

Supporting Information

for

RDC-based Determination of the Relative
Configuration of the Fungicidal Cyclopentenone 4,6-
Diacetylhygrophorone A¹²

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NMR Spectra of 4,6-Diacetylhygrophorone A¹² (3)

Spectra were acquired on a Bruker Avance II 700MHz US spectrometer equipped with a TXI 700S4 cryo probe. Relevant delays and durations are given, were appropriate. Unless otherwise noted experiments were performed at 300K.

Assignment of Chemical Shifts

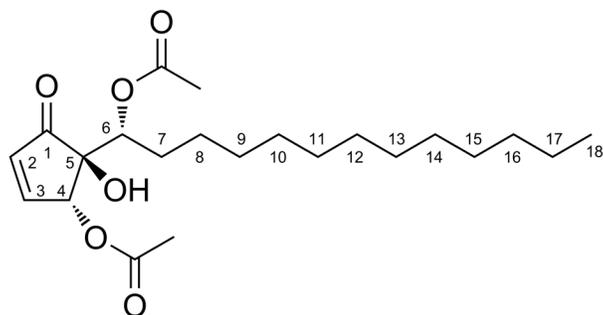
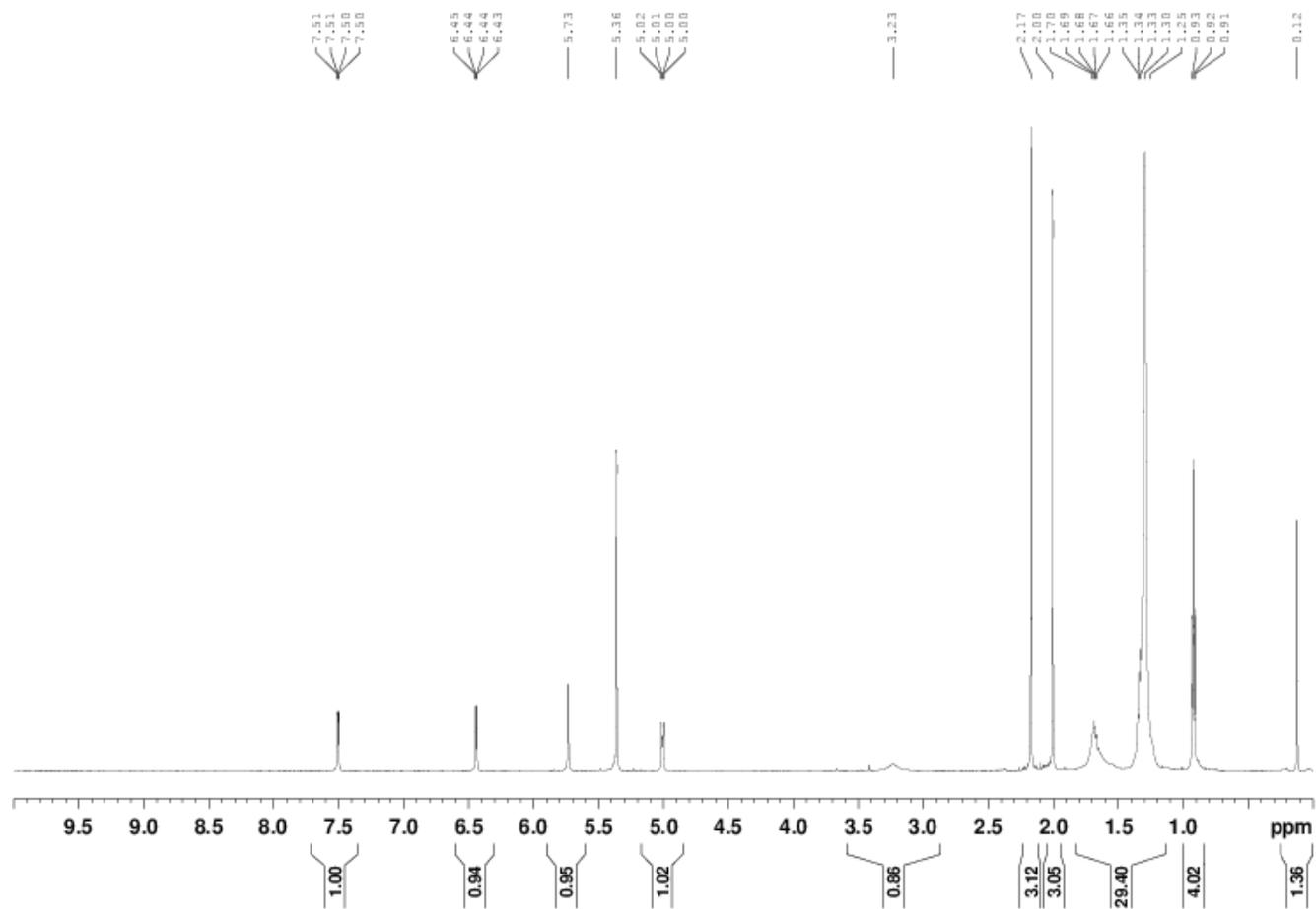


