



Crystal structure, Hirshfeld surface analysis and density functional theory study of 6-methyl-2-[(5-methylisoxazol-3-yl)methyl]-1*H*-benzimidazole

Ahlam Idrissi,^a Karim Chkirate,^a Nadeem Abad,^{b*} Bahia Djerrari,^a Redouane Achour,^a El Mokhtar Essassi^a and Luc Van Meervelt^c

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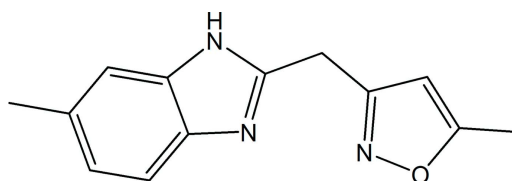
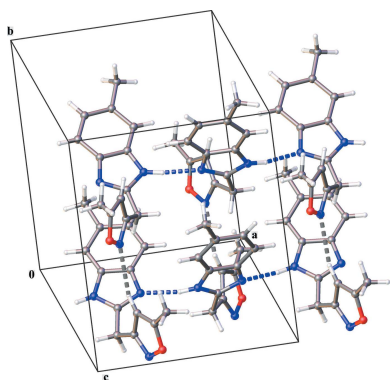
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^aLaboratory of Heterocyclic Organic Chemistry URAC 21, Pharmacochemistry Competence Center, Av. Ibn Battouta, BP 1014, Faculty of Sciences, Mohammed V University, Rabat, Morocco, ^bDepartment of Biochemistry, Faculty of Education & Science, Al-Baydha University, Yemen, and ^cKU Leuven, Chemistry Department, Celestijnenlaan 200F box 2404, Leuven, (Heverlee), B-3001, Belgium. *Correspondence e-mail: abadnadeem3@gmail.com

In the title molecule, C₁₃H₁₃N₃O, the isoxazole ring is inclined to the benzimidazole ring at a dihedral angle of 69.28 (14)°. In the crystal, N—H···N hydrogen bonds between neighboring benzimidazole rings form chains along the *a*-axis direction. Hirshfeld surface analysis indicates that the most important contributions to the crystal packing are from H···H (48.8%), H···C/C···H (20.9%) and H···N/N···H (19.3%) interactions. The optimized structure calculated using density functional theory at the B3LYP/6–311 G(d,p) level is compared with the experimentally determined structure in the solid state. The calculated highest occupied molecular orbital (HOMO) and lowest unoccupied molecular orbital (LUMO) energy gap is 4.9266 eV.

1. Chemical context

Nitrogen-based structures have attracted increased attention in structural and inorganic chemistry in recent years because of their interesting properties (Lahmidi *et al.*, 2018; Chkirate *et al.*, 2020a; Taia *et al.*, 2020; Al Ati *et al.*, 2021). The benzimidazole family, particularly compounds containing the 2-methyl benzimidazole moiety, is important in medicinal chemistry because of their wide range of pharmacological applications including as antimicrobial and antitubercular agents (Ranjith *et al.*, 2013), potential urease enzyme inhibitors (Menteşe *et al.*, 2019) and antibacterial agents (Chkirate *et al.*, 2020b). In particular, isoxazolyl benzimidazole derivatives are used as analgesic and anti-inflammatory agents (Kankala *et al.*, 2013). They are also potent and orally bioavailable bromodomain BET inhibitors (Sperandio *et al.*, 2019). Given the wide range of therapeutic applications for such compounds, and in a continuation of the work already carried out on the synthesis of compounds resulting from 1,5-benzodiazepine (Chkirate *et al.*, 2001, 2018, 2019a,b,c, 2021), a similar approach gave the title compound, 6-methyl-2-[(5-methylisoxazol-3-yl)methyl]-1*H*-benzimidazole C₁₃H₁₃N₃O (I).



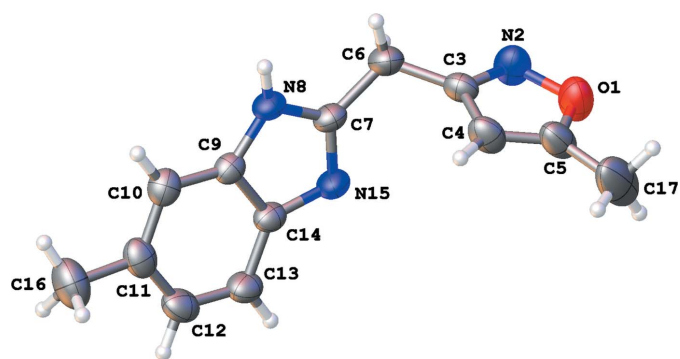


Figure 1
Molecular structure of the title molecule with the atom labeling scheme and 50% probability ellipsoids.

Besides the synthesis, we also report the molecular and crystal structures along with the results of a Hirshfeld surface analysis and density functional theory computational calculations carried out at the B3LYP/6-311 G(d,p) level.

