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Coexisting Ferromagnetic and Ferroelectric Order in a CuCl4-based Organic-Inorganic Hybrid

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Download date: 21-07-2025

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 Drs. A. Meetsma
                                # Address of author for correspondence
_publ_contact_author_address
 Crystal Structure Center, Chemical Physics,
 Zernike Institute for Advanced Materials,
 University of Groningen,
 Nijenborgh 4,
 NL-9747 AG Groningen, The Netherlands.
                                A.Meetsma@rug.nl
_publ_contact_author_email
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publ requested journal
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# 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: 2009-09-21 13:30:04
 Please consider this CIF submission for publication as a
 Regular Structural Paper in Acta Crystallographica E.
 All authors have seen and approved this submission.
 The CIF has passed the Chester CHECKCIF routines and gives
 a satisfactory PRINTCIF file.
 The Structure Factor Listing (in CIF format) and any artwork
 (Schemes, Figures) as HPGL, Postscript, TIFF or encapsulated Postscript files
 will be transferred to Chester by ftp as instructed in Notes for Authors.
 Consider this CIF submission for deposition of the Xray-structure quoted
 in Journal Ref: Inorganica Chimica Acta 259 (2000) ...
 "A remarkable Lewis acid-base adduct: preparation...
 Compound : ....
 Your Reference number : ....
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Base.
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 (Our Compound_Identification_Code : a0007)
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_journal_date_proofs_in
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_journal_coden_ASTM
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_journal_year
_journal_volume
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# 3. TITLE AND AUTHOR LIST
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# The loop structure below should contain the names and adresses of all
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Please consider this CIF submission for deposition at the Cambridge Data

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_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.
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?
_publ_section_abstract
?
# Insert blank lines between paragraphs
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_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\sim iso\sim (H) = 1.2U\sim eq\sim (C) or 1.5U\sim eq\sim (methyl C).
 The amine-group was refined as rigid groups, which were allowed
  to rotate freely.
 Assigned values of bond distances for secondary C-H\sim2\sim = 0.97 \%A,
  aromatic C-H = 0.93 \ and N-H = 0.89 \A
;
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Allen, F.H. (2002). Acta Cryst. B58, 380-388.
 Beurskens, P.T., Beurskens, G., Gelder, R. de, Smits, J.M.M.,
```

```
The DIRDIF08 program system, Technical Report of the Crystallography
 Laboratory, University of Nijmegen, The Netherlands.
 Boeyens, J.C.A. (1978). J. Cryst. Mol. Struct. 8, 317-320.
 Bondi, A. (1964). J. Phys. Chem. 68, 441-451.
 Bruker, (2007). SMART (Version 5.632), SAINT-Plus (Version 7.46a) and
 SADABS (Version 2.10). Bruker AXS Inc., Madison, Wisconsin, USA.
 Hahn, T. (1983). Ed. International Tables for Crystallography,
 Volume A, Space-group symmetry, Kluwer Academic Publishers,
 Dordrecht, The Netherlands.
Meetsma, A. (2009). Extended version of the program PLUTO.
 University of Groningen, The Netherlands. (unpublished).
 Sheldrick, G.M. (2008). Acta Cryst. A64, 112-122.
Spek, A.L. (2003). J. Appl. Cryst. 36, 7-13.
Wilson, A.J.C. (1992). Ed. International Tables for Crystallography,
 Volume C, Kluwer Academic Publishers, Dordrecht, The Netherlands.
_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.
          The H-atoms have been omitted to improve clarity.
          The H-atoms are drawn with an arbitrary radius.
  Fig. 4. Portion of the crystal packing, showing
          N-H...Cl hydrogen bonds shown as dashed lines.
_publ_section_acknowledgements
# 5. CHEMICAL DATA
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;
_chemical_name_common
_chemical_melting_point
_chemical_formula_moiety
'2(C8 H12 N), Cl4 Cu'
# 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
'C16 H24 C14 Cu N2'
_chemical_formula_iupac
_chemical_formula_weight
                                      449.72
```

Garc\'ia-Granda, S. & Gould, R.O. (2008).

