

AUSTRALIAN JOURNAL OF CHEMICAL SCIENCE

publishing research papers from all fields of chemical science, including synthesis, structure, new materials, macromolecules, supramolecular chemistry, biological chemistry, nanotechnology, surface chemistry, and analytical techniques. Volume 55, 2002 © CSIRO 2002

All enquiries and manuscripts should be directed to:

Dr Alison Green Australian Journal of Chemistry – an International Journal for Chemical Science



CSIRO PUBLISHING PO Box 1139 (150 Oxford St) Collingwood, Vic. 3066, Australia

Telephone: +61 3 9662 7630 Fax: +61 3 9662 7611 E-mail: publishing.ajc@csiro.au

Published by **CSIRO** PUBLISHING for CSIRO and the Australian Academy of Science

www.publish.csiro.au/journals/ajc

```
10.1071/CH02065_AC
© CSIRO 2002
Accessory Publication: Aust. J. Chem., 2002, 55(5), 311-313.
data_Mndca2H2O2Me4pyz2
                                  SHELXL-97
audit creation method
_chemical_name_systematic
;
?
;
_chemical_name_common
                                  ?
_chemical_melting_point
                                 ?
_chemical_formula_moiety
                                 ?
_chemical_formula_sum
 'C20 H28 Mn N10 O2'
                                 495.46
_chemical_formula_weight
loop_
_atom_type_symbol
_atom_type_description
_atom_type_scat_dispersion_real
 _atom_type_scat_dispersion_imag
 _atom_type_scat_source
 'C' 'C' 0.0033
                   0.0016
 'International Tables Vol C Tables 4.2.6.8 and 6.1.1.4'
 'H' 'H' 0.0000 0.0000
 'International Tables Vol C Tables 4.2.6.8 and 6.1.1.4'
 'N' 'N' 0.0061 0.0033
 'International Tables Vol C Tables 4.2.6.8 and 6.1.1.4'
 'O' 'O' 0.0106 0.0060
 'International Tables Vol C Tables 4.2.6.8 and 6.1.1.4'
 'Mn' 'Mn' 0.3368 0.7283
 'International Tables Vol C Tables 4.2.6.8 and 6.1.1.4'
_symmetry_cell_setting
                                  Triclinic
_symmetry_space_group_name_H-M
                                  P-1
loop_
 _symmetry_equiv_pos_as_xyz
 'x, y, z'
 '-x, -y, -z'
_cell_length_a
                                  7.7131(4)
_cell_length_b
                                  9.1907(6)
                                  10.4207(6)
_cell_length_c
_cell_angle_alpha
                                  66.339(3)
_cell_angle_beta
                                  80.242(3)
_cell_angle_gamma
                                  66.184(3)
_cell_volume
                                  618.97(6)
_cell_formula_units_Z
                                  1
_cell_measurement_temperature
                                  123(2)
_cell_measurement_reflns_used
                                  ?
_cell_measurement_theta_min
                                  ?
_cell_measurement_theta_max
                                  ?
_exptl_crystal_description
                                  plate
```

