

Supplemental Information

Novel Urea- and Piperazine-Functionalized Alepterolic Acid Derivatives: Synthesis, Anticancer Activity, and Mechanism of Action

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General Information for Chemical Synthesis

To a 500 mL round-bottom flask charged with alepterolic acid (4.99 mmol, 1.60 g, 1.00 eq) in CH₂Cl₂ (200 mL) were added *N,N*-diisopropylethylamine (DIPEA, 9.98 mmol, 1.74 mL, 2.00 eq) and HATU (7.49 mmol, 2.85 g, 1.50 eq). The mixture was stirred at room temperature with magnetic stirring for 1 h. The alepterolic acid was converted into the activated ester intermediate as confirmed by TLC (CH₂Cl₂: MeOH = 10:1). The resulting solution of activated ester intermediate was then added dropwise to a solution of piperazine (14.97 mmol, 1.29 g, 3.00 eq) in CH₂Cl₂ (200 mL). The resulted mixture was stirred at room temperature overnight. The reaction was quenched with water, extracted three times with CH₂Cl₂. The organic layer was washed once with saturated NaCl solution and then dried over anhydrous sodium sulfate and concentrated *in vacuo*. The residue was purified by column chromatography (200- 300 mesh) to yield the intermediate **13** as white solid (yield: 95%).

For synthesis of the target compounds **14a~14g** and **15a~15t**, intermediate **13** (0.15 mmol, 58.29 mg, 1.50 eq) and DIPEA (0.30 mmol, 0.05 mL, 3.00 eq) was added to the solution of the isocyanate (0.30 mmol, 3.00 eq). The reaction was carried out at room temperature with stirring for 1 h. The reaction was confirmed to be complete by TLC (CH₂Cl₂: MeOH = 10:1). The solution was purified by silica gel column chromatography (200- 300 mesh) to obtain the target products. The isocyanates were commercially purchased, except from **14a**, **15s** and **15t**. These three isocyanates prepared from the reaction of corresponding amines (0.30 mmol, 3.00 eq), triphosgene (0.10 mmol, 1.00 eq) and DIPEA (0.60 mmol, 0.10 mL, 6.00 eq) *in situ*. The reaction was conducted in ice water for 1h to obtain as crude products, which used without further purification (yield: 70-96%).

For synthesis of the target compounds **15u~15x**, compound **13** (0.15 mmol, 58.29 mg, 1.50 eq) and DIPEA (0.30 mmol, 0.05 mL, 3.00 eq) was added to the solution of the isocyanate (0.08 mmol, 0.75 eq). The next operations were similar to the synthesis of **14a** (yield: 91-96%).

Detailed Information of Cell Lines Used in This Study

A549 human non-small cell lung cancer cells, MCF-7 human breast cancer cells, and LX-2 human normal hepatic cells were purchased from Shanghai Fuheng Biotechnology Co., Ltd (Shanghai, China). HepG2 cells were purchased from National Collection of Authenticated Cell Cultures, Chinese Academy of Sciences (Shanghai, China). H460 human large cell lung cancer cells and MDA-MB-231 human triple-negative breast cancer cells were purchased from Procell Life Science & Technology Co., Ltd. (Wuhan, China). 16HBE cells are a gift from Prof. Zi Liu, originally provided

by the Cell Resource Center of the Chinese Academy of Medical Sciences (Beijing, China)

Cell Viability Assay

Cells in exponential growth phase with good viability were selected and processed to generate a homogeneous single-cell suspension. Compounds were added, followed by continued incubation. After 72 h of treatment, 20 μ L of MTT reagent was added to each well, and the incubation was prolonged for an additional 3 h. Subsequently, 100 μ L solution (10% sodium dodecyl sulfate [SDS], 5% iso-butanol, 0.01 M hydrochloric acid [HCl]) was introduced into each well. Following a 12-h incubation period, the absorbance was measured at 570 nm using a microplate reader. Cell viability, inhibition rate, and half-maximal inhibitory concentration (IC_{50}) values for each compound were subsequently calculated using Microsoft Excel and GraphPad Prism software.

Cloning Assay

A549 cells in the logarithmic growth phase were selected and processed to generate a homogeneous single-cell suspension. Following 24 h of initial incubation, compounds were added (1, 2, 4 μ M), and the incubation was continued for an additional 24 h. Subsequently, the culture medium was replaced with fresh medium, and the cells were further cultured. Fresh medium was renewed every 48 h until visible cell colonies emerged. After that, medium was discarded, and 1 mL of fixing agent (10% acetic acid, 10% methanol, 80% ultrapure water) was added to each well. After 15 min, the fixative was removed, and each well was rinsed with 1 mL PBS. Thereafter, 1 mL of crystal violet staining solution was added to each well. After 1 min, the crystal violet solution was recovered. The 6-well plate was gently rinsed for 30 s, then inverted on a clean bench and air-dried naturally. Images of the cell colonies were captured, and data analysis was performed using ImageJ and GraphPad Prism 8.0 software, followed by graph plotting.

Scratch Wound Healing Assay

A 6-well plate was prepared, and three parallel lines were marked at the bottom of each well. A549 cells in good growth status were selected, and the cells were plated at a density of 3.0×10^5 per well. A homogeneous single-cell suspension was prepared and seeded into the pre-marked 6-well plate. After 24 h, the scratch wound was made. Next, compounds were added to the corresponding wells, and images were captured at the pre-marked positions. The cells were continuously cultured for an additional 72 h, after which cell migration was observed and images were captured again at the same marked positions. Image analysis was performed using ImageJ software to measure scratch widths. Statistical analysis of the data was conducted using GraphPad Prism 8.0

software, and graphs were generated. The scratch healing rate was calculated using the following formula:

$$\text{Scratch healing rate} = (\text{Scratch width at 0 h} / \text{Scratch width at 72 h}) / \text{Scratch width at 0 h} \times 100\%$$

Cell Proliferation Assay

A549 cells in good growth status were selected and prepared to be a homogeneous single-cell suspension with a density of 1.0×10^5 cells/mL. The cell suspension was seeded at a volume of 2 mL per well. Experimental groups were set up with incubation time at 24 h, 48 h, and 72 h. After 24 h, compounds were added to the corresponding groups. At 24 h, 48 h, and 72 h, cell counts were determined using a cell counter. Finally, cell growth curves were plotted using GraphPad Prism software.

Cell Morphology Observation Assay

A549 cells in good growth status were selected and processed to a homogeneous single-cell suspension. The suspension was seeded into a 6-well plate at a density of 3.0×10^5 cells with a total volume of 2 mL per well. After 24 h of incubation, compounds at different concentrations were added to the respective wells. Following an additional 24 h of continuous incubation, the cells were observed under a phase-contrast microscope, and images were captured.

Nuclear Morphology Observation Assay

A549 cells in good growth status were selected and processed to a homogeneous single-cell suspension. The suspension was seeded into culture wells at a density of 2.0×10^5 cells with a total volume of 2 mL per well. After 24 h of incubation, compounds were added, and the incubation was continued for an additional 24 h. Subsequently, the supernatant was discarded, and 1 mL of cell staining buffer was added to each well, followed by the addition of 5 μ L of Hoechst staining solution. The mixture was gently mixed and incubated on ice or at 4 °C for 20–30 min. After incubation, the cells were washed with PBS. One drop of fluorescence enhancer was added, and the cell-bearing coverslip was then inverted onto the slide. The samples were observed under a fluorescence microscope, and images were captured.

Cell Cycle Detection Assay

A549 cells in good growth status were selected and processed to a homogeneous single-cell suspension. The suspension was seeded into wells at a density of 3.0×10^5 cells with a total volume of 2 mL per well. After 24 h, compounds at different concentrations were added, and the incubation was continued for an additional 24 h. Subsequently, cell cycle detection was performed strictly following the manufacturer's

instructions provided by the cell cycle detection kit (Beyotime Biotechnology Co., Ltd.). Data analysis was performed using GraphPad Prism 8.0 software, and graphs were generated.

Apoptosis Detection Assay

A549 cells in good growth status were selected and processed to a homogeneous single-cell suspension. The suspension was seeded into wells at a density of 3.0×10^5 cells with a total volume of 2 mL per well. After 24 h of initial incubation, compounds were added, and the incubation was cultured for an additional 24 h. Subsequently, the staining solution was prepared in strict accordance with the instructions provided with the Beyotime Apoptosis Detection Kit, followed by the performance of relevant subsequent staining and detection procedures.