2. Structural commentary

The title compound crystallizes in the orthorhombic space group *Pbca* with one molecule in the asymmetric unit (Fig. 1). The molecule is not planar, as indicated by the torsion angles C4–C3–C6–C7 [$-40.4(4)^\circ$] and C3–C6–C7–N15 [$-46.0(4)^\circ$]. The best plane of the isoxazole ring (O1/N2/C3–C5; r.m.s. deviation = 0.003 Å) makes a dihedral angle of $69.28(14)^\circ$ with the best plane of the benzimidazole ring (C7/N8/C9–C14/N15; r.m.s. deviation = 0.015 Å). Both methyl groups are in the same plane as the ring to which they are

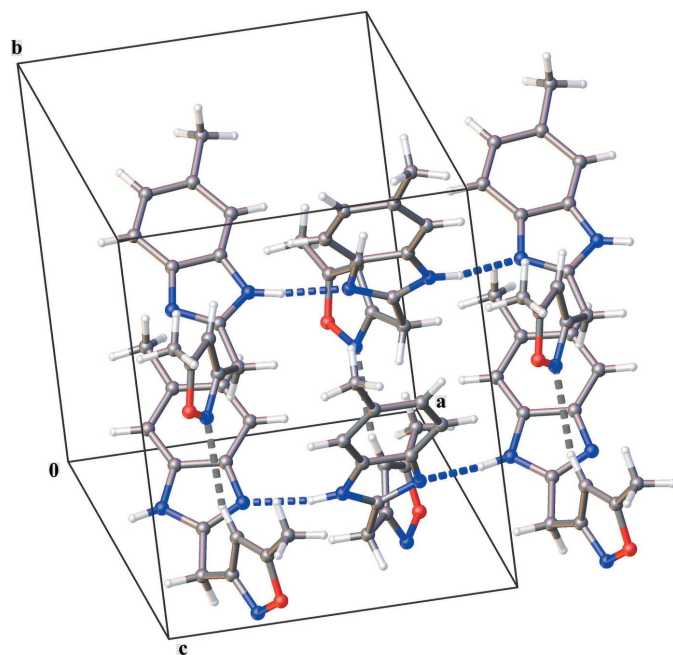


Figure 2
Partial crystal packing of the title compound. N–H...N hydrogen bonds are shown by blue dashed lines and C–H...N hydrogen bonds by gray dashed lines.

Table 1
Hydrogen-bond geometry (Å, °).

<i>D</i> –H... <i>A</i>	<i>D</i> –H	H... <i>A</i>	<i>D</i> ... <i>A</i>	<i>D</i> –H... <i>A</i>
N8–H8...N15 ⁱ	0.89 (3)	1.96 (3)	2.830 (3)	167 (2)
C4–H4...N2 ⁱⁱ	0.93	2.57	3.447 (3)	157

Symmetry codes: (i) $x - \frac{1}{2}, y, -z + \frac{3}{2}$; (ii) $-x + \frac{3}{2}, y + \frac{1}{2}, z$.

attached [deviation of C17 from the isoxazole plane = 0.016 (6) Å, deviation of C16 from the benzimidazole ring = 0.067 (4) Å].

3. Supramolecular features

The crystal packing is characterized by N–H...N and C–H...N interactions (Fig. 2, Table 1). Chains of molecules running in the *a*-axis direction are formed by N8–H8...N15ⁱ hydrogen bonds between neighboring benzimidazole rings [symmetry code: (i) $-\frac{1}{2} + x, y, 3/2 - z$]. Parallel chains interact through C4–H4...N2ⁱⁱ hydrogen bonds between neighboring isoxazole rings [symmetry code: (ii) $3/2 - x, \frac{1}{2} + y, z$] resulting in the three-dimensional structure. The crystal packing contains no voids.

4. Hirshfeld surface analysis

The *CrystalExplorer* program (Turner *et al.*, 2017) was used to investigate and visualize the intermolecular interactions of (I). The Hirshfeld surface plotted over d_{norm} in the range -0.6149 to 1.3177 a.u. is shown in Fig. 3*a*. The red spots are close

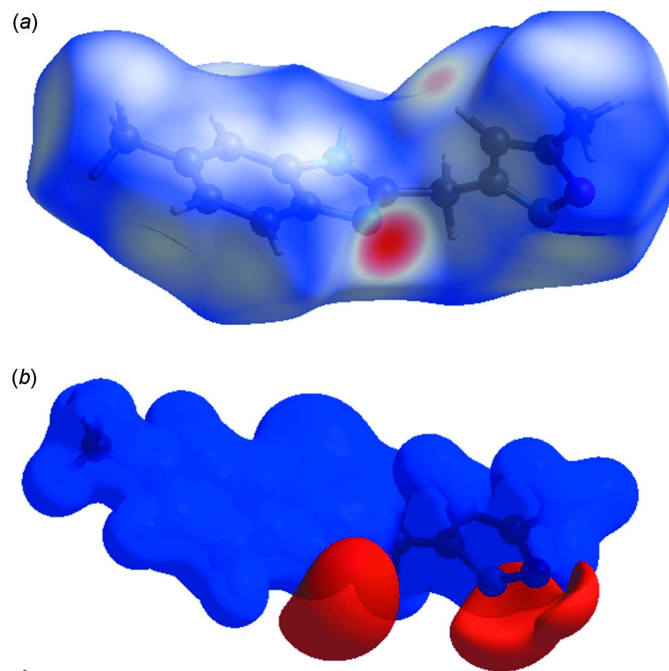


Figure 3
(*a*) View of the three-dimensional Hirshfeld surface of the title compound, plotted over d_{norm} in the range -0.6149 to 1.3177 a.u. (*b*) View of the three-dimensional Hirshfeld surface of the title compound plotted over electrostatic potential energy in the range -0.0500 to 0.0500 a.u. using the STO-3 G basis set at the Hartree–Fock level of theory.

