

ELECTRONIC SUPPORTING INFORMATION

A New Tool for Photoaffinity Labeling Studies: a Conformationally Rigidified, Benzophenone Based, α -Amino Acid

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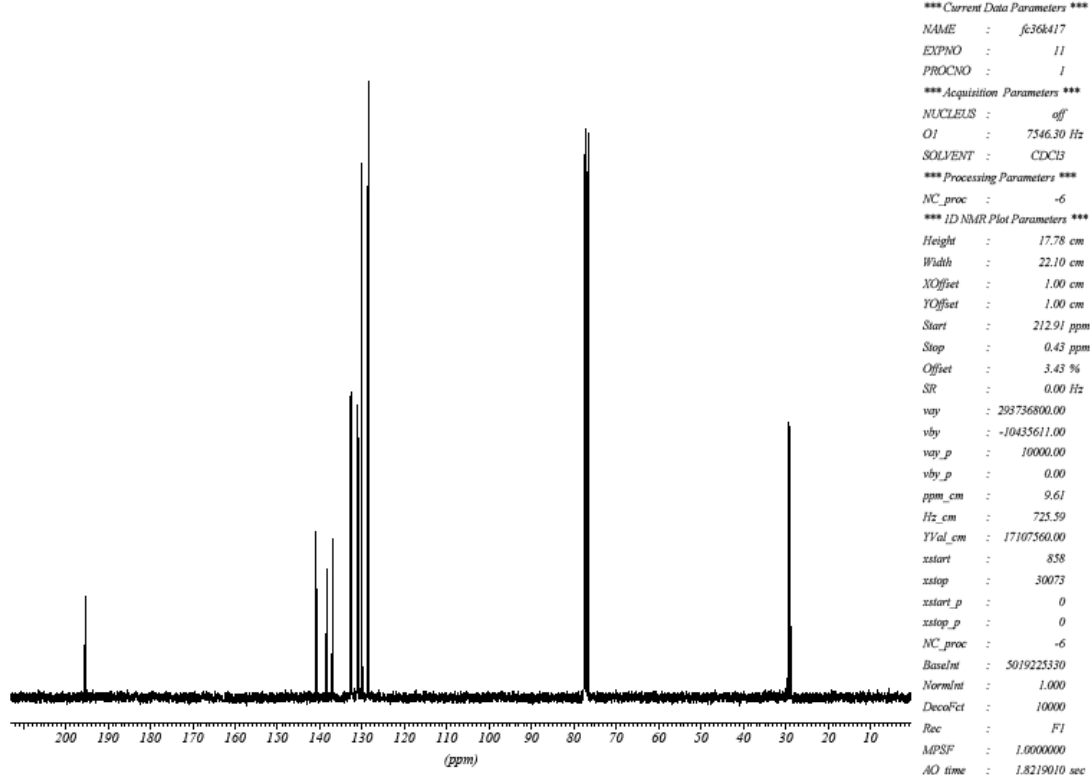
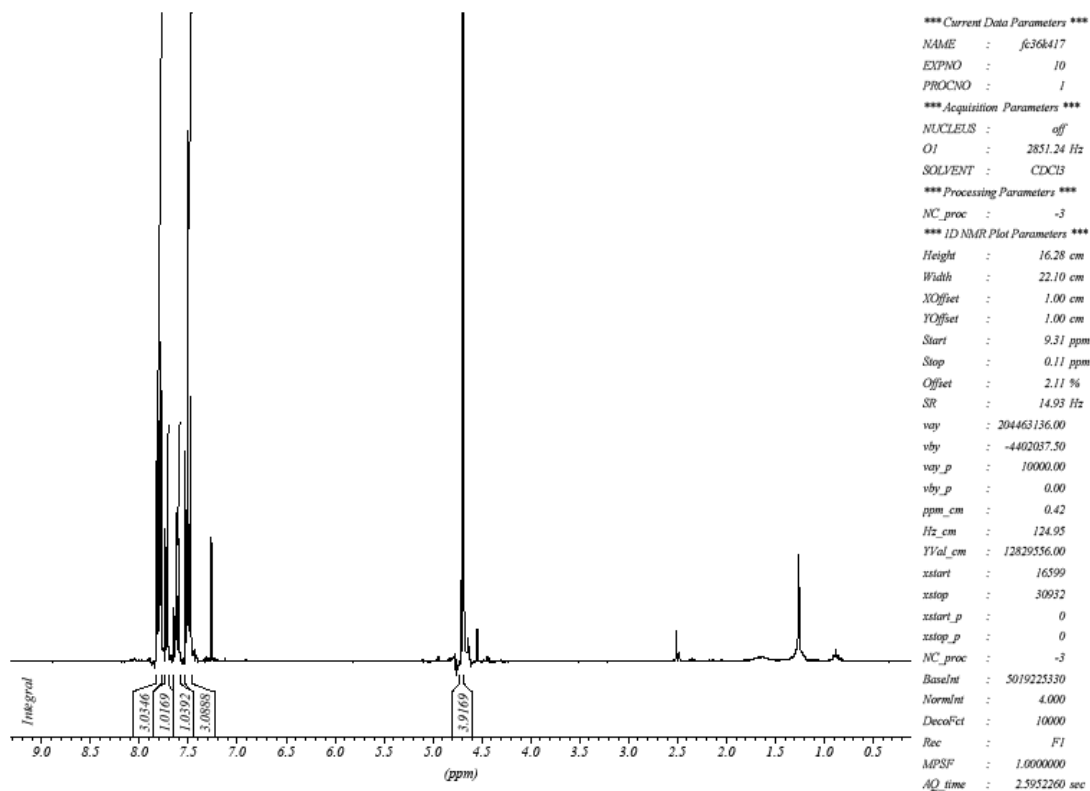
TLC



