

Ferrocene-1-carbaldehyde 4-ethylthiosemicarbazone

M. R. Vikneswaran,^a Siang Guan Teoh,^a Chin Sing Yeap^{b,‡} and Hoong-Kun Fun^{b,*§}

^aSchool of Chemical Sciences, Universiti Sains Malaysia, 11800 USM, Penang, Malaysia, and ^bX-ray Crystallography Unit, School of Physics, Universiti Sains Malaysia, 11800 USM, Penang, Malaysia

Correspondence e-mail: hkfun@usm.my

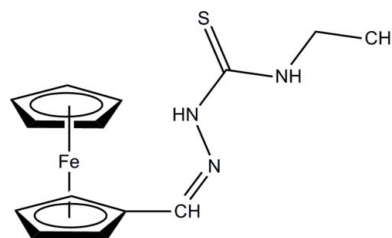
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Key indicators: single-crystal X-ray study; $T = 100$ K; mean $\sigma(\text{C}-\text{C}) = 0.007$ Å; R factor = 0.073; wR factor = 0.170; data-to-parameter ratio = 22.6.

The asymmetric unit of title compound, $[\text{Fe}(\text{C}_5\text{H}_5)(\text{C}_9\text{H}_{12}\text{N}_3\text{S})]$, contains two crystallographically independent molecules, *A* and *B*. The two cyclopentadienyl (Cp) rings are parallel to each other in both molecules, forming dihedral angles of 2.3 (3) and 1.0 (3)°, respectively, and adopt an eclipsed conformation. The mean plane of the semicarbazone group is twisted slightly away from the attached Cp ring in both molecules, the dihedral angles between the mean plane and the Cp ring being 15.3 (2) and 10.8 (2)°. The ethyl group in molecule *A* is coplanar with the mean plane of the semicarbazone group [C–N–C–C torsion angle = -175.2 (4)°], whereas it is nearly perpendicular in molecule *B* [C–N–C–C torsion angle = 84.8 (6)°]. In the crystal structure, intermolecular N–H...S hydrogen bonds link the molecules into dimers. These dimers are further linked into chains *via* intermolecular C–H...S hydrogen bonds. The crystal studied was a non-merohedral twin with a refined ratio of the twin components of 0.265 (2):0.735 (2).

Related literature

For related structures, see: Vikneswaran *et al.* (2009, 2010). For the preparation of the title compound, see: Casas *et al.* (2004). For the stability of the temperature controller used for the data collection, see: Cosier & Glazer (1986).



Experimental

Crystal data

$[\text{Fe}(\text{C}_5\text{H}_5)(\text{C}_9\text{H}_{12}\text{N}_3\text{S})]$
 $M_r = 315.22$
 Triclinic, $P\bar{1}$
 $a = 7.4432$ (3) Å
 $b = 10.6906$ (5) Å
 $c = 18.4616$ (9) Å
 $\alpha = 77.975$ (3)°
 $\beta = 83.807$ (3)°
 $\gamma = 78.076$ (3)°
 $V = 1402.56$ (11) Å³
 $Z = 4$
 Mo $K\alpha$ radiation
 $\mu = 1.21$ mm⁻¹
 $T = 100$ K
 $0.29 \times 0.16 \times 0.09$ mm

Data collection

Bruker SMART APEXII CCD area-detector diffractometer
 Absorption correction: multi-scan (SADABS; Bruker, 2009)
 $T_{\min} = 0.723$, $T_{\max} = 0.901$
 8184 measured reflections
 8184 independent reflections
 6947 reflections with $I > 2\sigma(I)$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.073$
 $wR(F^2) = 0.170$
 $S = 1.07$
 8184 reflections
 362 parameters
 H atoms treated by a mixture of independent and constrained refinement
 $\Delta\rho_{\max} = 3.94$ e Å⁻³
 $\Delta\rho_{\min} = -1.22$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

| <i>D</i> –H... <i>A</i> | <i>D</i> –H | H... <i>A</i> | <i>D</i> ... <i>A</i> | <i>D</i> –H... <i>A</i> |
|---|-------------|---------------|-----------------------|-------------------------|
| N2 <i>A</i> –H2 <i>NA</i> ...S1 <i>A</i> ⁱ | 0.82 (6) | 2.59 (6) | 3.387 (4) | 164 (5) |
| N2 <i>B</i> –H2 <i>NB</i> ...S1 <i>B</i> ⁱⁱ | 0.89 (9) | 2.55 (9) | 3.430 (5) | 170 (5) |
| C4 <i>A</i> –H4 <i>AA</i> ...S1 <i>B</i> ⁱⁱⁱ | 0.98 | 2.79 | 3.715 (4) | 157 |

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

Data collection: APEX2 (Bruker, 2009); cell refinement: SAINT (Bruker, 2009); data reduction: SAINT; program(s) used to solve structure: SHELXTL (Sheldrick, 2008); program(s) used to refine structure: SHELXTL; molecular graphics: SHELXTL; software used to prepare material for publication: SHELXTL (Sheldrick, 2008) and PLATON (Spek, 2009).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: RZ2450).

‡ Thomson Reuters ResearcherID: A-5523-2009.

§ Thomson Reuters ResearcherID: A-3561-2009.

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supplementary materials

Acta Cryst. (2010). E66, m697-m698 [doi:10.1107/S1600536810018209]

Ferrocene-1-carbaldehyde 4-ethylthiosemicarbazone

M. R. Vikneswaran, S. G. Teoh, C. S. Yeap and H.-K. Fun

Comment

As a continuation of our research related to ferrocenyl thiosemicarbazones and its metal complexes, herein we report the crystal structure of formylferrocene 4-ethylthiosemicarbazone.

The asymmetric unit of title compound consists of two crystallographically independent molecules, *A* and *B* (Fig. 1). The geometric parameter are comparable to those observed in its closely related structures (Vikneswaran *et al.*, 2009, 2010). The Cp rings of each ferrocene residue are parallel, with dihedral angles of Cp1/Cp2 [C1A–C5A/C6A–C10A] = 2.3 (3)° and Cp3/Cp4 [C1B–C5B/C6B–C10B] = 1.0 (3)°. The Cp rings in both molecules adopt an eclipsed conformation [average torsion angles for C–Cg–Cg–C of 5.89 and 6.14°]. The mean plane of the semicarbazone group is slightly twisted away from the attached Cp rings in both molecules, the dihedral angles between the mean plane and the Cp ring being 15.3 (2) and 10.8 (2)° respectively. The ethyl group in molecule *A* is coplanar with the mean plane of semicarbazone group [torsion angle of C12A–N3A–C13A–C14B = -175.2 (4)°] whereas it is nearly perpendicular to the semicarbazone group [torsion angle of C12B–N3B–C13B–C14B = 84.8 (6)°] in molecule *B*.

In the crystal structure, intermolecular N2A–H2NA...S1A and N2B–H2NB...S1B hydrogen bonds link the molecules into dimers. These dimers are linked into one-dimensional chain *via* intermolecular C4A–H4AA...S1B hydrogen bonds (Fig. 2, Table 1).

Experimental

Formylferrocene 4-ethylthiosemicarbazone was prepared as described by Casas *et al.* (2004). The single crystals were grown from a CH₂Cl₂/n-CH₆H₁₄ (1:1 v/v) solution at room temperature in the dark.

Refinement

N bound H-atoms were located from difference Fourier map and refined freely. The rest of H-atoms were placed in calculated positions, with C–H = 0.93–0.98 Å and refined using a riding model, with $U_{\text{iso}}(\text{H}) = 1.2$ or $1.5U_{\text{eq}}(\text{C})$. Rotating-group model were applied for methyl group. The highest residual density peak is located 0.88 Å from atom Fe1B and the deepest hole is located 1.32 Å from atom C12B. The crystal studied is a non-merohedral twin with the refined ratio of twin components of 0.265 (2):0.735 (2).

Figures

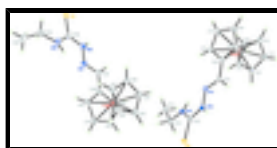


Fig. 1. The molecular structure of the title compound, showing 50% probability displacement ellipsoids and the atom-numbering scheme.

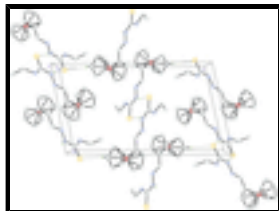


Fig. 2. The crystal packing of the title compound, viewed along the *a* axis, showing the molecules link into 1-D chains. Hydrogen atoms not involved in the hydrogen-bonding (dashed lines) are omitted for clarity.

