

## Supporting Information for

# Application of Residual Dipolar Couplings in Structural Analysis of Thiocalix[4]arene Derivatives

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## NMR SECTION

### USED CHEMICALS

Chloroform- $d_1$  ( $\text{CDCl}_3$ ) – degree of deuteration min. 99.8 % and dimethylsulfoxide- $d_6$  (DMSO) – degree of deuteration min. 99.8 % were purchased from Merck, Germany. Poly( $\gamma$ -ethyl-L-glutamate) –  $M > 100\,000$  g/mol (medium A) and poly( $\gamma$ -benzyl-L-glutamate) –  $150\,000 < M < 350\,000$  g/mol (medium B) were purchased from Sigma-Aldrich.

### MEASUREMENTS CONDITIONS

All NMR data were acquired on Bruker Avance<sup>III</sup> 600 MHz (proton frequency) NMR spectrometer equipped with triple resonance cryo-probe. Scalar coupling constants ( $^1J_{C-H}$ ) and total splitting ( $^1T_{C-H}$ ) were measured using heteronuclear ( $^1\text{H}$ - $^{13}\text{C}$ ) CLIP-HSQC pulse sequence without proton decoupling in direct (F2) domain. Nine heteronuclear one bond RDCs were sufficient to obtain structural information, so it was unnecessary to measure any other type of RDCs.

#### 1D | $^1\text{H}$ NMR spectra

$\pi/2$  pulse for  $^1\text{H}$  nuclei was approximately 9  $\mu\text{s}$ . Spectral width: 7 kHz, size of fid: 32k data points, relaxation period: 10 s, number of scans: 8.

#### 1D | $^2\text{H}$ NMR spectra

$\pi/2$  pulse for  $^2\text{H}$  nuclei was approximately 68  $\mu\text{s}$ . Spectral width: 4 kHz, size of fid: 16k data points, relaxation period: 10 s, number of scans: 8.

#### 1D | $^{13}\text{C}$ -APT NMR spectra

$^{13}\text{C}$  NMR spectra with proton decoupling - *waltz16* (decoupling pulse 100  $\mu\text{s}$ , power level 23.6 dB).  $\pi/2$  pulse for  $^{13}\text{C}$  nuclei was approximately 12  $\mu\text{s}$ . Spectral width: 31.5 kHz, size of fid: 64k data points, polarization transfer: 7 ms, relaxation period: 2.0 s, number of scans: 20 000.

#### 1D | $^{13}\text{C}$ (de)coupled NMR spectra (Z-Restored)

Modified version of classical 1D  $^{13}\text{C}$  NMR experiment with straight baseline free of distortions. Spectral width: 19.5 kHz, size of fid: 64k data points, relaxation period: 1 s, number of scans: 4k - 16k.

For pulse sequence details see ref.: Y. Xia, S. Moran, E. P. Nikonorowicz, X. Gao, *Magn. Res. Chem.*, **2007**, *46*, 432.

#### 2D | $^1\text{H}$ - $^1\text{H}$ COSY spectra

Spectral width: 3.8 kHz in both domains, size of fid: 1024 (F2) a 256 (F1) data points, relaxation period: 1.0 s, number of scans: 12.

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## 2D | $^1\text{H}$ - $^{13}\text{C}$ HMQC spectra

Spectral width: 4.0 kHz (F2) a 21.4 kHz (F1), size of fid: 2048 (F2) a 256 (F1) data points, polarization transfer: 3.5 ms, relaxation period: 1.0 s, number of scans: 16.

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## 2D | CLIP $^1\text{H}$ - $^{13}\text{C}$ HSQC spectra

2D  $^1\text{H}$ - $^{13}\text{C}$  CLIP HSQC is a modification of classical F2-coupled HSQC experiment providing clear spectra without antiphase artefacts and other spectral distortions. Residual dipolar coupling constants were measured from splitting in the direct domain (F2). Spectral width: 4.7 kHz (F2) a 19.6 kHz (F1), size of fid: 8k (F2) a 256 (F1) data points, polarization transfer: 3,125 ms, relaxation period: 0.1 s, number of scans: 16.

For pulse sequence details see ref.: A. Enthart, J. C. Freudenberger, J. Furrer, H. Kessler, B. Luy, *J. Magn. Reson.*, **2008**, 192, 314.

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## 2D | $^1\text{H}$ - $^{13}\text{C}$ HMBC spectra

Spectral width: 4,0 kHz (F2) a 27,7 kHz (F1), size of fid: 2k (F2) a 256 (F1) data points, polarization transfer: 70 ms, relaxation period: 1,0 s, number of scans: 64.

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## 1D | $^1\text{H}$ DPFGSE – NOE spectra

Nuclear Overhauser effect (NOE) based experiment with selective inversion – DPFGSE sequence. Selective inversion was performed via 80ms *q3-gaussian cascade*. Spectral width: 12.0 kHz, size of fid: 32k data points, mixing time: 0.2-1.6 s, relaxation period: 2.0 s, number of scans: 128.

## NMR CHARACTERIZATION

FIGURE 1:  $^1\text{H}$  NMR SPECTRUM OF 5C (600 MHZ,  $\text{CDCl}_3$ )

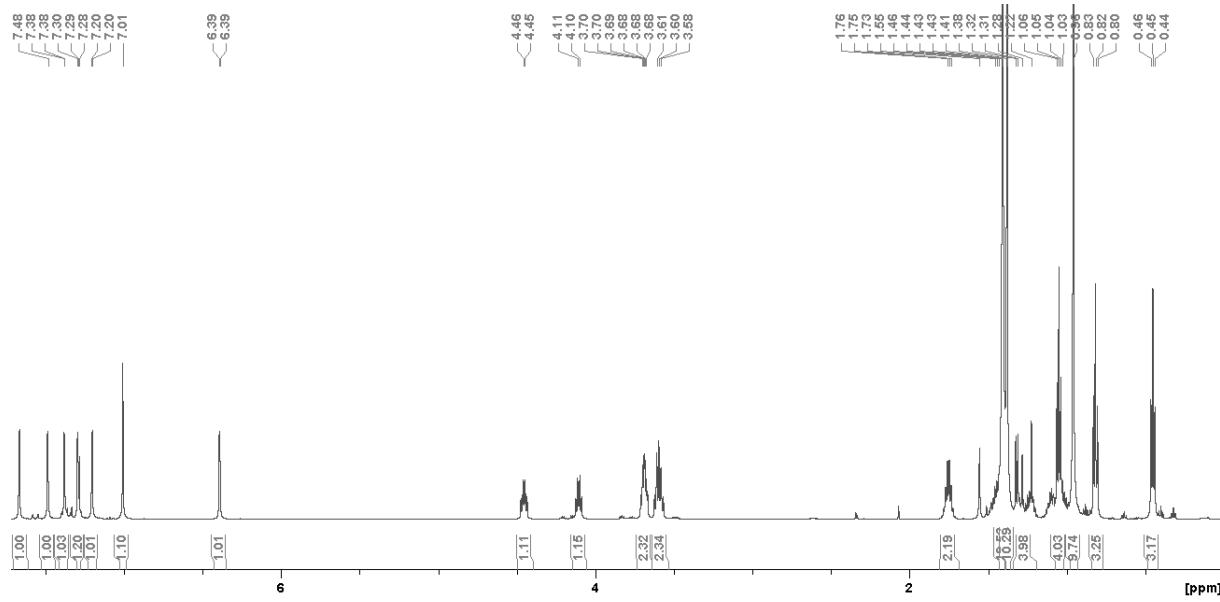


FIGURE 2:  $^{13}\text{C}$  NMR SPECTRUM OF 5C (600 MHZ,  $\text{CDCl}_3$ )

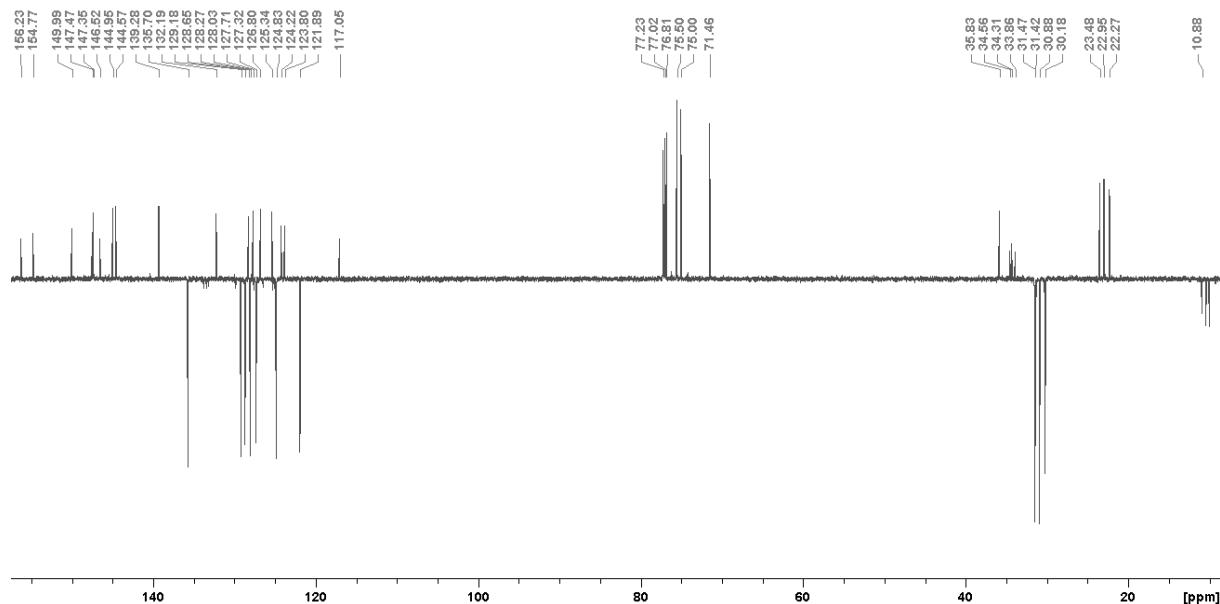


FIGURE 3:  $^1\text{H}$ - $^1\text{H}$  COSY SPECTRUM OF 5C (600 MHZ,  $\text{CDCl}_3$ )

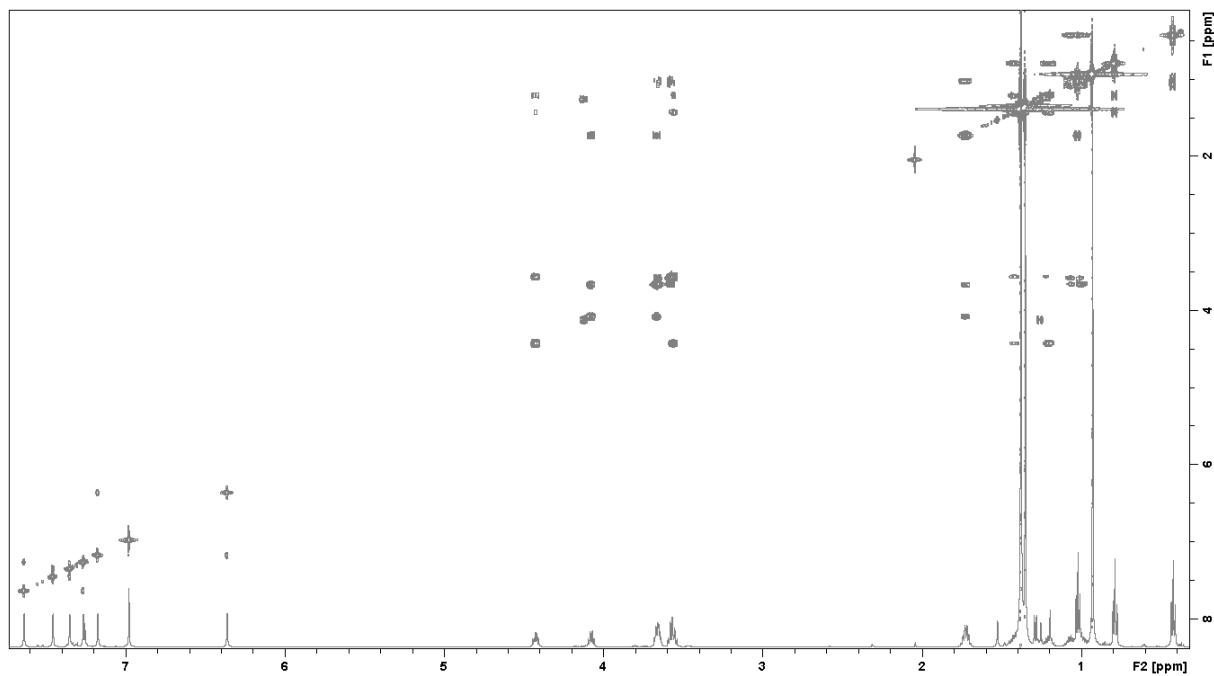


FIGURE 4:  $^1\text{H}$ - $^{13}\text{C}$  HMQC SPECTRUM OF 5C (600 MHZ,  $\text{CDCl}_3$ )

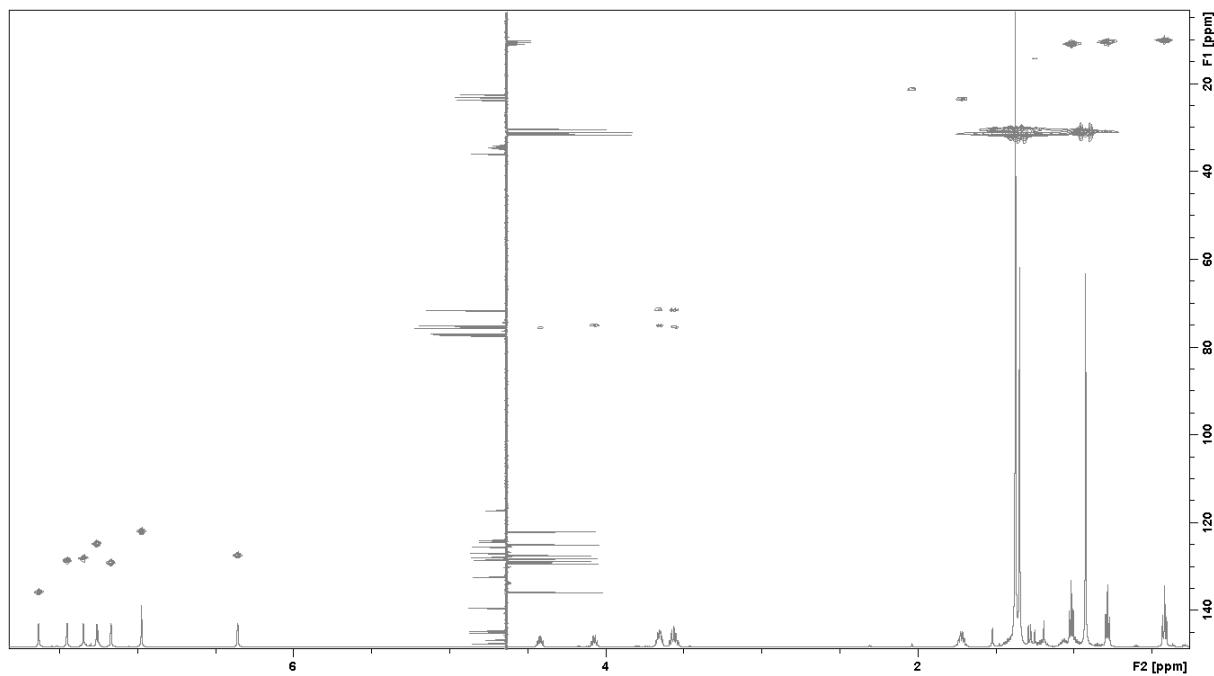


FIGURE 5:  $^1\text{H}$ - $^{13}\text{C}$  HMBC SPECTRUM OF 5C (600 MHZ,  $\text{CDCl}_3$ )