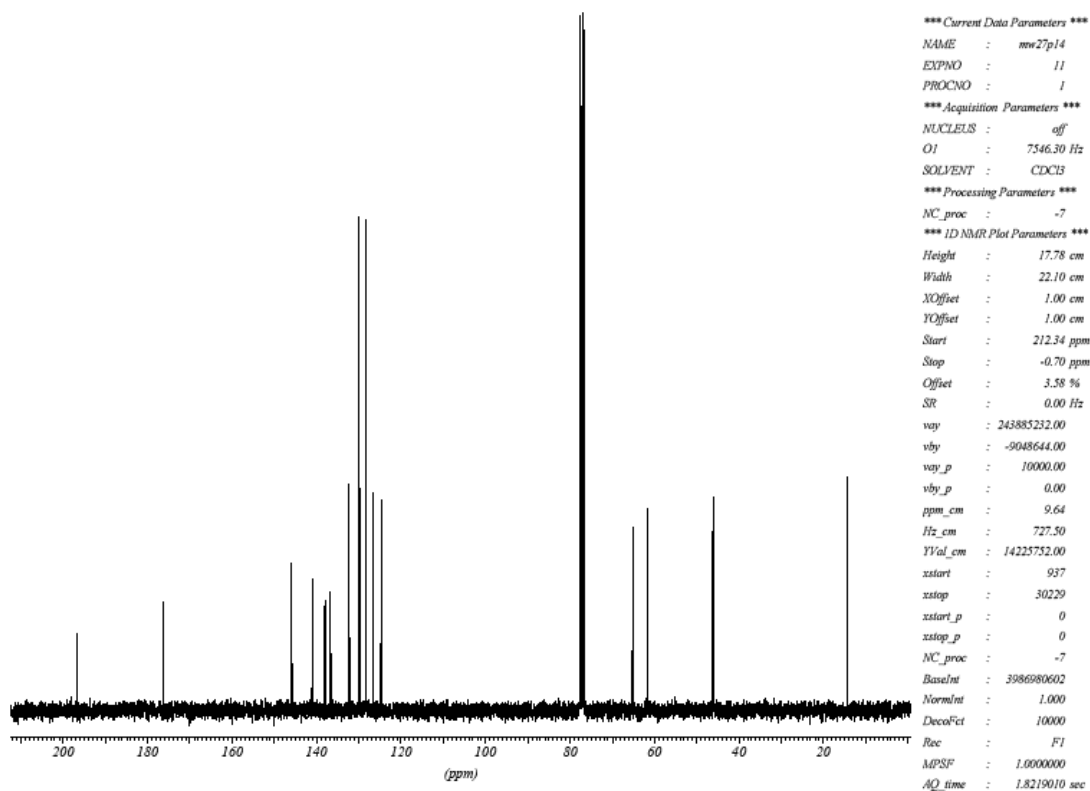
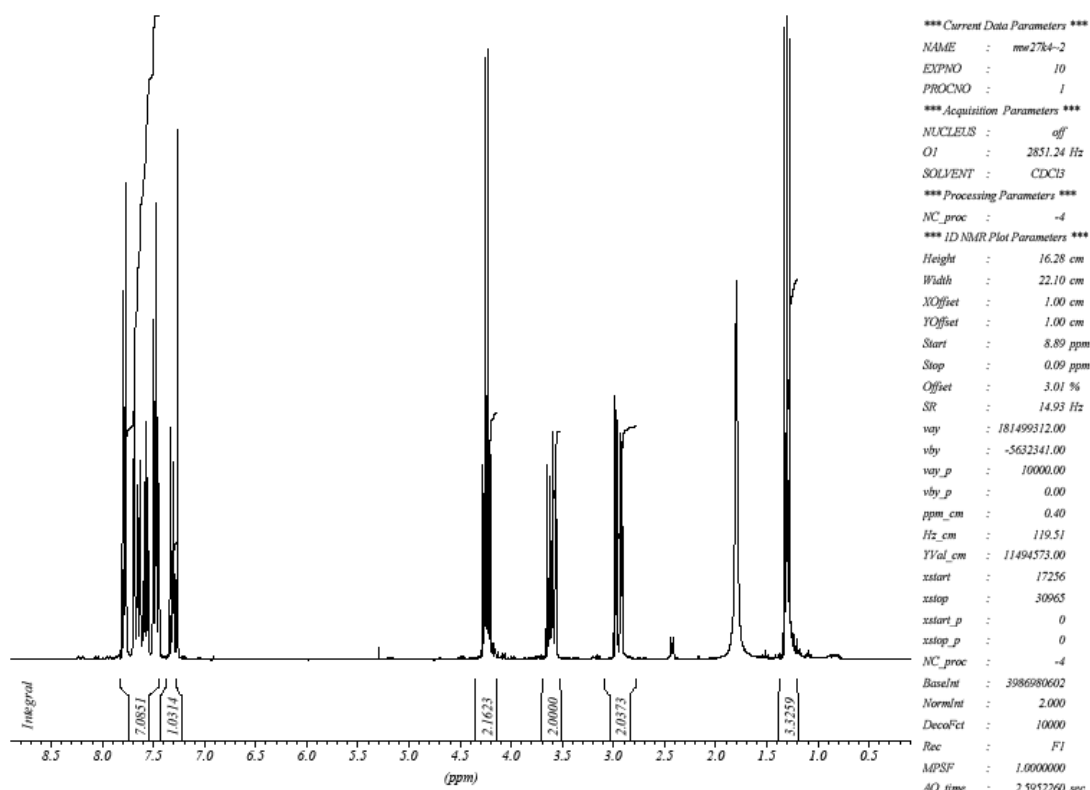
TLC of Z-(*RS*)-BpAib-(*S*)-Phe-NHChx separation. (Kieselgel F 254, Merck).
Eluent : Cyclohexane : Ethyl acetate 1 : 1.
Left lane : Z-(*S*)-BpAib-(*S*)-Phe-NHChx **7a** Rf 0.45
Middle lane : Z-(*RS*)-BpAib-(*S*)-Phe-NHChx (artificial mixture)
Right lane : Z-(*R*)-BpAib-(*S*)-Phe-NHChx **8a** Rf 0.37

