

Controlling Solvate Formation of a Schiff Base by Combining Mechanochemistry with Solution Synthesis[†]

Vladimir Stilinović, Dominik Cinčić, Marija Zbačnik, and Branko Kaitner*

Department of Chemistry, Laboratory of General and Inorganic Chemistry, Faculty of Science, Horvatovac 102a,
University of Zagreb, 10000 Zagreb, Croatia

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Abstract. A Schiff base dicarboxylic acid (**1**) was prepared by condensation of 2-hydroxy-1-naphthaldehyde with 5-aminoisophthalic acid. Its solvates with pyridine (**2a** and **2b**) and dimethylformamide (**3**) were prepared by liquid-assisted grinding and by conventional solvent-based methods. All products were characterised by FT-IR spectroscopy, thermogravimetric analysis and differential scanning calorimetry. The structures of **1**, **2b** and **3** were determined by single crystal X-ray diffraction. **1** was found to be a pure Schiff base, **2b** a pyridine solvate and **3** a dimethylformamide solvate monohydrate. In all three structures, the Schiff base molecule appears to be present as the ketoamine tautomer.(doi: [10.5562/cca2111](http://dx.doi.org/10.5562/cca2111))

Keywords: Schiff base, solvates, dicarboxylic acid, mechanochemistry, tautomerism

INTRODUCTION

Schiff bases are imines prepared by condensation of primary amines with carbonyl compounds.¹ During the last five decades a vast number of structural studies on Schiff bases derived from hydroxyaryl aldehydes and their coordination compounds have been undertaken.^{2,3} Schiff bases have become some of the most typical ligands for coordinating transition and inner transition metal ions.^{4–7} This most commonly involves Schiff base ligands derived from salicylaldehyde and 2-hydroxy-1-naphthaldehyde.^{8–12} Also, Schiff bases have been intensively used as synthetic intermediates¹³ and, more recently, as anion receptors.^{14–16} They have attracted particular interest due to their biological activity^{17,18} and as model systems for biological macromolecules.^{19,20} Solid-state thermochromism and photochromism are another characteristic of these compounds leading to their application in various areas of materials science.^{21–23} Schiff bases derived from *o*-hydroxyaromatic aldehydes and ketones have shown to be excellent models for the study of keto-enol tautomerism both in solution and in the solid state.^{24–28} Such Schiff bases have drawn attention due to their physical properties in the crystalline state which are greatly influenced by the topochemistry of the Schiff base molecules which in turn is highly affected by the crystal structure.^{29–33}

During the past decade it became well-recognised that solvent-free synthesis and mechanochemistry are effective for the efficient and rapid synthesis of a wide range of imines.^{34,35} In general, conventional synthesis by condensation of primary amines with carbonyl compounds in solution requires acid catalysis, removal of the generated water and removal of bulk solvent upon the isolation of the product. It is therefore of high interest that many of these reactions can be performed in a single step with quantitative yields by gas-solid, liquid-solid or solid-solid reactions.^{36–39} This approach circumvents the use of the bulk solvent, avoiding at the same time problems encountered in solution-based chemistry, such as low solubility of reactants and solvolysis or hydrolysis of products. Among these preparatory procedures, mechanochemical methods such as liquid-assisted grinding (LAG)^{40,41} or ion- and liquid-assisted grinding (ILAG)^{42,43} have shown a great potential as efficient methods for supramolecular and covalent synthesis of new materials.^{44–47}

Our group has recently reported mechanochemical synthesis of a Schiff base derived from 2-hydroxy-1-naphthaldehyde and 2-aminobenzonitrile⁴⁸ and described simultaneous covalent solvent-free synthesis and polymorphism control using neat grinding and seeding-assisted grinding (SEAG), *i.e.* neat grinding in the presence of seed crystals of the desired product.

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* Author to whom correspondence should be addressed. (E-mail: kaitner@chem.pmf.hr)

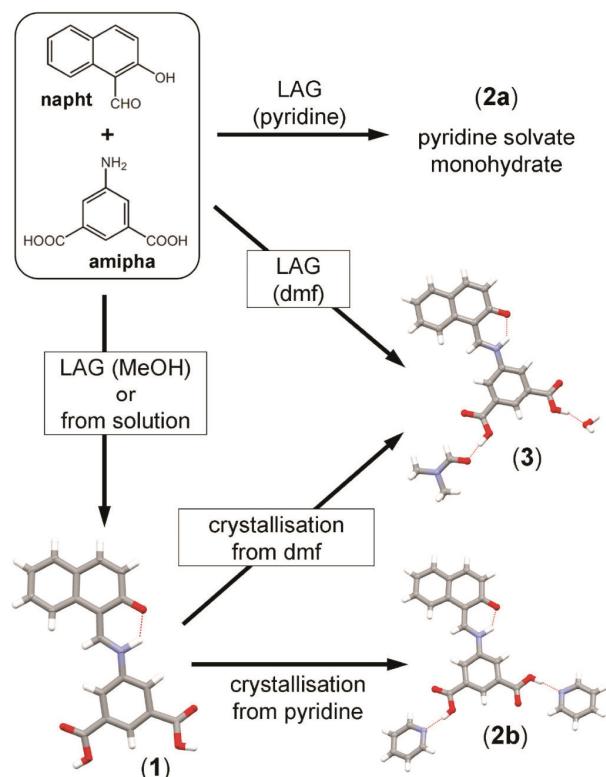


Figure 1. Summary of reactions to form Schiff base (**1**) and its solvates.

