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**SUPRAMOLECULAR ARCHITECTURE OF A 1:1 COMPLEX
OF 5-AMINOISOPHTHALIC ACID WITH 1,2-BIS(4-PYRIDYL)ETHANE**

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The crystal structure has been solved for a 1:1 molecular complex of 5-aminoisophthalic acid and 1,2-bis(4-pyridyl)ethane. Crystal data: $C_{20}H_{19}N_3O_4$, $M = 365.38$, orthorhombic, $P2_1/n$ space group; unit cell parameters: $a = 10.7521(6)$ Å, $b = 11.9151(6)$ Å, $c = 13.7645(8)$ Å, $V = 1763.40(17)$ Å³, $Z = 4$, $d_x = 1.376$ g/cm³, $R_1 = 0.0514$, $T = 110$ K. The 5-aminoisophthalic acid molecule is linked by O—H···N hydrogen bond to 1,2-bis(4-pyridyl)ethane, forming a linear hydrogen bonded chain running parallel to [0 3 0]. In the crystal, the intermolecular interactions link 5-aminoisophthalic acid molecules with head-to-head N—H···O hydrogen bonds generating $R_2^2(14)$ centrosymmetric dimers, which are extended head-to-tail *via* amine and carboxyl group N—H···O=C hydrogen bond. The supramolecular structure is consolidated by C—H···O hydrogen bonds and $\pi\cdots\pi$ interactions.

К e y w o r d s: 5-aminoisophthalic acid, 1,2-bis(4-pyridyl)ethane, hydrogen bonding, centrosymmetric, supramolecular.

The 1,2-bis(4-pyridyl)ethane is a versatile spacer and a useful component in the construction of supramolecular architectures both by means of protonation at pyridine nitrogen involving N⁺—H···O⁻ synthon and through formation of co-crystal involving N···H—O hydrogen bonds [1, 2]. Each of the pyridyl nitrogen atoms acts as a hydrogen bond acceptor, forming linear hydrogen-bonded associations [3–5]. The structures of a number of 1:1 adducts of benzoic acid derivatives with 1,2-bis(4-pyridyl)ethane have been reported [6, 8]. Not only benzoic acid derivatives are more effective for the promotion of hydrogen-bonding extensions, but also from various weak noncovalent interactions, such as $\pi\cdots\pi$ stacking and C—H···π interactions.

This study is the continuation of our research [9, 10] on the complexes formed by 1,2-bis(4-pyridyl)ethane (**Bpe**) with 5-aminoisophthalic acid (**ApA**). We present here the crystal structure of the 5-aminoisophthalic acid and 1,2-bis(4-pyridyl)ethane 1:1 adduct. The structure of the title compound comprises 1,2-bis(4-pyridyl)ethane molecule and 5-aminoisophthalic acid molecule, with no proton transfer. Neutral molecules connected via O—H···N hydrogen-bonds leading to chains is found in the 1:1 co-crystals.

EXPERIMENTAL

Materials and methods. All chemicals purchased were reagent grade and used without further purification. Elemental analysis (C, H and N) was performed on a Heraeus CHN—O-Rapid elemental analyzer. FT-IR spectra were recorded in the range 400—4000 cm⁻¹ on a JASCO 6200 FT-IR spectrophotometer using a KBr pellet.

Table 1

Crystal data, experimental details and refinement parameters for the structure of [C₁₂H₁₂N₂·C₈H₇NO₄]

Chemical formula	[C ₁₂ H ₁₂ N ₂ ·C ₈ H ₇ NO ₄]
Molecular weight	365.38
Temperature, K	110(2)
Wavelength, Å	0.71073
Crystal system	Orthorhombic
Space group	P2 ₁ /n
Unit cell parameters, <i>a</i> , <i>b</i> , <i>c</i> , Å	10.7521(6), 11.9151(6), 13.7645(8)
Volume, Å ³	1763.40(17)
<i>Z</i>	4
<i>d</i> _x , g/cm ³	1.376
Extinction coefficient, mm ⁻¹	0.098
<i>F</i> (000)	768
Crystal size, mm	0.28×0.26×0.08
<i>h</i> , <i>k</i> , <i>l</i> limits	-12 ≤ <i>h</i> ≤ 12, -13 ≤ <i>k</i> ≤ 14, -16 ≤ <i>l</i> ≤ 12
Data completeness on θ, deg.	2.95—25.20, 99.8 %
Measured / independent reflections	7358 / 3169 [<i>R</i> (int) = 0.0499]
Reflections with <i>I</i> > 2σ(<i>I</i>)	2387
Number of parameters	3399
Transmission	0.98517—1.00000
Refinement	Full-matrix least squares against <i>F</i> ²
GOOF against <i>F</i> ²	0.942
Final <i>R</i> factor <i>I</i> > 2σ(<i>I</i>)	<i>R</i> ₁ = 0.0514, <i>wR</i> ₂ = 0.1314
<i>R</i> factor (all data)	<i>R</i> ₁ = 0.0690, <i>wR</i> ₂ = 0.1372
Residual electron density, e/Å ⁻³	-0.473 and 0.061
CCDC deposition number	917169