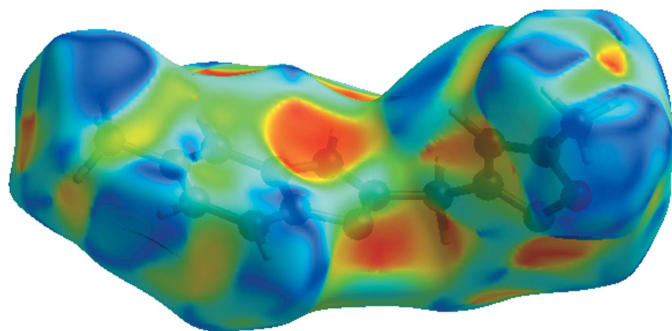


Figure 4
Hirshfeld surface of the title compound plotted over shape-index.

contacts with a negative d_{norm} value and represent N—H···N and C—H···N interactions. The white regions representing contacts equal to the van der Waals separation and a d_{norm} value of zero are indicative of the H···H interactions. The electrostatic potential using the STO-3G basis set at the Hartree–Fock level of theory and mapped on the Hirshfeld surface over the range ± 0.05 a.u. clearly shows the positions of close intermolecular contacts in the compound (Fig. 3*b*). The positive electrostatic potential (blue region) over the surface indicates hydrogen-donor potential, whereas the hydrogen-bond acceptors are represented by negative electrostatic potential (red region). The shape-index (Fig. 4) generated in the ranges -1 to 1 Å reveals that there are no significant π – π interactions (normally indicated by adjacent red and blue triangles).

The overall two-dimensional fingerprint plot (McKinnon *et al.*, 2007) is shown in Fig. 5*a*, while those delineated into

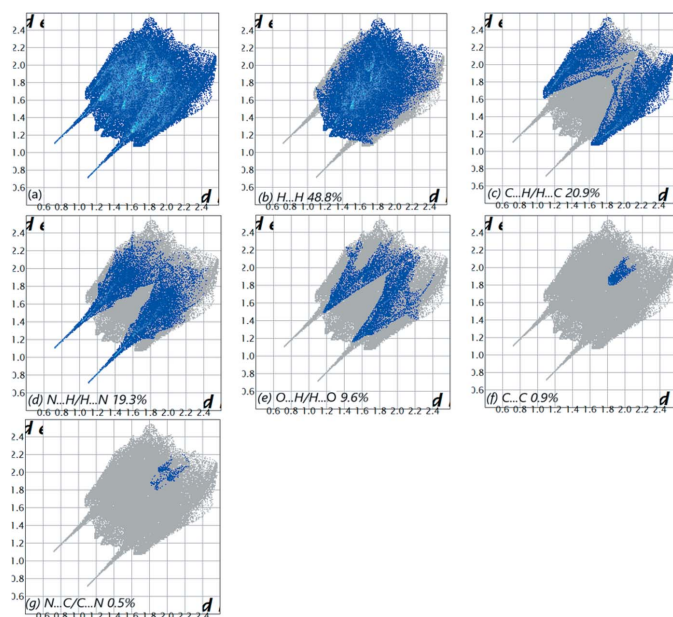


Figure 5
The full two-dimensional fingerprint plots for the title compound, showing (a) all interactions, and delineated into (b) H···H, (c) H···C/C···H, (d) H···N/N···H, (e) H···O/O···H, (f) C···C and (g) C···N/N···C interactions. The d_i and d_e values are the closest internal and external distances (in Å) from given points on the Hirshfeld surface.

H···H, H···C/C···H, H···N/N···H, H···O/O···H, C···C and C···N/N···C contacts are illustrated in Fig. 5*b–g*, respectively, together with their relative contributions to the Hirshfeld surface (HS). The most important interaction is H···H, contributing 48.8% to the overall crystal packing, which is reflected in Fig. 5*b* as widely scattered points of high density due to the large hydrogen content of the molecule, with the tip at $d_e = d_i = 1.28$ Å. In the presence of C—H interactions, the pair of characteristic wings in the fingerprint plot delineated into H···C/C···H contacts (20.9% contribution to the HS), Fig. 5*c*, has the tips at $d_e + d_i = 2.69$ Å. The pair of scattered points of spikes in the fingerprint plot delineated into H···N/N···H, Fig. 5*d* (19.3%), have the tips at $d_e + d_i = 1.81$ Å. The H···O/O···H contacts, Fig. 5*e* (9.6%), have the tips at $d_e + d_i = 2.65$ Å. The C···C contacts, Fig. 5*f*, contribute 0.9% to the HS and appear as a pair of scattered points of spikes with the tips at $d_e + d_i = 3.60$ Å. Finally, the C···N/N···C contacts, Fig. 5*g*, make only a 0.5% contribution to the HS and have a low-density distribution of points.

