

Supplementary Material

Discovery of Primarolides A and B from Marine Fungus *Asteromyces cruciatus* by Osmotic Stress and Treatment with Suberoylanilide Hydroxamic Acid

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Table S1. Summary of mass features shown in the heat map in **Figure 1**.

| Mass ID | <i>m/z</i> _RT |
|---------|----------------|
| MF_1 | 200.11_3.48 |
| MF_2 | 216.10_2.90 |
| MF_3 | 444.18_3.50 |
| MF_4 | 369.13_3.33 |
| MF_5 | 277.22_4.51 |
| MF_6 | 300.17_2.44 |
| MF_7 | 317.21_4.49 |
| MF_8 | 327.22_4.54 |
| MF_9 | 353.23_3.25 |
| MF_10 | 367.36_6.12 |
| MF_11 | 375.23_3.32 |
| MF_12 | 385.37_6.12 |
| MF_13 | 388.28_4.01 |
| MF_14 | 394.32_4.67 |
| MF_15 | 395.39_6.93 |
| MF_16 | 402.39_6.11 |
| MF_17 | 413.38_5.86 |
| MF_18 | 429.37_6.65 |
| MF_19 | 549.13_2.65 |
| MF_20 | 558.30_4.15 |
| MF_21 | 569.31_3.51 |
| MF_22 | 635.41_4.39 |
| MF_23 | 659.39_3.77 |
| MF_24 | 781.53_4.35 |
| MF_25 | 803.58_4.33 |
| MF_26 | 954.62_3.61 |
| MF_27 | 388.28_4.01 |
| MF_28 | 402.39_6.11 |
| MF_29 | 429.37_6.66 |
| MF_30 | 549.13_2.65 |
| MF_31 | 558.30_4.15 |
| MF_32 | 659.39_3.77 |
| MF_33 | 781.53_4.35 |
| MF_34 | 954.62_3.61 |
| MF_35 | 300.17_2.44 |
| MF_36 | 377.29_4.66 |

Abbreviations: MF: mass feature; *m/z*: mass to charge ratio; RT: retention time

Table S2. One-way ANOVA Test (Primarolide A).

| | DF | SUM SQ | MEAN SQ | F-VALUE | Pr>F |
|-----------|----|-----------|-----------|---------|--------------|
| Treatment | 4 | 1.347e+12 | 3.367e+11 | 29.44 | 3.94e-09 *** |
| Residuals | 25 | 2.859e+11 | 1.144e+10 | | |

Significant codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 '' 1. Abbreviations DF = degree of freedom, SUM SQ = sum of squares, Mean SQ = mean squares, F-VALUE = Fisher statistics value (F value = 1 indicates no variability between means), Pr > F = probability of the F statistics.

Table S3. One-way ANOVA test (Primarolide B).

| | DF | SUM SQ | MEAN SQ | F-VALUE | Pr>F |
|-----------|----|-----------|-----------|---------|--------------|
| Treatment | 4 | 9.660e+11 | 2.415e+11 | 20.16 | 1.58e-07 *** |
| Residuals | 25 | 2.995e+11 | 1.198e+10 | | |

Significant codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 '' 1. Abbreviations DF = degree of freedom, SUM SQ = sum of squares, Mean SQ = mean squares, F-VALUE = Fisher statistics value (F value = 1 no variability between means), Pr > F = probability of the F statistics.

Table S4. Tukey's HSD Test (Primarolide A).

| Treatment | Difference | Lower limit | Upper limit | Adjusted P-value |
|-----------------------------------|------------|-------------|-------------|------------------|
| Solvent control versus control | 28953.97 | -152381.56 | 210289.5 | 0.9894786 |
| SAHA versus control | 52078.87 | -129256.65 | 233414.4 | 0.9143093 |
| NaCl versus control | 89961.41 | -91374.12 | 271296.9 | 0.5983612 |
| SAHA+ NaCl versus control | 567321.92 | 385986.39 | 748657.4 | 0.0000000 |
| SAHA Versus Solvent control | 23124.90 | -158210.62 | 204460.4 | 0.9955347 |
| NaCl versus solvent control | 61007.44 | -120328.09 | 242343.0 | 0.8581530 |
| SAHA+ NaCl versus Solvent control | 538367.95 | 357032.42 | 719703.5 | 0.0000000 |
| NaCl versus SAHA | 37882.53 | -143452.99 | 219218.1 | 0.9715838 |
| SAHA+ NaCl versus SAHA | 515243.04 | 333907.52 | 696578.6 | 0.0000001 |
| SAHA+ NaCl versus NaCl | 477360.51 | 296024.98 | 658696.0 | 0.0000004 |

This analysis was done using normalized peak area from treatments with SAHA, NaCl and SAHA + NaCl (n = 6). Adjusted p-value < 0.005 indicates treatments with significant variability between means.

Table S5. Tukey's HSD Test (Primarolide B).

| Treatment | Difference | Lower limit | Upper limit | Adjusted P-value |
|-----------------------------------|---------------|-------------|-------------|------------------|
| Solvent control versus control | -5.335702e-11 | -185599.2 | 185599.2 | 1.0000000 |
| SAHA versus control | 7.715992e+04 | -108439.2 | 262759.1 | 0.7395279 |
| NaCl versus control | 1.358179e-10 | -185599.2 | 185599.2 | 1.0000000 |
| SAHA+ NaCl versus control | 4.616350e+05 | 276035.8 | 647234.1 | 0.0000011 |
| SAHA Versus Solvent control | 7.715992e+04 | -108439.2 | 262759.1 | 0.7395279 |
| NaCl versus solvent control | 1.891749e-10 | -185599.2 | 185599.2 | 1.0000000 |
| SAHA+ NaCl versus Solvent control | 4.616350e+05 | 276035.8 | 647234.1 | 0.0000011 |
| NaCl versus SAHA | -7.715992e+04 | -262759.1 | 108439.2 | 0.7395279 |
| SAHA+ NaCl versus SAHA | 3.844750e+05 | 198875.9 | 570074.2 | 0.0000216 |
| SAHA+ NaCl versus NaCl | 4.616350e+05 | 276035.8 | 647234.1 | 0.0000011 |

This analysis was done using normalized peak area from treatments with SAHA, NaCl and SAHA + NaCl (n = 6). Adjusted p-value < 0.005 indicates treatments with significant variability between means.

