

1 **Combined Virtual/Experimental Multicomponent Solid Forms Screening of Sildenafil: New Salts,**  
2 **Cocrystals, and Hybrid Salt–Cocrystals**

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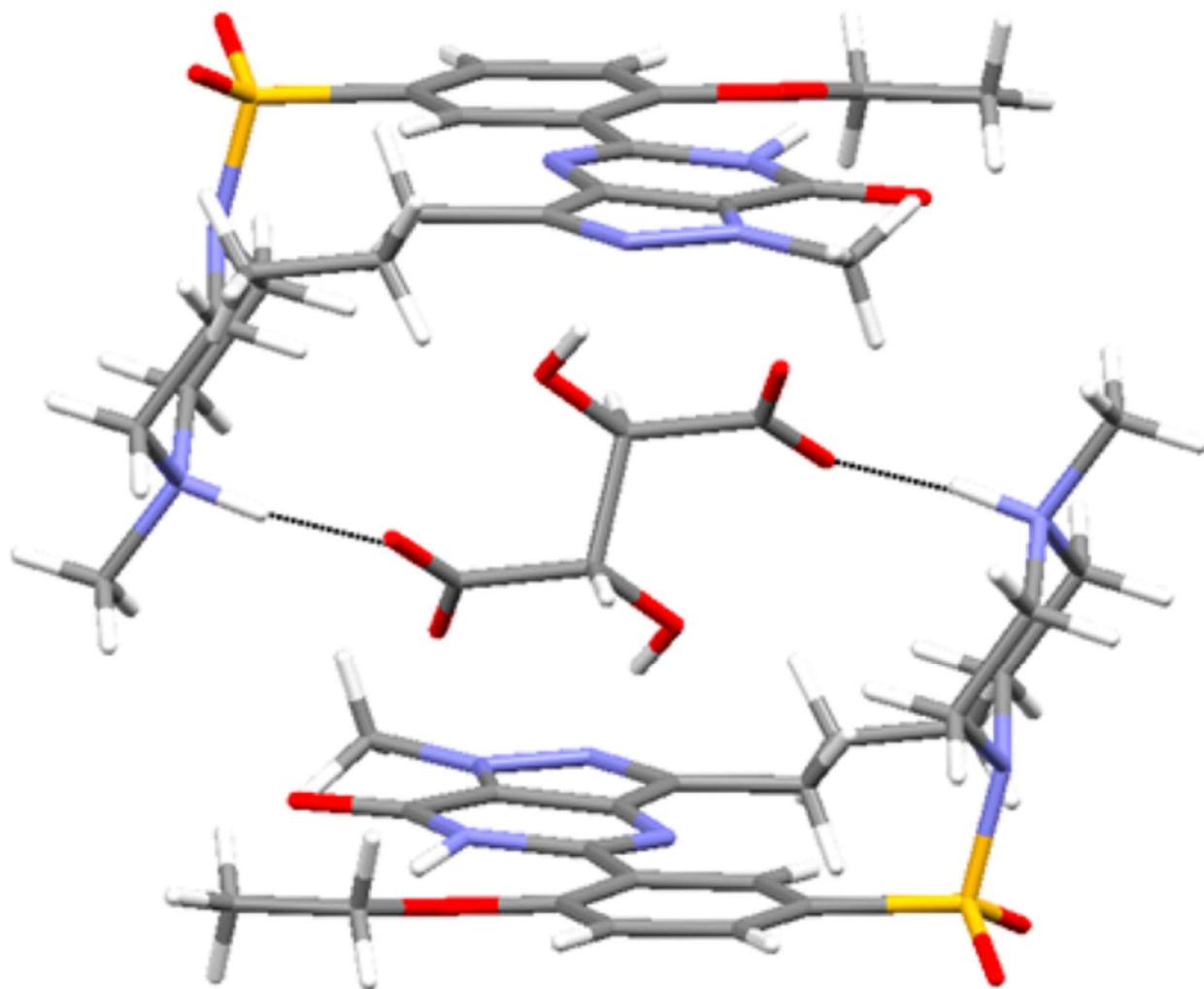
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43 **ABSTRACT:**

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45 New multicomponent solid forms of sildenafil have been discovered  
46 by means of a combined virtual/experimental cocrystal screening. Coformer  
47 selection of candidates was conducted based on an in silico screening method from  
48 a database of more than 2000 organic compounds, and the intensive experimental  
49 screen produced 23 new solid forms. Since the 12 cofomers chosen have a  
50 combination of phenol and carboxylic acid groups, a variety of cocrystals, salts, and  
51 hybrid salt-cocrystals were discovered and characterized.

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## 60 1. INTRODUCTION

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62 Cocrystals of active pharmaceutical ingredients (APIs) have received massive attention over the past  
63 decade because they offer many opportunities to improve physicochemical properties of drugs.<sup>1,2</sup>  
64 Indeed, solubility is one of the most important properties for a drug compound since it has a direct  
65 impact on bioavailability, and the cocrystal approach is a versatile toolbox to tune this and another  
66 important property<sup>3,4</sup> such as stability<sup>5</sup> because of the high number of available potential cofomers.  
67 On February 2018, the Food and Drug Administration (FDA) released a final guidance titled  
68 “Regulatory Classification of Pharmaceutical Co-Crystals Guidance for Industry”,<sup>6</sup> providing applicants  
69 planning to submit new drug applications with information on the regulatory classification of  
70 pharmaceutical cocrystals, classifying them as a new form of the API, comparable in that respect to  
71 polymorphs, hydrates, salts, etc. The FDA asks the applicants to provide evidence to demonstrate that  
72 “both the API and cofomers are present in the unit cell” and “the component API and cofomer co-exist  
73 in the cocrystal which interact nonionically”. The FDA guidance suggests that the applicant consider the  
74 difference of  $\Delta pK_a$  between the API and the cofomer or to provide evidence that proton transfer has not  
75 occurred in the lattice by means of spectroscopic tools or other orthogonal approaches. Thus, from a  
76 regulatory point of view, it is very important to assess the proton transfer in a multicomponent  
77 Sildenafil, the active principle of Viagra, is the first oral drug used for the medical treatment of erectile  
78 dysfunction in elderly patients, and it was initially used as an antihypertensive drug<sup>8,9</sup> but due to its  
79 poor aqueous solubility and low bioavailability it is generally formulated as sildenafil citrate.<sup>10</sup> This  
80 solid form still exhibits moderate bioavailability, and this is the reason why some efforts have been  
81 conducted to discover new salts and cocrystals with enhanced physicochemical properties. In this sense,  
82 a sildenafil/acetylsalicylic cocrystal exhibiting enhanced intrinsic dissolution rate compared to sildenafil  
83 citrate has been reported.<sup>11</sup> Moreover, pharmacokinetics of salts and cocrystals of sildenafil with  
84 dicarboxylic acids has been studied, and the glutarate salt was revealed to be a good candidate for  
85 alternative formulation of the citrate salt.<sup>12</sup> The crystal structures of sildenafil base, sildenafil citrate  
86 monohydrate, sildenafil saccharinate, and sildenafil acesulfamate have been published in the  
87 literature,<sup>13,14</sup> and some of us have described a polymorph of sildenafil free base and new  
88 solvates.<sup>15,16</sup> With the aim to discover new multicomponent forms and extend the solid state  
89 knowledge of this important API, we have conducted a combined virtual/experimental salt/cocrystal  
90 screening by using a broad set of thermodynamic and kinetic experimental conditions. Twenty-three  
91 new solid forms of sildenafil, including salts, cocrystals, and hybrid salt–cocrystals, have been  
92 discovered and some of their crystal structures solved.

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98 **2. MATERIALS AND METHODS**

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100 **2.1. Materials.** Sildenafil (SIL) used in this study was of reagent grade and used as received from  
101 Polpharma (form I). The cofomers quercetin (QUE), methyl gallate (MEG), tartaric acid (TAR), 3-  
102 hydroxybenzoic acid (3-HBA), 4-hydroxybenzoic acid (4-HBA), resorcinol (RES), 3,4-  
103 dihydroxybenzoic acid (3,4-DHBA), and caffeic acid (CAF) were purchased from Sigma-Aldrich.

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105 **2.2. Methods.** 2.2.1. Virtual Cocrystal Screening. For each compound, the molecule was drawn in an  
106 extended conformation and energy minimized using the molecular mechanics methods implemented in  
107 TorchLite.17 Gaussian 09 was used to optimize the geometry and calculate the MEPS on the 0.002 Bohr  
108 Å<sup>-3</sup> electron density isosurface using density functional theory (DFT) and a B3LYP/6-31G\* basis  
109 set.18 The MEPS was converted into SSIPs using in-house software.19

110 2.2.2. Cocrystal Screening. Screening through liquid assisted grinding experiments (LAG) was  
111 conducted by grinding 20–35 mg of a 1:1 mixture of SIL and each cofomer together with one drop of  
112 different solvents using a Retsch MM 2000 grinding mill. The samples were placed in 2 mL volume  
113 stainless steel jars, along with two stainless tungsten grinding balls of 3 mm diameter. Grinding was  
114 performed for 15–30 min, with a frequency of the mill of 30 Hz. Finally, the samples were collected  
115 immediately without prior drying for powder X-ray diffraction (PXRD) analysis. The formation of a  
116 new solid form was determined by comparing PXRD patterns of starting materials and products from  
117 cocrystal screening LAG experiments. Screening through reaction crystallization (RC) was conducted  
118 by preparing a saturated solution of the most soluble component (SIL or cofomer) in different solvents  
119 in a sealed vial under stirring. A small quantity of the less soluble component was added until it did not  
120 dissolve anymore. The suspension was stirred at different times, and the resulting solids were filtered  
121 and analyzed by PXRD. Screening through solvent mediated transformations (SMT) were conducted by  
122 preparing suspensions of SIL and cofomer in different molar ratios (40–1200mg of the final mixture) in  
123 selected solvents. The sealed vials were stirred for different times, and the resulting solids were filtered  
124 and analyzed by PXRD.

125 2.2.3. Solution Crystallization. Solutions of SIL/cofomer in a 1:1 molar ratio (10–20 mg of the final  
126 mixture) were prepared in different solvents and heated in a heating stainless steel block. The heater was  
127 switched off, and the solutions were allowed to slowly cool down to 25°C inside the heating block. The  
128 samples which did not crystallize were tightly sealed and kept at 25 °C until precipitation was observed.

129 2.2.4. Synthesis of the Different Crystal Forms of Sildenafil. Details of synthesis and characterization of  
130 each form can be found in Supporting Information (see section 1 and Table S1). Stoichiometry has been  
131 assessed based on NMR and thermogravimetric analysis (TGA) measurements when the crystal  
132 structure is not available. In those cases where the crystal structure has not been solved, the definition of  
133 the form as a salt or a cocrystal has been done based on the probability of proton transfer determined  
134 with eq 3.

