

Minutissamides E - L, Antiproliferative Cyclic Lipodecapeptides from the Cultured Freshwater Cyanobacterium cf. *Anabaena* sp.

*Hahk-Soo Kang, Megan Sturdy, Aleksej Krunic, Hyunjung Kim, Qi Shen, Steven M. Swanson and Jimmy
Orjala**

Department of Medicinal Chemistry and Pharmacognosy, University of Illinois at Chicago, 833 S.
Wood St., Chicago, Illinois, 60612

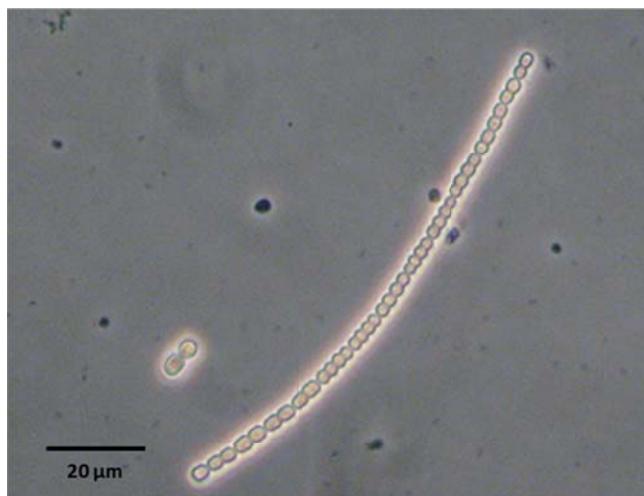
Supplementary Data

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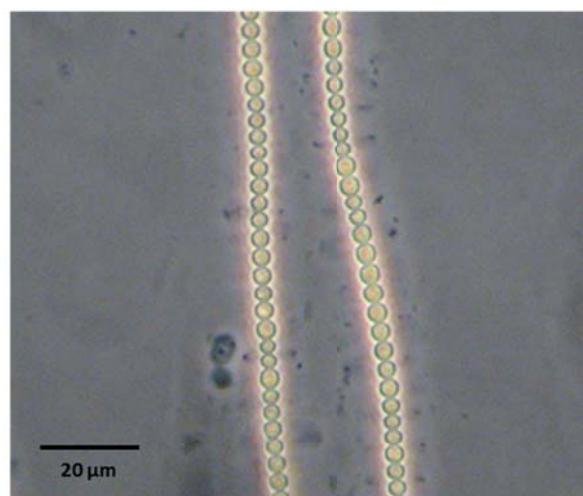
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S1. Morphological comparison between *Anabaena minutissima* UTEX 1613 and cf. *Anabaena* sp. UIC 10035

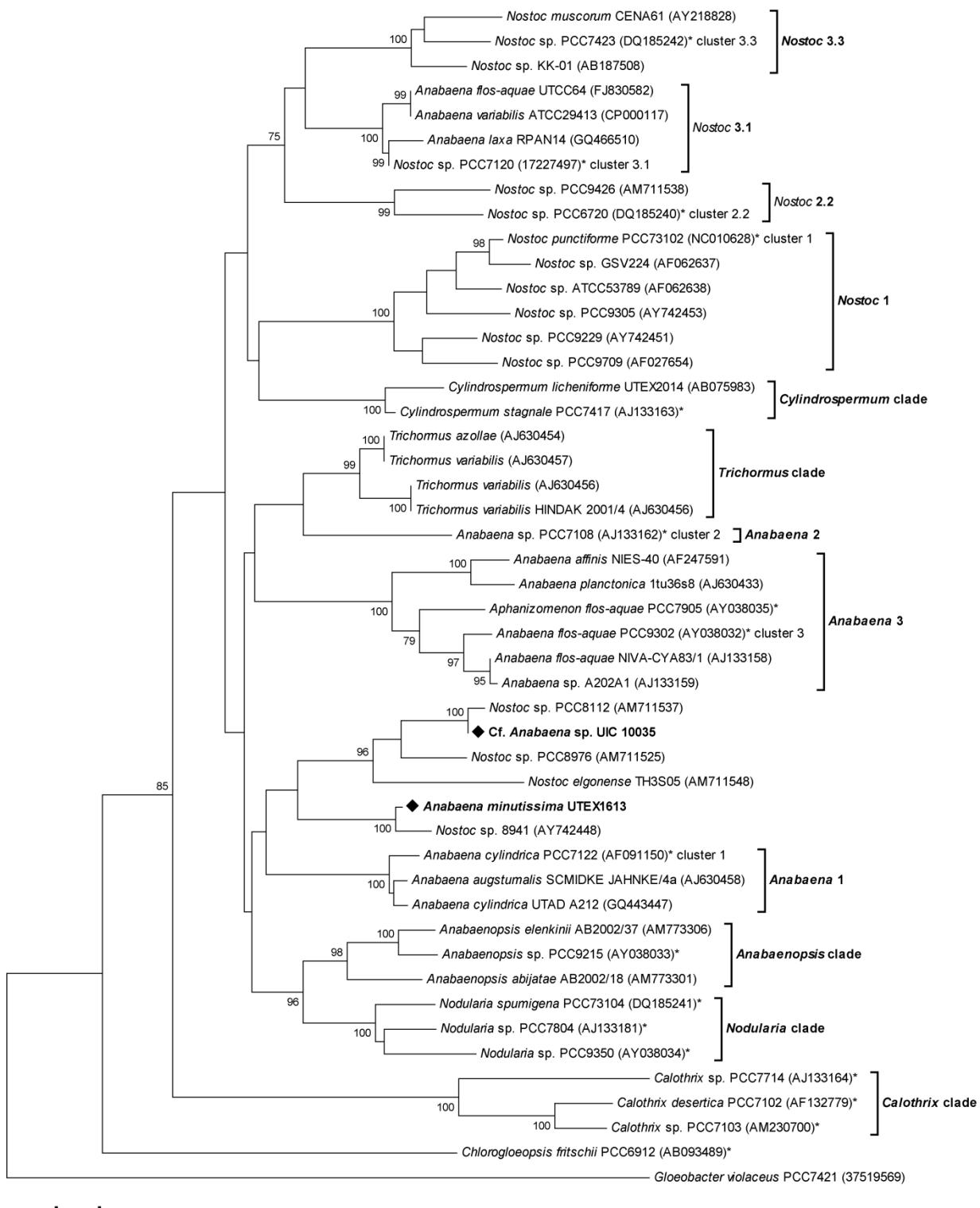


UTEX 1613



UIC 10035

S2. Complete phylogenetic tree for taxonomic identification of UIC 10035



Evolutionary distances were determined using the minimum evolution method with 1,000 replicate bootstrap re-sampling to construct the phylogenetic tree. Strains denoted with an asterisk (*) are “Bergey’s” reference strains. Strains were obtained from NCBI with the accession number given in parentheses. Only bootstrap values greater than or equal to 75% are displayed.

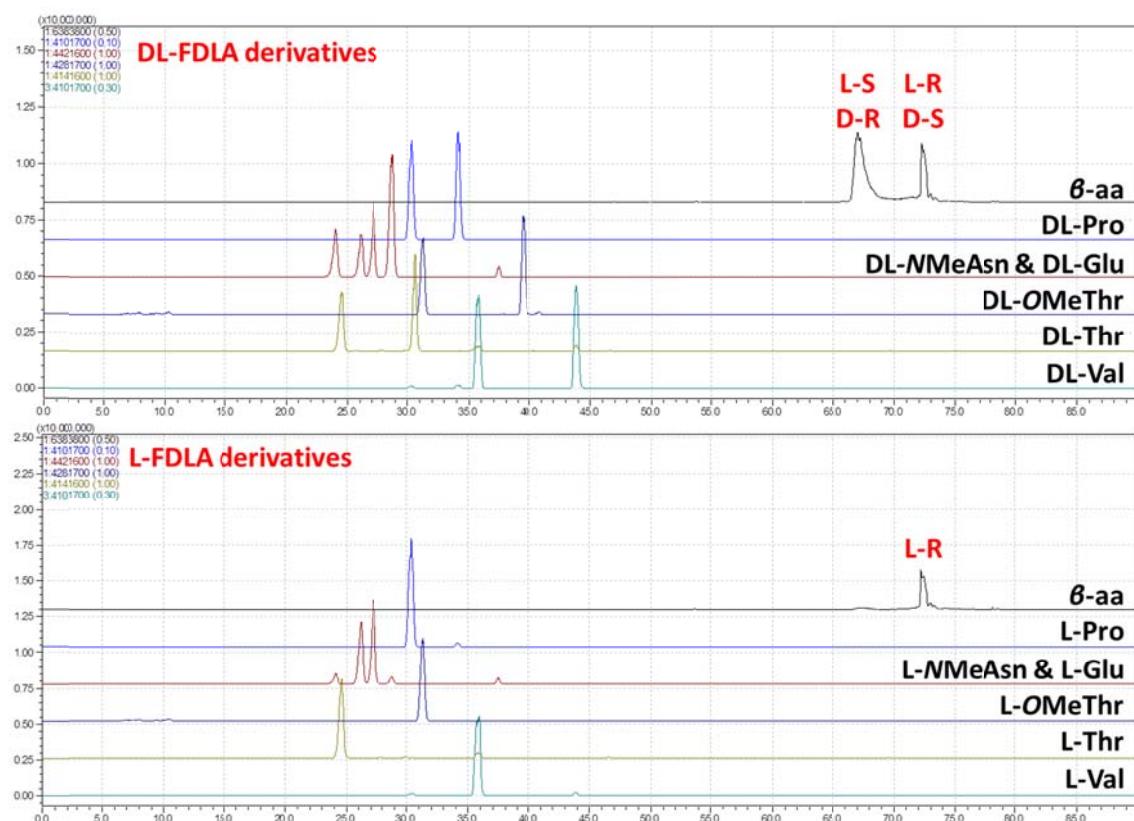
S3. Complete NMR data for minutissamide E (**1**) in DMSO-*d*₆

| | | Minutissamide E (1) | | | | | |
|--------|-----------------|------------------------------|-----------------------|----------------------------|-------------------|------------------------------------|--|
| | | δ_{C}^a | δ_{H}^b | mult. (<i>J</i> in Hz) | COSY ^b | HMBC ^b | ROESY ^b |
| Ahmoo | 1 | 169.6 | | | | | |
| | 2 | 69.6 | 4.17 | d (4.8) | H-3 | 1 | NH |
| | 3 | 56.0 | 3.92 | td (10.8, 4.8) | H-2, H-4, NH | 1, 2, 4, 4-Me | H-2, H ₂ -5, 4-Me |
| | 4 | 32.2 | 1.68 | m | H-3, H-5, 4-Me | | H-2 |
| | 5 | 33.4 | 1.17 | m | | | H-3, 4-Me |
| | | | 1.62 | m | | | |
| | 6 | 25.5 | 1.17 | m | | | |
| | | | 1.25 | m | | | |
| | 7 | 29.7 | 1.25 | m | | | |
| | 8 | 29.3 | 1.25 | m | | | |
| | 9 | 29.1 | 1.25 | m | overlapped | | overlapped |
| | 10 | 29.0 | 1.25 | m | | | |
| | 11 | 28.9 | 1.25 | m | | | |
| | 12 | 28.6 | 1.25 | m | | | |
| Pro | 13 | 23.3 | 1.44 | m | H-14, H-12 | 12, 14, 15 | |
| | 14 | 41.9 | 2.37 | t (7.2) | H-13 | 12, 13 | |
| | 15 | 210.6 | | | | | |
| | 16 | 43.8 | 2.36 | t (7.2) | H-17 | 15, 17, 18 | |
| | 17 | 16.7 | 1.47 | s (7.2) | H-16, H-18 | 15, 16, 18 | |
| | 18 | 13.6 | 0.83 | t (7.2) | H-17 | 16, 17 | |
| | 3-NH | | 6.77 | d (10.8) | | 3, 1 _{Pro} | H-2, H-4, 4-Me, H-2 _{NMeAsn} |
| | 4-Me | 16.0 | 0.57 | d (6.6) | | 3, 5 | H-3, H-5 |
| | | | | | | | |
| | | | | | | | |
| NMeAsn | 1 | 171.2 | | | | | |
| | 2 | 59.9 | 4.25 | dd (8.4, 2.4) | H-3 | 1, 3, 4 | H-3, H-4, NH _{Ahmoo} |
| | 3 | 30.1 | 1.94 | m | | | |
| | | | 1.99 | m | H-2, H-4 | | |
| | 4 | 23.4 | 1.71 | m | | | |
| | | | 1.84 | m | H-3, H-5 | | |
| OMeThr | 5 | 46.7 | 3.11 | m | | | |
| | | | 4.21 | m | H-4 | | H-2, H-3, H-4, H-2 _{NMeAsn} |
| | | | | | | | |
| | 1 | 167.4 | | | | | |
| | 2 | 49.7 | 5.52 | dd (11.4, 3.6) | H-3 | 1, 3, 4, N-Me, 1 _{OMeThr} | H-3, NH _{Thr2} , H-5 _{Pro} , H-2 _{Th2} , NH _{Thr2} , NH _{Ahmoo} |
| | 3 | 33.7 | 1.97 | overlapped | H-2 | 1, 2, 4 | H-2 |
| Ala | | | 3.00 | dd (15.6, 12.0) | | | |
| | 4 | 171.5 | | | | | |
| | N-Me | 30.5 | 2.93 | s | | 2, 1 _{OMeThr} | H-2 _{OMeThr} , H-3 _{OMeThr} |
| | NH ₂ | | 5.99 | s | | 3, 4 | H-3, H-3 _{Dhb} |
| | | | 7.49 | s | | | |
| | | | | | | | |
| Gln | 1 | 169.6 | | | | | |
| | 2 | 52.6 | 4.79 | dd (9.0, 1.8) | NH, H-3 | 3, 4, 1 _{Ala} | H-3, H-4, NH, NMe |
| | 3 | 75.1 | 3.71 | qd (6.0, 1.8) | H-2, H-4 | OMe | H-2, H-4, NMe, H ₂ -5 _{Pro} |
| | 4 | 14.7 | 0.95 | d (6.0) | H-3 | 2, 3 | H-2, H-3, OMe |
| | O-Me | 55.6 | 3.13 | s | | 3 | H-3 |
| | NH | | 6.74 | d (9.0) | H-2 | 1 _{Ala} | NH _{Ala} |
| Ala | 1 | 171.9 | | | | | |
| | 2 | 49.1 | 4.19 | p (7.8) | H-3, NH | 3 | NH |
| | 3 | 16.3 | 1.29 | d (7.8) | H-2 | 1, 2 | H-2 |
| | NH | | 7.58 | d (7.8) | H-2 | 2, 3, 1 _{Gln} | H-2, H-3, NH _{OMeThr} , H-2 _{Gln} |

| | | | | | | |
|------|----|-------|------|----------------|----------------|---|
| | | | 7.27 | s | | |
| Thr1 | 1 | 170.4 | | | | |
| | 2 | 61.3 | 3.90 | m | H-3, NH | 3 |
| | 3 | 65.2 | 4.17 | overlapped | H-2, H-4 | |
| | 4 | 20.6 | 1.25 | overlapped | H-3 | 2, 3 |
| | NH | | 8.84 | brs | H-2 | |
| | | | | | | H-3, H-4 H-2, H-3, H-2 _{Thr2} , H-3 _{Thr2} , NH _{Gln} |
| Thr2 | 1 | 174.2 | | | | |
| | 2 | 56.6 | 5.02 | dd (10.2, 2.4) | H-3, NH | 1, 3, 1 _{Dhb} |
| | 3 | 70.3 | 4.57 | brm | H-2, H-4 | H-3, H-4, NH, NH _{Thr1} H-2, H-4, H-2 _{OMeThr} , NH _{Gln} , NH _{Thr1} |
| | 4 | 19.0 | 1.25 | overlapped | H-3 | H-3, H-5 _{Pro} |
| | NH | | 8.37 | d (10.2) | H-2 | H-2, H-2 _{NMeAsn} , NH _{OMeThr} |
| Dhb | 1 | 163.9 | | | | |
| | 2 | 132.4 | | | | |
| | 3 | 117.3 | 5.38 | q (7.2) | H-4 | 1, 2, 4 |
| | 4 | 13.2 | 1.75 | d (7.2) | H-3 | 2, 3 |
| | NH | | 9.08 | s | | 1, 2, 3, 1 _{Val} |
| Val | 1 | 168.8 | | | | |
| | 2 | 55.5 | 4.31 | dd (9.0, 6.0) | H-3, NH | 3, 4, 4' |
| | 3 | 32.7 | 1.81 | m | H-2, H-4, H-4' | 1, 2, 4, 4' |
| | 4 | 18.4 | 0.83 | d (6.0) | H-3 | 2, 3, 4' |
| | 4' | 19.0 | 0.89 | d (6.0) | H-3 | 2, 3, 4 |
| | NH | | 6.85 | d (9.0) | | H-2, H-3, H-4, H-2 _{Ahmoo} |

^aCarbon chemical shifts were assigned from the DEPT-Q spectrum (226 MHz), ^bMeasured at 600 MHz

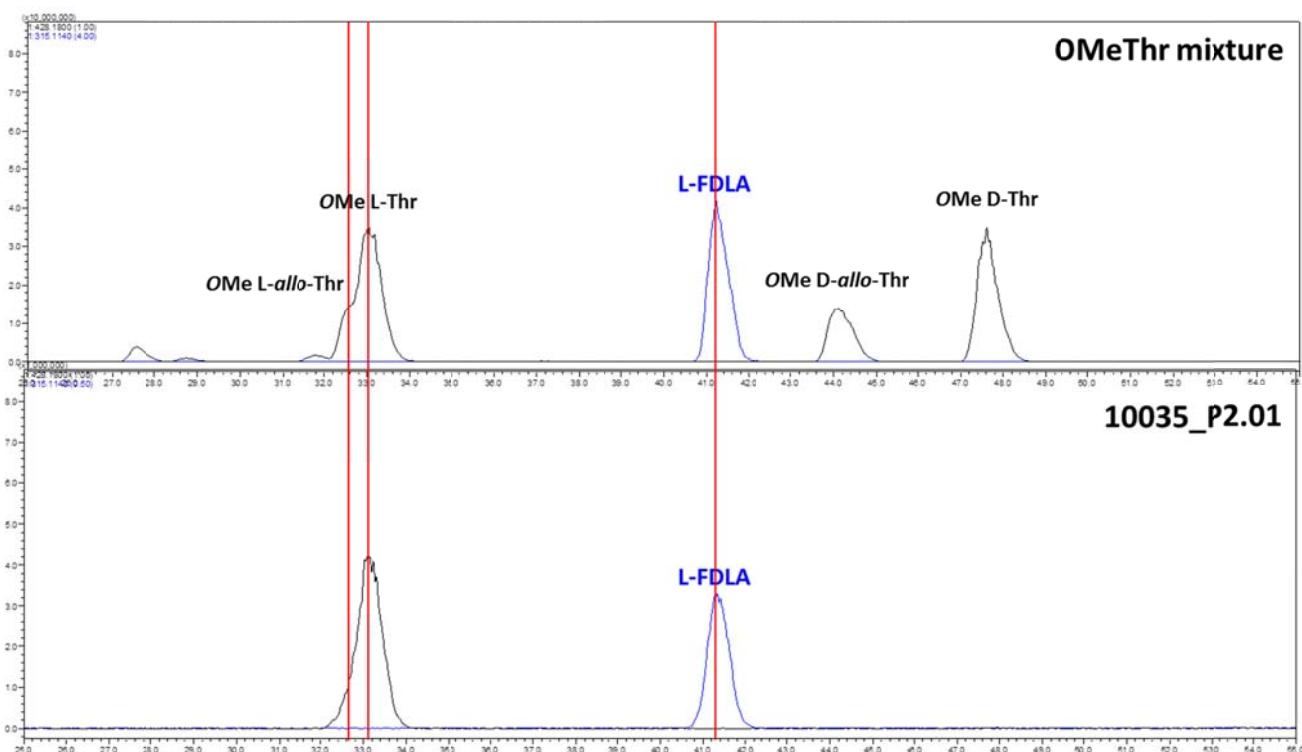
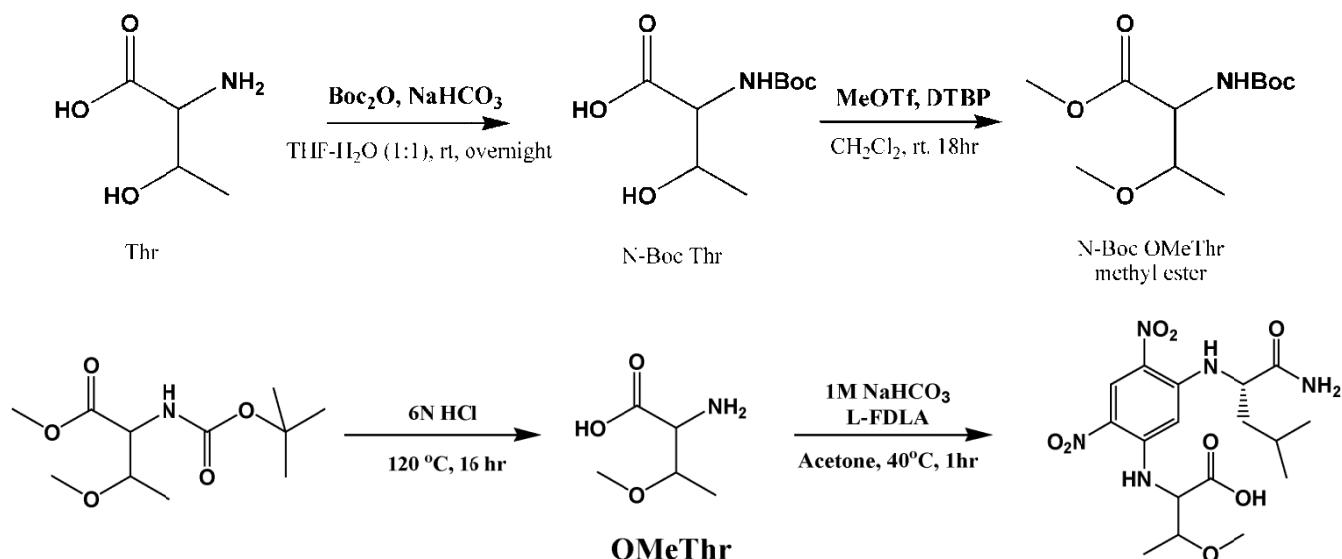
S4. Advanced Marfey's analysis of **1** for the determination of amino acid configurations