5. Density Functional Theory calculations

The structure in the gas phase of the title compound was optimized by means of density functional theory. The density functional theory calculation was performed by the hybrid B3LYP method and the 6–311 G(d,p) basis-set, which is based on Becke's model (Becke, 1993) and considers a mixture of the exact (Hartree–Fock) and density functional theory exchange utilizing the B3 functional, together with the LYP correlation functional (Lee *et al.*, 1988). After obtaining the converged geometry, the harmonic vibrational frequencies were calculated at the same theoretical level to confirm that the number of imaginary frequencies is zero for the stationary point. Both the geometry optimization and harmonic vibrational frequency analysis of the title compound were performed with the GAUSSIAN 09 program (Frisch *et al.*, 2009). The theoretical and experimental results related to bond lengths and angles are in good agreement, as well as with the results of the previous structural study of 5,6-dimethyl-2-[(5-methyl-1,2-oxazol-3-yl)methyl]-1-(prop-2-en-1-yl)-1*H*-benzimidazole, (III) (Benyahya *et al.*, 2017) and 5-methyl-3-(1-(2-pyridylmethyl)-1*H*-benzimidazol-2-ylmethyl)isoxazole, (IV) (Doubmia *et al.*, 2009), which are summarized in Table 2. Calculated numerical values for title compound including electronegativity (χ), hardness (η), ionization potential (I), dipole moment (μ), electron affinity (A), electrophilicity (ω) and softness (σ) are collated in Table 3. The electron transition from the highest occupied molecular orbital (HOMO) to the lowest unoccupied molecular orbital (LUMO) energy level is shown in Fig. 6. The HOMO and LUMO are localized in the plane extending over the whole 6-methyl-2-[(5-methyl-isoxazol-3-yl)methyl]-1*H*-benzimidazole system. The energy band gap [$\Delta E = E_{\text{LUMO}} - E_{\text{HOMO}}$] of the molecule is 4.9266 eV, and the frontier molecular orbital energies, E_{HOMO} and E_{LUMO} , are -5.8170 and -0.8904 eV, respectively.

Table 2

Comparison of selected (X-ray and DFT bond lengths and angles (Å, °) in the title compound and related structures.

	X-ray	B3LYP/6-311G(d,p)	(III) ^a	(IV) ^b
O1—N2	1.413 (3)	1.3949	1.417	1.4100
O1—C5	1.339 (4)	1.3481	1.356	1.3526
N2—C3	1.299 (3)	1.3115	1.304	1.3044
C3—C6	1.488 (4)	1.5065	1.501	1.504
C5—C17	1.485 (4)	1.4868	1.476	1.478
C6—C7	1.488 (4)	1.5026	1.498	1.494
C7—N8	1.349 (3)	1.3755	1.377	1.3720
C7—N15	1.320 (3)	1.3092	1.312	1.3079
N8—C9	1.371 (3)	1.3814	1.386	1.3840
C11—C16	1.500 (4)	1.5112	1.504	—
C14—N15	1.391 (3)	1.388	1.400	1.3880
C5—O1—N2	108.2 (2)	109.1398	108.37	108.57
C3—N2—O1	105.5 (2)	106.0707	105.15	105.28
N2—C3—C4	111.3 (2)	111.0906	112.00	111.51
N2—C3—C6	118.9 (2)	120.8172	120.16	119.88
O1—C5—C17	117.0 (3)	116.8621	116.33	115.90
C4—C5—O1	109.5 (3)	109.3513	109.34	109.15
N8—C7—C6	121.7 (2)	122.8089	123.02	122.62
N15—C7—C6	125.6 (2)	123.8733	123.28	124.10
N15—C7—N8	112.7 (2)	113.2373	113.69	113.28
C7—N8—C9	107.59 (19)	106.9514	106.09	106.49
N8—C9—C14	105.29 (19)	104.6015	105.63	105.05
C13—C14—N15	130.8 (2)	130.4265	129.98	129.63
N15—C14—C9	109.67 (19)	110.2891	110.23	110.42
C7—N15—C14	104.72 (19)	104.9141	104.36	104.75

Notes: (a) Results of the previous DFT-optimized geometry of 5,6-dimethyl-2-[(5-methyl-1,2-oxazol-3-yl)methyl]-1-(prop-2-en-1-yl)-1*H*-benzimidazole (Benyahya *et al.*, 2017); (b) results of the previous crystallographic study of 5-methyl-3-(1-(2-pyridylmethyl)-1*H*-benzimidazol-2-ylmethyl)isoxazole (Dombia *et al.*, 2009)

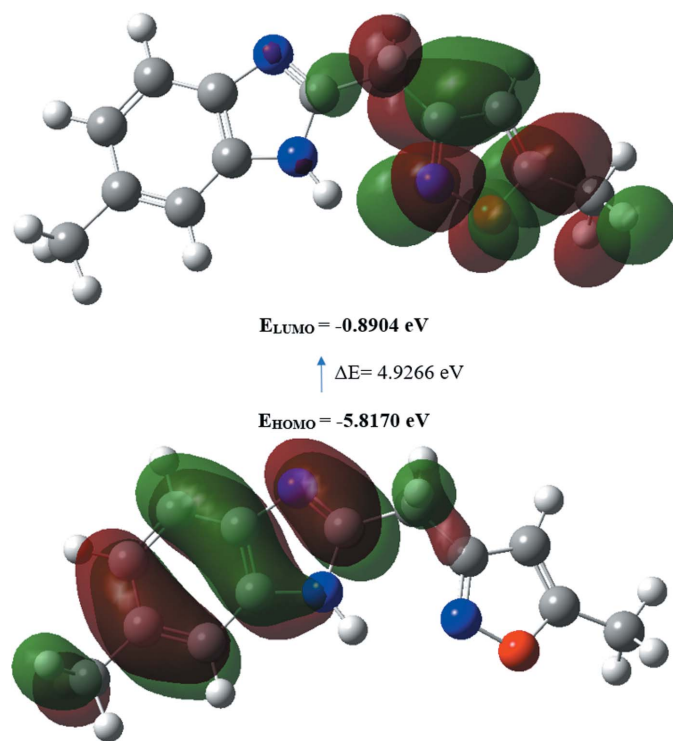


Figure 6

The energy band gap of 6-methyl-2-[(5-methylisoxazol-3-yl)methyl]-1*H*-benzimidazole.

