

## Supplementary Material

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**Table S1.** Cyclic heptapeptides in the sponge *Styliissa caribica*.

| Cyclic peptide    | [M+H] <sup>+</sup>   | Exact Mass | Retention time<br>RP-18 <sup>a</sup> | Retention time<br>PFP <sup>b</sup> | Citation  |
|-------------------|--|------------|--------------------------------------|------------------------------------|-----------|
| Styliissamide A   | C <sub>44</sub> H <sub>61</sub> N <sub>8</sub> O <sub>9</sub>    | 845.4556   | 1.72                                 | 26.34                              | c,d,e     |
| Styliissamide B   | C <sub>44</sub> H <sub>58</sub> N <sub>7</sub> O <sub>8</sub>    | 812.4341   | ND                                   | ND                                 | c,d,e     |
| Styliissamide C   | C <sub>48</sub> H <sub>60</sub> N <sub>7</sub> O <sub>8</sub>    | 862.4498   | 20.29                                | 32.66                              | c,d,e     |
| Styliissamide D   | C <sub>45</sub> H <sub>62</sub> N <sub>7</sub> O <sub>8</sub>    | 828.4654   | 19.55                                | 32.48                              | c,d,e     |
| Styliissamide E   | C <sub>39</sub> H <sub>59</sub> N <sub>8</sub> O <sub>9</sub>    | 783.4400   | 13.29                                | 28.76                              | d,e       |
| Styliissamide F   | C <sub>43</sub> H <sub>57</sub> N <sub>10</sub> O <sub>9</sub>   | 857.4304   | 13.94                                | 32.21                              | d,e       |
| Styliissamide G   | C <sub>45</sub> H <sub>62</sub> N <sub>7</sub> O <sub>7</sub>    | 812.4705   | ND                                   | ND                                 | e         |
| Styliissamide H   | C <sub>44</sub> H <sub>59</sub> N <sub>8</sub> O <sub>8</sub>    | 827.4450   | ND                                   | ND                                 | e         |
| Styliissamide L   | C <sub>41</sub> H <sub>53</sub> N <sub>8</sub> O <sub>10</sub>   | 817.3876   | 6.21                                 | 24.95                              | this work |
| Hymenamide C      | C <sub>43</sub> H <sub>55</sub> N <sub>8</sub> O <sub>9</sub>    | 827.4087   | 20.71                                | 32.58                              | f         |
| Hymenamide F      | C <sub>35</sub> H <sub>61</sub> N <sub>10</sub> O <sub>8</sub> S | 781.4389   | 1.67                                 | 26.40                              | f         |
| Phakellistatin 3  | C <sub>42</sub> H <sub>55</sub> N <sub>8</sub> O <sub>9</sub>    | 815.4087   | 18.50                                | 31.89                              | h         |
| Phakellistatin 13 | C <sub>42</sub> H <sub>55</sub> N <sub>8</sub> O <sub>8</sub>    | 799.4137   | 20.09                                | 32.47                              | g         |
| Stylisin 1        | C <sub>45</sub> H <sub>62</sub> N <sub>7</sub> O <sub>8</sub>    | 828.4654   | ND                                   | ND                                 | c,d,e,g   |
| Stylisin 2        | C <sub>44</sub> H <sub>58</sub> N <sub>7</sub> O <sub>8</sub>    | 812.4341   | ND                                   | ND                                 | c,d,e,g   |

- a. Experiments were performed with a Kinetex 5  $\mu\text{m}$ , 50 mm  $\times$  2.10 mm C18 column using a flow rate of 200  $\mu\text{L}/\text{min}$  and the following elution gradient: 10% MeOH for 1 min, 10%–100% MeOH over 30 min, and 100% MeOH for 10 min.
- b. Experiments were performed with a Kinetex 5  $\mu\text{m}$ , 100 mm  $\times$  2.1 mm PFP column using a flow rate of 200  $\mu\text{L}/\text{min}$  and the same elution gradient of H<sub>2</sub>O and MeOH described above.
- c. Schmidt, G.; Grube, A.; Köck, M. Styliissamides A-D - New proline-containing cyclic heptapeptides from the marine sponge *Styliissa caribica*. European J. Org. Chem. 2007, 2, 4103–4110, doi:10.1002/ejoc.200700013.
- d. Cychon C.; Köck, M. Styliissamides E and F, Cyclic Heptapeptides from the Caribbean Sponge *Styliissa caribica*. J. Nat. Prod. 2010, 73, 738–742, doi: 10.1021/np900664f.
- e. Wang, X.; Morinaka, B. I.; Molinski, T. F. Structures and solution conformational dynamics of styliissamides G and H from the Bahamian Sponge *Styliissa caribica*. J. Nat. Prod. 2014, 77, 625–630, doi:10.1021/np400891s.
- f. Grube, A.; Maier, T.; Köck, M. MS-guided Fractionation as a Fast Way to the Identification of New Natural Products – MALDI-TOF-MS Screening of the Marine Sponge *Styliissa caribica*. Zeitschrift für Naturforsch. B 2007, 62, 600–604, doi:10.1515/znb-2007-0420..
- g. Mohammed, R.; Peng, J.; Kelly, M.; Hamann, M.T. Cyclic heptapeptides from the Jamaican sponge *Styliissa caribica*. J. Nat. Prod. 2006, 69, 1739–1744, doi:10.1021/np060006n.
- h. This compound was not previously detected in specimens of *Styliissa caribica*. It was putatively identified by the Dereplicator tool in GNPS.

Table S2. Full NMR data of stylissamide L (1) ( $^1\text{H}$  700 MHz,  $^{13}\text{C}$  175 MHz, DMSO- $d_6$ ).

