

O(3)—H···O(8)	2.681 (4)	O(7)—H···O(4 ^v)	2.876 (4)
O(4)—H···O(7)	2.803 (4)	O(7)—H···O(6 ^{vi})	2.788 (4)
O(5)—H···O(1 ⁱⁱ)	2.576 (4)		

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

Calculations were performed using *SHELXS86* (Sheldrick, 1985).

The authors wish to thank Professor K. R. Seddon (The Queen's University of Belfast) for helpful discussions, and DRA (Fort Halstead) for generous financial support.

Lists of structure factors, anisotropic displacement parameters, H-atom coordinates and complete geometry have been deposited with the British Library Document Supply Centre as Supplementary Publication No. SUP 71713 (11 pp.). Copies may be obtained through The Technical Editor, International Union of Crystallography, 5 Abbey Square, Chester CH1 2HU, England. [CIF reference: HA1071]

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1-Methyl-5-nitro-2-phenylimidazole and 2-(*p*-Aminophenyl)-1-methyl-5-nitroimidazole

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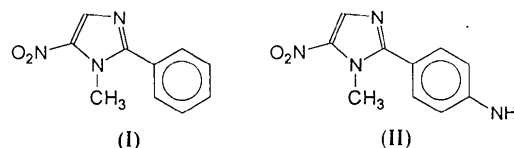
Abstract

1-Methyl-5-nitro-2-phenylimidazole (I), $C_{10}H_9N_3O_2$, and 2-(*p*-aminophenyl)-1-methyl-5-nitroimidazole (II), $C_{10}H_{10}N_4O_2$, show similar bond distances and angles.

Some differences are observed in the bond distances and angles in the phenyl rings due to the presence of the amino substituent and hydrogen bonds in (II).

Comment

Perspective views showing the atomic numbering scheme for (I) and (II) are given in Figs. 1 and 2. In both structures the imidazole and phenyl rings are planar within experimental error. The dihedral angles between the phenyl and imidazole rings in (I) and (II) are 138.58 (6) and 143.1 (1)°, respectively. The angle between the nitro group and the imidazole plane is 8.1 (2)° in (I) and 6.4 (3)° in (II).



The amino and nitro groups in (II) are involved in hydrogen bonding [N24—H241···O502ⁱ 2.11 (5) Å, 151 (3)°; N24—H242···O501ⁱⁱ 2.33 (4) Å, 167 (4)°; symmetry codes: (i) $x - 1, y, z$; (ii) $x - 1, -y + \frac{1}{2}, z - \frac{1}{2}$]. Each molecule of (II) is surrounded by four other molecules. This results in folded layers parallel to (010).

The bond lengths and bond angles in both compounds are similar, except for those in the phenyl rings. A comparison of the angles in the phenyl rings shows the influence of the amino substituent in (II). The differences found are

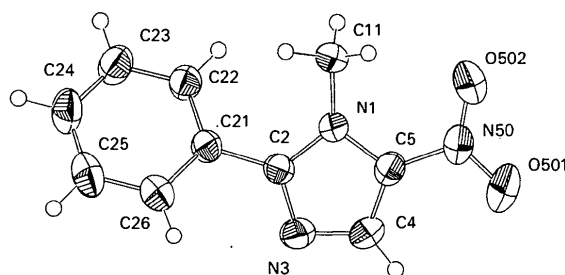


Fig. 1. A perspective view of molecule (I) with atomic numbering scheme. The displacement ellipsoids are plotted at the 50% probability level.

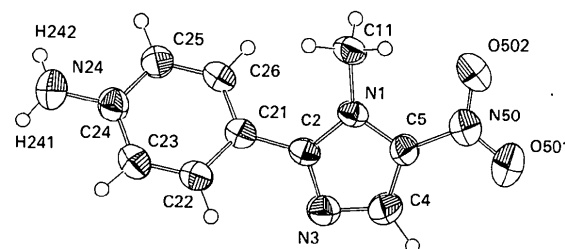


Fig. 2. A perspective view of molecule (II) with atomic numbering scheme. The displacement ellipsoids are plotted at the 50% probability level.

larger than those given by Domenicano & Murray-Rust (1979) and are caused by the presence of the hydrogen bonding in (II).