Herein we report the synthesis of 5-[(2-oxo-2*H*-naphthalen-1-ylidenemethyl)-amino]-bezene-1,3-dicarboxylic acid (**1**), Schiff base derived from 2-hydroxy-1-naphthaldehyde (**napht**) and 5-amino-isophthalic acid (**amiphtha**) as well as its solvates with pyridine (**2a** and **2b**) and dimethylformamide (**3**) (Figure 1). All products were characterised by FT-IR spectroscopy, thermogravimetric analysis (TGA), and differential scanning calorimetry (DSC). The structures of **1**, **2b** and **3** were determined by single crystal X-ray diffraction. To the best of our knowledge, this is the first structurally characterised Schiff base with two carboxylic groups bound to an aromatic ring of the *N*-substituent.²

EXPERIMENTAL

Materials and Synthesis

The starting materials, 2-hydroxy-1-naphthaldehyde and 5-aminoisophthalic acid were obtained from Acros Organics and Merck, respectively. **Napht** was recrystallised from methanol and **amiphtha** was used without further purification. Solvents were purchased from Kemika and T.T.T., Zagreb.

Compound **1** was synthesised by liquid-assisted grinding of a mixture of **napht** (0.103 g, 0.600 mmol) and **amiphtha** (0.109 g, 0.600 mmol) in the presence of 45

μL of methanol. The compound **2a** was synthesised by liquid-assisted grinding of a mixture of **napht** (0.103 g, 0.600 mmol) and **amiphtha** (0.109 g, 0.600 mmol) in the presence of 90 μL of pyridine. The compound **3** was synthesised by liquid-assisted grinding of a mixture of **napht** (0.103 g, 0.600 mmol) and **amiphtha** (0.109 g, 0.600 mmol) in the presence of 60 μL of dimethylformamide. Grinding was performed in a stainless steel jar of 10 mL volume, using two stainless steel grinding balls, 7 mm in diameter. A Retsch MM200 grinder mill operating at 25 Hz frequency was used for the synthesis. Analysis of the products using PXRD revealed complete conversion of reactants into **1**, **2a** and **3** after 50 min, 40 min and 20 min grinding, respectively.

For solution-based synthesis of **1**, **napht** (1.72 g, 0.010 mol) and **amiphtha** (1.81 g, 0.010 mol), were dissolved in hot methanol (30 mL and 20 mL, respectively). The solutions were mixed and the resulting mixture left at room temperature. Yellow precipitate began appearing almost immediately. After 30 min it was separated from the mother liquor by filtration, and washed with methanol. Single crystals of **1** suitable for X-ray diffraction were obtained by the liquid phase diffusion procedure from dichloromethane solutions of the reactants (**napht** and **amiphtha**), which yielded orange crystals. **2b** and **3** were prepared by crystallisation of **1** from pyridine and dimethylformamide, respectively. Single crystals of **2b** suitable for X-ray diffraction were obtained by dissolution of 100 mg of **1** in 3.0 mL of pyridine. The solution was allowed to evaporate slowly for 1 day at room temperature, yielding orange crystals. Single crystals of **3** suitable for X-ray diffraction were obtained by dissolution of 100 mg of **1** in 2.0 mL of dimethylformamide. The solution was allowed to evaporate slowly for 5 days at room temperature, yielding orange crystals.

Thermal and Spectroscopic Analysis

Thermal analysis was carried out on Mettler Toledo TGA/SDTA 851 and DSC823 modules in sealed aluminium pans (40 μL), heated in flowing nitrogen (200 mL min⁻¹) at 10 °C min⁻¹. The data collection and analysis was performed using the program package STARe Software 9.01.⁴⁹ Infrared spectra were recorded on an EQUINOX 55 FTIR spectrophotometer using the KBr pellet method. The data collection and analysis was performed using the program package OPUS 4.0.⁵⁰

X-ray Diffraction Experiments

Crystal and molecular structures of **1**, **2b** and **3** were determined by single crystal X-ray diffraction. Diffraction measurements were made on an Oxford Diffraction Xcalibur Kappa CCD X-ray diffractometer with graphite-monochromated MoKα ($\lambda = 0.71073 \text{ \AA}$) radiation.⁵¹

Table 1. Crystal data and summary of experimental details for compounds **1**, **2b** and **3**.

	1	2b	3
Molecular formula	C ₁₉ H ₁₃ NO ₅	C ₁₉ H ₁₃ NO ₅ ·2C ₅ H ₅ N	C ₁₉ H ₁₃ NO ₅ ·C ₃ H ₇ NO·H ₂ O
<i>M</i> _r	335.3	493.5	426.42
Crystal system	Triclinic	Monoclinic	Monoclinic
Space group	<i>P</i> $\overline{1}$	<i>P</i> 2 ₁	<i>P</i> 2 ₁ /c
Crystal data:			
<i>a</i> / Å	7.1802(6)	6.1453(4)	6.0236(2)
<i>b</i> / Å	8.3777(9)	25.5518(19)	22.0929(11)
<i>c</i> / Å	12.6430(12)	7.6712(6)	14.9471(7)
α / °	100.073(8)	90	90
β / °	106.148(7)	93.348(7)	91.365(4)
γ / °	90.593(8)	90	90
<i>V</i> / Å ³	717.85(12)	1202.50(15)	1988.58(15)
<i>Z</i>	2	2	4
ρ_{calc} /g cm ⁻³	1.551	1.363	1.424
$\lambda(\text{MoK}_\alpha)$ / Å, graphite monochromator	0.71073	0.71073	0.71073
<i>T</i> / K	150(2)	295(2)	150(2)
Crystal dimensions / mm ³	0.31 × 0.25 × 0.02	0.45 × 0.32 × 0.23	0.31 × 0.24 × 0.15
μ / mm ⁻¹	0.114	0.095	0.107
<i>F</i> (000)	348	516	896
<i>hkl</i> range	-7, 8; -9, 9; -13, 15	-7, 7; -32, 32; -9, 9	-5, 7; -27, 23; -14, 18
Number of measured reflections	3922	9063	10434
Number of independent reflections	2482	5093	3886
Number of reflections with <i>I</i> > 2σ(<i>I</i>)	1305	3182	2137
Number of parameters	238	345	324
Δρ _{max} , Δρ _{min} / e Å ⁻³	0.244, -0.265	0.262, -0.246	0.281, -0.257
<i>R</i> [<i>F</i> ² > 4σ(<i>F</i> ²)]	0.0443	0.0805	0.0399
<i>wR</i> (<i>F</i> ²)	0.1366	0.204	0.0980
Goodness-of-fit, <i>S</i>	0.823	1.089	0.831