X-ray crystallographic analysis. Diffraction data of the title compound were collected at 110 K on a Oxford Diffraction Gemini-S CCD diffractometer with graphite-monochromatized MoK_α radiation ($\lambda = 0.71073 \text{ \AA}$) in multi-scan mode. The structure was solved by direct methods using the program SHELXS-97 [11] and refined by full-matrix least-squares methods on *F*² using the program SHELXL-97. All non-hydrogen atoms were refined anisotropically. Hydrogen atoms were placed in geometrically calculated positions, with O—H = 0.84, C—H = 0.95 (aromatic), and 0.99 Å (methyl), and refined in the riding model with *U*_{iso}(H) = 1.5_{eq}(C) for methyl and *U*_{iso}(H) = 1.2_{eq}(C) for others. Besides, H atoms can also localize in the hydrogen bonds, but we found the two methods to localize hydrogen atoms are in the same position. The crystallographic data and experimental details for structural analyses are summarized in Table 1.

Structural CIF file containing complete information on the studied structure was deposited with CCDC, deposition number 917169, and is freely available upon request from the following web site: www.ccdc.cam.ac.uk/data_request/cif.

Synthesis. The 5-aminoisophthalic acid (0.0362 g, 0.20 mmol) and 1,2-bis(4-pyridyl)ethane (0.0369 g, 0.20 mmol) were dissolved in 20 mL methanol-water (1:1), the solution was refluxed for 30 min. The filtered solution was transferred to a 25 mL tube. After one week at room temperature colorless transparent crystals were obtained (yield 55 %). We have also attempted mixing the components in the solution at room temperature no crystallization occurred even in three weeks. Elemental analysis calcd. for C₂₀H₁₉N₃O₄ (*M*_r = 365.38): C 65.69, H 5.24, N 11.49 %. Found: C 66.03, H 5.29,

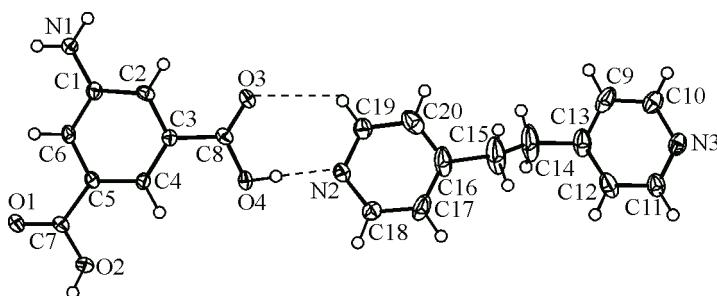


Fig. 1. View of the 1:1 molecular associate in the structure of the title compound with atomic labeling. Thermal ellipsoids are drawn at the 50 % probability level

N 11.75 %. IR (KBr, ν , cm^{-1}): 3446(s), 3366(s), 1701(s), 1645(w), 1608(s), 1459(m), 1421(s), 1415(s), 1359(m), 1279(s), 1204(m), 899(m), 818(m), 769(w).

RESULTS AND DISCUSSION

The ORTEP view of the title compound is shown in Fig. 1. The bond lengths and angles are listed in Table 2 and 3. The asymmetric unit is composed of one molecule of Bpe and one molecule of Apa, with no proton transfer. In the structure, the molecules associate 5-aminoisophthalic acid and 1,2-bis(4-pyridyl)ethane via carboxylic and pyridine group O—H \cdots N, forming linear hydrogen bonding

Table 2

Bond lengths d, Å in the structure of [C₁₂H₁₂N₂·C₈H₇NO₄]

