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## X-ray Crystallographic Structure of 3-(Propan-2-ylidene) benzofuran-2(3*H*)-one

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Abstract: 3-(Propan-2-ylidene)benzofuran-2(3H)-one, C<sub>11</sub>H<sub>10</sub>O<sub>2</sub>, crystallizes in the monoclinic space group  $P2_1/c$  with unit cell parameters a = 7.1869(3), b = 18.0636(10), c = 13.1656(7) Å,  $\beta$ = 96.763(3)°,  $V = 1697.28(15) \text{ Å}^3$ , Z = 8 (Z' = 2 independent molecules, A and B, per asymmetric unit),  $D_c = 1.363$  g cm<sup>-3</sup> and the linear absorption coefficient = 0.093 mm<sup>-1</sup>. The crystal structure determination was carried out using MoKa X-ray data measured at 120(2) K. In the final refinement cycle the data/restraints/parameter ratios were 3827/0/239, the goodness-of-fit on F<sup>2</sup> = 1.019. Final R indices for [I>2sigma(I)] were R1 = 0.0517, wR2 = 0.1115 and R indices (all data) R1 = 0.1007, wR2 =0.1354. The largest electron density difference peak and hole were 0.254 and -0.244 electrons Å<sup>-3</sup>, respectively. The two independent molecules A and B have essentially identical bond lengths and angles and are highly planar, with rms deviation for all 12 non-H atoms of 0.0292 Å for molecule A and 0.0592 Å for molecule B. The two molecules in the asymmetric unit are assembled parallel to each other within 1.58(4)° with each of the six atoms of benzene ring A overlapping with one of the atoms in benzene ring B at a mean distance of 3.84(3) Å). The closest contacts between molecules A and B are C(12A)H---O(4B) = 3.521 Å and C(11B)H---O(4A) = 3.441 Å. The crystal structure is formed by infinite sheets of these assemblies all lying parallel to the  $(1 \ 0 \ \overline{1})$  plane. The presence of two independent molecules in the asymmetric unit provides an opportunity to examine the molecular geometry in detail by comparison. The benzene ring in both molecules A and B exhibits distortions as a result of the presence of the furan moiety. In particular, C(1)-C(6) =1.369(2) Å in molecule A and 1.367(2) Å in molecule B are both significantly less than the average of the other 10 C-C bonds, 1.393(2), by about  $12\sigma$ . Other examples of crystal structures are discussed where this effect is observed.

**Keywords:** Knoevenagel condensation; X-ray crystal structure; molecular geometry

#### 1. Introduction

Recent interest in the Knoevenagel condensations of indol-2-one with aldehydes or ketones led to the report of a number of X-ray structures<sup>1, 2</sup> of the resulting 3-methylidene-1,3-dihydro-2*H*-indol-2-one compounds (**Figure 1**), which display intermolecular hydrogen bonding between the NH and C=O groups of neighbouring units. The aim, in this current study, is to compare the structure of the Knoevenagel condensation product of acetone with benzofuran-2(*3H*)-one as opposed to that with indolin-2-one since, in the former case, due to the lack of an NH group, no intermolecular H-bonding, of the type described above, is possible in the final product.



3-(Propan-2-ylidene)benzofuran-2(3H)-one 3-(Propan-2-ylidene)indolin-2-one

Figure 1. Knoevenagel condensation products.

#### 2. Results and Discussion

Figure 2 shows the chemical scheme and crystallographic atom numbering for  $C_{11}H_{10}O_2$  (I). Figure 3 shows molecule A determined in the structure analysis and drawn with Visualizer, Discovery Studio (Release 3.5, Accelrys Inc., 2012).<sup>3</sup> ORTEP<sup>4</sup> and

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RASTER3D,<sup>5</sup> as implemented in WinGX, were used to prepare **Figure 4** which shows the relative disposition of molecules A and B depicted with their thermal ellipsoids at 50% probability.



Figure 2. Schematic chemical structure and atom numbering for  $C_{11}H_{10}O_2$  (I). The bonding shown is discussed. X denotes the region of geometry discussed in more detail.



**Figure 3.** The 3-(propan-2-ylidene)benzofuran-2(*3H*)one,  $C_{11}H_{10}O_2$  (**I**) molecule A as determined by X-ray analysis (Drawn with Visualizer, Discovery Studio, Release 3.5, Accelrys Inc., 2012)<sup>3</sup> [http://accelrys.com/products/datasheets/whats-newin-discovery-studio.pdf].