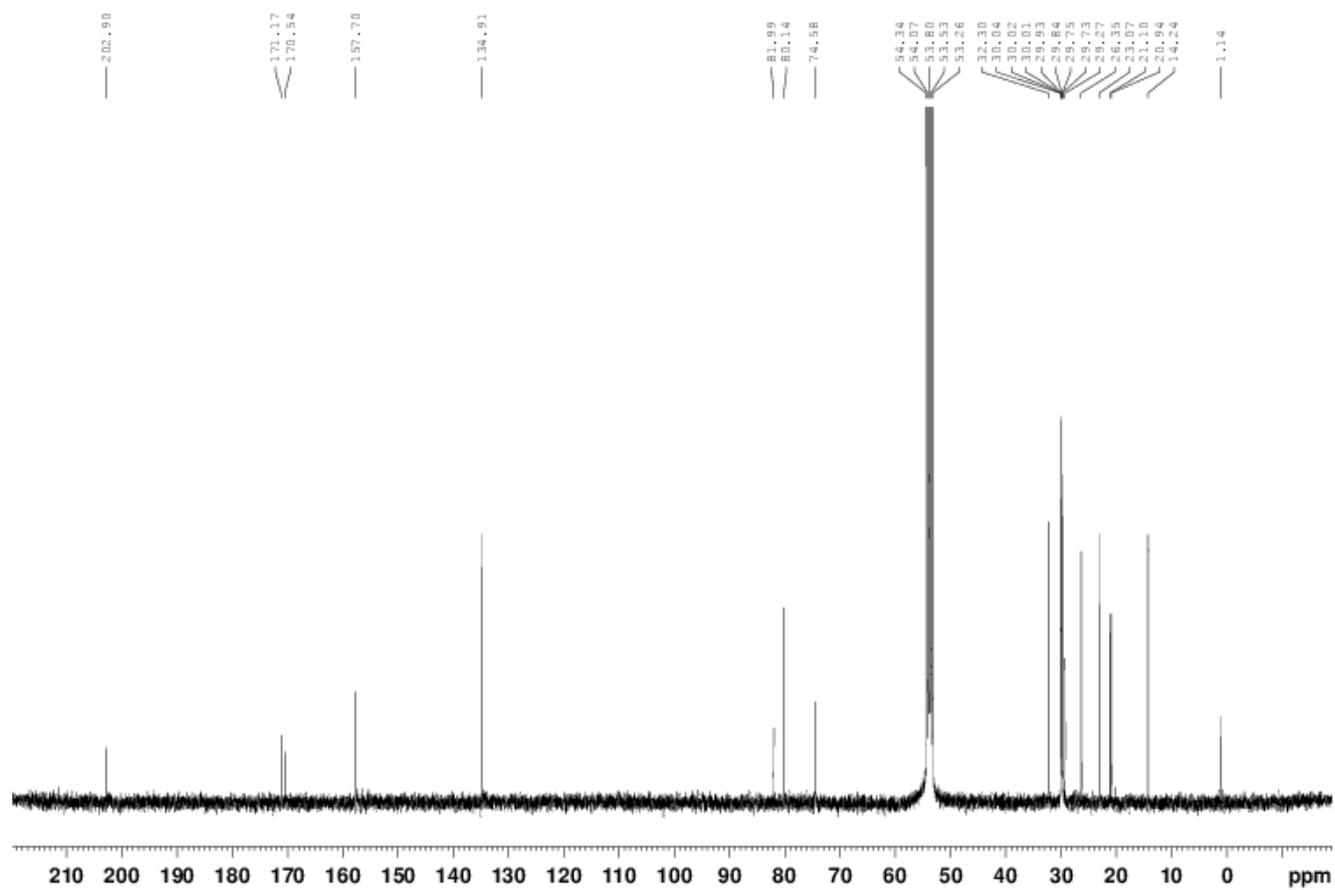
Table SI 1: Assignment of ¹H and ¹³C chemical shifts of 4,6-diacetylhygrophorone A¹² (3) in isotropic solution (CD₂Cl₂). Numbering scheme see above.

Position	¹ H (ppm)	¹³ C (ppm)
1	–	202.9
2	6.4 (dd)	134.9
3	7.5 (dd)	157.7
4	5.7 (dd)	80.2
4-OAc	–	171.2
4-OAc	2.1 (s)	21.1
5	–	80.2
6	5.0 (dd)	74.6
6-OAc	–	170.5
6-OAc	2.0	20.9
7	1.6 (m)	32.3
8-17	1.1 – 1.3 (m)	26.0 – 30.0
18	0.9 (t)	14.2