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'see text'
chemical compound source
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_atom_type_description
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_atom_type_scat_dispersion_imag
_atom_type_scat_source
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 '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'
Cl Cl 0.1484 0.1585
 'International Tables Vol C Tables 4.2.6.8 and 6.1.1.4'
 н н 0.0000 0.0000
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 C C 0.0033 0.0016
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_symmetry_space_group_name_Hall
                                    '-C 2bc 2'
                                    'C m c a'
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_symmetry_Int_Tables_number
                                            64
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_symmetry_equiv_pos_as_xyz
   1 x,y,z
   2 - x, 1/2 - y, 1/2 + z
   3 \times, -y, -z
   4 - x, 1/2 + y, 1/2 - z
   5 - x, -y, -z
   6 x, 1/2+y, 1/2-z
   7 -x,y,z
   8 \times 1/2 - y \cdot 1/2 + z
  9 \frac{1}{2} + x, \frac{1}{2} + y, z
  10 1/2-x, -y, 1/2+z
  11 1/2+x, 1/2-y, -z
  12 \ 1/2-x, y, 1/2-z
  13 1/2-x, 1/2-y, -z
  14 \ 1/2 + x, y, 1/2 - z
  15 \ 1/2-x, 1/2+y, z
  16 1/2+x, -y, 1/2+z
_cell_length_a
                                      39.021(8)
_cell_length_b
                                       7.3430(15)
_cell_length_c
                                      7.3939(15)
_cell_angle_alpha
                                      90
                                     90
_cell_angle_beta
_cell_angle_gamma
                                      90
_cell_volume
                                    2118.6(7)
_cell_formula_units_Z
_cell_measurement_temperature
                                 373(1)
_cell_measurement_reflns_used
                                 2107
_cell_measurement_theta_min
                                  3.13
_cell_measurement_theta_max
                                 23.02
_cell_special_details
 The final unit cell was obtained from the xyz centroids of
   2107 reflections after integration using the SAINTPLUS
   software package (Bruker, 2007).
```

```
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)
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_exptl_crystal_colour
                               'orange'
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_exptl_crystal_size_mid
                                0.10
                                0.10
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_exptl_crystal_size_rad
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                               1.410 'not measured'
_exptl_crystal_density_diffrn
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_exptl_crystal_F_000
                               924
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                               1.535
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                               '(SADABS, (Bruker, 2007))'
_exptl_absorpt_process_details
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# 7. EXPERIMENTAL DATA
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_diffrn_radiation_wavelength
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_diffrn_radiation_type
_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, 2007)).
_diffrn_detector_area_resol_mean
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_diffrn_standards_decay_%
_diffrn_standard_refln_index_h
_diffrn_standard_refln_index_k
_diffrn_standard_refln_index_l
# number of measured reflections (redundant set)
_diffrn_reflns_number
                                7895
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_diffrn_reflns_av_sigmaI/netI
                               0.0218
```

```
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                                _9
_diffrn_reflns_limit_k_min
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_diffrn_reflns_limit_l_min
                                -9
_diffrn_reflns_limit_l_max
_diffrn_reflns_theta_min
                                 3.13
_diffrn_reflns_theta_max
                                26.72
_diffrn_measured_fraction_theta_max
                                     0.998
_diffrn_reflns_theta_full
                                     25.00
_diffrn_measured_fraction_theta_full
                                     0.998
_diffrn_reflns_reduction_process
 Intensity data were corrected for Lorentz and polarization
 effects, decay and absorption and reduced to F~o~^2^
 using SAINT-Plus & SADABS (Bruker, 2007).
# number of unique reflections
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_reflns_number_gt
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                                  'SAINT-Plus (Bruker, 2007)'
_computing_cell_refinement
                                  'SAINT-Plus'
_computing_data_reduction
_computing_structure_solution
DIRDIF-08 (Beurskens et al., 2008)
                                   'SHELXL-97 (Sheldrick, 2008)'
_computing_structure_refinement
_computing_molecular_graphics
PLATON (Spek, 2003)
PLUTO (Meetsma, 2009)
_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 > 2sigma(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.
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_refine_ls_matrix_type
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_refine_ls_weighting_scheme
                                calc
_refine_ls_weighting_details
'calc w=1/[\s^2^(Fo^2^)+(0.1044P)^2^+0.6794P] 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
                                constr
_refine_ls_extinction_method
                                none
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_chemical_absolute_configuration '.'