The data sets were collected using the ω scan mode over the 2θ range up to 54°. The structures were solved by direct methods and refined using the SHELXS and SHELXL programs, respectively.⁵² The structural refinement was performed on *F*² using all data. The hydrogen atoms not involved in hydrogen bonding were placed in calculated positions and treated as riding on their parent atoms [C–H = 0.93 Å and *U*_{iso}(H) = 1.2 *U*_{eq}(C); C–H = 0.97 Å and *U*_{iso}(H) = 1.2 *U*_{eq}(C)] while the others were located from the electron difference map. All calculations were performed using the WinGX crystallographic suite of programs.⁵³ The crystal data are listed in Table 1. Further details are available from the Cambridge Crystallographic Centre.⁵⁴ Molecular structures of the compounds are presented by ORTEP-3⁵⁵ and their packing diagrams were prepared by Mercury.⁵⁶

The powder X-ray diffraction (PXRD) experiments on the samples were performed on a PHILIPS PW 1840 X-ray diffractometer with CuK_{α1} (1.54056 Å)

radiation at 40 mA and 40 kV. The scattered intensities were measured with a scintillation counter. The angular range (2θ) was from 3 to 50° with steps of 0.02°, and the measuring time was 0.5 s per step. The data collection and analysis were performed using the program package *Philips X'Pert*.^{57–59}

RESULTS AND DISCUSSION

We first attempted the solvent-free synthesis of **1** by neat grinding of **napht** and **amipha** in the stoichiometric ratio 1:1. Ball milling for 30 min resulted in a mixture containing solid reactants, as evidenced by PXRD. Even after 80 min of neat grinding, no traces of product were detectable. Guided by the observation that small amount of a liquid phase can significantly enhance the scope and rate of mechanochemical synthesis we turned to LAG.^{12,23,46} To observe LAG mechano-synthesis, as well as to facilitate the characterisation of new materials by single crystal X-ray diffraction, LAG

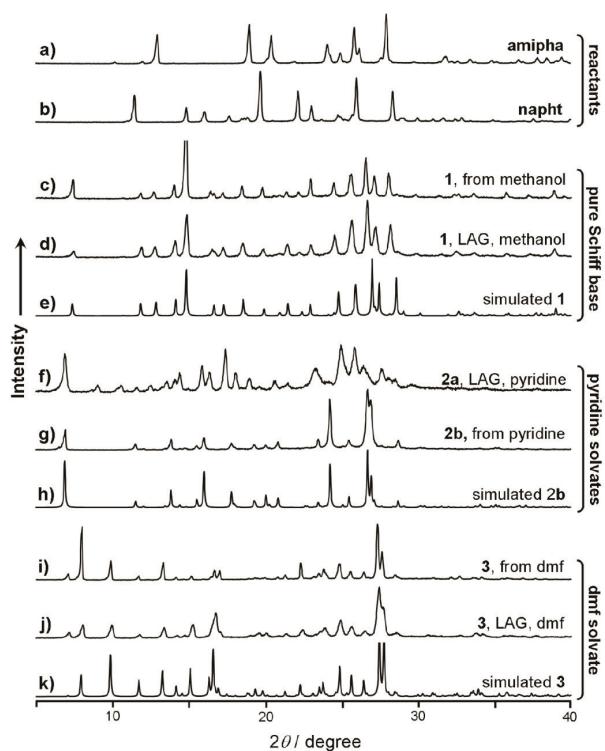


Figure 2. PXRD patterns of: a) 5-aminoisophthalic acid, b) 2-hydroxy-1-naphthaldehyde, c) **1** prepared from a methanol solution, d) **1** prepared by liquid-assisted grinding, e) simulated pattern for **1**, f) **2a** prepared by liquid-assisted grinding, g) **2b** prepared from a pyridine solution, h) simulated pattern for **2b**, i) **3** prepared from a dimethylformamide (**dmf**) solution, j) **3** prepared by liquid-assisted grinding and k) simulated pattern for **3**.

experiments were accompanied by conventional solution-based experiments. Addition of a small quantity of methanol to the reaction mixture, followed by grinding for 50 min, quantitatively provided **1** which was identified by comparing its PXRD pattern with the one calculated from single crystal data (Figure 2c–e). Liquid-assisted grinding of solid reactants in stoichiometric ratio and in the presence of a small amount of pyridine afforded a yellow powder, **2a**, of PXRD pattern not identical to that of crystals prepared by crystallisation of **1** from pyridine, **2b** (Figure 2f). TG curve of **2a** shows a mass loss of 25 % in the range between 45 and 230 °C which corresponds to a loss of one pyridine and one water per a Schiff base molecule (expected 22 %). This indicates that **2a** is a pyridine solvate monohydrate of **1**. Unfortunately the TG curve of **2b** did not enable the determination of its composition since the curve shows a weight loss in the range between 70 and 140 °C of 24 % (expected 19 % for one pyridine molecule and 32 % for two). Its correct composition was therefore determined by single-crystal X-ray structural analysis and found to be a pyridine solvate with two molecules of pyridine per a Schiff base molecule.