Bond	d	Bond	d	Bond	d	Bond	d
O1—C7	1.213(3)	C3—C4	1.389(4)	N2—C19	1.340(4)	C13—C14	1.503(5)
O3—C8	1.213(3)	C3—C8	1.494(3)	C5—C4	1.395(3)	C16—C20	1.367(6)
O4—C8	1.319(3)	N3—C10	1.338(4)	C5—C7	1.492(4)	C16—C17	1.391(5)
C2—C1	1.393(3)	N3—C11	1.337(3)	C18—C17	1.359(5)	C16—C15	1.510(5)
C2—C3	1.387(4)	O2—C7	1.319(3)	C10—C9	1.378(5)	C12—C11	1.382(4)
C6—C5	1.387(4)	N1—C1	1.389(4)	C13—C12	1.373(6)	C19—C20	1.362(4)
C6—C1	1.389(4)	N2—C18	1.327(4)	C13—C9	1.398(5)	C15—C14	1.408(6)

Table 3

Bond angles ω, deg. in the structure of [C₁₂H₁₂N₂·C₈H₇NO₄]

Angles	ω	Angles	ω	Angles	ω
C1—C2—C3	120.43(24)	C4—C5—C7	122.92(21)	C9—C13—C14	122.42(32)
C5—C6—C1	121.31(24)	N2—C18—C17	123.18(30)	C20—C16—C17	116.80(37)
C4—C3—C2	120.97(25)	N3—C10—C9	122.43(29)	C20—C16—C15	119.91(36)
C4—C3—C8	122.22(22)	O3—C8—O4	123.07(25)	C17—C16—C15	122.96(32)
C2—C3—C8	116.78(23)	O3—C8—C3	122.54(22)	C10—C9—C13	120.42(33)
C10—N3—C11	117.66(25)	O4—C8—C3	114.39(23)	C18—C17—C16	119.83(33)
N1—C1—C2	121.11(24)	C3—C4—C5	118.74(22)	C13—C12—C11	120.82(34)
N1—C1—C6	120.39(24)	O1—C7—O2	123.04(25)	N2—C19—C20	123.10(26)
C2—C1—C6	118.48(22)	O1—C7—C5	122.16(22)	N3—C11—C12	122.50(25)
C18—N2—C19	116.91(26)	O2—C7—C5	114.80(23)	C19—C20—C16	120.08(34)
C6—C5—C4	120.06(24)	C12—C13—C9	116.16(37)	C14—C15—C16	119.03(32)
C6—C5—C7	116.99(23)	C12—C13—C14	121.32(36)	C15—C14—C13	112.31(32)

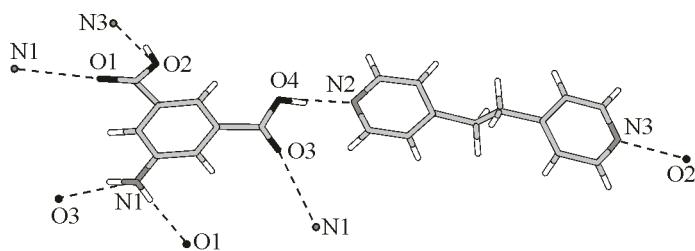


Fig. 2. Fragment of the structure that illustrates the hydrogen bonds

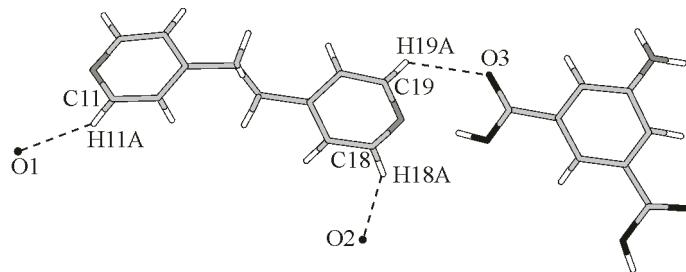


Fig. 3. Fragment of the structure that illustrates non-classical hydrogen bonds

Table 4

Parameters of the hydrogen bonds (\AA , deg.)

D—H···A	<i>d</i> (D—H)	<i>d</i> (H···A)	<i>d</i> (D···A)	$\angle\text{DHA}$	Symmetry operations
N1—H1A···O1	0.90	2.15	2.990(3)	156	$1/2-x, 1/2+y, 5/2-z$
N1—H1B···O3	0.89	2.25	3.041(3)	148	$1/2-x, -1/2+y, 5/2-z$
O2—H2B···N3	0.83	1.83	2.613(3)	157	$1/2-x, -3/2+y, 3/2-z$
O4—H4B···N2	0.83	1.81	2.640(3)	173	
C11—H11A···O1	0.93	2.58	3.223(3)	126	$1/2-x, 3/2+y, 3/2-z$
C18—H18A···O2	0.93	2.54	3.339(4)	145	$1/2-x, 1/2+y, 3/2-z$
C19—H19A···O3	0.93	2.60	3.245(3)	127	

