### **Supporting Information**

# Thermal and Pressure – Induced Spin Crossover in a Novel Three-Dimensional Hoffman-like Chathrate Complex

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The X-ray diffraction pattern of the powder sample is shown in (ESI Figure 1), along with that calculated from the single crystal pattern. One can conclude that the crystal structure obtained from single crystal methods is representative of the powder sample that was used for all other characterisation techniques.



**ESI Figure 1.** Powder diffraction patterns of  $Fe(bpac)_2[Ag(CN)_2]_2$ . red - calculated from single crystal data, and black - obtained from the bulk sample

Room temperature Raman spectra as a function of pressure are shown in ESI Figure 2. The luminescence spectra of ruby used for pressure determination are given in ESI Figure 3.



ESI Figure 2. Room temperature Raman spectra of Fe(bpac)<sub>2</sub>[Ag(CN)<sub>2</sub>]<sub>2</sub> as a function of pressure



ESI Figure 3. Ruby fluorescence on increasing pressure.

The thermogravimetric analysis is shown in ESI Figure 4.



Figure 4. Thermogravimetric analysis of Fe(bpac)<sub>2</sub>[Ag(CN)<sub>2</sub>]<sub>2</sub>

#### CIF for Fe(bpac)<sub>2</sub>[Ag(CN)<sub>2</sub>]<sub>2</sub>

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 (compiled 2010.09.30 svn.r1450, GUI svn.r3284)
_publ_requested_journal
                            '?'
                               ?
chemical name common
_chemical_name_systematic
                              191
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chemical formula sum
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chemical formula weight
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chemical melting point
chemical_oxdiff_usercomment
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loop
 atom type symbol
_atom_type_description
 _atom_type_scat_dispersion_real
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space group crystal system 'monoclinic' space group IT number 14 'P 1 21/n 1' space group name H-M alt space group name Hall '-P 2yn' loop space group symop id \_space\_group\_symop\_operation xyz 1 'x, y, z' 2 '-x+1/2, y+1/2, -z+1/2' 3 '-x, -y, -z' 4 'x-1/2, -y-1/2, z-1/2' cell length a 9.6204(4) \_cell\_length b 10.8948(5)cell length c 13.3105(5)\_cell\_angle\_alpha 90.00 cell angle beta 95.411(4) \_cell\_angle\_gamma 90.00 cell volume 1388.89(10) cell formula units Z 4 cell measurement reflns used 2158 cell measurement temperature 220 cell measurement theta max 29.0122 cell measurement theta min 3.5632 exptl absorpt coefficient mu 1.944 exptl absorpt correction T max 1.00000 \_exptl\_absorpt\_correction\_T min 0.97777 exptl absorpt correction type multi-scan exptl absorpt process details CrysAlisPro, Oxford Diffraction Ltd., Version 1.171.33.66 (release 28-04-2010 CrysAlis171 .NET) (compiled Apr 28 2010,14:27:37) Empirical absorption correction using spherical harmonics, implemented in SCALE3 ABSPACK scaling algorithm. ? exptl crystal colour exptl crystal colour modifier dark exptl crystal colour primary orange exptl crystal density diffrn 1.760 exptl crystal density meas ?

\_exptl\_crystal density method 'not measured' exptl crystal description irregular exptl crystal F 000 720 exptl crystal size max 0.11 \_exptl\_crystal size mid 0.07 exptl crystal size min 0.06 '?' exptl special details diffrn reflns av R equivalents 0.0292 diffrn reflns av unetI/netI 0.0792 diffrn reflns limit h max 13 diffrn reflns limit h min -11 diffrn reflns limit k max 14 diffrn reflns limit k min -14 diffrn reflns limit 1 max 17 diffrn reflns limit 1 min -17 diffrn reflns number 5937 diffrn reflns theta full 25.00 diffrn reflns theta max 29.0122 diffrn reflns theta min 3.5632 diffrn ambient temperature 220.0 diffrn detector area resol mean 16.1978 diffrn measured fraction theta full 0.997 diffrn measured fraction theta\_max 0.850 diffrn measurement details # type start end width exp.time 1 omega -90.00 23.00 1.0000 50.0000 omega theta kappa phi frames -18.3269 0.0000 -30.0000 113 \_ #\_\_\_type\_\_start\_\_\_end\_\_\_\_width\_\_\_\_exp.time\_ 2 omega -90.00 -64.00 1.0000 50.0000 theta kappa phi omega frames -18.3269 179.0000 -90.0000 26 # type\_start\_end\_width\_exp.time\_ 3 omega -40.00 -15.00 1.0000 50.0000 omega theta kappa phi frames -18.3269 -37.0000 -120.0000 25 -# type start end width exp.time 4 omega -12.00 38.00 1.0000 50.0000 omega theta kappa phi frames 18.5612 84.0000 -128.0000 50 diffrn measurement device type 'Xcalibur, Eos, Gemini ultra' diffrn measurement method '\w scans' diffrn orient matrix UB 11 0.0127488000 diffrn orient matrix UB 12 0.0566904000 diffrn orient matrix UB 13 0.0255021000