Table 3

Calculated energies.

Molecular Energy	Title Compound
Total Energy <i>TE</i> (eV)	−20214.1624
<i>E</i> _{HOMO} (eV)	−5.8170
<i>E</i> _{LUMO} (eV)	−0.8904
Gap, Δ <i>E</i> (eV)	4.9266
Dipole moment, μ (Debye)	4.4403
Ionization potential, <i>I</i> (eV)	5.8170
Electron affinity, <i>A</i>	0.8904
Electronegativity, χ	3.3537
Hardness, η	2.4633
Electrophilicity, index ω	2.2830
Softness, σ	0.4060
Fraction of electron transferred, Δ <i>N</i>	0.7401

6. Database survey

A search of the Cambridge Structural Database (CSD version 5.40, updated March 2020; Groom *et al.*, 2016) with the 2-methylbenzimidazole fragment yielded multiple matches. Of these, three had an isoxazol-3-yl substituent comparable to (I) and they are shown in Fig. 7. The first compound (II) (refcode REQZIW; Attar *et al.*, 2001) has no substituent on the phenyl ring. For the second one (III) (refcode FECPIP; Benyahya *et al.*, 2017) the phenyl ring is disubstituted with an allyl substituent on nitrogen 1. The third one (IV) (refcode PUGLAF; Dombia *et al.*, 2009) carries pyridin-2-ylmethyl on nitrogen 1. The benzimidazole and isoxazole moieties are planar and make a dihedral angle of 76,15 (4)° in REQZIW. In FECPIP, the benzimidazole moiety is slightly non-planar, as indicated by the dihedral angle of 1.3 (1)° between the five- and six-membered rings. The isoxazole ring is planar to within 0.005 (1) Å and makes a dihedral angle of 89.78 (8)° with the benzimidazole ring. On the other hand, in PUGLAF, the fused-ring system is essentially planar, with a maximum deviation of 0.019 (1) Å. It forms interplanar angles of 70.03 (7)° with the isoxazole ring and 81.68 (7)° with the pyridine ring. The two latter rings are also planar, the maximum deviations from the mean planes being 0.0028 (15)

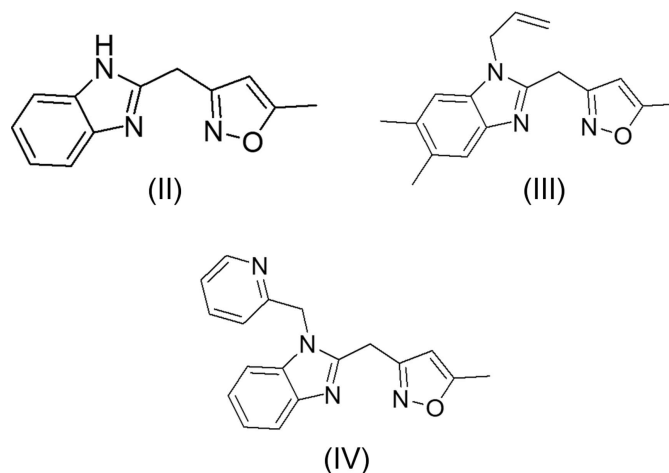


Figure 7

Structural fragments (II), (III) and (IV) used in the database survey.

and 0.0047 (12) Å. In (I), The isoxazole ring is inclined to the mean plane of the benzimidazole ring by 69.28 (14)° which is approximately the same as in PUGLAF, but less tilted than in REQZIW and FECPIP.

7. Synthesis and crystallization

(Z)-7-Methyl-4-(2-oxopropylidene)-1,5-benzodiazepin-2-one (2.3 g, 0.01 mol) and hydroxylamine hydrochloride (0.7 g, 0.01 mol) were brought to reflux in 40 ml of methanol for 2 h. After neutralization with NaHCO₃, the compound that precipitated was filtered and recrystallized from ethyl acetate. The product was dissolved to saturation in ethyl acetate and crystals were obtained by evaporation at room temperature. yield: 70%; m.p. 465–467 K; IR [KBr, $\nu(\text{cm}^{-1})$]: ν_{NH} = 3416; ν_{CH} = 3012–3263; $\nu_{\text{C=N-C=C}}$ = 1525–1672; ¹H NMR [300MHz, DMSO-*d*₆, $\delta(\text{ppm})$]: 2.32 (s, 3H, CH₃ isoxazole); 2.57 (s, 3H, CH₃ benzimidazole); 4.23 (s, 2H, CH₂); 6.22 (s, 1H, CH isoxazole); 7.00–7.60 (m, 3H, CH_{ar}); 5.0 (s, 1H, NH). ¹³C NMR [75MHz, DMSO-*d*₆, $\delta(\text{ppm})$]: 13.2 (CH₃ isoxazole); 24.3 (CH₃ benzimidazole); 26.7 (CH₂); 101.8 (CH isoxazole); 115.2–125.8 (CH aryl); 132.7–169.6 (C quaternary).