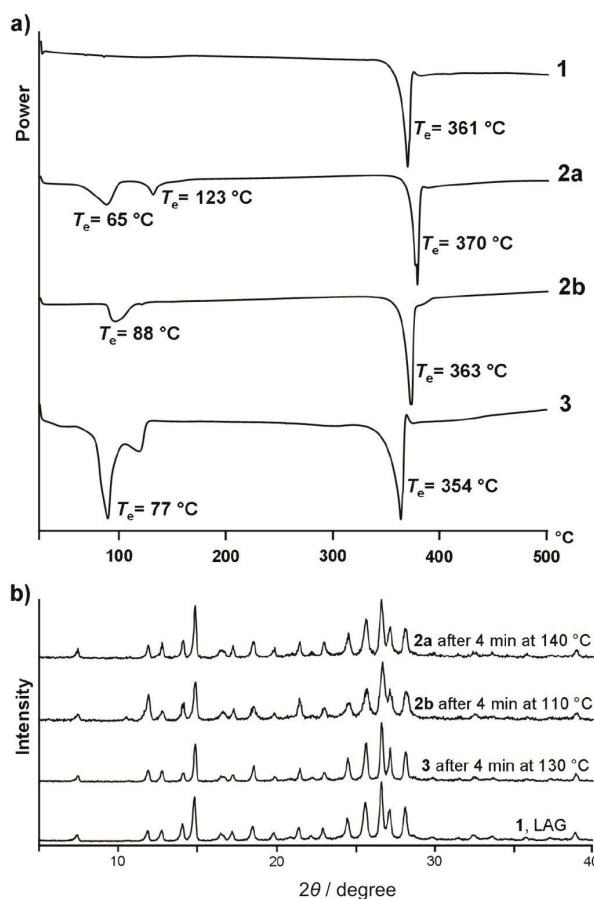


Figure 3. a) DSC curves of **1–3** and b) PXRD patterns of **2a**, **2b** and **3** after annealing.

Liquid-assisted grinding of **napht** and **amipha** in stoichiometric ratio and in the presence of a small amount of dimethylformamide afforded a yellow powder, **3**, identical to that prepared by crystallisation of **1** from dimethylformamide as shown by the comparison of their PXRD patterns (Figure 2i and j). TG curves of **3** obtained by LAG and from solution are almost identical. In the range between 30 and 140 °C they show a mass loss of 23 % (LAG) and 21 % (from solution) which corresponds to a loss of one dimethylformamide and one water per a Schiff base molecule (expected 23 %). The differences in mass loss can be attributed to the differences of particle size in the two samples since the product obtained by LAG is more finely grained than the product obtained from solution. DSC curve of **1** shows one endothermic peak at 361 °C, which corresponds to the melting point and decomposition (Figure 3a). Identical peaks are also observed in DSC curves of all the solvates as they were transformed into **1** by desolvation at lower temperatures. The TG curve of **1** presents no obvious weight loss from 25 to 340 °C. In the range between 350 and 550 °C, a mass loss of 62 % can be attributed to evaporation and pyrolytic decompo-

sition. **2a** and **2b** have remarkably different thermal properties as shown by corresponding DSC curves (Figure 3a). The DSC curve of **2a** shows two endothermic peaks in the range between 50 and 160 °C (39.8 kJ mol⁻¹ and 14.5 kJ mol⁻¹) which correspond to the desolvation. The DSC curve of **2b** shows only one endothermic peak (28.9 kJ mol⁻¹) in the range between 70 and 120 °C. The DSC curve of **3** shows two strong partly overlapping endothermic peaks (90.3 kJ mol⁻¹) in the range between 50 and 130 °C which correspond to desolvation (Figure 3a). All three solvates transform into **1** upon desolvation as shown by PXRD patterns of desolvated products (Figure 3b).

IR spectra of all four compounds show strong bands corresponding to the C–N stretching vibration at about 1620 cm⁻¹ (1618 cm⁻¹ for **1**, 1625 cm⁻¹ for **2a**, 1618 cm⁻¹ for **2b** and 1621 cm⁻¹ for **3**). The spectra of all compounds also show strong bands corresponding to carboxylic C–O stretching at 1705 cm⁻¹ and 1693 cm⁻¹ for **1**, at 1715 cm⁻¹ for **2a**, 1706 cm⁻¹ and 1693 cm⁻¹ for **2b** and 1698 cm⁻¹ for **3**. The IR spectrum of **3** also exhibits a band at 1645 cm⁻¹ corresponding to C–O stretching of the dimethylformamide carbonyl group. All four samples show wide signals in the range between 2800 cm⁻¹ to 3500 cm⁻¹ corresponding to N–H and O–H stretching vibrations. These wide signals are the result of the superposition of several N–H and O–H stretching bands (corresponding to several N–H and O–H bonds in each compound) which are expectedly very wide due to the involvement of N–H and O–H groups in strong intramolecular and intermolecular hydrogen bonding.