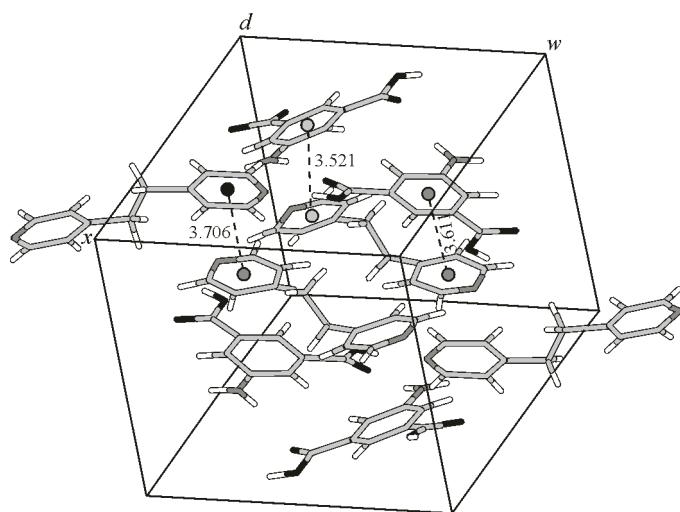


Fig. 4. Fragment of the structure of the title compound that illustrates the $\pi\cdots\pi$ interactions

Table 5

Parameters of $\pi \cdots \pi$ interactions for title compound (\AA , deg.)

Cg(I)	Cg(J)	Cg···Cg	Perpendicular distance	Dihedral angles	Symmetry operations
Cg(1)	Cg(2)	3.652	3.7057(18)	4.52(15)	$1/2-x, -1/2+y, 3/2-z$
Cg(1)	Cg(3)	3.415	3.6107(16)	9.47(13)	$-x, 1-y, 2-z$
Cg(2)	Cg(3)	3.442	3.5213(16)	5.37(13)	$-1/2+x, 3/2-y, -1/2+z$

Cg(1): N2/C16—C17—C18—C19—C20, Cg(2): N3/C9—C10—C11—C12—C13, Cg(3): C1—C2—C3—C4—C5—C6.

parallel to [0 3 0]. The intermolecular interactions link 5-aminoisophthalic acid molecules with head-to-head N—H···O hydrogen bonds, generating $R_2^2(14)$ [12] centrosymmetric dimers, which are extended head-to-tail *via* N—H···O=C hydrogen bond between the amine and carboxylic groups. The hydrogen bonding interactions (Fig. 2 and Fig. 3) are normal, based on H···A distances of 1.81—2.25 \AA , D···A distances of 2.613(3)—3.041(3) \AA , and bond angles of 100—173° (Table 4) [13].

In the title compound, the Apa molecule has usual geometry. The C—C_(ring), C—C_(carboxyl) bond lengths and CCC_(ring), CCC_(carboxyl) angles fall within 1.387(4)—1.395(3) \AA , 1.213(3)—1.492(4) \AA and 118.48(22)—121.31(24)°, 116.78(23)—116.99(23)° ranges with the average values of 1.390(3) \AA , 1.353(3) \AA , and 120.00(13)°, 116.89(23)°. There is no significant difference between the bond angles at the carbon atoms bearing the carboxylic groups and those at the rest of the carbon atoms. The former vary within 120.06—120.97°, and the latter vary within 118.74—121.31° with the average values of 120.52° and 120.16°, the difference being only 0.46°. Both carboxylic groups are nearly coplanar with the aryl ring, as indicated by the torsion angles C3—C8—O3—O4 and C5—C7—O1—O2 of 1.67(35) and −6.49(47)°, respectively [14].

The stability of the structure is additionally reinforced by $\pi \cdots \pi$ interactions of the 1,2-bis(4-pyridyl)ethane molecule with the centroid of the benzene ring of the 5-aminoisophthalic acid molecule with the distances between Cg1···Cg2 being 3.7057(18) \AA , Cg1···Cg3 being 3.6107(16) \AA , Cg2···Cg3 being 3.5213(16) \AA . The dihedral angles between the two rings are 4.52(15)°, 9.47(13)° and 5.37(13)°, respectively (Fig. 4 and Table 5). The stacks are joined into a 3D architecture by O—H···N and N—H···O hydrogen bonds, and by weak C—H···O hydrogen bonds and $\pi \cdots \pi$ stacking interactions [15, 16]. The packing of the molecules in the structure is illustrated in Fig. 5.