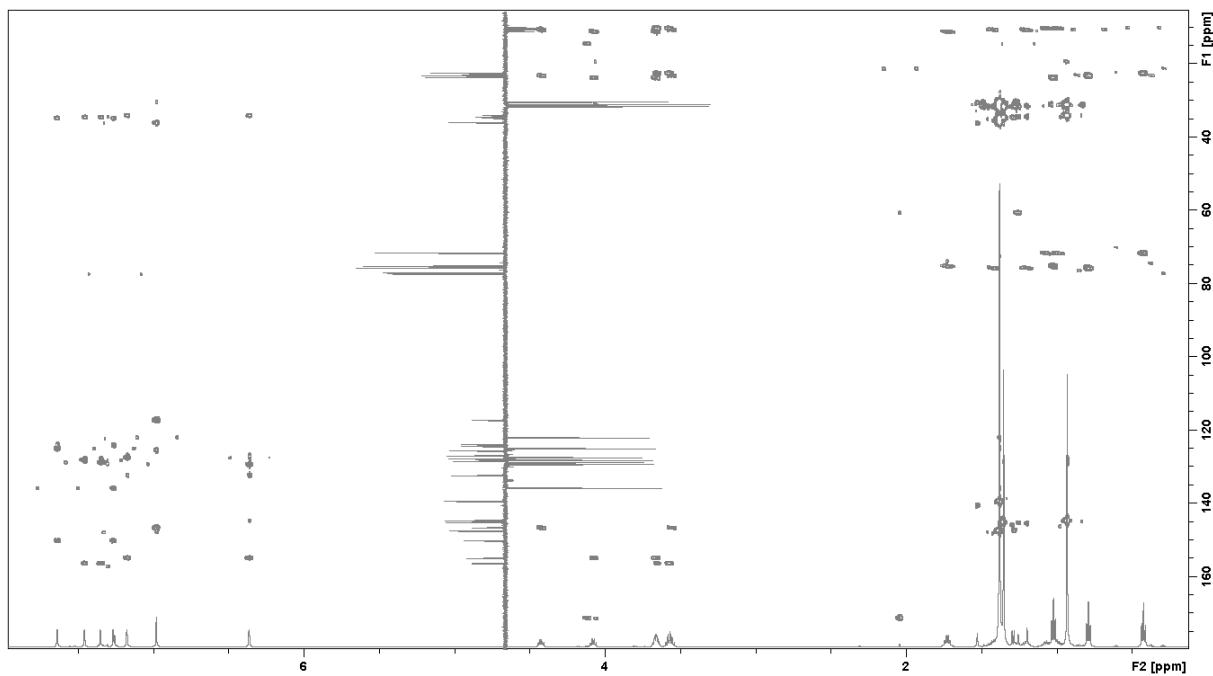
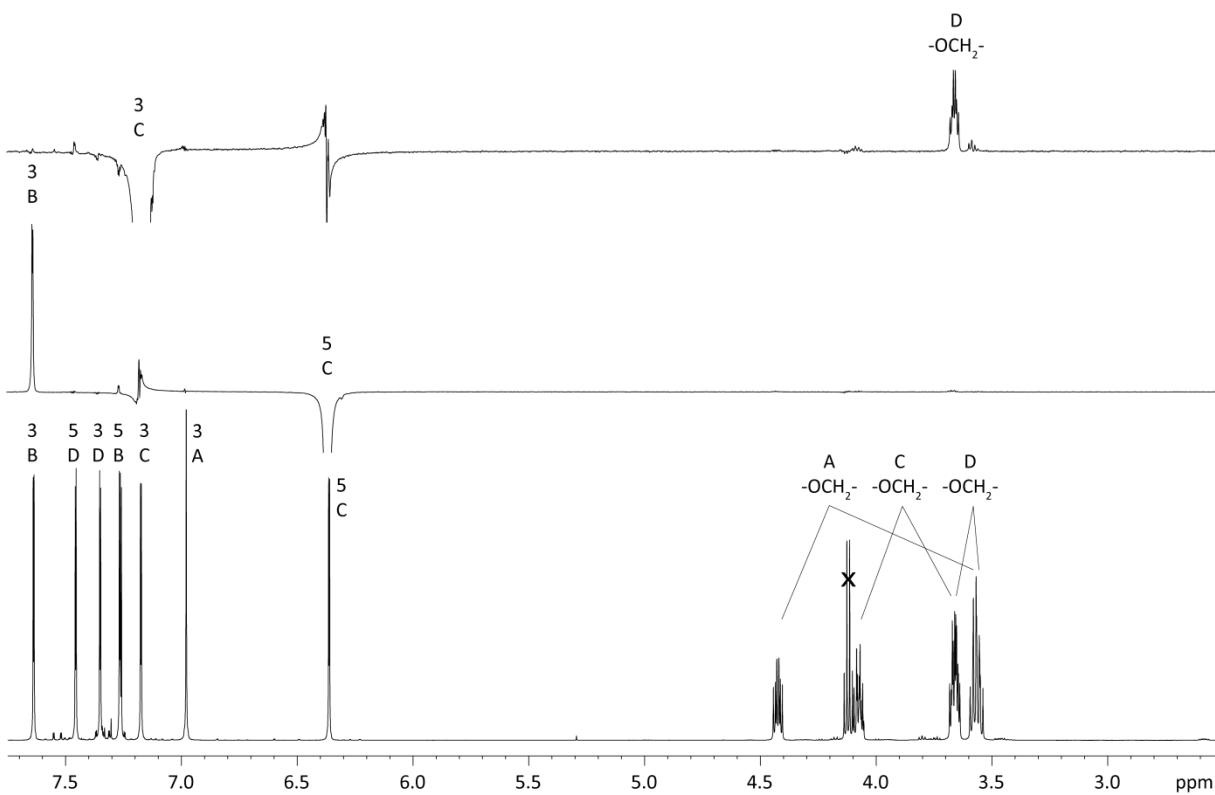
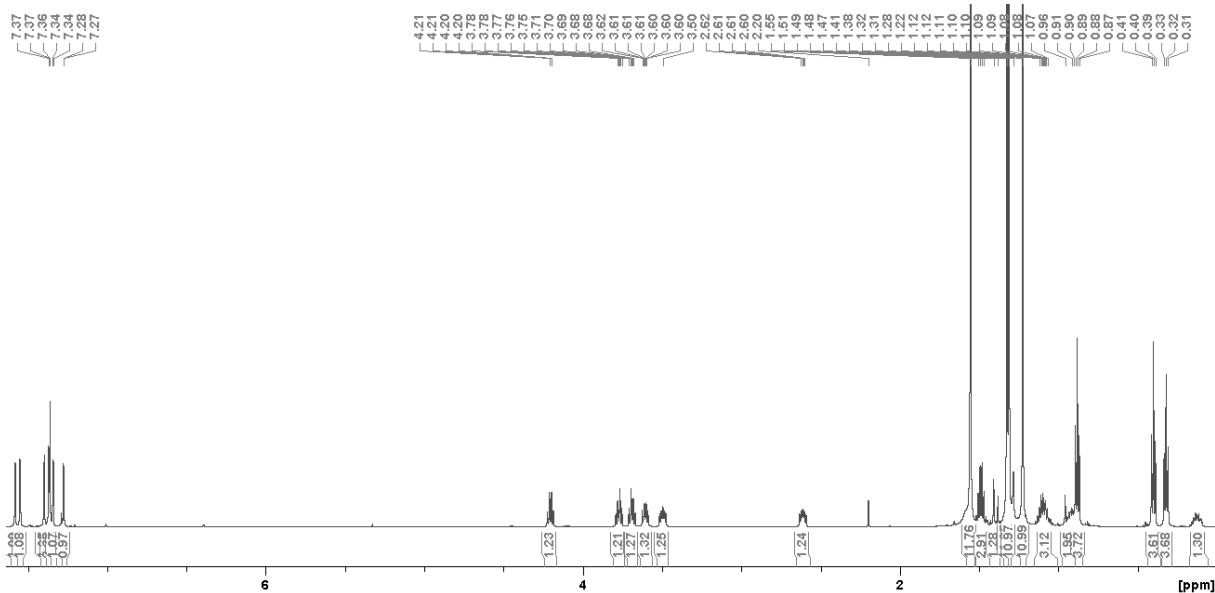


FIGURE 6: 1D  $^1\text{H}$ -DPFGSE NOE EXPERIMENTS OF 5C (600 MHZ,  $\text{CDCl}_3$ )



**FIGURE 7:  $^1\text{H}$  NMR SPECTRUM OF 5D (600 MHZ,  $\text{CDCl}_3$ )**



**FIGURE 8:  $^{13}\text{C}$  NMR SPECTRUM OF 5D (600 MHZ,  $\text{CDCl}_3$ )**

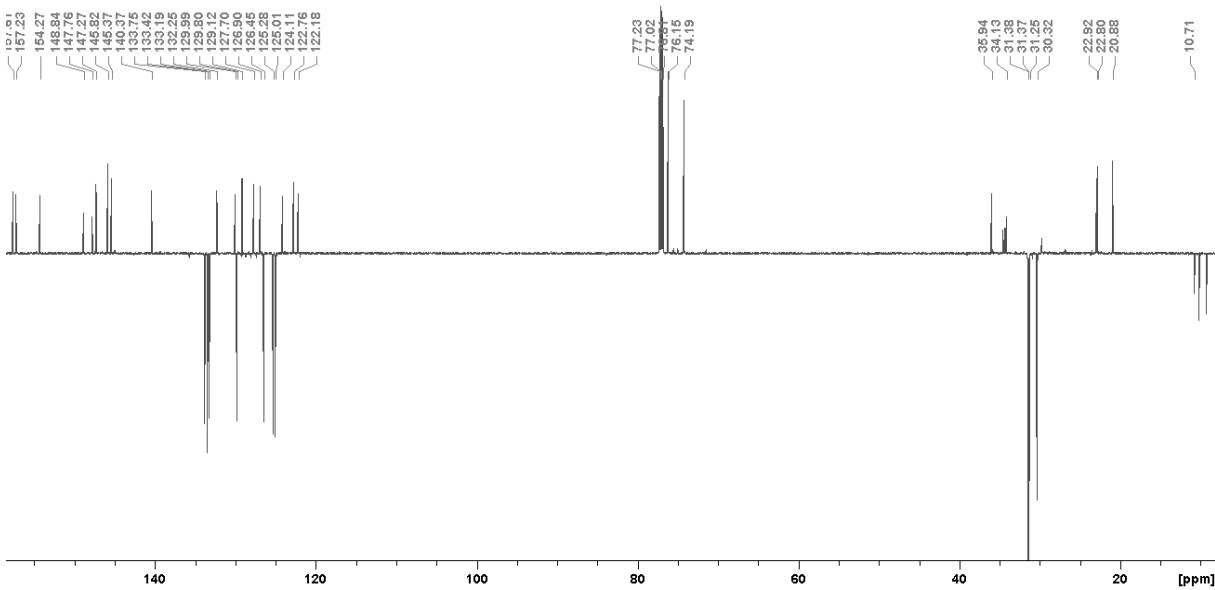


FIGURE 9:  $^1\text{H}$ - $^1\text{H}$  COSY SPECTRUM OF 5D (600 MHZ,  $\text{CDCl}_3$ )

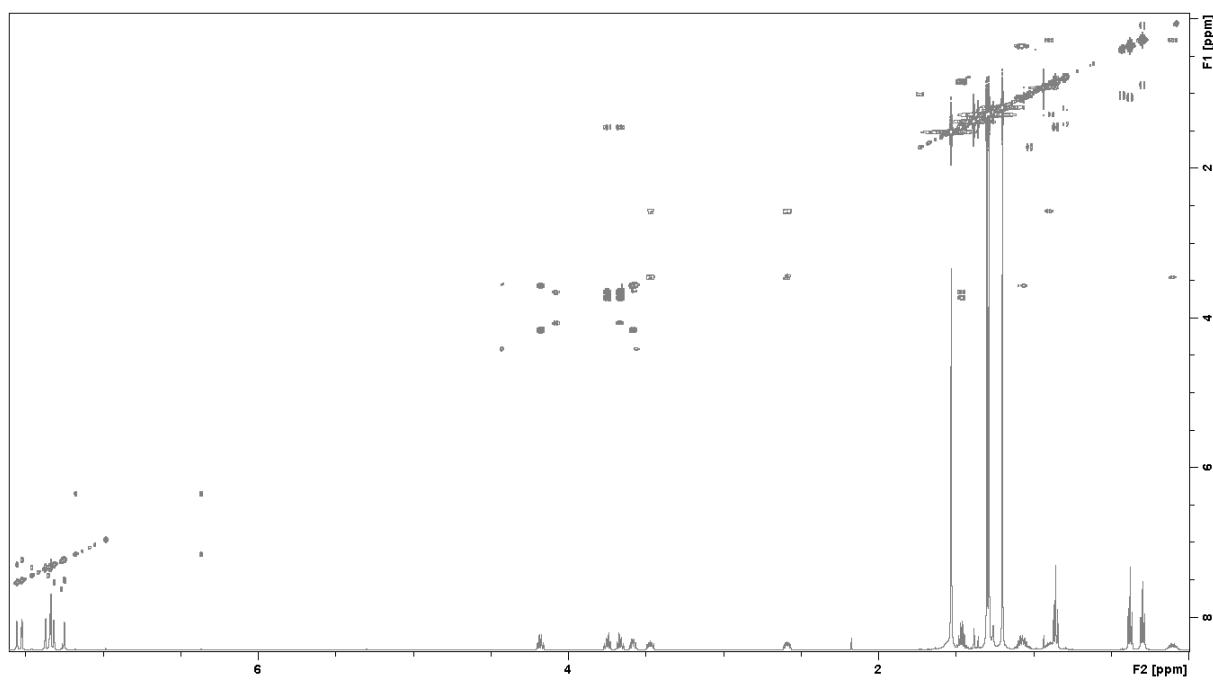


FIGURE 10:  $^1\text{H}$ - $^{13}\text{C}$  HMQC SPECTRUM OF 5D (600 MHZ,  $\text{CDCl}_3$ )

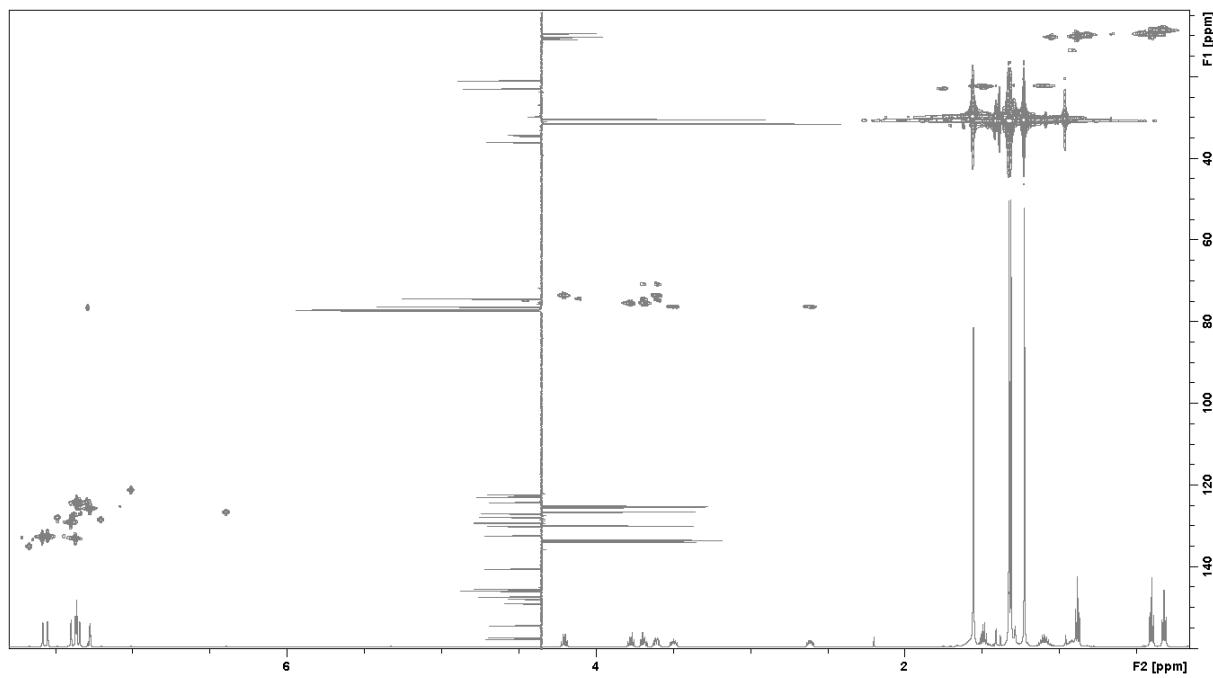


FIGURE 11:  $^1\text{H}$ - $^{13}\text{C}$  HMBC SPECTRUM OF 5D (600 MHZ,  $\text{CDCl}_3$ )

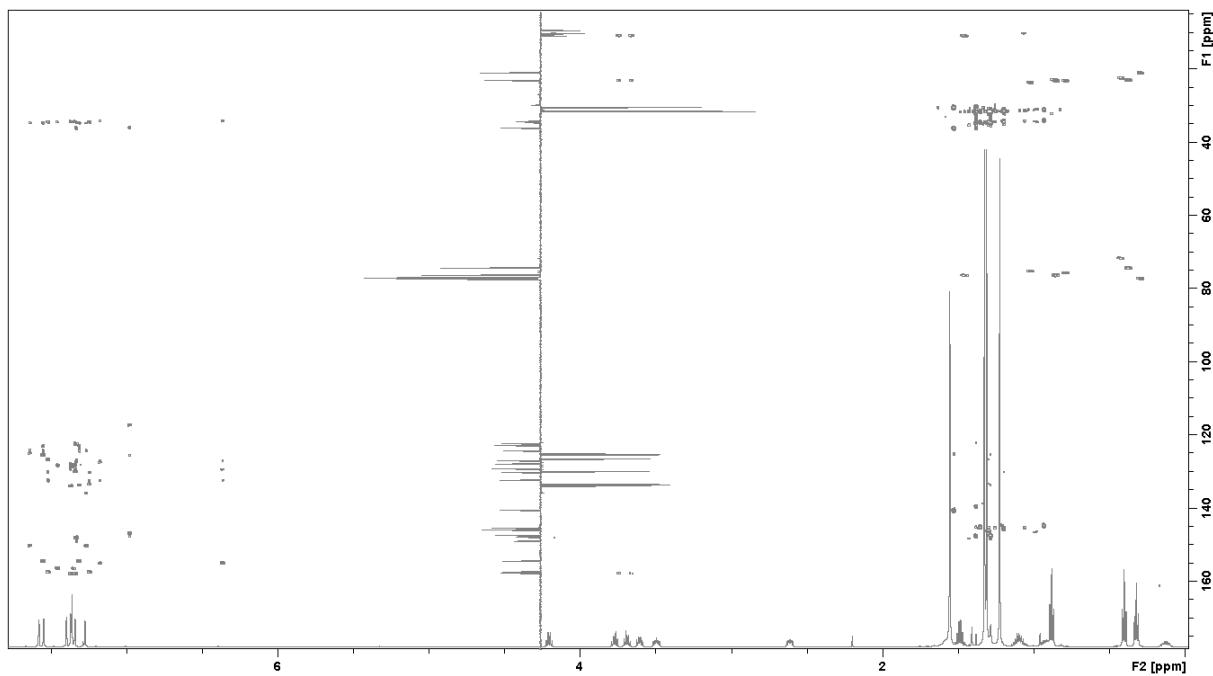


FIGURE 12: 1D  $^1\text{H}$ -DPFGSE NOE EXPERIMENTS OF 5D (600 MHZ,  $\text{CDCl}_3$ )

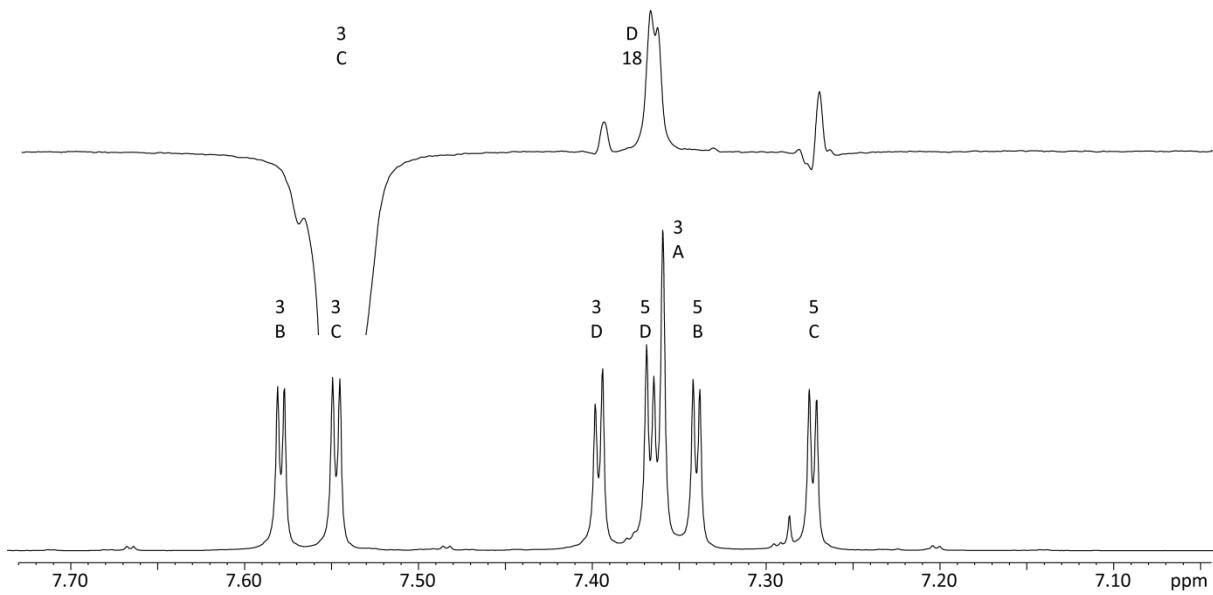


FIGURE 13:  $^1\text{H}$  NMR SPECTRUM OF 7C (600 MHZ,  $\text{CDCl}_3$ )