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\_computing\_structure\_refinement 'SHELXL-97 (Sheldrick, 1997)' computing structure solution 'A short history of SHELX (Sheldrick, 2007)' refine diff density max 2.579 refine diff density min -0.401refine diff density rms 0.133 refine ls extinction coef ? refine ls extinction method none \_refine\_ls\_goodness of fit ref 0.905 refine ls hydrogen treatment riding refine ls matrix type full refine ls number parameters 178 refine ls number reflns 3161 refine ls number restraints 0 refine ls R factor all 0.0803 refine ls R factor gt 0.0458 refine ls restrained S all 0.905 refine ls shift/su max 0.000 refine ls shift/su mean 0.000 refine ls structure factor coef Fsqd refine ls weighting details calc w=1/[ $s^2(Fo^2)$ +(0.0722P)<sup>2</sup>+0.0000P] where P=(Fo<sup>2</sup>+2Fc<sup>2</sup>)/3 refine ls weighting scheme calc refine ls wR factor gt 0.1190 refine ls wR factor ref 0.1264 refine special details

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Refinement of F<sup>2</sup> against ALL reflections. The weighted R-factor wR and goodness of fit S are based on F<sup>2</sup>, conventional R-factors R are based on F, with F set to zero for negative F<sup>2</sup>. The threshold expression of  $F^2^> 2 \text{sigma}(F^2^>)$  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<sup>2</sup> are statistically about twice as large as those based on F, and R-factors based on ALL data will be even larger.

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atom sites solution hydrogens
                               geom
atom sites solution primary
                               direct
atom sites solution secondary
                               difmap
loop
  atom site label
  _atom_site type symbol
  atom site fract x
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 atom site fract z
 atom site U iso or equiv
  atom site adp type
 _atom_site_occupancy
 atom site symmetry multiplicity
 atom site calc flag
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\_atom\_site refinement flags atom site disorder assembly atom site disorder group Ag Ag 0.24039(5) -0.33496(5) 0.68018(3) 0.03852(17) Uani 1 1 d . . . Fe Fe 0.5000 0.0000 0.5000 0.0284(3) Uani 1 2 d S . . N4 N -0.5077(5) 0.1670(5) -0.0684(4) 0.0435(13) Uani 1 1 d . . . C13 C -0.0455(6) 0.0866(6) 0.1849(5) 0.0421(15) Uani 1 1 d . . . C1 C 0.3644(6) -0.2481(6) 0.5810(4) 0.0365(14) Uani 1 1 d . . . N3 N 0.3104(5) 0.0320(5) 0.3931(3) 0.0336(11) Uani 1 1 d . . . C7 C 0.3193(6) 0.0890(7) 0.3055(4) 0.0440(16) Uani 1 1 d . . . H7 H 0.4074 0.1164 0.2899 0.053 Uiso 1 1 calc R . . C4 C 0.0662(6) 0.0073(7) 0.3472(5) 0.0460(17) Uani 1 1 d . . . H4 H -0.0198 -0.0238 0.3639 0.055 Uiso 1 1 calc R . . N2 N 0.4236(5) -0.1816(5) 0.5338(4) 0.0387(12) Uani 1 1 d . . . C9 C -0.2619(6) 0.1272(6) -0.0431(5) 0.0443(16) Uani 1 1 d ... H9 H -0.1773 0.1141 -0.0715 0.053 Uiso 1 1 calc R . . C8 C -0.3865(6) 0.1457(7) -0.1047(5) 0.0454(17) Uani 1 1 d . . . H8 H -0.3839 0.1430 -0.1751 0.054 Uiso 1 1 calc R . . C10 C -0.2667(6) 0.1285(5) 0.0605(4) 0.0361(14) Uani 1 1 d . . . C12 C -0.5100(7) 0.1703(7) 0.0302(5) 0.0491(17) Uani 1 1 d . . . H12 H -0.5952 0.1874 0.0565 0.059 Uiso 1 1 calc R . . C11 C -0.3926(6) 0.1499(6) 0.0978(5) 0.0474(17) Uani 1 1 d . . . H11 H -0.3996 0.1507 0.1677 0.057 Uiso 1 1 calc R . . C5 C 0.0750(6) 0.0670(6) 0.2576(4) 0.0372(14) Uani 1 1 d . . . C14 C -0.1431(6) 0.1066(6) 0.1266(5) 0.0391(14) Uani 1 1 d . . . C3 C 0.1852(6) -0.0067(7) 0.4127(4) 0.0444(16) Uani 1 1 d . . . H3 H 0.1770 -0.0460 0.4747 0.053 Uiso 1 1 calc R . . C6 C 0.2048(6) 0.1100(7) 0.2358(4) 0.0459(17) Uani 1 1 d . . . H6 H 0.2149 0.1523 0.1754 0.055 Uiso 1 1 calc R . . N1 N 0.3931(5) 0.0652(5) 0.6215(4) 0.0377(12) Uani 1 1 d . . . C2 C 0.3374(6) 0.0963(6) 0.6905(4) 0.0342(13) Uani 1 1 d . . .