_exptl_crystal_colour colourless _exptl_crystal_size_max 0.375 _exptl_crystal_size_mid 0.125 _exptl_crystal_size_min 0.025 _exptl_crystal_density_meas ? _exptl_crystal_density_diffrn 1.329 _exptl_crystal_density_method 'not measured' _exptl_crystal_F_000 259 0.570 _exptl_absorpt_coefficient_mu _exptl_absorpt_correction_type none _exptl_absorpt_correction_T_min ? _exptl_absorpt_correction_T_max ? _exptl_absorpt_process_details ? _exptl_special_details ; ? ; _diffrn_ambient_temperature 123(2)_diffrn_radiation_wavelength 0.71070 _diffrn_radiation_type MoK∖a _diffrn_radiation_source 'fine-focus sealed tube' _diffrn_radiation_monochromator graphite _diffrn_measurement_device_type 'Nonius KappaCCD' _diffrn_measurement_method 'phi scans' _diffrn_detector_area_resol_mean ? _diffrn_standards_number ? _diffrn_standards_interval_count ? _diffrn_standards_interval_time ? _diffrn_standards_decay_% ? 9078 _diffrn_reflns_number _diffrn_reflns_av_R_equivalents 0.0482 _diffrn_reflns_av_sigmaI/netI 0.0612 _diffrn_reflns_limit_h_min -10 _diffrn_reflns_limit_h_max 9 _diffrn_reflns_limit_k_min -12 _diffrn_reflns_limit_k_max 12 _diffrn_reflns_limit_l_min -13 _diffrn_reflns_limit_l_max 13 _diffrn_reflns_theta_min 2.61 _diffrn_reflns_theta_max 28.38 _reflns_number_total 2940 _reflns_number_gt 2486 _reflns_threshold_expression >2sigma(I) ? _computing_data_collection ? _computing_cell_refinement _computing_data_reduction ? _computing_structure_solution 'SHELXS-97 (Sheldrick, 1990)' _computing_structure_refinement 'SHELXL-97 (Sheldrick, 1997)' _computing_molecular_graphics ? computing publication material ? _refine_special_details 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 $F^{2^{,}} > 2 \text{sigma}(F^{2^{,}})$ is used only for calculating R-factors(gt) etc. and is not relevant to the choice of reflections for refinement. R-factors based on $F^{2^{,}}$ are statistically about twice as large as those based on F, and Rfactors based on ALL data will be even larger.

```
_refine_ls_structure_factor_coef Fsqd
_refine_ls_matrix_type
                                  full
_refine_ls_weighting_scheme
                                  calc
_refine_ls_weighting_details
 'calc w=1/[\s^2^(Fo^2^)+(0.0686P)^2^+2.5037P] where P=(Fo^2^+2Fc^2^)/3'
_atom_sites_solution_primary
                                  direct
_atom_sites_solution_secondary
                                  difmap
_atom_sites_solution_hydrogens
                                  qeom
_refine_ls_hydrogen_treatment
                                  mixed
_refine_ls_extinction_method
                                  none
_refine_ls_extinction_coef
                                  ?
_refine_ls_number_reflns
                                  2940
_refine_ls_number_parameters
                                  159
_refine_ls_number_restraints
                                  0
_refine_ls_R_factor_all
                                  0.0951
_refine_ls_R_factor_gt
                                  0.0795
_refine_ls_wR_factor_ref
                                  0.2187
_refine_ls_wR_factor_gt
                                  0.2116
_refine_ls_goodness_of_fit_ref
                                  1.206
_refine_ls_restrained_S_all
                                  1.206
_refine_ls_shift/su_max
                                  0.000
_refine_ls_shift/su_mean
                                  0.000
loop_
 _atom_site_label
 _atom_site_type_symbol
 _atom_site_fract_x
 _atom_site_fract_y