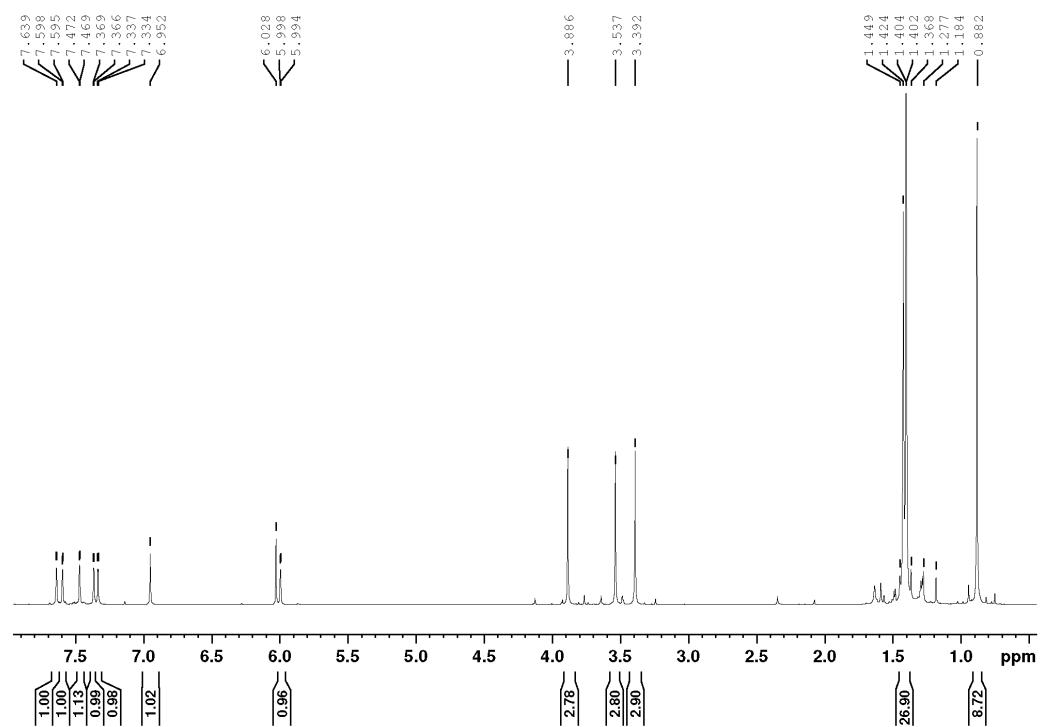


FIGURE 14:  $^{13}\text{C}$  NMR SPECTRUM OF 7C (600 MHZ,  $\text{CDCl}_3$ )

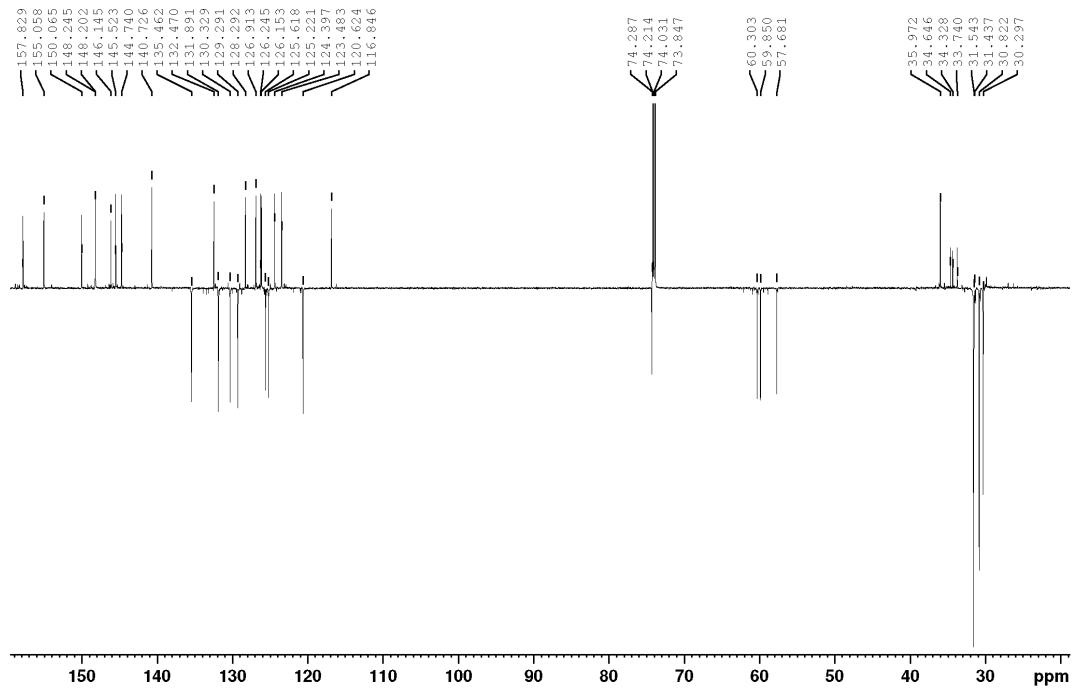


FIGURE 15:  $^1\text{H}$ - $^1\text{H}$  COSY SPECTRUM OF 7C (600 MHZ,  $\text{CDCl}_3$ )

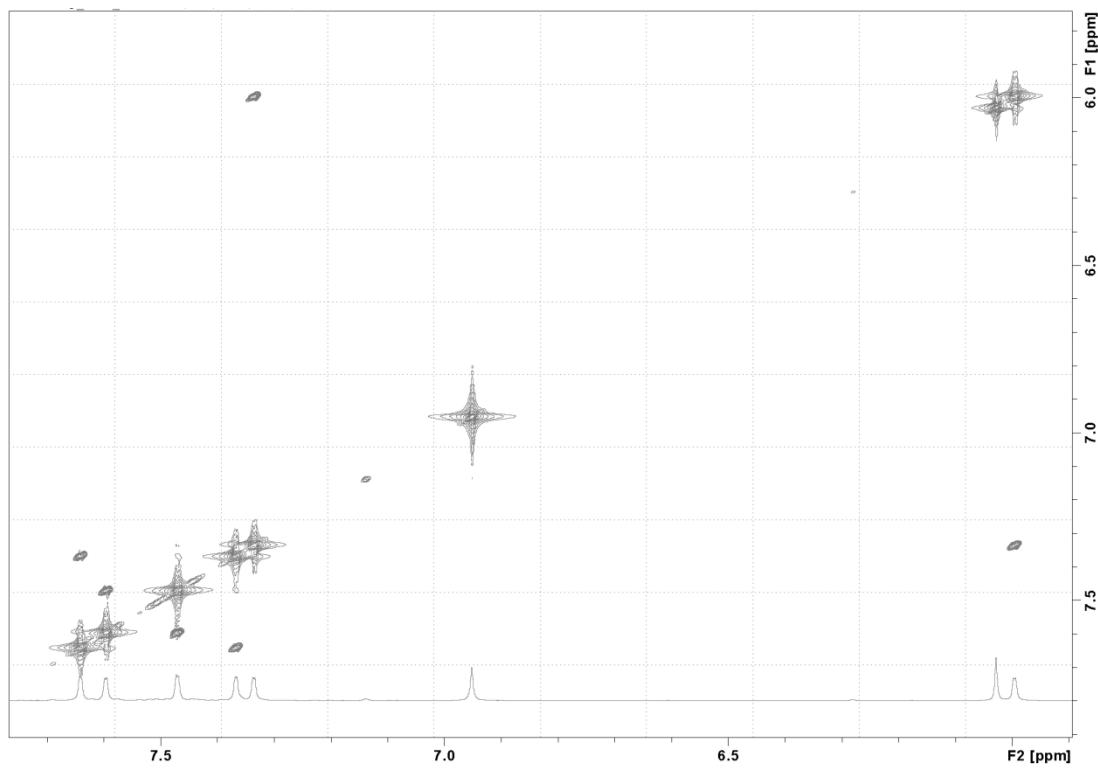


FIGURE 16:  $^1\text{H}$ - $^{13}\text{C}$  HMQC SPECTRUM OF 7C (600 MHZ,  $\text{CDCl}_3$ )

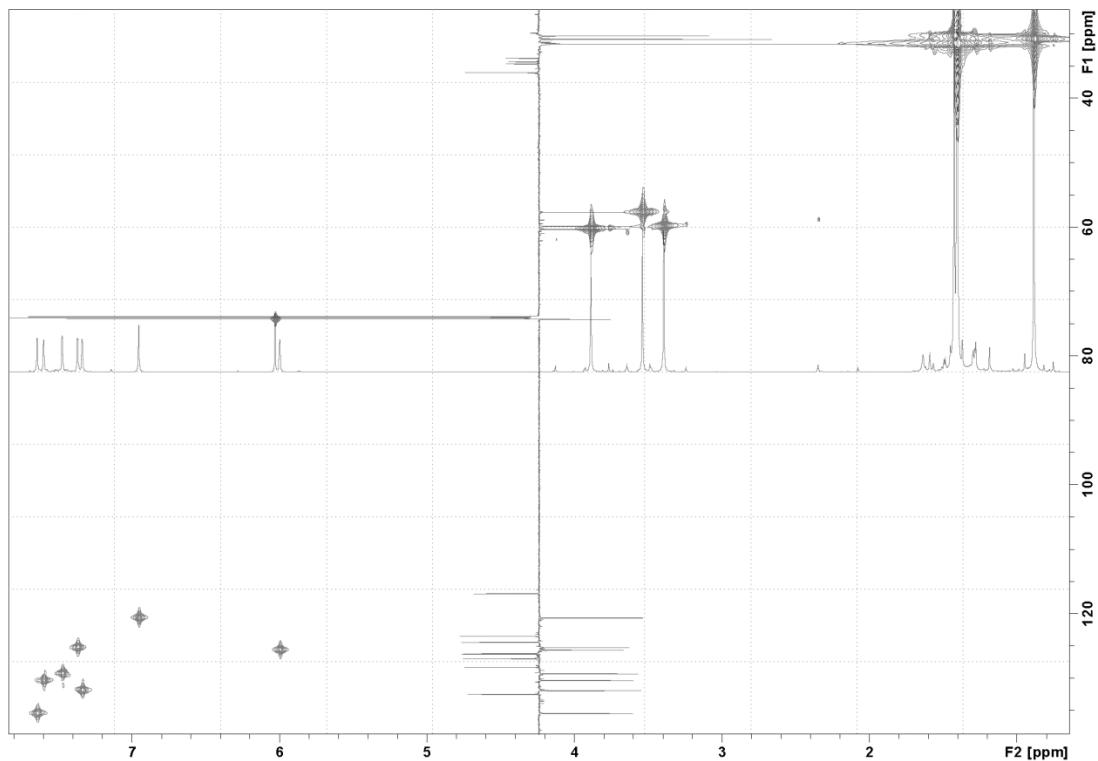


FIGURE 17:  $^1\text{H}$ - $^{13}\text{C}$  HMBC SPECTRUM OF 7C (600 MHZ,  $\text{CDCl}_3$ )

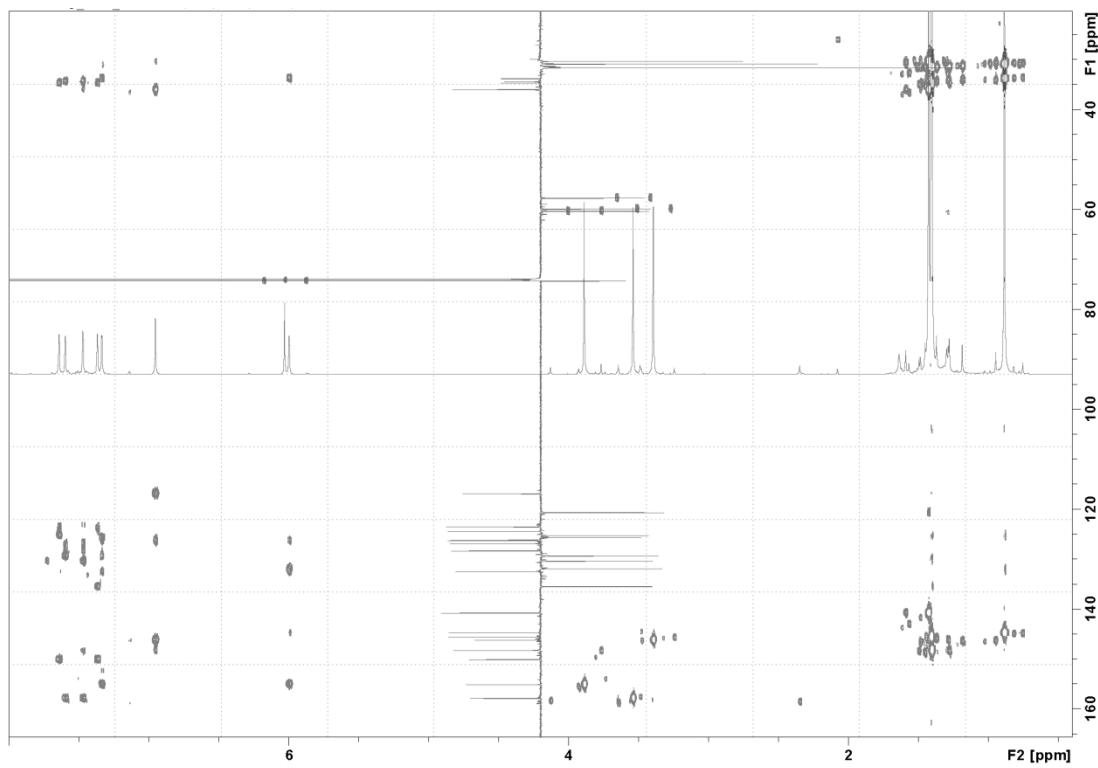


FIGURE 18: 1D  $^1\text{H}$ -DPFGSE NOE EXPERIMENTS OF 7C (600 MHZ,  $\text{CDCl}_3$ , 273 K)

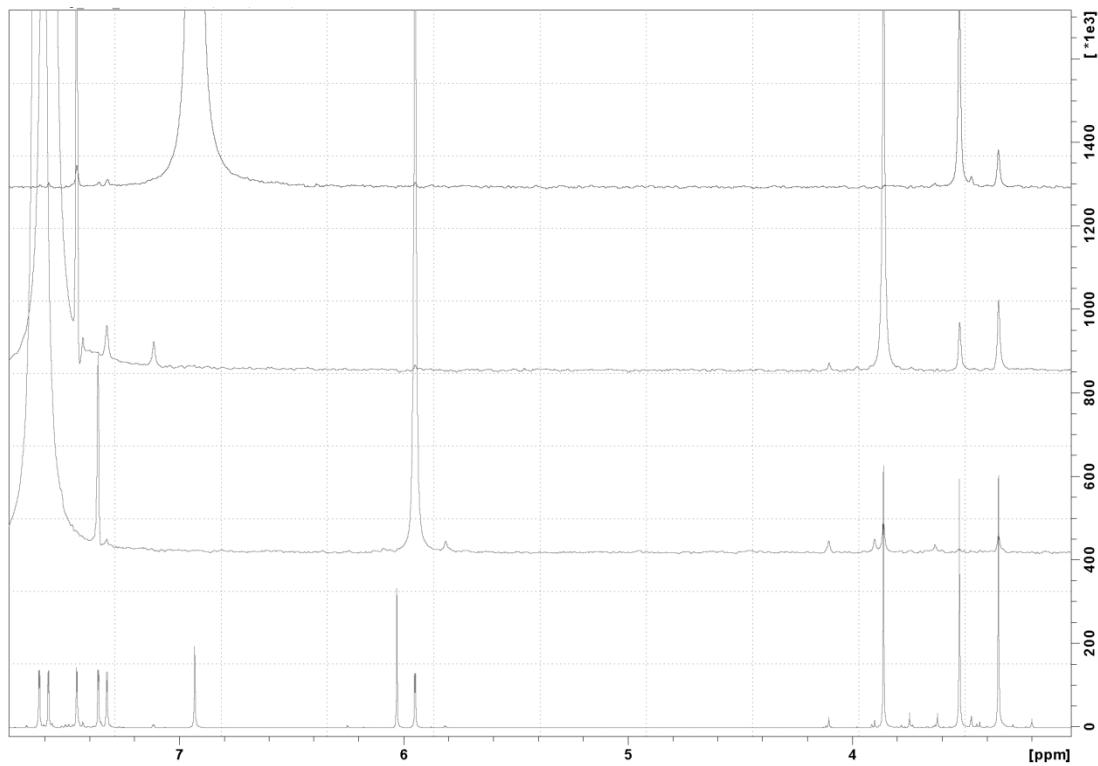


FIGURE 19:  $^1\text{H}$  NMR SPECTRUM OF 7D (600 MHZ,  $\text{CDCl}_3$ )

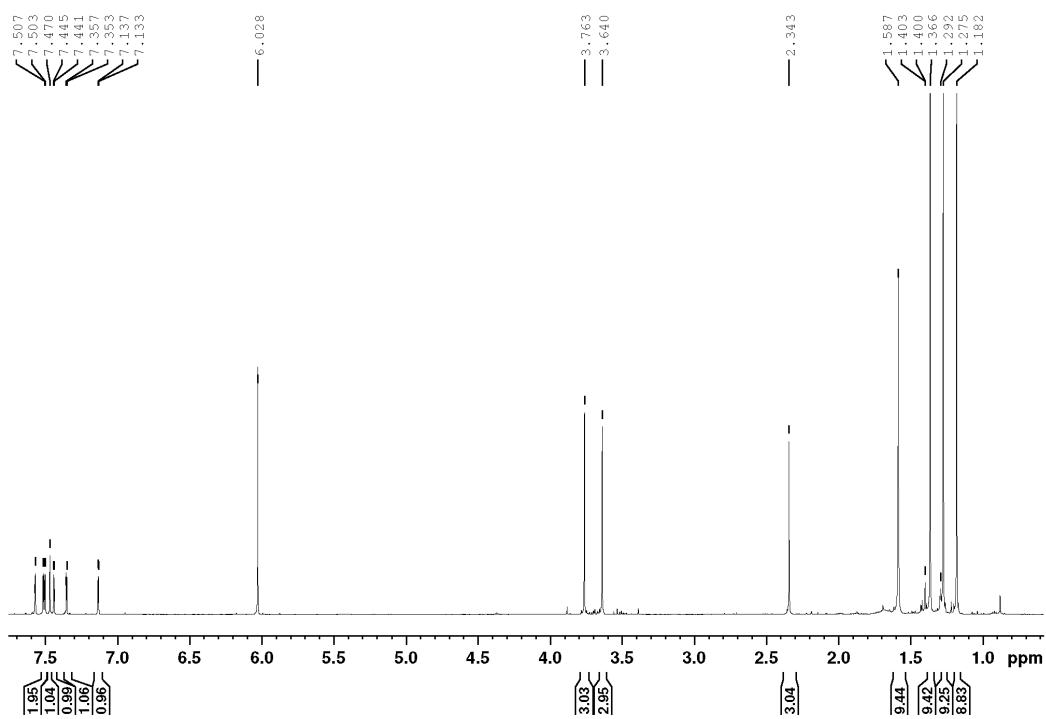


FIGURE 20:  $^{13}\text{C}$  NMR SPECTRUM OF 7D (600 MHZ,  $\text{CDCl}_3$ )