| AA                 | pos.              | $\delta_{\text{C}}$ , type | $\delta_{\text{H}}$ , mult (J in Hz)                          | NOESY  | HMBC                                       |
|--------------------|-------------------|----------------------------|---|--|--|
| Pro <sup>I</sup>   | 1                 | 170.3, C                   | -   |  |  |
|                    | 2                 | 59.1, CH                   | 4.34, dd (5.1, 8.6)   | Ser-NH, Tyr-NH, Phe-NH   | Phe-1                                      |
|                    | 3                 | 28.1, CH <sub>2</sub>      | a 2.15, m<br>b 1.75, m  | Pro <sup>II</sup> -4b, Pro <sup>II</sup> -3a   | Pro <sup>I</sup> -1                        |
|                    | 4                 | 24.3, CH <sub>2</sub>      | 1.87, m   | Pro <sup>II</sup> -2, Pro <sup>II</sup> -4b  | Pro <sup>I</sup> -1                        |
|                    | 5                 | 46.7, CH <sub>2</sub>      | a 3.45, m<br>b 3.36, m  |  | Phe-1, Pro <sup>I</sup> -2                 |
| Pro <sup>II</sup>  | 1                 | 171.8, C                   | -   |  |  |
|                    | 2                 | 60.1, CH                   | 4.28, dd (1.5, 8.8)   | Ser-NH, Pro <sup>I</sup> -2, Pro <sup>I</sup> -3b  | Pro <sup>II</sup> -1, Pro <sup>II</sup> -5 |
|                    | 3                 | 31.8, CH <sub>2</sub>      | a 2.16, m<br>b 2.00, m  | Ser-NH   | Pro <sup>II</sup> -1                       |
|                    | 4                 | 21.7, CH <sub>2</sub>      | a 1.77, m<br>b 1.57, m  | Pro <sup>III</sup> -3a   |  |
|                    | 5                 | 46.8, CH <sub>2</sub>      | a 3.60, ddd (1.5, 8.4, 10.8)<br>b 3.33, ddd (10.8, 10.8, 7.1) | Ser-NH<br>Pro <sup>III</sup> -4b   | Pro <sup>I</sup> -1                        |
| Ser                | NH                |                            | 7.65, d (5.9)   | Tyr-NH, Pro <sup>I</sup> -2, Pro <sup>II</sup> -2, Pro <sup>II</sup> -5a, Pro <sup>II</sup> -4b  | Pro <sup>II</sup> -1                       |
|                    | 1                 | 167.7, C                   | -   |  |  |
|                    | 2                 | 60.0, CH                   | 3.85, ddd (3.6, 5.9, 10.2)                                    | Tyr-NH   | Ser -1                                     |
|                    | 3                 | 60.9, CH <sub>2</sub>      | a 3.46, dd (10.2, 11.9)<br>b 3.14, dd (11.9, 3.6)             | Tyr-NH, Tyr-5/9  |  |
| Tyr                | NH                |                            | 7.34, d (9.1)   | Ser-NH, Ser-3a   | Ser-1                                      |
|                    | 1                 | 171.5 C                    | -   |  |  |
|                    | 2                 | 51.5 CH                    | 4.88 ddd (3.2, 9.1, 10.9)                                     | Pro <sup>III</sup> -5a/b   | Tyr- 1                                     |
|                    | 3                 | 37.0 CH <sub>2</sub>       | a 3.35, dd (3.2, 13.5)<br>b 2.42, dd (10.9, 13.5)             | Pro <sup>III</sup> -5a/b, Phe-NH, Gln-NH<br>Gln-NH, Phe-NH, Pro <sup>III</sup> -5a               | Tyr-5/9<br>Tyr-1, Tyr-4                    |
|                    | 4                 | 126.6 C                    | -   |  |  |
|                    | 5/9               | 130.5 CH                   | 7.08, d (8.5)   | Tyr-3a, Tyr-3b   | Tyr-7                                      |
|                    | 6/8               | 114.9 CH                   | 6.66, d (8.5)   |  | Tyr-7                                      |
| Pro <sup>III</sup> | 7                 | 156.0 C                    | -   |  |  |
|                    | 7-OH              |                            | 7.42, s   |  |  |
|                    | 1                 | 171.9, C                   | -   |  |  |
|                    | 2                 | 63.1, CH                   | 4.06, t (8.7)   | Phe-NH,  | Pro <sup>III</sup> -1                      |
|                    | 3                 | 28.7, CH <sub>2</sub>      | a 2.22 m<br>b 1.81, m   | Gln-NH   | Pro <sup>III</sup> - 1                     |
| Gln                | 4                 | 25.0, CH <sub>2</sub>      | a 2.11, m<br>b 1.98, m  | Gln-NH, Gln-3a, Gln-3b   |  |
|                    | 5                 | 46.9, CH <sub>2</sub>      | a 3.93, ddd (6.8, 9.8, 9.8)<br>b 3.82, m                      | Tyr-2, Tyr-3a<br>Tyr-2, Gln-NH, Tyr-3a   | Pro <sup>III</sup> -3                      |
|                    | NH                |                            | 8.17, d (7.0)   | Phe-NH, Tyr-3a, Pro <sup>III</sup> -2,<br>Pro <sup>III</sup> -3b, Pro <sup>III</sup> -4a         | Pro <sup>III</sup> -1                      |
|                    | 1                 | 170.7, C                   | -   |  |  |
|                    | 2                 | 52.8, CH                   | 4.05, ddd (4.3, 7.0, 10.0)                                    | Phe-NH, Pro <sup>III</sup> -4b   | Gln-1                                      |
| Phe                | 3                 | 25.9, CH <sub>2</sub>      | a 1.85, m<br>b 1.73, m  | Pro <sup>III</sup> -4a<br>Pro <sup>III</sup> - 4a  | Gln-1, Gln-5<br>Gln-5                      |
|                    | 4                 | 31.5, CH <sub>2</sub>      | a 2.13, ddd (7.2, 15.7, 7.2)<br>b 2.04, ddd (7.2, 15.7, 7.2)  | Pro <sup>III</sup> -5b, Pro <sup>III</sup> -3a, Pro <sup>III</sup> -3b<br>Pro <sup>III</sup> -4a | Gln-5<br>Gln-5                             |
|                    | 5                 | 174.5, C                   | -   |  |  |
|                    | 5-NH <sub>2</sub> |                            | 6.92, s   | Gln-4a, Gln-4b   | Gln-5                                      |
|                    | NH                |                            | 7.11, d (7.2)   | Phe-2, Gln-NH, Pro <sup>III</sup> -2, Pro <sup>I</sup> -5b,<br>Tyr-NH                            | Gln-1                                      |
|                    | 1                 | 167.5 C                    | -   |  |  |
|                    | 2                 | 51.5, CH                   | 4.69, ddd (5.8, 7.2, 8.0)                                     | Pro <sup>I</sup> -5a, Pro <sup>I</sup> -5b   | Phe-1, Phe-4                               |
|                    | 3                 | 36.9 CH <sub>2</sub>       | a 3.18, dd (8.0, 14.2)<br>b 2.71 (5.8, 14.2)                  |  | Phe-1, Phe-4                               |
|                    | 4                 | 138.0, C                   | -   |  | Phe-1, Phe-5/9                             |
|                    | 5/9               | 128.9, CH                  | 7.16, d (7.5)   |  |  |
|                    | 6/8               | 126.0, CH                  | 7.18, t (7.3)   |  |  |
|                    | 7                 | 128.0, CH                  | 7.22, t (7.5)   |  |  |

Table S3. Links to LC-MS data and molecular networks.

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| Description  | Link  |
|--|---|
| Classical Molecular Networking (Metabolomics) workflow | <a href="https://gnps.ucsd.edu/ProteoSAFe/status.jsp?task=4acb49028d7041c39fccf03dd6e8c195">https://gnps.ucsd.edu/ProteoSAFe/status.jsp?task=4acb49028d7041c39fccf03dd6e8c195</a>   |
| Feature-Based Molecular Networking workflow            | <a href="https://gnps.ucsd.edu/ProteoSAFe/status.jsp?task=db7aa9e8bec64d6290cc040254f00b87">https://gnps.ucsd.edu/ProteoSAFe/status.jsp?task=db7aa9e8bec64d6290cc040254f00b87</a>   |
| Dereplicator   | <a href="https://gnps.ucsd.edu/ProteoSAFe/status.jsp?task=f8b123b5ea89494481549c1a519caf5d">https://gnps.ucsd.edu/ProteoSAFe/status.jsp?task=f8b123b5ea89494481549c1a519caf5d</a>   |
| LC-MS data on Massive                                  | <a href="https://massive.ucsd.edu/ProteoSAFe/dataset.jsp?task=f77c49cb5d73489583c08d3e66b558e1">https://massive.ucsd.edu/ProteoSAFe/dataset.jsp?task=f77c49cb5d73489583c08d3e66b558e1</a><br>FTP access: ftp://MSV000085904@massive.ucsd.edu<br>password: Sty_reviewers |

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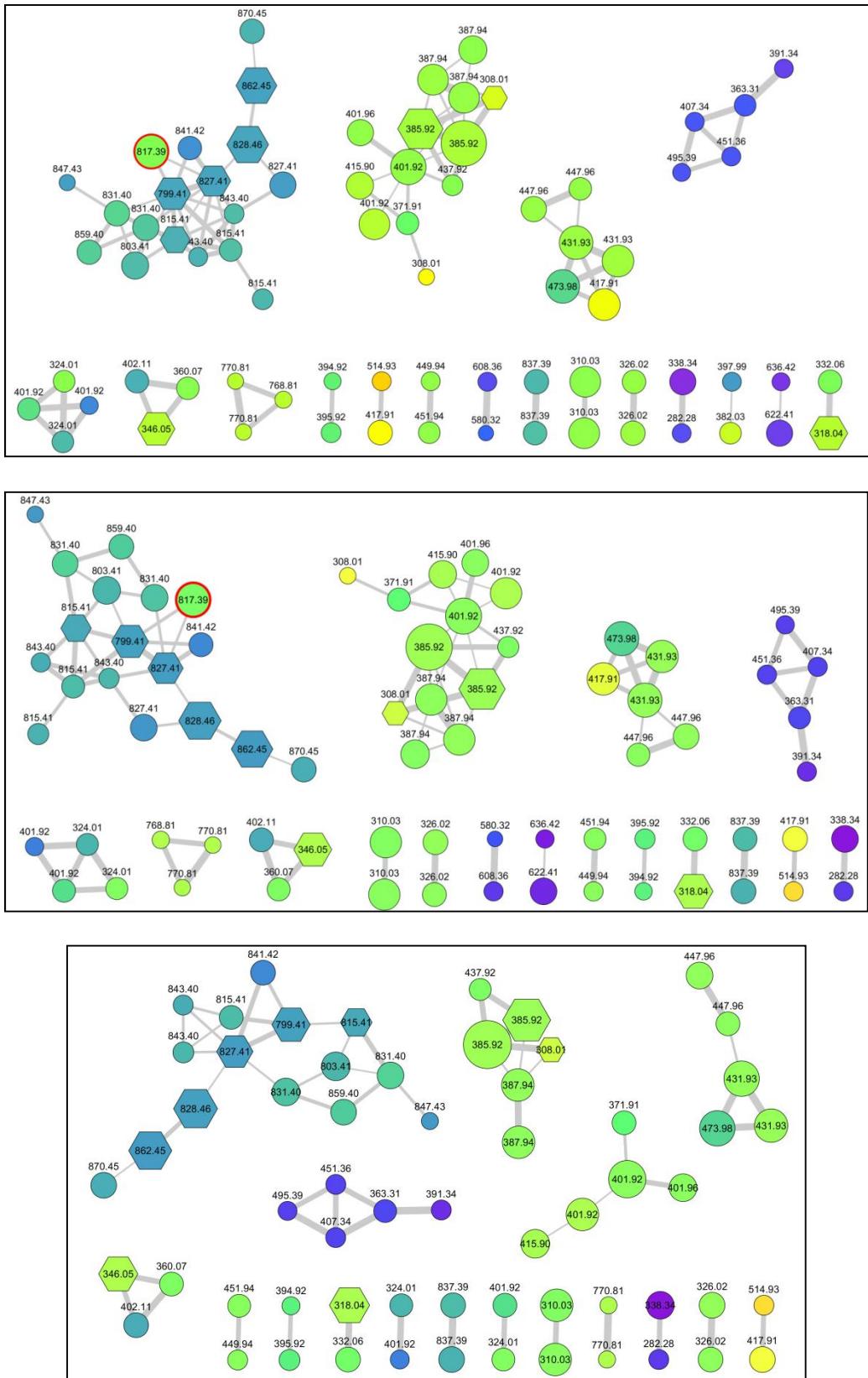


