# organic compounds

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# 1,3-Dinitrosoimidazolidine

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Key indicators: single-crystal X-ray study; T = 120 K; mean  $\sigma(O-N) = 0.009$  Å; disorder in main residue; R factor = 0.039; wR factor = 0.126; data-to-parameter ratio = 6.0.

The title compound, C<sub>3</sub>H<sub>6</sub>N<sub>4</sub>O<sub>2</sub>, exhibits partial disorder with the refined occupancy ratios of the two components being 0.582 (5):0.418 (5). In the major component, the nitroso groups have a relative syn spatial arrangement  $[O=N\cdots N=O$  pseudo-torsion angle = 1.1 (4)°], whereas the other component has an *anti* disposition  $[177.6 (1)^{\circ}]$ . The N–N=O moieties are almost coplanar with a dihedral angle of 5.3  $(3)^{\circ}$ , while in the minor occupied set of atoms, this angle is 8  $(1)^{\circ}$ . In both components, the imidazolidine ring adopts a twisted conformation on the C-C bond and the crystal structure shows the strain of this ring according to the N- $CH_2 - CH_2 - N$  torsion angles [25.9 (5) and -23.8 (7)°]. In the crystal, molecules are linked by weak C-H···O hydrogen bonds.

#### **Related literature**

For a related structure, see: Rivera et al. (2011). For the synthesis of the title compound, see: Rivera et al. (1997). For ring conformations, see Cremer & Pople (1975). For chemical background on the synthesis and uses of N-nitroso amines, see: Di Salvo et al. (2008).



a = 9.5154 (2) Å b = 5.4338(1) Å c = 10.7104 (2) Å

**Experimental** 

V = 553.78 (2) Å<sup>3</sup> 7 - 4Cu Ka radiation

#### Data collection

Agilent Xcalibur diffractometer	5160 measured reflections
with an Atlas (Gemini Ultra Cu)	522 independent reflections
detector	514 reflections with $I > 3\sigma(I)$
Absorption correction: multi-scan	$R_{\rm int} = 0.031$
(CrysAlis PRO; Agilent, 2010)	
$T_{\min} = 0.636, T_{\max} = 1$	

Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.039$ 87 parameters  $wR(F^2) = 0.126$ H-atom parameters constrained S = 2.86 $\Delta \rho_{\rm max} = 0.12 \text{ e } \text{\AA}^ \Delta \rho_{\rm min} = -0.19 \text{ e} \text{ Å}^{-3}$ 522 reflections

 $\mu = 1.14 \text{ mm}^{-1}$ T = 120 K

 $0.39 \times 0.20 \times 0.14 \text{ mm}$ 

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

$D - H \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
$C3y - H3ya \cdots O1y^i$	0.96	1.85	2.681 (12)	143

Symmetry code: (i) x, y + 1, z.

Data collection: CrysAlis PRO (Agilent, 2010); cell refinement: CrysAlis PRO; data reduction: CrysAlis PRO; program(s) used to solve structure: SIR2002 (Burla et al., 2003); program(s) used to refine structure: JANA2006 (Petříček et al., 2006); molecular graphics: DIAMOND (Brandenburg & Putz, 2005); software used to prepare material for publication: JANA2006.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: BT5956).

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# supporting information

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# 1,3-Dinitrosoimidazolidine

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# S1. Comment

*N*-nitrosamines are interesting molecules due their strong carcinogenic and mutagenic properties and their utility as synthetic intermediates for the preparation of various *N*,*N*-bonded functionalities (Di Salvo *et al.*, 2008). Our group has previously explored the reaction of nitrous acid with cyclic aminals which actually are tertiary amines (Rivera *et al.*, 1997, 2011). Earlier we reported the synthesis and complete characterization by NMR of the title compound 1,3-dinitro-soimidazolidine, obtained by the nitrosation reaction of the cyclic aminal 1,3,6,8-tetraazatricyclo[4.4.1.1.<sup>3,8</sup>]dodecane (Rivera *et al.*, 1997). NMR experiments of this compound evidenced the existence of a mixture of three isomers: *syn-cis*, *anti*, and *cis-trans* with a ratio of 31:50:19 respectively (Rivera, *et al.* 1997). However, a recently investigation of (3aRS,7aRS)-1,3-dinitrosooctahydro-1*H*-benzimidazole, we found that the nitroso groups of this analogous X-ray crystal structure are on a *syn-cis* disposition (Rivera, *et al.* 2011). This result suggests that the orientation of the nitroso groups on the imidazolidine ring is largely influenced by their molecular skeletons. To identify the orientation of nitroso groups, we synthesized the title compound and investigated its crystal structure.