 _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
_atom_site_refinement_flags
_atom_site_disorder_assembly
_atom_site_disorder_group
Mn1 Mn -0.5000 0.5000 0.5000 0.0198(3) Uani 1 2 d S . .
01 0 -0.5680(5) 0.5199(6) 0.7037(4) 0.0355(9) Uani 1 1 d . . .
H1 H -0.497(13) 0.527(11) 0.751(9) 0.08(3) Uiso 1 1 d . . .
H2 H -0.669(10) 0.536(8) 0.739(7) 0.048(19) Uiso 1 1 d . . .
N1 N -0.3604(5) 0.6889(5) 0.4385(4) 0.0279(9) Uani 1 1 d . . .
C1 C -0.2437(6) 0.7467(5) 0.4122(5) 0.0231(9) Uani 1 1 d . . .
N2 N -0.1292(5) 0.8284(5) 0.3766(5) 0.0333(10) Uani 1 1 d . . .
C2 C 0.0516(6) 0.7471(5) 0.4052(5) 0.0231(9) Uani 1 1 d . . .
N3 N 0.2122(5) 0.6918(5) 0.4235(5) 0.0295(9) Uani 1 1 d . . .
N4 N -0.2928(5) 0.5426(5) 0.8311(4) 0.0275(8) Uani 1 1 d . . .
C3 C -0.2578(6) 0.6893(6) 0.7699(5) 0.0226(9) Uani 1 1 d . . .
C4 C -0.0733(6) 0.6850(6) 0.7602(5) 0.0221(9) Uani 1 1 d . . .
```

```
N5 N 0.0736(5) 0.5352(5) 0.8106(4) 0.0223(8) Uani 1 1 d . . .
C5 C 0.0390(6) 0.3903(6) 0.8708(4) 0.0219(9) Uani 1 1 d . . .
C6 C -0.1459(6) 0.3944(6) 0.8818(5) 0.0238(9) Uani 1 1 d . . .
C7 C -0.4255(7) 0.8531(6) 0.7155(6) 0.0337(11) Uani 1 1 d . . .
H7A H -0.5411 0.8280 0.7324 0.051 Uiso 1 1 calc R . .
H7B H -0.4093 0.9100 0.6146 0.051 Uiso 1 1 calc R . .
H7C H -0.4354 0.9282 0.7639 0.051 Uiso 1 1 calc R .
C8 C -0.0308(7) 0.8463(6) 0.6956(5) 0.0323(11) Uani 1 1 d . . .
H8A H 0.1059 0.8169 0.6987 0.048 Uiso 1 1 calc R . .
H8B H -0.0980 0.9211 0.7482 0.048 Uiso 1 1 calc R . .
H8C H -0.0729 0.9056 0.5980 0.048 Uiso 1 1 calc R .
C9 C 0.2073(7) 0.2265(6) 0.9243(5) 0.0303(10) Uani 1 1 d . . .
H9A H 0.3228 0.2520 0.9066 0.045 Uiso 1 1 calc R . .
H9B H 0.2166 0.1515 0.8761 0.045 Uiso 1 1 calc R . .
H9C H 0.1924 0.1693 1.0253 0.045 Uiso 1 1 calc R . .
C10 C -0.1925(8) 0.2361(7) 0.9487(6) 0.0380(12) Uani 1 1 d . . .
H10A H -0.3296 0.2678 0.9452 0.057 Uiso 1 1 calc R . .
H10B H -0.1516 0.1786 1.0466 0.057 Uiso 1 1 calc R . .
H10C H -0.1267 0.1588 0.8981 0.057 Uiso 1 1 calc R . .
loop
_atom_site_aniso_label
_atom_site_aniso_U_11
_atom_site_aniso_U_22
_atom_site_aniso_U_33
_atom_site_aniso_U_23
_atom_site_aniso_U_13
 _atom_site_aniso_U_12
Mn1 0.0107(4) 0.0244(5) 0.0277(5) -0.0114(4) 0.0028(3) -0.0094(3)
01 0.0196(17) 0.066(3) 0.0336(19) -0.0275(19) 0.0098(15) -0.0231(18)
N1 0.0201(18) 0.029(2) 0.038(2) -0.0125(17) -0.0002(16) -0.0116(16)
\texttt{C1} \ \texttt{0.0164(19)} \ \texttt{0.019(2)} \ \texttt{0.030(2)} \ \texttt{-0.0080(17)} \ \texttt{-0.0020(16)} \ \texttt{-0.0024(16)}
N2 \ 0.0177(18) \ 0.0198(19) \ 0.058(3) \ -0.0064(18) \ -0.0071(18) \ -0.0077(15)
C2 0.019(2) 0.019(2) 0.033(2) -0.0098(18) 0.0013(17) -0.0088(17)
N3 0.0179(18) 0.027(2) 0.044(2) -0.0131(18) -0.0004(16) -0.0090(16)
N4 0.0195(18) 0.029(2) 0.038(2) -0.0138(17) -0.0075(16) -0.0077(16)
\texttt{C3} \ \texttt{0.020(2)} \ \texttt{0.022(2)} \ \texttt{0.025(2)} \ \texttt{-0.0080(17)} \ \texttt{-0.0007(16)} \ \texttt{-0.0070(17)}
C4 0.022(2) 0.024(2) 0.023(2) -0.0110(17) 0.0036(16) -0.0116(17)
N5 0.0187(17) 0.0280(19) 0.0235(18) -0.0113(15) 0.0027(14) -0.0111(15)
C5 0.020(2) 0.025(2) 0.0181(19) -0.0072(17) 0.0012(15) -0.0084(17)
C6 0.024(2) 0.025(2) 0.022(2) -0.0029(17) -0.0006(16) -0.0135(18)
C7 \quad 0.026(2) \quad 0.027(2) \quad 0.038(3) \quad -0.010(2) \quad -0.004(2) \quad -0.0002(19)
C8 \ 0.038(3) \ 0.029(2) \ 0.038(3) \ -0.013(2) \ 0.007(2) \ -0.021(2)
C9 0.029(2) 0.026(2) 0.027(2) -0.0033(19) -0.0029(19) -0.0068(19)
C10 \ 0.038(3) \ 0.031(3) \ 0.041(3) \ 0.002(2) \ -0.003(2) \ -0.022(2)
```