The crystal structures of **1**, **2b** and **3** were determined by X-ray diffraction on single crystal samples (Figure 4) and their identity confirmed by comparing the calculated powder diffraction patterns with those obtained by measurement of the bulk material (Figure 2). The composition of **1** and **3** was found to be in accordance with the one deduced from the thermogravimetric measurements and **2b** was found to be a pyridine solvate with two pyridine per a Schiff base molecule. The Schiff base molecule is in all three structures approximately planar with the dihedral angle between the mean planes of the naphthyl and isophtalyl groups of 8.7° in **1**, 0.7° in **2b** and 3.1° in **3**. This planar conformation is in part stabilised by an intramolecular hydrogen bond between the atoms O1 and N1 (2.574 Å in **1**, 2.545 Å in **2b** and 2.543 Å in **3**). As in *o*-hydroxynaphthaleneamine Schiff bases the hydrogen atom may travel along this hydrogen bond, and thus be bonded either to the oxygen or to the nitrogen atom, such molecules can exist in two tautomeric forms.

Here presented Schiff base appears to be present as the keto-amine tautomer in all three structures. This is evidenced by the position of the electron density maximum corresponding to the hydrogen atom partici-

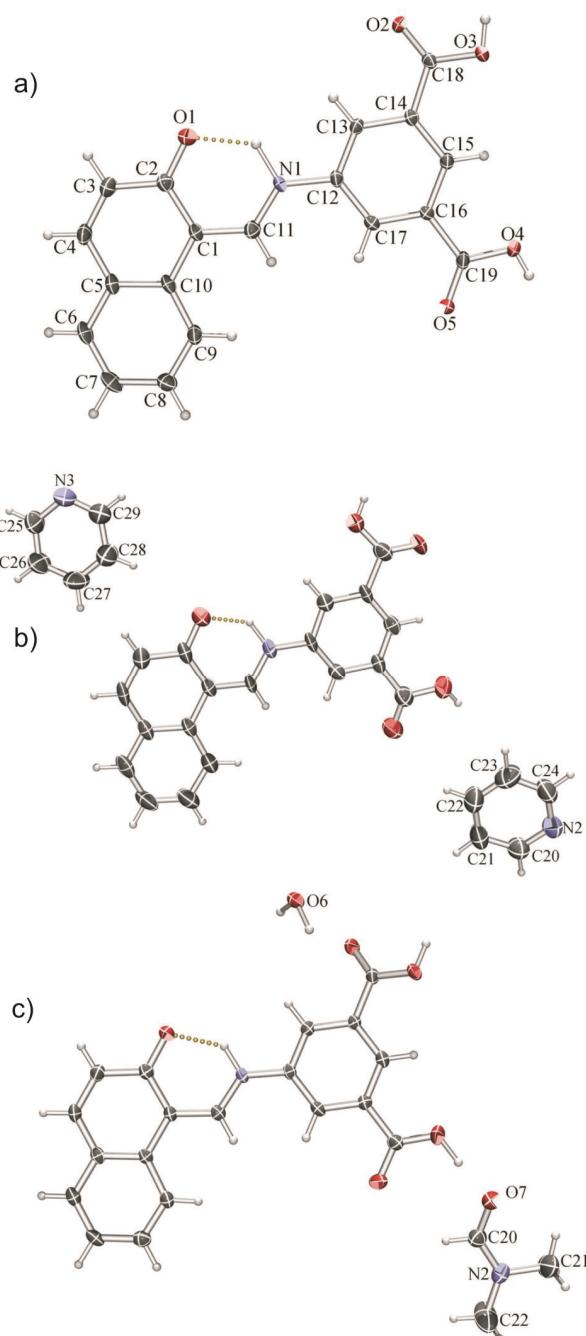


Figure 4. Ortep plots of the asymmetric units in a) **1**, b) **2b** and c) **3**. The atom labelling scheme is the same for the Schiff base molecule and is shown in a). Thermal ellipsoids are plotted at 50 % probability level, and the hydrogen atoms are shown as small spheres of arbitrary radius.

pating in the intramolecular hydrogen bond, which is in the vicinity of the nitrogen atom (Figure 5). However, while in **1** and **3** this position is rather unequivocal (Figure 5a and b) with the maximal electron density of *ca* 0.8 e/Å³, in the case of **2b** the positioning of the hydrogen atom from the electron difference map is somewhat dubious, since the maximal electron density (expectedly