Experimental

Compound (I)

Crystal data

C₁₀H₉N₃O₂

M_r = 203.20

Monoclinic

*P*2₁/*c*

a = 11.127 (1) Å

b = 7.575 (1) Å

c = 12.046 (1) Å

β = 108.85 (1)°

V = 960.9 (2) Å³

Z = 4

D_x = 1.4046 Mg m⁻³

Data collection

Stoe Stadi-4 four-circle diffractometer

ω scans

Absorption correction: none

4660 measured reflections

2224 independent reflections

1580 observed reflections

[*I* > 2.0σ(*I*)]

Refinement

Refinement on *F*

R = 0.041

wR = 0.063

S = 1.2695

1580 reflections

172 parameters

All H-atom parameters refined

w = 1.0179/[σ²(*F*) + 0.002064*F*²]

Mo *K*α radiation

λ = 0.71069 Å

Cell parameters from 38 reflections

θ = 10.03–12.48°

μ = 0.0951 mm⁻¹

T = 293 K

Needle

0.30 × 0.10 × 0.10 mm

Colourless

*R*_{int} = 0.016

θ_{max} = 27.89°

h = -14 → 14

k = -10 → 10

l = 0 → 16

6 standard reflections

frequency: 60 min

intensity variation: 5%

(Δ/σ)_{max} = 0.003

Δρ_{max} = 0.22 e Å⁻³

Δρ_{min} = -0.21 e Å⁻³

Extinction correction: none

Atomic scattering factors from *CRYSRULER* (Rizzoli, Sangermano, Calestani & Andreetti, 1989)

Table 2. Selected geometric parameters (Å, °) for (I)

N1—C11	1.465 (1)	C23—C24	1.380 (3)
N1—C2	1.357 (1)	C24—C25	1.378 (3)
N1—C5	1.372 (1)	C25—C26	1.385 (2)
C2—C21	1.472 (1)	N3—C4	1.352 (2)
C2—N3	1.334 (1)	C4—C5	1.358 (2)
C21—C22	1.392 (1)	C5—N50	1.413 (2)
C21—C26	1.395 (2)	N50—O501	1.219 (2)
C22—C23	1.385 (2)	N50—O502	1.223 (3)
C2—N1—C5	105.0 (1)	C23—C24—C25	120.2 (2)
C11—N1—C5	128.1 (1)	C24—C25—C26	120.4 (2)
C11—N1—C2	126.2 (1)	C21—C26—C25	119.9 (2)
N1—C2—N3	112.2 (1)	C2—N3—C4	105.4 (1)
N1—C2—C21	124.8 (1)	N3—C4—C5	109.8 (1)
C21—C2—N3	123.0 (1)	N1—C5—C4	107.7 (1)
C2—C21—C26	119.0 (1)	C4—C5—N50	128.0 (1)
C2—C21—C22	121.7 (1)	N1—C5—N50	124.2 (1)
C22—C21—C26	119.1 (1)	C5—N50—O502	119.4 (1)
C21—C22—C23	120.4 (1)	C5—N50—O501	117.0 (2)
C22—C23—C24	120.0 (2)	O501—N50—O502	123.5 (2)

Compound (II)

Crystal data

C₁₀H₁₀N₄O₂

M_r = 218.21

Monoclinic

*P*2₁/*c*

a = 11.864 (1) Å

b = 7.434 (1) Å

c = 12.216 (2) Å

β = 110.05 (1)°

V = 1012.1 (2) Å³

Z = 4

D_x = 1.4321 Mg m⁻³

Mo *K*α radiation

λ = 0.71069 Å

Cell parameters from 47 reflections

θ = 10.07–12.94°

μ = 0.0978 mm⁻¹

T = 293 K

Needle

0.30 × 0.10 × 0.10 mm

Orange

Data collection

Stoe Stadi-4 four-circle diffractometer

ω scans

Absorption correction: none

2686 measured reflections

2334 independent reflections

790 observed reflections

[*I* > 2.0σ(*I*)]