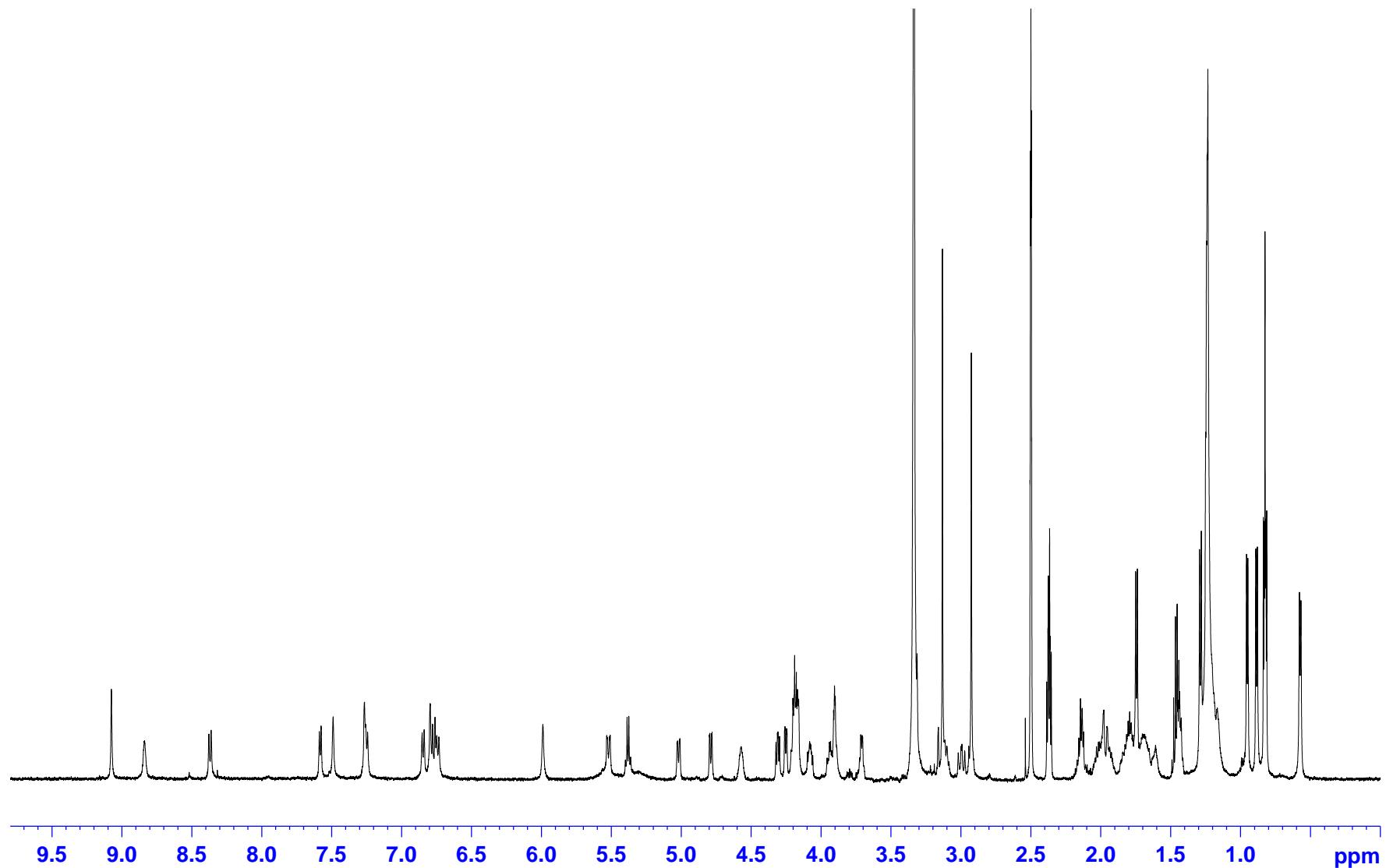
| Amino acid | HPLC retention times of DL-FDLA derivative of acid hydrolysate of 1 | | HPLC retention times of L-FDLA derivative of acid hydrolysate of 1 | Assignment |
|------------|--|--------------|---|------------|
| | L | D | | |
| Ahmoo | 67.0 min (S) | 72.4 min (R) | 72.4 min | R |
| Pro | 30.4 min | 34.1 min | 30.5 min | L |
| NMeAsn | 26.3 min | 24.0 min | 26.3 min | L |
| Ala | 30.9 min | 35.4 min | 35.6 min | D |
| Gln | 27.2 min | 28.7 min | 27.4 min | L |
| Thr | 24.5 min | 30.6 min | 24.7 min | L |
| Val | 35.8 min | 43.8 min | 35.9 min | L |

- L-FDLA derivative of Thr was further compared with L-FDLA derivatives of four Thr amino acid standards L-Thr, L-*allo*-Thr, D-Thr and D-*allo*-Thr, and corresponded with the retention time of L-Thr.

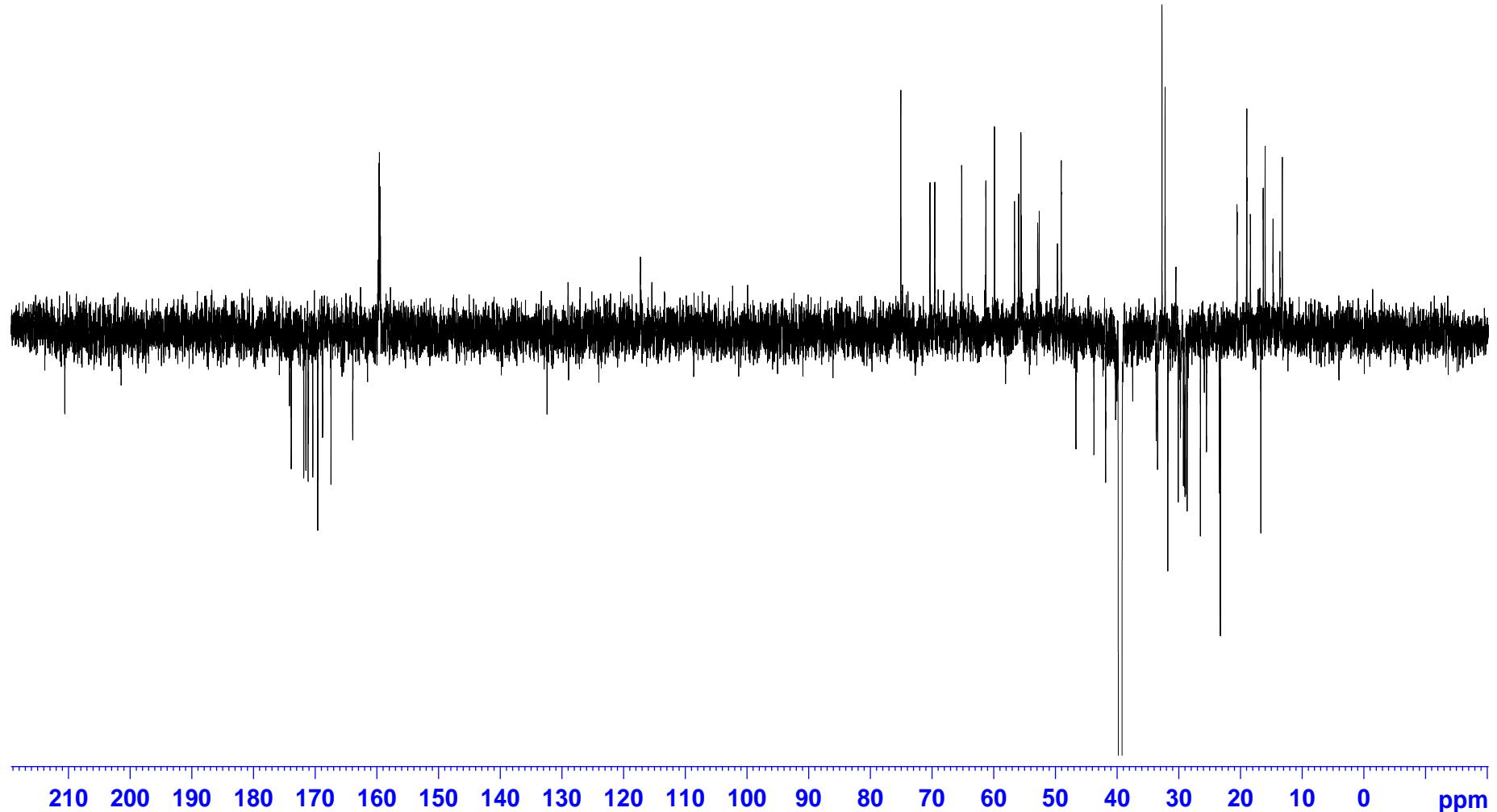
S5. Synthesis of OMeThr standards and Marfey's analysis of **1** for the absolute configuration of OMeThr



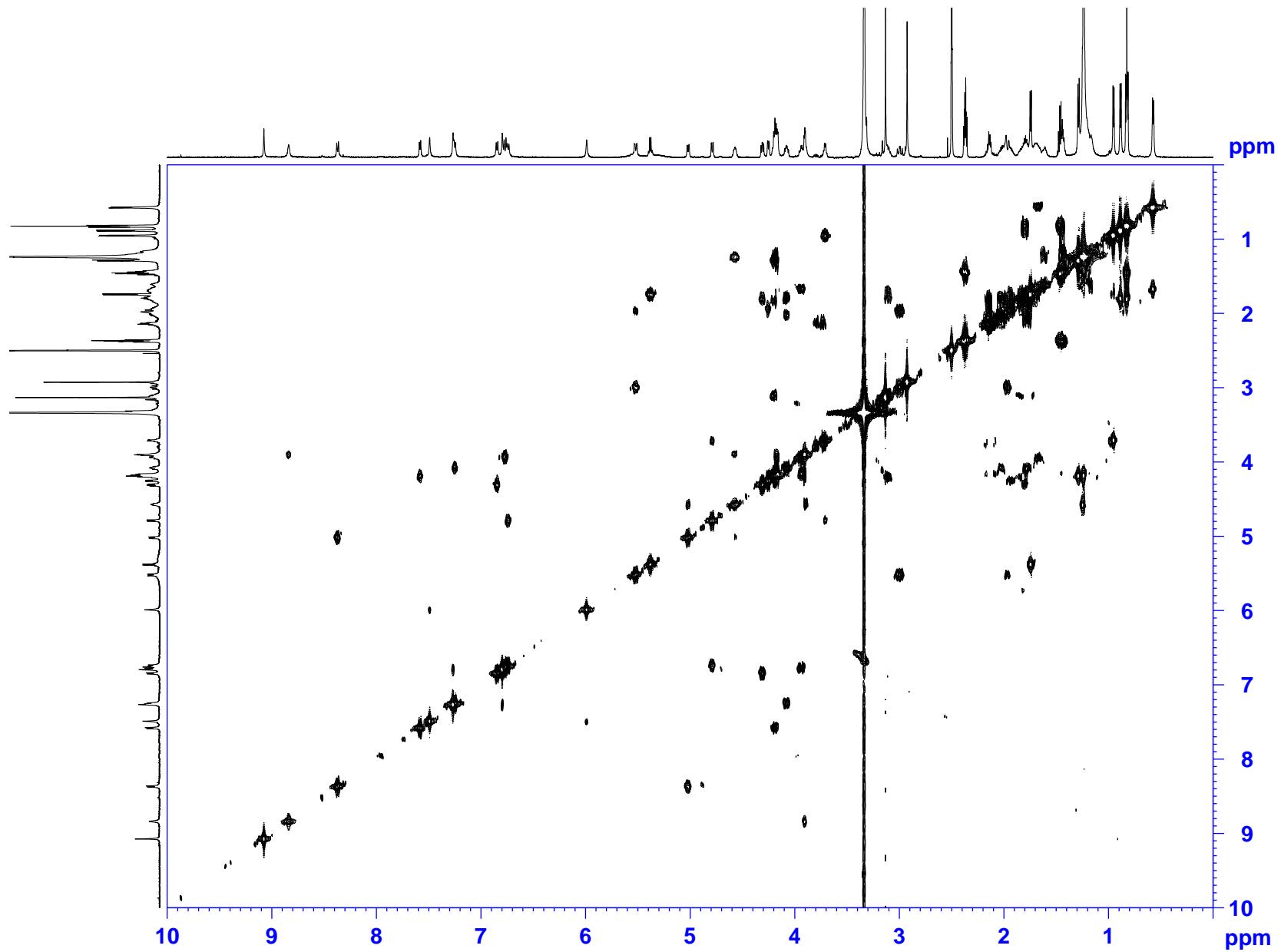
S6. ^1H NMR spectrum (600 MHz, DMSO- d_6) of **1**



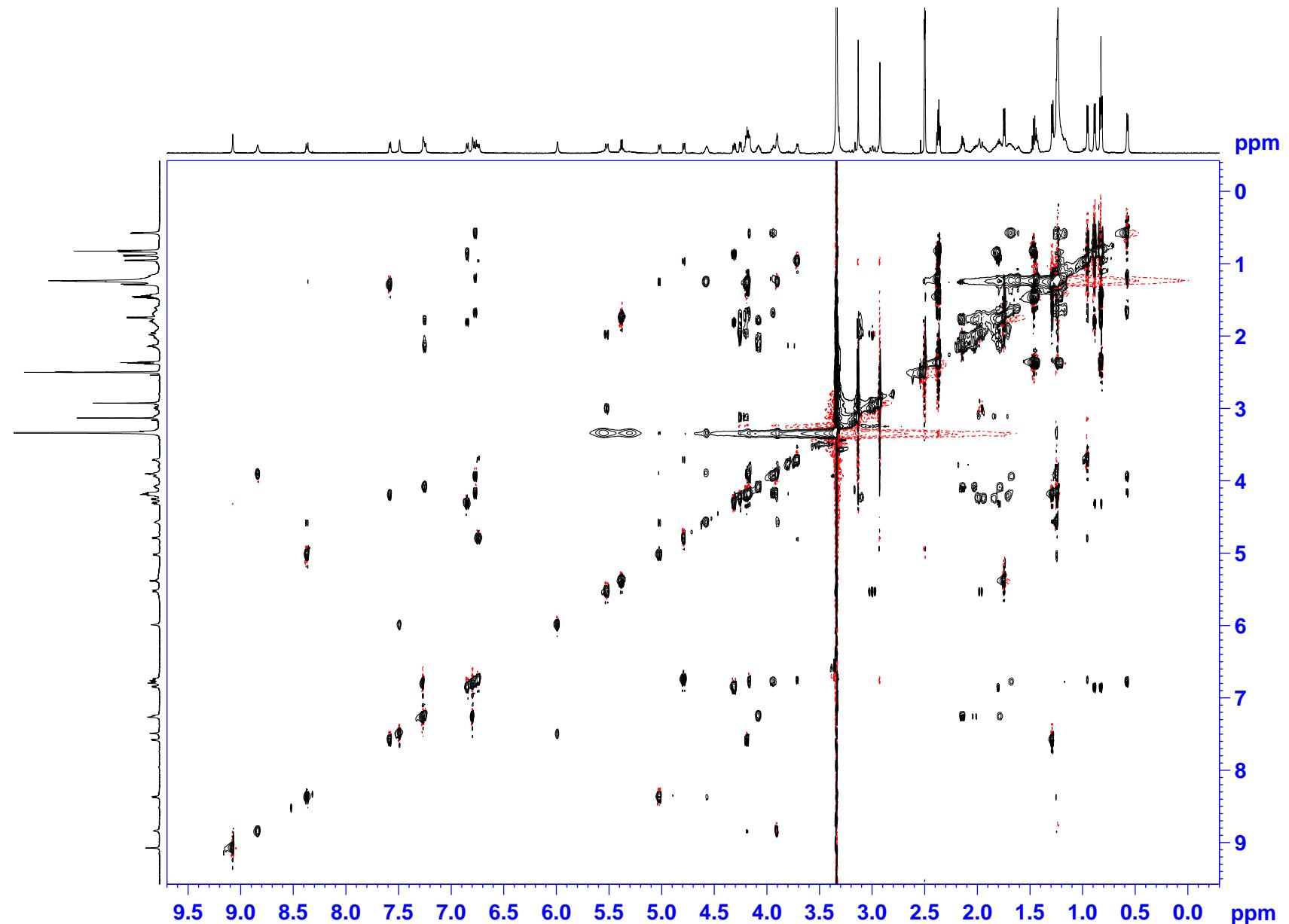
S7. DEPTQ spectrum (226 MHz, DMSO-*d*₆) of **1**



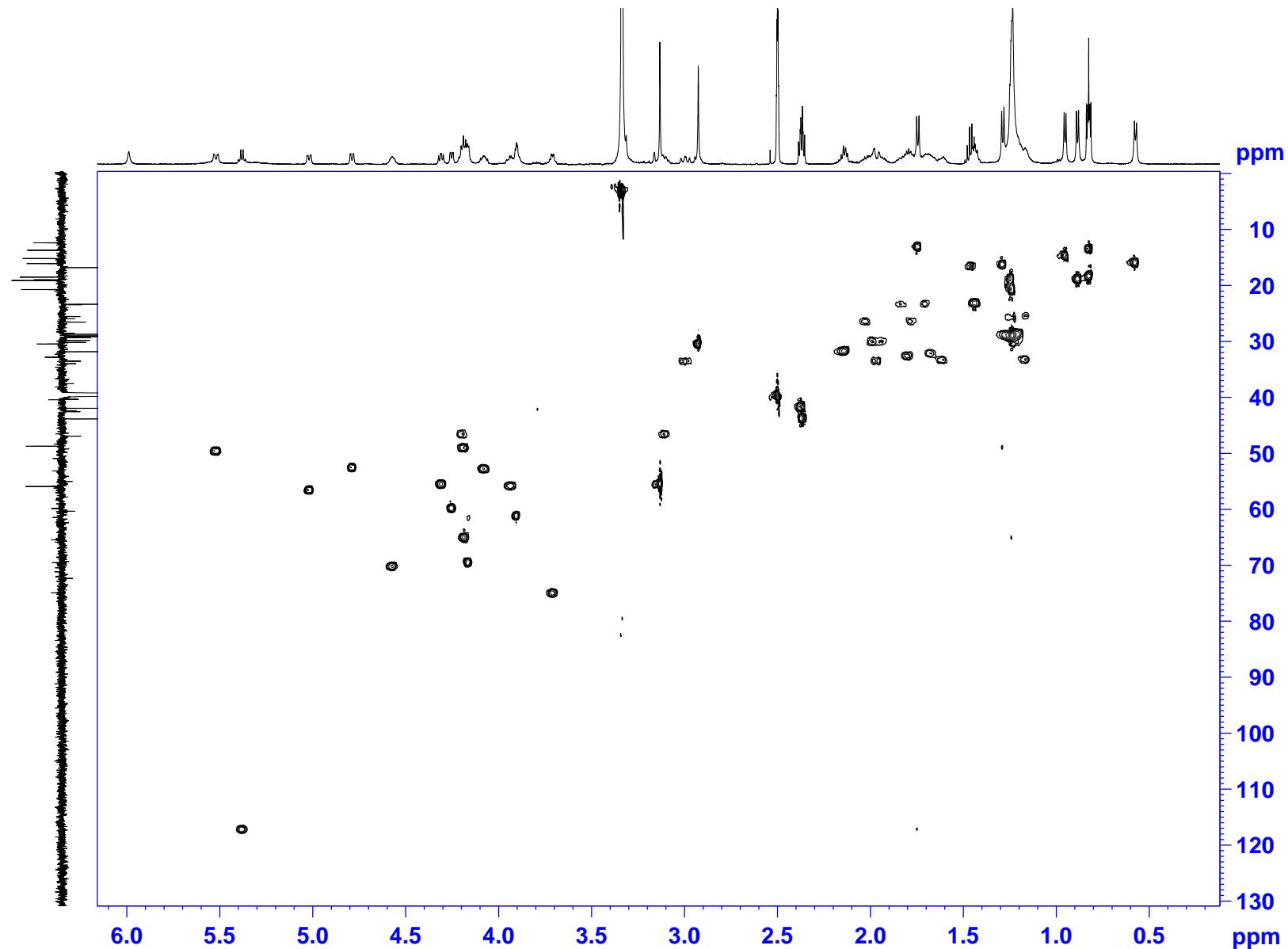
S8. COSY spectrum (600 MHz, DMSO-*d*₆) of **1**