¹H and ¹³C NMR spectra

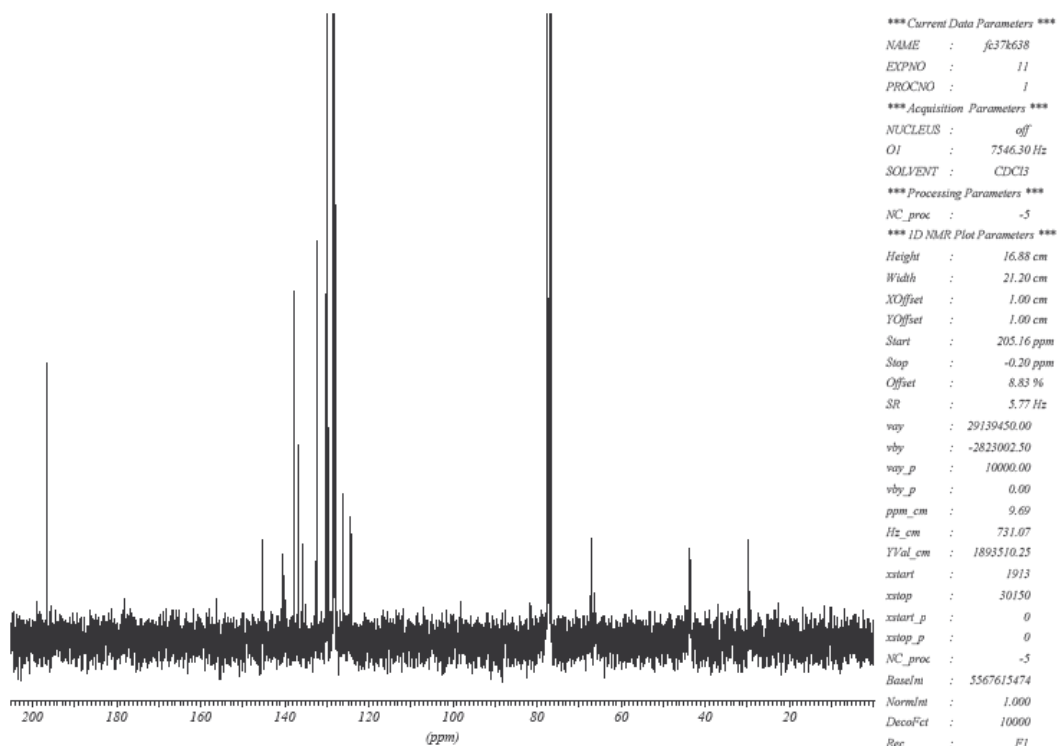
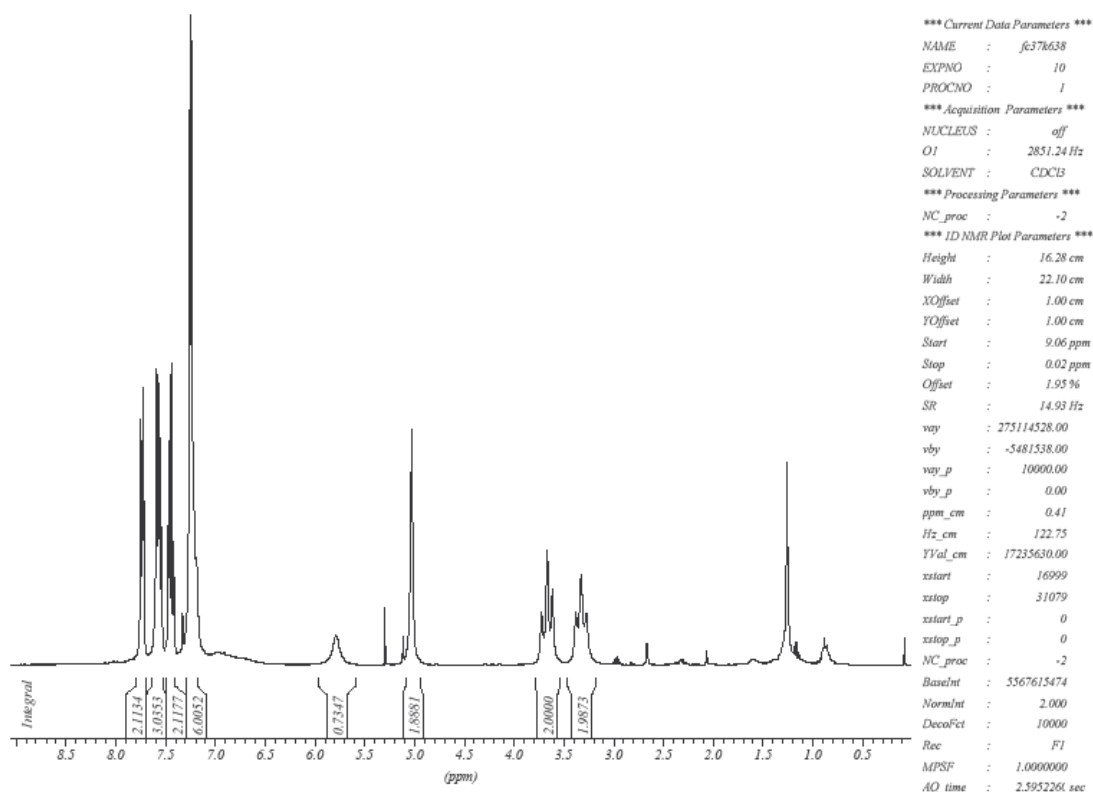
3,4-*bis*-(bromomethyl)benzophenone **3**