**Figure 4**. 3-(Propan-2-ylidene)benzofuran-2(3*H*)-one (Ortep/Raster)<sup>4,5</sup>, molecules A and B. The two molecules in the asymmetric unit are parallel within 1.67(6)° and the benzene rings overlap with a mean intermolecular distance of 3.84(3) Å between corresponding atoms. Thermal ellipsoids are shown at 50% probability.

#### 2.1. Molecular Geometry

#### 2.1.1. Bond lengths

Crystal data are summarized in **Table 1**. Bond lengths are determined to a precision of  $\pm$  0.002Å and bond

angles to  $\pm$  (0.2°). **Table 2** lists corresponding bond lengths in molecules A and B, respectively. These data indicate that there is a complete one to one agreement in bond length values. In the benzene ring all bond lengths are standard aromatic C-C bonds<sup>6</sup> except C(1)-C(6) which is 1.369(2)Å in molecule A and 1.367(2)Å in molecule B, which are both shorter by about  $12\sigma$  than the standard value of 1.40 Å. This is possibly a result of the proximity of O(5) in the adjoining ring. In fact in the chain extending to the furan ring, i.e., C(6)-C(1)-O(5)-C(4), C(1)-O(5) = 1.391(2) Å in molecule A and 1.396(2) Å in molecule B, and O(5)-C(4) = 1.388(2) Å in molecule A and 1.395(2) Å in molecule B, this reduction in bond length is consistent with the presence of a delocalisation effect. This region is designated X in Figure 2.

Table 1. Crystal data and structure refinement for (I).

Identification code	2008src0248 CCDC804780
Empirical formula	$C_{11} H_{10} O_2$
Formula weight	174.20
Temperature	120(2) K
Wavelength (wtd. mean)	0.71073 Å
Crystal system	Monoclinic
Space group	P21/c
Unit cell dimensions	a = 7.1869(3) Å; α = 90°
	b= 18.0636(10)Å;
	$\beta = 96.763(3)^{\circ}$
	c = 13.1656(7) Å; γ = 90°
Volume	1697.28(15) Å <sup>3</sup>
Z	8 (2 molecules/asymmetric
	unit)
D <sub>c</sub>	1.363 g cm <sup>-3</sup>
Absorption coefficient	$0.093 \text{ mm}^{-1}$
F(000)	736
Crystal size	$0.44 \times 0.20 \times 0.08 \text{ mm}^3$
Thota range for data	0.44 X 0.50 X 0.06 IIIII 2 07 to 27 49°
collection	5.07 to 27.40 .
Index ranges	0 = b = 0 $22 = b = 22$
linuex ranges	-5 < -11 < -5, -25 < -K < -25, 17 < -1 < -17
Pofloctions collected in total	-1/<-1/
sphore	20300
Independent reflections	3827 [P(int) = 0.0800]
Completeness to theta =	98.3.0%
27.48°	90.3 %
Max. and min. transmission	0.9926 and 0.9601
Refinement method	Full-matrix least-squares on
	F <sup>2</sup>
Data/ restraints/ parameters	3827/0/239
Goodness-of-fit on F <sup>2</sup>	1.019
Final R indices [I>2sigma(I)]	R1 = 0.0517, wR2 = 0.1149
R indices (all data)	R1 = 0.1007, $wR2 = 0.1354$
Largest diff. peak and hole	0.254 and $-0.244$ electrons Å <sup>-3</sup>

#### 2.1.2. Geometry Optimization

To investigate molecule A further its geometry was optimized in the program Visualizer, Discovery Studio (Release Accelrys 4.0, Inc.. 2013).7[http://accelrys.com/products/discoverystudio/] using the Smart Minimizer algorithm and conjugate gradient minimization criteria. The molecule was typed with the CHARMM forcefield<sup>8</sup> using the Partial Charge method.<sup>9</sup> After 2000 steps convergence was reached and the final potential energy of the molecule I was 61.5 kcal/mol. The results for molecules A and B bond lengths are listed in Table 3. These results show that the energy minimized molecular geometry is consistent with the shortening of bond C(1)-C(6) in both molecules A and B. Further examples of structures where such a short bond length feature may be present are noted below.

