

Supplementary materials.

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Charge-density analysis and electrostatic properties of 2-Carboxy-4-methylanilinium chloride monohydrate obtained using a multipolar model and a spherical charges model.

Table S1. Wave function coefficients and orbital exponents for the virtual atom described as a combination of Slater functions.

$$\text{Slater}(N_l, \xi, r) = \text{coeff} * r^{N_l} * \exp(-\xi r)$$

ATOM	ξ	Coeff.
0	6.26445	0.07360
1	6.06204	0.13954
2	3.73854	0.87683

Figure S1. Analysis of experimental vs. expected $\Delta\rho$.

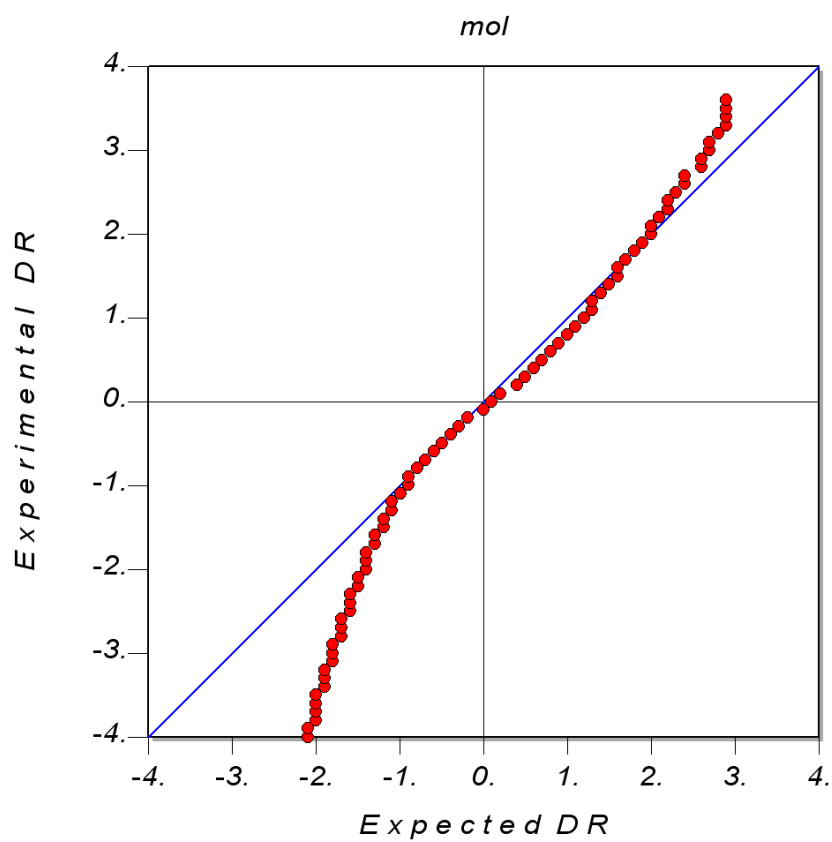


Figure S2. Analysis of $\langle F_o \rangle / \langle F_c \rangle$ as a function of reciprocal resolution.