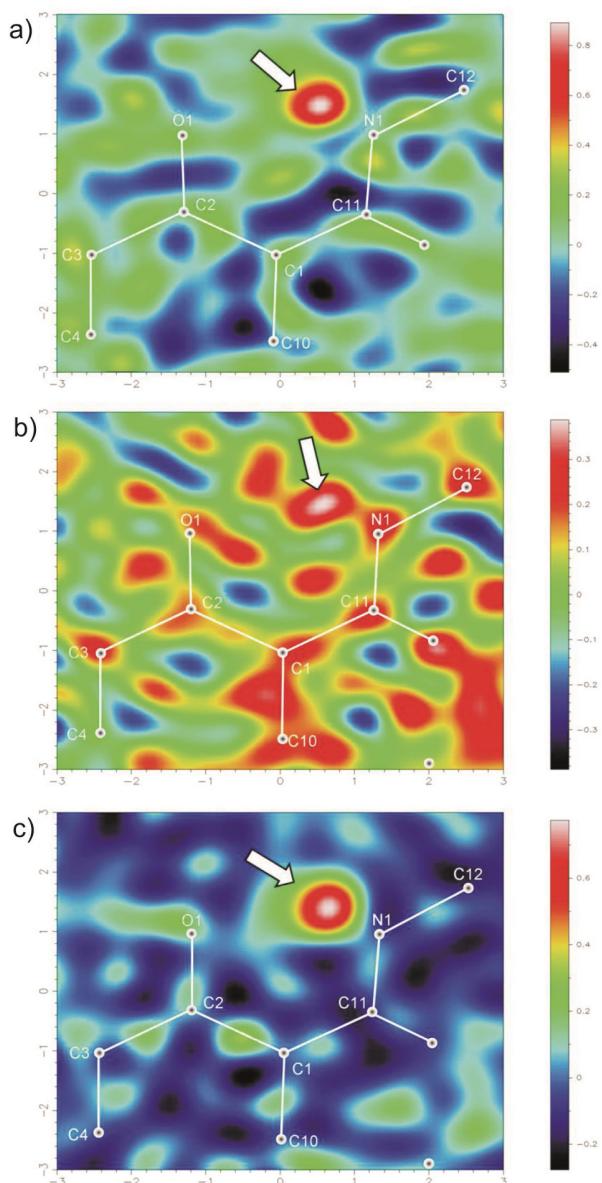


Figure 5. Electron difference maps of the section of Schiff base molecule participating in the intramolecular hydrogen bonding in a) **1**, b) **2b** and c) **3**. The maximum corresponding to the location of the hydrogen atom is indicated by an arrow.

marking the position of the hydrogen atom) is only *ca* 0.4 e/Å³, which is scarcely more than the artificial maxima which comprise the inherent unevenness of the electron difference map (Figure 5c). It was therefore necessary to study carefully whether the bond lengths and angles are in agreement with those expected for keto-amine tautomer.³ For **1** and **3** these parameters do confirm the tautomeric form, with the C2=O1 bond length of 1.267 Å in **1** (1.279 Å in **3**), C1–C11 of 1.400 Å in **1** (1.395 Å in **3**) and C11–N1 of 1.321 Å in **1** (1.323 Å in **3**). Both structures also show the shortening of C3–C4 bond (1.334 Å in **1** and 1.334 Å in **3**) as well

as lengthening of other bonds within the same aromatic ring, consistent with the quinoid effect which is characteristic for keto-amine tautomeric forms of *o*-hydroxynaphthalideneamines. **2b** also shows the same structural characteristics of the keto-amine tautomer, although with the C=O and C1–C11 bonds somewhat longer (1.281 Å and 1.401 Å), C11–N1 somewhat shorter (1.317 Å), and the quinoid effect slightly less pronounced (C3–C4 bond of 1.343 Å). This could indicate that although the Schiff base molecules in **2b** exist predominately as the keto-amine tautomer, a minute amount of enol-imino tautomer might be present. This would also account for the lower residual electron density in the position of the keto-amine hydrogen.

The complete absence of the enol-imino tautomer in **1** and **3** could be attributed to the specific intermolecular interactions that the Schiff base molecule is involved in their crystal structures. Namely, in both these structures atom O1 participates in hydrogen bonding as a hydrogen acceptor – for a carboxylic group of a neighbouring molecule in **1** (O3–H1o···O1 of 2.686 Å) and a water molecule in **3** (O6–H1w···O1 of 2.658 Å). As a carbonyl oxygen atom is much stronger hydrogen acceptor than a hydroxyl one, the participation of O1 in such hydrogen bonding would expectedly reduce the probability of proton transfer from N1 to O1, thus stabilising the keto-amine tautomer. In **2b** pyridine molecules are hydrogen bonded to both carboxyl groups, thus leaving no strong hydrogen donors free to bind to O1. This absence of hydrogen bond renders it more likely that in some molecules hydrogen atom is transferred from N1 to O1, and doing so, transforming the keto-amine into enol-imino tautomer.

Differences in the intermolecular hydrogen bonding in the structures of **1**, **2b** and **3** lead to significant differences in their crystal packing (Figure 6). In **1** each molecule acts as a donor and an acceptor of O3–H1o···O1 hydrogen bonds with the same neighbour which leads to the formation of centrosymmetric dimers through a $R_2^2(22)$ hydrogen bonding motif (Figure 7). This leaves the other carboxyl group free to bond with a carboxyl group of another neighbouring molecule *via* the centrosymmetric $R_2^2(8)$ homosynthron characteristic for carboxylic acids. This combination of hydrogen bonds leads to the formation of chains of molecules along the [01-1] crystallographic direction. The chains are further connected *via* weak C–H···O hydrogen bonds into approximately planar sheets which are stacked along the *a* axis. This hydrogen bonding network can be identified as the cause of relatively high melting point of **1** in comparison to other Schiff bases of similar molecular mass.