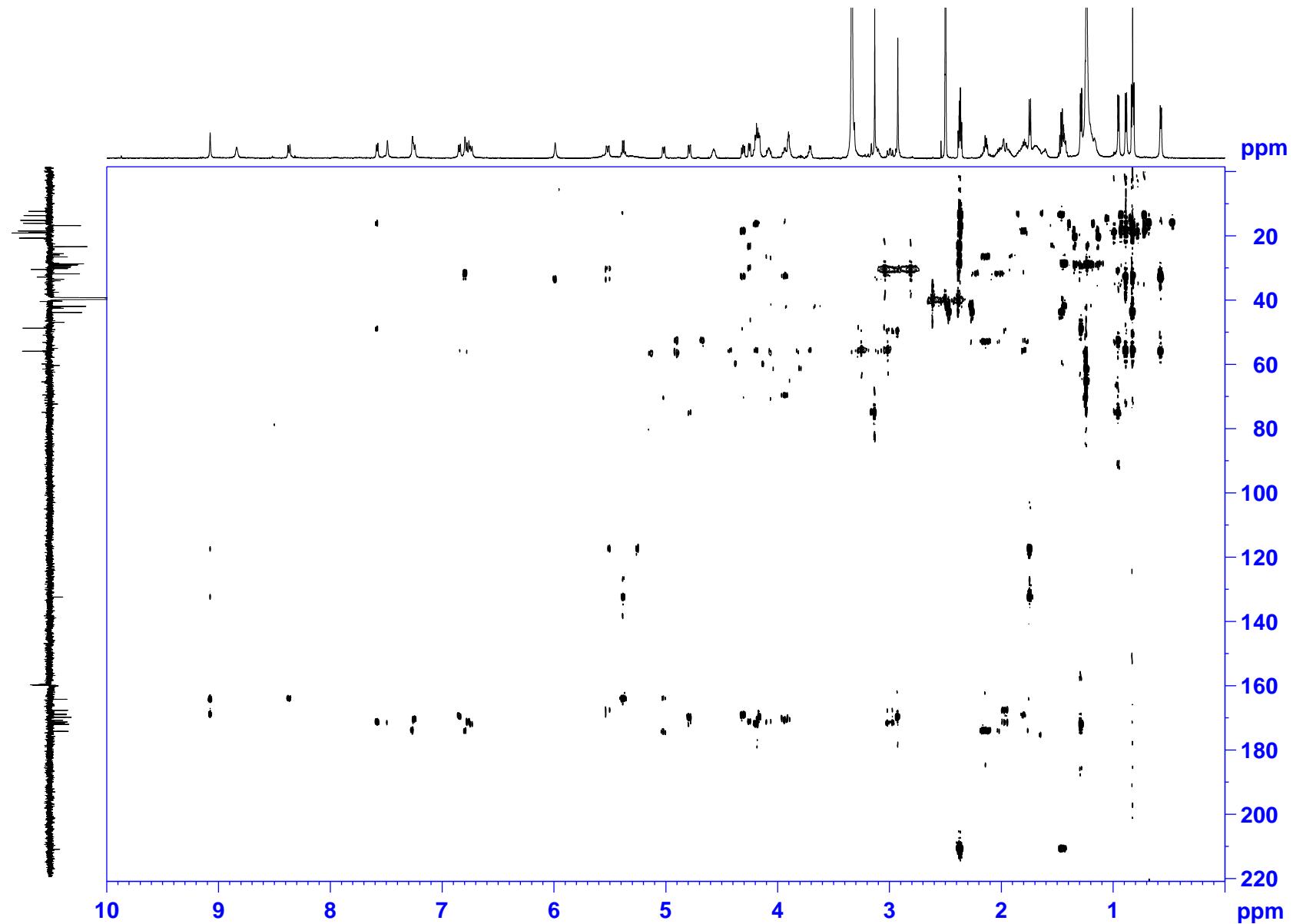
S9. TOCSY spectrum (600 MHz, DMSO-*d*₆) of **1**



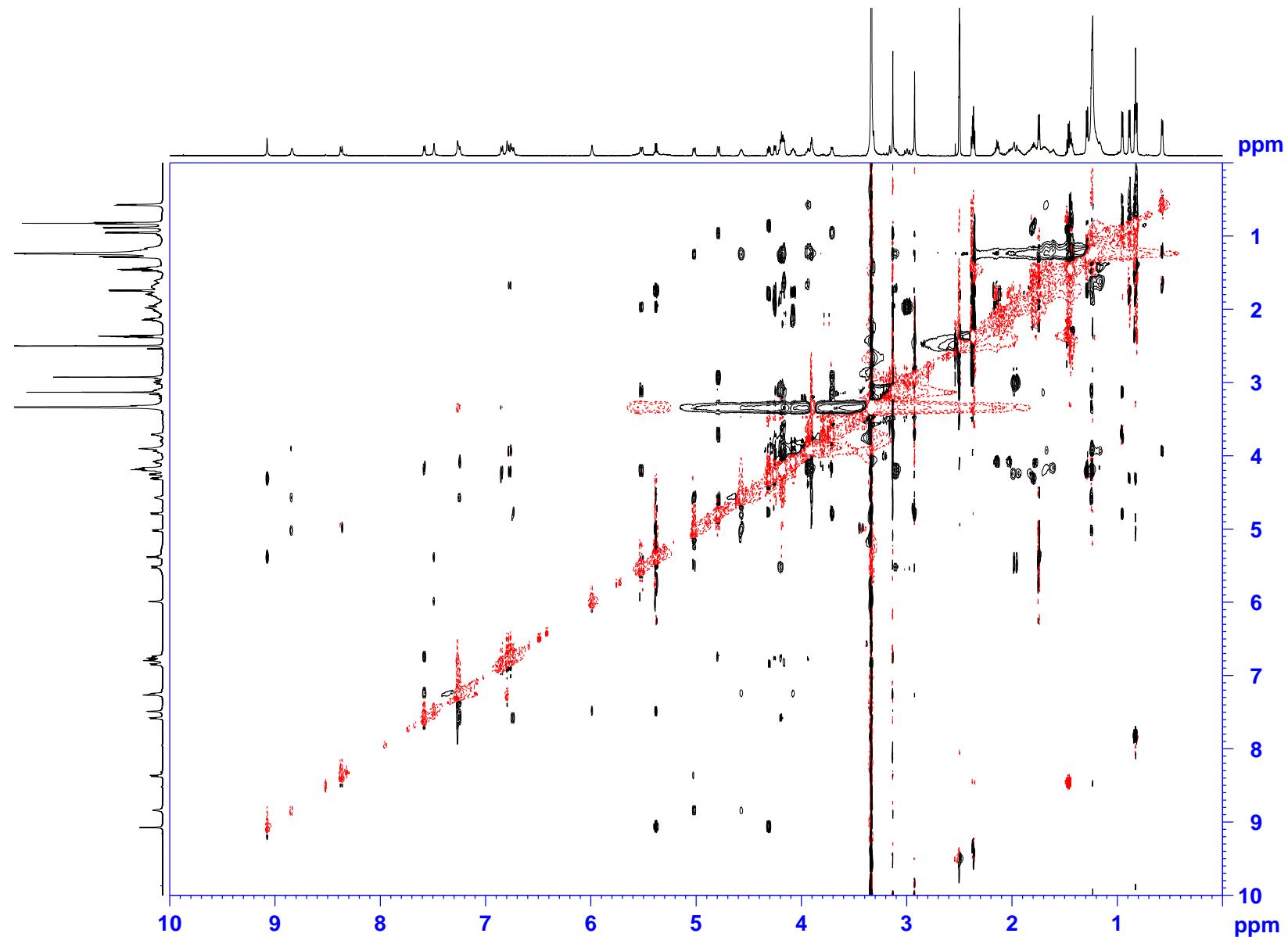
S10. HSQC spectrum (600 MHz, DMSO-*d*₆) of **1**



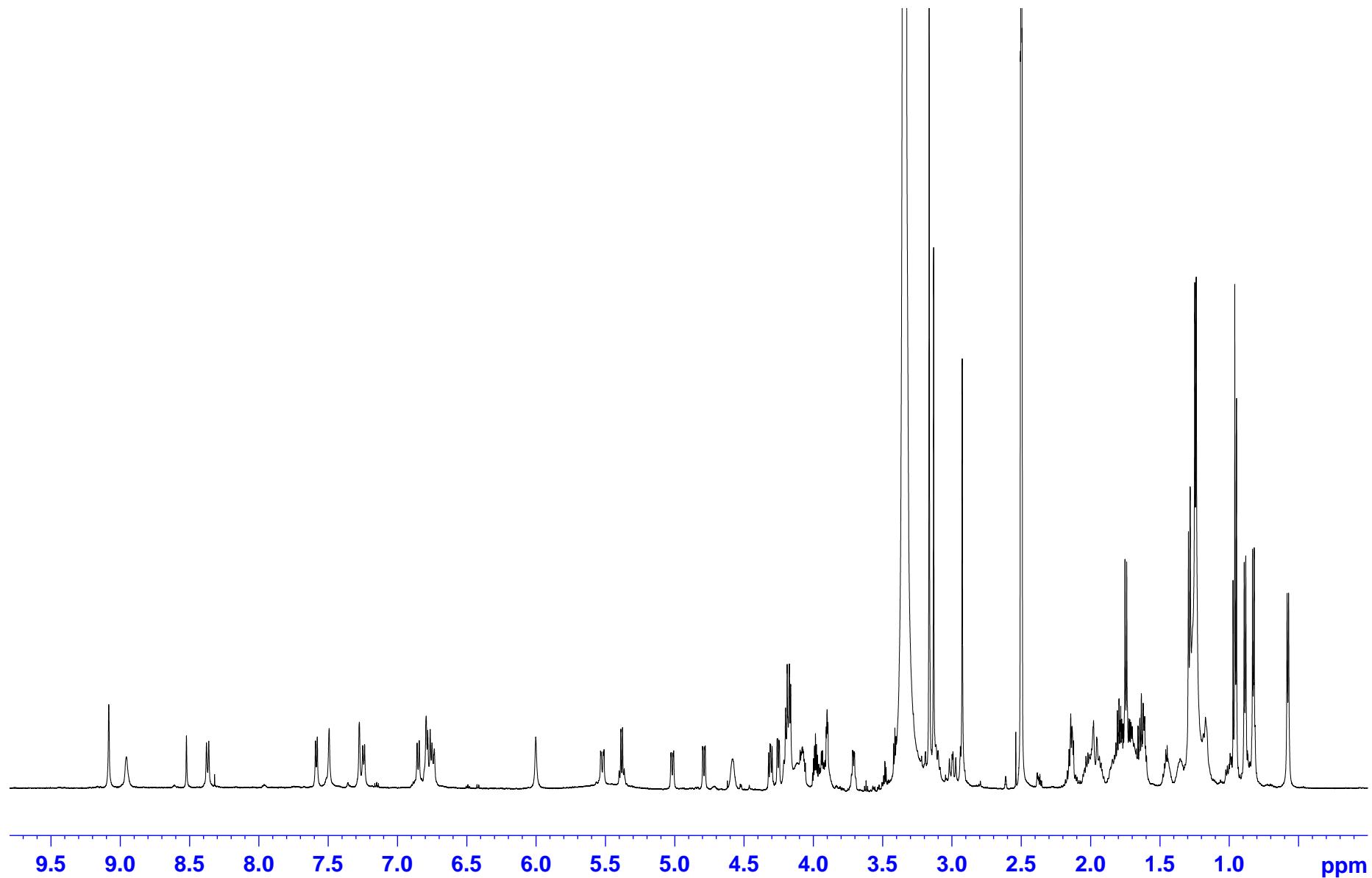
S11. HMBC spectrum (600 MHz, DMSO-*d*₆) of **1**



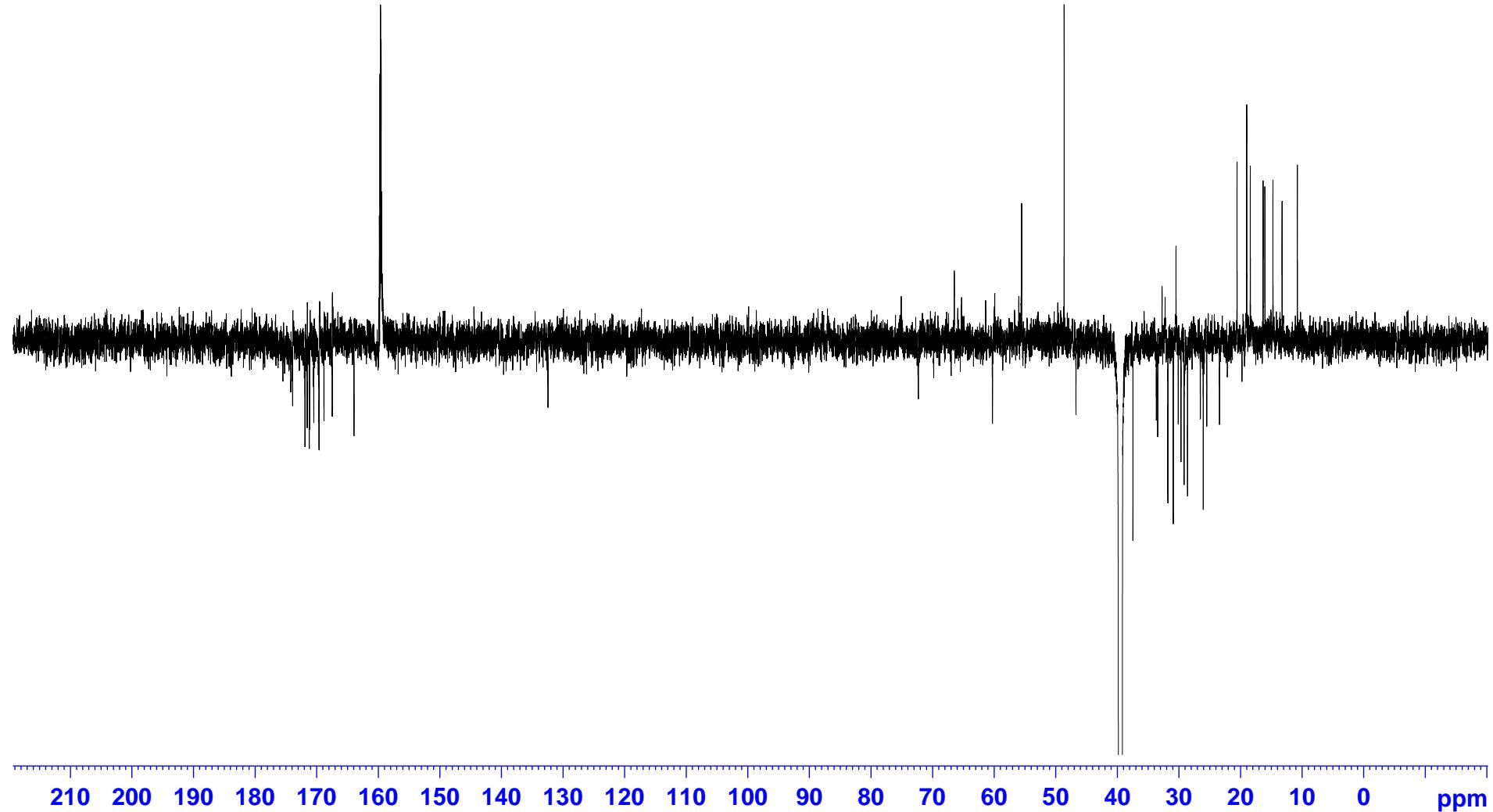
S12. T-ROESY spectrum (600 MHz, DMSO-*d*₆) of **1**



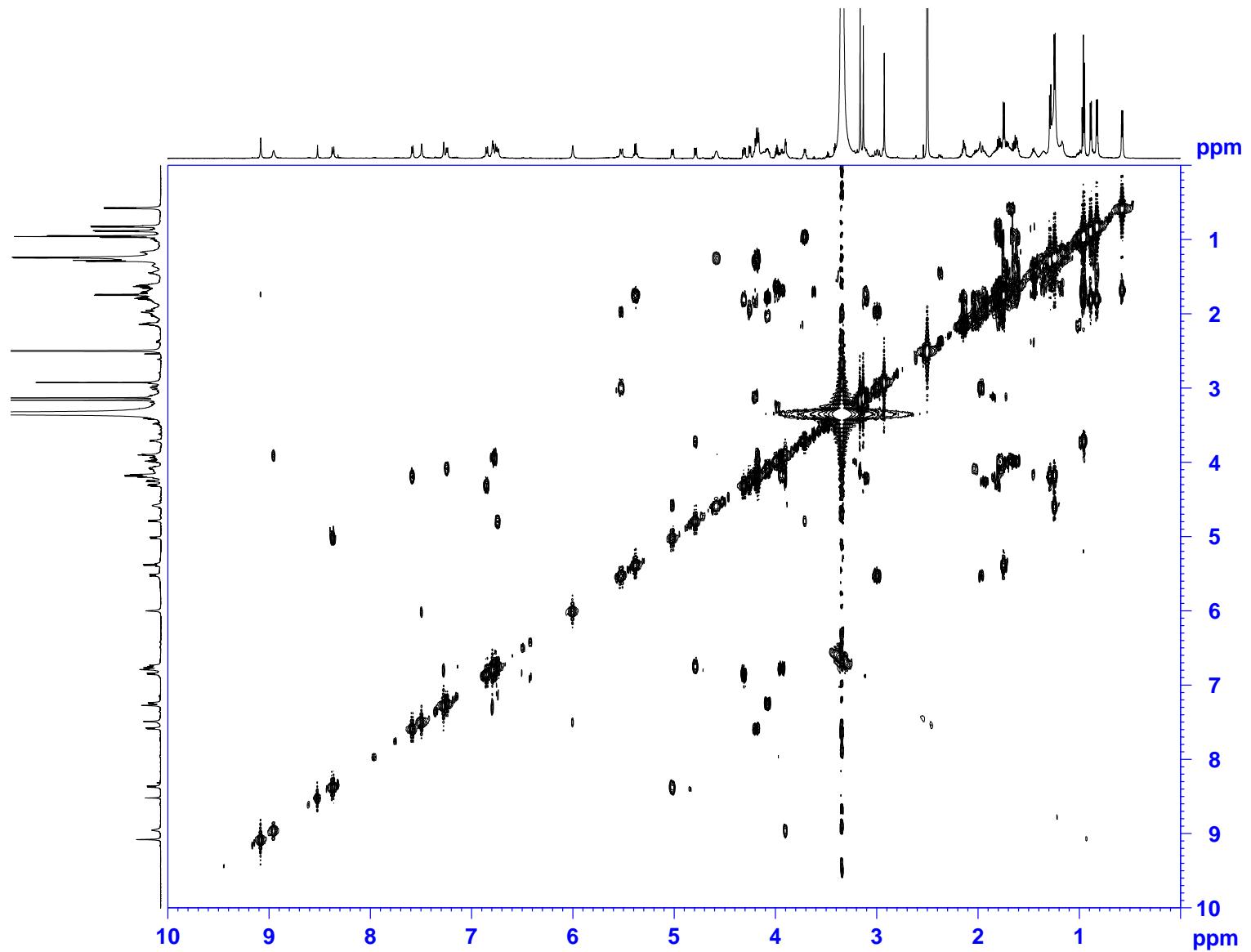
S13. ^1H NMR spectrum (600 MHz, DMSO- d_6) of **2**



