

## 2,3-Dimethylquinoxaline–dimethylglyoxime (1/1)

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## Key indicators

Single-crystal X-ray study

$T = 294$  K

Mean  $\sigma(\text{C}-\text{C}) = 0.002$  Å

$R$  factor = 0.046

$wR$  factor = 0.133

Data-to-parameter ratio = 14.9

For details of how these key indicators were automatically derived from the article, see <http://journals.iucr.org/e>.

The title compound,  $\text{C}_{10}\text{H}_{10}\text{N}_2 \cdot \text{C}_4\text{H}_8\text{N}_2\text{O}_2$ , consists of hydrogen-bonded 2,3-dimethylquinoxaline and dimethylglyoxime molecules. Both dimethylglyoximes are located on an inversion centre and there are two half-molecules in the asymmetric unit. The dimethylglyoxime molecules are linked about inversion centres to the 2,3-dimethylquinoxaline moiety by  $\text{O}-\text{H} \cdots \text{N}$  hydrogen bonds ( $\text{O} \cdots \text{N}$  2.898 and 2.805 Å) to form polymeric chains.

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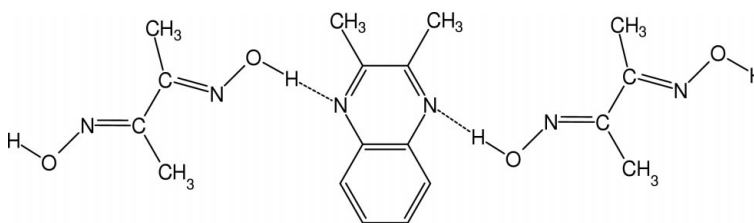
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## Comment

Intermolecular hydrogen bonding has received considerable attention among the directional non-covalent intermolecular interactions (Etter *et al.*, 1990), which combines moderate strength and directionality (Karle *et al.*, 1996) in designing compounds to form supramolecular structures.

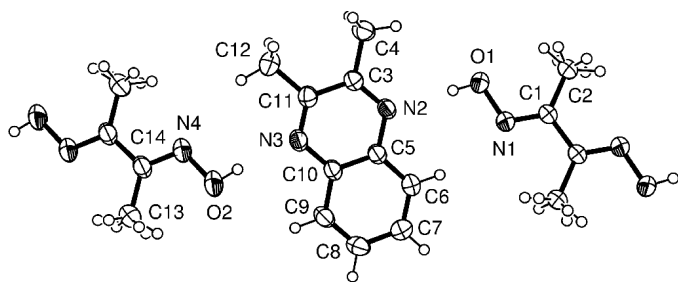
Oxime ( $-\text{C}=\text{N}-\text{OH}$ ) groups possess stronger hydrogen-bonding capabilities than alcohols, phenols and carboxylic acids (Marsman *et al.*, 1999). The hydrogen-bond systems in the crystals of oximes have been analysed and a correlation between a pattern of hydrogen bonding and  $\text{N}-\text{O}$  bond lengths has been suggested (Bertolasi *et al.*, 1982). The configurational and/or conformational isomers of glyoxime derivatives (dioximes) have also been analysed (Chertanova *et al.*, 1994).

Oxime/oximato metal complexes have been investigated actively since the beginning of the last century of the millennium (Kukushkin & Pombeiro, 1999). In general, oxime and dioxime derivatives are very important compounds in the chemical industry and medicine. Dimethylglyoxime is actually the first selective organic reagent applied in the analysis of metals (Tshugaeff, 1905).

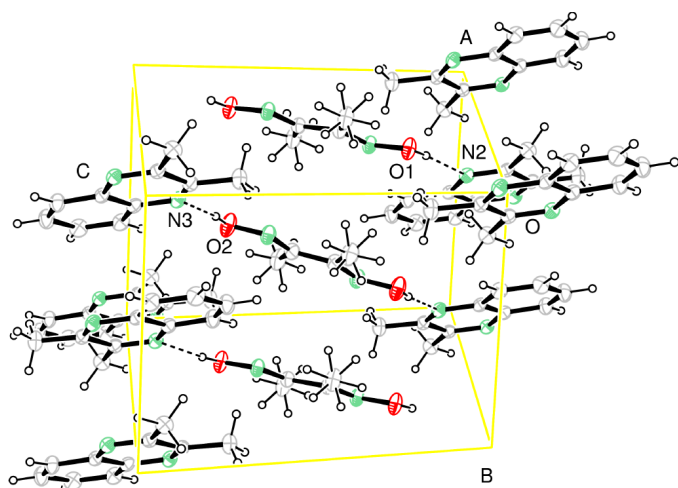


(I)

Crystals of dimethylglyoxime were reported by McCrone (1949) to be triclinic and its structure was undertaken by Merritt & Lanterman (1952). A neutron diffraction refinement of the crystal structure of dimethylglyoxime clarified the existence of the  $\text{O}-\text{H}$  (classical structure) bonding rather



**Figure 1**  
An ORTEP (Johnson, 1976) drawing of the title molecule with the atom-numbering scheme. The displacement ellipsoids are drawn at the 50% probability level.