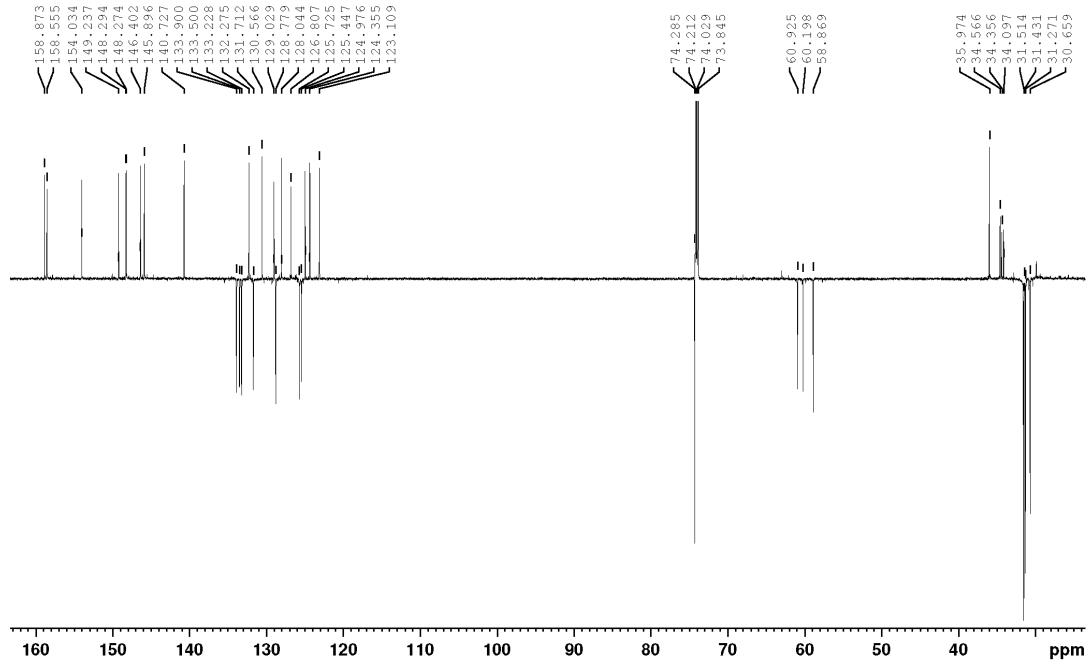


FIGURE 21:  $^1\text{H}$ - $^1\text{H}$  COSY SPECTRUM OF 7D (600 MHZ,  $\text{CDCl}_3$ )

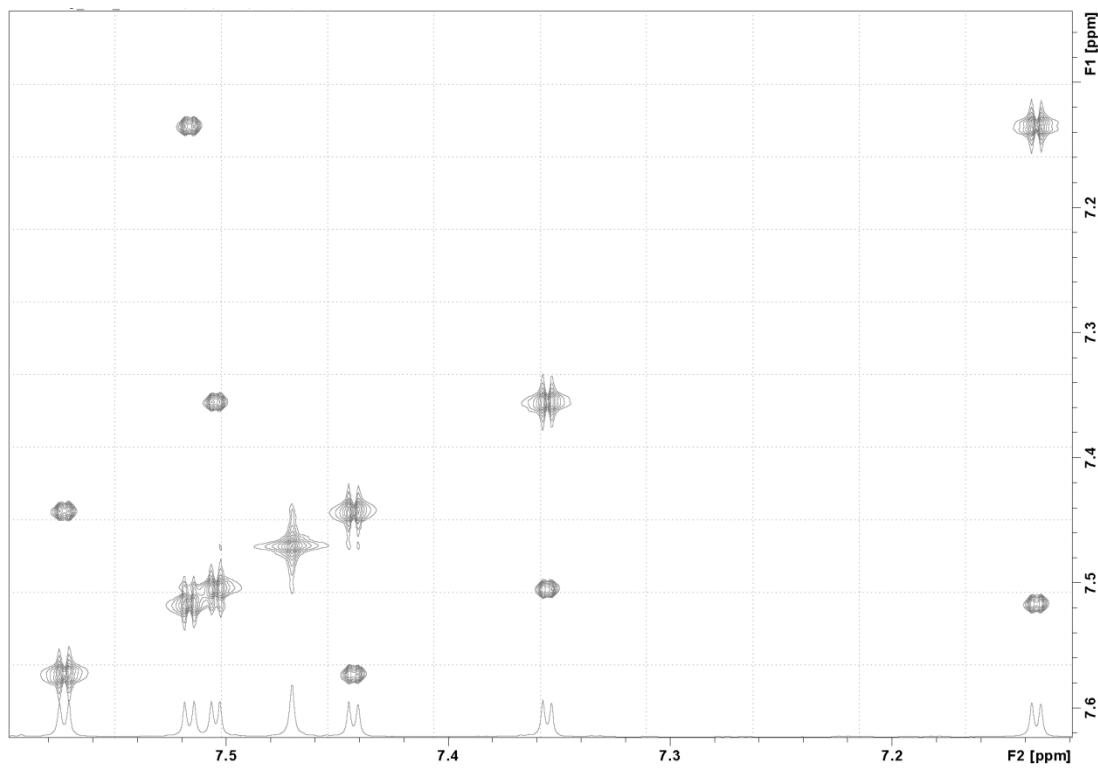


FIGURE 22:  $^1\text{H}$ - $^{13}\text{C}$  HMQC SPECTRUM OF 7D (600 MHZ,  $\text{CDCl}_3$ )

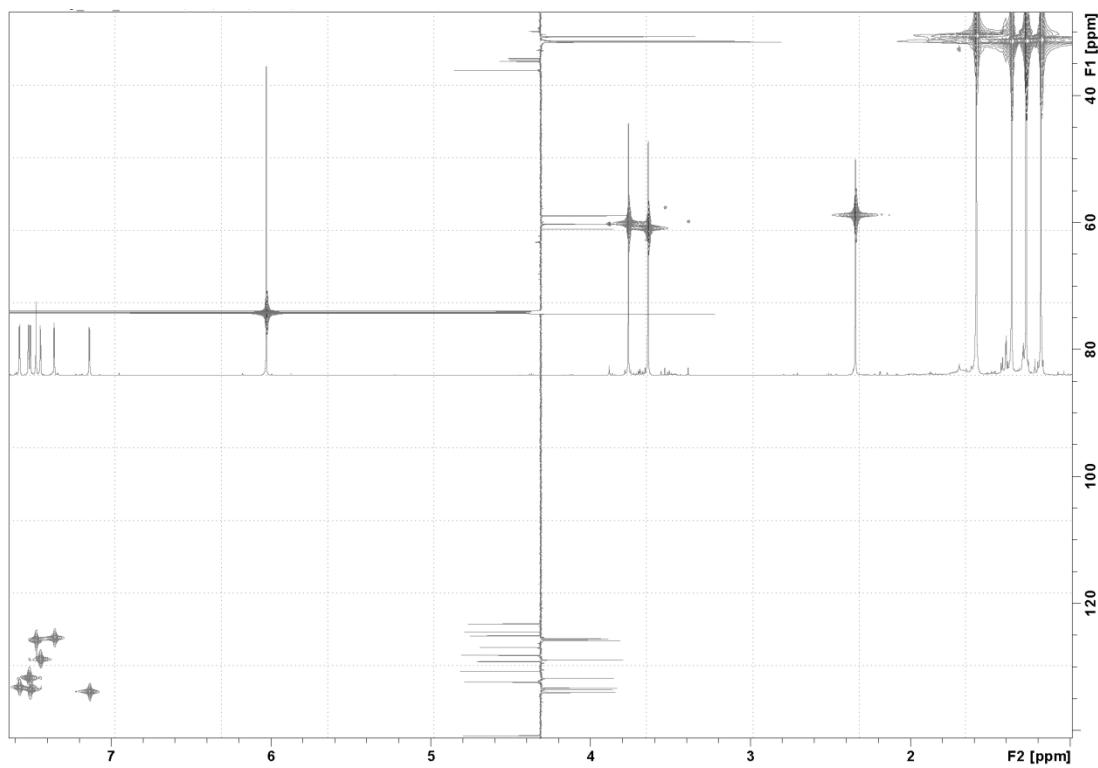


FIGURE 23:  $^1\text{H}$ - $^{13}\text{C}$  HMBC SPECTRUM OF 7D (600 MHZ,  $\text{CDCl}_3$ )

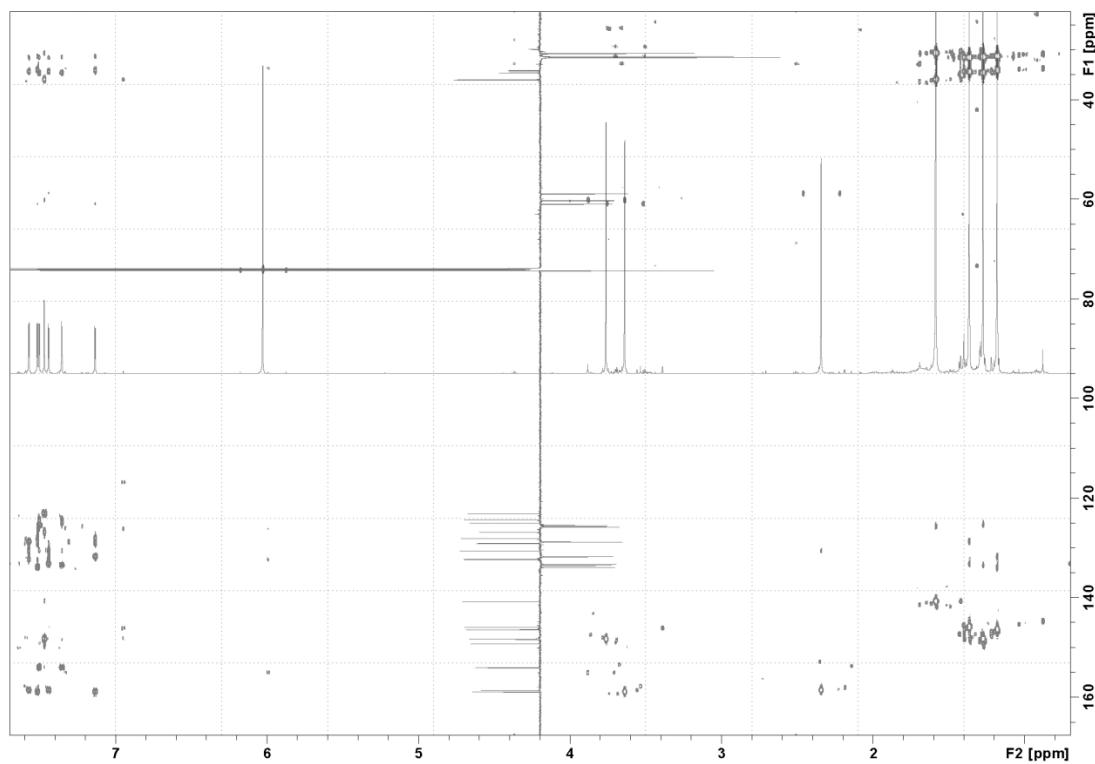
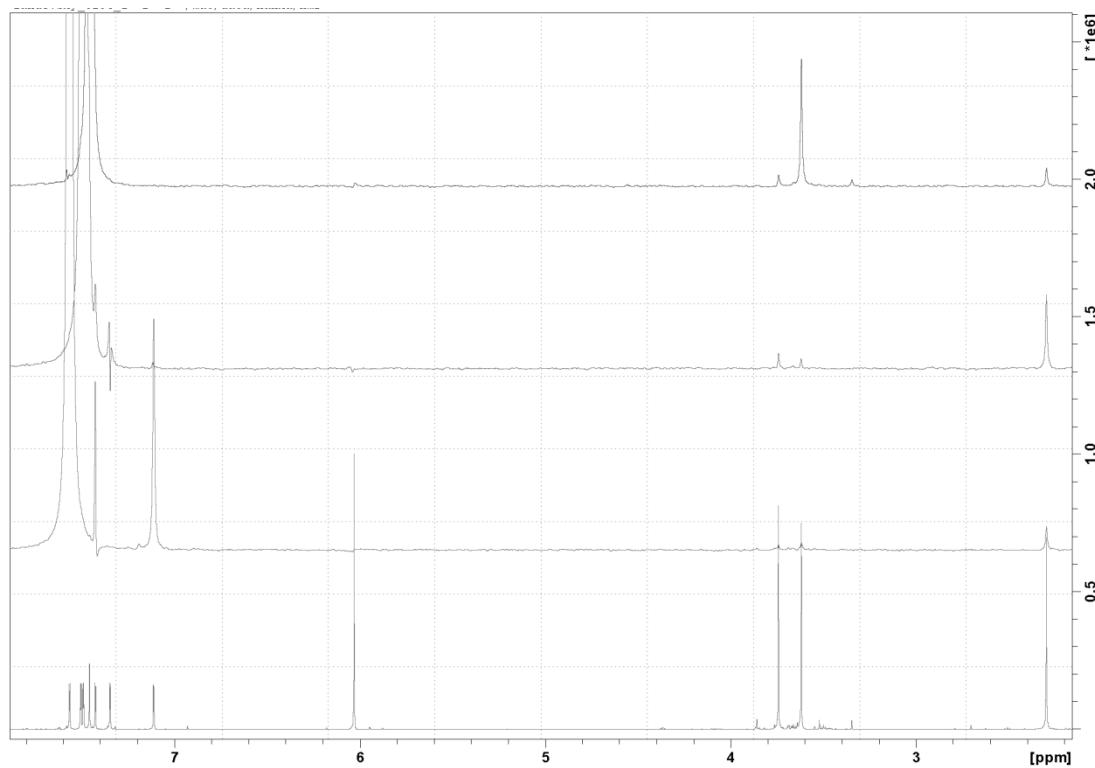


FIGURE 24: 1D  $^1\text{H}$ -DPFGSE NOE EXPERIMENTS OF 7D (600 MHZ,  $\text{CDCl}_3$ , 273 K)



## RDC SECTION

### STATISTICAL FORMULAS

$$x_{\text{rms}} = \sqrt{\frac{1}{n} (x_1^2 + x_2^2 + \dots + x_n^2)}$$

$$r_{xy} = \frac{\sum_{i=1}^n (x_i - \bar{x})(y_i - \bar{y})}{(n-1)s_x s_y} = \frac{\sum_{i=1}^n (x_i - \bar{x})(y_i - \bar{y})}{\sqrt{\sum_{i=1}^n (x_i - \bar{x})^2} \sqrt{\sum_{i=1}^n (y_i - \bar{y})^2}},$$

$$r_{xy} = \frac{\sum x_i y_i - n \bar{x} \bar{y}}{(n-1)s_x s_y} = \frac{n \sum x_i y_i - \sum x_i \sum y_i}{\sqrt{n \sum x_i^2 - (\sum x_i)^2} \sqrt{n \sum y_i^2 - (\sum y_i)^2}}.$$

5C (D-PACO) @ POLY( $\gamma$ -ETHYL-L-GLUTAMATE), MEDIUM A

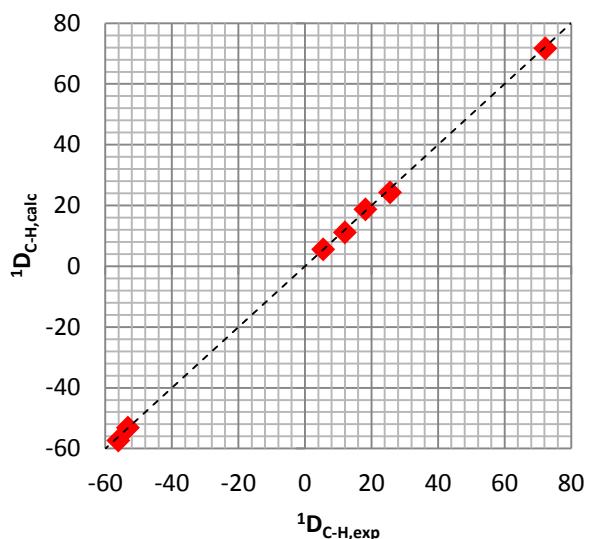
EVALUATION OF RDCs

	$^1J_{C-H}$ [Hz]	$^1T_{C-H}$ [Hz]	RDC [Hz]	$^1D_{C-H}$ [Hz]
CH-A3	159.8	184.0	24.1	12.1
CH-B8	160.3	48.4	-111.9	-56.0
CH-B10	161.1	197.4	36.3	18.2
CH-C13	157.0	50.8	-106.2	-53.1
CH-C15	159.7	304.4	144.7	72.3
CH-D18	159.4	170.4	11.1	5.5
CH-D20	158.0	209.3	51.3	25.6

FITTING RESULTS

Correlation between calculated  
and experimental RDCs

$^1D_{C-H}$	$^1D_{C-H,exp}$	$^1D_{C-H,calc}$
CH-A3	12.1	11.2
CH-B8	-56.0	-57.3
CH-B10	18.2	18.7
CH-C13	-53.1	-53.2
CH-C15	72.3	71.6
CH-D18	5.5	5.5
CH-D20	25.6	24.3



n =	7
RMS =	0,9
R(xy) =	1,000
$R^2$ =	99,98%

**PALES output file (fitting procedure results)**

REMARK Molecular Alignment Simulation.

REMARK Simulation parameters.

DATA PALES\_MODE DC

DATA TENSOR\_MODE SVD (Order Matrix Method)

REMARK Order matrix.

DATA SAUPE\_MATRIX S(zz) S(xx-yy) S(xy) S(xz) S(yz)  
DATA SAUPE 1.2979e-003 -2.1603e-003 3.3227e-005 -9.8188e-004 -6.7164e-004

DATA IRREDUCIBLE REP A0 A1R A1I A2R A2I  
DATA IRREDUCIBLE 2.0577e-003 1.2710e-003 -8.6938e-004 -1.3981e-003 -4.3009e-005  
DATA IRREDUCIBLE GENERAL\_MAGNITUDE 3.5902e-003

REMARK Mapping of coordinates.