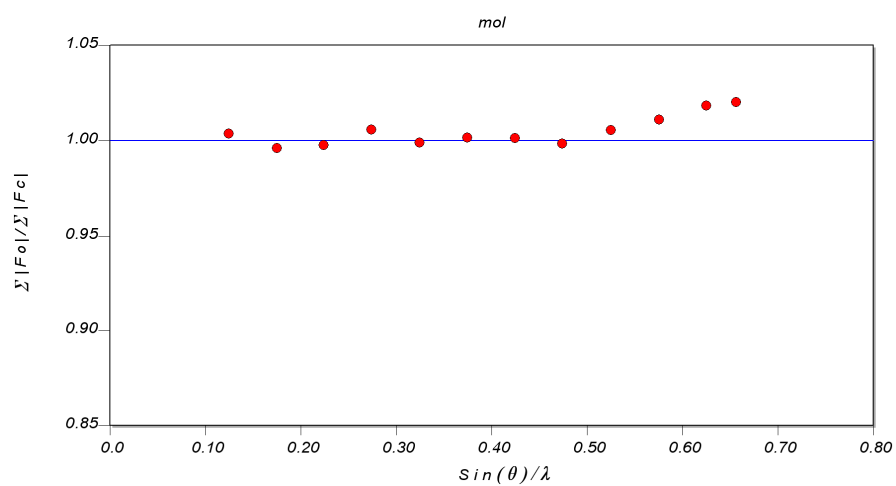
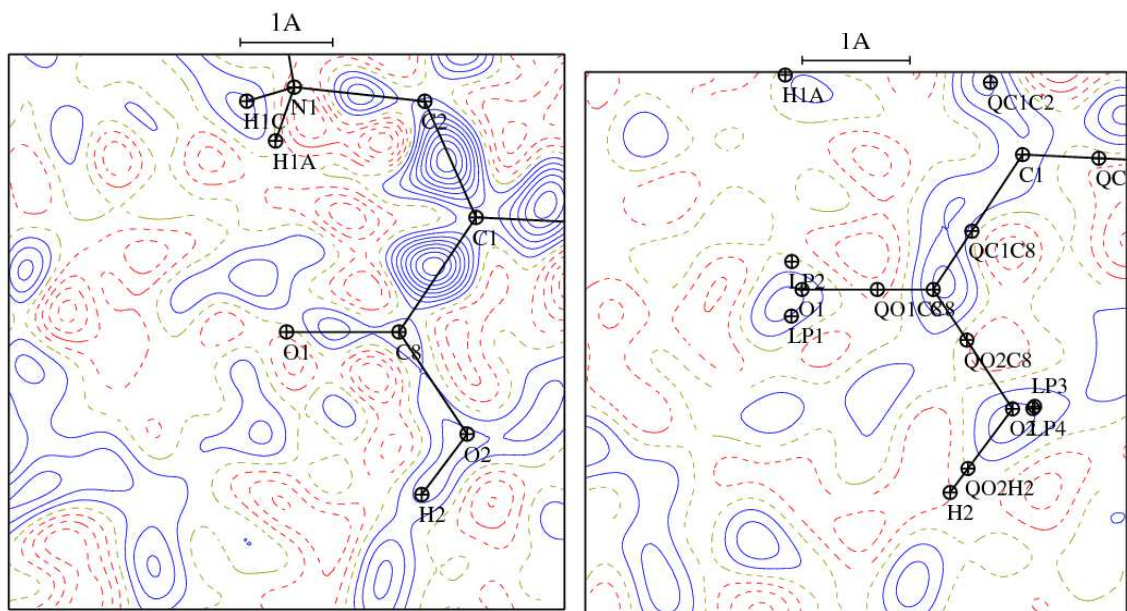


Figure S3.

Experimental residual electron density in the lone pairs planes.

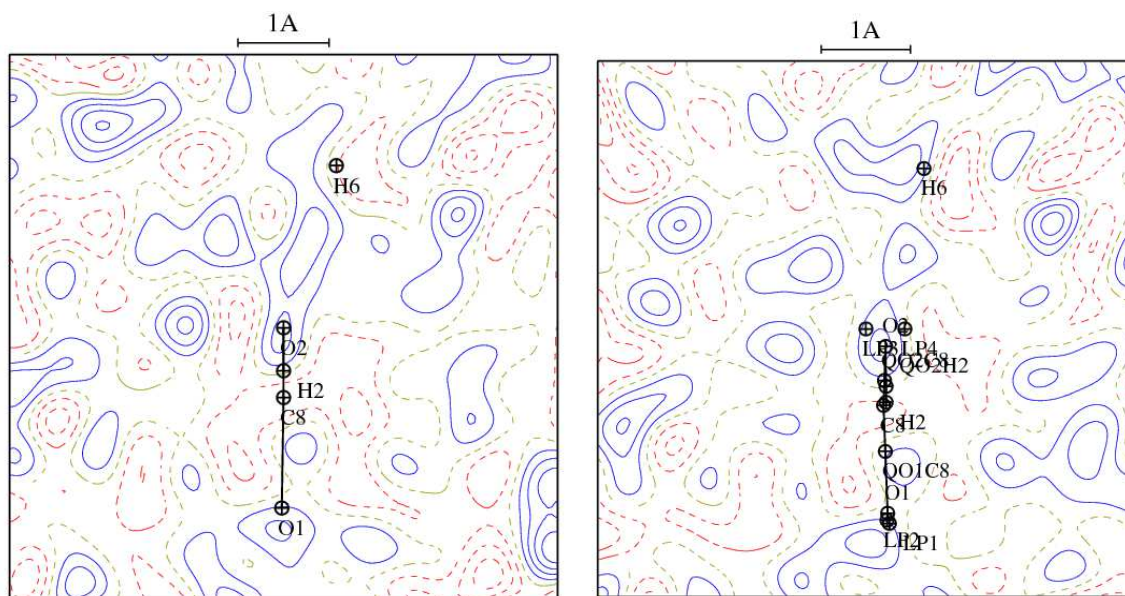
Contour level: $\pm 0.05 \text{ e}/\text{\AA}^3$. Blue solid lines and red dashed lines denote positive and negative contours, respectively. The zero contours is shown as yellow line.