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_refine_ls_number_constraints
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                                0.1597
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_refine_ls_wR_factor_gt
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_refine_ls_restrained_S_all
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_refine_ls_shift/su_mean
                                0.000
_refine_diff_density_max
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_refine_diff_density_min
                              -0.723
_refine_diff_density_rms
                                0.104
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_vrn_publ_code_number_frames
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9. ATOMIC COORDINATES AND DISPLACEMENT PARAMETERS

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C2 C Uani 0.1196(2) 0.455(2) -0.1085(16) 0.5 0.185(10) . .
C3 C Uani 0.1505(3) 0.500 0.00000 1.000 0.201(9) . .
                                               0.210(7) . .
C4 C Uani 0.1699(2) 0.6185(16) -0.1079(16) 1.0
C5 C Uani 0.2045(2) 0.6148(16) -0.0941(17) 1.0
                                               0.183(6) . .
C6 C Uani 0.2197(4) 0.500 0.00000 1.000 0.180(10) . .
Cul Cu Uani 0.000 0.50000 0.50000 1.000 0.0495(3) .
Cl1 Cl Uani -0.05866(4) 0.500 0.50000 1.000 0.0764(6) .
C12 C1 Uani 0.000 0.27901(16) 0.28124(16) 1.000 0.0671(5) . .
H1 H Uiso 0.09425 0.43231 0.10982 0.5 0.1945 . .
                                       0.1945 . .
H1' H Uiso 0.09425 0.62571 0.03151 0.5
                                        0.2200 . .
H2 H Uiso 0.11946 0.52552 -0.21974 0.5
H2' H Uiso 0.11946 0.32705 -0.13938 0.5
                                        0.2200 . .
H4 H Uiso 0.15919 0.69821 -0.18753 1.0
                                        0.2522 . .
H5 H Uiso 0.21733 0.70086 -0.15703 1.0 0.2197 . .
H6 H Uiso 0.24355 0.500 0.00000 1.000 0.2171 . .
H9 H Uiso 0.04704 0.54756 0.10319 0.5 0.1152 . .
H9' H Uiso 0.04704 0.38624 -0.01069 0.5 0.1152 . .
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_atom_site_aniso_U_23
_atom_site_aniso_U_13
_atom_site_aniso_U_12
N1 0.070(4) 0.078(4) 0.082(4) -0.004(3) 0.036(8) -0.017(10)
C1 \quad 0.080(7) \quad 0.163(13) \quad 0.24(2) \quad 0.074(10) \quad 0.085(15) \quad 0.043(15)
C2 \ 0.152(15) \ 0.232(18) \ 0.17(2) \ -0.088(16) \ -0.050(14) \ 0.061(14)
C3 0.050(5) 0.262(19) 0.29(2) 0.134(14) 0.00
                                               0.0000
C4 0.133(8) 0.260(14) 0.237(13) 0.153(13) 0.014(6) 0.003(7)
C5 0.097(7) 0.202(12) 0.250(13) 0.033(11) 0.038(7) -0.025(6)
C6 0.081(8) 0.22(2) 0.24(2) -0.036(12) 0.00 0.0000
Cul 0.0626(6) 0.0447(5) 0.0412(5) -0.0076(3) 0.00
                                                     0.0000
Cl1 0.0604(9) 0.0866(11) 0.0822(11) 0.0013(6) 0.00
C12 0.1076(10) 0.0485(6) 0.0451(7) -0.0088(5) 0.00 0.0000
# 10. MOLECULAR GEOMETRY
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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
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_geom_bond_site_symmetry_1
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Cul Cl2 2.9192(13)
Cul Cl1 2.2890(16)
Cul Cl2 2.2912(13)
                                                         2_555
                                                                                yes
                                                                                yes
                                                         •
                                                                                yes
                 2.2890(16)
        C11
                                                         5 566
 Cu1
                                                                                yes
 Cu1
        C12
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 N1
         Н9"
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                                                                                no
 N1
         Н9
                      0.89
                                                                                no
         Н9'
 N1
                      0.89
        Н9
 Ν1
                      0.89
                                                         3 565
                                                                                no
        Н9'
                                                         3_565
 N1
                      0.89
                                                                                nο
        C2
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 C1
                                                                                no
 С1
        C2
                  1.460(13)
                                                         3 565
        С3
 C2
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                                                                                no
         C4
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                                                                                no
