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## Key indicators

Single-crystal X-ray study
$T=296 \mathrm{~K}$
Mean $\sigma(\mathrm{C}-\mathrm{C})=0.003 \AA$
$R$ factor $=0.033$
$w R$ factor $=0.076$
Data-to-parameter ratio $=13.5$

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.
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## trans-4-(Ferrocenylideneamino)-1,5-dimethyl-2-phenyl-1H-pyrazol-3(2H)-one

The title compound, $\left[\mathrm{Fe}\left(\mathrm{C}_{5} \mathrm{H}_{5}\right)\left(\mathrm{C}_{17} \mathrm{H}_{16} \mathrm{~N}_{3} \mathrm{O}\right)\right]$, a new Schiff base containing a ferrocenyl group, has been characterized structurally. The central $\mathrm{C}-\mathrm{N}=\mathrm{C}-\mathrm{C}$ linkage has a nearplanar geometry, indicating extensive conjugation.

## Comment

Some Schiff bases bearing the ferrocenyl group and their complexes are excellent non-linear optical materials and liquid crystals (Colbert et al., 1995) because of their strong electron donors and electron-flow bridges. In the course of our investigation of the coordination of Schiff bases with transition metal salts, we observed that the title compound, (I), coordinates readily with $\mathrm{Ni}^{\mathrm{II}}$ and $\mathrm{Cu}^{\mathrm{II}}$ salts as an $N, O$-bidentate ligand. The crystal structure of (I) is reported here (Fig. 1).

(I)

In (I), all the bond lengths are within normal range (Allen et al., 1987). The N3-C12 bond length [1.277 (3) Å] confirms that it is a $\mathrm{C}=\mathrm{N}$ double bond. The dihedral angle between the five-membered pyrazole ring ( $\mathrm{N} 1 / \mathrm{N} 2 / \mathrm{C} 9 / \mathrm{C} 8 / \mathrm{C} 7$ ) and the plane


Figure 1
The structure of (I), showing the atom-labelling scheme. Displacement ellipsoids are drawn at the $30 \%$ probability level. H atoms are shown as small spheres of arbitrary radius.

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formed by atoms C8/N3/C12/C13 is $28.0(3)^{\circ}$. This deviation may be caused by the steric hindrance between atom O 1 and the H atom attached directly to C 12 . The $\mathrm{C} 8-\mathrm{N} 3-\mathrm{C} 12-\mathrm{C} 13$ torsion angle is $178.56(19)^{\circ}$.

## Experimental

A solution of ferrocenylaldehyde ( $0.214 \mathrm{~g}, 1 \mathrm{mmol}$ ) in absolute ethanol ( 10 ml ) was added dropwise to a solution of 4 -amino-1,5-dimethyl-2-phenyl-1,2-dihydropyrazol-3-one ( $0.203 \mathrm{~g}, 1 \mathrm{mmol}$ ) in absolute ethanol $(10 \mathrm{ml})$. The mixture was refluxed and stirred for 2 h and a yellow solid precipitated. The solid was isolated, washed three times with cold absolute ethanol and dried in a vacuum desiccator with anhydrous $\mathrm{CaCl}_{2}$ (yield: $81 \%$ ). A yellow single crystal suitable for X-ray analysis was obtained by slow evaporation of an ethyl acetate solution at room temperature over a period of a month. Analysis calculated for $\mathrm{C}_{22} \mathrm{H}_{21} \mathrm{FeN}_{3} \mathrm{O}: \mathrm{C} 66.18, \mathrm{H} 5.30, \mathrm{~N} 10.52 \%$; found: C 66.10, H 5.41, N $10.69 \%$.

## Crystal data

$\left[\mathrm{Fe}\left(\mathrm{C}_{5} \mathrm{H}_{5}\right)\left(\mathrm{C}_{17} \mathrm{H}_{16} \mathrm{~N}_{3} \mathrm{O}\right)\right.$ ]
$M_{r}=399.27$
Monoclinic, $P 2_{\mathrm{d}} / n$
$a=7.4749$ (6) А
$b=17.426$ (2) $\AA$
$c=14.443$ (2) $\AA$
$\beta=101.087$ (8) ${ }^{\circ}$
$V=1846.1$ (4) $\AA^{3}$
$Z=4$
Data collection
Siemens $P 4$ diffractometer $\omega$ scans
Absorption correction: multi-scan
(SADABS; Sheldrick, 1996)
$T_{\text {min }}=0.694, T_{\text {max }}=0.819$
3870 measured reflections
3335 independent reflections
2530 reflections with $I>2 \sigma(I)$