(a) EXP_IAM

(b) EXP_VIR

in the C8O1O2 plane



(c) EXP_IAM

(d) EXP_VIR

in the oxygen atom O2 lone pairs plane.

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#-----#
# ELMAM2 MODEL  CHARGE DENSITY PARAMETERS  #
#-----#
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N1 C2 Z N1 H1A X
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C5 C4 bX C5 C6 Y
C6 C5 bX C6 C1 Y
C7 C5 Z C7 H7B X
C8 O2 bX C8 O1 Y
H2 O2 Z H2 C8 X
H3 C3 Z H3 C2 X
H4 C4 Z H4 C5 X
H6 C6 Z H6 C5 X
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H7C C7 Z H7C C5 X
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THEO_MUL MODEL: CHARGE DENSITY PARAMETERS #
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H7B 1.000 0. 0. 0. 0.135
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0.0010(64)
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0.983(2) 0.98(3) 2 4.466 2 4.466 2 4.466 3 4.466 4 4.466

H1W 0.86(1) 0. 0. 0. 0.14(1)
0.08(1) 0. 0. 0. 0.
0. 0.
0. 0. 0. 0.
1.160(2) 1.48(7) 1 2.000 1 2.000 2 2.000 2 2.000 2 2.000
H2W 0.85(1) 0. 0. 0. 0.14(1)
0.08(1) 0. 0. 0. 0.
0. 0.
0. 0. 0. 0.
1.160(2) 1.46(6) 1 2.000 1 2.000 2 2.000 2 2.000 2 2.000

loop_
_atom_local_axes_atom_label
_atom_local_axes_atom0
_atom_local_axes_ax1
_atom_local_axes_atom1
_atom_local_axes_atom2
_atom_local_axes_ax2

CL1 OW bZ CL1 N1 X
O1 C8 X O1 O2 Y
O2 C8 bX O2 H2 Y
N1 C2 Z N1 H1A X
C1 C2 bX C1 C6 Y
C2 C1 bX C2 C3 Y
C3 C2 bX C3 C4 Y
C4 C5 bX C4 C3 Y
C5 C4 bX C5 C6 Y
C6 C5 bX C6 C1 Y
C7 C5 Z C7 H7B X
C8 O2 bX C8 O1 Y
H2 O2 Z H2 C8 X
H3 C3 Z H3 C2 X
H4 C4 Z H4 C5 X
H6 C6 Z H6 C5 X
H1A N1 Z H1A C2 X
H1B N1 Z H1B C2 X
H1C N1 Z H1C C2 X
H7A C7 Z H7A C5 X
H7B C7 Z H7B C5 X
H7C C7 Z H7C C5 X
OW H1W bX OW H2W Y
H1W OW Z H1W H2W X
H2W OW Z H2W H1W X

###-THEO_VIR_MODEL CHARGE DENSITY PARAMETERS#####

loop_
_atom_site_type_symbol
_atom_site_label
_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_calc_flag
_atom_site_refinement_flags_posn
_atom_site_refinement_flags_adp
_atom_site_disorder_assembly
_atom_site_disorder_group

