

4,4'-[Oxalylbis(azanediyl)]dipyridinium bis(perchlorate)

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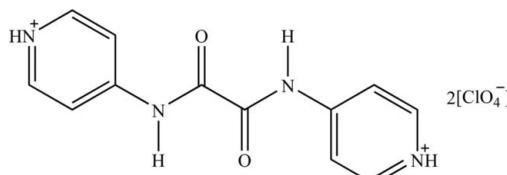
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Key indicators: single-crystal X-ray study; $T = 295\text{ K}$; mean $\sigma(\text{C-C}) = 0.005\text{ \AA}$; R factor = 0.051; wR factor = 0.119; data-to-parameter ratio = 11.4.

In the title molecular salt, $\text{C}_{12}\text{H}_{12}\text{N}_4\text{O}_2^{2+} \cdot 2\text{ClO}_4^-$, the complete cation is generated by crystallographic inversion symmetry. In the crystal, the cations and anions are linked via $\text{N}-\text{H}\cdots\text{O}$ and $\text{N}-\text{H}\cdots(\text{O},\text{O})$ hydrogen bonds, forming a three-dimensional framework.

Related literature

For the applications of N,N' -bis(pyridyl)oxamides, see: Hsu *et al.* (2004); Hu *et al.* (2004).



Experimental

Crystal data

$\text{C}_{12}\text{H}_{12}\text{N}_4\text{O}_2^{2+} \cdot 2\text{ClO}_4^-$	$V = 832.4(2)\text{ \AA}^3$
$M_r = 443.16$	$Z = 2$
Monoclinic, $P2_1/n$	Mo $K\alpha$ radiation
$a = 7.873(1)\text{ \AA}$	$\mu = 0.46\text{ mm}^{-1}$
$b = 9.3728(15)\text{ \AA}$	$T = 295\text{ K}$
$c = 11.3205(16)\text{ \AA}$	$0.6 \times 0.4 \times 0.2\text{ mm}$
$\beta = 94.827(10)^\circ$	

Data collection

Bruker P4 diffractometer
Absorption correction: ψ scan (*XSCANS*; Siemens, 1995)
 $T_{\min} = 0.919$, $T_{\max} = 0.982$
2017 measured reflections
1450 independent reflections

921 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.038$
3 standard reflections every 97 reflections
intensity decay: none

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.051$
 $wR(F^2) = 0.119$
 $S = 1.03$
1450 reflections

127 parameters
H-atom parameters constrained
 $\Delta\rho_{\max} = 0.31\text{ e \AA}^{-3}$
 $\Delta\rho_{\min} = -0.30\text{ e \AA}^{-3}$

Table 1
Hydrogen-bond geometry (\AA , $^\circ$).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
N1—H1A \cdots O4	0.86	2.21	2.950 (4)	144
N1—H1A \cdots O3 ⁱ	0.86	2.35	2.966 (5)	129
N2—H2A \cdots O2 ⁱⁱ	0.86	2.14	2.975 (5)	162

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

Data collection: *XSCANS* (Siemens, 1995); cell refinement: *XSCANS*; data reduction: *SHELXTL* (Sheldrick, 2008); program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *SHELXTL*; software used to prepare material for publication: *SHELXTL*.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: GK2310).

References

- Hsu, Y.-F. & Chen, J.-D. (2004). *Eur. J. Inorg. Chem.* pp. 1488–1493.
- Hu, H.-L., Yeh, C.-W. & Chen, J.-D. (2004). *Eur. J. Inorg. Chem.* pp. 4696–4701.
- Sheldrick, G. M. (2008). *Acta Cryst. A* **64**, 112–122.
- Siemens (1995). *XSCANS*. Siemens Analytical X-ray Instruments Inc., Madison, Wisconsin, USA.

supporting information

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4,4'-[Oxalylbis(azanediyl)]dipyridinium bis(perchlorate)

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S1. Comment

Several Ag(I) complexes containing *N,N'*-bis(2-pyridyl)oxamide ligands have been prepared, which show one-dimensional and two-dimensional structures (Hsu, *et al.*, 2004; Hu, *et al.*, 2004). To investigate the effect of ligand-isomerism on the structural type of such complexes, the ligand *N,N'*-bis(4-pyridyl)oxamide was synthesized and reacted with AgClO₄ in CH₂Cl₂. The reaction resulted unexpectedly in the perchlorate salt of the organic ligand. Within this project the crystal structure of the title compound was determined.

