

Article

Vibrational Circular Dichroism Detects Symmetry Breaking due to Conformational Mobility in C₂-Symmetry Chiral Molecules and Provides Further Insight into Inter-Chromophoric Interactions

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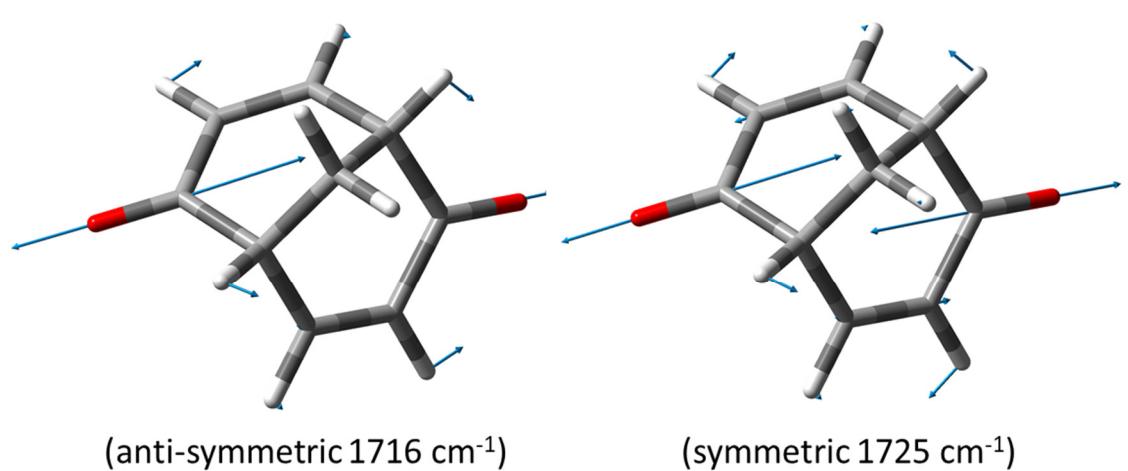
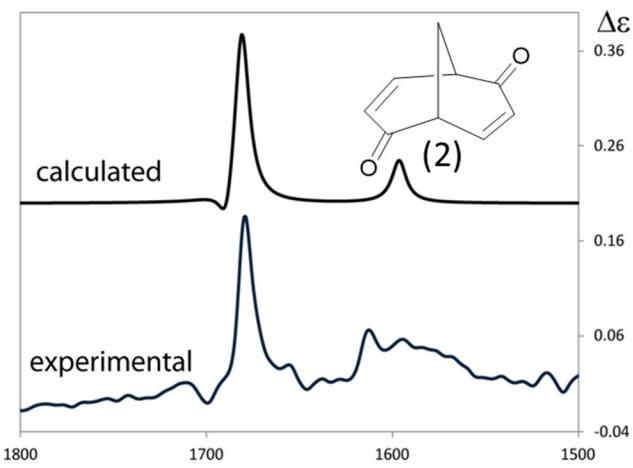
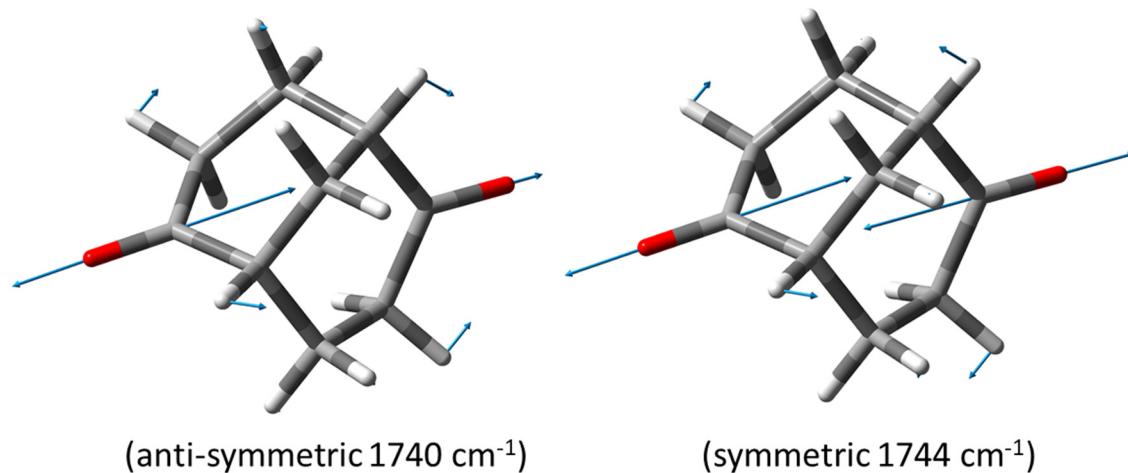
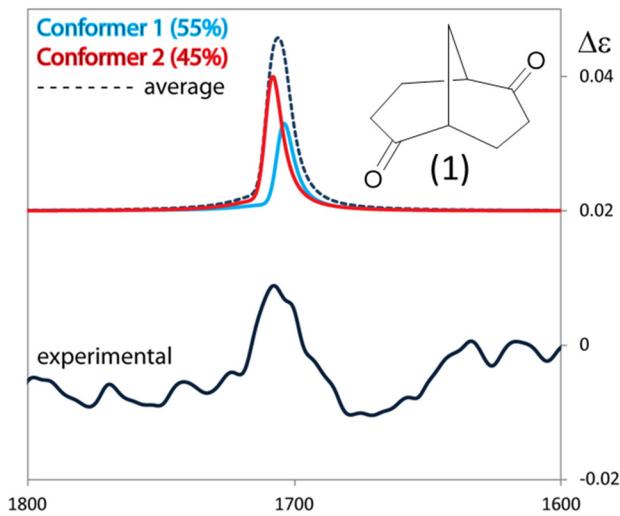