8. Refinement

Crystal data, data collection and structure details refinement are given in Table 4. Hydrogen atoms were located in the first difference-Fourier map. C-bound H atoms were positioned geometrically (C–H = 0.93–0.97 Å) and included as riding contributions with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$ (1.5 for methyl groups). At the end of the refinement, the final difference Fourier map showed no residual peaks of chemical significance.

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Table 4

Experimental details.

Crystal data	
Chemical formula	C ₁₃ H ₁₃ N ₃ O
<i>M_r</i>	227.26
Crystal system, space group	Orthorhombic, <i>Pbca</i>
Temperature (K)	294
<i>a</i> , <i>b</i> , <i>c</i> (Å)	9.6545 (6), 11.2437 (6), 22.9108 (14)
<i>V</i> (Å ³)	2487.0 (3)
<i>Z</i>	8
Radiation type	Mo <i>K</i> α
μ (mm ⁻¹)	0.08
Crystal size (mm)	0.35 × 0.2 × 0.2
Data collection	
Diffractometer	Rigaku Oxford Diffraction Super-Nova, Single source at offset/far, Eos
Absorption correction	Multi-scan (<i>CrysAlis PRO</i> ; Rigaku OD, 2018)
<i>T_{min}</i> , <i>T_{max}</i>	0.883, 1.000
No. of measured, independent and observed [<i>I</i> > 2σ(<i>I</i>)] reflections	13352, 2519, 1723
<i>R_{int}</i>	0.024
(<i>sin</i> θ/ <i>λ</i>) _{max} (Å ⁻¹)	0.625
Refinement	
<i>R</i> [<i>F</i> ² > 2σ(<i>F</i> ²)], <i>wR</i> (<i>F</i> ²), <i>S</i>	0.064, 0.203, 1.05
No. of reflections	2519
No. of parameters	160
H-atom treatment	H atoms treated by a mixture of independent and constrained refinement
$\Delta\rho_{\text{max}}$, $\Delta\rho_{\text{min}}$ (e Å ⁻³)	0.33, −0.26

Computer programs: *CrysAlis PRO* (Rigaku OD, 2018), *SHELXT* (Sheldrick, 2015a), *SHELXL2016/4* (Sheldrick, 2015b) and *OLEX2* (Dolomanov *et al.*, 2009).

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supporting information

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Crystal structure, Hirshfeld surface analysis and density functional theory study of 6-methyl-2-[(5-methylisoxazol-3-yl)methyl]-1*H*-benzimidazole

Ahlam Idrissi, Karim Chkirate, Nadeem Abad, Bahia Djerrari, Redouane Achour, El Mokhtar Essassi and Luc Van Meervelt

Computing details

Data collection: *CrysAlis PRO* (Rigaku OD, 2018); cell refinement: *CrysAlis PRO* (Rigaku OD, 2018); data reduction: *CrysAlis PRO* (Rigaku OD, 2018); program(s) used to solve structure: SHELXT (Sheldrick, 2015a); program(s) used to refine structure: *SHELXL2016/4* (Sheldrick, 2015b); molecular graphics: *OLEX2* (Dolomanov *et al.*, 2009); software used to prepare material for publication: *OLEX2* (Dolomanov *et al.*, 2009).

6-Methyl-2-[(5-methylisoxazol-3-yl)methyl]-1*H*-benzimidazole

Crystal data

C₁₃H₁₃N₃O

M_r = 227.26

Orthorhombic, *Pbca*

a = 9.6545 (6) Å

b = 11.2437 (6) Å

c = 22.9108 (14) Å

V = 2487.0 (3) Å³

Z = 8

F(000) = 960

D_x = 1.214 Mg m⁻³

Mo *Kα* radiation, λ = 0.71073 Å

Cell parameters from 3531 reflections

θ = 2.9–23.3°

μ = 0.08 mm⁻¹

T = 294 K

Prism, brown

0.35 × 0.2 × 0.2 mm

Data collection

Rigaku Oxford Diffraction SuperNova, Single source at offset/far, Eos diffractometer

Radiation source: micro-focus sealed X-ray tube, SuperNova (Mo) X-ray Source

Mirror monochromator

Detector resolution: 15.9631 pixels mm⁻¹

ω scans

Absorption correction: multi-scan (CrysAlisPro; Rigaku OD, 2018)

T_{min} = 0.883, *T_{max}* = 1.000

13352 measured reflections

2519 independent reflections

1723 reflections with *I* > 2σ(*I*)

R_{int} = 0.024

θ_{max} = 26.4°, θ_{min} = 2.8°

h = -10→12

k = -14→13

l = -28→28

Refinement

Refinement on *F*²

Least-squares matrix: full

R [*F*² > 2σ(*F*²)] = 0.064

wR (*F*²) = 0.203

S = 1.05

2519 reflections

160 parameters

0 restraints

Primary atom site location: dual

Hydrogen site location: mixed

H atoms treated by a mixture of independent and constrained refinement

w = 1/[σ²(*F_o*²) + (0.0963*P*)² + 0.6658*P*]

where *P* = (*F_o*² + 2*F_c*²)/3

$$(\Delta/\sigma)_{\max} < 0.001$$

$$\Delta\rho_{\max} = 0.33 \text{ e } \text{\AA}^{-3}$$

$$\Delta\rho_{\min} = -0.26 \text{ e } \text{\AA}^{-3}$$

Special details

Geometry. All esds (except the esd in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell esds are taken into account individually in the estimation of esds in distances, angles and torsion angles; correlations between esds in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell esds is used for estimating esds involving l.s. planes.