*R*_{int} = 0.018

θ_{max} = 27.49°

h = -15 → 15

k = 0 → 10

l = 0 → 16

3 standard reflections

frequency: 60 min

intensity variation: 7%

Refinement

Refinement on *F*

R = 0.037

wR = 0.040

S = 0.9902

790 reflections

185 parameters

All H-atom parameters refined

w = 0.6504/[σ²(*F*) + 0.000440*F*²]

(Δ/σ)_{max} = 0.002

Δρ_{max} = 0.13 e Å⁻³

Δρ_{min} = -0.22 e Å⁻³

Extinction correction: none

Atomic scattering fac-

tors from *CRYSRULER* (Rizzoli, Sangermano, Calestani & Andreetti, 1989)

Table 1. Fractional atomic coordinates and equivalent isotropic displacement parameters (Å²) for (I)

$$U_{eq} = (1/3)\sum_i\sum_j U_{ij}a_i^*a_j^*a_j$$

	<i>x</i>	<i>y</i>	<i>z</i>	<i>U</i> _{eq}
N1	0.6610 (1)	0.0941 (2)	0.7529 (1)	0.0364 (4)
C11	0.6372 (2)	0.0142 (2)	0.6370 (1)	0.0435 (5)
C2	0.5730 (1)	0.1261 (2)	0.8064 (1)	0.0375 (4)
C21	0.4347 (1)	0.1169 (2)	0.7485 (1)	0.0389 (4)
C22	0.3772 (1)	0.1849 (2)	0.6365 (1)	0.0452 (5)
C23	0.2462 (2)	0.1863 (3)	0.5876 (2)	0.0568 (6)
C24	0.1718 (2)	0.1214 (3)	0.6504 (2)	0.0629 (7)
C25	0.2276 (2)	0.0540 (3)	0.7615 (2)	0.0602 (8)
C26	0.3586 (2)	0.0512 (2)	0.8111 (2)	0.0487 (6)
N3	0.6259 (1)	0.1744 (2)	0.9181 (1)	0.0507 (5)
C4	0.7524 (2)	0.1753 (2)	0.9367 (1)	0.0537 (6)
C5	0.7758 (1)	0.1261 (2)	0.8372 (1)	0.0443 (6)
N50	0.8940 (1)	0.1195 (2)	0.8171 (2)	0.0623 (6)
O501	0.9887 (1)	0.1454 (3)	0.9012 (2)	0.1055 (9)
O502	0.8965 (1)	0.0952 (3)	0.7175 (2)	0.0971 (9)

Table 3. Fractional atomic coordinates and equivalent isotropic displacement parameters (Å²) for (II)

$$U_{eq} = (1/3)\sum_i\sum_j U_{ij}a_i^*a_j^*a_j$$

	<i>x</i>	<i>y</i>	<i>z</i>	<i>U</i> _{eq}
N1	0.6626 (2)	0.0962 (4)	0.2523 (2)	0.043 (1)
C11	0.6331 (4)	0.0214 (6)	0.1343 (3)	0.050 (2)

C2	0.5842 (3)	0.1271 (4)	0.3089 (2)	0.042 (1)
C21	0.4545 (3)	0.1193 (5)	0.2560 (3)	0.041 (1)
C22	0.3854 (3)	0.0582 (5)	0.3210 (3)	0.048 (1)
C23	0.2628 (3)	0.0585 (5)	0.2770 (3)	0.055 (2)
C24	0.2017 (3)	0.1215 (5)	0.1645 (3)	0.054 (2)
N24	0.0795 (3)	0.1208 (7)	0.1189 (4)	0.090 (2)
C25	0.2690 (3)	0.1835 (5)	0.0981 (3)	0.052 (2)
C26	0.3927 (3)	0.1819 (5)	0.1440 (3)	0.046 (2)
N3	0.6414 (3)	0.1681 (4)	0.4214 (2)	0.056 (1)
C4	0.7593 (3)	0.1646 (5)	0.4363 (3)	0.058 (2)
C5	0.7748 (3)	0.1216 (5)	0.3348 (3)	0.048 (1)
N50	0.8830 (3)	0.1151 (5)	0.3106 (3)	0.071 (2)
O501	0.9777 (2)	0.1343 (5)	0.3921 (3)	0.093 (1)
O502	0.8786 (2)	0.0923 (6)	0.2105 (3)	0.116 (2)