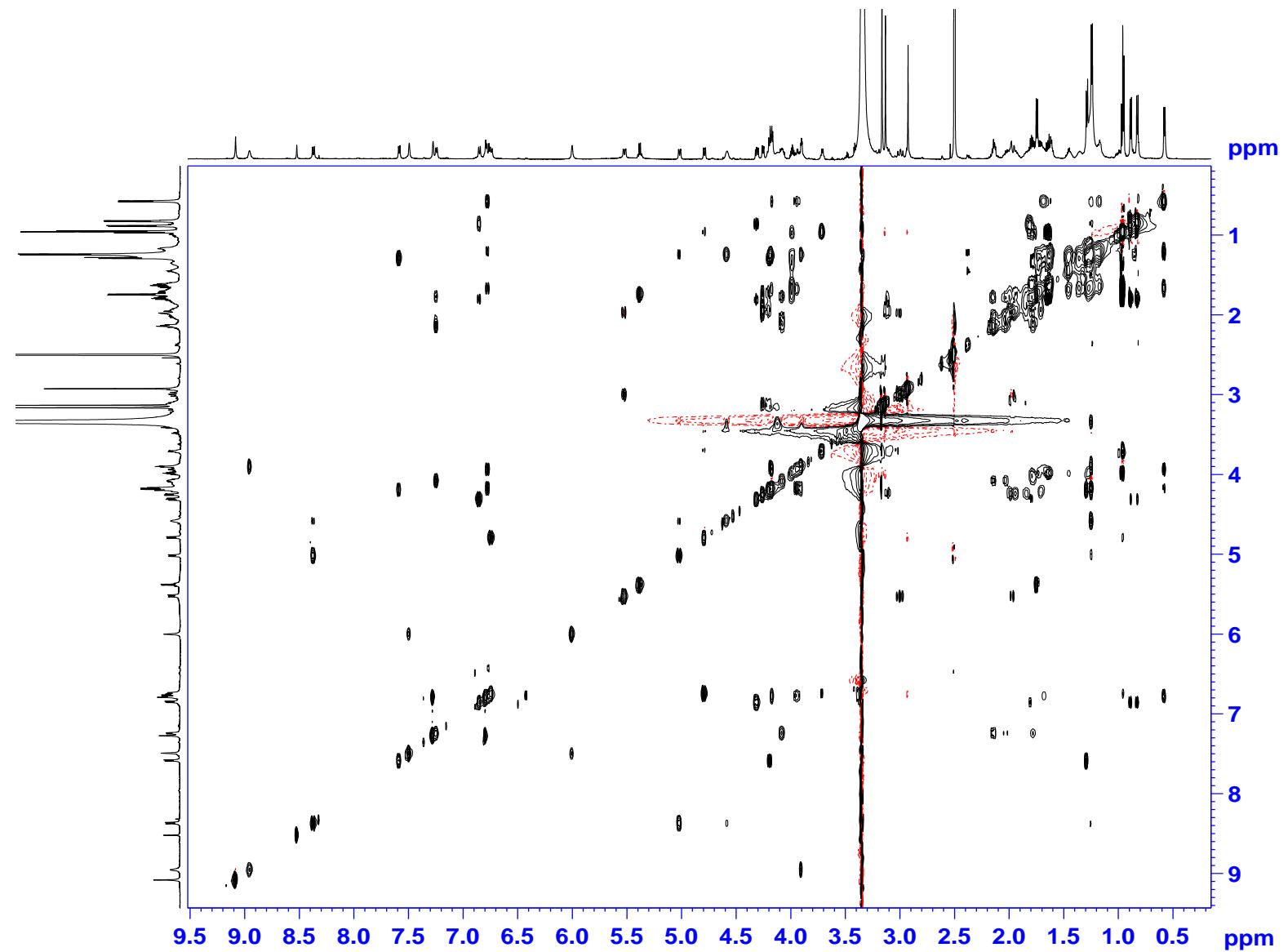
S14. DEPTQ spectrum (226 MHz, DMSO-*d*₆) of **2**



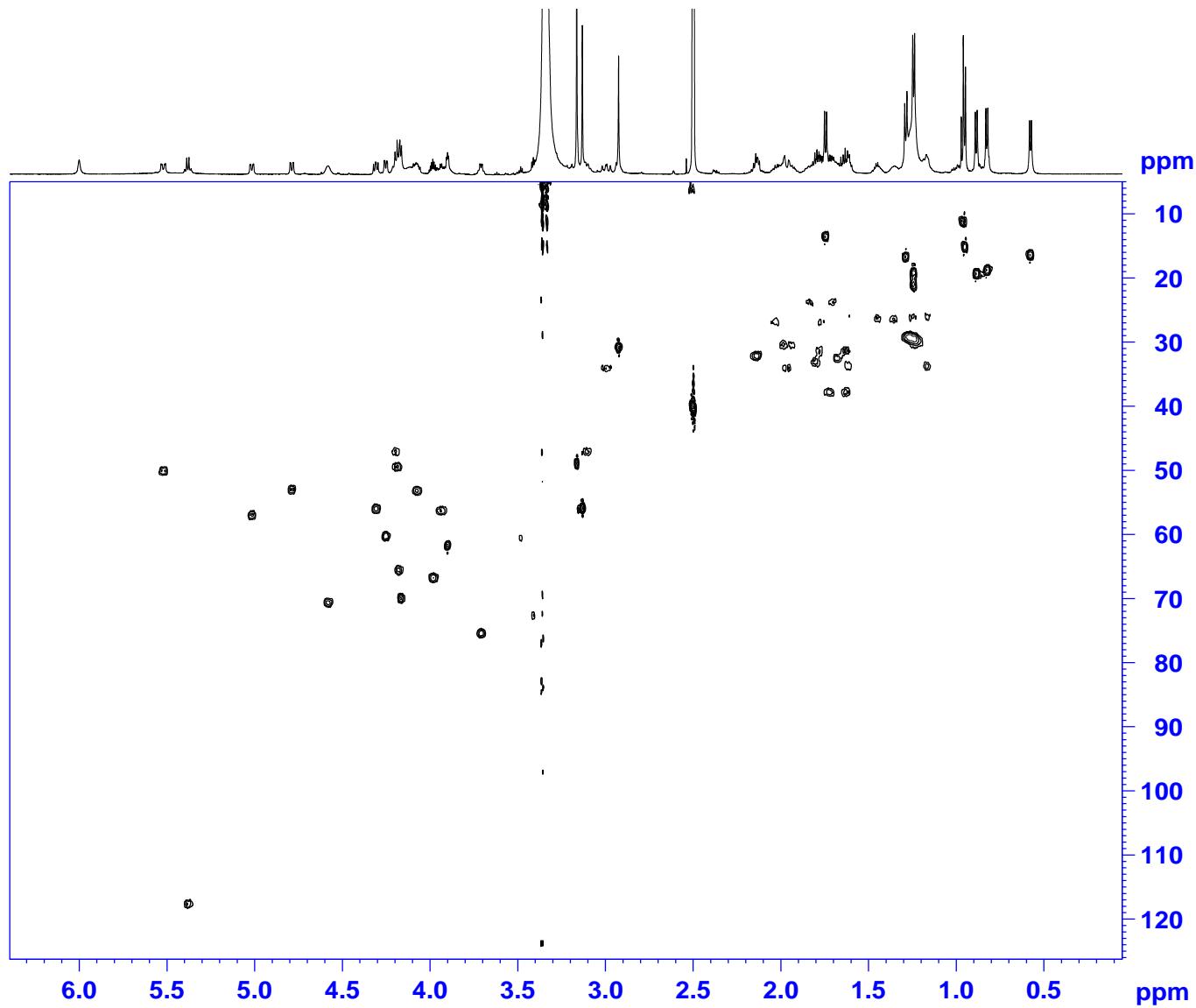
S15. COSY spectrum (600 MHz, DMSO-*d*₆) of **2**



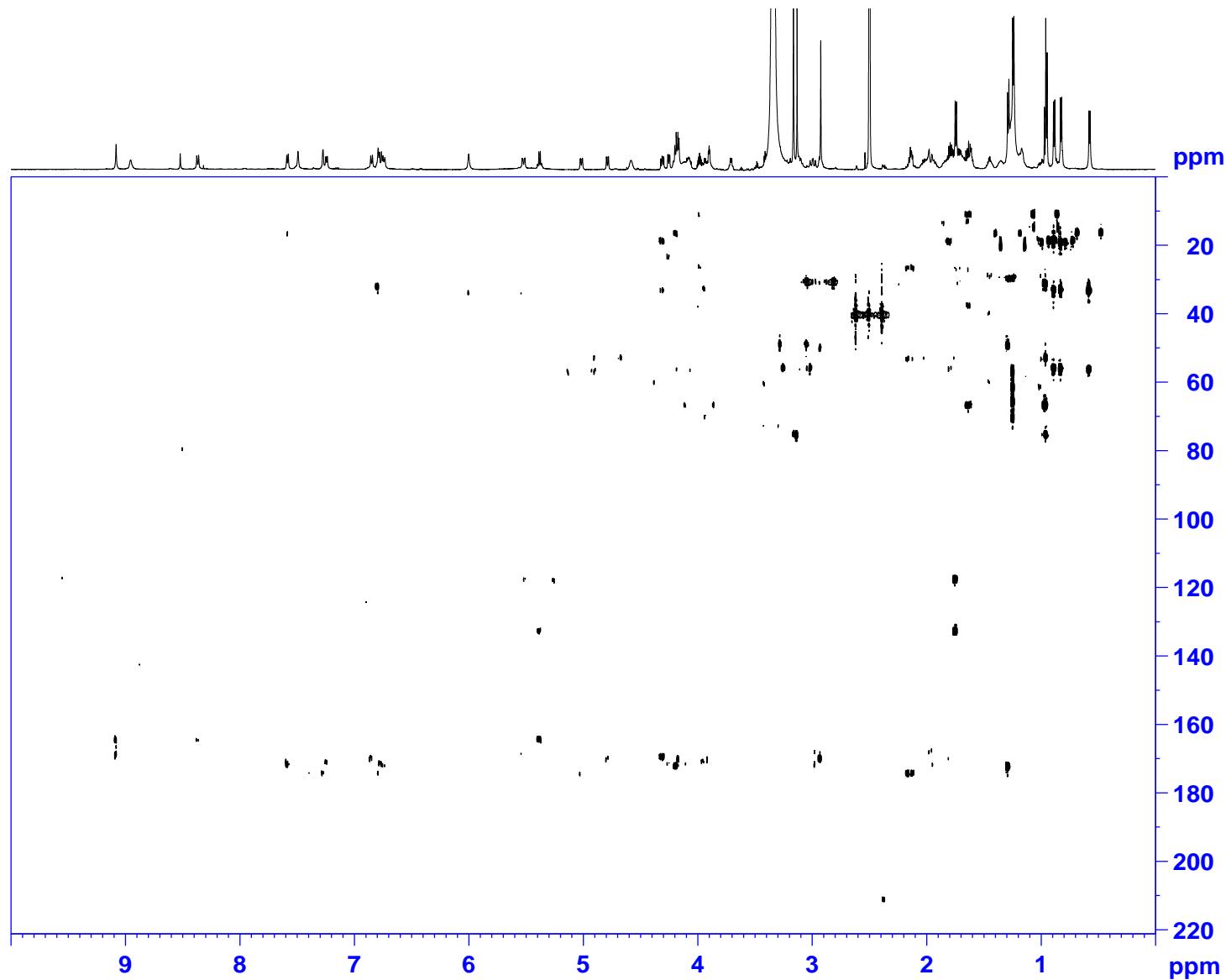
S16. TOCSY spectrum (600 MHz, DMSO-*d*₆) of 2



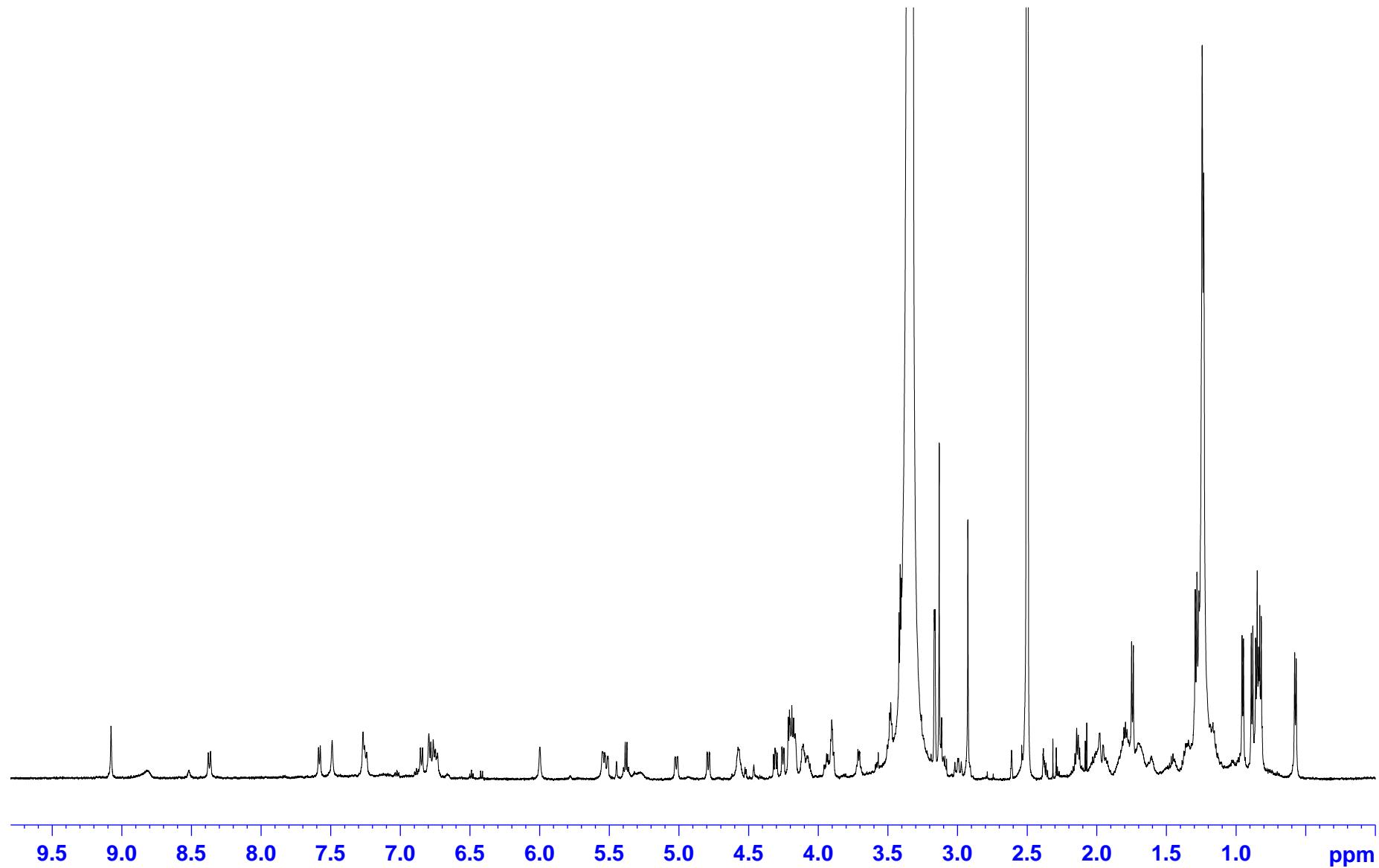
S17. HSQC spectrum (600 MHz, DMSO-*d*₆) of **2**



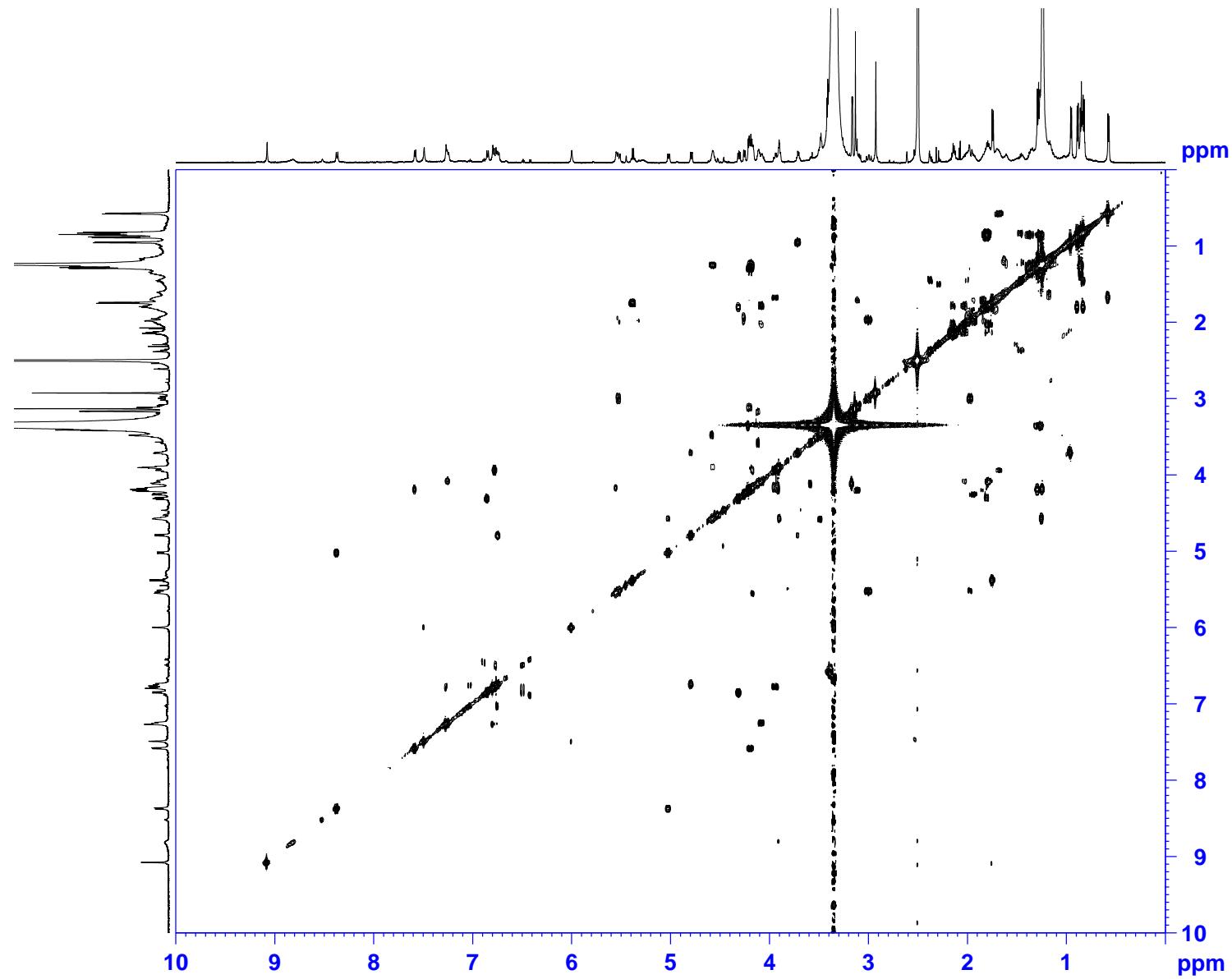
S18. HMBC spectrum (600 MHz, DMSO-*d*₆) of **2**



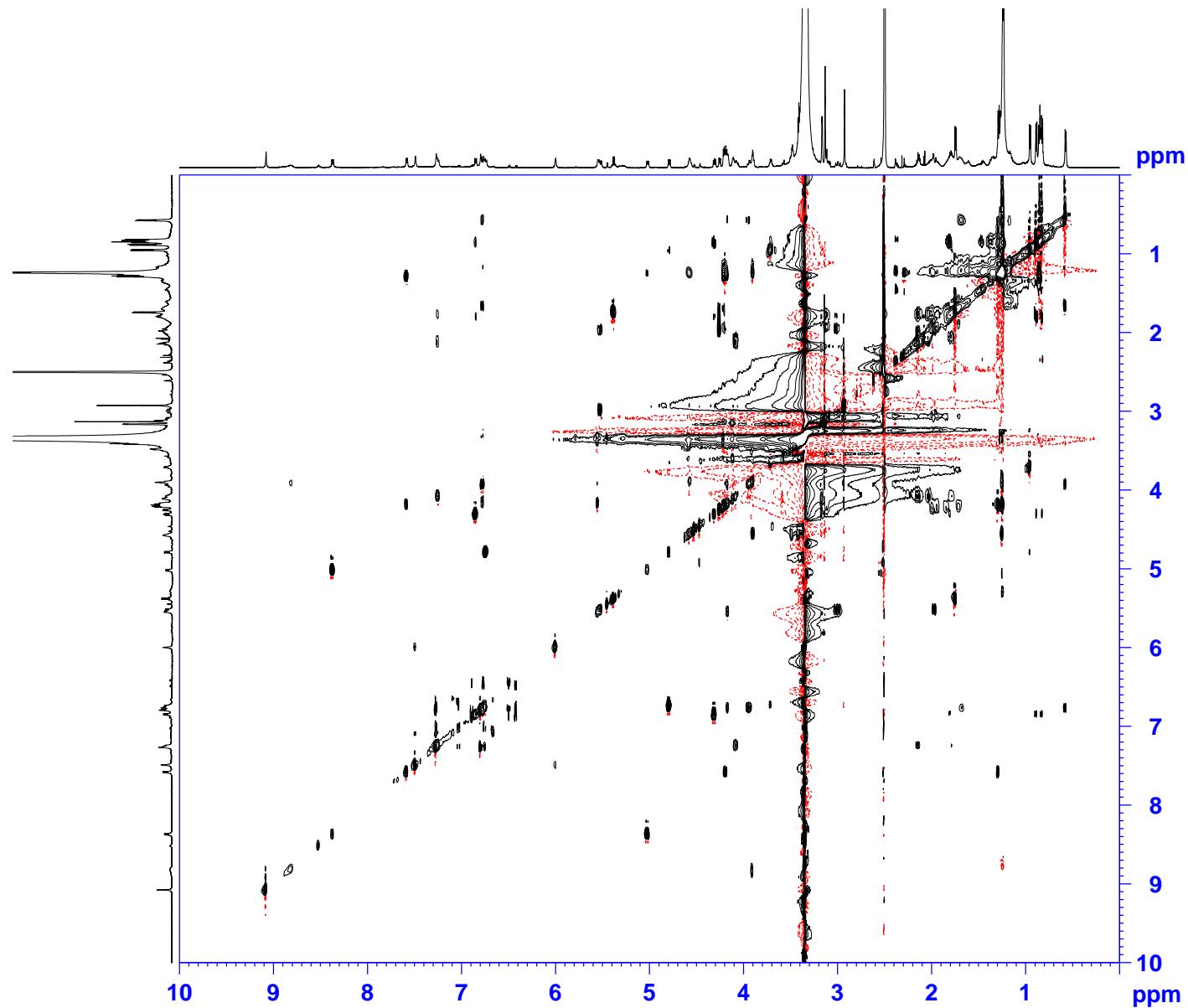
S19. ^1H NMR spectrum (600 MHz, DMSO- d_6) of **3**