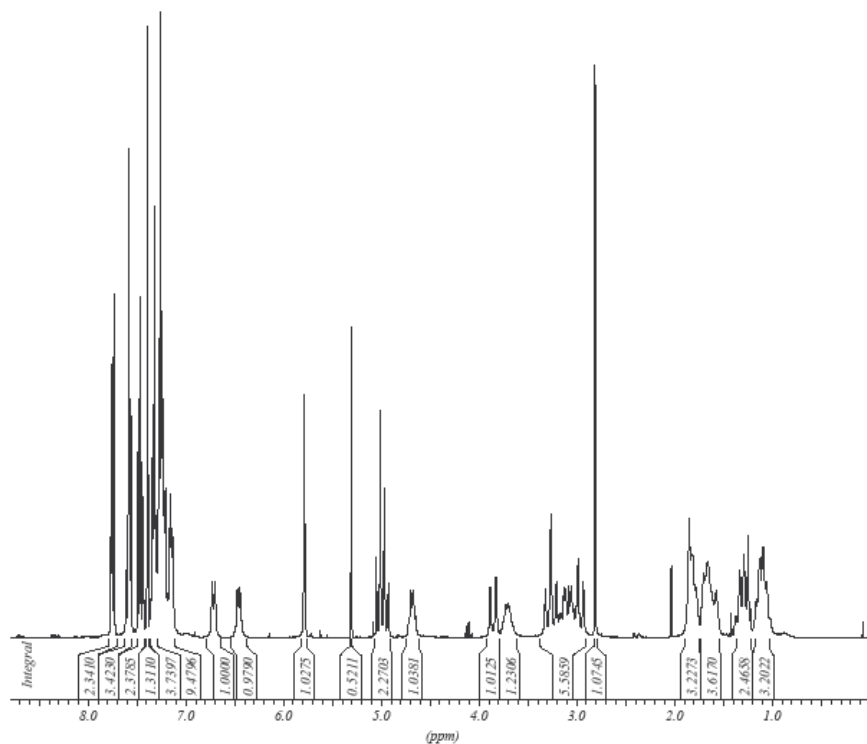
H-(RS)-BpAib-OEt 4



Z-(RS)-BpAib-OH 5

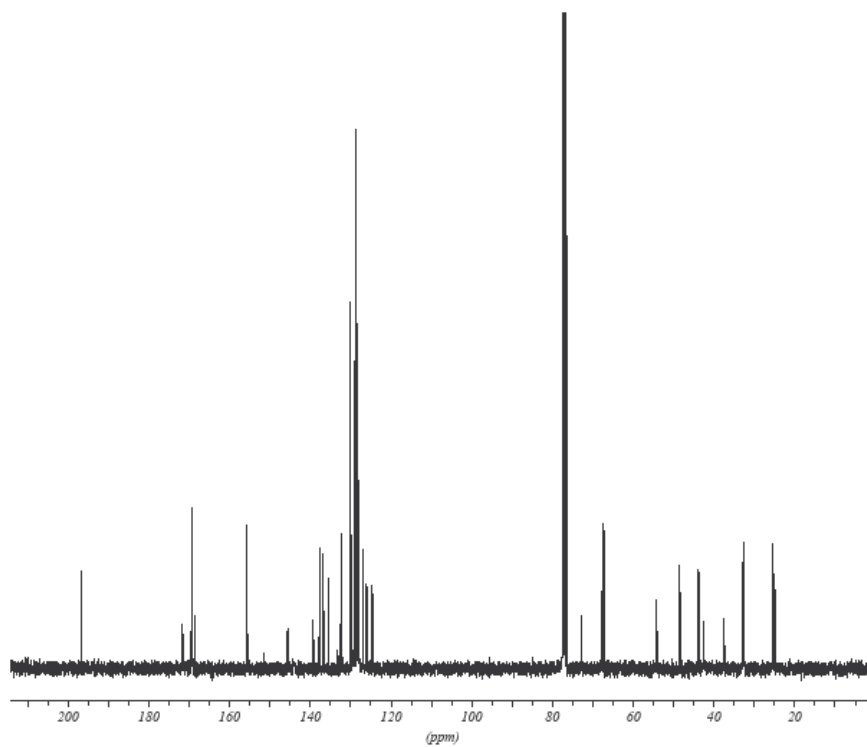


Z-(S)-BpAib-(S)-Phe-NHChx **7a**



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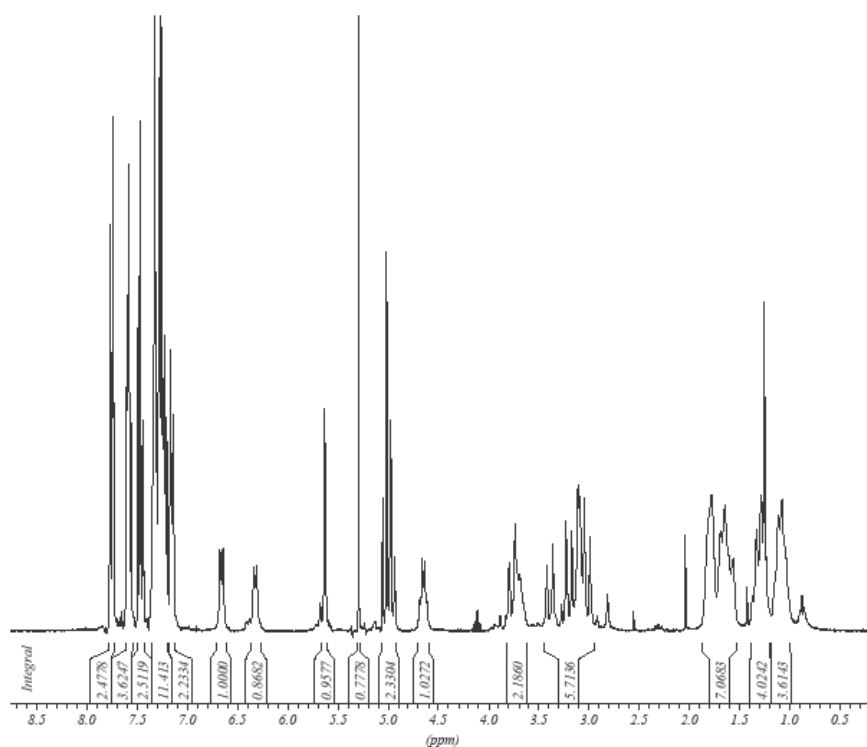
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SOLVENT  : CDCl3
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Start    : 8.79 ppm
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SR       : 14.93 Hz
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YVal_cm  : 17021338.00
xstart   : 17412
xstop    : 31168
xstart_p : 0
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NormInt  : 1.000
DecoFct  : 10000
Rec      : F1
    
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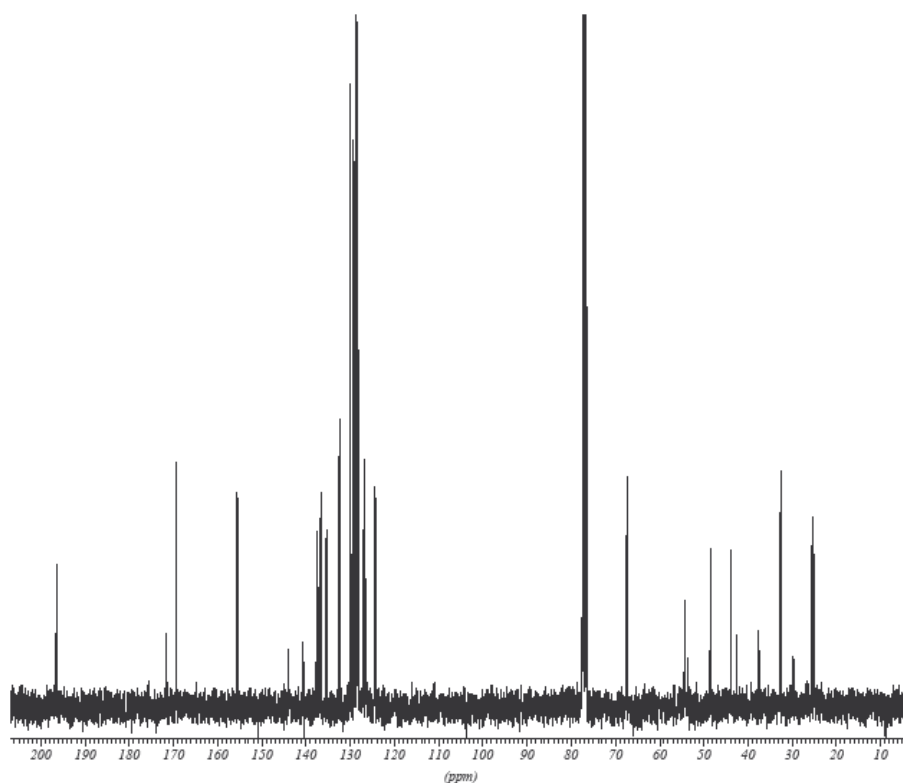
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SOLVENT  : CDCl3
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YVal_cm  : 13518571.00
xstart   : 663
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Z-(R)-BpAib-(S)-Phe-NHChx **8a**



