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Synthetic, Crystallographic, and Computational Study of Copper(II) Complexes of Ethylenediaminetetracarboxylate Ligands

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# CIF-file generated for C13H34BaCuN2O16 P-1 q1106
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data_global
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# 0. AUDIT DETAILS

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_audit_creation_method
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PLATON <TABLE ACC> option      (version :: 260511)
SHELXL97-2 & Manual Editing
;
_audit_update_record
;
First refinement : '2006-04-21 12:23:22'
2012-09-10      updated : H in riding mode
;

=====
# 1. SUBMISSION DETAILS

_publ_contact_author_name      # Name of author for correspondence
;
Drs. A. Meetsma
;
_publ_contact_author_address    # Address of author for correspondence
;
Crystal Structure Center, Chemical Physics,
Zernike Institute for Advanced Materials,
University of Groningen,
Nijenborgh 4,
NL-9747 AG Groningen, The Netherlands.
;
_publ_contact_author_email      A.Meetsma@rug.nl
_publ_contact_author_fax        '+31 50 3634441'
_publ_contact_author_phone      '+31 50 3634368'

_publ_requested_journal         'Inorganic Chemistry.'
# Publication choise FI, CI or EI for Inorganic
#           FM, CM or EM for Metal-organic
#           FO, CO or EO for Organic
_publ_requested_category        ?
_publ_requested_coeditor_name   ?

_publ_contact_letter            # Include date of submission
;
Date of submission : 2012-09-10 14:12:07

Consider this CIF submission for deposition of the first
X-ray structure of a manuscript to be submitted to : 'Inorganic Chemistry.'
(Our Compound_Identification_Code : q1106)
;

=====
# 2. PROCESSING SUMMARY (JOURNAL OFFICE ONLY)

_journal_date_recd_electronic   ?
_journal_date_to_coeditor       ?
_journal_date_from_coeditor     ?
_journal_date_accepted         ?
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```
_journal_date_printers_first      ?
_journal_date_printers_final     ?
_journal_date_proofs_out        ?
_journal_date_proofs_in         ?

_journal_coeditor_name          ?
_journal_coeditor_code          ?
_journal_coeditor_notes         ;
;

_journal_techeditor_code        ?
_journal_techeditor_notes       ;
;

_journal_coden_ASTM            ?
_journal_name_full              ?
_journal_year                   ?
_journal_volume                 ?
_journal_issue                  ?
_journal_page_first             ?
_journal_page_last              ?
```

```
_journal_suppl_publ_number      ?
_journal_suppl_publ_pages       ?
```

```
#=====
```

```
# 3. TITLE AND AUTHOR LIST
```

```
_publ_section_title
;
;
_publ_section_title_footnote
;
;
```

```
# The loop structure below should contain the names and addresses of all
# authors, in the required order of publication. Repeat as necessary.
```

```
loop_
_publ_author_name
_publ_author_footnote
_publ_author_address
'?' # author name
;   # author related footnote
;   # Address of this author
;
'Meetsma, Auke'
;   ? # author related footnote
;
;
Crystal Structure Center, Chemical Physics,
Zernike Institute for Advanced Materials,
University of Groningen,
Nijenborgh 4,
NL-9747 AG Groningen, The Netherlands.
;
```

```
#=====
```

```

# 4. TEXT

_publ_section_synopsis
;
?
;
_publ_section_abstract
;
?
;

# Insert blank lines between paragraphs

_publ_section_comment
;
;
_publ_section_exptl_prep
;
;
_publ_section_exptl_refinement
;

The hydrogen atoms were generated by geometrical considerations,
constrained to idealized geometries, and allowed to ride on the
carrier atom with an isotropic displacement parameter related to the
equivalent displacement parameter of their carrier atoms,
with U~iso~(H) = 1.2U~eq~(C) or 1.5U~eq~(methyl C).
The methyl-groups were refined as rigid groups, which were allowed
to rotate freely.
Assigned values of bond distances for secondary C-H~2~ = 0.99 \%\AA,
The hydrogen atoms of the O-H bonds were leave free to a target value of
and their isotropic displacement parameters (IDPs) were set
to 1.5 times the U~eq~ value of their parent atom.
;

_publ_section_related_literature
;
;

# Insert blank lines between references

_publ_section_references
;

Beurskens, P.T., Beurskens, G., Gelder, R. de, Garc\'ia-Granda, S. Gould, R.O.,
Isra\"el, & Smits, J.M.M. (1999). The <i>DIRDIF99</i> program system,
Technical Report of the Crystallography Laboratory, University of Nijmegen,
The Netherlands.