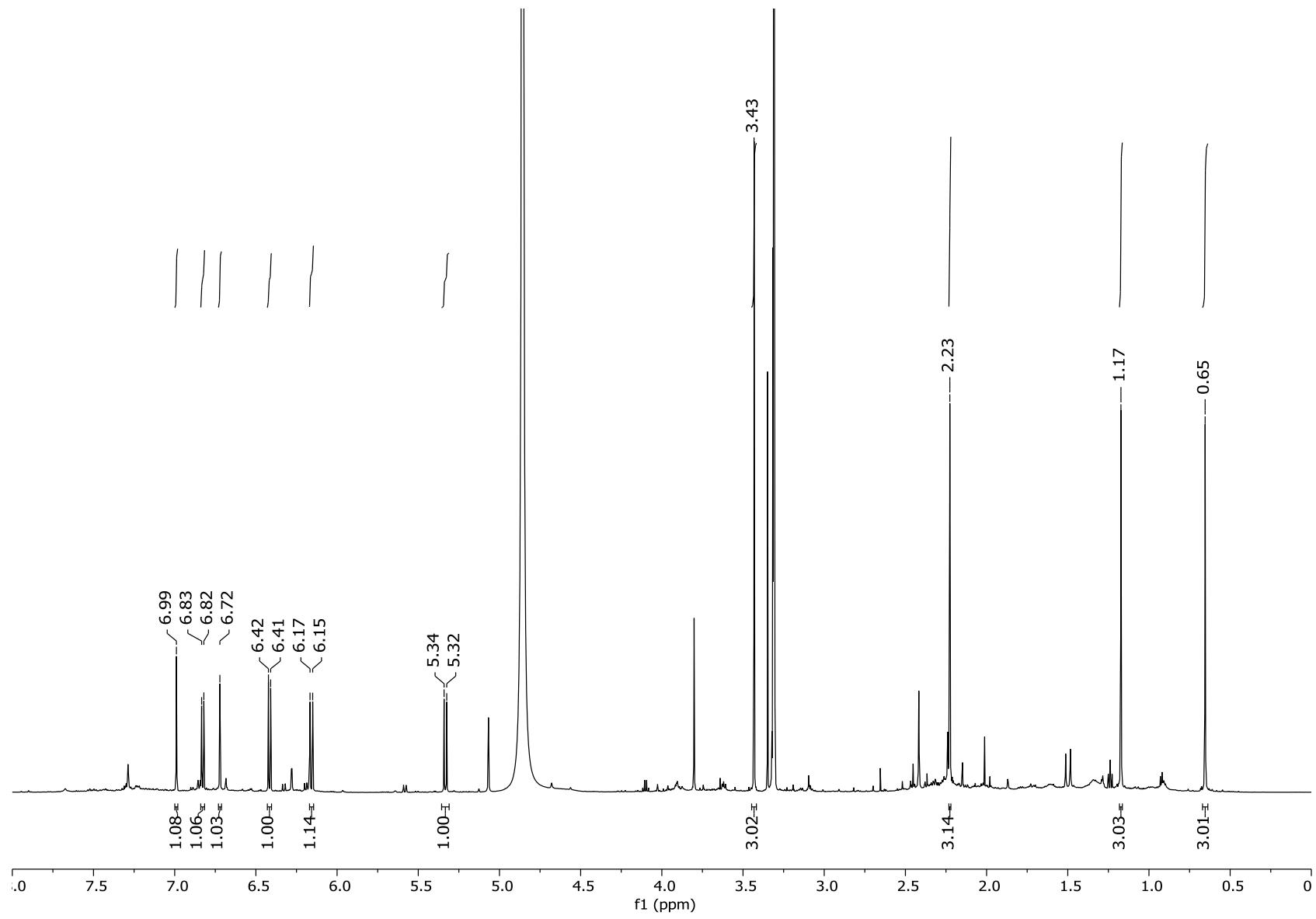


Figure S1. ^1H NMR spectrum (600 MHz, CD_3OD) of Primarolide A (**1**).

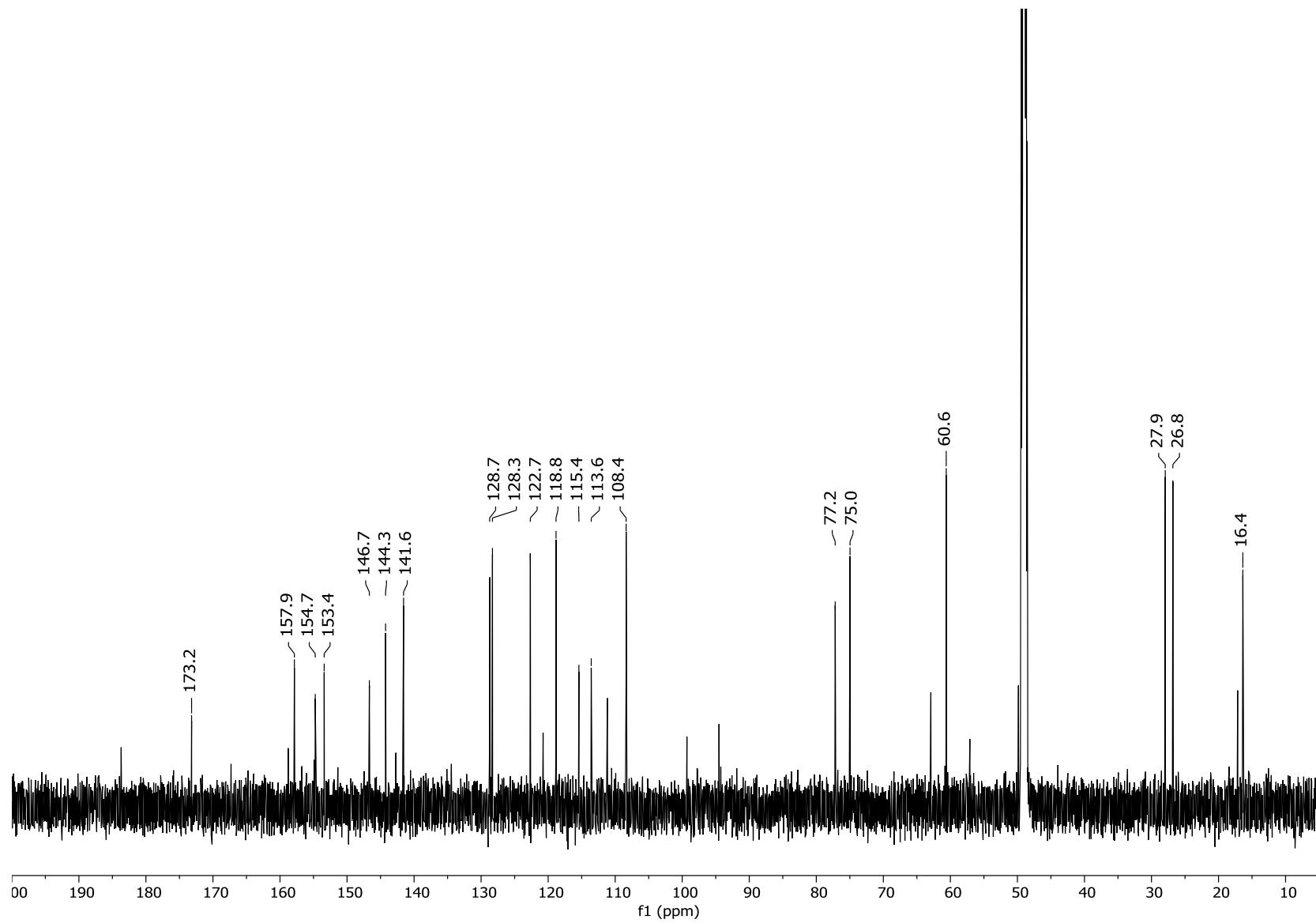


Figure S2. ^{13}C NMR spectrum (150 MHz, CD_3OD) of Primarolide A (**1**).