Figure SI-1. Calculated spectra and carbonyl stretching normal modes description for compounds 1-cc conformer (top) and 2 (bottom).

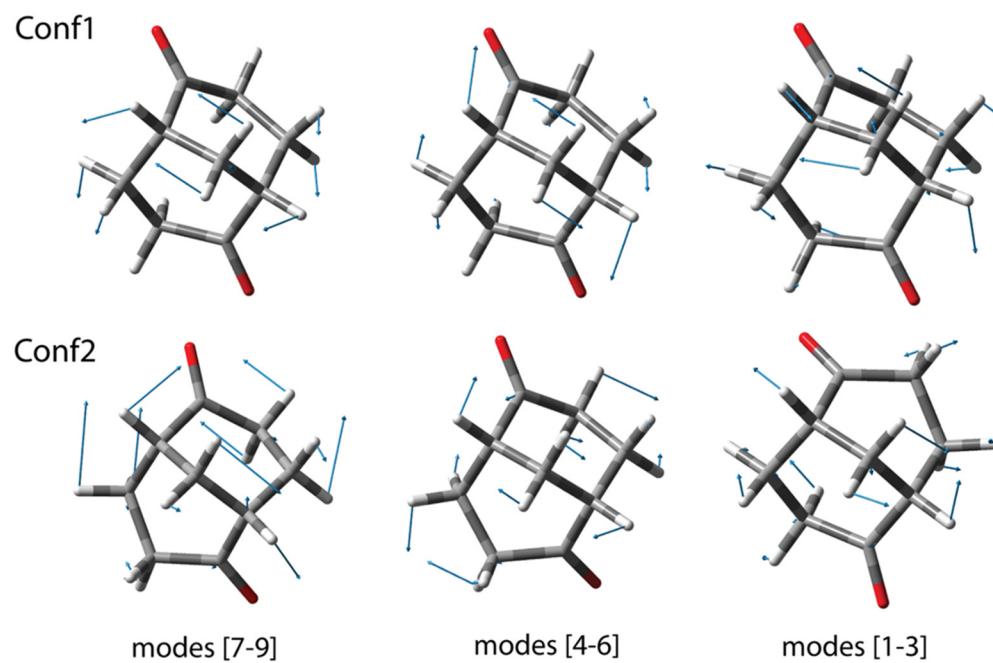
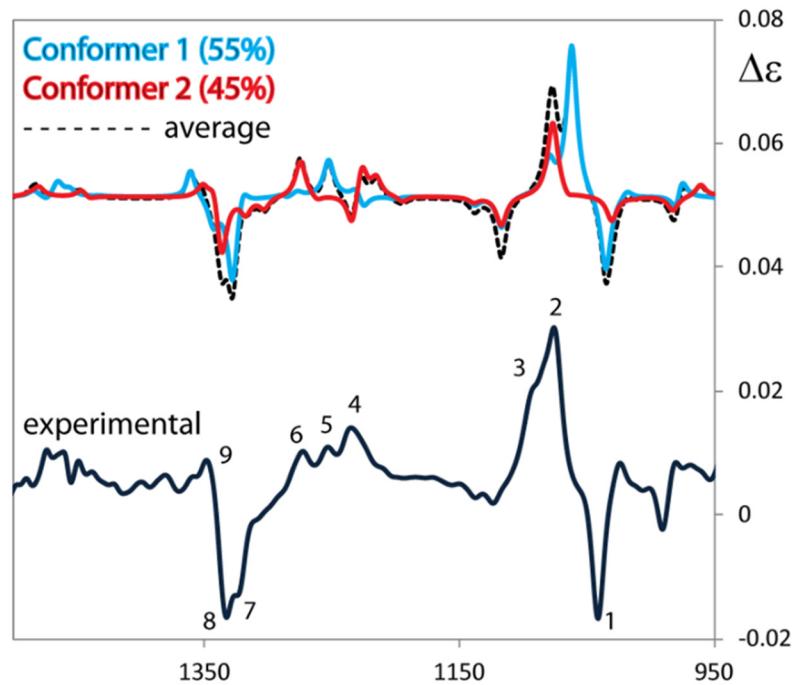


