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A critical analysis of dipole moment calculations as obtained from experimental and theoretical structure factors.

Constrains/restraints imposed on the symmetry and simmilarity of the equivalent atoms:

a) symmetry:

	constrained
atom	symemtry
C1	mymz
C2	mymz
C3	mymz
C4	mymz
C5	mymz
C6	mymz
C7	mz
C8	mz
C9	mz
C71	3m
C91	су
N1	mymz
N2	mymz
N8	mymz
N91	cy
O81	mz
O82	mz

where m - mirror plane, 3m - threefold axe, cy - cyllindrical symmetry.

b) equivalent atoms for similarity constraints/restraints on Plm, Pval, κ , κ ':

C2 and C6 C3 and C5 O81 and O82 H2 and H6 H3 and H5 H71, H72 and H73

Model	Multipolar	Treatment of	Parameter	No	Restraints/	RF	Goof	Dipole moment	Residual
	expansion level	κ and κ ' of	refined in order	reflections/	Constraints	factor		value	density
	of non-hydrogen	hydrogen	to remove the	No		wR2F		(module charge	$[e/A^3]$ up to
	/ hydrogen	atoms	negative electron	parameters		factor		dipole/atomic	1.2Å ⁻¹
	atoms		density					dipole) [D]	
1m	HEX / QUA	refined	SCA	15074 /	-	0.548	0.064	14.85	0.287(15)/
				532		0.526		(13.82/2.29)	-0.260(15)
1n	HEX / QUA	1.16	SCA	15074 /	-	0.554	0.064	15.32	0.291(15)/
				532		0.531		(14.49/2.12)	-0.258(15)
10	OCT / DIP	refined	SCA	15074 /	-	0.629	0.076	12.92	0.287(18)/
				339		0.636		(12.94/1.45)	-0.248(18)
1p	OCT / DIP	1.16	SCA	15074 /	-	0.630	0.078	13.04	0.283(18)/
				339		0.646		(12.62/1.40)	-0.252(18)
1q	HEX / QUA	refined	SCA	15074 /	R _{free}	0.560	0.053	15.17	0.288(16)/
				532	restraints	0.542		(13.63/2.39)	-0.261(16)
1r	HEX / QUA	1.16	SCA	15074 /	R _{free}	0.565	0.060	15.64	0.292(16)/
				532	restraints	0.546		(14.29/2.22)	-0.258(16)
1s	OCT / DIP	refined	SCA	15074 /	R _{free}	0.638	0.068	13.50	0.287(18)/
				339	restraints	0.649		(12.96/1.32)	-0.248(18)
1t	OCT / DIP	1.16	SCA	15074 /	R _{free}	0.637	0.076	13.28	0.282(19)/
				339	restraints	0.660		(12.26/1.49)	-0.252(19)

Table S1. MM of theoretical data with scale factor refined to reduce the effect of different wave functions.



Figure S1. Example of residual Fourier density map for model 1c (*cf.* Table 1). a), b) without proper core scaling and c), d) after κ_{core} refinement; contours 0.05 e/Å³, $s_{max} = 1$ Å⁻¹, red negative, blue positive.











Figure S2. Static deformation and Fourier residual maps for models 1c and 1g; contours $\pm 0.05e/A^3$, reciprocal resolution up to $s = 1Å^{-1}$.



Figure S3. Dipole moment directions (grey – total, red – contribution from atomic dipoles, blue – contribution from atomic charges) for models 1a-11.





Figure S5. Electron density of a spherical virtual atom with $P_{vir}=1$ as a function of the distance *r* to the nucleus. For comparison, the same curve is shown for a hydrogen atom.

Table S2. Wave function coefficients and orbital exponents for the virtual atom described as a combination of Slater functions. Slater(N_1 , ξ ,r) = coeff * r^{N_1} * exp(- ξr)

Sidei (i i), i	5,,,) = 00011	, exp(g)
N_1	ξ	Coeff.
0	6.26445	0.07360
1	6.06204	0.13954
2	3.73854	0.87683

Table S3. κ , P_{val} , P_{10} , P_{11+} and P_{11-} parameters for models 1a-h.

	kp1							
	1a	1b	1c	1d	1e	1f	1g	1h
C1	0.990	0.990	0.989	0.989	0.997	0.997	0.996	0.997
C2	0.991	0.989	0.993	0.995	0.990	0.989	0.993	0.995
C3	0.988	0.987	0.991	0.993	0.985	0.984	0.987	0.990
C4	0.987	0.986	0.990	0.992	0.987	0.985	0.990	0.992
C5	0.988	0.987	0.991	0.993	0.986	0.985	0.988	0.991
C6	0.989	0.989	0.991	0.995	0.990	0.988	0.992	0.994
C7	0.987	0.987	0.993	0.993	0.985	0.985	0.992	0.993
C8	1.002	1.002	1.010	1.011	1.004	1.003	1.011	1.011
C9	0.993	0.993	1.003	1.003	0.997	0.997	1.007	1.007
C71	0.997	0.994	1.006	1.012	0.998	0.995	1.007	1.013
C91	0.982	0.981	0.988	0.988	0.982	0.981	0.988	0.989
N1	0.995	0.995	0.993	0.993	0.995	0.995	0.993	0.993
N2	0.987	0.987	0.991	0.991	0.988	0.988	0.991	0.991
N8	0.986	0.986	0.988	0.988	0.986	0.986	0.988	0.988
N91	0.991	0.991	0.994	0.994	0.992	0.991	0.994	0.994
O81	0.988	0.988	0.988	0.988	0.987	0.987	0.988	0.988
O82	0.988	0.988	0.988	0.988	0.987	0.987	0.988	0.988
H6	1.153	1.160	1.198	1.160	1.150	1.160	1.196	1.160
H2	1.149	1.160	1.192	1.160	1.149	1.160	1.195	1.160
H3	1.150	1.160	1.195	1.160	1.153	1.160	1.200	1.160
H4	1.148	1.160	1.192	1.160	1.150	1.160	1.192	1.160