Figure S1. Molecular networks of the extract of *S. caribica* obtained with the same data and the same parameters ( $m/z$  tolerance 0.01 Da, cosine score  $> 0.55$ , matched peaks  $> 8$ , maximum number of neighbor nodes = 10) using (a) the program MetGem (b) the Metabolomics workflow on GNPS, and (c) the Feature-Based Molecular Network workflow on GNPS.

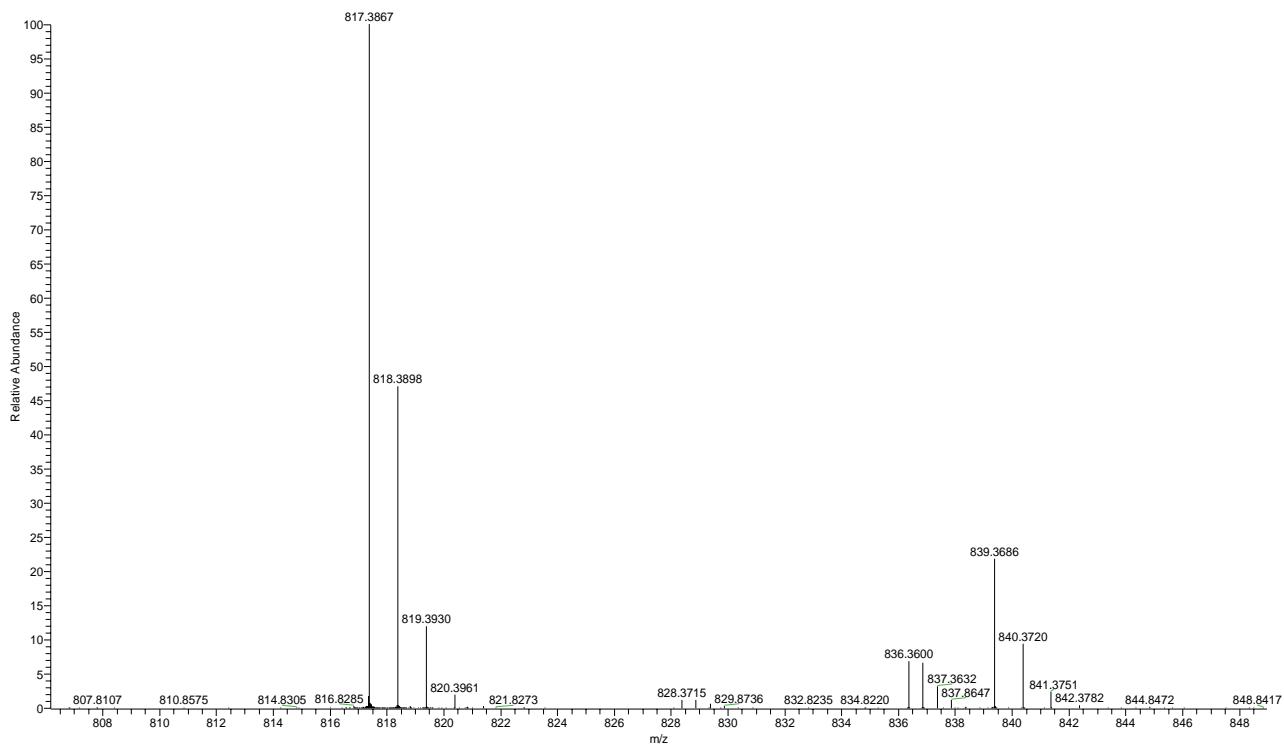


Figure S2. Positive ion mode high-resolution ESI mass spectrum of stylissamide L (1).

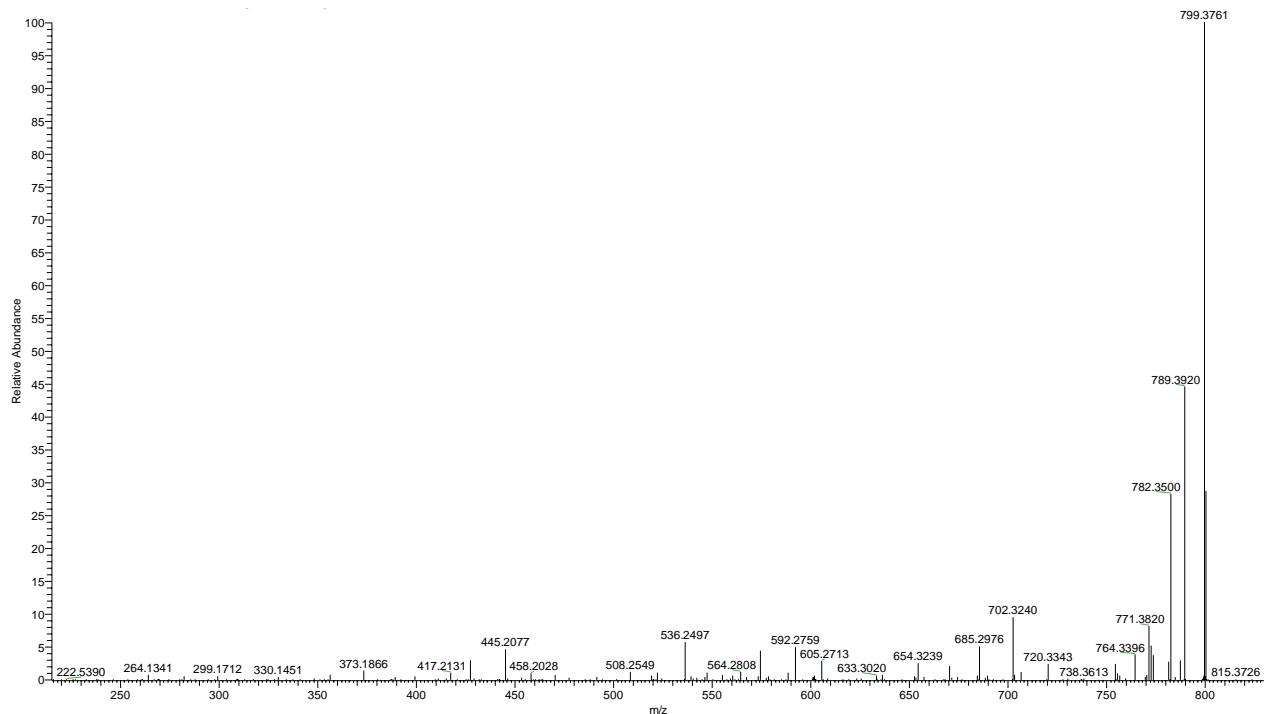


Figure S3. Positive ion mode high-resolution ESI MS/MS spectrum of stylissamide L (1).