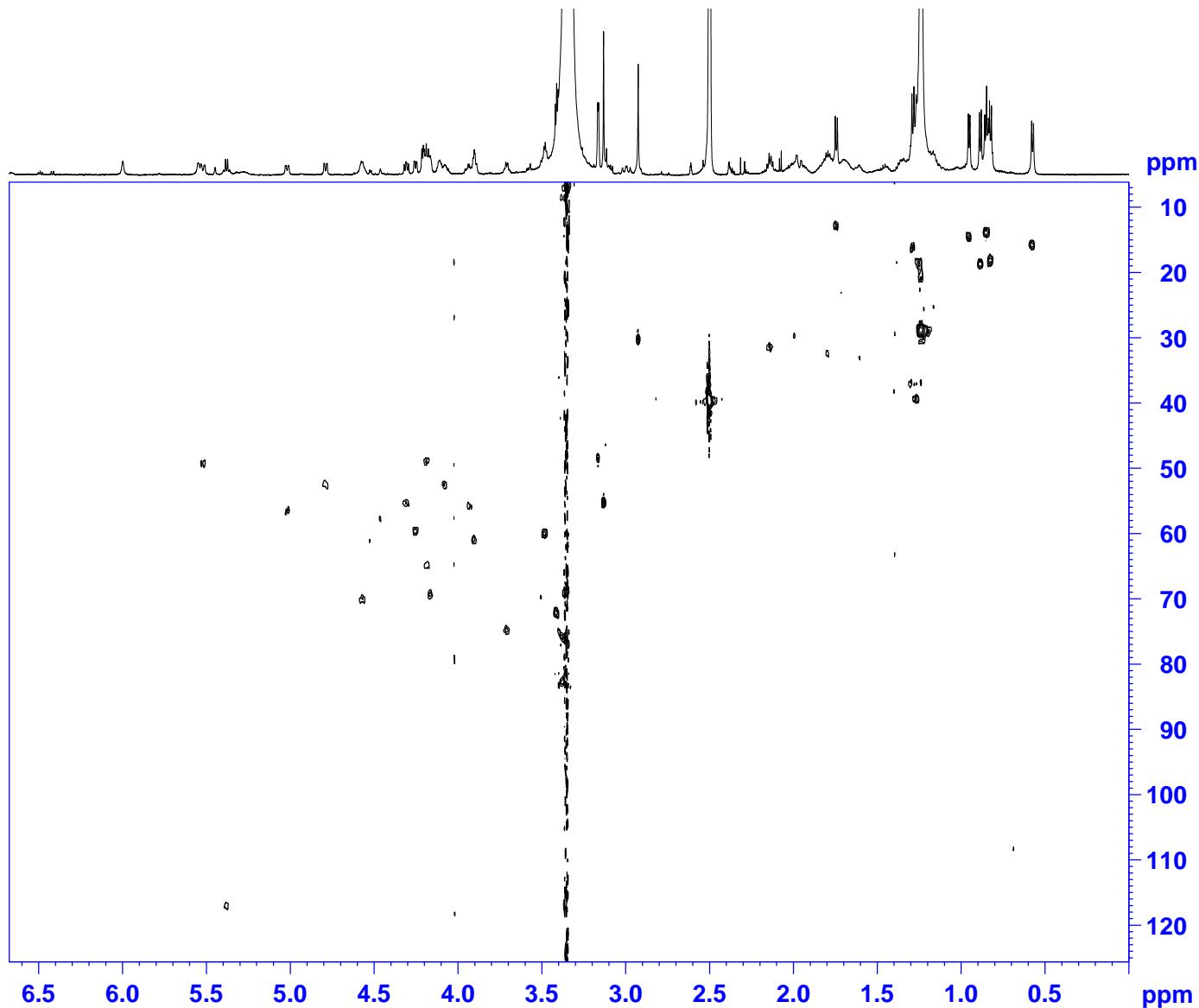
S20. COSY spectrum (600 MHz, DMSO-*d*₆) of **3**



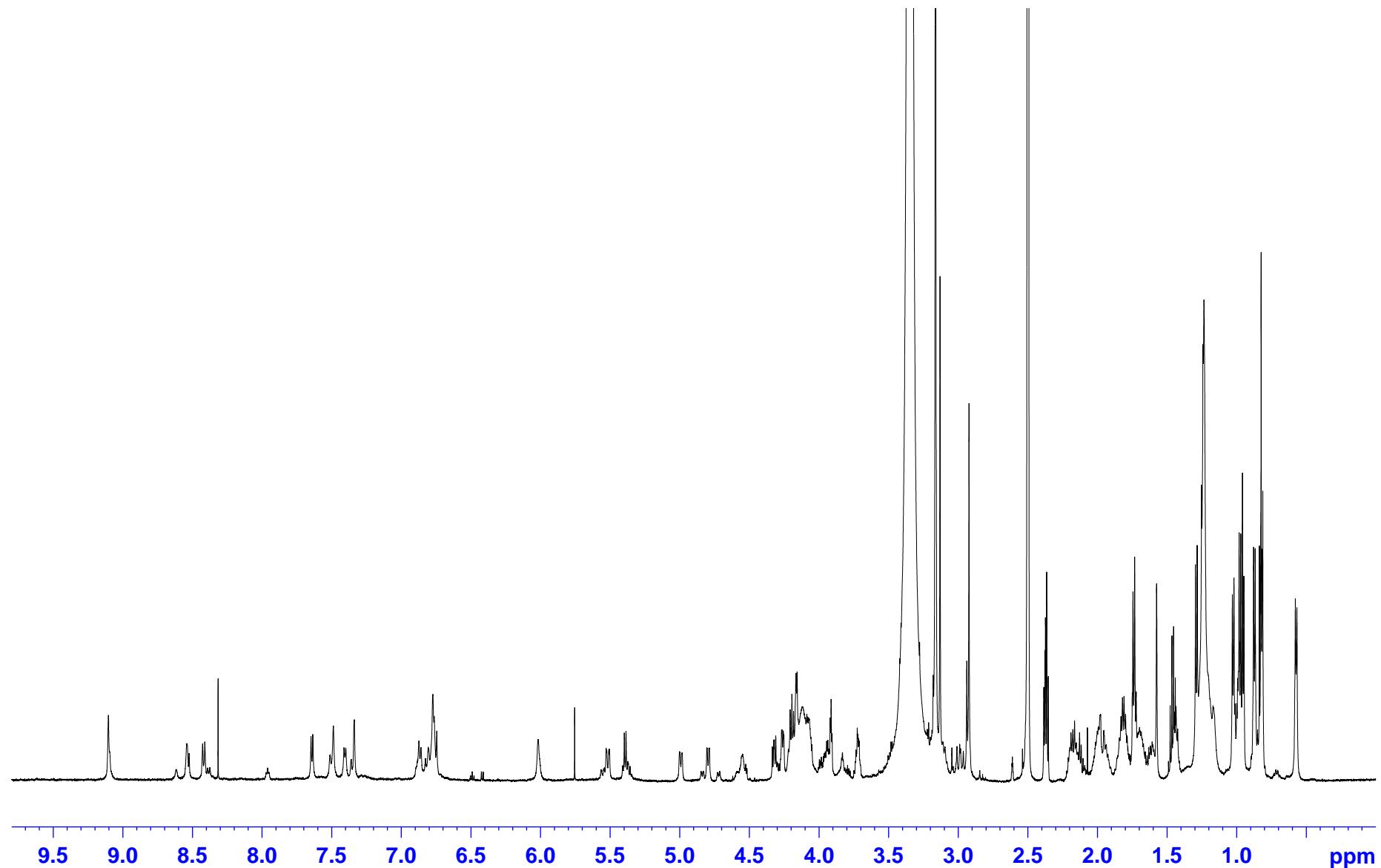
S21. TOCSY spectrum (600 MHz, DMSO-*d*₆) of **3**



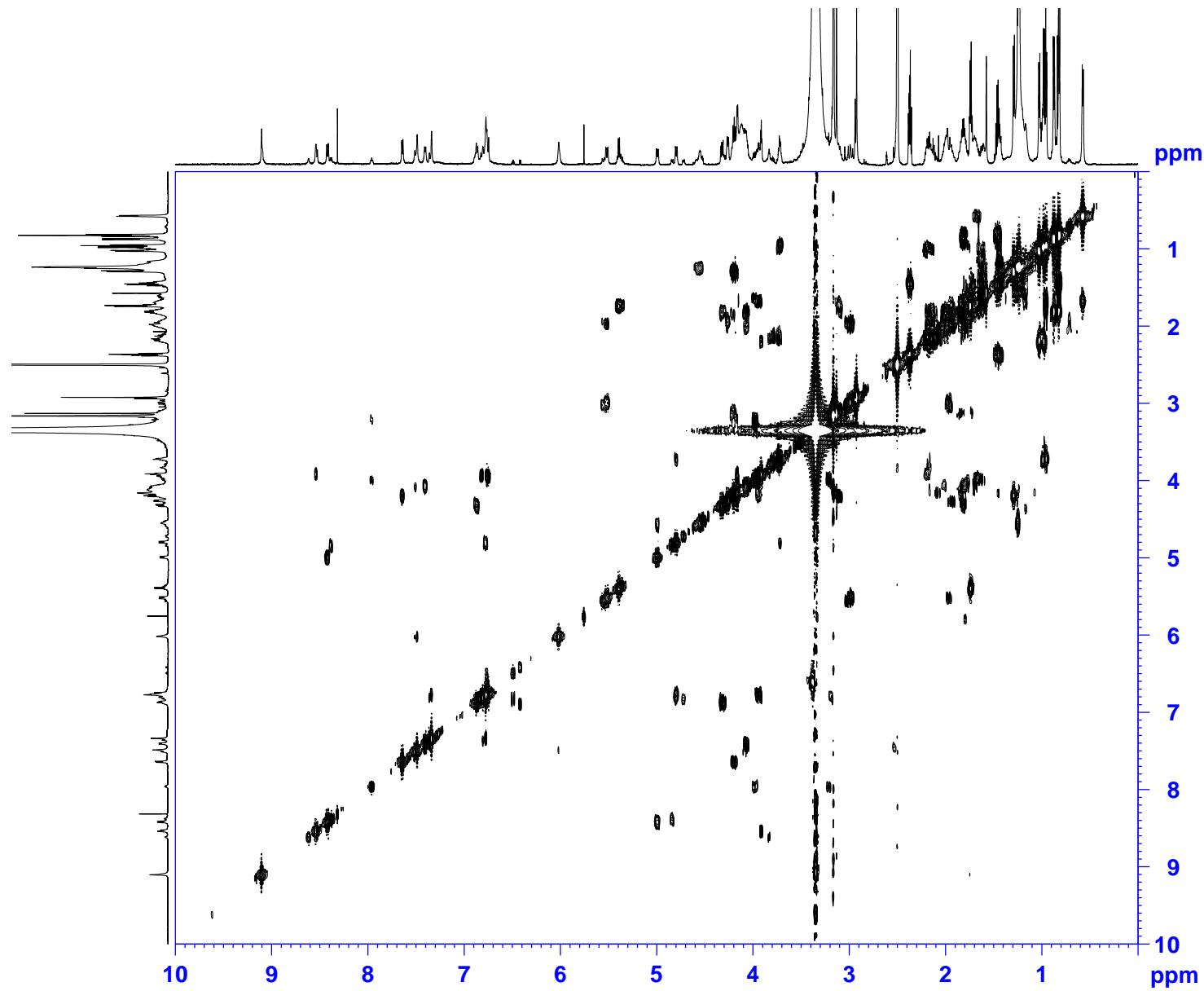
S22. HSQC spectrum (600 MHz, DMSO-*d*₆) of **3**



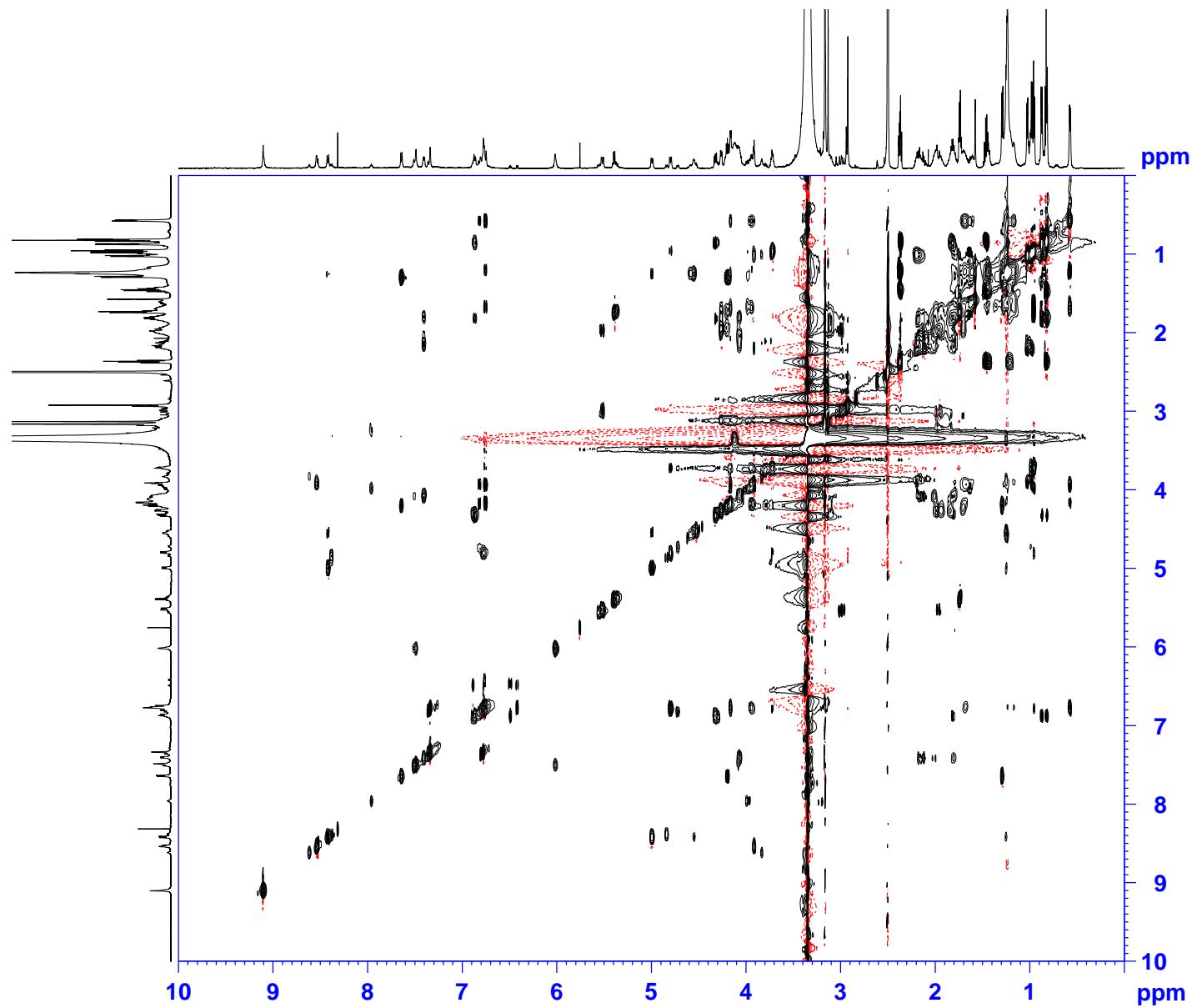
S23. ^1H NMR spectrum (600 MHz, DMSO- d_6) of **4**



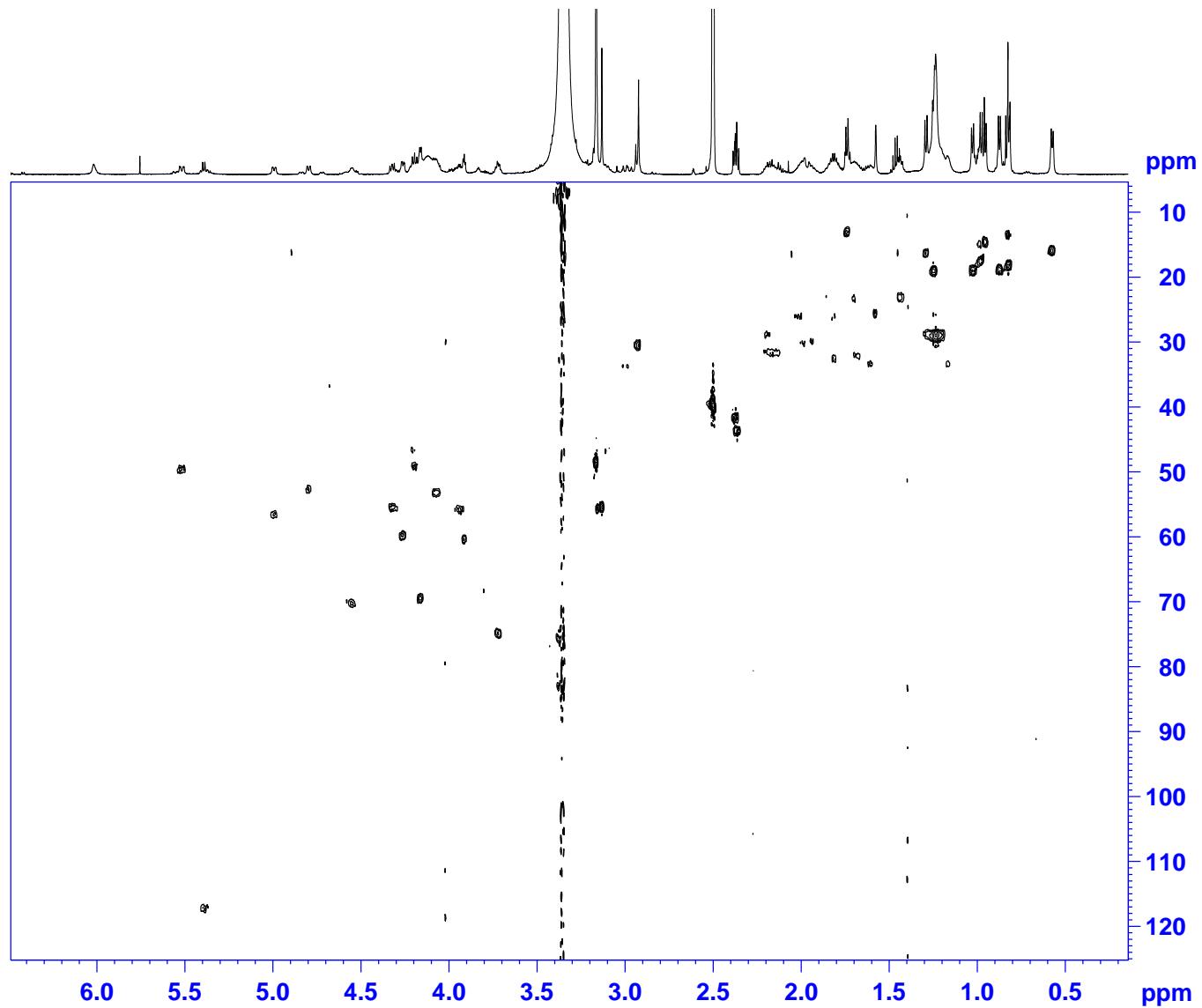
S24. COSY spectrum (600 MHz, DMSO-*d*₆) of **4**



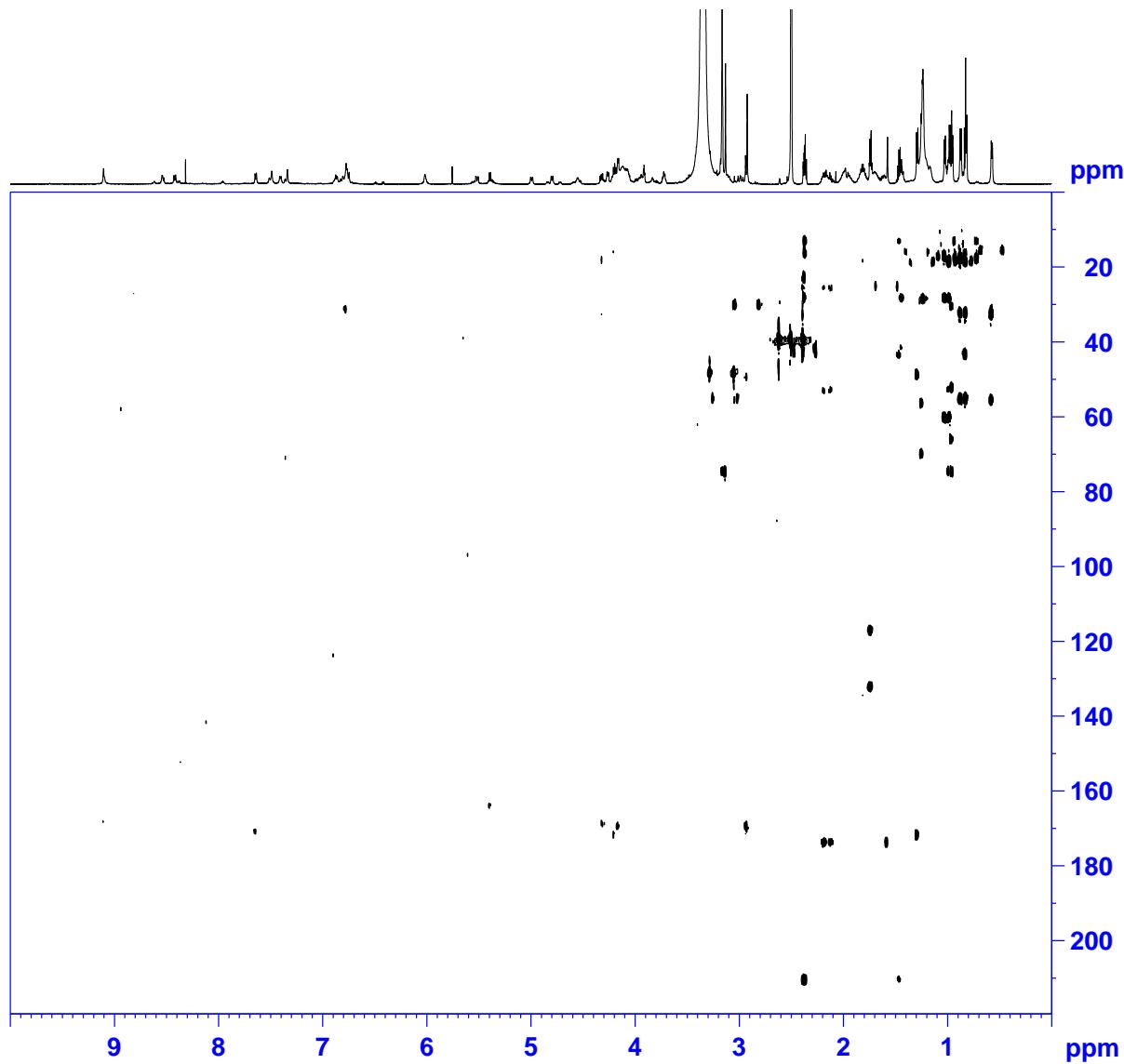
S25. TOCSY spectrum (600 MHz, DMSO-*d*₆) of **4**