H5	1.149	1.160	1.200	1.160	1.151	1.160	1.199	1.160
H73	1.149	1.160	1.184	1.160	1.151	1.160	1.187	1.160
H72	1.148	1.160	1.183	1.160	1.150	1.160	1.187	1.160
H71	1.160	1.160	1.196	1.160	1.151	1.160	1.187	1.160
	P _{val}							
	1a	1b	1c	1d	1e	1f	1g	1h
C1	4.05	4.04	4.08	4.09	3.98	3.97	3.97	3.98
C2	4.01	4.03	4.05	4.00	4.03	4.05	4.08	4.02
C3	4.04	4.04	4.11	4.06	4.05	4.05	4.16	4.11
C4	4.04	4.05	4.12	4.07	4.02	4.03	4.09	4.04
C5	4.02	4.03	4.12	4.07	4.05	4.05	4.16	4.11
C6	4.01	4.03	4.06	4.00	4.03	4.05	4.08	4.02
C7	4.21	4.21	4.16	4.16	4.26	4.26	4.17	4.17
C8	3.96	3.96	3.89	3.89	3.96	3.96	3.91	3.91
C9	3.96	3.97	3.98	3.97	3.93	3.93	3.92	3.92
C71	3.88	3.94	4.05	3.91	3.88	3.94	4.03	3.90
C91	4.52	4.52	4.35	4.35	4.53	4.53	4.36	4.35
N1	4.95	4.95	5.04	5.05	4.94	4.95	5.07	5.07
N2	4.96	4.97	4.97	4.97	4.94	4.95	4.97	4.97
N8	5.03	5.03	5.01	5.01	5.03	5.03	5.01	5.01
N91	4.69	4.70	4.82	4.82	4.67	4.68	4.82	4.83
O81	6.14	6.14	6.14	6.15	6.14	6.15	6.14	6.14
O82	6.13	6.14	6.14	6.14	6.14	6.15	6.14	6.14
H6	0.92	0.90	0.84	0.90	0.92	0.90	0.84	0.89
H2	0.92	0.90	0.85	0.89	0.92	0.90	0.84	0.89
H3	0.93	0.91	0.85	0.90	0.91	0.90	0.83	0.88
H4	0.94	0.92	0.86	0.90	0.94	0.92	0.86	0.90
H5	0.91	0.89	0.82	0.88	0.91	0.90	0.83	0.88
H73	0.95	0.93	0.91	0.95	0.94	0.92	0.90	0.95
H72	0.93	0.91	0.90	0.94	0.94	0.92	0.90	0.95
H71	0.91	0.91	0.88	0.94	0.94	0.92	0.90	0.95
dip x								
C1	0.121	0.117	0.171	0.173	0.104	0.102	0.146	0.148

C2	0.013	0.009	-0.030	-0.017	0.022	0.017	-0.024	-0.010
C3	0.021	0.014	-0.010	0.003	0.013	0.008	-0.013	0.003
C4	0.022	0.016	-0.010	0.003	0.020	0.013	-0.015	-0.002
C5	0.002	-0.005	-0.028	-0.012	0.013	0.007	-0.014	0.002
C6	0.026	0.023	-0.027	-0.013	0.023	0.017	-0.023	-0.009
C7	0.018	0.018	-0.010	-0.012	0.022	0.023	-0.006	-0.008
C8	-0.079	-0.077	-0.066	-0.065	-0.085	-0.084	-0.072	-0.071
C9	0.016	0.016	0.036	0.038	0.015	0.014	0.038	0.038
C71	0.018	0.013	0.017	0.013	0.001	0.001	0.002	0.002
C91	-0.003	-0.004	0.012	0.013	0.000	0.000	0.001	0.001
N1	0.003	0.003	0.014	0.016	-0.002	-0.002	0.015	0.015
N2	-0.138	-0.138	-0.126	-0.126	-0.140	-0.140	-0.125	-0.125
N8	-0.027	-0.027	-0.021	-0.021	-0.026	-0.027	-0.020	-0.021
N91	-0.001	-0.001	0.000	0.000	0.000	0.000	0.000	0.000
O81	-0.116	-0.116	-0.115	-0.115	-0.116	-0.116	-0.115	-0.115
O82	-0.116	-0.116	-0.116	-0.115	-0.116	-0.115	-0.115	-0.115
dip y								
C1	0.002	0.003	0.002	0.003	0.000	0.000	0.000	0.000
C2	0.030	0.028	0.032	0.032	0.001	0.001	0.001	0.001
C3	0.009	0.009	0.009	0.007	0.000	0.000	0.000	0.000
C4	-0.002	-0.001	-0.002	-0.001	0.000	0.000	0.000	0.000
C5	-0.015	-0.015	-0.010	-0.007	0.000	0.000	0.000	0.000
C6	0.031	0.028	0.033	0.031	0.001	0.001	0.001	0.001
C7	0.080	0.079	0.096	0.098	0.070	0.069	0.087	0.088
C8	0.164	0.161	0.162	0.162	0.156	0.154	0.159	0.159
C9	-0.077	-0.078	-0.088	-0.089	-0.078	-0.079	-0.080	-0.081
C71	-0.006	-0.006	-0.004	-0.004	0.000	0.000	0.000	0.000
C91	0.008	0.008	0.007	0.008	0.001	0.001	0.001	0.001
N1	-0.017	-0.017	-0.014	-0.013	-0.001	-0.001	-0.001	-0.001
N2	-0.004	-0.005	-0.002	-0.001	-0.001	-0.001	0.000	0.000
N8	0.005	0.005	0.000	-0.002	0.000	0.000	0.000	0.000
N91	0.002	0.002	0.003	0.002	0.001	0.001	0.001	0.001
O81	-0.002	-0.002	-0.005	-0.005	-0.002	-0.002	-0.005	-0.005
O82	-0.002	0.002	-0.005	-0.005	-0.002	-0.002	-0.005	-0.005