Bond Lengths		
Bond	(I)A	(I)B
C(1)-C(2)	1.392(2)	1.395(2)
C(2)-C(3)	1.471(2)	1.469(2)
C(3)-C(10)	1.350(2)	1.349(2)
C(10)-C(11)	1.492(2)	1.494(2)
C(10)-C(12)	1.503(2)	1.494(2)
C(3)-C(4)	1.489(2)	1.479(2)
0(4)-C(4)	1.204(2)	1.204(2)
C(4)-O(5)	1.388(2)	1.395(2)
0(5)-C(1)	1.391(2)	1.396(2)
C(1)-C(6)	1.369(2)	1.367(2)
C(6)-C(7)	1.393(2)	1.398(2)
C(7)-C(8)	1.390(2)	1.391(2)
C(8)-C(9)	1.389(2)	1.389(2)
C(9)-C(2)	1.398(2)	1.397(2)
Bond Angles		
C(1)-C(2)-C(3)	106.5(2)	106.7(2)
C(2)-C(3)-C(10)	130.8(2)	129.8(2)
C(3)-C(10)-C(11)	122.3(2)	121.4 (2)
C(3)-C(10)-C(12)	122.8(2)	123.4(2)
C(11)-C(10)-C(12)	115.0(2)	115.2(2)
C(2)-C(3)-C(4)	104.9(2)	105.1(2)
C(3)-C(4)-O(4)	132.5 (2)	132.6(2)
C(3)-C(4)-O(5)	108.4(2)	108.6(2)
0(4)-C(4)-O(5)	119.2(2)	118.7(2)
C(4)-O(5)-C(1)	108.2(2)	108.4(2)
C(6)-C(1)-O(5)	123.5(2)	123.7(2)
C(6)-C(1)-C(2)	124.6(2)	124.6(2)
C(1)-C(6)-C(7)	116.7(2)	116.7(2)
C(6)-C(7)-C(8)	120.9(2)	120.6(2)
C(7)-C(8)-C(9)	121.1(2)	121.4 (2)
C(8)-C(9)-C(2)	119.1(2)	118.9(2)
Features in region X. Fig	gure 2, are in bold.	

**Table 2.** 1-Benzofuran-2(3*H*)-one: bond lengths [Å] and angles [°] for molecules (**I**) A and B.

reatures in region A, **Figure 2**, are in r

#### 2.1.3. Bond angles

**Table 2** lists corresponding bond angles in molecule A and B, respectively. These data indicate that there is a complete one to one agreement in bond angle values. Of the bond angles in the benzene ring C(1)-C(6)- $C(7) = 116.7(2)^{\circ}$  in both molecules, is significantly less than 120° and may be associated with the unusually short bond length C(1)-C(6) discussed above. The molecular geometry of the two independent molecules A and B is practically identical, as revealed by application of the AUTOFIT routine in PLATON<sup>10</sup> using Molfit with Quaternion Transformation.<sup>11</sup>

**Table 3.** 3-(Propan-2-ylidene) benzofuran-2(*3H*)-one (**I**): X-ray bond lengths [Å] for molecules A and B and molecules A and B after minimization in the Visualizer (Discovery Studio, Relase 4.0, Accelrys Inc., 2013).

( )				,
Bond	X-ray Structure		Minimize	ed Structure
	(I)A	(I)B	А	В
C(1)-C(2)	1.392(2)	1.395(2)	1.402	1.402
C(2)-C(3)	1.471(2)	1.469(2)	1.489	1.488
C(3)-C(10)	1.350(2)	1.349(2)	1.359	1.358
C(10)-C(11)	1.492(2)	1.494(2)	1.515	1.514
C(10)-C(12)	1.503(2)	1.494(2)	1.516	1.516
C(3)-C(4)	1.489(2)	1.479(2)	1.503	1.501
O(4)-C(4)	1.204(2)	1.204(2)	1.225	1.224
C(4)-O(5)	1.388(2)	1.395(2)	1.337	1.337
0(5)-C(1)	1.391(2)	1.396(2)	1.461	1.460
C(1)-C(6)	1.369(2)	1.367(2)	1.367	1.366
C(6)-C(7)	1.393(2)	1.398(2)	1.394	1.394
C(7)-C(8)	1.390(2)	1.391(2)	1.396	1.394
C(8)-C(9)	1.389(2)	1.389(2)	1.398	1.398
C(9)-C(2)	1.398(2)	1.397(2)	1.375	1.375

#### 2.2. Crystal Packing

The two molecules in the asymmetric unit are assembled parallel to each other within  $1.58(4)^{\circ}$  with each of the six atoms of benzene ring A overlapping with one of the atoms in benzene ring B at a mean distance of 3.84(3) Å); **Table 4(a)** and **Figure 5(a)**. The closest contacts between molecules A and B are C(12A)H---O(4B) = 3.521 Å and C(11B)H---O(4A) = 3.441 Å, **Table 4(b)** and **Figures 5(b)** and **(c)**. The crystal structure is formed by infinite sheets of these assemblies all lying parallel to the  $(1 \ 0 \ \overline{1})$  plane, **Figure 6**.