Table 4. Selected geometric parameters (Å, °) for (II)

N1—C11	1.471 (4)	C24—N24	1.363 (4)
N1—C2	1.355 (4)	C24—C25	1.397 (5)
N1—C5	1.380 (3)	C25—C26	1.380 (4)
C2—C21	1.452 (4)	N3—C4	1.347 (5)
C2—N3	1.343 (3)	C4—C5	1.353 (5)
C21—C22	1.398 (5)	C5—N50	1.413 (5)
C21—C26	1.392 (4)	N50—O501	1.228 (3)
C22—C23	1.367 (4)	N50—O502	1.218 (5)
C23—C24	1.397 (4)		
C2—N1—C5	105.3 (2)	C23—C24—N24	121.4 (3)
C11—N1—C5	127.8 (3)	N24—C24—C25	120.2 (3)
C11—N1—C2	126.1 (3)	C24—C25—C26	120.1 (3)
N1—C2—N3	111.5 (3)	C21—C26—C25	122.1 (3)
N1—C2—C21	125.1 (3)	C2—N3—C4	105.7 (3)
C21—C2—N3	123.5 (3)	N3—C4—C5	110.0 (3)
C2—C21—C26	123.0 (3)	N1—C5—C4	107.6 (3)
C2—C21—C22	119.9 (3)	C4—C5—N50	128.2 (3)
C22—C21—C26	116.9 (3)	N1—C5—N50	124.1 (3)
C21—C22—C23	121.9 (3)	C5—N50—O502	119.0 (3)
C22—C23—C24	120.7 (3)	C5—N50—O501	118.1 (3)
C23—C24—C25	118.3 (3)	O501—N50—O502	122.9 (4)

Data collection: *DIF4* (Stoe & Cie, 1992a). Cell refinement: *DIF4*. Data reduction: *REDU4* (Stoe & Cie, 1992b). Program(s) used to solve structure: *SHELXS86* (Sheldrick, 1985). Program(s) used to refine structure: *SHELX76* (Sheldrick, 1976). Molecular graphics: *ORTEPII* (Johnson, 1976). Software used to prepare material for publication: *PARST* (Nardelli, 1983).

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Lists of structure factors, anisotropic displacement parameters, H-atom coordinates and complete geometry have been deposited with the British Library Document Supply Centre as Supplementary Publication No. SUP 71819 (23 pp.). Copies may be obtained through The Technical Editor, International Union of Crystallography, 5 Abbey Square, Chester CH1 2HU, England. [CIF reference: NA1071]

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Methyl DL-3-Benzyl-2,2-dioxo-1,2,3-oxathiazolidine-4-carboxylate – an Intermediate for Amino Acid Synthesis

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Abstract

The first structural measurements on a cyclic sulfamidate, DL-3-benzyl-2,2-dioxo-1,2,3-oxathiazolidine-4-carboxylate, C₁₁H₁₃NO₅S, show that the ring N atom has almost planar bonding geometry (sum of angles at N = 357.8°), in contrast to that found in simple dialkyl sulfonamides and acyclic sulfamate esters. The structure contains a close 1,4 N···O contact [2.764 (3) Å].

Comment

The methyl ester of DL-serine can be converted to the 1,2,3-oxathiazolidine dioxide (1) by three steps: benzylation, cyclization with thionyl chloride, and oxidation with sodium periodate catalysed by ruthenium trichloride. This cyclic sulfamidate reacts with nucleophiles by ring opening at C-5 to produce precursors of new amino acids (Gritsonie, Pilkington & Wallis, 1994; Baldwin, Spivey & Schofield, 1990). We were interested in the bonding geometry at the N atom in (1), and how this might be affected by the sizes of the N-atom substituent and the adjacent carboxylic ester group. By analogy with *N,N*-dimethyl sulfonamides in which the N atom is typically *circa* 0.27 Å out of the plane defined by the atoms to which it is bonded (Häkkinen, Ruostesuo & Kivekäs, 1988; Cromer, Ryan, Karthikeyan & Paine, 1990), pyramidal geometry around the N atom was expected. The only *N,N*-dialkyl sulfamate esters recorded in the Cam-