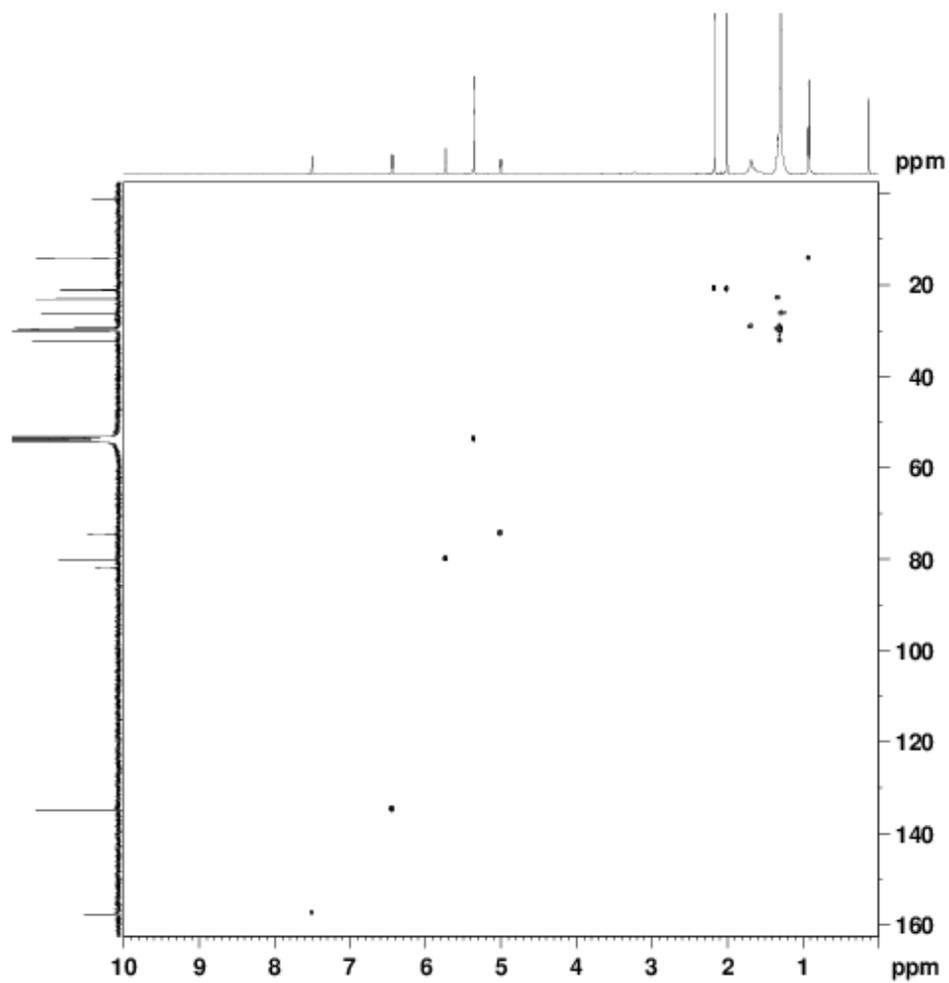
^1H , 300K, isotropic



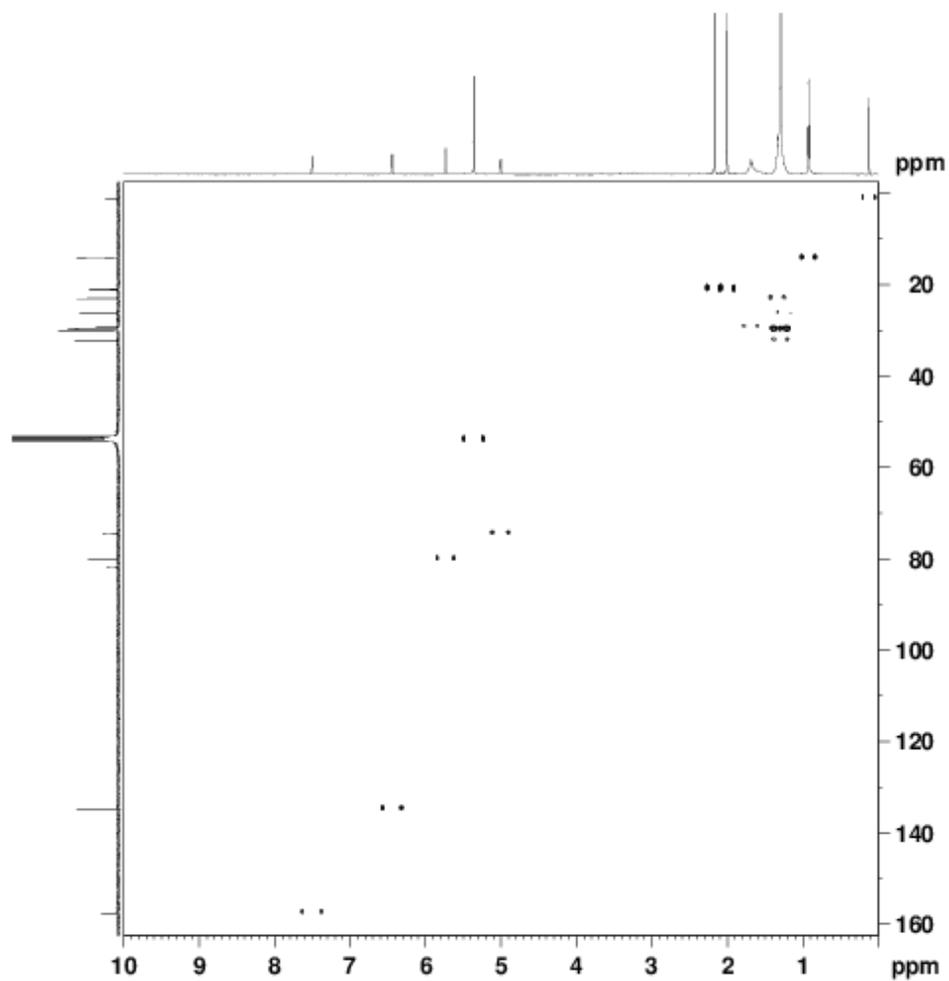
^{13}C , 300K, isotropic



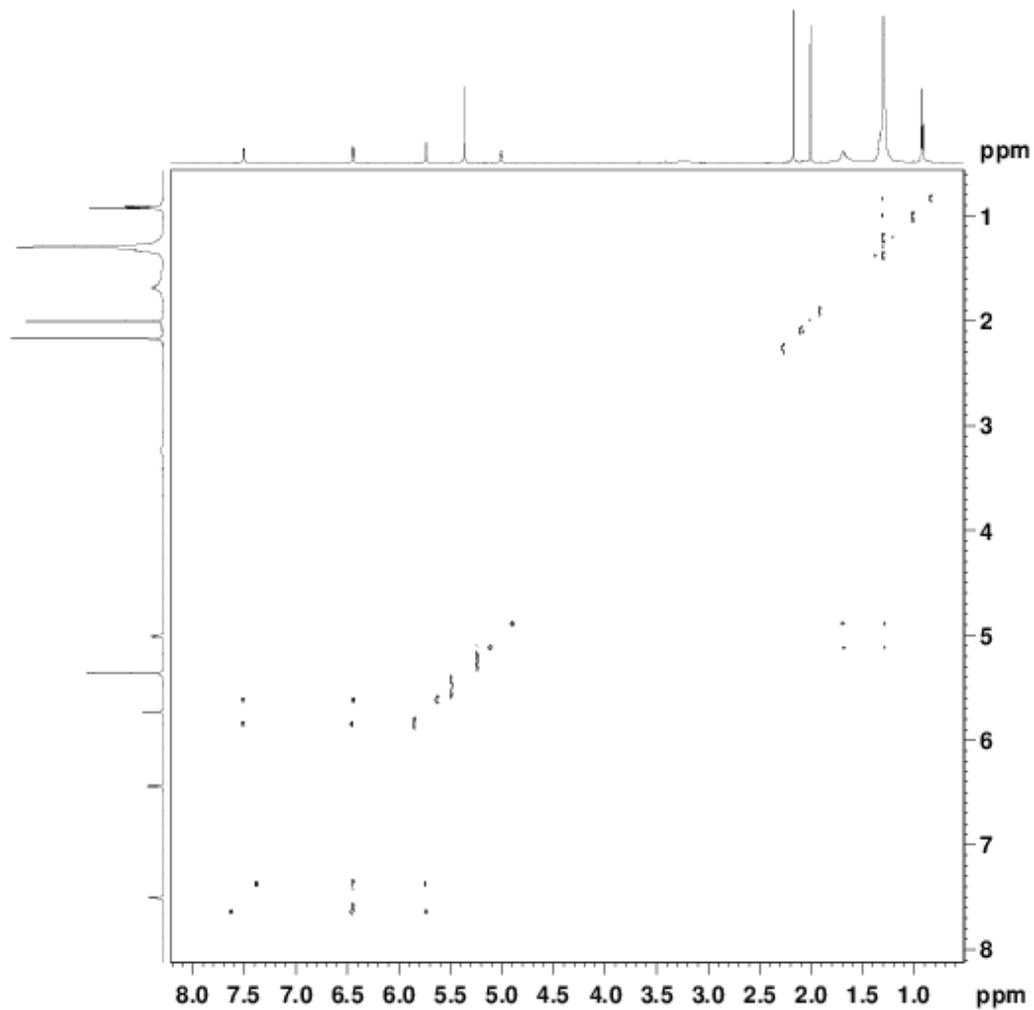