DATA MAPPING\_COOR Szz\_d(x) Szz\_d(y) Syy\_d(x) Syy\_d(y) Sxx\_d(x) Sxx\_d(y)  
DATA MAPPING 0.06763 0.29927 -1.00494 1.07262 1.63742 0.38551  
DATA MAPPING INV -2.93433 -0.29927 0.49620 -1.07262 -1.27360 -0.38551

REMARK Eigensystem & Euler angles for clockwise rotation about z, y', z''.

DATA EIGENVALUES (Sxx\_d,Syy\_d,Szz\_d) 1.4665e-004 1.8838e-003 -2.0304e-003  
DATA EIGENVECTORS (x\_coor y\_coor z\_coor)  
DATA EIGENVECTORS X\_AXIS -1.8074e-001 9.0881e-001 3.7603e-001  
DATA EIGENVECTORS Y\_AXIS -2.4252e-001 -4.1170e-001 8.7845e-001  
DATA EIGENVECTORS Z\_AXIS 9.5316e-001 6.7575e-002 2.9482e-001

DATA Q\_EULER\_SOLUTIONS ALPHA BETA GAMMA  
DATA Q\_EULER\_ANGLES 1 66.83 72.85 175.94  
DATA Q\_EULER\_ANGLES 2 246.83 72.85 175.94  
DATA Q\_EULER\_ANGLES 3 113.17 107.15 355.94  
DATA Q\_EULER\_ANGLES 4 293.17 107.15 355.94

REMARK Euler angles (psi(theta/phi) for rotation about x, y, z.

DATA EULER\_SOLUTIONS 2  
DATA EULER\_ANGLES 12.91 -72.39 53.31  
DATA EULER\_ANGLES 192.91 252.39 233.31

DATA Da -1.015225e-003  
DATA Dr -5.790510e-004

DATA Aa -2.030449e-003  
DATA Ar -1.158102e-003

DATA Da\_HN -2.191383e+001  
DATA Rhombicity 5.703673e-001

REMARK Dipolar couplings.

DATA N 7  
DATA RMS 0.854

```

DATA Chi2          5.103
DATA CORR R       1.000
DATA Q SAUPE      0.012
DATA REGRESSION OFFSET -0.544 +/- 0.296 [Hz]
DATA REGRESSION SLOPE  1.001 +/- 0.007 [Hz]
DATA REGRESSION BAX SLOPE 1.001 +/- 0.005 [Hz]

VARS   RESID_I RESNAME_I ATOMNAME_I RESID_J RESNAME_J ATOMNAME_J DI D_OBS D D_DIFF
DD W
FORMAT %4d %4s %4s %4d %4s %4s %9.2f %9.3f %9.3f %9.3f %.2f %.2f

 1 CAL C7    1 CAL H6  47734.91  12.1000  11.1586  0.9414  1.0000  1.00
 1 CAL C5    1 CAL H1  47198.19 -56.0000 -57.3349  1.3349  1.0000  1.00
 1 CAL C6    1 CAL H5  47580.15  18.2000  18.7039 -0.5039  1.0000  1.00
 1 CAL C4    1 CAL H3  47355.66 -53.1000 -53.1602  0.0602  1.0000  1.00
 1 CAL C3    1 CAL H4  47434.36  72.3000  71.6497  0.6503  1.0000  1.00
 1 CAL C2    1 CAL H2  47449.07  5.5000   5.5168 -0.0168  1.0000  1.00
 1 CAL C1    1 CAL H7  47166.41  25.6000  24.2754  1.3246  1.0000  1.00

```

5B (C-PACO) @ POLY( $\gamma$ -ETHYL-L-GLUTAMATE), MEDIUM A

EVALUATION OF RDCs

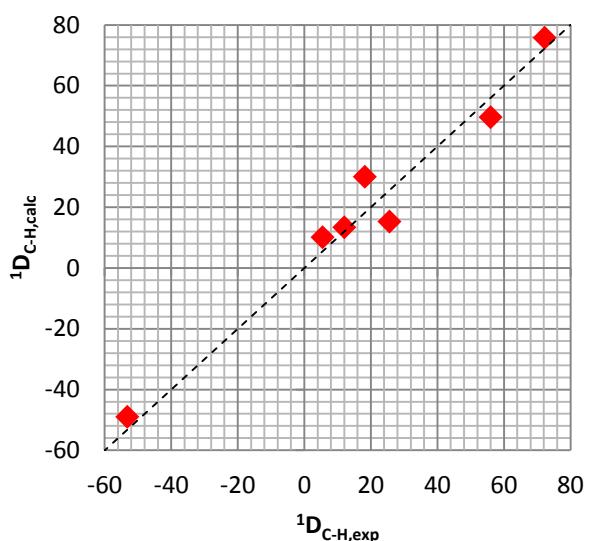
	$^1J_{C-H}$ [Hz]	$^1T_{C-H}$ [Hz]	RDC [Hz]	$^1D_{C-H}$ [Hz]
CH-A3	159.8	184.0	24.1	12.1
CH-B8	160.3	48.4	-111.9	-56.0
CH-B10	161.1	197.4	36.3	18.2
CH-C13	157.0	50.8	-106.2	-53.1
CH-C15	159.7	304.4	144.7	72.3
CH-D18	159.4	170.4	11.1	5.5
CH-D20	158.0	209.3	51.3	25.6

FITTING RESULTS

$^1D_{C-H}$	$^1D_{C-H,exp}$	$^1D_{C-H,calc}$
CH-A3	12.1	13.4
CH-B8	56.0	49.6
CH-B10	18.2	29.9
CH-C13	-53.1	-49.0
CH-C15	72.3	75.7
CH-D18	5.5	10.0
CH-D20	25.6	15.3

n =	7
RMS =	6,9
R(xy) =	0,983
R <sup>2</sup> =	96,67%

Correlation between calculated and experimental RDCs



5C (D-PACO) @ POLY( $\gamma$ -BENZYL-L-GLUTAMATE), MEDIUM B

EVALUATION OF RDCs

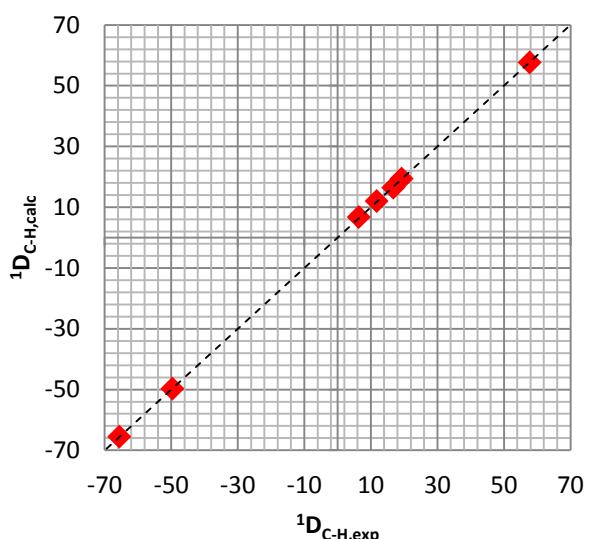
	$^1J_{C-H}$ [Hz]	$^1T_{C-H}$ [Hz]	RDC [Hz]	$^1D_{C-H}$ [Hz]
CH-A3	159.8	183.5	23.7	11.8
CH-B8	160.3	61.1	-99.2	-49.6
CH-B10	161.1	194.7	33.6	16.8
CH-C13	157.0	26.1	-130.9	-65.5
CH-C15	159.7	275.2	115.5	57.8
CH-D18	159.4	172.2	12.8	6.4
CH-D20	158.0	196.5	38.5	19.3

FITTING RESULTS

$^1D_{C-H}$	$^1D_{C-H,exp}$	$^1D_{C-H,calc}$
CH-A3	11.8	12.0
CH-B8	-49.6	-49.8
CH-B10	16.8	16.4
CH-C13	-65.5	-65.6
CH-C15	57.8	57.6
CH-D18	6.4	6.7
CH-D20	19.3	19.3

n =	7
RMS =	0.2
R(xy) =	1.000
R <sup>2</sup> =	100.00%

Correlation between calculated and experimental RDCs



**PALES output file (fitting procedure results)**

REMARK Molecular Alignment Simulation.

REMARK Simulation parameters.

DATA PALES\_MODE DC

DATA TENSOR\_MODE SVD (Order Matrix Method)

REMARK Order matrix.

DATA SAUPE\_MATRIX S(zz) S(xx-yy) S(xy) S(xz) S(yz)  
DATA SAUPE 1.0384e-003 -2.5764e-003 2.1279e-004 -4.7530e-004 -6.7312e-004

DATA IRREDUCIBLE REP A0 A1R A1I A2R A2I  
DATA IRREDUCIBLE 1.6462e-003 6.1523e-004 -8.7129e-004 -1.6675e-003 -2.7544e-004  
DATA IRREDUCIBLE GENERAL\_MAGNITUDE 3.2708e-003

REMARK Mapping of coordinates.

DATA MAPPING\_COOR Szz\_d(x) Szz\_d(y) Syy\_d(x) Syy\_d(y) Sxx\_d(x) Sxx\_d(y)  
DATA MAPPING -0.04085 0.15156 -1.13990 0.88494 1.30011 0.66248  
DATA MAPPING INV 3.06473 -0.15156 0.84977 -0.88494 -1.17694 -0.66248

REMARK Eigensystem & Euler angles for clockwise rotation about z, y', z''.

DATA EIGENVALUES (Sxx\_d,Syy\_d,Szz\_d) 2.2567e-004 1.6632e-003 -1.8889e-003  
DATA EIGENVECTORS (x\_coor y\_coor z\_coor)  
DATA EIGENVECTORS X\_AXIS -6.1522e-002 7.8607e-001 6.1507e-001  
DATA EIGENVECTORS Y\_AXIS -1.4380e-001 -6.1679e-001 7.7388e-001  
DATA EIGENVECTORS Z\_AXIS 9.8769e-001 -4.0837e-002 1.5098e-001

DATA Q\_EULER\_SOLUTIONS ALPHA BETA GAMMA  
DATA Q\_EULER\_ANGLES 1 51.52 81.32 182.37  
DATA Q\_EULER\_ANGLES 2 231.52 81.32 182.37  
DATA Q\_EULER\_ANGLES 3 128.48 98.68 2.37  
DATA Q\_EULER\_ANGLES 4 308.48 98.68 2.37

REMARK Euler angles (psi(theta/phi) for rotation about x, y, z.

DATA EULER\_SOLUTIONS 2  
DATA EULER\_ANGLES -15.13 -81.00 66.84  
DATA EULER\_ANGLES 164.87 261.00 246.84

DATA Da -9.444267e-004  
DATA Dr -4.791692e-004

DATA Aa -1.888853e-003  
DATA Ar -9.583384e-004

DATA Da\_HN -2.038564e+001  
DATA Rhombicity 5.073652e-001

REMARK Dipolar couplings.

DATA N 7  
DATA RMS 0.221  
DATA Chi2 0.343

```

DATA CORR R          1.000
DATA Q SAUPE        0.004
DATA REGRESSION OFFSET -0.052 +/- 0.096 [Hz]
DATA REGRESSION SLOPE 1.000 +/- 0.002 [Hz]
DATA REGRESSION BAX SLOPE 1.000 +/- 0.002 [Hz]

VARS    RESID_I RESNAME_I ATOMNAME_I RESID_J RESNAME_J ATOMNAME_J DI D_OBS D D_DIFF
DD W
FORMAT %4d %4s %4s %4d %4s %4s %9.2f %9.3f %9.3f %9.3f %.2f %.2f

1 CAL C1      1 CAL H7   47758.57   11.8000   12.0062   -0.2062   1.0000 1.00
1 CAL C29     1 CAL H33  47176.79   -49.6000  -49.7862   0.1862   1.0000 1.00
1 CAL C27     1 CAL H54  47536.98   16.8000   16.4300   0.3700   1.0000 1.00
1 CAL C83     1 CAL H87  47668.52   -65.5000  -65.5755   0.0755   1.0000 1.00
1 CAL C81     1 CAL H85  47711.00   57.8000   57.5835   0.2165   1.0000 1.00
1 CAL C95     1 CAL H97  47612.93   6.4000    6.6769   -0.2769   1.0000 1.00
1 CAL C93     1 CAL H96  47258.25   19.3000   19.3005   -0.0005   1.0000 1.00

```

5B (C-PACO) @ POLY( $\gamma$ -BENZYL-L-GLUTAMATE), MEDIUM B

EVALUATION OF RDCs

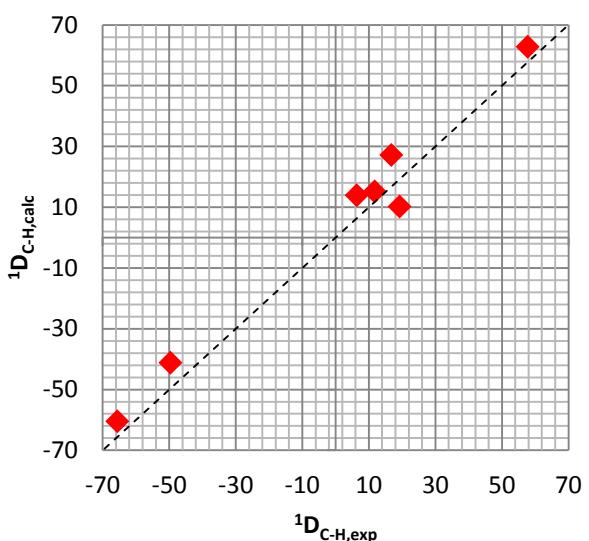
	$^1J_{C-H}$ [Hz]	$^1T_{C-H}$ [Hz]	RDC [Hz]	$^1D_{C-H}$ [Hz]
CH-A3	159.8	183.5	23.7	11.8
CH-B8	160.3	61.1	-99.2	-49.6
CH-B10	161.1	194.7	33.6	16.8
CH-C13	157.0	26.1	-130.9	-65.5
CH-C15	159.7	275.2	115.5	57.8
CH-D18	159.4	172.2	12.8	6.4
CH-D20	158.0	196.5	38.5	19.3

FITTING RESULTS

$^1D_{C-H}$	$^1D_{C-H,exp}$	$^1D_{C-H,calc}$
CH-A3	11.8	15.1
CH-B8	-49.6	-41.2
CH-B10	16.8	27.1
CH-C13	-65.5	-60.5
CH-C15	57.8	62.7
CH-D18	6.4	13.8
CH-D20	19.3	10.2

n =	7
RMS =	7.3
R(xy) =	0.989
R <sup>2</sup> =	97.78%

Correlation between calculated and experimental RDCs



5D (1,2-ALT) @ POLY( $\gamma$ -ETHYL-L-GLUTAMATE), MEDIUM A

EVALUATION OF RDCs

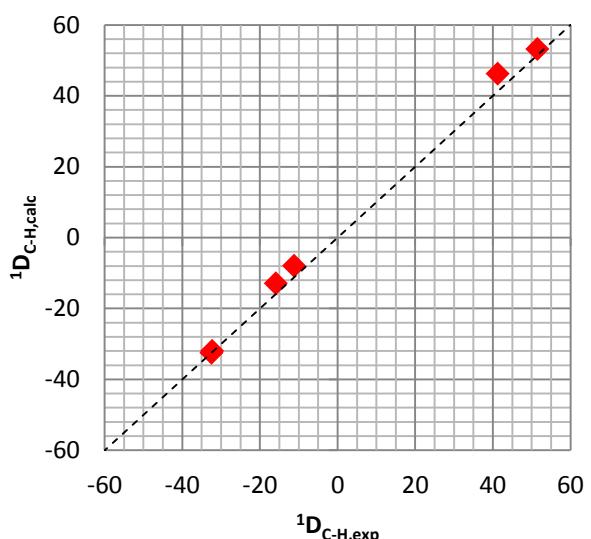
	$^1J_{C-H}$ [Hz]	$^1T_{C-H}$ [Hz]	RDC [Hz]	$^1D_{C-H}$ [Hz]
CH-A3	160.4	138.0	-22.4	-11.2
CH-B8	159.3	32.0	-127.3	-63.6
CH-B10	160.8	96.0	-64.8	-32.4
CH-C13	157.0	260.0	103.0	51.5
CH-C15	159.8	128.0	-31.8	-15.9
CH-D18	158.6	82.0	-76.6	-38.3
CH-D20	158.3	241.0	82.7	41.3

FITTING RESULTS

$^1D_{C-H}$	$^1D_{C-H,exp}$	$^1D_{C-H,calc}$
CH-A3	-11.2	-8.0
CH-B8	-63.6	-59.5
CH-B10	-32.4	-32.6
CH-C13	51.5	53.1
CH-C15	-15.9	-13.0
CH-D18	-32.3	-32.0
CH-D20	41.3	46.2

n =	7
RMS =	3.0
R(xy) =	0.999
R <sup>2</sup> =	99.80%

Correlation between calculated and experimental RDCs



**PALES output file (fitting procedure results)**

REMARK Molecular Alignment Simulation.

REMARK Simulation parameters.

DATA PALES\_MODE DC

DATA TENSOR\_MODE SVD (Order Matrix Method)

REMARK Order matrix.

DATA SAUPE\_MATRIX S(zz) S(xx-yy) S(xy) S(xz) S(yz)  
DATA SAUPE 3.2052e-03 -9.9240e-04 7.8311e-04 -2.1170e-04 -1.0380e-04

DATA IRREDUCIBLE REP A0 A1R A1I A2R A2I  
DATA IRREDUCIBLE 5.0812e-03 2.7403e-04 -1.3436e-04 -6.4229e-04 -1.0137e-03  
DATA IRREDUCIBLE GENERAL\_MAGNITUDE 5.3745e-03

REMARK Mapping of coordinates.

DATA MAPPING\_COOR Szz\_d(x) Szz\_d(y) Syy\_d(x) Syy\_d(y) Sxx\_d(x) Sxx\_d(y)  
DATA MAPPING -0.13797 1.51597 -0.50041 0.02368 1.07011 0.04944  
DATA MAPPING INV 0.03420 -1.51597 2.64030 -0.02368 -2.06764 -0.04944

REMARK Eigensystem & Euler angles for clockwise rotation about z, y', z'.

DATA EIGENVALUES (Sxx\_d,Syy\_d,Szz\_d) -6.8506e-04 -2.5329e-03 3.2179e-03  
DATA EIGENVECTORS (x\_coor y\_coor z\_coor)  
DATA EIGENVECTORS X\_AXIS 4.7829e-01 8.7681e-01 4.9423e-02  
DATA EIGENVECTORS Y\_AXIS 8.7707e-01 -4.7977e-01 2.3680e-02  
DATA EIGENVECTORS Z\_AXIS -4.4475e-02 -3.2022e-02 9.9850e-01

DATA Q\_EULER\_SOLUTIONS ALPHA BETA GAMMA  
DATA Q\_EULER\_ANGLES 1 154.40 3.14 324.25  
DATA Q\_EULER\_ANGLES 2 334.40 3.14 324.25  
DATA Q\_EULER\_ANGLES 3 25.60 176.86 144.25  
DATA Q\_EULER\_ANGLES 4 205.60 176.86 144.25

REMARK Euler angles (psi(theta/phi) for rotation about x, y, z.