Ferrocene-1-carbaldehyde 4-ethylthiosemicarbazone

Crystal data

| | |
|---|---|
| [Fe(C ₅ H ₅)(C ₉ H ₁₂ N ₃ S)] | <i>Z</i> = 4 |
| <i>M_r</i> = 315.22 | <i>F</i> (000) = 656 |
| Triclinic, <i>P</i> $\bar{1}$ | <i>D_x</i> = 1.493 Mg m ⁻³ |
| Hall symbol: -P 1 | Mo <i>K</i> α radiation, λ = 0.71073 Å |
| <i>a</i> = 7.4432 (3) Å | Cell parameters from 9974 reflections |
| <i>b</i> = 10.6906 (5) Å | θ = 2.3–30.0° |
| <i>c</i> = 18.4616 (9) Å | μ = 1.21 mm ⁻¹ |
| α = 77.975 (3)° | <i>T</i> = 100 K |
| β = 83.807 (3)° | Block, brown |
| γ = 78.076 (3)° | 0.29 × 0.16 × 0.09 mm |
| <i>V</i> = 1402.56 (11) Å ³ | |

Data collection

| | |
|---|---|
| Bruker SMART APEXII CCD area-detector diffractometer | 8184 independent reflections |
| Radiation source: fine-focus sealed tube graphite | 6947 reflections with <i>I</i> > 2σ(<i>I</i>) |
| φ and ω scans | <i>R</i> _{int} = 0.0000 |
| Absorption correction: multi-scan (<i>SADABS</i> ; Bruker, 2009) | θ _{max} = 30.1°, θ _{min} = 1.1° |
| <i>T</i> _{min} = 0.723, <i>T</i> _{max} = 0.901 | <i>h</i> = -10→10 |
| 8184 measured reflections | <i>k</i> = -14→15 |
| | <i>l</i> = -11→25 |

Refinement

| | |
|-------------------------------------|--|
| Refinement on <i>F</i> ² | Primary atom site location: structure-invariant direct methods |
| Least-squares matrix: full | Secondary atom site location: difference Fourier map |
| $R[F^2 > 2\sigma(F^2)] = 0.073$ | Hydrogen site location: inferred from neighbouring sites |
| $wR(F^2) = 0.170$ | H atoms treated by a mixture of independent and constrained refinement |
| <i>S</i> = 1.07 | $w = 1/[\sigma^2(F_o^2) + (0.0564P)^2 + 6.6396P]$ |
| 8184 reflections | where $P = (F_o^2 + 2F_c^2)/3$ |
| 362 parameters | (Δ/σ) _{max} < 0.001 |
| | Δρ _{max} = 3.94 e Å ⁻³ |

0 restraints

$$\Delta\rho_{\min} = -1.22 \text{ e \AA}^{-3}$$

Special details

Experimental. The crystal was placed in the cold stream of an Oxford Cryosystems Cobra open-flow nitrogen cryostat (Cosier & Glazer, 1986) operating at 100.0 (1) K.

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

| | <i>x</i> | <i>y</i> | <i>z</i> | $U_{\text{iso}}^*/U_{\text{eq}}$ |
|------|--------------|--------------|-------------|----------------------------------|
| Fe1A | 0.13896 (8) | -0.05245 (6) | 0.63951 (3) | 0.01124 (13) |
| S1A | 0.29669 (15) | 0.61964 (10) | 0.42316 (6) | 0.0175 (2) |
| N1A | 0.2762 (5) | 0.2535 (3) | 0.5056 (2) | 0.0145 (7) |
| N2A | 0.3209 (5) | 0.3757 (3) | 0.4940 (2) | 0.0150 (7) |
| N3A | 0.0898 (5) | 0.4487 (3) | 0.4141 (2) | 0.0140 (7) |
| C1A | -0.1290 (6) | 0.0406 (4) | 0.6505 (2) | 0.0171 (8) |
| H1AA | -0.2002 | 0.0985 | 0.6108 | 0.021* |
| C2A | -0.1151 (6) | -0.0966 (4) | 0.6707 (2) | 0.0174 (8) |
| H2AA | -0.1754 | -0.1498 | 0.6474 | 0.021* |
| C3A | 0.0008 (6) | -0.1433 (4) | 0.7305 (2) | 0.0201 (9) |
| H3AA | 0.0348 | -0.2343 | 0.7555 | 0.024* |
| C4A | 0.0590 (7) | -0.0348 (5) | 0.7480 (2) | 0.0211 (9) |
| H4AA | 0.1403 | -0.0376 | 0.7869 | 0.025* |
| C5A | -0.0221 (6) | 0.0788 (5) | 0.6976 (3) | 0.0203 (9) |
| H5AA | -0.0052 | 0.1679 | 0.6959 | 0.024* |
| C6A | 0.2245 (5) | -0.0166 (4) | 0.5289 (2) | 0.0137 (7) |
| H6AA | 0.1491 | 0.0332 | 0.4883 | 0.016* |
| C7A | 0.2465 (6) | -0.1531 (4) | 0.5565 (2) | 0.0151 (8) |
| H7AA | 0.1878 | -0.2136 | 0.5384 | 0.018* |
| C8A | 0.3655 (6) | -0.1867 (4) | 0.6163 (2) | 0.0168 (8) |
| H8AA | 0.4035 | -0.2740 | 0.6458 | 0.020* |
| C9A | 0.4196 (5) | -0.0705 (4) | 0.6252 (2) | 0.0154 (8) |
| H9AA | 0.5011 | -0.0638 | 0.6620 | 0.018* |
| C10A | 0.3323 (5) | 0.0348 (4) | 0.5711 (2) | 0.0123 (7) |
| C11A | 0.3566 (5) | 0.1695 (4) | 0.5581 (2) | 0.0146 (8) |
| H11A | 0.4304 | 0.1947 | 0.5879 | 0.018* |
| C12A | 0.2306 (5) | 0.4739 (4) | 0.4442 (2) | 0.0125 (7) |
| C13A | -0.0243 (6) | 0.5431 (4) | 0.3617 (2) | 0.0151 (8) |
| H13A | 0.0495 | 0.5685 | 0.3164 | 0.018* |

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|------|--------------|---------------|--------------|--------------|
| H13B | -0.0741 | 0.6203 | 0.3825 | 0.018* |
| C14A | -0.1810 (6) | 0.4851 (5) | 0.3440 (3) | 0.0223 (9) |
| H14A | -0.2522 | 0.5466 | 0.3073 | 0.033* |
| H14B | -0.2583 | 0.4652 | 0.3883 | 0.033* |
| H14C | -0.1314 | 0.4067 | 0.3253 | 0.033* |
| Fe1B | 0.30485 (8) | 0.46654 (6) | 0.14382 (3) | 0.01438 (14) |
| S1B | 0.24370 (18) | -0.06556 (11) | -0.06872 (7) | 0.0224 (2) |
| N1B | 0.2135 (5) | 0.2488 (4) | 0.0132 (2) | 0.0195 (8) |
| N2B | 0.1830 (6) | 0.1303 (4) | 0.0031 (2) | 0.0223 (8) |
| N3B | 0.4293 (6) | 0.1273 (4) | -0.0817 (2) | 0.0252 (9) |
| C1B | 0.5691 (6) | 0.3613 (5) | 0.1531 (2) | 0.0197 (9) |
| H1BA | 0.6511 | 0.3301 | 0.1127 | 0.024* |
| C2B | 0.5528 (6) | 0.4837 (4) | 0.1738 (3) | 0.0189 (9) |
| H2BA | 0.6230 | 0.5516 | 0.1507 | 0.023* |
| C3B | 0.4170 (6) | 0.4906 (4) | 0.2346 (3) | 0.0187 (8) |
| H3BA | 0.3781 | 0.5635 | 0.2609 | 0.022* |
| C4B | 0.3493 (6) | 0.3726 (4) | 0.2500 (2) | 0.0190 (8) |
| H4BA | 0.2539 | 0.3502 | 0.2887 | 0.023* |
| C5B | 0.4432 (7) | 0.2922 (4) | 0.2007 (3) | 0.0209 (9) |
| H5BA | 0.4231 | 0.2052 | 0.1987 | 0.025* |
| C6B | 0.2535 (6) | 0.5058 (5) | 0.0337 (2) | 0.0195 (9) |
| H6BA | 0.3413 | 0.4832 | -0.0072 | 0.023* |
| C7B | 0.2245 (7) | 0.6241 (4) | 0.0616 (3) | 0.0221 (9) |
| H7BA | 0.2895 | 0.6967 | 0.0433 | 0.026* |
| C8B | 0.0860 (6) | 0.6177 (4) | 0.1209 (3) | 0.0226 (9) |
| H8BA | 0.0391 | 0.6850 | 0.1506 | 0.027* |
| C9B | 0.0287 (6) | 0.4950 (5) | 0.1302 (3) | 0.0205 (9) |
| H9BA | -0.0648 | 0.4638 | 0.1672 | 0.025* |
| C10B | 0.1320 (6) | 0.4262 (4) | 0.0763 (2) | 0.0171 (8) |
| C11B | 0.1143 (6) | 0.2982 (4) | 0.0654 (3) | 0.0193 (9) |
| H11B | 0.0320 | 0.2527 | 0.0960 | 0.023* |
| C12B | 0.2896 (7) | 0.0721 (5) | -0.0494 (3) | 0.0204 (9) |
| C13B | 0.5572 (7) | 0.0852 (5) | -0.1416 (3) | 0.0284 (11) |
| H13C | 0.5809 | -0.0091 | -0.1340 | 0.034* |
| H13D | 0.6731 | 0.1122 | -0.1398 | 0.034* |
| C14B | 0.4841 (9) | 0.1411 (6) | -0.2179 (3) | 0.0374 (13) |
| H14D | 0.5742 | 0.1130 | -0.2553 | 0.056* |
| H14E | 0.4593 | 0.2345 | -0.2256 | 0.056* |
| H14F | 0.3727 | 0.1109 | -0.2208 | 0.056* |
| H2NA | 0.416 (8) | 0.390 (5) | 0.507 (3) | 0.016 (13)* |
| H2NB | 0.073 (12) | 0.109 (8) | 0.015 (5) | 0.07 (3)* |
| H3NB | 0.458 (11) | 0.186 (7) | -0.053 (4) | 0.05 (2)* |
| H3NA | 0.046 (8) | 0.384 (6) | 0.437 (3) | 0.026 (15)* |