^1H - ^{13}C HSQC, 300K, INEPT delay optimized for 145 Hz, isotropic



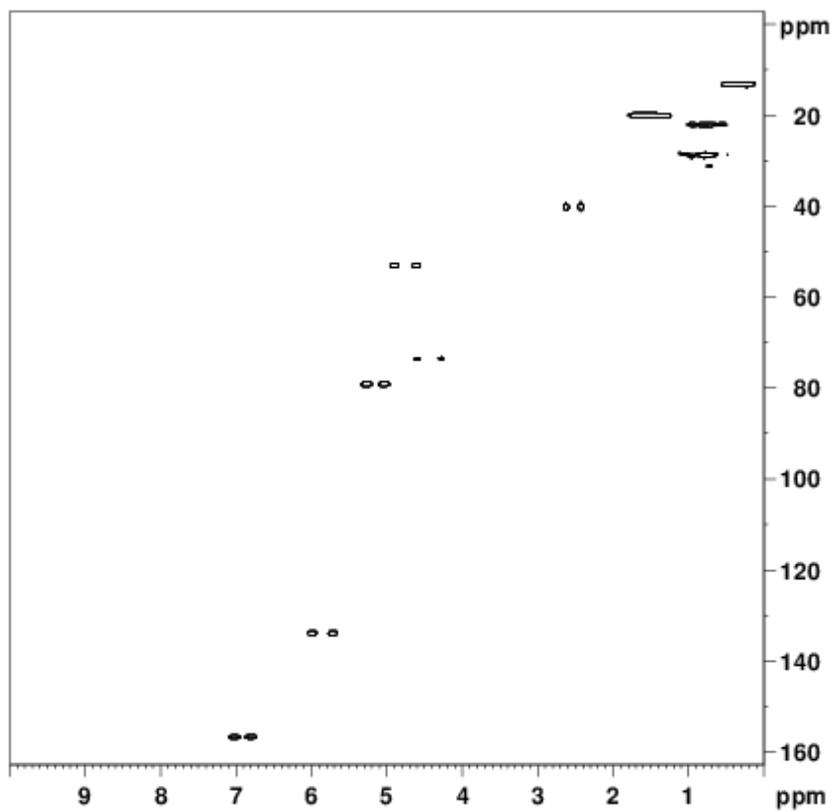
^1H - ^{13}C CLIP-HSQC, 1 300K, INEPT delay optimized for 145 Hz, isotropic