_geom_special_details

;

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_

```
_geom_bond_atom_site_label_1
 _geom_bond_atom_site_label_2
 _geom_bond_distance
 _geom_bond_site_symmetry_2
 _geom_bond_publ_flag
Mn1 01 2.162(4) 2_466 ?
Mn1 O1 2.162(4) . ?
Mn1 N1 2.229(4) . ?
Mn1 N1 2.229(4) 2_466 ?
Mn1 N3 2.232(4) 1_455 ?
Mn1 N3 2.232(4) 2_566 ?
O1 H1 0.84(9) . ?
O1 H2 0.79(7) . ?
N1 C1 1.163(6) . ?
C1 N2 1.297(6) . ?
N2 C2 1.309(6) . ?
C2 N3 1.150(6) . ?
N3 Mn1 2.232(4) 1_655 ?
N4 C6 1.341(6) . ?
N4 C3 1.357(6) . ?
C3 C4 1.396(6) . ?
C3 C7 1.500(6) . ?
C4 N5 1.347(6) . ?
C4 C8 1.508(6) . ?
N5 C5 1.341(6) . ?
C5 C6 1.397(6) . ?
C5 C9 1.501(6) . ?
C6 C10 1.501(6) . ?
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
O1 Mn1 O1 179.998(1) 2_466 . ?
O1 Mn1 N1 89.78(15) 2_466 . ?
O1 Mn1 N1 90.22(15) . . ?
O1 Mn1 N1 90.22(15) 2_466 2_466 ?
O1 Mn1 N1 89.78(15) . 2_466 ?
N1 Mn1 N1 179.999(1) . 2_466 ?
O1 Mn1 N3 89.69(15) 2_466 1_455 ?
O1 Mn1 N3 90.32(15) . 1_455 ?
N1 Mn1 N3 96.01(14) . 1_455 ?
N1 Mn1 N3 83.99(14) 2_466 1_455 ?
O1 Mn1 N3 90.32(15) 2_466 2_566 ?
O1 Mn1 N3 89.68(15) . 2_566 ?
N1 Mn1 N3 83.99(14) . 2_566 ?
N1 Mn1 N3 96.01(14) 2_466 2_566 ?
N3 Mn1 N3 179.998(2) 1_455 2_566 ?
H1 O1 H2 109(7) . . ?
H1 O1 Mn1 125(5) . . ?
H2 O1 Mn1 125(5) . . ?
C1 N1 Mn1 161.1(4) . .
N1 C1 N2 173.3(5) . . ?
```

C1 N2 C2 121.0(4) . . ? N3 C2 N2 173.0(5) . . ? C2 N3 Mn1 159.3(4) . 1_655 ? C6 N4 C3 118.3(4) . . ? N4 C3 C4 120.6(4) . . ? N4 C3 C7 117.0(4) . . ? C4 C3 C7 122.4(4) . . ? N5 C4 C3 120.7(4) . . ? N5 C4 C8 117.7(4) . . ? C3 C4 C8 121.7(4) . . ? C5 N5 C4 118.8(4) . . ? N5 C5 C6 120.7(4) . . ? N5 C5 C9 116.8(4) . . ? C6 C5 C9 122.5(4) . . ? N4 C6 C5 121.0(4) . . ? N4 C6 C10 116.1(4) . . ? C5 C6 C10 122.8(4) . . ? loop_ _geom_hbond_atom_site_label_D _geom_hbond_atom_site_label_H _geom_hbond_atom_site_label_A _geom_hbond_distance_DH _geom_hbond_distance_HA _geom_hbond_distance_DA _geom_hbond_angle_DHA _geom_hbond_site_symmetry_A _geom_hbond_publ_flag # # Short contacts # ===== ========= # # Η D А D-H Н...А D...A D-H...A symm publ ____ # _ _ ___ ____ _____ ___ _ ____ 01 H1 0.84(9) 1.98(10) 2.801(5) 169(8) 1_555 no N4O1 H2 N5 0.79(7) 2.00(7) 2.770(5) 166(7) 1_455 NO 0.950 _diffrn_measured_fraction_theta_max _diffrn_reflns_theta_full 27.50 _diffrn_measured_fraction_theta_full 0.982 _refine_diff_density_max 1.662 _refine_diff_density_min -0.404 _refine_diff_density_rms 0.117 _publ_contact_author ; Dr. Stuart R. Batten School of Chemistry PO Box 23 Monash University 3800 AUSTRALIA ; _publ_contact_author_phone ' +61 3 9905 4606 ' ′ +61 3 9905 4597 ′ _publ_contact_author_fax _publ_contact_author_email

```
Keith.S.Murray@sci.monash.edu.au '
loop_
_publ_author_name
_publ_author_address
'Kutasi, Anna M.'
School of Chemistry
PO Box 23
Monash University 3800
Australia
;
'Batten, Stuart R.'
;
School of Chemistry
PO Box 23
Monash University 3800
Australia
;
'Harris, Alexander R.'
;
School of Chemistry
PO Box 23
Monash University 3800
Australia
;
'Moubaraki, Boujemaa'
;
School of Chemistry
PO Box 23
Monash University 3800
Australia
;
'Murray, Keith S.'
;
School of Chemistry
PO Box 23
Monash University 3800
Australia
;
_publ_section_title
;
 Structural and Magnetic Properties of the Coordination
 Polymer Mn(dca)2(H2O)2.2Me4pyz, dca =
 Dicyanamide (N(CN)2-), Me4pyz = Tetramethylpyrazine
;
```

#=END OF CIF