Atomic displacement parameters (\AA^2)

| | U^{11} | U^{22} | U^{33} | U^{12} | U^{13} | U^{23} |
|------|------------|------------|------------|-------------|-------------|-------------|
| Fe1A | 0.0099 (3) | 0.0130 (3) | 0.0111 (3) | -0.0036 (2) | -0.0009 (2) | -0.0015 (2) |

| | | | | | | |
|------|-------------|-------------|-------------|--------------|--------------|--------------|
| S1A | 0.0198 (5) | 0.0117 (5) | 0.0231 (5) | -0.0066 (4) | -0.0057 (4) | -0.0027 (4) |
| N1A | 0.0116 (15) | 0.0160 (17) | 0.0175 (17) | -0.0050 (13) | -0.0004 (13) | -0.0046 (13) |
| N2A | 0.0147 (16) | 0.0125 (16) | 0.0189 (17) | -0.0042 (13) | -0.0027 (13) | -0.0028 (13) |
| N3A | 0.0161 (16) | 0.0100 (15) | 0.0162 (17) | -0.0061 (13) | -0.0021 (13) | 0.0009 (13) |
| C1A | 0.0122 (17) | 0.018 (2) | 0.020 (2) | -0.0018 (15) | 0.0021 (15) | -0.0048 (16) |
| C2A | 0.0162 (18) | 0.018 (2) | 0.021 (2) | -0.0094 (15) | 0.0027 (16) | -0.0045 (16) |
| C3A | 0.025 (2) | 0.018 (2) | 0.016 (2) | -0.0071 (17) | 0.0021 (17) | 0.0007 (16) |
| C4A | 0.025 (2) | 0.028 (2) | 0.0121 (19) | -0.0105 (18) | 0.0030 (16) | -0.0041 (17) |
| C5A | 0.026 (2) | 0.020 (2) | 0.018 (2) | -0.0067 (17) | 0.0034 (17) | -0.0086 (17) |
| C6A | 0.0102 (17) | 0.0179 (19) | 0.0135 (18) | -0.0039 (14) | 0.0000 (14) | -0.0032 (15) |
| C7A | 0.0132 (17) | 0.0169 (19) | 0.0153 (19) | -0.0026 (14) | -0.0007 (14) | -0.0037 (15) |
| C8A | 0.0136 (18) | 0.0140 (19) | 0.021 (2) | 0.0005 (14) | -0.0020 (15) | -0.0006 (15) |
| C9A | 0.0098 (16) | 0.018 (2) | 0.0181 (19) | -0.0022 (14) | -0.0029 (14) | -0.0026 (15) |
| C10A | 0.0086 (16) | 0.0145 (18) | 0.0136 (18) | -0.0033 (14) | 0.0015 (13) | -0.0023 (14) |
| C11A | 0.0114 (17) | 0.0159 (19) | 0.0175 (19) | -0.0048 (14) | -0.0015 (14) | -0.0031 (15) |
| C12A | 0.0135 (17) | 0.0125 (18) | 0.0133 (18) | -0.0047 (14) | 0.0006 (14) | -0.0053 (14) |
| C13A | 0.0137 (17) | 0.0115 (18) | 0.019 (2) | -0.0021 (14) | -0.0043 (15) | -0.0001 (15) |
| C14A | 0.018 (2) | 0.025 (2) | 0.026 (2) | -0.0062 (17) | -0.0062 (17) | -0.0051 (18) |
| Fe1B | 0.0136 (3) | 0.0156 (3) | 0.0144 (3) | -0.0045 (2) | -0.0029 (2) | -0.0013 (2) |
| S1B | 0.0350 (6) | 0.0162 (5) | 0.0199 (5) | -0.0113 (4) | -0.0027 (4) | -0.0051 (4) |
| N1B | 0.0208 (18) | 0.0182 (18) | 0.0225 (19) | -0.0076 (14) | -0.0043 (15) | -0.0056 (15) |
| N2B | 0.029 (2) | 0.0195 (19) | 0.023 (2) | -0.0120 (16) | -0.0028 (16) | -0.0053 (15) |
| N3B | 0.0226 (19) | 0.031 (2) | 0.029 (2) | -0.0131 (17) | -0.0025 (16) | -0.0129 (18) |
| C1B | 0.019 (2) | 0.024 (2) | 0.016 (2) | -0.0011 (17) | -0.0039 (16) | -0.0049 (17) |
| C2B | 0.0118 (18) | 0.021 (2) | 0.024 (2) | -0.0075 (16) | -0.0022 (15) | -0.0016 (17) |
| C3B | 0.0182 (19) | 0.017 (2) | 0.023 (2) | -0.0021 (16) | -0.0045 (16) | -0.0076 (16) |
| C4B | 0.0179 (19) | 0.021 (2) | 0.017 (2) | -0.0045 (16) | -0.0027 (16) | -0.0009 (16) |
| C5B | 0.026 (2) | 0.016 (2) | 0.021 (2) | -0.0027 (17) | -0.0045 (17) | -0.0033 (16) |
| C6B | 0.021 (2) | 0.023 (2) | 0.0134 (19) | -0.0066 (17) | -0.0042 (16) | 0.0020 (16) |
| C7B | 0.031 (2) | 0.0121 (19) | 0.022 (2) | -0.0031 (17) | -0.0130 (19) | 0.0039 (16) |
| C8B | 0.023 (2) | 0.015 (2) | 0.029 (2) | 0.0032 (17) | -0.0095 (18) | -0.0050 (17) |
| C9B | 0.0165 (19) | 0.024 (2) | 0.021 (2) | -0.0053 (17) | -0.0042 (16) | -0.0028 (17) |
| C10B | 0.0169 (19) | 0.018 (2) | 0.0171 (19) | -0.0042 (15) | -0.0068 (15) | -0.0019 (16) |
| C11B | 0.019 (2) | 0.022 (2) | 0.020 (2) | -0.0095 (17) | -0.0035 (16) | -0.0021 (17) |
| C12B | 0.025 (2) | 0.020 (2) | 0.018 (2) | -0.0074 (17) | -0.0062 (17) | -0.0029 (16) |
| C13B | 0.023 (2) | 0.030 (3) | 0.035 (3) | -0.0052 (19) | 0.003 (2) | -0.014 (2) |
| C14B | 0.041 (3) | 0.046 (3) | 0.033 (3) | -0.018 (3) | 0.014 (2) | -0.024 (3) |

Geometric parameters (Å, °)

| | | | |
|-----------|-----------|-----------|-----------|
| Fe1A—C2A | 2.037 (4) | Fe1B—C3B | 2.036 (4) |
| Fe1A—C3A | 2.046 (4) | Fe1B—C4B | 2.036 (4) |
| Fe1A—C1A | 2.046 (4) | Fe1B—C2B | 2.037 (4) |
| Fe1A—C8A | 2.046 (4) | Fe1B—C10B | 2.048 (4) |
| Fe1A—C9A | 2.051 (4) | Fe1B—C9B | 2.050 (4) |
| Fe1A—C10A | 2.051 (4) | Fe1B—C6B | 2.051 (4) |
| Fe1A—C5A | 2.052 (5) | Fe1B—C7B | 2.053 (4) |
| Fe1A—C7A | 2.052 (4) | Fe1B—C8B | 2.054 (4) |
| Fe1A—C6A | 2.053 (4) | Fe1B—C1B | 2.060 (4) |