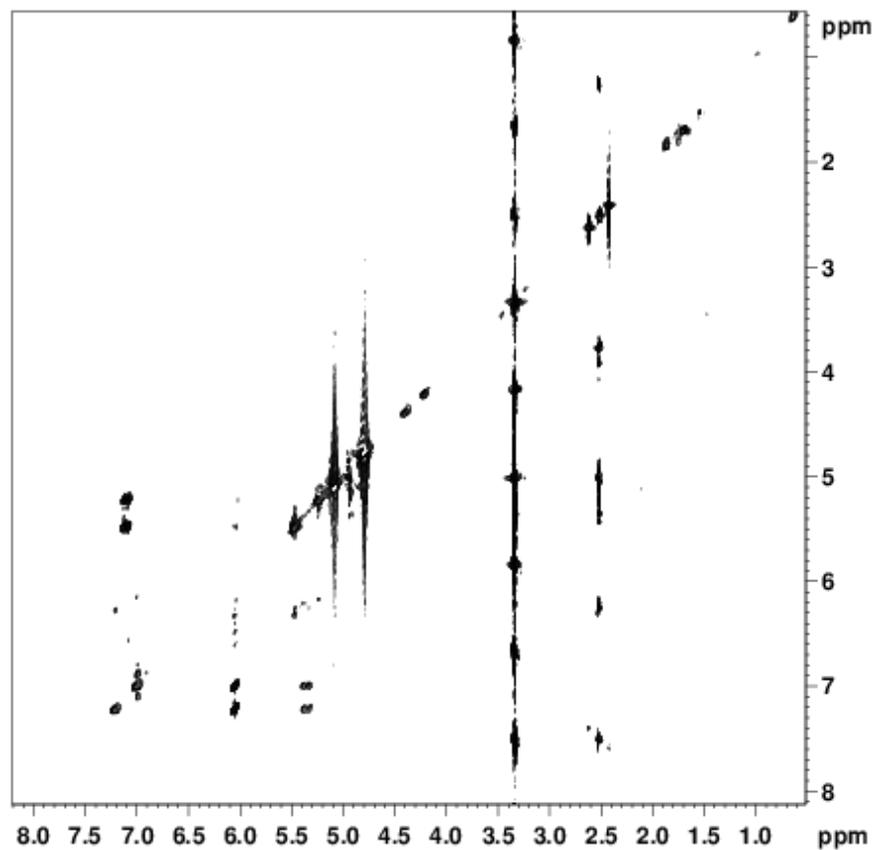
^1H - ^{13}C HETLOC,^{2,3} 300K, INEPT delay optimized for 145 Hz, TOCSY mixing 50ms, isotropic



^1H - ^{13}C CLIP-HSQC, 1 300K, INEPT delay optimized for 145 Hz, anisotropic



^1H - ^{13}C HETLOC, 2,3 300K, INEPT delay optimized for 145 Hz, TOCSY mixing 50ms, anisotropic



Cartesian coordinates of calculated structure models

Structure calculations were performed on a model system, where the alkyl chain on C5 was truncated after the third carbon atom. Using conventional force field methods (SYBYL, Tripos 7.0) several low energy conformers were found and further optimized by DFT methods (Gaussian 03,⁴ B3LYP/6-311+G**). The coordinates of these optimized geometries were then used as structure models in the SVD-based RDC fit with the software hotFCHT.^{5,6}

4R5R6R

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4r5r6r

C3	-0.470	-2.478	-1.139
H3	-0.949	-2.938	-1.995
C4	-0.713	-1.029	-0.804
H4	-0.526	-0.351	-1.639
C5	0.231	-0.739	0.413
C1	0.789	-2.153	0.774
C2	0.331	-3.097	-0.266
H2	0.627	-4.138	-0.280
O1	1.485	-2.391	1.733
O4	-2.103	-0.932	-0.443
C6	1.432	0.174	0.054
H6	1.988	0.341	0.981
C7	2.380	-0.368	-1.018
H7a	2.712	-1.364	-0.711
H7b	1.837	-0.490	-1.961
O6	0.963	1.458	-0.437
Oh	-0.514	-0.228	1.494
HOh	-0.032	0.507	1.905
C8	3.605	0.524	-1.237
H8a	3.316	1.518	-1.583
H8b	4.180	0.639	-0.313
H8c	4.267	0.086	-1.987
C6O	0.662	2.436	0.442
C6OC	0.242	3.691	-0.265
H6OCa	0.950	3.945	-1.056
H6OCb	-0.729	3.506	-0.730
H6OCc	0.162	4.505	0.453
O6CO	0.713	2.291	1.646
C4O	-2.682	0.295	-0.477
C4OC	-4.062	0.242	0.120
H4OCa	-4.611	-0.624	-0.251
H4OCb	-3.967	0.135	1.204
H4OCc	-4.596	1.162	-0.109
O4CO	-2.146	1.275	-0.931