In **2b** both carboxyl groups of a Schiff base molecule are hydrogen bonded to pyridine molecules. The location of the hydrogen atom as well as the geometric

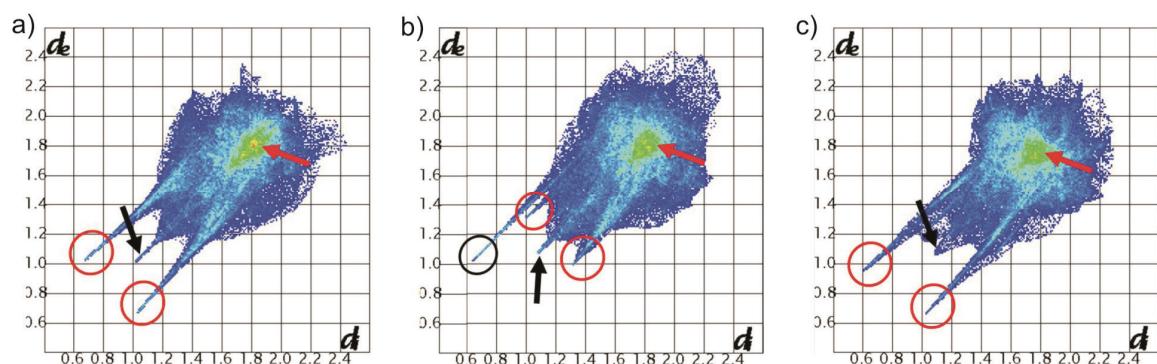


Figure 6. Two-dimensional fingerprint plot for: a) compound **1**, b) a pyridine solvate **2b**, c) a dimethylformamide solvate **3** (marked corresponding regions: red circle for O···H contacts, black circle for N···H contacts, red arrow for C···C contacts and black arrow for short H···H contacts).

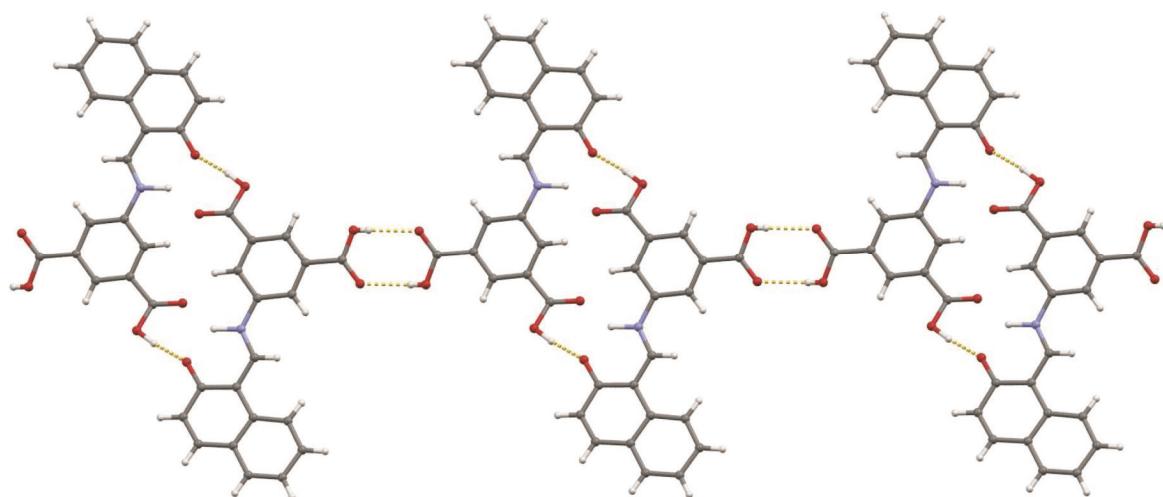


Figure 7. Chains of hydrogen bonded molecules in the crystal structure of **1**.

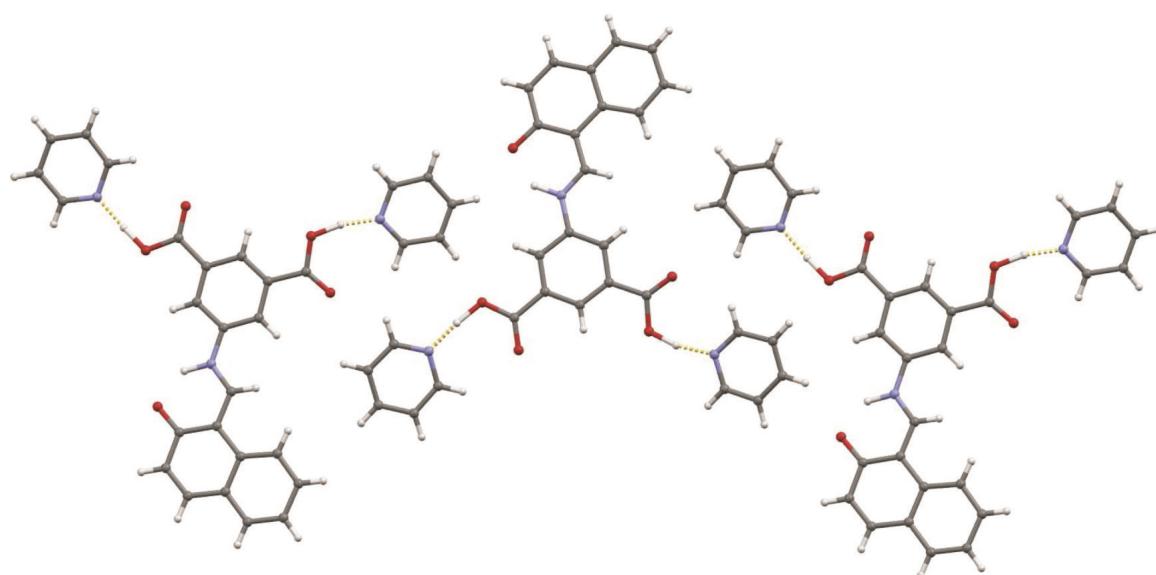


Figure 8. Packing of molecular adducts in the crystal structure of **2b**.