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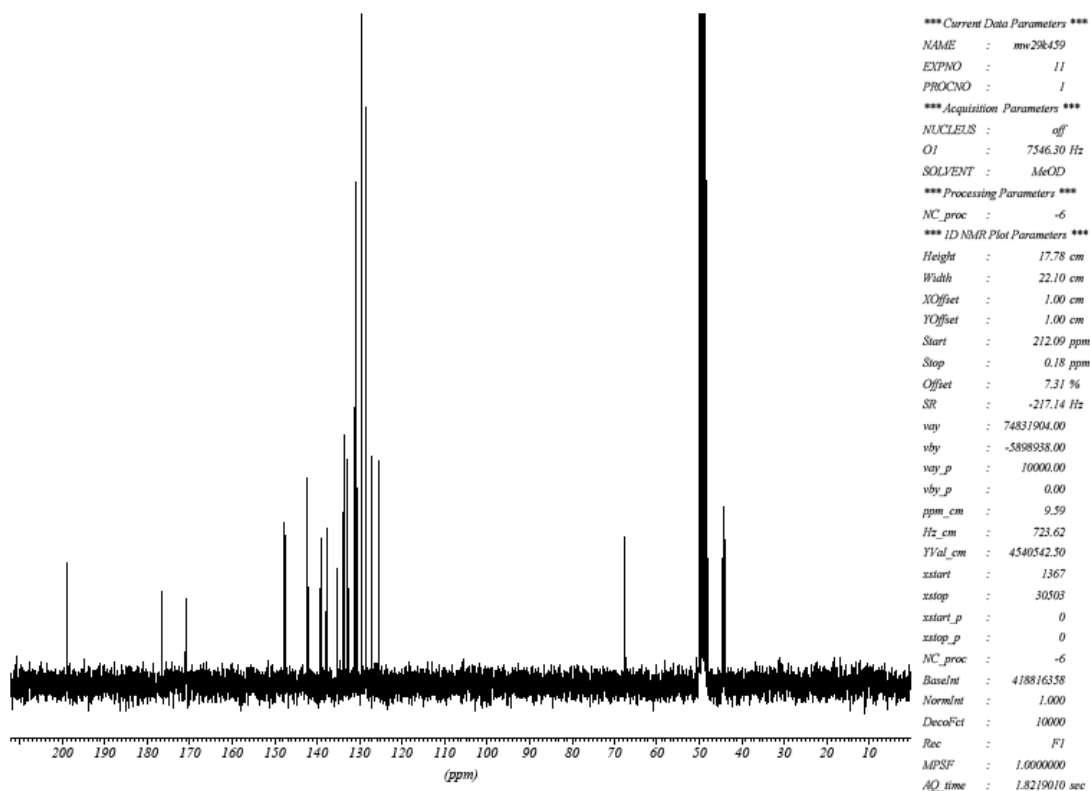
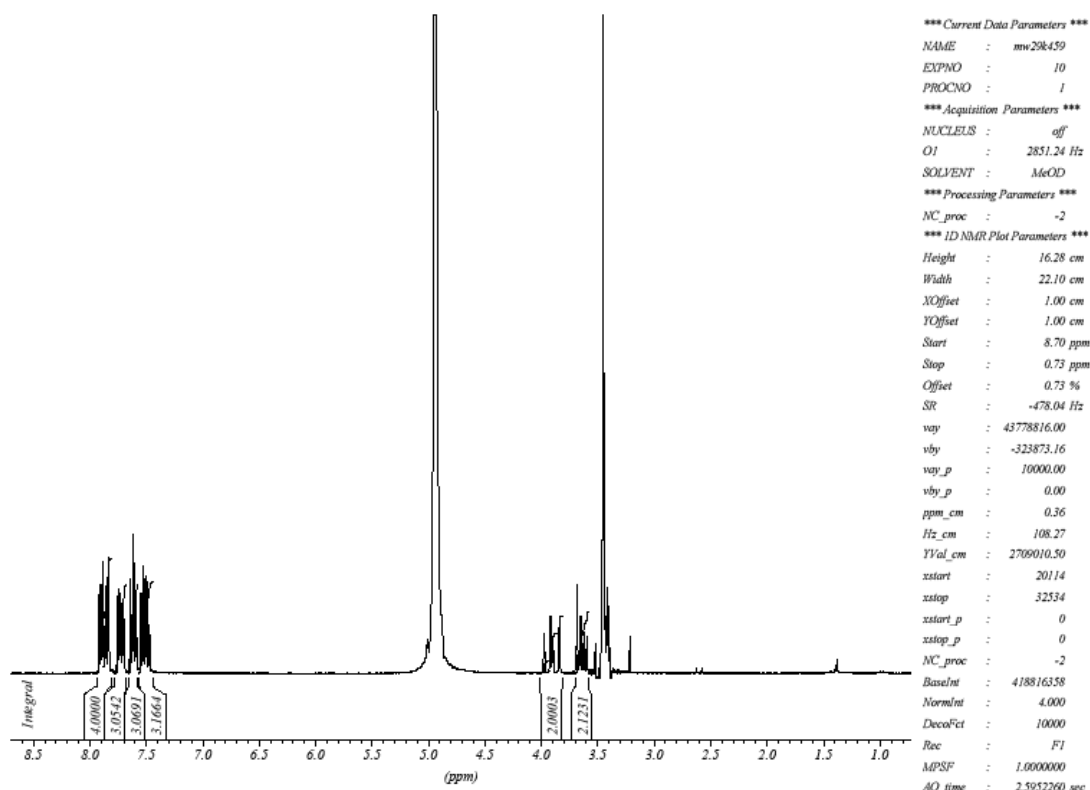
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EXPNO    : 10
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vby      : -5529252.00
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ppm_cm   : 0.41
Hz_cm    : 121.75
YVal_cm  : 19066386.00
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xstart_p : 0
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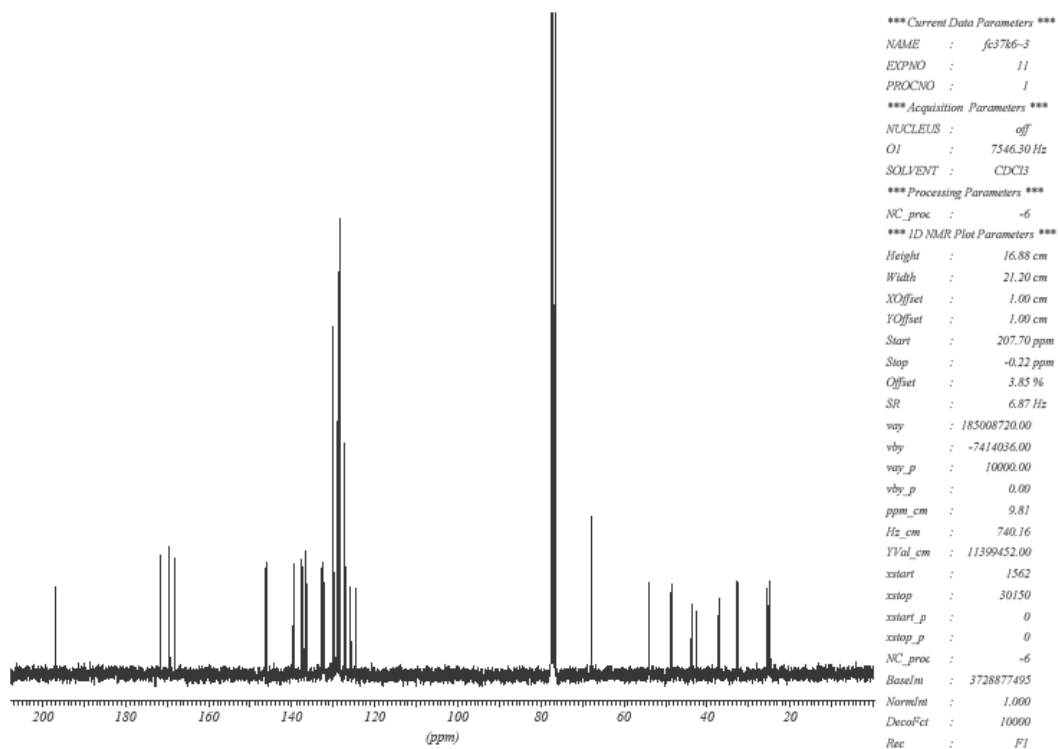
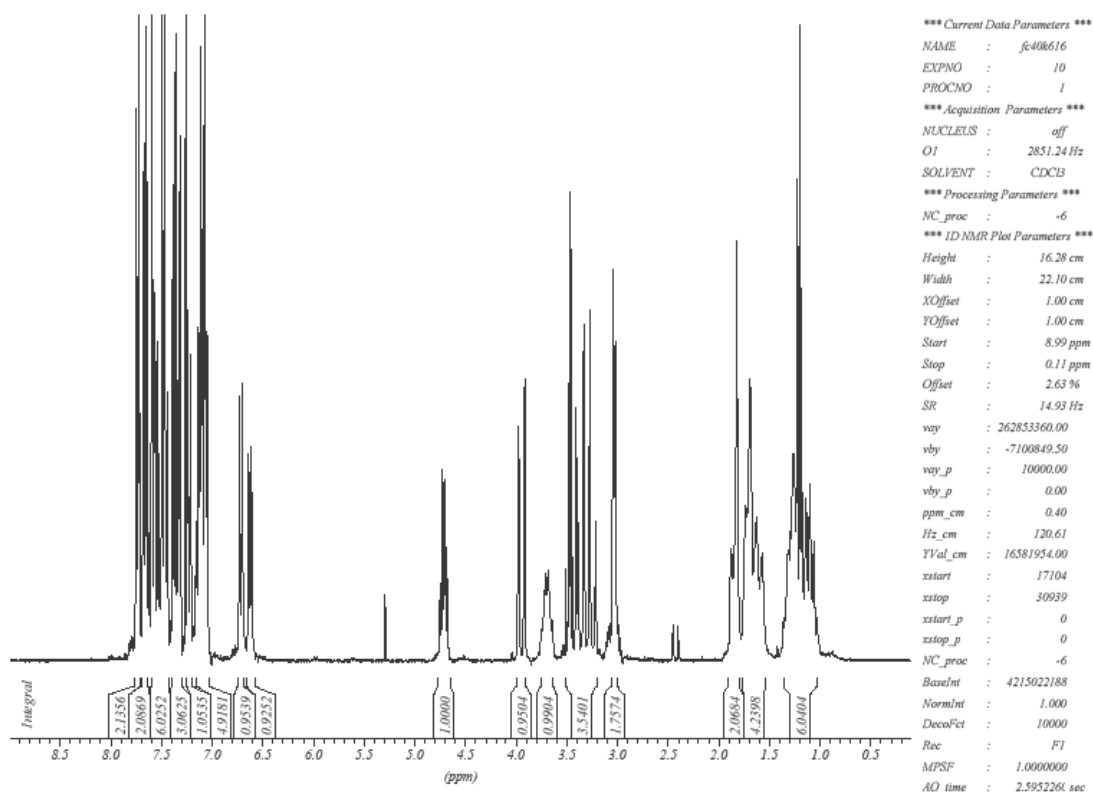