supplementary materials

| | | | |
|---------------|-------------|---------------|-------------|
| Fe1A—C4A | 2.063 (4) | Fe1B—C5B | 2.064 (5) |
| S1A—C12A | 1.687 (4) | S1B—C12B | 1.691 (5) |
| N1A—C11A | 1.283 (5) | N1B—C11B | 1.285 (6) |
| N1A—N2A | 1.383 (5) | N1B—N2B | 1.384 (5) |
| N2A—C12A | 1.356 (5) | N2B—C12B | 1.357 (7) |
| N2A—H2NA | 0.82 (6) | N2B—H2NB | 0.89 (9) |
| N3A—C12A | 1.335 (5) | N3B—C12B | 1.326 (6) |
| N3A—C13A | 1.447 (5) | N3B—C13B | 1.461 (7) |
| N3A—H3NA | 0.84 (6) | N3B—H3NB | 0.97 (8) |
| C1A—C5A | 1.410 (6) | C1B—C2B | 1.417 (7) |
| C1A—C2A | 1.421 (6) | C1B—C5B | 1.426 (6) |
| C1A—H1AA | 0.9800 | C1B—H1BA | 0.9800 |
| C2A—C3A | 1.418 (6) | C2B—C3B | 1.431 (6) |
| C2A—H2AA | 0.9800 | C2B—H2BA | 0.9800 |
| C3A—C4A | 1.425 (7) | C3B—C4B | 1.418 (6) |
| C3A—H3AA | 0.9800 | C3B—H3BA | 0.9800 |
| C4A—C5A | 1.433 (7) | C4B—C5B | 1.414 (7) |
| C4A—H4AA | 0.9800 | C4B—H4BA | 0.9800 |
| C5A—H5AA | 0.9800 | C5B—H5BA | 0.9800 |
| C6A—C7A | 1.424 (6) | C6B—C10B | 1.431 (6) |
| C6A—C10A | 1.430 (5) | C6B—C7B | 1.431 (7) |
| C6A—H6AA | 0.9800 | C6B—H6BA | 0.9800 |
| C7A—C8A | 1.429 (6) | C7B—C8B | 1.421 (7) |
| C7A—H7AA | 0.9800 | C7B—H7BA | 0.9800 |
| C8A—C9A | 1.428 (6) | C8B—C9B | 1.433 (6) |
| C8A—H8AA | 0.9800 | C8B—H8BA | 0.9800 |
| C9A—C10A | 1.434 (6) | C9B—C10B | 1.424 (7) |
| C9A—H9AA | 0.9800 | C9B—H9BA | 0.9800 |
| C10A—C11A | 1.456 (6) | C10B—C11B | 1.459 (6) |
| C11A—H11A | 0.9300 | C11B—H11B | 0.9300 |
| C13A—C14A | 1.523 (6) | C13B—C14B | 1.522 (8) |
| C13A—H13A | 0.9700 | C13B—H13C | 0.9700 |
| C13A—H13B | 0.9700 | C13B—H13D | 0.9700 |
| C14A—H14A | 0.9600 | C14B—H14D | 0.9600 |
| C14A—H14B | 0.9600 | C14B—H14E | 0.9600 |
| C14A—H14C | 0.9600 | C14B—H14F | 0.9600 |
| C2A—Fe1A—C3A | 40.63 (18) | C3B—Fe1B—C4B | 40.76 (18) |
| C2A—Fe1A—C1A | 40.72 (17) | C3B—Fe1B—C2B | 41.13 (18) |
| C3A—Fe1A—C1A | 68.45 (18) | C4B—Fe1B—C2B | 68.65 (18) |
| C2A—Fe1A—C8A | 123.50 (18) | C3B—Fe1B—C10B | 162.68 (18) |
| C3A—Fe1A—C8A | 106.06 (18) | C4B—Fe1B—C10B | 125.67 (17) |
| C1A—Fe1A—C8A | 161.08 (18) | C2B—Fe1B—C10B | 154.96 (19) |
| C2A—Fe1A—C9A | 161.10 (17) | C3B—Fe1B—C9B | 125.40 (19) |
| C3A—Fe1A—C9A | 124.71 (18) | C4B—Fe1B—C9B | 108.12 (18) |
| C1A—Fe1A—C9A | 157.02 (18) | C2B—Fe1B—C9B | 162.68 (19) |
| C8A—Fe1A—C9A | 40.79 (17) | C10B—Fe1B—C9B | 40.67 (19) |
| C2A—Fe1A—C10A | 155.75 (17) | C3B—Fe1B—C6B | 155.00 (18) |
| C3A—Fe1A—C10A | 162.98 (17) | C4B—Fe1B—C6B | 162.63 (19) |
| C1A—Fe1A—C10A | 121.71 (17) | C2B—Fe1B—C6B | 119.64 (18) |

| | | | |
|---------------|-------------|---------------|-------------|
| C8A—Fe1A—C10A | 68.66 (16) | C10B—Fe1B—C6B | 40.86 (17) |
| C9A—Fe1A—C10A | 40.93 (16) | C9B—Fe1B—C6B | 68.61 (18) |
| C2A—Fe1A—C5A | 68.06 (18) | C3B—Fe1B—C7B | 120.00 (18) |
| C3A—Fe1A—C5A | 68.28 (19) | C4B—Fe1B—C7B | 155.5 (2) |
| C1A—Fe1A—C5A | 40.26 (18) | C2B—Fe1B—C7B | 106.91 (18) |
| C8A—Fe1A—C5A | 156.34 (18) | C10B—Fe1B—C7B | 68.64 (18) |
| C9A—Fe1A—C5A | 122.29 (18) | C9B—Fe1B—C7B | 68.55 (19) |
| C10A—Fe1A—C5A | 109.67 (17) | C6B—Fe1B—C7B | 40.81 (19) |
| C2A—Fe1A—C7A | 106.16 (17) | C3B—Fe1B—C8B | 107.44 (19) |
| C3A—Fe1A—C7A | 119.01 (18) | C4B—Fe1B—C8B | 121.0 (2) |
| C1A—Fe1A—C7A | 124.63 (17) | C2B—Fe1B—C8B | 125.02 (19) |
| C8A—Fe1A—C7A | 40.82 (16) | C10B—Fe1B—C8B | 68.52 (18) |
| C9A—Fe1A—C7A | 68.53 (17) | C9B—Fe1B—C8B | 40.87 (18) |
| C10A—Fe1A—C7A | 68.29 (16) | C6B—Fe1B—C8B | 68.4 (2) |
| C5A—Fe1A—C7A | 162.30 (18) | C7B—Fe1B—C8B | 40.5 (2) |
| C2A—Fe1A—C6A | 119.68 (17) | C3B—Fe1B—C1B | 68.50 (18) |
| C3A—Fe1A—C6A | 154.10 (18) | C4B—Fe1B—C1B | 68.19 (18) |
| C1A—Fe1A—C6A | 107.78 (17) | C2B—Fe1B—C1B | 40.47 (18) |
| C8A—Fe1A—C6A | 68.80 (17) | C10B—Fe1B—C1B | 120.59 (18) |
| C9A—Fe1A—C6A | 68.88 (16) | C9B—Fe1B—C1B | 155.71 (19) |
| C10A—Fe1A—C6A | 40.79 (15) | C6B—Fe1B—C1B | 107.35 (19) |
| C5A—Fe1A—C6A | 126.39 (18) | C7B—Fe1B—C1B | 125.0 (2) |
| C7A—Fe1A—C6A | 40.58 (16) | C8B—Fe1B—C1B | 161.93 (19) |
| C2A—Fe1A—C4A | 68.27 (18) | C3B—Fe1B—C5B | 68.25 (18) |
| C3A—Fe1A—C4A | 40.57 (18) | C4B—Fe1B—C5B | 40.33 (19) |
| C1A—Fe1A—C4A | 68.32 (19) | C2B—Fe1B—C5B | 68.20 (18) |
| C8A—Fe1A—C4A | 120.07 (19) | C10B—Fe1B—C5B | 108.19 (18) |
| C9A—Fe1A—C4A | 108.26 (18) | C9B—Fe1B—C5B | 121.11 (19) |
| C10A—Fe1A—C4A | 126.87 (18) | C6B—Fe1B—C5B | 125.56 (19) |
| C5A—Fe1A—C4A | 40.75 (19) | C7B—Fe1B—C5B | 162.4 (2) |
| C7A—Fe1A—C4A | 154.51 (18) | C8B—Fe1B—C5B | 156.1 (2) |
| C6A—Fe1A—C4A | 163.95 (18) | C1B—Fe1B—C5B | 40.47 (18) |
| C11A—N1A—N2A | 115.2 (3) | C11B—N1B—N2B | 116.6 (4) |
| C12A—N2A—N1A | 119.4 (3) | C12B—N2B—N1B | 118.6 (4) |
| C12A—N2A—H2NA | 115 (4) | C12B—N2B—H2NB | 116 (5) |
| N1A—N2A—H2NA | 124 (4) | N1B—N2B—H2NB | 121 (6) |
| C12A—N3A—C13A | 124.1 (3) | C12B—N3B—C13B | 126.0 (4) |
| C12A—N3A—H3NA | 116 (4) | C12B—N3B—H3NB | 111 (5) |
| C13A—N3A—H3NA | 117 (4) | C13B—N3B—H3NB | 121 (5) |
| C5A—C1A—C2A | 107.9 (4) | C2B—C1B—C5B | 107.9 (4) |
| C5A—C1A—Fe1A | 70.1 (3) | C2B—C1B—Fe1B | 68.9 (2) |
| C2A—C1A—Fe1A | 69.3 (2) | C5B—C1B—Fe1B | 69.9 (3) |
| C5A—C1A—H1AA | 126.1 | C2B—C1B—H1BA | 126.0 |
| C2A—C1A—H1AA | 126.1 | C5B—C1B—H1BA | 126.0 |
| Fe1A—C1A—H1AA | 126.1 | Fe1B—C1B—H1BA | 126.0 |
| C3A—C2A—C1A | 108.3 (4) | C1B—C2B—C3B | 108.1 (4) |
| C3A—C2A—Fe1A | 70.0 (2) | C1B—C2B—Fe1B | 70.6 (2) |
| C1A—C2A—Fe1A | 70.0 (2) | C3B—C2B—Fe1B | 69.4 (2) |
| C3A—C2A—H2AA | 125.8 | C1B—C2B—H2BA | 126.0 |