dip z								
C1	0.000	0.000	0.003	0.003				
C2	0.005	0.005	0.002	0.002				
C3	-0.005	-0.005	-0.002	-0.003				
C4	-0.001	-0.001	0.002	0.002				
C5	0.000	0.000	0.000	0.000				
C6	-0.001	-0.001	0.001	0.000				
C7	0.006	0.006	0.006	0.006				
C8	-0.002	-0.002	-0.001	-0.001				
C9	0.003	0.003	0.000	-0.001				
C71	-0.065	-0.068	-0.091	-0.082	-0.067	-0.071	-0.086	-0.078
C91	0.385	0.382	0.316	0.312	0.387	0.384	0.306	0.302
N1	-0.004	-0.004	-0.003	-0.002	-0.001	-0.001	0.000	0.000
N2	0.001	0.001	0.001	0.001	0.000	0.000	0.000	0.000
N8	-0.002	-0.002	-0.003	-0.003	0.000	0.000	0.000	0.000
N91	-0.133	-0.133	-0.120	-0.119	-0.133	-0.133	-0.119	-0.118
O81	0.000	0.000	0.001	0.001	0.000	0.000	0.000	0.000
O82	-0.001	-0.001	-0.001	-0.001	0.000	0.000	0.000	0.000
H6	0.168	0.167	0.130	0.154	0.167	0.169	0.131	0.158
H2	0.173	0.175	0.134	0.162	0.168	0.170	0.132	0.159
H3	0.163	0.165	0.125	0.151	0.167	0.170	0.125	0.154
H4	0.161	0.166	0.121	0.151	0.164	0.169	0.122	0.151
H5	0.166	0.171	0.127	0.157	0.168	0.171	0.126	0.155
H73	0.157	0.150	0.112	0.133	0.151	0.146	0.112	0.135
H72	0.149	0.140	0.106	0.128	0.150	0.146	0.111	0.134
H71	0.156	0.155	0.117	0.142	0.151	0.147	0.112	0.136

Model	Restraints/Constraints	No	RF	Goof	Dipole	Residual
		reflections/	factor		moment	density [e/A ³]
		No	wR2F		value [D]	up to 1.2\AA^{-1}
		parameters	factor			
2j	Default restraints*	15074 /	0.986	0.134	17.30	0.605(33)/
		213	1.123			-0.688(33)
2k	Modified rest**	15074 /	0.840	0.118	17.33	0.329(29)/
		213	0.987			-0.644(29)
21	No restraints	15074 /	0.819	0.117	18.70	0.297(29)/
		213	0.977			-0.578(29)
2m	Modified rest κ_{HYD} =	15074 /	0.863	0.122	11.30	0.322(30)/
	1.1	205	1.018			-0.646(30)
2n	Modified rest κ_{HYD} =	15074 /	0.853	0.120	12.84	0.324(30)/
	1.13	205	1.004			-0.644(30)
20	Modified rest	15074 /	0.847	0.119	14.31	0.326(29)/
	$\kappa_{\rm HYD} = 1.16$	205	0.995			-0.644(29)

Table S4 Results of the VIRAT_{theo} refinement (no additional atoms on N=O bonds)

***Default restraints:** Linearity 0.001; Distances: H-Q 0.37 0.01, LP 0.28 0.01; Distance similarities: LP 0.01; Angles LP 0.1; Planarity (LP) 0.001.

****Modified restraints:** Linearity 0.1; Distances: H-Q 0.37 0.01, LP 0.28 0.01; Distance similarities: LP 0.01; Planarity (LP) 0.001













Figure S6. Static deformation and Fourier residual density maps for VIRTAT 3j, 3k, 3l, contour 0.05e/A, resolution 1.2A⁻¹.



Figure S7. Dipole moment directions in sub-models 2.