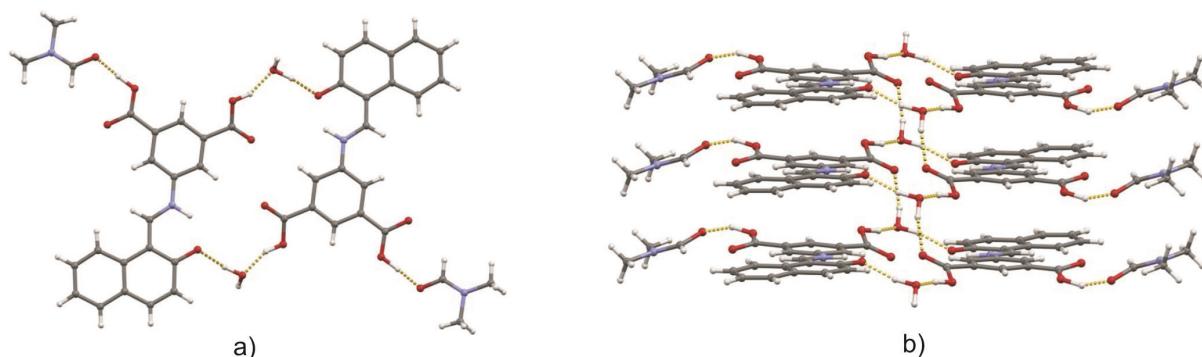


Figure 9. a) Centrosymmetric hexamers in the crystal structure of **3** and b) their bonding into chains.

parameters of carboxyl groups (carbonyl bond lengths C18–O3 of 1.206 Å, C19–O5 of 1.189 Å, hydroxyl C18–O2 of 1.321 Å, C19–O4 of 1.340 Å) and pyridine rings (C20–N2–C24 of 116.5° and C26–N3–C30 of 116.7°), are in accordance with the position of the hydrogen atom on the carboxyl oxygen, thus proving that **2b** is indeed a pyridine solvate, rather than a pyridinium salt. The pyridine rings are approximately coplanar with the isophthalyl group (dihedral angles of 21.3° and 11.2°), due to additional C–H···O hydrogen bonds between the pyridine molecules (C20–H20···O3 of 3.293 Å and C30–H30···O5 of 3.280 Å). The connection between the formation of planarity of supramolecular aggregates of pyridines and acids has already been well documented both with carboxylic acids^{60,61} and substituted phenols.⁶² These three-molecular aggregates are further connected by weak interactions into layers which are stacked along the [101] direction (Figure 8).

In **3** the water molecule which is a hydrogen donor to the O1 atom of a Schiff base molecule, also acts as an acceptor of a O3–H1o···O6 hydrogen bond (2.538 Å) with a carboxyl group from a neighbouring Schiff base molecule. The same neighbour is also an acceptor of a O6–H1w···O1 hydrogen bond with a second water molecule which in turn acts as an acceptor of a carboxyl hydrogen from the first Schiff base molecule. This leads to a hydrogen bonded centrosymmetric ring of $R_4^4(26)$ topology, which corresponds to the $R_2^2(22)$ ring in the structure of **1**, with a water molecule interpolated between the carboxylic group and O1 carbonyl oxygen. Unlike in **1**, however, the remaining carboxyl groups do not bind to carboxyl groups of neighbouring molecules, but rather to solvent **dmf** molecules, forming thus a centrosymmetric hexamer (Figure 9). The **dmf** molecules are disordered so that the formyl groups in two components are twisted by 180°. In the major component (56 %) the formyl carboxyl oxygen of the **dmf** molecule is hydrogen bonded to O4 carboxyl atom (O4–H2o···O7 of 3.485 Å) while in the minor component is hydrogen bonded to O5 carboxyl atom

(O5–H2oa···O7a of 3.566 Å). This rotation of the formyl group does not influence the position of the dimethylamino group, so both components occupy the same position in the crystal structure. Hydrogen bonding of the formyl carboxyl oxygen however requires the O4–C19–O5 carboxyl groups also to be disordered as changing of the position of the carboxyl hydrogen from O4 to O5 atom implies the change of bond orders (C19–O4 becomes a double bond, and C19–O5 a single). This is in accordance with the measured bond lengths of 1.269 Å (C19–O4) and 1.267 Å (C19–O5), which is intermediate between a carboxyl C–O single (1.21 Å) and C=O double bond (1.31 Å). This is due to the fact that the measured bond lengths are an average throughout the structure.

CONCLUSION

Although the Schiff base **1** and the dimethylformamide/water solvate **3** can be prepared both by LAG and in solution, in the case of pyridine solvates, the mechanochemical and conventional synthesis have shown to be complementary; **2a** was obtained only by LAG, whereas **2b** formed quantitatively from solution. A possible reason for this is larger water/pyridine ratio in the LAG experiment (as water is produced by the condensation reaction). The presence of two carboxyl groups on the Schiff base molecule leads to a specific hydrogen bonding pattern in the crystal structure. The high stability of this hydrogen bonding pattern was evidenced not only by a high melting/decomposition temperature of **1**, but also by the fact that upon the loss of solvent molecules all the solvates transform into **1**. The specific hydrogen bonding patterns in **1** and its solvates have some effect on the keto-enol equilibrium in the solid state, as a hydrogen bond to the carbonyl oxygen apparently stabilises the keto-amine tautomer.

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