

Supplementary Information

An NmrA-like enzyme-catalysed redox-mediated Diels-Alder cycloaddition with *anti*-selectivity

Zhiwen Liu^{1,11}, Sebastian Rivera^{2,11}, Sean A. Newmister^{2,11}, Jacob N. Sanders^{3, 11}, Qiuyue Nie¹, Shuai Liu¹, Fanglong Zhao¹, Joseph D. Ferrara⁴, Hao-Wei Shih¹, Siddhant Patil⁵, Weijun Xu⁵, Mitchell D. Miller⁵, George N. Phillips, Jr.^{5,6}, K. N. Houk^{3*}, David H. Sherman^{2,7,8,9*} and Xue Gao^{1,6,10*}

¹Department of Chemical and Biomolecular Engineering, Rice University, Houston, TX, USA.

²Life Sciences Institute, University of Michigan, Ann Arbor, MI, USA.

³Department of Chemistry and Biochemistry, University of California, Los Angeles, CA, USA.

⁴Rigaku Americas Corporation, 9009 New Trails Drive, The Woodlands, TX, USA.

⁵Department of Biosciences, Rice University, Houston, TX, USA.

⁶Department of Chemistry, Rice University, Houston, TX, USA.

⁷Department of Medicinal Chemistry, University of Michigan, Ann Arbor, MI, USA.

⁸Department of Microbiology & Immunology, University of Michigan, Ann Arbor, MI, USA.

⁹Department of Chemistry, University of Michigan, Ann Arbor, MI, USA.

¹⁰Department of Bioengineering, Rice University, Houston, TX, USA.

¹¹These authors contributed equally.

*Corresponding authors: xue.gao@rice.edu; davidhs@umich.edu; houk@chem.ucla.edu

Table of Contents

Supplementary Tables	S4
Supplementary Table 1 Strains and plasmids in this study.	S4
Supplementary Table 2 Primers used in this study.	S5
Supplementary Table 3 BLASTP CtdN homologs in NCBI databases.	S6
Supplementary Table 4 BLASTP CtdP homologs in NCBI databases.	S7
Supplementary Table 5 NMR Data of compound 3 (δ in ppm, J in Hz).	S8
Supplementary Table 6 NMR Data of compound 5 (δ in ppm, J in Hz).	S9
Supplementary Table 7 NMR Data of compound 6 (δ in ppm, J in Hz)	S10
Supplementary Table 8 NMR Data of compound 7 (δ in ppm, J in Hz).	S11
Supplementary Table 9 NMR Data of compound 10 (δ in ppm, J in Hz).	S12
Supplementary Table 10 NMR Data of compound R1 (δ in ppm, J in Hz).	S13
Supplementary Table 11 Crystal data and structure refinement for compound 5 .	S14
Supplementary Table 12 Crystal data and structure refinement for compound 10 .	S15
Supplementary Table 13 Crystallographic data collection and structure refinement statistics.	S16
Supplementary Table 14 Structural homologs of CtdP identified by DALI server.	S17
Supplementary Figures	S18
Supplementary Fig. 1 Biomimetic synthesis of the CtdP and MalC substrates.	S18
Supplementary Fig. 2 SDS-PAGE (12%) analysis of purified proteins.	S19
Supplementary Fig. 3 PCR confirmation of <i>ctd</i> mutants.	S20
Supplementary Fig. 4 UV and mass spectra of compounds 1-6 .	S21
Supplementary Fig. 5 UV and mass spectra of compounds 7-11 .	S22
Supplementary Fig. 6 Electronic circular dichroism spectra of 2 , 5 , and 6 .	S23
Supplementary Fig. 7 Kinetic analysis of CtdP, NADP ⁺ content in the purified CtdP protein, and refold CtdP assays.	S24
Supplementary Fig. 8 The proposed pathways of spontaneous Diels-Alder reactions of compound 3 .	S25
Supplementary Fig. 9-1 ¹ H NMR spectrum of 3 in DMSO- <i>d</i> ₆ .	S26
Supplementary Fig. 9-2 ¹³ C NMR spectrum of 3 in DMSO- <i>d</i> ₆ .	S27
Supplementary Fig. 9-3 DEPT135 and ¹³ C NMR spectra of 3 in DMSO- <i>d</i> ₆ .	S28
Supplementary Fig. 9-4 ¹ H- ¹ H COSY NMR spectrum of 3 in DMSO- <i>d</i> ₆ .	S29
Supplementary Fig. 9-5 HSQC NMR spectrum of 3 in DMSO- <i>d</i> ₆ .	S30
Supplementary Fig. 9-6 HMBC NMR spectrum of 3 in DMSO- <i>d</i> ₆ .	S31
Supplementary Fig. 9-7 HRMS spectrum of 3 .	S32
Supplementary Fig. 10-1 ¹ H NMR spectrum of 5 in CDCl ₃ .	S33
Supplementary Fig. 10-2 ¹³ C NMR spectrum of 5 in CDCl ₃ .	S34
Supplementary Fig. 10-3 DEPT135 and ¹³ C NMR spectra of 5 in CDCl ₃ .	S35
Supplementary Fig. 10-4 ¹ H- ¹ H COSY NMR spectrum of 5 in CDCl ₃ .	S36
Supplementary Fig. 10-5 HSQC NMR spectrum of 5 in CDCl ₃ .	S37
Supplementary Fig. 10-6 HMBC NMR spectrum of 5 in CDCl ₃ .	S38
Supplementary Fig. 10-7 NOESY NMR spectrum of 5 in CDCl ₃ .	S39
Supplementary Fig. 10-8 HRMS spectrum of 5 .	S40
Supplementary Fig. 11-1 ¹ H NMR spectrum of 6 in CD ₃ CN.	S41
Supplementary Fig. 11-2 ¹³ C NMR spectrum of 6 in CD ₃ CN.	S42
Supplementary Fig. 11-3 DEPT135 and ¹³ C NMR spectra of 6 in CD ₃ CN.	S43
Supplementary Fig. 11-4 ¹ H- ¹ H COSY NMR spectrum of 6 in CD ₃ CN.	S44
Supplementary Fig. 11-5 HSQC NMR spectrum of 6 in CD ₃ CN.	S45
Supplementary Fig. 11-6 HMBC NMR spectrum of 6 in CD ₃ CN.	S46

Supplementary Fig. 11-7 NOESY NMR spectrum of 6 in CD ₃ CN.	S47
Supplementary Fig. 11-8 HRMS spectrum of 6 .	S48
Supplementary Fig. 12-1 ¹ H NMR spectrum of 7 in DMSO- <i>d</i> ₆ .	S49
Supplementary Fig. 12-2 ¹³ C NMR spectrum of 7 in DMSO- <i>d</i> ₆ .	S50
Supplementary Fig. 12-3 DEPT135 and ¹³ C NMR spectra of 7 in DMSO- <i>d</i> ₆ .	S51
Supplementary Fig. 12-4 ¹ H- ¹ H COSY NMR spectrum of 7 in DMSO- <i>d</i> ₆ .	S52
Supplementary Fig. 12-5 HSQC NMR spectrum of 7 in DMSO- <i>d</i> ₆ .	S53
Supplementary Fig. 12-6 HMBC NMR spectrum of 7 in DMSO- <i>d</i> ₆ .	S54
Supplementary Fig. 12-7 HRMS spectrum of 7 .	S55
Supplementary Fig. 13-1 ¹ H NMR spectrum of 9 in DMSO- <i>d</i> ₆ .	S56
Supplementary Fig. 13-2 ¹³ C NMR spectrum of 9 in DMSO- <i>d</i> ₆ .	S57
Supplementary Fig. 14-1 ¹ H NMR spectrum of 10 in CD ₃ CN.	S58
Supplementary Fig. 14-2 ¹³ C NMR spectrum of 10 in CD ₃ CN.	S59
Supplementary Fig. 14-3 DEPT135 and ¹³ C NMR spectra of 10 in CD ₃ CN.	S60
Supplementary Fig. 14-4 ¹ H- ¹ H COSY NMR spectrum of 10 in CD ₃ CN.	S61
Supplementary Fig. 14-5 HSQC NMR spectrum of 10 in CD ₃ CN.	S62
Supplementary Fig. 14-6 HMBC NMR spectrum of 10 in CD ₃ CN.	S63
Supplementary Fig. 14-7 NOESY NMR spectrum of 10 in CD ₃ CN.	S64
Supplementary Fig. 14-8 HRMS spectrum of 10 .	S65
Supplementary Fig. 15-1 ¹ H NMR spectrum of 11 in CD ₃ OD/CDCl ₃ (10/1).	S66
Supplementary Fig. 15-2 ¹³ C NMR spectrum of 11 in CD ₃ OD/CDCl ₃ (10/1).	S67
Supplementary Fig. 16-1 ¹ H NMR spectrum of S7 in DMSO- <i>d</i> ₆ .	S68
Supplementary Fig. 16-2 ¹³ C NMR spectrum of S7 in DMSO- <i>d</i> ₆ .	S69
Supplementary Fig. 16-3 DEPT135 and ¹³ C NMR spectra of S7 in DMSO- <i>d</i> ₆ .	S70
Supplementary Fig. 16-4 ¹ H- ¹ H COSY NMR spectrum of S7 in DMSO- <i>d</i> ₆ .	S71
Supplementary Fig. 16-5 HSQC NMR spectrum of S7 in DMSO- <i>d</i> ₆ .	S72
Supplementary Fig. 16-6 HMBC NMR spectrum of S7 in DMSO- <i>d</i> ₆ .	S73
Supplementary Fig. 17-1 ¹ H NMR spectrum of R1 in CDCl ₃ .	S74
Supplementary Fig. 17-2 ¹³ C NMR spectrum of R1 in CDCl ₃ .	S75
Supplementary Fig. 17-3 DEPT135 and ¹³ C NMR spectra of R1 in CDCl ₃ .	S76
Supplementary Fig. 17-4 ¹ H- ¹ H COSY NMR spectrum of R1 in CDCl ₃ .	S77
Supplementary Fig. 17-5 HSQC-TOCSY NMR spectrum of R1 in CDCl ₃ .	S78
Supplementary Fig. 17-6 HMBC NMR spectrum of R1 in CDCl ₃ .	S79
Supplementary Fig. 17-7 NOESY NMR spectrum of R1 in CDCl ₃ .	S80
Supplementary Fig. 17-8 HRMS spectrum of R1 .	S81
Energies and molecular coordinates of calculated structures	S82
Supplementary References	S102

Supplementary Tables

Supplementary Table 1 | Strains and plasmids in this study.

Strain or plasmid	Characteristics	Source
Strains		
<i>E.coli</i> TOP 10	General cloning host strain	Invitrogen
<i>E.coli</i> BL21 (DE3)	Protein production host strain	StrataGene
<i>Penicillin citrinum</i> ATCC 9849	Wild-type <i>P. citrinum</i> used in this study	ATCC
Δ <i>ctdP</i>	The <i>ctdP</i> knockout mutant of <i>P. citrinum</i> ATCC 9849	This study
Δ <i>ctdR</i>	The <i>ctdR</i> knockout mutant of <i>P. citrinum</i> ATCC 9849	This study
Δ <i>ctdO</i>	The <i>ctdO</i> knockout mutant of <i>P. citrinum</i> ATCC 9849	This study
Δ <i>ctdN</i>	The <i>ctdN</i> knockout mutant of <i>P. citrinum</i> ATCC 9849	This study
Plasmids		
Modified pETDuet-1	Amp ^r , vector for protein expression	This study
pUC57	Amp ^r , vector for gene cloning	Addgene
<i>ctdP-KO-P</i>	Amp ^r , gene knockout plasmid used for Δ <i>ctdP</i> mutant construction	This study
<i>ctdR-KO-P</i>	Amp ^r , gene knockout plasmid used for Δ <i>ctdR</i> mutant construction	This study
<i>ctdO-KO-P</i>	Amp ^r , gene knockout plasmid used for Δ <i>ctdO</i> mutant construction	This study
pET- <i>ctdR</i>	Amp ^r , vector for CtdR expression	This study
pET- <i>ctdO</i>	Amp ^r , vector for CtdO expression	This study
pET- <i>malC</i>	Amp ^r , vector for MalC expression	This study
pET- <i>ctdP</i>	Amp ^r , vector for CtdP expression	This study
pET- <i>ctdP</i> (Q118A)	Amp ^r , vector for CtdP (Q118A) expression	This study
pET- <i>ctdP</i> (V133A)	Amp ^r , vector for CtdP (V133A) expression	This study
pET- <i>ctdP</i> (L134A)	Amp ^r , vector for CtdP (L134A) expression	This study
pET- <i>ctdP</i> (W160A)	Amp ^r , vector for CtdP (W160A) expression	This study
pET- <i>ctdP</i> (Y161A)	Amp ^r , vector for CtdP (Y161A) expression	This study
pET- <i>ctdP</i> (N164A)	Amp ^r , vector for CtdP (N164A) expression	This study
pET- <i>ctdP</i> (F170A)	Amp ^r , vector for CtdP (F170A) expression	This study
pET- <i>ctdP</i> (E173A)	Amp ^r , vector for CtdP (E173A) expression	This study
pET- <i>ctdP</i> (S273A)	Amp ^r , vector for CtdP (S273A) expression	This study
pET- <i>ctdP</i> (F277A)	Amp ^r , vector for CtdP (F277A) expression	This study
pET- <i>ctdP</i> (Y269A)	Amp ^r , vector for CtdP (Y269A) expression	This study
pET- <i>ctdP</i> (Y280A)	Amp ^r , vector for CtdP (Y280A) expression	This study
pET- <i>ctdP</i> (Y280F)	Amp ^r , vector for CtdP (Y280F) expression	This study
pET- <i>ctdP</i> (S340A)	Amp ^r , vector for CtdP (S340A) expression	This study
pET- <i>ctdP</i> (P342A)	Amp ^r , vector for CtdP (P342A) expression	This study
pET-SUMO- <i>ctdP</i> (Δ 330-367)	Amp ^r , vector for SUMO-CtdP (Δ 330-367) expression	This study
pET-SUMO- <i>ctdP</i> (Δ 335-367)	Amp ^r , vector for SUMO-CtdP (Δ 335-367) expression	This study
pET-SUMO- <i>ctdP</i> (Δ 342-367)	Amp ^r , vector for SUMO-CtdP (Δ 342-367) expression	This study
pET-SUMO- <i>ctdP</i> (Δ 345-367)	Amp ^r , vector for SUMO-CtdP (Δ 345-367) expression	This study
pET- <i>BmGDH</i>	Amp ^r , vector for BmGDH expression	This study

Supplementary Table 2 | Primers used in this study.

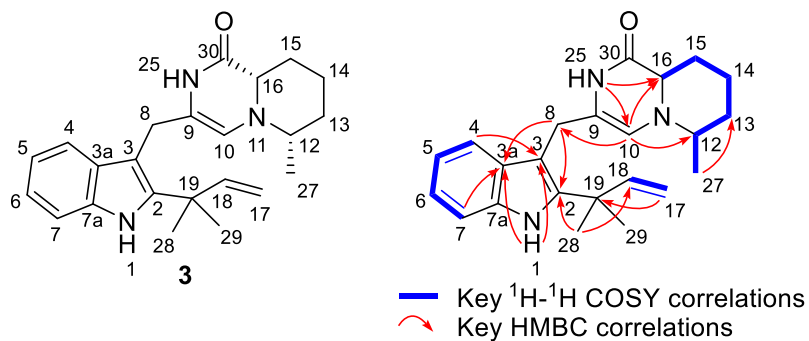
Primer	Sequence (5'-3')	Description
<i>bar-R</i>	TCATCGCAAGACCGCAACAGGATTCATC	For colony PCR verification
<i>bar-F</i>	CATACCTTCTTAAGTTCGCCCTTCCTCCCT	mutant
<i>ctdP-up-F</i>	ATGGCGGGTTCCTTTTACTTAAGAGTCCG	For colony PCR verification of Δ <i>ctdP</i>
<i>ctdP-dn-R</i>	AGTCTGTCCGTTTGAGACTGGGAGCAGATG	
<i>ctdO-up-F</i>	GGGGGAAAACCTGCGGAGCTTTTCAAGACTG	For colony PCR verification of Δ <i>ctdO</i>
<i>ctdO-dn-R</i>	GTGACTCCTGGAGGTAAGGACGGTCCGCGAG	
<i>ctdR-up-F</i>	CTGATGTTGATTAGACCCAGCCTGGAGGTC	For colony PCR verification of Δ <i>ctdR</i>
<i>ctdR-dn-R</i>	CCAGGGTACCGTTCAGGGCATTCCAATCCG	
<i>ctdN-up-F</i>	TACATGGGGATCACCAGGAG	For colony PCR verification of Δ <i>ctdN</i>
<i>ctdN-dn-R</i>	CAACGAACTGTAATGGAGCG	
<i>ctdN-Exp-F</i>	CTTTAAGAAGGAGATATACCATGACCCAGGGTGTAAAGAAA	Plasmid construction for CtdN expression
<i>ctdN-Exp-R</i>	TTAGTGATGGTGGTGGTGTAGTCAAAAAGGAACCCGCCATT	
<i>ctdP-Exp-F</i>	CTTTAAGAAGGAGATATACCATGACACACGAGATTAACAAAC	Plasmid construction for CtdP expression
<i>ctdP-Exp-R</i>	TTAGTGATGGTGGTGGTGTAGTTCACCAGGATCTTCTCTGAC	
<i>ctdO-Exp-F</i>	CTTTAAGAAGGAGATATACCATGACAAATCCAACAAAAATC	Plasmid construction for CtdO expression
<i>ctdO-Exp-R</i>	TTAGTGATGGTGGTGGTGTAGTGTGTCAAGTTACCCTTA	
<i>ctdR-Exp-F</i>	CTTTAAGAAGGAGATATACCATGACTGTTGAAAGAAAGATTG	Plasmid construction for CtdR expression
<i>ctdR-Exp-R</i>	TAGTGATGGTGGTGGTGTAGGGAAGCATTGAGACCGCTTAAAA	
<i>ctdP(Q118A)-Exp-F</i>	TCCGATGCGCTGTCTGGCGGCAAAATTAACACCCAGTTCTG	Plasmid construction for CtdP (Q118A) expression
<i>ctdP(Q118A)-Exp-R</i>	GCCGCCAGACACGCGCATGCGGATGAGGGTCCGAGCTGAACACGAC	
<i>ctdP(V133A)-Exp-F</i>	GTGAAAAGCCTGGGGCGAGTCTTGGGGACGAGCTTGTCCACACA	Plasmid construction for CtdP (V133A) expression
<i>ctdP(V133A)-Exp-R</i>	AGACTCGCCCCAGGCTTTCACGTCCAGTGTGGGGTGTAAATTT	
<i>ctdP(L134A)-Exp-F</i>	GTGAAAAGCCTGGGGCGAGTCTTGGGGACGAGCTTGTCCACACA	Plasmid construction for CtdP (L134A) expression
<i>ctdP(L134A)-Exp-R</i>	AGACTCGCCCCAGGCTTTCACGTCTGCAACTGGGGTGTAAATTT	
<i>ctdP(W160A)-Exp-F</i>	TACTTCCAGAACTTTTTTTCATTCCATCTTTCGTGGCCGAGTTTGGTGGC	Plasmid construction for CtdP (W160A) expression
<i>ctdP(W160A)-Exp-R</i>	AATGAAAAAGTCTGGAAGTATGCCGATGCCATGATCGGCGTGAAGCT	
<i>ctdP(Y161A)-Exp-F</i>	TTCCAGAACTTTTTTTCATTCCATCTTTCGTGGCCGAGTTTGGTGGCTTT	Plasmid construction for CtdP (Y161A) expression
<i>ctdP(Y161A)-Exp-R</i>	AATGAAAAAGTCTGGAATGCCACGATGCCATGATCGGCGTGAAGCT	
<i>ctdP(N164A)-Exp-F</i>	ATTCCATCTTTCGTGGCCGAGTTTGGTGGCTTTCCG	Plasmid construction for CtdP (N164A) expression
<i>ctdP(N164A)-Exp-R</i>	CTCGGCCACGAAAGATGGAATGAAAAATGCCCTGGAAGTACCACGA	
<i>ctdP(F170A)-Exp-F</i>	GAGTTTGGTGGCTTTCGGTGAATCAAGACGATGAA	Plasmid construction for CtdP (F170A) expression
<i>ctdP(F170A)-Exp-R</i>	CCACGAAAGCCACCAAACCTCGGCCACTGCAGATGGAATGAAAAA	
<i>ctdP(E173A)-Exp-F</i>	GGCTTTCGGTGAATCAAGACGATGAAAGTTATCTGACTTTGCGT	Plasmid construction for CtdP (E173A) expression
<i>ctdP(E173A)-Exp-R</i>	ATCGTCTTGATTCACGGAAGCCACAAATGCGGCCACGAAAAGATGG	
<i>ctdP(E173A)-Exp-F</i>	GGCTTTCGGTGAATCAAGACGATGAAAGTTATCTGACTTTGCGT	Plasmid construction for CtdP (E173A) expression
<i>ctdP(F174A)-Exp-R</i>	GTCTTGATTCCACGGAAGCCACTGCTCGGCCACGAAAAGATGGAAT	
<i>ctdP(S273A)-Exp-F</i>	TTCGCGTTCTATCAAATGCGTGATGGCGAACTCTTC	Plasmid construction for CtdP (S273A) expression
<i>ctdP(S273A)-Exp-R</i>	ACGCATTTGATAGAACGCGAATACCTGACGTGCCTCTGGAGGTA	
<i>ctdP(F277A)-Exp-F</i>	GCGTTCTATCAAATGCGTGATGGCGAACTC	Plasmid construction for CtdP (F277A) expression
<i>ctdP(F277A)-Exp-R</i>	ATCACGATTTGATAGAACGCTGTCTACTGACGTGACTCCTGGAGGTA	
<i>ctdP(Y269A)-Exp-F</i>	CTCCAGGATCACGTACAGTATTCGCGTTCTATCAAATGCGTGAT	Plasmid construction for CtdP (Y269A) expression
<i>ctdP(Y269A)-Exp-R</i>	GAATACCTGACGTGACTCCTGGAGTGCAGGACGGTCCGACGGCATATC	
<i>ctdP(P280A)-Exp-F</i>	CGTGATGGCGAACTTTCGGTAATGGAATTACGGAATAAAGA	Plasmid construction for CtdP (Y280A and Y280F) expression
<i>ctdP(P280A)-Exp-R</i>	CCGAAAGAGTTCGCCATCACGCATTTGTGCGAACGCGAATACC	
<i>ctdP(P280F)-Exp-R</i>	ACCGAAGAGTTCGCCATCACGCATTTGGAAGAACGCGAATAC	
<i>ctdP(S340A)-Exp-F</i>	ATCGTCAAGAGAAGATCCTGGTGAACATCACACCACCATCAC	Plasmid construction for CtdP (S340A) expression
<i>ctdP(S340A)-Exp-R</i>	ACCAGGATCTTCTGACGATGGGACCTGCTCTCTCGATCTT	
<i>ctdP(P342A)-Exp-F</i>	GAAGATCCTGGTGAACATCACACCACCATCACTAATGA	Plasmid construction for CtdP (P342A) expression
<i>ctdP(P342A)-Exp-R</i>	TTCACCAGGATCTTCTGACGATGGACCCGATCTCTCGATCTT	
<i>ctdP(A330-367)-Exp-F</i>	GAGGCTCACAGAGAACAGATTTGGTGGATCCACACAGATTAACAACTCTCC	Plasmid construction for Sumo-CtdP(Δ 330-367)-expression
<i>ctdP(A330-367)-Exp-R</i>	CTCAGCTTCTTTCGGGCCCTCGAGTGGCGCCGCTTATGCCCGGCAATGCCTTTCGAAC	
<i>ctdP(A335-367)-Exp-R</i>	AACTCAGCTTCTTTCGGGCCCTCGAGTGGCGCCGCTTAGGAAGTTTGTGTTGCCCGGC	Plasmid construction for Sumo-CtdP(Δ 335-367)-expression
<i>ctdP(A342-367)-Exp-R</i>	CTCAGCTTCTTTCGGGCCCTCGAGTGGCGCCGCTTAACCCGATCTCTCGATCTTCTCG	Plasmid construction for Sumo-CtdP(Δ 342-367)-expression
<i>ctdP(A348-367)-Exp-R</i>	ACTCAGCTTCTTTCGGGCCCTCGAGTGGCGCCGCTTAATCTTCTGACGATGGGACC	Plasmid construction for Sumo-CtdP(Δ 348-367)-expression
<i>BmGDH-Exp-F</i>	CTTTAAGAAGGAGATATACCATGTATAAAGATTTAGAAGGAA	Plasmid construction for BmGDH expression
<i>BmGDH-Exp-R</i>	TTAGTGATGGTGGTGGTGTATCTCCGCGTCTGCTTGGAAATGA	
<i>pET-Exp-F</i>	CATCACCACCACCATCACTAATGATAATTTGAACGCCAGCACA	Plasmid construction for proteins expression
<i>pET-Exp-R</i>	GGTATATCTCCTTCTTAAAGTTAAACAAAATTTTCTAGAGGGG	

Supplementary Table 3 | BLASTP CtdN homologs in NCBI databases.

Description	Max Score	Total Score	Query Cover	E value	Per. Ident	Accession
Short-chain dehydrogenase/reductase PhqE [<i>Penicillium fellutanum</i>]	296	296	99%	2e ⁻¹⁰⁰	50.57%	L0E2Z4.1
Short-chain dehydrogenase/reductase MalC [<i>Malbranchea aurantiaca</i>]	251	251	97%	1e ⁻⁸²	45.77%	L0E4F8.1
Short-chain dehydrogenase/reductase ATR9 [<i>Stachybotrys chlorohalonata</i> IBT 40285]	169	169	95%	9e ⁻⁵¹	37.89%	A0A084R1K2.1
Short-chain dehydrogenase/reductase UcsE [<i>Acremonium</i> sp.]	142	142	95%	7e ⁻⁴⁰	30.08%	A0A411KUU9.1
Short-chain dehydrogenase/reductase Fsr5 [<i>Fusarium fujikuroi</i> IMI 58289]	127	127	97%	3e ⁻³⁴	31.82%	S0DRI2.1
Uncharacterized oxidoreductase YkvO [<i>Bacillus subtilis</i> subsp. <i>subtilis</i> str. 168]	87.8	87.8	95%	2e ⁻¹⁹	27.03%	O31680.1
3alpha-hydroxysteroid dehydrogenase [<i>Ruminococcus gnavus</i> ATCC 29149]	67.8	67.8	94%	4e ⁻¹²	25.77%	A7B3K3.1
Tropinone reductase homolog P29X [<i>Datura stramonium</i>]	67.4	67.4	97%	5e ⁻¹²	26.60%	P50165.1
Dihydroantipyrin 7-dehydrogenase BacC [<i>Bacillus subtilis</i>]	65.5	65.5	94%	2e ⁻¹¹	26.54%	Q8KWT4.1
Tropinone reductase TR-II [<i>Hyoscyamus niger</i>]	64.7	64.7	95%	5e ⁻¹¹	25.77%	P50164.1
Noroxomaritidine/norcrasgoline reductase NorRed [<i>Narcissus pseudonarcissus</i>]	64.3	64.3	94%	6e ⁻¹¹	27.84%	A0A1A9TAK5.1
Tropinone reductase TR-I [<i>Datura stramonium</i>]	63.2	63.2	94%	2e ⁻¹⁰	25.95%	P50162.1
3alpha-hydroxy bile acid-CoA-ester 3-dehydrogenase 1/3 [<i>Clostridium scindens</i>]	62.4	62.4	95%	2e ⁻¹⁰	25.97%	P07914.3
Peroxisomal trans-2-enoyl-CoA reductase [<i>Cavia porcellus</i>]	40.4	40.4	25%	0.010	33.80%	Q9JIF5.1
Peroxisomal trans-2-enoyl-CoA reductase [<i>Pongo abelii</i>]	39.7	39.7	29%	0.017	29.76%	Q5RCH8

Supplementary Table 4 | BLASTP CtdP homologs in NCBI databases.

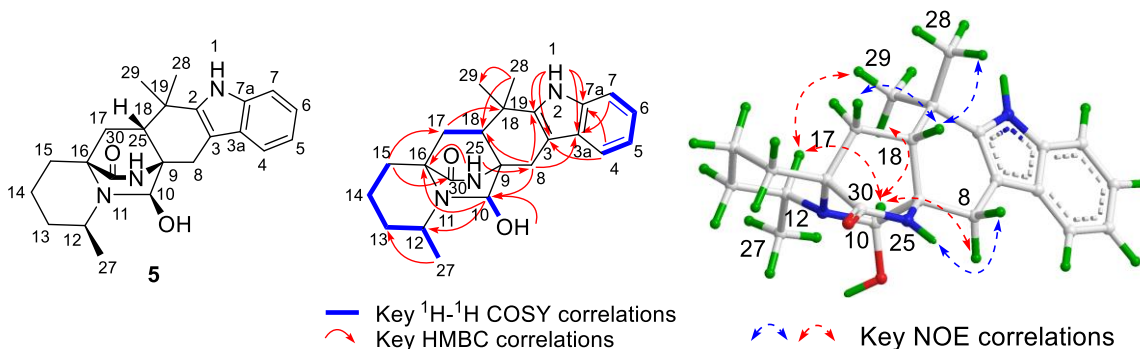
Description	Max Score	Total Score	Query Cover	E value	Per. Ident	Accession
NmrA-like family domain-containing oxidoreductase HimF [<i>Aspergillus japonicus</i>]	266	266	85%	3e ⁻⁸⁶	45.62%	A0A2Z5TWF0.1
NmrA-like family domain-containing oxidoreductase LnaB [<i>Aspergillus flavus</i> NRRL3357]	245	245	85%	1e ⁻⁷⁷	41.69%	B8NU00.1
NmrA-like family domain-containing oxidoreductase PtmS [<i>Penicillium simplicissimum</i>]	207	207	88%	5e ⁻⁶³	35.87%	A0A140JWT5.1
NmrA-like family domain-containing oxidoreductase PhqG [<i>Penicillium fellutanum</i>]	200	200	85%	2e ⁻⁶⁰	38.56%	L0E2U6.1
NmrA-like family domain-containing oxidoreductase NotA' [<i>Aspergillus versicolor</i>]	195	195	87%	3e ⁻⁵⁸	34.97%	L7WRQ4.1
NmrA-like family domain-containing protein DDB_G0286605 [<i>Dictyostelium discoideum</i>]	104	104	74%	2e ⁻²⁴	29.02%	Q54LJ8.1
NmrA-like family domain-containing protein 1 [<i>Bos taurus</i>]	82.0	82.0	74%	2e ⁻¹⁶	27.30%	Q0VCN1.1
NmrA-like family domain-containing protein 1 [<i>Gallus gallus</i>]	81.3	81.3	67%	4e ⁻¹⁶	26.88%	Q5ZID0.1
NmrA-like family domain-containing protein 1 [<i>Mus musculus</i>]	80.5	80.5	45%	8e ⁻¹⁶	31.64%	Q8K2T1.1
NmrA-like family domain-containing protein 1 [<i>Homo sapiens</i>]	78.6	78.6	42%	3e ⁻¹⁵	33.33%	Q9HBL8.1
NmrA-like family domain-containing protein 1 [<i>Rattus norvegicus</i>]	40.0	40.0	13%	0.007	42.31%	P86172.1
CtdP aligned sequence with MalC and PhqE						
Short-chain dehydrogenase/reductase MalC [<i>Malbranchea aurantiaca</i>]	24.6	41.2	30%	0.004	30.00%	L0E4F8.1
Short-chain dehydrogenase/reductase PhqE [<i>Penicillium fellutanum</i>]	16.2	16.2	6%	2.0	20.83%	L0E2Z4.1



Supplementary Table 5 | NMR Data of compound 3 (δ in ppm, J in Hz)^a.

Position	^1H	^{13}C		^1H - ^1H COSY	HMBC
1(NH)	10.53 s				C2, 3, 3a
2		140.7	C		
3		104.5	C		
3a		129.1	C		
4	7.32 d (7.5)	117.7	CH	H5	C3, 6, 7a
5	6.92 dd (7.5, 7.5)	118.3	CH	H4, 6	C3a, 7
6	7.00 dd (7.5, 7.5)	120.4	CH	H5, 7	C4, 7a
7	7.32 d (7.5)	110.9	CH	H6	C3a, 5
7a		134.7	C		
8	3.49 s	25.2	CH ₂		C2, 3a, 10
9		115.9	C		
10	4.95 s	112.8	CH		C3, 12, 16
12	2.46 m	54.4	CH	H13, 27	
13	a 1.41 m	32.5	CH ₂	H12, 14	
	b 0.98 m			H12, 14	
14	a 1.68 m	23.1	CH ₂	H13, 15	C12
	b 1.30 m			H13, 15	
15	a 1.75 m	24.6	CH ₂	H14	C13
	b 1.35 m			H14	
16	3.01 m	60.7	CH		
17	a 5.05 dd (17.4, 1.1)	111.1	CH ₂	H18	C19
	b 5.02 dd (10.5, 1.1)			H18	
18	6.16 dd (17.4, 10.5)	146.0	CH	H17	C28, 29
19		38.9	C		
25 (NH)	9.03 s				
27	0.75 d (6.3)	20.1	CH ₃	H12	C13
28	1.48 s	27.5	CH ₃		C2, 18, 29
29	1.48 s	27.5	CH ₃		C2, 18, 28
30		166.8	C		

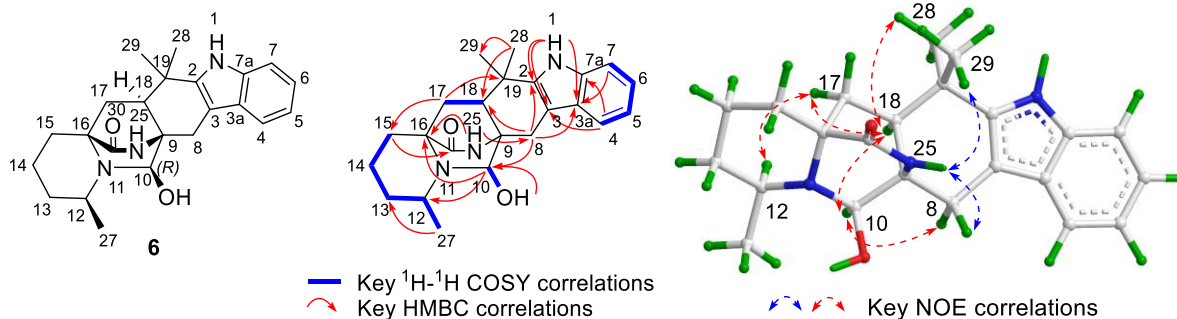
^aMeasured in DMSO-*d*₆, 600 MHz for ^1H and 150 MHz for ^{13}C NMR. Overlapped signals are reported without designating multiplicity.



Supplementary Table 6 | NMR Data of compound 5 (δ in ppm, J in Hz)^a

Position	^1H	^{13}C		^1H - ^1H COSY	HMBC	NOESY
1	7.83 s		NH		C2, 3, 3a, 7a	
2		140.0	C			
3		104.3	C			
3a		127.1	C			
4	7.48 d (7.8)	118.6	CH	H5	C3, 6, 7a	
5	7.18 dd (7.8, 7.8)	119.7	CH	H4, 6	C3a, 7	
6	7.11 dd (7.8, 7.8)	122.2	CH	H5, 7	C4, 7a	
7	7.32 d (7.8)	110.7	CH	H6	C3a, 5	
7a		136.5	C			
8	α 3.78 d (15.5)	26.8	CH_2		C2, 3a, 10, 18	H10
	β 2.53 d (15.5)					C2, 3a, 10, 18
9		58.4	C			
10	4.44 d (7.6)	81.5	CH	OH	C12, 16	H8 α , 17, 29
12	3.01 m	50.9	CH	H13, 27		H10, 29
13	a 1.64 overlapped	35.1	CH_2	H12, 14	C15	
	b 1.18 m					H12, 14
14	a 1.73 overlapped	19.8	CH_2	H13, 15		
	b 1.50 m					H13, 15
15	a 2.07 m	29.6	CH_2	H14	C17, 30	
	b 1.73 overlapped					H14
16		58.3	C			
17	α 2.43 m	24.5	CH_2	H18	C19, 30	H12
	β 1.60 overlapped					H18
18	2.41 m	48.1	CH	H17	C8, 10, 28, 29	H28
19		34.9	C			
25	5.91 s		NH			H8 β
27	1.12 d (6.1)	23.1	CH_3	H12	C13	
28	1.38 s	31.1	CH_3		C3, 18, 29	H18
29	1.42 s	27.1	CH_3		C3, 18, 28	H10, 12
30		175.0	C			
OH	1.57 d (7.6)			H10	C10	

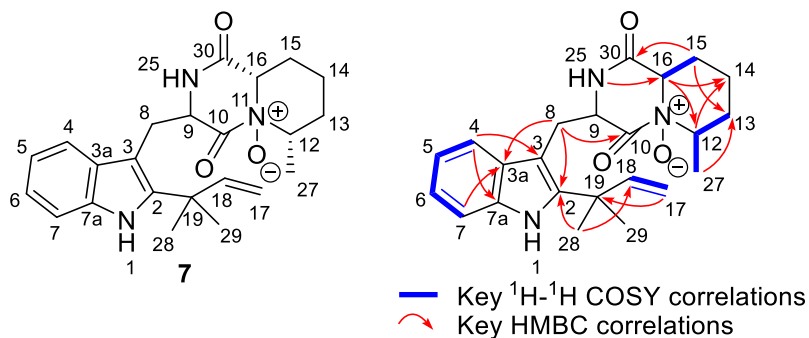
^aMeasured in CDCl_3 , 600 MHz for ^1H and 150 MHz for ^{13}C NMR. Overlapped signals are reported without designating multiplicity.



Supplementary Table 7 | NMR Data of compound 6 (δ in ppm, J in Hz)^a.

Position	^1H	^{13}C		^1H - ^1H COSY	HMBC	NOESY
1	9.10 s		NH		C2, 3, 3a	
2		142.5	C			
3		104.6	C			
3a		128.4	C			
4	7.63 d (7.7)	118.7	CH	H5	C3, 6, 7a	
5	7.08 dd (7.7, 7.7)	119.6	CH	H4, 6	C3a, 7	
6	7.01 dd (7.7, 7.7)	121.9	CH	H5, 7	C4, 7a	
7	7.31 d (7.7)	111.5	CH	H6	C3a, 5	
7a		137.5	C			
8	β 2.99 d (17.4)	27.9	CH_2		C2, 3a, 18	H25
	α 2.91 d (17.4)				C2, 10	H10
9		57.6	C			
10	3.87 d (8.1)	91.0	CH	OH	C12	H8 α , 18
12	2.81 m	53.1	CH	H13, 27	C10, 16	H17 α
13	a 1.64 m	35.7	CH_2	H12, 14	C15	
	b 1.13 m			H12, 14		
14	a 1.61 overlapped	20.0	CH_2	H13, 15		
	b 1.54 m			H13, 15		
15	a 1.84 m	30.2	CH_2	H14	C17	
	b 1.61 overlapped			H14	C13	
16		58.9	C			
17	α 2.62 m	24.7	CH_2	H18	C19, 30	H12, 18
	β 1.28 m			H18		
18	1.97 m	44.2	CH	H17	C8, 29	H28
19		35.0	C			
25	6.33 s		NH			H8 β , 29
27	1.15 d (6.1)	23.4	CH_3	H12	C13	
28	1.28 s	28.4	CH_3		C3, 18, 29	H18
29	1.17 s	24.7	CH_3		C3, 18, 28	H25
30		174.9	C			
OH	2.61 overlapped			H10	C10	

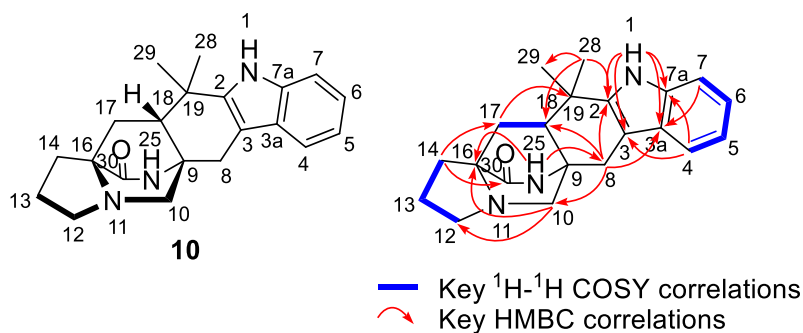
^aMeasured in CD_3CN , 600 MHz for ^1H and 150 MHz for ^{13}C NMR. Overlapped signals are reported without designating multiplicity.



Supplementary Table 8 | NMR Data of compound 7 (δ in ppm, J in Hz)^a.

Position	^1H	^{13}C		^1H - ^1H COSY	HMBC
1	10.64 s		NH		C2, 3, 3a
2		141.3	C		
3		102.5	C		
3a		129.4	C		
4	7.29 d (7.8)	117.5	CH	H5	C3, 6, 7a
5	6.92 dd (7.8, 7.8)	118.4	CH	H4, 6	C3a, 7
6	7.02 dd (7.8, 7.8)	120.5	CH	H5, 7	C4, 7a
7	7.31 d (7.8)	110.9	CH	H6	C3a, 5
7a		134.6	C		
8	3.98 s	33.4	CH ₂		C2, 3a, 10
9	Missing	Missing	C		
10		172.2	C		
12	4.00 overlapped	48.5	CH	H13, 27	C14
13	1.62 overlapped	27.5	CH ₂	H12	
	1.52 overlapped				
14	a 1.63 overlapped	15.9	CH ₂		
	b 1.51 overlapped				
15	a 2.00 m	24.7	CH ₂	H16	C2, 12
	b 1.56 m				
16	4.91 m	51.2	CH		
17	a 5.06 d (17.4)	111.2	CH ₂	H18	C19
	b 5.03 dd (10.6)				
18	6.08 dd (17.4, 10.6)	145.9	CH	H17	C2, 28/29
19		39.5	C		
25	8.09 s		NH		C16
27	1.21 d (7.0)	21.5	CH ₃	H12	C13
28	1.44 s	27.5	CH ₃		C2, 18
29	1.44 s	27.5	CH ₃		C2, 18
30		171.7	C		

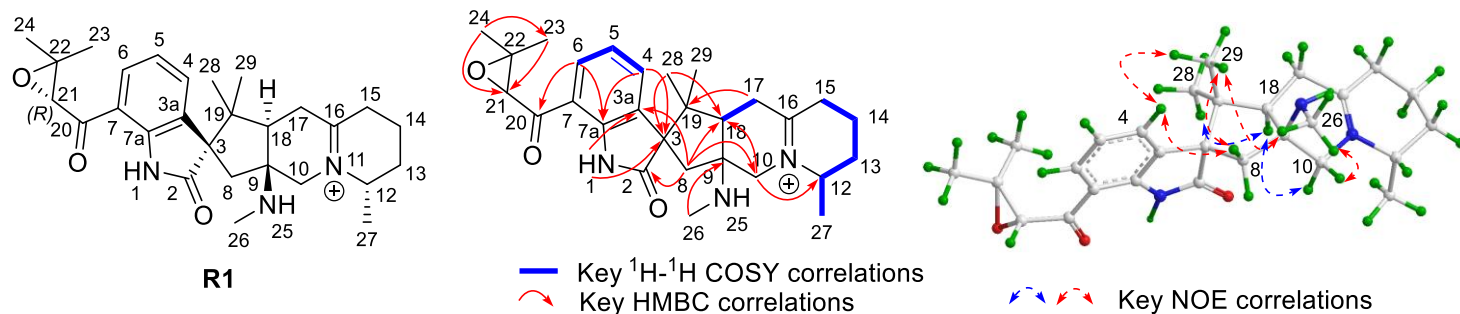
^aMeasured in DMSO-*d*₆, 600 MHz for ^1H and 150 MHz for ^{13}C NMR. Overlapped signals are reported without designating multiplicity.