Table S4. κ and P_{val} for models 2a-o.

	kp1														
	2a	2b	2c	2d	2e	2f	2g	2h	2i	2j	2k	21	2m	2n	20
C1	1.0387	1.0382	1.0377	1.0387	1.0349	1.0344	1.0386	1.0349	1.0346	1.0373	1.0379	1.0381	1.0396	1.0390	1.0385
C2	1.0312	1.0290	1.0268	1.0311	1.0268	1.0246	1.0302	1.0263	1.0240	1.0195	1.0236	1.0241	1.0331	1.0308	1.0287
C3	1.0294	1.0275	1.0257	1.0294	1.0252	1.0233	1.0293	1.0255	1.0237	1.0219	1.0229	1.0221	1.0300	1.0281	1.0264
C4	1.0257	1.0236	1.0215	1.0257	1.0232	1.0212	1.0256	1.0232	1.0211	1.0163	1.0171	1.0172	1.0256	1.0236	1.0215
C5	1.0293	1.0273	1.0254	1.0294	1.0259	1.0240	1.0293	1.0255	1.0237	1.0203	1.0234	1.0212	1.0309	1.0288	1.0270
C6	1.0291	1.0267	1.0246	1.0293	1.0258	1.0236	1.0302	1.0263	1.0240	1.0197	1.0214	1.0216	1.0303	1.0280	1.0258
C7	1.0528	1.0516	1.0506	1.0529	1.0509	1.0500	1.0528	1.0510	1.0503	1.0504	1.0487	1.0462	1.0526	1.0513	1.0502
C8	1.0530	1.0520	1.0512	1.0530	1.0556	1.0548	1.0532	1.0555	1.0546	1.0566	1.0519	1.0508	1.0556	1.0547	1.0538
C9	1.0434	1.0430	1.0426	1.0434	1.0430	1.0426	1.0433	1.0430	1.0428	1.0374	1.0424	1.0426	1.0437	1.0434	1.0431
C71	1.0599	1.0520	1.0450	1.0600	1.0536	1.0466	1.0602	1.0536	1.0465	1.0396	1.0380	1.0392	1.0612	1.0535	1.0472
C91	1.0072	1.0061	1.0051	1.0070	1.0055	1.0045	1.0071	1.0056	1.0045	0.9983	0.9976	0.9973	1.0013	1.0003	0.9993
N1	1.0117	1.0116	1.0116	1.0118	1.0108	1.0107	1.0118	1.0108	1.0108	1.0101	1.0110	1.0114	1.0114	1.0113	1.0112
N2	0.9549	0.9543	0.9536	0.9549	0.9554	0.9546	0.9551	0.9553	0.9547	0.9527	0.9523	0.9624	0.9549	0.9543	0.9537
N8	1.0436	1.0431	1.0424	1.0434	1.0444	1.0437	1.0424	1.0452	1.0445	1.0280	1.0293	1.0337	1.0302	1.0301	1.0296
N91	0.9382	0.9373	0.9364	0.9381	0.9364	0.9355	0.9381	0.9364	0.9354	0.9315	0.9332	0.9388	0.9368	0.9358	0.9350
O81	0.9665	0.9653	0.9652	0.9664	0.9660	0.9653	0.9670	0.9660	0.9655	0.9520	0.9408	0.9252	0.9439	0.9428	0.9423
O82	0.9660	0.9657	0.9652	0.9663	0.9669	0.9667	0.9670	0.9660	0.9655	0.9518	0.9432	0.9287	0.9458	0.9453	0.9449
H6	1.1000	1.1300	1.1600	1.1000	1.1300	1.1600	1.1000	1.1300	1.1600	1.2311	1.2335	1.2338	1.1000	1.1300	1.1600
H2	1.1000	1.1300	1.1600	1.1000	1.1300	1.1600	1.1000	1.1300	1.1600	1.2470	1.2501	1.2559	1.1000	1.1300	1.1600
H3	1.1000	1.1300	1.1600	1.1000	1.1300	1.1600	1.1000	1.1300	1.1600	1.2361	1.2266	1.2277	1.1000	1.1300	1.1600
H4	1.1000	1.1300	1.1600	1.1000	1.1300	1.1600	1.1000	1.1300	1.1600	1.2461	1.2433	1.2454	1.1000	1.1300	1.1600
H5	1.1000	1.1300	1.1600	1.1000	1.1300	1.1600	1.1000	1.1300	1.1600	1.2394	1.2271	1.2311	1.1000	1.1300	1.1600
H73	1.1000	1.1300	1.1600	1.1000	1.1300	1.1600	1.1000	1.1300	1.1600	1.2186	1.2213	1.2320	1.1000	1.1300	1.1600
H72	1.1000	1.1300	1.1600	1.1000	1.1300	1.1600	1.1000	1.1300	1.1600	1.2071	1.1930	1.2013	1.1000	1.1300	1.1600
H71	1.1000	1.1300	1.1600	1.1000	1.1300	1.1600	1.1000	1.1300	1.1600	1.2042	1.1953	1.1960	1.1000	1.1300	1.1600
QC1N1	1.1877	1.1900	1.1919	1.1877	1.2061	1.2088	1.1877	1.2061	1.2084	1.2531	1.2002	1.1971	1.1903	1.1925	1.1953
QC1C6	0.8202	0.8239	0.8269	0.8276	0.8353	0.8385	0.8280	0.8357	0.8389	0.8537	0.8304	0.8281	0.8189	0.8225	0.8259
QC1C2	0.8452	0.8487	0.8523	0.8368	0.8402	0.8435	0.8365	0.8401	0.8434	0.8629	0.8533	0.8527	0.8410	0.8450	0.8482
QC2C3	0.8347	0.8404	0.8454	0.8248	0.8389	0.8439	0.8247	0.8392	0.8441	0.8657	0.8564	0.8543	0.8371	0.8431	0.8488
QC2H2	0.9182	0.9123	0.9053	0.9153	0.9169	0.9093	0.9151	0.9169	0.9094	0.8713	0.8760	0.8750	0.9152	0.9098	0.9031