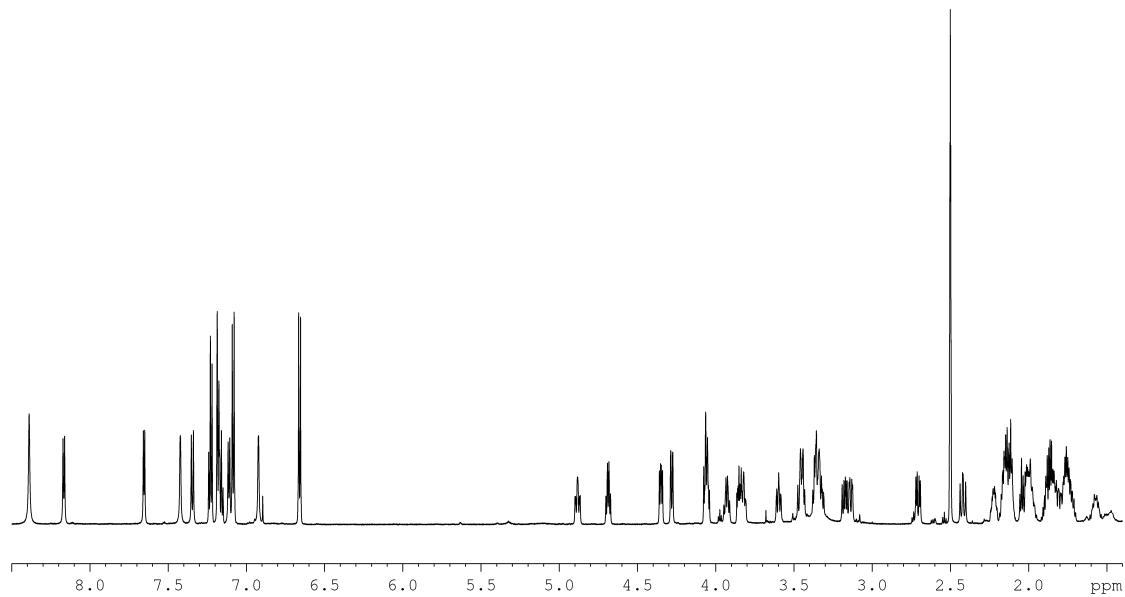


Figure S4. <sup>1</sup>H-NMR spectrum of stylissamide L (1) (700 MHz, DMSO-*d*<sub>6</sub>).

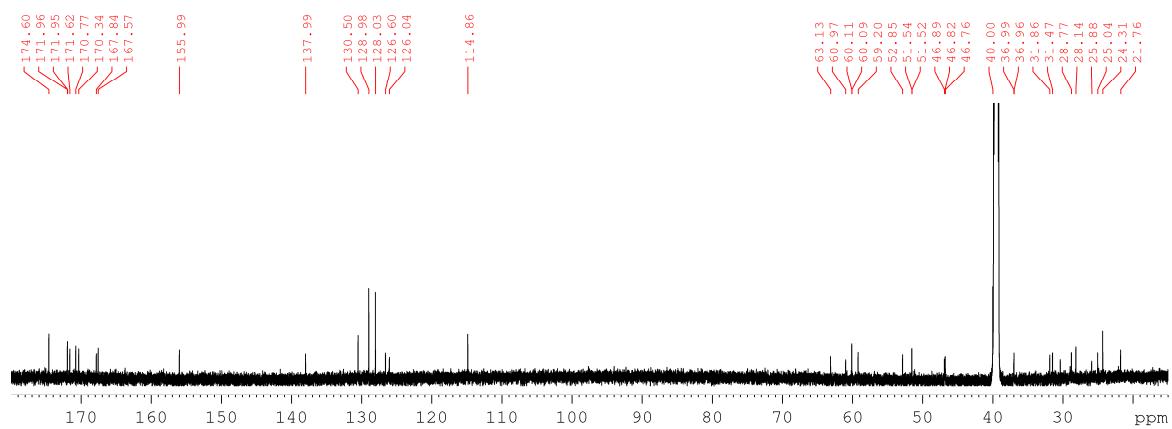


Figure S5. <sup>13</sup>C-NMR spectrum of stylissamide L (1) (175 MHz, DMSO-*d*<sub>6</sub>).

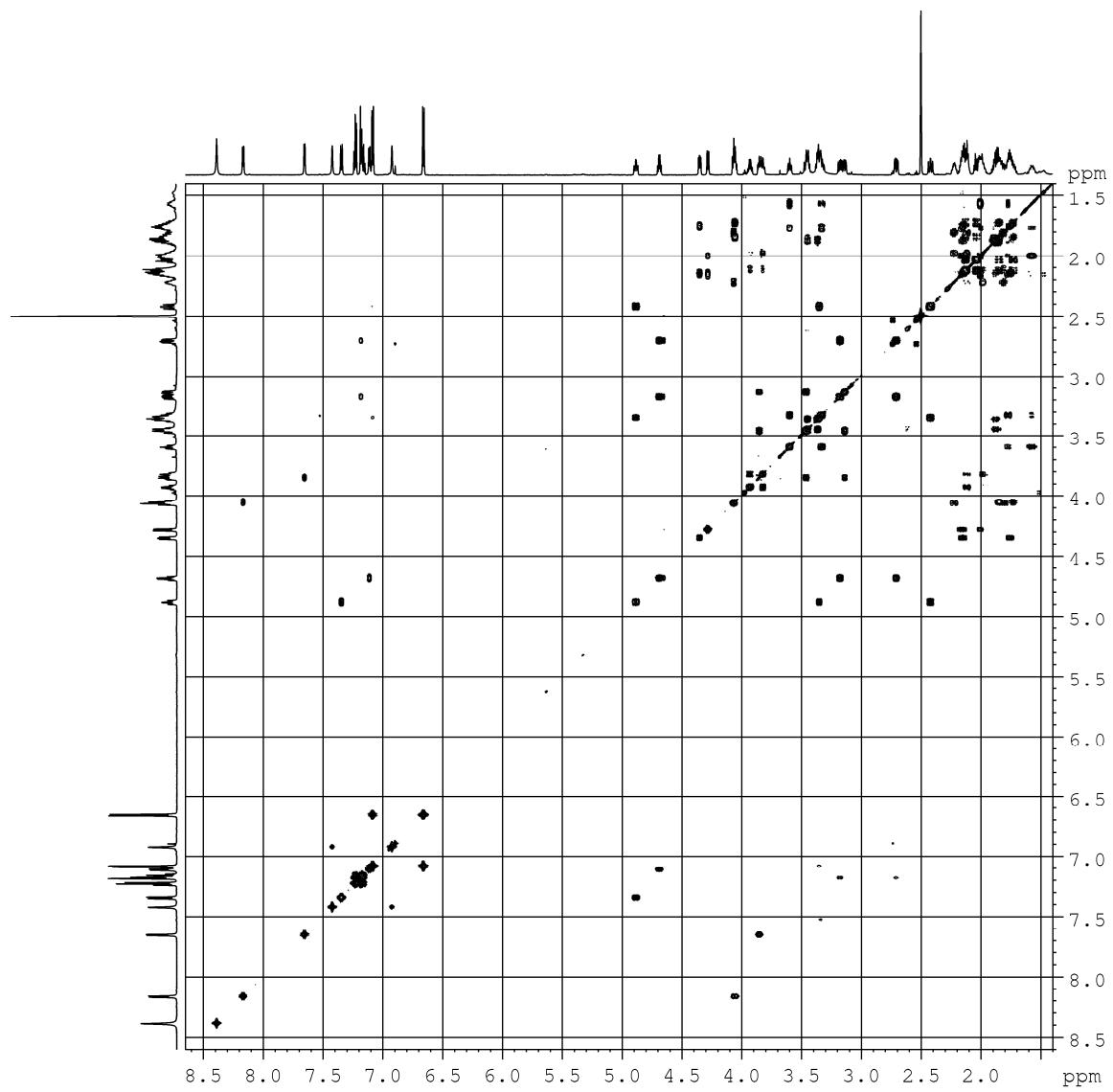


Figure S6. COSY spectrum of stylissamide L (1) (700 MHz,  $\text{DMSO}-d_6$ ).