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*** Current Data Parameters ***
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NUCLEUS  : off
OI       : 7546.30 Hz
SOLVENT  : CDCl3
*** Processing Parameters ***
NC_proc  : -6
*** 1D NMR Plot Parameters ***
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Stop     : 3.18 ppm
Offset   : 4.55 %
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Hz_cm    : 695.51
YVal_cm  : 5445058.00
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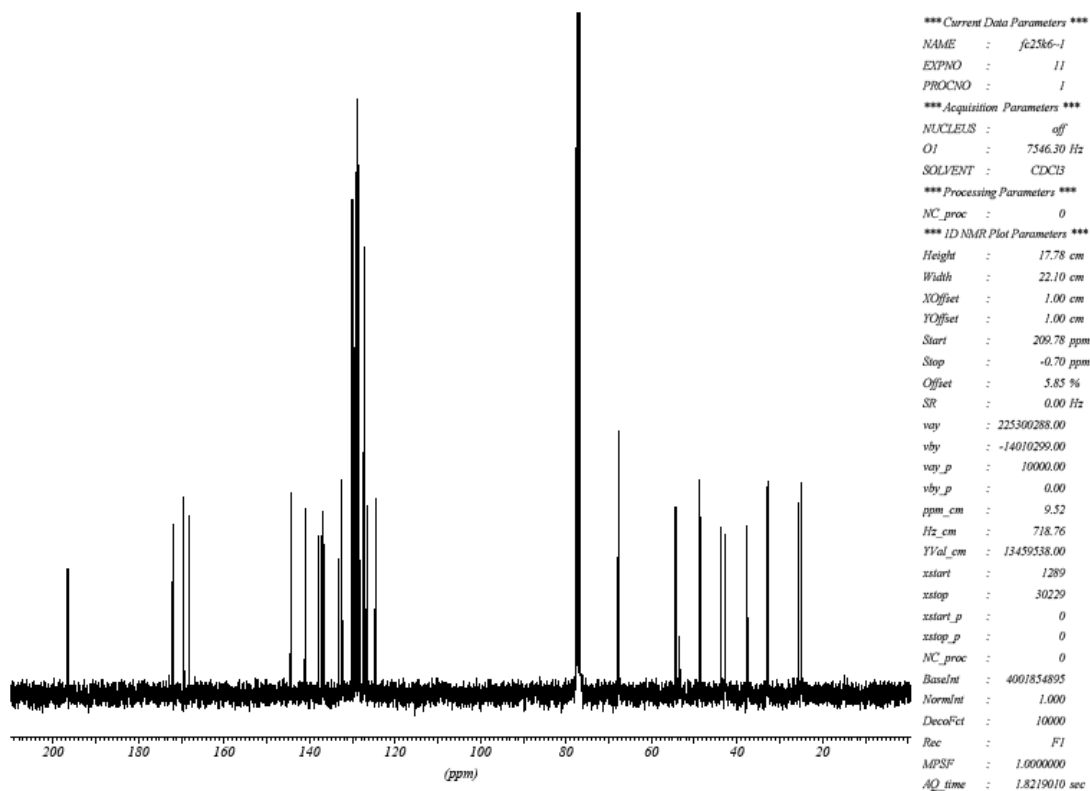
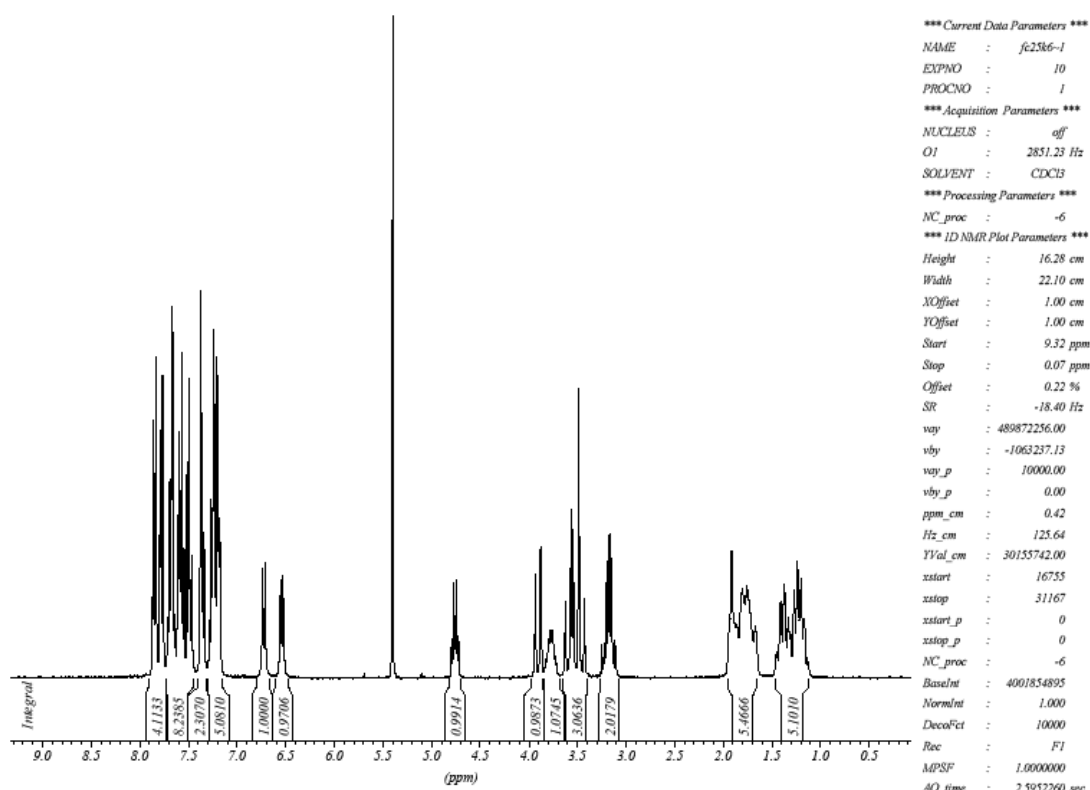
Bz-(*RS*)-BpAib-OH 6