Mitochondrial Membrane Potential Detection Assay

A549 cells in good growth status were selected and processed to a homogeneous single-cell suspension. The suspension was seeded into wells at a density of 3.0×10^5 cells with a total volume of 2 mL per well, followed by initial incubation. After 24 h, test compounds were added, and incubation was continued for 24 h. Subsequently, carbonyl cyanide 3-chlorophenylhydrazone (CCCP), a positive control agent, was added to the positive control wells at a dilution ratio of 1:1000 for 20 min. Concurrently, relevant staining solutions were prepared and subsequent staining operations were performed in strict accordance with the instructions provided with the Beyotime Mitochondrial Membrane Potential Detection Kit.

Western blotting

Proteins were extracted using SDS lysis buffer and subjected to sodium dodecyl sulfate-polyacrylamide gel electrophoresis (SDS-PAGE). Subsequently, proteins separated on the gel were transferred onto a nitrocellulose membrane (NC membrane). The membrane was blocked with 5% non-fat milk in Tris-buffered saline containing Tween 20 (TBST). The aforementioned antibodies were used as primary antibodies for membrane probing, followed by incubation with a horseradish peroxidase (HRP)-conjugated secondary antibody. Target proteins on the membrane were visualized using an enhanced chemiluminescence (ECL) kit (Proteintech Group, Inc.). The density of the protein bands was quantitatively analyzed using ImageJ software.

ADMET Analysis

The physicochemical properties and pharmacokinetic profiles of the target compounds were evaluated using two online web databases: ADMET Lab 3.0 (<https://admetlab3.scbdd.com/>) and SwissADME (<http://www.swissadme.ch/>). These

properties primarily include five key aspects of drug disposition: absorption, distribution, metabolism, excretion, and toxicity (ADMET).

Protein Docking

The apoptotic protein structures were retrieved on online databases, including UniProt (<https://www.uniprot.org/>) and RCSB PDB (<https://www.rcsb.org/>). Compound structures were drawn using ChemDraw 20.0 software. AutoDock Tools software was employed to perform pre-docking optimization of both compound and apoptotic proteins, involving water molecule removal and hydrogenation; subsequently, the docking free energy was calculated. The binding modes of compound with apoptotic proteins were visualized using PyMOL software, and the resulting images were edited using PowerPoint software.

- 1□. Protein and PDB ID: Caspase-3 (1GFW) 、Bax (4BD6) 、Bcl-2 (1G5M) 、Caspase-9 (2AR9) 、PARP-1 (4OPX)
- 2□. Software: AutoDockTools-1.5.7
- 3□. Scoring function: Vina default scoring
- 4□. Exhaustiveness: 8
- 5□. Flexible handling strategy: Rigid frame, flexible side chain
- 6□. Website for protein structures: <https://www.uniprot.org/>;<https://www.rcsb.org/>
- 7□. Treatment of small molecule ligands: Minimum binding free energy (ChemDraw 20.0; Chem3D 20.0)
- 8□. Grid spacing: 0.375 Å
- 9□. Clustering algorithms and clustering thresholds: No built-in clustering
- 10□. CPU model: Intel® Core (TM) i5-3210M CPU @ 2.50 GHZ 2.50GHZ
- 11□. OS version: Win_64 flagship version
- 12□. Connect pocket parameters (See Table 1 and Table 2)

Table 1. the connect pocket parameters of compound **15o**

Caspase-3	Bax	Bcl-2	Caspase-9	PARP-1
center x = 26.71	center x = -16.34	center x = 3.061	center x = 23.957	center x =
center y = 22.591	center y = -9.58	center y = -9.58	center y = 33.011	17.644
center z = 37.128	center z = 8.335	center z = 2.734	center z = 27.123	center y =
size x = 66.15	size x = 70.7	size x = 68.46	size x = 111.16	33.011
size y = 66.15	size y = 70.7	size y = 68.46	size y = 111.16	center z =
size z = 66.15	size z = 70.7	size z = 68.46	size z = 111.16	31.626
		size z = 68.46	size z = 111.16	size x = 126.0

size y = 126.0

size z = 126.0

Table 2. the connect pocket parameters of compound **15u**

Caspase-3	Bax	Bcl-2	Caspase-9	PARP-1
center x = 26.71	center x = -16.94	center x = 1.954	center x = 19.547	center x = 19.547
center y = 22.591	center y = -16.647	center y = -3.058	center y = 38.445	center y = 51.59
center z = 37.128	center z = -0.94	center z = 3.202	center z = 23.65	center z = 36.625
size x = 53.9	size x = 69.3	size x = 54.46	size x = 105.0	size x = 126.0
size y = 53.9	size y = 69.3	size y = 54.46	size y = 105.0	size y = 126.0
size z = 53.9	size z = 69.3	size z = 54.46	size z = 105.0	size z = 126.0

Data Processing and Statistical Analysis

All experiments were performed in triplicate at minimum. Image processing was conducted using Photoshop software, while cell count and quantitative analysis of band densities were performed using ImageJ software. Data collation and statistical analysis were carried out using GraphPad Prism 8.0 software, with statistical graphs generated simultaneously. Statistical comparisons were performed using the t-test. In the graphs, statistical significance was denoted as follows: * $p < 0.05$, ** $p < 0.01$, and *** $p < 0.001$.

Structural characterization of synthetic compounds

N-Butyl-4-((*E*)-5-((1*R*,4*aS*,6*R*,8*aS*)-6-hydroxy-5,5,8*a*-trimethyl-2-methylenedecahydronaphthalen-1-yl)-3-methylpent-2-enoyl)

piperazine-1-

carboxamide (**14a**)

Colorless oil, yield 70%, ¹H NMR (400 MHz, CDCl₃) δ 5.72 (s, 1H), 4.86 (s, 1H), 4.51 (s, 1H), 4.49 (t, *J* = 5.2 Hz, 1H), 3.71 – 3.59 (m, 2H), 3.53 – 3.46 (m, 2H), 3.46 – 3.38 (m, 2H), 3.38 – 3.28 (m, 2H), 3.28 – 3.18 (m, 3H), 2.40 (ddd, *J* = 12.8, 4.2, 2.4 Hz, 1H), 2.29 – 2.20 (m, 1H), 1.99 – 1.89 (m, 2H), 1.88 (s, 3H), 1.80 – 1.55 (m, 7H), 1.49 (t, *J* = 6.6 Hz, 3H), 1.40 – 1.30 (m, 3H), 1.15 (td, *J* = 13.1, 3.8 Hz, 1H), 1.06 (dd, *J* = 12.5, 2.8 Hz, 1H), 0.98 (s, 3H), 0.92 (t, *J* = 7.3 Hz, 3H), 0.76 (s, 3H), 0.68 (s, 3H). ¹³C NMR (101 MHz, CDCl₃) δ 167.54, 157.60, 150.94, 147.75, 116.78, 106.80, 78.72, 55.92, 54.67, 45.94, 43.91, 43.60, 40.75, 40.65, 39.41, 39.13, 38.57, 38.17, 37.09, 32.30, 28.30, 27.87, 23.99, 21.57, 20.09, 18.72, 15.41, 14.52, 13.81. HRMS calc for C₂₉H₅₀N₃O₃ [M+H]⁺: 488.3852, found 488.3854. HPLC purity: 97.86%, retention time: 12.640 min.

N-(tert-Butyl)-4-((*E*)-5-((1*R*,4*aS*,6*R*,8*aS*)-6-hydroxy-5,5,8*a*-trimethyl-2-methylenedecahydronaphthalen-1-yl)-3-methylpent-2-enoyl) piperazine-1-carboxamide (**14b**)

Colorless oil, yield 95%, ¹H NMR (400 MHz, CDCl₃) δ 5.70 (s, 1H), 4.84 (s, 1H), 4.49 (s, 1H), 4.33 (s, 1H), 3.67 – 3.58 (m, 2H), 3.50 – 3.42 (m, 2H), 3.38 – 3.31 (m, 2H), 3.30 – 3.23 (m, 2H), 3.21 (dd, *J* = 11.6, 4.4 Hz, 1H), 2.38 (dt, *J* = 12.3, 3.2 Hz, 1H), 2.26 – 2.18 (m, 1H), 1.97 – 1.89 (m, 2H), 1.86 (s, 3H), 1.77 – 1.51 (m, 7H), 1.39 (qd, *J* = 12.9, 4.2 Hz, 1H), 1.32 (s, 9H), 1.12 (td, *J* = 13.1, 3.6 Hz, 1H), 1.04 (dd, *J* = 12.5, 2.7 Hz, 1H), 0.96 (s, 3H), 0.74 (s, 3H), 0.66 (s, 3H). ¹³C NMR (101 MHz, Chloroform-*d*) δ 167.53, 156.74, 150.83, 147.75, 116.80, 106.79, 78.68, 55.90, 54.67, 50.98, 45.92, 44.07, 43.42, 40.74, 39.40, 39.13, 38.55, 38.16, 37.10, 29.40, 28.33, 27.86, 23.99, 21.56, 18.71, 15.43, 14.50. HRMS calc for C₂₉H₅₀N₃O₃ [M+H]⁺: 488.3852, found 488.3856. HPLC purity: 98.66%, retention time: 12.953 min.