Bruker, (2001). <i>SMART</i>, SAINTPLUS and <i>XPREP</i>. Software Reference
Manual Bruker AXS Inc. Madison, Wisconsin, USA.

Sheldrick, G. M. (1997). <i>SHELXL97</i>. Program for Crystal Structure
Refinement. University of G\"ottingen, Germany.

Spek, A. L. (2003). <i>J. Appl. Cryst.</i> <b>36</b>, 7--13.

;

_publ_section_figure_captions
;
Fig. 1. Perspective PLUTO drawings of the molecule illustrating the

```

configuration and the adopted numbering scheme.

Fig. 2. Molecular packing viewed down unit cell axes.

Fig. 3. Perspective ORTEP drawing of the title compound.

Displacement ellipsoids for non-H atoms are represented at the 50% probability level.

;

_publ_section_acknowledgements

;

;

#=====

5. CHEMICAL DATA

_chemical_name_systematic

;

;

_chemical_name_common ?

_chemical_melting_point ?

_chemical_formula_moiety

'C13 H18 Ba Cu N2 O8, 8(H2 O)'

Ex: 'C12 H16 N2 O6, H2 O' and '(Cd 2+)3, (C6 N6 Cr 3-)2, 2(H2 O)'

_chemical_formula_structural ?

_chemical_formula_sum

'C13 H34 Ba Cu N2 O16'

_chemical_formula_iupac ?

_chemical_formula_weight 675.29

_chemical_compound_source 'see text'

loop_

_atom_type_symbol

_atom_type_description

_atom_type_scat_dispersion_real

_atom_type_scat_dispersion_imag

_atom_type_scat_source

O O 0.0106 0.0060

'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'

Cu Cu 0.3201 1.2651

'International Tables Vol C Tables 4.2.6.8 and 6.1.1.4'

Ba Ba -0.3244 2.2819

'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'

C C 0.0033 0.0016

'International Tables Vol C Tables 4.2.6.8 and 6.1.1.4'

#=====

6. CRYSTAL DATA

_symmetry_cell_setting

triclinic

_symmetry_space_group_name_Hall '-P 1'

_symmetry_space_group_name_H-M 'P -1'

_symmetry_Int_Tables_number 2

loop_

_symmetry_equiv_pos_site_id

_symmetry_equiv_pos_as_xyz

1 x,y,z

2 -x,-y,-z

```

_cell_length_a          9.0778(4)
_cell_length_b          11.8362(5)
_cell_length_c          11.9806(5)
_cell_angle_alpha       101.056(1)
_cell_angle_beta        108.737(1)
_cell_angle_gamma       95.195(1)
_cell_volume            1180.29(9)
_cell_formula_units_Z   2

_cell_measurement_temperature 100(1)
_cell_measurement_reflns_used 9337
_cell_measurement_theta_min    2.24
_cell_measurement_theta_max    29.65
_cell_special_details
;

The final unit cell was obtained from the xyz centroids of
9337 reflections after integration using the SAINTPLUS
software package (Bruker, 2000).

Reduced cell calculations did not indicate any higher metric lattice symmetry
and examination of the final atomic coordinates of the structure did not
yield extra symmetry elements (Spek, 1988; Le Page 1987, 1988)
;

_exptl_crystal_description      'Block'
_exptl_crystal_colour          'blue'
_exptl_crystal_size_max         0.52
_exptl_crystal_size_mid         0.37
_exptl_crystal_size_min         0.31
_exptl_crystal_size_rad         ?
_exptl_crystal_density_meas    ?
_exptl_crystal_density_diffrn  1.900
_exptl_crystal_density_method   'not measured'
_exptl_crystal_F_000            678
_exptl_absorpt_coefficient_mu   2.637
_exptl_absorpt_correction_type  multi-scan
_exptl_absorpt_process_details  '(SADABS, Bruker, 2001))'
_exptl_absorpt_correction_T_min 0.4069
_exptl_absorpt_correction_T_max 0.4953

#=====
# 7. EXPERIMENTAL DATA

_exptl_special_details
;
;
_diffrn_ambient_temperature    100(1)
_diffrn_radiation_wavelength   0.71073
_diffrn_radiation_type          'MoK\alpha'
_diffrn_radiation_source        'fine focus sealed Siemens Mo tube '
_diffrn_radiation_monochromator 'parallel mounted graphite'
_diffrn_radiation_detector
;
    CCD area-detector
;
_diffrn_measurement_device_type
;
    Bruker Smart Apex; CCD area detector
;
_diffrn_measurement_method      '\f and \w scans'
_diffrn_special_details
;
    Crystal into the cold nitrogen stream of the low-temparature unit

```

```

(KRYOFLEX, (Bruker, 2000)).  