Cl	CL1	0.16000	0.49786	0.67712	0.	Uiso	1	d
O	O1	0.17413	0.46924	0.30185	0.	Uiso	1	d
O	O2	0.37352	0.52449	0.36549	0.	Uiso	1	d
N	N1	0.03425	0.01228	0.32497	0.	Uiso	1	d
C	C1	0.22773	0.20083	0.40329	0.	Uiso	1	d
C	C2	0.12204	0.01647	0.39089	0.	Uiso	1	d
C	C3	0.09716	-0.16926	0.44079	0.	Uiso	1	d
C	C4	0.17781	-0.17376	0.50395	0.	Uiso	1	d
C	C5	0.28423	0.00540	0.51798	0.	Uiso	1	d
C	C6	0.30797	0.19008	0.46727	0.	Uiso	1	d
C	C7	0.37128	0.00175	0.58712	0.	Uiso	1	d
C	C8	0.25417	0.40907	0.35124	0.	Uiso	1	d
H	H2	0.38760	0.65111	0.32947	0.	Uiso	1	d	D	U	.	.
H	H3	0.01517	-0.31080	0.43076	0.	Uiso	1	d	D	U	.	.
H	H4	0.15733	-0.31896	0.54269	0.	Uiso	1	d	D	U	.	.
H	H6	0.39072	0.32982	0.47734	0.	Uiso	1	d	D	U	.	.
H	H1A	0.09223	0.01198	0.28473	0.	Uiso	1	d	D	U	.	.
H	H1B	-0.02454	-0.15896	0.32245	0.	Uiso	1	d	D	U	.	.
H	H1C	-0.02642	0.18091	0.32153	0.	Uiso	1	d	D	U	.	.
H	H7A	0.31820	0.09034	0.62504	0.	Uiso	1	d	D	U	.	.
H	H7B	0.39595	-0.19986	0.60110	0.	Uiso	1	d	D	U	.	.
H	H7C	0.46024	0.11225	0.58333	0.	Uiso	1	d	D	U	.	.
O	OW	0.42920	0.67298	0.76040	0.	Uiso	1	d
H	H1W	0.48164	0.50919	0.76268	0.	Uiso	1	d	D	U	.	.
H	H2W	0.34648	0.62584	0.73260	0.	Uiso	1	d	D	U	.	.
Q	QO1C8	0.22014	0.43465	0.33024	0.	Uiso	1	d
Q	QO2C8	0.30466	0.45790	0.35727	0.	Uiso	1	d
Q	QO2H2	0.38365	0.61552	0.33959	0.	Uiso	1	d
Q	QN1C2	0.07290	0.01413	0.35399	0.	Uiso	1	d
Q	Q5	-0.00138	-0.09151	0.32344	0.	Uiso	1	d
Q	Q6	-0.00197	0.11295	0.32291	0.	Uiso	1	d
Q	Q7	0.06982	0.01210	0.30028	0.	Uiso	1	d
Q	QC1C8	0.24276	0.31919	0.37371	0.	Uiso	1	d
Q	QC1C2	0.17167	0.10304	0.39671	0.	Uiso	1	d
Q	QC1C6	0.26934	0.19526	0.43647	0.	Uiso	1	d
Q	QC2C3	0.11008	-0.07284	0.41488	0.	Uiso	1	d
Q	QC3C4	0.13791	-0.17154	0.47270	0.	Uiso	1	d
Q	QC3H3	0.04479	-0.25967	0.43438	0.	Uiso	1	d
Q	QC4C5	0.23062	-0.08486	0.51091	0.	Uiso	1	d
Q	QC4H4	0.16510	-0.26384	0.52798	0.	Uiso	1	d
Q	QC5C6	0.29630	0.09926	0.49221	0.	Uiso	1	d
Q	QC5C7	0.32689	0.00361	0.55186	0.	Uiso	1	d
Q	QC6H6	0.35976	0.27753	0.47357	0.	Uiso	1	d
Q	Q19	0.42842	0.07272	0.58469	0.	Uiso	1	d
Q	Q20	0.38680	-0.12507	0.59591	0.	Uiso	1	d
Q	Q21	0.33811	0.05712	0.61082	0.	Uiso	1	d
Q	Q22	0.47033	0.54453	0.76218	0.	Uiso	1	d
Q	Q23	0.36482	0.63629	0.73877	0.	Uiso	1	d
Q	LP1	0.18338	0.50772	0.29455	0.	Uiso	1	d
Q	LP2	0.15143	0.43792	0.30186	0.	Uiso	1	d
Q	LP3	0.39373	0.48213	0.36776	0.	Uiso	1	d
Q	LP4	0.37644	0.54653	0.37915	0.	Uiso	1	d
Q	LP5	0.44205	0.70647	0.75197	0.	Uiso	1	d
Q	LP6	0.42135	0.67782	0.77349	0.	Uiso	1	d