Supplementary Table 9 | NMR Data of compound 10 (δ in ppm, J in Hz)^a.

Position	^1H	^{13}C		^1H - ^1H COSY	HMBC	NOESY
1	9.12 s		NH			
2		142.9	C			
3		104.3	C			
3a		128.2	C			
4	7.40 d (7.7)	118.5	CH	H5	C3, 6, 7a	H8 α
5	7.01 dd (7.7, 7.7)	119.7	CH	H4, 6	C3a, 7	
6	7.02 dd (7.7, 7.7)	121.9	CH	H5, 7	C4, 7a	
7	7.31 d (7.7)	111.5	CH	H6	C3a, 5	H1
7a		137.6	C			
8	α 2.90 d (17.0)	29.0	CH ₂		C2, 3a, 18	H4, 25
	β 2.82 d (17.0)				C2, 3a, 10	H10 β , 18
9		56.4	C			
10	α 3.07 d (10.1)	62.5	CH ₂		C12, 16	H12
	β 2.62 d (10.1)				C8, 12, 16, 18	H8 β
12	a 3.02 m	53.7	CH ₂	H13		H10 α
	b 2.31 dd (16.6, 8.5)			H13		
13	a 1.83 m	23.5	CH ₂	H12, 14	C15	
	b 1.80 m			H12, 14	C27	
14	a 2.42 m	28.0	CH ₂	H13	C12	
	b 1.36 m			H13	C16	
16		65.8	C			
17	β 2.06 m	32.2	CH ₂	H18	C15, 19, 30	
	α 1.89 m			H18	C15, 19, 30	
18	2.17 overlapped	46.9	CH	H17	C8, 10, 16, 28, 29	H8 β , 28
19		35.1	C			
25	6.44 s		NH		C10, 16	H8 α , 29
28	1.26 s	28.0	CH ₃		C2, 18, 29	H18
29	1.16 s	24.5	CH ₃		C2, 18, 28	H25
30		174.2	C			

^aMeasured in CD₃CN, 600 MHz for ^1H and 150 MHz for ^{13}C NMR. Overlapped signals are reported without designating multiplicity.



Supplementary Table 10 | NMR Data of compound R1 (δ in ppm, J in Hz)^a.

Position	^1H	^{13}C		^1H - ^1H COSY	HMBC	NOESY
1	9.64 s		NH		C3, 3a	
2		182.5	C			
3		64.4	C			
3a		131.8	C			
4	7.46 d (7.7)	131.1	CH	H5	C3, 6, 7a	H8 β , 29
5	7.10 dd (7.7, 7.7)	121.0	CH	H4, 6	C3a, 7	
6	7.76 d (7.7)	128.0	CH	H5	C4, 7a	
7		116.8	C			
7a		144.0	C			
8	β 2.34 d (14.8)	32.9	CH ₂		C2, 3a, 10, 18	H4, 29
	α 2.05 d (14.8)				C2, 3a, 10, 19	H10, 12
9		76.0	C			
10	β 4.00 m	57.9	CH ₂			H18
	α 2.82 m				C12, 18	H26
12	2.81 m	63.2	CH	H13, 27	C14	
13	a 1.85 overlapped	30.7	CH ₂	H12, 14		
	b 1.69 overlapped			H12, 14		
14	1.86 overlapped	20.4	CH ₂	H13, 15	C12, 16	
	1.69 overlapped					
15	a 2.26 overlapped	27.7	CH ₂	H14, 16		
	b 2.24 overlapped			H14, 16	C13	
16	missing	missing	C			
17	a 2.42 overlapped	38.9	CH ₂	H18		
	b 1.66 overlapped			H18	C19	
18	3.05 m	52.5	CH	H17	C10	H-28
19		47.3	C			
20		194.9	C			
21	4.02 s	64.4	CH		C23, 24	
22		61.6	C			
23	1.60 s	24.4	CH ₃		C21, 24	
24	1.25 s	18.8	CH ₃		C21, 23	
26	2.47 s	30.7	CH ₃		C9	H29
27	1.35 d (3.5)	20.4	CH ₃	H12	C13	
28	0.72 s	23.5	CH ₃		C3, 18, 29	H18
29	1.06 s	22.1	CH ₃		C3, 18, 28	H4, 8 β , 26

^aMeasured in CDCl₃, 600 MHz for ^1H and 150 MHz for ^{13}C NMR. Overlapped signals are reported without designating multiplicity.

Supplementary Table 11 | Crystal data and structure refinement for compound 5.

Empirical formula	C ₂₃ H ₂₉ N ₃ O ₂
Formula weight	379.49
Temperature/K	100.15
Crystal system	monoclinic
Space group	P2 ₁
a/Å	9.17410(10)
b/Å	22.2451(2)
c/Å	11.6697(2)
α/°	90
β/°	106.2410(10)
γ/°	90
Volume/Å ³	2286.50(5)
Z	4
ρ _{calc} /g/cm ³	1.102
μ/mm ⁻¹	0.562
F(000)	816.0
Crystal size/mm ³	0.38 × 0.28 × 0.14
Radiation	Cu Kα (λ = 1.54184)
2θ range for data collection/°	7.89 to 149
Index ranges	-11 ≤ h ≤ 11, -22 ≤ k ≤ 27, -14 ≤ l ≤ 14
Reflections collected	25927
Independent reflections	7483 [R _{int} = 0.0289, R _{sigma} = 0.0242]
Data/restraints/parameters	7483/1/582
Goodness-of-fit on F ²	1.043
Final R indexes [I ≥ 2σ (I)]	R ₁ = 0.0357, wR ₂ = 0.0980
Final R indexes [all data]	R ₁ = 0.0362, wR ₂ = 0.0984
Largest diff. peak/hole / e Å ⁻³	0.29/-0.22
Flack parameter	0.04(9)
CCDC number	2127333

Supplementary Table 12 | Crystal data and structure refinement for compound 10.

Empirical formula	C ₂₃ H ₂₉ N ₃ O ₂
Formula weight	379.49
Temperature/K	99.99(10)
Crystal system	monoclinic
Space group	P2 ₁
a/Å	9.17410(10)
b/Å	22.2451(2)
c/Å	11.6697(2)
α/°	90
β/°	106.2410(10)
γ/°	90
Volume/Å ³	2286.50(5)
Z	4
ρ _{calc} /cm ³	1.102
μ/mm ⁻¹	0.562
F(000)	816.0
Crystal size/mm ³	0.38 × 0.28 × 0.14
Radiation	Cu Kα (λ = 1.54184)
2θ range for data collection/°	7.89 to 149
Index ranges	-11 ≤ h ≤ 11, -22 ≤ k ≤ 27, -14 ≤ l ≤ 14
Reflections collected	25927
Independent reflections	7483 [R _{int} = 0.0289, R _{sigma} = 0.0242]
Data/restraints/parameters	7483/1/741
Goodness-of-fit on F ²	1.052
Final R indexes [I ≥ 2σ (I)]	R1 = 0.0344, wR2 = 0.0942
Final R indexes [all data]	R1 = 0.0348, wR2 = 0.0946
Largest diff. peak/hole / e Å ⁻³	0.26/-0.21
Flack parameter	0.05(9)
CCDC number	2127332

Supplementary Table 13 | Crystallographic data collection and structure refinement statistics.

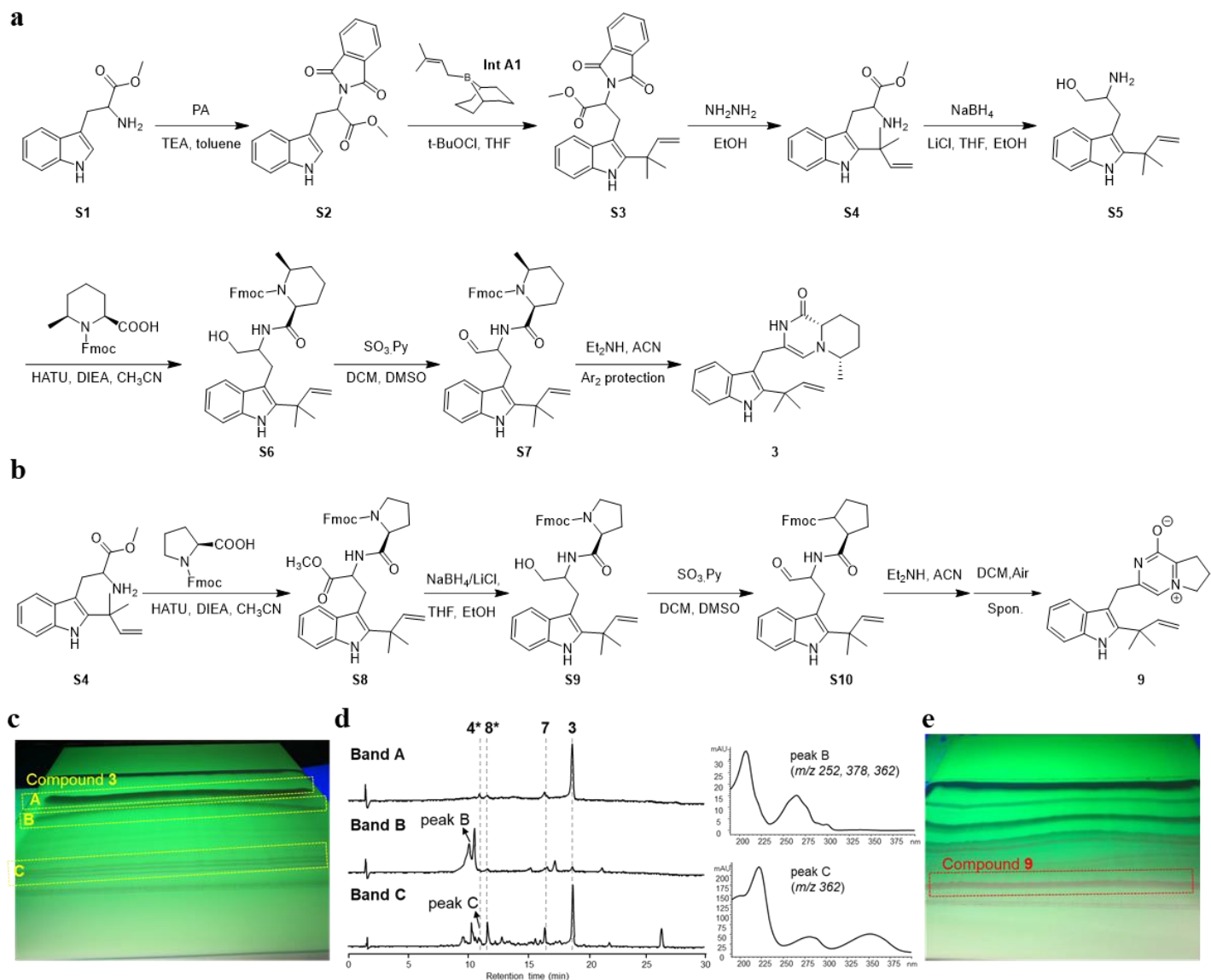
	SeMet CtdP
PDB code	7UF8
Data Collection*	
Space group	P 6 ₁ 2 2
<u>Cell Dimensions</u>	
<i>a</i> , <i>b</i> , <i>c</i> (Å)	166.8, 166.8, 195.4
α , β , γ (°)	90, 90, 120
Wavelength (Å)	0.979
Resolution range (Å)	48.14 - 2.5 (2.589 - 2.5)
R _{meas} (%)	18.45 (3.161)
<i>I</i> / σ <i>I</i>	24.52 (2.07)
Completeness (%)	99.77 (99.18)
Redundancy	39.8 (39.3)
Refinement	
Resolution range (Å)	48.14 - 2.5
No. Reflections	55726
R _{work} / R _{free}	0.214 / 0.245
<u>No. Atoms</u>	
Protein	10397
Ligand	462
Water	1325
<u>B-factors (Å²)</u>	
Protein	71.97
Ligand	74.51
Water	58.56
<u>R.M.S. deviations</u>	
Bond lengths (Å)	0.011
Bond angles (°)	1.26

*Values in parentheses are for highest-resolution shell

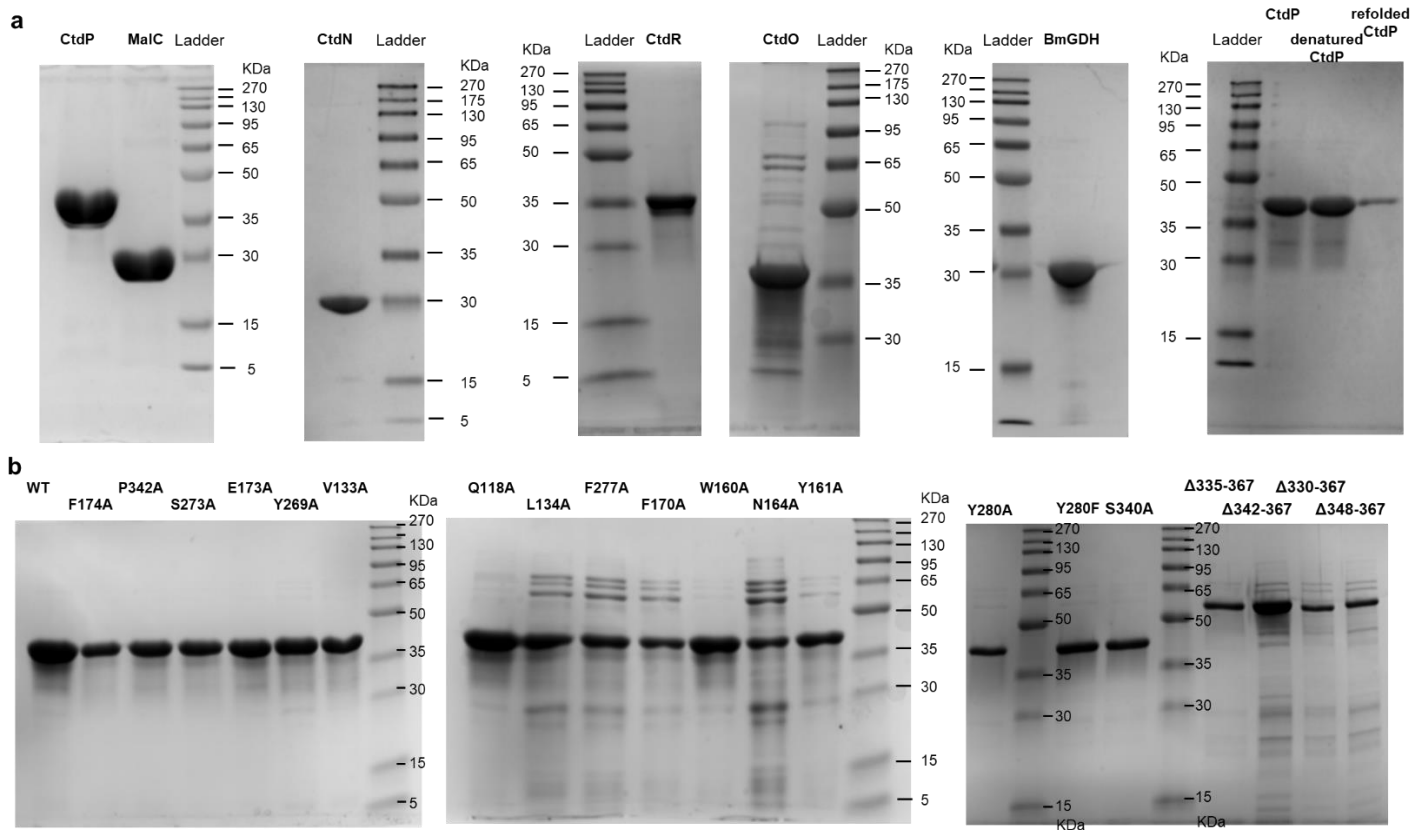
Supplementary Table 14 | Structural homologs of CtdP identified by DALI server¹.

No	Protein	PDB-Chain	Z score	rmsd	Number of structurally equivalent residues	Total number of amino acids	%identity
1	NADP(H) sensor HSCARG protein (<i>Homo sapiens</i>)	2exx-A	31.8	2	290	305	24
2	NmrA-like family domain containing protein 1 (<i>Homo sapiens</i>)	2wmd-A	31.9	2.1	289	295	24
3		2wm3-A	31.8	2.1	289	296	24
4		3dxf-A	31.5	2.1	288	292	24
5	NmrA-like family domain containing protein 1	3e5m-A	31.4	2	287	294	24
6		3e5m-B	31.1	2	287	295	24
7	NmrA (<i>Aspergillus nidulans</i>)	1k6i-A	30.2	2.6	294	318	20
8	NmrA	1xgk-A	30.2	2.6	295	325	21
9	NmrA-like family domain containing protein 1	3dxf-B	30.1	2.1	280	282	24
10	NmrA	1k6j-B	30.1	2.7	296	321	20
11		1ti7-A	30	2.6	294	324	20
12		2vuu-A	29.8	2.7	294	318	20
13		2vus-E	29.8	2.9	297	318	20
14		2vus-D	29.8	2.9	297	318	20
15		2vus-B	29.7	2.7	294	318	20
16		2vuu-F	29.7	2.7	294	318	20
17		2vus-F	29.7	2.7	294	318	20
18		2vuu-G	29.7	2.7	294	319	20
19		2vuu-E	29.7	2.7	294	318	20
20		2vus-G	29.7	2.7	294	318	20
21		1k6x-A	29.6	2.6	293	324	21
22		2vuu-D	29.6	2.7	295	319	20
23		1k6j-A	29.5	2.6	294	322	21
24		2vut-G	29.5	2.8	295	319	20
25		2vut-D	29.4	2.8	294	320	20
26		2vuu-B	29.4	2.8	293	319	20
27		2vus-A	29.3	2.6	291	318	21
28		2vus-C	29.2	2.6	292	318	21
29		2vuu-C	29.2	2.6	292	318	21
30		2vus-H	29.1	2.6	292	318	21
31		2vuu-H	29.1	2.7	292	319	21
32		2vut-F	29.1	2.6	291	318	21
33		2vut-A	29.1	2.7	293	318	21
34		2vut-C	29.1	2.6	291	318	21
35		2vut-E	29.0	2.6	291	318	21
36		2vut-H	29.0	2.7	292	319	21
37	NmrA-like protein KstA11 (<i>Micromonospora sp.</i> TP-A0468)	5f5n-A	28.9	2.5	281	289	26

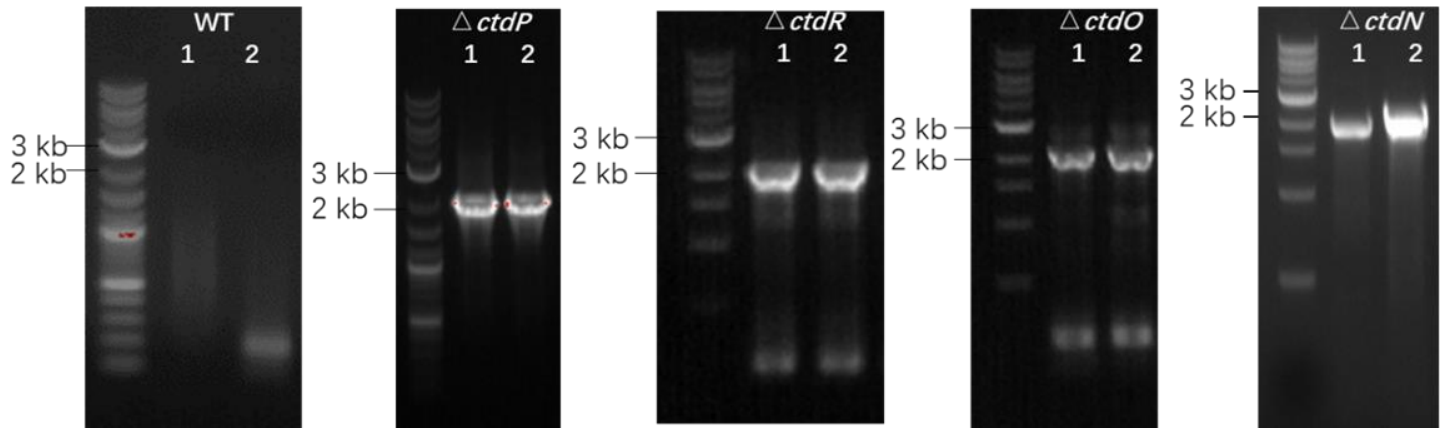
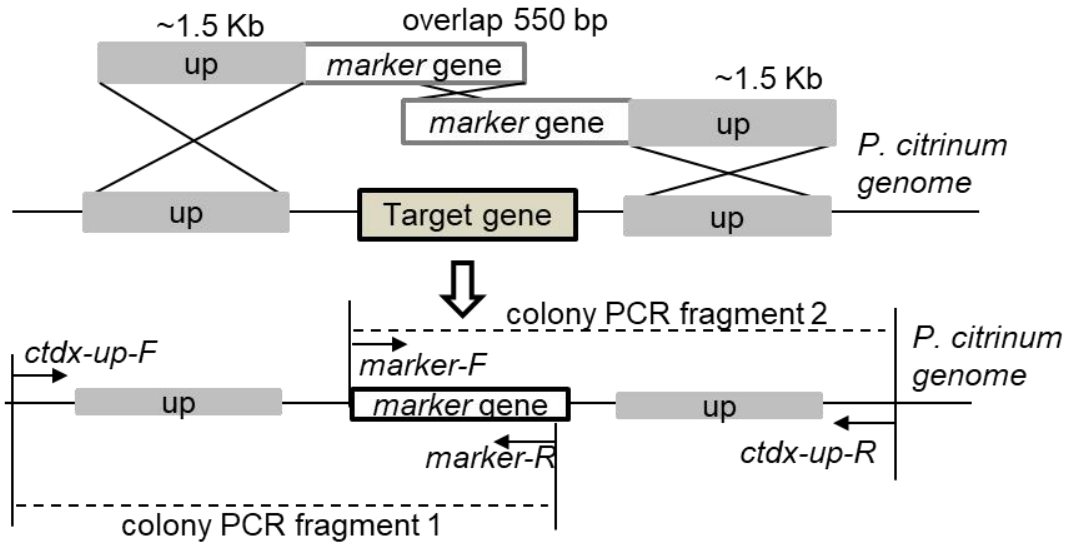
Supplementary Figures



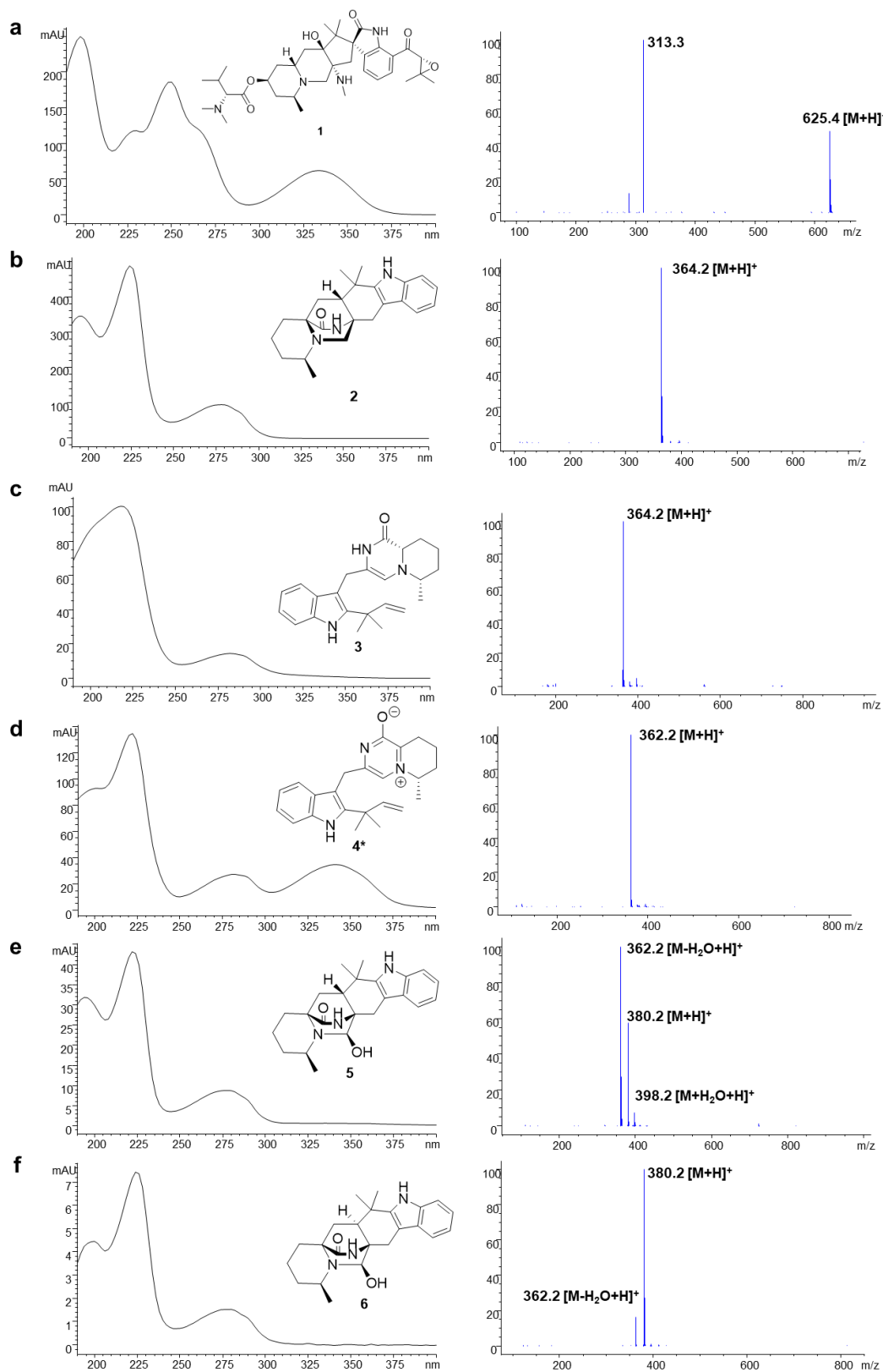
Supplementary Fig. 1 | Biomimetic synthesis of the CtdP and MalC substrates. a, Compound **3** synthetic pathway. **b**, Compound **9** synthetic pathway. **c**, Thin-layer chromatography (TLC) purification of compound **3** after the reaction from **S7**. **d**, LCMS analysis of bands A-C in **c**. Band A is pure **3**, while band C is presumed to be tautomerized to major **3** and proposed to contain compound **4** based on its UV and MS data. **e**, TLC purification of compound **9** after the reaction from **S10**. The symbol * represents the compound identified by MS and UV spectra.



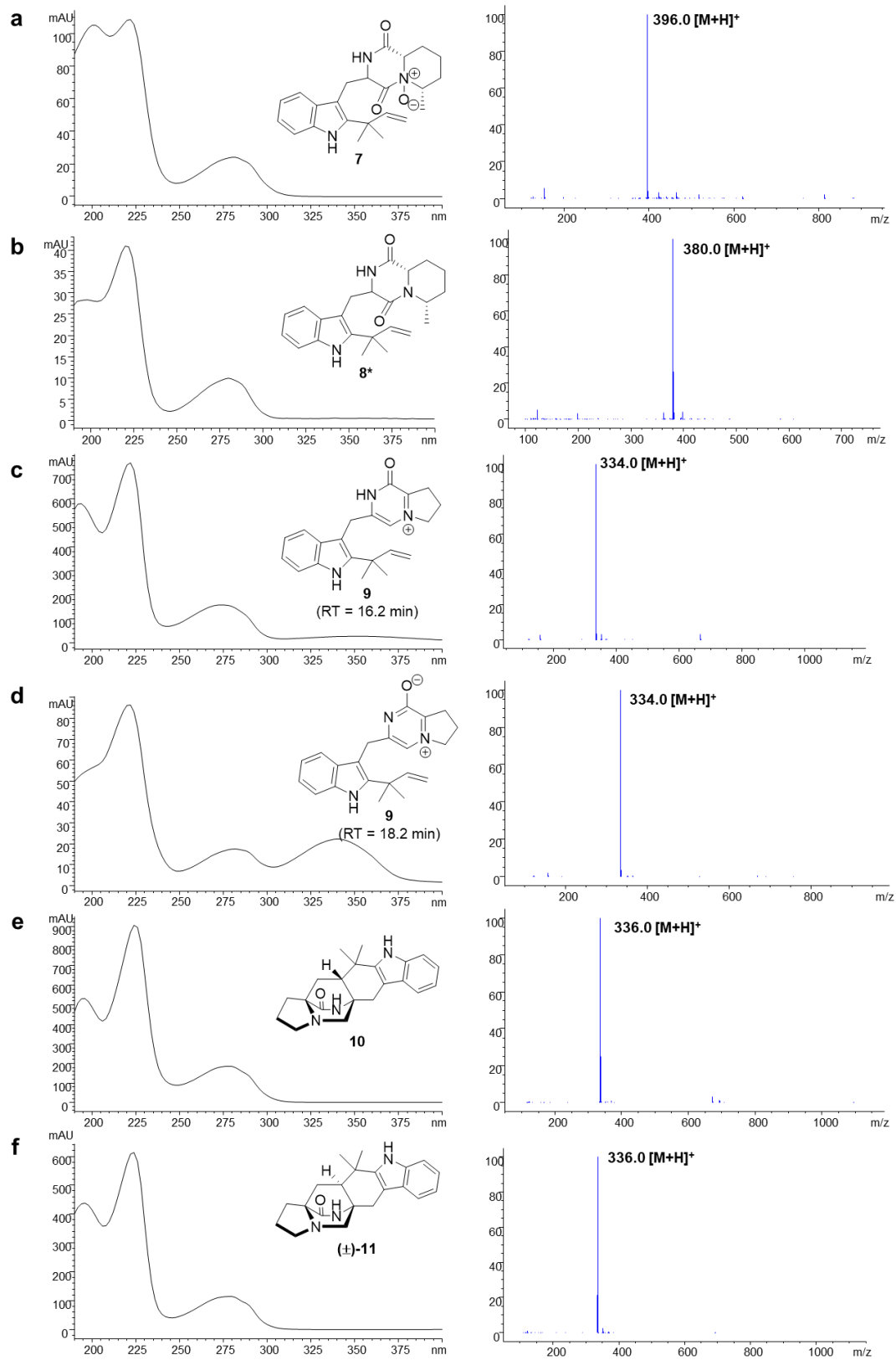
Supplementary Fig. 2 | SDS-PAGE (12%) analysis of purified proteins. a. C-His-tagged proteins of CtdP, MalC, CtdN, CtdR, CtdO, BmGDH, and refolded CtdP enzymes. **b.** C-His-tagged proteins of CtdP mutants. Experiments in **a** and **b** were repeated independently with similar results for three times.



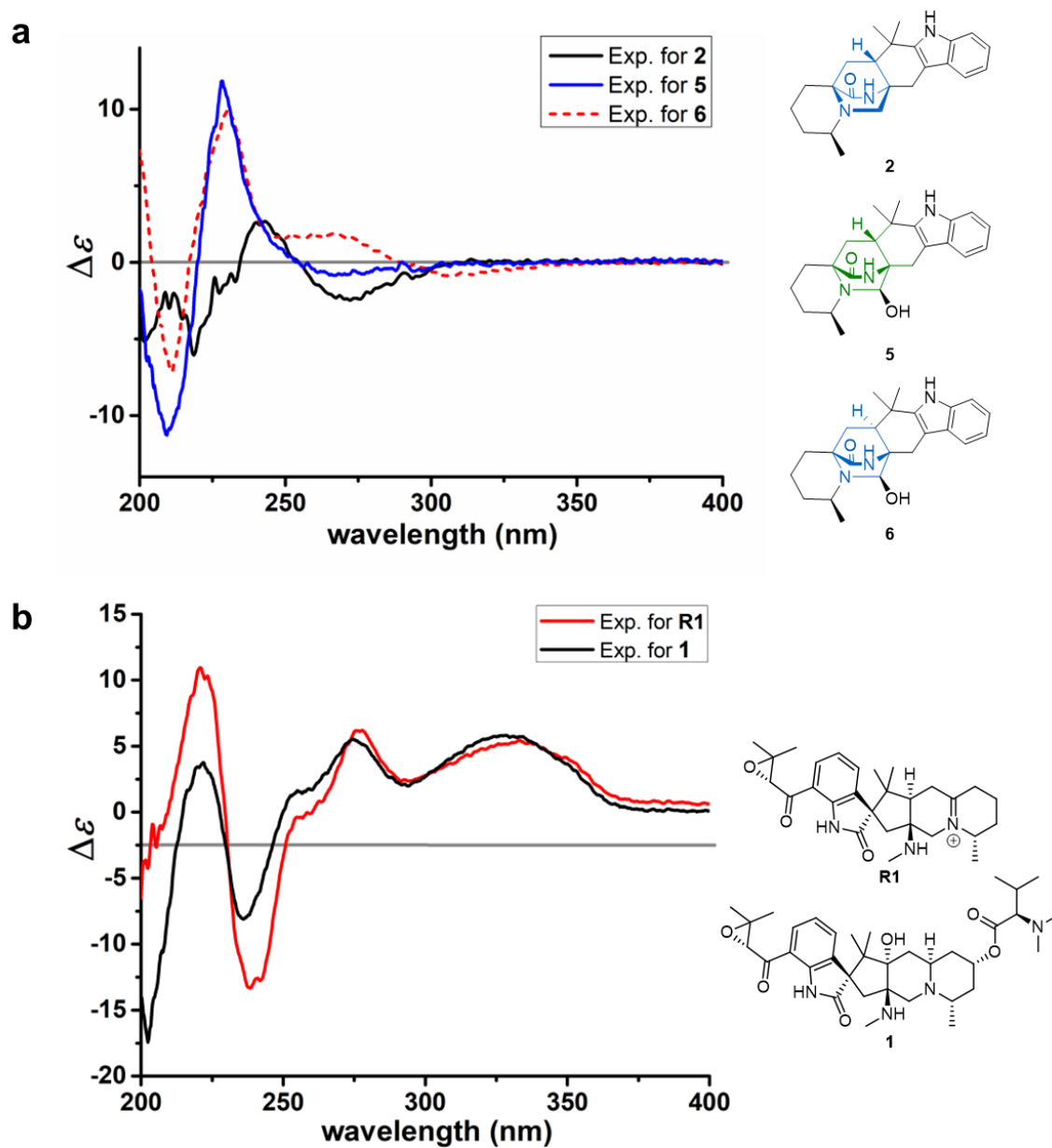
Supplementary Fig. 3 | PCR confirmation of *ctd* mutants. *ctd* transformants were screened by PCR using primers *ctdx-up-F* and *bar-R* for up-stream screening, primers *bar-F* and *ctdx-dn-R* for down-stream screening.



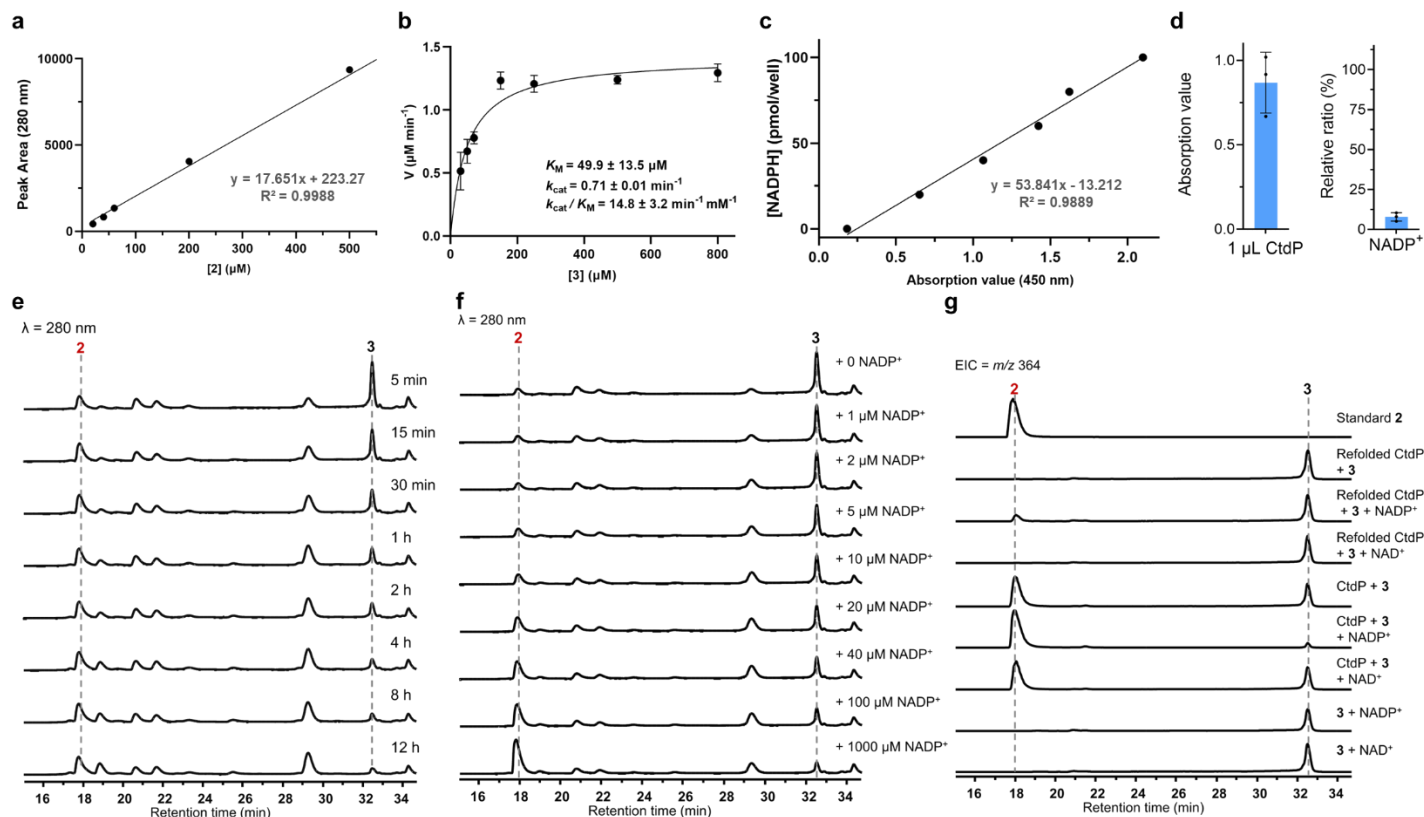
Supplementary Fig. 4 | UV and mass spectra of compounds 1-6.



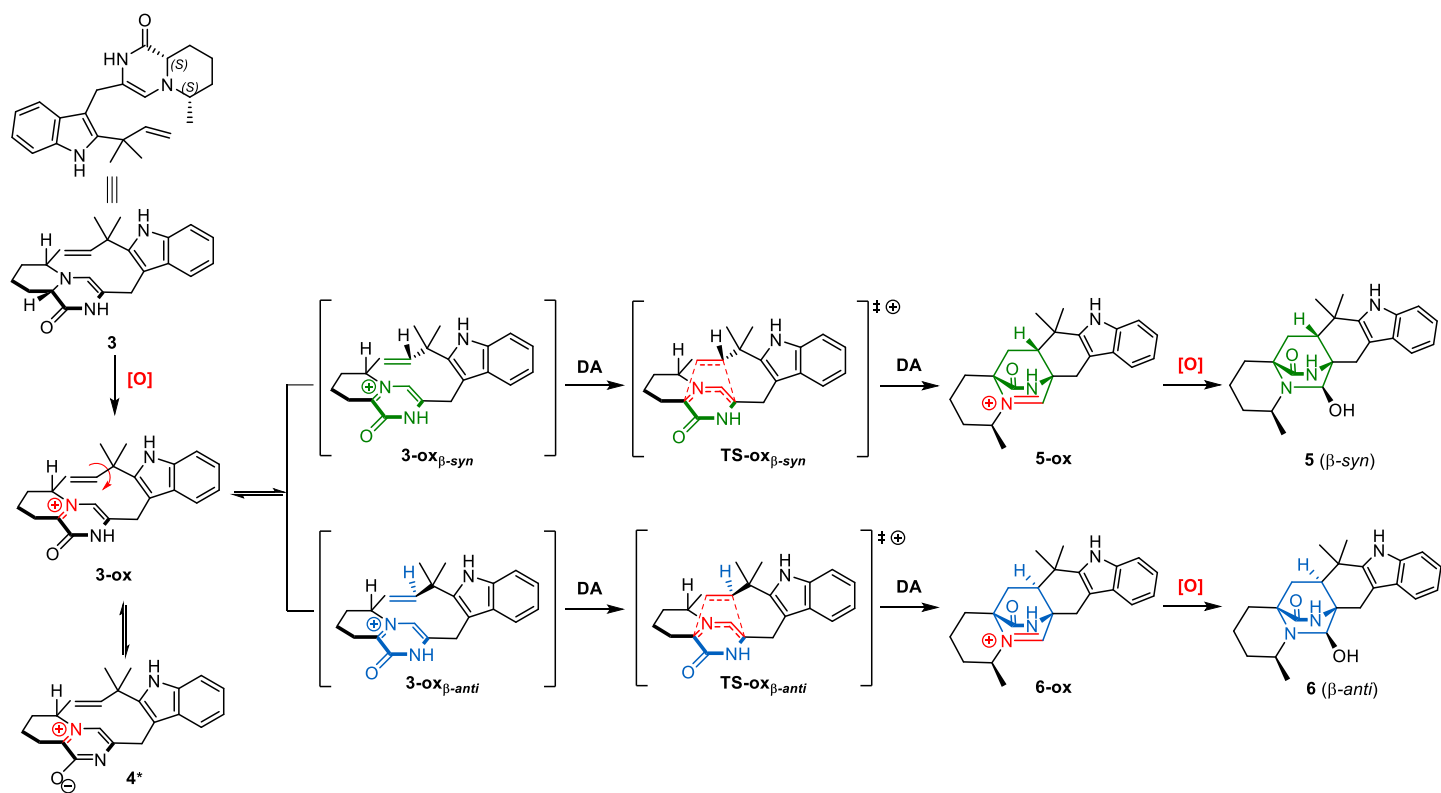
Supplementary Fig. 5 | UV and mass spectra of compounds 7-11.