**Figure 2**  
Packing diagram for (I). Hydrogen bonds are shown as dotted lines.

than N—H (zwitterion structure) (Hamilton, 1961).

Quinoxaline derivatives are very useful compounds with well known biological activities (Cheeseman & Werstiuk, 1978; Sato, 1996). Some of the quinoxaline derivatives act as *in vitro* anticancer compounds (Corona *et al.*, 1998). On the other hand, 2,3-dimethylquinoxaline is a polarographically active compound (Rodrigues *et al.*, 1997).

The structure of 2,3-dimethylquinoxaline was reported by Wozniak *et al.* (1990) to be monoclinic. The crystal structures containing 2,3-disubstituted derivatives of quinoxaline have aroused considerable interest because of the repulsions between the neighbouring substituents (Visser *et al.*, 1968; Visser & Vos, 1971; Lipkowski *et al.*, 1985; Krigier *et al.*, 1985).

The crystal structure determination of the title molecule, (I), was carried out in order to understand the strength of the hydrogen-bonding capabilities of the oxime groups having the classical (—C=N—O—H) structures. As shown in Fig. 1, the title compound, (I), consists of hydrogen-bonded 2,3-dimethylquinoxaline and dimethylglyoxime moieties. Both dimethylglyoximes are located on an inversion centre and there are two half-molecules in the asymmetric unit.

The bond lengths and angles of the 2,3-dimethylquinoxaline moiety are in accordance with the reported values (Woźniak *et al.*, 1990).

In the dimethylglyoxime moieties, the O—N, C=N, C—C bond lengths and C=N—O bond angles are larger, while the O—H, C—C' bonds and N—O—H, C—C=N, C'—C=N angles are smaller than the corresponding ones reported in dimethylglyoxime (Hamilton, 1961) (Table 2). A comparison of the bond lengths and angles of the dimethylglyoxime moieties in the title compound, (I), with the corresponding ones in dimethylglyoxime, (II) (Hamilton, 1961), nickel dimethylglyoxime, (III) (Godycki & Rundle, 1953), copper dimethylglyoxime, (IV) (Frasson *et al.*, 1959) and acetoxime, (V) (Bierlein & Lingafelter, 1951), are given in Table 2.

As can be seen from the packing diagram (Fig. 2), there are intermolecular hydrogen bonds between the hydroxy H atoms of the dimethylglyoxime and N atoms of the 2,3-dimethylquinoxaline moieties (Table 3). The intermolecular hydrogen bonds are highly effective in forming the polymeric chains. Dipole-dipole and van der Waals interactions are also effective in the molecular packing.

An examination of the deviations from the least-squares planes through the individual rings shows that rings A (C5—C10) and B (C5/N2/C3/C11/N3/C10) are nearly planar with maximum deviations for the C5 [−0.007 (2) Å] and C5 [0.016 (2) Å] atoms, respectively.

## Experimental

The title compound, (I), was prepared from a mixture of 2,3-butaneidone monoxime (2.02 g, 20.0 mmol), *o*-phenylenediamine (1.08 g, 10.0 mmol) and Na<sub>2</sub>SO<sub>4</sub> (2.84 g, 20.0 mmol) in ethanol (30 ml). The mixture was heated at 343 K for 6 h. The solution was filtered and allowed to stand overnight in a refrigerator. The resulting precipitate was filtered and then recrystallized from 1,4-dioxane.

### Crystal data

C<sub>10</sub>H<sub>10</sub>N<sub>2</sub>·C<sub>4</sub>H<sub>8</sub>N<sub>2</sub>O<sub>2</sub>  
*M<sub>r</sub>* = 274.32  
 Triclinic, *P* $\bar{1}$   
*a* = 7.9814 (10) Å  
*b* = 8.2719 (10) Å  
*c* = 12.168 (3) Å  
 $\alpha$  = 91.779 (18)°  
 $\beta$  = 106.668 (14)°  
 $\gamma$  = 111.018 (10)°  
*V* = 710.3 (2) Å<sup>3</sup>

*Z* = 2  
*D<sub>x</sub>* = 1.283 Mg m<sup>−3</sup>  
 Mo *K*α radiation  
 Cell parameters from 25 reflections  
 $\theta$  = 10–11°  
 $\mu$  = 0.09 mm<sup>−1</sup>  
*T* = 294 (2) K  
 Rod, yellow  
 0.30 × 0.25 × 0.20 mm

### Data collection

Enraf–Nonius CAD-4 diffractometer  
 $\omega/2\theta$  scans  
 Absorption correction:  $\psi$  scan (Fair, 1990)  
*T<sub>min</sub>* = 0.974, *T<sub>max</sub>* = 0.982  
 2884 measured reflections  
 2884 independent reflections