Figure SI-2. Calculated spectra and description of vibrational modes in the MID-IR region of compound 1-cc conformer. Modes are grouped and displayed for vibration transition type (only relative phases change within each group).

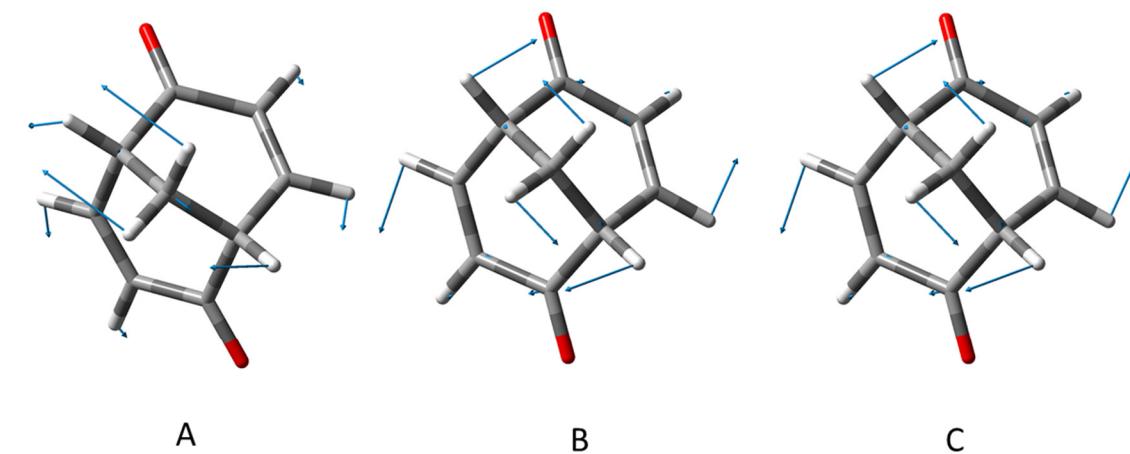
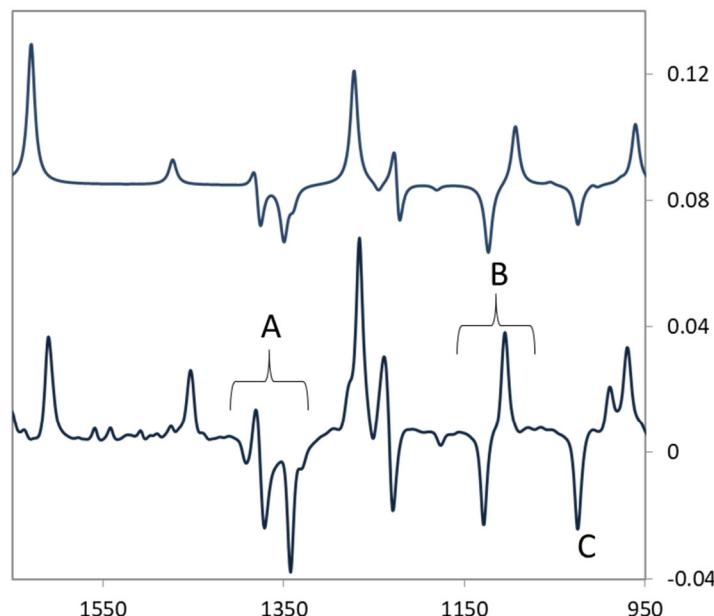
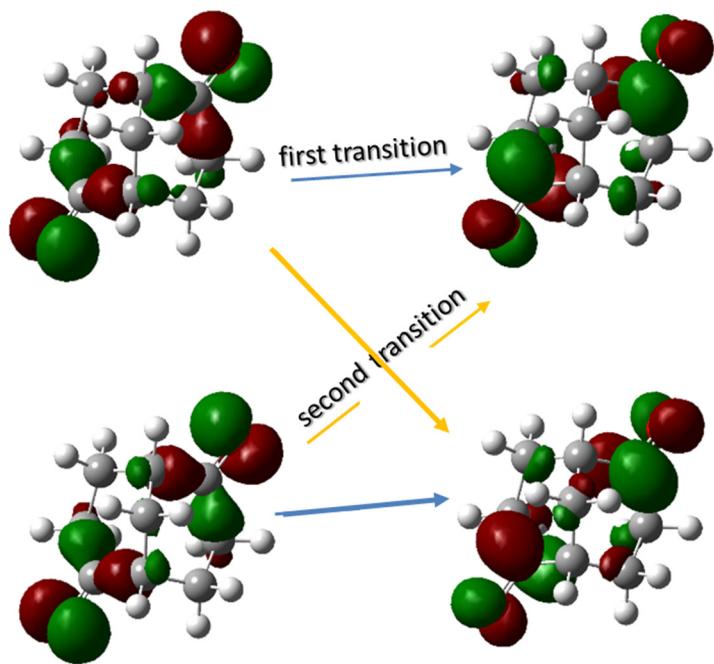