4R5R6S

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4r5r6s

C3	-1.749	2.243	0.036
H3	-2.599	2.790	0.427
C4	-1.525	0.801	0.400
H4	-1.467	0.650	1.479
C5	-0.182	0.402	-0.326
C1	0.156	1.682	-1.158
C2	-0.824	2.729	-0.796
H2	-0.777	3.732	-1.202
O1	1.051	1.774	-1.963
O4	-2.651	0.055	-0.082
C6	0.927	0.092	0.716
H6	0.624	-0.820	1.239
C7	1.230	1.209	1.717
H7a	1.662	2.057	1.177
H7b	0.296	1.560	2.162
O6	2.198	-0.165	0.051
Oh	-0.383	-0.669	-1.214
HOh	0.103	-1.451	-0.902
C8	2.179	0.764	2.835
H8a	1.756	-0.070	3.403
H8b	3.143	0.447	2.432
H8c	2.361	1.585	3.532
C6O	2.484	-1.397	-0.404
C6OC	3.884	-1.464	-0.947
H6OCa	3.947	-0.830	-1.835
H6OCb	4.598	-1.079	-0.216
H6OCc	4.124	-2.492	-1.210
O6CO	1.698	-2.321	-0.389
C4O	-2.808	-1.192	0.434
C4OC	-3.934	-1.917	-0.248
H4OCa	-4.811	-1.272	-0.336
H4OCb	-3.616	-2.183	-1.259
H4OCc	-4.177	-2.820	0.307
O4CO	-2.117	-1.629	1.322

4R5S6R (OH5 and H6 antiperiplanar)

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4r5s6r antiperiplanar

C3	-2.565	0.143	-1.095
H3	-3.272	0.826	-1.553
C4	-1.089	0.465	-1.072
H4	-0.705	0.619	-2.080
C5	-0.417	-0.834	-0.499
C1	-1.647	-1.630	0.065
C2	-2.864	-1.013	-0.503
H2	-3.843	-1.455	-0.384
O1	-1.594	-2.585	0.800
O4	-0.832	1.760	-0.455
C6	0.686	-0.680	0.573
H6	0.193	-0.408	1.505
C7	1.518	-1.944	0.792
H7a	2.115	-2.155	-0.097
H7b	0.810	-2.769	0.896
O6	1.575	0.487	0.379
Oh	-0.022	-1.630	-1.609
HOh	0.820	-1.256	-1.927
C8	2.411	-1.861	2.034
H8a	1.817	-1.701	2.939
H8b	3.138	-1.048	1.963
H8c	2.967	-2.793	2.163
C6O	2.268	0.673	-0.746
C6OC	3.171	1.875	-0.650
H6OCa	2.572	2.774	-0.824
H6OCb	3.939	1.814	-1.418
H6OCc	3.618	1.955	0.341
O6CO	2.169	-0.014	-1.746
C4O	-1.165	1.995	0.837
C4OC	-0.882	3.428	1.206
H4OCa	0.199	3.579	1.252
H4OCb	-1.316	3.641	2.182
H4OCc	-1.282	4.109	0.453
O4CO	-1.605	1.156	1.582

4R5S6R (OH5 and H6 (+)-synclinal)

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4r5s6r plus synclinal			
C3	1.832	1.759	0.585
H3	2.858	1.878	0.912
C4	1.440	0.674	-0.383
H4	1.795	0.920	-1.385
C5	-0.117	0.693	-0.411
C1	-0.434	2.081	0.238
C2	0.804	2.557	0.894
H2	0.832	3.449	1.506
O1	-1.497	2.647	0.183
O4	1.965	-0.630	-0.035
C6	-0.751	-0.461	0.418
H6	-0.450	-1.401	-0.050
C7	-0.425	-0.487	1.912
H7a	-0.863	0.397	2.385
H7b	0.656	-0.415	2.034
O6	-2.201	-0.365	0.340
Oh	-0.592	0.698	-1.739
HOh	-0.951	-0.172	-1.968
C8	-0.935	-1.754	2.604
H8a	-0.486	-2.652	2.168
H8b	-2.020	-1.842	2.523
H8c	-0.681	-1.738	3.667
C6O	-2.853	-0.970	-0.669
C6OC	-4.341	-0.823	-0.519
H6OCa	-4.660	-1.146	0.473
H6OCb	-4.844	-1.404	-1.290
H6OCc	-4.602	0.233	-0.622
O6CO	-2.296	-1.544	-1.583
C4O	3.244	-0.900	-0.415
C4OC	3.613	-2.322	-0.085
H4OCa	3.388	-2.543	0.960
H4OCb	4.671	-2.477	-0.284
H4OCc	3.021	-3.004	-0.701
O4CO	3.959	-0.089	-0.947

4R5S6R (OH5 and H6 (-)-synclinal)