X-Ray analysis confirms the molecular structure and atom connectivity as illustrated in Fig. 1. The bond lengths N—C1 and N—NO are normal and comparable to the corresponding values observed in the related structure (Rivera, *et al.* 2011). The title compound are disordered over two sets of sites [site occupancies = 0.588 (6) and 0.412 (6)]. In both components, the N—N=O moieties are almost coplanar showing dihedral angles of 5.277 (340)° for the major component and 7.81 (97)° for the minor. The nitroso substituents in the major component are on a *syn* spatial arrangement as can be seen from O1x=N3x…N4=O2 pseudo torsion angle of = 1.119 (410)°, whereas the other component have an *anti* disposition [pseudo torsion angle O1y=N3y…N4=O2 = 177.662 (129)°] (Figure 2). Both imidazole ring system are twisted on CH<sub>2</sub>CH<sub>2</sub> fragment as seen in the puckering parameters Q(2) = 0.255 (5) Å and  $\varphi 2$  = 122.7 (12)° for major component and Q(2) = 0.236 (8) Å and  $\varphi 2$  = 90.9 (16)° for minor component (Cremer & Pople, 1975). The crystal structure shows the strain of this ring according to the NCH<sub>2</sub>CH<sub>2</sub>N torsion angles [N1xC2xC3xN2x = 25.874 (534)° and N1yC2yC3yN2y = -23.808 (735)°].

The crystal packing displays a weak intermolecular C—H···O [C···O = 2.681 (12) Å] non-conventional hydrogen bonding interactions between oxygen atoms in the nitroso moiety and hydrogen atoms in methylene carbons of the heterocyclic ring (Figure 3).

## **S2. Experimental**

For the originally reported synthesis, see: Rivera *et al.* (1997). Single crystals of the title compound were obtained by recrystallization from EtOH solution (m.p 318 K).

# **S3. Refinement**

All hydrogen atoms were positioned geometrically and treated as riding on their parent atoms. The isotropic atomic displacement parameters of hydrogen atoms were set to  $1.2 \times U_{eq}$  of the parent atom.

The molecule is disordered over two positions with occupancies 0.588 (6):0.412 (6). Selected atoms of both components were refined isotropically as anisotropic refinement lead to unreasonable ADPs.

As the structure contains only light atoms, the Friedel-pairs were merged and the Flack parameter has not been determined.



## Figure 1

A view of the title compound with the numbering scheme. Only the major disordered component is shown. Displacement ellipsoids are drawn at the 50% probability level.





Overlapped structures of disordered components, showing structural differences.



Figure 3

The packing of the title compound, viewed along the *a* axis. The dashed lines indicate the hydrogen bonds. H atoms not involved in hydrogen bonding have been omitted for clarity.

1,3-Dinitrosoimidazolidine

Crystal data  $C_3H_6N_4O_2$  $M_r = 130.1$ 

Orthorhombic, *Pna2*<sub>1</sub> Hall symbol: P 2c -2n Cu *K* $\alpha$  radiation,  $\lambda = 1.5418$  Å

 $\theta = 4.1 - 66.8^{\circ}$ 

 $\mu = 1.14 \text{ mm}^{-1}$ 

Prism, colourless

 $0.39 \times 0.20 \times 0.14 \text{ mm}$ 

T = 120 K

Cell parameters from 4445 reflections

a = 9.5154 (2) Å b = 5.4338 (1) Å c = 10.7104 (2) Å  $V = 553.78 (2) \text{ Å}^3$  Z = 4 F(000) = 272 $D_x = 1.56 \text{ Mg m}^{-3}$ 

Data collection

Data conection	
Agilent Xcalibur	$T_{\min} = 0.636, \ T_{\max} = 1$
diffractometer with an Atlas (Gemini Ultra Cu)	5160 measured reflections
detector	522 independent reflections
Radiation source: Enhance Ultra (Cu) X-ray	514 reflections with $I > 3\sigma(I)$
Source	$R_{\rm int} = 0.031$
Mirror monochromator	$\theta_{\rm max} = 67.0^{\circ},  \theta_{\rm min} = 9.2^{\circ}$
Detector resolution: 10.3784 pixels mm <sup>-1</sup>	$h = -11 \rightarrow 11$
Rotation method data acquisition using $\omega$ scans	$k = -6 \rightarrow 6$
Absorption correction: multi-scan	$l = -12 \rightarrow 12$
(CrysAlis PRO; Agilent, 2010)	

## Refinement

Refinement on $F^2$	86 constraints
$R[F > 3\sigma(F)] = 0.039$	H-atom parameters constrained
wR(F) = 0.126	Weighting scheme based on measured s.u.'s $w =$
<i>S</i> = 2.86	$1/(\sigma^2(I) + 0.0016I^2)$
522 reflections	$(\Delta/\sigma)_{\rm max} = 0.013$
87 parameters	$\Delta \rho_{\rm max} = 0.12 \text{ e } \text{\AA}^{-3}$
0 restraints	$\Delta \rho_{\min} = -0.19 \text{ e} \text{ Å}^{-3}$

## Special details

**Experimental**. CrysAlisPro (Agilent Technologies, 2010) Empirical absorption correction using spherical harmonics, implemented in SCALE3 ABSPACK scaling algorithm.