supplementary materials

| | | | |
|---------------|-----------|---------------|-----------|
| C1A—C2A—H2AA | 125.8 | C3B—C2B—H2BA | 126.0 |
| Fe1A—C2A—H2AA | 125.8 | Fe1B—C2B—H2BA | 126.0 |
| C2A—C3A—C4A | 108.1 (4) | C4B—C3B—C2B | 107.5 (4) |
| C2A—C3A—Fe1A | 69.4 (2) | C4B—C3B—Fe1B | 69.6 (2) |
| C4A—C3A—Fe1A | 70.4 (2) | C2B—C3B—Fe1B | 69.5 (2) |
| C2A—C3A—H3AA | 125.9 | C4B—C3B—H3BA | 126.3 |
| C4A—C3A—H3AA | 125.9 | C2B—C3B—H3BA | 126.3 |
| Fe1A—C3A—H3AA | 125.9 | Fe1B—C3B—H3BA | 126.3 |
| C3A—C4A—C5A | 107.2 (4) | C5B—C4B—C3B | 108.6 (4) |
| C3A—C4A—Fe1A | 69.1 (3) | C5B—C4B—Fe1B | 70.9 (3) |
| C5A—C4A—Fe1A | 69.2 (3) | C3B—C4B—Fe1B | 69.6 (3) |
| C3A—C4A—H4AA | 126.4 | C5B—C4B—H4BA | 125.7 |
| C5A—C4A—H4AA | 126.4 | C3B—C4B—H4BA | 125.7 |
| Fe1A—C4A—H4AA | 126.4 | Fe1B—C4B—H4BA | 125.7 |
| C1A—C5A—C4A | 108.5 (4) | C4B—C5B—C1B | 107.9 (4) |
| C1A—C5A—Fe1A | 69.6 (3) | C4B—C5B—Fe1B | 68.8 (3) |
| C4A—C5A—Fe1A | 70.0 (3) | C1B—C5B—Fe1B | 69.6 (3) |
| C1A—C5A—H5AA | 125.8 | C4B—C5B—H5BA | 126.1 |
| C4A—C5A—H5AA | 125.8 | C1B—C5B—H5BA | 126.1 |
| Fe1A—C5A—H5AA | 125.8 | Fe1B—C5B—H5BA | 126.1 |
| C7A—C6A—C10A | 107.6 (3) | C10B—C6B—C7B | 107.8 (4) |
| C7A—C6A—Fe1A | 69.7 (2) | C10B—C6B—Fe1B | 69.5 (2) |
| C10A—C6A—Fe1A | 69.5 (2) | C7B—C6B—Fe1B | 69.7 (2) |
| C7A—C6A—H6AA | 126.2 | C10B—C6B—H6BA | 126.1 |
| C10A—C6A—H6AA | 126.2 | C7B—C6B—H6BA | 126.1 |
| Fe1A—C6A—H6AA | 126.2 | Fe1B—C6B—H6BA | 126.1 |
| C6A—C7A—C8A | 108.5 (4) | C8B—C7B—C6B | 108.1 (4) |
| C6A—C7A—Fe1A | 69.7 (2) | C8B—C7B—Fe1B | 69.8 (3) |
| C8A—C7A—Fe1A | 69.4 (3) | C6B—C7B—Fe1B | 69.5 (2) |
| C6A—C7A—H7AA | 125.7 | C8B—C7B—H7BA | 126.0 |
| C8A—C7A—H7AA | 125.7 | C6B—C7B—H7BA | 126.0 |
| Fe1A—C7A—H7AA | 125.7 | Fe1B—C7B—H7BA | 126.0 |
| C9A—C8A—C7A | 107.9 (4) | C7B—C8B—C9B | 108.1 (4) |
| C9A—C8A—Fe1A | 69.8 (2) | C7B—C8B—Fe1B | 69.7 (3) |
| C7A—C8A—Fe1A | 69.8 (2) | C9B—C8B—Fe1B | 69.4 (3) |
| C9A—C8A—H8AA | 126.0 | C7B—C8B—H8BA | 125.9 |
| C7A—C8A—H8AA | 126.0 | C9B—C8B—H8BA | 125.9 |
| Fe1A—C8A—H8AA | 126.0 | Fe1B—C8B—H8BA | 125.9 |
| C8A—C9A—C10A | 107.7 (3) | C10B—C9B—C8B | 107.9 (4) |
| C8A—C9A—Fe1A | 69.4 (2) | C10B—C9B—Fe1B | 69.6 (2) |
| C10A—C9A—Fe1A | 69.6 (2) | C8B—C9B—Fe1B | 69.7 (2) |
| C8A—C9A—H9AA | 126.1 | C10B—C9B—H9BA | 126.1 |
| C10A—C9A—H9AA | 126.1 | C8B—C9B—H9BA | 126.1 |
| Fe1A—C9A—H9AA | 126.1 | Fe1B—C9B—H9BA | 126.1 |
| C6A—C10A—C9A | 108.2 (4) | C9B—C10B—C6B | 108.1 (4) |
| C6A—C10A—C11A | 125.5 (4) | C9B—C10B—C11B | 126.0 (4) |
| C9A—C10A—C11A | 126.2 (4) | C6B—C10B—C11B | 125.9 (4) |
| C6A—C10A—Fe1A | 69.7 (2) | C9B—C10B—Fe1B | 69.8 (2) |
| C9A—C10A—Fe1A | 69.5 (2) | C6B—C10B—Fe1B | 69.7 (2) |

| | | | |
|-------------------|------------|-------------------|------------|
| C11A—C10A—Fe1A | 129.0 (3) | C11B—C10B—Fe1B | 127.2 (3) |
| N1A—C11A—C10A | 120.0 (4) | N1B—C11B—C10B | 119.4 (4) |
| N1A—C11A—H11A | 120.0 | N1B—C11B—H11B | 120.3 |
| C10A—C11A—H11A | 120.0 | C10B—C11B—H11B | 120.3 |
| N3A—C12A—N2A | 116.8 (4) | N3B—C12B—N2B | 116.3 (4) |
| N3A—C12A—S1A | 123.0 (3) | N3B—C12B—S1B | 123.8 (4) |
| N2A—C12A—S1A | 120.1 (3) | N2B—C12B—S1B | 119.9 (4) |
| N3A—C13A—C14A | 110.0 (3) | N3B—C13B—C14B | 112.6 (4) |
| N3A—C13A—H13A | 109.7 | N3B—C13B—H13C | 109.1 |
| C14A—C13A—H13A | 109.7 | C14B—C13B—H13C | 109.1 |
| N3A—C13A—H13B | 109.7 | N3B—C13B—H13D | 109.1 |
| C14A—C13A—H13B | 109.7 | C14B—C13B—H13D | 109.1 |
| H13A—C13A—H13B | 108.2 | H13C—C13B—H13D | 107.8 |
| C13A—C14A—H14A | 109.5 | C13B—C14B—H14D | 109.5 |
| C13A—C14A—H14B | 109.5 | C13B—C14B—H14E | 109.5 |
| H14A—C14A—H14B | 109.5 | H14D—C14B—H14E | 109.5 |
| C13A—C14A—H14C | 109.5 | C13B—C14B—H14F | 109.5 |
| H14A—C14A—H14C | 109.5 | H14D—C14B—H14F | 109.5 |
| H14B—C14A—H14C | 109.5 | H14E—C14B—H14F | 109.5 |
| C11A—N1A—N2A—C12A | -174.2 (4) | C11B—N1B—N2B—C12B | -177.4 (4) |
| C2A—Fe1A—C1A—C5A | 119.1 (4) | C3B—Fe1B—C1B—C2B | -38.2 (3) |
| C3A—Fe1A—C1A—C5A | 81.4 (3) | C4B—Fe1B—C1B—C2B | -82.2 (3) |
| C8A—Fe1A—C1A—C5A | 157.8 (5) | C10B—Fe1B—C1B—C2B | 158.2 (3) |
| C9A—Fe1A—C1A—C5A | -48.1 (6) | C9B—Fe1B—C1B—C2B | -168.1 (4) |
| C10A—Fe1A—C1A—C5A | -83.3 (3) | C6B—Fe1B—C1B—C2B | 115.6 (3) |
| C7A—Fe1A—C1A—C5A | -167.4 (3) | C7B—Fe1B—C1B—C2B | 74.1 (3) |
| C6A—Fe1A—C1A—C5A | -125.8 (3) | C8B—Fe1B—C1B—C2B | 42.1 (7) |
| C4A—Fe1A—C1A—C5A | 37.6 (3) | C5B—Fe1B—C1B—C2B | -119.5 (4) |
| C3A—Fe1A—C1A—C2A | -37.6 (3) | C3B—Fe1B—C1B—C5B | 81.3 (3) |
| C8A—Fe1A—C1A—C2A | 38.8 (6) | C4B—Fe1B—C1B—C5B | 37.3 (3) |
| C9A—Fe1A—C1A—C2A | -167.2 (4) | C2B—Fe1B—C1B—C5B | 119.5 (4) |
| C10A—Fe1A—C1A—C2A | 157.7 (3) | C10B—Fe1B—C1B—C5B | -82.3 (3) |
| C5A—Fe1A—C1A—C2A | -119.1 (4) | C9B—Fe1B—C1B—C5B | -48.6 (5) |
| C7A—Fe1A—C1A—C2A | 73.5 (3) | C6B—Fe1B—C1B—C5B | -124.9 (3) |
| C6A—Fe1A—C1A—C2A | 115.1 (3) | C7B—Fe1B—C1B—C5B | -166.4 (3) |
| C4A—Fe1A—C1A—C2A | -81.4 (3) | C8B—Fe1B—C1B—C5B | 161.6 (6) |
| C5A—C1A—C2A—C3A | 0.0 (5) | C5B—C1B—C2B—C3B | 0.3 (5) |
| Fe1A—C1A—C2A—C3A | 59.7 (3) | Fe1B—C1B—C2B—C3B | 59.5 (3) |
| C5A—C1A—C2A—Fe1A | -59.7 (3) | C5B—C1B—C2B—Fe1B | -59.3 (3) |
| C1A—Fe1A—C2A—C3A | -119.3 (4) | C3B—Fe1B—C2B—C1B | 118.9 (4) |
| C8A—Fe1A—C2A—C3A | 74.8 (3) | C4B—Fe1B—C2B—C1B | 81.0 (3) |
| C9A—Fe1A—C2A—C3A | 45.2 (7) | C10B—Fe1B—C2B—C1B | -49.0 (5) |
| C10A—Fe1A—C2A—C3A | -171.2 (4) | C9B—Fe1B—C2B—C1B | 163.4 (6) |
| C5A—Fe1A—C2A—C3A | -81.8 (3) | C6B—Fe1B—C2B—C1B | -82.1 (3) |
| C7A—Fe1A—C2A—C3A | 116.0 (3) | C7B—Fe1B—C2B—C1B | -124.6 (3) |
| C6A—Fe1A—C2A—C3A | 157.8 (3) | C8B—Fe1B—C2B—C1B | -165.3 (3) |
| C4A—Fe1A—C2A—C3A | -37.7 (3) | C5B—Fe1B—C2B—C1B | 37.5 (3) |
| C3A—Fe1A—C2A—C1A | 119.3 (4) | C4B—Fe1B—C2B—C3B | -37.9 (3) |
| C8A—Fe1A—C2A—C1A | -165.9 (3) | C10B—Fe1B—C2B—C3B | -167.9 (4) |