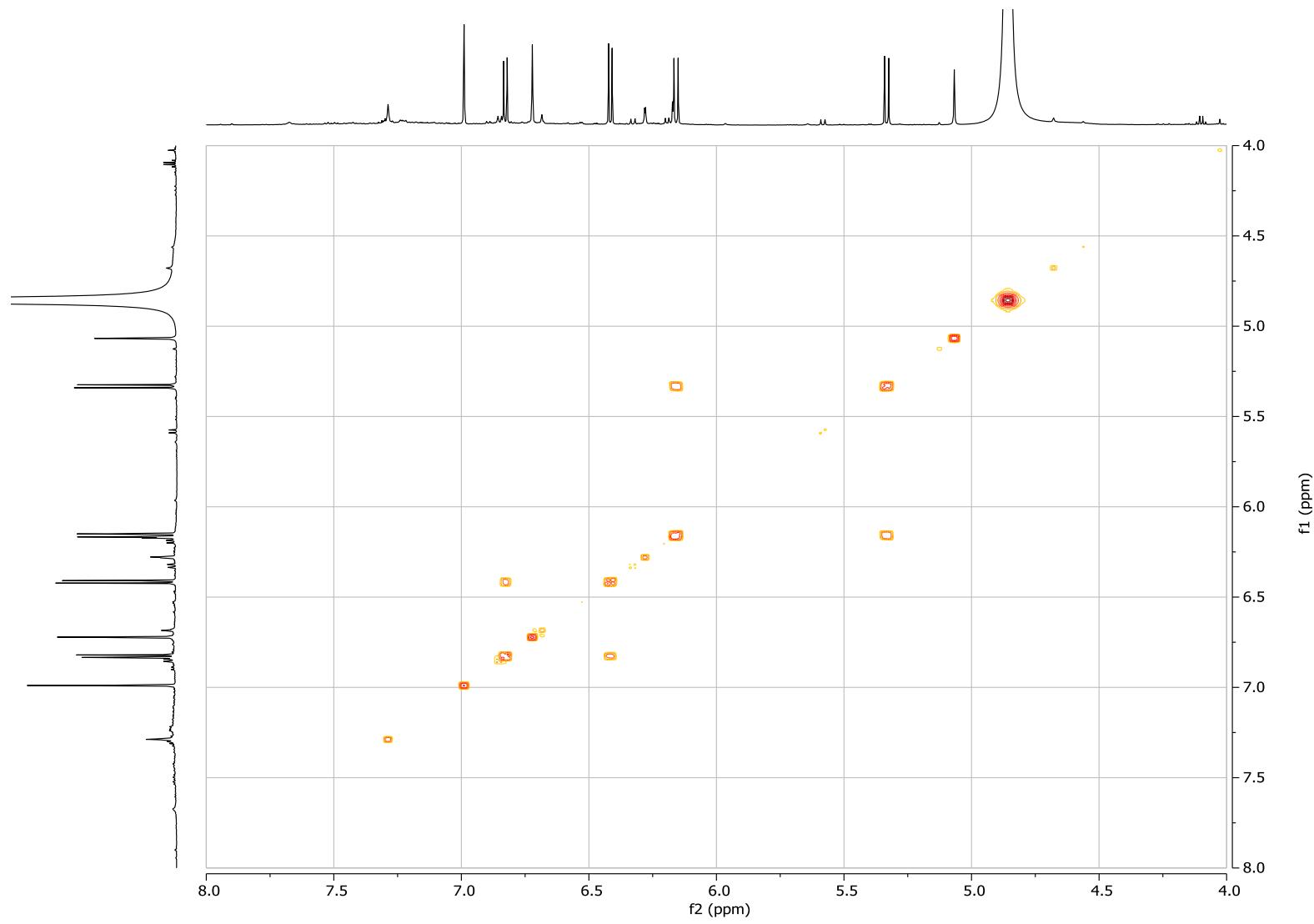


Figure S3. COSY NMR spectrum of Primarolide A (1).