Figure SI-3. Calculated spectra and description of vibrational modes in the MID-IR region of compound 2.

Conf1



Conf2

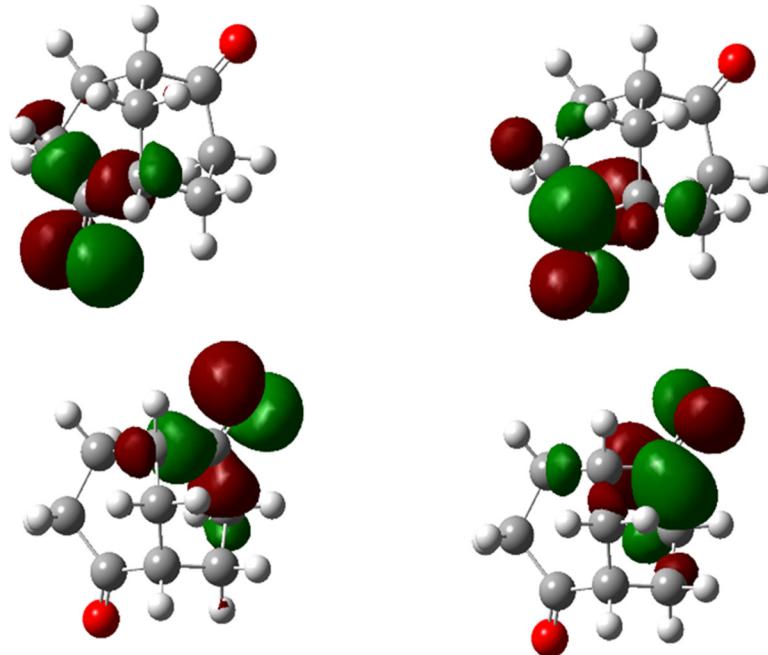


Figure SI-4. Natural Transition Orbital investigation of first and second electronic transitions of compound 1 for both the cc and cb conformers. Level of theory CAM-B3LYP/aug-cc-pVDZ.