N-Cyclopentyl-4-((*E*)-5-((1*R*,4*aS*,6*R*,8*aS*)-6-hydroxy-5,5,8*a*-trimethyl-2-methylenedecahydronaphthalen-1-yl)-3-methylpent-2-enoyl) piperazine-1-carboxamide (**14c**)

Colorless oil, yield 91%, ¹H NMR (400 MHz, CDCl₃) δ 5.69 (s, 1H), 4.83 (s, 1H), 4.53 (s, 1H), 4.48 (s, 1H), 4.07 – 4.01 (m, 1H), 3.69 – 3.57 (m, 2H), 3.51 – 3.43 (m, 2H), 3.42 – 3.36 (m, 2H), 3.33 – 3.24 (m, 2H), 3.21 (dd, *J* = 11.6, 4.3 Hz, 1H), 2.37 (ddd, *J* = 12.8, 4.2, 2.4 Hz, 1H), 2.26 – 2.18 (m, 1H), 1.99 – 1.86 (m, 4H), 1.84 (s, 3H), 1.77 – 1.49 (m, 11H), 1.39 – 1.27 (m, 3H), 1.12 (td, *J* = 13.2, 3.9 Hz, 1H), 1.03 (dd, *J* = 12.6, 2.7 Hz, 1H), 0.95 (s, 3H), 0.73 (s, 3H), 0.65 (s, 3H). ¹³C NMR (101 MHz, CDCl₃) δ 167.60, 157.39, 150.91, 147.78, 116.79, 106.82, 78.67, 55.93, 54.70, 52.70, 45.98, 43.91, 43.60, 40.79, 39.43, 39.16, 38.57, 38.20, 37.14, 33.53, 28.36, 27.87, 24.03, 23.72, 21.59, 18.77, 15.49, 14.54. HRMS calc for C₃₀H₅₀N₃O₃ [M+H]⁺: 500.3852, found 500.3855. HPLC purity: 98.66%, retention time: 12.953 min.

N-Cyclohexyl-4-((*E*)-5-((1*R*,4*aS*,6*R*,8*aS*)-6-hydroxy-5,5,8*a*-trimethyl-2-methylenedecahydronaphthalen-1-yl)-3-methylpent-2-enoyl) piperazine-1-carboxamide (**14d**)

Colorless oil, yield, 92%, ^1H NMR (400 MHz, CDCl_3) δ 5.69 (s, 1H), 4.82 (s, 1H), 4.47 (s, 1H), 4.44 (s, 1H), 3.68 – 3.54 (m, 3H), 3.52 – 3.43 (m, 2H), 3.42 – 3.34 (m, 2H), 3.33 – 3.25 (m, 2H), 3.20 (dd, $J = 11.7, 4.4$ Hz, 1H), 2.37 (ddd, $J = 12.8, 4.2, 2.4$ Hz, 1H), 2.26 – 2.15 (m, 1H), 1.96 – 1.85 (m, 4H), 1.84 (s, 3H), 1.78 – 1.44 (m, 10H), 1.38 – 1.25 (m, 3H), 1.16 – 1.06 (m, 3H), 1.03 (dd, $J = 12.5, 2.3$ Hz, 2H), 0.95 (s, 3H), 0.73 (s, 3H), 0.65 (s, 3H). ^{13}C NMR (101 MHz, CDCl_3) δ 167.63, 156.92, 150.92, 147.77, 116.77, 106.82, 78.67, 55.91, 54.68, 49.66, 45.97, 43.88, 43.59, 40.78, 39.42, 39.16, 38.55, 38.19, 37.12, 33.92, 28.36, 27.84, 25.65, 25.11, 24.01, 21.57, 18.78, 15.50, 14.54. HRMS calc for $\text{C}_{31}\text{H}_{52}\text{N}_3\text{O}_3$ $[\text{M}+\text{H}]^+$: 514.4009, found 514.4008. HPLC purity: 100.00%, retention time: 13.993 min.

N-Dodecyl-4-((*E*)-5-((1*R*,4*aS*,6*R*,8*aS*)-6-hydroxy-5,5,8*a*-trimethyl-2-methylenedecahydronaphthalen-1-yl)-3-methylpent-2-enoyl) piperazine-1-carboxamide (**14e**)

Light yellow oil, yield 88%, ^1H NMR (400 MHz, CDCl_3) δ 5.70 (s, 1H), 4.84 (s, 1H), 4.58 (t, $J = 5.5$ Hz, 1H), 4.50 (s, 1H), 3.67 – 3.58 (m, 2H), 3.51 – 3.44 (m, 2H), 3.44 – 3.37 (m, 2H), 3.35 – 3.27 (m, 2H), 3.26 – 3.16 (m, 3H), 2.39 (ddd, $J = 12.8, 4.2, 2.4$ Hz, 1H), 2.28 – 2.19 (m, 1H), 1.95 – 1.89 (m, 2H), 1.86 (s, 3H), 1.78 – 1.47 (m, 9H), 1.37 (qd, $J = 12.9, 4.2$ Hz, 1H), 1.29 – 1.21 (m, 18H), 1.13 (td, $J = 13.1, 3.9$ Hz, 1H), 1.05 (dd, $J = 12.5, 2.7$ Hz, 1H), 0.97 (s, 3H), 0.85 (t, $J = 6.7$ Hz, 3H), 0.75 (s, 3H), 0.67 (s, 3H). ^{13}C NMR (101 MHz, CDCl_3) δ 167.61, 157.68, 150.99, 147.83, 116.86, 106.87, 78.76, 56.00, 54.76, 46.02, 44.00, 43.68, 41.16, 40.83, 39.49, 39.22, 38.65, 38.25, 37.19, 31.99, 30.31, 29.69, 29.43, 28.41, 27.95, 27.06, 24.08, 22.76, 21.66, 18.80, 15.51, 14.58, 14.20. HRMS calc for $\text{C}_{37}\text{H}_{65}\text{N}_3\text{O}_3\text{Na}$ $[\text{M} + \text{Na}]^+$: 622.4924, found 622.4926. HPLC purity: 97.06%, retention time: 16.613 min.

N-(2-Chloroethyl)-4-((*E*)-5-((1*R*,4*aS*,6*R*,8*aS*)-6-hydroxy-5,5,8*a*-trimethyl-2-methylenedecahydronaphthalen-1-yl)-3-methylpent-2-enoyl) piperazine-1-carboxamide (**14f**)

Colorless oil, yield 90%, ^1H NMR (400 MHz, CDCl_3) δ 5.70 (s, 1H), 4.83 (s, 1H), 4.48 (s, 1H), 4.01 (s, 1H), 3.77 – 3.59 (m, 4H), 3.61 – 3.32 (m, 8H), 3.21 (dd, $J = 11.7, 4.4$ Hz, 1H), 2.38 (d, $J = 13.9$ Hz, 1H), 2.27 – 2.17 (m, 1H), 1.96 – 1.87 (m, 2H), 1.85 (s, 3H), 1.77 – 1.45 (m, 7H), 1.36 (qd, $J = 12.9, 4.2$ Hz, 1H), 1.13 (td, $J = 13.1, 3.9$ Hz, 1H), 1.04 (dd, $J = 12.5, 2.7$ Hz, 1H), 0.96 (s, 3H), 0.74 (s, 3H), 0.66 (s, 3H). ^{13}C NMR (101 MHz, CDCl_3) δ 167.68, 157.26, 151.20, 147.78, 116.70, 106.86, 78.74, 55.93, 54.68, 45.91, 44.78, 43.85, 43.69, 42.73, 40.82, 39.44, 39.17, 38.59, 38.20, 37.13, 28.38, 27.86, 24.02, 21.60, 18.82, 15.52, 14.57. HRMS calc for $\text{C}_{27}\text{H}_{44}\text{ClN}_3\text{O}_3\text{Na}$ $[\text{M} + \text{Na}]^+$: 516.2969, found 516.2972. HPLC purity: 95.89%, retention time: 11.217 min.