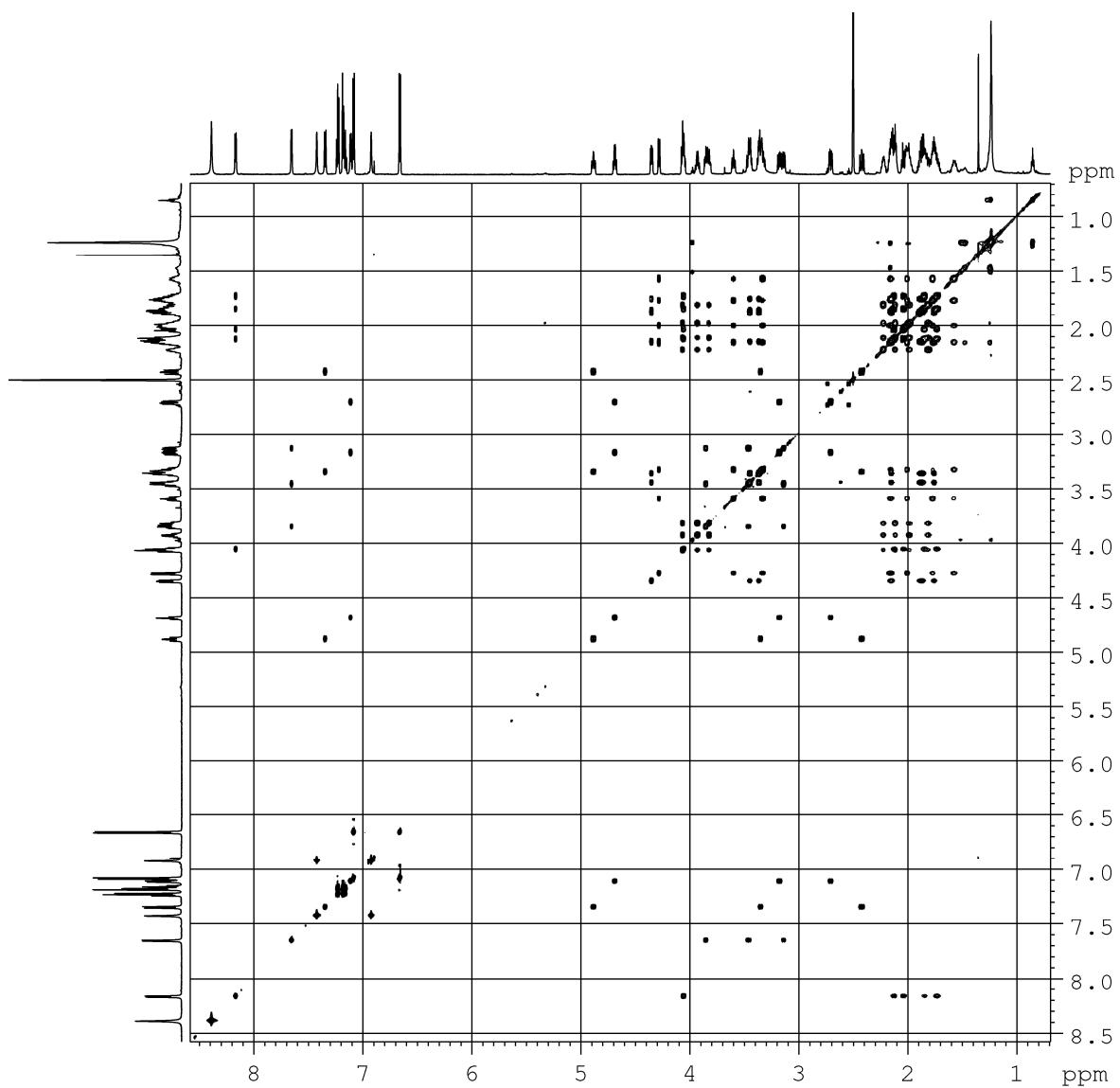


Figure S7. TOCSY spectrum of stylissamide L (1) (700 MHz, DMSO-*d*<sub>6</sub>).

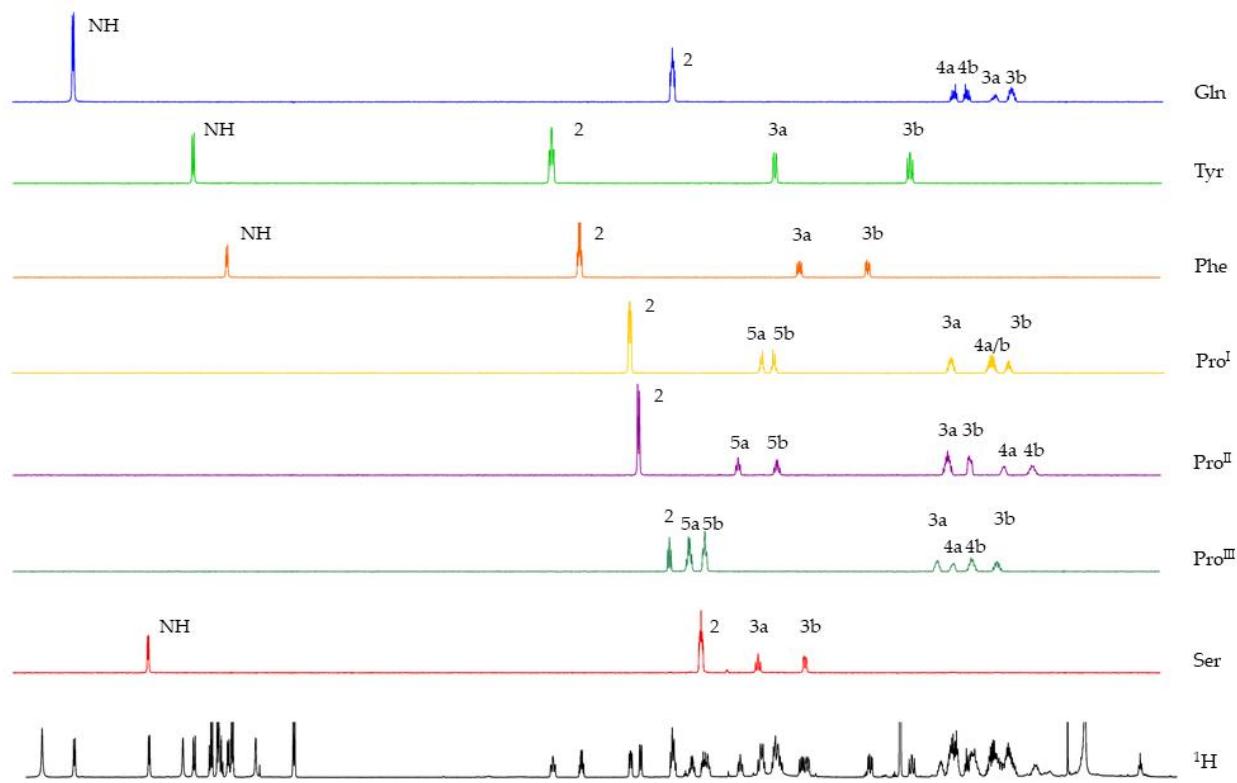


Figure S8. Spin systems of stylissamide L (1) from the sections of the TOCSY spectrum.