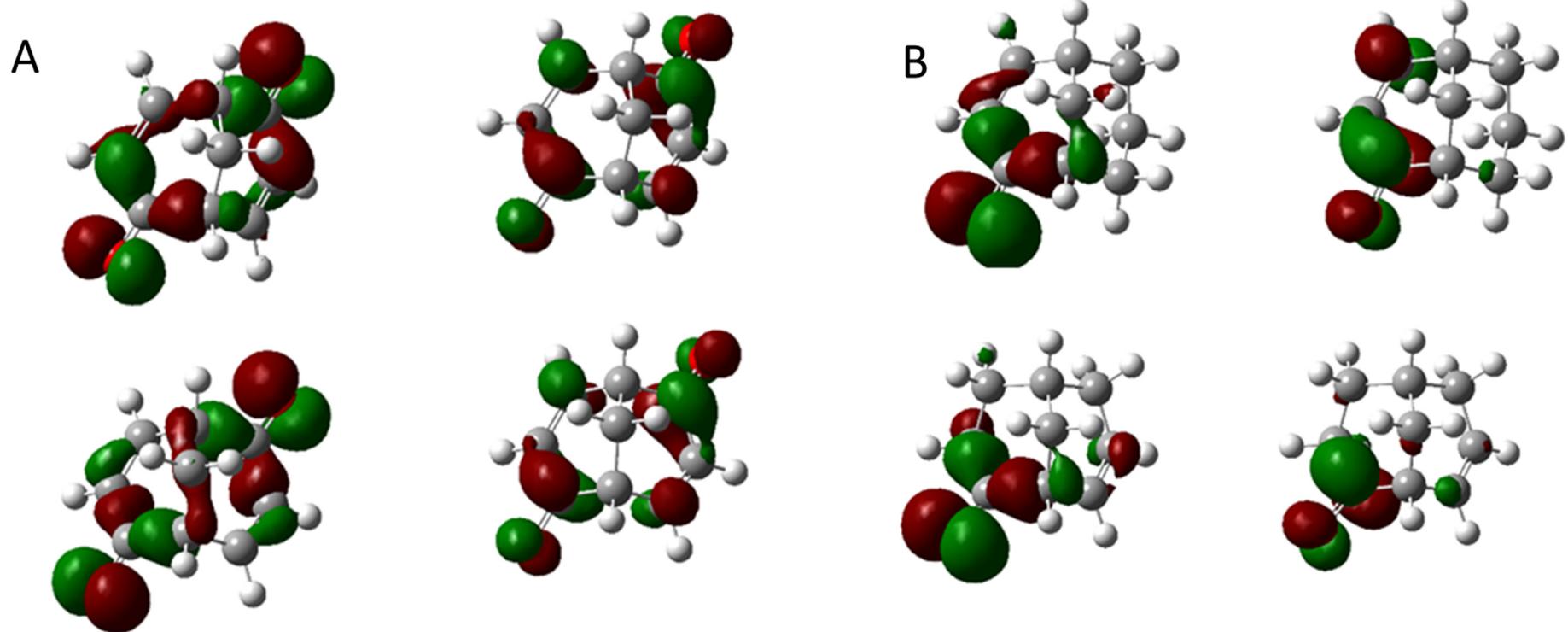
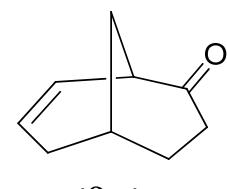
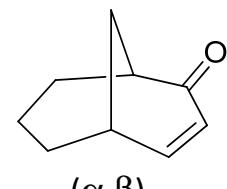
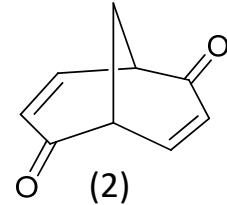


Figure SI-5. Molecular orbitals involved in the first and second electronic transitions of compound 2. Level of theory CAM-B3LYP/aug-cc-pVDZ.

Table SI-1. Compound **1** in **cc** and **cb** conformation, and mono-ketone model. APT and AAT (a.u) of C and O atoms are reported in the reference cartesian system of each carbonyl group, z axis pointing towards the O atom, y-axis pointing towards the nearby asymmetric carbon C*.