                   1.402(12)
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 C3
                                                                                no
 C4
         C5
                   1.354(11)
                                                                                no
 C5
         С6
                   1.244(14)
                                                                                no
 С1
        Н1'
                      0.97
                                                                                no
 C1
        Н1
                      0.97
                                                                                no
 C1
                     0.97
                                                        3 565
        Н1
                                                                                nο
         Н1'
 С1
                     0.97
                                                         3_565
 C2
         H2
                      0.97
                                                                                no
         н2'
 C2
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                                                                                nο
 C4
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                                                                                nο
 C5
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no

С6

Н6

0.93

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 C11
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 C11
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                  C12
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                                                                  3 566
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                 C12
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 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
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# 2. PROCESSING SUMMARY (JOURNAL OFFICE ONLY)
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   'Meetsma, Auke'
; ? # author related footnote
  Crystal Structure Center, Chemical Physics,
  Zernike Institute for Advanced Materials,
  University of Groningen,
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_journal_date_to_coeditor

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Nijenborgh 4,
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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 <i>U</i><iso~(H) = 1.2<i>U</i><eq~(C)
or 1.5 < i > U < /i > \sim eq \sim (methyl C). The methyl-groups were refined as rigid groups,
which were allowed to rotate freely. Assigned values of bond distances:
secondary C---H\sim2\sim = 0.99 \%A, methyl C---H\sim3\sim = 0.98 \%A, aromatic C---H=
0.95 \%A.
C---H distances were in the range of 0.95 -- 0.99(3) \%A.
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Beurskens, P. T., Beurskens, G., de Gelder, R., Garc\'ia-Granda, S., Gould, R.
O., Isra\"el, R. & Smits, J. M. M. (1999). The <i>DIRDIF99</i> program system,
Technical Report of the Crystallography Laboratory, University of Nijmegen,
The Netherlands.
Brandenburg, K. (2006). <i>DIAMOND</i>. Release 3.1. Crystal Impact GbR, Bonn,
Germany
        (2006). <i>SMART</i> (Version 5.632), <i>SAINT-Plus</i> (Version 6.45)
and <i>SADABS</i> (Version 2.10). Bruker AXS Inc., Madison, Wisconsin, USA.
Le Page, Y. (1987). <i>J. Appl. Cryst.</i> <b>20</b>, 264--269.
Le Page, Y. (1988). <i>J. Appl. Cryst.</i> <b>21</b>, 983--984.
Meetsma, A. (2007). Extended version of the program <i>PLUTO</i>. University of
Groningen, The Netherlands. (unpublished).
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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 <i>PLUTO</i> drawings of the moieties of the asymmetric
unit illustrating the configuration and the adopted numbering scheme.
Fig. 2. Molecular packing viewed down unit cell axes.
Fig. 3. Perspective <i>ORTEP</i> drawing of the anion and cation of the title
compound. Displacement ellipsoids for non-H atoms are represented at the 50%
probability level. The H-atoms have been omitted to improve clarity.
Fig. 4. Portion of the crystal packing, showing N---H...Cl hydrogen bonds shown
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_diffrn_reflns_av_sigmaI/netI 0.0251
_diffrn_reflns_limit_h_min
_diffrn_reflns_limit_h_max
_diffrn_reflns_limit_k_min
                                 _9
_diffrn_reflns_limit_k_max
_diffrn_reflns_limit_l_min
                                 -50
_diffrn_reflns_limit_l_max
_diffrn_reflns_theta_min
                                  3.02
_diffrn_reflns_theta_max
_diffrn_measured_fraction_theta_max
                                      0.939
                                       25.00
_diffrn_reflns_theta_full
_diffrn_measured_fraction_theta_full
                                       0.976
_diffrn_reflns_reduction_process
 Intensity data were corrected for Lorentz and polarization
  effects, decay and absorption and reduced to F~o~^2^
  using SAINT-Plus & SADABS (Bruker, 2006).