**Refinement**. The refinement was carried out against all reflections. The conventional *R*-factor is always based on *F*. The goodness of fit as well as the weighted *R*-factor are based on *F* and  $F^2$  for refinement carried out on *F* and  $F^2$ , respectively. The threshold expression is used only for calculating *R*-factors *etc*. and it is not relevant to the choice of reflections for refinement.

The program used for refinement, Jana2006, uses the weighting scheme based on the experimental expectations, see \_refine\_ls\_weighting\_details, that does not force S to be one. Therefore the values of S are usually larger than the ones from the *SHELX* program.

Fractional atomic coordinates and isotropic	c or equivalent	isotropic displaceme	ent parameters (2	$(A^2)$
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	x	У	Ζ	$U_{ m iso}$ */ $U_{ m eq}$	Occ. (<1)
O1x	0.3214 (4)	-0.3809 (8)	0.266995	0.0436 (13)	0.582 (5)
02	0.0084 (2)	0.3676 (5)	0.3407 (6)	0.0527 (8)	
N1x	0.2304 (5)	-0.1122 (9)	0.1438 (8)	0.0333 (9)*	0.582 (5)
N1y	0.2125 (8)	-0.1576 (12)	0.1559 (9)	0.0333 (9)*	0.418 (5)
N2x	0.0781 (4)	0.1899 (8)	0.1754 (7)	0.0292 (8)*	0.582 (5)
N2y	0.1109 (6)	0.2145 (12)	0.1820 (9)	0.0292 (8)*	0.418 (5)
N3x	0.3056 (5)	-0.3111 (9)	0.1569 (10)	0.0342 (15)	0.582 (5)
N3y	0.2972 (18)	-0.338 (3)	0.2034 (16)	0.063 (5)*	0.418 (5)
O1y	0.3425 (9)	-0.4441 (19)	0.1083 (12)	0.081 (2)*	0.418 (5)
N4	0.0159 (3)	0.3748 (4)	0.2258 (6)	0.0469 (9)	

C1	0.1576 (2)	0.0075 (6)	0.2498 (6)	0.0348 (8)	
C2x	0.2145 (5)	0.0048 (10)	0.0225 (8)	0.0369 (11)*	0.582 (5)
C2y	0.1820 (8)	-0.0681 (14)	0.0312 (9)	0.0369 (11)*	0.418 (5)
C3x	0.0820 (5)	0.1482 (9)	0.0410 (8)	0.0338 (13)	0.582 (5)
C3y	0.1527 (8)	0.2091 (13)	0.0512 (10)	0.0338 (13)	0.418 (5)
H2xa	0.199648	-0.11884	-0.0401	0.0443*	0.582 (5)
H2xb	0.291244	0.116195	0.008783	0.0443*	0.582 (5)
H2ya	0.098714	-0.146931	0.000258	0.0443*	0.418 (5)
H2yb	0.263608	-0.087618	-0.020601	0.0443*	0.418 (5)
H3xa	0.089387	0.303307	-0.0015	0.0406*	0.582 (5)
H3xb	0.00295	0.049365	0.016793	0.0406*	0.582 (5)
H3ya	0.23834	0.300237	0.040787	0.0406*	0.418 (5)
H3yb	0.074938	0.258704	-0.000047	0.0406*	0.418 (5)
H1ax	0.225236	0.091148	0.301089	0.0417*	0.582 (5)
H1bx	0.093385	-0.107021	0.28725	0.0417*	0.582 (5)
Hlay	0.232011	0.055714	0.30525	0.0417*	0.418 (5)
H1by	0.079178	-0.068441	0.29095	0.0417*	0.418 (5)

