

Fourth stable radical species in X-irradiated solid-state sucrose

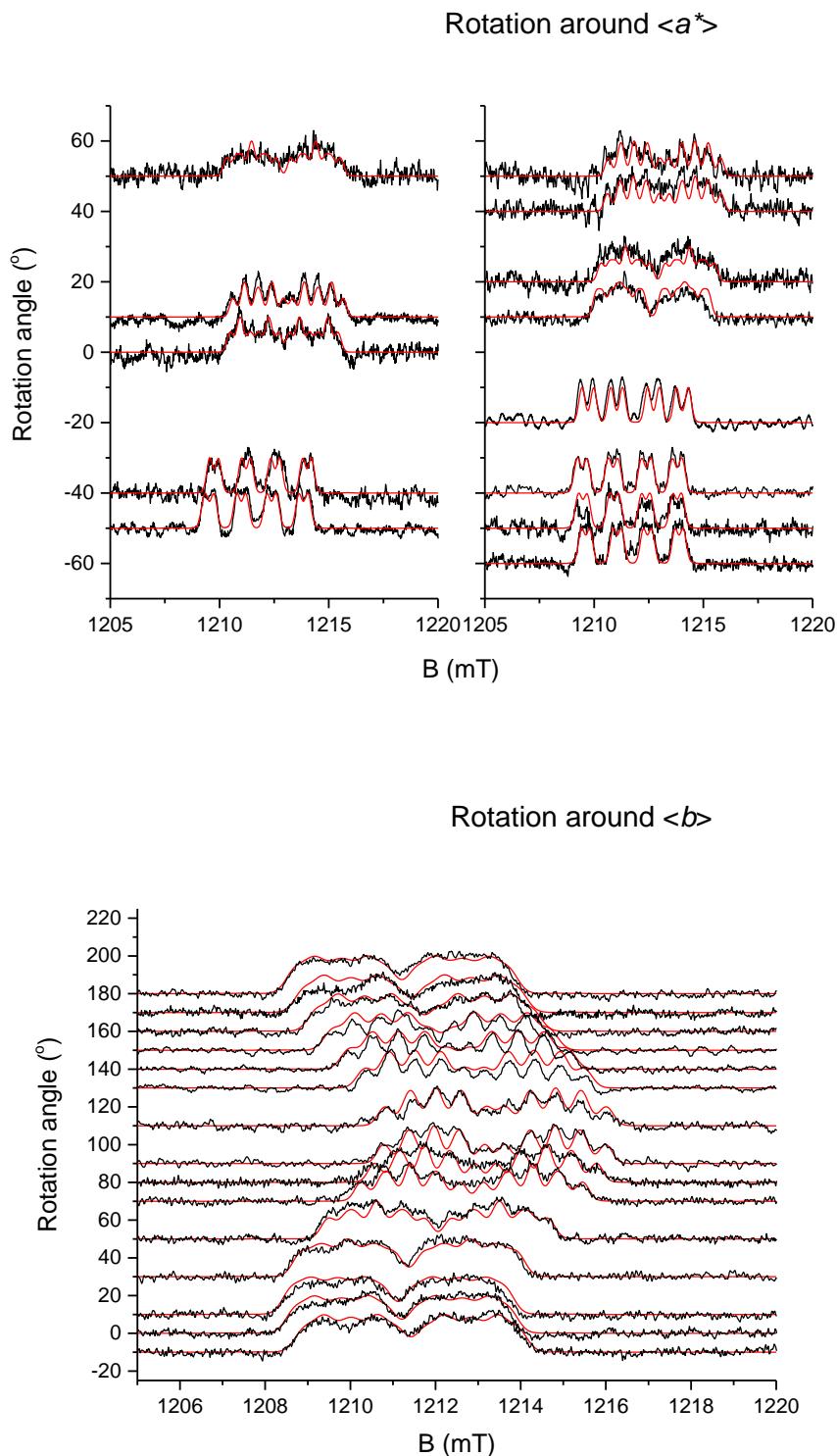
Jevgenij Kusakovskij¹, Ignacio Caretti², Sabine Van Doorslaer², Freddy Callens¹ and Henk Vrielinck¹

¹ Ghent University, Department of Solid State Sciences, Krijgslaan 281-S1, 9000 Ghent, Belgium

²Antwerp University, Department of Physics, Universiteitsplein 1, 2610 Wilrijk, Belgium

Supporting information

Figure S1. Angular dependence of the Q-band EIE spectra of the fourth radical species recorded at 110 K: experiment (black) and simulation (red) of the absorption EPR spectra using the experimental tables from Table 1 and Figure 1.