;  

_diffrn_detector_area_resol_mean      66.06  

  

_diffrn_standards_number            0  

_diffrn_standards_interval_count    ?  

_diffrn_standards_interval_time     ?  

_diffrn_standards_decay_%          0  

  

loop_  

_diffrn_standard_refln_index_h  

_diffrn_standard_refln_index_k  

_diffrn_standard_refln_index_l  

? ? ?  

  

# number of measured reflections (redundant set)  

_diffrn_reflns_number              10786  

_diffrn_reflns_av_R_equivalents    0.0133  

_diffrn_reflns_av_sigmaI/netI     0.0213  

_diffrn_reflns_limit_h_min        -11  

_diffrn_reflns_limit_h_max        12  

_diffrn_reflns_limit_k_min        -15  

_diffrn_reflns_limit_k_max        15  

_diffrn_reflns_limit_l_min        -15  

_diffrn_reflns_limit_l_max        15  

_diffrn_reflns_theta_min          2.24  

_diffrn_reflns_theta_max          28.28  

_diffrn_measured_fraction_theta_max 0.960  

_diffrn_reflns_theta_full         25.00  

_diffrn_measured_fraction_theta_full 0.982  

  

_diffrn_reflns_reduction_process  

;  

Intensity data were corrected for Lorentz and polarization  

effects, decay and absorption and reduced to F~o~^2~  

using SAINT (Bruker, 2000) and SADABS (Sheldrick, 2001)  

;  

# number of unique reflections  

_reflns_number_total              5613  

_reflns_number_gt                 5496  

_reflns_threshold_expression      I>2\s(I)  

  

_computing_data_collection        'SMART, (Bruker, 2001)'  

_computing_cell_refinement        'SAINT-Plus, (Bruker, 2001)'  

_computing_data_reduction         'SAINT-Plus, (Bruker, 2001)'  

_computing_structure_solution  

;  

DIRDIF-99 (Beurskens et al., 1999)  

;  

_computing_structure_refinement   'SHELXL-97 (Sheldrick, 1997)'  

_computing_molecular_graphics  

;  

PLUTO (Meetsma, 2006)  

PLATON (Spek, 2003)  

;  

_computing_publication_material   'PLATON (Spek, 2003)'  

#=====
# 8. REFINEMENT DATA  

  

_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 R-factors 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^)+(0.0174P)^2^+1.8242P] where P=(Fo^2^+2Fc^2^)/3'
_atom_sites_solution_primary heavy
_atom_sites_solution_secondary direct
_atom_sites_solution_hydrogens geom
_refine_ls_hydrogen_treatment mixed
_refine_ls_extinction_method none
_refine_ls_extinction_coef ?
_refine_ls_abs_structure_details
;
?
;
_chemical_absolute_configuration '.'
_refine_ls_abs_structure_Flack ?
_refine_ls_number_reflns 5613
_refine_ls_number_parameters 347
_refine_ls_number_restraints 0
_refine_ls_number_constraints ?
_refine_ls_R_factor_all 0.0231
_refine_ls_R_factor_gt 0.0224
_refine_ls_wR_factor_ref 0.0529
_refine_ls_wR_factor_gt 0.0526
_refine_ls_goodness_of_fit_ref 1.118
_refine_ls_restrained_S_all 1.118
_refine_ls_shift/su_max 0.001
_refine_ls_shift/su_mean 0.000
_refine_diff_density_max 1.161
_refine_diff_density_min -0.386
_refine_diff_density_rms 0.084

_vrn_publ_code_frame_time_sec 10.0
_vrn_publ_code_meas_time_hour 8.0
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```