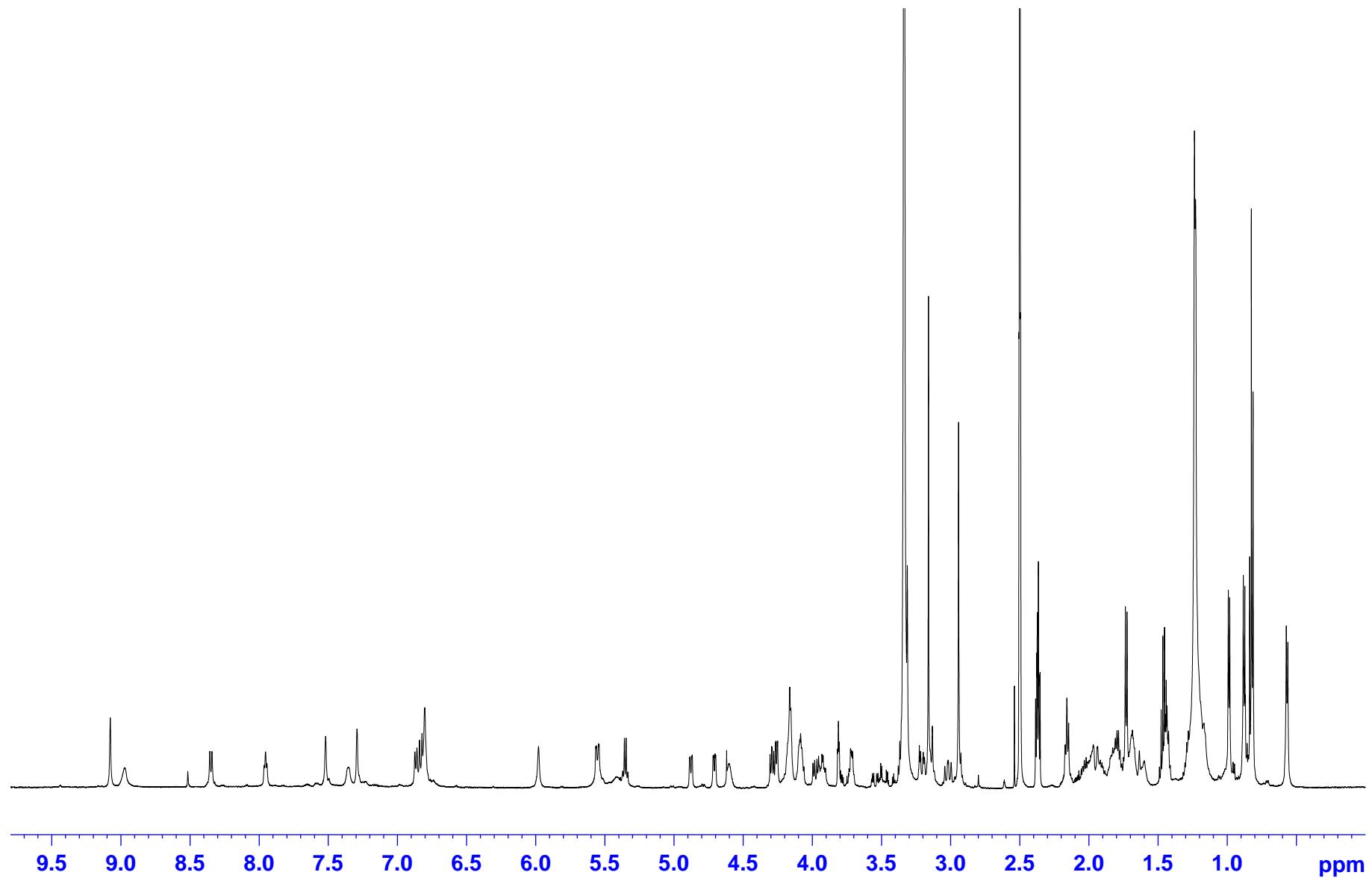
S26. HSQC spectrum (600 MHz, DMSO-*d*₆) of **4**



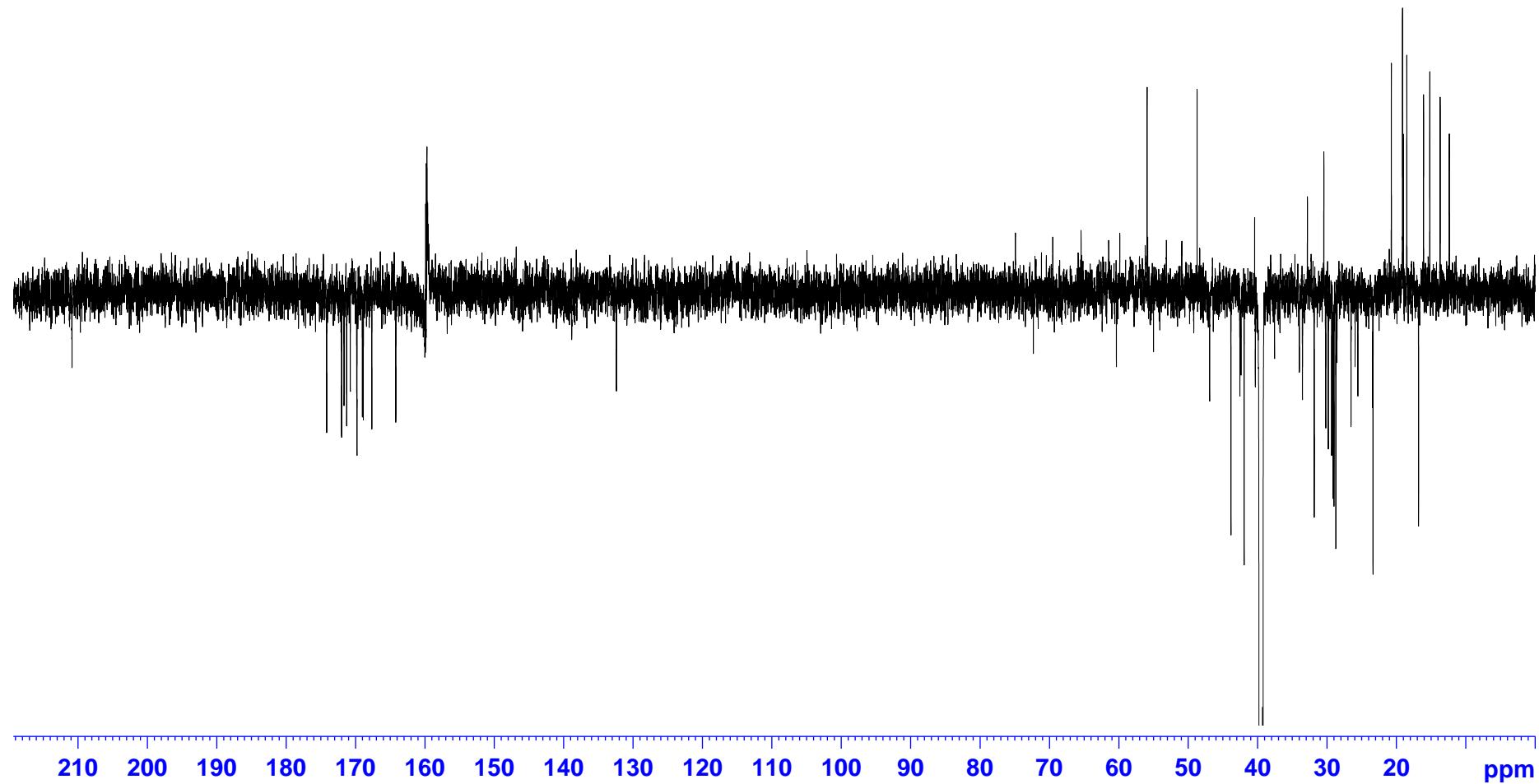
S27. HMBC spectrum (600 MHz, DMSO-*d*₆) of **4**



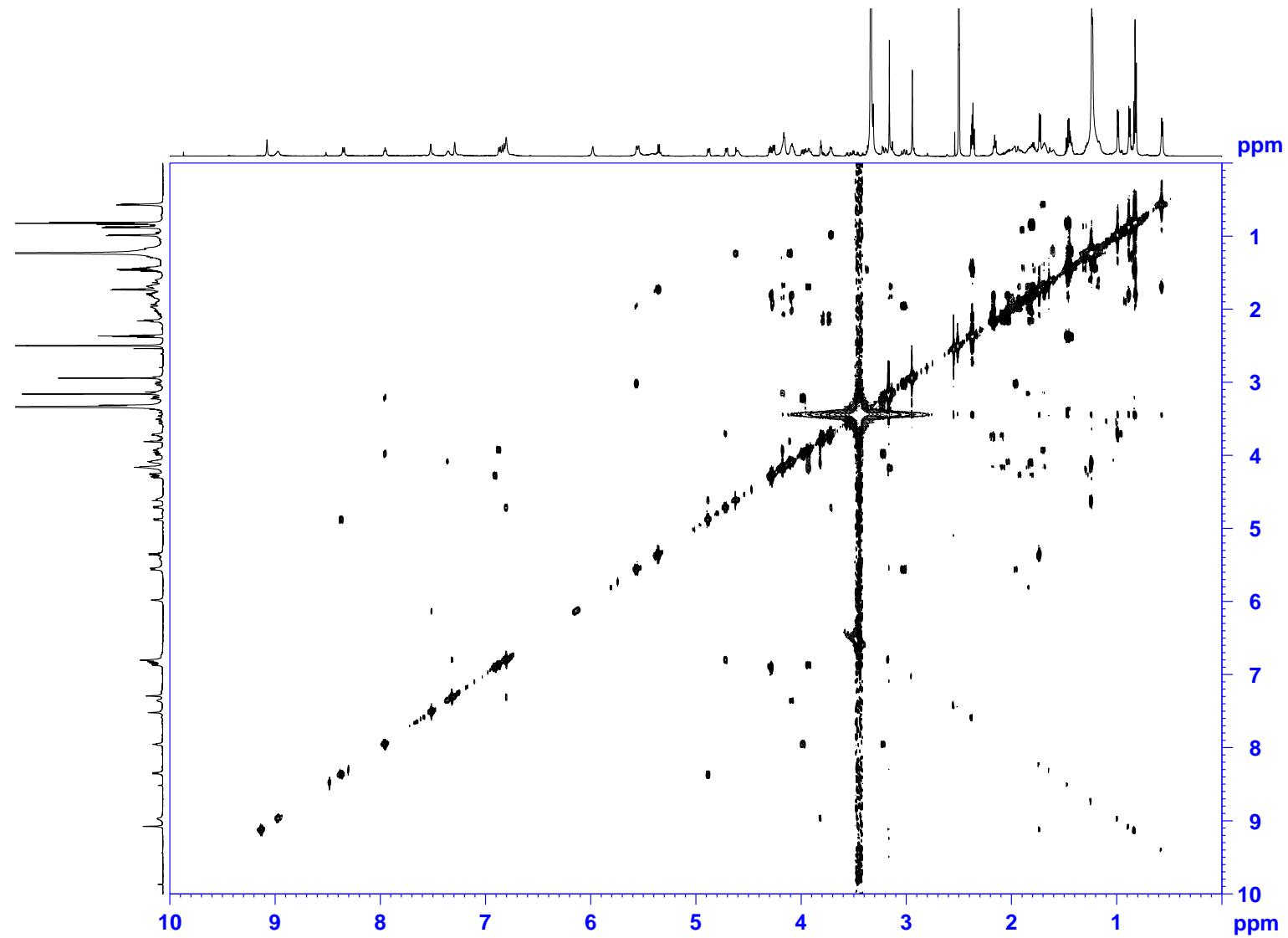
S28. ^1H NMR spectrum (600 MHz, DMSO- d_6) of **5**



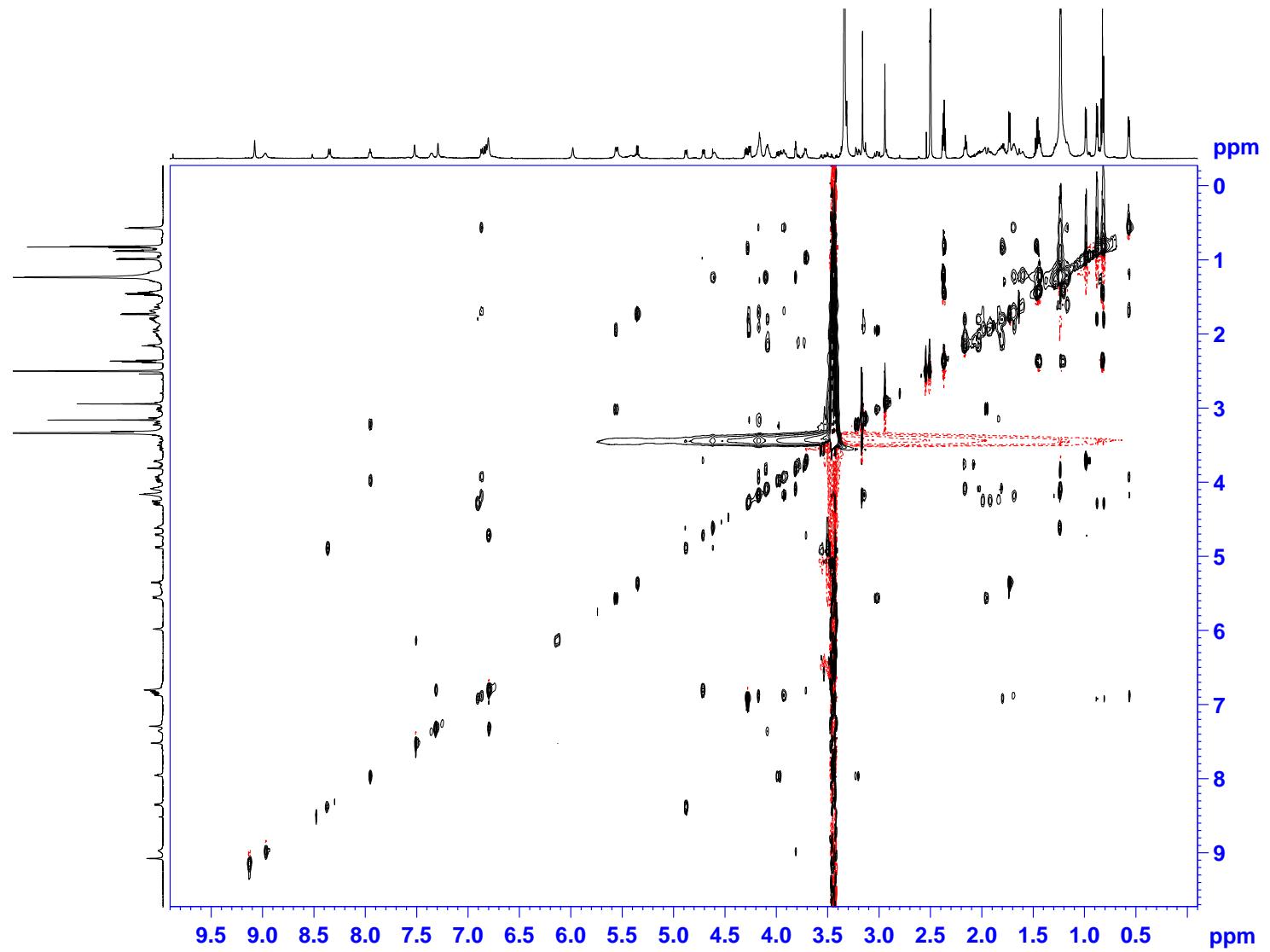
S29. DEPTQ spectrum (226 MHz, DMSO-*d*₆) of **5**



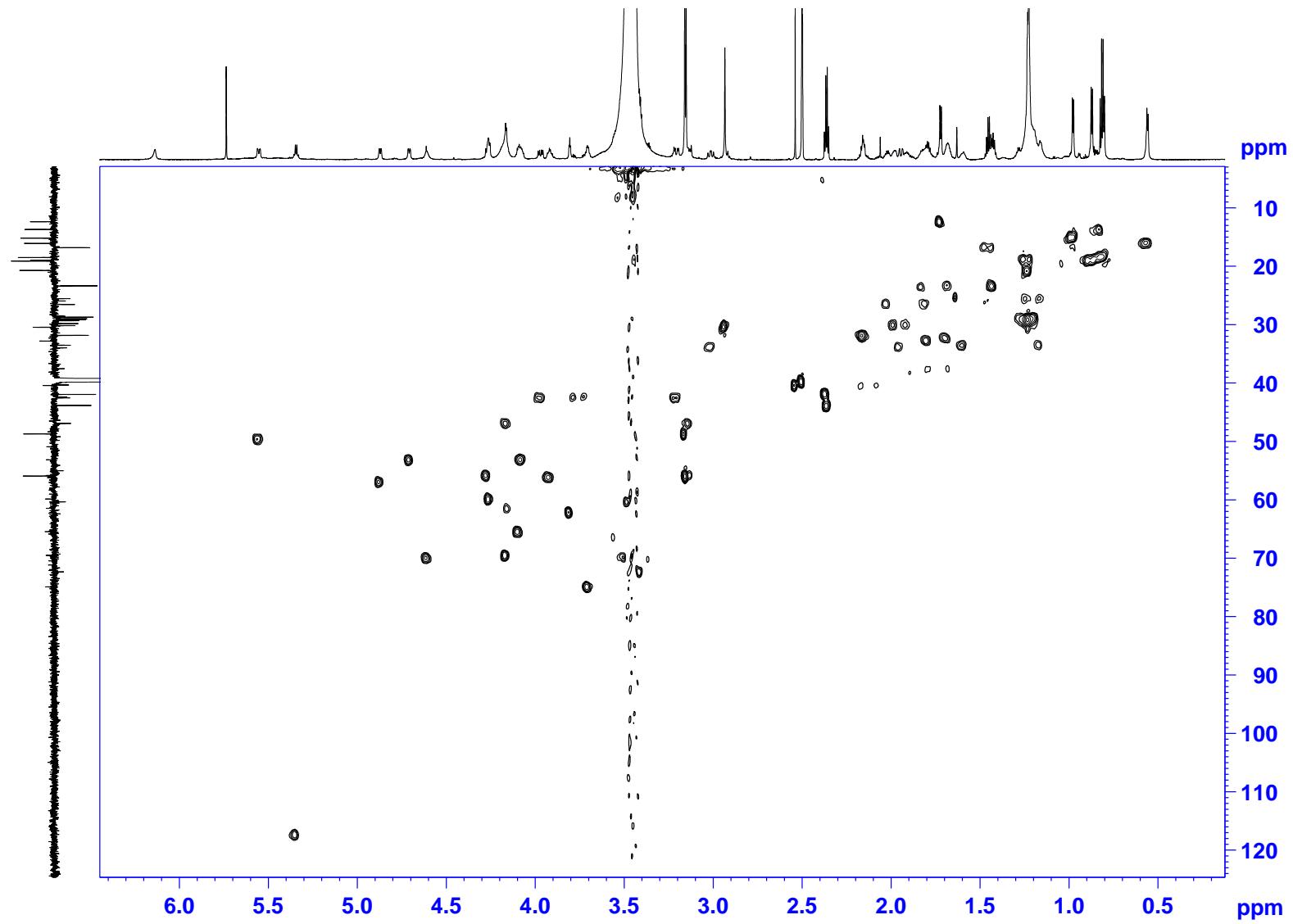
S30. COSY spectrum (600 MHz, DMSO-*d*₆) of **5**