QC3C4	0.8225	0.8280	0.8324	0.8207	0.8338	0.8385	0.8207	0.8341	0.8388	0.8471	0.8380	0.8354	0.8207	0.8258	0.8313
QC3H3	0.9208	0.9147	0.9070	0.9133	0.9049	0.8975	0.9133	0.9051	0.8976	0.8764	0.8883	0.8886	0.9233	0.9175	0.9095
QC4C5	0.8141	0.8194	0.8241	0.8159	0.8323	0.8365	0.8161	0.8324	0.8365	0.8509	0.8287	0.8246	0.8118	0.8172	0.8218
QC4H4	0.9134	0.9075	0.9004	0.9134	0.9135	0.9066	0.9133	0.9135	0.9067	0.8717	0.8711	0.8715	0.9083	0.9025	0.8958
QC5C6	0.8134	0.8189	0.8236	0.8228	0.8320	0.8367	0.8232	0.8324	0.8370	0.8566	0.8347	0.8302	0.8169	0.8224	0.8277
QC5H5	0.9160	0.9107	0.9041	0.9232	0.9180	0.9111	0.9233	0.9182	0.9112	0.8817	0.8860	0.8869	0.9152	0.9101	0.9039
QC6H6	0.8987	0.8923	0.8845	0.9012	0.9056	0.8982	0.9014	0.9059	0.8985	0.8681	0.8616	0.8628	0.9002	0.8940	0.8852
QC7N1	0.9882	0.9900	0.9918	0.9880	0.9942	0.9965	0.9877	0.9942	0.9963	1.0141	0.9974	0.9957	0.9883	0.9901	0.9935
QC7N2	0.9596	0.9626	0.9655	0.9593	0.9628	0.9665	0.9586	0.9632	0.9659	0.9702	0.9652	0.9693	0.9544	0.9577	0.9599
Q15	0.8078	0.8154	0.8214	0.8077	0.8231	0.8299	0.8076	0.8231	0.8298	0.8226	0.8254	0.8267	0.8055	0.8127	0.8178
QC8N8	1.0415	1.0425	1.0436	1.0419	1.0276	1.0285	1.0431	1.0266	1.0275	1.0269	1.0254	1.0185	1.0236	1.0243	1.0249
QC8N2	1.0082	1.0115	1.0142	1.0080	1.0144	1.0163	1.0090	1.0138	1.0176	1.0307	1.0175	1.0245	1.0044	1.0075	1.0114
QC8C9	0.8671	0.8683	0.8697	0.8671	0.8697	0.8714	0.8670	0.8699	0.8715	0.8754	0.8675	0.8672	0.8625	0.8637	0.8644
QC9N1	1.1080	1.1105	1.1129	1.1080	1.1140	1.1168	1.1076	1.1142	1.1168	1.1443	1.1226	1.1191	1.1153	1.1179	1.1183
Q20	0.8771	0.8791	0.8808	0.8774	0.8927	0.8943	0.8771	0.8928	0.8943	0.8853	0.8799	0.8808	0.8736	0.8754	0.8770
Q21	0.8687	0.8654	0.8601	0.8747	0.8670	0.8616	0.8745	0.8670	0.8617	0.8331	0.8507	0.8533	0.8689	0.8656	0.8590
Q22	0.8695	0.8660	0.8596	0.8667	0.8554	0.8496	0.8666	0.8552	0.8495	0.8299	0.8509	0.8506	0.8691	0.8657	0.8583
Q23	0.8676	0.8649	0.8598	0.8642	0.8602	0.8549	0.8641	0.8601	0.8549	0.8426	0.8404	0.8353	0.8674	0.8644	0.8587
Q24	0.7969	0.7994	0.8016	0.7970	0.8000	0.8025	0.7968	0.8001	0.8026	0.8040	0.8031	0.8021	0.7937	0.7962	0.7986
Q25	1.2254	1.2266	1.2309	1.2254	1.2060	1.2091	1.2215	1.2072	1.2096	1.2431	1.2523	1.2306	1.2345	1.2429	1.2469
Q26	1.2129	1.2191	1.2230	1.2141	1.2081	1.2123	1.2154	1.2057	1.2104	1.2378	1.2303	1.2121	1.2084	1.2128	1.2193
LP1	1.0658	1.0688	1.0717	1.0655	1.0482	1.0501	1.0675	1.0475	1.0512	1.0614	1.0817	1.1330	1.0695	1.0727	1.0757
LP2	1.4787	1.4828	1.4866	1.4790	1.4873	1.4935	1.4872	1.4765	1.4808	1.5221	1.4424	1.3996	1.4370	1.4374	1.4388
LP3	1.4436	1.4472	1.4508	1.4434	1.4533	1.4607	1.4518	1.4429	1.4484	1.4990	1.4239	1.3844	1.4160	1.4182	1.4205
LP4	1.4973	1.5005	1.5043	1.4975	1.4751	1.4762	1.4874	1.4877	1.4895	1.5117	1.4583	1.4138	1.4500	1.4522	1.4552
LP5	1.4696	1.4730	1.4753	1.4702	1.4425	1.4426	1.4584	1.4552	1.4568	1.4727	1.4289	1.3942	1.4240	1.4252	1.4266
LP6	1.0295	1.0339	1.0379	1.0300	1.0309	1.0356	1.0297	1.0310	1.0356	1.0362	1.0412	1.0580	1.0256	1.0300	1.0335
Q81a	1.3201	1.3570	1.3269	1.3225	1.3709	1.3810	1.3210	1.3500	1.3531						
Q82a	1.4370	1.4179	1.4093	1.4223	1.3607	1.3424	1.3431	1.4394	1.4282						

	val														
	2a	2b	2c	2d	2e	2f	2g	2h	2i	2j	2k	21	2m	2n	20
C1	3.3128	3.3231	3.3328	3.3119	3.3359	3.3456	3.3121	3.3361	3.3457	3.4059	3.3300	3.3251	3.2904	3.3014	3.3118