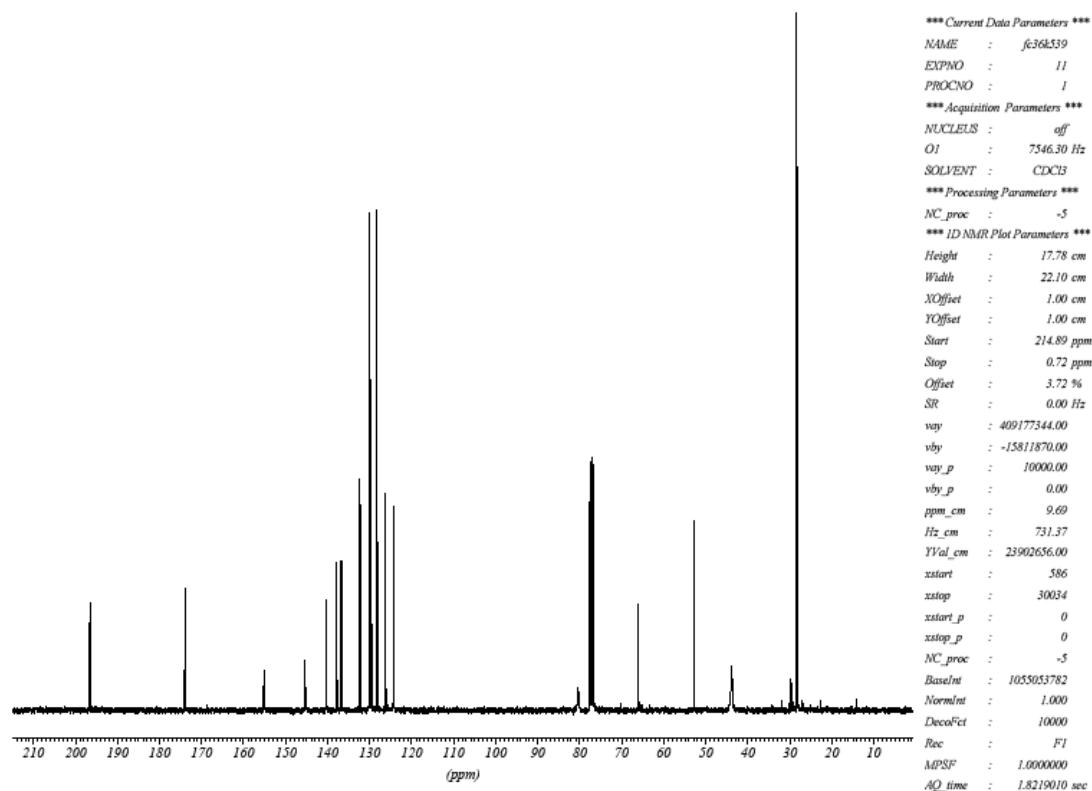
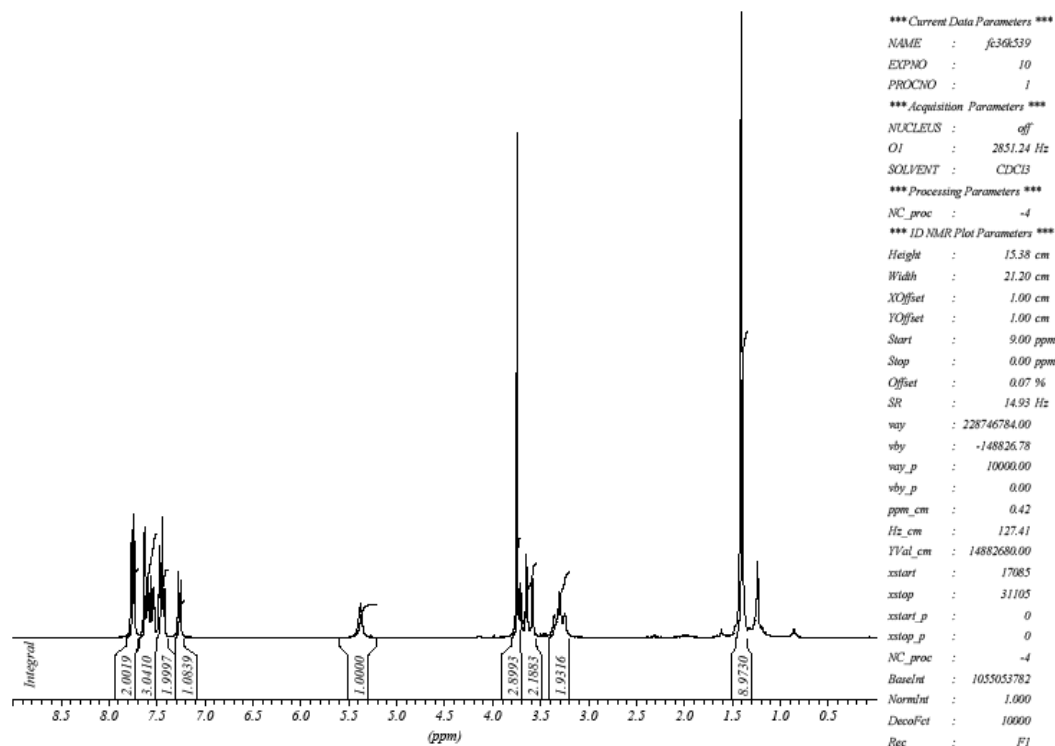
Bz-(S)-BpAib-(S)-Phe-NHChx **7b**