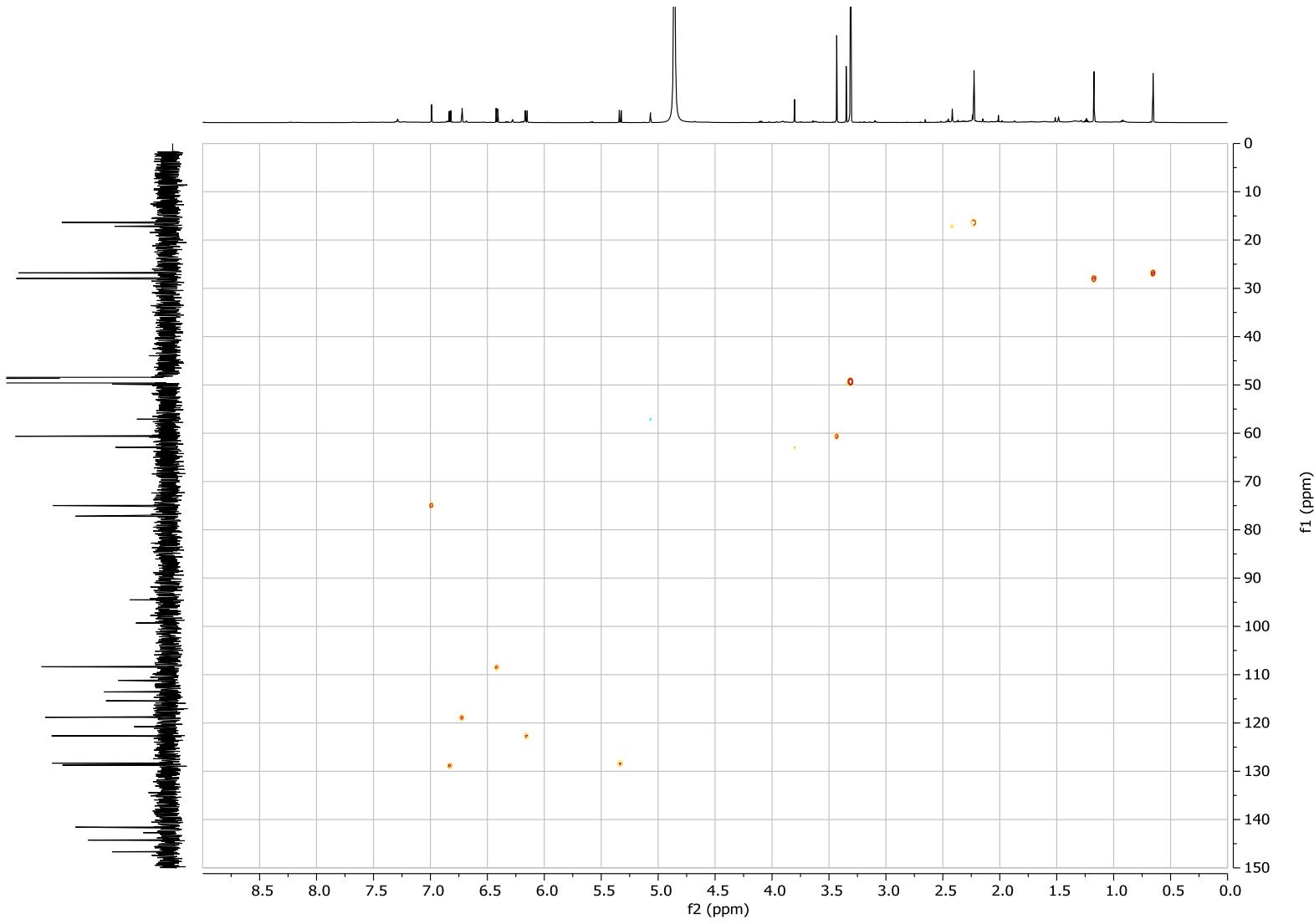


Figure S4. HSQC NMR spectrum of Primarolide A (1).

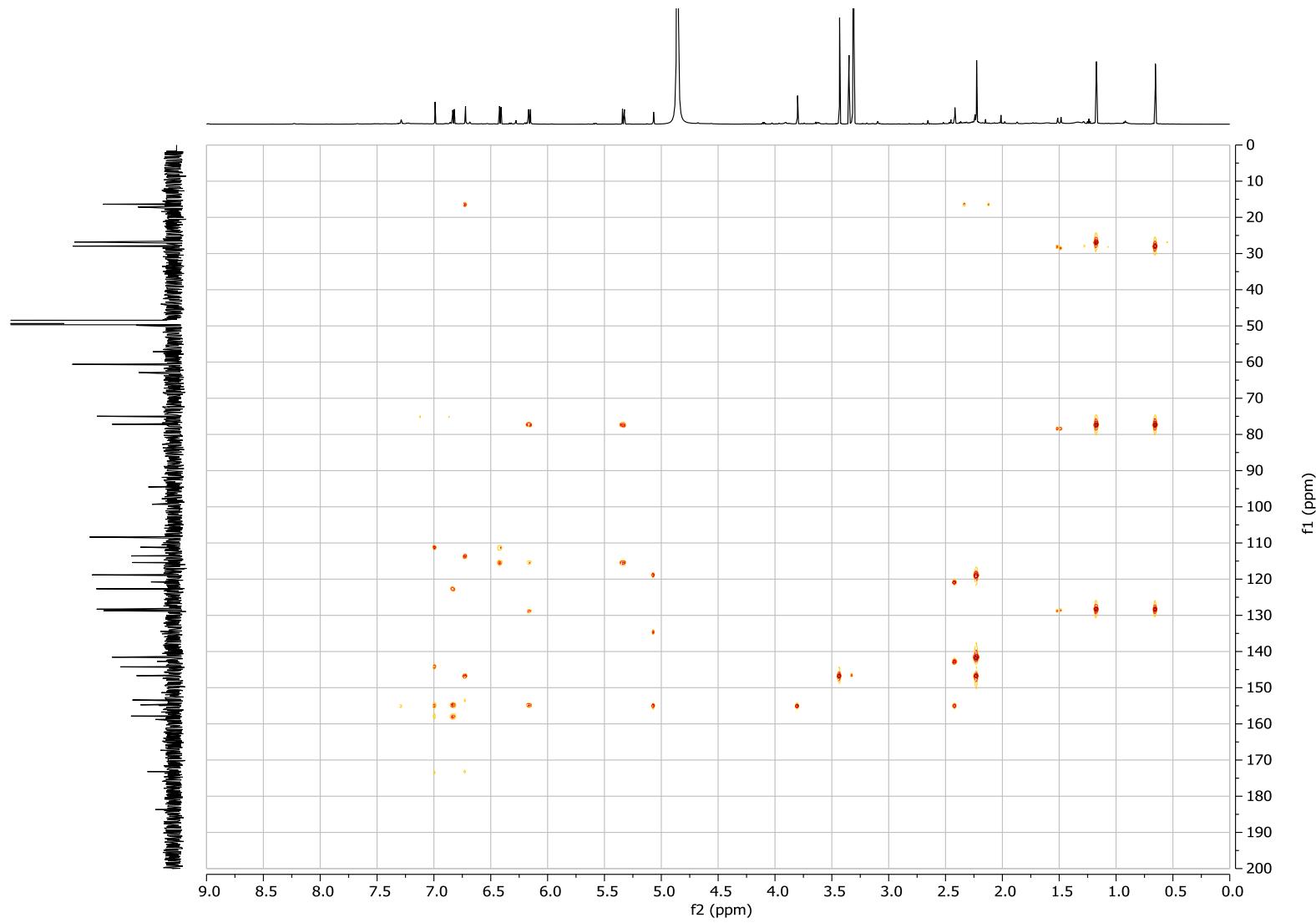


Figure S5. HMBC NMR spectrum of Primarolide A (**1**).