Rotation around $\langle c \rangle$

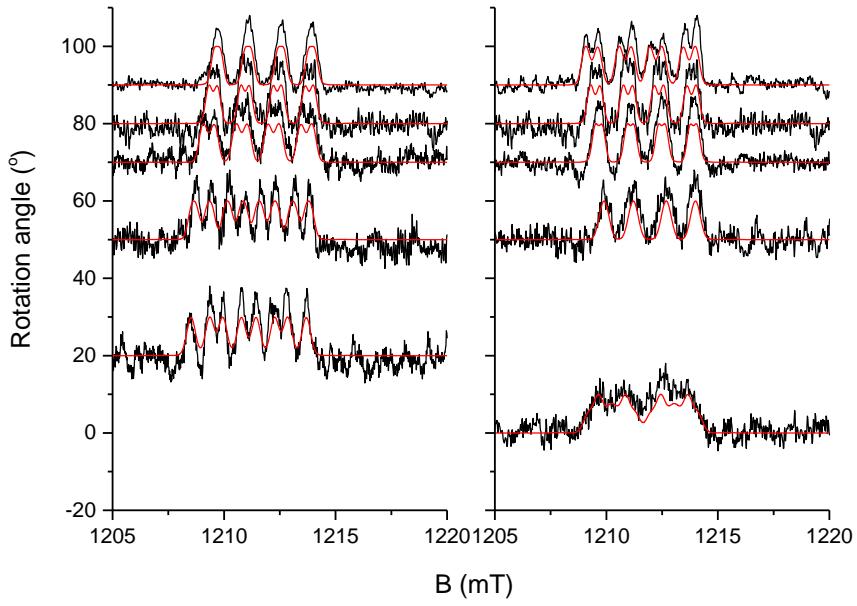


Table S1. Schonland conjugates of the experimental tensors (Table 1) of the fourth radical species in X-irradiated sucrose. Principal values of the g tensor are also presented as shifts δg_i ($i = \{1,2,3\}$) from the free electron $g_e = 2.002319$ and expressed in ppt. HF tensors are expressed in MHz.

Tensor	g-value	Iso	δg_i / Aniso	Eigenvectors		
				a^*	b	c
g	2.00173		-0.59	0.397	0.223	0.890
	2.00484		2.52	-0.313	-0.879	0.360
	2.00596		3.64	0.863	-0.422	-0.278
$\beta 1$ -HF		80.23	-5.48	0.505	0.640	0.579
			0.82	-0.839	0.204	0.505
			4.66	0.205	-0.740	0.640
$\beta 2$ -HF		36.22	-3.84	0.432	-0.279	0.858
			-2.42	-0.794	0.334	0.508
			6.26	-0.428	-0.901	-0.077
$\beta 3$ -HF		-15.78	-7.96	0.799	-0.593	0.098
			-3.36	-0.123	-0.001	0.992
			11.32	-0.588	-0.805	-0.074
hydroxy-HF		-6.11	-11.27	0.645	-0.341	0.684
			-5.48	0.650	-0.227	-0.725
			16.75	0.402	0.912	0.075

Table S2. Comparison of the experimental HF tensors, (expressed in MHz,) in the reference frame of the **g** tensor to the calculated tensors, using the structure in Figure 5B.

					β1-HF			β2-HF			β3-HF			hydroxy-HF			
					Iso→	80.12		36.22		-15.78		-7.07					
					Aniso→	-4.31	-1.45	5.76	-3.84	-2.42	6.26	-8.09	-3.07	11.17	-7.96	-6.56	14.52
					a*	0.600	-0.686	0.412	0.931	0.204	-0.302	0.053	0.998	0.045	0.996	-0.077	0.040
					b	-0.800	-0.505	0.324	0.346	-0.754	0.558	-0.071	-0.041	0.997	0.086	0.835	-0.543
					c	-0.014	-0.524	-0.852	-0.114	-0.624	-0.773	0.996	-0.056	0.068	0.009	0.545	0.839
H1												5					
		-13.02	-6.71	0.003	-0.142	0.990						4					
				-3.52	1.000	-0.002	-0.003					5					
				10.23	0.002	0.990	0.142										
H3b		38.03	-3.56	0.964	0.006	-0.265				22							
				-2.38	-0.111	-0.899	-0.424			23							
				5.94	-0.241	0.438	-0.866			9							
H3a		67.35	-3.50	0.591	-0.806	0.027	2										
				-2.71	-0.701	-0.530	-0.476	3									
				6.21	0.399	0.262	-0.879		4								
H2		-9.61	-10.74	1.000	-0.008	-0.006								5		5	
				-3.00	0.010	0.834	0.552										
				13.74	0.001	-0.552	0.834								2		