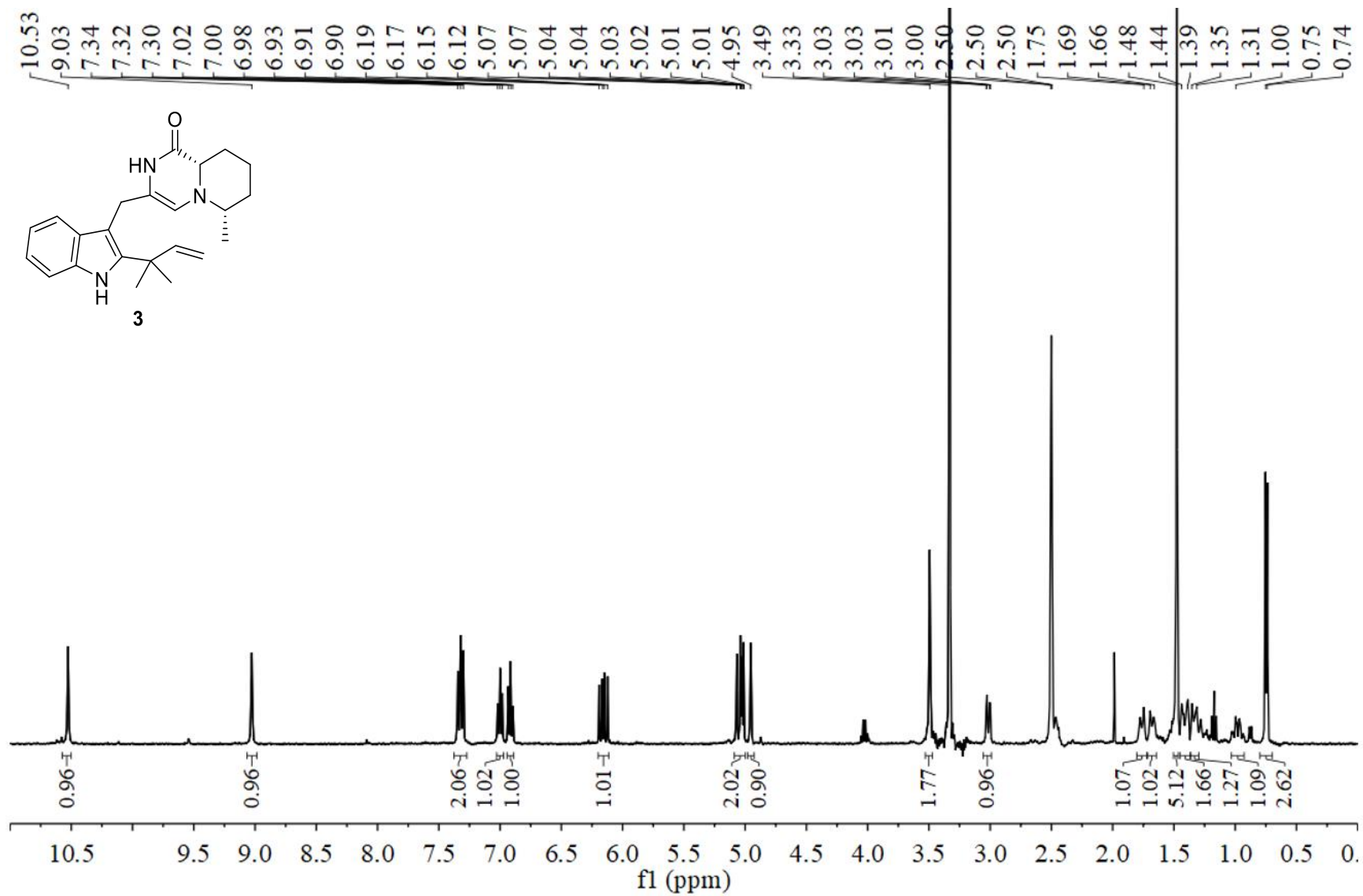
Supplementary Fig. 6 | Electronic circular dichroism spectra. a, Compounds 2, 5, and 6. b, Compounds R1 and 1.



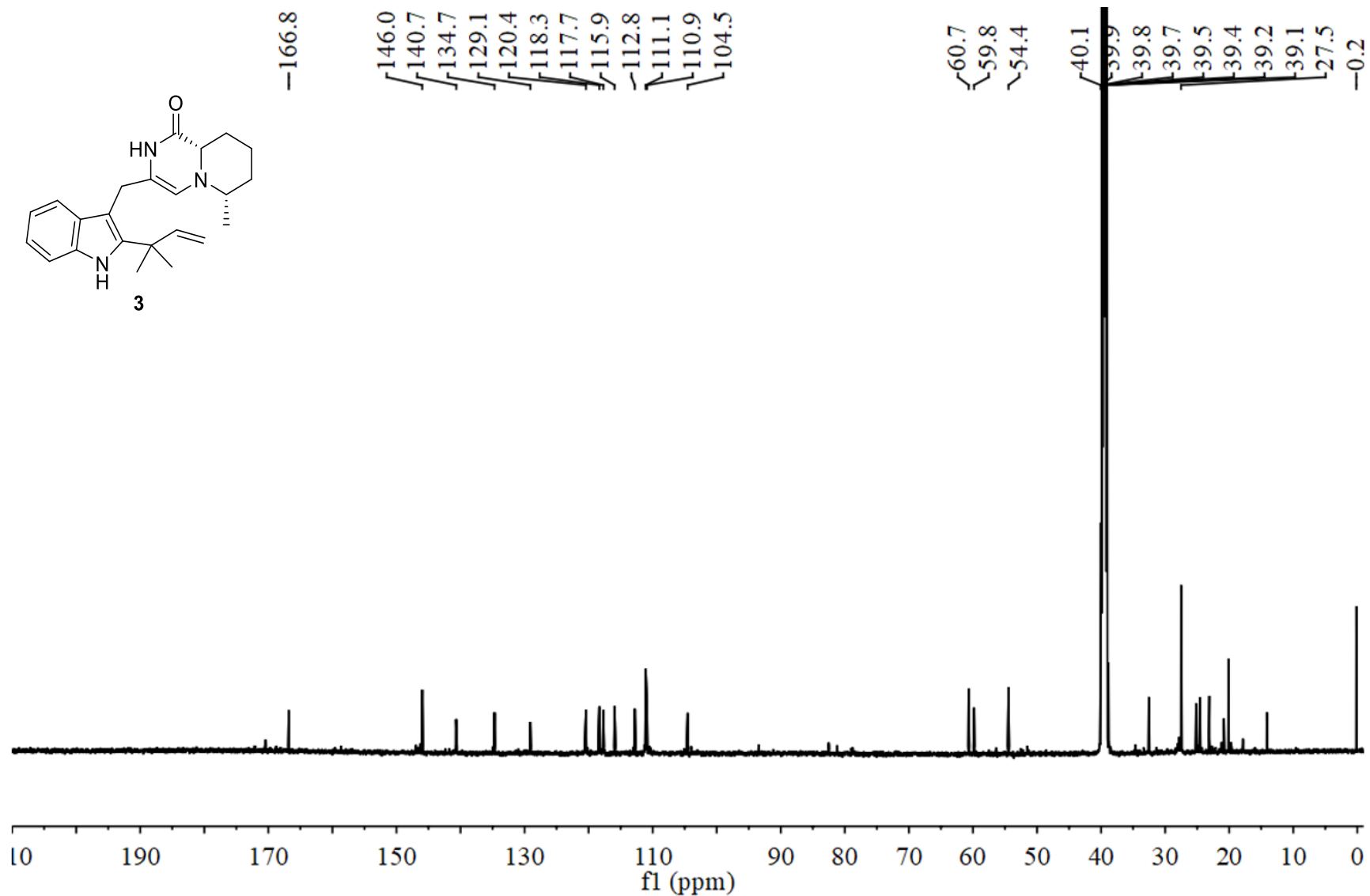
Supplementary Fig. 7 | Kinetic analysis of CtdP, NADP⁺ content in the purified CtdP protein, and refold CtdP assays. **a**, Standard curve of compound **2**. **b**, Michaelis-Menton kinetic analysis of CtdP with and **3**. Data was presented as mean \pm s.d. from triplicate independent experiments ($n = 3$). **c**, Standard curve of NADPH. **d**, Left: absorption value of NADP⁺ in 1 μL CtdP (0.88 ± 0.18). NADP⁺ amount = total NADP – NADPH. Right: the relative ratio of NADP⁺ in CtdP protein ($7.8 \pm 2.6\%$). Data was presented as mean \pm s.d. from triplicate independent experiments ($n = 3$). **e**, 20 μM CtdP (no addition of extra NADP⁺) reacts with 200 μM **3** in time course. The conversion rate reaches a maximum of 33.8% at 30 min. **f**, 10 μM CtdP and 200 μM **3** react with the addition of different concentrations of NADP⁺ for 15 min. The conversion rate of **2** increased from 10.4% to 63.9% with the increased amount of additional NADP⁺ from 0 to 1000 μM . **g**, CtdP (40 μM) and refolded CtdP (40 μM) react with **3** (200 μM) for 1 hour, respectively. The reaction buffer is 50 mM Tris-HCl (pH 7.0) and the reaction temperature is 28 $^\circ\text{C}$. The 100% conversion of **2** as the control.



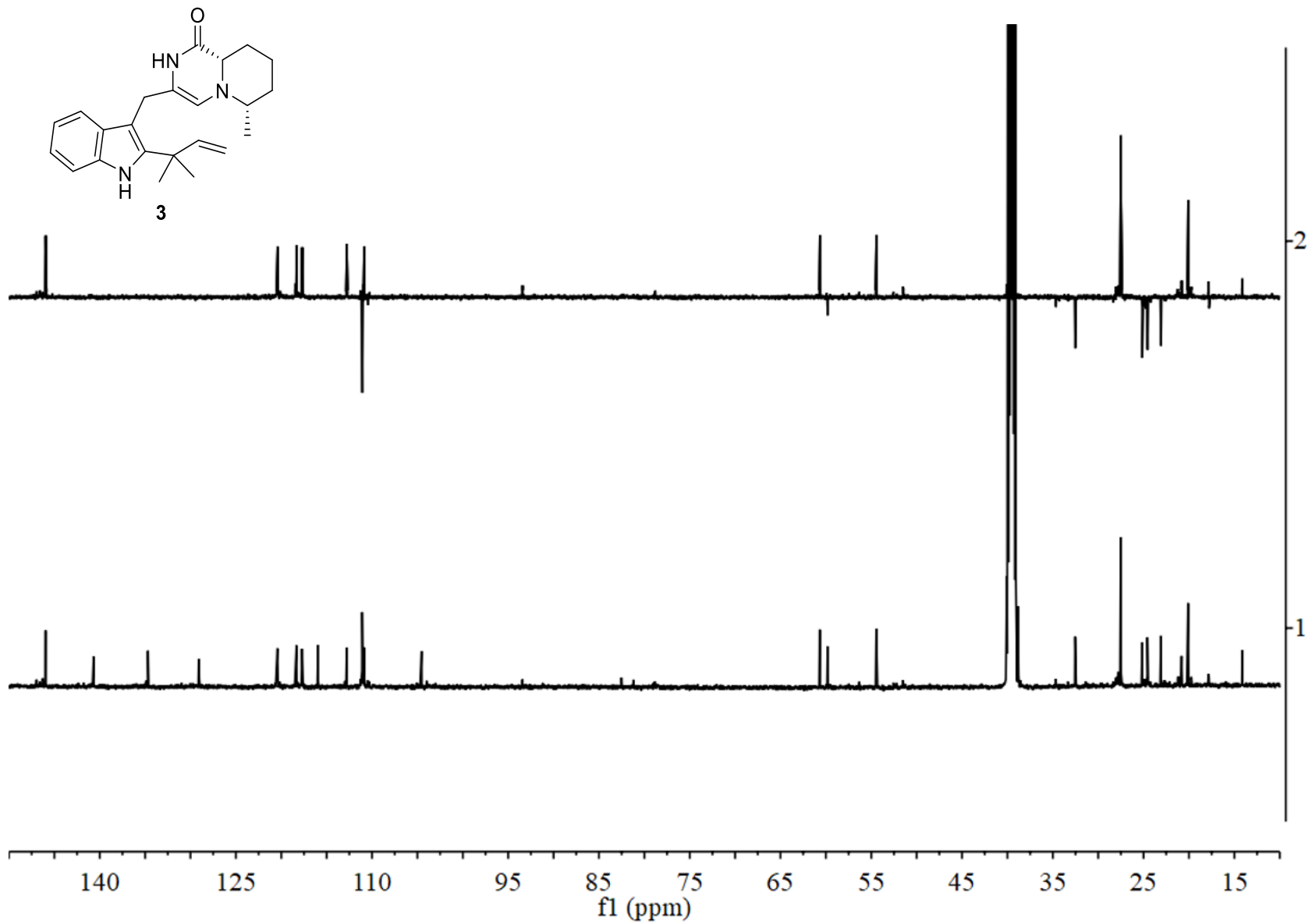
Supplementary Fig. 8 | The proposed pathways of spontaneous Diels-Alder reactions of compound 3.



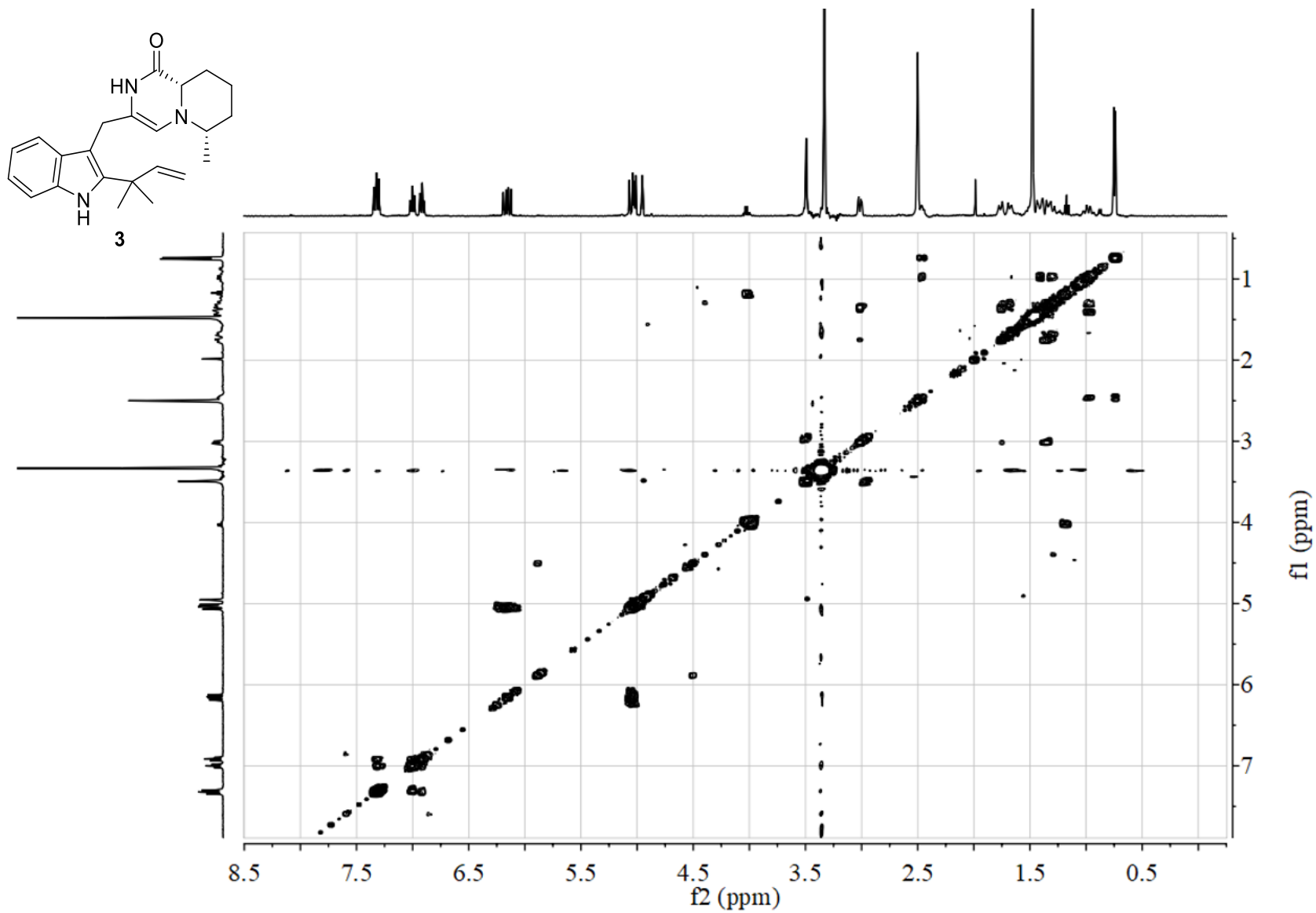
Supplementary Fig. 9-1 | ¹H NMR spectrum of 3 in DMSO-*d*₆.



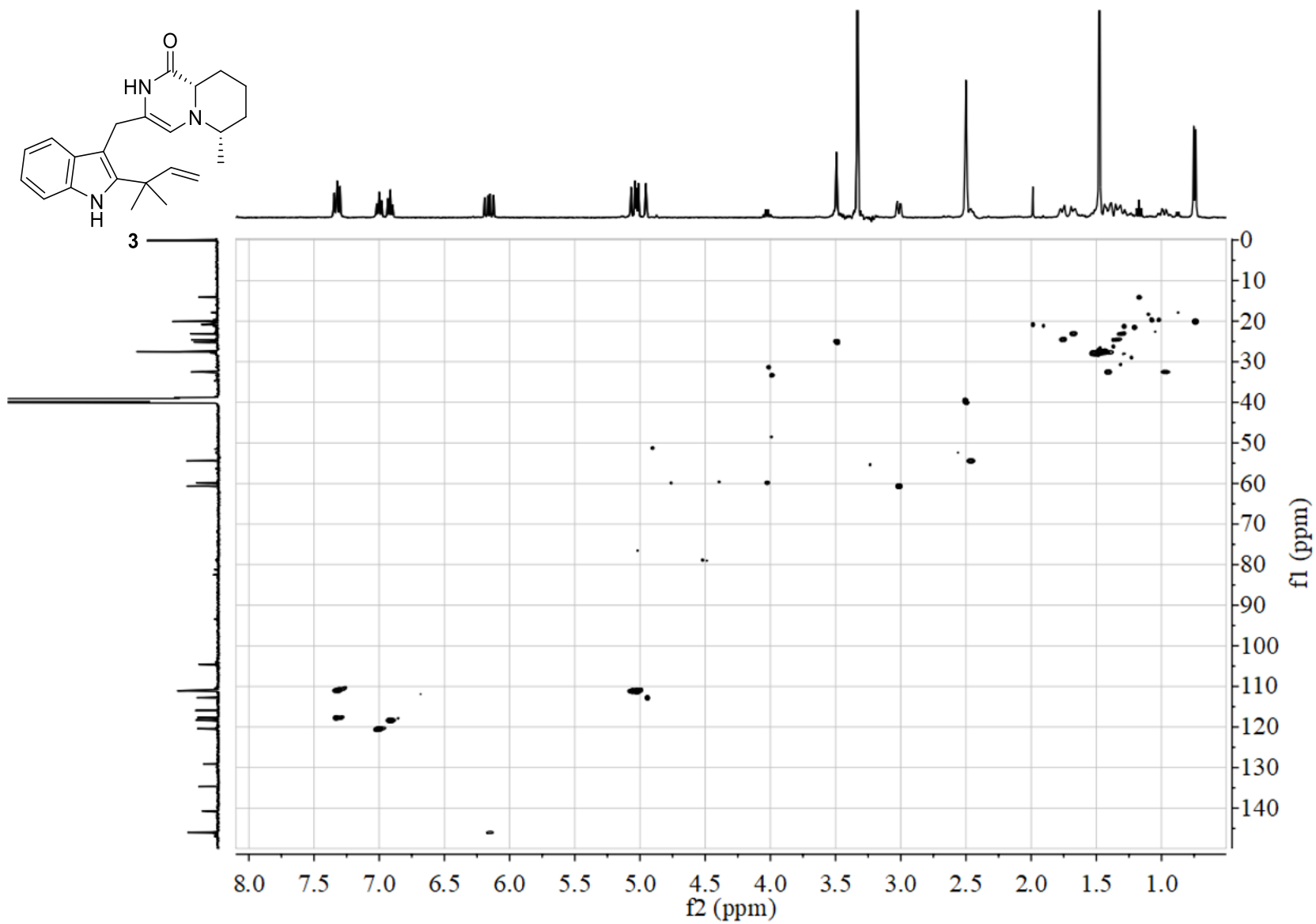
Supplementary Fig. 9-2 | ^{13}C NMR spectrum of 3 in $\text{DMSO-}d_6$.



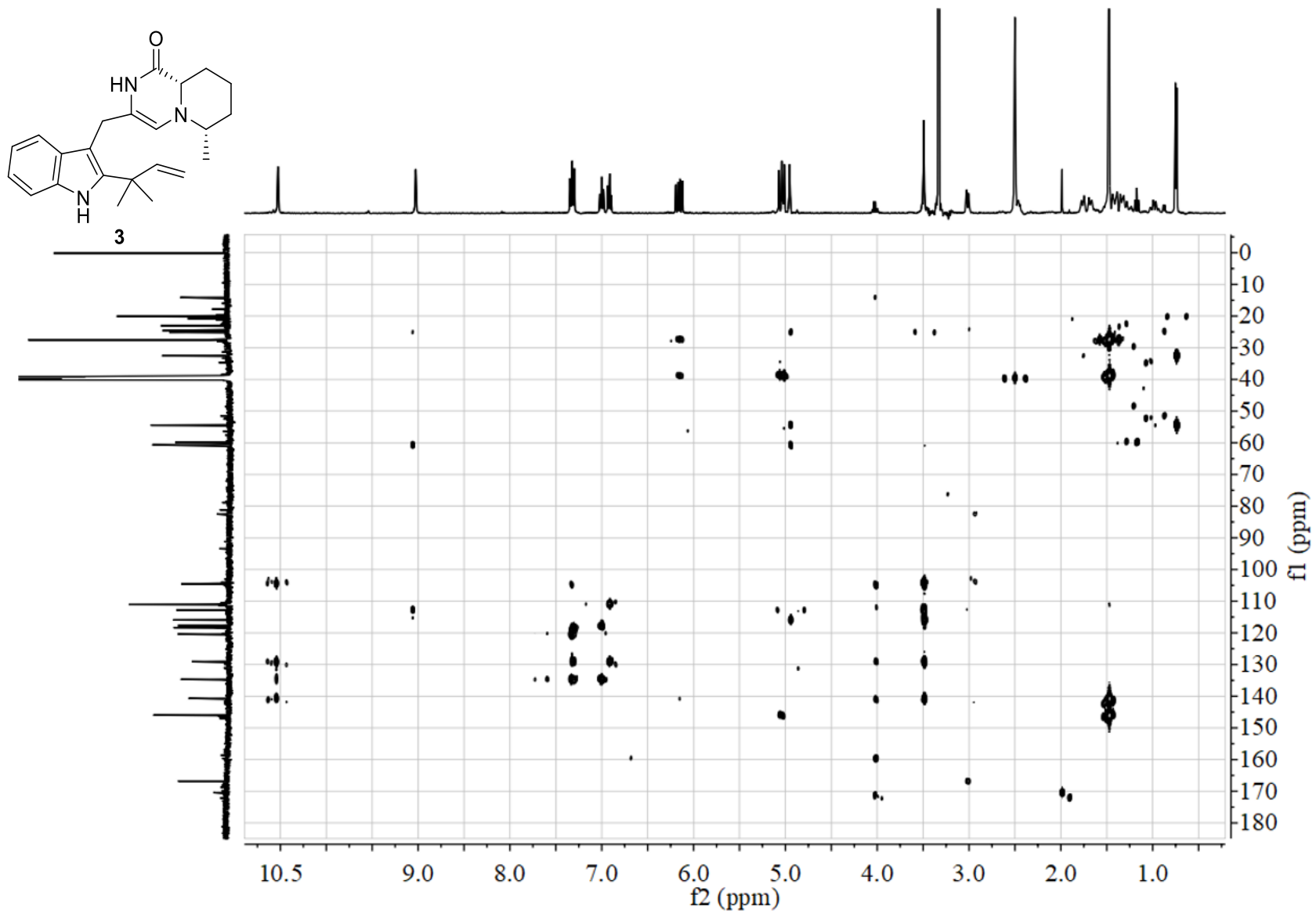
Supplementary Fig. 9-3 | DEPT135 and ¹³C NMR spectra of 3 in DMSO-*d*₆.



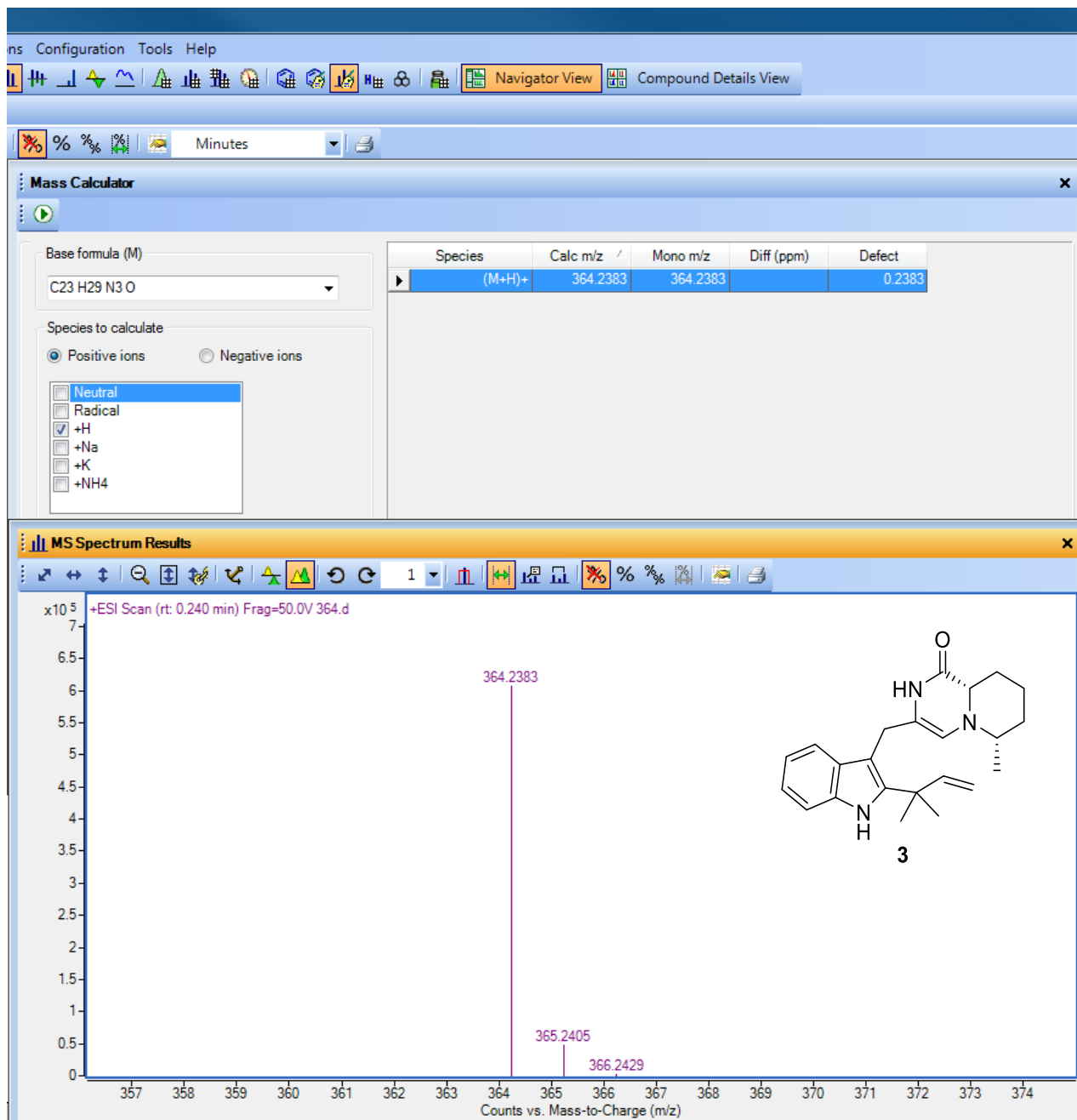
Supplementary Fig. 9-4 | ^1H - ^1H COSY NMR spectrum of **3** in $\text{DMSO-}d_6$.



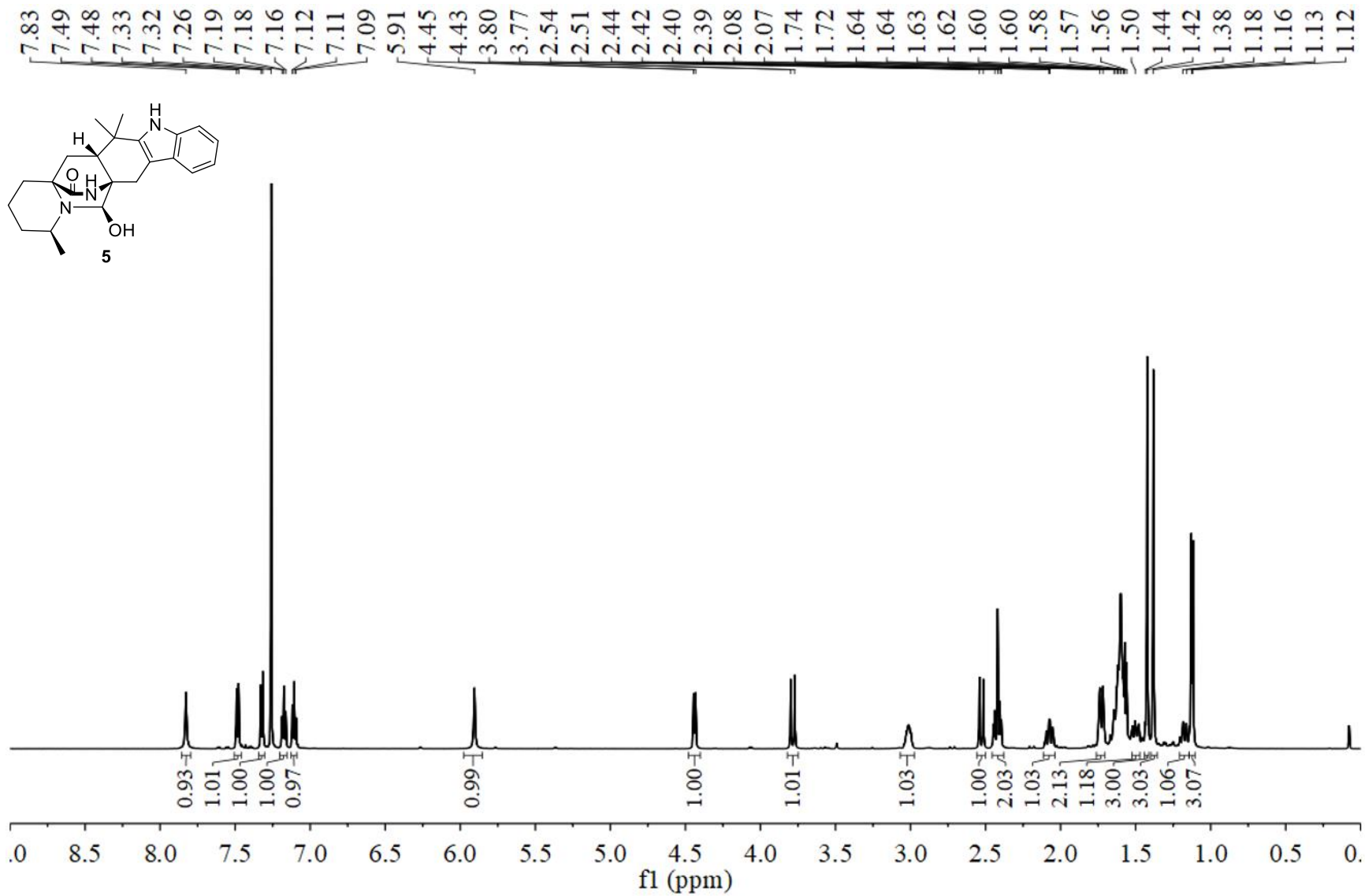
Supplementary Fig. 9-5 | HSQC NMR spectrum of 3 in DMSO-*d*₆.



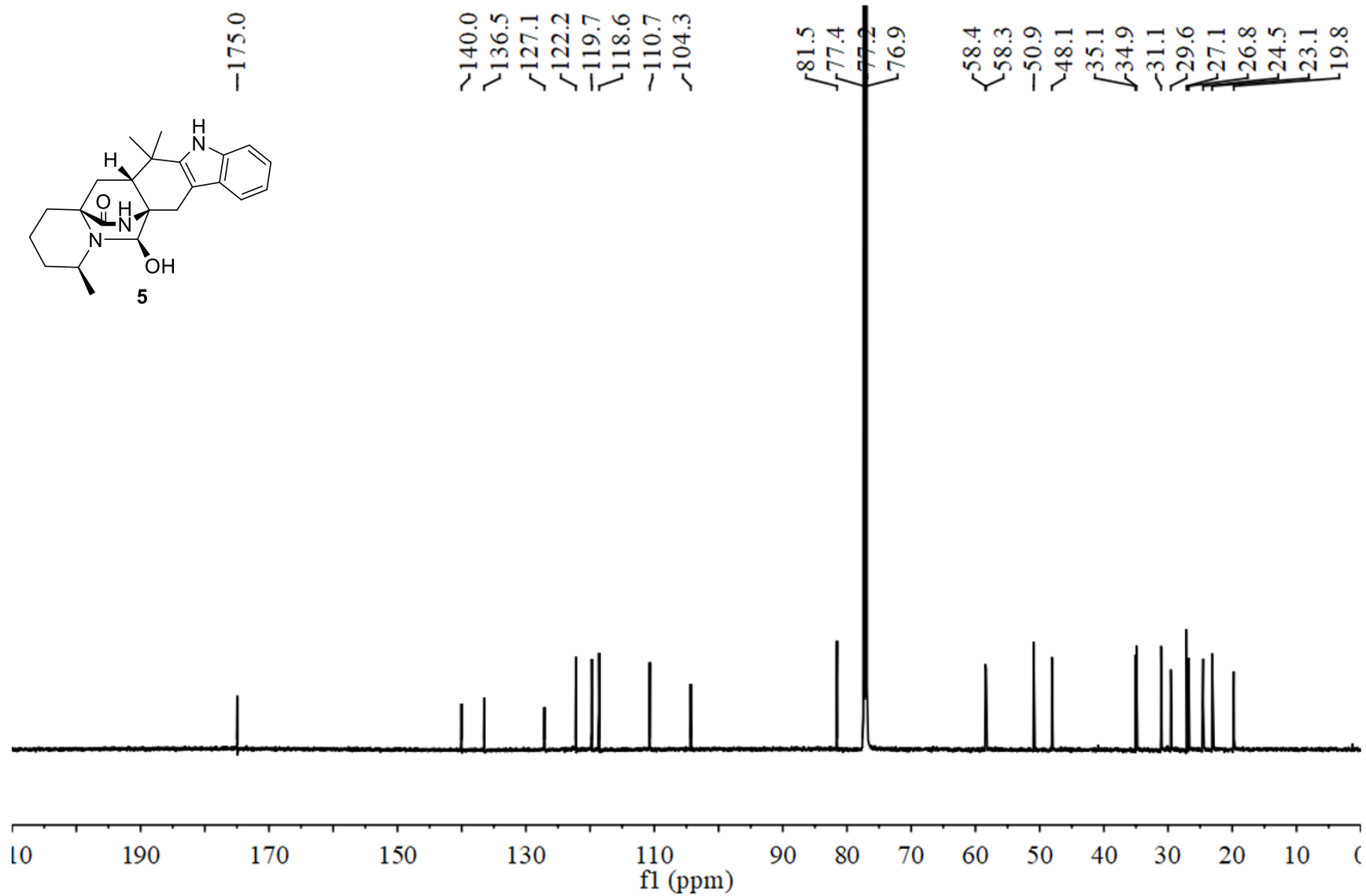
Supplementary Fig. 9-6 | HMBC NMR spectrum of **3** in DMSO-*d*₆.



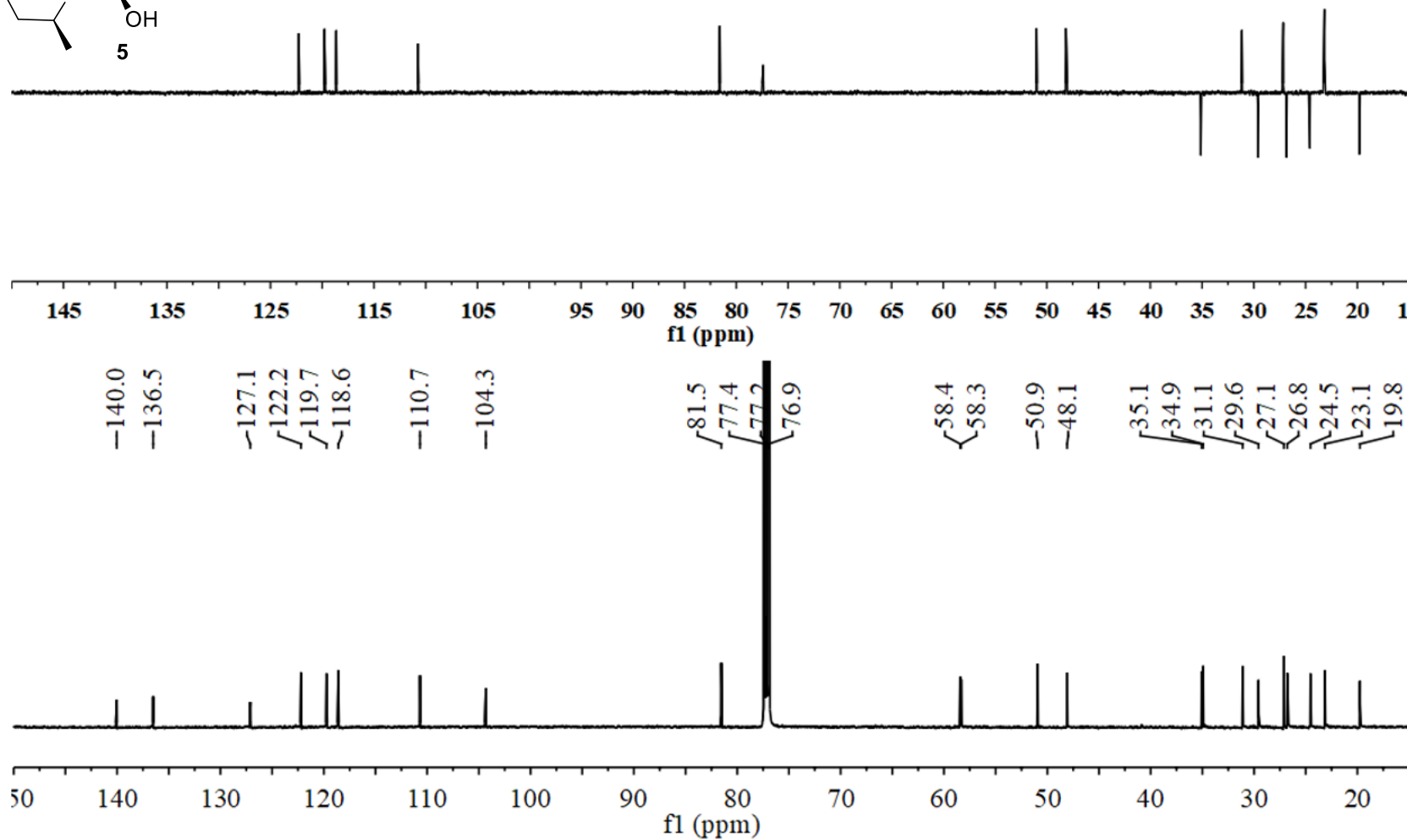
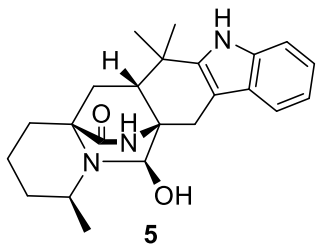
Supplementary Fig. 9-7 | HRMS spectrum of 3.



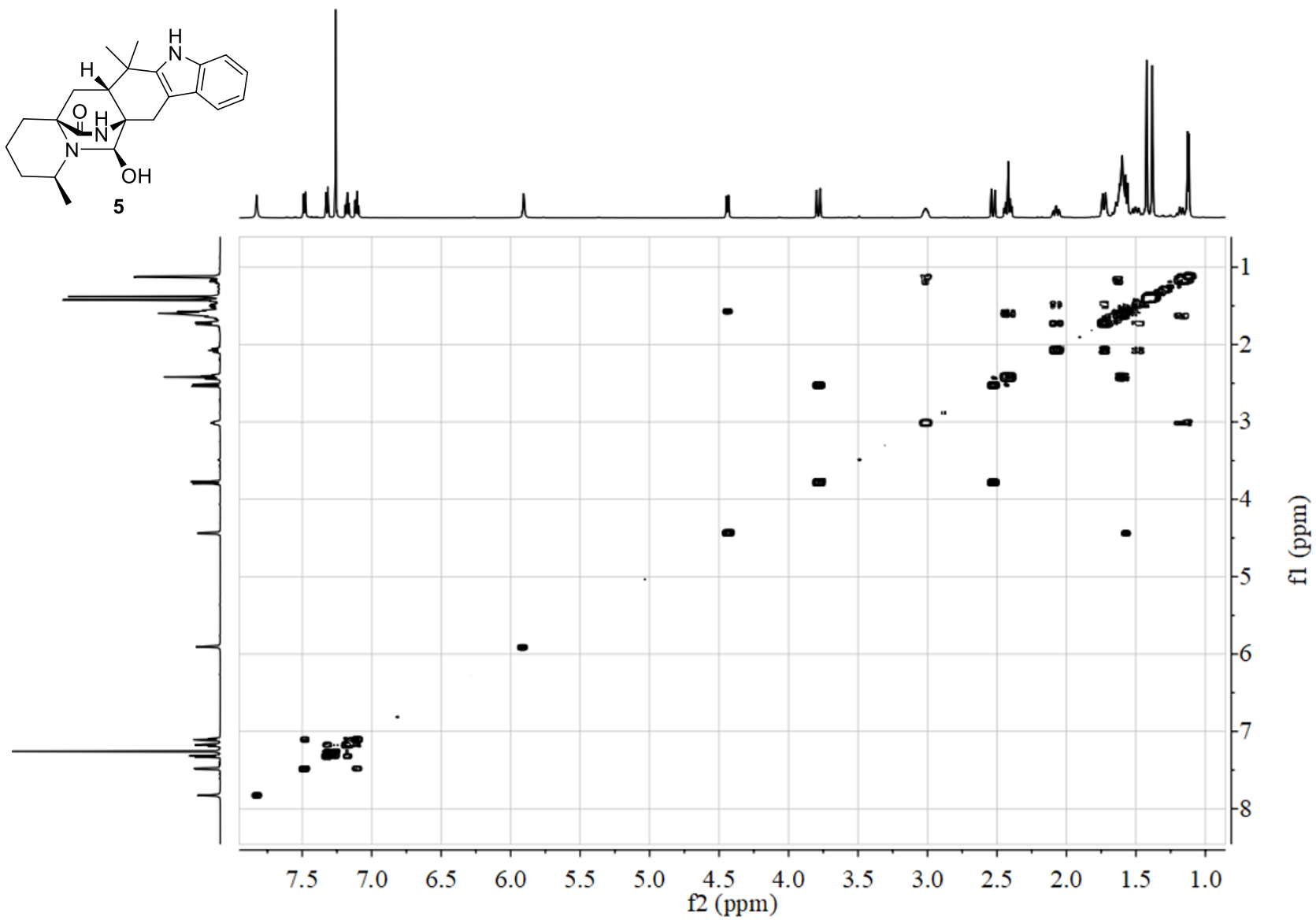
Supplementary Fig. 10-1 | ^1H NMR spectrum of 5 in CDCl_3 .