135 Twenty-three multicomponent forms of SIL (cocrystals, salts, and hybrid salt–cocrystal) have been  
136 obtained through a cocrystal screening with 8 out of the 12 cofomers used. Five cocrystal forms of SIL  
137 have been obtained with three cofomers: two forms with quercetin in a 1:1 stoichiometry (one as an  
138 isopropanol solvate, SILQUE I, and one as a tetrahydrofuran solvate, SIL-QUE II); two with resorcinol  
139 in two different stoichiometries (one in a 1:1 molar ratio, SIL-RES I, and one in a 1:2 molar ratio, SIL-  
140 RES II); one form with methyl gallate in a 1:1 stoichiometry, SIL-MEG. Fourteen salts of SIL have been  
141 obtained with five cofomers: one form with 3,4-dihydroxybenzoic acid in a 1:1 stoichiometry as an  
142 isopropanol solvate, SIL-3,4-DHBA I; four forms with tartaric acid in two different stoichiometries: two  
143 in a 1:1 molar ratio (an anhydrous form, SIL-TAR I, and an isopropanol solvate, SIL-TAR III); two in a  
144 2:1 molar ratio (an anhydrous form, SIL-TAR II, and an isopropanol solvate, SIL-TAR IV); two forms  
145 with caffeic acid in a 2:3 stoichiometry (one as an anhydrous form, SIL-CAF I, and one as a  
146 monohydrate, SIL-CAF II); three forms with 3-hydroxybenzoic acid in a 1:1 stoichiometry (one as an  
147 acetonitrile solvate, SIL-3-HBA I, one as a tetrahydrofuran solvate sesquihydrate, SIL-3-HBA II, and  
148 one as an anhydrous form, SIL-3-HBA III); four forms with 4-hydroxybenzoic acid in a 1:1  
149 stoichiometry (two as anhydrous forms, SIL-4-HBA I and SIL-4-HBA III, one as a hemiisopropanol  
150 solvate, SIL-4-HBA II, and one as a tetrahydrofuran solvate, SIL-4-HBA IV). Four hybrid salt-cocrystal  
151 forms of SIL have been obtained with two cofomers: three forms with 3,4-dihydroxybenzoic acid in  
152 two different stoichiometries (two of them in a 1:2 molar ratio as acetonitrile solvates, SIL-3,4-DHBA II

153 and SIL-3,4-DHBA III, and one in a 2:3 molar ratio as a dehydrate, SIL-3,4-DHBA IV) and finally one  
154 form with 3-hydroxybenzoic acid in a 2:3 stoichiometry as a dehydrate, SIL-3-HBA IV.

155 2.2.5. X-ray Crystallographic Analysis. Single crystal X-ray diffraction intensity data of the different  
156 crystal forms of sildenafil were collected using a D8 Venture system equipped with a multilayer  
157 monochromator and a Mo microfocus ( $\lambda = 0.71073 \text{ \AA}$ ). Frames were integrated with the Bruker SAINT  
158 software package using a SAINT algorithm. Data were corrected for absorption effects using the  
159 multiscan method (SADABS).<sup>20</sup> The structures were solved and refined using the Bruker SHELXTL  
160 Software Package, a computer program for automatic solution of crystal structures and refined by  
161 fullmatrix least-squares method with ShelXle Version 4.8.0, a Qt graphical user interface for SHELXL  
162 computer program.<sup>21</sup>

163 Powder X-ray diffraction patterns were obtained on a PANalytical X'Pert PRO MPD diffractometer in  
164 transmission configuration using Cu K $\alpha$ 1 + 2 radiation ( $\lambda = 1.5406 \text{ \AA}$ ) with a focusing elliptic mirror  
165 and a PIXcel detector working at a maximum detector's active length of 3.347°. Configuration of  
166 convergent beam with a focalizing mirror and a transmission geometry with flat sample sandwiched  
167 between low absorbing films measuring from 2 to 40° in 2 $\theta$ , with a step size of 0.026° or from 2 to 70°  
168 in 2 $\theta$ , with a step size of 0.013° with measuring times of 30 min to 4 h. The powder diffractograms were  
169 indexed, and the lattice parameters were refined by means of LeBail fits by means of Dicvol04,<sup>22</sup> and  
170 the space groups were determined from the systematic absences. A summary of crystal data and relevant  
171 refinement parameters are given in Tables 1 and 2.

172 2.2.6. Differential Scanning Calorimetry (DSC). Differential scanning calorimetry analysis were carried  
173 out by means of a Mettler-Toledo DSC-822e calorimeter. Experimental conditions: aluminium crucibles  
174 of 40  $\mu\text{L}$  volume, atmosphere of dry nitrogen with a 50 mL/min flow rate, and heating rate of 10 °C/min.  
175 The calorimeter was calibrated with indium of 99.99% purity (m.p.: 156.4 °C,  $\Delta H$ : 28.55 J/g).

176 2.2.7. Thermogravimetric Analysis (TGA). Thermogravimetric analyses were performed on a Mettler-  
177 Toledo TGA-851e thermobalance. Experimental conditions: alumina crucibles of 70  $\mu\text{L}$  volume,  
178 atmosphere of dry nitrogen with 50 mL/min flow rate, and a heating rate of 10 °C/min.

179 2.2.8. Nuclear Magnetic Resonance (NMR). Proton nuclear magnetic resonance (1H NMR) spectra was  
180 recorded on a Varian Mercury 400 (400 MHz). Chemical shifts for proton are reported in parts per  
181 million (ppm) downfield from tetramethylsilane and are referenced to residual proton in the NMR  
182 solvent (DMSO-d<sub>6</sub>:  $\delta$  2.50). Experimental conditions: delay: 1; pulse: 45°; scans: 32 or 64.

183 2.2.9. Dissolution Study. The dissolution measurements were carried out only with the solid forms that  
184 could be obtained pure in amounts sufficient to perform a dissolution study: pure sildenafil, salts of  
185 sildenafil with citric acid, tartaric acid, 3-HBA and 3,4-DHBA, a cocrystals with RES, QUE, and two  
186 hybrid salt-cocrystals of sildenafil with 3,4-DHBA. The dissolution was determined in 0.1 N HCl (pH  
187 1.2), phosphate buffer pH 6.5, and a biorelevant dissolution medium fasted state simulated intestinal  
188 fluid (FaSSIF) at 25 °C. For dissolution studies 40 mg of crystalline compounds were added to the  
189 dissolution medium stirred at 100 rpm over 24 h, and samples were withdrawn at 1 and 24 h. The  
190 amount of SIL dissolved in 1 h (D1h) and 24 h (D24h) was determined using the HPLC technique. The  
191 details about dissolution medium and HPLC method are provided in Supporting Information.

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### 195 3. RESULTS AND DISCUSSION

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197 **3.1. Virtual Cocrystal Screen.** We selected the cofomers for experimental screening based on the  
198 virtual cocrystal screening methodology developed by some of us to predict the probability of cocrystal  
199 formation.<sup>23</sup> This computational tool has been validated using experimental data extracted from the  
200 literature.

201 The difference between the calculated energy of the cocrystal and the pure components was used to rank  
202 potential cofomers.<sup>24</sup> This approach uses surface site interaction points (SSIPs) calculated from the ab  
203 initio molecular electrostatic potential surface (MEPS) of the isolated molecule in the gas phase.<sup>19,25</sup>  
204 The interaction of a molecule with its environment is described by a discrete set of SSIPs, each  
205 represented by an interaction parameter,  $\epsilon_i$ , which is positive for a H-bond donor site (or positive region  
206 on the MEPS) and negative for a H-bond acceptor site (or negative region on the MEPS). The energy of  
207 interaction between two SSIPs,  $i$  and  $j$ , is given by the product  $\epsilon_i\epsilon_j$ . We assume that pairwise interactions  
208 between SSIPs are optimized in a solid, and this provides a method for evaluating the interaction site  
209 pairing energy of a solid without knowledge of the crystal structure.<sup>26</sup> The most positive SSIP is paired  
210 with the most negative SSIP, the next most positive SSIP with the next most negative, and so on, giving  
211 a hierarchical list of interactions.<sup>27,28</sup> This interaction site pairing strategy provides a straightforward  
212 method for estimating the energy of a solid,  $E$  (eq 1). The same approach can be used to estimate the  
213 energy of a cocrystal, and the difference between the interaction site pairing energies of the cocrystal  
214 and the pure components,  $\Delta E$ , can be used to estimate the probability of cocrystal formation (eq 2).

215

$$216 E = \sum \epsilon_i \epsilon_j \quad (1)$$

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$$218 \Delta E = -(E_{cc} - E_1 - E_2) \quad (2)$$

219

220 where  $E_1$ ,  $E_2$ , and  $E_{cc}$  are the interaction site pairing energies of the pure solids, 1 and 2, and a 1:1  
221 cocrystal respectively. Note that this definition means that  $\Delta E$  is always positive, and a large value  
222 indicates a high probability of cocrystal formation.

223 Some of us have previously applied the method to successfully predict the formation of new  
224 cocrystals,<sup>29,30</sup> and in this work we have followed this theoretical approach to guide the selection of a  
225 limited number of cofomers to test experimentally. Thus, the difference between the interaction site  
226 pairing energies of the 1:1 cocrystal and the pure components was calculated for each  
227 sildenafil/coformer combination using a coformer database which contains more than 2000 organic  
228 compounds (including 860 products from the GRAS list). The cofomers were ranked in order of  
229 decreasing  $\Delta E$ , and only 12 cofomers were chosen from the top 100 compounds according to toxicity  
230 criteria and probability of success in a cocrystallization experiment (Table 3). This theoretical approach

231 defines an energy threshold of 11kJ/mol where the probability of cocrystal formation is higher than  
232 50%. Thus, only coformers with  $\Delta E > 11$  kJ/mol were chosen for experimental screening  
233 Since sildenafil has a strong basic group, the formation of salts with strong carboxylic acids is expected.  
234 In fact, the formation of a salt or a cocrystal can be assessed based on the “rule of thumb”<sup>31</sup> which  
235 states that salts are formed when  $\Delta pK_a$  [ $pK_a(\text{base}) - pK_a(\text{acid})$ ]  $\geq 3$ , and a cocrystal is expected when  
236 this value is  $\leq 0$ , the combinations with a value  $0 \leq [pK_a(\text{base}) - pK_a(\text{acid})] \leq 3$  being much less reliable  
237 and falling around a “salt-cocrystal continuum” region.<sup>32</sup> This uncertainty motivated the analysis and  
238 correlation by Cruz-Cabeza<sup>33</sup> of a big set of experimental cocrystal/salt data in order to develop a more  
239 reliable equation to predict the salt/cocrystal outcome. According to this statistical analysis, eq 3 allows  
240 prediction of the probability of proton transfer around the region of  $\Delta pK_a$  values between  $-1$  and  $4$ .