S2. Experimental

N,N'-bis(4-pyridyl)oxamide (0.24 g, 1.0 mmol) and AgClO₄ (0.21 g, 1.0 mmol) were placed in a flask containing 10 ml CH₂Cl₂. The mixture was then reflux for 12 h. The resulting solution was then filtered and then layered with diethyl ether to afford colorless plate crystals of the title compound suitable for X-ray crystallography.

S3. Refinement

All the hydrogen atoms were placed in idealized positions and refined using the riding model approximation with C—H = 0.93 — 0.96 Å, N—H = 0.86 Å and *U*_{iso}(H) = 1.2 *U*_{eq}(C, N).

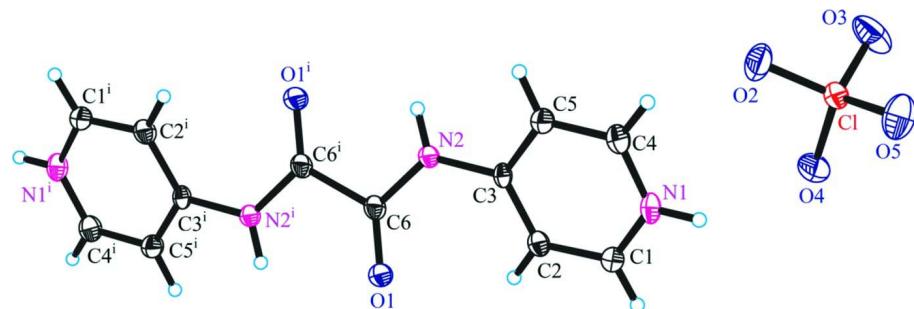
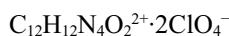


Figure 1

Crystal structure of the title compound with atom labeling and displacement ellipsoids drawn at the 30% probability level. Symmetry code: i = -x + 1, -y + 2, -z + 2.

4,4'-(Oxalylbis(azanediyl))dipyridinium bis(perchlorate)

Crystal data



*M*_r = 443.16

Monoclinic, P2₁/n

Hall symbol: -P 2yn

a = 7.873 (1) Å

b = 9.3728 (15) Å

c = 11.3205 (16) Å

β = 94.827 (10)°

V = 832.4 (2) Å³

Z = 2

$F(000) = 452$
 $D_x = 1.768 \text{ Mg m}^{-3}$
Mo $K\alpha$ radiation, $\lambda = 0.71073 \text{ \AA}$
Cell parameters from 25 reflections
 $\theta = 5.6\text{--}14.2^\circ$

$\mu = 0.46 \text{ mm}^{-1}$
 $T = 295 \text{ K}$
Plate, colorless
 $0.6 \times 0.4 \times 0.2 \text{ mm}$

Data collection

Bruker P4
diffractometer
Radiation source: fine-focus sealed tube
Graphite monochromator
 ω scans
Absorption correction: ψ scan
(*XSCANS*; Siemens, 1995)
 $T_{\min} = 0.919$, $T_{\max} = 0.982$
2017 measured reflections

1450 independent reflections
921 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.038$
 $\theta_{\max} = 25.0^\circ$, $\theta_{\min} = 2.8^\circ$
 $h = -9 \rightarrow 1$
 $k = -1 \rightarrow 11$
 $l = -13 \rightarrow 13$
3 standard reflections every 97 reflections
intensity decay: none