**Table 4(a).** Intermolecular C...C distances in the asymmetric unit (Å).

C(1A)C(9B)	3.853
C(2A)C(2B)	3.848
C(6A)C(8B)	3.798
C(7A)C(7B)	3.804
C(8A)C(6B)	3.837
C(9A)C(1B)	3.886



**Figure 5(a).** 3-(Propan-2-ylidene)benzofuran-2(3*H*)one mode of overlapping of molecules A and B  $(Ortep/Raster)^{4.5}$ . The two molecules in the asymmetric unit are parallel within 1.58(4)°



**Figure 5(b).** 3-(Propan-2-ylidene)benzofuran-2(3*H*)one, overlapping of molecules A and B. The short CH---O contacts as listed in Table 4 are indicated. Benzene – benzene distances range from 3.597 Å for C3(A) ---C3(B) to 3.886 Å for C1(A) --- C1(B); (Drawn with Mercury)<sup>5</sup>.

Further Examples of Structures where Chain X Exhibits Delocalisation

In order to further investigate the delocalization effect observed in (I), molecules A and B, the corresponding bond in 2,3-diphenyl-5-methyl-1-benzofuran  $C_{21}H_{16}O_1$ 

**(II)**,<sup>12</sup> **Figure 7a** and 3-(1-phenylmeth-(*E*)-ylidene)-*3H*-benzofuran-2-one **(III)**,<sup>13</sup> **Figure 7b** was noted. Bond lengths in chain X for molecules **(I)**, **(II)** and **(III)** are listed in **Table 5**. The delocalization in bond C(1)-C(6), evident in all 4 molecules listed here, is less prominent in molecule **(II)**.



**Figure 5(c).** 3-(Propan-2-ylidene)benzofuran-2(3*H*)one, details of the intra molecular CH---O contact in Molecule A and between Molecule A and Molecule B. Essentially equivalent contacts occur within Molecule B and between Molecule B---A (Drawn with Visualizer, Discovery Studio, Release 3.5, Accelrys Inc., 2012)<sup>3</sup> [http://accelrys.com/products/datasheets/whats-newin-discovery-studio.pdf].) see also **Table 4**.



**Figure 6.** 3-(Propan-2-ylidene)benzofuran-2(3*H*)-one crystal packing viewed along **b**. The molecular planes are parallel to  $(1 \ 0 \ \overline{1})$ . (Drawn with Mercury)<sup>5</sup>.

Table 4(b)	. Possible	hydrogen	bonds [A	Å and o	leg.]
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#### 3. Experimental

3-(Propan-2-ylidene)benzofuran-2(3H)-one(I),

 $C_{11}H_{10}O_2,$  was provided by the Trost laboratory  $^{14}\,$  and crystallized from dichloromethane/hexane by vapour diffusion.

#### 3.1. Data Collection

A colourless crystal plate fragment of size  $0.44 \times 0.30 \times$ 0.08 mm<sup>3</sup> was mounted on a glass fibre and flash frozen to 120 K. Intensities were collected, using monochromated MoK $\alpha$  radiation,  $\lambda = 0.71073$  Å, with a Bruker-Nonius *Kappa* CCD camera, employing  $\phi$  and  $\omega$ scans to cover an asymmetric unit. Programs used were: unit cell determination with the program DirAx;<sup>15</sup> data collection was controlled by Collect;<sup>16</sup> data reduction and cell refinement were carried out using the program Denzo;17 an absorption correction was made with SADABS.<sup>18, 19</sup> An Oxford Cryosystems "Cryostreams" 700,<sup>20</sup> enabled the data to be collected at 120 K. The crystals are monoclinic, space group P2<sub>1</sub>/c with unit cell dimensions a = 7.1869(3), b = 18.0636(10), c =13.1656(7) Å,  $\beta$  = 96.763(3)°, V = 1697.28(15) Å<sup>3</sup>, Z = 8 (Z' = 2), D<sub>c</sub> = 1.363 g cm<sup>-3</sup> and linear absorption coefficient = 0.093 mm<sup>-1</sup>. The crystal structure determination was carried out using MoKa X-ray data measured at 120(2) K. In total 20306 integrated reflections were collected, reducing to an asymmetric unit data set of 3827 [R(int) = 0.080], and completeness of data to theta = 27.48° of 98.3 % corresponding to a resolution of 0.770 Å. There was no significant variation in intensity of reference reflections during the course of data collection.