4-((*E*)-5-((1*R*,4*aS*,6*R*,8*aS*)-6-Hydroxy-5,5,8*a*-trimethyl-2-methylenedecahydronaphthalen-1-yl)-3-methylpent-2-enoyl)-*N*-(2,2,2-trifluoroethyl) piperazine-1-carboxamide (**14g**)

Colorless oil, yield 87%, ^1H NMR (400 MHz, CDCl_3) δ 5.71 (s, 1H), 5.23 (t, $J = 6.5$ Hz, 1H), 4.85 (s, 1H), 4.50 (s, 1H), 3.89 (qd, $J = 9.1, 6.3$ Hz, 2H), 3.71 – 3.60 (m, 2H), 3.55 – 3.34 (m, 6H), 3.22 (dd, $J = 11.7, 4.4$ Hz, 1H), 2.40 (dd, $J = 12.8, 2.4$ Hz, 1H), 2.29 – 2.19 (m, 1H), 1.99 – 1.90 (m, 2H), 1.87 (s, 3H), 1.78 – 1.52 (m, 7H), 1.38 (qd, $J = 12.9, 4.0$ Hz, 1H), 1.14 (td, $J = 13.2, 3.9$ Hz, 1H), 1.05 (dd, $J = 12.5, 2.7$ Hz, 1H), 0.98 (s, 3H), 0.76 (s, 3H), 0.68 (s, 3H). ^{13}C NMR (101 MHz, CDCl_3) δ 167.67, 156.60, 151.49, 147.85, 124.56 (q, $J = 278.4$ Hz), 116.67, 106.90, 78.84, 56.05, 54.79, 45.87, 44.04, 43.93, 43.85, 42.09 (q, $J = 34.3$ Hz), 40.77, 39.52, 39.24, 38.72, 38.27, 37.21, 28.41, 27.97, 24.10, 21.71, 18.83, 15.51, 14.60. HRMS calc for $\text{C}_{27}\text{H}_{43}\text{F}_3\text{N}_3\text{O}_3$ $[\text{M}+\text{H}]^+$: 514.3257, found 514.3255. HPLC purity: 100.00%, retention time: 11.797 min. ^{19}F NMR (376 MHz, CDCl_3) δ -72.99.

4-((*E*)-5-((1*R*,4*aS*,6*R*,8*aS*)-6-Hydroxy-5,5,8*a*-trimethyl-2-methylenedecahydronaphthalen-1-yl)-3-methylpent-2-enoyl)-*N*-phenylpiperazine-1-carboxamide (**15a**)

Colorless oil, yield 92%, ^1H NMR (400 MHz, CDCl_3) δ 7.24 – 7.14 (m, 4H), 6.96 (t, $J = 7.2$ Hz, 1H), 6.38 (s, 1H), 5.65 (s, 1H), 4.78 (s, 1H), 4.43 (s, 1H), 3.68 – 3.59 (m, 2H), 3.48 – 3.35 (m, 6H), 3.15 (dd, $J = 11.7, 4.4$ Hz, 1H), 2.32 (ddd, $J = 12.7, 4.2, 2.3$ Hz, 1H), 2.20 – 2.14 (m, 1H), 1.89 – 1.83 (m, 2H), 1.81 (s, 3H), 1.72 – 1.50 (m, 7H), 1.30 (qd, $J = 13.0, 4.2$ Hz, 3H), 1.08 (td, $J = 13.2, 3.7$ Hz, 1H), 0.98 (dd, $J = 12.5, 2.7$ Hz, 1H), 0.90 (s, 3H), 0.68 (s, 3H), 0.60 (s, 3H). ^{13}C NMR (101 MHz, CDCl_3) δ 167.70, 155.12, 151.53, 147.92, 138.81, 129.13, 123.65, 120.31, 116.79, 106.98, 78.92, 56.10, 54.84, 46.04, 44.50, 43.98, 40.96, 39.59, 39.30, 38.80, 38.34, 37.27, 28.48, 28.05, 24.16, 21.77, 18.93, 15.57, 14.68. HRMS calc for $\text{C}_{31}\text{H}_{45}\text{N}_3\text{O}_3\text{Na}$ $[\text{M}+\text{Na}]^+$: 530.3359, found 530.3358. HPLC purity: 99.69%, retention time: 12.207 min.

4-((*E*)-5-((1*R*,4*aS*,6*R*,8*aS*)-6-Hydroxy-5,5,8*a*-trimethyl-2-methylenedecahydronaphthalen-1-yl)-3-methylpent-2-enoyl)-*N*-(*p*-tolyl) piperazine-1-carboxamide (**15b**)

Colorless oil, yield 96%, ^1H NMR (400 MHz, CDCl_3) δ 7.20 (d, $J = 8.3$ Hz, 2H), 7.05 (d, $J = 8.1$ Hz, 2H), 6.91 (s, 1H), 5.71 (s, 1H), 4.85 (s, 1H), 4.50 (s, 1H), 3.70 – 3.58 (m, 2H), 3.52 – 3.38 (m, 6H), 3.21 (dd, $J = 11.6, 4.4$ Hz, 1H), 2.39 (d, $J = 2.4$ Hz, 1H), 2.27 (s, 3H), 2.25 – 2.19 (m, 1H), 1.97 – 1.88 (m, 2H), 1.86 (s, 3H), 1.80 – 1.45 (m, 7H), 1.35 (qd, $J = 12.9, 4.2$ Hz, 1H), 1.13 (td, $J = 13.1, 3.9$ Hz, 1H), 1.04 (dd, $J = 12.5, 2.7$ Hz, 1H), 0.96 (s, 3H), 0.75 (s, 3H), 0.67 (s, 3H). ^{13}C NMR (101 MHz, CDCl_3) δ 167.67, 155.52, 151.16, 147.78, 136.28, 133.00, 129.40, 120.72, 116.72, 106.86, 78.71, 55.97, 54.71, 45.99, 44.32, 43.84, 40.94, 39.45, 39.17, 38.61, 38.22, 37.16, 28.38, 27.88, 24.04, 21.62, 20.84, 18.83, 15.52, 14.56. HRMS calc for $\text{C}_{32}\text{H}_{47}\text{N}_3\text{O}_3\text{Na}$ $[\text{M}+\text{Na}]^+$: 544.3515, found 544.3514. HPLC purity: 98.96%, retention time: 14.637 min.

4-((*E*)-5-((1*R*,4*aS*,6*R*,8*aS*)-6-Hydroxy-5,5,8*a*-trimethyl-2-methylenedecahydronaphthalen-1-yl)-3-methylpent-2-enoyl)-*N*-(*o*-tolyl) piperazine-1-carboxamide (**15c**)

Colorless oil, yield 94%, ¹H NMR (400 MHz, CDCl₃) δ 7.40 (d, 1H), 7.18 – 7.10 (m, 2H), 7.02 (td, *J* = 7.4, 1.3 Hz, 1H), 6.51 (s, 1H), 5.70 (s, 1H), 4.85 (s, 1H), 4.50 (s, 1H), 3.69 – 3.59 (m, 2H), 3.50 – 3.36 (m, 6H), 3.21 (dd, *J* = 11.7, 4.4 Hz, 1H), 2.44 – 2.35 (m, 1H), 2.30 – 2.21 (m, 1H), 2.20 (s, 3H), 1.98 – 1.89 (m, 2H), 1.87 (s, 3H), 1.78 – 1.46 (m, 7H), 1.37 (qd, *J* = 12.9, 4.2 Hz, 1H), 1.10 (td, *J* = 3.9 Hz, 1H), 1.04 (dd, *J* = 12.5, 2.7 Hz, 1H), 0.96 (s, 3H), 0.74 (s, 3H), 0.67 (s, 3H). ¹³C NMR (101 MHz, CDCl₃) δ 167.62, 155.75, 151.27, 147.76, 136.66, 130.51, 130.44, 126.63, 124.87, 123.93, 116.62, 106.80, 78.67, 55.91, 54.66, 45.87, 44.32, 43.93, 40.81, 39.41, 39.12, 38.59, 38.17, 37.11, 28.33, 27.80, 23.99, 21.58, 18.78, 17.89, 15.47, 14.53. HRMS calc for C₃₂H₄₈N₃O₃ [M+H]⁺: 522.3696, found 522.3700. HPLC purity: 98.34%, retention time: 13.540 min.

