1	Theoretical Study on Isomerization of α-Acids:
2	A DFT Calculation
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#### 15 **ABSTRACT**:

16 The  $\alpha$ -acids contained in hops are one of the ingredients of beer. The isomerization of  $\alpha$ -acids produces iso- $\alpha$ -acids, the main source of bitterness in beer. In this study, the 17 18 isomerization mechanism of the  $\alpha$ -acid, cohumulone, was elucidated by using density 19 functional theory in conjunction with the polarizable continuum model or 3D-RISM integral 20 equation theory of liquids. The calculated reaction diagram is consistent with experimental 21 results; the activation free energy difference between the *cis* and *trans* isomers is in good 22 agreement with the experimental estimate. The activation energy difference results from 23 solvation energy. Additionally, a calculation of NMR chemical shifts showed that the proton position of isocohumulone is different from that proposed previously. The effect of Mg<sup>2+</sup> cation 24 25 on the isomerization was also investigated. Both the activation and reaction free energy are 26 stabilized by the presence of Mg<sup>2+</sup>, which is consistent with experimental results. Water 27 solvation reduces the activation free energy.

28 KEYWORDS: α-acid; iso-α-acid; isomerization; cohumulone; metal cation effect; solvent
29 effect

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#### 31 **1. Introduction**

Hops (*Humulus lupulus* L.) have been cultivated around the world, such as in Europe,
North America, Australia, and Asia (Delyser & Kasper, 1994). Among the many uses of hops,
its biological activities, such as its anti-inflammatory and antioxidant properties, have been

exploited for medical purposes (Zanoli & Zavatti, 2008). Hops are a well-known raw material
for beer and are one of the four essential ingredients of beer: barley malt, brewing water, yeast,
and hops (Denis De Keukeleire, 2000). Hops impart a characteristic aroma and taste to beer,
especially bitterness (Schönberger & Kostelecky, 2011).

The important chemical compounds in hop flowers are  $\alpha$ -acids. Unique side chains are found in some types of  $\alpha$ -acids, such as humulone, cohumulone, and adhumulone (Scheme 1).  $\alpha$ -acids do not have a bitter taste. Heating  $\alpha$ -acids results in thermal isomerization via an acyloin-type ring contraction. The isomerization products are iso- $\alpha$ -acids, the main bittertasting components of beer. These iso- $\alpha$ -acids have *cis/trans* isomers (Scheme 1) (Jaskula-Goiris et al., 2010).

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50 Many experimental studies have been performed to elucidate the isomerization
51 mechanism of α-acids (Askew, 1964; Jaskula et al., 2008; Malowicki & Shellhammer, 2005;

52 Ocvirk & Košir, 2020; Verzele & van Boven, 1971). For example, the isomerization rate was 53 measured in an aqueous buffer at pH 5.2 and 90–130°C (Malowicki & Shellhammer, 2005). The 54 isomerization rate was found to be first order in the  $\alpha$ -acid concentration, and the activation 55 energy for the overall *cis* and *trans* isomerization was found to be 23.6 kcal/mol. Jaskula et al. 56 (Jaskula et al., 2008) subsequently measured the *cis* and *trans* isomerization rates and proposed 57 the difference in reaction profiles between the *cis* and *trans* isomers. The activation energy for 58 the isomerization of *cis*-iso- $\alpha$ -acids is several kcal/mol higher than that of *trans*-iso- $\alpha$ -acids. 59 There is a 3.4-kcal/mol difference in the activation energies of *cis* and *trans* cohumulone. By contrast, *cis*-iso- $\alpha$ -acids have a lower reaction energy than *trans*-iso- $\alpha$ -acids. The 60 61 aforementioned studies determined the kinetics and thermodynamics of isomerization. 62 However, the molecular mechanism for isomerization has not been elucidated.