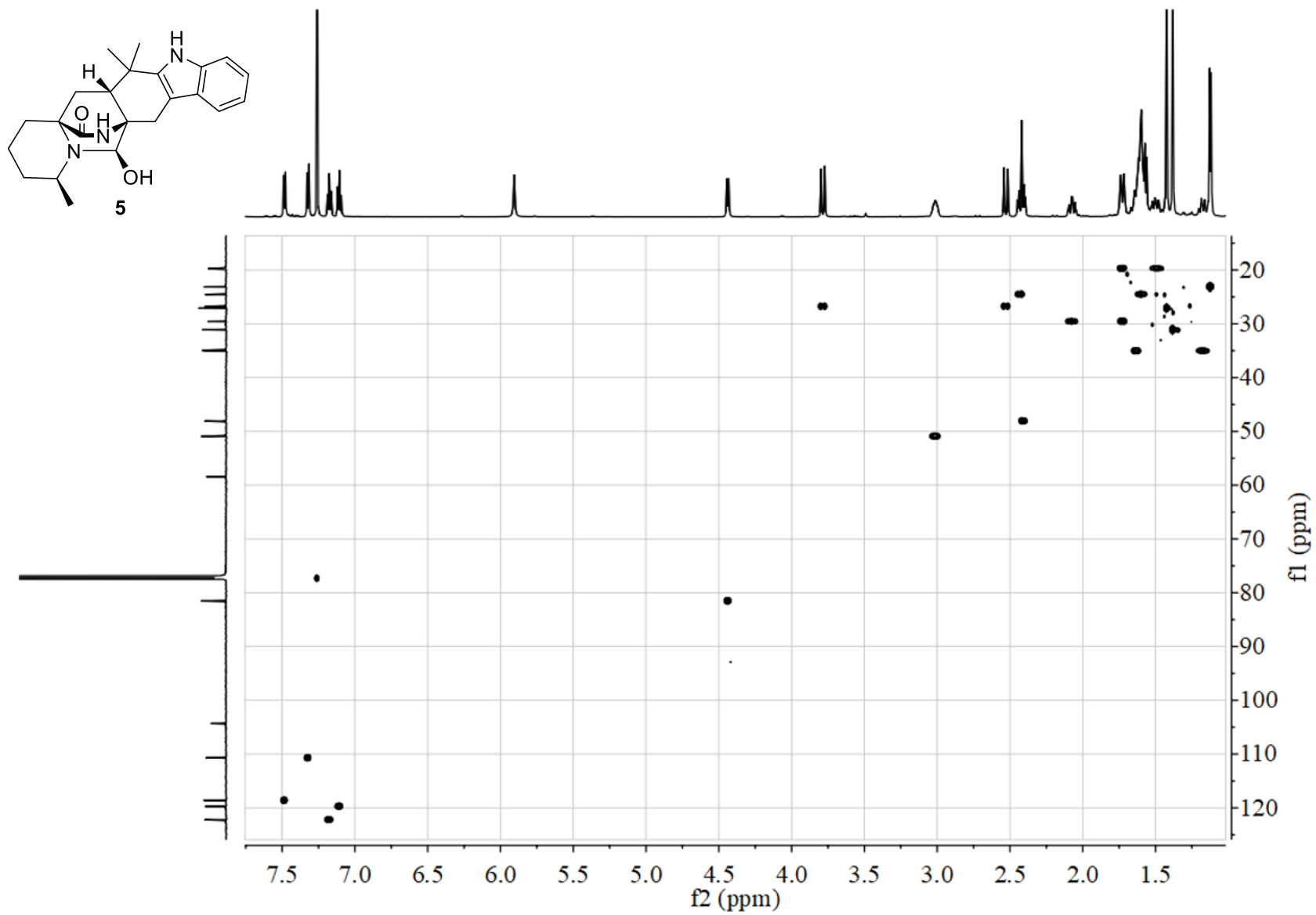
Supplementary Fig. 10-2 | ¹³C NMR spectrum of 5 in CDCl₃.



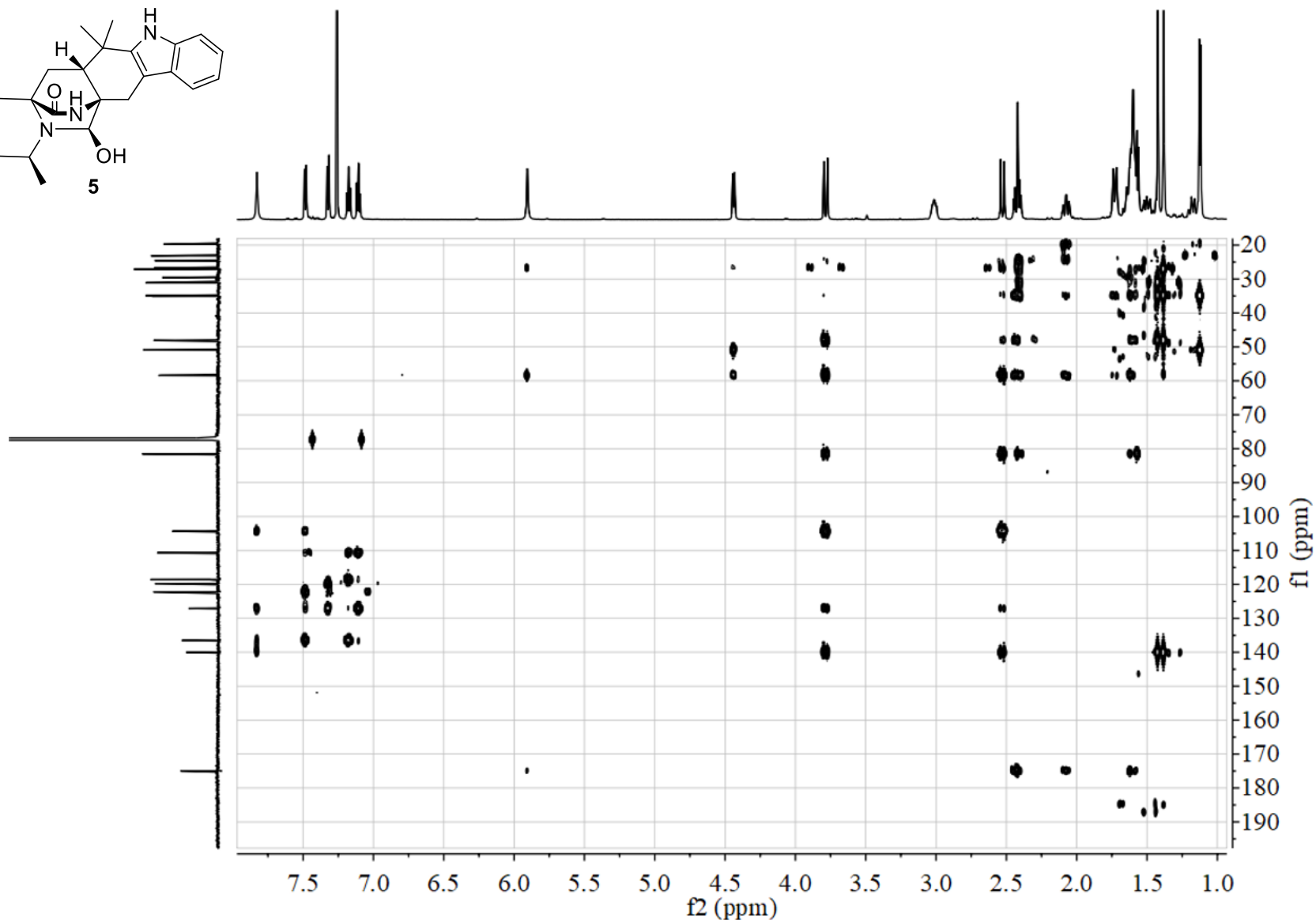
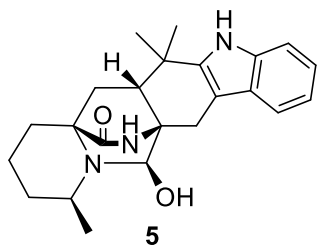
Supplementary Fig. 10-3 | DEPT135 and ^{13}C NMR spectra of 5 in CDCl_3 .



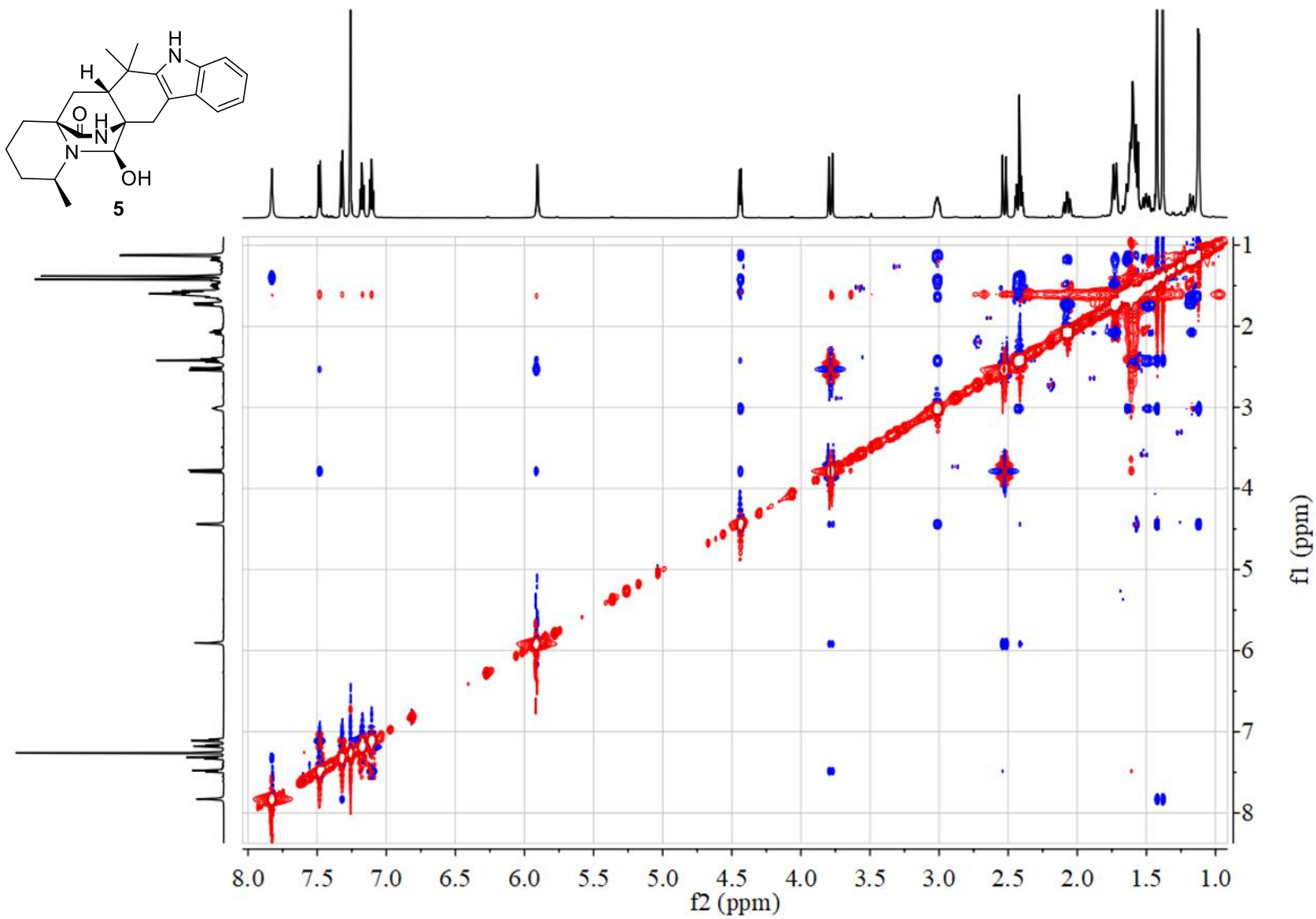
Supplementary Fig. 10-4 | ^1H - ^1H COSY NMR spectrum of 5 in CDCl_3 .



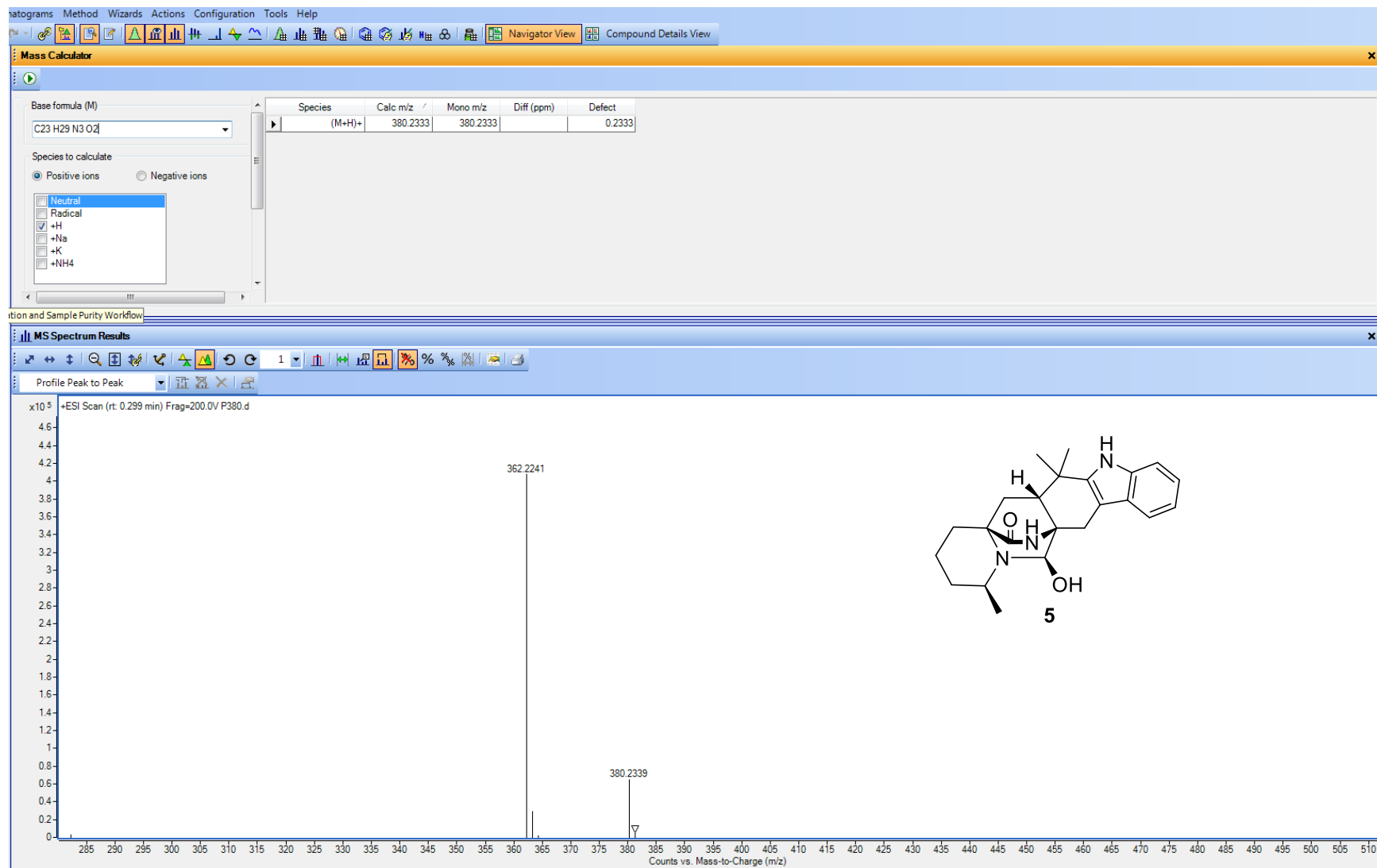
Supplementary Fig. 10-5 | HSQC NMR spectrum of 5 in CDCl₃.



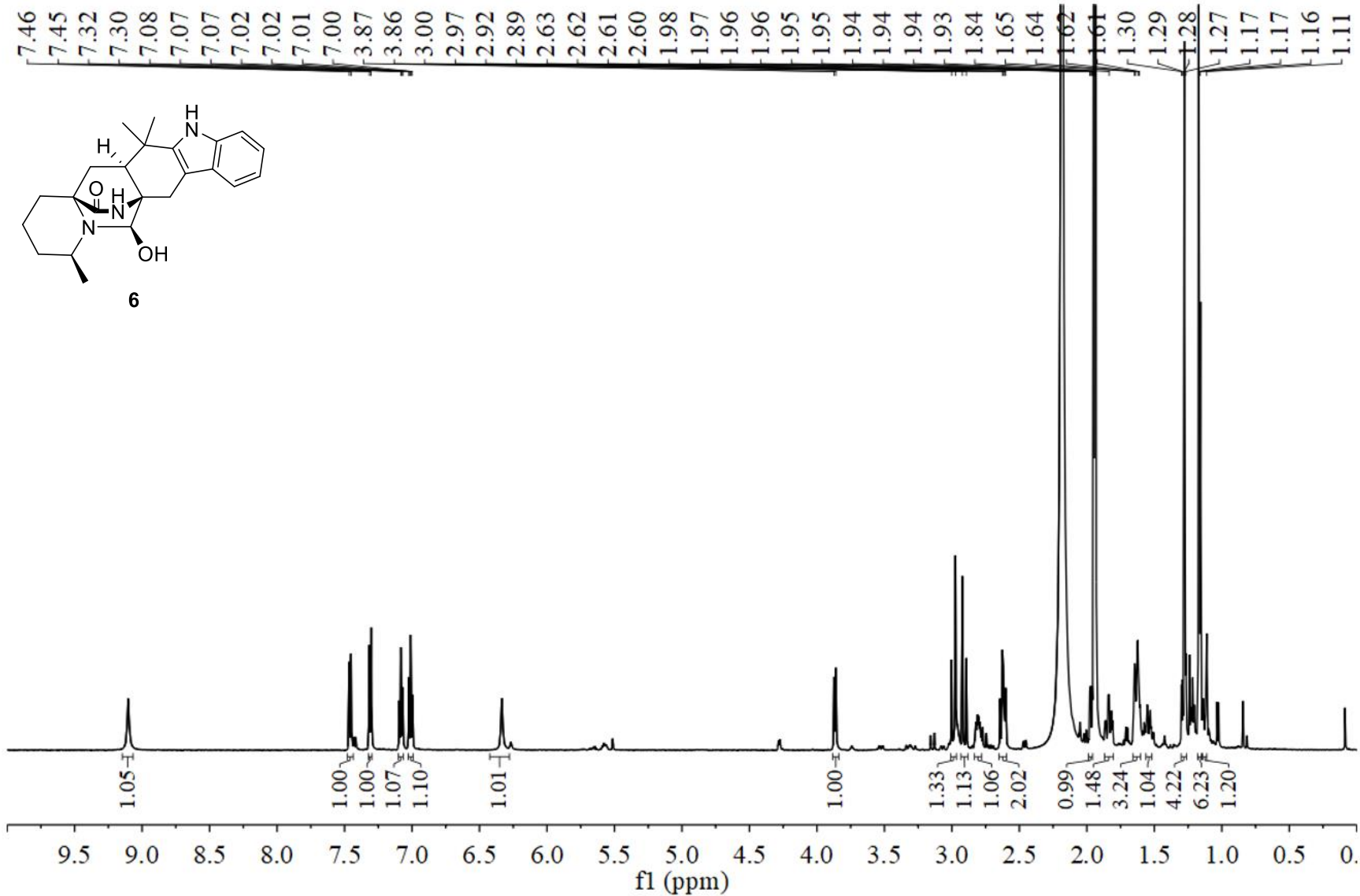
Supplementary Fig. 10-6 | HMBC NMR spectrum of 5 in CDCl₃.



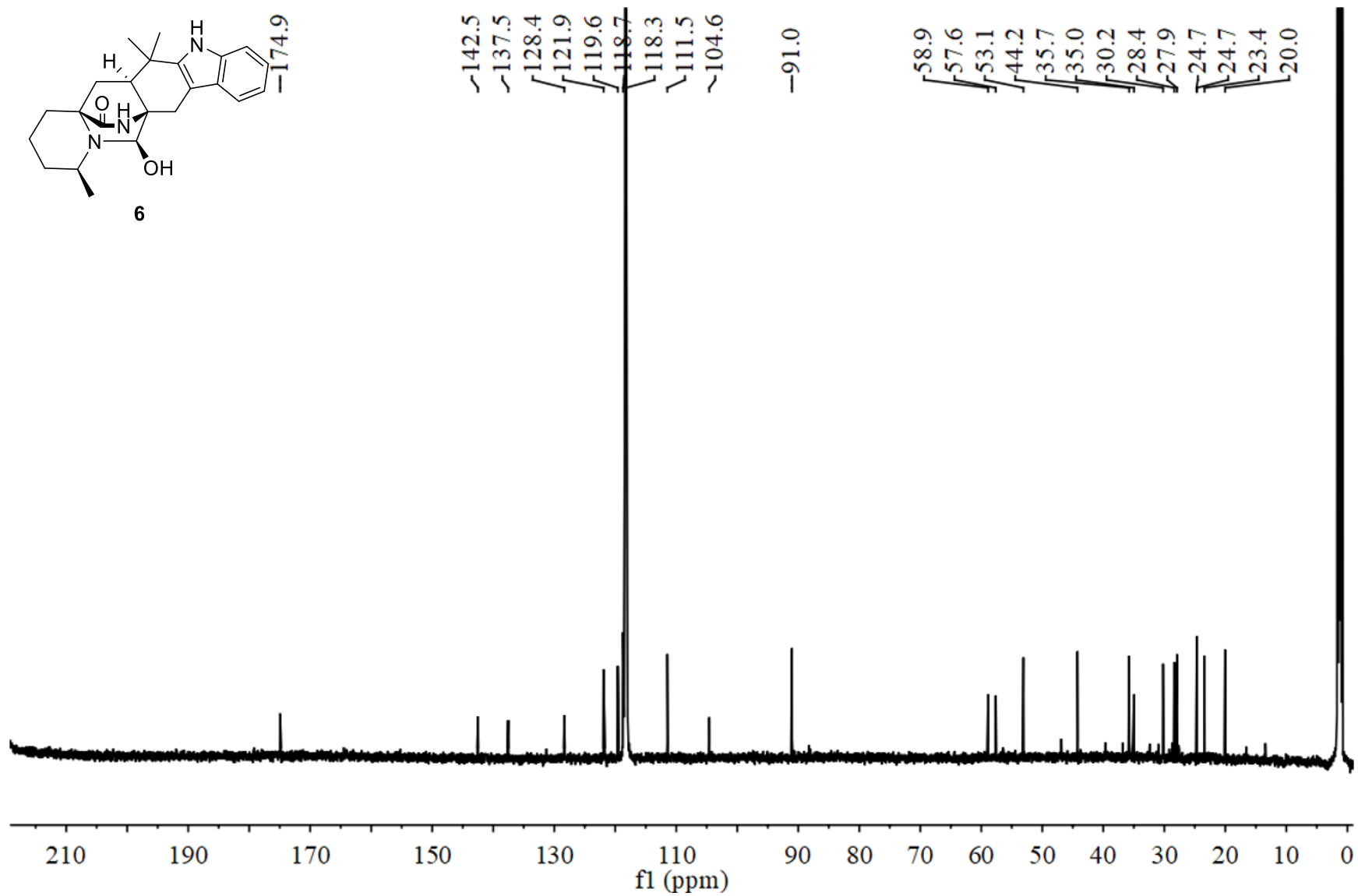
Supplementary Fig. 10-7 | NOESY NMR spectrum of 5 in CDCl₃.



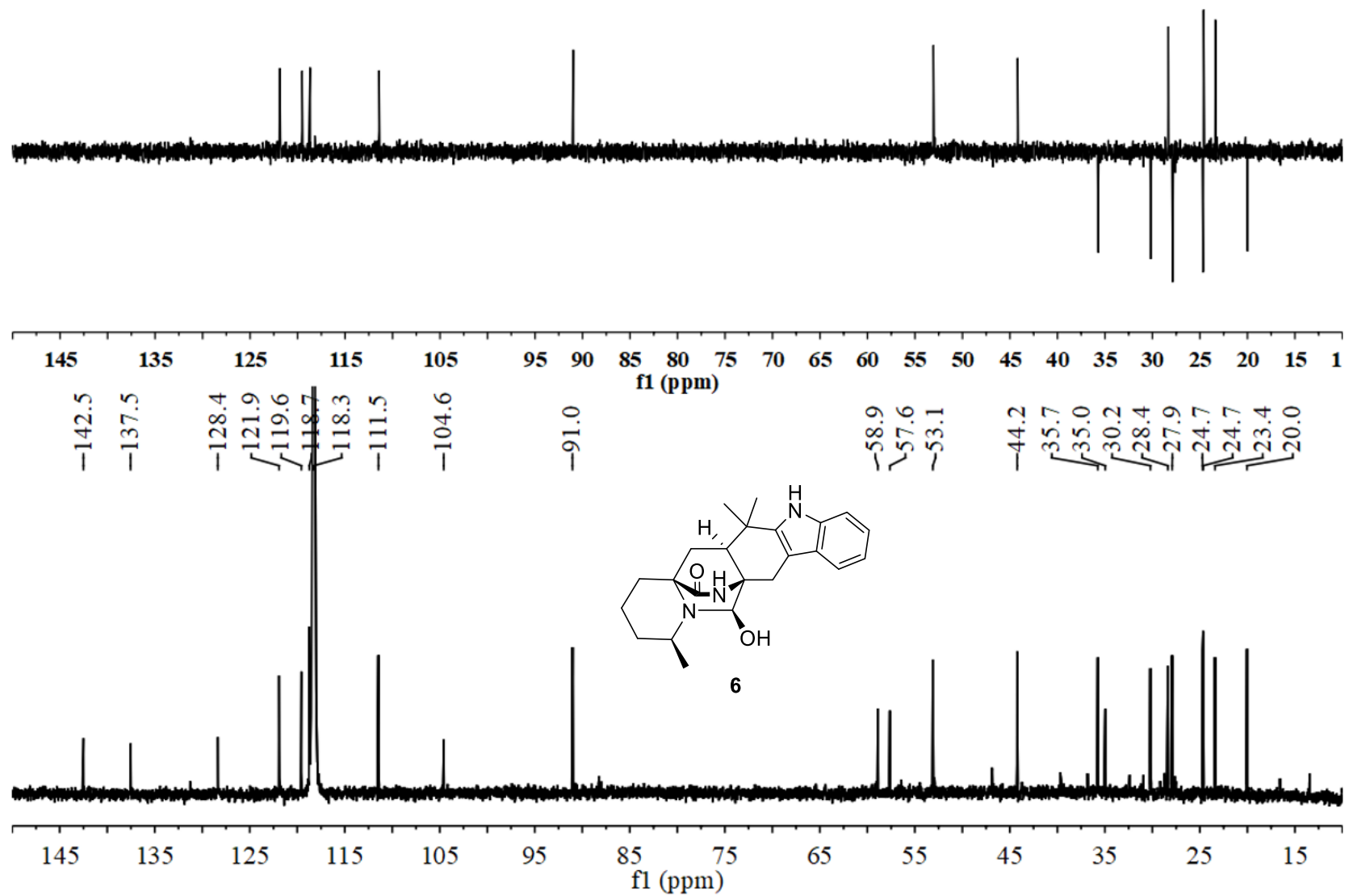
Supplementary Fig. 10-8 | HRMS spectrum of 5.



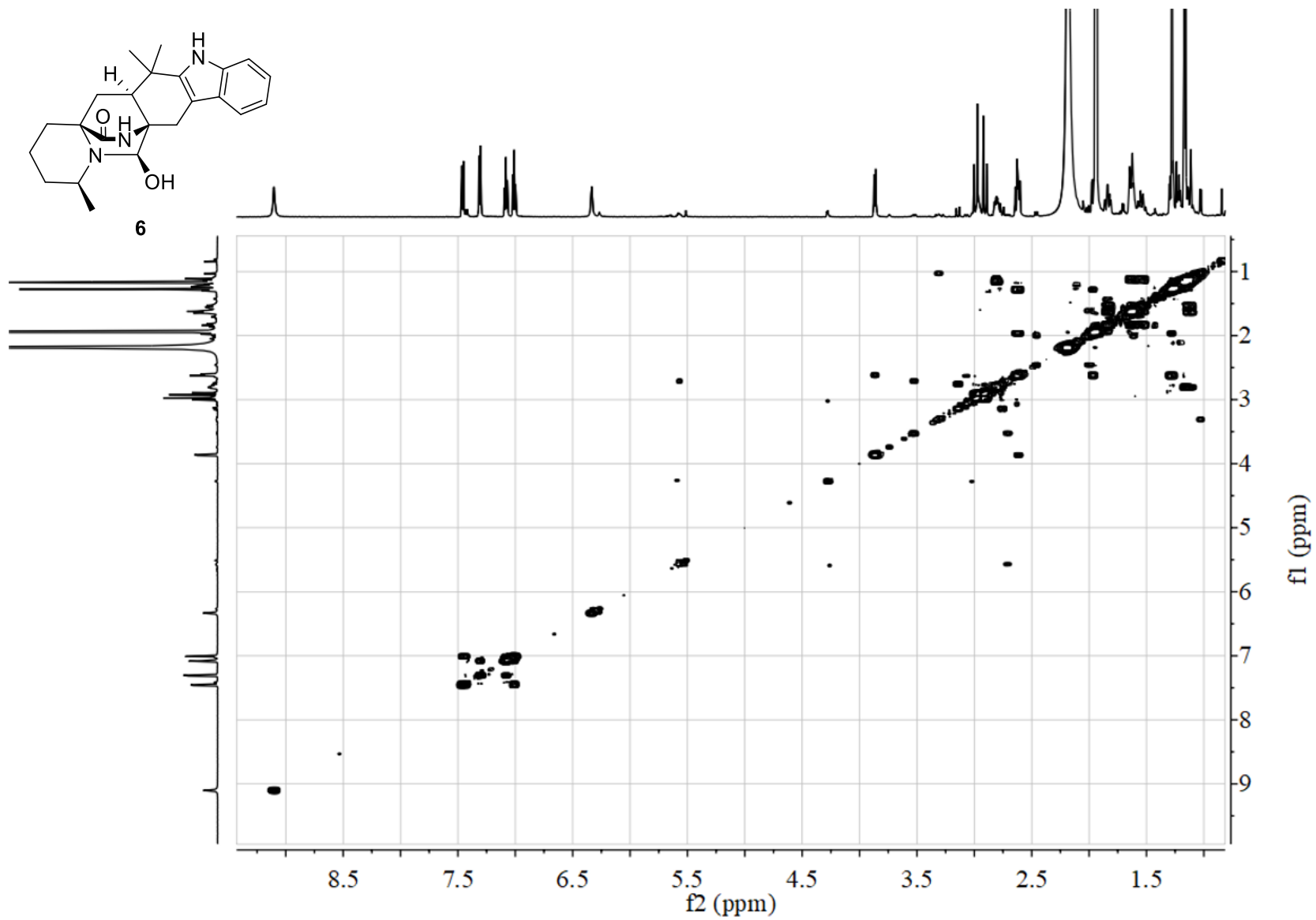
Supplementary Fig. 11-1 | ¹H NMR spectrum of 6 in CD₃CN.



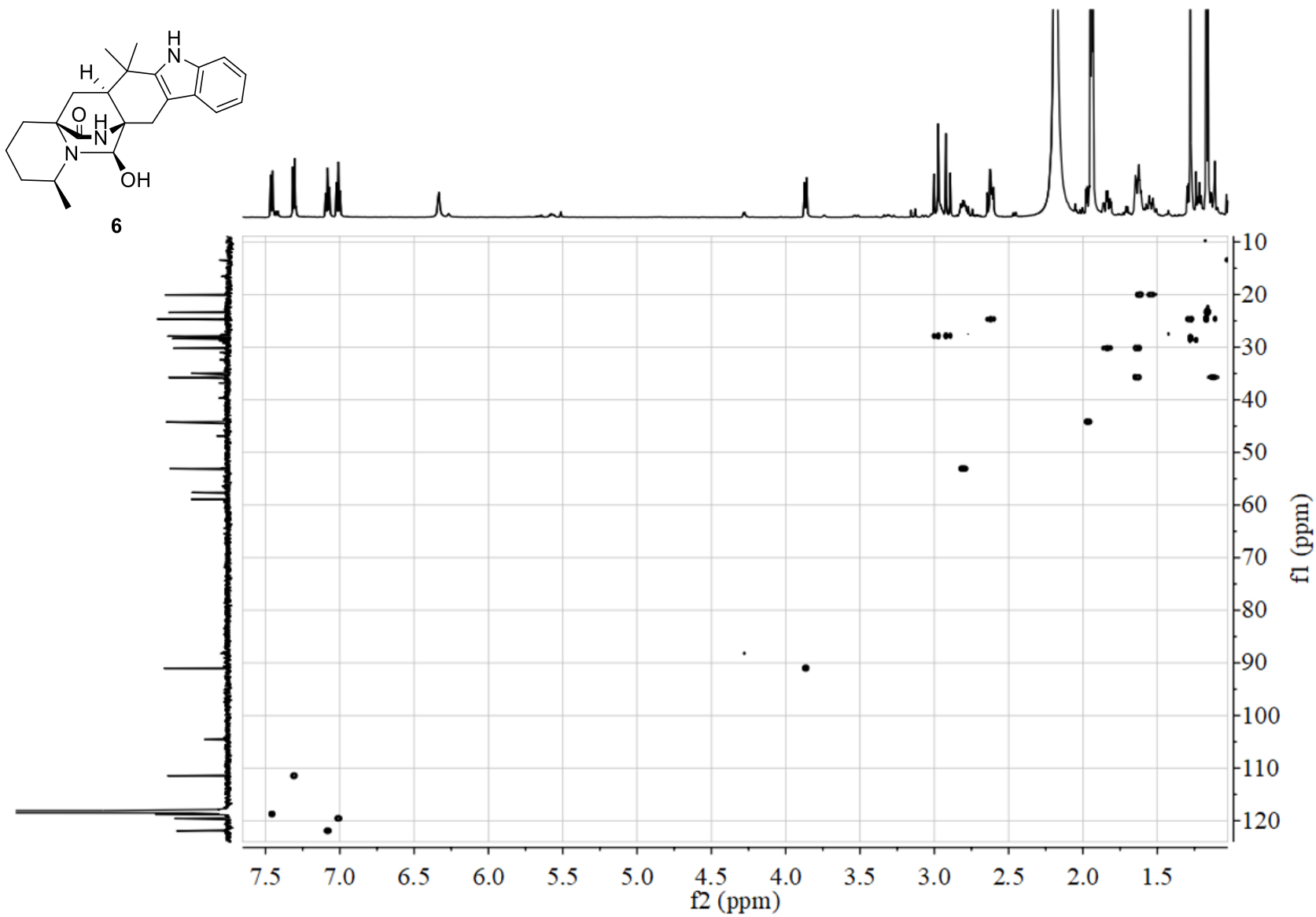
Supplementary Fig. 11-2 | ^{13}C NMR spectrum of 6 in CD_3CN .



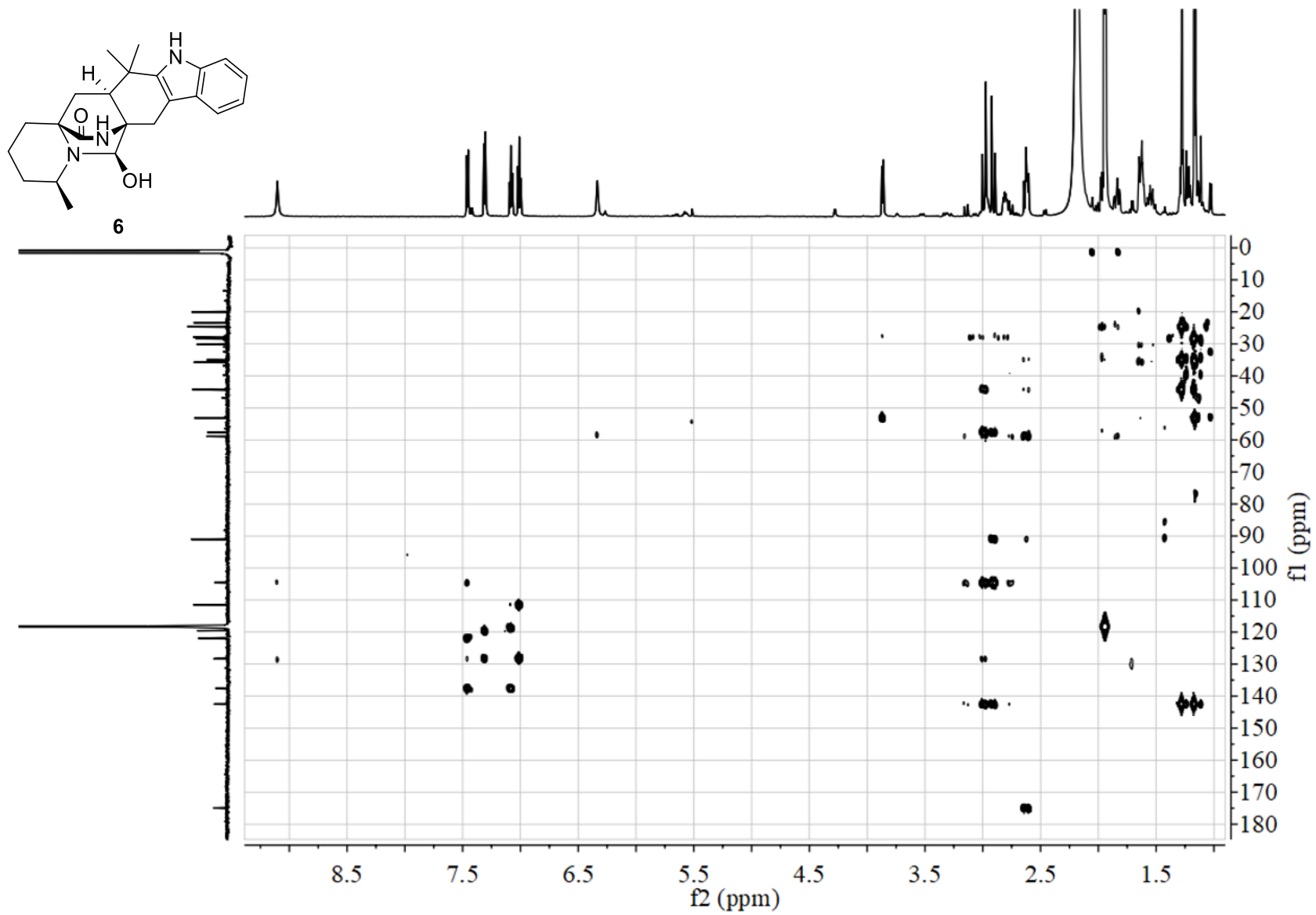
Supplementary Fig. 11-3 | DEPT135 and ¹³C NMR spectra of 6 in CD₃CN.



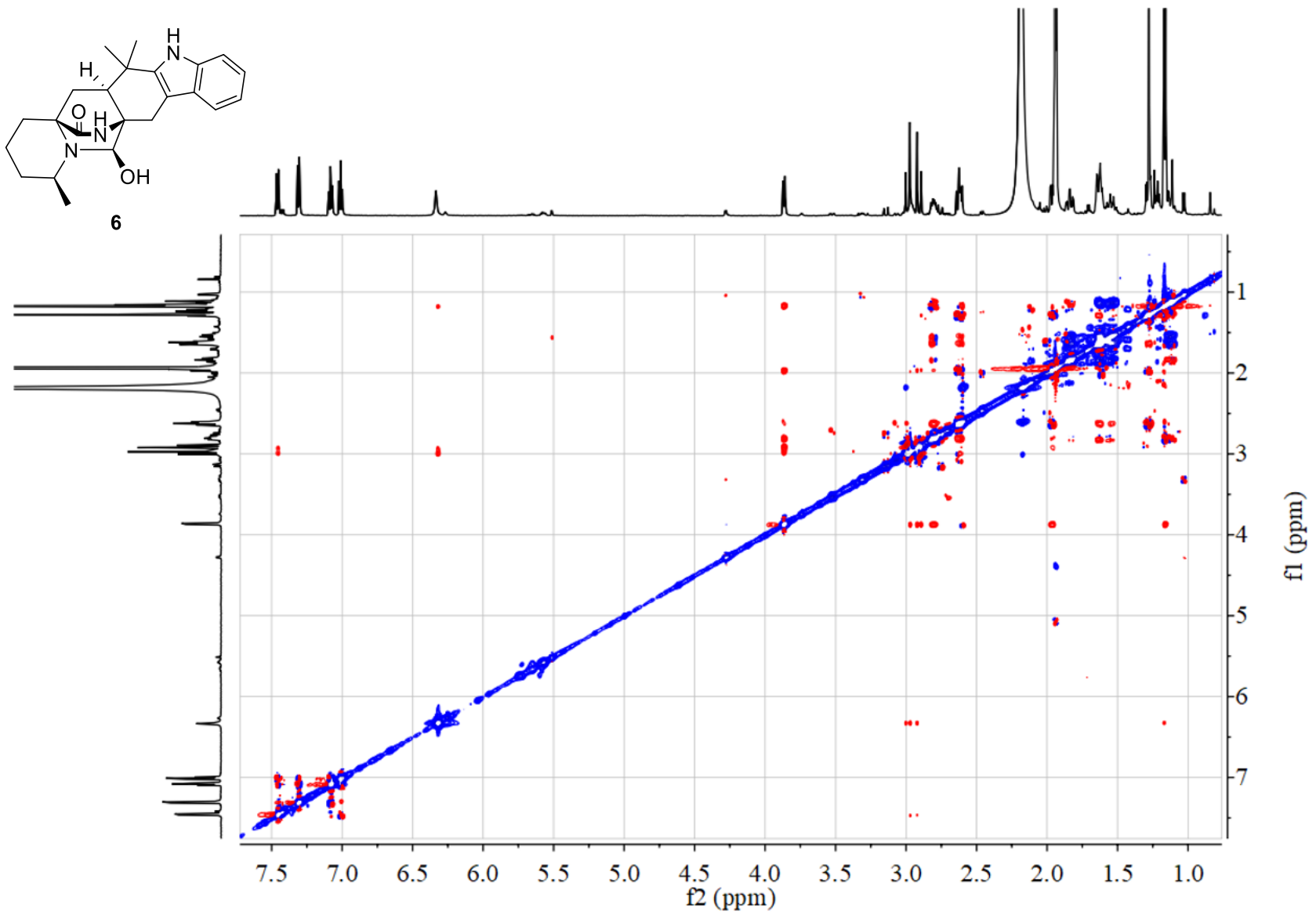
Supplementary Fig. 11-4 | ^1H - ^1H COSY NMR spectrum of 6 in CD_3CN .