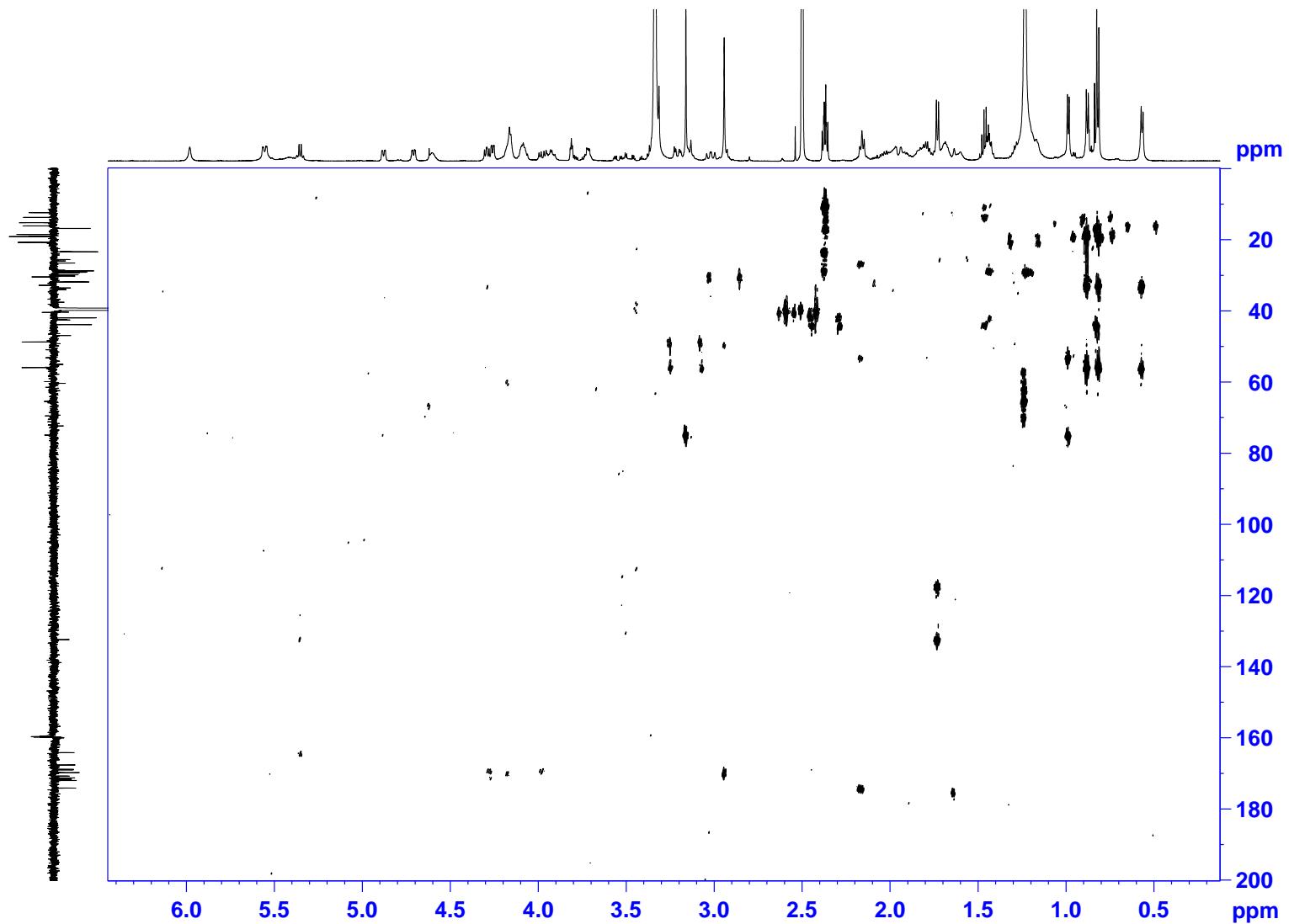
S31. TOCSY spectrum (600 MHz, DMSO-*d*₆) of **5**



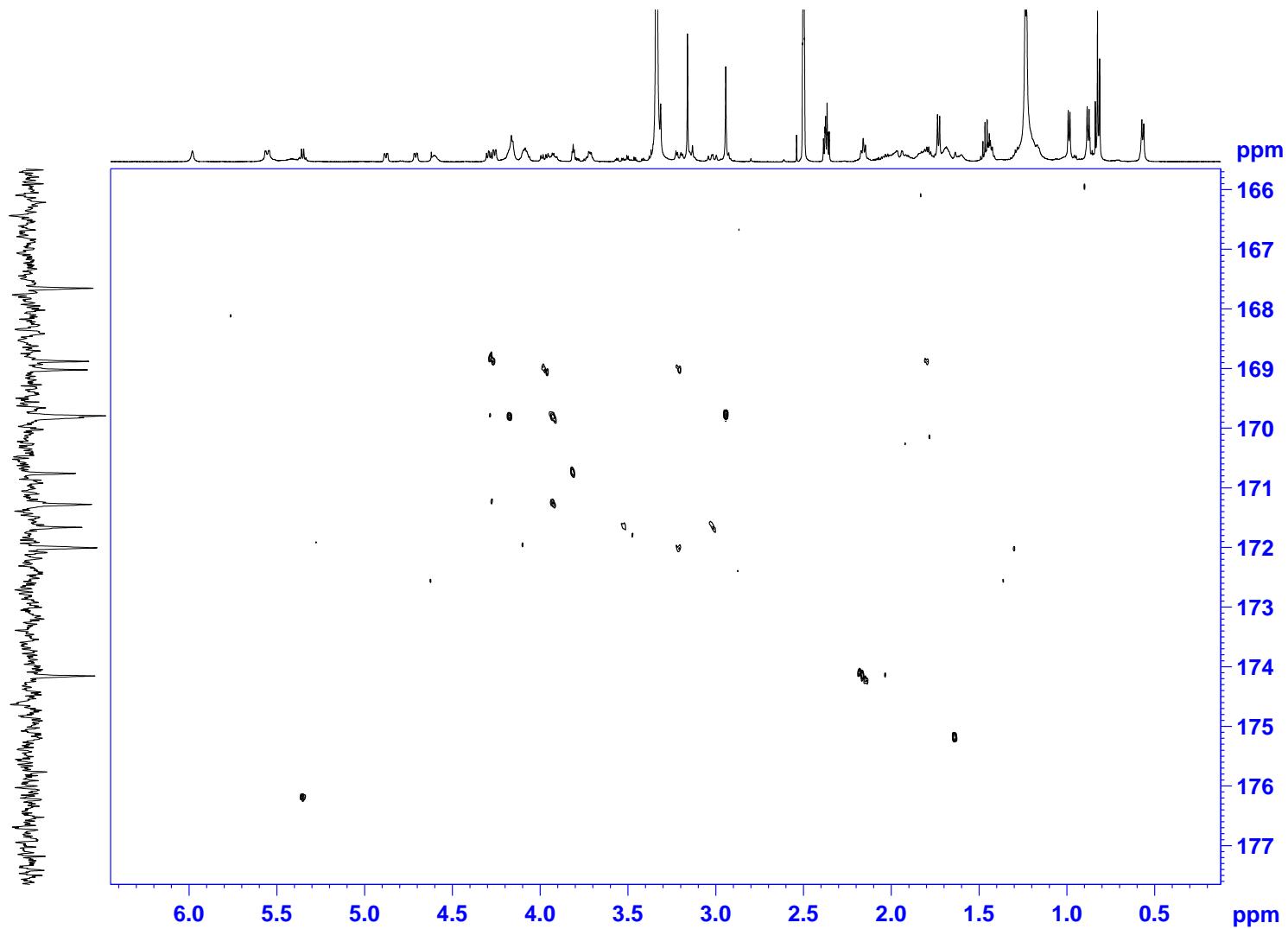
S32. HSQC spectrum (600 MHz, DMSO-*d*₆) of **5**



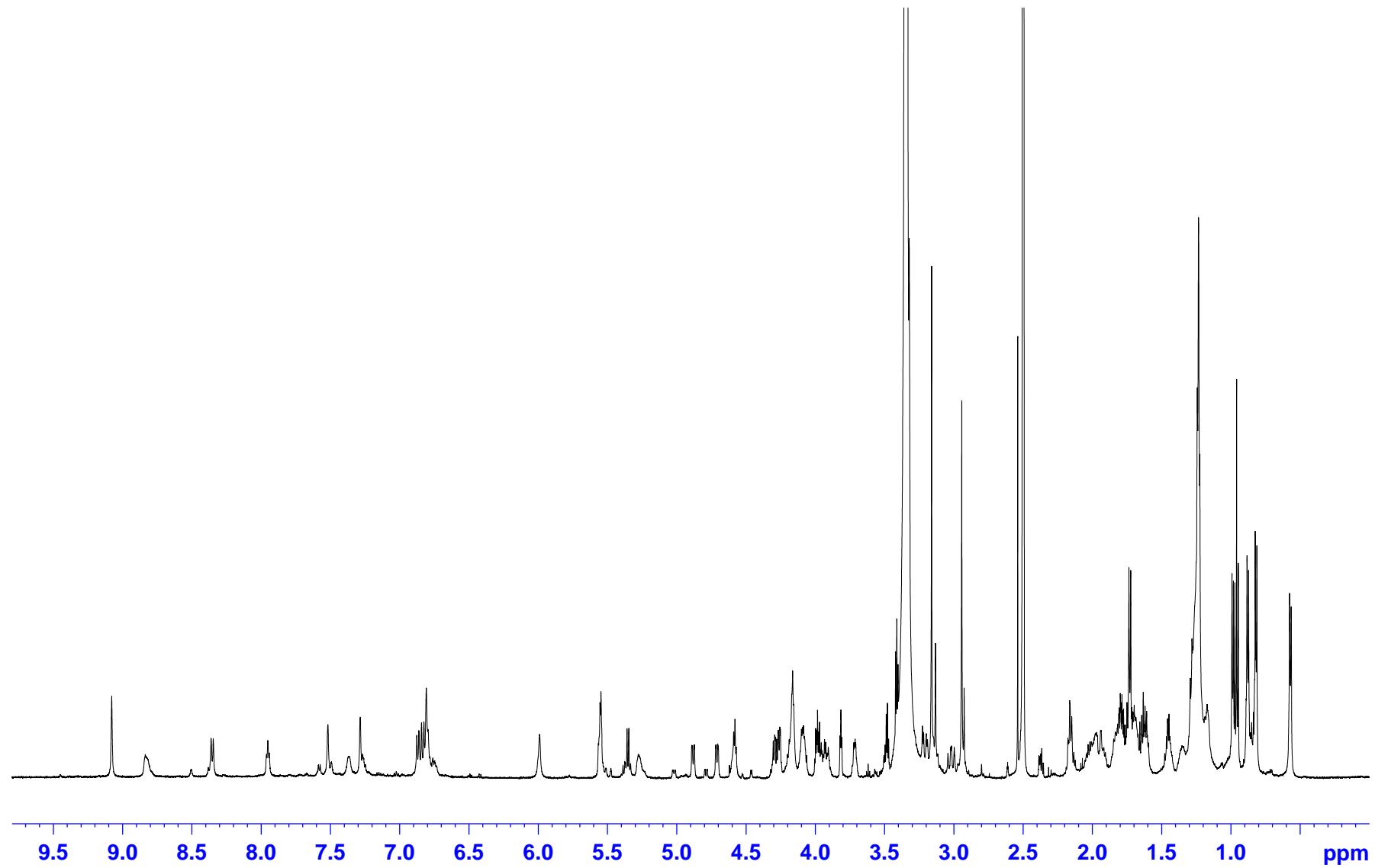
S33. HMBC spectrum (600 MHz, DMSO-*d*₆) of **5**



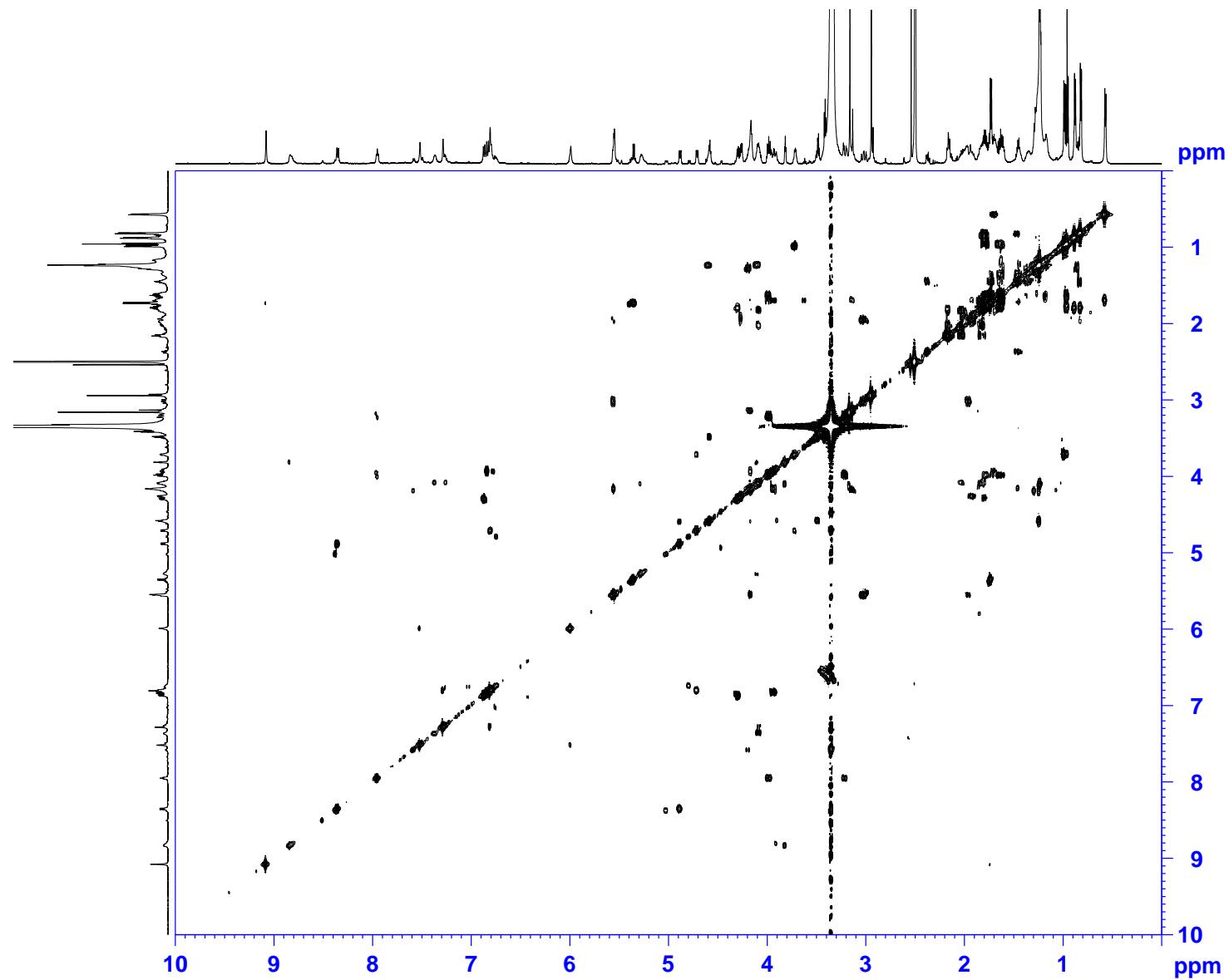
S34. Selective HMBC spectrum (900 MHz, DMSO-*d*₆) of **5**



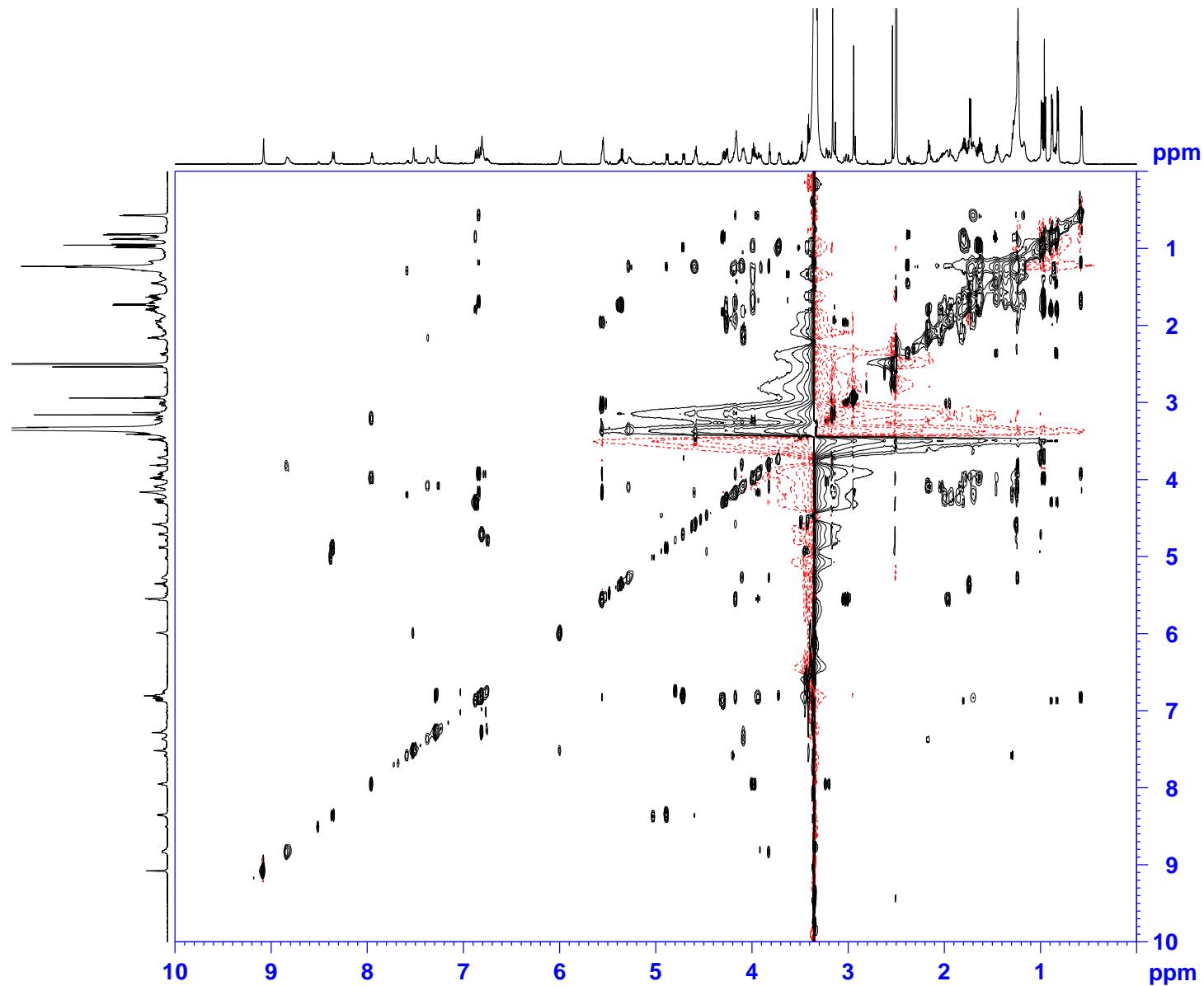
S35. ^1H NMR spectrum (600 MHz, DMSO- d_6) of **6**



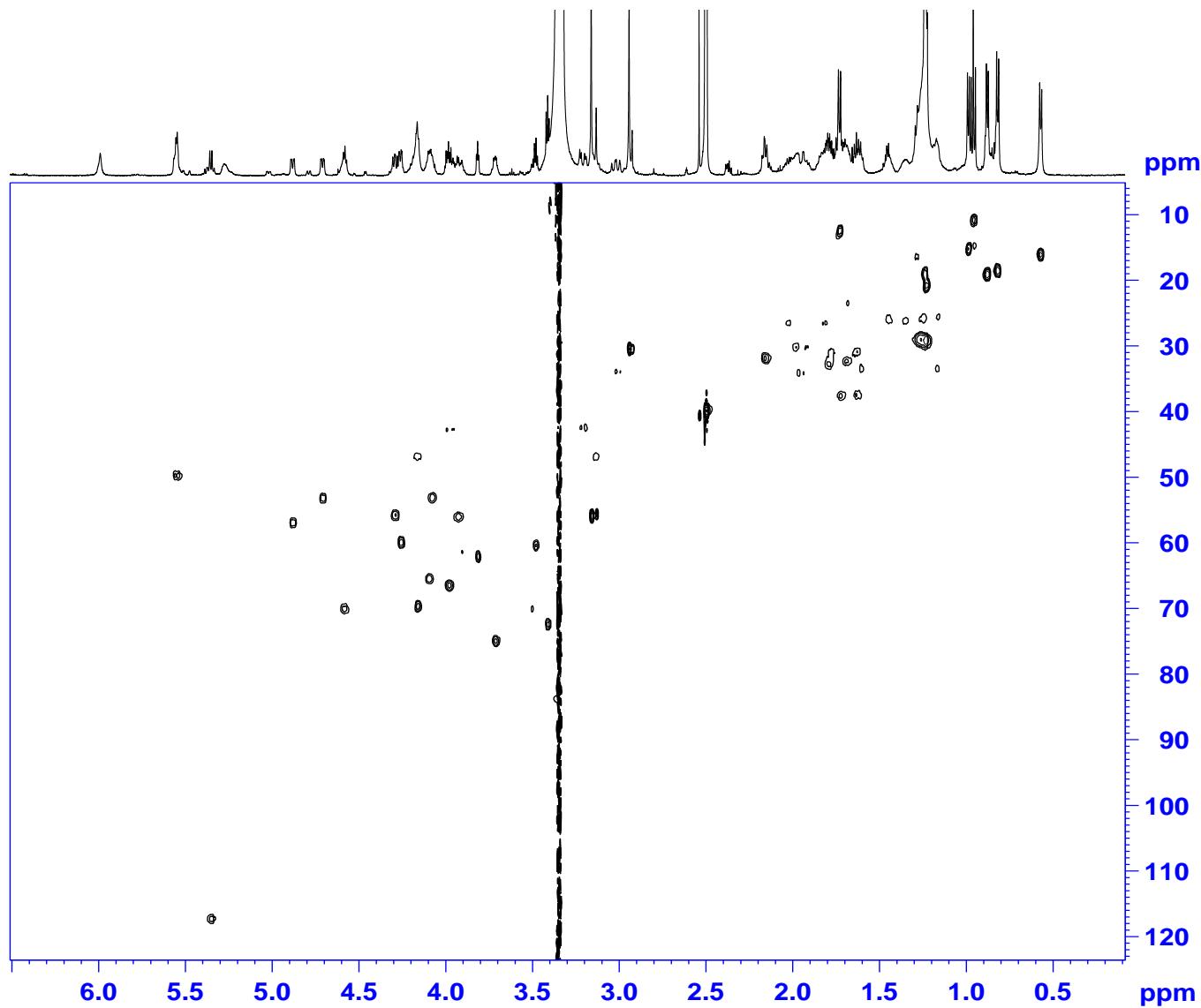
S36. COSY spectrum (600 MHz, DMSO-*d*₆) of **6**