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4r5s6r minus synclinal

C3	-1.399	-1.100	-1.795
H3	-1.143	-1.423	-2.798
C4	-0.499	-1.406	-0.632
H4	-0.606	-2.460	-0.359
C5	-1.066	-0.576	0.558
C1	-2.489	-0.191	0.022
C2	-2.527	-0.487	-1.423
H2	-3.369	-0.227	-2.051
O1	-3.362	0.286	0.702
O4	0.871	-1.195	-1.015
C6	-0.383	0.771	0.975
H6	-1.167	1.249	1.569
C7	0.853	0.707	1.874
H7a	1.723	0.313	1.356
H7b	0.598	0.010	2.676
O6	-0.314	1.646	-0.194
Oh	-1.246	-1.377	1.711
HOh	-0.421	-1.865	1.849
C8	1.192	2.065	2.498
H8a	0.336	2.475	3.044
H8b	1.499	2.794	1.746
H8c	2.018	1.959	3.206
C6O	0.800	1.968	-0.890
C6OC	0.437	2.917	-2.008
H6OCa	-0.319	2.464	-2.653
H6OCb	1.329	3.154	-2.585
H6OCc	0.007	3.831	-1.593
O6CO	1.917	1.576	-0.660
C4O	1.801	-1.790	-0.230
C4OC	3.185	-1.585	-0.769
H4OCa	3.259	-2.005	-1.776
H4OCb	3.908	-2.063	-0.111
H4OCc	3.379	-0.513	-0.842
O4CO	1.509	-2.391	0.781

4S5R6R

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4s5r6r

C3	2.248	1.811	-0.001
H3	2.436	2.855	0.221
C4	0.881	1.354	-0.434
H4	0.624	1.732	-1.426
C5	0.992	-0.193	-0.524
C1	2.514	-0.454	-0.315
C2	3.153	0.826	0.037
H2	4.206	0.915	0.269
O1	3.034	-1.535	-0.473
O4	-0.088	1.857	0.506
C6	0.181	-1.037	0.487
H6	0.404	-2.080	0.244
C7	0.465	-0.803	1.971
H7a	1.545	-0.895	2.123
H7b	0.191	0.217	2.243
O6	-1.228	-0.817	0.239
Oh	0.639	-0.552	-1.853
HOh	0.789	-1.501	-1.967
C8	-0.256	-1.809	2.874
H8a	-1.340	-1.731	2.767
H8b	0.033	-2.837	2.636
H8c	-0.010	-1.627	3.923
C6O	-1.890	-1.703	-0.555
C6OC	-3.301	-1.240	-0.798
H6OCa	-3.793	-0.995	0.146
H6OCb	-3.277	-0.332	-1.406
H6OCc	-3.854	-2.018	-1.320
O6CO	-1.397	-2.711	-0.995
C4O	-1.272	2.324	0.016
C4OC	-2.177	2.736	1.145
H4OCa	-2.488	1.844	1.695
H4OCb	-1.648	3.387	1.843
H4OCc	-3.053	3.242	0.744
O4CO	-1.532	2.388	-1.157

RDC Fits with hotFCHT

Experimental RDCs are fitted using the SVD-based approach outlined by Losonczi et al.,⁷ as implemented in a self-written module of the software hotFCHT.^{5,6} Based on a proposed (rigid) geometry, an alignment tensor is calculated, and used to back-calculate the RDCs. Through a user controlled flag an automated normalization of RDCs before SVD can be performed within hotFCHT. Comparison of experimental and back-calculated RDCs give quality factors, showing the goodness-of-fit.⁸ Derived parameters describing the magnitude, eigenvalues and orientation of the alignment tensor⁹ are also given below.

The weighting factor is determined by normalizing the magnetogyric ratios and distances of each coupling to the respective values of the largest coupling. Each row of the coefficient matrix is multiplied by the weighting factor w_i of the respective coupling before SVD, and then renormalized after the alignment tensor is calculated.

$$w_i = \left| \frac{\max(\gamma_j \gamma_k / r^3)}{\gamma_{i,j} \gamma_{i,k} / r_i^3} \right| \quad (1)$$

Table SI 2: Fits of experimental RDCs to the calculated structure models without additional weighting.

RDC	w_i	D_{exp} (Hz)	Structure Models						
			D_{calc} (Hz)						
			4R5R6R	4R5R6S	4R5S6R	4R5S6R antiperip.	4R5S6R (+)-syncl.	4R5S6R (-)-syncl.	4S5R6R
C2 – H2	1.00	5.85	4.52	8.62	5.61	7.59	4.41	2.67	
C3 – H3	1.00	-15.02	-13.91	-15.04	-15.05	-15.14	-14.78	-14.63	
C4 – H4	1.00	4.93	5.97	4.86	4.93	4.10	4.74	3.35	
C6 – H6	1.00	44.39	16.97	26.60	43.31	27.50	35.58	29.96	
C4 – H3	1.00	0.52	-5.75	-3.31	0.41	-2.90	-3.33	-3.44	
C4 – H2	1.00	0.09	1.00	0.76	0.01	0.81	0.76	0.72	
C3 – H2	1.00	-0.88	8.09	5.14	-1.25	5.18	5.30	5.25	
C3 – H4	1.00	4.12	2.38	3.88	4.46	4.03	4.26	5.65	
Quality Factor		Q	0.62	0.41	0.03	0.39	0.24	0.35	
Condition Number		cond.	11.54	10.45	4.12	10.77	7.05	2.95	
Tensor Magnitude		GDO ($\cdot 10^{-3}$)	4.35	3.57	3.30	3.22	3.58	5.23	
Axial Component		D_a ($\cdot 10^{-3}$)	1.94	1.77	-1.56	1.59	1.78	2.57	
Rhombic Component		D_r ($\cdot 10^{-3}$)	1.13	0.24	-0.63	0.25	0.22	0.55	
Tensor Orientation		α ($^\circ$)	66.58	162.42	82.70	161.18	158.52	120.64	
–		β ($^\circ$)	33.22	51.76	59.61	97.79	50.57	116.88	
Euler Angles		γ ($^\circ$)	52.35	83.14	33.14	132.77	123.96	127.36	

Table SI 3: Fits of experimental RDCs to the calculated structure models with additional weighting according to eq. (1).