63 Many experimental studies on the structural analysis of iso- $\alpha$ -acids have also been carried out (D. De Keukeleire & Verzele, 1971; Hoek et al., 2001; Khatib et al., 2007; Urban et 64 65 al., 2013; Verzele & van Boven, 1971). The absolute configuration of *cis*-isohumulone was first determined as (4*R*, 5*S*) by Horeau's method in 1971 (D. De Keukeleire & Verzele, 1971; Verzele 66 67 & van Boven, 1971). This structure has been used for over 40 years, including to assign <sup>13</sup>C and <sup>1</sup>H NMR chemical shifts (Hoek et al., 2001; Khatib et al., 2007). In 2013, the absolute 68 configuration of *cis*-iso- $\alpha$ -acids was redetermined as (4*S*, 5*R*) by X-ray crystallography (Urban 69 70 et al., 2013). These measurements were used to identify almost the entire molecular 71 configuration. However, it is still difficult to accurately determine the positions of hydroxyl protons using these measurements. An alternative molecular configuration, 1-OH-PRD
(shown below), to the traditional configuration, 3-OH-PRD, is possible.

74 Isomerization yields are affected by various factors, such as the pH and metal cations 75 (Jaskula et al., 2010; Steenackers et al., 2015). Several metal cations have long been known to affect isomerization (Lance et al., 1975). In particular, the acceleration effect of Mg<sup>2+</sup> on 76 77 isomerization has been well studied (Köller, 1969). Isomerization with the addition of 78 magnesium oxide, MgO, at ambient temperature produces high yields with no side products. 79 Increasing the quantity of MgO added has been shown to increase the yield over fewer days of reaction (Kostrzewa et al., 2016). This effect of Mg<sup>2+</sup> on isomerization could be attributed to 80 81 complex formation between an  $\alpha$ -acid and a Mg<sup>2+</sup> cation, because  $\alpha$ -acids have a 1,4-diketone 82 structure. This O-Mg-O structure is also found in the RNA folding, where Mg<sup>2+</sup> cation is more 83 suitable than other cations such as Na<sup>+</sup> and Ca<sup>2+</sup> for its size and charge density (Petrov et al., 84 2011). However, Wietstock et al. recently carried out UV-VIS spectroscopy on  $\alpha$ -acid and 85 metal cation mixtures and proposed no complexation between  $\alpha$ -acids and Mg<sup>2+</sup> (Wietstock et al., 2016). The spectrum of equimolar mixtures of  $\alpha$ -acids and Fe<sup>2+</sup> was found to differ from 86 87 that of  $\alpha$ -acids alone, indicating complexation between  $\alpha$ -acids and Fe<sup>2+</sup>, whereas equimolar mixtures of  $\alpha$ -acids and Mg<sup>2+</sup> exhibited a similar spectrum to that of  $\alpha$ -acids alone. Thus, the 88 mechanism for the effect of Mg<sup>2+</sup> on isomerization remains controversial. 89

90 The purposes of this study are to theoretically examine the isomerization mechanism
91 of cohumulone (R=isopropyl), the hydroxyl proton position for isocohumulone, and the effect
92 of Mg<sup>2+</sup> addition on isomerization. There are some theoretical studies on the chemical reaction

93 of ingredients of foods and beverages (D. Milenković et al., 2018; D. A. Milenković et al., 2020; 94 Tošović et al., 2017). However, to the best of our knowledge, no theoretical studies on hop  $\alpha$ -95 acid isomerization have been performed to date. First, the transition state structure and the 96 activation free energy are analyzed to elucidate the isomerization mechanism. Next, the 97 position of the hydroxyl proton is determined by comparing calculated and experimentally 98 measured NMR chemical shifts. Finally, the activation free energy of isomerization with Mg<sup>2+</sup> 99 is decomposed into several components to identify the main effect of the metal cation on the 100 stability of isomerization.

101 **2.** Material and methods

102 We used density functional theory (DFT) in conjunction with the polarizable 103 continuum model (PCM) or three-dimensional reference interaction site model (3D-RISM) 104 integral equation theory (Beglov & Roux, 1997; Kovalenko & Hirata, 1998) to investigate the 105 isomerization mechanism of cohumulone, the NMR spectra of cohumulone and 106 isocohumulone, and the absorption spectra of cohumulone and complexes between 107 cohumulone and metal cations. The combination of quantum chemical method and 3D-RISM 108 theory is referred to as 3D-RISM-SCF method (Sato et al., 2000; Yoshida et al., 2011; Yoshida 109 & Hirata, 2006), which enables us to obtain molecular solvation structure around a solute. The 110 geometries of the reactants, intermediates, transition states, and products in solution were 111 optimized at the B3LYP-GD3(BJ)/6-31+G(d,p) level with the SMD solvent model (Marenich 112 et al., 2009). Notably, the dispersion correction, such as the D3 version of Grimme's dispersion