loop_
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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
N1 H1A LP1 1.033 2.664 3.014 99.6 1_545
N1 H1B LP2 1.033 2.653 2.947 96.0 2_545
N1 H1A CL1 1.033 2.288 3.291 163.1 4_554
C6 H6 O2 1.083 2.379 2.732 97.0 1_555
C6 H6 LP3 1.083 2.277 2.651 97.8 1_555
C6 H6 LP4 1.083 2.191 2.618 100.7 1_555
N1 H1A O1 1.033 2.424 2.734 96.1 1_555
N1 H1A QO1C8 1.033 2.564 2.802 92.2 1_555
N1 H1A LP1 1.033 2.624 2.976 99.6 1_555
N1 H1A LP2 1.033 2.211 2.486 92.8 1_555
N1 H1B QC3H3 1.033 2.274 2.522 91.5 1_555
OW H2W CL1 0.967 2.148 3.113 175.4 1_555
N1 H1B CL1 1.033 2.165 3.195 174.7 3_556
O2 H2 OW 0.967 1.702 2.658 169.4 4_564
O2 H2 Q22 0.967 2.228 3.173 165.5 4_564
O2 H2 Q23 0.967 2.054 2.989 162.3 4_564
O2 H2 LP5 0.967 1.815 2.748 161.2 4_564
O2 H2 LP6 0.967 1.457 2.420 173.5 4_564
N1 H1C CL1 1.033 2.086 3.115 173.7 3_566
OW H1W OW 0.967 1.971 2.913 163.9 2_646
OW H1W Q22 0.967 2.417 3.324 156.0 2_646
OW H1W Q23 0.967 2.412 3.372 172.1 2_646
OW H1W LP5 0.967 1.726 2.677 167.0 2_646
OW H1W LP6 0.967 2.078 2.995 157.6 2_646
C7 H7B LP3 1.059 2.549 3.411 138.0 3_656
O2 H2 Q22 0.967 2.608 3.121 113.5 3_666
C7 H7C O2 1.059 2.579 3.522 148.1 3_666
C7 H7C QO2H2 1.059 2.450 3.301 136.7 3_666
C7 H7C LP3 1.059 2.606 3.525 144.9 3_666
C7 H7C LP4 1.059 2.412 3.391 153.2 3_666
OW H1W QO2H2 0.967 2.619 3.207 119.5 3_666

#-----#
MULTIPOLE PARAMETERS #
#-----#

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_atom_rho_multipole_coeff_P10
_atom_rho_multipole_coeff_P20
_atom_rho_multipole_coeff_P21
_atom_rho_multipole_coeff_P2-1
_atom_rho_multipole_coeff_P22
_atom_rho_multipole_coeff_P2-2
_atom_rho_multipole_coeff_P30
_atom_rho_multipole_coeff_P31
_atom_rho_multipole_coeff_P3-1
_atom_rho_multipole_coeff_P32
_atom_rho_multipole_coeff_P3-2
_atom_rho_multipole_coeff_P33
_atom_rho_multipole_coeff_P3-3
_atom_rho_multipole_coeff_P40