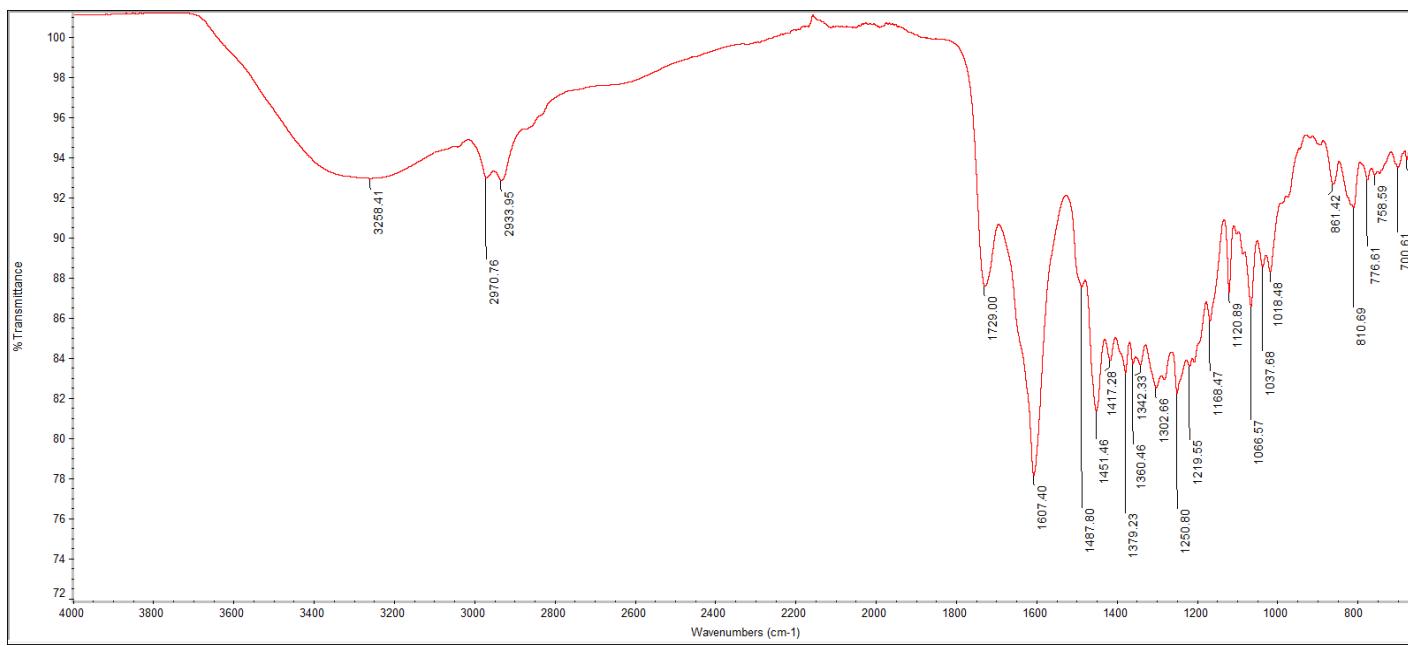


Figure S6. FTIR spectrum of Primarolide A (**1**).

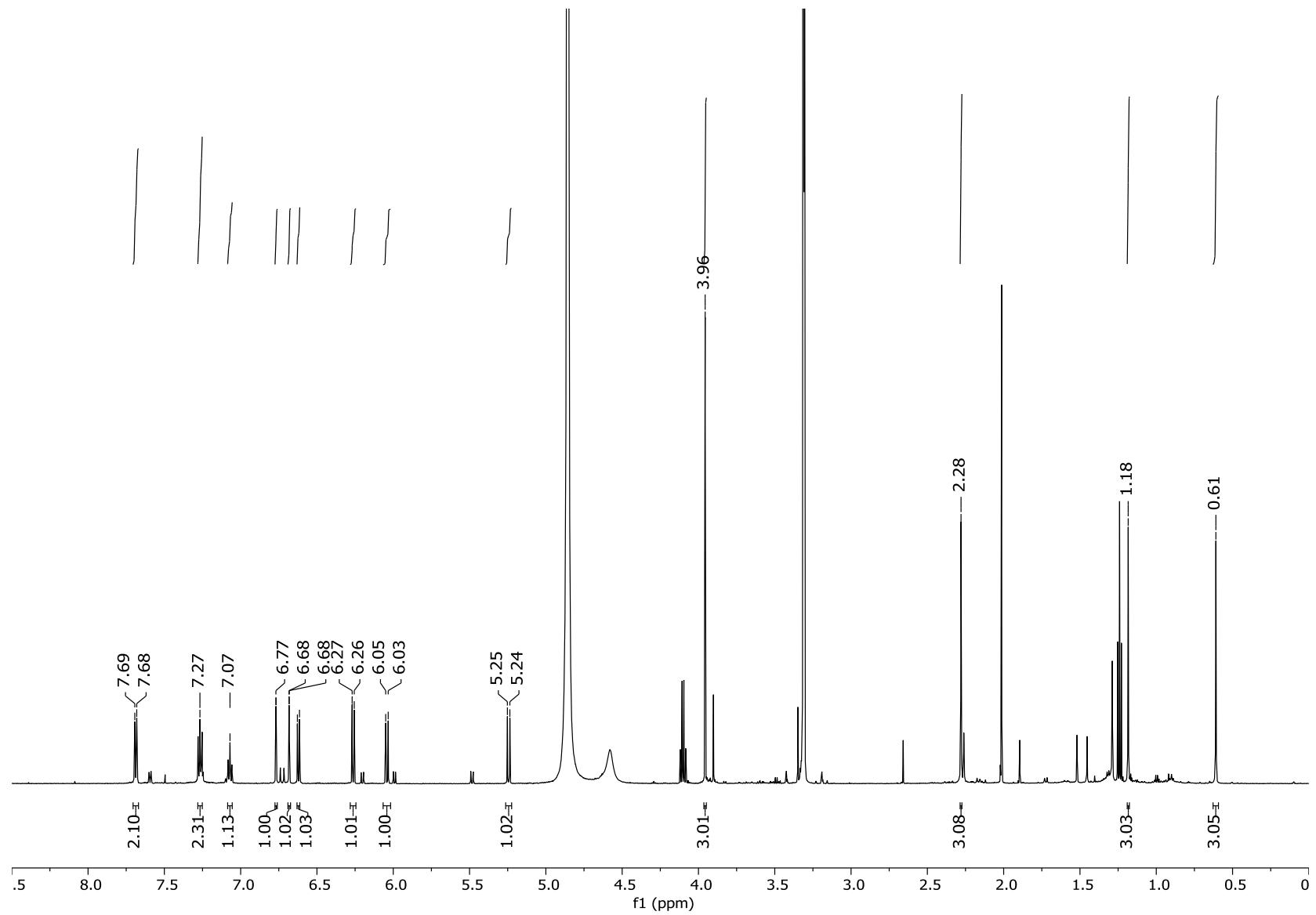


Figure S7. ^1H NMR spectrum (600 MHz, CD_3OD) of Primarolide B (2).