$$241 \\ 242 P(\%) = 17\Delta pK_a + 28 \quad (3)$$

243  
244 Sildenafil has a basic functional group (piperazine) with a  $pK_a$  value of 6.78,<sup>34</sup> and we have applied this  
245 statistical approach to the coformers with acidic groups selected from the virtual cocrystal screening to  
246 assess the probability of salt formation (Table 4). Coformers with acidic groups such as  
247 (3-hydroxybenzoic acid, 4-hydroxybenzoic acid, caffeic acid, 3,4-dihydroxybenzoic acid, and tartaric  
248 acid) were expected to form salts. However, salt stoichiometry is an important outcome not always easy  
249 to predict because hybrid salt–cocrystal forms are also possible. In this sense, there are interesting  
250 examples in the literature with unexpected stoichiometries due to the presence of nonionized molecules  
251 in the crystal structure such as the *p*-coumaric acid/quinine<sup>39</sup> or the *trans*-*N,N'*-  
252 dibenzyl-diaminocyclohexane/2,3-dichlorophenylacetic acid<sup>40</sup> hybrid salt–cocrystals. Moreover,  
253 Aakeröy et al.<sup>41</sup> suggested in a structural analysis of more than 80 cocrystals and salts formed between  
254 carboxylic acids and *N*-heterocycles that the formation of unexpected hybrid salt–cocrystals could be  
255 because carboxylate moieties are not readily satisfied by a single hydrogen-bond donor making  
256 necessary the presence of neutral carboxylic acids in the crystal structure. We have examined the  
257 Cambridge Structural Database (version 5.39, 2018) in order to assess the formation of hybrid  
258 salt–cocrystal forms in multicomponent crystals containing a piperazine ring (the basic group of  
259 sildenafil) and a carboxylic acid (Figure 3).

260 A total of 247 crystal structures containing atomic coordinates were found and classified as salt,  
261 cocrystal, or hybrid salt–cocrystal according to the C–O bond lengths of the carboxylate moiety.  
262 Although 184 structures showed total proton transfer between donor and acceptor, 63 of them revealed  
263 that cocrystals or mixed salt-cocrystals were formed. This encouraged us to test the carboxylic acids  
264 previously chosen in the virtual cocrystal prediction. Table 5 summarizes the results of this structural  
265 analysis.

266 **3.2. Salt/Cocrystal Screening.** With the aim to discover new salts or cocrystals of sildenafil, an  
267 extensive multicomponent solid forms screening was conducted by using a broad set of thermodynamic

268 and kinetic experimental conditions from a variety of 54 solvents,<sup>42</sup> which produced 194 individual  
269 crystalline solids (see Supporting Information for experimental and characterization details).

270 **3.3. Crystal Structures Analysis.** The crystal structures of 5 out of the new 23 forms of sildenafil have  
271 been solved by single crystal X-ray diffraction, and the following analysis shows that in all cases salts or  
272 hybrid salt-cocrystals have been formed with tartaric acid, 3-hydroxybenzoic acid, and 3,4-  
273 dihydroxybenzoic acid.

274 3.3.1. Tartaric Acid Salt Isopropanol Hemisolvate (SIL-TAR IV). Tartaric acid salt isopropanol  
275 hemisolvate crystallizes with one molecule of sildenafil cation, half molecule of tartrate dianion, and  
276 half disordered molecule of isopropanol in the asymmetric unit. Transfer of both protons of tartaric acid  
277 has been deduced since tartrate C–O distances are 1.183(9) and 1.232(8) Å. The dianion, which shows  
278 disorder between two conformations (in a 1:1 ratio), is encapsulated between two molecules of sildenafil  
279 establishing strong charge-assisted hydrogen bonds. Sildenafil/tartrate cages are packed with a  
280 combination of electrostatic interactions between sulphonamide moieties in a self-association fashion  
281 and weak hydrogen bonds between N-methylpyrazole rings (Figures 4 and 5). Molecular cavities are  
282 present and occupied by disordered molecules of isopropanol.

283 3.3.2. Hybrid 3-Hydroxybenzoic Acid Salt–Cocrystal Monohydrate (SIL-3-HBA IV). The hybrid  
284 salt–cocrystal formed by 3-hydroxybenzoic acid and sildenafil crystallizes with one molecule of  
285 sildenafil cation, one molecule of the carboxylate, half molecule of the carboxylic acid, and one  
286 molecule of water in the asymmetric unit. Chains of selfassembled sildenafil cations are formed through  
287 strong hydrogen bonds between the piperazinium ring and the carbonylic oxygen. As expected, strong  
288 charge-assisted hydrogen bonds are formed between the carboxylate anion and the piperazinium cation,  
289 but one molecule of the nonionized carboxylic acid interacts with the carboxylate anion via the phenol  
290 and carboxylic hydrogen in an alternate manner (Figure 9). Weak antiparallel dipole–dipole interactions  
291 between stacked pyrimidinone rings are established conferring extra stabilization to the crystal (Figure  
292 8).

293 In addition, one molecule of water is also present acting as a bridge between carboxylates (Figure 6).  
294 Nonionized 3-hydroxybenzoic acid molecules are located in channels establishing strong hydrogen  
295 bonds with other 3-hydroxybenzoate molecules (Figure 7).

296 3.3.3. 3-Hydroxybenzoic Acid Salt THF Hemisolvate Sesquihydrate (SIL-3-HBA II). The salt formed  
297 by 3-hydroxybenzoic acid and sildenafil crystallizes with one molecule of sildenafil cation, one  
298 molecule of the carboxylate, half disordered molecule of THF, and 1.5 molecules of water in the  
299 asymmetric unit. In spite of the different degree of proton transfer, this solid form is isostructural to the  
300 hybrid 3-hydroxybenzoic acid saltcocrystal, and the same interactions between sildenafil and 3-  
301 hydroxybenzoate molecules are established. Moreover, identical channels are formed but filled by  
302 disordered tetrahydrofuran and water molecules instead of molecules of 3-hydroxybenzoic acid. Only  
303 small differences between both structures are present like, for instance, centroid–centroid distances  
304 measured between pyrimidinone rings and torsion angles of propyl groups (Figure 8).

305 3.3.4. Hybrid 3,4-Dihydroxybenzoic Acid Salt–Cocrystal Monohydrate (SIL-3,4-DHBA IV). The  
306 hybrid salt-cocrystal formed by 3,4-dihydroxybenzoic acid and sildenafil crystallizes with one molecule  
307 of sildenafil cations, one molecule of the carboxylate, half molecule of the carboxylic acid, and one  
308 molecule of water in the asymmetric unit. This solid form is isostructural to the hybrid 3-  
309 hydroxybenzoic acid salt–cocrystal. The presence of an extra phenol group in the 3,4-dihydroxybenzoic  
310 acid only reinforces the same packing without disrupting any of the main observed interactions in the  
311 hybrid 3-hydroxybenzoic acid salt–cocrystal. Figure 9 shows chains of carboxylate molecules linked by  
312 water molecules in both structures.

313 3.3.5. Hybrid 3,4-Dihydroxybenzoic Acid Salt–Cocrystal Acetonitrile Disolvate (SIL-3,4-DHBA II).  
314 The hybrid 3,4-dihydroxybenzoic acid salt–cocrystal acetonitrile disolvate crystallizes with one  
315 molecule of the sildenafil cation, one molecule of the carboxylate, one molecule of the carboxylic acid,  
316 and two molecules of acetonitrile in the asymmetric unit. In a similar way that the tartaric acid salt,  
317 instead of catemeric chains of sildenafil cations, self-assembled dimers are formed through charge-  
318 assisted hydrogen bonds (Figure 10).

319 However, the antiparallel dipole–dipole interactions between stacked pyrimidinone rings are not  
320 observed in this form. This is the only structure of this family of hybrid salts–cocrystals where water is  
321 not present, and this produces a different architecture of the cofomer self-assembling, which consists of  
322 layers of alternate carboxylic/carboxylate interactions (Figure 11).

323 **3.4. Dissolution Study.** The dissolution studies were carried out at pH 1.2, pH 6.5, and FaSSIF (pH  
324 6.5), which represent the average pH values of the fast state stomach and intestine, respectively. SIL has  
325 pH dependent solubility which decreases with an increase in pH. One of the major challenges in the  
326 dissolution study of multicomponent entities is continuous change in the solution composition due to  
327 precipitation of either of the component over the dissolution testing period. The solubility data generated  
328 may be erroneous due to limitations of the analytical method; for example, estimations carried out by  
329 UV spectrophotometry are subject to the overlap in the absorption spectra of the two components. We  
330 have used the HPLC method to quantify the amount of SIL dissolved; hence we see some difference in  
331 reported dissolutions compared to the previous SIL salt dissolution data reported.<sup>43</sup> Figures 12, 13, and  
332 14 (and Figures S75–S77 of the Supporting Information) show the dissolution data.

333 At pH 1.2 the amount dissolved from SIL salts was significantly higher than the cocrystals and the  
334 hybrid salt–cocrystal forms. The hybrid salt–cocrystals showed poor dissolution performance compared  
335 to cocrystals. The D1h and D24h values for SIL-TAR I were higher than for SIL-CIT, a commercially  
336 used salt of SIL. At pH 1.2 in the salt category SILTAR > SIL-3-HBA I > SIL-CIT > SIL-3,4-DHBA I.  
337 On the other hand D1h for SIL-RES II > SIL-QUE I and SIL-QUE II. As expected, the amount  
338 dissolved at pH 6.5 was at least 10 times lower than the amount dissolved at pH 1.2. But in both  
339 conditions, the amount of SIL dissolved was significantly higher for salts than for cocrystals and hybrid  
340 salt–cocrystals. Most of the cocrystals and hybrid salt–cocrystals did not provide any release of SIL at  
341 pH 6.5 or even in FaSSIF, which contains an additive like lecithin included in the dissolution media.

342 Thus, although an exhaustive solubility study of the new forms was not the main objective of this  
343 research, our data suggest that SILTAR could be a potential alternative (in terms of bioavailability) to  
344 the commercial citrate salt of sildenafil.

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350 **4. CONCLUSION**

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352 In summary, we have revisited the multicomponent solid form landscape of sildenafil by conducting a  
353 combined virtual and experimental screening. Twenty-three new solid forms have been discovered and  
354 characterized, and dissolution data have been measured for some of the solid forms suggesting the new  
355 tartrate salt as a potential alternative to the marketed citrate salt. The analysis of the five crystal  
356 structures solved by SXRD showed a variety of salts and hybrid salt–cocrystals with different hydrogen  
357 bond architectures and presence of solvent channels. This study extends the knowledge about the solid  
358 state of this important drug compound, contributes with new cases to the body of data of unexpected  
359 stoichiometric hybrid salt–cocrystals, and it is a new example of successful application of combined  
360 virtual/experimental methodologies for the discovery of new solid forms.