supplementary materials

| | | | |
|-------------------|------------|-------------------|------------|
| C9A—Fe1A—C2A—C1A | 164.5 (5) | C9B—Fe1B—C2B—C3B | 44.5 (7) |
| C10A—Fe1A—C2A—C1A | -51.9 (5) | C6B—Fe1B—C2B—C3B | 159.0 (3) |
| C5A—Fe1A—C2A—C1A | 37.5 (3) | C7B—Fe1B—C2B—C3B | 116.5 (3) |
| C7A—Fe1A—C2A—C1A | -124.8 (3) | C8B—Fe1B—C2B—C3B | 75.8 (3) |
| C6A—Fe1A—C2A—C1A | -82.9 (3) | C1B—Fe1B—C2B—C3B | -118.9 (4) |
| C4A—Fe1A—C2A—C1A | 81.6 (3) | C5B—Fe1B—C2B—C3B | -81.4 (3) |
| C1A—C2A—C3A—C4A | 0.3 (5) | C1B—C2B—C3B—C4B | -0.8 (5) |
| Fe1A—C2A—C3A—C4A | 60.0 (3) | Fe1B—C2B—C3B—C4B | 59.5 (3) |
| C1A—C2A—C3A—Fe1A | -59.7 (3) | C1B—C2B—C3B—Fe1B | -60.3 (3) |
| C1A—Fe1A—C3A—C2A | 37.7 (3) | C2B—Fe1B—C3B—C4B | -118.7 (4) |
| C8A—Fe1A—C3A—C2A | -123.1 (3) | C10B—Fe1B—C3B—C4B | 43.9 (7) |
| C9A—Fe1A—C3A—C2A | -163.8 (3) | C9B—Fe1B—C3B—C4B | 76.1 (3) |
| C10A—Fe1A—C3A—C2A | 167.6 (5) | C6B—Fe1B—C3B—C4B | -166.2 (4) |
| C5A—Fe1A—C3A—C2A | 81.2 (3) | C7B—Fe1B—C3B—C4B | 159.9 (3) |
| C7A—Fe1A—C3A—C2A | -80.9 (3) | C8B—Fe1B—C3B—C4B | 117.6 (3) |
| C6A—Fe1A—C3A—C2A | -48.7 (5) | C1B—Fe1B—C3B—C4B | -81.1 (3) |
| C4A—Fe1A—C3A—C2A | 119.1 (4) | C5B—Fe1B—C3B—C4B | -37.4 (3) |
| C2A—Fe1A—C3A—C4A | -119.1 (4) | C4B—Fe1B—C3B—C2B | 118.7 (4) |
| C1A—Fe1A—C3A—C4A | -81.4 (3) | C10B—Fe1B—C3B—C2B | 162.7 (5) |
| C8A—Fe1A—C3A—C4A | 117.7 (3) | C9B—Fe1B—C3B—C2B | -165.2 (3) |
| C9A—Fe1A—C3A—C4A | 77.1 (3) | C6B—Fe1B—C3B—C2B | -47.4 (5) |
| C10A—Fe1A—C3A—C4A | 48.4 (7) | C7B—Fe1B—C3B—C2B | -81.4 (3) |
| C5A—Fe1A—C3A—C4A | -37.9 (3) | C8B—Fe1B—C3B—C2B | -123.7 (3) |
| C7A—Fe1A—C3A—C4A | 159.9 (3) | C1B—Fe1B—C3B—C2B | 37.6 (3) |
| C6A—Fe1A—C3A—C4A | -167.9 (4) | C5B—Fe1B—C3B—C2B | 81.3 (3) |
| C2A—C3A—C4A—C5A | -0.4 (5) | C2B—C3B—C4B—C5B | 1.0 (5) |
| Fe1A—C3A—C4A—C5A | 58.9 (3) | Fe1B—C3B—C4B—C5B | 60.4 (3) |
| C2A—C3A—C4A—Fe1A | -59.3 (3) | C2B—C3B—C4B—Fe1B | -59.4 (3) |
| C2A—Fe1A—C4A—C3A | 37.8 (3) | C3B—Fe1B—C4B—C5B | -119.3 (4) |
| C1A—Fe1A—C4A—C3A | 81.7 (3) | C2B—Fe1B—C4B—C5B | -81.0 (3) |
| C8A—Fe1A—C4A—C3A | -79.4 (3) | C10B—Fe1B—C4B—C5B | 75.4 (3) |
| C9A—Fe1A—C4A—C3A | -122.4 (3) | C9B—Fe1B—C4B—C5B | 117.0 (3) |
| C10A—Fe1A—C4A—C3A | -164.1 (3) | C6B—Fe1B—C4B—C5B | 40.9 (7) |
| C5A—Fe1A—C4A—C3A | 118.9 (4) | C7B—Fe1B—C4B—C5B | -165.1 (4) |
| C7A—Fe1A—C4A—C3A | -44.2 (5) | C8B—Fe1B—C4B—C5B | 160.0 (3) |
| C6A—Fe1A—C4A—C3A | 160.6 (5) | C1B—Fe1B—C4B—C5B | -37.4 (3) |
| C2A—Fe1A—C4A—C5A | -81.2 (3) | C2B—Fe1B—C4B—C3B | 38.3 (3) |
| C3A—Fe1A—C4A—C5A | -118.9 (4) | C10B—Fe1B—C4B—C3B | -165.3 (3) |
| C1A—Fe1A—C4A—C5A | -37.2 (3) | C9B—Fe1B—C4B—C3B | -123.6 (3) |
| C8A—Fe1A—C4A—C5A | 161.7 (3) | C6B—Fe1B—C4B—C3B | 160.2 (6) |
| C9A—Fe1A—C4A—C5A | 118.6 (3) | C7B—Fe1B—C4B—C3B | -45.7 (6) |
| C10A—Fe1A—C4A—C5A | 77.0 (3) | C8B—Fe1B—C4B—C3B | -80.7 (3) |
| C7A—Fe1A—C4A—C5A | -163.1 (4) | C1B—Fe1B—C4B—C3B | 81.9 (3) |
| C6A—Fe1A—C4A—C5A | 41.7 (7) | C5B—Fe1B—C4B—C3B | 119.3 (4) |
| C2A—C1A—C5A—C4A | -0.2 (5) | C3B—C4B—C5B—C1B | -0.8 (5) |
| Fe1A—C1A—C5A—C4A | -59.5 (3) | Fe1B—C4B—C5B—C1B | 58.8 (3) |
| C2A—C1A—C5A—Fe1A | 59.2 (3) | C3B—C4B—C5B—Fe1B | -59.6 (3) |
| C3A—C4A—C5A—C1A | 0.4 (5) | C2B—C1B—C5B—C4B | 0.3 (5) |
| Fe1A—C4A—C5A—C1A | 59.2 (3) | Fe1B—C1B—C5B—C4B | -58.3 (3) |