DATA EULER\_SOLUTIONS 2  
DATA EULER\_ANGLES -1.84 2.55 118.60  
DATA EULER\_ANGLES 178.16 177.45 -61.40

DATA Da 1.608961e-03  
DATA Dr 6.159366e-04

DATA Aa 3.217922e-03  
DATA Ar 1.231873e-03

DATA Da\_HN 3.472975e+01  
DATA Rhombicity 3.828164e-01

REMARK Dipolar couplings.

DATA N 7  
DATA RMS 3.000  
DATA Chi2 62.993

```

DATA CORR R          0.999
DATA Q SAUPE        0.029
DATA REGRESSION OFFSET   2.499 +/- 0.799 [Hz]
DATA REGRESSION SLOPE    1.009 +/- 0.020 [Hz]
DATA REGRESSION BAX SLOPE 1.010 +/- 0.014 [Hz]

VARS    RESID_I RESNAME_I ATOMNAME_I RESID_J RESNAME_J ATOMNAME_J DI D_OBS D D_DIFF
DD W
FORMAT %4d %4s %4s %4d %4s %4s %9.2f %9.3f %9.3f %9.3f %.2f %.2f

1  CAL  C14      1  CAL  H33  47786.71  -11.2000  -7.9838  -3.2162  1.0000  1.00
1  CAL  C40      1  CAL  H37  47458.52  -63.6000  -59.4735  -4.1265  1.0000  1.00
1  CAL  C13      1  CAL  H15  47274.22  -32.4000  -32.5502  0.1502  1.0000  1.00
1  CAL  C12      1  CAL  H18  47421.91  51.5000   53.1087  -1.6087  1.0000  1.00
1  CAL  C6       1  CAL  H20  47278.38  -15.9000  -12.9653  -2.9347  1.0000  1.00
1  CAL  C21      1  CAL  H22  47586.48  -32.3000  -32.0141  -0.2859  1.0000  1.00
1  CAL  C11      1  CAL  H26  47145.48  41.3000   46.2311  -4.9311  1.0000  1.00

```

5A (CONE) @ POLY( $\gamma$ -ETHYL-L-GLUTAMATE), MEDIUM A

EVALUATION OF RDCs

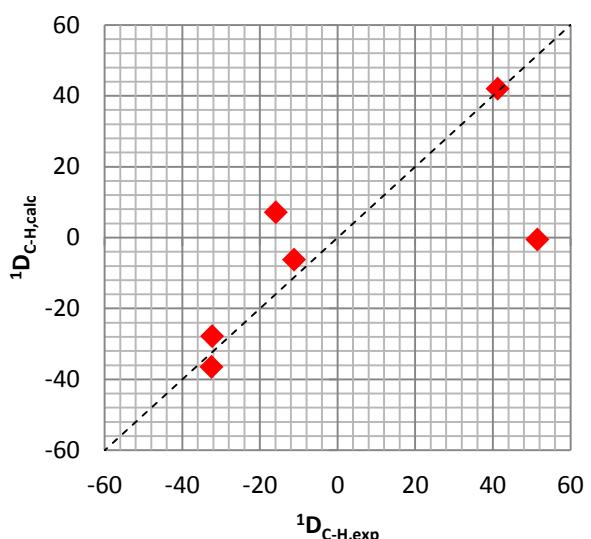
	$^1J_{C-H}$ [Hz]	$^1T_{C-H}$ [Hz]	RDC [Hz]	$^1D_{C-H}$ [Hz]
CH-A3	160.4	138.0	-22.4	-11.2
CH-B8	159.3	32.0	-127.3	-63.6
CH-B10	160.8	96.0	-64.8	-32.4
CH-C13	157.0	260.0	103.0	51.5
CH-C15	159.8	128.0	-31.8	-15.9
CH-D18	158.6	82.0	-76.6	-38.3
CH-D20	158.3	241.0	82.7	41.3

FITTING RESULTS

$^1D_{C-H}$	$^1D_{C-H,exp}$	$^1D_{C-H,calc}$
CH-A3	-11.2	-6.3
CH-B8	-63.6	-60.0
CH-B10	-32.4	-36.5
CH-C13	51.5	-0.6
CH-C15	-15.9	7.0
CH-D18	-32.3	-27.8
CH-D20	41.3	41.9

n =	7
RMS =	21.8
R(xy) =	0.827
R <sup>2</sup> =	68.47%

Correlation between calculated and experimental RDCs



5D (1,2-ALT) @ POLY( $\gamma$ -BENZYL-L-GLUTAMATE), MEDIUM B

EVALUATION OF RDCs

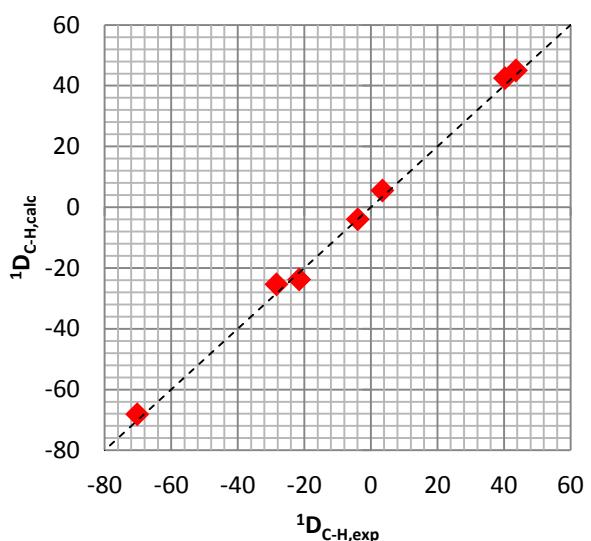
	$^1J_{C-H}$ [Hz]	$^1T_{C-H}$ [Hz]	RDC [Hz]	$^1D_{C-H}$ [Hz]
CH-A3	160.4	152.5	-7.9	-3.9
CH-B8	159.3	18.8	-140.5	-70.2
CH-B10	160.8	104.1	-56.7	-28.3
CH-C13	157.0	244.3	87.3	43.6
CH-C15	159.8	166.9	7.1	3.5
CH-D18	158.6	115.6	-43.0	-21.5
CH-D20	158.3	239.2	80.9	40.4

FITTING RESULTS

$^1D_{C-H}$	$^1D_{C-H,exp}$	$^1D_{C-H,calc}$
CH-A3	-3.9	-4.0
CH-B8	-70.2	-68.2
CH-B10	-28.3	-25.4
CH-C13	43.6	44.9
CH-C15	3.5	5.4
CH-D18	-21.5	-23.9
CH-D20	40.4	42.4

n =	7
RMS =	2.0
R(xy) =	0.999
R <sup>2</sup> =	99.80%

Correlation between calculated and experimental RDCs



**PALES output file (fitting procedure results)**

REMARK Molecular Alignment Simulation.

REMARK Simulation parameters.

DATA PALES\_MODE DC

DATA TENSOR\_MODE SVD (Order Matrix Method)

REMARK Order matrix.

DATA SAUPE\_MATRIX S(zz) S(xx-yy) S(xy) S(xz) S(yz)  
DATA SAUPE 2.6191e-003 -1.1043e-003 5.0729e-004 2.6151e-004 2.1389e-005

DATA IRREDUCIBLE REP A0 A1R A1I A2R A2I  
DATA IRREDUCIBLE 4.1521e-003 -3.3851e-004 2.7687e-005 -7.1470e-004 -6.5665e-004  
DATA IRREDUCIBLE GENERAL\_MAGNITUDE 4.3994e-003

REMARK Mapping of coordinates.

DATA MAPPING\_COOR Szz\_d(x) Szz\_d(y) Syy\_d(x) Syy\_d(y) Sxx\_d(x) Sxx\_d(y)  
DATA MAPPING 0.01539 1.50908 -0.36725 -0.05035 1.20411 -0.03566  
DATA MAPPING INV -0.17836 -1.50908 2.77036 0.05035 -1.93548 0.03566

REMARK Eigensystem & Euler angles for clockwise rotation about z, y', z''.

DATA EIGENVALUES (Sxx\_d,Syy\_d,Szz\_d) -5.6384e-004 -2.0712e-003 2.6351e-003  
DATA EIGENVECTORS (x\_coor y\_coor z\_coor)  
DATA EIGENVECTORS X\_AXIS 3.5758e-001 9.3320e-001 -3.5650e-002  
DATA EIGENVECTORS Y\_AXIS 9.3197e-001 -3.5903e-001 -5.0325e-002  
DATA EIGENVECTORS Z\_AXIS 5.9763e-002 1.5230e-002 9.9810e-001

DATA Q\_EULER\_SOLUTIONS ALPHA BETA GAMMA  
DATA Q\_EULER\_ANGLES 1 305.31 3.54 165.70  
DATA Q\_EULER\_ANGLES 2 125.31 3.54 165.70  
DATA Q\_EULER\_ANGLES 3 234.69 176.46 345.70  
DATA Q\_EULER\_ANGLES 4 54.69 176.46 345.70

REMARK Euler angles (psi(theta/phi) for rotation about x, y, z.

DATA EULER\_SOLUTIONS 2  
DATA EULER\_ANGLES 0.87 -3.43 110.99  
DATA EULER\_ANGLES 180.87 183.43 -69.01

DATA Da 1.317540e-003  
DATA Dr 5.024638e-004

DATA Aa 2.635081e-003  
DATA Ar 1.004928e-003

DATA Da\_HN 2.843937e+001  
DATA Rhombicity 3.813650e-001

REMARK Dipolar couplings.

DATA N 7  
DATA RMS 1.998

```

DATA Chi2          27.946
DATA CORR R       0.999
DATA Q SAUPE      0.024
DATA REGRESSION OFFSET   1.101 +/- 0.755 [Hz]
DATA REGRESSION SLOPE    1.001 +/- 0.020 [Hz]
DATA REGRESSION BAX SLOPE 1.002 +/- 0.014 [Hz]

VARS   RESID_I RESNAME_I ATOMNAME_I RESID_J RESNAME_J ATOMNAME_J DI D_OBS D D_DIFF
DD W
FORMAT %4d %4s %4s %4d %4s %4s %9.2f %9.3f %9.3f %9.3f %.2f %.2f

 1 CAL C14     1 CAL H33  47786.72   -3.9000   -4.0434   0.1434  1.0000  1.00
 1 CAL C40     1 CAL H37  47458.52   -70.2000  -68.1616  -2.0384  1.0000  1.00
 1 CAL C13     1 CAL H15  47274.22   -28.3000  -25.3818  -2.9182  1.0000  1.00
 1 CAL C12     1 CAL H18  47421.91   43.6000   44.9232  -1.3232  1.0000  1.00
 1 CAL C6      1 CAL H20  47278.38   3.5000    5.4229  -1.9229  1.0000  1.00
 1 CAL C21     1 CAL H22  47586.48  -21.5000  -23.9042  2.4042  1.0000  1.00
 1 CAL C11     1 CAL H26  47145.48   40.4000   42.4065  -2.0065  1.0000  1.00

```

5A (CONE) @ POLY( $\gamma$ -BENZYL-L-GLUTAMATE), MEDIUM B

EVALUATION OF RDCs

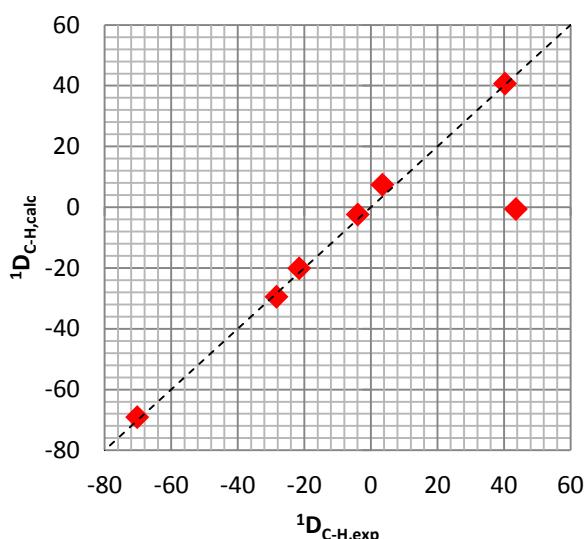
	$^1J_{C-H}$ [Hz]	$^1T_{C-H}$ [Hz]	RDC [Hz]	$^1D_{C-H}$ [Hz]
CH-A3	160.4	152.5	-7.9	-3.9
CH-B8	159.3	18.8	-140.5	-70.2
CH-B10	160.8	104.1	-56.7	-28.3
CH-C13	157.0	244.3	87.3	43.6
CH-C15	159.8	166.9	7.1	3.5
CH-D18	158.6	115.6	-43.0	-21.5
CH-D20	158.3	239.2	80.9	40.4

FITTING RESULTS

$^1D_{C-H}$	$^1D_{C-H,exp}$	$^1D_{C-H,calc}$
CH-A3	-3.9	-2.4
CH-B8	-70.2	-69.1
CH-B10	-28.3	-29.6
CH-C13	43.6	-0.6
CH-C15	3.5	7.4
CH-D18	-21.5	-20.2
CH-D20	40.4	40.6

n =	7
RMS =	16.8
R(xy) =	0.904
R <sup>2</sup> =	81.65%

Correlation between calculated and experimental RDCs



---

**SUMMARY (COMPARISON OF THE RESULTS)**

Table 1: Comparison of RDC values of structure 5c (*D-paco*) in various polypeptide alignment media.

	$^1J_{CH}$ (CDCl <sub>3</sub> )	$^1D_{CH}$ (PELG)	$^1D_{CH}$ (PBLG)
<b>CH-3A</b>	159.8	12.1	11.8
<b>CH-8B</b>	160.3	-56.0	-49.6
<b>CH-10B</b>	161.1	18.2	16.8
<b>CH-13C</b>	157.0	-53.1	-65.5
<b>CH-15C</b>	159.7	72.3	57.8
<b>CH-18D</b>	159.4	5.5	6.4
<b>CH-20D</b>	158.0	25.6	19.3

Table 2: Comparison of RDC values of structure 5d (1,2-*alt*) in various polypeptide alignment media.

	$^1J_{CH}$ (CDCl <sub>3</sub> )	$^1D_{CH}$ (PELG)	$^1D_{CH}$ (PBLG)
<b>CH-3A</b>	160.4	-11.2	-3.9
<b>CH-8B</b>	159.3	-63.6	-70.2
<b>CH-10B</b>	160.8	-32.4	-28.3
<b>CH-13C</b>	157.0	51.5	43.6
<b>CH-15C</b>	159.8	-15.9	3.5
<b>CH-18D</b>	158.6	-32.3	-21.5
<b>CH-20D</b>	158.3	41.3	40.4

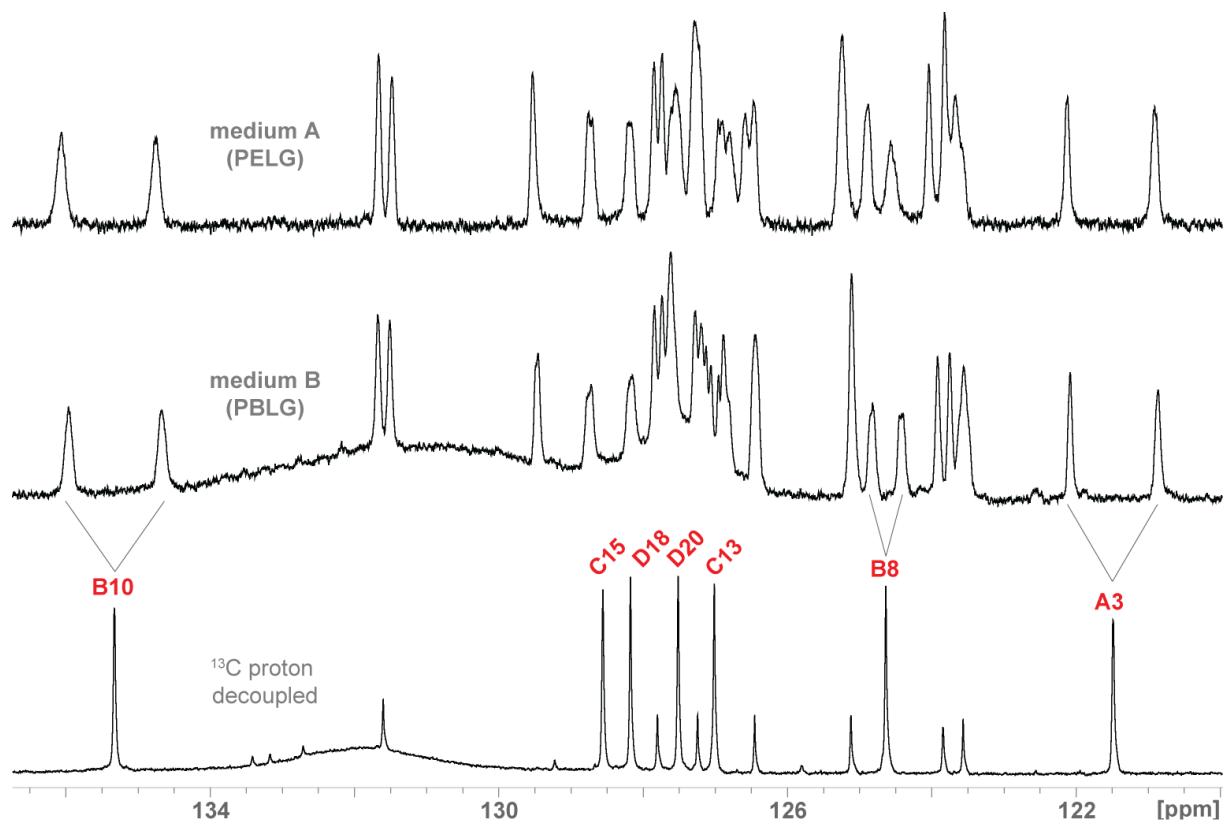


Figure 25: Comparison of selected region of <sup>13</sup>C Z-restored proton coupled spectra of 5c (D-paco) in polypeptide alignment media.

Table 3: Comparison of fitting quality of structures 5c (D-paco) and 5b (C-paco) in various alignment media.