Bz-(R)-BpAib-(S)-Phe-NHChx **8b**



Boc-(S)-BpAib-OMe 9



X-Ray diffraction data of 8a

Single crystals of *Z-(R)*-BpAib-*(S)*-Phe-NHChx were grown by slow evaporation from a dichloromethane – methanol solution. Diffraction data were collected at $T = 293(2)$ K with $\text{CuK}\alpha$ radiation ($\lambda = 1.54178$ Å) using a Philips PW 1100 diffractometer in the $\theta - 2\theta$ scan mode up to $\theta = 55^\circ$. The crystal did not significantly diffract at higher resolution. Cell parameters were obtained by least-squares refinement of the angular settings of 48 carefully centered reflections in the $12 - 18^\circ$ θ range. Three standard reflections, periodically monitored, showed an intensity decay that reached 15% at the end of data collection. Intensities were re-scaled accordingly. Intensities were corrected for Lorentz and polarization effects, but not for absorption. The structure was solved by direct methods with the SIR 2002 program [S1]. The asymmetric unit is composed of one peptide molecule and one co-crystallized dichloromethane molecule. Refinement was carried out by full-matrix block least-squares procedures on F^2 , using all data, by application of the SHELXL 97 program [S2], and allowing the positional parameters and the anisotropic displacement parameters of the non-hydrogen atoms to refine at alternate cycles. All non-hydrogen atoms were refined anisotropically. A common population parameter for the atoms of the co-crystallized dichloromethane molecule refined to the value of 0.712(7). Hydrogen atoms were calculated at idealized positions and refined using a riding model.

Formula: $\text{C}_{40}\text{H}_{41}\text{N}_3\text{O}_5 \times 0.71 \text{CH}_2\text{Cl}_2$; formula weight: 703.2; orthorhombic, space group $\text{P}2_12_12_1$; unit cell parameters: $a = 10.949(3)$, $b = 14.354(3)$, $c = 24.633(5)$ Å; $V = 3871.4(15)$ Å³; $Z = 4$; $D_{\text{calcd}} = 1.207$ Mg m⁻³; crystal size: $0.50 \times 0.20 \times 0.10$ mm³; 3104 reflections collected, of which 3061 independent ($R_{\text{int}} = 0.045$); index ranges: $-1 \leq h \leq 11$, $0 \leq k \leq 15$, $0 \leq l \leq 26$; data / parameters: 3061/414; $R_1 = 0.0678$ [on $F \geq 4\sigma(F)$]; $wR_2 = 0.2070$ (on F^2 , all data); goodness of fit on F^2 : 1.019; Flack parameter 0.95(7); largest peak and hole in the final difference Fourier map: 0.413 and -0.222 e Å⁻³.

Crystallographic data (including atomic coordinates, bond distances, bond angles, torsion angles, intra- and intermolecular H-bonds parameters) have been deposited at The Cambridge Crystallographic Data Centre with deposition number CCDC-746110, and may be found in the Supporting Information as a CIF file.

Supporting references

[S1] Burla, M. C.; Camalli, M.; Carrozzini, B.; Cascarano, G. L.; Giacovazzo, C.; Polidori, G.; Spagna, R. *J. Appl. Crystallogr.* **2003**, *36*, 1103.

[S2] Sheldrick, G. M. *Acta Crystallogr.* **2008**, *A64*, 112-122.

Mass spectrum of the mixture of the two diastereomeric products (**11A** and **11B**) isolated after the photoreaction.