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_atom_rho_multipole_coeff_P44
_atom_rho_multipole_coeff_P4-4
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_atom_rho_multipole_radial_slater_n2
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_atom_rho_multipole_radial_slater_zeta4
CL1 4.60(4) 0. 0. 0. 0.
0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0. 0. 0.
1.032(3) 0. 4 4.259 4 4.259 4 4.259 6 4.259 8 4.259
O1 5.7(1) 0. 0. 0. 0.
0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0. 0.
0.962(4) 0. 2 4.466 2 4.466 2 4.466 3 4.466 4 4.466
O2 6.3(1) 0. 0. 0. 0.
0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0. 0.
0.976(4) 0. 2 4.466 2 4.466 2 4.466 3 4.466 4 4.466
N1 4.1(1) 0. 0. 0. 0.
0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0. 0.
1.058(7) 0. 2 3.839 2 3.839 2 3.839 3 3.839 4 3.839
C1 2.98(9) 0. 0. 0. 0.
0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0. 0.
1.063(7) 0. 2 3.176 2 3.176 2 3.176 3 3.176 4 3.176
C2 2.92(10) 0. 0. 0. 0.
0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0. 0.
1.059(8) 0. 2 3.176 2 3.176 2 3.176 3 3.176 4 3.176
C3 2.7(1) 0. 0. 0. 0.
0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0. 0.
1.054(8) 0. 2 3.176 2 3.176 2 3.176 3 3.176 4 3.176
C4 2.69(8) 0. 0. 0. 0.
0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0. 0.
1.063(8) 0. 2 3.176 2 3.176 2 3.176 3 3.176 4 3.176
C5 2.9(1) 0. 0. 0. 0.

```

0. 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0. 0. 0. 0. 0.
1.064(9) 0. 2 3.176 2 3.176 2 3.176 3 3.176 4 3.176
C6 2.78(9) 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0. 0. 0. 0. 0.
1.051(8) 0. 2 3.176 2 3.176 2 3.176 3 3.176 4 3.176
C7 2.4(1) 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0. 0. 0. 0. 0.
1.08(1) 0. 2 3.176 2 3.176 2 3.176 3 3.176 4 3.176
C8 2.5(1) 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0. 0. 0. 0. 0.
1.046(9) 0. 2 3.176 2 3.176 2 3.176 3 3.176 4 3.176
H2 0.71(8) 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0. 0. 0. 0. 0.
1.02(8) 0. 1 2.000 1 2.000 2 2.000 2 2.000 2 2.000
H3 0.80(8) 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0. 0. 0. 0. 0.
1.14(5) 0. 1 2.000 1 2.000 2 2.000 2 2.000 2 2.000
H4 0.78(6) 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0. 0. 0. 0. 0.
1.17(3) 0. 1 2.000 1 2.000 2 2.000 2 2.000 2 2.000
H6 0.779 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0. 0. 0. 0. 0.
1.17(3) 0. 1 2.000 1 2.000 2 2.000 2 2.000 2 2.000
H1A 1.16(5) 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0. 0. 0. 0. 0.
0.77(4) 0. 1 2.000 1 2.000 2 2.000 2 2.000 2 2.000
H1B 1.162 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0. 0. 0. 0. 0.
0.77(4) 0. 1 2.000 1 2.000 2 2.000 2 2.000 2 2.000
H1C 1.162 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0. 0. 0. 0. 0.
0.77(4) 0. 1 2.000 1 2.000 2 2.000 2 2.000 2 2.000
H7A 0.73(5) 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0. 0. 0. 0. 0.