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (\AA^2)

	x	y	z	$U_{\text{iso}}^*/U_{\text{eq}}$
O1	0.8450 (3)	0.0193 (2)	0.57078 (9)	0.1116 (8)
N2	0.7873 (3)	-0.0264 (2)	0.62282 (11)	0.0916 (8)
C3	0.7336 (2)	0.0646 (2)	0.64943 (11)	0.0649 (6)
C4	0.7542 (3)	0.1693 (2)	0.61779 (12)	0.0852 (9)
H4	0.726197	0.245615	0.628050	0.102*
C5	0.8227 (4)	0.1368 (3)	0.56964 (12)	0.0962 (10)
C6	0.6621 (3)	0.0459 (3)	0.70624 (12)	0.0802 (8)
H6A	0.562991	0.042286	0.699489	0.096*
H6B	0.690721	-0.030027	0.722308	0.096*
C7	0.6914 (2)	0.1410 (2)	0.74970 (10)	0.0628 (6)
N8	0.5894 (2)	0.19185 (18)	0.78133 (8)	0.0616 (5)
C9	0.6490 (2)	0.2765 (2)	0.81631 (9)	0.0578 (6)
C10	0.5943 (3)	0.3565 (2)	0.85619 (10)	0.0727 (7)
H10	0.499999	0.357606	0.864518	0.087*
C11	0.6850 (3)	0.4350 (3)	0.88328 (11)	0.0815 (8)
C12	0.8243 (3)	0.4291 (3)	0.87071 (12)	0.0855 (9)
H12	0.883742	0.481816	0.889438	0.103*
C13	0.8799 (3)	0.3491 (3)	0.83190 (11)	0.0785 (8)
H13	0.974628	0.347148	0.824563	0.094*
C14	0.7902 (2)	0.2713 (2)	0.80401 (10)	0.0615 (6)
N15	0.81476 (19)	0.18510 (19)	0.76175 (9)	0.0678 (6)
C16	0.6344 (5)	0.5270 (4)	0.92565 (15)	0.1269 (13)
H16A	0.638258	0.604086	0.907763	0.190*
H16B	0.691872	0.526214	0.959831	0.190*
H16C	0.540500	0.509387	0.936487	0.190*
C17	0.8775 (7)	0.2031 (4)	0.51859 (15)	0.174 (2)
H17A	0.975882	0.191938	0.516094	0.261*
H17B	0.857462	0.286270	0.523070	0.261*
H17C	0.834513	0.174111	0.483589	0.261*
H8	0.501 (3)	0.182 (2)	0.7723 (11)	0.079 (8)*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
O1	0.153 (2)	0.0868 (15)	0.0948 (15)	-0.0016 (13)	0.0324 (14)	-0.0116 (12)
N2	0.118 (2)	0.0609 (13)	0.0961 (16)	0.0068 (12)	0.0172 (14)	0.0016 (12)
C3	0.0657 (15)	0.0504 (12)	0.0785 (15)	-0.0017 (10)	-0.0074 (12)	-0.0070 (11)

C4	0.121 (2)	0.0538 (14)	0.0809 (18)	-0.0009 (14)	-0.0090 (17)	-0.0022 (12)
C5	0.150 (3)	0.0717 (18)	0.0673 (17)	-0.0246 (18)	-0.0082 (17)	-0.0019 (14)
C6	0.0720 (17)	0.0801 (18)	0.0885 (18)	-0.0208 (13)	0.0074 (13)	-0.0092 (14)
C7	0.0463 (13)	0.0698 (14)	0.0724 (14)	-0.0068 (11)	0.0015 (10)	0.0054 (11)
N8	0.0404 (11)	0.0732 (13)	0.0711 (12)	-0.0031 (9)	-0.0037 (9)	0.0051 (10)
C9	0.0494 (12)	0.0671 (14)	0.0569 (12)	0.0038 (10)	-0.0035 (10)	0.0105 (10)
C10	0.0704 (16)	0.0799 (17)	0.0678 (14)	0.0137 (13)	-0.0001 (12)	0.0093 (13)
C11	0.103 (2)	0.0806 (18)	0.0608 (15)	0.0105 (16)	-0.0089 (14)	0.0014 (12)
C12	0.094 (2)	0.093 (2)	0.0687 (16)	-0.0185 (16)	-0.0220 (15)	0.0013 (14)
C13	0.0649 (16)	0.099 (2)	0.0717 (15)	-0.0169 (14)	-0.0119 (13)	0.0017 (15)
C14	0.0528 (13)	0.0738 (15)	0.0580 (12)	-0.0059 (10)	-0.0080 (10)	0.0101 (11)
N15	0.0453 (11)	0.0817 (14)	0.0763 (13)	-0.0062 (9)	0.0034 (9)	-0.0031 (10)
C16	0.157 (3)	0.126 (3)	0.097 (2)	0.025 (3)	-0.003 (2)	-0.029 (2)
C17	0.311 (7)	0.140 (3)	0.072 (2)	-0.080 (4)	0.008 (3)	0.011 (2)

Geometric parameters (Å, °)