9. ATOMIC COORDINATES AND DISPLACEMENT PARAMETERS

```

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_atom_site_label
_atom_site_type_symbol
_atom_site_thermal_displace_type
_atom_site_fract_x
_atom_site_fract_y
_atom_site_fract_z
_atom_site_occupancy
_atom_site_U_iso_or_equiv
_atom_site_calc_flag
_atom_site_refinement_flags

_atom_site_disorder_assembly
_atom_site_disorder_group

```

Ba	Ba	Uani	1.23683(1)	0.04164(1)	0.47151(1)	1.0	0.0107(1)
Cu	Cu	Uani	0.80510(3)	0.21778(2)	0.29180(2)	1.0	0.0113(1)


```

loop_
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_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
Ba 0.0107(1) 0.0102(1) 0.0123(1) 0.0030(1) 0.0052(1) 0.0026(1)
Cu 0.0121(1) 0.0116(1) 0.0125(1) 0.0059(1) 0.0049(1) 0.0044(1)
O1 0.0143(8) 0.0200(8) 0.0144(7) 0.0034(6) 0.0051(6) 0.0054(7)
O2 0.0203(9) 0.0222(9) 0.0194(8) 0.0068(7) 0.0099(7) 0.0097(7)
O3 0.0215(9) 0.0168(8) 0.0175(8) 0.0040(6) 0.0057(7) 0.0011(7)
O4 0.0361(11) 0.0172(9) 0.0226(9) 0.0063(7) 0.0101(8) 0.0072(8)
O5 0.0153(8) 0.0131(7) 0.0166(8) 0.0023(6) 0.0035(6) 0.0019(6)
O6 0.0192(8) 0.0257(9) 0.0172(8) 0.0058(7) 0.0054(7) 0.0070(7)
O7 0.0156(8) 0.0126(7) 0.0172(8) 0.0028(6) 0.0092(6) 0.0006(6)
O8 0.0152(8) 0.0148(8) 0.0201(8) 0.0078(6) 0.0093(6) 0.0045(6)
O24 0.0240(9) 0.0145(8) 0.0206(8) 0.0045(7) 0.0081(7) 0.0052(7)
O25 0.0177(9) 0.0159(8) 0.0197(8) 0.0023(7) 0.0046(7) 0.0058(7)
O27 0.0153(8) 0.0170(8) 0.0186(8) 0.0057(6) 0.0077(7) 0.0027(7)
O28 0.0170(8) 0.0157(8) 0.0161(8) 0.0039(7) 0.0054(6) 0.0046(7)
N1 0.0254(11) 0.0181(10) 0.0146(9) 0.0050(8) 0.0052(8) -0.0012(8)
N2 0.0172(10) 0.0158(9) 0.0196(10) 0.0035(8) 0.0021(8) 0.0051(8)
C1 0.0180(11) 0.0167(11) 0.0129(10) 0.0021(8) 0.0037(8) 0.0040(9)
C2 0.0267(13) 0.0207(12) 0.0221(12) 0.0091(10) 0.0065(10) 0.0050(10)
C3 0.0261(13) 0.0194(12) 0.0228(12) 0.0079(10) 0.0079(10) 0.0035(10)
C4 0.0314(14) 0.0188(11) 0.0113(10) 0.0041(9) 0.0048(9) -0.0001(10)
C5 0.0258(12) 0.0179(11) 0.0131(10) 0.0029(8) 0.0053(9) 0.0009(10)
C6 0.0217(12) 0.0148(11) 0.0127(10) 0.0019(8) 0.0009(9) -0.0021(9)
C7 0.0232(12) 0.0168(11) 0.0208(11) 0.0119(9) 0.0031(9) 0.0034(9)
C8 0.0219(12) 0.0199(12) 0.0218(12) 0.0074(9) 0.0050(10) 0.0048(10)
C9 0.0184(11) 0.0132(11) 0.0222(11) 0.0014(9) 0.0020(9) 0.0066(9)
C10 0.0154(11) 0.0137(10) 0.0202(11) 0.0003(8) 0.0036(9) 0.0037(9)
C11 0.0149(11) 0.0104(10) 0.0205(11) 0.0025(8) 0.0043(9) 0.0024(8)
C12 0.0193(11) 0.0129(10) 0.0218(11) 0.0069(9) 0.0036(9) 0.0021(9)
C13 0.0185(11) 0.0088(9) 0.0143(10) 0.0008(8) 0.0054(8) 0.0011(8)
O21 0.0219(9) 0.0184(9) 0.0252(9) 0.0011(7) 0.0047(7) 0.0049(8)
O22 0.0225(9) 0.0205(9) 0.0218(9) 0.0037(7) 0.0098(7) 0.0068(7)
O23 0.0258(10) 0.0172(9) 0.0277(10) 0.0021(8) 0.0107(8) 0.0061(8)
O26 0.0264(11) 0.0249(11) 0.0343(11) -0.0079(9) 0.0068(9) 0.0031(8)