# number of unique reflections
_reflns_number_total
                                  2340
_reflns_number_gt
                                  2163
_reflns_threshold_expression
                               I>2\s(I)
_computing_data_collection
                                    'SMART (Bruker, 2006)'
{\tt \_computing\_cell\_refinement}
                                    'SAINT-Plus (Bruker, 2006)'
                                    'SAINT-Plus (Bruker, 2006)'
_computing_data_reduction
_computing_structure_solution
```

```
DIRDIF-99 (Beurskens et al., 1999)
_computing_structure_refinement
                                   'SHELXL-97 (Sheldrick, 1997)'
_computing_molecular_graphics
DIAMOND (Brandenburg, 2006)
 PLATON (Spek, 2003)
PLUTO (Meetsma, 2007)
_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 > 2sigma(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^)+(0.0P)^2^+9.9068P] where P=(Fo^2^+2Fc^2^)/3'
_atom_sites_solution_primary
                                heavv
_atom_sites_solution_secondary
_atom_sites_solution_hydrogens
                                 geom
_refine_ls_hydrogen_treatment
                                 constr
_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
                                  2340
                                  107
_refine_ls_number_parameters
_refine_ls_number_restraints
_refine_ls_number_constraints
_refine_ls_R_factor_all
                                  0.0437
_refine_ls_R_factor_gt
                                 0.0400
_refine_ls_wR_factor_ref
                                 0.1120
_refine_ls_wR_factor_gt
                                 0.1106
_refine_ls_goodness_of_fit_ref
                                 1.329
                                  1.329
_refine_ls_restrained_S_all
_refine_ls_shift/su_max
                                  0.000
_refine_ls_shift/su_mean
                                 0.000
_refine_diff_density_max
                                 0.733
_refine_diff_density_min
                                 -0.605
_refine_diff_density_rms
                                 0.121
_vrn_publ_code_void_volume
                                 70.6
_vrn_publ_code_frame_time_sec
                                  5.0
_vrn_publ_code_meas_time_hour
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9. ATOMIC COORDINATES AND DISPLACEMENT PARAMETERS

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_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
N N Uani 1.0138(4) 0.0064(4) 0.55580(7) 1.000 0.0165(8) . .
C1 C Uani 0.9605(6) -0.0433(5) 0.59223(9) 1.000 0.0176(10) . .
C2 C Uani 1.0436(6) 0.0920(5) 0.61820(9) 1.000 0.0212(10) . .
C3 C Uani 1.0184(6) 0.0273(5) 0.65541(9) 1.000 0.0179(10) . .
C4 C Uani 0.8791(6) 0.0978(5) 0.67676(10) 1.000 0.0219(11) . .
C5 C Uani 0.8605(6) 0.0357(6) 0.71109(10) 1.000 0.0265(11) . .
C6 C Uani 0.9804(7) -0.0957(6) 0.72429(10) 1.000 0.0265(11) . .
C7 C Uani 1.1182(7) -0.1673(6) 0.70311(11) 1.000 0.0283(14) . .
C8 C Uani 1.1375(6) -0.1052(6) 0.66918(10) 1.000 0.0241(11) . .
Cu Cu Uani 0.00000 0.00000 0.00000 1.000 0.0123(2)
Cl1 Cl Uani 0.03423(12) 0.01365(12) 0.05984(2) 1.000 0.0160(2) . .
C12 C1 Uani 0.21493(11) 0.22950(11) -0.00597(2) 1.000 0.0149(2) . .
H1 H Uiso 1.00485 -0.16910 0.59760 1.000 0.0211 . .
H1' H Uiso 0.82362 -0.04248 0.59440 1.000 0.0211 . .
H2 H Uiso 1.17755 0.10695 0.61326 1.000 0.0254 . .
H2' H Uiso 0.98343 0.21355 0.61530 1.000 0.0254 . .
H4 H Uiso 0.79652 0.18842 0.66792 1.000 0.0264 . .
H5 H Uiso 0.76482 0.08388 0.72551 1.000 0.0315 . .
H6 H Uiso 0.96822 -0.13651 0.74778 1.000 0.0315 . .
H7 H Uiso 1.19945 -0.25918 0.71191 1.000 0.0340 . .
H8 H Uiso 1.23375 -0.15377 0.65495 1.000 0.0291 . .
H9 H Uiso 0.97516 0.12294 0.55104 1.000 0.0247 . .