N-Benzyl-4-((*E*)-5-((1*R*,4*aS*,6*R*,8*aS*)-6-hydroxy-5,5,8*a*-trimethyl-2-methylenedecahydronaphthalen-1-yl)-3-methylpent-2-enoyl) piperazine-1-carboxamide (**15d**)

Colorless oil, yield 91%, ¹H NMR (400 MHz, CDCl₃) δ 7.32 – 7.24 (m, 5H), 5.69 (s, 1H), 4.87 (t, *J* = 5.1 Hz, 1H), 4.84 (s, 1H), 4.49 (s, 1H), 4.40 (d, *J* = 5.3 Hz, 2H), 3.67 – 3.57 (m, 2H), 3.51 – 3.37 (m, 4H), 3.37 – 3.25 (m, 2H), 3.21 (dd, *J* = 11.7, 4.4 Hz, 1H), 2.38 (ddd, *J* = 12.8, 4.3, 2.4 Hz, 1H), 2.26 – 2.17 (m, 1H), 1.96 – 1.87 (m, 2H), 1.86 (s, 3H), 1.77 – 1.50 (m, 7H), 1.36 (qd, *J* = 13.0, 4.3 Hz, 1H), 1.12 (td, *J* = 13.0, 3.9 Hz, 1H), 1.03 (dd, *J* = 12.5, 2.6 Hz, 1H), 0.96 (s, 3H), 0.74 (s, 3H), 0.66 (s, 3H). ¹³C NMR (101 MHz, CDCl₃) δ 167.63, 157.48, 151.15, 147.84, 139.20, 128.79, 127.91, 127.56, 116.82, 106.91, 78.82, 56.00, 54.75, 45.99, 45.17, 44.02, 43.78, 40.84, 39.51, 39.23, 38.67, 38.27, 28.43, 27.96, 24.09, 21.66, 18.84, 15.53, 14.61. HRMS calc for C₃₂H₄₈N₃O₃ [M+H]⁺: 522.3696, found 522.3698. HPLC purity: 100.00%, retention time: 13.213 min.

N-(3,5-Dimethylphenyl)-4-((*E*)-5-((1*R*,4*aS*,6*R*,8*aS*)-6-hydroxy-5,5,8*a*-trimethyl-2-methylenedecahydronaphthalen-1-yl)-3-methylpent-2-enoyl) piperazine-1-carboxamide (**15e**)

White powder solid, melt point: 119 °C, yield 89%, ¹H NMR (400 MHz, CDCl₃) δ 6.96 (s, 2H), 6.75 (s, 1H), 6.67 (s, 1H), 5.72 (s, 1H), 4.85 (s, 1H), 4.50 (s, 1H), 3.69 – 3.59 (m, 2H), 3.53 – 3.39 (m, 6H), 3.21 (dd, *J* = 11.6, 4.4 Hz, 1H), 2.39 (ddd, *J* = 12.6, 4.2, 2.4 Hz, 1H), 2.27 – 2.18 (m, 1H), 2.24 (s, 6H), 1.97 – 1.89 (m, 2H), 1.87 (s, 3H), 1.78 – 1.47 (m, 7H), 1.37 (qd, *J* = 13.0, 4.2 Hz, 1H), 1.13 (td, *J* = 13.1, 3.8 Hz, 1H), 1.04 (dd, *J* = 12.5, 2.7 Hz, 1H), 0.97 (s, 3H), 0.75 (s, 3H), 0.67 (s, 3H). ¹³C NMR (101 MHz, CDCl₃) δ 167.64, 155.33, 151.22, 147.79, 138.67, 138.54, 125.20, 118.15, 116.72, 106.87, 78.73, 55.97, 54.72, 46.00, 44.40, 43.87, 40.93, 39.46, 39.18, 38.64, 38.23, 37.16, 28.40, 27.91, 24.05, 21.63, 21.45, 18.83, 15.51, 14.56. HRMS calc for C₃₃H₅₀N₃O₃ [M+H]⁺: 536.3852, found 536.3853. HPLC purity: 100.00%, retention time: 15.820 min.

N-(4-Ethylphenyl)-4-((*E*)-5-((1*R*,4*aS*,6*R*,8*aS*)-6-hydroxy-5,5,8*a*-trimethyl-2-

methylenedecahydronaphthalen-1-yl)-3-methylpent-2-enoyl) piperazine-1-carboxamide (**15f**)

White solid, melt point: 184 °C, yield 90%, ¹H NMR (400 MHz, CDCl₃) δ 7.23 (d, *J* = 8.4 Hz, 2H), 7.08 (d, *J* = 8.5 Hz, 2H), 6.85 (s, 1H), 5.71 (s, 1H), 4.85 (s, 1H), 4.50 (s, 1H), 3.69 – 3.59 (m, 2H), 3.53 – 3.39 (m, 6H), 3.21 (dd, *J* = 11.6, 4.4 Hz, 1H), 2.57 (q, *J* = 7.6 Hz, 2H), 2.39 (dd, *J* = 12.7, 2.4 Hz, 1H), 2.28 – 2.20 (m, 1H), 1.97 – 1.89 (m, 2H), 1.87 (s, 3H), 1.78 – 1.46 (m, 7H), 1.37 (qd, *J* = 13.1, 4.3 Hz, 1H), 1.18 (t, *J* = 7.6 Hz, 3H), 1.12 (td, *J* = 13.1, 3.8 Hz, 1H), 1.04 (dd, *J* = 12.4, 2.7 Hz, 1H), 0.97 (s, 3H), 0.75 (s, 3H), 0.67 (s, 3H). ¹³C NMR (101 MHz, CDCl₃) δ 167.64, 155.48, 151.18, 147.80, 139.51, 136.44, 128.25, 120.74, 116.74, 106.87, 78.72, 55.98, 54.72, 46.00, 44.35, 43.86, 40.93, 39.46, 39.19, 38.63, 38.23, 37.16, 28.40, 28.38, 28.28, 27.90, 24.05, 21.63, 18.84, 15.80, 15.53, 14.59. HRMS calc for C₃₃H₄₉N₃O₃Na [M+ Na]⁺: 558.3672, found 558.3669. HPLC purity: 100.00%, retention time: 15.873 min.

4-((*E*)-5-((1*R*,4*aS*,6*R*,8*aS*)-6-Hydroxy-5,5,8*a*-trimethyl-2-methylenedecahydronaphthalen-1-yl)-3-methylpent-2-enoyl)-*N*-phenethylpiperazine-1-carboxamide (**15g**)

Colorless oil, yield 89%, ¹H NMR (400 MHz, CDCl₃) δ 7.29 (t, *J* = 6.6 Hz, 2H), 7.25 – 7.19 (m, 1H), 7.17 (d, *J* = 7.1 Hz, 2H), 5.70 (s, 1H), 4.85 (s, 1H), 4.50 (s, 1H), 3.68 – 3.54 (m, 2H), 3.52 – 3.42 (m, 4H), 3.41 – 3.32 (m, 2H), 3.30 – 3.18 (m, 3H), 2.81 (t, *J* = 6.9 Hz, 2H), 2.39 (ddd, *J* = 12.7, 4.3, 2.4 Hz, 1H), 2.28 – 2.19 (m, 1H), 1.99 – 1.88 (m, 2H), 1.87 (s, 3H), 1.78 – 1.45 (m, 7H), 1.37 (qd, *J* = 12.9, 4.2 Hz, 1H), 1.14 (td, *J* = 13.1, 3.8 Hz, 1H), 1.05 (dd, *J* = 12.5, 2.7 Hz, 1H), 0.97 (s, 3H), 0.75 (s, 3H), 0.67 (s, 3H). ¹³C NMR (101 MHz, CDCl₃) δ 167.63, 157.50, 151.12, 147.81, 139.26, 128.89, 128.70, 126.55, 116.77, 106.87, 78.75, 55.97, 54.72, 45.96, 43.86, 43.62, 42.16, 40.75, 39.47, 39.20, 38.64, 38.23, 37.16, 36.30, 28.40, 27.90, 24.06, 21.63, 18.81, 15.53, 14.59. HRMS calc for C₃₃H₅₀N₃O₃ [M+H]⁺: 536.3852, found 536.3854. HPLC purity: 100.00%, retention time: 13.837 min.

