## metal-organic papers

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#### **Key indicators**

Single-crystal X-ray study T = 100 KMean  $\sigma$ (C–C) = 0.002 Å R factor = 0.026 wR factor = 0.075 Data-to-parameter ratio = 25.9

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

# 1,12-Diferrocenyldodecane at 100 K

1,12-Diferrocenyldodecane,  $[Fe_2(C_5H_5)_2(C_{22}H_{32})]$ , was synthesized from ferrocene and 1,12-dodecanedioyl chloride, followed by Clemmensen reduction. The single-crystal structure was determined at 100 K by X-ray diffraction and the spectroscopic and cyclic voltammetric data of 1,12diferrocenyldodecane and its precursor are reported. Received 11 April 2005 Accepted 28 April 2005 Online 14 May 2005

#### Comment

Ferrocene-containing compounds have been extensively studied due to their unique structural and electrochemical properties. Ferrocenes have also been investigated with regard to their potential applications in chemical sensing, redox catalysts, ferromagnetism and self-assembled monolayers (Togni & Hayashi, 1995; Sawamura & Ito, 1992; Nicolosi *et al.*, 1994; Sammakia *et al.*, 1995; Beer *et al.*, 1993; Zhang *et al.*, 1995; Chidsey *et al.*, 1990; Creager & Rowe, 1997) As an intermediate in the synthesis of self-assembled monolayers with multiple redox centers, 1,12-diferrocenyldodecane, (1), was synthesized *via* Clemmensen reduction of its precursor 1,12-diferrocenyldodecane-1,12-dione, (2), and its solid-state structure was established by X-ray diffraction at 100 K.



1,12-Diferrocenyldodecane crystallizes in the monoclinic space group  $P2_1/c$  with two molecules in the unit cell (Fig. 1). The center of each molecule is located on a crystallographic inversion center halfway between C16 and C16<sup>i</sup> [symmetry code: (i) -x + 1, -y + 1, -z + 2]. The dodecane chain forms an extended zigzag chain. All torsion angles in the chain are close to  $180^{\circ}$  [between 175.70 (6)° for C12···C14···C16 and 179.39 (6)° for C14···C16···C15<sup>i</sup>].

The C–C single bonds of the dodecane chain close to the ferrocene unit alternate slightly; the C10–C11 bond length between the ferrocene unit and the first methylene C atom is 1.501 (1) Å and that of C11–C12 is 1.529 (1) Å. All other C–C bond distances within the dodecane chain are identical within experimental error: 1.525 (1), 1.525 (1), 1.526 (1), 1.523 (1) and 1.525 (2) Å for C12–C13, C13–C14, C14–C15, C15–C16, and C16–C16<sup>i</sup>, respectively. The bond distances within the ferrocene units are unexceptional, with the unsubstituted ring having basically identical C–C bond distances of 1.424 (2) Å, and C–C distances of the substituted

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### Figure 1

The molecular structure of (1), showing 50% probability displacement ellipsoids. [symmetry code: (i) -x + 1, -y + 1, -z + 2]

ring being between 1.429 (2) and 1.433 (2) Å.

The Fe1...Fe1<sup>i</sup> distance within one molecule is 18.7196 (9) Å in the solid state. The closest  $Fe \cdot \cdot \cdot Fe$  distances between different molecules are 6.3972 (3) Å for Fe1...Fe1<sup>ii</sup> [symmetry code: (ii)  $x, \frac{1}{2} - y, z + \frac{1}{2}$ ] and 7.1275 (3) Å for Fe1...Fe1<sup>iii</sup> [symmetry code: (iii)  $x, \frac{3}{2} - y, z + \frac{1}{2}$ ].

### **Experimental**

1,12-Diferrocenyldodecane-1,12-dione, (2), was synthesized from ferrocene and 1,12-dodecanedioyl chloride via the method of Chidsey et al. (1990) and was reduced to the title compound, (1), by Clemmensen reduction, as described by Creager & Rowe (1997). Ferrocene (35.00 g, 0.188 mmol) was dissolved at 273 K in CH<sub>2</sub>Cl<sub>2</sub> (250 ml). 1,12-Dodecanedioyl chloride (25.00 g, 0.0935 mmol) was added and the resulting solution was stirred for 5 min. Anhydrous AlCl<sub>3</sub> (27.50 g, 0.207 mmol) was added and the solution was allowed to warm to room temperature and stirred for 1.5 h. De-ionized water (300 ml) was added dropwise and the mixture was stirred for an additional 1.5 h. The organic phase was washed with five portions of de-ionized water, then three times with brine, and dried over anhydrous sodium sulfate. The solvent was removed in vacuo. The crude product was purified via column chromatography on silica with 80/20% hexanes/ethyl acetate as the eluent. The second band to elute  $(R_F = 0.75)$  contained the desired product, (2), in 12.2% yield. <sup>1</sup>H NMR (400 MHz, CDCl<sub>3</sub>): δ 1.322 [m, 12H, (CH<sub>2</sub>)<sub>6</sub>], 1.701 (m, 4H, COCH<sub>2</sub>CH<sub>2</sub>), 2.692 (t, 4H, COCH<sub>2</sub>), 4.194 (s, 10H, C<sub>5</sub>H<sub>5</sub>), 4.487 (t, 4H, C<sub>5</sub>H<sub>4</sub>), 4.780 (t, 4H, C<sub>5</sub>H<sub>4</sub>). MS: calculated for  $(1^+)$  566 m/z, found 566 m/z. CV [1.6 mm diameter Pt disc working, Pt wire counter, nonaqueous Ag/Ag<sup>+</sup> reference electrodes, 1.0 mM (2), 0.1 M {(*n*-C<sub>4</sub>H<sub>9</sub>)N}ClO<sub>4</sub> in CH<sub>3</sub>CN]: reversible single, two-electron oxidation