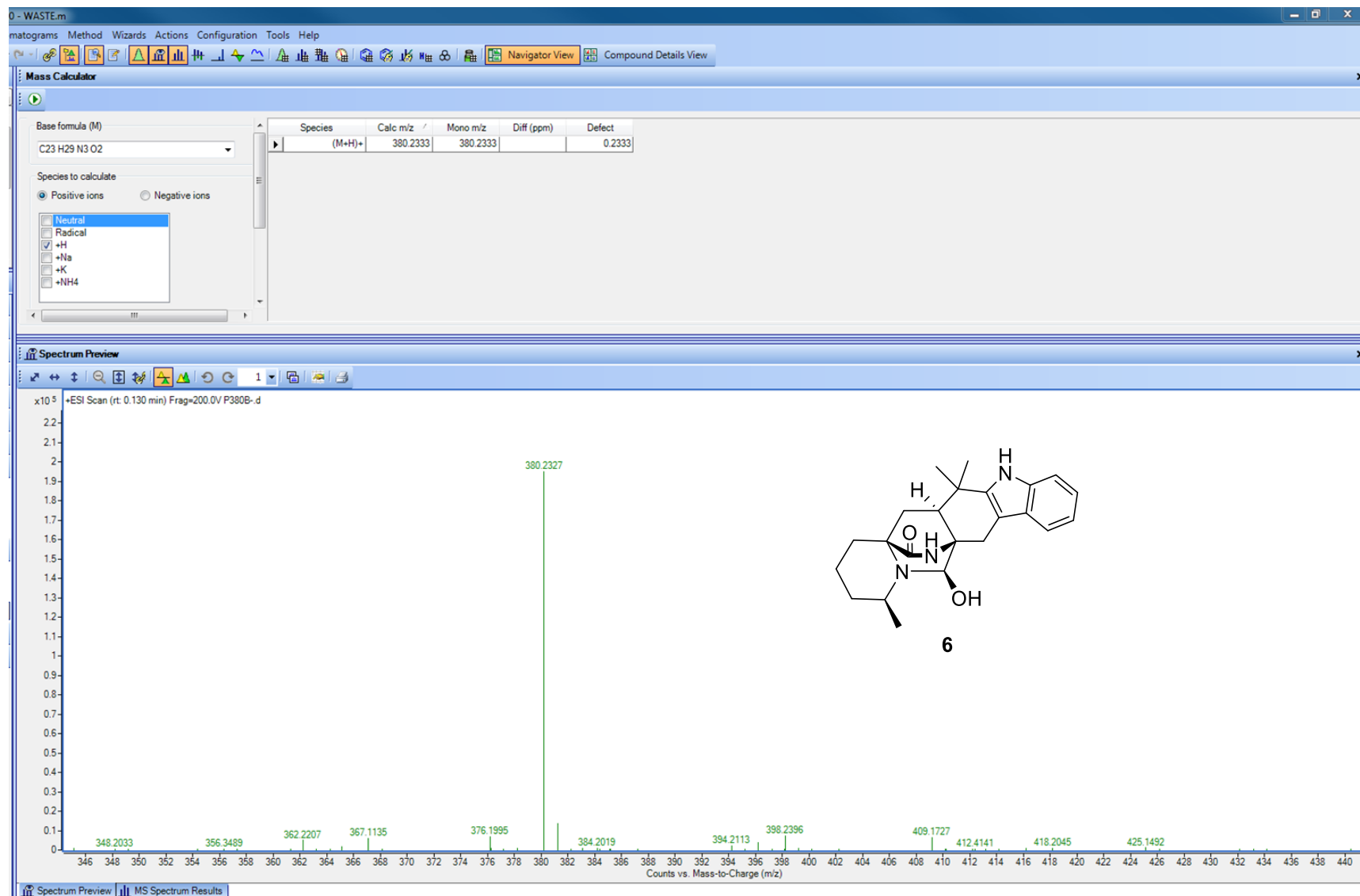
Supplementary Fig. 11-5 | HSQC NMR spectrum of 6 in CD₃CN.



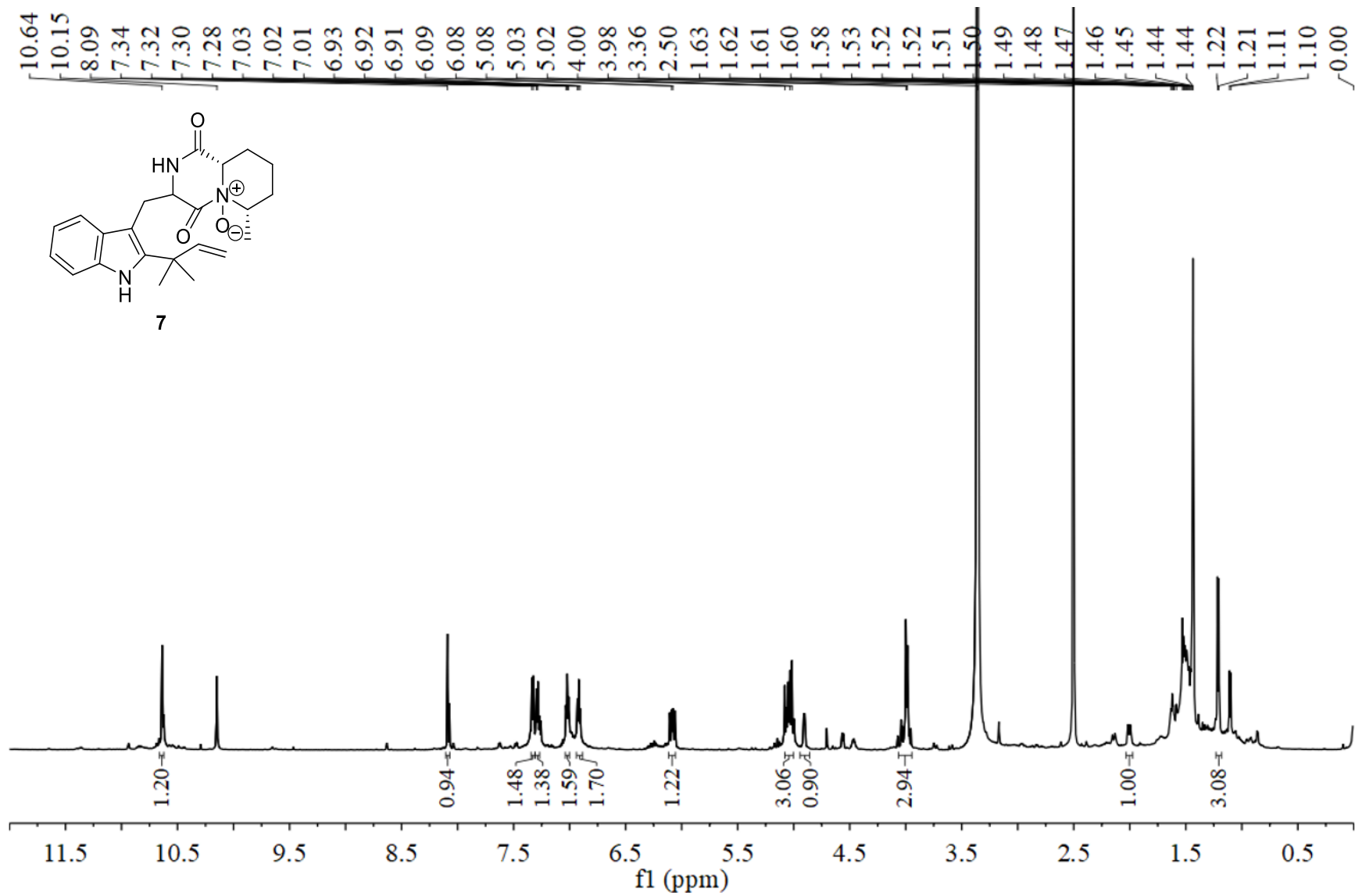
Supplementary Fig. 11-6 | HMBC NMR spectrum of **6** in CD₃CN.



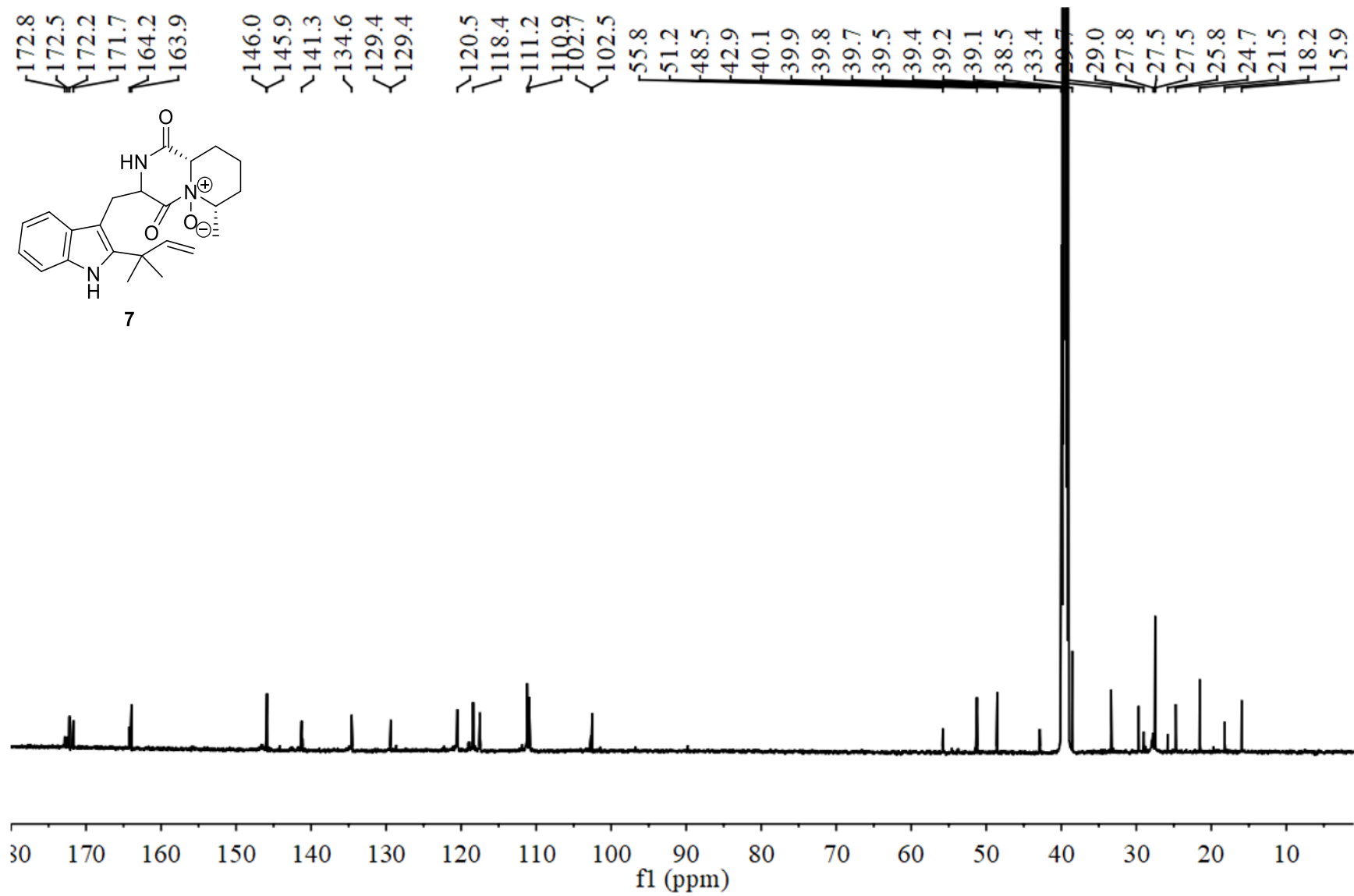
Supplementary Fig. 11-7 | NOESY NMR spectrum of 6 in CD₃CN.



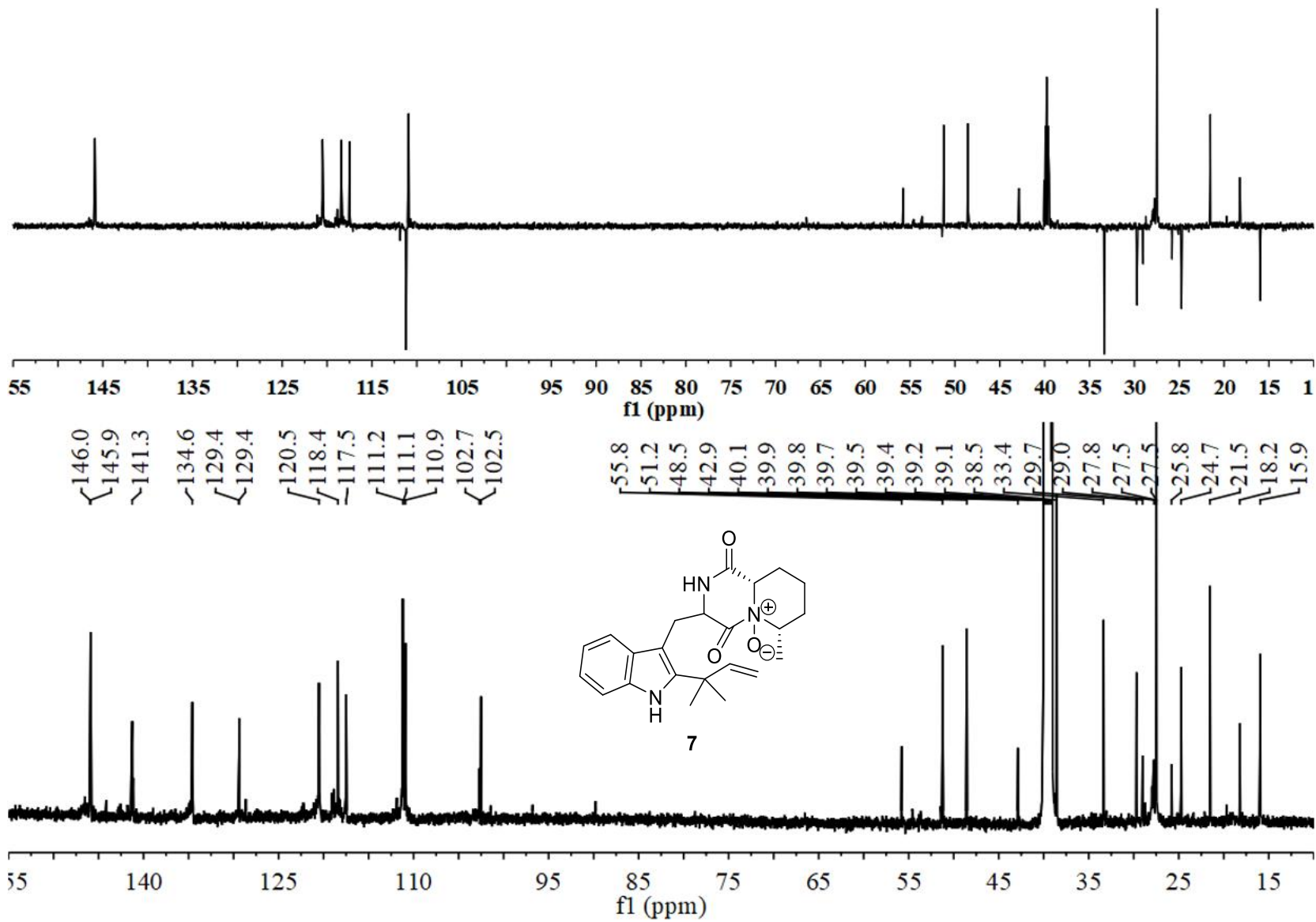
Supplementary Fig. 11-8 | HRMS spectrum of 6.



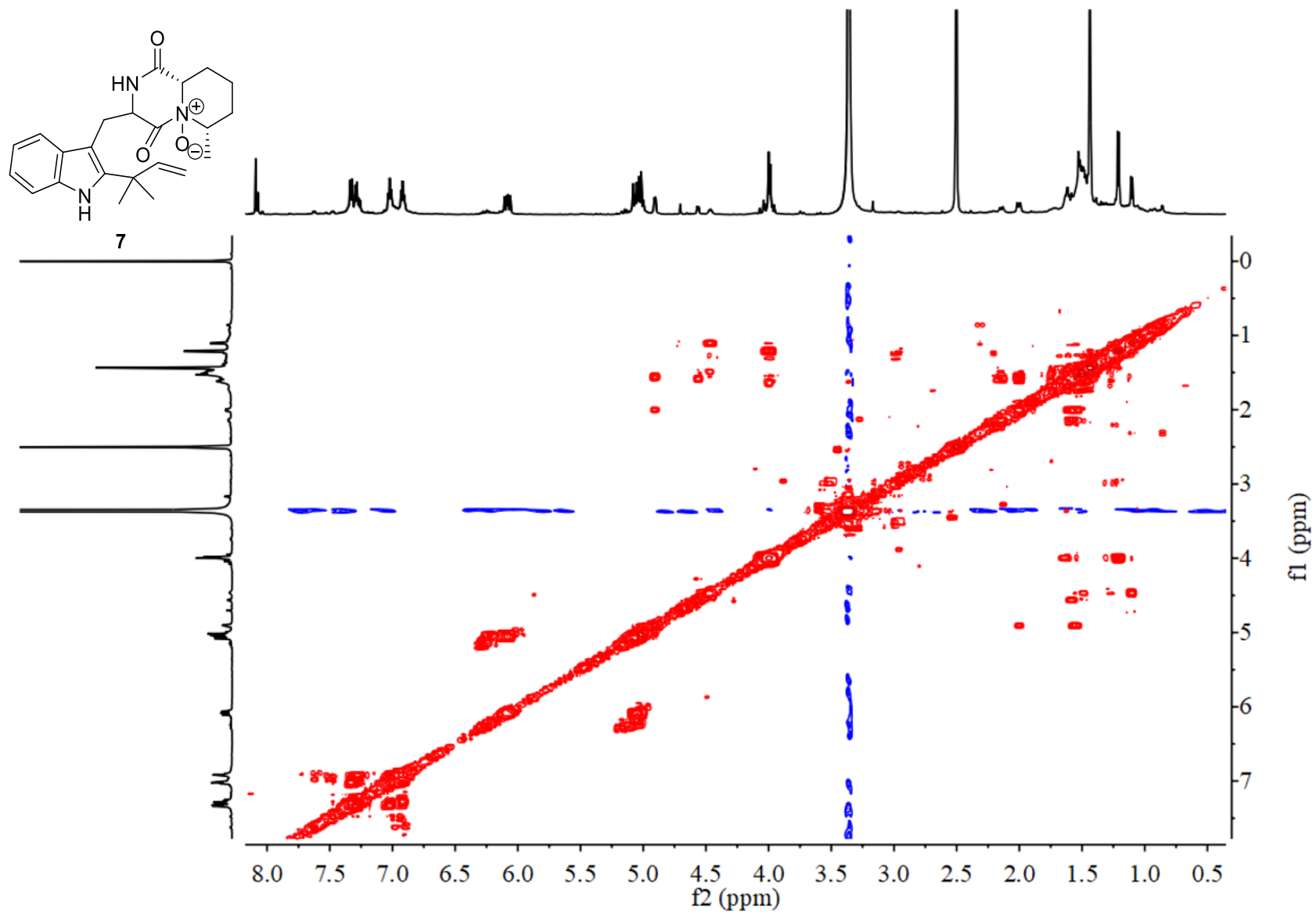
Supplementary Fig. 12-1 | ¹H NMR spectrum of 7 in DMSO-*d*₆.



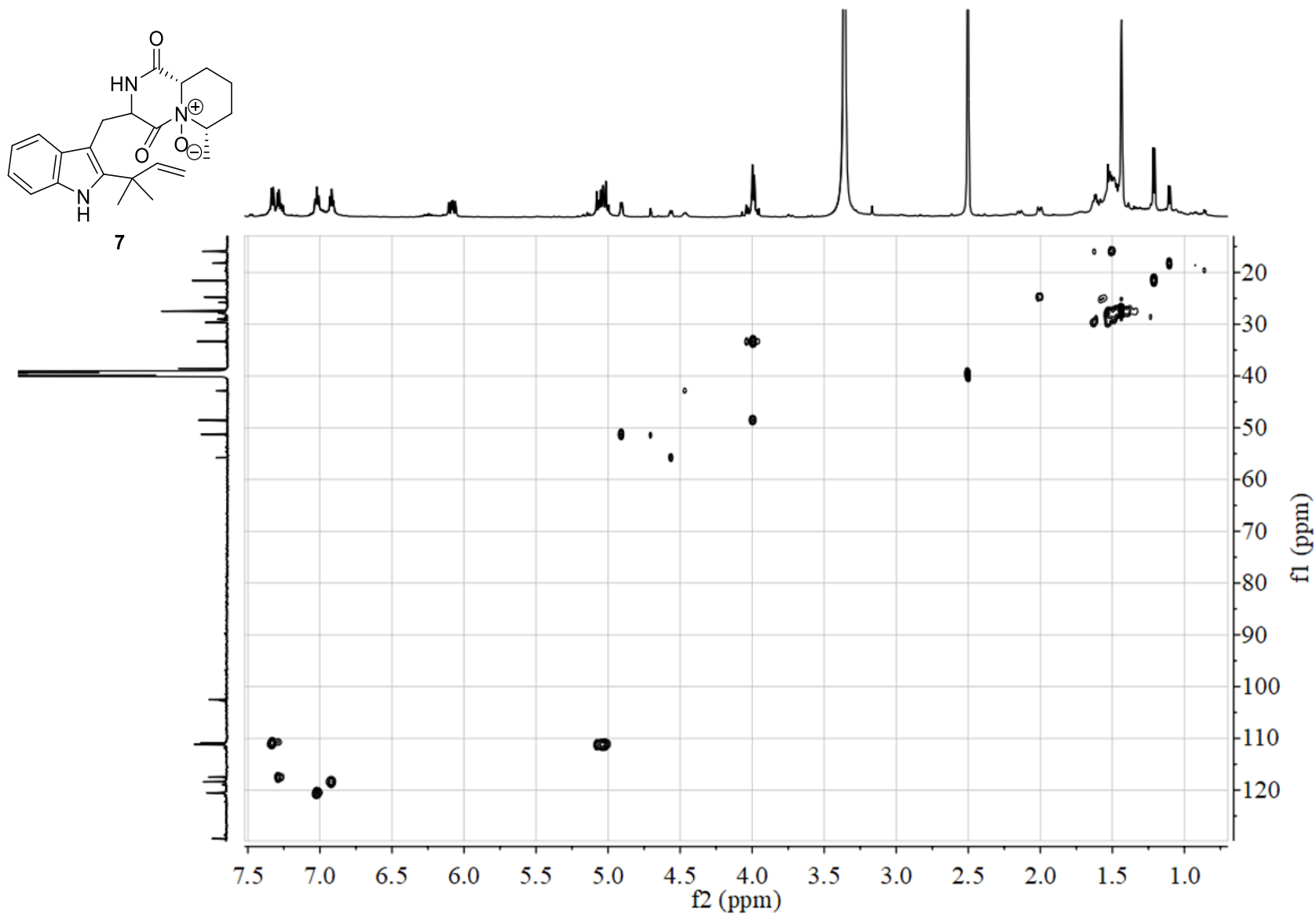
Supplementary Fig. 12-2 | ^{13}C NMR spectrum of 7 in $\text{DMSO-}d_6$.



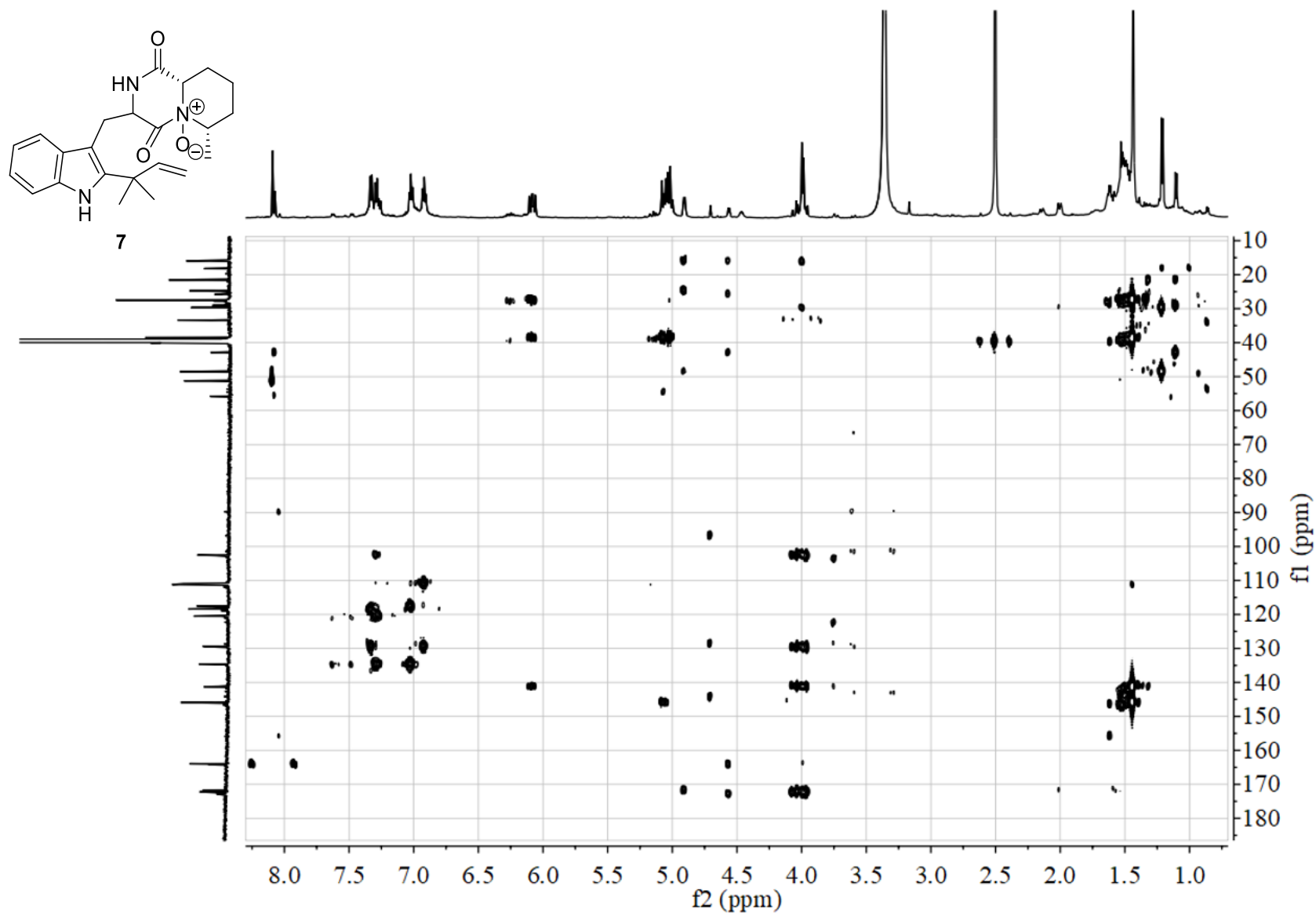
Supplementary Fig. 12-3 | DEPT135 and ¹³C NMR spectra of 7 in DMSO-*d*₆.



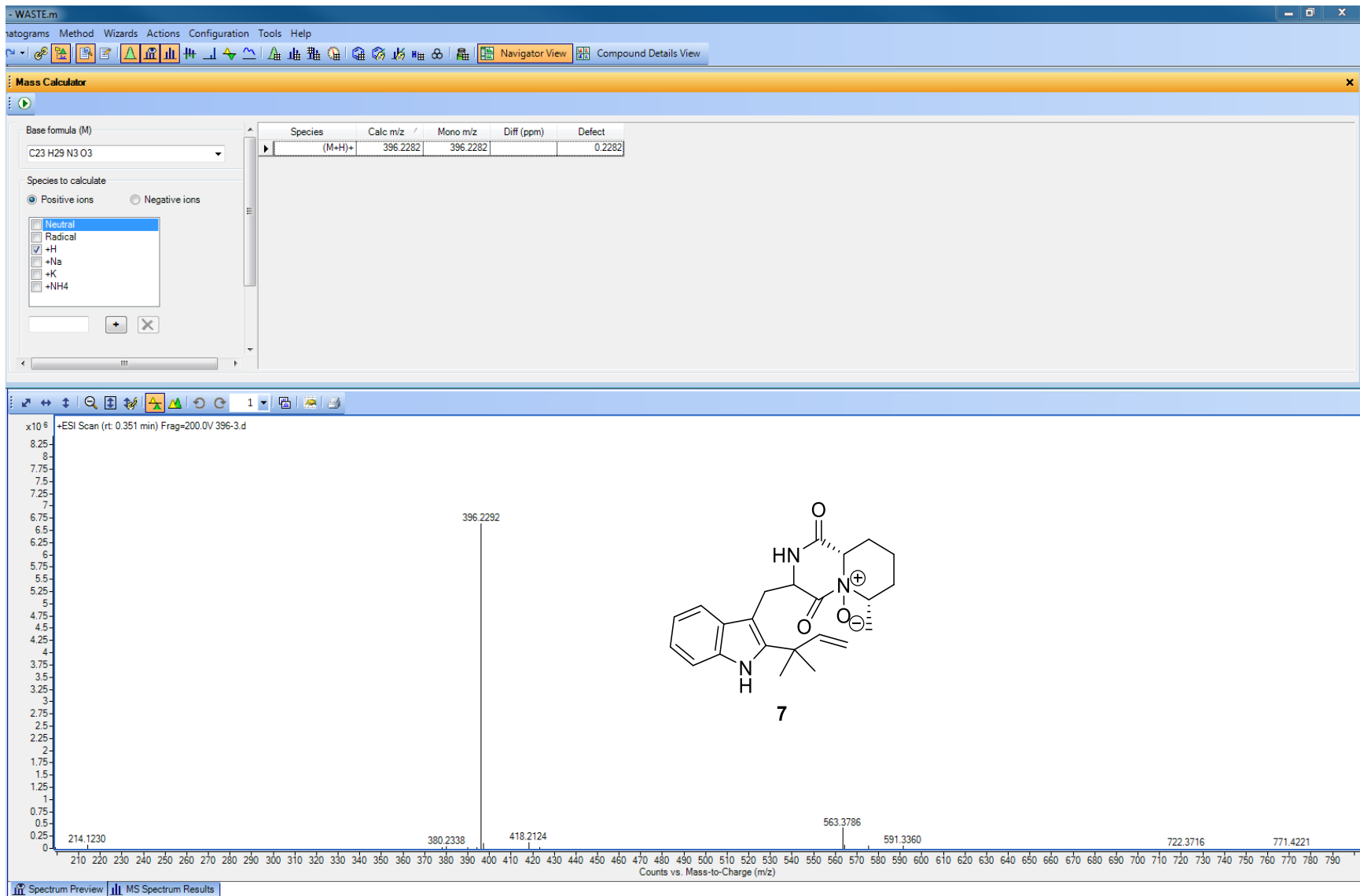
Supplementary Fig. 12-4 | ^1H - ^1H COSY NMR spectrum of **7** in $\text{DMSO-}d_6$.



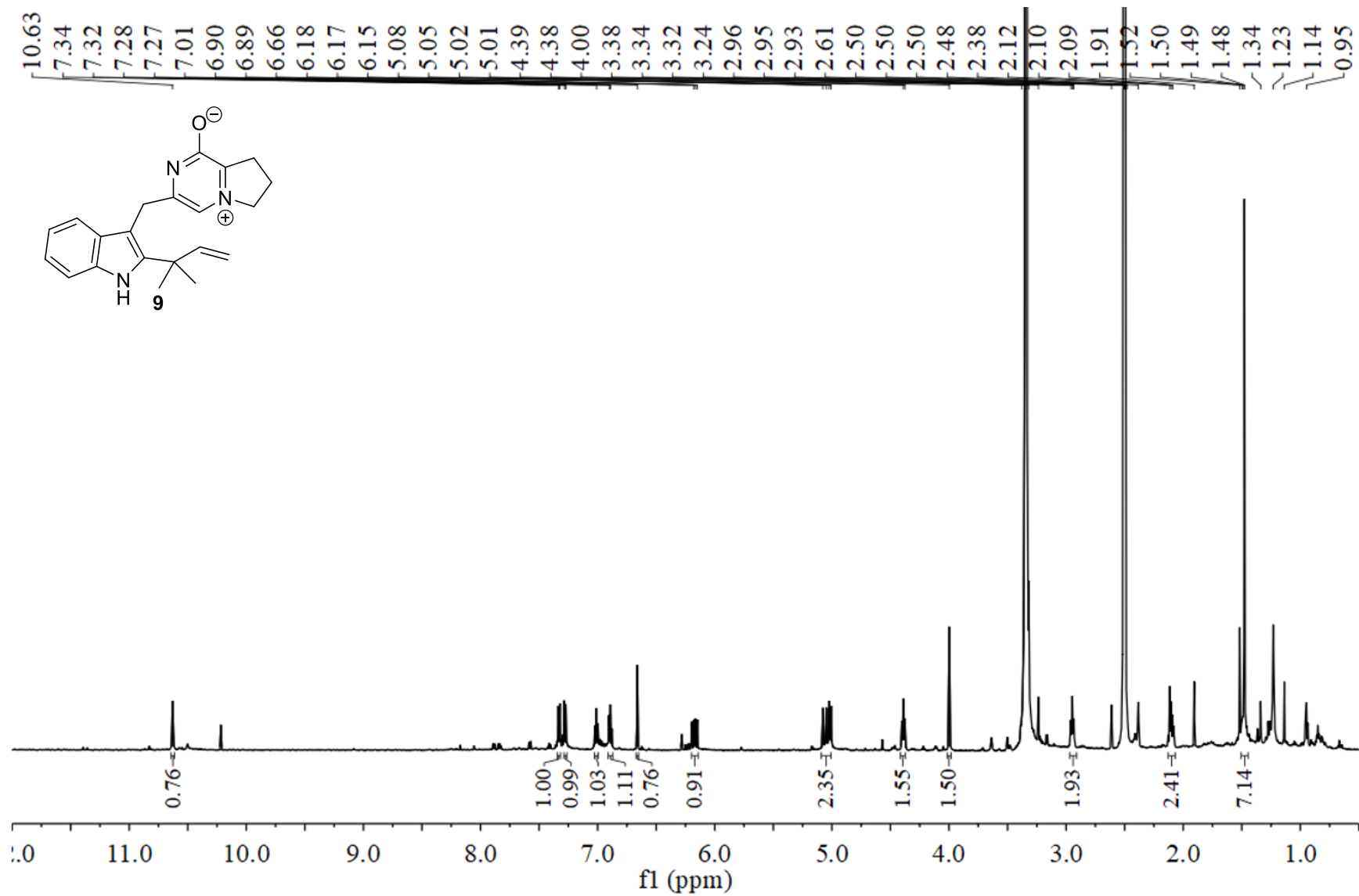
Supplementary Fig. 12-5 | HSQC NMR spectrum of 7 in DMSO-*d*₆.



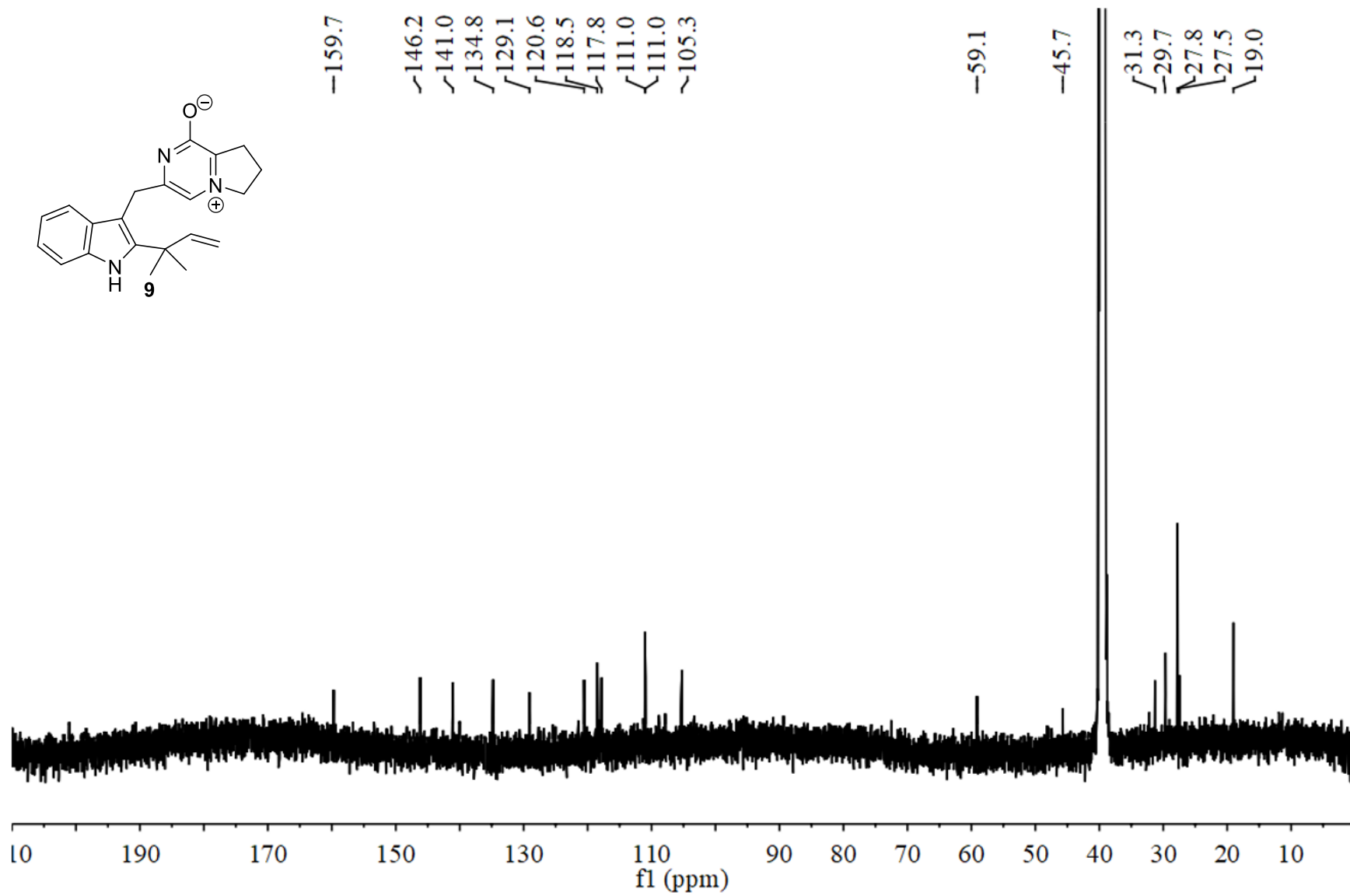
Supplementary Fig. 12-6 | HMBC NMR spectrum of 7 in DMSO-*d*₆.



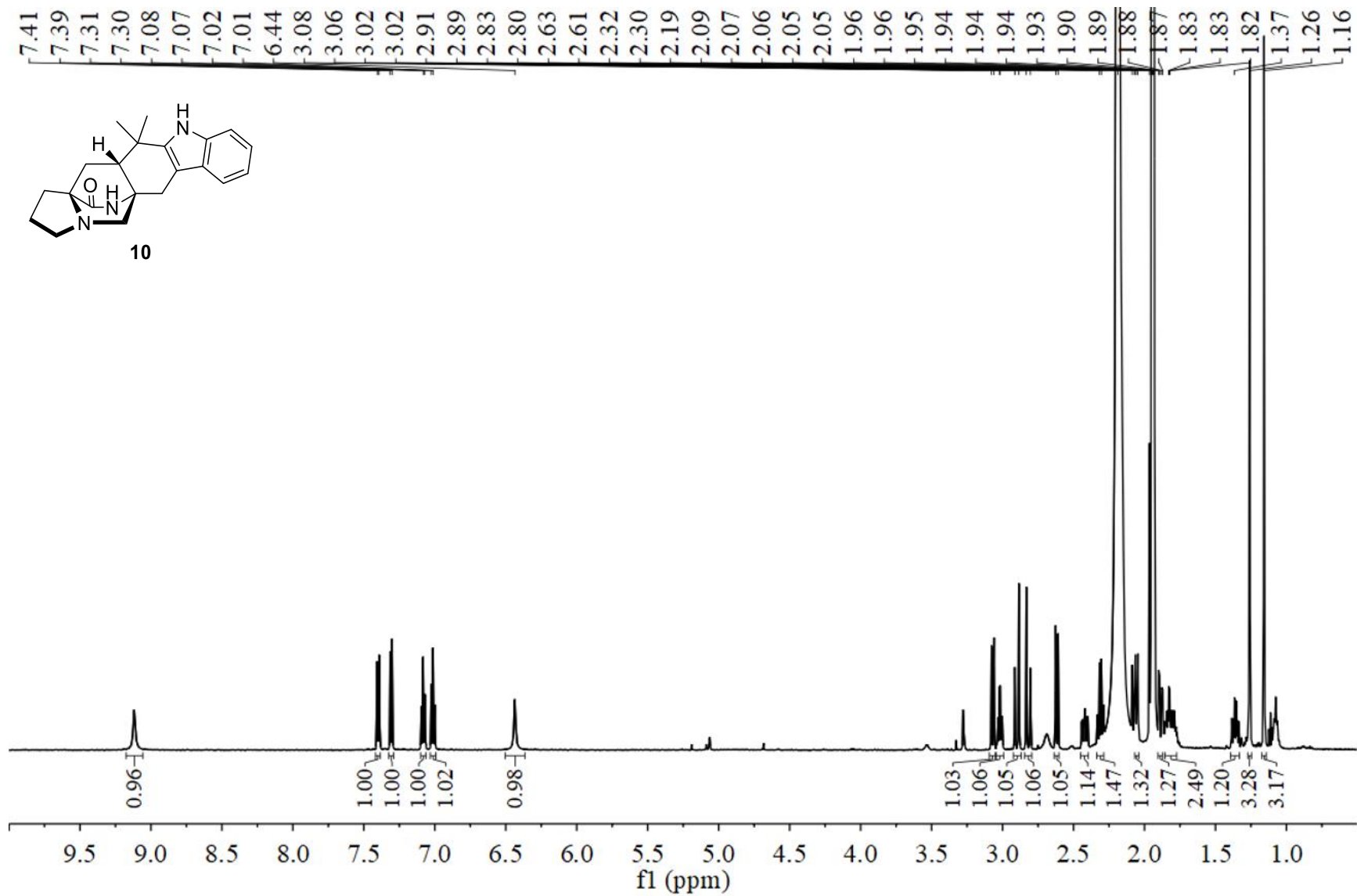
Supplementary Fig. 12-7 | HRMS spectrum of 7.