#### 3.2. X-ray Structure Analysis

The crystal structure was solved by Direct Methods (SHELXS-86) and refined using SHELXL-97<sup>21, 22</sup> both implemented in the WinGX system of programs.<sup>23</sup> Nonhydrogen atoms were refined anisotropically by fullmatrix least squares methods. All H atoms were set geometrically and refined in riding mode. Geometrical calculations were made with the programs PARST and PLATON<sup>10</sup> as implemented in WinGX. In the final refinement cycle there were 3827 data to 239 parameters, resulting in a final goodness-of-fit on F<sup>2</sup> of 1.019. Final R indices for [I>2sigma(I)] were R1 = 0.0517, wR2 = 0.1149 and R indices (all data) R1 = 0.1007, wR2 = 0.1354. The largest and smallest difference electron density regions were +0.254 and -0.244 electrons Å-3, respectively. Crystal data are summarized in Table 1.

D-HA	d(D-H) Å	d(HA) Å	d(DA) Å	<(DHA) deg
C(12A)-H(11A)O(4A)*	0.98	2.20	2.957(2)*	132.6
С(12А)-Н(11С)О(4В)	0.98	2.80	3.521(2)	131.4
C(11B)-H(11D)O(4A)	0.98	2.73	3.441(2)	130.2
C(12B)-H(11G)O(4B)*	0.98	2.25	3.001(2)*	132.3
C(8A)-H(7)O(4B)#1	0.95	2.75	3.412(2)	127.5
C(9A)-H(8)O(5B)#1	0.95	2.86	3.798(2)	168.4
C(9A)-H(8)O(4B)#1	0.95	2.84	3.452(2)	122.8
C(11A)-H(10A)O(5B)#1	0.98	3.00	3.459(2)	110.1
C(7B)-H(106)O(4B)#2	0.95	2.85	3.798(2)	176.8
C(11B)-H(11F)O(5A)#3	0.98	2.75	3.241(2)	111.7
C(9B)-H(108)O(4A)#3	0.95	2.70	3.414(2)	132.3

\*C...O distance is less than the sum of the van der Waals radii; Symmetry transformations used to generate equivalent atoms: #1 -x,-y,-z; #2 -x,y+1/2,-z+1/2; #3 -x+1,-y,-z+1



**Figure 7a.** 2,3-Diphenyl-5-methyl-1-benzofuran  $C_{21}H_{16}O_1$  (**II**)<sup>6</sup> determined by X-ray analysis (Drawn with Visualizer, Discovery Studio, Release 3.5, Accelrys Inc.,2012)<sup>3</sup>

[http://accelrys.com/products/datasheets/whats-newin-discovery-studio.pdf].

**Table 5.** Bond lengths [Å] in the chain C(4)-O(5)-C(1)-C(6) in 1-benzofuran-2-one (I) molecules A and B and the corresponding values in molecules (II) and (III).

Bond	(I)A	(I)B	(II)	(III)
C(4)-O(5)	1.388(2)	1.395(2)	1.398(4)	1.381(3)
0(5)-C(1)	1.391(2)	1.396(2)	1.380(4)	1.398(3)
C(1) C(6)	1.369(2)	1.367(2)	1.378(4)	1.362(3)
C(6)-C(7)	1.393(2)	1.398(2)	1.386(4)	1.387(3)



**Figure 7b.** 3-(1-Phenylmeth-(E)-ylidene)-3*H*benzofuran-2-one *C*<sub>21</sub>*H*<sub>16</sub>*O*<sub>1</sub> (**III**)<sup>7</sup> determined in the X-ray analysis (Drawn with Visualizer, Discovery Studio, Release 3.5, Accelrys Inc., 2012)<sup>3</sup>

[http://accelrys.com/products/datasheets/whats-newin-discovery-studio.pdf].

#### **Supplementary Material**

Crystallographic data for the structure reported in this paper have been deposited with the Cambridge Crystallographic Data Centre as supplementary publication no. CCDC 804780. Copies of available material

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