											1		1		
C2	3.2614	3.2956	3.3263	3.2085	3.2778	3.3076	3.2079	3.2787	3.3088	3.3586	3.3548	3.3515	3.2425	3.2778	3.3080
C3	3.2394	3.2716	3.2971	3.2207	3.2763	3.3021	3.2211	3.2774	3.3029	3.3480	3.3453	3.3317	3.2479	3.2799	3.3076
C4	3.1908	3.2227	3.2492	3.1907	3.2656	3.2921	3.1903	3.2661	3.2931	3.3130	3.2595	3.2467	3.1621	3.1936	3.2213
C5	3.2013	3.2350	3.2635	3.2207	3.2764	3.3021	3.2211	3.2774	3.3029	3.3589	3.3020	3.2686	3.1950	3.2288	3.2588
C6	3.1563	3.1923	3.2214	3.2083	3.2777	3.3075	3.2079	3.2787	3.3088	3.3479	3.2619	3.2478	3.1580	3.1932	3.2223
C7	3.0578	3.0712	3.0833	3.0569	3.1086	3.1216	3.0568	3.1086	3.1209	3.1406	3.0897	3.0936	3.0413	3.0549	3.0671
C8	3.3705	3.3813	3.3908	3.3703	3.3628	3.3726	3.3690	3.3638	3.3739	3.4067	3.3846	3.3889	3.3425	3.3527	3.3632
C9	3.4298	3.4363	3.4429	3.4301	3.4555	3.4619	3.4299	3.4559	3.4615	3.4417	3.4388	3.4374	3.4159	3.4219	3.4266
C71	2.8256	2.9224	2.9988	2.8239	2.8905	2.9657	2.8223	2.8900	2.9667	2.9534	3.0561	3.0706	2.8067	2.9006	2.9681
C91	3.0149	3.0304	3.0442	3.0169	3.0653	3.0787	3.0153	3.0659	3.0795	3.0968	3.0616	3.0576	3.0048	3.0197	3.0343
N1	4.9256	4.9276	4.9289	4.9253	4.9245	4.9265	4.9246	4.9246	4.9259	4.9674	4.9441	4.9375	4.9363	4.9377	4.9403
N2	4.8015	4.8187	4.8346	4.8004	4.7831	4.7987	4.8031	4.7821	4.8009	4.8204	4.8646	4.9286	4.7988	4.8162	4.8319
N8	4.0976	4.1036	4.1121	4.0994	4.0763	4.0836	4.1078	4.0694	4.0768	4.0512	4.0400	3.9799	4.0173	4.0237	4.0307
N91	4.4474	4.4721	4.4946	4.4498	4.4554	4.4796	4.4487	4.4559	4.4798	4.4947	4.5173	4.5507	4.4255	4.4500	4.4734
O81	6.0630	6.0617	6.0802	6.0619	6.0675	6.0787	6.0720	6.0619	6.0717	6.1029	5.9665	5.9179	5.9297	5.9389	5.9490
O82	6.0586	6.0709	6.0812	6.0619	6.0675	6.0787	6.0720	6.0619	6.0717	6.0920	5.9725	5.9271	5.9424	5.9502	5.9586
H6	0.7667	0.7372	0.7074	0.7547	0.7327	0.7039	0.7544	0.7327	0.7038	0.6376	0.6312	0.6391	0.7636	0.7345	0.7045
H2	0.7428	0.7160	0.6888	0.7544	0.7324	0.7036	0.7544	0.7327	0.7038	0.6068	0.6023	0.5955	0.7388	0.7125	0.6859
H3	0.7813	0.7504	0.7195	0.7658	0.7303	0.7002	0.7656	0.7303	0.7002	0.6369	0.6485	0.6557	0.7770	0.7466	0.7162
H4	0.7617	0.7315	0.7012	0.7617	0.7366	0.7065	0.7613	0.7368	0.7066	0.6175	0.6179	0.6227	0.7602	0.7303	0.7002
H5	0.7501	0.7212	0.6919	0.7656	0.7300	0.6998	0.7656	0.7303	0.7002	0.6044	0.6247	0.6438	0.7452	0.7165	0.6885
H73	0.8045	0.7759	0.7453	0.7952	0.7646	0.7346	0.7950	0.7647	0.7347	0.6798	0.6792	0.6476	0.8018	0.7729	0.7427
H72	0.8026	0.7739	0.7433	0.7952	0.7649	0.7348	0.7950	0.7647	0.7347	0.6851	0.7058	0.6894	0.7946	0.7665	0.7362
H71	0.7792	0.7500	0.7195	0.7951	0.7646	0.7346	0.7950	0.7647	0.7347	0.6659	0.6856	0.6978	0.7787	0.7495	0.7187
QC1N1	0.1287	0.1277	0.1268	0.1288	0.1203	0.1194	0.1288	0.1203	0.1195	0.1049	0.1241	0.1253	0.1280	0.1270	0.1259
QC1C6	0.5687	0.5584	0.5501	0.5507	0.5248	0.5164	0.5498	0.5238	0.5156	0.4843	0.5440	0.5491	0.5752	0.5648	0.5561
QC1C2	0.5106	0.5015	0.4932	0.5289	0.5152	0.5070	0.5298	0.5154	0.5072	0.4725	0.4931	0.4946	0.5236	0.5137	0.5055
QC2C3	0.5348	0.5203	0.5082	0.5591	0.5191	0.5073	0.5598	0.5187	0.5070	0.4694	0.4853	0.4897	0.5288	0.5140	0.5012
QC2H2	0.4367	0.4388	0.4436	0.4413	0.4269	0.4328	0.4417	0.4268	0.4328	0.4853	0.4842	0.4868	0.4447	0.4461	0.4506
QC3C4	0.5649	0.5501	0.5387	0.5705	0.5349	0.5234	0.5703	0.5342	0.5228	0.5070	0.5274	0.5349	0.5702	0.5558	0.5427
QC3H3	0.4155	0.4183	0.4243	0.4306	0.4381	0.4445	0.4305	0.4378	0.4442	0.4608	0.4453	0.4428	0.4134	0.4157	0.4213
QC4C5	0.5818	0.5665	0.5540	0.5760	0.5273	0.5162	0.5758	0.5272	0.5162	0.4941	0.5461	0.5600	0.5921	0.5765	0.5634
QC4H4	0.4422	0.4452	0.4512	0.4421	0.4342	0.4397	0.4424	0.4341	0.4395	0.4872	0.4925	0.4907	0.4514	0.4545	0.4601
QC5C6	0.5817	0.5663	0.5538	0.5571	0.5292	0.5181	0.5561	0.5284	0.5173	0.4830	0.5284	0.5417	0.5725	0.5574	0.5443