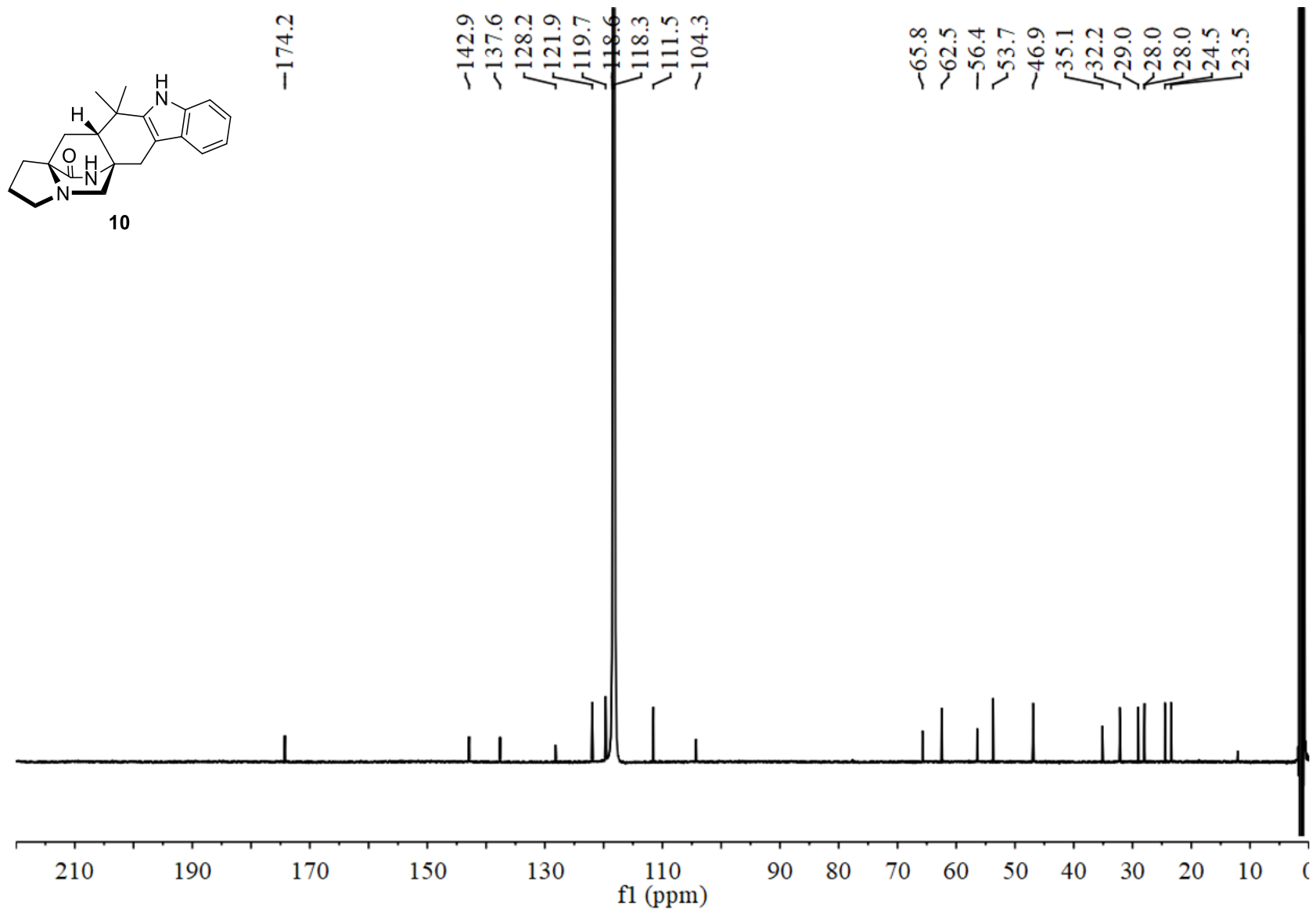
Supplementary Fig. 13-1 | ¹H NMR spectrum of 9 in DMSO-d₆.



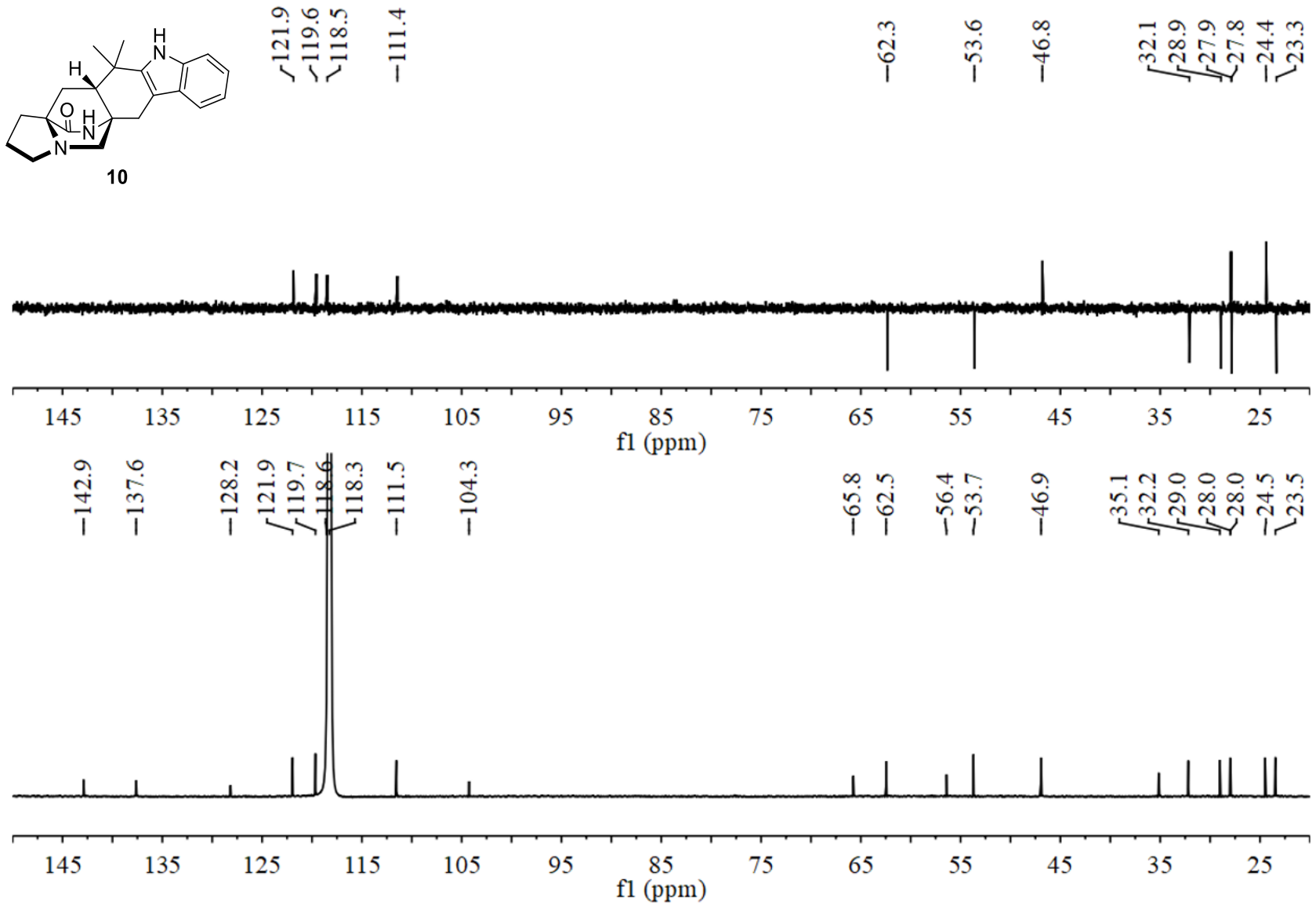
Supplementary Fig. 13-2 | ^{13}C NMR spectrum of 9 in $\text{DMSO-}d_6$.



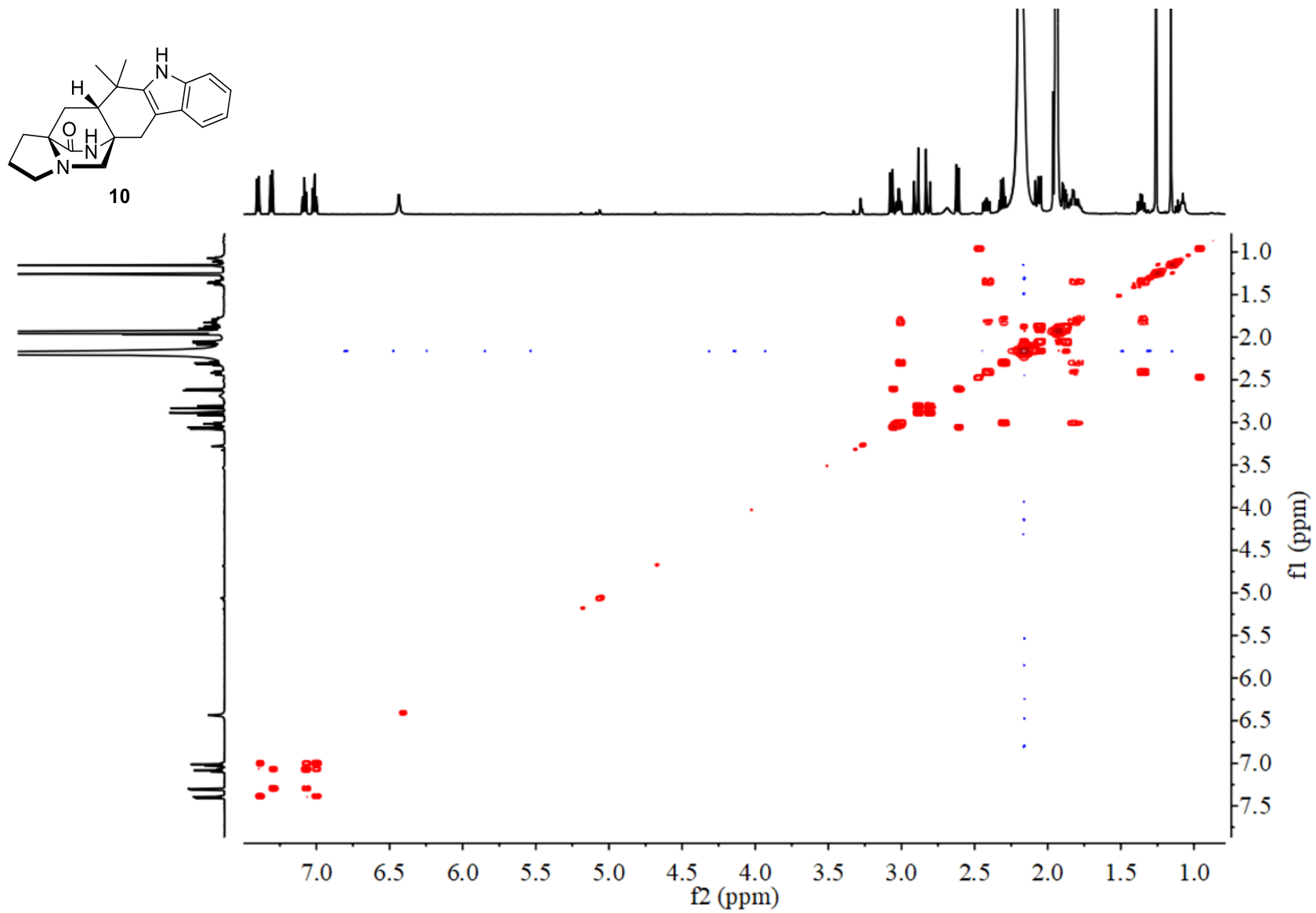
Supplementary Fig. 14-1 | ^1H NMR spectrum of 10 in CD_3CN .



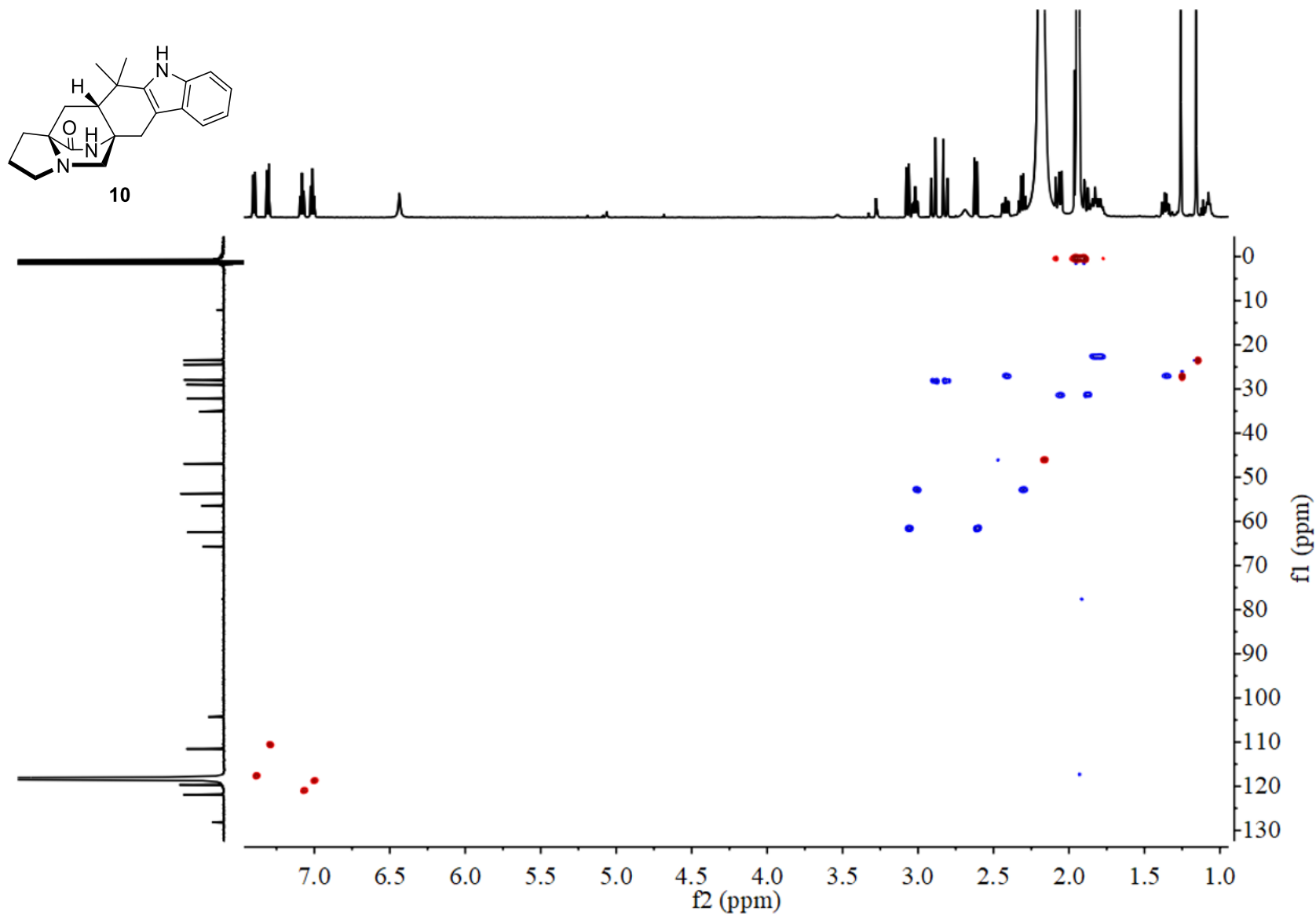
Supplementary Fig. 14-2 | ¹³C NMR spectrum of 10 in CD₃CN.



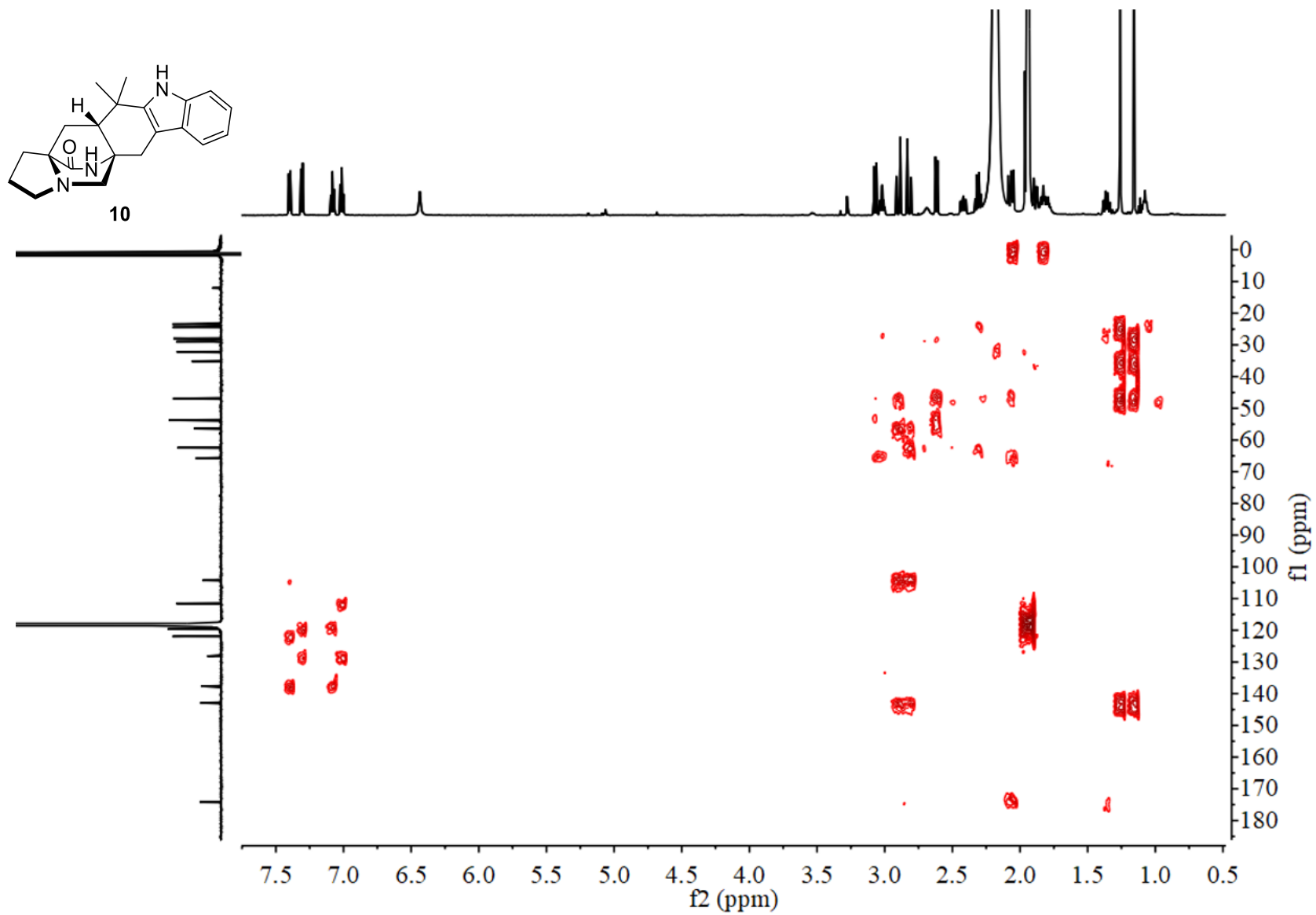
Supplementary Fig. 14-3 | DEPT135 and ¹³C NMR spectra of 10 in CD₃CN.



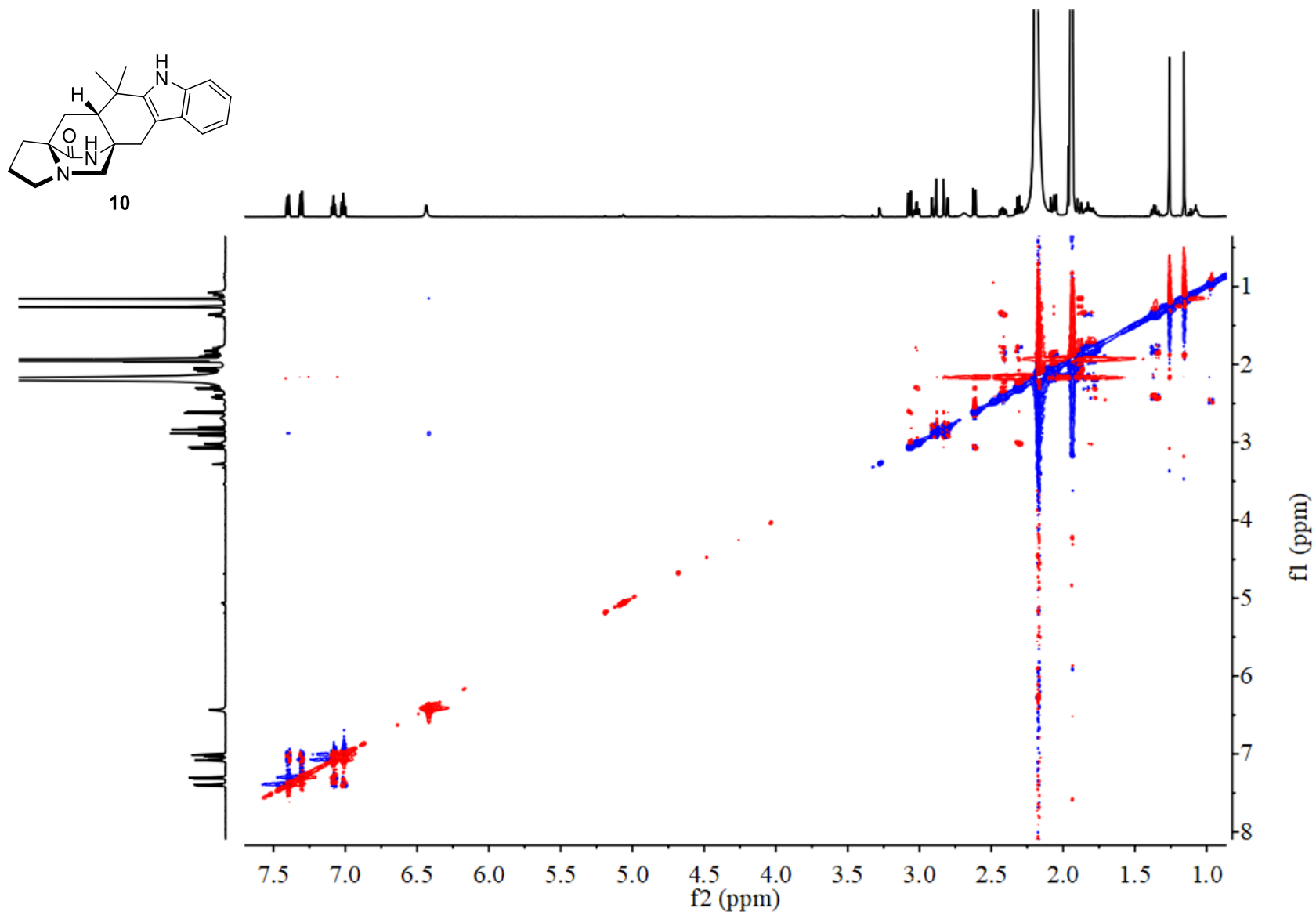
Supplementary Fig. 14-4 | ^1H - ^1H COSY NMR spectrum of 10 in CD_3CN .



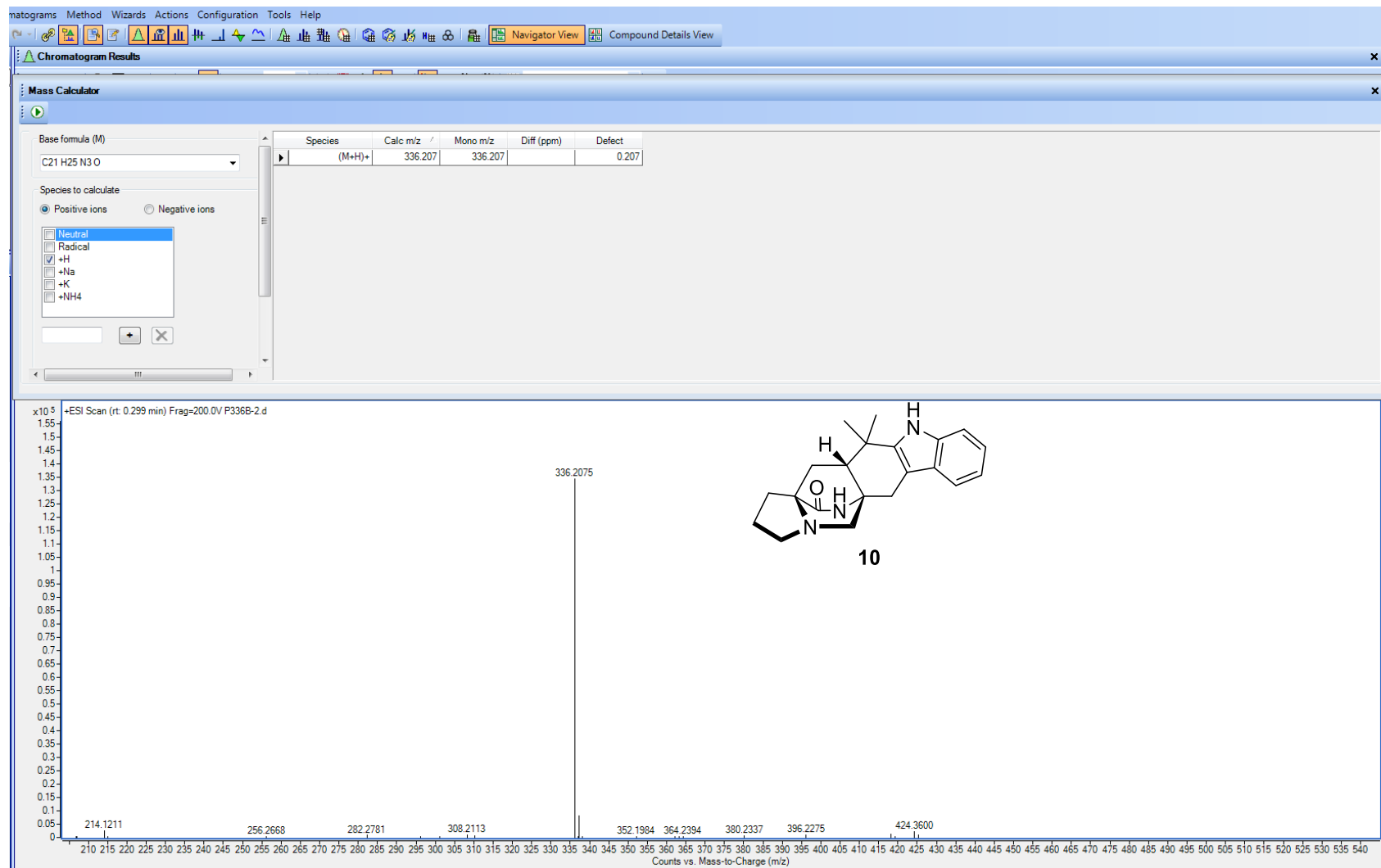
Supplementary Fig. 14-5 | HSQC NMR spectrum of 10 in CD₃CN.



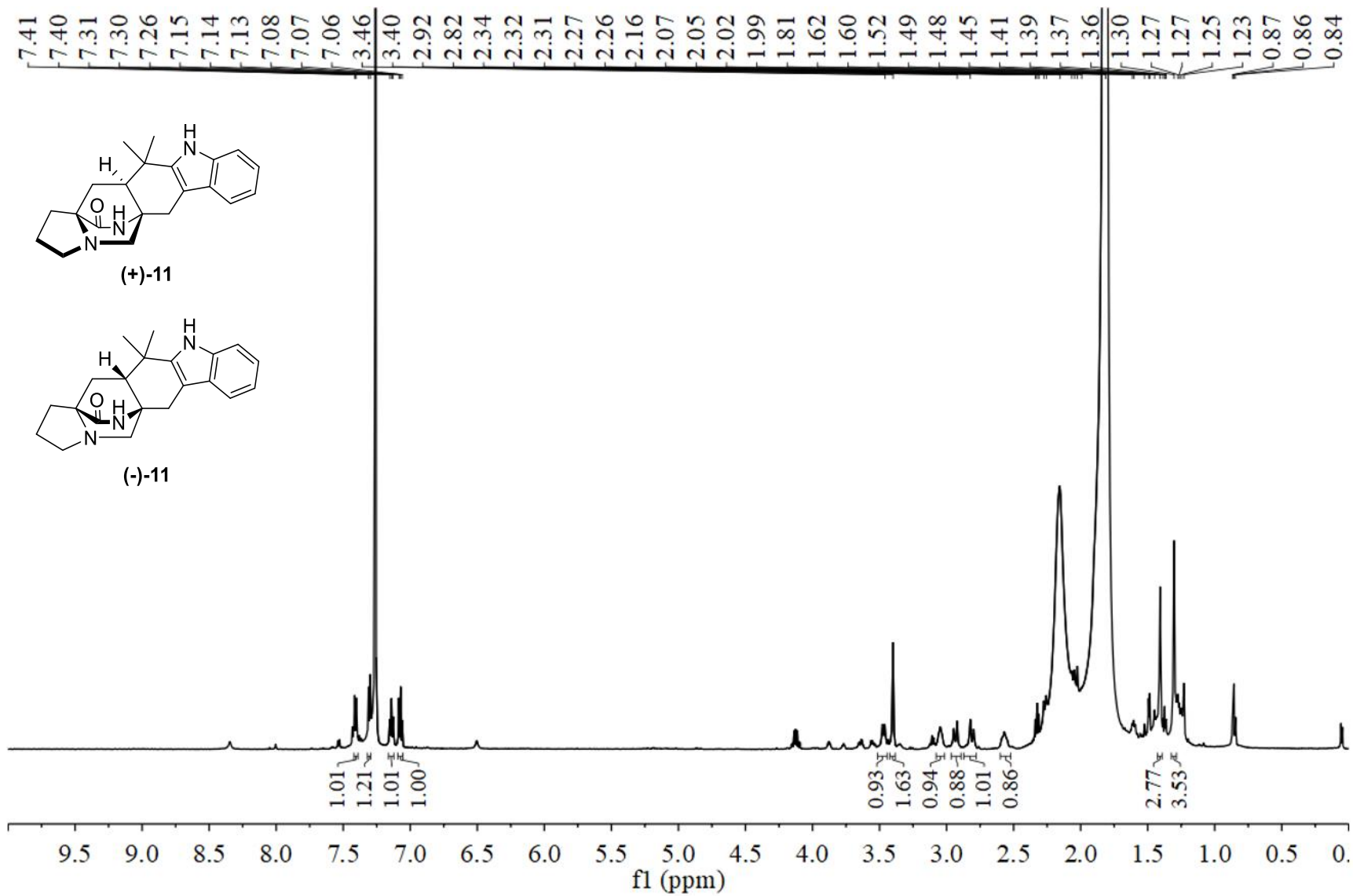
Supplementary Fig. 14-6 | HMBC NMR spectrum of 10 in CD₃CN.



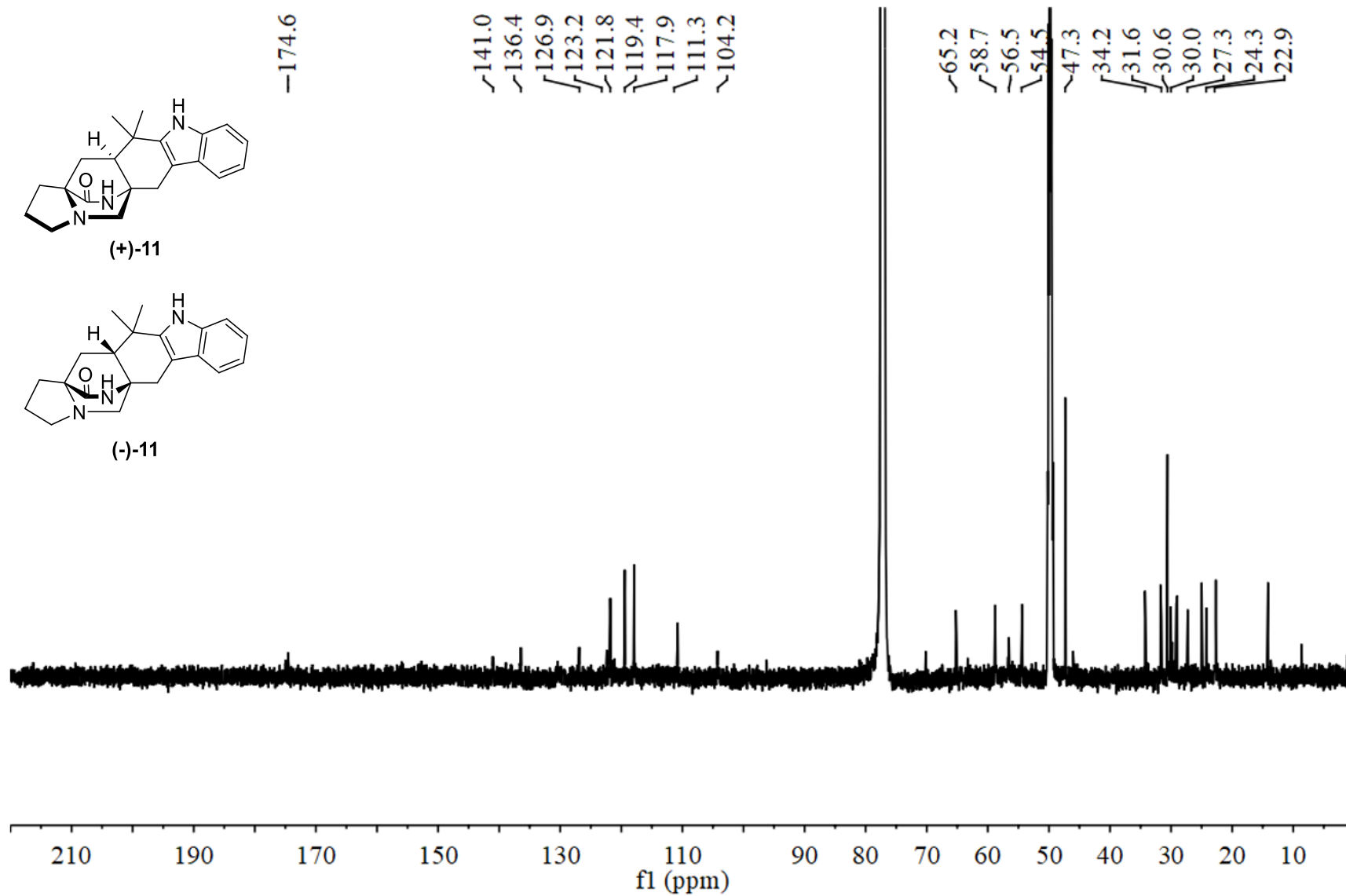
Supplementary Fig. 14-7 | NOESY NMR spectrum of 10 in CD₃CN.



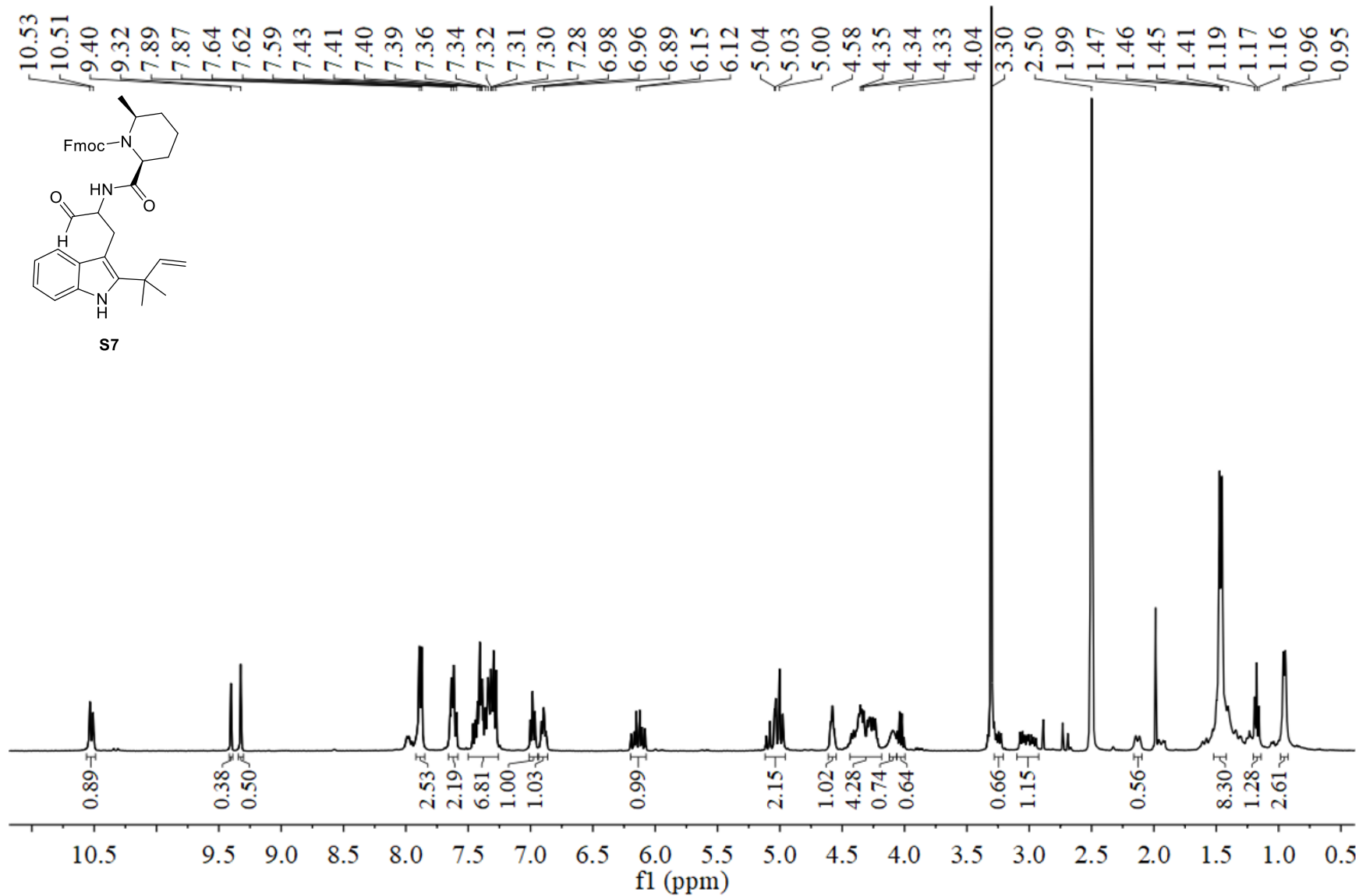
Supplementary Fig. 14-8 | HRMS spectrum of 10.



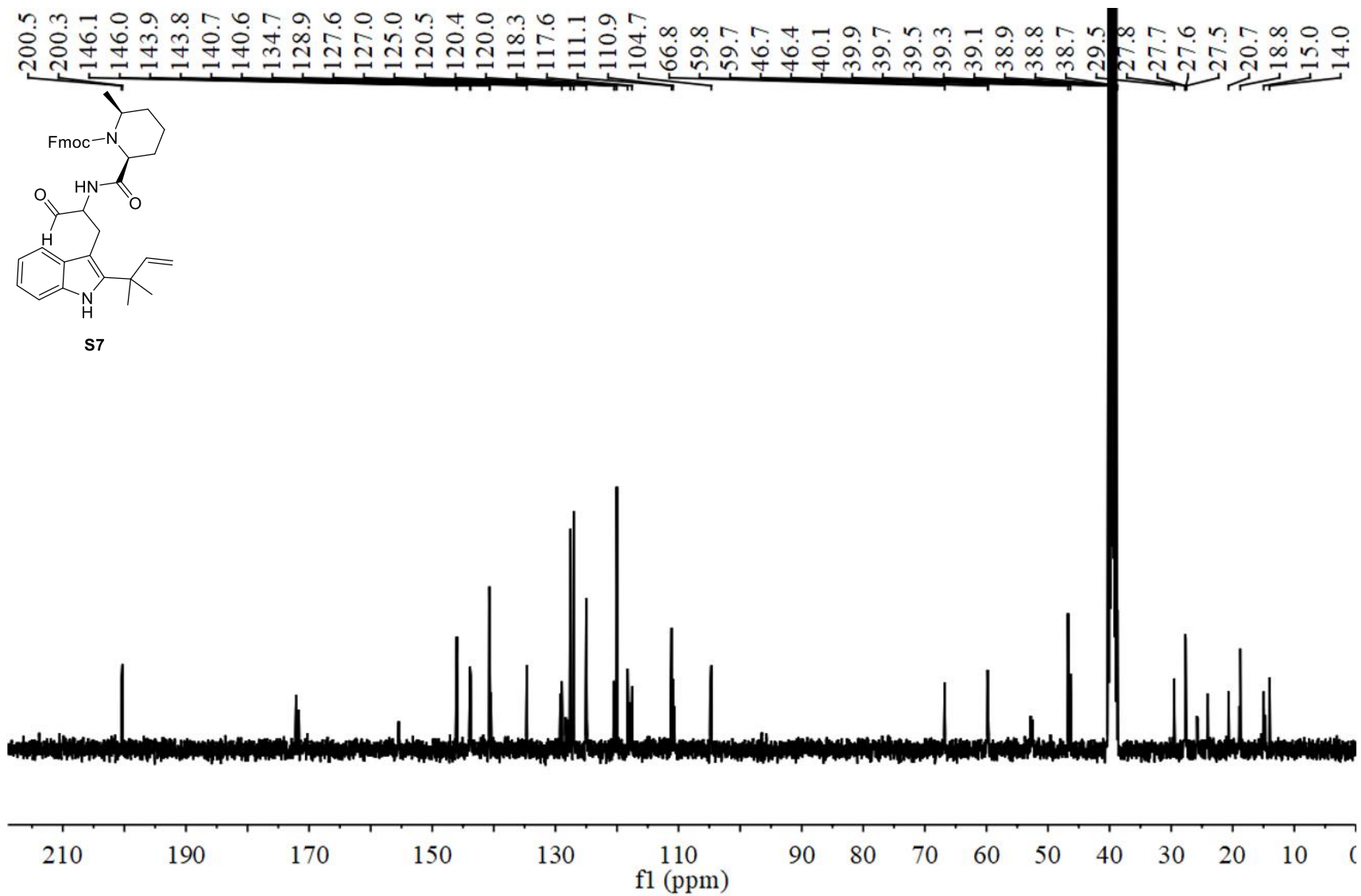
Supplementary Fig. 15-1 | ¹H NMR spectrum of 11 in CD₃CD/CDCl₃ (1/10).



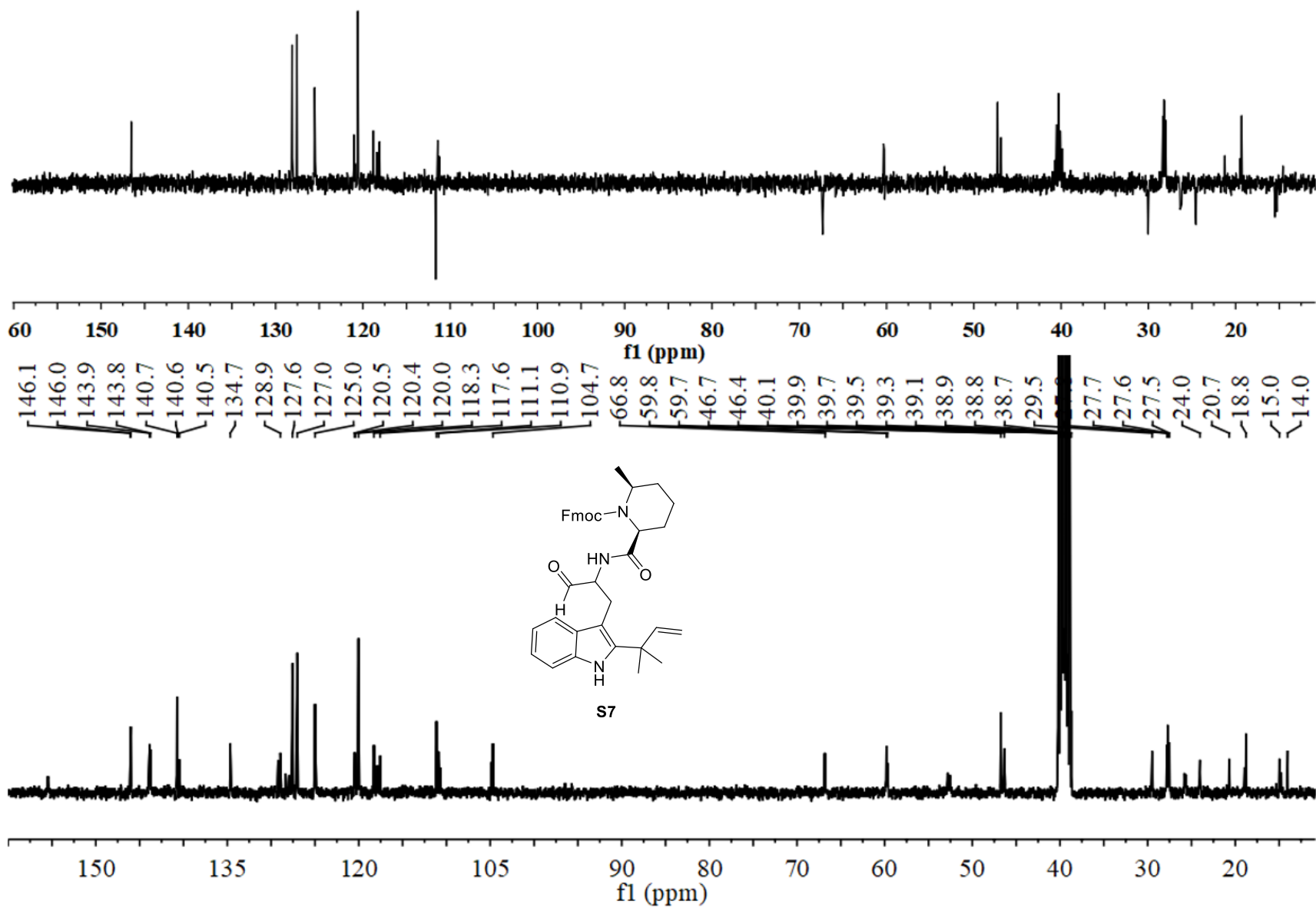
Supplementary Fig. 15-2 | ¹³C NMR spectrum of 11 in CD₃CD/CDCl₃ (1/10)².