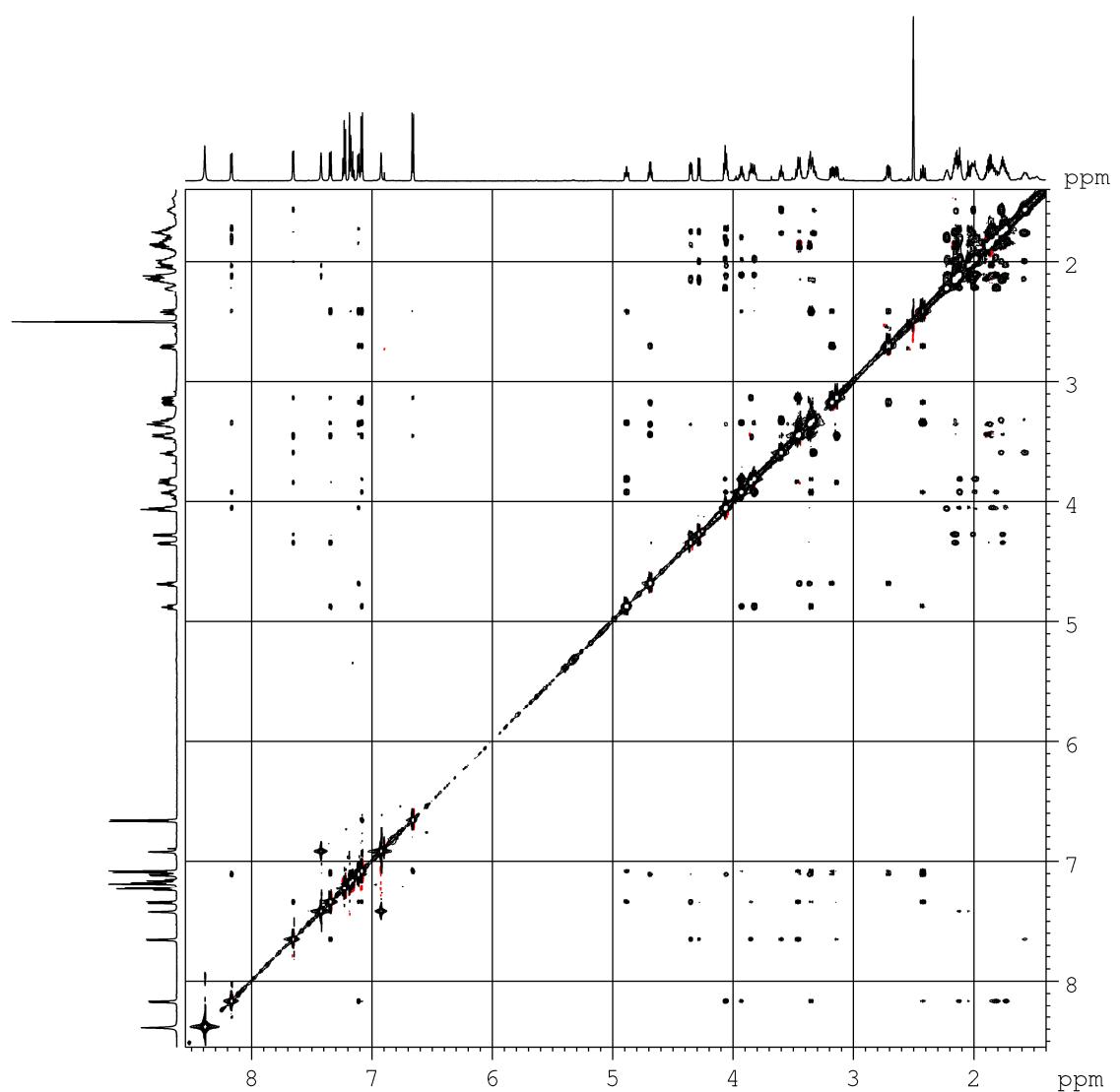


Figure S9. NOESY spectrum of stylissamide L (1) (700 MHz,  $\text{DMSO}-d_6$ ).

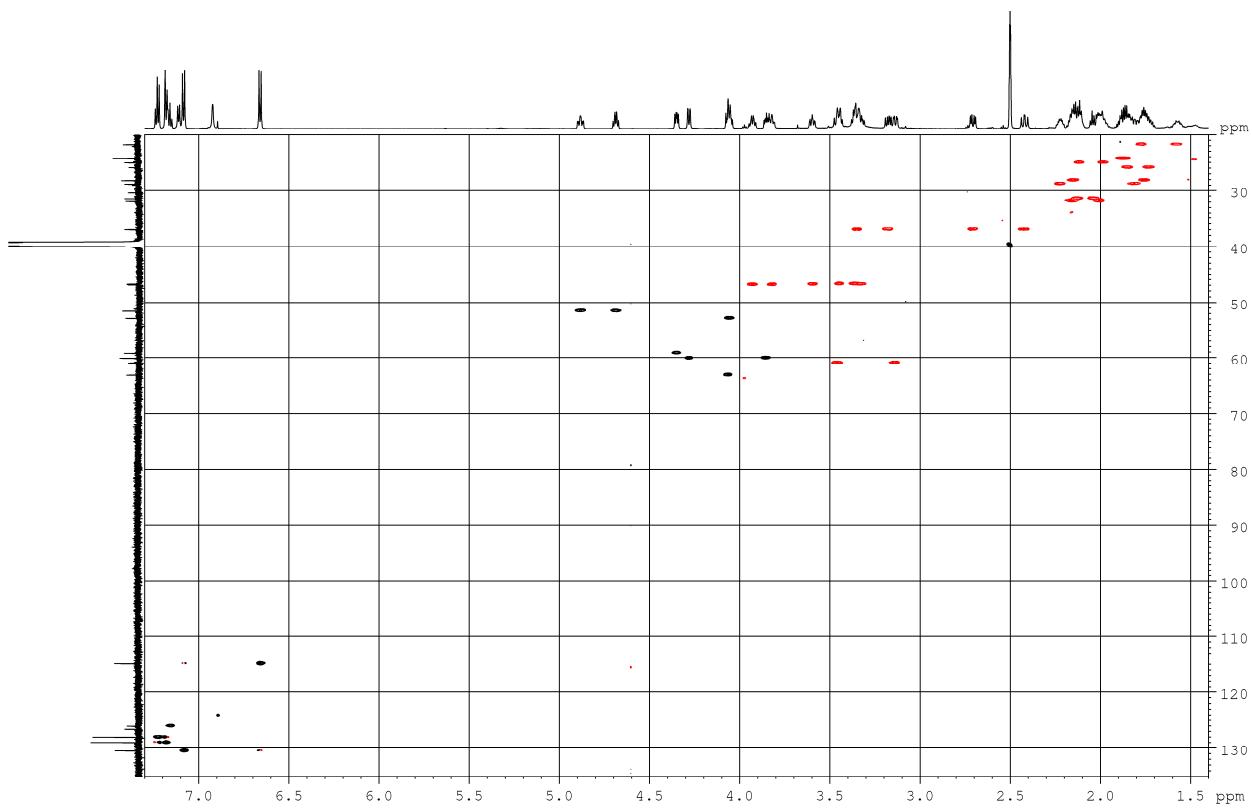


Figure S10. HSQC spectrum of stylissamide L (1) (700 MHz,  $\text{DMSO}-d_6$ ).

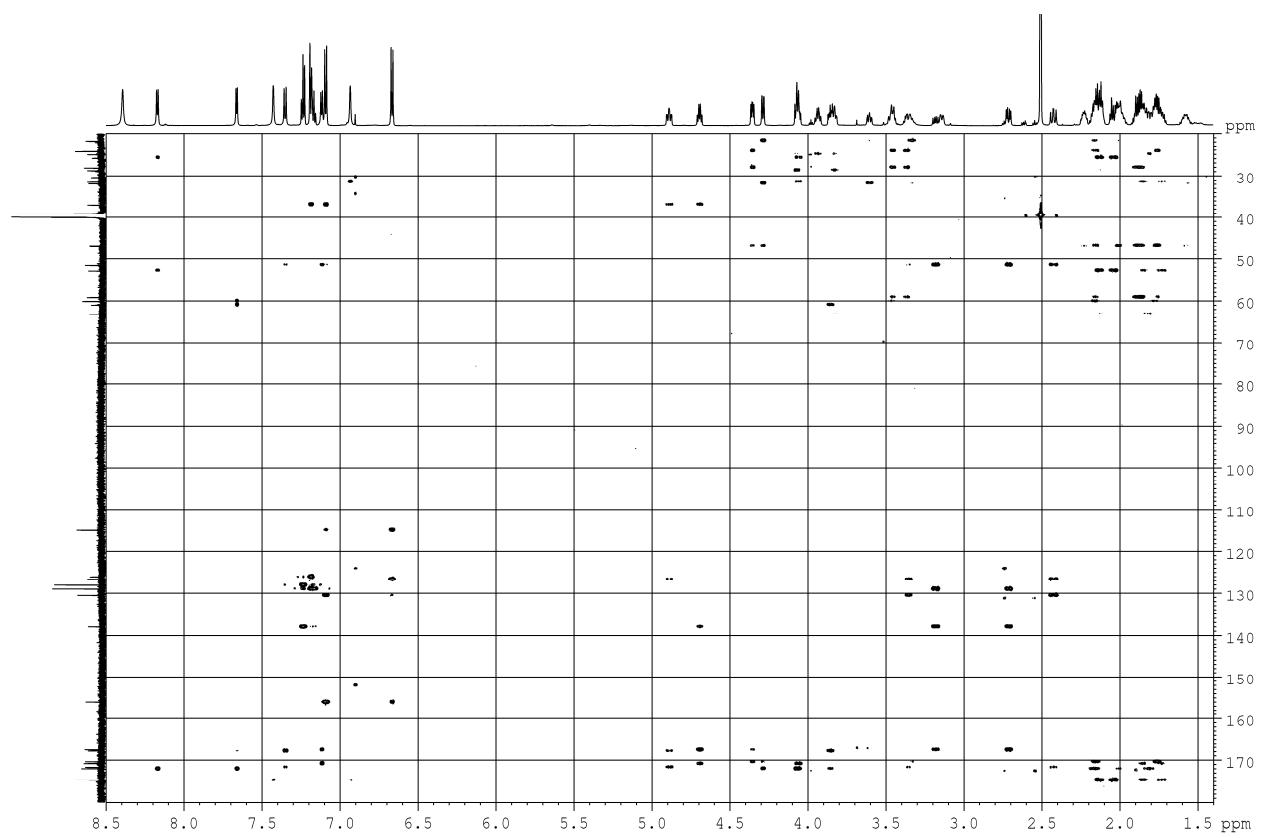


Figure S11. HMBC spectrum of stylissamide L (1) (700 MHz,  $\text{DMSO}-d_6$ ).