113 with Becke-Johnson damping, GD3(BJ) (Grimme et al., 2011), is needed to reproduce the 114 experimental reaction profile. To match the experimental conditions, the reaction profile and 115 absorption spectra were calculated in aqueous solution, whereas the NMR chemical shifts were 116 calculated in chloroform solution. Frequency calculations were used to obtain thermal 117 corrections to the Gibbs free energies at 25°C (Ochterski & Ph, 2000). Truhlar's quasiharmonic 118 approximation was adopted for the free energy calculations to correct for the spurious 119 overestimation of vibrational entropies introduced by treating low-frequency vibrational 120 modes as harmonic oscillators. This uses the same usual harmonic oscillator approximation in 121 the calculations of the vibrational partition functions, except that all of the vibrational 122 frequencies that are lower than 100 cm<sup>-1</sup> are set equal to 100 cm<sup>-1</sup>(Lam & Houk, 2014; Ribeiro 123 et al., 2011). The same geometries optimized at SMD-B3LYP-GD3(BJ)/6-31+G(d,p) level were 124 used for 3D-RISM-SCF calculations. The Kovalenko-Hirata closure is employed to solve the 125 3D-RISM calculation. The Lennard-Jones parameters for solutes were taken from the general AMBER force field (GAFF) (Wang et al., 2004). The simple point charge (SPC) (Berendsen et 126 127 al., 1981) model with modified hydrogen parameters ( $\sigma = 1.0$  Å and  $\varepsilon = 0.056$  kcal/mol) was 128 used for water solvent. The grid points in the 3D-RISM-SCF calculations were  $128 \times 128 \times 128$ 129 with a spacing of 0.5 Å. The gauge-independent atomic orbital (GIAO) (Ditchfield, 1974) 130 method and time-dependent DFT (TDDFT) were employed to calculate the NMR and 131 absorption spectra, respectively. NMR spectra are calculated by using the SMD model because 132 the solvent effect of non-polar chloroform is considered to be small. All the calculations were

- 133 performed using Gaussian 16 (Frisch et al., 2016), and a modified version of GAMESS (Schmidt
- 134 et al., 1993) where the 3D-RISM-SCF method has been implemented (Yoshida & Hirata, 2006).

# 135 **3.** Results and Discussion

# 136 **3-1. Isomerization reaction profile**

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139 Figure 1. Cohumulone isomerization reaction profile

We first analyzed the mechanism of cohumulone isomerization. Following previous studies, (D. De Keukeleire & Verzele, 1971; Steenackers et al., 2015) the reaction path in Scheme S1 (Supplementary Materials) was assumed. In the first step, keto-enol tautomerization generates the *cis* and *trans* intermediates, INT*cis* and INT*trans*, from the reactant

145	cohumulone (REACT). Next, INT cis and INT trans pass through two transition states, TS cis and
146	TS <sub>trans</sub> , and are converted into two products that both have a hydroxy group at the C1 position,
147	1-OH-PRD <sub>cis/trans</sub> . Finally, the equilibrium between 1-OH-PRD <sub>cis/trans</sub> and 3-OH-PRD <sub>cis/trans</sub> with
148	a hydroxy group at the C3 position is investigated. Note that 3-OH-PRD <sub>cisl trans</sub> are traditionally
149	regarded as the structures of <i>cis/trans</i> isocohumulone (Urban et al., 2013). It is also noted that
150	no direct involvement of water in the reaction was assumed because intramolecular ring
151	contraction and proton transfer are found to proceed at the TSs in a concerted manner.
152	The reaction profile calculated with the 3D-RISM-SCF method is shown in Figure 1.
153	Both INT cis and INT trans are unstable compared with REACT by $\sim$ 3 kcal/mol. The activation free
154	energy of TS <sub>cis</sub> (31.62 kcal/mol) is higher than that of TS <sub>trans</sub> (29.84 kcal/mol), whereas the <i>cis</i>
155	products, 1-OH-PRD <sub>cis</sub> and 3-OH-PRD <sub>cis</sub> , have lower reaction free energies (-1.35 and 1.42
156	kcal/mol, respectively) than the <i>trans</i> products, 1-OH-PRD <sub>trans</sub> and 3-OH-PRD <sub>trans</sub> (-0.58 and
157	1.73 kcal/mol, respectively). These calculated results are consistent with experimental results
158	(Jaskula et al., 2008). The calculated activation free energy difference of 1.78 kcal/mol between
159	the <i>cis</i> and <i>trans</i> products is in good agreement with the experimental value of 3.4 kcal/mol
160	(Malowicki & Shellhammer, 2005). Furthermore, 1-OH-PRDs are found to be more stable than
161	3-OH-PRDs, which suggests that 1-OH-PRDs are more realistic candidates for the structure
162	of isocohumulone (see discussion below).