A	1-cc	APT			AAT			cm-1	Dip. str.	Rot. str.	E-M
	C						antis.	1775	1015	59	24
		0.031	-0.013	-0.106	0.055	0.574	0.080	symm.	1779	10	-20
		0.055	0.890	-0.052	-1.206	0.046	-0.126				
		-0.105	-0.006	1.313	-0.154	0.118	0.041				
O											
		-0.299	0.003	0.037	-0.009	-1.555	-0.063				
		-0.023	-0.533	0.015	1.279	-0.013	0.030				
		0.040	0.021	-1.224	0.173	-0.150	-0.061				
B	1-cb	APT			AAT			cm-1	Dip. str.	Rot. str.	E-M
	C						antis.	1779	864	112	59
		0.096	-0.009	0.015	-0.129	0.675	0.107	symm.	1783	116	-39
		0.083	0.929	-0.001	-1.016	0.049	-0.740				
		-0.150	-0.040	1.193	-0.409	0.634	0.106				
O							Average		1005	54	
		-0.337	-0.001	0.023	0.051	-1.576	-0.063				
		-0.024	-0.544	0.032	1.168	-0.011	0.405				
		0.130	0.052	-1.168	0.321	-0.651	-0.042				
C											
		0.017	0.011	-0.064	0.035	0.558	0.025				
		0.069	0.848	-0.081	-1.206	0.099	0.073				
		-0.033	-0.024	1.360	-0.142	0.048	0.031				
O											
		-0.286	0.000	0.022	-0.016	-1.529	-0.051				
		-0.028	-0.520	0.025	1.287	-0.029	-0.040				
		-0.009	0.023	-1.234	0.133	-0.043	-0.063				
C	Mono-ketone	APT			AAT			cm-1	Dip. str.	Rot. str.	E-M
	C										
		0.032	-0.013	-0.101	0.060	0.574	0.087				
		0.061	0.888	-0.048	-1.128	0.018	-0.134				
		-0.101	-0.004	1.307	-0.162	0.091	0.026				
O											
		-0.304	0.003	0.037	-0.016	-1.532	-0.069				
		-0.025	-0.537	0.013	1.241	-0.003	0.028				
		0.040	0.017	-1.243	0.192	-0.126	-0.050				

Table SI-2. Compound 2 and mono-ketone diene models. APT and AAT (a.u.) of C and O atoms are reported in the reference cartesian system of each carbonyl group, z axis pointing towards the O atom, y-axis pointing towards the nearby asymmetric carbon C*.



A	2	APT			AAT				cm-1	Dip. str.	Rot. str.	E-M
	C							antis.	1750	1334	388	15
		0.110	0.025	0.014	0.006	0.535	0.433	symm.	1759	44	-95	180
		0.144	0.871	0.094	-2.093	0.482	-0.406	.				
		-0.195	-0.053	1.542	-0.309	0.260	0.318	sum		1378	293	
	O											
		-0.313	0.002	-0.004	0.013	-1.474	-0.130					
		-0.026	-0.500	-0.017	1.427	-0.104	0.154					
		0.128	0.056	-1.358	0.305	-0.311	-0.225					
B	α,β	APT			AAT				cm-1	Dip. str.	Rot. str.	E-M
	C								1742	603	51	80
		0.137	-0.012	-0.060	-0.028	0.475	0.359					
		0.035	0.865	0.267	-1.498	0.176	-0.342					
		-0.152	0.018	1.369	-0.421	0.353	0.040					
	O											
		-0.343	0.005	0.022	0.037	-1.473	-0.091					
		-0.006	-0.528	-0.046	1.288	-0.051	0.181					
		0.100	0.027	-1.283	0.357	-0.375	-0.038					
C	β,γ	APT			AAT				cm-1	Dip. str.	Rot. str.	E-M
	C								1781	612	67	39
		0.009	0.020	-0.026	0.103	0.544	0.159					
		0.098	0.865	-0.209	-1.245	0.161	-0.046					
		-0.052	-0.050	1.458	-0.060	-0.115	0.174					
	O											
		-0.280	0.000	0.009	-0.041	-1.484	-0.096					
		-0.037	-0.519	0.057	1.256	-0.041	-0.034					
		0.010	0.030	-1.318	0.082	0.061	-0.158					

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