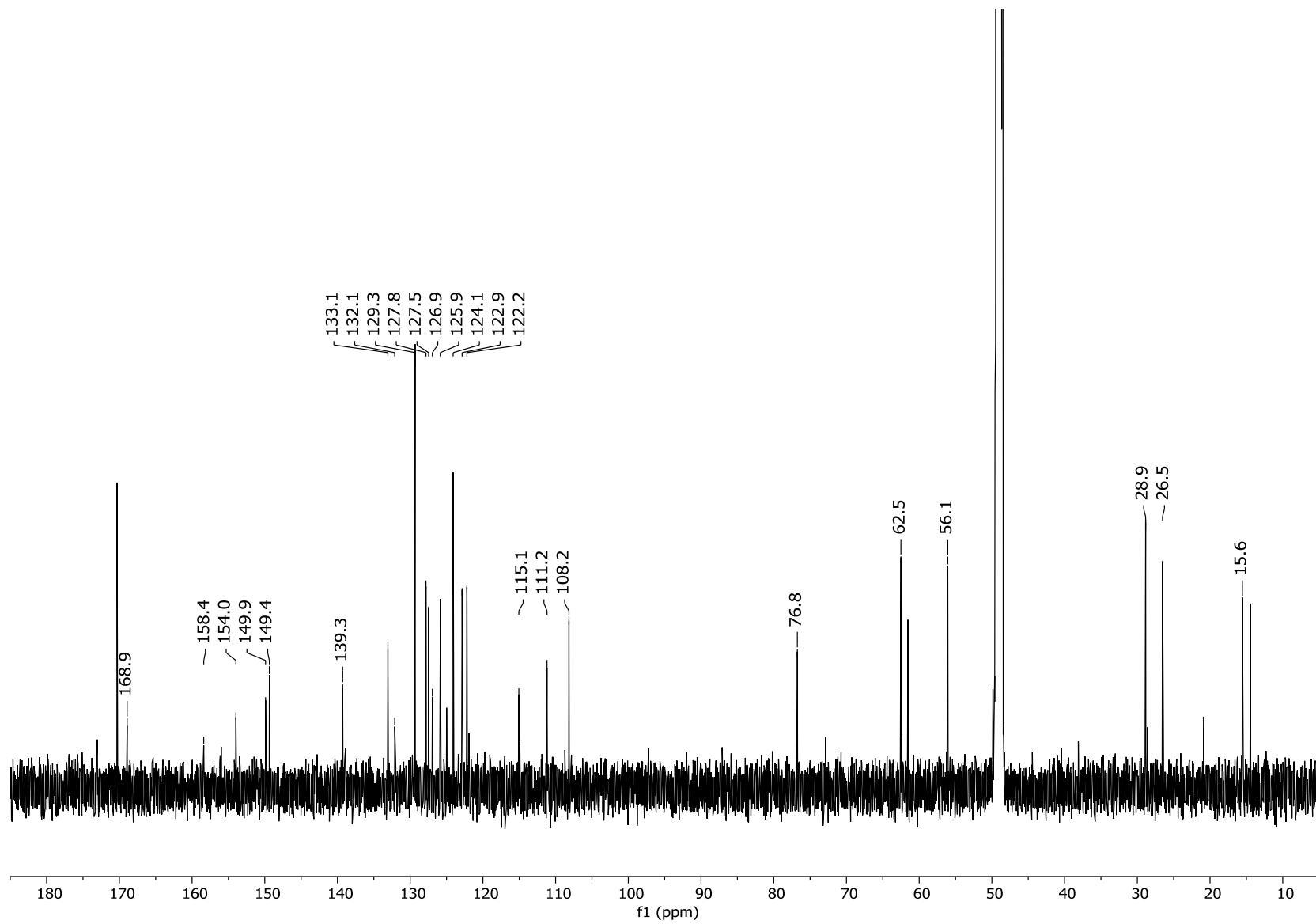


Figure S8. ^{13}C NMR spectrum (150 MHz, CD_3OD) of Primarolide B (2).

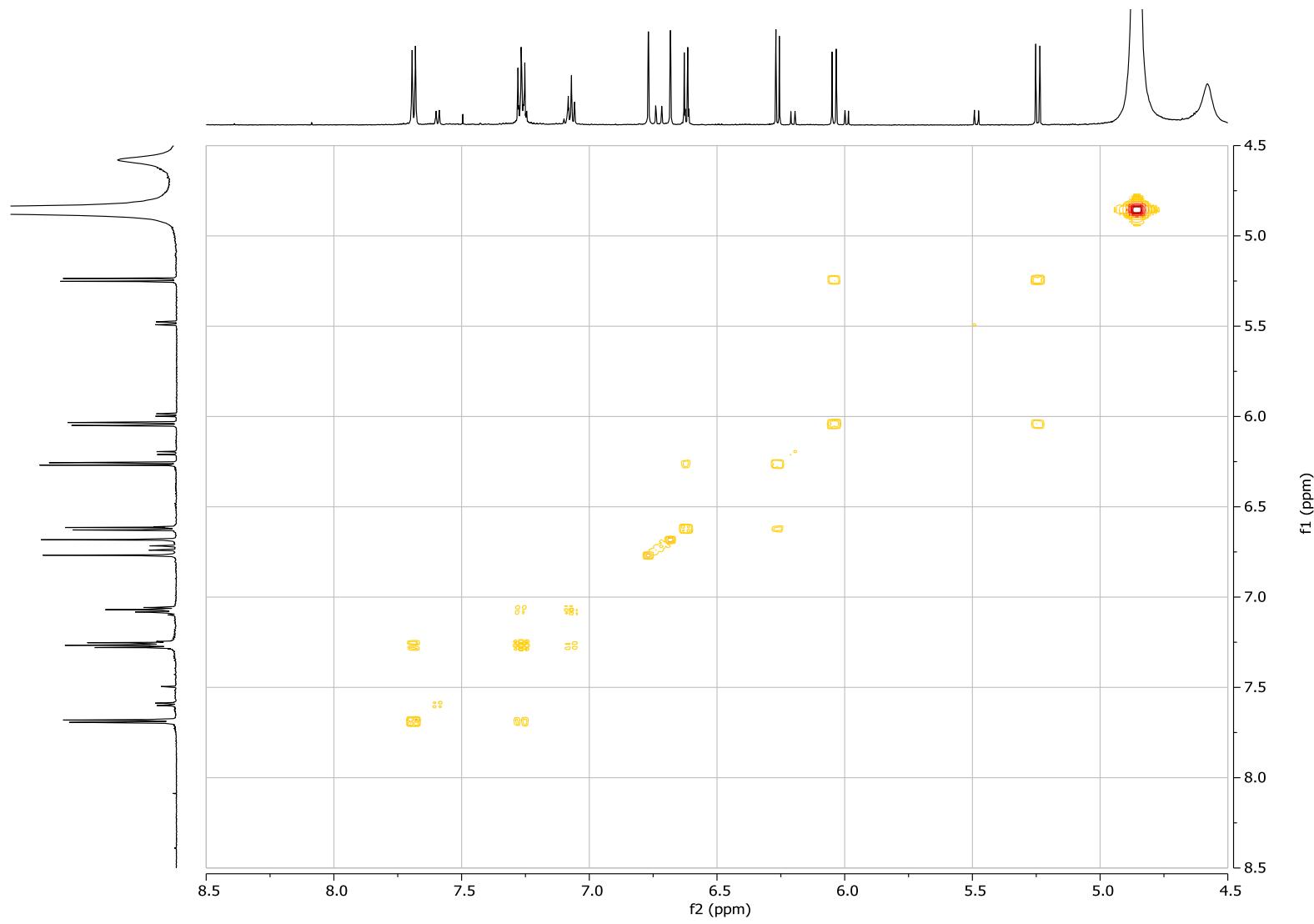


Figure S9. COSY NMR spectrum of Primarolide B (2).