4-((*E*)-5-((1*R*,4*aS*,6*R*,8*aS*)-6-Hydroxy-5,5,8*a*-trimethyl-2-methylenedecahydronaphthalen-1-yl)-3-methylpent-2-enoyl)-*N*-(2-isopropylphenyl)piperazine-1-carboxamide (**15h**)

Colorless oil, yield 87%, ¹H NMR (400 MHz, CDCl₃) δ 7.42 – 7.36 (m, 1H), 7.27 – 7.24 (m, 1H), 7.17 – 7.11 (m, 2H), 6.38 (s, 1H), 5.72 (s, 1H), 4.86 (s, 1H), 4.51 (s, 1H), 3.71 – 3.61 (m, 2H), 3.54 – 3.40 (m, 6H), 3.22 (dd, *J* = 11.7, 4.4 Hz, 1H), 3.02 (p, *J* = 6.9 Hz, 1H), 2.40 (ddd, *J* = 12.7, 4.2, 2.3 Hz, 1H), 2.29 – 2.20 (m, 1H), 1.97 – 1.90 (m, 2H), 1.89 (s, 3H), 1.79 – 1.53 (m, 7H), 1.38 (qd, *J* = 12.9, 4.2 Hz, 1H), 1.22 (d, *J* = 6.8 Hz, 6H), 1.15 (td, 1H), 1.05 (dd, *J* = 12.5, 2.7 Hz, 1H), 0.97 (s, 3H), 0.75 (s, 3H), 0.68 (s, 3H). ¹³C NMR (101 MHz, CDCl₃) δ 167.59, 156.14, 151.32, 147.86, 141.38, 135.24, 126.40, 125.73, 125.45, 116.75, 106.86, 78.74, 56.02, 54.76, 45.94, 44.41, 44.03, 40.87, 39.50, 39.21, 38.72, 38.26, 37.20, 28.40, 28.12, 27.94, 24.08, 23.15, 21.69, 18.82, 15.51, 14.61. HRMS calc for C₃₄H₅₁N₃O₃Na [M +Na]⁺: 572.3828, found 572.3825.

HPLC purity: 100.00%, retention time: 15.267 min.

N-(4-Chlorophenyl)-4-((*E*)-5-((1*R*,4*aS*,6*R*,8*aS*)-6-hydroxy-5,5,8*a*-trimethyl-2-methylenedecahydronaphthalen-1-yl)-3-methylpent-2-enoyl) piperazine-1-carbothioamide (**15i**)

Colorless oil, yield 89%, ¹H NMR (400 MHz, CDCl₃) δ 7.89 (s, 1H), 7.25 (d, *J* = 9.4 Hz, 2H), 7.12 (d, *J* = 8.6 Hz, 2H), 5.69 (s, 1H), 4.84 (s, 1H), 4.47 (s, 1H), 3.99 – 3.91 (m, 2H), 3.79 – 3.65 (m, 4H), 3.61 – 3.50 (m, 2H), 3.21 (dd, *J* = 11.6, 4.3 Hz, 1H), 2.38 (d, *J* = 13.0 Hz, 1H), 2.22 (t, *J* = 9.1 Hz, 1H), 1.95 – 1.88 (m, 2H), 1.86 (s, 3H), 1.75 – 1.49 (m, 7H), 1.36 (qd, *J* = 12.7, 3.8 Hz, 1H), 1.12 (td, *J* = 13.5, 3.9 Hz, 1H), 1.03 (dd, *J* = 12.5, 1.8 Hz, 1H), 0.95 (s, 3H), 0.73 (s, 3H), 0.66 (s, 3H). ¹³C NMR (101 MHz, CDCl₃) δ 167.71, 152.34, 147.71, 138.47, 130.90, 129.01, 125.74, 116.21, 106.85, 78.73, 55.92, 54.63, 48.81, 48.18, 45.30, 40.38, 39.42, 39.15, 38.72, 38.17, 37.12, 28.36, 27.84, 23.99, 21.63, 18.87, 15.51, 14.56. HRMS calc for C₃₁H₄₄ClN₃O₃K [M+K]⁺: 580.2708, found 580.2705. HPLC purity: 100.00%, retention time: 17.027 min.

N-(2-Chlorophenyl)-4-((*E*)-5-((1*R*,4*aS*,6*R*,8*aS*)-6-hydroxy-5,5,8*a*-trimethyl-2-methylenedecahydronaphthalen-1-yl)-3-methylpent-2-enoyl) piperazine-1-carboxamide (**15j**)

Colorless oil, yield 90%, ¹H NMR (400 MHz, CDCl₃) δ 8.12 (dd, *J* = 8.3, 1.5 Hz, 1H), 7.32 (dd, *J* = 8.0, 1.5 Hz, 1H), 7.23 (td, *J* = 8.0, 1.5 Hz, 1H), 7.00 (s, 1H), 6.96 (td, *J* = 7.7, 1.6 Hz, 1H), 5.74 (s, 1H), 4.86 (s, 1H), 4.51 (s, 1H), 3.81 – 3.67 (m, 2H), 3.63 – 3.48 (m, 6H), 3.22 (dd, *J* = 11.7, 4.4 Hz, 1H), 2.40 (ddd, *J* = 12.8, 4.2, 2.4 Hz, 1H), 2.30 – 2.19 (m, 1H), 1.98 – 1.90 (m, 2H), 1.90 (s, 3H), 1.81 – 1.45 (m, 7H), 1.37 (qd, *J* = 12.9, 4.2 Hz, 1H), 1.14 (td, *J* = 13.1, 3.9 Hz, 1H), 1.05 (dd, *J* = 12.5, 2.7 Hz, 1H), 0.97 (s, 3H), 0.75 (s, 3H), 0.68 (s, 3H). ¹³C NMR (101 MHz, CDCl₃) δ 167.59, 154.13, 151.62, 147.82, 135.46, 128.90, 127.81, 123.62, 122.66, 121.23, 116.60, 106.87, 78.74, 55.97, 54.72, 45.81, 44.15, 43.87, 40.79, 39.48, 39.20, 38.71, 38.24, 37.16, 28.41, 27.92, 24.06, 21.65, 18.86, 15.51, 14.58. HRMS calc for C₃₁H₄₄ClN₃O₃Na [M + Na]⁺: 564.2969, found 564.2968. HPLC purity: 96.77%, retention time: 14.910 min.

N-(3-Chlorophenyl)-4-((*E*)-5-((1*R*,4*aS*,6*R*,8*aS*)-6-hydroxy-5,5,8*a*-trimethyl-2-methylenedecahydronaphthalen-1-yl)-3-methylpent-2-enoyl) piperazine-1-carboxamide (**15k**)

Colorless oil, yield 93%, ¹H NMR (400 MHz, CDCl₃) δ 7.43 – 7.38 (m, 1H), 7.33 – 7.26 (m, 1H), 7.26 – 7.22 (m, 1H), 7.15 (t, *J* = 8.0 Hz, 1H), 7.01 – 6.93 (m, 1H), 5.72 (s, 1H), 4.85 (s, 1H), 4.49 (s, 1H), 3.72 – 3.60 (m, 2H), 3.56 – 3.41 (m, 6H), 3.21 (dd, *J* = 11.7, 4.3 Hz, 1H), 2.39 (ddd, *J* = 12.7, 4.2, 2.4 Hz, 1H), 2.30 – 2.18 (m, 1H), 1.96 – 1.89 (m, 2H), 1.86 (s, 3H), 1.77 – 1.45 (m, 7H), 1.37 (qd, *J* = 13.1, 4.3 Hz, 1H), 1.12 (td, *J* = 13.1, 3.9 Hz, 1H), 1.04 (dd, *J* = 12.5, 2.7 Hz, 1H), 0.97 (s, 3H), 0.75 (s, 3H), 0.67 (s, 3H). ¹³C NMR (101 MHz, CDCl₃) δ 167.77, 154.99, 151.44, 147.79, 140.43, 134.36, 129.87, 123.21, 120.28, 118.31, 116.64, 106.90, 78.77, 56.00, 54.74, 46.02,

44.41, 43.84, 40.99, 39.48, 39.20, 38.63, 38.24, 37.18, 28.42, 27.92, 24.05, 21.65, 18.90, 15.54, 14.59. HRMS calc for C₃₁H₄₄ClN₃O₃Na [M +Na]⁺: 564.2969, found 564.2966. HPLC purity: 98.76%, retention time: 16.310 min.