Refinement

Refinement on F^2
Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.051$
 $wR(F^2) = 0.119$
 $S = 1.03$
1450 reflections
127 parameters
0 restraints
Primary atom site location: structure-invariant
direct methods

Secondary atom site location: difference Fourier
map
Hydrogen site location: inferred from
neighbouring sites
H-atom parameters constrained
 $w = 1/[\sigma^2(F_o^2) + (0.0424P)^2 + 0.6192P]$
where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} < 0.001$
 $\Delta\rho_{\max} = 0.31 \text{ e \AA}^{-3}$
 $\Delta\rho_{\min} = -0.30 \text{ e \AA}^{-3}$

Special details

Experimental. Refinement of F^2 against ALL reflections. The weighted R -factor wR and goodness of fit S are based on F^2 , conventional R -factors R are based on F , with F set to zero for negative F^2 . The threshold expression of $F^2 > \sigma(F^2)$ is used only for calculating R -factors(gt) etc. and is not relevant to the choice of reflections for refinement. R -factors based on F^2 are statistically about twice as large as those based on F , and R -factors based on ALL data will be even larger.

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

Refinement. Refinement of F^2 against ALL reflections. The weighted R -factor wR and goodness of fit S are based on F^2 , conventional R -factors R are based on F , with F set to zero for negative F^2 . The threshold expression of $F^2 > \sigma(F^2)$ is used only for calculating R -factors(gt) etc. and is not relevant to the choice of reflections for refinement. R -factors based on F^2 are statistically about twice as large as those based on F , and R -factors based on ALL data will be even larger.

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

	x	y	z	$U_{\text{iso}}^*/U_{\text{eq}}$
C1	0.3752 (6)	1.1939 (5)	0.5833 (3)	0.0381 (11)
H1B	0.3929	1.2868	0.5582	0.046*
C2	0.4314 (5)	1.1532 (5)	0.6960 (3)	0.0326 (10)
H2C	0.4865	1.2182	0.7482	0.039*
C3	0.4053 (5)	1.0143 (4)	0.7310 (3)	0.0248 (9)
C4	0.2689 (6)	0.9652 (5)	0.5391 (3)	0.0382 (11)
H4B	0.2127	0.9029	0.4851	0.046*

C5	0.3260 (5)	0.9182 (5)	0.6505 (3)	0.0327 (10)
H5A	0.3118	0.8234	0.6718	0.039*
C6	0.4772 (5)	1.0472 (4)	0.9445 (3)	0.0288 (9)
N1	0.2943 (5)	1.0996 (4)	0.5092 (3)	0.0376 (9)
H1A	0.2575	1.1276	0.4394	0.045*
N2	0.4560 (4)	0.9647 (4)	0.8451 (2)	0.0285 (8)
H2A	0.4754	0.8748	0.8533	0.034*
O1	0.4640 (4)	1.1740 (3)	0.9511 (2)	0.0426 (8)
Cl	0.07220 (14)	0.92622 (11)	0.21648 (8)	0.0338 (3)
O2	0.0153 (5)	0.8460 (4)	0.3141 (2)	0.0579 (10)
O3	0.1889 (4)	0.8433 (4)	0.1564 (3)	0.0653 (11)
O4	0.1543 (4)	1.0522 (3)	0.2623 (2)	0.0539 (9)
O5	-0.0718 (4)	0.9610 (4)	0.1369 (3)	0.0598 (10)