361

362

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371 **Notes**

372 The authors declare no competing financial interest.

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492 1-propanol, 1-pentanol, 1-octanol, 2,2,2-trifluoroethanol, benzyl alcohol, ACN, propionitrile,  
493 MEK, acetone, MiBK, water, DMF, DMSO, pentane, heptane, cyclohexane, hexane,  
494 methylcyclohexane, toluene, xylene, mesitylene, anisole, 2-nitrotoluene, nitrobenzene, AcOEt,

495 isopropyl acetate, diethylether, THF, 1-methyl-2-pyrrolidone, dimethyl ethylene glycol,  
496 diisopropyl ether, dioxane, iodomethane, dichloromethane, 1,2-dichloroethane, chloroform, 1,1,1-  
497 trichloroethane, 1,1,2-trichloroethane, formic acid, acetic acid, trifluoroacetic acid, propanoic  
498 acid, NH<sub>3</sub> (32%) in water, diethylamine, trimethylamine and pyridine.

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502

503 **Legends to figures**

504

505 **Figure. 1.** Molecular structure of sildenafil.

506

507 **Figure. 2** SSIPs calculated for sildenafil. Blue spheres correspond to Hbond donors and red spheres to  
508 H-bond acceptors.

509

510 **Figure. 3** Fragments searched in multicomponent crystals in the CSD.

511

512 **Figure. 4** Crystal structure of tartaric acid salt of sildenafil. The most relevant interactions are  
513 highlighted. Channels filled by solvent molecules are highlighted with gray circles.

514

515 **Figure. 5** Sildenafil/tartrate cage in the crystal structure of tartaric acid salt of sildenafil.

516

517 **Figure. 6** Chains of carboxylate/carboxylic acid molecules linked by molecules of water in the hybrid 3-  
518 hydroxybenzoic salt-cocrystal monohydrate.

519

520 **Figure. 7** Representation of the crystal structures of 3-hydroxybenzoic acid salt and hybrid  
521 salt-cocrystal of sildenafil. The most relevant interactions have been highlighted, and hydrogens have  
522 been partially omitted for clarity. Channels filled by THF and water molecules in the salt or 3-  
523 hydroxybenzoic molecules in the hybrid salt-cocrystal are highlighted with gray circles.

524

525 **Figure. 8** Antiparallel dipole-dipole interactions established between stacked pyrimidinone rings in the  
526 crystal structures of 3-hydroxybenzoic acid salt (right) and hybrid salt-cocrystal (left). Differences in  
527 centroid-centroid distances measured between pyrimidinone rings and torsion angles of propyl groups  
528 are shown for each structure.

529

530 **Figure. 9** Chains of carboxylate molecules linked by molecules of water in the hybrid 3-hydroxybenzoic  
531 acid salt-cocrystal (left) and in the hybrid 3,4-dihydroxybenzoic acid salt-cocrystal (right).

532

533 **Figure. 10** Crystal structure of 3,4-dihydroxybenzoic acid hybrid salt-cocrystal. Self-assembled dimers  
534 formed through charge-assisted hydrogen bonds are highlighted. Hydrogens have been partially omitted  
535 for clarity.

536

537 **Figure. 11** Layers of alternate carboxylic (blue)/carboxylate (red) interactions.

538

539 **Figure. 12** Comparative solubility of SIL salts, cocrystals, and hybrid salt–cocrystals in 0.1 N HCl (pH  
540 1.2).

541

542

543 **Figure. 13** Comparative solubility of SIL salts, cocrystals, and hybrid salt–cocrystals in phosphate  
544 buffer pH 6.5.

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546

547 **Figure. 14** Comparative solubility of SIL salts, cocrystals, and hybrid salt–cocrystals in FaSSIF.

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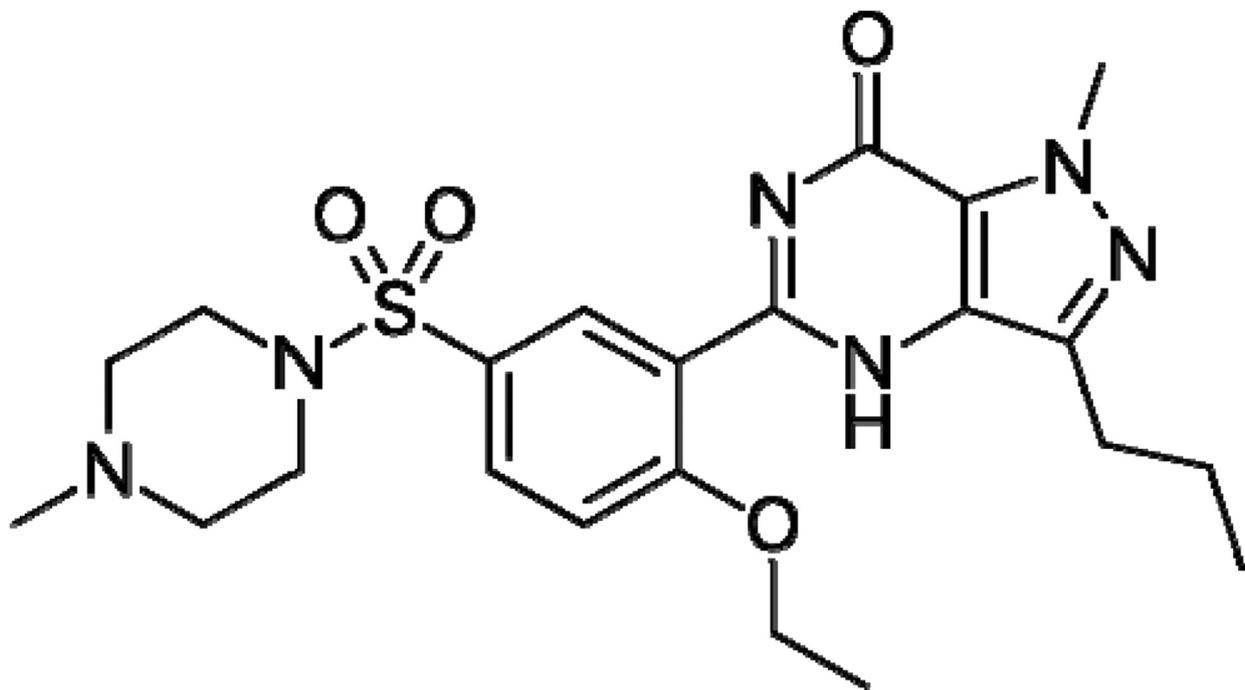
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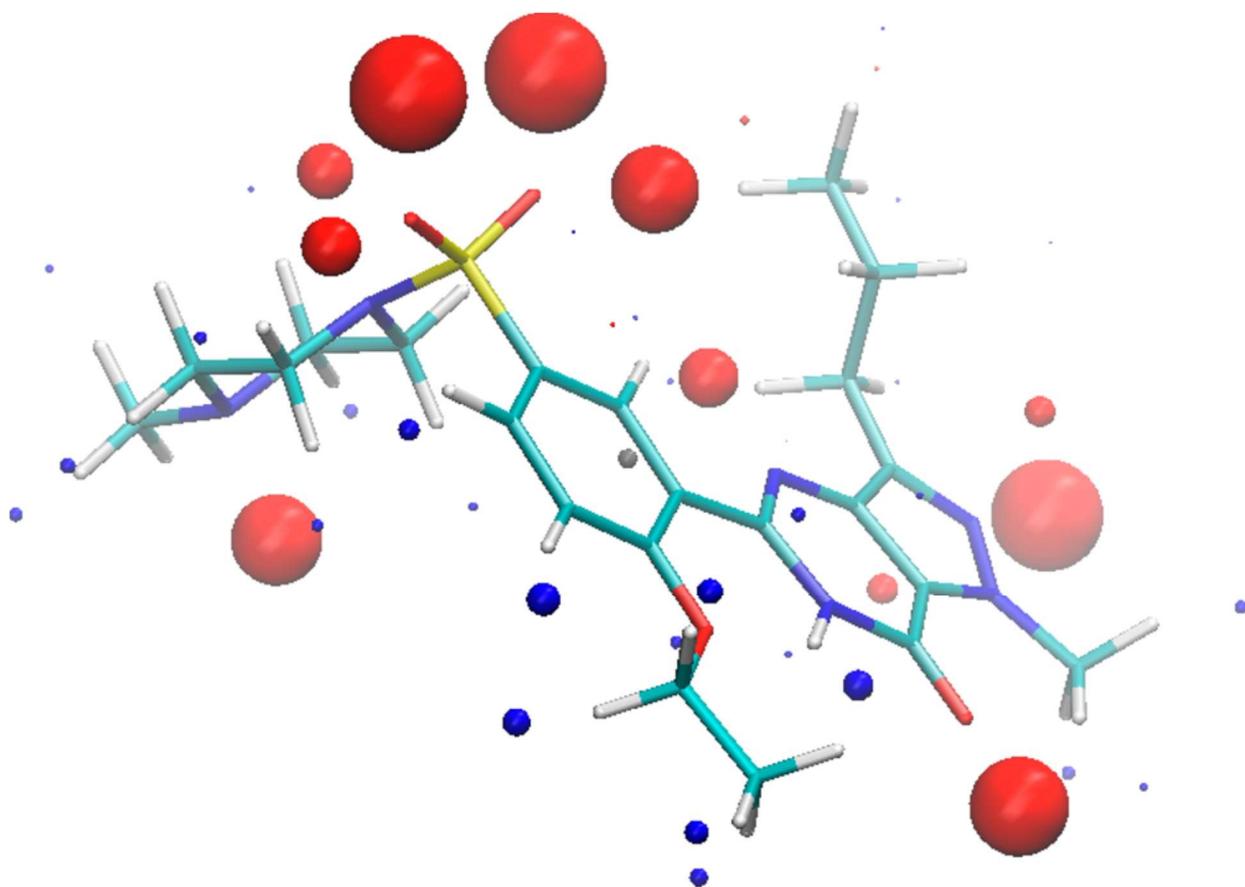
FIGURE 1



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FIGURE 2



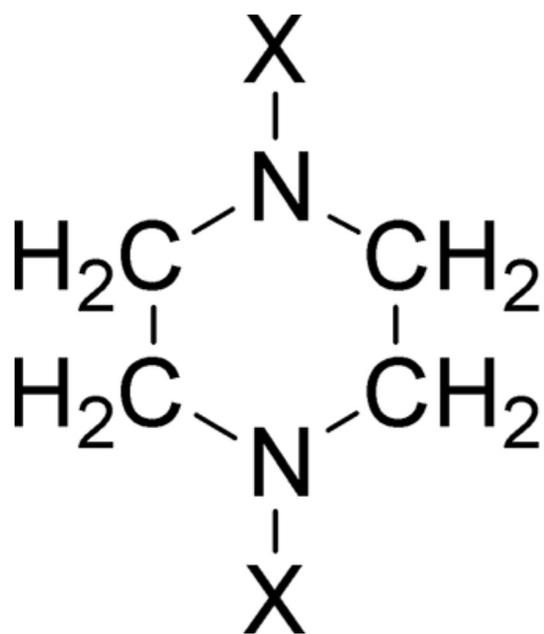
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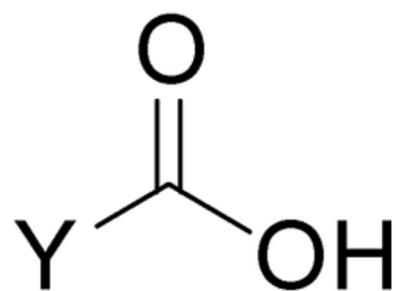
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FIGURE 3