O1—N2	1.413 (3)	C9—C14	1.393 (3)
O1—C5	1.339 (4)	C10—H10	0.9300
N2—C3	1.299 (3)	C10—C11	1.390 (4)
C3—C4	1.396 (4)	C11—C12	1.377 (4)
C3—C6	1.488 (4)	C11—C16	1.500 (4)
C4—H4	0.9300	C12—H12	0.9300
C4—C5	1.337 (4)	C12—C13	1.374 (4)
C5—C17	1.485 (4)	C13—H13	0.9300
C6—H6A	0.9700	C13—C14	1.386 (3)
C6—H6B	0.9700	C14—N15	1.391 (3)
C6—C7	1.488 (4)	C16—H16A	0.9600
C7—N8	1.349 (3)	C16—H16B	0.9600
C7—N15	1.320 (3)	C16—H16C	0.9600
N8—C9	1.371 (3)	C17—H17A	0.9600
N8—H8	0.88 (3)	C17—H17B	0.9600
C9—C10	1.387 (3)	C17—H17C	0.9600
C5—O1—N2	108.2 (2)	C9—C10—C11	117.8 (3)
C3—N2—O1	105.5 (2)	C11—C10—H10	121.1
N2—C3—C4	111.3 (2)	C10—C11—C16	121.4 (3)
N2—C3—C6	118.9 (2)	C12—C11—C10	119.4 (3)
C4—C3—C6	129.7 (2)	C12—C11—C16	119.2 (3)
C3—C4—H4	127.2	C11—C12—H12	118.4
C5—C4—C3	105.6 (3)	C13—C12—C11	123.3 (3)
C5—C4—H4	127.2	C13—C12—H12	118.4
O1—C5—C17	117.0 (3)	C12—C13—H13	121.1
C4—C5—O1	109.5 (3)	C12—C13—C14	117.8 (3)
C4—C5—C17	133.5 (3)	C14—C13—H13	121.1
C3—C6—H6A	108.9	C13—C14—C9	119.5 (2)
C3—C6—H6B	108.9	C13—C14—N15	130.8 (2)
H6A—C6—H6B	107.7	N15—C14—C9	109.67 (19)

C7—C6—C3	113.3 (2)	C7—N15—C14	104.72 (19)
C7—C6—H6A	108.9	C11—C16—H16A	109.5
C7—C6—H6B	108.9	C11—C16—H16B	109.5
N8—C7—C6	121.7 (2)	C11—C16—H16C	109.5
N15—C7—C6	125.6 (2)	H16A—C16—H16B	109.5
N15—C7—N8	112.7 (2)	H16A—C16—H16C	109.5
C7—N8—C9	107.59 (19)	H16B—C16—H16C	109.5
C7—N8—H8	121.4 (17)	C5—C17—H17A	109.5
C9—N8—H8	129.2 (17)	C5—C17—H17B	109.5
N8—C9—C10	132.5 (2)	C5—C17—H17C	109.5
N8—C9—C14	105.29 (19)	H17A—C17—H17B	109.5
C10—C9—C14	122.2 (2)	H17A—C17—H17C	109.5
C9—C10—H10	121.1	H17B—C17—H17C	109.5
O1—N2—C3—C4	-0.7 (3)	N8—C7—N15—C14	-0.4 (3)
O1—N2—C3—C6	179.3 (2)	N8—C9—C10—C11	-177.5 (2)
N2—O1—C5—C4	0.0 (4)	N8—C9—C14—C13	178.6 (2)
N2—O1—C5—C17	179.1 (3)	N8—C9—C14—N15	0.5 (2)
N2—C3—C4—C5	0.7 (3)	C9—C10—C11—C12	-1.3 (4)
N2—C3—C6—C7	139.7 (3)	C9—C10—C11—C16	178.2 (3)
C3—C4—C5—O1	-0.4 (4)	C9—C14—N15—C7	-0.1 (3)
C3—C4—C5—C17	-179.3 (4)	C10—C9—C14—C13	-0.6 (3)
C3—C6—C7—N8	133.7 (2)	C10—C9—C14—N15	-178.7 (2)
C3—C6—C7—N15	-46.0 (4)	C10—C11—C12—C13	0.5 (4)
C4—C3—C6—C7	-40.4 (4)	C11—C12—C13—C14	0.3 (4)
C5—O1—N2—C3	0.4 (3)	C12—C13—C14—C9	-0.3 (4)
C6—C3—C4—C5	-179.3 (3)	C12—C13—C14—N15	177.4 (2)
C6—C7—N8—C9	-179.1 (2)	C13—C14—N15—C7	-177.9 (3)
C6—C7—N15—C14	179.3 (2)	C14—C9—C10—C11	1.4 (3)
C7—N8—C9—C10	178.4 (2)	N15—C7—N8—C9	0.7 (3)
C7—N8—C9—C14	-0.7 (2)	C16—C11—C12—C13	-179.0 (3)

Hydrogen-bond geometry (Å, °)

<i>D</i> —H... <i>A</i>	<i>D</i> —H	H... <i>A</i>	<i>D</i> ... <i>A</i>	<i>D</i> —H... <i>A</i>
N8—H8...N15 ⁱ	0.89 (3)	1.96 (3)	2.830 (3)	167 (2)
C4—H4...N2 ⁱⁱ	0.93	2.57	3.447 (3)	157

Symmetry codes: (i) $x-1/2, y, -z+3/2$; (ii) $-x+3/2, y+1/2, z$.