## $D_{x}=1.436 \mathrm{Mg} \mathrm{m}^{-3}$

Mo K $\alpha$ radiation
Cell parameters from 34 reflections
$\theta=6.2-14.5^{\circ}$
$\mu=0.83 \mathrm{~mm}^{-1}$
$T=296$ (2) K
Prism, yellow
$0.48 \times 0.32 \times 0.24 \mathrm{~mm}$

$$
\begin{aligned}
& R_{\text {int }}=0.022 \\
& \theta_{\max }=25.3^{\circ} \\
& h=0 \rightarrow 8 \\
& k=0 \rightarrow 20 \\
& l=-17 \rightarrow 17 \\
& 3 \text { standard reflections } \\
& \quad \text { every } 97 \text { reflections } \\
& \quad \text { intensity decay: } 5.6 \%
\end{aligned}
$$

## Refinement

Refinement on $F^{2}$
$R\left[F^{2}>2 \sigma\left(F^{2}\right)\right]=0.033$
$w R\left(F^{2}\right)=0.076$
$S=1.00$
3335 reflections
247 parameters
H -atom parameters constrained

$$
\begin{gathered}
w=1 /\left[\sigma^{2}\left(F_{o}^{2}\right)+(0.0366 P)^{2}\right] \\
\text { where } P=\left(F_{o}^{2}+2 F_{c}^{2}\right) / 3 \\
(\Delta / \sigma)_{\max }=0.001 \\
\Delta \rho_{\max }=0.27 \mathrm{e}^{-3} \\
\Delta \rho_{\min }=-0.27 \mathrm{e} \AA^{-3}
\end{gathered}
$$

Extinction correction: SHELXL
Extinction coefficient: 0.0066 (6)

Table 1
Selected geometric parameters ( $\left(\AA,{ }^{\circ}\right)$.

| $\mathrm{O} 1-\mathrm{C} 7$ | $1.232(3)$ | $\mathrm{C} 7-\mathrm{C} 8$ | $1.438(3)$ |
| :--- | :---: | :--- | :---: |
| $\mathrm{N} 3-\mathrm{C} 12$ | $1.277(3)$ | $\mathrm{C} 8-\mathrm{C} 9$ | $1.359(3)$ |
| $\mathrm{N} 3-\mathrm{C} 8$ | $1.407(3)$ | $\mathrm{C} 12-\mathrm{C} 13$ | $1.459(3)$ |
|  |  |  |  |
| $\mathrm{C} 12-\mathrm{N} 3-\mathrm{C} 8$ | $119.7(2)$ | $\mathrm{N} 3-\mathrm{C} 8-\mathrm{C} 7$ | $129.0(2)$ |
| $\mathrm{O} 1-\mathrm{C} 7-\mathrm{C} 8$ | $131.8(2)$ | $\mathrm{N} 3-\mathrm{C} 12-\mathrm{C} 13$ | $120.4(2)$ |
| $\mathrm{C} 9-\mathrm{C} 8-\mathrm{N} 3$ | $122.5(2)$ |  |  |
| $\mathrm{C} 7-\mathrm{N} 1-\mathrm{N} 2-\mathrm{C} 9$ | $-9.6(2)$ | $\mathrm{O} 1-\mathrm{C} 7-\mathrm{C} 8-\mathrm{N} 3$ | $-2.0(4)$ |
| $\mathrm{C} 6-\mathrm{N} 1-\mathrm{N} 2-\mathrm{C} 9$ | $-163.45(19)$ | $\mathrm{C} 8-\mathrm{N} 3-\mathrm{C} 12-\mathrm{C} 13$ | $178.56(19)$ |
| $\mathrm{C} 6-\mathrm{N} 1-\mathrm{C} 7-\mathrm{C} 8$ | $160.0(2)$ | $\mathrm{N} 3-\mathrm{C} 12-\mathrm{C} 13-\mathrm{C} 14$ | $26.9(3)$ |
| $\mathrm{O} 1-\mathrm{C} 7-\mathrm{C} 8-\mathrm{C} 9$ | $176.4(3)$ |  |  |



Figure 2
The crystal packing of (I), viewed along the $a$ axis. H atoms have been omitted.

All H atoms were initially located in a difference Fourier map. All H atoms were then constrained to an ideal geometry, with $\mathrm{C}-\mathrm{H}$ distances of $0.93-0.96 \AA$ and with $U_{\text {iso }}(\mathrm{H})=1.2 U_{\text {eq }}(\mathrm{C})$.

Data collection: XSCANS (Siemens, 1996); cell refinement: XSCANS; data reduction: SHELXTL (Sheldrick, 1997b); program(s) used to solve structure: SHELXS97 (Sheldrick, 1997a); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997a); molecular graphics: SHELXTL (Sheldrick, 1997b); software used to prepare material for publication: SHELXTL.

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