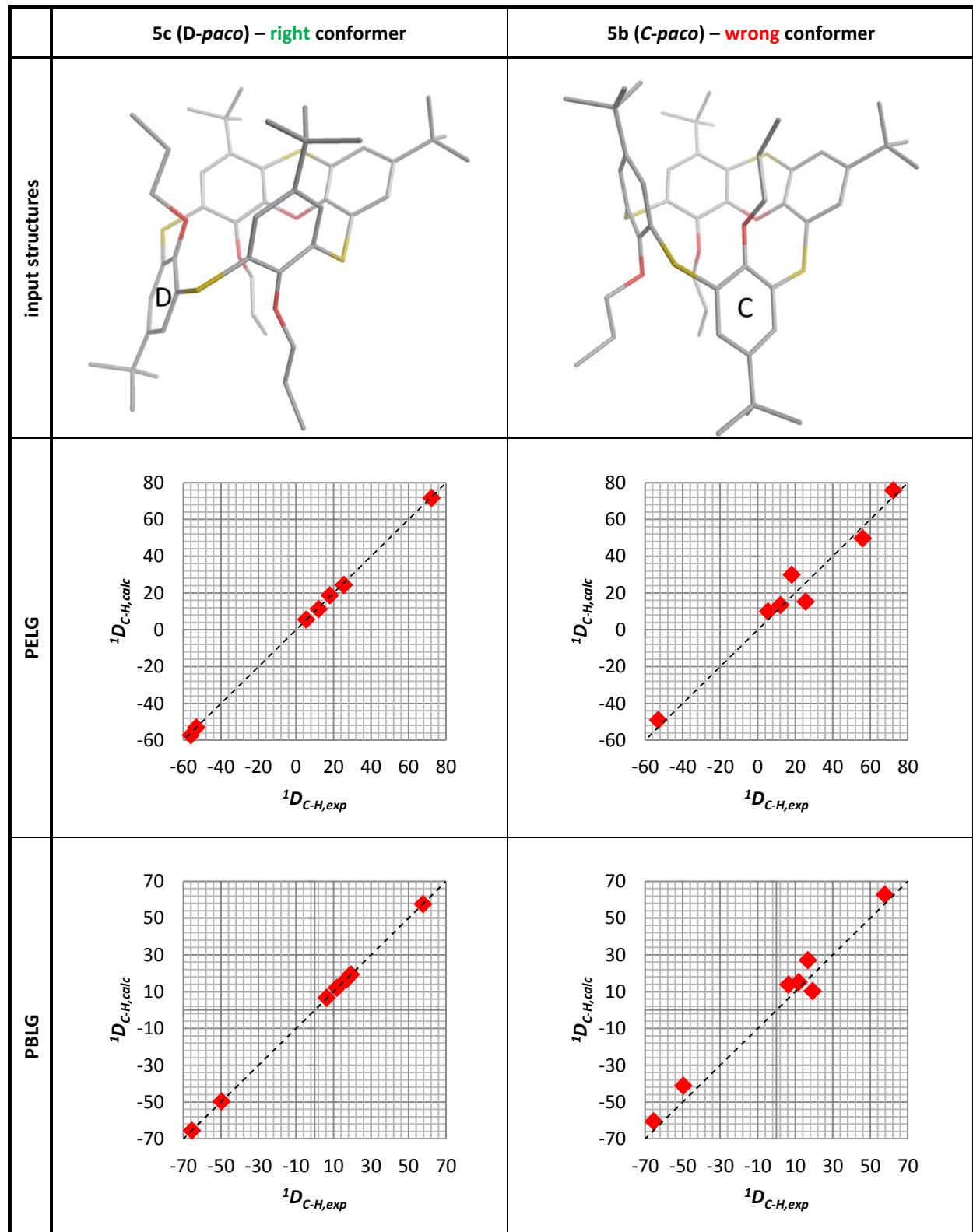


Table 4: Comparison of fitting quality of structures K2( CD\_down) and K2( cone) in various alignment media.

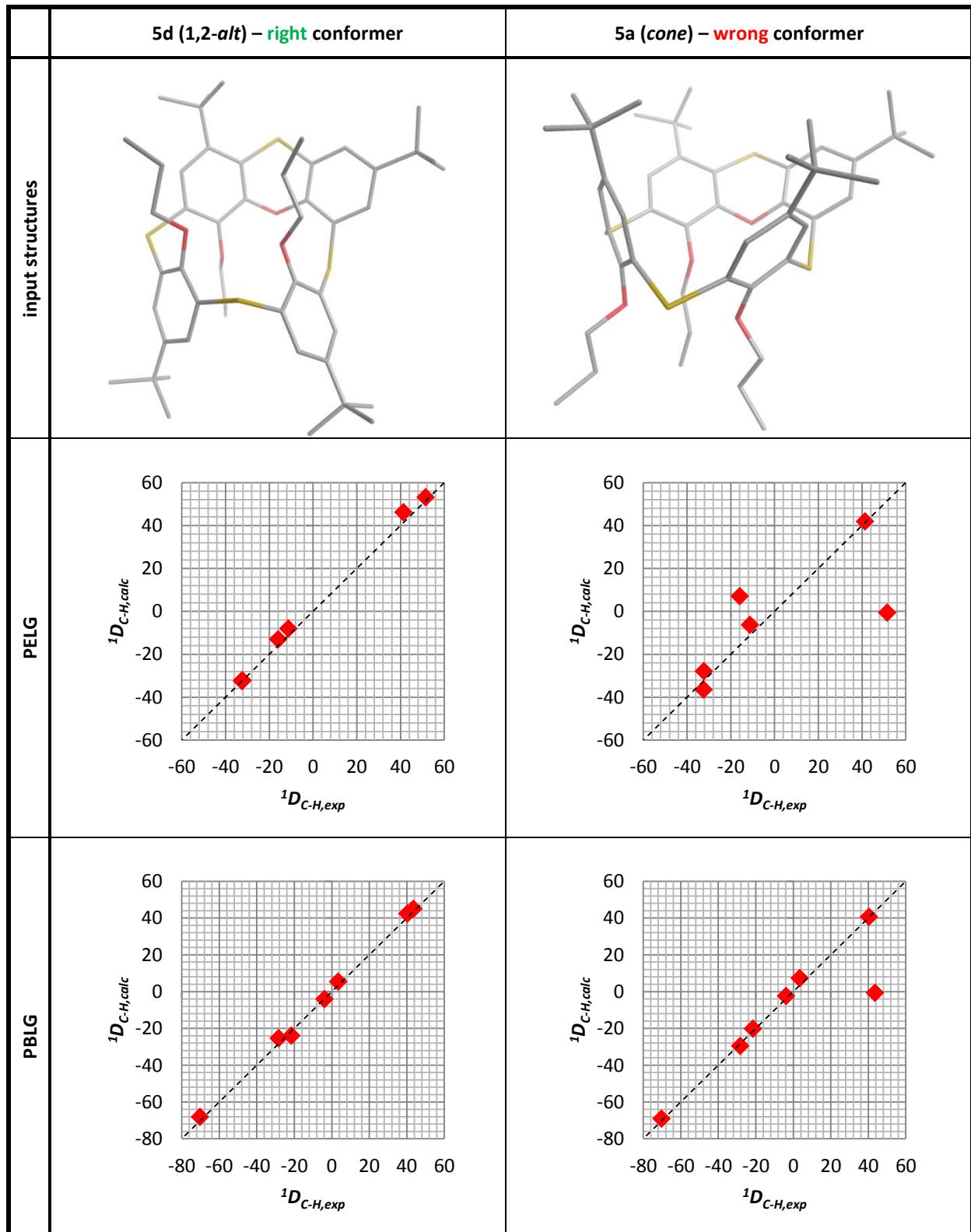


Table 5: Comparison of molecular orientations of 5c (D-paco) within the magnetic field.

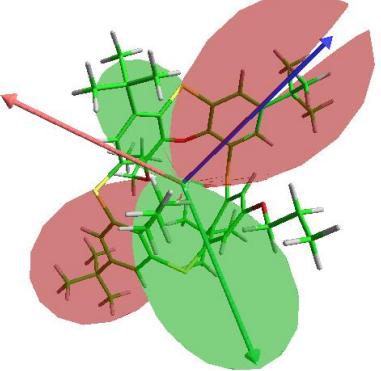
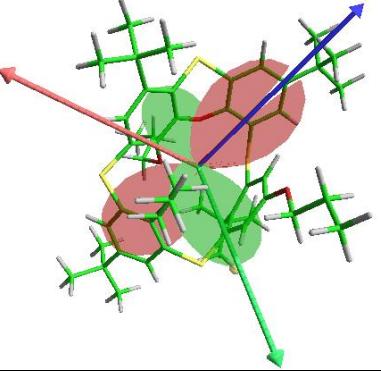
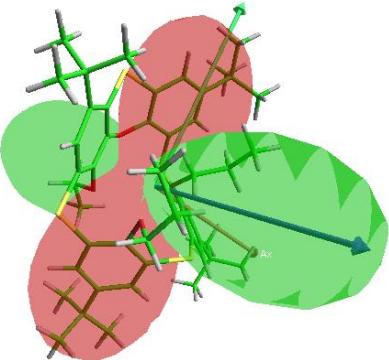
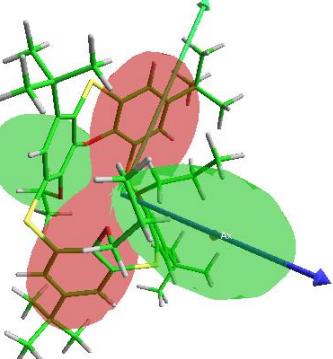
Alignment medium	Orientation of 5c (D-paco) within the magnetic field
<b>PELG (medium A)</b>	 A 3D molecular model of compound 5c (D-paco) showing its orientation within a magnetic field. The molecule is represented by green sticks and red spheres, with a blue arrow indicating the direction of the magnetic field. The molecule is tilted, with its long axis approximately parallel to the blue field vector. A red arrow points to the left, and a green arrow points downwards.
<b>PBLG (medium B)</b>	 A 3D molecular model of compound 5c (D-paco) showing its orientation within a magnetic field. The molecule is represented by green sticks and red spheres, with a blue arrow indicating the direction of the magnetic field. The molecule is tilted, with its long axis approximately parallel to the blue field vector. A red arrow points to the left, and a green arrow points downwards.

Table 6: Comparison of molecular orientations of 5d (1,2-*alt*) within the magnetic field.

Alignment medium	Orientation of 5d (1,2- <i>alt</i> ) within the magnetic field
<b><i>PELG</i> (medium A)</b>	
<b><i>PBLG</i> (medium B)</b>	

## DFT OPTIMIZATION OF CONFORMERS

**Ab initio geometry optimizations were performed by program Gaussian03** (M. J. Frisch, H. B. Schlegel, G. E. Scuseria, M. A. Robb, J. R. Cheeseman, J. A. Montgomery, Jr., T. Vreven, K. N. Kudin, J. C. Burant, J. M. Millam, S. S. Iyengar, J. Tomasi, V. Barone, B. Mennucci, M. Cossi, G. Scalmani, N. Rega, G. A. Petersson, H. Nakatsuji, M. Hada, M. Ehara, K. Toyota, R. Fukuda, J. Hasegawa, M. Ishida, T. Nakajima, Y. Honda, O. Kitao, H. Nakai, M. Klene, X. Li, J. E. Knox, H. P. Hratchian, J. B. Cross, V. Bakken, C. Adamo, J. Jaramillo, R. Gomperts, R. E. Stratmann, O. Yazyev, A. J. Austin, R. Cammi, C. Pomelli, J. W. Ochterski, P. Y. Ayala, K. Morokuma, G. A. Voth, P. Salvador, J. J. Dannenberg, V. G. Zakrzewski, S. Dapprich, A. D. Daniels, M. C. Strain, O. Farkas, D. K. Malick, A. D. Rabuck, K. Raghavachari, J. B. Foresman, J. V. Ortiz, Q. Cui, A. G. Baboul, S. Clifford, J. Cioslowski, B. B. Stefanov, G. Liu, A. Liashenko, P. Piskorz, I. Komaromi, R. L. Martin, D. J. Fox, T. Keith, M. A. Al-Laham, C. Y. Peng, A. Nanayakkara, M. Challacombe, P. M. W. Gill, B. Johnson, W. Chen, M. W. Wong, C. Gonzalez, and J. A. Pople, Gaussian, Inc., Wallingford CT, **2004.**)

**Table 7: Sums of electronic and thermal free energies (on computational level RB3LYP/6-31G\*) of conformers 5a, 5b, 5c and 5d.** 5c (D-paco) was selected due to the lowest energy state as the reference.

structure	GFE [Hartree]	GFE [kJ/mol]	ΔE [kJ/mol]	population
<b>5a (cone)</b>	-3798,460693	-9972858,55	29,78	6,06E-06
<b>5b (C-paco)</b>	-3798,460455	-9972857,92	30,40	4,71E-06
<b>5c (D-paco)</b>	<b>-3798,472035</b>	<b>-9972888,33</b>	<b>0,00</b>	<b>1,00E+00</b>
<b>5d (1,2-alt)</b>	-3798,462869	-9972864,26	24,07	6,08E-05





-7.8501	-0.5873	-2.8596	H	0	0	0	0	0	0	0	0	0	0	0	0	0	0
1.9994	6.1711	-2.1977	H	0	0	0	0	0	0	0	0	0	0	0	0	0	0
3.4346	5.2188	-1.7938	H	0	0	0	0	0	0	0	0	0	0	0	0	0	0
3.1974	6.7968	-1.0492	H	0	0	0	0	0	0	0	0	0	0	0	0	0	0
-0.6321	1.0798	-3.8932	H	0	0	0	0	0	0	0	0	0	0	0	0	0	0
-0.6562	-0.5911	-3.2870	H	0	0	0	0	0	0	0	0	0	0	0	0	0	0
2.3142	-2.9011	0.7535	H	0	0	0	0	0	0	0	0	0	0	0	0	0	0
3.1793	-2.0102	2.6891	H	0	0	0	0	0	0	0	0	0	0	0	0	0	0
2.0067	-1.2365	3.7634	H	0	0	0	0	0	0	0	0	0	0	0	0	0	0
3.1870	-2.3890	4.4195	H	0	0	0	0	0	0	0	0	0	0	0	0	0	0
5.9097	-1.3387	1.7738	H	0	0	0	0	0	0	0	0	0	0	0	0	0	0
7.6379	-1.4099	1.3764	H	0	0	0	0	0	0	0	0	0	0	0	0	0	0
6.7197	0.0723	1.0773	H	0	0	0	0	0	0	0	0	0	0	0	0	0	0
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1 20 4	0 0 0	0															
1 27 1	0 0 0	0															
2 4 4	0 0 0	0															
2 8 4	0 0 0	0															
2 57 1	0 0 0	0															
3 6 4	0 0 0	0															
3 18 4	0 0 0	0															
3 24 1	0 0 0	0															
4 14 4	0 0 0	0															
4 24 1	0 0 0	0															
5 8 4	0 0 0	0															
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5 46 1	0 0 0	0															
6 58 1	0 0 0	0															
7 9 4	0 0 0	0															
7 17 4	0 0 0	0															
7 46 1	0 0 0	0															
8 45 1	0 0 0	0															
9 16 4	0 0 0	0															
9 59 1	0 0 0	0															
10 11 4	0 0 0	0															
10 17 4	0 0 0	0															
10 31 1	0 0 0	0															
11 16 4	0 0 0	0															
11 60 1	0 0 0	0															
12 13 4	0 0 0	0															
12 30 1	0 0 0	0															
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13 15 4	0 0 0	0															
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14 19 4	0 0 0	0															
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18 33 1	0 0 0	0															
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20 21 4	0 0 0	0															
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22 47 1	0 0 0	0															
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22 64 1	0 0 0	0															
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25 43 4	0 0 0	0															

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35	78	1	0	0	0	0
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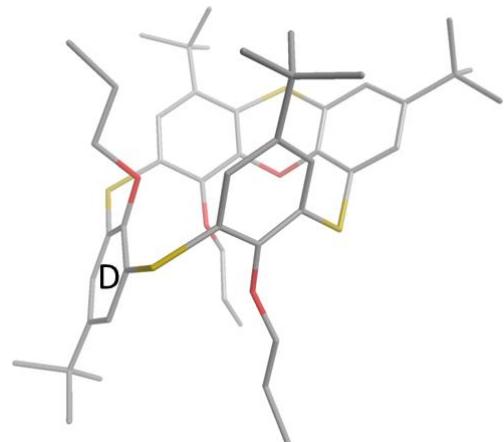
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-0.3695	-5.6704	3.9190 H	0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0
-2.1180	-5.8437	4.1702 H	0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0
5.2142	4.7337	0.0230 H	0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0
5.7507	4.1374	1.6005 H	0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0
6.9369	4.4499	0.3175 H	0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0
-3.0927	-4.0102	1.8483 H	0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0
-3.0965	-2.4904	2.7539 H	0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0
-3.7105	-3.9765	3.5057 H	0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0
2.4157	1.1816	-2.9690 H	0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0
3.3293	-0.1317	-2.2369 H	0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0
3.0664	0.3542	-5.2803 H	0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0
4.4959	0.7136	-4.3033 H	0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0
3.9892	-0.9648	-4.5438 H	0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0
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2 32 4 0 0 0 0 0			
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3 6 1 0 0 0 0 0			
3 10 1 0 0 0 0 0			
3 13 1 0 0 0 0 0			
3 27 1 0 0 0 0 0			
4 27 4 0 0 0 0 0			
4 33 4 0 0 0 0 0			
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6 61 1 0 0 0 0 0			
6 62 1 0 0 0 0 0			
7 12 4 0 0 0 0 0			
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8 32 4 0 0 0 0 0			
8 41 1 0 0 0 0 0			
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26	80	1	0	0	0	0
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28	82	1	0	0	0	0
28	83	1	0	0	0	0
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38	94	1	0	0	0	0
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38	96	1	0	0	0	0
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42	51	1	0	0	0	0
42	54	1	0	0	0	0
43	97	1	0	0	0	0
43	98	1	0	0	0	0
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44	100	1	0	0	0	0
44	101	1	0	0	0	0
45	102	1	0	0	0	0
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**FINAL GEOMETRY OF 5C (D-PACO)**