N-(2-Bromophenyl)-4-((*E*)-5-((1*R*,4*aS*,6*R*,8*aS*)-6-hydroxy-5,5,8*a*-trimethyl-2-methylenedecahydronaphthalen-1-yl)-3-methylpent-2-enoyl) piperazine-1-carboxamide (**15l**)

White solid, melt point: 199 °C, yield 90%, ¹H NMR (400 MHz, CDCl₃) δ 8.11 (dd, *J* = 8.4, 1.6 Hz, 1H), 7.48 (dd, *J* = 8.0, 1.5 Hz, 1H), 7.27 (td, *J* = 15.7, 1.3 Hz, 1H), 7.02 (s, 1H), 6.89 (td, *J* = 8.0, 1.5 Hz, 1H), 5.73 (s, 1H), 4.85 (s, 1H), 4.50 (s, 1H), 3.78 – 3.67 (m, 2H), 3.61 – 3.47 (m, 6H), 3.21 (dd, *J* = 11.6, 4.4 Hz, 1H), 2.39 (ddd, *J* = 12.8, 4.2, 2.4 Hz, 1H), 2.25 (td, *J* = 11.0, 6.2 Hz, 1H), 2.01 – 1.90 (m, 2H), 1.89 (s, 3H), 1.79 – 1.51 (m, 7H), 1.37 (qd, *J* = 12.9, 4.2 Hz, 1H), 1.14 (td, *J* = 13.1, 3.9 Hz, 1H), 1.05 (dd, *J* = 12.5, 2.7 Hz, 1H), 0.97 (s, 3H), 0.75 (s, 3H), 0.67 (s, 3H). ¹³C NMR (101 MHz, CDCl₃) δ 167.56, 154.12, 151.60, 147.79, 136.51, 132.08, 128.45, 124.15, 121.44, 116.58, 113.55, 106.86, 78.69, 55.93, 54.68, 45.79, 44.12, 43.83, 40.76, 39.45, 39.18, 38.68, 38.21, 37.13, 28.39, 27.90, 24.03, 21.61, 18.85, 15.51, 14.57. HRMS calc for C₃₁H₄₄BrN₃O₃Na [M+Na]⁺: 610.2443, found 610.2454. HPLC purity: 99.20%, retention time: 15.273 min.

4-((*E*)-5-((1*R*,4*aS*,6*R*,8*aS*)-6-Hydroxy-5,5,8*a*-trimethyl-2-methylenedecahydronaphthalen-1-yl)-3-methylpent-2-enoyl)-*N*-(4-iodophenyl) piperazine-1-carboxamide (**15m**)

Colorless oil, yield 94%, ¹H NMR (400 MHz, CDCl₃) δ 7.53 (d, *J* = 8.8 Hz, 1H), 7.14 (d, *J* = 8.7 Hz, 2H), 7.10 (s, 1H), 5.72 (s, 1H), 4.85 (s, 1H), 4.49 (s, 1H), 3.70 – 3.60 (m, 2H), 3.55 – 3.40 (m, 6H), 3.21 (dd, *J* = 11.7, 4.4 Hz, 1H), 2.39 (ddd, *J* = 12.8, 4.2, 2.3 Hz, 1H), 2.30 – 2.18 (m, 1H), 1.98 – 1.87 (m, 3H), 1.87 (s, 3H), 1.79 – 1.47 (m, 7H), 1.37 (qd, *J* = 13.0, 4.2 Hz, 1H), 1.13 (td, *J* = 13.1, 3.8 Hz, 1H), 1.04 (dd, *J* = 12.5, 2.7 Hz, 1H), 0.97 (s, 3H), 0.75 (s, 3H), 0.67 (s, 3H). ¹³C NMR (101 MHz, CDCl₃) δ 167.70, 154.75, 151.69, 147.87, 138.79, 137.90, 122.08, 116.65, 106.94, 86.39, 78.87, 56.07, 54.80, 45.99, 44.46, 43.84, 40.92, 39.55, 39.26, 38.77, 38.30, 37.24, 28.45, 28.01, 24.12, 21.73, 18.92, 15.54, 14.65. HRMS calc for C₃₁H₄₅IN₃O₃ [M+H]⁺: 634.2506, found 634.2504. HPLC purity: 100.00%, retention time: 17.343 min.

N-(4-Cyanophenyl)-4-((*E*)-5-((1*R*,4*aS*,6*R*,8*aS*)-6-hydroxy-5,5,8*a*-trimethyl-2-methylenedecahydronaphthalen-1-yl)-3-methylpent-2-enoyl) piperazine-1-carboxamide (**15n**)

Colorless oil, yield 95%, ¹H NMR (400 MHz, CDCl₃) δ 7.57 (s, 1H), 7.55 – 7.48 (m, 4H), 5.72 (s, 1H), 4.85 (s, 1H), 4.48 (s, 1H), 3.72 – 3.62 (m, 2H), 3.59 – 3.48 (m, 6H), 3.21 (dd, *J* = 11.6, 4.4 Hz, 1H), 2.39 (dt, *J* = 12.4, 3.2 Hz, 1H), 2.29 – 2.19 (m, 1H), 1.96 – 1.89 (m, 2H), 1.86 (s, 3H), 1.77 – 1.49 (m, 7H), 1.37 (qd, *J* = 13.0, 4.2 Hz, 1H), 1.12 (td, *J* = 13.1, 3.8 Hz, 1H), 1.04 (dd, *J* = 12.4, 2.7 Hz, 1H), 0.96 (s, 3H), 0.75 (s,

3H), 0.66 (s, 3H). ¹³C NMR (101 MHz, CDCl₃) δ 167.76, 154.38, 151.73, 147.78, 143.88, 133.09, 119.51, 119.34, 116.51, 106.86, 105.33, 78.75, 56.03, 54.75, 46.00, 44.51, 43.83, 40.98, 39.48, 39.20, 38.68, 38.22, 37.20, 28.41, 27.91, 24.05, 21.70, 18.88, 15.51, 14.58. HRMS calc for C₃₂H₄₅N₄O₃ [M+H]⁺: 533.3492, found 533.3489. HPLC purity: 100.00%, retention time: 13.930 min.

4-((*E*)-5-((1*R*,4*aS*,6*R*,8*aS*)-6-Hydroxy-5,5,8*a*-trimethyl-2-methylenedecahydronaphthalen-1-yl)-3-methylpent-2-enoyl)-*N*-(2-nitrophenyl)piperazine-1-carboxamide (**15o**)

Yellow oil, yield 92%, ¹H NMR (400 MHz, CDCl₃) δ 10.22 (s, 1H), 8.59 (dd, *J* = 8.7, 1.3 Hz, 1H), 8.18 (dd, *J* = 8.5, 1.6 Hz, 1H), 7.60 (ddd, *J* = 8.8, 7.2, 1.6 Hz, 1H), 7.07 (ddd, *J* = 8.5, 7.2, 1.3 Hz, 1H), 5.74 (s, 1H), 4.86 (s, 1H), 4.51 (s, 1H), 3.79 – 3.67 (m, 2H), 3.65 – 3.55 (m, 6H), 3.23 (dd, *J* = 11.7, 4.4 Hz, 1H), 2.40 (ddd, *J* = 12.7, 4.2, 2.3 Hz, 1H), 2.32 – 2.20 (m, 1H), 1.99 – 1.91 (m, 2H), 1.90 (s, 3H), 1.80 – 1.54 (m, 7H), 1.38 (qd, *J* = 13.0, 4.3 Hz, 1H), 1.15 (td, *J* = 13.2, 3.9 Hz, 1H), 1.06 (dd, *J* = 12.5, 2.8 Hz, 1H), 0.98 (s, 3H), 0.76 (s, 3H), 0.68 (s, 3H). ¹³C NMR (101 MHz, CDCl₃) δ 167.59, 153.62, 151.73, 147.82, 137.01, 136.20, 135.80, 125.89, 121.90, 121.54, 116.59, 106.91, 78.77, 56.01, 54.77, 45.76, 43.93, 41.82, 40.97, 39.51, 39.22, 38.73, 38.26, 37.20, 28.42, 27.95, 24.08, 21.67, 18.89, 15.51, 14.60. HRMS calc for C₃₁H₄₅N₄O₅ [M+H]⁺: 553.3390, found 553.3392. HPLC purity: 100.00%, retention time: 15.457 min.