H9' H Uiso 0.95965 -0.07338 0.54056 1.000 0.0247 . .
H9" H Uiso 1.13931 -0.00002 0.55354 1.000 0.0247 . .
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
 \hbox{N 0.0180(15) 0.0196(14) 0.0119(13) -0.0003(12) -0.0003(12) 0.0007(13) } \\
C1 \quad 0.0199(19) \quad 0.0210(17) \quad 0.0120(15) \quad 0.0011(13) \quad 0.0006(14) \quad -0.0029(15)
C2 \ 0.032(2) \ 0.0196(17) \ 0.0120(16) \ -0.0007(14) \ 0.0014(15) \ -0.0050(16)
C4 \ 0.024(2) \ 0.0201(18) \ 0.0217(19) \ -0.0007(15) \ 0.0007(16) \ 0.0026(16)
C5 0.030(2) 0.031(2) 0.0184(18) -0.0044(16) 0.0072(16) -0.0008(18)
C6 \ 0.037(2) \ 0.028(2) \ 0.0144(17) \ 0.0008(15) \ -0.0011(17) \ -0.0066(19)
C7 0.035(3) 0.028(2) 0.022(2) 0.0021(16) -0.0064(18) 0.0061(19)
C8 0.026(2) 0.027(2) 0.0193(18) -0.0031(16) 0.0014(16) 0.0056(17)
Cu 0.0132(3) 0.0118(3) 0.0120(3) -0.0002(2) 0.0001(2) -0.0022(2)
C11 \quad 0.0183(4) \quad 0.0171(4) \quad 0.0126(4) \quad -0.0003(3) \quad 0.0000(3) \quad -0.0003(3)
\texttt{C12} \ \ \textbf{0.0133(4)} \ \ \textbf{0.0136(4)} \ \ \textbf{0.0179(4)} \ \ \textbf{0.0003(3)} \ \ \textbf{0.0001(3)} \ \ -\textbf{0.0017(3)}
# 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 _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 C12 2.2879(9) yes C12 2.8531(9) 3_455 Cu yes Cu Cl1 2.3036(8) 5_555 yes Cu C12 2.2879(9) 5 555 yes Cu C12 2.8531(9) 7_545 yes Cu Cl1 2.3036(8) yes Ν 1.490(4) C1 yes Н9" Ν 0.9100 no Н9' Ν 0.9100 no Ν Н9 0.9100 no C1 C2 1.520(5) no C2 C3 1.510(5) no СЗ С8 1.393(6) no C3 C4 1.392(6) no C5 C4 1.395(5) no C5 С6 1.383(6) no С6 C7 1.383(7) no C7 С8 1.381(6) no Н1' 0.9900 C1 no C10.9900 Н1 no C2 0.9900 Н2 no C2 Н2' 0.9900 no C4 H4 0.9500 no C5 Н5 0.9500 no С6 Н6 0.9500 no С7 Н7 0.9500 no С8 Н8 0.9500 no _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_2 $_geom_angle_site_symmetry_3$ _geom_angle_publ_flag C12 90.36(3) 3 455 5 555 C12 Cu ves C12 Cu C12 180.00 3_455 7_545 yes 5_555 5_555 C11 Cu C12 89.72(3) yes 88.17(3) C12 5_555 7_545 C11 Cu yes 5_555 C12 Cu C12 89.64(3) 7_545 yes Cl1Cu C12 89.72(3) yes Cl1 88.17(3) 3_455 Cu C12 yes C11 5_555 Cu Cl1 180.00 yes 5_555 90.28(3) C11 Cu C12 ves C11 Cu C12 91.83(3) 7 545 ves 3_455 C12 Cu C12 89.64(3) yes Cl1 Cu C12 90.28(3) 5 555 yes 5 555 yes C12 Cu C12 180.00 . C12 Cu C12 90.36(3) 7_545 yes 3 455 5 555 Cl1 Cu C12 91.83(3) yes

Cu

C12

Cu

169.15(3)

3_555

yes

Н9	N	H9 "	1	09.00					no
Н9'	N	Н9"		09.00		•	•		no
C1	N	Н9 "		09.00	•	•	•		no
C1	N	Н9		09.00	•	•	•		no
C1	N	Н9'		09.00	•	•	•		no
Н9	N G1	Н9'		09.00	•	•	•		no
N C1	C1 C2	C2 C3		.6(3)	•	•	•		yes
C2	C3	C8		.5(3) .8(4)	•	•	•		no
C4	C3	C8		.5(3)	•	•	•		no no
C2	C3	C4		.7(3)	•	•	•		no
C3	C4	C5		.2(4)		·			no
C4	C5	C6		.4(4)			•		no
C5	С6	C7		.7(4)					no
C6	С7	C8	120	.0(4)	•	•	•		no
С3	C8	C7		.3(4)	•		•		no
N	C1	Н1		10.00	•	•	•		no
N	C1	H1'		10.00	•	•	•		no
C2	C1	Н1		10.00	•	•	•		no
C2	C1	H1'		10.00	•	•	•		no
H1	C1	H1'		08.00	•	•	•		no
C1	C2	H2'		09.00	•	•	•		no
C3	C2 C2	H2 H2 '		09.00	•	•	•		no
H2	C2	H2'		09.00 08.00	•	•	•		no
C1	C2	H2		09.00	•	•	•		no no
C5	C4	H4		20.00	•	•	•		no
C3	C4	H4		20.00					no
C4	C5	Н5		20.00			•		no
C6	C5	Н5		20.00	•	•	•		no
C7	С6	Н6		20.00					no
C5	С6	Н6	1.	20.00	•		•		no
C6	С7	Н7	1.	20.00	•	•	•		no
C8	С7	Н7		20.00					no
C3	C8	Н8		19.00	•	•	•		no
C7	C8	Н8	1	19.00	•	•	•		no
loop_									
		_atomsite							
		_atom_site							
geom	torsion_	_atom_site _atom_site	e_label_3						