164 **3-2.** Analysis of TS<sub>cis/trans</sub>

165 The difference in the activation free energies of the *cis* and *trans* products was analyzed. 166 We decomposed the free energy difference between the TS<sub>*cis*</sub> and TS<sub>*trans*</sub>,  $\Delta G = 1.78$  kcal/mol, 167 into three contributions, the structural energy difference in the gas phase  $\Delta E_{gas}$ , the solvation 168 free energy difference  $\Delta \mu_{sol}$ , and thermal free energy difference  $\Delta G_{thermal}$ :

169 
$$\Delta G = \Delta E_{\rm gas} + \Delta \mu_{\rm sol} + \Delta G_{\rm thermal} \tag{1}$$

170 where  $\Delta E_{gas} = E_{gas}(TS_{cis}) - E_{gas}(TS_{trans})$ , and  $\Delta \mu_{sol}$ , and  $\Delta G_{thermal}$  are defined analogously.  $\Delta E_{gas}$ 171 is calculated to be -4.71 kcal/mol, indicating that TScis has lower structural energy than TStrans. 172 On the other hand, TS<sub>trans</sub> has lower solvation free energy than TS<sub>cis</sub>,  $\Delta \mu_{sol} = 6.33$  kcal/mol.  $\Delta G_{\rm thermal}$  is found to be as small as 0.16 kcal/mol. These results demonstrate that solvation plays 173 174 a major role in the stabilization of TS<sub>trans</sub>. Notably, the sophisticated 3D-RISM-SCF method was 175 needed for the proper description of solvation. The SMD solvation model failed to reproduce 176 the experimental result; the free energy of TS<sub>trans</sub> (28.43 kcal/mol) calculated with the SMD 177 model is similar to that of TS<sub>cis</sub> (28.40 kcal/mol).

178Further analysis decomposing the solvation free energy into each atom contribution179showed that the main reason for the difference is the C2 atom, the center of the β-triketo180system. TS<sub>cis</sub> has low structural energy due to the interaction between the C2 atom and side181chain whereas TS<sub>trans</sub> has low solvation free energy due to the strong solvation of the C2 atom.182The importance of the stabilization of the β-triketo system for the isomerization was suggested183in the previous study (Jaskula-Goiris et al., 2010).

# 185 **3-3.** Proton position

186 Experimental studies have reported 3-OH-PRD as the isomerization end product 187 (Urban et al., 2013). However, the calculated reaction profile showed that 1-OH-PRD is more 188 stable than 3-OH-PRD (Figure 1). We analyzed the proton position in more detail by 189 calculating the <sup>13</sup>C and <sup>1</sup>H NMR chemical shifts of REACT, PRD<sub>cis</sub>, and PRD<sub>trans</sub> for comparison 190 with experimental data (Hoek et al., 2001; Khatib et al., 2007). Although direct comparison is 191 precluded by the unavailability of NMR data for hydroxyl protons, a useful comparison can be 192 made for NMR data for <sup>13</sup>C and other protons. The atomic numbering of REACT, PRD<sub>cis</sub>, and 193 PRD*trans* is shown in Figure S1.

First, the NMR spectra of REACT were used as a reference. The calculated and experimental NMR chemical shifts of REACT and the correlation between these results are shown in Table S1 and Figure 2. Reasonably good correlations are obtained for both <sup>13</sup>C and <sup>1</sup>H, with coefficients of determination, R<sup>2</sup>, of 0.9980 and 0.9947, respectively. This result validates our NMR calculation.