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C1	0.057 (3)	0.031 (2)	0.027 (2)	0.002 (2)	0.005 (2)	0.0014 (19)
C2	0.040 (3)	0.034 (3)	0.0221 (18)	0.000 (2)	-0.0045 (19)	-0.0037 (18)
C3	0.027 (2)	0.031 (2)	0.0155 (18)	0.000 (2)	-0.0032 (16)	0.0006 (17)
C4	0.047 (3)	0.041 (3)	0.025 (2)	0.000 (2)	-0.0038 (19)	-0.003 (2)
C5	0.040 (2)	0.031 (2)	0.0267 (19)	0.001 (2)	-0.0010 (18)	0.001 (2)
C6	0.032 (2)	0.032 (3)	0.0213 (19)	0.001 (2)	-0.0044 (17)	-0.0023 (19)
N1	0.047 (2)	0.046 (2)	0.0185 (15)	0.002 (2)	-0.0052 (15)	0.0025 (17)
N2	0.039 (2)	0.0258 (19)	0.0195 (16)	0.0027 (16)	-0.0035 (14)	-0.0014 (14)
O1	0.071 (2)	0.0297 (18)	0.0259 (15)	0.0076 (17)	-0.0039 (14)	-0.0017 (13)
Cl	0.0438 (6)	0.0314 (6)	0.0252 (5)	0.0015 (6)	-0.0035 (4)	-0.0029 (5)
O2	0.087 (3)	0.051 (2)	0.0344 (15)	-0.015 (2)	-0.0001 (17)	0.0134 (16)
O3	0.061 (2)	0.077 (3)	0.058 (2)	0.019 (2)	0.0025 (18)	-0.034 (2)
O4	0.082 (2)	0.035 (2)	0.0436 (17)	-0.0117 (18)	0.0001 (17)	-0.0096 (15)
O5	0.0471 (19)	0.080 (3)	0.0489 (18)	0.005 (2)	-0.0158 (16)	0.0135 (19)

Geometric parameters (\AA , $^\circ$)

C1—N1	1.342 (5)	C5—H5A	0.9300
C1—C2	1.368 (5)	C6—O1	1.196 (5)
C1—H1B	0.9300	C6—N2	1.363 (4)
C2—C3	1.382 (6)	C6—C6 ⁱ	1.555 (7)
C2—H2C	0.9300	N1—H1A	0.8600
C3—C5	1.391 (5)	N2—H2A	0.8600
C3—N2	1.399 (4)	Cl—O3	1.419 (3)
C4—N1	1.324 (5)	Cl—O4	1.423 (3)
C4—C5	1.375 (5)	Cl—O5	1.425 (3)
C4—H4B	0.9300	Cl—O2	1.439 (3)
N1—C1—C2	119.9 (4)	O1—C6—N2	127.6 (4)
N1—C1—H1B	120.0	O1—C6—C6 ⁱ	122.0 (4)
C2—C1—H1B	120.0	N2—C6—C6 ⁱ	110.4 (4)

C1—C2—C3	119.1 (4)	C4—N1—C1	122.7 (3)
C1—C2—H2C	120.5	C4—N1—H1A	118.6
C3—C2—H2C	120.5	C1—N1—H1A	118.6
C2—C3—C5	119.4 (3)	C6—N2—C3	125.3 (3)
C2—C3—N2	122.7 (3)	C6—N2—H2A	117.3
C5—C3—N2	117.9 (4)	C3—N2—H2A	117.3
N1—C4—C5	119.6 (4)	O3—Cl—O4	109.7 (2)
N1—C4—H4B	120.2	O3—Cl—O5	109.6 (2)
C5—C4—H4B	120.2	O4—Cl—O5	110.7 (2)
C4—C5—C3	119.2 (4)	O3—Cl—O2	109.7 (2)
C4—C5—H5A	120.4	O4—Cl—O2	108.32 (18)
C3—C5—H5A	120.4	O5—Cl—O2	108.8 (2)

Symmetry code: (i) $-x+1, -y+2, -z+2$.

Hydrogen-bond geometry (\AA , $^\circ$)

$D\text{—H}\cdots A$	$D\text{—H}$	$H\cdots A$	$D\cdots A$	$D\text{—H}\cdots A$
N1—H1A…O4	0.86	2.21	2.950 (4)	144
N1—H1A…O3 ⁱⁱ	0.86	2.35	2.966 (5)	129
N2—H2A…O2 ⁱⁱⁱ	0.86	2.14	2.975 (5)	162

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