumaroyl-Glc	62 ± 22	(-3.447/-3.911)
40	delphinidin 3-O-coumaroyl-Glc	60 ± 26	(-3.370/-3.937)
41	malvidin 3-O-coumaroyl-Glc	34 ± 13	(-3.467/-3.543)
42	peonidin 3-O-coumaroyl-Glc	44 ± 16	(-3.376/-3.892)
43	petunidin 3-O-coumaroyl-Glc	56 ± 30	(-3.413/-3.854)
44	p-hydroxyphenylpyranomalvidin-3-O-Glc	19 ± 21	(-3.297/-3.438)
45	carboxypyranomalvidin 3-O-Glc = vitisin A	13 ± 18	(-3.344/-3.432)
46	tryptophan	14 ± 8	-2.793
47	tryptophol	3 ± 14	-2.481
48	tyrosine	2 ± 3	-2.574
49	tyrosol	2 ± 3	-2.237

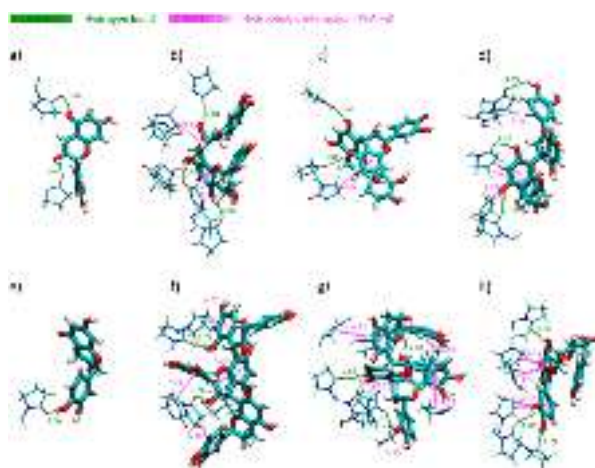
All-atom MDS were performed in order to understand the  
 molecular behavior between PVPP polymer and the 30  
 polyphenols that had the best interaction energies (Table 1)  
 immersed in a solvent box that simulates the main components  
 of the rosé wine (ethanol/water). Figure 4a shows the initial



**Figure 4.** Snapshots of MDS stages. (a) The initial state of PVPP–polyphenols system and their behavior at 25 and 50 ns. The polyphenols colored green are those that have been captured by PVPP, both in (b) the micropockets of the surface and in (c) their interior cavities.

state of PVPP–polyphenols system, and its behavior at 25 and 50 ns. The polyphenols colored with green are those that have been captured by PVPP. The PVPP microparticle has micropockets on the surface, which allow the interaction and capture of large polyphenols, such as anthocyanins and flavanols (dimers and trimers) (Figure 4b). The PVPP porous structure generates small cavities capable of capturing and retaining the smaller polyphenols (Figure 4c).

The studies with molecular simulation allowed observing the binding interactions that govern the affinity of PVPP for flavanols and coumaroyl anthocyanins, focusing on six polyphenols with great absorption capacity: procyanidin B1, procyanidin B2, procyanidin B3, procyanidin C1, procyanidin C2, and cyanidin-3-*O*-coumaroyl-Glc (Figure 5). Catechin and epicatechin were incorporated into the analysis by way of



**Figure 5.** Snapshots of PVPP–polyphenols binding interactions. These are the polyphenols with greater absorption capacity for the flavanols and anthocyanins families: (a) catechin, (b) procyanidin B1, (c) procyanidin B2, (d) procyanidin B3, (e) epicatechin, (f) procyanidin C1, (g) procyanidin C2, and (h) cyanidin-3-*O*-coumaroyl-Glc.

comparison. There is a presence of characteristic hydrogen bonds in all studied systems, the difference being in the number of bonds and their stability throughout the simulation (Appendix 8 of the SI). The catechin and epicatechin monomers (Figure 5a, e) can form only between one and two hydrogen bonds mainly with the regions of PVPP internal cavities, whereas their dimers, trimers, and cyanidin-3-*O*-coumaroyl-Glc can form between two and four hydrogen bonds (Appendix 8 of the SI) in the surface pockets of the PVPP particle (Figure 5b–d, f–h). Comparing the dimers and trimers of flavanols with the cyanidin-3-*O*-coumaroyl-Glc, it can be observed that hydrophobic interactions play an important role in their differentiation. The trimers are able to generate more  $\pi$ -alkyl type interactions between their aromatic rings and the PVPP backbone; this could enhance the affinity for the polymer, improving the absorption capacity of PVPP.

As discussed previously, the anthocyanins in their hydrated form showed higher interaction energies, compared to their flavylium ion form, and a strong correlation with adsorption percentage was evidenced. Anthocyanins in their hydrated form showed a better affinity for PVPP mainly because of the presence of hydrogen bonds that remained stable for more than 80% of the simulation time (Appendix 9 of the SI). These computationally studied models would indicate that the hydrated structures of the anthocyanins are the preferred forms for adsorption of phenolic compounds by PVPP.

Altogether, our results showed the high selectivity for the polyphenol adsorption in the process of treating rosé wines by PVPP. Further research is needed to include other polyphenols involved such as tannin-anthocyanin adducts or oxidized forms of polyphenols.

## ■ ASSOCIATED CONTENT

### ● Supporting Information

The Supporting Information is available free of charge on the ACS Publications website at DOI: 10.1021/acs.jafc.7b04461.

Appendix 1. Evolution of CIEL\**a*\**b*\* parameters of two wines exposed to PVPP for 7 h and statistics associated (Tukey (HSD):  $p < 0.05$ ). Appendix 2. Structures of the 49 targets considering their protonation state at the wine pH. Appendix 3. Methodology for design and study by SDM of the intermolecular properties of PVPP–polyphenols system. Appendix 4. Complete L\**a*\**b*\* results. Appendix 5. Complete UPLC-QqQ-MS results: concentrations before (C) and after treatment (PVPP) (mean  $\pm$  standard deviation,  $n = 3$ ). Appendix 6. Correlation between PVPP/polyphenols interaction energy calculations ( $\text{kcal mol}^{-1}$ ) and adsorption percentage measured in rosé wines for all the polyphenols. Appendix 7. Correlation between PVPP/polyphenols interaction energy calculations ( $\text{kcal mol}^{-1}$ ) and adsorption percentage measured in rosé wines for flavanols and anthocyanins. Appendix 8. Number of hydrogen bonds between PVPP and polyphenols with better affinity: flavanols (monomers, dimers, trimers) and cyanidin-3-*O*-coumaroyl-Glc. Appendix 9. Comparative graphs of the number of hydrogen bonds between PVPP and the five coumaroylated anthocyanins considering (a) their flavylium ion form and (b) their hydrated form (DOCX)

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