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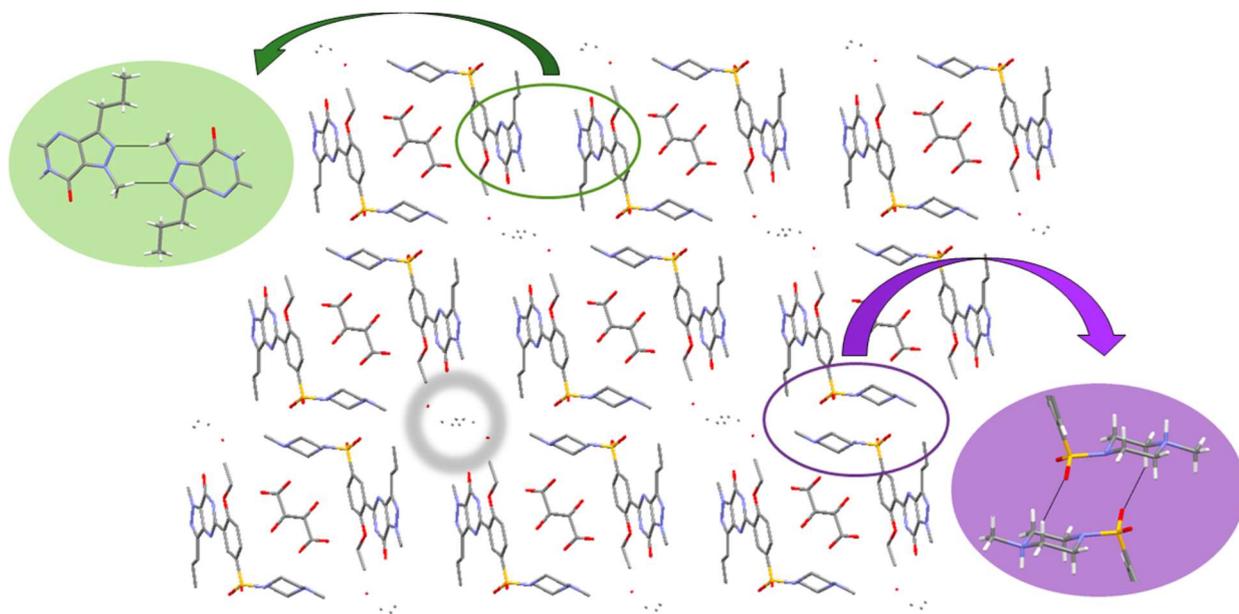


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FIGURE 4

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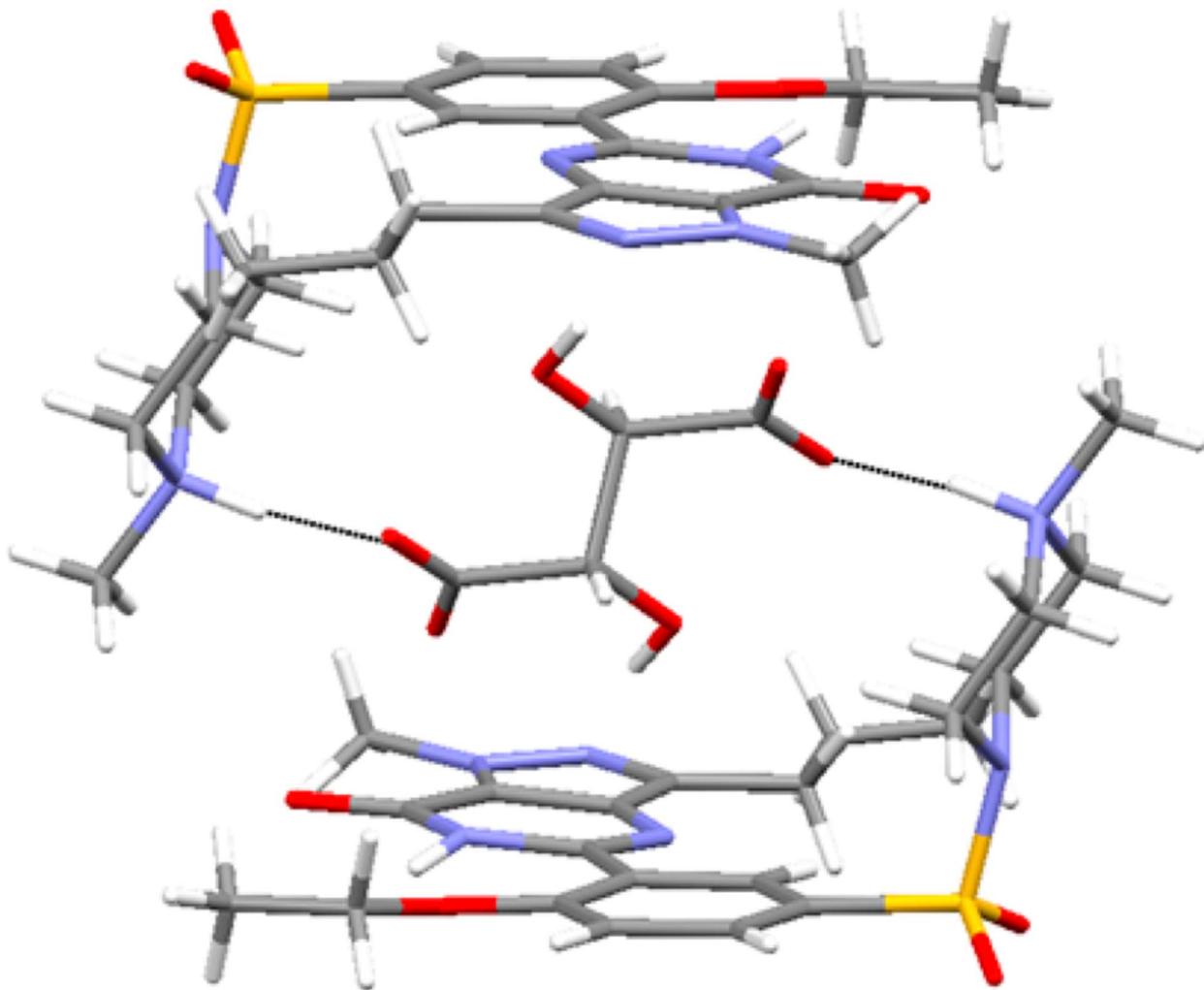
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FIGURE 5.

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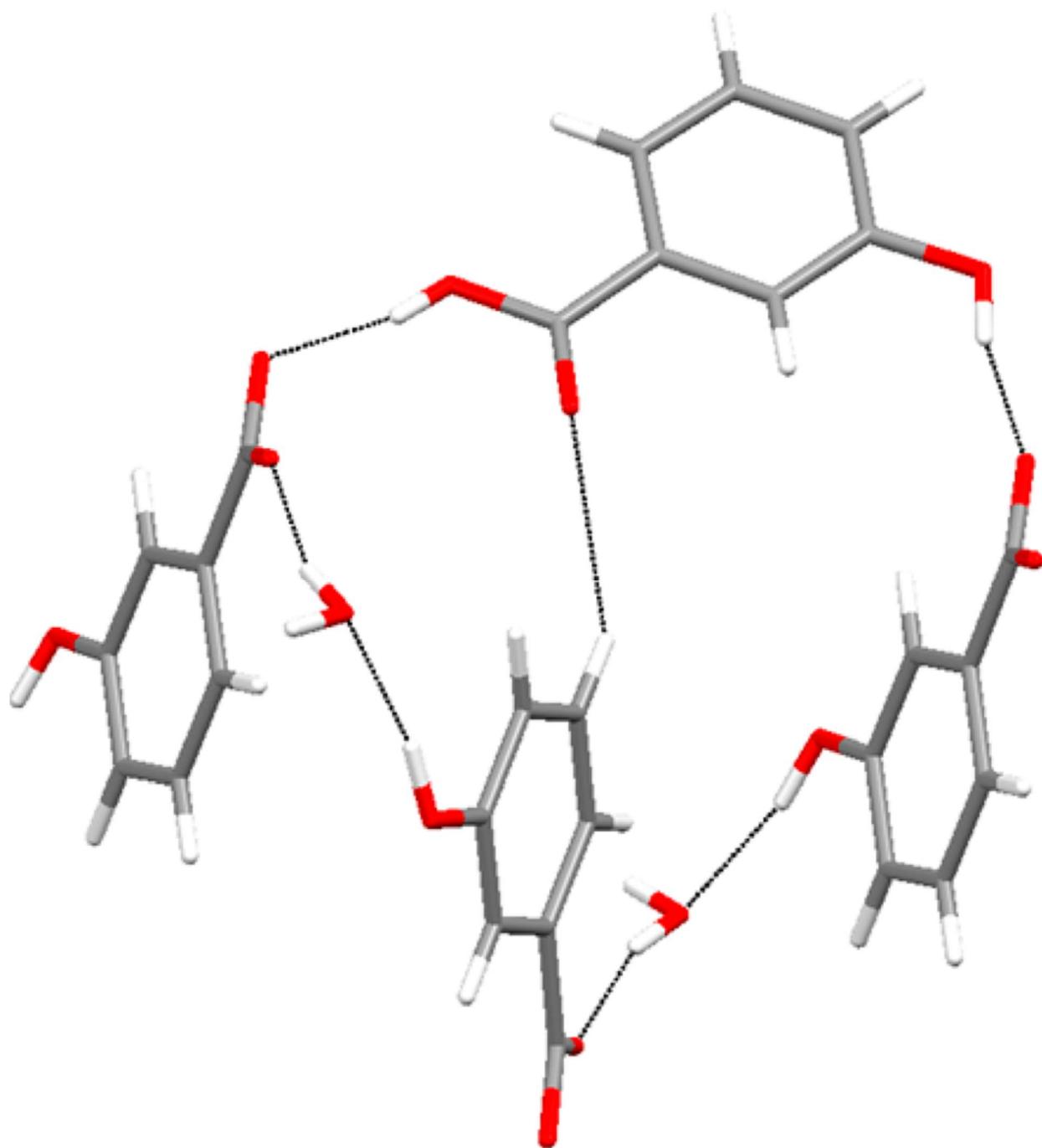
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FIGURE 6.