QC5H5	0.4494	0.4515	0.4568	0.4344	0.4348	0.4410	0.4342	0.4346	0.4408	0.4822	0.4809	0.4745	0.4534	0.4554	0.4596
QC6H6	0.4517	0.4554	0.4625	0.4478	0.4279	0.4338	0.4475	0.4273	0.4333	0.4677	0.4944	0.4905	0.4513	0.4546	0.4627
QC7N1	0.2558	0.2538	0.2521	0.2559	0.2478	0.2456	0.2561	0.2478	0.2458	0.2308	0.2479	0.2502	0.2558	0.2539	0.2515
QC7N2	0.3389	0.3344	0.3301	0.3394	0.3365	0.3313	0.3398	0.3362	0.3315	0.3251	0.3284	0.3172	0.3457	0.3409	0.3370
Q15	0.5681	0.5463	0.5295	0.5687	0.5247	0.5081	0.5691	0.5248	0.5080	0.5163	0.5201	0.5169	0.5766	0.5553	0.5405
QC8N8	0.2734	0.2722	0.2708	0.2730	0.2891	0.2878	0.2717	0.2902	0.2890	0.3021	0.2998	0.3096	0.3034	0.3022	0.3010
QC8N2	0.2571	0.2536	0.2506	0.2574	0.2543	0.2517	0.2566	0.2547	0.2508	0.2391	0.2465	0.2358	0.2603	0.2567	0.2531
QC8C9	0.4456	0.4425	0.4395	0.4455	0.4437	0.4403	0.4459	0.4434	0.4402	0.4357	0.4455	0.4460	0.4577	0.4547	0.4522
QC9N1	0.1629	0.1616	0.1604	0.1629	0.1588	0.1574	0.1630	0.1587	0.1575	0.1449	0.1549	0.1562	0.1592	0.1580	0.1572
Q20	0.4667	0.4626	0.4591	0.4661	0.4355	0.4322	0.4666	0.4352	0.4321	0.4456	0.4618	0.4615	0.4759	0.4721	0.4686
Q21	0.4971	0.4908	0.4909	0.4834	0.4915	0.4910	0.4838	0.4917	0.4909	0.5466	0.4989	0.4868	0.4987	0.4928	0.4953
Q22	0.4611	0.4542	0.4546	0.4679	0.4795	0.4801	0.4682	0.4799	0.4803	0.5122	0.4634	0.4653	0.4663	0.4594	0.4616
Q23	0.4871	0.4792	0.4785	0.4952	0.4921	0.4912	0.4956	0.4923	0.4912	0.5095	0.5018	0.5174	0.4890	0.4819	0.4826
Q24	0.9224	0.9106	0.9004	0.9213	0.9039	0.8934	0.9223	0.9034	0.8930	0.8939	0.8997	0.8915	0.9425	0.9311	0.9197
Q25	0.4407	0.3501	0.4242	0.4338	0.3157	0.3027	0.4125	0.3488	0.3433	0.1306	0.1326	0.1418	0.1393	0.1366	0.1351
Q26	0.2634	0.2768	0.2803	0.2725	0.3277	0.3548	0.3512	0.2657	0.2696	0.1314	0.1387	0.1475	0.1467	0.1447	0.1427
LP1	0.3285	0.3251	0.3225	0.3289	0.3466	0.3447	0.3265	0.3475	0.3433	0.3379	0.3144	0.2533	0.3261	0.3226	0.3199
LP2	0.1725	0.1714	0.1704	0.1724	0.1691	0.1672	0.1693	0.1733	0.1721	0.1527	0.2240	0.2726	0.2236	0.2248	0.2240
LP3	0.1861	0.1850	0.1841	0.1862	0.1825	0.1798	0.1827	0.1869	0.1849	0.1619	0.2342	0.2834	0.2344	0.2351	0.2342
LP4	0.1623	0.1614	0.1608	0.1620	0.1705	0.1703	0.1666	0.1653	0.1651	0.1553	0.2116	0.2578	0.2122	0.2116	0.2106
LP5	0.1695	0.1690	0.1685	0.1696	0.1803	0.1808	0.1752	0.1742	0.1745	0.1649	0.2228	0.2665	0.2223	0.2220	0.2215
LP6	0.4329	0.4270	0.4221	0.4323	0.4320	0.4260	0.4327	0.4319	0.4260	0.4316	0.4228	0.3894	0.4435	0.4378	0.4327
Q81a	-0.4045	-0.3047	-0.3883	-0.3967	-0.2603	-0.2466	-0.3801	-0.2966	-0.2914						
Q82a	-0.1974	-0.2156	-0.2232	-0.2097	-0.2767	-0.3106	-0.3098	-0.1950	-0.2032						