Atomic displacement parameters  $(Å^2)$ 

	$U^{11}$	$U^{22}$	$U^{33}$	$U^{12}$	$U^{13}$	$U^{23}$
O1x	0.0389 (19)	0.053 (2)	0.039 (3)	0.0103 (15)	-0.0008 (19)	0.0147 (19)
O2	0.0448 (12)	0.0675 (15)	0.0458 (16)	0.0025 (10)	0.0059 (10)	-0.0244 (11)
N3x	0.031 (2)	0.036 (2)	0.036 (3)	0.0200 (15)	0.008 (2)	0.002 (2)
N4	0.0526 (17)	0.0417 (14)	0.0463 (17)	0.0098 (11)	0.0081 (13)	-0.0109 (11)
C1	0.0288 (13)	0.0449 (14)	0.0306 (14)	0.0034 (9)	-0.0021 (11)	-0.0035 (11)
C3x	0.038 (2)	0.031 (2)	0.033 (2)	0.0003 (18)	0.007 (2)	0.0029 (16)
C3y	0.038 (2)	0.031 (2)	0.033 (2)	0.0003 (18)	0.007 (2)	0.0029 (16)

Geometric parameters (Å, °)

O1x—N3x	1.248 (10)	N2y—C1	1.411 (9)
O2—N4	1.233 (9)	N2y—C3y	1.457 (14)
N1x—N3x	1.303 (7)	N3y—O1y	1.25 (2)
N1x—C1	1.481 (9)	C1—H1bx	0.96
N1x—C2x	1.454 (11)	C1—H1ay	0.96
N1y—N3y	1.366 (17)	C1—H1by	0.96
N1y—C1	1.445 (10)	C2x—C3x	1.495 (7)
N1y—C2y	1.451 (13)	C2x—H2xa	0.96
N2x—N4	1.285 (6)	C2x—H2xb	0.96
N2x—C1	1.480 (7)	C2y—C3y	1.547 (11)
N2x—C3x	1.458 (12)	C3x—H3xb	0.96
N2y—N4	1.340 (8)	C3y—H3yb	0.96
N3x—N1x—C1	122.6 (7)	N1v—C1—H1av	109.4716
N3x—N1x—C2x	121.0 (7)	N1v—C1—H1bv	109.4714
C1—N1x—C2x	116.4 (4)	N2x—C1—H1bx	109.4709
N3y—N1y—C1	113.5 (10)	N2y—C1—H1ay	109.4711

134.5 (10)	N2y—C1—H1by	109.4712
111.1 (6)	N1x— $C2x$ — $C3x$	101.4 (6)
122.2 (7)	N1x—C2x—H2xa	109.4711
123.2 (5)	N1x—C2x—H2xb	109.4713
114.5 (4)	C3x—C2x—H2xa	109.4718
123.4 (8)	C3x—C2x—H2xb	109.4715
122.3 (7)	H2xa—C2x—H2xb	116.4713
113.1 (6)	N1y—C2y—C3y	103.6 (7)
114.8 (8)	N2x— $C3x$ — $C2x$	103.4 (6)
103.4 (13)	N2x—C3x—H3xb	109.4711
114.9 (5)	C2x—C3x—H3xb	109.4711
111.6 (5)	N2y—C3y—C2y	101.7 (7)
96.9 (5)	N2y—C3y—H3yb	109.4711
109.4713	C2y—C3y—H3yb	109.4709
104.5 (7)		
	134.5 (10) 111.1 (6) 122.2 (7) 123.2 (5) 114.5 (4) 123.4 (8) 122.3 (7) 113.1 (6) 114.8 (8) 103.4 (13) 114.9 (5) 111.6 (5) 96.9 (5) 109.4713 104.5 (7)	134.5 (10)N2y—C1—H1by $111.1 (6)$ N1x—C2x—C3x $122.2 (7)$ N1x—C2x—H2xa $123.2 (5)$ N1x—C2x—H2xb $114.5 (4)$ C3x—C2x—H2xb $123.4 (8)$ C3x—C2x—H2xb $122.3 (7)$ H2xa—C2x—H2xb $113.1 (6)$ N1y—C2y—C3y $114.8 (8)$ N2x—C3x—C2x $103.4 (13)$ N2x—C3x—H3xb $114.9 (5)$ C2x—C3y—H3xb $111.6 (5)$ N2y—C3y—H3yb $109.4713$ C2y—C3y—H3yb $104.5 (7)$ C3x

# Hydrogen-bond geometry (Å, °)

<i>D</i> —H··· <i>A</i>	D—H	Н…А	D····A	<i>D</i> —H··· <i>A</i>
C2y—H2 $ya$ ···O2 <sup>i</sup>	0.96	2.32	3.177 (10)	148
C3 <i>y</i> —H3 <i>ya</i> ···O1 <i>y</i> <sup>ii</sup>	0.96	1.85	2.681 (12)	143

Symmetry codes: (i) –*x*, –*y*, *z*–1/2; (ii) *x*, *y*+1, *z*.