1.19(3) 0. 1 2.000 1 2.000 2 2.000 2 2.000 2 2.000
H7B 0.734 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0.

0. 0. 0. 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0. 0. 0. 0.
1.19(3) 0. 1 2.000 1 2.000 2 2.000 2 2.000 2 2.000
H7C 0.734 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0. 0. 0. 0.
1.19(3) 0. 1 2.000 1 2.000 2 2.000 2 2.000 2 2.000
OW 6.1(1) 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0. 0. 0. 0.
0.958(4) 0. 2 4.466 2 4.466 2 4.466 3 4.466 4 4.466
H1W 1.09(7) 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0. 0. 0. 0.
0.57(6) 0. 1 2.000 1 2.000 2 2.000 2 2.000 2 2.000
H2W 1.089 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0. 0. 0. 0.
0.57(6) 0. 1 2.000 1 2.000 2 2.000 2 2.000 2 2.000
QO1C8 0.38(9) 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0. 0. 0. 0.
1.26(8) 0. 1 2.000 1 2.000 2 2.000 2 2.000 2 2.000
QO2C8 -0.029(64) 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0. 0. 0. 0.
1.8(1) 0. 1 2.000 1 2.000 2 2.000 2 2.000 2 2.000
QO2H2 -0.085(87) 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0. 0. 0. 0.
2.0(2) 0. 1 2.000 1 2.000 2 2.000 2 2.000 2 2.000
QN1C2 0.22(7) 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0. 0. 0. 0.
1.30(10) 0. 1 2.000 1 2.000 2 2.000 2 2.000 2 2.000
Q5 0.28(7) 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0. 0. 0. 0.
1.26(5) 0. 1 2.000 1 2.000 2 2.000 2 2.000 2 2.000
Q6 0.284 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0. 0. 0. 0.
1.26(5) 0. 1 2.000 1 2.000 2 2.000 2 2.000 2 2.000
Q7 0.284 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0. 0. 0. 0.
1.26(5) 0. 1 2.000 1 2.000 2 2.000 2 2.000 2 2.000
QC1C8 1.05(9) 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0. 0. 0.

0. 0. 0. 0. 0. 0. 0. 0. 0. 0.
0.83(5) 0. 1 2.000 1 2.000 2 2.000 2 2.000 2 2.000
QC1C2 1.00(9) 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0. 0. 0. 0. 0.
0.83(5) 0. 1 2.000 1 2.000 2 2.000 2 2.000 2 2.000
QC1C6 1.06(9) 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0. 0. 0. 0. 0.
0.84(5) 0. 1 2.000 1 2.000 2 2.000 2 2.000 2 2.000
QC2C3 0.84(10) 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0. 0. 0. 0. 0.
0.97(5) 0. 1 2.000 1 2.000 2 2.000 2 2.000 2 2.000
QC3C4 1.12(10) 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0. 0. 0. 0. 0.
0.89(4) 0. 1 2.000 1 2.000 2 2.000 2 2.000 2 2.000
QC3H3 0.9(1) 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0. 0. 0. 0. 0.
0.89(7) 0. 1 2.000 1 2.000 2 2.000 2 2.000 2 2.000
QC4C5 0.70(8) 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0. 0. 0. 0. 0.
1.03(3) 0. 1 2.000 1 2.000 2 2.000 2 2.000 2 2.000
QC4H4 0.71(8) 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0. 0. 0. 0. 0.
0.98(5) 0. 1 2.000 1 2.000 2 2.000 2 2.000 2 2.000
QC5C6 0.698 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0. 0. 0. 0. 0.
1.03(3) 0. 1 2.000 1 2.000 2 2.000 2 2.000 2 2.000
QC5C7 0.15(10) 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0. 0. 0. 0. 0.