Figure 2 shows the correlations between the experimental and calculated NMR chemical shifts for 3-OH-PRDs (the proposed isocohumulone structure) and 1-OH-PRDs (the lower free energy). The calculated and experimental NMR chemical shifts of 3-OH-PRDs and 1-OH-PRDs are summarized in Tables S2 and S3. The R<sup>2</sup> of the aforementioned correlations for 1-OH-PRDs are larger than those for 3-OH-PRDs for both the <sup>13</sup>C and <sup>1</sup>H NMR chemical shifts of the *cis* and *trans* products. For example, the correlation for <sup>13</sup>C NMR of 1-OH-PRD<sub>cis</sub>

has a R<sup>2</sup> of 0.9987, which is larger than the R<sup>2</sup> of 0.9944 for the corresponding 3-OH-PRD<sub>cis</sub> correlation. These results show that better correlations between the calculated and experimental results were obtained for the stable 1-OH-PRDs than for 3-OH-PRDs in both the *cis* and *trans* configurations. Therefore, 1-OH-PRD is considered to be the actual  $\alpha$ -acid isomerization product, not 3-OH-PRD.

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Figure 2. Correlations between experimental and calculated NMR chemical shifts for 0) cohumulone, 1) 3-OH-PRD<sub>cis</sub>, 2) 1-OH-PRD<sub>cis</sub>, 3) 3-OH-PRD<sub>trans</sub>, and 4) 1-OH-PRD<sub>trans</sub>: a) <sup>13</sup>C and b) <sup>1</sup>H

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### 216 3-4. Reaction profile with Mg<sup>2+</sup>

The effect of the magnesium cation on the isomerization reaction profile was investigated. The DFT calculation was used to determine the most stable structures of the Mg complexes of cohumulone and isocohumulone, which were used to propose the reaction scheme with magnesium cation shown in Scheme S2. First, cohumulone becomes an anion by
protonating a water molecule and forms the Mg-REACT complex, which includes a 6membered ring structure with O-Mg-O coordination. Because Mg<sup>2+</sup> cation is generally hexacoordinated, 4 water molecules were additionally coordinated to Mg<sup>2+</sup> cation. Next, MgREACT transforms into the Mg-INT intermediates, which pass through transition states, MgTSs, and finally generate the products, Mg-PRDs.

226 The reaction profile of the isomerization of cohumulone with the Mg cation calculated

227 with the 3D-RISM-SCF method is shown in Figure 3b. The reaction profile of anionic

228 cohumulone is included in Figure 3a for comparison.



**Figure 3.** Isomerization reaction profiles for a) anionic cohumulone and b) Mg<sup>2+</sup> complex.

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Table 1 shows the activation free energy and the reaction free energy for the aforementioned isomerization reactions. The activation and reaction free energies of the Mg complex are approximately 5 kcal/mol lower than those of the neutral states. For example, the activation free energy of the Mg complex in the *trans* configuration (22.43 kcal/mol) is lower

than for anionic and neutral cohumulone (23.36 and 29.84 kcal/mol, respectively). This result
is consistent with experimental results (Jaskula et al., 2010; Köller, 1969; Kostrzewa et al., 2016).

	$\Delta G^{\ddagger}$		$\Delta_{ m r} G$	
	trans	cis	trans	cis
$Mg^{2+}$	22.43	26.62	-8.41	-5.05
anion	23.36	25.32	-5.61	-4.85
neutral	29.84	31.62	-0.58	-1.35

**Table 1.** Activation and reaction free energy differences (in kcal/mol)

We determined the cause of the decrease in the activation free energy of the Mg complex by decomposing the activation free energy  $\Delta G^{\ddagger}$  into three contributions, the structural energy in the gas phase  $\Delta E_{gas}$ , the solvation free energy difference  $\Delta \mu_{sol}$  and the thermal free energy difference  $\Delta G_{thermal}$  between TS and REACT:

244 
$$\Delta G^{\ddagger} = \Delta E_{\text{gas}} + \Delta \mu_{\text{sol}} + \Delta G_{\text{thermal}}$$
(2)

where  $\Delta E_{gas} = E_{gas}(TS) - E_{gas}(REACT)$ , and  $\Delta \mu_{sol}$  and  $\Delta G_{thermal}$  are defined analogously. We evaluated the activation free energy differences between the Mg complexes and neutral cohumulone to compare the relative size of each contribution:

248 
$$\Delta\Delta G^{\ddagger} = \Delta\Delta E_{\text{gas}} + \Delta\Delta \mu_{\text{sol}} + \Delta\Delta G_{\text{thermal}}$$
(3)

where  $\Delta\Delta G^{\ddagger} = \Delta G^{\ddagger}(Mg^{2+}) - \Delta G^{\ddagger}(neutral)$ , and  $\Delta\Delta E_{gas}$ ,  $\Delta\Delta\mu_{sol}$ , and  $\Delta\Delta G_{thermal}$  are defined analogously. The calculated results for the *cis* and *trans* activation free energies are shown in Table 2. The solvation energy difference contributes considerably to the decreases in the activation free energies of –13.61 and –27.47 kcal/mol for the *cis* and *trans* configurations, respectively. This result shows that one of the major factors that stabilize the TS of Mg complexes is the large difference in the solvation free energies between REACT and the TS of Mg complexes.