centered at  $E^{\circ} = +266$  mV. 1,12-Diferrocenyldodecane-1,12-dione, (2) (9.02 g, 0.0158 mol), dissolved in toluene (800 ml), was mixed with freshly prepared zinc/mercury amalgam [mercuric chloride (1.06 g, 0.039 mol), zinc dust (15.79 g, 0.024 mol), de-ionized water (30 ml) and 12.1 M HCl (40 ml)] and was refluxed under static argon for 20 h. Two additional 30 ml portions of 12.1 M HCl were added with an addition funnel after 6 and 12 h under reflux. Upon cooling, the organic phase was washed three times with de-ionized water and three times with brine, and was dried over anhydrous sodium sulfate. The solvent was removed in vacuo. The crude product was purified via column chromatography on silica with 75/25% hexanes/ethyl acetate as the eluent. The first band to elute  $(R_F = 0.9)$  contained the desired product, viz. 1,12-diferrocenyldodecane, (1). <sup>1</sup>H NMR (400 MHz, CDCl<sub>3</sub>): δ 1.199 [m, 16H, (CH<sub>2</sub>)<sub>8</sub>], 1.417 (m, 4H, CpCH<sub>2</sub>CH<sub>2</sub>), 2.228 (t, 4H, CpCH<sub>2</sub>), 3.976 (s, 4H, C<sub>5</sub>H<sub>4</sub>), 3.996 (s, 4H,  $C_5H_4$ ), 4.032 (s, 10H,  $C_5H_5$ ). MS: calculated for (2<sup>+</sup>) 538 m/z, found 538 m/z. CV [1.6 mm diameter Pt disc working, Pt wire counter, nonaqueous Ag/Ag<sup>+</sup> reference electrodes, 1.0 mM (1), 0.1 M {(n- $C_4H_9$ NClO<sub>4</sub> in CH<sub>3</sub>CN]: reversible single, two-electron oxidation centered at  $E^{\circ} = -25$  mV. Single crystals of (1) were grown from a 75/ 25% hexanes/ethyl acetate solution by slow evaporation.

> $D_x = 1.363 \text{ Mg m}^{-3}$ Mo  $K\alpha$  radiation

> > reflections

T = 100 (2) K

Block, red

 $R_{\rm int} = 0.018$  $\theta_{\text{max}} = 30.5^{\circ}$  $h = -22 \rightarrow 22$ 

 $k = -10 \rightarrow 10$ 

 $l = -15 \rightarrow 15$ 

 $\theta = 2.7 - 30.5^{\circ}$  $\mu = 1.12~\mathrm{mm}^{-1}$ 

Cell parameters from 6072

 $0.47 \times 0.29 \times 0.25 \text{ mm}$ 

3988 independent reflections

3769 reflections with  $I > 2\sigma(I)$ 

 $(0.0461P)^2$ 

 $+ 2F_c^2)/3$ 

#### Crystal data

[Fe<sub>2</sub>(C<sub>5</sub>H<sub>5</sub>)<sub>2</sub>(C<sub>22</sub>H<sub>32</sub>)]  $M_r = 538.36$ Monoclinic,  $P2_1/c$ a = 16.0203 (9) Åb = 7.5367 (4) Åc = 11.1773 (6) Å  $\beta = 103.597 (1)^{\circ}$  $V = 1311.73 (12) \text{ Å}^3$ Z = 2

#### Data collection

Bruker SMART APEX CCD diffractometer  $\omega$  scans Absorption correction: multi-scan (SADABS in SAINT-Plus; Bruker, 2003)  $T_{\min} = 0.644, \ T_{\max} = 0.75$ 15 135 measured reflections

#### Refinement

Refinement on $F^2$	$w = 1/[\sigma^2(F_o^2) +$
$R[F^2 > 2\sigma(F^2)] = 0.026$	+ 0.3363P]
$wR(F^2) = 0.075$	where $P = (F_{i})$
S = 1.07	$(\Delta/\sigma)_{\rm max} = 0.001$
3988 reflections	$\Delta \rho_{\rm max} = 0.45 \ {\rm e}$
154 parameters	$\Delta \rho_{\rm min} = -0.37  {\rm e}$
H-atom parameters constrained	

All H atoms were placed in calculated positions, with a C-H bond distance of 0.99 (methylene) or 1.00 Å (Cp), and were refined with isotropic displacement parameters 1.2 times  $U_{eq}$  of the parent C atom. The s.u. values of the cell parameters are taken from the software, recognizing that the values are unreasonably small (Herbstein. 2000).

Data collection: SMART (Bruker, 2002); cell refinement: SAINT-Plus (Bruker, 2003); data reduction: SAINT-Plus; program(s) used to solve structure: SHELXTL (Bruker, 2000); program(s) used to refine structure: SHELXTL; molecular graphics: SHELXTL; software used to prepare material for publication: SHELXTL.

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