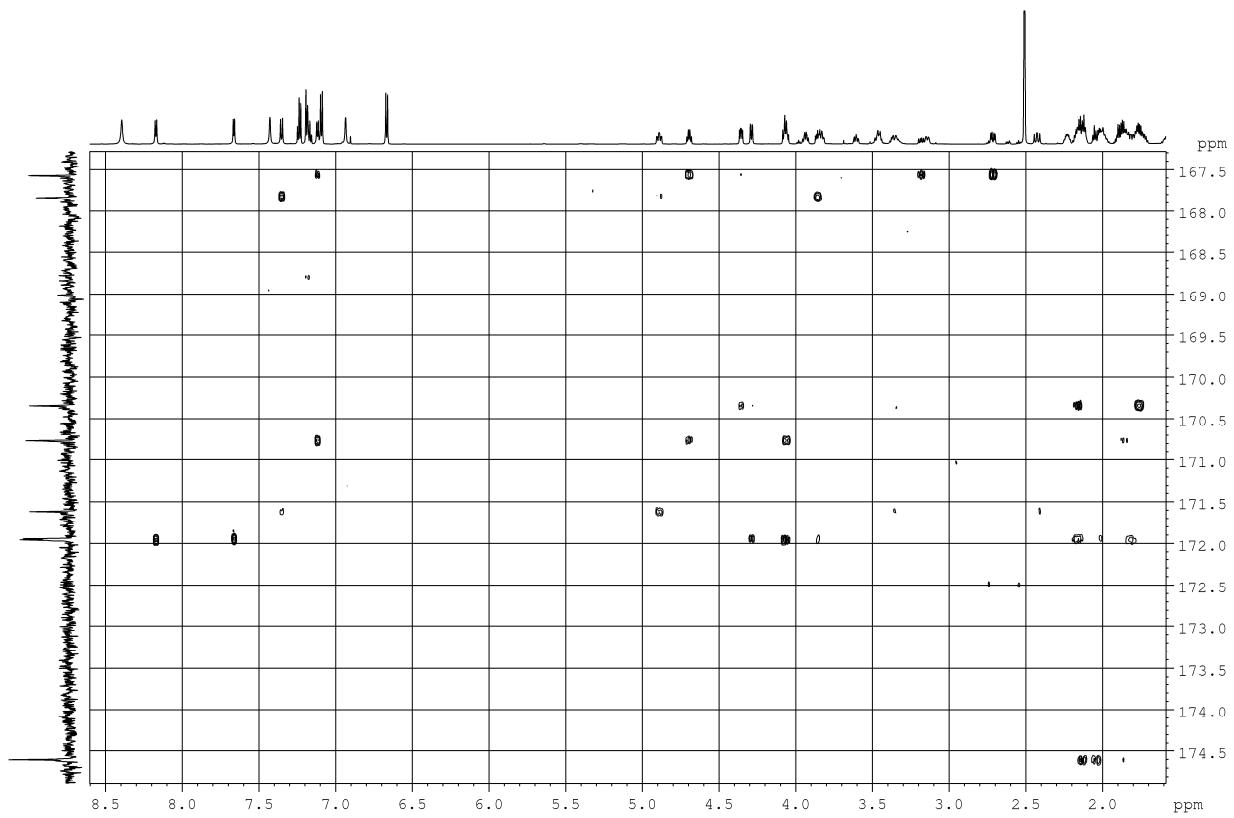


Figure S12. Band-selective HMBC spectrum of stylissamide L (1) (700 MHz, DMSO-*d*<sub>6</sub>).