RDC	w_i	D_{exp} (Hz)	$4R5R6R$	$4R5R6S$	$4R5S6R$	$4R5S6R$	$4R5S6R$	$4S5R6R$
			D_{calc} (Hz)	D_{calc} (Hz)	D_{calc} (Hz)	antiperip.	(+)-syncl.	(-)-syncl.
C2 – H2	1.00	5.85	6.19	9.95	6.76	7.01	-0.03	0.23
C3 – H3	1.01	-15.02	-12.01	-15.17	-15.03	-15.58	-13.26	-13.45
C4 – H4	1.02	4.93	7.56	4.75	4.99	3.13	3.65	-0.20
C6 – H6	1.02	44.39	-20.75	2.29	38.67	1.62	-9.81	-0.93
C4 – H3	9.17	0.52	0.06	0.55	0.36	0.72	-0.01	0.00
C4 – H2	31.82	0.09	0.06	0.16	0.06	0.13	0.01	-0.01
C3 – H2	8.04	-0.88	-0.58	-0.63	-0.94	-0.56	-0.26	-0.60
C3 – H4	7.93	4.12	4.05	4.11	4.15	4.10	4.15	4.21
Quality Factor		Q	1.37	0.89	0.12	0.90	1.15	0.96
Condition Number		cond.	2.86	13.33	3.78	1.33	3.43	3.31
Tensor Magnitude		GDO ($\cdot 10^{-3}$)	3.26	2.76	2.93	3.41	4.24	4.22
Axial Component		D_a ($\cdot 10^{-3}$)	-1.51	-1.29	-1.38	1.51	1.84	-1.85
Rhombic Component		D_r ($\cdot 10^{-3}$)	-0.70	-0.57	-0.58	0.92	1.22	-1.17
Tensor Orientation		α ($^\circ$)	114.84	68.89	84.31	66.12	117.60	99.75
–		β ($^\circ$)	93.95	125.92	59.18	109.83	55.61	28.30
Euler Angles		γ ($^\circ$)	130.66	29.85	32.02	161.17	92.90	96.00

Back-Calculation of the C6-H6 RDC

To cross-validate the rigid model of the exocyclic C5-C6 bond, RDC fits were performed without the C6-H6 RDC for each conformer of the 4R5S5R relative configuration. The three conformers are denoted according to their OH5-C5-C6-H6 torsion: antiperiplanar, (+)-synclinal and (-)-synclinal. The resulting tensors were then used to back-calculate the C6-H6 RDC. In the absence of conformational flexibility this approach should give a value close to the experimental one, whereas conformational averaging would strongly reduce the observed RDC.

Table SI 4: Fits of experimental RDCs without the C6-H6 RDC with and without additional weighting before SVD. The resulting tensor is used to back-calculate the value for the C6-H6 RDC.

RDC	D_{exp} (Hz)	no weighting			with SVD weighting		
		<i>4R5S6R</i>	<i>4R5S6R</i>	<i>4R5S6R</i>	<i>4R5S6R</i>	<i>4R5S6R</i>	<i>4R5S6R</i>
		antiperip.	(+)-syncl.	(-)-syncl.	antiperip.	(+)-syncl.	(-)-syncl.
	D_{exp} (Hz)	D_{calc} (Hz)	D_{calc} (Hz)	D_{calc} (Hz)	D_{calc} (Hz)	D_{calc} (Hz)	D_{calc} (Hz)
C2 – H2	5.85	5.89	5.82	5.82	7.30	4.99	4.97
C3 – H3	-15.02	-15.02	-15.02	-15.02	-14.93	-15.15	-15.17
C4 – H4	4.93	4.93	4.93	4.93	4.94	5.16	5.25
C4 – H3	0.52	0.25	0.82	0.78	0.41	0.68	0.70
C4 – H2	0.09	0.04	0.07	0.08	0.08	0.06	0.06
C3 – H2	-0.88	-0.94	-0.80	-0.81	-0.90	-0.81	-0.80
C3 – H4	4.12	4.12	4.11	4.11	4.12	4.12	4.11
Condition Number	cond.	20.62	14.59	15.92	3.78	11.60	3.61
Tensor Magnitude	GDO ($\cdot 10^{-3}$)	2.99	3.62	3.58	2.85	3.41	3.69
Axial Component	D_a ($\cdot 10^{-3}$)	-1.40	1.59	1.45	-1.34	1.48	1.62
Rhombic Component	D_r ($\cdot 10^{-3}$)	-0.60	1.00	0.94	-0.56	0.97	1.03
Tensor Orientation	α ($^\circ$)	82.78	118.14	67.40	85.26	66.52	118.32
–	β ($^\circ$)	59.21	57.42	109.69	59.17	109.45	57.14
Euler Angles	γ ($^\circ$)	32.90	84.56	161.87	31.45	161.18	82.26
C6 – H6	44.39	39.26	-0.40	-20.61	37.73	-0.67	-19.94

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