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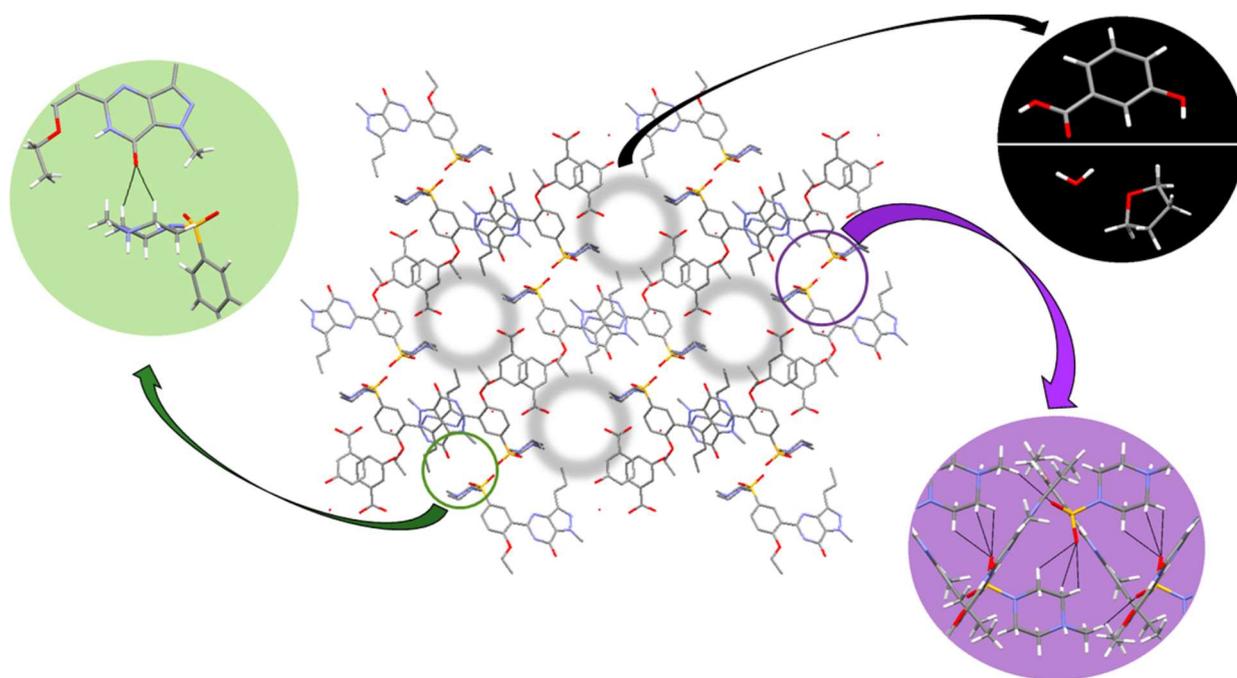
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FIGURE 7.

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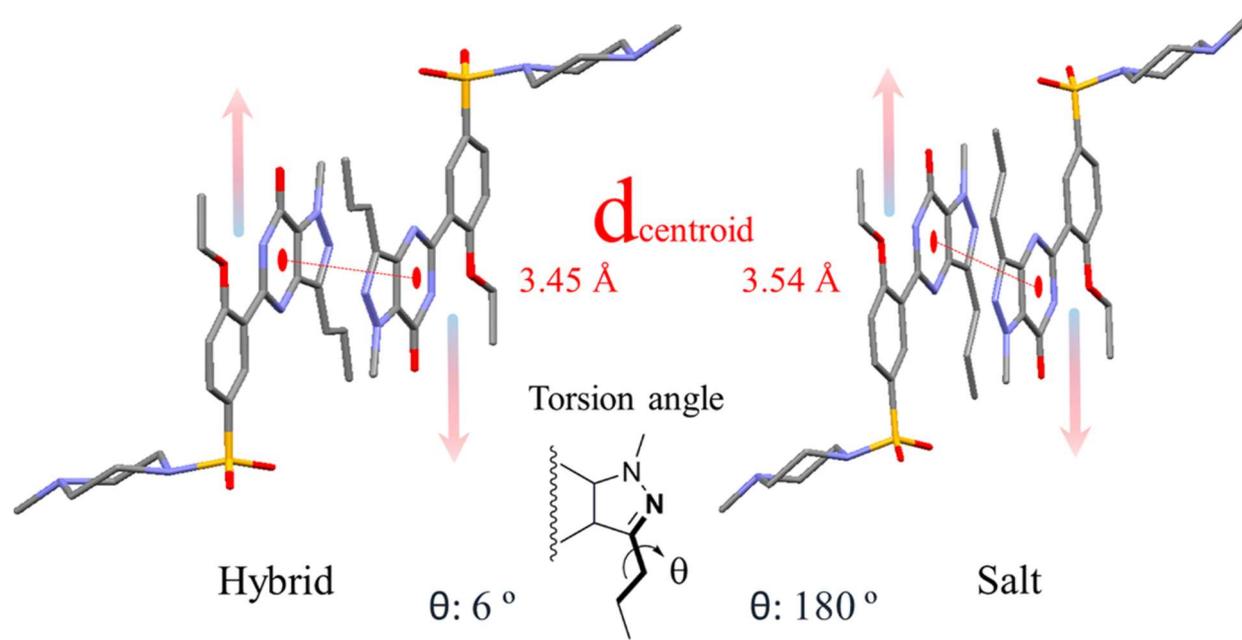
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FIGURE 8.

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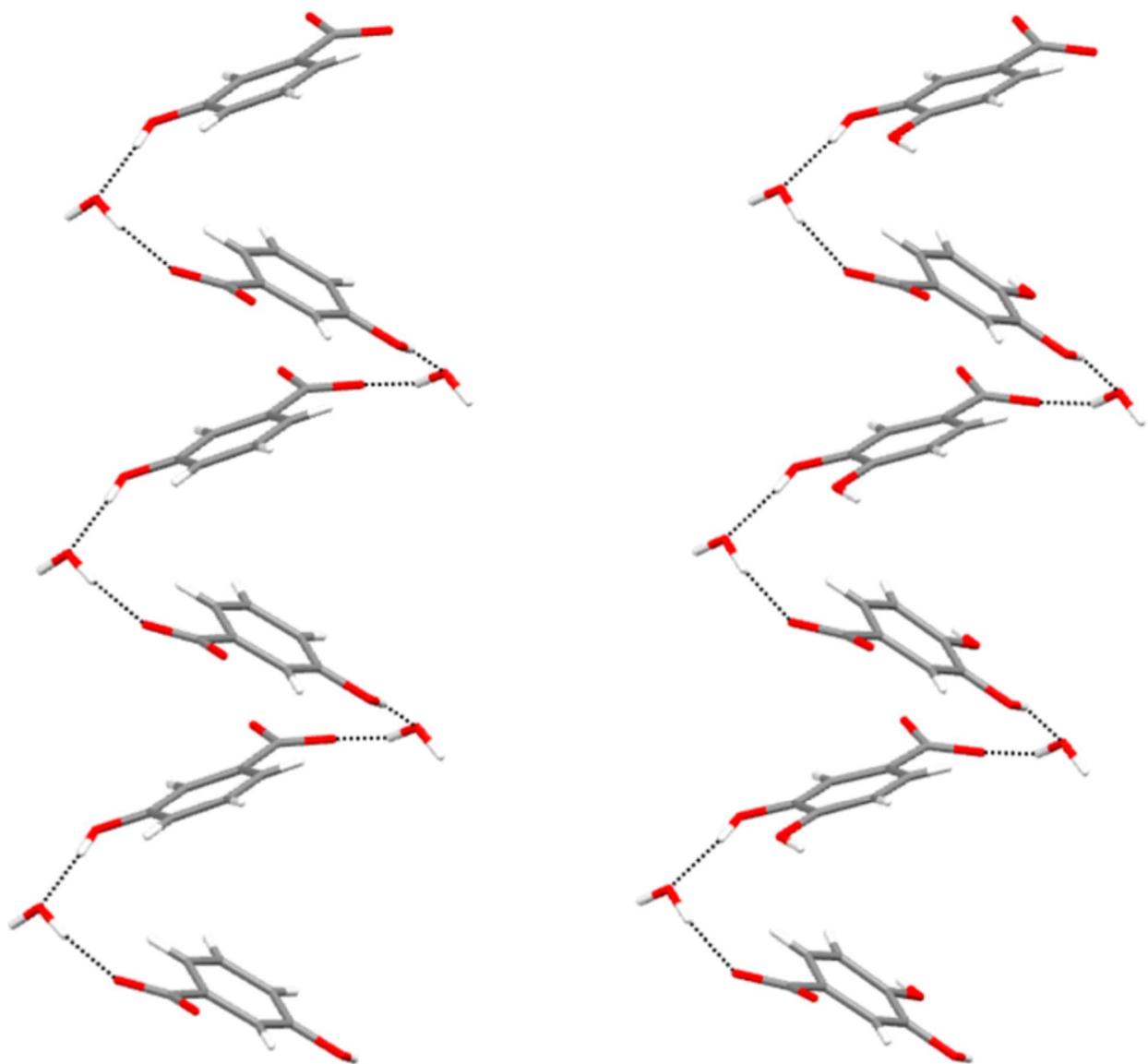
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FIGURE 9.

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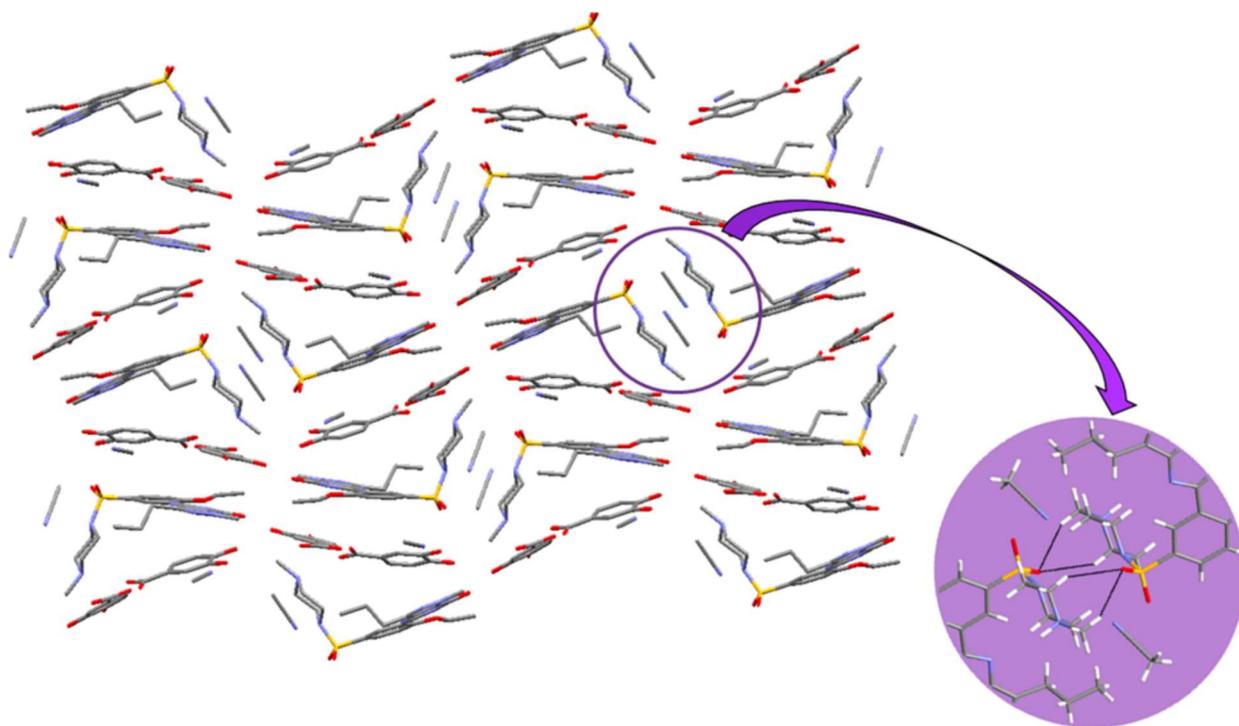
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FIGURE 10.