geom	torsion_ torsion	_acom_sice	e_raber_4						
		_site_symr	metrv 1						
		_site_symr _site_symr							
		_site_symr							
		 _site_symr							
		_publ_flag							
N	C1	C2	C3	-170.6(3)	•	•	•		no
C1	C2	C3	C4	-100.5(4)	•		•		no
C1	C2	C3	C8	80.0(5)	•	•	•	•	no
C2	C3	C4	C5	-179.6(4)	•	•	•	•	no
C8	C3	C4	C5	-0.1(6)	•	•	•	•	no
C2	C3	C8	C7	180.0(4)	•	•	ē	•	no
C4	C3	C8	C7	0.5(6)	•	•	•	•	no
C3	C4	C5	C6	0.3(6)	•	•	•	•	no
C4 C5	C5 C6	C6 C7	C7 C8	-0.8(7)	•	•	•	•	no
C5 C6	C6 C7	C 7	C3	1.1(7) -1.0(7)	•	•	•	•	no no
	C /	00	0.5	1.0(/)	•	•	•	•	110
loop_ geom	contact	_atom_site	e label 1						

```
__geom_contact_atom_site_label_1
__geom_contact_atom_site_label_2
__geom_contact_distance
__geom_contact_site_symmetry_1
__geom_contact_site_symmetry_2
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Cu	Н9'	3.7000	•	2_554	no
Cu	Н9"	3.3100	•	2_654	no
Cu	H9	3.3700	•	4_645	no
Cu	H9'	3.4800	•	4_655	no
Cu	Н9'	3.7000	•	6_455	no
Cu	Н9"	3.3100	•	6_355	no
Cu	H9	3.3700	•	8_454	no
Cl1 Cl1	Cl2 N	3.2388(12) 3.266(3)	•	• 2_654	no
Cl1	Cl2	3.6092(12)	•	3_455	no no
Cl1	C12	3.2545(12)	•	5_555	no
Cl1	N	3.494(3)	•	8_454	no
Cl1	C2	3.633(4)	•	8_454	no
C12	C11	3.2388(12)	•	0_101	no
C12	C11	3.6092(12)	-	3_555	no
C12	N	3.302(3)		6_455	no
C12	C12	3.6459(12)	<u>.</u>	3_455	no
C12	C12	3.6459(12)	•	3_555	no
C12	N	3.225(3)		4_655	no
C12	N	3.371(3)		8 <u>4</u> 54	no
C12	C11	3.2545(12)		5 <u>5</u> 55	no
C12	C12	3.6682(12)		7_545	no
C12	C12	3.6682(12)	•	7_555	no
Cl1	H1'	2.9100	•	2_554	no
Cl1	Н9"	2.3700	•	2_654	no
Cl1	H2	3.0400	•	2_654	no
Cl1	Н2'	2.9300		8_454	no
Cl1	Н9	2.7000	•	8_454	no
C12	Н9'	3.1200	•	6_455	no
C12	Н9	2.6600	•	6_455	no
C12	Н9 '	2.3200	•	4_655	no
C12	Н9 "	3.0100	•	2_654	no
C12	Н9"	3.0600	•	8_454	no
C12	H9	2.9800	•	8_454	no
N	C12	3.302(3)	•	6_555	no
N	C11	3.494(3)	•	8_655	no
N N	Cl2 Cl1	3.225(3) 3.266(3)	•	4_645 2_655	no
N	C11	3.371(3)	•	8_655	no no
C2	C12	3.633(4)	•	8_655	no
C2	Н8	2.8200	•	7_755	no
C3	Н8	2.9300	•	7_755 7_755	no
C6	Н5	2.9200	•	7_645	no
C6	Н5	3.1000	-	6_556	no
C8	Н1	2.9400			no
Н1	C8	2.9400		•	no
Н1	C11	2.9000		8_645	no
H1'	Cl1	2.9100	•	2_555	no
Н2	Н8	2.4400		7_755	no
Н2	Cl1	3.0400	•	2_655	no
Н2	Н9"	2.4300	•		no
Н2	Н8	2.5100	•		no
H2'	Cl1	2.9300	•	8_655	no
Н2'	Н9	2.5400	•	•	no
Н2'	Н4	2.4300	•	•	no
H4	H2'	2.4300	•		no
H5	C6	3.1000	•	6_456	no
H5	C6	2.9200	•	7_655	no
Н8	H2	2.4400	•	7_745	no
H8	H2	2.5100	•	•	no
H8	C2	2.8200	•	7_745	no
Н8	C3	2.9300	•	7_745	no
H9	H2'	2.5400	•	1 655	no
Н9 Н9	Cu Cl2	3.3700 2.6600	•	4_655	no
117	CIZ	2.0000	•	6_555	no

```
8_655
 Н9
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                       3.3700
                                                                                 no
                                                          8_655
 Н9
         Cl1
                       2.7000
                                                                                 no
                                                          4_645
 Н9'
         Cu
                       3.4800
                                                                                 no
 Н9'
         C12
                       3.1200
                                                          6_555
                                                                                 no
 Н9'
         C12
                       2.3200
                                                          4_645
