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## Structure Reports

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Chad Eller, Bradley W. Smucker, Robert Kirgan, David M. Eichhorn and D. Paul Rillema*

Department of Chemistry, Wichita State
University, KS 67226, USA
Correspondence e-mail:
paul.rillema@wichita.edu

## Key indicators

Single-crystal X-ray study
$T=298 \mathrm{~K}$
Mean $\sigma(\mathrm{C}-\mathrm{C})=0.003 \AA$
$R$ factor $=0.048$
$\omega R$ factor $=0.136$
Data-to-parameter ratio $=12.1$
For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.
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## 5,5'-Dimethyl-2,2'-bipyrazine

The crystal structure of the title compound, $\mathrm{C}_{10} \mathrm{H}_{10} \mathrm{~N}_{4}$, a molecule synthesized as a dye, was determined by X-ray diffraction at room temperature. The molecule has a center of inversion located midway between the pyrazine units.

## Comment

5,5'-Dimethyl-2,2'-bipyrazine, (I), was prepared for attachment to $\mathrm{Ru}^{\mathrm{II}}$ and $\mathrm{Re}^{\mathrm{I}}$ atoms, for use as dyes in solid-state regenerative solar-energy cells. The compound acts as a conduit of electrons between the metal centers and the semiconducting dye anchors. The molecule is centrosymmetric, and thus essentially planar.

(I)

## Experimental

The title compound was prepared according to the procedures described by Lafferty \& Case (1967). 5-Methyl-2-pyrazinoic acid was dissolved in concentrated $\mathrm{NH}_{4} \mathrm{OH}$ and the solvent was removed by rotary evaporation, resulting in isolation of the ammonium pyrazinate salt. The solid was dried at 313 K in a vacuum oven and then dissolved in water. A saturated solution of copper(II) acetate was added in the ratio of one copper cation to two pyrazinate anions. The resulting blue-green solid was removed by vacuum filtration, washed with ether and dried at 373 K in a vacuum oven. The copper salt was placed in a pyrolysis vessel and heated between 558 and 584 K , displacing a yellow product that precipitated on the walls of the pyrolysis vessel. The product was dissolved in acetone and recrystallized from ether (yield 6\%; m.p. 420 K ). ${ }^{1} \mathrm{H}$ NMR $\left(\mathrm{CDCl}_{3}\right): \delta 2.595$ $(s, 6 \mathrm{H}), 8.688(s, 2 \mathrm{H}), 9.062(d, 2 \mathrm{H})$. Crystals of (I) suitable for X-ray diffraction were grown from an ether solution. The crystal of the compound does not contain any solvent molecules.


Figure 1
ORTEP-3 (Farrugia, 1997) representation of (I), with displacement ellipsoids at the $50 \%$ probability level.

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## Crystal data

| $\mathrm{C}_{10} \mathrm{H}_{10} \mathrm{~N}_{4}$ | $Z=1$ |
| :--- | :--- |
| $M_{r}=186.14$ | $D_{x}=1.334 \mathrm{Mg} \mathrm{m}^{-3}$ |
| Triclinic, $P \overline{1}$ | Mo $K \alpha$ radiation |
| $a=3.9580(7) \AA$ | Cell parameters from 24 |
| $b=5.9476(16) \AA$ | reflections |
| $c=10.107(2) \AA$ | $\theta=10-12^{\circ}$ |
| $\alpha=102.828(19)^{\circ}$ | $\mu=0.09 \mathrm{~mm}^{-1}$ |
| $\beta=92.100(17)^{\circ}$ | $T=298(2) \mathrm{K}$ |
| $\gamma=91.27(2)^{\circ}$ | Needle, white |
| $V=231.72(9) \AA^{3}$ | $0.4 \times 0.1 \times 0.1 \mathrm{~mm}$ |

Data collection

| Enraf-Nonius CAD-4 | $R_{\text {int }}=0.055$ |
| :--- | :--- |
| $\quad$ diffractometer | $\theta_{\max }=25.0^{\circ}$ |
| Non-profiled $\omega / 2 \theta$ scans | $h=-4 \rightarrow 4$ |
| Absorption correction: $\psi$-scan | $k=-7 \rightarrow 7$ |
| (North et al., 1968 ) | $l=-11 \rightarrow 11$ |
| $T_{\min }=0.802, T_{\max }=0.989$ | 3 standard reflections |
| 1613 measured reflections | frequency: 60 min |
| 811 independent reflections | intensity decay: $10 \%$ |
| 425 reflections with $I>2 \sigma(I)$ |  |

## Refinement

Refinement on $F^{2}$
H -atom parameters constrained
$R\left[F^{2}>2 \sigma\left(F^{2}\right)\right]=0.048$
$w=1 /\left[\sigma^{2}\left(F_{o}{ }^{2}\right)+(0.103 P)^{2}\right]$
where $P=\left(F_{o}{ }^{2}+2 F_{c}{ }^{2}\right) / 3$
$w R\left(F^{2}\right)=0.136$
$S=0.96$
811 reflections
67 parameters
$0.96 \AA$. The $U_{\text {iso }}(\mathrm{H})$ values of the methyl H atoms were taken as $1.5 U_{\text {eq }}(\mathrm{C})$; the $U_{\text {iso }}(\mathrm{H})$ values of the other H atoms were refined. The torsion angle of the methyl H atoms was estimated from the electron density, and was allowed to refine.

Data collection: CAD-4 EXPRESS (Enraf-Nonius, 1994); cell refinement: CAD-4 EXPRESS; data reduction: XCAD4 (Harms \& Wocadlo, 1995); program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: ORTEP-3 for Windows (Farrugia, 1997); software used to prepare material for publication: WinGX (Farrugia, 1999).

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