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#### \_geom\_special\_details

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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.

## loop\_

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C12 C11 1.393(9) . ? C5 C6 1.389(9) . ? N1 C2 1.158(7) . ? C2 Ag 2.080(6) 2\_556 ?

loop\_

geom angle atom site label 1 \_geom\_angle\_atom\_site\_label 2 \_geom\_angle\_atom\_site label 3 \_geom\_angle geom angle site symmetry 1 \_geom\_angle\_site\_symmetry\_3 \_geom\_angle\_publ\_flag N4 C8 C9 123.1(6) . . ? N4 C12 C11 123.4(6) . . ? C13 C14 C10 176.5(7) . . ? C1 Ag N4 98.6(2) . 2 445 ? C1 N2 Fe 152.2(5) . . ? N3 Fe N3 180.00(14) 3 656 . ? N3 C7 C6 123.3(6) . . ? N3 C3 C4 123.8(5) . . ? C7 N3 Fe 121.0(4) . . ? C4 C5 C13 122.6(6) . . ? C4 C5 C6 118.2(5) . . ? N2 Fe N3 90.12(18) 3\_656 3\_656 ? N2 Fe N3 90.12(18) . . ? N2 Fe N3 89.88(18) 3 656 . ? N2 Fe N3 89.88(18) . 3 656 ? N2 Fe N2 180.0 3 656 .? N2 C1 Ag 167.7(6) . . ? C9 C10 C14 120.5(6) . . ? C8 N4 Ag 122.4(4) . 2 455 ? C10 C9 C8 118.3(6) . . ? C10 C11 C12 118.9(6) . . ? C12 N4 Ag 118.9(4) . 2\_455 ? C12 N4 C8 117.9(5) . . ? C11 C10 C9 118.4(5) . . ? C11 C10 C14 121.1(6) . . ? C5 C4 C3 119.1(6) . . ? C5 C6 C7 118.5(6) . . ? C14 C13 C5 177.8(7) . . ? C3 N3 Fe 121.9(4) . . ? C3 N3 C7 117.1(5) . . ? C6 C5 C13 119.2(6) . . ? N1 Fe N3 89.56(17) 3 656.? N1 Fe N3 90.44(17) . . ? N1 Fe N3 90.44(17) 3\_656 3\_656 ? N1 Fe N3 89.56(17) . 3\_656 ? N1 Fe N2 92.9(2). 3 656? N1 Fe N2 92.9(2) 3 656 . ?

N1 Fe N2 87.1(2) . . ? N1 Fe N2 87.1(2) 3\_656 3\_656 ? N1 Fe N1 179.999(1) 3\_656 . ? N1 C2 Ag 172.8(5) . 2\_556 ? C2 Ag N4 96.90(19) 2\_546 2\_445 ? C2 Ag C1 163.4(2) 2\_546 . ? C2 N1 Fe 176.8(5) . . ?