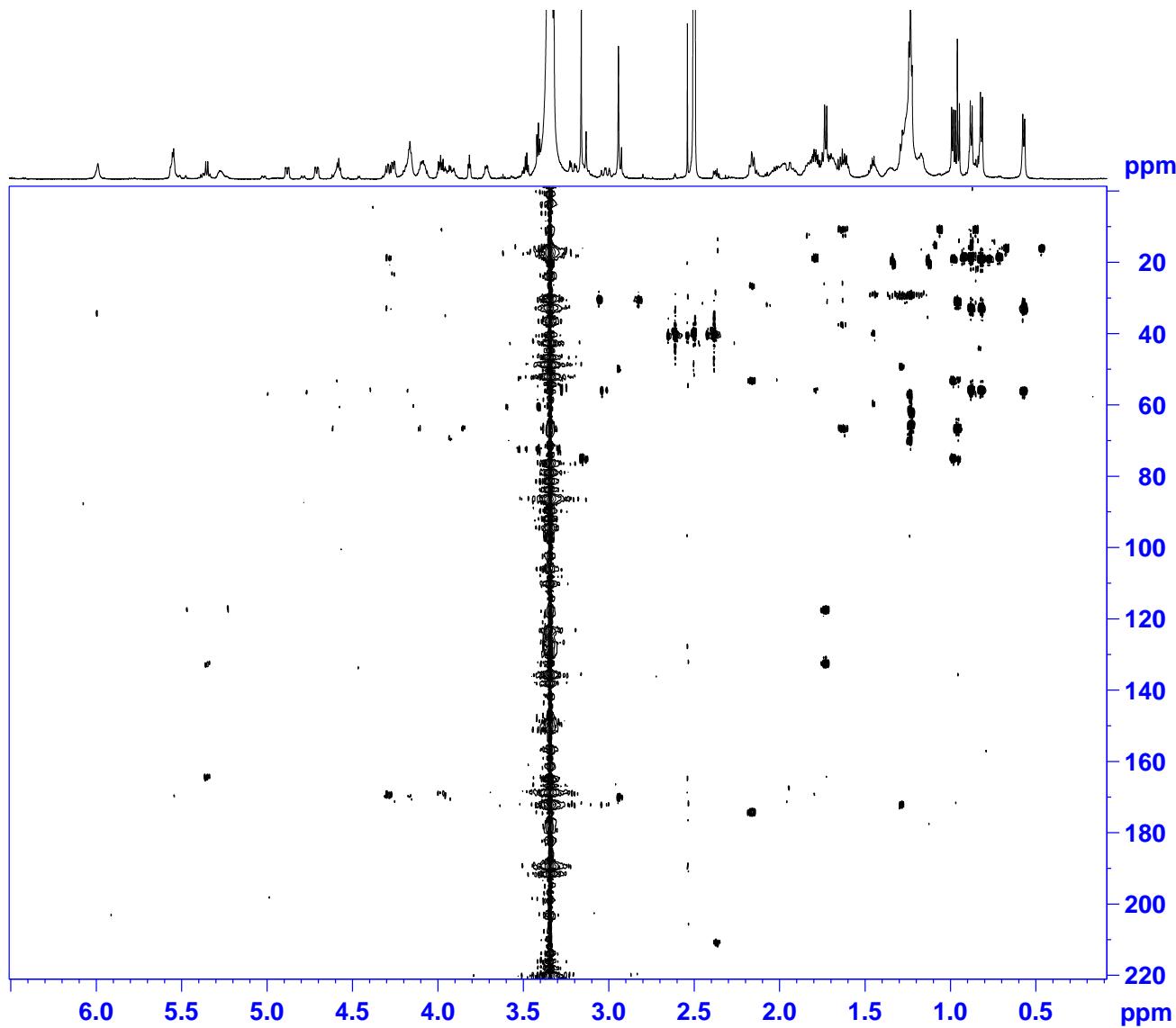
S37. TOCSY spectrum (600 MHz, DMSO-*d*₆) of **6**



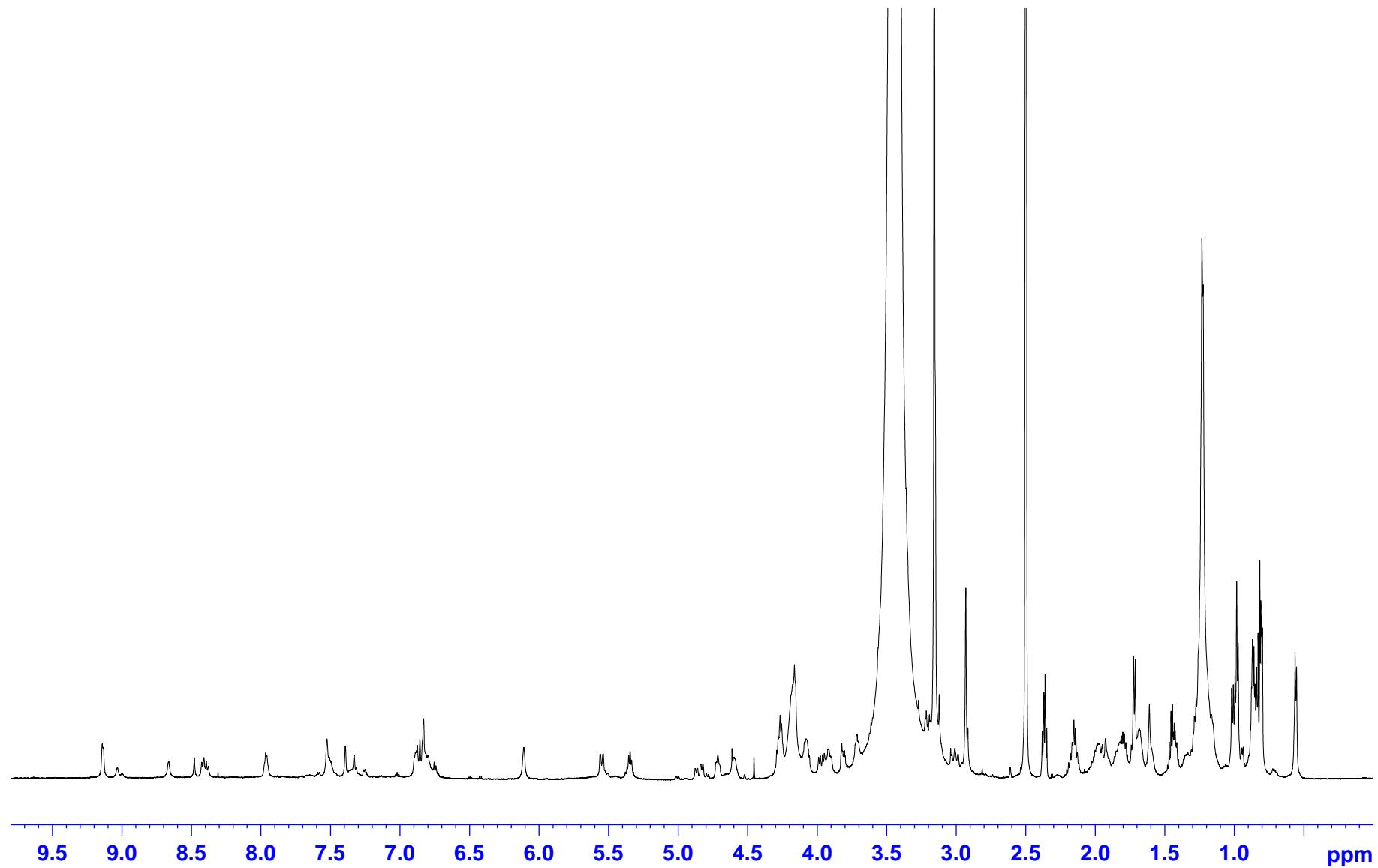
S38. HSQC spectrum (600 MHz, DMSO-*d*₆) of **6**



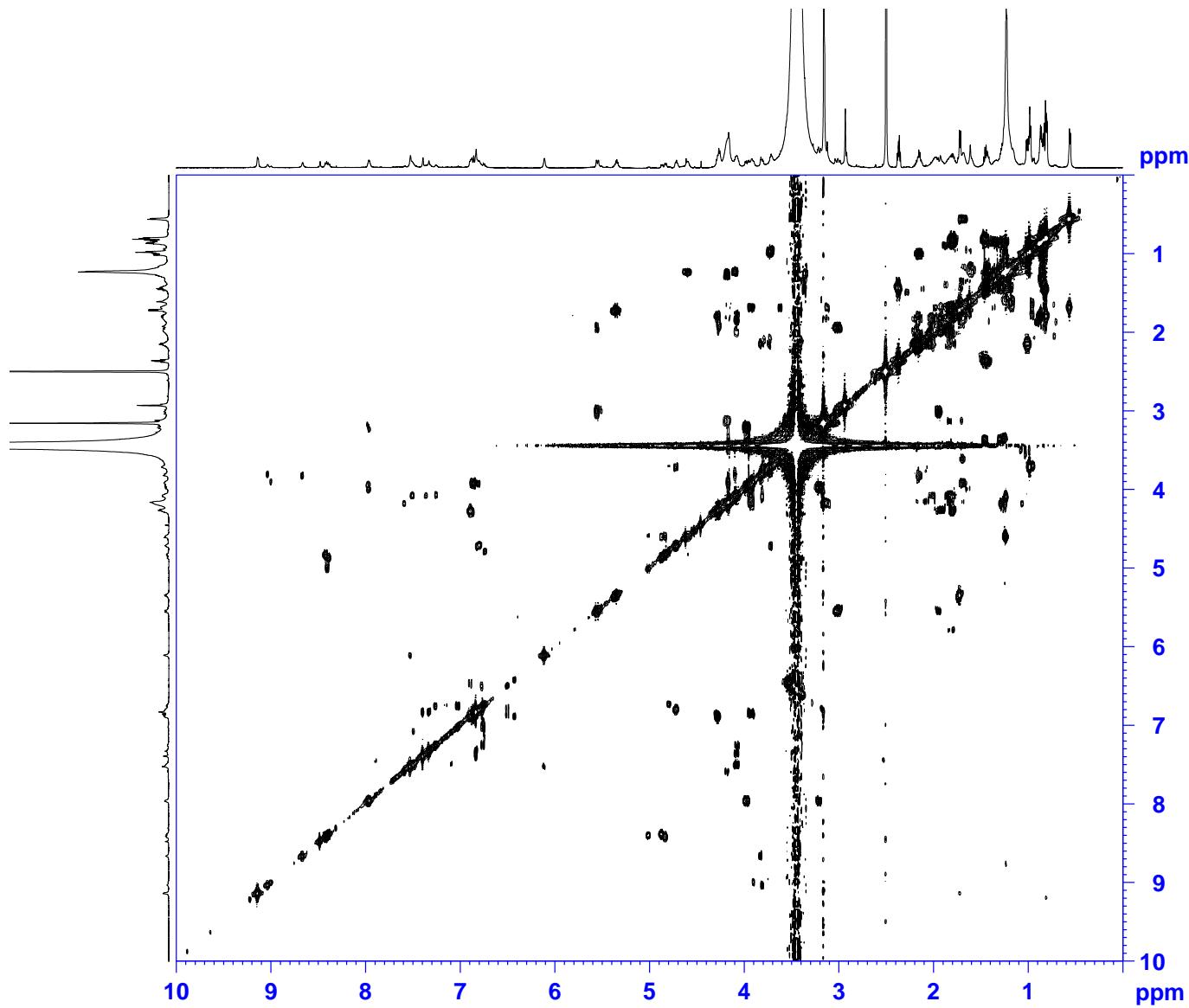
S39. HMBC spectrum (600 MHz, DMSO-*d*₆) of **6**



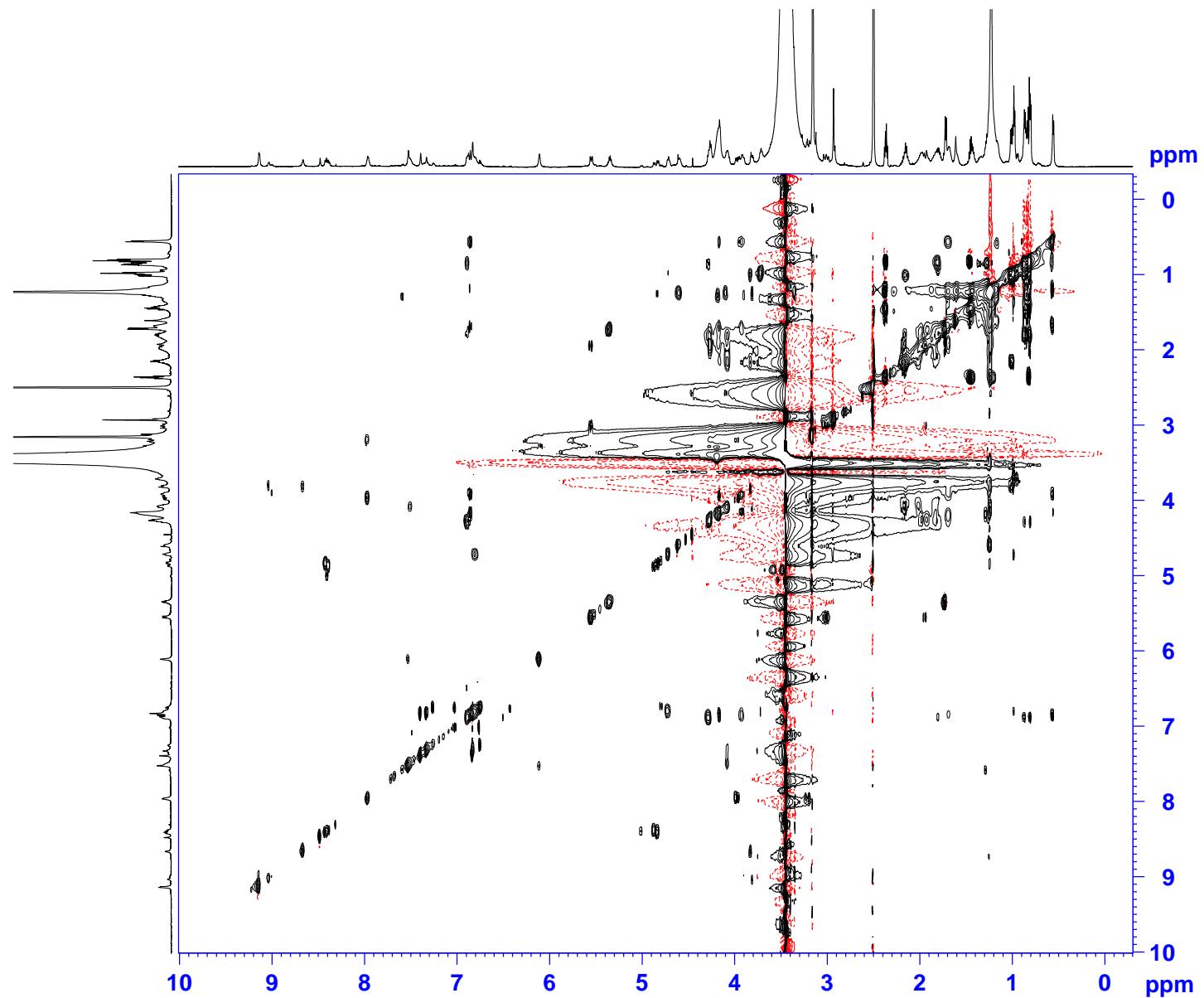
S40. ^1H NMR spectrum (600 MHz, DMSO- d_6) of inseparable mixture of **7** and **8**



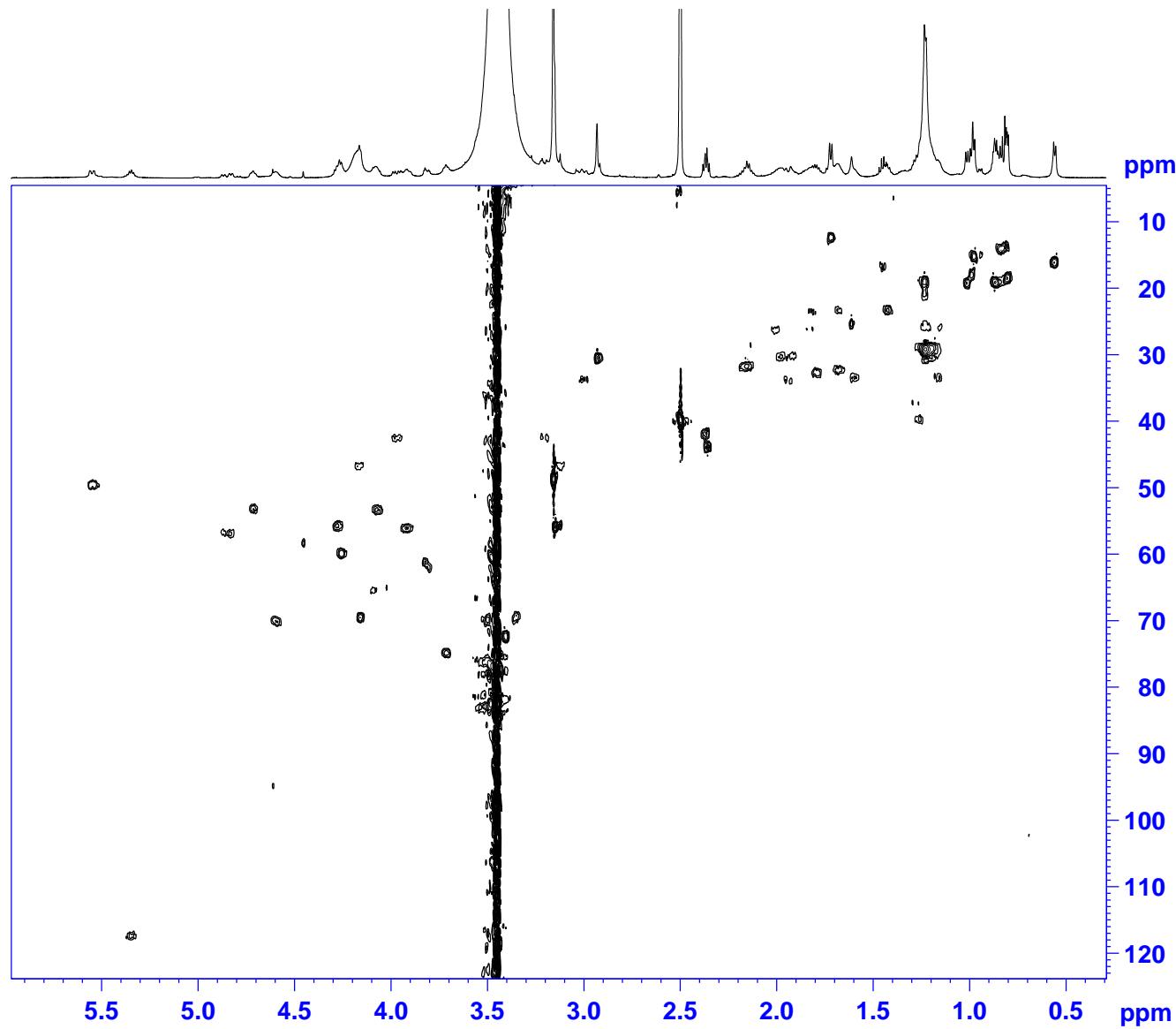
S41. COSY spectrum (600 MHz, DMSO-*d*₆) of inseparable mixture of **7** and **8**



S42. TOCSY spectrum (600 MHz, DMSO-*d*₆) of inseparable mixture of **7** and **8**



S43. HSQC spectrum (600 MHz, DMSO-*d*₆) of inseparable mixture of **7** and **8**



S44. HMBC spectrum (600 MHz, DMSO-*d*₆) of inseparable mixture of **7** and **8**