1.27(6) 0. 1 2.000 1 2.000 2 2.000 2 2.000 2 2.000
QC6H6 0.708 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0. 0. 0. 0. 0.
0.98(5) 0. 1 2.000 1 2.000 2 2.000 2 2.000 2 2.000
Q19 1.01(8) 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0. 0. 0. 0. 0.
0.81(3) 0. 1 2.000 1 2.000 2 2.000 2 2.000 2 2.000
Q20 1.008 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0. 0. 0.
0. 0. 0. 0. 0. 0. 0. 0. 0. 0.

0.81(3) 0. 1 2.000 1 2.000 2 2.000 2 2.000 2 2.000
 Q21 1.008 0. 0. 0. 0.
 0. 0. 0. 0. 0.
 0. 0. 0. 0. 0. 0. 0.
 0. 0. 0. 0. 0. 0. 0. 0. 0.
 0.81(3) 0. 1 2.000 1 2.000 2 2.000 2 2.000 2 2.000
 Q22 0.012(70) 0. 0. 0. 0.
 0. 0. 0. 0. 0.
 0. 0. 0. 0. 0. 0. 0.
 0. 0. 0. 0. 0. 0. 0. 0. 0.
 2.0(1) 0. 1 2.000 1 2.000 2 2.000 2 2.000 2 2.000
 Q23 0.012 0. 0. 0. 0.
 0. 0. 0. 0. 0.
 0. 0. 0. 0. 0. 0. 0.
 0. 0. 0. 0. 0. 0. 0. 0. 0.
 2.0(1) 0. 1 2.000 1 2.000 2 2.000 2 2.000 2 2.000
 LP1 0.25(5) 0. 0. 0. 0.
 0. 0. 0. 0. 0.
 0. 0. 0. 0. 0. 0. 0.
 0. 0. 0. 0. 0. 0. 0. 0. 0.
 1.7(1) 0. 1 2.000 1 2.000 2 2.000 2 2.000 2 2.000
 LP2 0.255 0. 0. 0. 0.
 0. 0. 0. 0. 0.
 0. 0. 0. 0. 0. 0. 0.
 0. 0. 0. 0. 0. 0. 0. 0. 0.
 1.7(1) 0. 1 2.000 1 2.000 2 2.000 2 2.000 2 2.000
 LP3 0.06(3) 0. 0. 0. 0.
 0. 0. 0. 0. 0.
 0. 0. 0. 0. 0. 0. 0.
 0. 0. 0. 0. 0. 0. 0. 0. 0.
 2.3(2) 0. 1 2.000 1 2.000 2 2.000 2 2.000 2 2.000
 LP4 0.061 0. 0. 0. 0.
 0. 0. 0. 0. 0.
 0. 0. 0. 0. 0. 0. 0.
 0. 0. 0. 0. 0. 0. 0. 0. 0.
 2.3(2) 0. 1 2.000 1 2.000 2 2.000 2 2.000 2 2.000
 LP5 0.19(4) 0. 0. 0. 0.
 0. 0. 0. 0. 0.
 0. 0. 0. 0. 0. 0. 0.
 0. 0. 0. 0. 0. 0. 0. 0. 0.
 1.8(2) 0. 1 2.000 1 2.000 2 2.000 2 2.000 2 2.000
 LP6 0.195 0. 0. 0. 0.
 0. 0. 0. 0. 0.
 0. 0. 0. 0. 0. 0. 0.
 0. 0. 0. 0. 0. 0. 0. 0. 0.
 1.8(2) 0. 1 2.000 1 2.000 2 2.000 2 2.000 2 2.000

loop_
 _atom_local_axes_atom_label
 _atom_local_axes_atom0
 _atom_local_axes_ax1
 _atom_local_axes_atom1
 _atom_local_axes_atom2
 _atom_local_axes_ax2
 CL1 OW bZ CL1 N1 X
 O1 C8 X O1 O2 Y
 O2 C8 bX O2 H2 Y
 N1 C2 Z N1 H1A X
 C1 C2 bX C1 C6 Y
 C2 C1 bX C2 C3 Y
 C3 C2 bX C3 C4 Y

C4 C5 bX C4 C3 Y
C5 C4 bX C5 C6 Y
C6 C5 bX C6 C1 Y
C7 C5 Z C7 H7B X
C8 O2 bX C8 O1 Y
H2 O2 Z H2 C8 X
H3 C3 Z H3 C2 X
H4 C4 Z H4 C5 X
H6 C6 Z H6 C5 X
H1A N1 Z H1A C2 X
H1B N1 Z H1B C2 X
H1C N1 Z H1C C2 X
H7A C7 Z H7A C5 X
H7B C7 Z H7B C5 X
H7C C7 Z H7C C5 X
OW H1W bX OW H2W Y
H1W OW Z H1W H2W X
H2W OW Z H2W H1W X
QO1C8 O1 X QO1C8 C8 Y
QO2C8 O2 X QO2C8 C8 Y
QO2H2 O2 X QO2H2 H2 Y
QN1C2 N1 X QN1C2 C2 Y
Q5 N1 X Q5 H1B Y
Q6 N1 X Q6 H1C Y
Q7 N1 X Q7 H1A Y
QC1C8 C1 X QC1C8 C8 Y
QC1C2 C1 X QC1C2 C2 Y
QC1C6 C1 X QC1C6 C6 Y
QC2C3 C2 X QC2C3 C3 Y
QC3C4 C3 X QC3C4 C4 Y
QC3H3 C3 X QC3H3 H3 Y
QC4C5 C4 X QC4C5 C5 Y
QC4H4 C4 X QC4H4 H4 Y
QC5C6 C5 X QC5C6 C6 Y
QC5C7 C5 X QC5C7 C7 Y
QC6H6 C6 X QC6H6 H6 Y
Q19 C7 X Q19 H7C Y
Q20 C7 X Q20 H7B Y
Q21 C7 X Q21 H7A Y
Q22 OW X Q22 H1W Y
Q23 OW X Q23 H2W Y
LP1 O1 X LP1 C8 Y
LP2 O1 X LP2 C8 Y
LP3 O2 X LP3 C8 Y
LP4 O2 X LP4 C8 Y
LP5 OW X LP5 H1W Y
LP6 OW X LP6 H1W Y