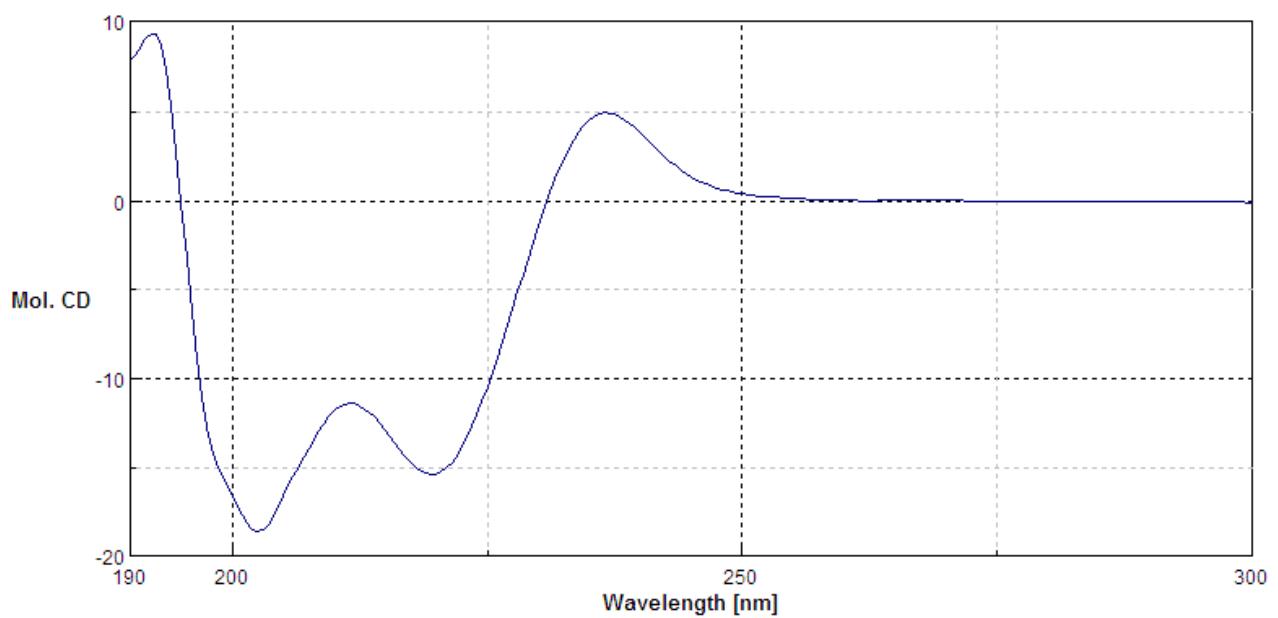
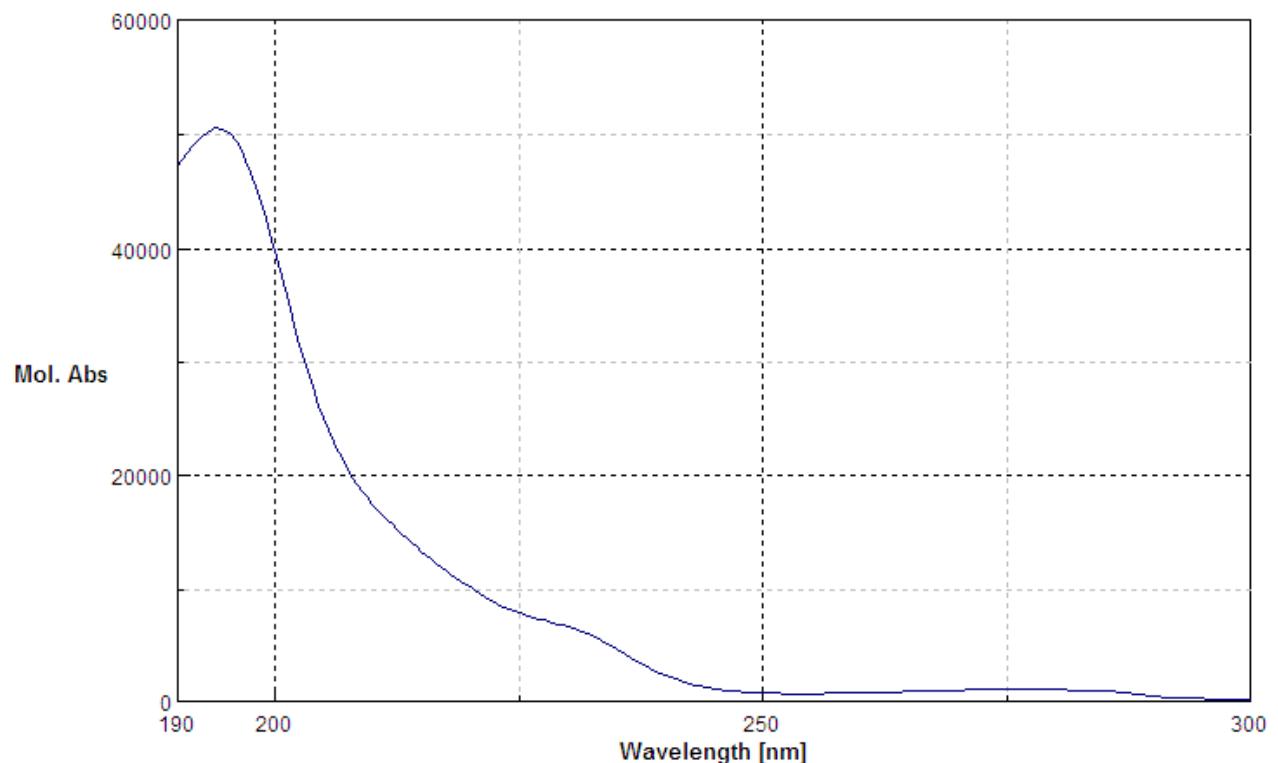
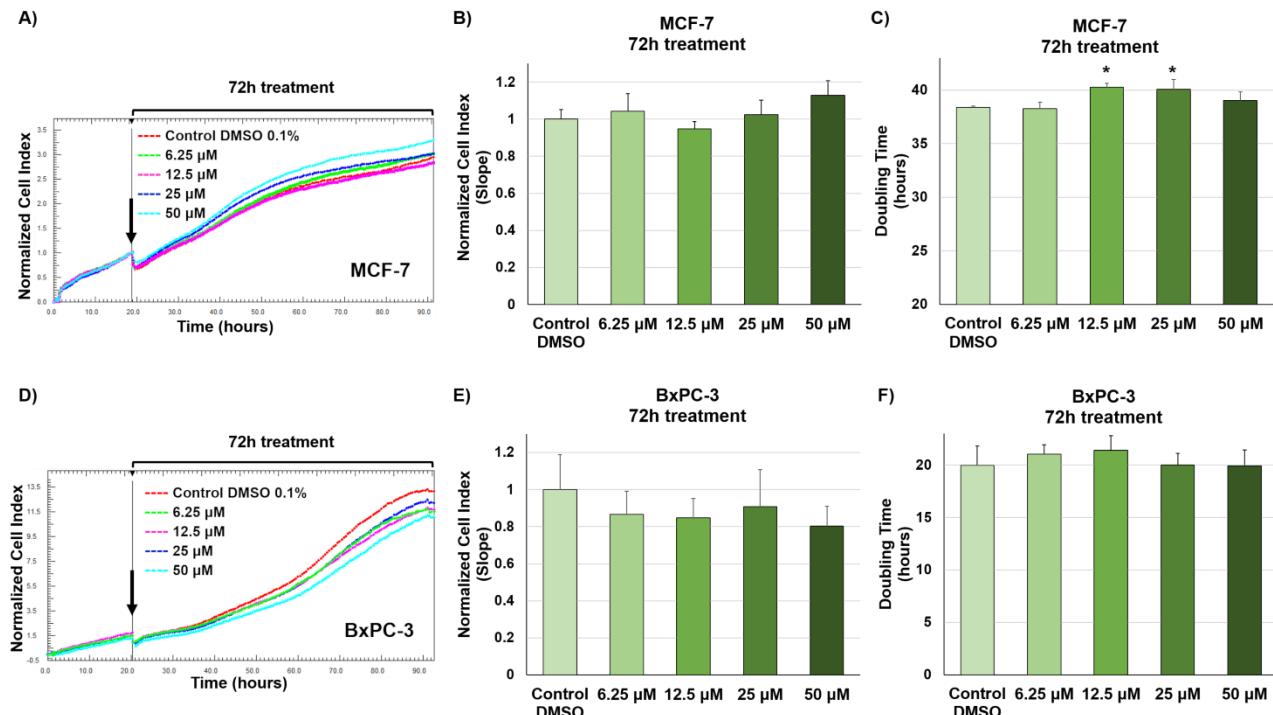
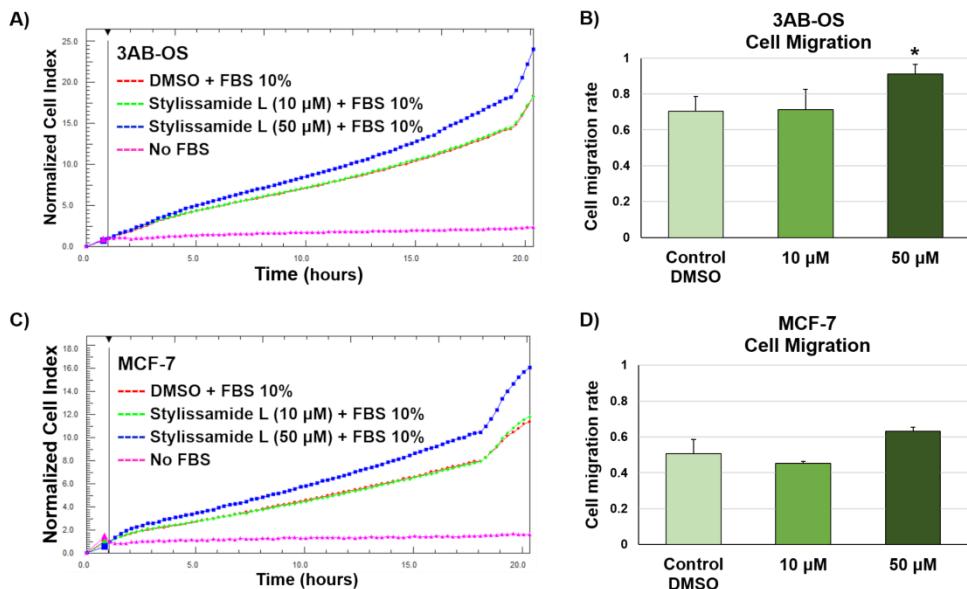


Figure S13. UV and ECD spectra of stylissamide L (1) in ACN.



**Figure S14.** Real time monitoring of cancer cell proliferation after exposure to stylissamide L (**1**) and DMSO vehicle, by using the xCELLigence System Real-Time Cell Analyzer. **(A,D)** Normalized cell index (NCI) traces of MCF7 (**A**) and BxPC-3 (**D**) cells treated with different concentrations (6.25, 12.5, 25, and 50 µM) of stylissamide L (**1**) and DMSO vehicle (0.1%) for 72 hours. Black arrow shows the start of drug treatment. Each cell index value was normalized at this time. **(B,E)** Slope values of growth curves of MCF-7 (**B**) and BxPC-3 (**E**) cells after 72 h exposure to different concentrations (6.25, 12.5, 25, and 50 µM) of stylissamide L (**1**) and DMSO vehicle (0.1%). NCI slope values are relative to controls treated with DMSO vehicle. **(C,F)** Doubling times of NCI of MCF-7 (**C**) and BxPC-3 (**F**) cells after 72 h treatment with different concentrations (6.25, 12.5, 25, and 50 µM) of stylissamide L (**1**) and DMSO (0.1%). Data are presented as mean ± SD; n=3. Statistical significances are referred to the DMSO control. One-way analysis of variance (ANOVA) was applied to compare means of groups and Dunnett's method was used as a post-hoc test to compare multiple groups versus the control group. *p*-values < 0.05 were considered to be statistically significant. Statistical analysis was performed using the GraphPad Prism Software Version 5. \* *p* < 0.05.



**Figure S15.** Real-time monitoring of 3AB-OS and MCF-7 cell migration after exposure to stylissamide L (**1**). (A,C) NCI traces of 3AB-OS (A) and MCF-7 (C) cells seeded with compound **1** or DMSO (0.1%) vehicle, in presence of 10% Fetal Bovine Serum (FBS) as the chemoattractant. Migration was monitored for 20 hours, using the xCELLigence System equipped with specially designed 16-well plates (CIM-plate 16). (B,D) Migration activity of 3AB-OS (B) and MCF-7 (D) cells seeded with compound **1** or DMSO (0.1%) vehicle, in presence of 10% Fetal Bovine Serum (FBS) as the chemoattractant. Cell migration rates were recorded for 20 hours and expressed as slope values of NCI curves. Data are presented as mean  $\pm$  SD; n=3. Statistical significances are referred to the DMSO control. Two-group comparisons were performed using Student's t-test. P-values < 0.05 were considered to be statistically significant. Statistical analysis was performed using the GraphPad Prism Software Version 5.