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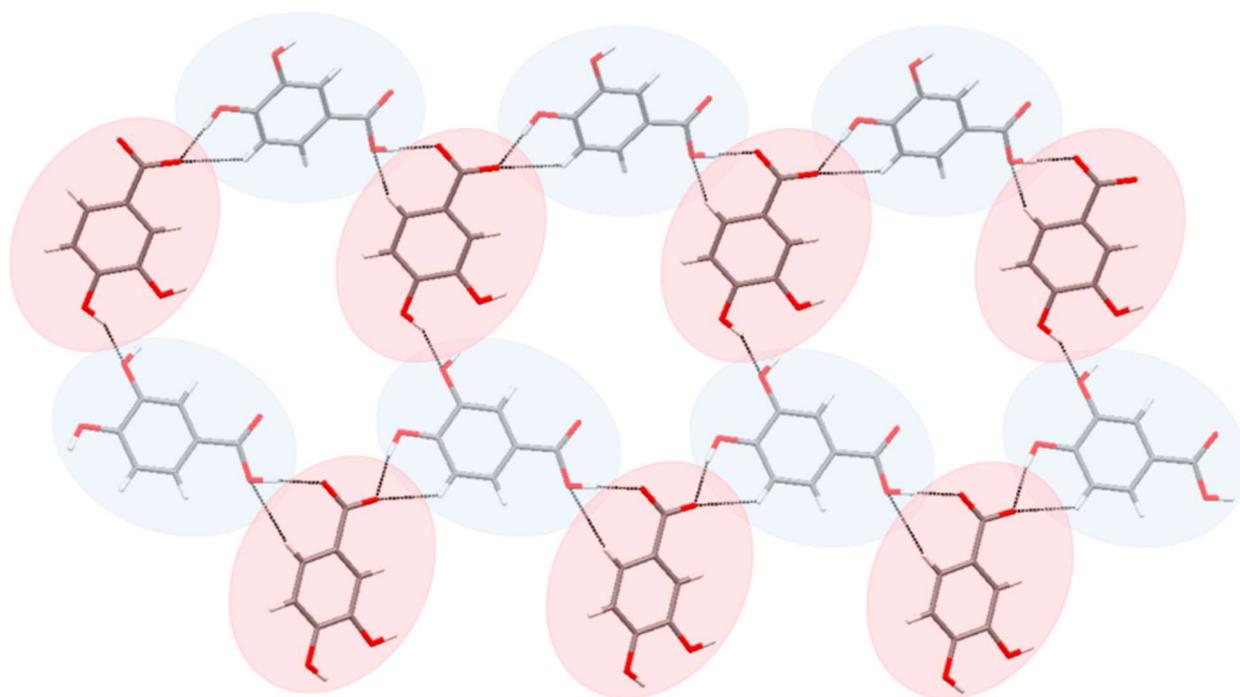
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**FIGURE 11.**

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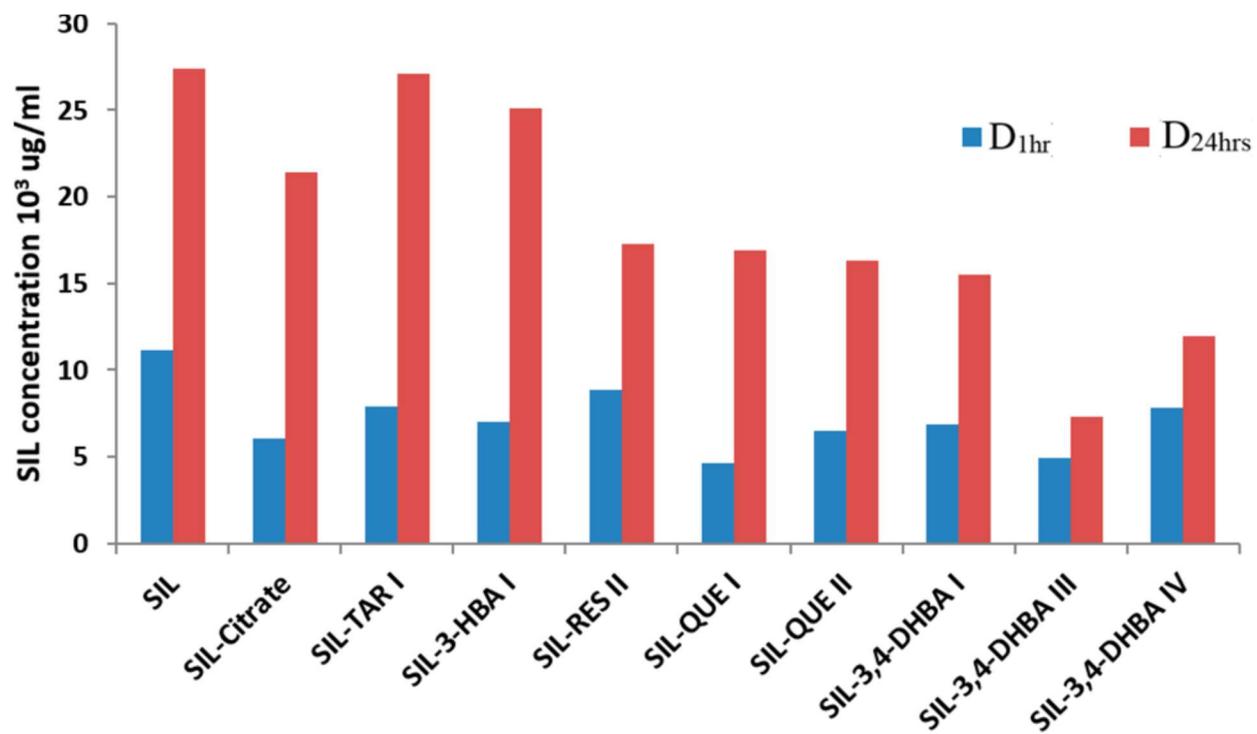
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FIGURE 12.

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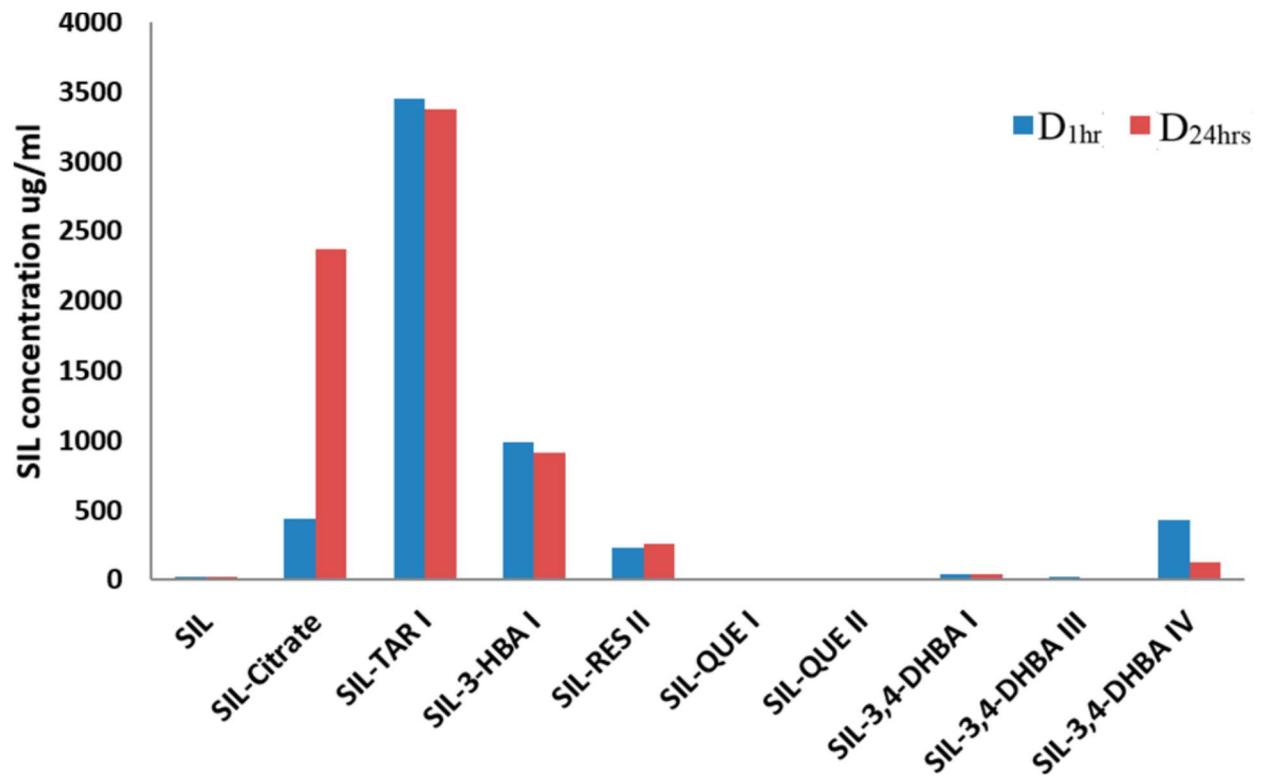
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FIGURE 13.

