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## Structure Reports

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## Key indicators

Single-crystal X-ray study
$T=294 \mathrm{~K}$
Mean $\sigma(\mathrm{C}-\mathrm{C})=0.005 \AA$
$R$ factor $=0.039$
$w R$ factor $=0.100$
Data-to-parameter ratio $=17.0$
For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.
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## $N$-Methyl- $N$-(ferrocenylmethyl)aniline

The title compound, $\left[\mathrm{Fe}\left(\mathrm{C}_{5} \mathrm{H}_{5}\right)\left(\mathrm{C}_{13} \mathrm{H}_{14} \mathrm{~N}\right)\right]$, a tertiary arylamine bearing a ferrocenyl group was synthesized by reductive methylation of amines. The result reveals that the cyclopentadiene ring carrying the methylaniline residue is nearly perpendicular to the benzene plane [dihedral angle $=$ 92.1 (3) ${ }^{\circ}$.

## Comment

Ferrocene-containing compounds have been widely studied because of their potential application in catalysis, materials science, hydrometallurgy and as molecular devices (Beer et al., 1997). As a part of our ongoing investigations on cyclometalation of tertiary amines, the title compound, (I), has been prepared and we present here its crystal structure.

(I)

The molecular structure of (I) is shown in Fig. 1. The methyl C atom ( C 12 ) and the methylene C atom ( C 11 ) are displaced from the plane of the phenyl ring by 0.1309 (3) and


Figure 1
The structure of (I), showing the atom-labelling scheme. Displacement ellipsoids are drawn at the $35 \%$ probability level. H atoms are shown as small spheres of arbitrary radii.

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0.3913 (2) $\AA$, respectively. The cyclopentadiene ring carrying the methylaniline residue is nearly perpendicular to the benzene plane [dihedral angle $=92.1(3)^{\circ}$ ].

## Experimental

A solution of sodium cyanoborohydride ( $378 \mathrm{mg}, 6 \mathrm{mmol}$ ) in acetonitrile ( 20 ml ) was added to a strongly stirred solution of N (ferrocenylmethylidene) aniline ( $582 \mathrm{mg}, 2 \mathrm{mmol}$ ) and $37 \%$ aqueous formaldehyde ( $2 \mathrm{ml}, 25 \mathrm{mmol}$ ) in acetonitrile ( 30 ml ). The reaction mixture was stirred for 30 min . Glacial acetic acid was then added dropwise until the solution was neutral. Stirring was continued for a further 4 h . During this period, glacial acetic acid was added occasionally to maintain the pH at nearly neutral. The reaction was monitored by thin-layer chromatography to establish when it was complete. The reaction was quenched by pouring it into diethyl ether $(80 \mathrm{ml})$ and the organic layer was washed with $1 M \mathrm{KOH}$ and saturated aqueous sodium chloride ( 50 ml ). The diethyl ether solution was dried with $\mathrm{K}_{2} \mathrm{CO}_{3}$ and the solvent was removed in vacuo. The organic solid was purified by silica-gel column chromatography using ethyl acetate-hexane (3:1) giving a yield of $78 \%$. A yellow singlecrystal suitable for X-ray analysis was obtained by slow evaporation of a dichloromethane and petroleum solution at room temperature over a period of a week. Analysis calculated for $\mathrm{C}_{18} \mathrm{H}_{19} \mathrm{FeN}$ : C 70.84, H 6.27, N $4.59 \%$. found: C 63.65 , H 5.34, N $4.12 \%$.

## Crystal data

$\left[\mathrm{Fe}\left(\mathrm{C}_{5} \mathrm{H}_{5}\right)\left(\mathrm{C}_{13} \mathrm{H}_{14} \mathrm{~N}\right)\right]$
$M_{r}=305.19$
Monoclinic, $P 2_{1} / c$
$a=9.626(2) \AA$ 。
$b=11.675$ (2) A
$c=14.062(3) \AA$
$\beta=104.762(4)^{\circ}$
$V=1528.1(5) \AA^{3}$
$Z=4$

## Data collection

Bruker SMART CCD area-detector diffractometer
$\varphi$ and $\omega$ scans
Absorption correction: multi-scan
(SADABS; Sheldrick, 1996)
$T_{\text {min }}=0.704, T_{\text {max }}=0.807$
8371 measured reflections