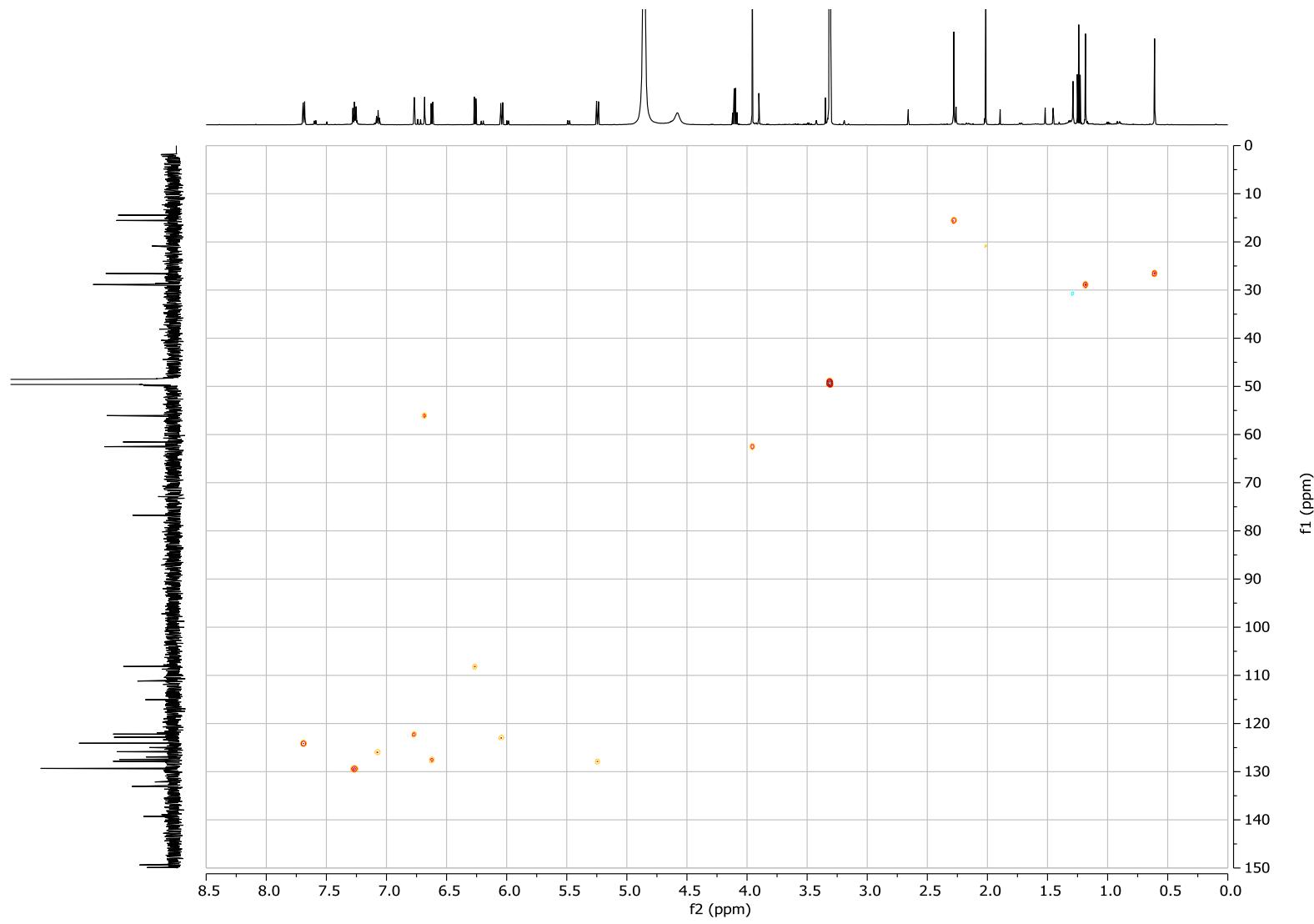


Figure S10. HSQC NMR spectrum of Primarolide B (**2**).

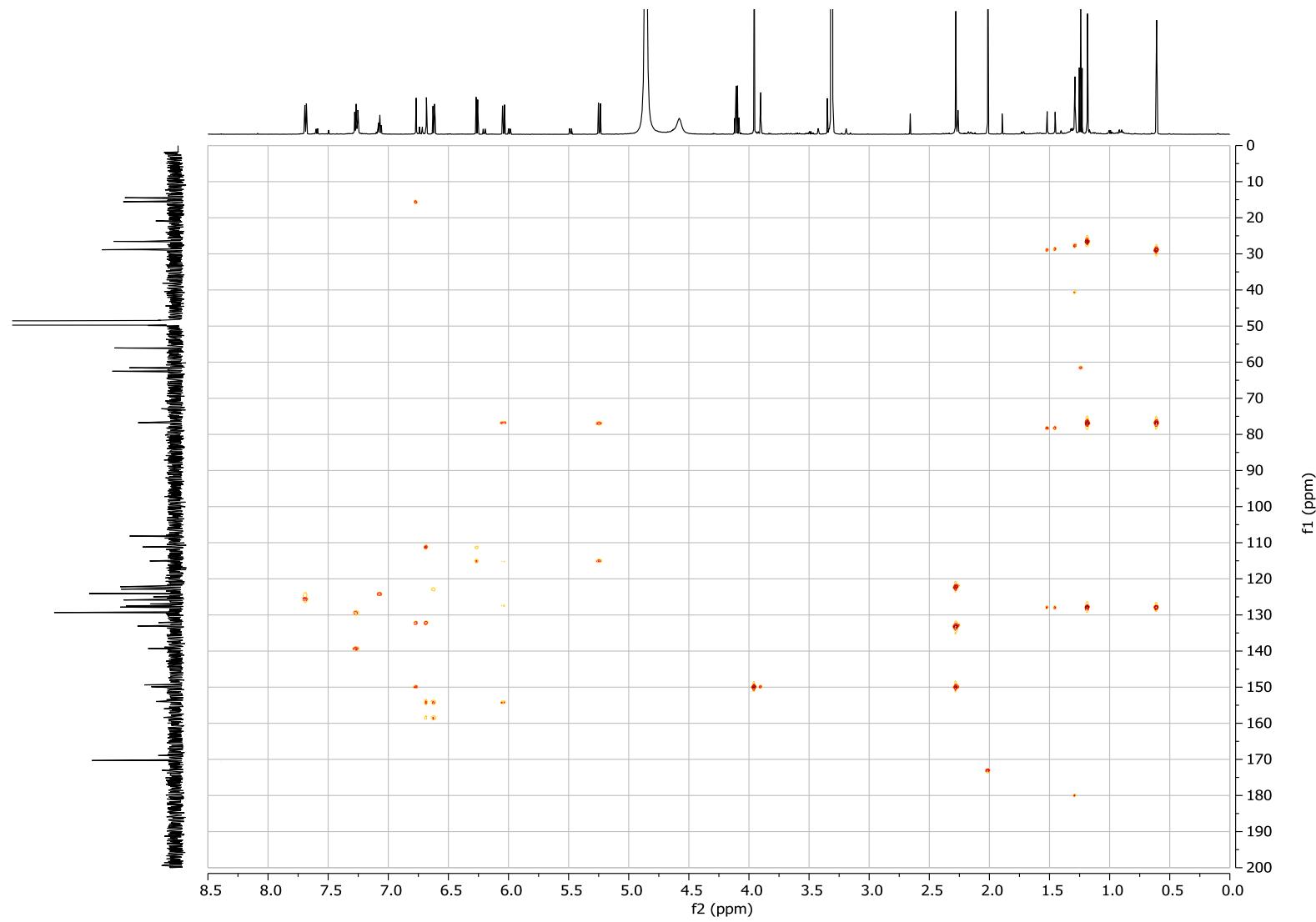


Figure S11. HMBC (8 Hz) NMR spectrum of Primarolide B (2).