                                                                                 no
 Н9'
                       3.7000
                                                          6_555
         Cu
                                                                                 no
 Н9'
                                                          2 555
         Cu
                       3.7000
                                                                                 no
 Н9'
                       3.4800
                                                          8 645
         Cu
                                                                                 no
 Н9"
                                                          2_655
         Cl1
                       2.3700
                                                                                 no
 Н9"
                                                          6_655
                       3.3100
         Cu
                                                                                 no
 Н9"
         C12
                       3.0600
                                                          8_655
                                                                                 no
                                                          2_655
 Н9"
         C12
                       3.0100
                                                                                 no
 Н9"
         Н2
                       2.4300
                                                                                 no
 Н9"
                                                          2_655
                       3.3100
         Cu
                                                                                 no
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_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
            D - H H...A
#D
                           D...A
                                     D - H...A symm(A)
     H A
#
Ν
        Н9
                C12
                             0.9100
                                         2.6600
                                                  3.302(3)
                                                                128.00
                                                                         6_555 yes
Ν
        Н9
                 C11
                             0.9100
                                         2.7000
                                                  3.494(3)
                                                                147.00
                                                                         8_655 yes
                                                                         4_645 yes
        Н9'
                             0.9100
                                                  3.225(3)
                C12
                                         2.3200
                                                                172.00
Ν
        Н9"
                                                                         2_655 yes
                C11
                             0.9100
                                         2.3700
                                                  3.266(3)
                                                                169.00
Ν
```

8 655

no

Н9

C12

2.9800

#===END of Crystallographic Information File