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\(D_{x}=1.327 \mathrm{Mg} \mathrm{m}^{-3}\)
Mo \(K \alpha\) radiation
Cell parameters from 2691 reflections
\(\theta=2.2-24.4^{\circ}\)
\(\mu=0.98 \mathrm{~mm}^{-1}\)
\(T=294 \mathrm{~K}\)
Block, yellow
\(0.34 \times 0.30 \times 0.22 \mathrm{~mm}\)
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> 3089 independent reflections
> 2166 reflections with $I>2 \sigma(I)$
> $R_{\text {int }}=0.028$
> $\theta_{\max }=26.3^{\circ}$
> $h=-11 \rightarrow 11$
> $k=-14 \rightarrow 13$
> $l=-10 \rightarrow 17$

## Refinement

Refinement on $F^{2}$
$R\left[F^{2}>2 \sigma\left(F^{2}\right)\right]=0.039$
$w R\left(F^{2}\right)=0.100$
$S=1.05$
3089 reflections
182 parameters
H -atom parameters constrained

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

| N1-C13 | 1.388 (4) | C13-C14 | 1.401 (4) |
| :---: | :---: | :---: | :---: |
| N1-C12 | 1.451 (4) | C14-C15 | 1.371 (5) |
| N1-C11 | 1.459 (4) | C15-C16 | 1.367 (5) |
| C10-C11 | 1.516 (4) | C16-C17 | 1.370 (5) |
| C13-C18 | 1.401 (4) | C17-C18 | 1.378 (5) |
| C13-N1-C12 | 119.7 (3) | C6-C10-C9 | 106.7 (3) |
| C13-N1-C11 | 120.4 (3) | C6-C10-C11 | 126.5 (3) |
| C12-N1-C11 | 113.6 (3) | C9-C10-C11 | 126.8 (3) |
| C2-C1-C5 | 107.1 (4) | N1-C11-C10 | 113.4 (3) |
| C3-C2-C1 | 109.4 (3) | N1-C13-C18 | 122.3 (3) |
| C4-C3-C2 | 107.9 (4) | N1-C13-C14 | 121.0 (3) |
| C3-C4-C5 | 109.1 (4) | C18-C13-C14 | 116.6 (3) |
| C4-C5-C1 | 106.5 (3) | C15-C14-C13 | 120.9 (3) |
| C7-C6-C10 | 109.0 (3) | C16-C15-C14 | 121.9 (4) |
| C8-C7-C6 | 108.0 (3) | C15-C16-C17 | 118.1 (4) |
| C7-C8-C9 | 108.1 (3) | C16-C17-C18 | 121.5 (4) |
| C8-C9-C10 | 108.2 (3) | C17-C18-C13 | 120.9 (3) |
| C13-N1-C11-C10 | -86.0 (4) | C11-N1-C13-C18 | -29.5 (5) |
| C12-N1-C11-C10 | 66.1 (4) | C12-N1-C13-C14 | 4.0 (5) |
| C6-C10-C11-N1 | 71.4 (4) | C11-N1-C13-C14 | 154.5 (3) |
| C9-C10-C11-N1 | -107.5 (3) | N1-C13-C14-C15 | 173.9 (3) |
| C12-N1-C13-C18 | -180.0 (3) | N1-C13-C18-C17 | -174.6 (3) |

All H atoms were located in a difference Fourier map. The methyl H atoms were constrained to an ideal geometry, with $\mathrm{C}-\mathrm{H}$ distances of $0.96 \AA$ and with $U_{\text {iso }}(\mathrm{H})=1.5 U_{\text {eq }}(\mathrm{C})$, but each group was allowed to rotate freely about its $\mathrm{C}-\mathrm{C}$ bond. All other H atoms were placed in geometrically idealized positions and constrained to ride on their parent atoms, with $\mathrm{C}-\mathrm{H}$ distances in the range $0.93-0.97 \AA$ and with $U_{\text {iso }}(\mathrm{H})=1.2 U_{\text {eq }}(\mathrm{C})$.

Data collection: SMART (Bruker, 1997); cell refinement: SAINT (Bruker, 1997); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: SHELXTL (Bruker, 1997); software used to prepare material for publication: SHELXTL.

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## References

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