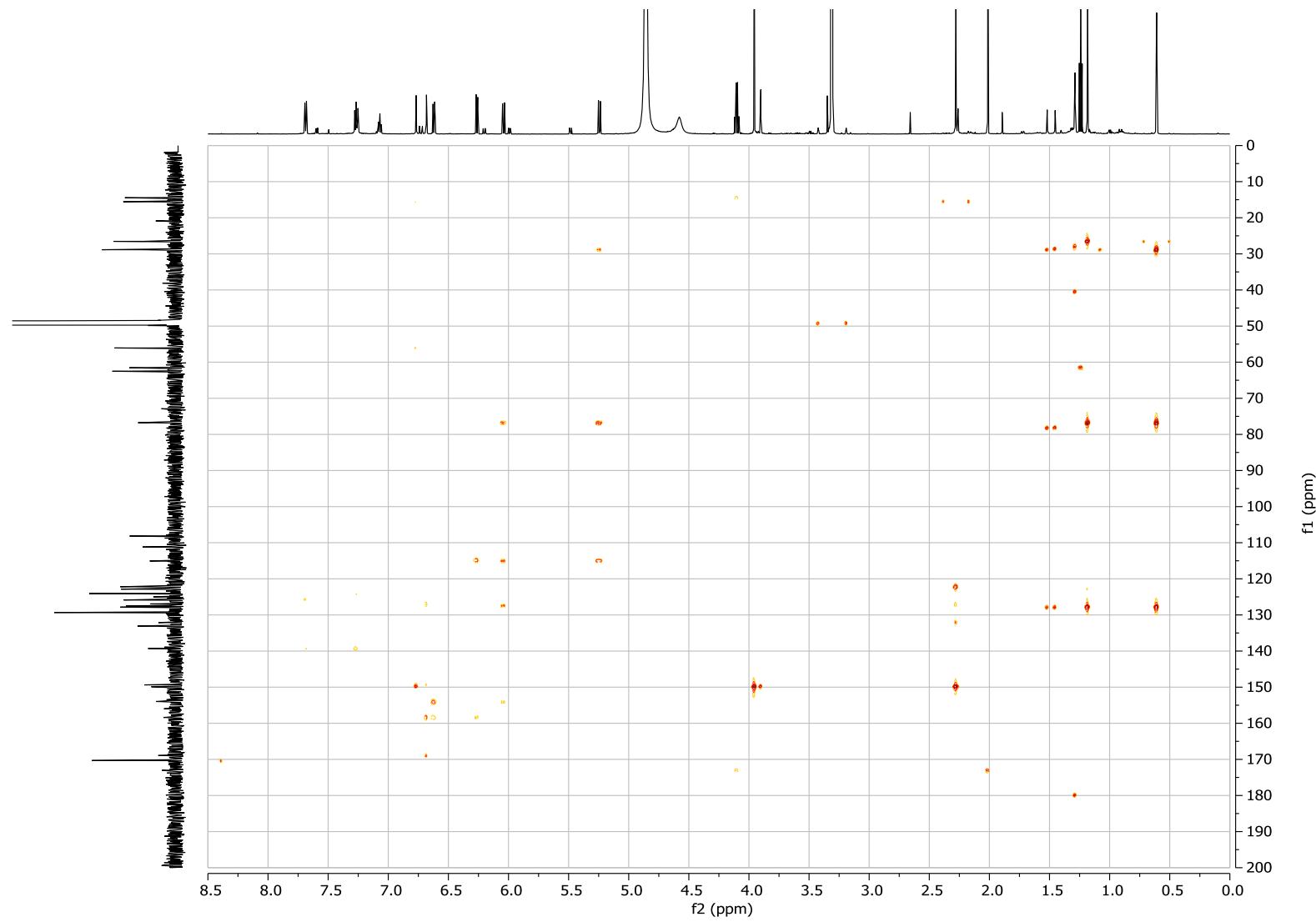


Figure S12. HMBC (3 Hz) NMR spectrum of Primarolide B (2).

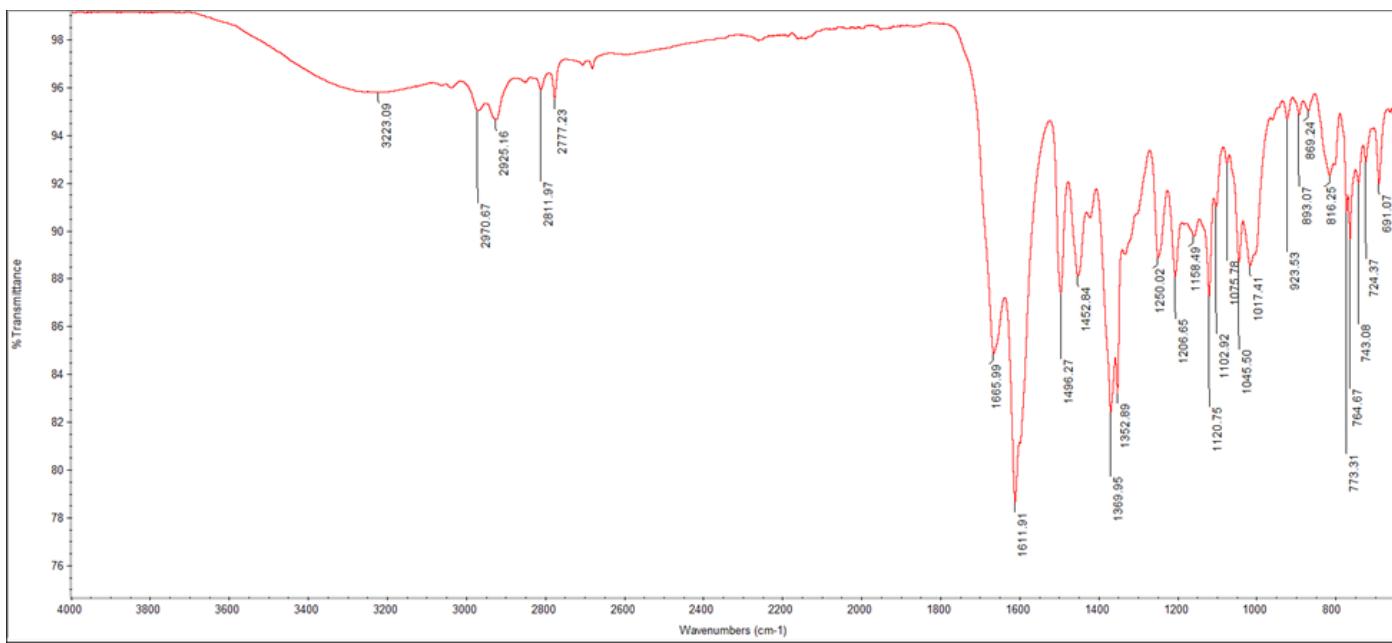


Figure S13. FTIR spectrum of Primarolide B (2).