2244 reflections with *I* > 2σ(*I*)  
 $\theta_{\max}$  = 26.3°  
*h* = 0 → 9  
*k* = −10 → 9  
*l* = −15 → 14  
 3 standard reflections  
 frequency: 120 min  
 intensity decay: 1%

### Refinement

Refinement on *F*<sup>2</sup>  
*R*[*F*<sup>2</sup> > 2σ(*F*<sup>2</sup>)] = 0.046  
*wR*(*F*<sup>2</sup>) = 0.133  
*S* = 1.08  
 2884 reflections  
 193 parameters  
 H atoms treated by a mixture of independent and constrained refinement

$w = 1/[\sigma^2(F_o^2) + (0.0745P)^2 + 0.0913P]$   
 where  $P = (F_o^2 + 2F_c^2)/3$   
 $(\Delta/\sigma)_{\max} < 0.001$   
 $\Delta\rho_{\max} = 0.20 \text{ e } \text{Å}^{-3}$   
 $\Delta\rho_{\min} = -0.23 \text{ e } \text{Å}^{-3}$

**Table 1**  
Selected geometric parameters (Å, °).

C1—N1	1.2811 (18)	C14—N4	1.2813 (19)
C1—C1 <sup>i</sup>	1.477 (3)	C14—C14 <sup>ii</sup>	1.473 (3)
C1—C2	1.491 (2)	C14—C13	1.499 (2)
C3—N2	1.3119 (18)	N1—O1	1.4033 (16)
C5—N2	1.3723 (18)	N4—O2	1.3956 (18)
C10—N3	1.3676 (19)	O1—H1	0.85 (2)
C11—N3	1.3123 (19)	O2—H2	0.82 (2)
N1—C1—C1 <sup>i</sup>	115.18 (15)	C14 <sup>ii</sup> —C14—C13	120.72 (16)
N1—C1—C2	124.09 (13)	C1—N1—O1	112.39 (12)
C1 <sup>i</sup> —C1—C2	120.73 (15)	C3—N2—C5	118.23 (11)
N2—C3—C11	121.25 (12)	C11—N3—C10	118.35 (12)
N3—C11—C3	121.11 (13)	C14—N4—O2	112.15 (13)
N4—C14—C14 <sup>ii</sup>	115.03 (17)	N1—O1—H1	104.5 (14)
N4—C14—C13	124.25 (15)	N4—O2—H2	102.4 (16)
N2—C5—C10—N3	2.0 (2)	C2—C1—N1—O1	0.8 (2)
N2—C3—C11—N3	2.0 (2)	C13—C14—N4—O2	−0.8 (2)

Symmetry codes: (i)  $1-x, -y, 1-z$ ; (ii)  $1-x, 1-y, -1-z$ .**Table 2**  
Hydrogen-bonding geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
O1—H1 $\cdots$ N2 <sup>i</sup>	0.85	2.05	2.898	176
O2—H2 $\cdots$ N3	0.82	1.99	2.805	173

Symmetry code: (i)  $x, y-1, 1+z$ .**Table 3**  
Comparison of the bond lengths (Å) and angles (°) in dimethylglyoxime moieties of (I) with the corresponding values in the related compounds (II), (III), (IV) and (V).

	(I)	(II)	(III)	(IV)	(V)
C1—C <sub>trans</sub>	1.477 (3)	1.473 (3)	1.562 (18)	1.53	1.52
C1—C <sub>cis</sub>	1.491 (2)	1.499 (2)	1.479 (15)	1.49	1.47
C1—N1	1.281 (2)	1.281 (2)	1.253 (11)	1.23	1.25
N1—O1	1.403 (2)	1.396 (2)	1.321 (21)	1.38	1.35
O1—H1	0.85 (2)	0.82 (2)	1.02 (4)	—	—
C <sub>cis</sub> —C1—C <sub>trans</sub>	120.7 (2)	120.7 (2)	120	123	122
N1—C1—C <sub>trans</sub>	115.2 (2)	115.0 (2)	114	111	113
N1—C1—C <sub>cis</sub>	124.1 (2)	124.3 (2)	126	127	125
C1—N1—O1	112.4 (2)	112.2 (2)	111	121	120
N1—O1—H1	104.5 (14)	102.4 (16)	110	102	—

The oxime H atoms were located from a difference map and refined isotropically; the positions of the remaining H atoms were calculated geometrically at distances of 0.93 (CH) and 0.96 Å (CH<sub>3</sub>) from the corresponding C atoms, and a riding model was used during the refinement process. The H atoms of the C2 and C13 methyl groups are disordered.

Data collection: *MolEN* (Fair, 1990); cell refinement: *MolEN*; data reduction: *MolEN*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1990); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEPII* (Johnson, 1976); software used to prepare material for publication: *MolEN*.

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