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Mg <sup>2+</sup> –neutral	$\Delta\Delta G^{\sharp}$	$\Delta\!\Delta E_{ m gas}$	$\Delta\Delta\mu_{ m sol}$	$\Delta\!\Delta G_{ m thermal}$
cis	-5.00	9.39	-13.61	-0.79
trans	-7.41	20.35	-27.47	-0.30

257 **Table 2.** Decomposition analysis of activation free energy differences (in kcal/mol)

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259 Finally, the effect of complex formation on the absorption spectra was investigated by 260 calculating UV-VIS absorption spectra at the TD-B3LYP-GD3(BJ)/6-31+G(d,p) level for the 261 neutral and anionic states and the  $Mg^{2+}$ ,  $Fe^{2+}$ , and  $Fe^{3+}$  complexes (Figure S2). As in the case of 262 the Mg<sup>2+</sup> complex, 4 water molecules were additionally coordinated to Fe<sup>2+</sup> and Fe<sup>3+</sup> complexes. 263 In Fe<sup>2+</sup> and Fe<sup>3+</sup> complexes, only SMD results are shown because of the difficulty of calculating 264 the higher excited states using 3D-RISM-SCF. It is noted that there is no qualitative difference 265 between 3D-RISM-SCF and SMD results in the lower excited states. Considering the long-266 wavelength tail, the spectrum of the Mg<sup>2+</sup> complex (Figure S2c) is similar to those of the neutral 267 and anionic states (Figures S2a and S2b). In these cases, no peak is found at >400 nm, and the

268 longest-wavelength (lowest-energy) peak corresponding to the  $\pi$ - $\pi$ \* excitation is located at 269 approximately 350 nm. By contrast, the spectra of the Fe<sup>2+</sup> and Fe<sup>3+</sup> complexes (Figures S2d and 270 S2e) are very different from that of REACT. The spectra of these complexes exhibit several 271 peaks at >400 nm and long-wavelength tails. The complexes have an empty low-energy *d*-type 272 molecular orbital that enables ligand-to-metal charge transfer (LMCT) transitions to proceed 273 at a low excitation energy. These results are consistent with experimental results (Wietstock 274 et al., 2016) and indicate that it is difficult to determine the complexation of Mg<sup>2+</sup> cation with 275  $\alpha$ -acids using UV-VIS spectra alone.

#### 276 **5.** Conclusions

277 In summary, the mechanism for the isomerization of cohumulone, the hydroxyl proton 278 position of isocohumulone, and the effect of Mg<sup>2+</sup> addition on isomerization were theoretically 279 investigated. The calculated reaction profile and the activation free energy difference between 280 cis and trans configurations are consistent with experimental results. The activation free 281 energy difference between the *cis* and *trans* configurations is attributed to the solvation energy. 282 An analysis of the calculated reaction profile and the NMR spectra showed that 1-OH-PRD is 283 the actual isomerization product, not 3-OH-PRD. The isomerization reaction profile for cohumulone with Mg<sup>2+</sup> demonstrated that Mg<sup>2+</sup> increases iso- $\alpha$ -acids yields and accelerates 284 285 isomerization, which is in good agreement with experimental results. The solvent effect of 286 water contributes considerably to the decrease in the activation free energy of the 287 isomerization reaction with Mg<sup>2+</sup>. Similar spectra were obtained for the Mg<sup>2+</sup> complex and

288	cohumulone, showing that it is difficult to determine complexation using UV-VIS spectra
289	alone.
290	
291	Declaration of Competing Interests
292	The authors declare no competing financial interest.
293	
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298	
299	Appendix A. Supplementary data
300	Supplementary data associated with this article can be found, in the online version, at
301	http://dx.doi.org.
302	
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