```

```
#=====
```

```
# 10. MOLECULAR GEOMETRY
```

```

_geom_special_details
;
Bond distances, angles etc. have been calculated using the
rounded fractional coordinates. All su's are estimated
from the variances of the (full) variance-covariance matrix.
The cell esds are taken into account in the estimation of
distances, angles and torsion angles
;

loop_
_geom_bond_atom_site_label_1
_geom_bond_atom_site_label_2
_geom_bond_distance
_geom_bond_site_symmetry_1
_geom_bond_site_symmetry_2
_geom_bond_publ_flag
Ba      O1      3.0481(17)          .          .          yes
Ba      O2      2.8337(17)          .          .          yes
Ba      O24     2.8828(18)          .          .          yes
Ba      O25     2.7744(18)          .          .          yes

```

Ba	O27	2.8170(18)	.	.	yes
Ba	O28	2.9044(18)	.	.	yes
Ba	O8	2.7778(18)	.	1_655	yes
Ba	O7	2.8321(16)	.	2_756	yes
Ba	O8	2.8346(18)	.	2_756	yes
Ba	O28	2.9135(19)	.	2_756	yes
Cu	O1	1.9515(18)	.	.	yes
Cu	O3	2.5177(17)	.	.	yes
Cu	O5	2.2991(17)	.	.	yes
Cu	O7	2.0126(16)	.	.	yes
Cu	N1	2.026(2)	.	.	yes
Cu	N2	2.022(2)	.	.	yes
O1	C1	1.277(3)	.	.	yes
O2	C1	1.246(3)	.	.	yes
O3	C6	1.260(3)	.	.	yes
O4	C6	1.262(3)	.	.	yes
O5	C11	1.267(3)	.	.	yes
O6	C11	1.261(3)	.	.	yes
O7	C13	1.286(3)	.	.	yes
O8	C13	1.244(3)	.	.	yes
O24	H24'	0.73(4)	.	.	no
O24	H24	0.89(4)	.	.	no
O25	H25'	0.87(4)	.	.	no
O25	H25	0.80(3)	.	.	no
O27	H27	0.77(4)	.	.	no
O27	H27'	0.64(4)	.	.	no
O28	H28	0.79(3)	.	.	no
O28	H28'	0.77(4)	.	.	no
O21	H21	0.83(4)	.	.	no
O21	H21'	0.78(4)	.	.	no
O22	H22	0.66(4)	.	.	no
O22	H22'	0.85(3)	.	.	no
O23	H23'	0.66(4)	.	.	no
O23	H23	0.77(4)	.	.	no
O26	H26'	0.67(5)	.	.	no
O26	H26	0.80(4)	.	.	no
N1	C7	1.526(4)	.	.	yes
N1	C3	1.479(4)	.	.	yes
N1	C4	1.504(3)	.	.	yes
N2	C9	1.501(3)	.	.	yes
N2	C12	1.498(3)	.	.	yes
N2	C8	1.483(3)	.	.	yes
C1	C2	1.534(3)	.	.	no
C2	C3	1.513(4)	.	.	no
C4	C5	1.532(3)	.	.	no
C5	C6	1.527(3)	.	.	no
C7	C8	1.504(4)	.	.	no
C9	C10	1.516(3)	.	.	no
C10	C11	1.529(4)	.	.	no
C12	C13	1.512(3)	.	.	no
C2	H2'	0.99	.	.	no
C2	H2	0.99	.	.	no
C3	H3	0.99	.	.	no
C3	H3'	0.99	.	.	no
C4	H4	0.99	.	.	no
C4	H4'	0.99	.	.	no
C5	H5	0.99	.	.	no
C5	H5'	0.99	.	.	no
C7	H7	0.99	.	.	no
C7	H7'	0.99	.	.	no
C8	H8'	0.99	.	.	no
C8	H8	0.99	.	.	no
C9	H9'	0.99	.	.	no
C9	H9	0.99	.	.	no
C10	H10	0.99	.	.	no
C10	H10'	0.99	.	.	no