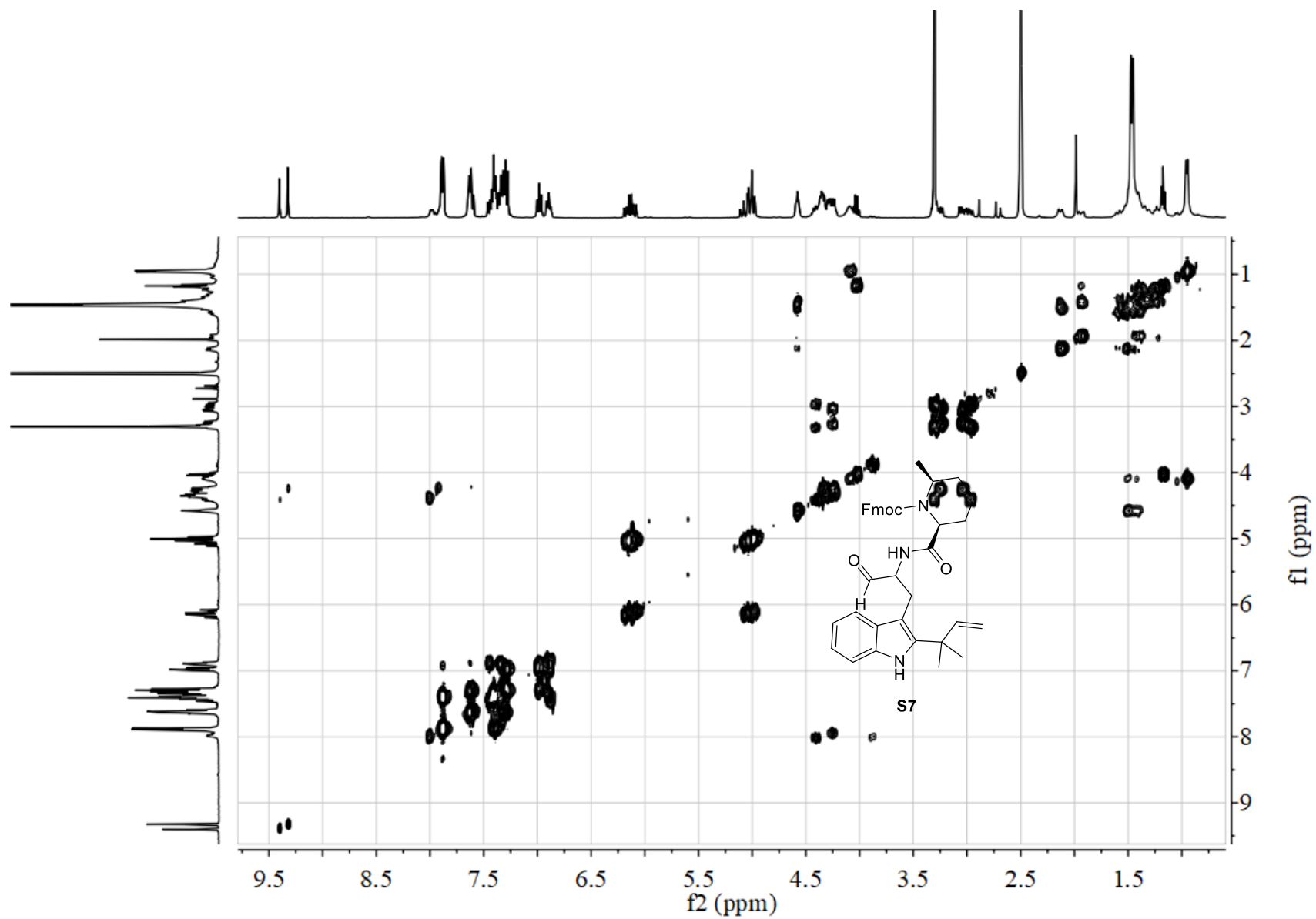
Supplementary Fig. 16-1 | ¹H NMR spectrum of S7 in DMSO-*d*₆.



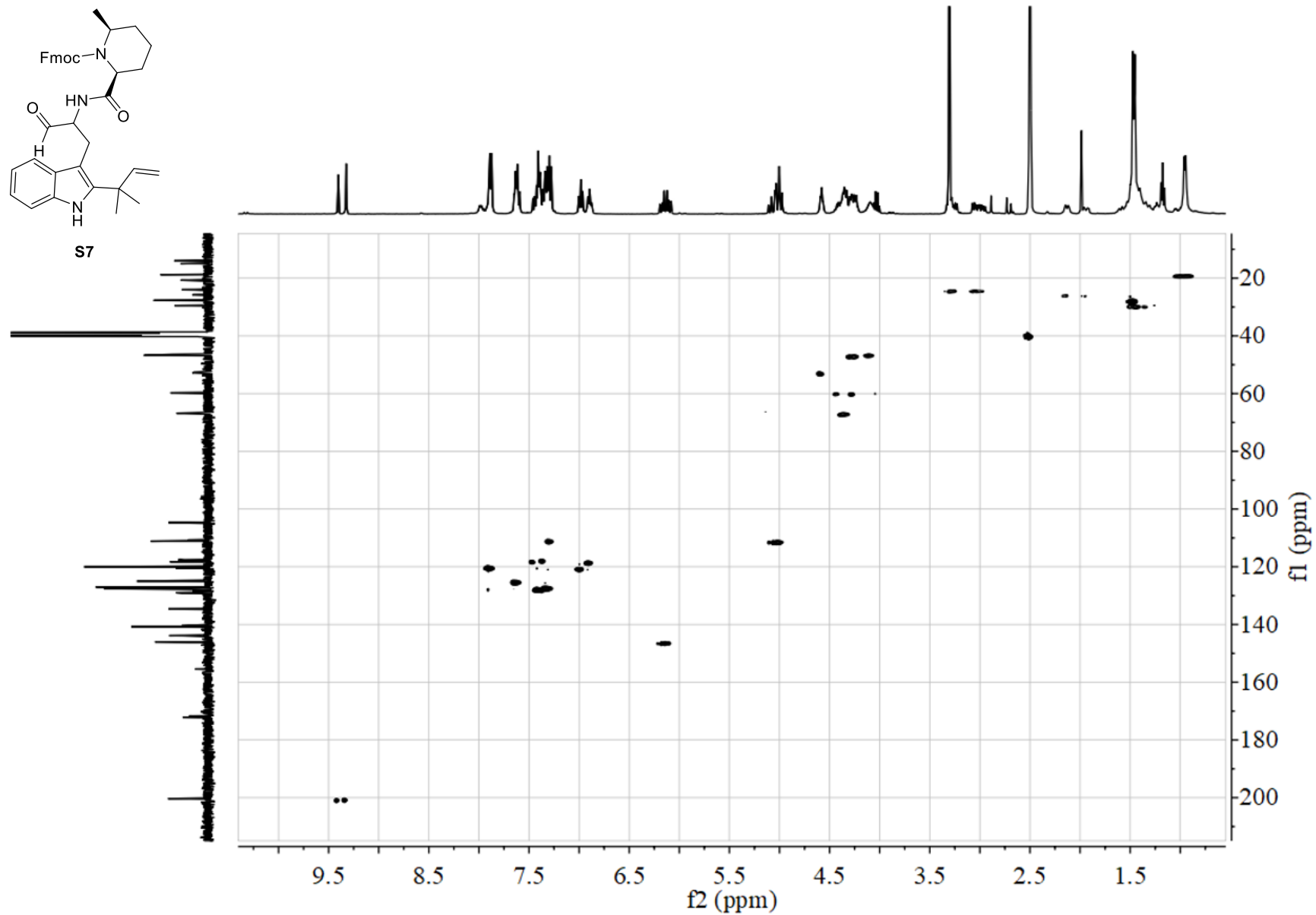
Supplementary Fig. 16-2 | ^{13}C NMR spectrum of S7 in $\text{DMSO-}d_6$.



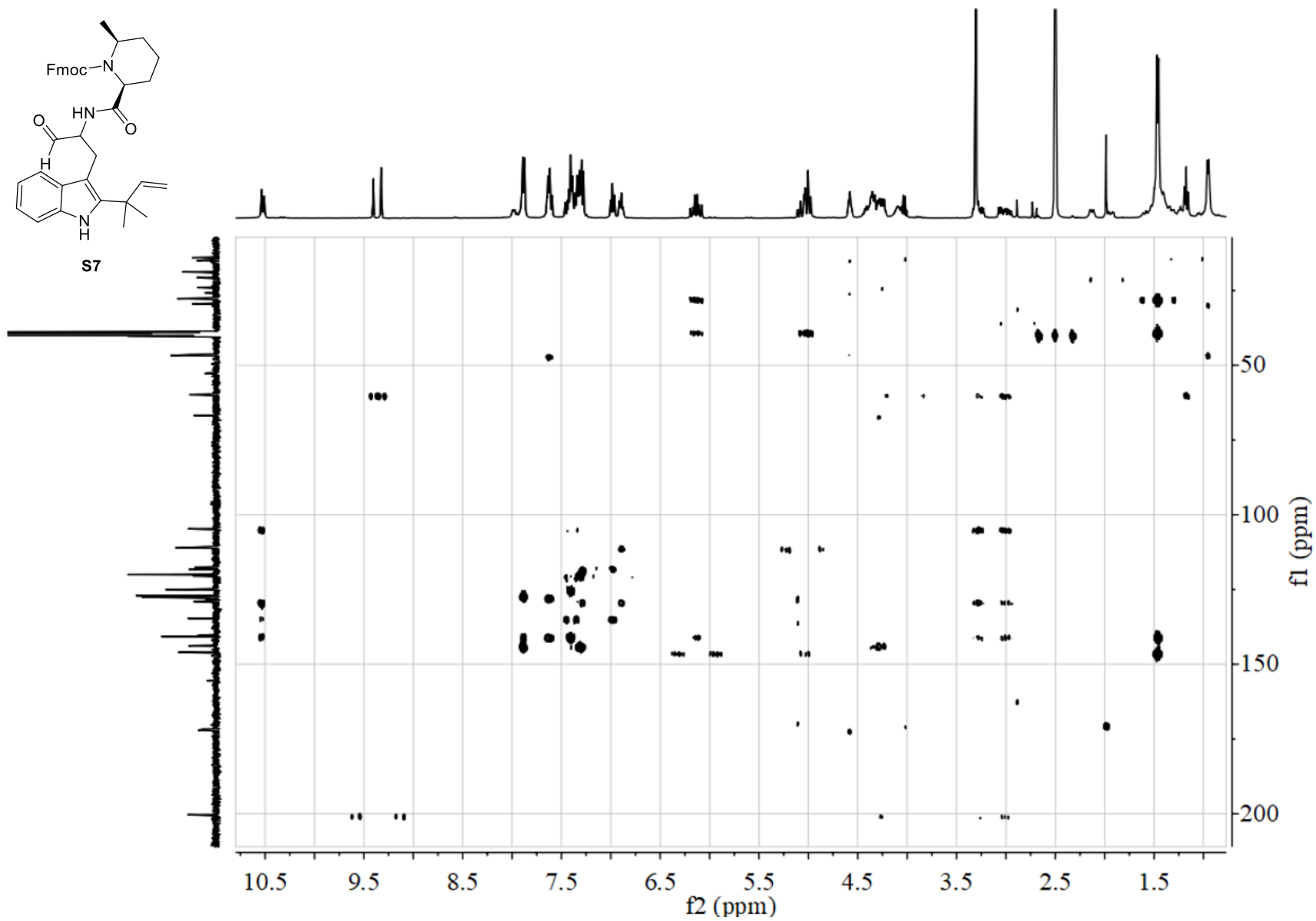
Supplementary Fig. 16-3 | DEPT135 and ¹³C NMR spectra of S7 in DMSO-*d*₆.



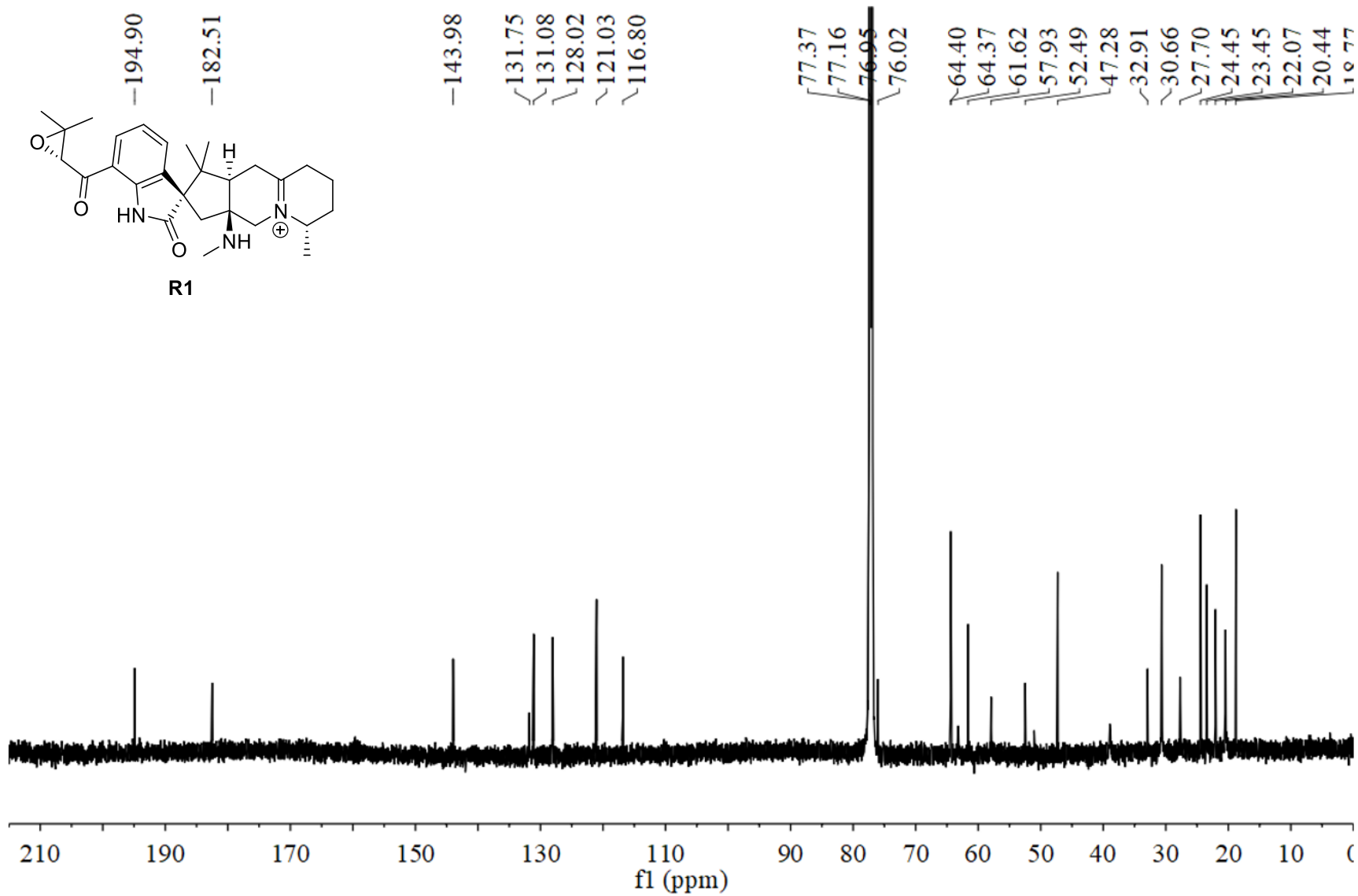
Supplementary Fig. 16-4 | ^1H - ^1H COSY NMR spectrum of S7 in $\text{DMSO-}d_6$.



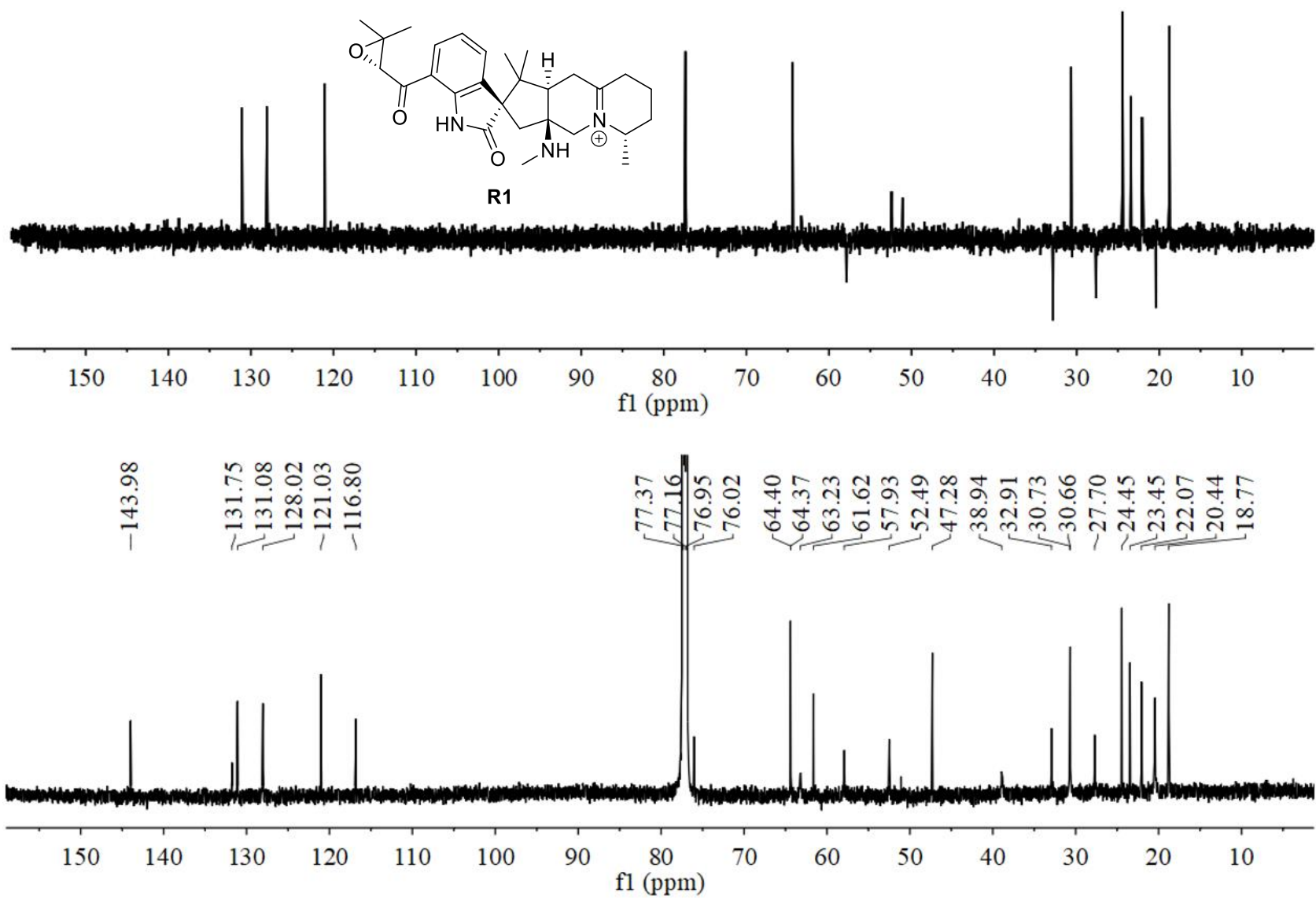
Supplementary Fig. 16-5 | HSQC NMR spectrum of S7 in DMSO-*d*₆.



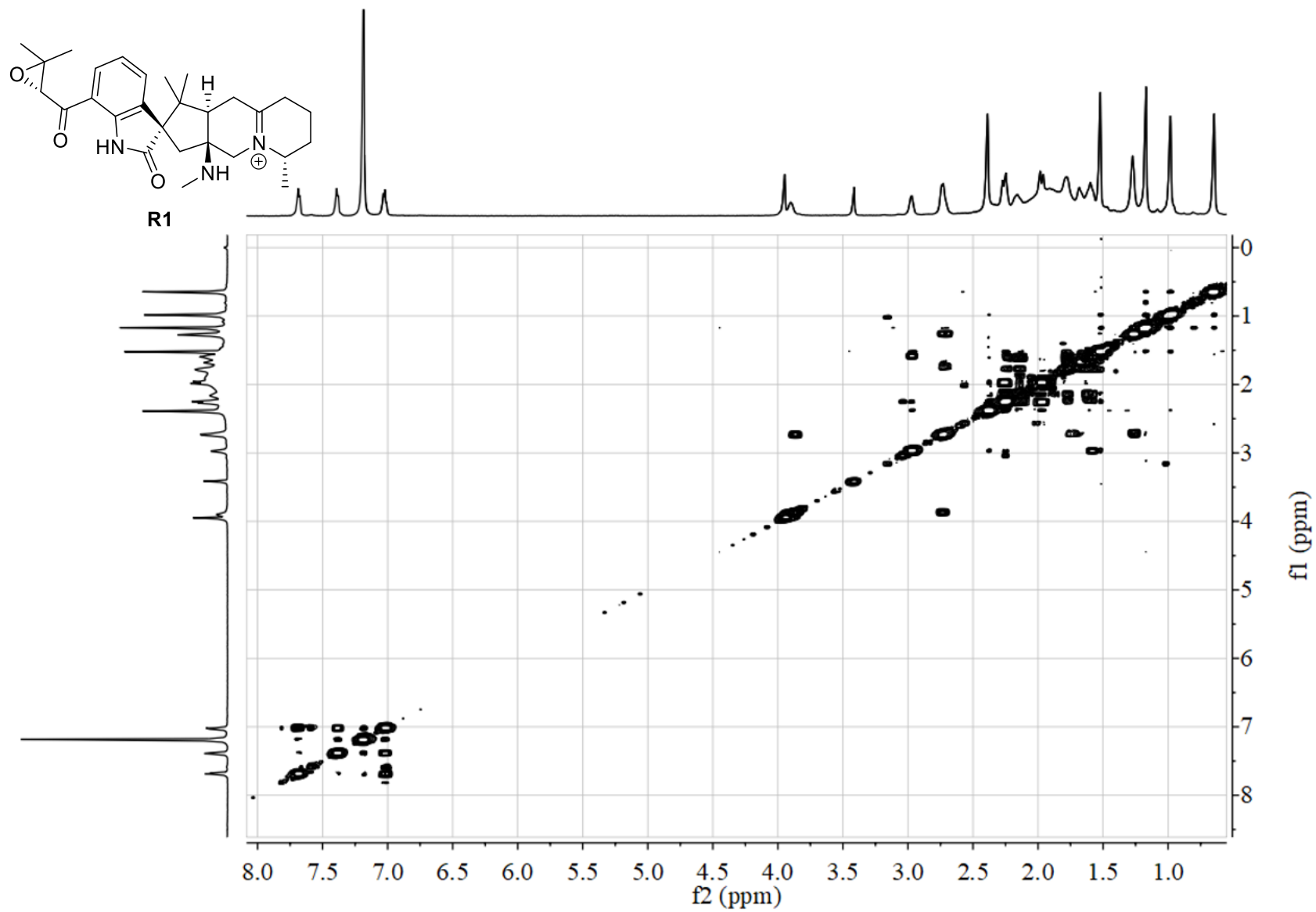
Supplementary Fig. 16-6 | HMBC NMR spectrum of S7 in DMSO-*d*₆.



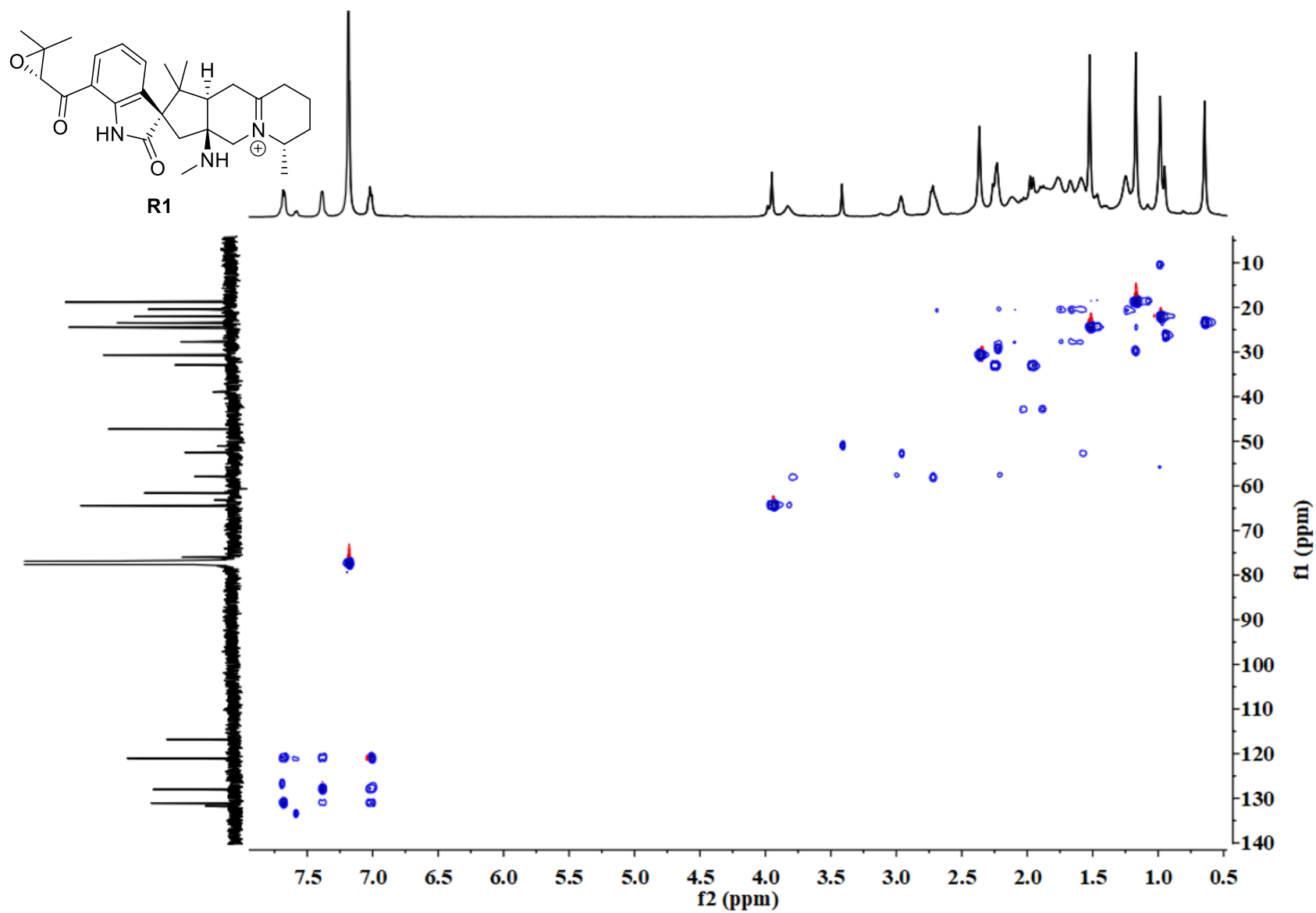
Supplementary Fig. 17-2 | ^{13}C NMR spectrum of R1 in CDCl_3 .



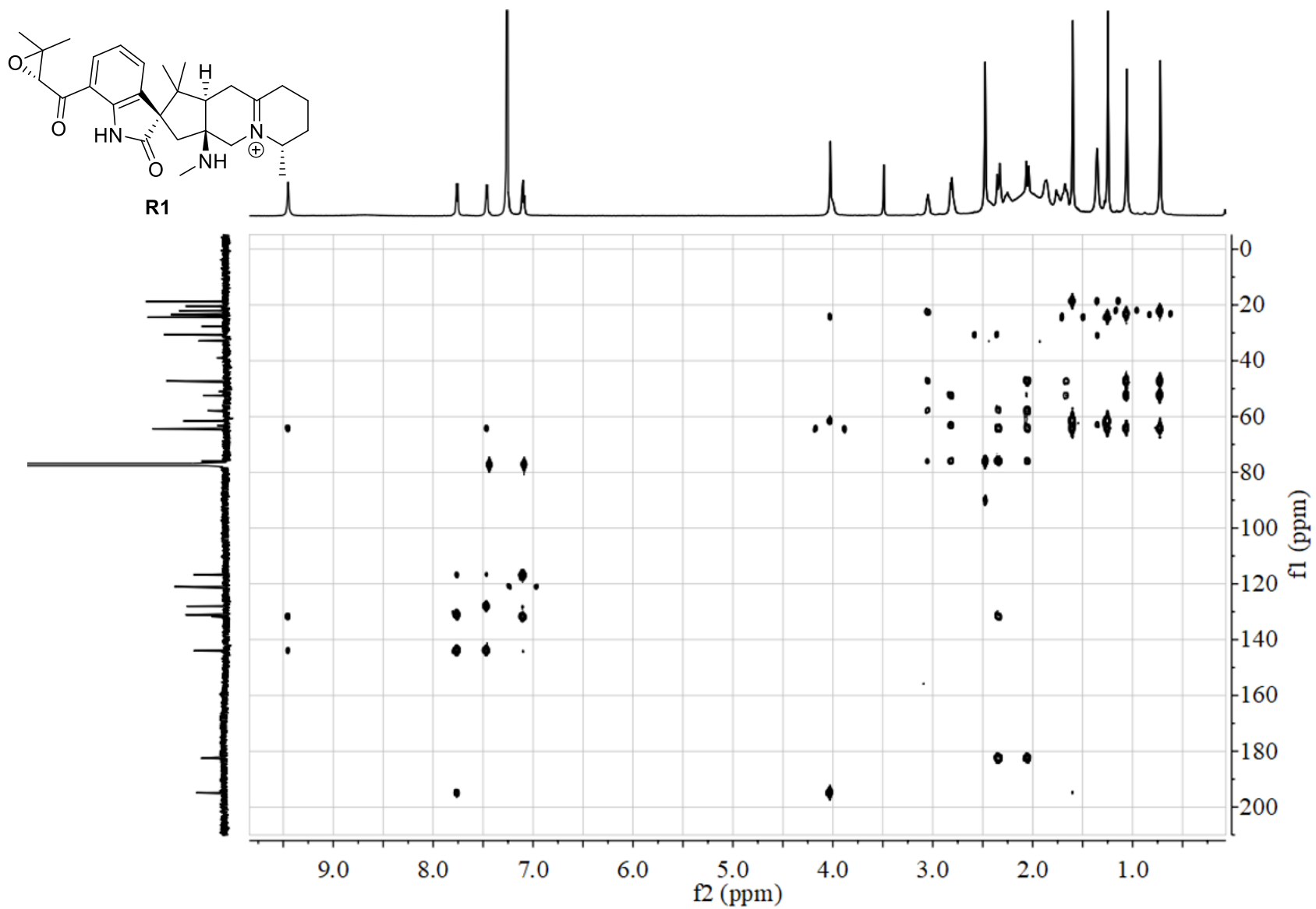
Supplementary Fig. 17-3 | DEPT135 and ^{13}C NMR spectra of R1 in CDCl_3 .



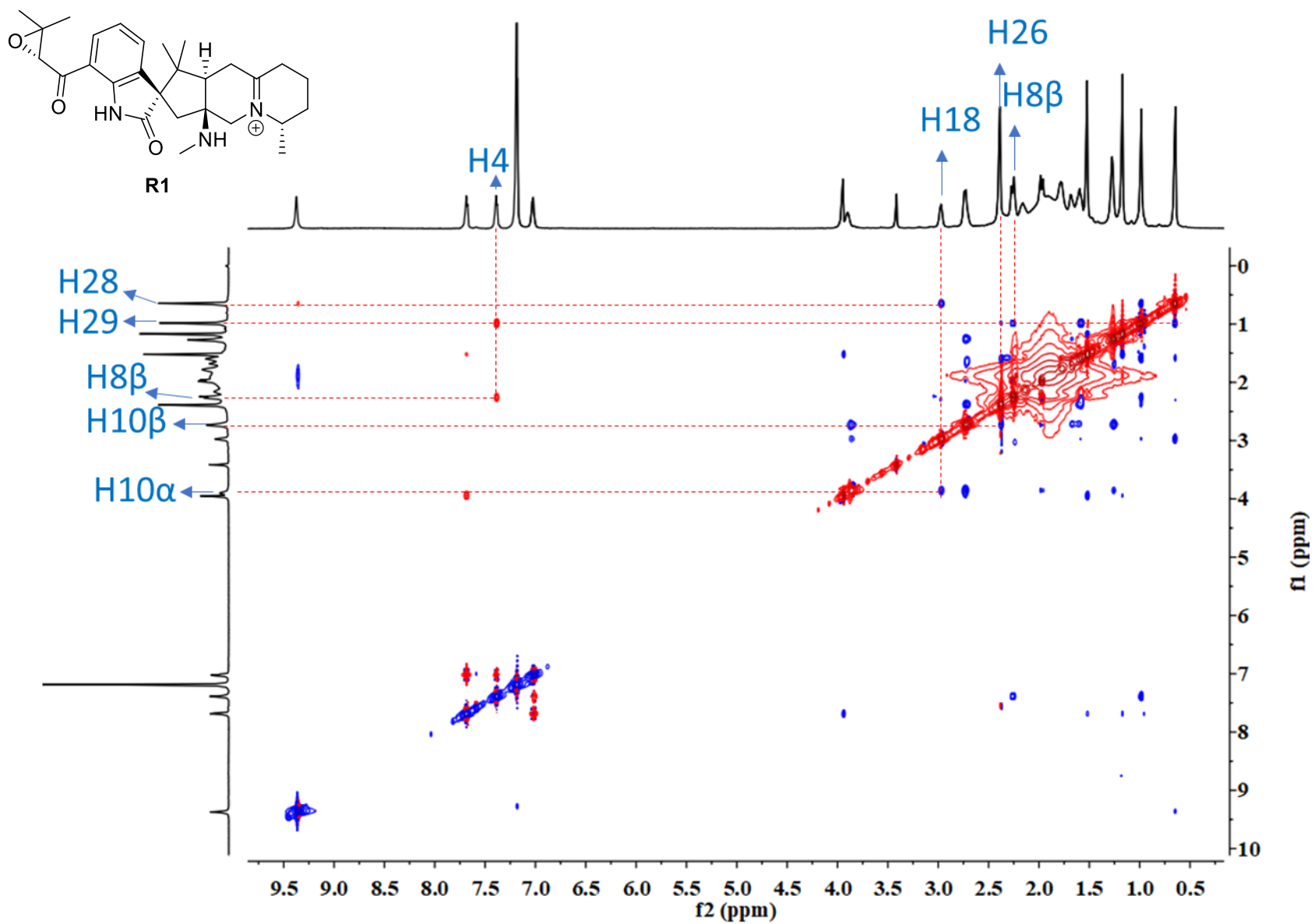
Supplementary Fig. 17-4 | ^1H - ^1H COSY NMR spectrum of R1 in CDCl_3 .



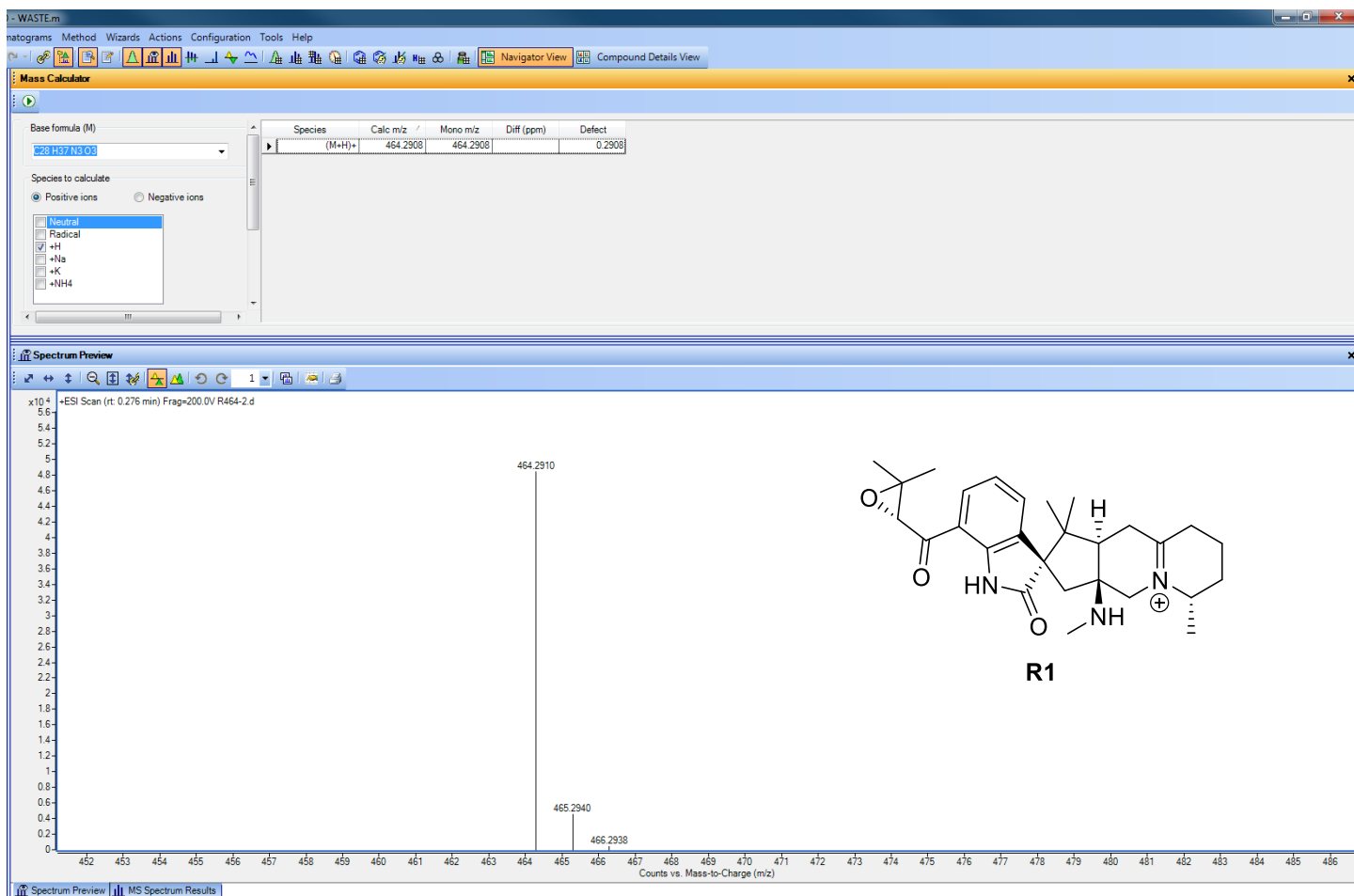
Supplementary Fig. 17-5 | HSQC-TOCSY NMR spectrum of R1 in CDCl₃.



Supplementary Fig. 17-6 | HMBC NMR spectrum of R1 in CDCl₃.



Supplementary Fig. 17-7 | NOESY NMR spectrum of R1 in CDCl₃.



Supplementary Fig. 17-8 | HRMS spectrum of R1 in CDCl₃.

Energies and Molecular Coordinates of Calculated Structures.