| | | | |
|-------------------|------------|-------------------|------------|
| C3A—C4A—C5A—Fe1A | -58.8 (3) | C2B—C1B—C5B—Fe1B | 58.6 (3) |
| C2A—Fe1A—C5A—C1A | -37.9 (3) | C3B—Fe1B—C5B—C4B | 37.8 (3) |
| C3A—Fe1A—C5A—C1A | -81.9 (3) | C2B—Fe1B—C5B—C4B | 82.3 (3) |
| C8A—Fe1A—C5A—C1A | -162.2 (4) | C10B—Fe1B—C5B—C4B | -124.1 (3) |
| C9A—Fe1A—C5A—C1A | 159.9 (3) | C9B—Fe1B—C5B—C4B | -81.4 (3) |
| C10A—Fe1A—C5A—C1A | 116.2 (3) | C6B—Fe1B—C5B—C4B | -166.1 (3) |
| C7A—Fe1A—C5A—C1A | 36.1 (7) | C7B—Fe1B—C5B—C4B | 159.3 (5) |
| C6A—Fe1A—C5A—C1A | 73.5 (3) | C8B—Fe1B—C5B—C4B | -46.3 (6) |
| C4A—Fe1A—C5A—C1A | -119.7 (4) | C1B—Fe1B—C5B—C4B | 119.7 (4) |
| C2A—Fe1A—C5A—C4A | 81.7 (3) | C3B—Fe1B—C5B—C1B | -81.9 (3) |
| C3A—Fe1A—C5A—C4A | 37.8 (3) | C4B—Fe1B—C5B—C1B | -119.7 (4) |
| C1A—Fe1A—C5A—C4A | 119.7 (4) | C2B—Fe1B—C5B—C1B | -37.5 (3) |
| C8A—Fe1A—C5A—C4A | -42.6 (5) | C10B—Fe1B—C5B—C1B | 116.1 (3) |
| C9A—Fe1A—C5A—C4A | -80.5 (3) | C9B—Fe1B—C5B—C1B | 158.9 (3) |
| C10A—Fe1A—C5A—C4A | -124.1 (3) | C6B—Fe1B—C5B—C1B | 74.2 (3) |
| C7A—Fe1A—C5A—C4A | 155.7 (5) | C7B—Fe1B—C5B—C1B | 39.5 (7) |
| C6A—Fe1A—C5A—C4A | -166.8 (2) | C8B—Fe1B—C5B—C1B | -166.0 (4) |
| C2A—Fe1A—C6A—C7A | -80.0 (3) | C3B—Fe1B—C6B—C10B | -166.8 (4) |
| C3A—Fe1A—C6A—C7A | -45.8 (5) | C4B—Fe1B—C6B—C10B | 44.7 (7) |
| C1A—Fe1A—C6A—C7A | -122.9 (2) | C2B—Fe1B—C6B—C10B | 159.3 (3) |
| C8A—Fe1A—C6A—C7A | 37.4 (2) | C9B—Fe1B—C6B—C10B | -37.6 (3) |
| C9A—Fe1A—C6A—C7A | 81.3 (3) | C7B—Fe1B—C6B—C10B | -119.1 (4) |
| C10A—Fe1A—C6A—C7A | 118.9 (3) | C8B—Fe1B—C6B—C10B | -81.7 (3) |
| C5A—Fe1A—C6A—C7A | -163.5 (2) | C1B—Fe1B—C6B—C10B | 117.0 (3) |
| C4A—Fe1A—C6A—C7A | 163.9 (6) | C5B—Fe1B—C6B—C10B | 76.1 (3) |
| C2A—Fe1A—C6A—C10A | 161.1 (2) | C3B—Fe1B—C6B—C7B | -47.7 (6) |
| C3A—Fe1A—C6A—C10A | -164.6 (4) | C4B—Fe1B—C6B—C7B | 163.9 (6) |
| C1A—Fe1A—C6A—C10A | 118.3 (2) | C2B—Fe1B—C6B—C7B | -81.6 (3) |
| C8A—Fe1A—C6A—C10A | -81.5 (3) | C10B—Fe1B—C6B—C7B | 119.1 (4) |
| C9A—Fe1A—C6A—C10A | -37.6 (2) | C9B—Fe1B—C6B—C7B | 81.5 (3) |
| C5A—Fe1A—C6A—C10A | 77.6 (3) | C8B—Fe1B—C6B—C7B | 37.5 (3) |
| C7A—Fe1A—C6A—C10A | -118.9 (3) | C1B—Fe1B—C6B—C7B | -123.9 (3) |
| C4A—Fe1A—C6A—C10A | 45.0 (7) | C5B—Fe1B—C6B—C7B | -164.8 (3) |
| C10A—C6A—C7A—C8A | 0.7 (5) | C10B—C6B—C7B—C8B | -0.1 (5) |
| Fe1A—C6A—C7A—C8A | -58.7 (3) | Fe1B—C6B—C7B—C8B | -59.4 (3) |
| C10A—C6A—C7A—Fe1A | 59.4 (3) | C10B—C6B—C7B—Fe1B | 59.2 (3) |
| C2A—Fe1A—C7A—C6A | 117.0 (2) | C3B—Fe1B—C7B—C8B | -81.8 (3) |
| C3A—Fe1A—C7A—C6A | 159.0 (2) | C4B—Fe1B—C7B—C8B | -49.1 (5) |
| C1A—Fe1A—C7A—C6A | 76.4 (3) | C2B—Fe1B—C7B—C8B | -124.6 (3) |
| C8A—Fe1A—C7A—C6A | -120.1 (3) | C10B—Fe1B—C7B—C8B | 81.5 (3) |
| C9A—Fe1A—C7A—C6A | -82.2 (2) | C9B—Fe1B—C7B—C8B | 37.7 (3) |
| C10A—Fe1A—C7A—C6A | -38.0 (2) | C6B—Fe1B—C7B—C8B | 119.4 (4) |
| C5A—Fe1A—C7A—C6A | 48.8 (6) | C1B—Fe1B—C7B—C8B | -165.3 (3) |
| C4A—Fe1A—C7A—C6A | -169.7 (4) | C5B—Fe1B—C7B—C8B | 164.4 (6) |
| C2A—Fe1A—C7A—C8A | -122.9 (3) | C3B—Fe1B—C7B—C6B | 158.8 (3) |
| C3A—Fe1A—C7A—C8A | -80.9 (3) | C4B—Fe1B—C7B—C6B | -168.5 (4) |
| C1A—Fe1A—C7A—C8A | -163.6 (2) | C2B—Fe1B—C7B—C6B | 116.0 (3) |
| C9A—Fe1A—C7A—C8A | 37.9 (2) | C10B—Fe1B—C7B—C6B | -37.9 (3) |
| C10A—Fe1A—C7A—C8A | 82.1 (3) | C9B—Fe1B—C7B—C6B | -81.7 (3) |

supplementary materials

| | | | |
|--------------------|------------|--------------------|------------|
| C5A—Fe1A—C7A—C8A | 168.9 (5) | C8B—Fe1B—C7B—C6B | -119.4 (4) |
| C6A—Fe1A—C7A—C8A | 120.1 (3) | C1B—Fe1B—C7B—C6B | 75.3 (3) |
| C4A—Fe1A—C7A—C8A | -49.7 (5) | C5B—Fe1B—C7B—C6B | 45.0 (7) |
| C6A—C7A—C8A—C9A | -0.7 (5) | C6B—C7B—C8B—C9B | 0.2 (5) |
| Fe1A—C7A—C8A—C9A | -59.6 (3) | Fe1B—C7B—C8B—C9B | -59.0 (3) |
| C6A—C7A—C8A—Fe1A | 58.9 (3) | C6B—C7B—C8B—Fe1B | 59.2 (3) |
| C2A—Fe1A—C8A—C9A | -165.8 (2) | C3B—Fe1B—C8B—C7B | 116.0 (3) |
| C3A—Fe1A—C8A—C9A | -125.0 (3) | C4B—Fe1B—C8B—C7B | 158.5 (3) |
| C1A—Fe1A—C8A—C9A | 164.8 (5) | C2B—Fe1B—C8B—C7B | 74.1 (3) |
| C10A—Fe1A—C8A—C9A | 37.9 (2) | C10B—Fe1B—C8B—C7B | -81.8 (3) |
| C5A—Fe1A—C8A—C9A | -52.6 (5) | C9B—Fe1B—C8B—C7B | -119.6 (4) |
| C7A—Fe1A—C8A—C9A | 119.0 (4) | C6B—Fe1B—C8B—C7B | -37.8 (3) |
| C6A—Fe1A—C8A—C9A | 81.8 (3) | C1B—Fe1B—C8B—C7B | 42.0 (7) |
| C4A—Fe1A—C8A—C9A | -83.3 (3) | C5B—Fe1B—C8B—C7B | -168.4 (4) |
| C2A—Fe1A—C8A—C7A | 75.2 (3) | C3B—Fe1B—C8B—C9B | -124.4 (3) |
| C3A—Fe1A—C8A—C7A | 116.0 (3) | C4B—Fe1B—C8B—C9B | -81.9 (3) |
| C1A—Fe1A—C8A—C7A | 45.9 (6) | C2B—Fe1B—C8B—C9B | -166.3 (3) |
| C9A—Fe1A—C8A—C7A | -119.0 (4) | C10B—Fe1B—C8B—C9B | 37.7 (3) |
| C10A—Fe1A—C8A—C7A | -81.1 (3) | C6B—Fe1B—C8B—C9B | 81.8 (3) |
| C5A—Fe1A—C8A—C7A | -171.6 (4) | C7B—Fe1B—C8B—C9B | 119.6 (4) |
| C6A—Fe1A—C8A—C7A | -37.2 (2) | C1B—Fe1B—C8B—C9B | 161.6 (5) |
| C4A—Fe1A—C8A—C7A | 157.7 (2) | C5B—Fe1B—C8B—C9B | -48.9 (6) |
| C7A—C8A—C9A—C10A | 0.4 (5) | C7B—C8B—C9B—C10B | -0.2 (5) |
| Fe1A—C8A—C9A—C10A | -59.2 (3) | Fe1B—C8B—C9B—C10B | -59.3 (3) |
| C7A—C8A—C9A—Fe1A | 59.6 (3) | C7B—C8B—C9B—Fe1B | 59.2 (3) |
| C2A—Fe1A—C9A—C8A | 39.1 (6) | C3B—Fe1B—C9B—C10B | -165.9 (3) |
| C3A—Fe1A—C9A—C8A | 73.3 (3) | C4B—Fe1B—C9B—C10B | -124.1 (3) |
| C1A—Fe1A—C9A—C8A | -167.5 (4) | C2B—Fe1B—C9B—C10B | 159.6 (6) |
| C10A—Fe1A—C9A—C8A | -119.1 (4) | C6B—Fe1B—C9B—C10B | 37.8 (3) |
| C5A—Fe1A—C9A—C8A | 157.8 (3) | C7B—Fe1B—C9B—C10B | 81.8 (3) |
| C7A—Fe1A—C9A—C8A | -37.9 (2) | C8B—Fe1B—C9B—C10B | 119.1 (4) |
| C6A—Fe1A—C9A—C8A | -81.6 (3) | C1B—Fe1B—C9B—C10B | -47.1 (5) |
| C4A—Fe1A—C9A—C8A | 115.2 (3) | C5B—Fe1B—C9B—C10B | -81.8 (3) |
| C2A—Fe1A—C9A—C10A | 158.2 (5) | C3B—Fe1B—C9B—C8B | 75.0 (3) |
| C3A—Fe1A—C9A—C10A | -167.6 (3) | C4B—Fe1B—C9B—C8B | 116.8 (3) |
| C1A—Fe1A—C9A—C10A | -48.4 (5) | C2B—Fe1B—C9B—C8B | 40.5 (7) |
| C8A—Fe1A—C9A—C10A | 119.1 (4) | C10B—Fe1B—C9B—C8B | -119.1 (4) |
| C5A—Fe1A—C9A—C10A | -83.1 (3) | C6B—Fe1B—C9B—C8B | -81.3 (3) |
| C7A—Fe1A—C9A—C10A | 81.2 (3) | C7B—Fe1B—C9B—C8B | -37.4 (3) |
| C6A—Fe1A—C9A—C10A | 37.5 (2) | C1B—Fe1B—C9B—C8B | -166.2 (4) |
| C4A—Fe1A—C9A—C10A | -125.7 (3) | C5B—Fe1B—C9B—C8B | 159.1 (3) |
| C7A—C6A—C10A—C9A | -0.5 (5) | C8B—C9B—C10B—C6B | 0.1 (5) |
| Fe1A—C6A—C10A—C9A | 59.0 (3) | Fe1B—C9B—C10B—C6B | -59.3 (3) |
| C7A—C6A—C10A—C11A | 176.4 (4) | C8B—C9B—C10B—C11B | -178.7 (4) |
| Fe1A—C6A—C10A—C11A | -124.1 (4) | Fe1B—C9B—C10B—C11B | 121.9 (4) |
| C7A—C6A—C10A—Fe1A | -59.5 (3) | C8B—C9B—C10B—Fe1B | 59.4 (3) |
| C8A—C9A—C10A—C6A | 0.1 (5) | C7B—C6B—C10B—C9B | 0.0 (5) |
| Fe1A—C9A—C10A—C6A | -59.1 (3) | Fe1B—C6B—C10B—C9B | 59.4 (3) |
| C8A—C9A—C10A—C11A | -176.8 (4) | C7B—C6B—C10B—C11B | 178.8 (4) |