CONCLUSIONS

In this study, 1,2-bis(4-pyridyl)ethane and 5-aminoisophthalic acid have been used to construct supramolecular architectures based on crystal engineering principles. It has been observed that supramolecular hydrogen-bonded polymers are formed through self-assembly via O—H···N, O—H···O or C—H···O hydrogen bonding between 1,2-bis(4-pyridyl)ethane and 5-aminoisophthalic acid molecules. The conformations of bipyridines play an important role in the design and construction of desirable frameworks. The change of length, symmetry, and flexibility of the bipyridines could result in a class of materials containing diverse architectures and some related experiments are in progress.

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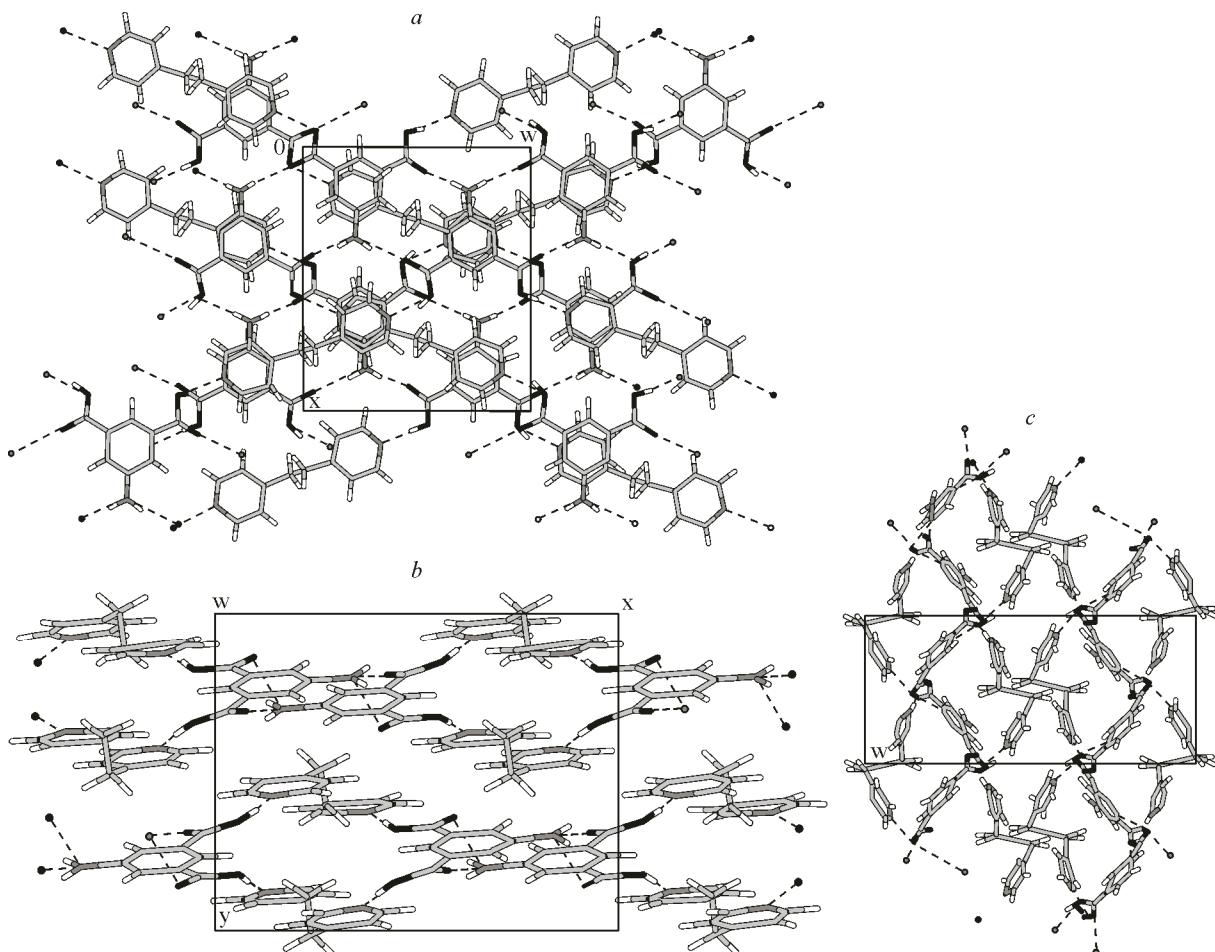


Fig. 5. Packing of the molecules in the title crystal: viewed along the *a* axis (*a*), viewed along the *b* axis (*b*), viewed along the *c* axis (*c*). Dashed lines denote hydrogen bonds

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