Structure	Single-Point Energy (Hartree) M06-2X/ 6-311++G(2d,2p) CPCM water solvent	Enthalpy Correction (Hartree) M06-2X/6-31+G(d) CPCM water solvent	Gibbs Free Energy Correction (Hartree) M06-2X/6-31+G(d) CPCM water solvent
Substrate 3	-1133.27763795	0.512919	0.432077
Substrate 3-taut	-1133.24220759	0.512060	0.431172
Substrate 3-ox	-1132.51988010	0.503054	0.423583
TS-taut _{<i>α-anti</i>}	-1133.20758299	0.510622	0.436393
TS-ox _{<i>α-anti</i>}	-1132.48695913	0.502384	0.429517
Product 2-taut	-1133.30637638	0.515543	0.443782
Product 2-ox	-1132.54357524	0.505760	0.433204
Product 2	-1133.32726952	0.515808	0.443338
Substrate 3-ox _{<i>α-syn</i>}	-1132.51871022	0.503050	0.423882
Substrate 3-ox _{<i>β-syn</i>}	-1132.51660234	0.503183	0.424415
Substrate 3-ox _{<i>β-anti</i>}	-1132.51549387	0.503060	0.425197
TS-ox _{<i>α-syn</i>}	-1132.48734470	0.502322	0.430120
TS-ox _{<i>β-syn</i>}	-1132.49035735	0.502212	0.429479
TS-ox _{<i>β-anti</i>}	-1132.48763403	0.502498	0.429897
NADP ⁺ Model	-456.702066786	0.169217	0.124487
NADPH Model	-457.446769424	0.178720	0.131205
Substrate 4	-1132.06562844	0.488953	0.408388
Substrate 9	-1053.44990317	0.428902	0.353768
Substrate 9-red	-1054.65441754	0.453013	0.376677

Substrate 3

0 1

N	-1.707056	-1.291350	-1.810658
C	-2.999269	-1.405467	-1.416401
O	-3.839078	-2.009447	-2.077905
C	-3.318343	-0.771696	-0.058276
N	3.297103	0.430707	0.880042
C	3.580603	-2.855172	-1.129870
N	-2.525273	0.454711	0.113926
C	3.099662	-3.046384	0.098080
C	-0.727432	-0.631251	-1.038922
C	0.704299	-0.943344	-1.387339
C	1.705779	-0.171221	-0.582238
C	2.002182	1.223259	-0.769252
C	1.505957	2.213515	-1.636554
C	2.020547	3.498543	-1.549285
C	3.023993	3.819523	-0.607850
C	3.529536	2.863045	0.261120
C	3.005978	1.567551	0.166459
C	2.513840	-0.620209	0.441014
C	2.610946	-1.991868	1.077538
C	3.596712	-1.990267	2.262194
C	1.226192	-2.406332	1.619648
C	-1.152630	0.218713	-0.095708
H	3.975810	0.377227	1.626435
H	3.630500	-1.866177	-1.580001
H	3.933645	-3.692115	-1.725305
H	3.067252	-4.060260	0.502322
H	0.855675	-0.720489	-2.455976
H	0.877879	-2.021615	-1.285739
H	0.726951	1.974344	-2.357103
H	1.647177	4.274800	-2.210944
H	3.405786	4.835261	-0.562293
H	4.299621	3.108185	0.986868
H	3.271723	-1.296572	3.046063
H	3.637334	-2.991208	2.702533
H	4.611855	-1.729524	1.944470
H	0.886956	-1.685716	2.371538
H	0.472546	-2.450595	0.828958
H	1.290735	-3.393732	2.090387
C	-4.815556	-0.514956	0.057646
H	-5.132730	0.111887	-0.785724
C	-2.823058	1.212077	1.348728
H	-2.505806	0.607499	2.220363
C	-5.136771	0.183290	1.372528
H	-6.208357	0.398426	1.437630
H	-4.887283	-0.477330	2.214478
C	-4.324833	1.468589	1.469711
H	-4.627446	2.158454	0.669065
H	-4.508513	1.975005	2.423610
H	-5.339672	-1.470179	-0.030184
C	-2.079865	2.547254	1.362549
H	-0.995863	2.445112	1.451799
H	-2.300889	3.104066	0.444981
H	-2.422105	3.139485	2.216701
H	-3.019373	-1.516981	0.710864
H	-1.440270	-1.774992	-2.662344
H	-0.434737	0.751667	0.513516

Substrate 3-taut

0 1

C	0.459260	-0.629075	-0.864877
C	-0.750694	-0.079799	-1.582561
C	-1.945099	0.172469	-0.709773
C	-2.137760	1.183512	0.210396
C	-1.225570	2.311206	0.663786
C	-0.434324	2.826272	-0.522193
C	0.893086	2.879288	-0.628438
C	2.722359	-1.206661	0.578036
C	1.509850	-1.760620	0.809606
N	0.339251	-1.327199	0.203816
C	-3.983600	-0.117287	0.313656
C	-5.227325	-0.685564	0.615272
C	-5.584497	-1.838983	-0.068701
C	-4.725861	-2.414484	-1.031120
C	-3.497431	-1.842564	-1.328103
C	-3.110094	-0.670033	-0.653713
N	-3.365938	0.998406	0.820605
C	1.823217	-0.440748	-1.480362
N	2.810111	-0.219933	-0.429447
C	4.135323	0.097518	-0.995887
C	5.242627	0.009480	0.051612
C	5.219672	-1.326852	0.778389
C	3.883318	-1.455126	1.501070
O	1.330104	-2.661936	1.835327
C	4.140817	1.487913	-1.629471
C	-2.057022	3.501149	1.188558
C	-0.318756	1.805389	1.797661
H	-1.048909	3.209075	-1.339943
H	-5.883678	-0.241411	1.358131
H	-6.540865	-2.308908	0.141348
H	-5.035663	-3.320291	-1.544242
H	-2.844911	-2.297529	-2.069596
H	-3.772420	1.617012	1.508095
H	4.355069	-0.654033	-1.780097
H	-0.450150	0.824807	-2.117451
H	-1.022066	-0.806345	-2.363694
H	1.556035	2.493732	0.141541
H	1.363324	3.298614	-1.514310
H	2.085045	-1.350544	-2.068030
H	1.809670	0.409443	-2.162750
H	6.198802	0.169010	-0.458159
H	5.119119	0.827149	0.776502
H	6.044994	-1.398215	1.493863
H	5.337785	-2.145052	0.055429
H	3.852343	-0.716640	2.317075
H	3.761379	-2.438595	1.961002
H	3.428909	1.592698	-2.451691
H	5.137291	1.695994	-2.031941
H	3.909597	2.243704	-0.871407
H	-2.780965	3.845431	0.442463
H	-1.386314	4.332054	1.426644
H	-2.593341	3.245177	2.109136
H	-0.932856	1.425470	2.620885
H	0.301497	2.622942	2.182051
H	0.334169	0.996615	1.456567
H	0.383324	-2.873196	1.858087

Substrate 3-ox

I 1

C	0.529772	-0.354241	-0.932042
C	-0.720620	0.066585	-1.658313
C	-1.915192	0.183780	-0.753813
C	-2.224274	1.182511	0.151307
C	-1.472454	2.441897	0.543732
C	-0.825718	3.046922	-0.686888
C	0.456524	3.386501	-0.816306
C	2.784237	-1.242056	0.427477
C	1.457417	-1.798626	0.781275
N	0.409535	-1.277339	0.063629
C	-3.848359	-0.376646	0.348926
C	-4.982191	-1.111550	0.715315
C	-5.186105	-2.325456	0.076570
C	-4.286886	-2.797822	-0.904638
C	-3.168860	-2.062401	-1.267030
C	-2.937656	-0.825033	-0.636048
N	-3.387626	0.835152	0.801806
C	1.767577	0.113531	-1.229887
N	2.871842	-0.352578	-0.533567
C	4.190420	0.262471	-0.937380
C	5.342061	-0.647958	-0.533057
C	5.254444	-1.021566	0.941851
C	3.950411	-1.768849	1.195083
O	1.316125	-2.645738	1.649149
C	4.260238	1.669808	-0.349015
C	-2.446701	3.501183	1.101717
C	-0.447188	2.089103	1.634000
H	-1.515506	3.245392	-1.509579
H	-5.669844	-0.745208	1.471820
H	-6.052993	-2.925426	0.336612
H	-4.477382	-3.754492	-1.381737
H	-2.485733	-2.440150	-2.024155
H	-3.854097	1.402385	1.496177
H	4.134737	0.318743	-2.027010
H	-0.504465	0.993761	-2.192264
H	-0.924535	-0.695730	-2.421229
H	1.183831	3.215880	-0.026774
H	0.820920	3.853830	-1.726860
H	5.321426	-1.553470	-1.150408
H	6.273507	-0.121068	-0.756823
H	5.304708	-0.126415	1.569191
H	6.095546	-1.657746	1.225629
H	3.666522	-1.771000	2.252674
H	4.040011	-2.828825	0.916043
H	3.450152	2.299414	-0.725681
H	5.208714	2.123986	-0.646634
H	4.206089	1.650799	0.742524
H	-3.246104	3.726484	0.388452
H	-1.894862	4.423513	1.303493
H	-2.895853	3.178717	2.047569
H	-0.951279	1.614211	2.481382
H	0.053100	2.993286	1.996621
H	0.317571	1.399904	1.260170
H	1.950012	0.842792	-2.006360
H	-0.524825	-1.606832	0.313884

TS-taut_{a-anti}

0 1

C	0.428085	-0.781343	-0.034476
C	-0.880332	-1.446132	-0.390417
C	-2.070004	-0.548121	-0.258990
C	-2.102003	0.814559	-0.426143
C	-1.008545	1.807348	-0.753842
C	0.272615	1.109271	-1.214764
C	1.517475	1.666667	-1.010278
C	2.769787	0.373842	0.462571
C	1.753959	0.324433	1.402995
N	0.568505	-0.259457	1.172656
C	-4.219360	0.192045	0.063938
C	-5.591154	0.145308	0.337831
C	-6.147980	-1.098871	0.602827
C	-5.360883	-2.270438	0.595930
C	-4.001044	-2.217050	0.322547
C	-3.409946	-0.970465	0.051987
N	-3.401684	1.255193	-0.230432
C	1.630910	-1.390039	-0.733500
N	2.886760	-0.670550	-0.505038
C	4.025899	-1.583303	-0.271610
C	5.318414	-0.797683	-0.045785
C	5.177230	0.267310	1.037111
C	4.010063	1.189976	0.670976
O	1.837040	1.111276	2.520426
C	4.196534	-2.523068	-1.462410
C	-1.475695	2.685360	-1.940423
C	-0.775986	2.718069	0.465265
H	0.142388	0.473730	-2.092818
H	-6.195784	1.047711	0.343003
H	-7.209803	-1.171503	0.819457
H	-5.830493	-3.226710	0.807377
H	-3.404175	-3.125969	0.316964
H	-3.708616	2.216371	-0.285990
H	3.816077	-2.185223	0.636014
H	-0.806932	-1.847133	-1.410063
H	-1.000816	-2.319579	0.267002
H	1.646096	2.567414	-0.416459
H	2.338067	1.409087	-1.669779
H	1.702564	-2.412777	-0.315173
H	1.431925	-1.501128	-1.806564
H	6.112004	-1.509838	0.207002
H	5.601799	-0.317098	-0.993076
H	6.104394	0.842016	1.135925
H	4.978596	-0.206674	2.007600
H	4.251464	1.708455	-0.267699
H	3.834636	1.946511	1.439215
H	3.333053	-3.176205	-1.613750
H	5.071833	-3.160837	-1.304091
H	4.352510	-1.938739	-2.376415
H	-1.705287	2.069554	-2.816273
H	-0.681806	3.389316	-2.208086
H	-2.368353	3.268486	-1.687528
H	-1.724975	3.138156	0.813320
H	-0.121554	3.557149	0.210682
H	-0.328601	2.157423	1.290034
H	0.950162	1.135476	2.918431

TS-oxa-anti

I 1

C	0.357901	-1.105925	0.271298
C	-0.993548	-1.642493	-0.075817
C	-2.048015	-0.580765	-0.156731
C	-1.911260	0.768164	-0.391316
C	-0.710998	1.673711	-0.601367
C	0.565103	0.940412	-1.007963
C	1.824753	1.386019	-0.598861
C	2.719403	0.091810	0.716184
C	1.671619	0.204073	1.778020
N	0.520733	-0.465752	1.482620
C	-4.127821	0.382472	-0.149443
C	-5.520740	0.488864	-0.057072
C	-6.231590	-0.677229	0.187575
C	-5.577755	-1.919655	0.336211
C	-4.197345	-2.017264	0.242436
C	-3.453607	-0.849914	-0.002201
N	-3.172034	1.339930	-0.383962
C	1.495842	-1.557268	-0.368928
N	2.695295	-1.051620	-0.044956
C	3.885339	-1.556938	-0.816058
C	5.176797	-1.118722	-0.134128
C	5.171627	0.382369	0.137451
C	4.041519	0.703969	1.105662
O	1.820013	0.885395	2.781651
C	3.812323	-1.130013	-2.282918
C	-0.994185	2.588797	-1.826879
C	-0.535078	2.578760	0.635492
H	0.488266	0.395744	-1.946857
H	-6.022469	1.445041	-0.172521
H	-7.313867	-0.633508	0.265498
H	-6.168707	-2.810780	0.525358
H	-3.703832	-2.979450	0.354172
H	-3.370821	2.322443	-0.511439
H	3.792845	-2.644374	-0.756842
H	-0.912435	-2.196949	-1.017250
H	-1.275358	-2.374986	0.693459
H	1.885748	2.254003	0.052030
H	2.647575	1.290620	-1.300500
H	5.295510	-1.661806	0.810722
H	6.008767	-1.405253	-0.783500
H	5.062961	0.950409	-0.793151
H	6.120167	0.691087	0.583752
H	3.896595	1.780297	1.230842
H	4.282535	0.313834	2.103335
H	2.854740	-1.389687	-2.741612
H	4.597868	-1.660127	-2.827873
H	3.980178	-0.058447	-2.413096
H	-1.240189	1.997491	-2.713472
H	-0.114130	3.201202	-2.041904
H	-1.825076	3.270361	-1.620435
H	-1.498589	3.018948	0.906207
H	0.149201	3.404887	0.421442
H	-0.163746	2.037933	1.510031
H	1.442993	-2.279792	-1.174622
H	-0.286493	-0.319616	2.085234

Product 2-taut

0 1

C	-0.374464	0.577250	-0.155862
C	0.920966	1.395286	-0.191594
C	2.133656	0.520714	-0.137266
C	2.107702	-0.836254	-0.322207
C	0.892276	-1.697449	-0.527430
C	-0.266551	-0.747003	-0.960642
C	-1.663192	-1.403192	-0.935223
C	-2.626848	-0.599823	-0.031644
C	-1.871020	-0.343117	1.260418
N	-0.752409	0.265300	1.229896
C	4.270316	-0.305226	0.006537
C	5.660171	-0.312460	0.168743
C	6.284612	0.906289	0.401847
C	5.546391	2.106825	0.471200
C	4.167596	2.108737	0.307604
C	3.508030	0.890320	0.071922
N	3.393311	-1.338295	-0.232566
C	-1.513482	1.416943	-0.769405
N	-2.808817	0.715065	-0.701517
C	-3.870590	1.568741	-0.138571
C	-5.187696	0.800860	-0.039591
C	-5.039687	-0.486143	0.761669
C	-3.946504	-1.344937	0.133220
O	-2.411123	-0.779470	2.415452
C	-4.069053	2.799402	-1.022430
C	1.122628	-2.702775	-1.669828
C	0.649020	-2.489200	0.776390
H	-0.030577	-0.447969	-1.991338
H	6.228282	-1.236872	0.115043
H	7.362399	0.935076	0.533393
H	6.068603	3.041329	0.654996
H	3.605482	3.037982	0.361897
H	3.655255	-2.309668	-0.329951
H	-3.582489	1.904500	0.879277
H	0.937974	2.001178	-1.109020
H	0.903962	2.102427	0.647329
H	-1.626866	-2.436909	-0.578435
H	-2.109775	-1.420747	-1.934153
H	-1.554156	2.351836	-0.192051
H	-1.267776	1.674498	-1.807228
H	-5.939297	1.457954	0.412747
H	-5.529167	0.566144	-1.058076
H	-5.984594	-1.039922	0.778708
H	-4.781755	-0.250349	1.800717
H	-4.268934	-1.662219	-0.867421
H	-3.764507	-2.252191	0.719435
H	-3.179381	3.431597	-1.080036
H	-4.886867	3.409526	-0.625861
H	-4.332742	2.484054	-2.038465
H	1.410088	-2.191038	-2.594284
H	0.206693	-3.273051	-1.862066
H	1.906420	-3.425236	-1.417385
H	1.509050	-3.141431	0.965662
H	-0.232714	-3.134398	0.706141
H	0.529815	-1.823770	1.633242
H	-1.793686	-0.553793	3.135725

Product 2-ox

1 1

C	0.349574	-0.690078	0.206323
C	-0.959165	-1.471386	0.121065
C	-2.117943	-0.543788	-0.045596
C	-2.001445	0.767573	-0.422966
C	-0.736393	1.550457	-0.649951
C	0.435867	0.525136	-0.795011
C	1.838929	1.156615	-0.633036
C	2.659290	0.413548	0.461708
C	1.791658	0.446939	1.747732
N	0.618287	-0.185526	1.547948
C	-4.207147	0.373311	-0.220199
C	-5.603538	0.460977	-0.192026
C	-6.308103	-0.683980	0.153931
C	-5.643930	-1.889950	0.464397
C	-4.259209	-1.972199	0.434089
C	-3.520677	-0.827533	0.088949
N	-3.259510	1.324602	-0.525177
C	1.533426	-1.549557	-0.143785
N	2.688781	-1.014373	-0.013171
C	3.916942	-1.723372	-0.529475
C	5.185722	-1.057167	-0.008854
C	5.131083	0.457733	-0.179627
C	4.038594	1.008711	0.728655
O	2.129390	1.023053	2.768264
C	3.847950	-1.754467	-2.057067
C	-0.813370	2.332031	-1.974978
C	-0.587411	2.571005	0.501029
H	0.344622	0.093376	-1.798851
H	-6.114590	1.388843	-0.431764
H	-7.393130	-0.650764	0.185743
H	-6.229552	-2.765146	0.729678
H	-3.754566	-2.904961	0.672870
H	-3.463839	2.279400	-0.787766
H	3.836533	-2.739181	-0.135404
H	-0.894341	-2.156908	-0.733643
H	-1.061236	-2.100439	1.014110
H	1.784618	2.203458	-0.331297
H	2.405329	1.121024	-1.568032
H	5.324930	-1.299098	1.050872
H	6.026446	-1.494398	-0.554725
H	4.961389	0.734469	-1.226566
H	6.087071	0.900598	0.111366
H	3.946965	2.094911	0.636423
H	4.304120	0.802353	1.769779
H	2.942237	-2.256167	-2.406938
H	4.711140	-2.310562	-2.431272
H	3.874153	-0.748764	-2.484688
H	-1.053438	1.669789	-2.812360
H	0.144883	2.821505	-2.181180
H	-1.574593	3.117305	-1.924068
H	-1.500458	3.171707	0.558421
H	0.235882	3.269321	0.326232
H	-0.447437	2.093498	1.472714
H	1.444656	-2.556672	-0.547721
H	-0.080039	-0.239207	2.282925

Product 2

0 1

C	-0.381757	0.587523	-0.269892
C	0.910257	1.402316	-0.332063
C	2.114402	0.524834	-0.192899
C	2.082911	-0.839536	-0.318982
C	0.872707	-1.705778	-0.540796
C	-0.282576	-0.768160	-1.013723
C	-1.687420	-1.411144	-0.948900
C	-2.630549	-0.608651	-0.022575
C	-1.890201	-0.406406	1.314777
N	-0.768953	0.324822	1.118583
C	4.236272	-0.305166	0.062241
C	5.619507	-0.314536	0.273998
C	6.244631	0.909282	0.474342
C	5.514231	2.116669	0.463631
C	4.142337	2.120652	0.252047
C	3.482532	0.896623	0.048450
N	3.360322	-1.342775	-0.162523
C	-1.542421	1.392832	-0.874258
N	-2.833959	0.705137	-0.680533
C	-3.851748	1.571450	-0.059239
C	-5.169891	0.814156	0.103517
C	-4.996974	-0.492171	0.867848
C	-3.946351	-1.350447	0.170217
O	-2.242638	-0.846482	2.409038
C	-4.092834	2.805957	-0.927878
C	1.133654	-2.721771	-1.667879
C	0.600818	-2.495366	0.759687
H	-0.051847	-0.515636	-2.057420
H	-0.116745	0.458441	1.885235
H	6.181462	-1.244091	0.281864
H	7.317309	0.937251	0.642274
H	6.037437	3.054990	0.623122
H	3.586431	3.055028	0.244148
H	3.619403	-2.318856	-0.208133
H	-3.507646	1.901476	0.942872
H	0.934495	1.936877	-1.291663
H	0.883494	2.176181	0.446674
H	-1.648791	-2.447837	-0.602570
H	-2.157665	-1.420540	-1.937256
H	-1.545491	2.363562	-0.362749
H	-1.352468	1.572708	-1.940125
H	-5.886314	1.470845	0.610378
H	-5.570874	0.605167	-0.898827
H	-5.948193	-1.032824	0.922302
H	-4.678910	-0.284849	1.896232
H	-4.311860	-1.643894	-0.822771
H	-3.747137	-2.267349	0.734753
H	-3.212368	3.446172	-1.022205
H	-4.895793	3.408855	-0.491937
H	-4.400442	2.494227	-1.932604
H	1.454075	-2.218635	-2.585901
H	0.219954	-3.286787	-1.884692
H	1.904066	-3.446654	-1.383908
H	1.473029	-3.116935	0.989698
H	-0.252949	-3.171816	0.655556
H	0.424357	-1.842479	1.616631

Substrate **3-oxa-syn**

1 1

C	-0.614527	0.237283	-1.304445
C	-3.217008	-0.253574	-0.458409
C	-2.978838	0.726914	-1.541163
N	-1.658551	0.891558	-1.885780
C	-0.886961	-0.659306	-0.322073
N	-2.195363	-0.890097	0.065747
C	-2.365886	-1.912736	1.164263
C	-3.784975	-2.464666	1.153509
C	-4.811423	-1.338659	1.181907
C	-4.638575	-0.471343	-0.059568
O	-3.887908	1.336699	-2.082350
C	-1.946652	-1.271985	2.485191
H	-1.663743	-2.706312	0.897808
H	-3.928007	-3.078388	0.256519
H	-3.889639	-3.122845	2.020187
H	-4.693596	-0.733361	2.085951
H	-5.824906	-1.745209	1.199533
H	-5.097748	0.516616	0.051144
H	-5.131705	-0.925050	-0.931481
H	-0.896494	-0.968933	2.467695
H	-2.073013	-2.008070	3.283109
H	-2.557267	-0.395094	2.715979
H	-0.114559	-1.220924	0.184957
H	-1.474402	1.569002	-2.624861
N	3.190737	0.394549	0.996384
C	-0.324337	3.525240	-0.284787
C	0.944816	3.127335	-0.196417
C	0.774139	0.521685	-1.813852
C	1.846740	0.279698	-0.791438
C	2.708105	-0.874237	-0.787397
C	2.848538	-1.990928	-1.631914
C	3.807878	-2.944122	-1.325844
C	4.636075	-2.808186	-0.190005
C	4.516985	-1.719957	0.661481
C	3.542452	-0.764526	0.348958
C	2.168483	1.031204	0.323811
C	1.552212	2.289958	0.910573
C	0.516726	1.893547	1.975101
C	2.642873	3.159973	1.572983
H	3.634270	0.737199	1.837530
H	-1.079791	3.236798	0.441791
H	-0.651531	4.159110	-1.104443
H	1.656447	3.451364	-0.958829
H	0.795104	1.545446	-2.196234
H	0.942887	-0.131022	-2.680030
H	2.216281	-2.110437	-2.508389
H	3.926633	-3.811872	-1.967843
H	5.378756	-3.571624	0.021646
H	5.149919	-1.613232	1.537406
H	0.981737	1.259743	2.736813
H	0.117038	2.785945	2.468625
H	-0.318405	1.340682	1.533291
H	3.075102	2.671355	2.452745
H	3.448454	3.392213	0.869170
H	2.196646	4.099566	1.911761

Substrate **3-ox β -syn**

I 1

C	-3.020343	0.515190	1.665661
C	1.818988	0.333673	0.781782
C	2.145266	1.121119	-0.306943
C	2.698727	-0.807066	0.757123
C	-0.645704	0.219419	1.293792
C	3.548691	-0.651990	-0.361875
C	-0.898623	-0.510131	0.178540
C	-0.597328	3.328300	0.155476
C	-2.345669	-1.635659	-1.449330
C	1.526904	2.384134	-0.882527
C	0.715244	3.097322	0.179181
C	0.737114	0.524532	1.804476
C	-4.710200	-0.749044	-1.534289
C	-1.551277	-2.924670	-1.249958
C	-3.805439	-1.950690	-1.751510
C	-4.643802	-0.365016	-0.063899
C	0.678041	2.010241	-2.108760
O	-3.946928	0.956714	2.327197
N	-1.703649	0.685861	2.014589
N	-2.203585	-0.756096	-0.219087
N	3.189932	0.520932	-0.979738
C	4.540901	-1.584478	-0.687345
C	-3.236308	-0.250693	0.416143
C	3.819284	-2.876891	1.251563
C	2.842126	-1.945990	1.570545
C	2.638781	3.362910	-1.320114
C	4.662272	-2.696079	0.133258
H	-3.844812	-2.302598	-2.786019
H	-5.127195	0.593780	0.147084
H	1.293653	1.501927	-2.857141
H	0.257964	2.912182	-2.566436
H	-0.146702	1.343240	-1.839316
H	3.631331	0.887881	-1.812070
H	5.185692	-1.443374	-1.549633
H	3.940348	-3.761602	1.869543
H	2.198502	-2.100253	2.433169
H	5.418669	-3.442731	-0.089480
H	-1.238904	2.976785	-0.648980
H	-1.078365	3.879986	0.958550
H	0.729724	1.539364	2.212000
H	0.923082	-0.142429	2.656457
H	-1.912991	-1.031148	-2.253344
H	1.303814	3.478190	1.016401
H	2.183810	4.293712	-1.672344
H	3.236719	2.959462	-2.143760
H	3.308646	3.596690	-0.486593
H	-5.740693	-0.985795	-1.807645
H	-4.386285	0.092754	-2.157224
H	-0.471020	-2.776828	-1.227008
H	-1.776478	-3.583388	-2.092842
H	-1.866535	-3.425445	-0.329770
H	-4.134084	-2.780766	-1.113019
H	-5.158031	-1.106257	0.564017
H	-1.533882	1.225579	2.862432
H	-0.105338	-0.898166	-0.440890

Substrate 3-ox β -anti

1 1

C	-0.529385	-0.917047	-0.383022
C	0.759043	-1.163204	-1.122443
C	1.942433	-0.392719	-0.599941
C	2.126387	0.968052	-0.409759
C	1.249657	2.198865	-0.601046
C	-0.026841	1.867173	-1.345980
C	-1.262977	2.214915	-0.982314
C	-2.844028	-0.186157	0.943839
C	-1.544711	-0.075421	1.640641
N	-0.464218	-0.492313	0.904577
C	4.055623	-0.032882	0.228250
C	5.348542	-0.310305	0.688942
C	5.738781	-1.640382	0.719598
C	4.866429	-2.669118	0.301683
C	3.588524	-2.384819	-0.154674
C	3.165075	-1.043689	-0.194993
N	3.399716	1.163758	0.085140
C	-1.752159	-1.079527	-0.950953
N	-2.892852	-0.731444	-0.252953
C	-4.033750	0.378602	1.644477
O	-1.444202	0.361788	2.777614
C	2.007705	3.217335	-1.488435
C	0.964351	2.838720	0.768208
H	0.116915	1.365376	-2.304794
H	6.015406	0.485085	1.008244
H	6.734079	-1.896061	1.070547
H	5.205801	-3.699908	0.338652
H	2.932575	-3.190121	-0.475096
H	3.793623	2.062080	0.330101
H	0.572821	-0.978002	-2.185030
H	0.984242	-2.233159	-1.040163
H	-1.477977	2.740941	-0.055278
H	-2.111517	1.989370	-1.623546
H	-3.694017	1.281442	2.161210
H	2.264536	2.776253	-2.456511
H	1.372167	4.091164	-1.659801
H	2.930416	3.562985	-1.012058
H	1.896732	3.069757	1.291159
H	0.414918	3.776696	0.641383
H	0.372232	2.177372	1.410802
C	-4.196735	-0.974058	-0.992022
H	-4.131159	-0.310375	-1.860611
C	-5.176905	0.647958	0.677767
H	-6.076707	0.918547	1.234384
H	-4.923454	1.491218	0.024122
C	-5.408302	-0.610629	-0.142285
H	-6.251762	-0.495362	-0.828612
H	-5.645461	-1.447634	0.526861
H	-4.336527	-0.325248	2.432478
C	-4.263478	-2.435708	-1.428427
H	-4.149263	-3.094430	-0.562537
H	-5.252326	-2.606928	-1.862040
H	-3.520822	-2.701501	-2.181501
H	0.457706	-0.376100	1.329132
H	-1.877733	-1.403491	-1.972439

TS-ox α -syn

1 1

N	1.360779	2.373123	-0.379573
C	2.676280	2.008306	-0.481812
O	3.535686	2.707717	-0.996900
C	2.937210	0.645962	0.044154
N	-2.787496	-1.334119	-0.710377
C	2.012997	-0.275617	-1.677385
N	2.114208	0.199733	1.038982
C	0.644108	-0.041813	-1.664538
C	0.404605	1.503277	0.091013
C	-1.039200	1.876291	-0.079849
C	-1.927965	0.673581	-0.213294
C	-3.289236	0.627946	0.252265
C	-4.123157	1.547637	0.912630
C	-5.419219	1.166843	1.226276
C	-5.903093	-0.116640	0.891514
C	-5.102443	-1.041937	0.238141
C	-3.795585	-0.650166	-0.076029
C	-1.655690	-0.546225	-0.796849
C	-0.419301	-1.109504	-1.471508
C	0.089889	-2.331695	-0.694249
C	-0.800573	-1.556569	-2.907753
C	0.812659	0.523681	0.984825
H	-2.864917	-2.285629	-1.043537
H	2.391100	-1.266171	-1.440819
H	2.638355	0.308230	-2.347866
H	0.291938	0.825426	-2.224102
H	-1.129211	2.537367	-0.950483
H	-1.350089	2.469554	0.789424
H	-3.763439	2.540061	1.172274
H	-6.076117	1.865000	1.736351
H	-6.923287	-0.384389	1.149947
H	-5.472703	-2.029337	-0.021060
H	-0.707993	-3.069378	-0.574335
H	0.909339	-2.824453	-1.226228
H	0.432611	-2.055569	0.306649
H	-1.548952	-2.354704	-2.881997
H	-1.209063	-0.720037	-3.482366
H	0.085056	-1.940178	-3.422647
C	4.360847	0.184788	0.069331
H	4.838347	0.497107	-0.861742
C	2.687597	-0.785664	2.013778
H	3.428651	-0.205540	2.575545
C	4.503934	-1.328021	0.305991
H	4.487515	-1.852566	-0.653399
H	5.488012	-1.509944	0.743851
C	3.399361	-1.886241	1.224328
H	2.648014	-2.437662	0.648698
H	3.817214	-2.592083	1.946866
H	4.862947	0.747164	0.869664
C	1.650594	-1.320391	2.984165
H	1.151433	-0.520000	3.536475
H	0.900348	-1.945260	2.489902
H	2.174074	-1.947463	3.709672
H	1.074752	3.221698	-0.862344
H	0.100423	-0.011111	1.598729

TS-ox β -syn

1 1

C	2.645427	1.101914	1.666884
C	-1.963821	0.382474	0.549866
C	-1.733512	-0.945709	0.254115
C	-3.313166	0.679168	0.145300
C	0.393606	1.148706	0.867381
C	-3.854528	-0.509845	-0.393786
C	0.822253	0.885937	-0.421424
C	1.910032	-1.398121	1.184940
C	2.531605	0.315054	-2.076909
C	-0.524499	-1.843611	0.435553
C	0.542311	-1.162736	1.273384
C	-1.051663	1.385247	1.197716
C	4.586218	-0.762844	-1.102739
C	2.028630	1.418501	-3.001492
C	4.041568	0.153846	-2.187576
C	4.338062	-0.104217	0.245202
C	-0.022450	-2.325479	-0.933825
O	3.498085	1.298234	2.518370
N	1.345522	1.511949	1.789669
N	2.119953	0.613235	-0.656877
N	-2.878123	-1.472097	-0.312742
C	-5.161037	-0.580648	-0.891658
C	2.907644	0.322066	0.424602
C	-5.405905	1.774035	-0.299626
C	-4.110551	1.836587	0.191388
C	-0.942500	-3.084062	1.270921
C	-5.925254	0.575570	-0.835936
H	4.246686	-0.242134	-3.186095
H	4.602474	-0.753669	1.084590
H	-0.808412	-2.886105	-1.447574
H	0.835659	-2.995697	-0.829851
H	0.255618	-1.489960	-1.581774
H	-2.982988	-2.420359	-0.647028
H	-5.558312	-1.504022	-1.302508
H	-6.034684	2.658877	-0.271621
H	-3.724104	2.764472	0.605333
H	-6.944026	0.558486	-1.211501
H	2.296411	-2.053680	0.408612
H	2.495301	-1.376022	2.100480
H	-1.167150	1.381898	2.288374
H	-1.320878	2.395615	0.864474
H	2.054936	-0.640001	-2.328118
H	0.173050	-0.815274	2.239144
H	-0.075850	-3.733605	1.424946
H	-1.710912	-3.663550	0.750386
H	-1.337883	-2.785177	2.246106
H	5.658347	-0.926884	-1.235329
H	4.098430	-1.744257	-1.155967
H	0.940356	1.465943	-3.069218
H	2.410170	1.218057	-4.006166
H	2.409387	2.390208	-2.673221
H	4.519944	1.139796	-2.120844
H	4.962174	0.793414	0.348182
H	1.061809	1.900217	2.686145
H	0.117488	0.834927	-1.240957

TS-ox β -anti

1 1

C	-0.391625	-1.029075	0.217560
C	0.946037	-1.605762	-0.119853
C	2.034481	-0.577342	-0.179927
C	1.942621	0.773127	-0.427723
C	0.774026	1.708542	-0.680006
C	-0.510452	1.001257	-1.102571
C	-1.768704	1.482873	-0.737014
C	-2.706932	0.262638	0.623029
C	-1.666691	0.348428	1.693633
N	-0.540926	-0.372500	1.420601
C	4.139797	0.327149	-0.123179
C	5.532197	0.395381	0.006597
C	6.203010	-0.788138	0.278973
C	5.510596	-2.010558	0.418824
C	4.131056	-2.070344	0.288078
C	3.427187	-0.884710	0.014684
N	3.218058	1.309306	-0.389922
C	-1.537185	-1.436071	-0.435662
N	-2.726959	-0.897058	-0.115532
C	-3.996563	0.958184	0.972127
O	-1.797883	1.048889	2.686244
C	1.110168	2.592924	-1.914780
C	0.594407	2.639434	0.536547
H	-0.425462	0.427279	-2.022827
H	6.063766	1.336155	-0.101992
H	7.283589	-0.774104	0.386155
H	6.070804	-2.916324	0.630611
H	3.608149	-3.017632	0.393262
H	3.448297	2.284137	-0.523171
H	0.857266	-2.151095	-1.065941
H	1.195477	-2.352208	0.646991
H	-1.829751	2.379173	-0.125800
H	-2.576995	1.363374	-1.453926
H	-3.773960	2.013601	1.147517
H	1.356109	1.978988	-2.785862
H	0.253747	3.227685	-2.158532
H	1.957345	3.252305	-1.702296
H	1.561322	3.065960	0.816903
H	-0.069333	3.474793	0.295505
H	0.197214	2.121857	1.413646
C	-3.923059	-1.316622	-0.930911
H	-3.748928	-0.902035	-1.932688
C	-5.079107	0.756822	-0.077031
H	-6.023404	1.161113	0.296019
H	-4.838850	1.300105	-0.998890
C	-5.208164	-0.732499	-0.358676
H	-6.000781	-0.931312	-1.085483
H	-5.469057	-1.267129	0.564052
H	-4.328471	0.553159	1.937553
C	-4.002372	-2.838643	-0.994481
H	-4.036772	-3.258087	0.015391
H	-4.926624	-3.107400	-1.512596
H	-3.177733	-3.296625	-1.542964
H	0.265666	-0.248365	2.028991
H	-1.489160	-2.128691	-1.265037

NADP⁺ Model

I 1

C	2.790063	-1.392824	0.158797
N	1.705279	-0.391857	0.053557
C	0.421878	-0.781114	0.056584
C	-0.603546	0.149820	-0.025245
C	-0.278816	1.499882	-0.133451
C	1.061863	1.881738	-0.145606
C	2.039401	0.913533	-0.045207
C	-2.007053	-0.404586	-0.034389
N	-2.985189	0.434961	0.347536
O	-2.199639	-1.560682	-0.393977
H	0.219375	-1.843532	0.123012
H	-1.050201	2.257244	-0.230332
H	1.351904	2.921105	-0.235254
H	3.099418	1.138415	-0.043286
H	-3.933108	0.079769	0.380652
H	-2.807902	1.316283	0.809492
H	2.346753	-2.383747	0.217120
H	3.420956	-1.315322	-0.725881
H	3.366557	-1.185181	1.059742

NADPH Model

0 1

C	2.804645	-1.420812	0.046659
N	1.737492	-0.438670	-0.066344
C	0.424910	-0.795876	-0.034847
C	-0.601348	0.090465	-0.008902
C	-0.368709	1.590423	-0.004480
C	1.113469	1.880448	0.020453
C	2.048215	0.923566	-0.010146
C	-1.961838	-0.473088	0.003684
N	-2.977515	0.430043	0.022826
O	-2.204118	-1.690726	0.003519
H	0.218394	-1.861939	-0.037528
H	-0.823865	2.065153	-0.888780
H	1.436491	2.915861	0.056101
H	3.110202	1.146716	0.002971
H	-3.929717	0.090575	0.010576
H	-2.832420	1.428218	-0.012397
H	2.405103	-2.408712	-0.185188
H	3.599379	-1.188959	-0.666868
H	3.225411	-1.434775	1.058130
H	-0.851952	2.063304	0.864925

Substrate 4

0 1

N	-1.653566	-1.443146	-1.806768
C	-2.974416	-1.335511	-1.516315
O	-3.863219	-1.895274	-2.195457
C	-3.379938	-0.528575	-0.354653
N	3.224166	0.526881	0.844709
C	3.680073	-2.739551	-1.185734
N	-2.452993	0.129521	0.328734
C	3.219104	-2.958944	0.045106
C	-0.763580	-0.798956	-1.042950
C	0.696335	-0.980887	-1.412691
C	1.664462	-0.163215	-0.613394
C	1.867801	1.249866	-0.786884
C	1.295475	2.214793	-1.635757
C	1.721676	3.531002	-1.535107
C	2.711499	3.906894	-0.599910
C	3.290239	2.976094	0.251309
C	2.854098	1.649673	0.145274
C	2.506527	-0.567940	0.401165
C	2.686444	-1.933632	1.032258
C	3.682696	-1.880065	2.206715
C	1.331788	-2.423128	1.588084
C	-1.126610	-0.003219	0.021925
H	3.913169	0.509447	1.583500
H	3.677444	-1.747703	-1.632196
H	4.068947	-3.555903	-1.787388
H	3.239554	-3.974623	0.445496
H	0.787614	-0.726848	-2.477719
H	0.932398	-2.048263	-1.344642
H	0.528515	1.933406	-2.354220
H	1.289436	4.288487	-2.182471
H	3.024376	4.945443	-0.544784
H	4.049535	3.263785	0.972719
H	3.324518	-1.212678	2.998926
H	3.788677	-2.879586	2.639224
H	4.676555	-1.556026	1.880236
H	0.951811	-1.714061	2.331314
H	0.580146	-2.528729	0.801248
H	1.458307	-3.397962	2.072151
C	-4.826315	-0.519207	0.025387
H	-5.371037	0.117244	-0.686448
C	-2.774861	1.053792	1.480893
H	-2.375200	0.539455	2.361469
C	-5.046279	-0.040126	1.452257
H	-6.112736	0.110556	1.638453
H	-4.690775	-0.796087	2.163336
C	-4.276837	1.257081	1.639300
H	-4.620604	1.997977	0.904986
H	-4.439055	1.686134	2.632424
H	-5.200256	-1.534173	-0.139497
C	-2.076443	2.398421	1.284264
H	-0.990263	2.345173	1.375399
H	-2.331534	2.816475	0.305239
H	-2.440568	3.083123	2.055296
H	-0.414112	0.517378	0.641646

Substrate 9

0 1

N	-2.174768	-0.794634	-1.699387
C	-3.479053	-0.638782	-1.330516
O	-4.439116	-0.993530	-2.048699
C	-3.723065	-0.027350	-0.036473
N	3.039611	0.251152	0.841516
C	3.014969	-2.664727	-1.679730
N	-2.701278	0.365695	0.703932
C	2.600086	-3.036108	-0.469125
C	-1.185127	-0.376414	-0.902387
C	0.224019	-0.609766	-1.413295
C	1.320114	-0.031563	-0.571604
C	1.683556	1.359762	-0.557268
C	1.188548	2.498558	-1.218360
C	1.782175	3.727244	-0.969221
C	2.864795	3.844393	-0.069461
C	3.371662	2.737939	0.597211
C	2.767316	1.501031	0.341175
C	2.169263	-0.677375	0.302921
C	2.233445	-2.138059	0.701038
C	3.291920	-2.373192	1.796103
C	0.866458	-2.578783	1.266501
C	-1.406805	0.225986	0.327607
H	3.763545	0.046527	1.515891
H	3.093912	-1.617797	-1.964210
H	3.282285	-3.405786	-2.427402
H	2.534213	-4.100567	-0.234661
H	0.269647	-0.182732	-2.424270
H	0.358563	-1.689014	-1.544756
H	0.352354	2.416871	-1.909326
H	1.411550	4.616405	-1.470899
H	3.308397	4.820259	0.104957
H	4.203333	2.826786	1.290196
H	3.053937	-1.807845	2.704398
H	3.310007	-3.434443	2.061982
H	4.296405	-2.103923	1.452821
H	0.603268	-1.966991	2.135930
H	0.067084	-2.483650	0.526833
H	0.914881	-3.626859	1.582310
H	-0.636492	0.575763	1.000905
C	-3.142864	1.028586	1.957634
C	-4.608184	0.593662	2.071577
C	-5.028695	0.279279	0.621431
H	-3.023786	2.105445	1.811898
H	-2.505167	0.700021	2.778495
H	-5.220880	1.369098	2.531872
H	-4.676883	-0.310205	2.681887
H	-5.723034	-0.560388	0.541614
H	-5.496351	1.139855	0.126150

Substrate 9-red

0 1

N	-2.214608	-0.574435	-1.688337
C	-3.508914	-0.646115	-1.272715
O	-4.436927	-0.943849	-2.019499
C	-3.670206	-0.378290	0.211961
N	3.113313	0.158812	0.850912
C	2.938336	-2.739603	-1.666171
N	-2.751139	0.681541	0.606993
C	2.516116	-3.096380	-0.453663
C	-1.128125	-0.209988	-0.853350
C	0.238801	-0.559956	-1.379389
C	1.365474	-0.034940	-0.541591
C	1.802898	1.334712	-0.536295
C	1.367493	2.495596	-1.200769
C	2.033020	3.689896	-0.966833
C	3.128776	3.751615	-0.077151
C	3.578017	2.622377	0.592482
C	2.902903	1.419841	0.349195
C	2.189019	-0.723723	0.323971
C	2.188261	-2.186676	0.719048
C	3.235994	-2.470679	1.813151
C	0.804209	-2.570184	1.284287
C	-1.413681	0.426002	0.294738
H	3.831243	-0.082059	1.519441
H	3.054870	-1.696015	-1.950155
H	3.174255	-3.489381	-2.415730
H	2.412208	-4.157999	-0.219905
H	0.334270	-0.150414	-2.397767
H	0.318042	-1.648403	-1.492803
H	0.519640	2.456076	-1.881125
H	1.708927	4.595390	-1.471621
H	3.628523	4.701893	0.086411
H	4.420245	2.667159	1.277000
H	3.025763	-1.893321	2.720707
H	3.203368	-3.531248	2.080517
H	4.252311	-2.250299	1.469529
H	0.576106	-1.960329	2.165044
H	0.006313	-2.420104	0.552401
H	0.804020	-3.624383	1.583279
H	-3.406578	-1.321845	0.736951
H	-2.030489	-0.835975	-2.651949
H	-0.639317	0.780754	0.967025
C	-3.149632	1.065048	1.955465
C	-4.686918	1.079456	1.864110
C	-5.032922	0.129805	0.683705
H	-2.718238	2.031600	2.227335
H	-2.810573	0.311144	2.686440
H	-5.044978	2.090485	1.655468
H	-5.139660	0.753189	2.802801
H	-5.683349	-0.694738	0.981367
H	-5.527975	0.671309	-0.126761

Supplementary References

1. Holm, L. & Rosenström, P. Dali server: conservation mapping in 3D. *Nucleic Acids Res.* **38**, W545-W549 (2010).
2. Ding, Y., Greshock, T.J., Miller, K.A., Sherman, D.H. & Williams, R.M. Premalbrancheamide: synthesis, isotopic labeling, biosynthetic incorporation, and detection in cultures of *Malbranchea aurantiaca*. *Org. Lett.* **10**, 4863-4866 (2008).