Figure S8. Fourier residual maps for model 3a).

Table S5 . κ and P_{va}	al for models 3a-f
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	3a	3b	3c	3d	3e	3f
	kp1					
C1	1.0376	1.0362	1.0376	1.0382	1.0369	1.0384
C2	1.0218	1.0253	1.0246	1.0227	1.0258	1.0259
C3	1.0217	1.0236	1.0243	1.0225	1.0245	1.0256
C4	1.0201	1.0183	1.0214	1.0212	1.0200	1.0228
C5	1.0217	1.0235	1.0216	1.0225	1.0241	1.0229
C6	1.0218	1.0225	1.0209	1.0227	1.0230	1.0219
C7	1.0371	1.0365	1.0356	1.0385	1.0381	1.0371
C8	1.0375	1.0372	1.0377	1.0393	1.0390	1.0401
C9	1.0384	1.0380	1.0382	1.0392	1.0388	1.0395
C71	1.0101	1.0104	1.0138	1.0113	1.0116	1.0153
C91	1.0411	1.0407	1.0413	1.0421	1.0415	1.0432
N1	1.0164	1.0160	1.0159	1.0170	1.0167	1.0163
N2	0.9988	0.9985	0.9995	0.9993	0.9992	1.0005
N8	1.0192	1.0188	1.0166	1.0199	1.0198	1.0174
N91	0.9943	0.9942	0.9951	0.9948	0.9947	0.9963
O81	0.9952	0.9963	0.9961	0.9956	0.9967	0.9965
O82	0.9952	0.9951	0.9957	0.9956	0.9956	0.9960
H6	1.4000	1.4000	1.4000	1.3353	1.3300	1.3206
H2	1.4000	1.4000	1.4000	1.3353	1.3292	1.3179
H3	1.4000	1.4000	1.4000	1.3330	1.3283	1.3221
H4	1.4000	1.4000	1.4000	1.3298	1.3302	1.3173
H5	1.4000	1.4000	1.4000	1.3330	1.3276	1.3195
H73	1.4000	1.4000	1.4000	1.3454	1.3391	1.3336
H72	1.4000	1.4000	1.4000	1.3454	1.3384	1.3246
H71	1.4000	1.4000	1.4000	1.3454	1.3386	1.3290

3a	3b	3c	3d	3e	3f
val					

C1	3.979	3.998	3.987	3.971	3.987	3.984
C2	4.226	4.206	4.226	4.197	4.178	4.192
C3	4.208	4.191	4.187	4.186	4.171	4.154
C4	4.218	4.236	4.211	4.195	4.211	4.176
C5	4.208	4.191	4.214	4.186	4.171	4.184
C6	4.226	4.206	4.211	4.197	4.178	4.181
C7	3.869	3.873	3.884	3.855	3.859	3.867
C8	4.043	4.045	4.038	4.022	4.021	4.005
C9	3.996	4.000	4.000	3.989	3.992	3.985
C71	4.614	4.610	4.576	4.559	4.548	4.509
C91	4.008	4.013	4.005	3.993	3.992	3.978
N1	5.009	5.011	5.009	5.003	5.003	5.004
N2	5.226	5.229	5.219	5.214	5.215	5.196
N8	4.660	4.665	4.687	4.651	4.656	4.678
N91	5.221	5.226	5.219	5.208	5.211	5.195
O81	6.181	6.174	6.183	6.173	6.166	6.175
O82	6.181	6.173	6.166	6.173	6.166	6.160
H6	0.748	0.752	0.759	0.788	0.795	0.810
H2	0.748	0.751	0.747	0.788	0.794	0.799
H3	0.747	0.751	0.760	0.787	0.794	0.809
H4	0.744	0.743	0.749	0.788	0.788	0.803
H5	0.747	0.751	0.748	0.787	0.794	0.798
H73	0.731	0.736	0.750	0.763	0.772	0.794
H72	0.731	0.734	0.732	0.763	0.771	0.782
<u>H7</u> 1	0.731	0.735	0.736	0.763	0.771	0.783









Figure S9. Fourier residual density maps for sub-models 4 not included in the main article, contour 0.05, resolution 1A⁻¹