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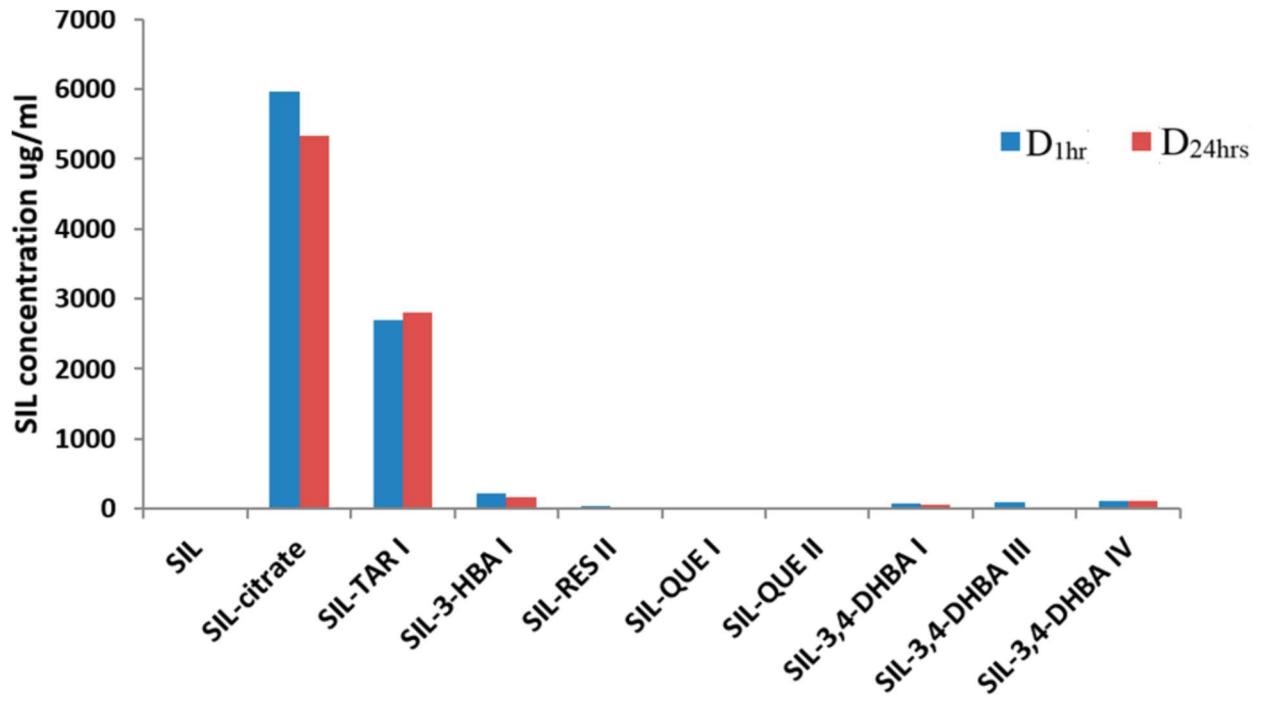
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FIGURE 14

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structure	SIL-TAR IV	SIL-3A-DHBA II	SIL-3A-DHBA IV	SIL-3-HBA II	SIL-3-HBA IV
empirical formula	C <sub>21</sub> H <sub>26</sub> N <sub>12</sub> O <sub>15</sub> S <sub>2</sub>	C <sub>40</sub> H <sub>48</sub> N <sub>8</sub> O <sub>12</sub> S	C <sub>40</sub> H <sub>42</sub> N <sub>12</sub> O <sub>12</sub> S <sub>2</sub>	C <sub>31</sub> H <sub>43</sub> N <sub>6</sub> O <sub>9</sub> S	C <sub>40</sub> H <sub>42</sub> N <sub>12</sub> O <sub>19</sub> S <sub>2</sub>
formula weight	1159.34	864.92	1447.54	675.77	1399.54
temperature (K)	293(2)	100(2)	100(2)	100(2)	100(2)
crystal system	triclinic	monoclinic	monoclinic	monoclinic	monoclinic
space group	$\bar{P}1$	$P2_1/n$	$P2_1/c$	$P2_1/n$	$P2_1/n$
<i>a</i> , <i>b</i> , <i>c</i> (Å)	6.3796(8) 13.4449(18) 17.620(2)	11.6418(8) 15.3849(13) 24.7913(19)	17.6070(17) 8.6938(8) 24.942(2)	17.7205(18) 8.2574(7) 22.991(2)	17.5329(10) 8.6545(4) 22.9653(13)
$\alpha$ , $\beta$ , $\gamma$ (deg)	108.607(5) 98.363(6) 92.814(6)	90 102.749(3) 90	90 117.101(5) 90	90 102.748(3) 90	90 105.591(2) 90
volume (Å <sup>3</sup> )	1409.7(3)	4330.8(6)	3398.7(5)	3281.2(5)	3356.5(3)
Z, density (calc.) (Mg/m <sup>3</sup> )	1, 1.366	4, 1.327	2, 1.414	4, 1.368	2, 1.385
crystal size (mm <sup>3</sup> )	0.344 × 0.144 × 0.044	0.358 × 0.196 × 0.074	0.324 × 0.152 × 0.112	0.129 × 0.119 × 0.109	0.336 × 0.197 × 0.086
reflections collected/unique	53107/6455 [R(int) = 0.1123]	66326/8875 [R(int) = 0.0949]	69444/7838 [R(int) = 0.0722]	18923/4567 [R(int) = 0.082]	33777/5726 [R(int) = 0.0482]
data/restraints/parameters	6455/8/396	8875/0/570	7838/2/460	4567/6/454	5726/0/502
goodness-of-fit on F <sup>2</sup>	1.105	1.070	1.024	1.017	1.038
final R indices [I > 2σ(I)]	R <sub>1</sub> = 0.0678 wR <sub>2</sub> = 0.1740	R <sub>1</sub> = 0.0537 wR <sub>2</sub> = 0.1151	R <sub>1</sub> = 0.0476 wR <sub>2</sub> = 0.1084	R <sub>1</sub> = 0.0504 wR <sub>2</sub> = 0.1168	R <sub>1</sub> = 0.0382 wR <sub>2</sub> = 0.0886
CCDC	1858573	1858576	1858577	1858574	1858575

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Table 2. Comparative Cell Parameters Data<sup>a</sup> from SXRD and PXRD

crystal form	a (Å)	b (Å)	c (Å)	$\alpha$ (deg)	$\beta$ (deg)	$\gamma$ (deg)	V (Å <sup>3</sup> )	Z	R (%)	space group
SIL-QUE I	PXRD 42.39(2)	14.854(4)	15.059(6)	82.41(3)	138.08(2)	120.01(2)	4992(3)	5	11.1	P $\bar{1}$
SIL-QUE II	PXRD 22.528(8)	13.536(4)	8.007(2)	57.95(2)	95.99(3)	95.96(2)	2054(1)	2	6.69	P1
SIL-3,4-DHBA I	PXRD 12.765(2)	13.463(1)	12.106(2)	112.03(1)	84.0929(9)	114.28(1)	1754.7(3)	2	8.41	P $\bar{1}$
SIL-3,4-DHBA II	SXRD 11.6418(8)	15.3849(13)	24.7913(19)	90	102.749(3)	90	4330.8(6)	4	5.37	P21/m
SIL-3,4-DHBA III	PXRD 24.947(6)	15.718(3)	11.642(3)	90	103.96(2)	90	4430(2)	4	7.76	P21/m
SIL-3,4-DHBA IV	SXRD 17.6070(17)	8.6938(8)	24.942(2)	90	117.101(5)	90	3398.7(5)	2	4.76	P21/c
SIL-RES I	PXRD 11.266(2)	14.835(2)	14.178(2)	38.099(7)	94.79(1)	96.09(9)	1453.9(3)	2	12.0	P $\bar{1}$
SIL-RES II	PXRD 14.2705(4)	26.083(2)	9.9925(5)	90	108.398(3)	90	3530.0(3)	4	6.20	P21/a
SIL-TAR I	PXRD 18.07(1)	13.609(6)	7.590(4)	85.98(3)	92.33(3)	110.09(4)	1749(1)	2	7.31	P $\bar{1}$
SIL-TAR II	PXRD 33.57(9)	15.070(3)	11.547(3)	90	90	90	5445(2)	4	8.11	P21 21 21
SIL-TAR III	PXRD 18.313(4)	15.070(3)	6.3764(9)	63.85(1)	101.56(1)	112.86(2)	1454.5(5)	2	7.64	P1
SIL-TAR IV	SXRD 63796(8)	13.4449(18)	17.620(2)	108.607(5)	98.263(6)	92.814(6)	1409.7(3)	1	6.78	P $\bar{1}$
SIL-CAF I	PXRD 25.845(7)	8.236(1)	20.32(6)	90	121.19(1)	90	3744(2)	2	5.76	P21/m
SIL-MEG	PXRD 12.9869(8)	13.4289(8)	13.4551(8)	53.033(3)	118.580(4)	111.246(4)	1637.1(1)	2	5.04	P $\bar{1}$
SIL-3-HBA II	SXRD 17.7205(18)	8.2574(7)	22.991(2)	90	102.748(3)	90	3281.2(5)	4	5.04	P21/m
SIL-3-HBA IV	SXRD 17.5329(10)	8.6545(4)	22.9653(13)	90	105.591(2)	90	3556.5(3)	2	3.82	P21/m

<sup>a</sup>R-factor for SXRD and R<sub>wp</sub> for PXRD.

635 **Table 3** Coformers Chosen in This Study Based on the Difference between the Interaction Site Pairing  
636 Energies of Sildenafil and the Pure Components,  $\Delta E$

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coformer	$\Delta E/\text{kJ mol}^{-1}$
quercetin	28.1
resveratrol	24.0
phloroglucinol	23.8
3,4-dihydroxybenzoic acid	21.7
resorcinol	19.3
tartaric acid	18.2
caffeic acid	17.8
myo-inositol	15.7
tert-butylhydroquinone	13.8
methyl galate	13.5
3-hydroxybenzoic acid	13.1
4-hydroxybenzoic acid	12.9

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640 **Table 4** Cocrystal Screening Coformers pKa's and Estimated Probability of Proton Transfe

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coformer	reported pK <sub>a</sub>	ΔpK <sub>a</sub>	P (%)
quercetin	8.45 <sup>35</sup>	-1.67	0
resveratrol	8.49 <sup>36</sup>	-1.71	0
phloroglucinol	7.97 <sup>37</sup>	-1.19	8
3,4-dihydroxybenzoic acid	4.40 <sup>37</sup>	2.38	68
resorcinol	9.44 <sup>37</sup>	-2.66	0
tartaric acid	3.03 <sup>37</sup>	3.75	92
caffeic acid	4.47 <sup>36</sup>	2.31	67
myo-inositol	12.29 <sup>36</sup>	-5.51	0
t-butylhydroquinone	9.94 <sup>36</sup>	-3.16	0
methyl gallate	8.11 <sup>35</sup>	-1.33	5
3-hydroxybenzoic acid	4.08 <sup>38</sup>	2.7	74
4-hydroxybenzoic acid	4.57 <sup>38</sup>	2.21	66

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644 **Table 5** Classification of Multicomponent Crystals with a Piperazine and a Carboxylic Group in the  
645 CSD

class	no. of structures	%
salt	184	75
cocrystal	38	15
hybrid	25	10

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