| | | | |
|--------------------|------------|--------------------|------------|
| Fe1A—C9A—C10A—C11A | 124.0 (4) | Fe1B—C6B—C10B—C11B | -121.8 (4) |
| C8A—C9A—C10A—Fe1A | 59.2 (3) | C7B—C6B—C10B—Fe1B | -59.3 (3) |
| C2A—Fe1A—C10A—C6A | -43.3 (5) | C3B—Fe1B—C10B—C9B | 41.8 (7) |
| C3A—Fe1A—C10A—C6A | 156.7 (6) | C4B—Fe1B—C10B—C9B | 75.6 (3) |
| C1A—Fe1A—C10A—C6A | -80.4 (3) | C2B—Fe1B—C10B—C9B | -165.8 (4) |
| C8A—Fe1A—C10A—C6A | 81.9 (3) | C6B—Fe1B—C10B—C9B | -119.3 (4) |
| C9A—Fe1A—C10A—C6A | 119.6 (3) | C7B—Fe1B—C10B—C9B | -81.5 (3) |
| C5A—Fe1A—C10A—C6A | -123.4 (3) | C8B—Fe1B—C10B—C9B | -37.9 (3) |
| C7A—Fe1A—C10A—C6A | 37.8 (2) | C1B—Fe1B—C10B—C9B | 159.5 (3) |
| C4A—Fe1A—C10A—C6A | -165.8 (3) | C5B—Fe1B—C10B—C9B | 116.9 (3) |
| C2A—Fe1A—C10A—C9A | -162.9 (4) | C3B—Fe1B—C10B—C6B | 161.1 (6) |
| C3A—Fe1A—C10A—C9A | 37.0 (7) | C4B—Fe1B—C10B—C6B | -165.0 (3) |
| C1A—Fe1A—C10A—C9A | 159.9 (3) | C2B—Fe1B—C10B—C6B | -46.5 (5) |
| C8A—Fe1A—C10A—C9A | -37.8 (3) | C9B—Fe1B—C10B—C6B | 119.3 (4) |
| C5A—Fe1A—C10A—C9A | 117.0 (3) | C7B—Fe1B—C10B—C6B | 37.8 (3) |
| C7A—Fe1A—C10A—C9A | -81.8 (3) | C8B—Fe1B—C10B—C6B | 81.4 (3) |
| C6A—Fe1A—C10A—C9A | -119.6 (3) | C1B—Fe1B—C10B—C6B | -81.2 (3) |
| C4A—Fe1A—C10A—C9A | 74.5 (3) | C5B—Fe1B—C10B—C6B | -123.8 (3) |
| C2A—Fe1A—C10A—C11A | 76.5 (6) | C3B—Fe1B—C10B—C11B | -78.6 (8) |
| C3A—Fe1A—C10A—C11A | -83.5 (7) | C4B—Fe1B—C10B—C11B | -44.8 (5) |
| C1A—Fe1A—C10A—C11A | 39.4 (4) | C2B—Fe1B—C10B—C11B | 73.8 (6) |
| C8A—Fe1A—C10A—C11A | -158.3 (4) | C9B—Fe1B—C10B—C11B | -120.4 (5) |
| C9A—Fe1A—C10A—C11A | -120.5 (5) | C6B—Fe1B—C10B—C11B | 120.3 (5) |
| C5A—Fe1A—C10A—C11A | -3.5 (4) | C7B—Fe1B—C10B—C11B | 158.1 (5) |
| C7A—Fe1A—C10A—C11A | 157.6 (4) | C8B—Fe1B—C10B—C11B | -158.3 (5) |
| C6A—Fe1A—C10A—C11A | 119.8 (5) | C1B—Fe1B—C10B—C11B | 39.1 (5) |
| C4A—Fe1A—C10A—C11A | -46.0 (4) | C5B—Fe1B—C10B—C11B | -3.5 (5) |
| N2A—N1A—C11A—C10A | -174.7 (4) | N2B—N1B—C11B—C10B | -176.5 (4) |
| C6A—C10A—C11A—N1A | 0.7 (6) | C9B—C10B—C11B—N1B | 178.3 (4) |
| C9A—C10A—C11A—N1A | 177.0 (4) | C6B—C10B—C11B—N1B | -0.3 (7) |
| Fe1A—C10A—C11A—N1A | -91.0 (5) | Fe1B—C10B—C11B—N1B | -91.0 (5) |
| C13A—N3A—C12A—N2A | 178.3 (4) | C13B—N3B—C12B—N2B | -177.3 (5) |
| C13A—N3A—C12A—S1A | -1.4 (6) | C13B—N3B—C12B—S1B | 4.3 (7) |
| N1A—N2A—C12A—N3A | 6.2 (6) | N1B—N2B—C12B—N3B | 7.1 (6) |
| N1A—N2A—C12A—S1A | -174.1 (3) | N1B—N2B—C12B—S1B | -174.4 (3) |
| C12A—N3A—C13A—C14A | -175.2 (4) | C12B—N3B—C13B—C14B | 84.8 (6) |

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

| $D-H\cdots A$ | $D-H$ | $H\cdots A$ | $D\cdots A$ | $D-H\cdots A$ |
|--------------------------------------|----------|-------------|-------------|---------------|
| N2A—H2NA \cdots S1A ⁱ | 0.82 (6) | 2.59 (6) | 3.387 (4) | 164 (5) |
| N2B—H2NB \cdots S1B ⁱⁱ | 0.89 (9) | 2.55 (9) | 3.430 (5) | 170 (5) |
| C4A—H4AA \cdots S1B ⁱⁱⁱ | 0.98 | 2.79 | 3.715 (4) | 157 |

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

Fig. 1

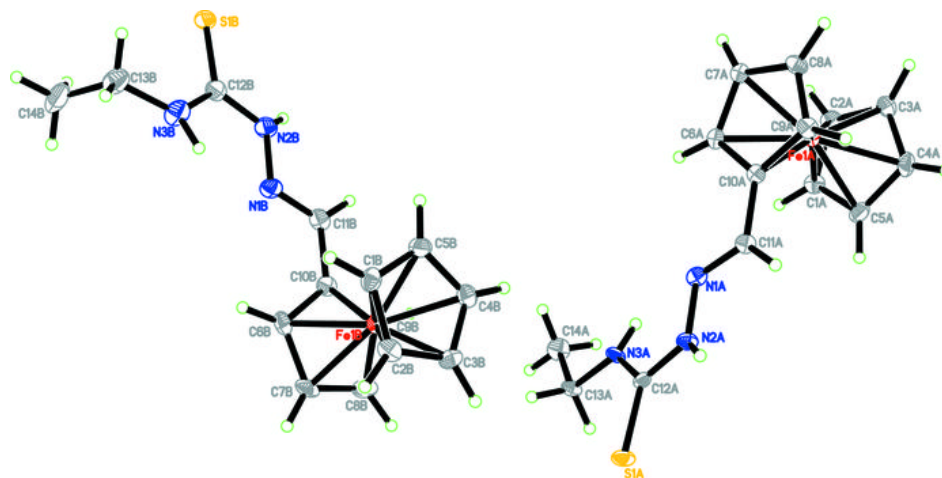


Fig. 2