DFT method: RB3LYP/6-31G\*



**Created by GaussView 3.09**

121126	0	0	0	0	0	0	0	0	0	0	0	0	0	0
-4.5839		1.4321		-0.3617	C	0	0	0	0	0	0	0	0	0
7.3160		-1.8173		-2.3143	C	0	0	0	0	0	0	0	0	0
3.1063		5.4884		-1.4578	C	0	0	0	0	0	0	0	0	0
2.4490		4.8902		-0.1892	C	0	0	0	0	0	0	0	0	0
2.3486		-1.7635		3.6940	C	0	0	0	0	0	0	0	0	0
-2.5166		-0.5374		-3.2269	C	0	0	0	0	0	0	0	0	0
1.2403		-1.8989		2.6325	C	0	0	0	0	0	0	0	0	0
-0.3661		3.0358		4.7072	C	0	0	0	0	0	0	0	0	0
-2.8159		-1.7223		2.4423	S	0	0	0	0	0	0	0	0	0
-0.3295		2.1914		3.4290	C	0	0	0	0	0	0	0	0	0
-2.5244		3.2161		-0.2380	S	0	0	0	0	0	0	0	0	0
3.2939		-0.5990		3.3153	C	0	0	0	0	0	0	0	0	0
6.3323		-3.7010		-0.9963	C	0	0	0	0	0	0	0	0	0
4.9251		-0.1226		-1.4024	C	0	0	0	0	0	0	0	0	0
-3.1984		-4.0715		-1.4098	C	0	0	0	0	0	0	0	0	0
0.6263		-2.3294		0.3009	C	0	0	0	0	0	0	0	0	0
2.5323		-0.2695		-1.7265	C	0	0	0	0	0	0	0	0	0
-0.3869		1.6747		-1.2396	C	0	0	0	0	0	0	0	0	0
-1.7253		1.8335		2.9415	C	0	0	0	0	0	0	0	0	0
6.4166		-2.1669		-1.1047	C	0	0	0	0	0	0	0	0	0
-1.1019		-1.9281		1.9438	C	0	0	0	0	0	0	0	0	0
-4.7080		-0.6761		0.7600	C	0	0	0	0	0	0	0	0	0
-6.6537		0.1234		-2.2091	C	0	0	0	0	0	0	0	0	0
1.3427		0.3208		-2.0686	O	0	0	0	0	0	0	0	0	0
-0.1188		-1.7554		2.9253	C	0	0	0	0	0	0	0	0	0
-1.6883		-2.5470		-0.3229	O	0	0	0	0	0	0	0	0	0
-0.8087		2.8352		-0.5846	C	0	0	0	0	0	0	0	0	0
-1.5953		1.0327		1.7499	O	0	0	0	0	0	0	0	0	0
-3.3303		1.6740		0.2028	C	0	0	0	0	0	0	0	0	0
-3.4538		-0.4509		1.3443	C	0	0	0	0	0	0	0	0	0
3.8447		-2.2595		-1.3425	C	0	0	0	0	0	0	0	0	0
1.4983		3.7243		-0.5639	C	0	0	0	0	0	0	0	0	0
3.1595		-3.0802		3.7553	C	0	0	0	0	0	0	0	0	0
-7.6460		1.2168		-0.1634	C	0	0	0	0	0	0	0	0	0
1.9167		2.5360		-1.2170	C	0	0	0	0	0	0	0	0	0
-2.1381		-3.9167		-0.3305	C	0	0	0	0	0	0	0	0	0
3.5817		2.2573		-1.8535	S	0	0	0	0	0	0	0	0	0
-6.7241		0.0792		-0.6647	C	0	0	0	0	0	0	0	0	0
5.0327		-1.5190		-1.2966	C	0	0	0	0	0	0	0	0	0
7.0665		-1.6266		0.1911	C	0	0	0	0	0	0	0	0	0
1.7811		-1.4886		5.0995	C	0	0	0	0	0	0	0	0	0
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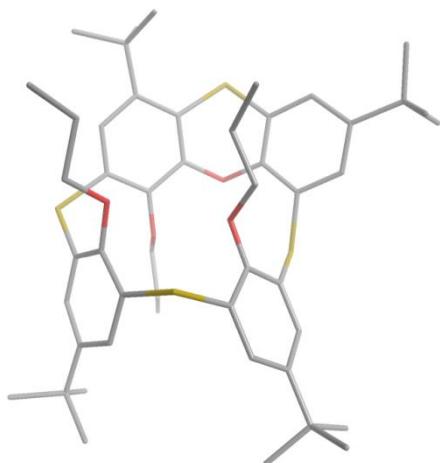
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0.9260	6.4860	-0.1462	H	0	0	0	0	0	0	0	0	0	0	0	0
2.4136	6.8458	0.7359	H	0	0	0	0	0	0	0	0	0	0	0	0
-0.2327	4.6859	0.2762	H	0	0	0	0	0	0	0	0	0	0	0	0
3.0621	4.0324	1.7273	H	0	0	0	0	0	0	0	0	0	0	0	0
4.1490	3.6062	0.3954	H	0	0	0	0	0	0	0	0	0	0	0	0
4.1888	5.2385	1.0784	H	0	0	0	0	0	0	0	0	0	0	0	0
-1.8415	1.5200	-3.4229	H	0	0	0	0	0	0	0	0	0	0	0	0
-0.5420	0.3036	-3.5133	H	0	0	0	0	0	0	0	0	0	0	0	0
-4.2228	-5.7820	-0.5130	H	0	0	0	0	0	0	0	0	0	0	0	0
-2.9693	-6.2286	-1.6812	H	0	0	0	0	0	0	0	0	0	0	0	0
-4.5181	-5.5893	-2.2470	H	0	0	0	0	0	0	0	0	0	0	0	0
1 29	4	0	0	0	0										
1 50	4	0	0	0	0										
1 58	1	0	0	0	0										
2 20	1	0	0	0	0										
2 59	1	0	0	0	0										
2 60	1	0	0	0	0										
2 61	1	0	0	0	0										
3 4	1	0	0	0	0										
3 62	1	0	0	0	0										
3 63	1	0	0	0	0										
3 64	1	0	0	0	0										
4 32	1	0	0	0	0										
4 47	1	0	0	0	0										
4 51	1	0	0	0	0										
5 7	1	0	0	0	0										
5 12	1	0	0	0	0										
5 33	1	0	0	0	0										
5 41	1	0	0	0	0										
6 45	1	0	0	0	0										
6 52	1	0	0	0	0										
6 65	1	0	0	0	0										
6 66	1	0	0	0	0										
7 25	4	0	0	0	0										
7 44	4	0	0	0	0										
8 10	1	0	0	0	0										
8 67	1	0	0	0	0										
8 68	1	0	0	0	0										
8 69	1	0	0	0	0										
9 21	1	0	0	0	0										
9 30	1	0	0	0	0										
10 19	1	0	0	0	0										
10 70	1	0	0	0	0										
10 71	1	0	0	0	0										
11 27	1	0	0	0	0										
11 29	1	0	0	0	0										
12 72	1	0	0	0	0										
12 73	1	0	0	0	0										
12 74	1	0	0	0	0										
13 20	1	0	0	0	0										
13 75	1	0	0	0	0										
13 76	1	0	0	0	0										
13 77	1	0	0	0	0										
14 39	4	0	0	0	0										
14 46	4	0	0	0	0										
14 78	1	0	0	0	0										
15 36	1	0	0	0	0										
15 53	1	0	0	0	0										
15 79	1	0	0	0	0										
15 80	1	0	0	0	0										
16 42	1	0	0	0	0										
16 44	4	0	0	0	0										
16 56	4	0	0	0	0										
17 24	1	0	0	0	0										

17	46	4	0	0	0	0
17	49	4	0	0	0	0
18	27	4	0	0	0	0
18	54	4	0	0	0	0
18	57	1	0	0	0	0
19	28	1	0	0	0	0
19	81	1	0	0	0	0
19	82	1	0	0	0	0
20	39	1	0	0	0	0
20	40	1	0	0	0	0
21	25	4	0	0	0	0
21	56	4	0	0	0	0
22	30	4	0	0	0	0
22	50	4	0	0	0	0
22	83	1	0	0	0	0
23	38	1	0	0	0	0
23	84	1	0	0	0	0
23	85	1	0	0	0	0
23	86	1	0	0	0	0
24	54	1	0	0	0	0
25	87	1	0	0	0	0
26	36	1	0	0	0	0
26	56	1	0	0	0	0
27	48	4	0	0	0	0
28	55	1	0	0	0	0
29	55	4	0	0	0	0
30	55	4	0	0	0	0
31	39	4	0	0	0	0
31	49	4	0	0	0	0
31	88	1	0	0	0	0
32	35	4	0	0	0	0
32	48	4	0	0	0	0
33	89	1	0	0	0	0
33	90	1	0	0	0	0
33	91	1	0	0	0	0
34	38	1	0	0	0	0
34	92	1	0	0	0	0
34	93	1	0	0	0	0
34	94	1	0	0	0	0
35	37	1	0	0	0	0
35	54	4	0	0	0	0
36	95	1	0	0	0	0
36	96	1	0	0	0	0
37	46	1	0	0	0	0
38	43	1	0	0	0	0
38	50	1	0	0	0	0
40	97	1	0	0	0	0
40	98	1	0	0	0	0
40	99	1	0	0	0	0
41	100	1	0	0	0	0
41	101	1	0	0	0	0
41	102	1	0	0	0	0
42	49	1	0	0	0	0
43	103	1	0	0	0	0
43	104	1	0	0	0	0
43	105	1	0	0	0	0
44	106	1	0	0	0	0
45	107	1	0	0	0	0
45	108	1	0	0	0	0
45	109	1	0	0	0	0
47	110	1	0	0	0	0
47	111	1	0	0	0	0
47	112	1	0	0	0	0
48	113	1	0	0	0	0
51	114	1	0	0	0	0
51	115	1	0	0	0	0

51116	1	0	0	0	0
52 57	1	0	0	0	0
52117	1	0	0	0	0
52118	1	0	0	0	0
53119	1	0	0	0	0
53120	1	0	0	0	0
53121	1	0	0	0	0

FINAL GEOMETRY OF 5D (1,2-ALT)

DFT method: RB3LYP/6-31G\*



Created by GaussView 3.09

121126	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
-4.1313		4.7334		1.7829	C	0	0	0	0	0	0	0	0	0	0	0	0	0	0
-3.9651		3.2153		-0.2235	C	0	0	0	0	0	0	0	0	0	0	0	0	0	0
1.7414		-2.0955		1.1364	C	0	0	0	0	0	0	0	0	0	0	0	0	0	0
-3.7462		0.9747		-1.1519	C	0	0	0	0	0	0	0	0	0	0	0	0	0	0
-0.5680		-1.7627		-1.4761	C	0	0	0	0	0	0	0	0	0	0	0	0	0	0
3.1480		2.4532		1.4625	C	0	0	0	0	0	0	0	0	0	0	0	0	0	0
2.0816		1.5583		1.5617	C	0	0	0	0	0	0	0	0	0	0	0	0	0	0
-5.0955		-2.9709		0.7809	C	0	0	0	0	0	0	0	0	0	0	0	0	0	0
-4.3957		-3.6877		-0.4000	C	0	0	0	0	0	0	0	0	0	0	0	0	0	0
3.3261		-2.8871		-0.5246	C	0	0	0	0	0	0	0	0	0	0	0	0	0	0
2.0655		3.5743		-0.3452	C	0	0	0	0	0	0	0	0	0	0	0	0	0	0
-2.6141		3.4242		-0.5463	C	0	0	0	0	0	0	0	0	0	0	0	0	0	0
-1.8499		-3.6069		-0.5816	C	0	0	0	0	0	0	0	0	0	0	0	0	0	0
-0.6129		-3.0448		-0.9166	C	0	0	0	0	0	0	0	0	0	0	0	0	0	0
-3.0169		-1.6407		-1.2879	C	0	0	0	0	0	0	0	0	0	0	0	0	0	0
3.1565		3.4947		0.5231	C	0	0	0	0	0	0	0	0	0	0	0	0	0	0
3.9505		-1.1061		0.9385	C	0	0	0	0	0	0	0	0	0	0	0	0	0	0
0.9469		1.7015		0.7310	C	0	0	0	0	0	0	0	0	0	0	0	0	0	0
2.0515		-2.9248		0.0411	C	0	0	0	0	0	0	0	0	0	0	0	0	0	0
-6.2389		3.9305		0.7035	C	0	0	0	0	0	0	0	0	0	0	0	0	0	0
0.9775		2.6882		-0.2708	C	0	0	0	0	0	0	0	0	0	0	0	0	0	0
-4.7946		5.5785		-0.5012	C	0	0	0	0	0	0	0	0	0	0	0	0	0	0
2.6734		-1.1185		1.5161	C	0	0	0	0	0	0	0	0	0	0	0	0	0	0
-5.3382		-3.7636		-1.6268	C	0	0	0	0	0	0	0	0	0	0	0	0	0	0
0.8046		-0.9875		-3.2524	C	0	0	0	0	0	0	0	0	0	0	0	0	0	0
-4.1448		-5.1453		0.0452	C	0	0	0	0	0	0	0	0	0	0	0	0	0	0
-3.0787		-2.9599		-0.7702	C	0	0	0	0	0	0	0	0	0	0	0	0	0	0
-4.7802		4.3453		0.4333	C	0	0	0	0	0	0	0	0	0	0	0	0	0	0
4.3215		-2.0150		-0.0565	C	0	0	0	0	0	0	0	0	0	0	0	0	0	0
-1.8184		2.4361		-1.1362	C	0	0	0	0	0	0	0	0	0	0	0	0	0	0
-2.4060		1.2026		-1.4570	C	0	0	0	0	0	0	0	0	0	0	0	0	0	0
-4.5131		1.9633		-0.5286	C	0	0	0	0	0	0	0	0	0	0	0	0	0	0
-1.7798		-1.0672		-1.6050	C	0	0	0	0	0	0	0	0	0	0	0	0	0	0
6.6763		-1.0173		-0.0295	C	0	0	0	0	0	0	0	0	0	0	0	0	0	0
-0.9590		1.1893		2.0160	C	0	0	0	0	0	0	0	0	0	0	0	0	0	0
5.7449		-2.0805		-0.6420	C	0	0	0	0	0	0	0	0	0	0	0	0	0	0
0.5454		-2.2373		1.7905	O	0	0	0	0	0	0	0	0	0	0	0	0	0	0
-2.0538		0.1434		2.1351	C	0	0	0	0	0	0	0	0	0	0	0	0	0	0
-0.7564		-4.3317		4.6605	C	0	0	0	0	0	0	0	0	0	0	0	0	0	0
-0.1504		2.8213		-1.6671	S	0	0	0	0	0	0	0	0	0	0	0	0	0	0
4.3427		4.4757		0.4690	C	0	0	0	0	0	0	0	0	0	0	0	0	0	0



5.0936	-2.6107	-2.6805 H	0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0
5.3026	-0.8722	-2.4163 H	0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0
6.7204	-1.9244	-2.5892 H	0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0
1.0262	-4.1093	2.5607 H	0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0
1.3218	-2.7380	3.6546 H	0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0
1.6367	0.5960	-5.4076 H	0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0
2.5092	-0.9436	-5.4793 H	0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0
3.3674	0.5417	-5.0469 H	0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0
-2.2626	0.2453	4.3112 H	0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0
-3.3638	1.3197	3.4359 H	0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0
-3.6749	-0.4202	3.4761 H	0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0
4.6352	4.4916	2.6525 H	0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0
3.5666	5.7710	2.0581 H	0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0
5.3205	5.8930	1.8156 H	0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0
1 28	1 0 0 0 0		
1 58	1 0 0 0 0		
1 59	1 0 0 0 0		
1 60	1 0 0 0 0		
2 12	4 0 0 0 0		
2 28	1 0 0 0 0		
2 32	4 0 0 0 0		
3 19	4 0 0 0 0		
3 23	4 0 0 0 0		
3 37	1 0 0 0 0		
4 31	4 0 0 0 0		
4 32	4 0 0 0 0		
4 52	1 0 0 0 0		
5 14	4 0 0 0 0		
5 33	4 0 0 0 0		
5 45	1 0 0 0 0		
6 7	4 0 0 0 0		
6 16	4 0 0 0 0		
6 61	1 0 0 0 0		
7 18	4 0 0 0 0		
7 46	1 0 0 0 0		
8 9	1 0 0 0 0		
8 62	1 0 0 0 0		
8 63	1 0 0 0 0		
8 64	1 0 0 0 0		
9 24	1 0 0 0 0		
9 26	1 0 0 0 0		
9 27	1 0 0 0 0		
10 19	4 0 0 0 0		
10 29	4 0 0 0 0		
10 65	1 0 0 0 0		
11 16	4 0 0 0 0		
11 21	4 0 0 0 0		
11 66	1 0 0 0 0		
12 30	4 0 0 0 0		
12 67	1 0 0 0 0		
13 14	4 0 0 0 0		
13 27	4 0 0 0 0		
13 68	1 0 0 0 0		
14 48	1 0 0 0 0		
15 27	4 0 0 0 0		
15 33	4 0 0 0 0		
15 52	1 0 0 0 0		
16 41	1 0 0 0 0		
17 23	4 0 0 0 0		
17 29	4 0 0 0 0		
17 69	1 0 0 0 0		
18 21	4 0 0 0 0		
18 53	1 0 0 0 0		
19 48	1 0 0 0 0		
20 28	1 0 0 0 0		
20 70	1 0 0 0 0		

20	71	1	0	0	0	0
20	72	1	0	0	0	0
21	40	1	0	0	0	0
22	28	1	0	0	0	0
22	73	1	0	0	0	0
22	74	1	0	0	0	0
22	75	1	0	0	0	0
23	46	1	0	0	0	0
24	76	1	0	0	0	0
24	77	1	0	0	0	0
24	78	1	0	0	0	0
25	42	1	0	0	0	0
25	45	1	0	0	0	0
25	79	1	0	0	0	0
25	80	1	0	0	0	0
26	81	1	0	0	0	0
26	82	1	0	0	0	0
26	83	1	0	0	0	0
29	36	1	0	0	0	0
30	31	4	0	0	0	0
30	40	1	0	0	0	0
31	57	1	0	0	0	0
32	84	1	0	0	0	0
33	57	1	0	0	0	0
34	36	1	0	0	0	0
34	85	1	0	0	0	0
34	86	1	0	0	0	0
34	87	1	0	0	0	0
35	38	1	0	0	0	0
35	53	1	0	0	0	0
35	88	1	0	0	0	0
35	89	1	0	0	0	0
36	43	1	0	0	0	0
36	50	1	0	0	0	0
37	51	1	0	0	0	0
38	55	1	0	0	0	0
38	90	1	0	0	0	0
38	91	1	0	0	0	0
39	44	1	0	0	0	0
39	92	1	0	0	0	0
39	93	1	0	0	0	0
39	94	1	0	0	0	0
41	47	1	0	0	0	0
41	49	1	0	0	0	0
41	56	1	0	0	0	0
42	54	1	0	0	0	0
42	95	1	0	0	0	0
42	96	1	0	0	0	0
43	97	1	0	0	0	0
43	98	1	0	0	0	0
43	99	1	0	0	0	0
44	51	1	0	0	0	0
44	100	1	0	0	0	0
44	101	1	0	0	0	0
47	102	1	0	0	0	0
47	103	1	0	0	0	0
47	104	1	0	0	0	0
49	105	1	0	0	0	0
49	106	1	0	0	0	0
49	107	1	0	0	0	0
50	108	1	0	0	0	0
50	109	1	0	0	0	0
50	110	1	0	0	0	0
51	111	1	0	0	0	0
51	112	1	0	0	0	0
54	113	1	0	0	0	0

54114	1	0	0	0	0
54115	1	0	0	0	0
55116	1	0	0	0	0
55117	1	0	0	0	0
55118	1	0	0	0	0
56119	1	0	0	0	0
56120	1	0	0	0	0
56121	1	0	0	0	0