4-((*E*)-5-((1*R*,4*aS*,6*R*,8*aS*)-6-Hydroxy-5,5,8*a*-trimethyl-2-methylenedecahydronaphthalen-1-yl)-3-methylpent-2-enoyl)-*N*-(3-methoxyphenyl)piperazine-1-carboxamide (**15p**)

Colorless oil, yield 90%, ¹H NMR (400 MHz, CDCl₃) δ 7.13 (t, *J* = 8.1 Hz, 1H), 7.08 – 7.01 (m, 2H), 6.84 (ddd, *J* = 8.1, 2.0, 0.9 Hz, 1H), 6.56 (ddd, *J* = 8.3, 2.5, 0.8 Hz, 1H), 5.70 (s, 1H), 4.85 (s, 1H), 4.49 (s, 1H), 3.74 (s, 3H), 3.68 – 3.58 (m, 2H), 3.53 – 3.38 (m, 6H), 3.21 (dd, *J* = 11.6, 4.4 Hz, 1H), 2.39 (ddd, *J* = 12.8, 4.2, 2.4 Hz, 1H), 2.24 (td, *J* = 10.7, 5.7 Hz, 1H), 2.00 – 1.88 (m, 2H), 1.87 (s, 3H), 1.80 – 1.47 (m, 7H), 1.37 (qd, *J* = 12.9, 4.2 Hz, 1H), 1.12 (td, *J* = 13.1, 3.8 Hz, 1H), 1.04 (dd, *J* = 12.5, 2.7 Hz, 1H), 0.96 (s, 3H), 0.74 (s, 3H), 0.66 (s, 3H). ¹³C NMR (101 MHz, CDCl₃) δ 167.71, 160.14, 155.26, 151.29, 147.78, 140.27, 129.54, 116.68, 112.48, 109.08, 106.86, 106.09, 78.73, 55.96, 55.30, 54.71, 45.99, 44.37, 43.86, 40.96, 39.45, 39.17, 38.61, 38.21, 37.15, 28.37, 27.87, 24.03, 21.62, 18.84, 15.52, 14.57. HRMS calc for C₃₂H₄₇N₃O₄K [M+K]⁺: 576.3204, found 576.3201. HPLC purity: 99.22%, retention time: 13.760 min

4-((*E*)-5-((1*R*,4*aS*,6*R*,8*aS*)-6-Hydroxy-5,5,8*a*-trimethyl-2-methylenedecahydronaphthalen-1-yl)-3-methylpent-2-enoyl)-*N*-(4-methoxybenzyl)piperazine-1-carboxamide (**15q**)

Colorless oil, yield 85%, ¹H NMR (400 MHz, CDCl₃) δ 7.19 (d, *J* = 8.5 Hz, 2H), 6.82 (d, *J* = 8.6 Hz, 2H), 5.69 (s, 1H), 5.05 (t, *J* = 5.4 Hz, 1H), 4.84 (s, 1H), 4.49 (s, 1H),

4.31 (d, $J = 5.2$ Hz, 2H), 3.76 (s, 3H), 3.65 – 3.56 (m, 2H), 3.49 – 3.37 (m, 4H), 3.35 – 3.27 (m, 2H), 3.20 (dd, $J = 11.7, 4.4$ Hz, 1H), 2.38 (ddd, $J = 12.7, 4.3, 2.4$ Hz, 1H), 2.22 (ddd, $J = 13.7, 10.2, 3.6$ Hz, 1H), 1.96 – 1.86 (m, 2H), 1.85 (s, 3H), 1.77 – 1.45 (m, 7H), 1.36 (qd, $J = 13.0, 4.3$ Hz, 1H), 1.12 (td, $J = 13.0, 3.8$ Hz, 1H), 1.03 (dd, $J = 12.4, 2.7$ Hz, 1H), 0.95 (s, 3H), 0.74 (s, 3H), 0.66 (s, 3H). ^{13}C NMR (101 MHz, CDCl_3) δ 167.65, 158.94, 157.49, 151.06, 147.76, 131.24, 129.18, 116.71, 114.02, 106.84, 78.69, 55.90, 55.35, 54.66, 45.95, 44.52, 43.88, 43.66, 40.81, 39.42, 39.15, 38.56, 38.18, 37.11, 28.36, 27.82, 24.00, 21.57, 18.78, 15.49, 14.55. HRMS calc for $\text{C}_{33}\text{H}_{50}\text{N}_3\text{O}_4$ $[\text{M}+\text{H}]^+$: 552.3801, found 552.3802. HPLC purity: 98.39%, retention time: 12.950 min.

N-(4-Acetylphenyl)-4-((*E*)-5-((1*R*,4*aS*,6*R*,8*aS*)-6-hydroxy-5,5,8*a*-trimethyl-2-methylenedecahydronaphthalen-1-yl)-3-methylpent-2-enoyl) piperazine-1-carboxamide (**15r**)

White powder solid, melt point: 189 °C, yield 86%, ^1H NMR (400 MHz, CDCl_3) δ 7.83 (d, $J = 8.8$ Hz, 2H), 7.77 (s, 1H), 7.47 (d, $J = 8.8$ Hz, 2H), 5.71 (s, 1H), 4.83 (s, 1H), 4.47 (s, 1H), 3.70 – 3.60 (m, 2H), 3.57 – 3.48 (m, 6H), 3.20 (dd, $J = 11.6, 4.4$ Hz, 1H), 2.52 (s, 3H), 2.37 (dt, $J = 12.8, 2.6$ Hz, 1H), 2.28 – 2.16 (m, 1H), 1.95 – 1.86 (m, 2H), 1.85 (s, 3H), 1.76 – 1.44 (m, 7H), 1.35 (qd, $J = 13.0, 4.2$ Hz, 1H), 1.11 (td, $J = 13.0, 3.6$ Hz, 1H), 1.02 (dd, $J = 12.5, 2.7$ Hz, 1H), 0.95 (s, 3H), 0.73 (s, 3H), 0.65 (s, 3H). ^{13}C NMR (101 MHz, CDCl_3) δ 197.50, 167.79, 154.72, 151.60, 147.73, 144.20, 131.54, 129.69, 118.83, 116.50, 106.85, 78.73, 55.93, 54.67, 46.01, 44.50, 43.88, 41.05, 39.42, 39.15, 38.59, 38.18, 37.13, 28.36, 27.81, 26.48, 24.00, 21.60, 18.87, 15.50, 14.55. HRMS calc for $\text{C}_{33}\text{H}_{47}\text{N}_3\text{O}_4\text{Na}$ $[\text{M}+\text{Na}]^+$: 572.3464, found 572.3463. HPLC purity: 99.40%, retention time: 12.800 min.

4-((*E*)-5-((1*R*,4*aS*,6*R*,8*aS*)-6-Hydroxy-5,5,8*a*-trimethyl-2-methylenedecahydronaphthalen-1-yl)-3-methylpent-2-enoyl)-*N*-(3-(trifluoromethoxy)phenyl) piperazine-1-carboxamide (**15s**)

Colorless oil, yield 97%, ^1H NMR (400 MHz, CDCl_3) δ 7.38 (s, 1H), 7.33 – 7.24 (m, 3H), 6.88 (d, $J = 7.7$ Hz, 1H), 5.74 (s, 1H), 4.88 (s, 1H), 4.52 (s, 1H), 3.73 – 3.63 (m, 2H), 3.59 – 3.46 (m, 6H), 3.24 (dd, $J = 11.7, 4.3$ Hz, 1H), 2.42 (dt, $J = 13.3, 3.1$ Hz, 1H), 2.32 – 2.21 (m, 1H), 2.00 – 1.90 (m, 3H), 1.89 (s, 3H), 1.82 – 1.50 (m, 7H), 1.40 (qd, $J = 13.1, 4.3$ Hz, 1H), 1.16 (td, $J = 13.1, 3.8$ Hz, 1H), 1.07 (dd, $J = 12.5, 2.7$ Hz, 1H), 0.99 (s, 3H), 0.78 (s, 3H), 0.70 (s, 3H). ^{13}C NMR (101 MHz, CDCl_3) δ 167.76, 154.85, 151.56, 149.61, 147.83, 140.70, 129.87, 120.54 (q, $J = 257.2$ Hz), 118.14, 116.61, 115.28, 112.77, 106.88, 78.81, 56.06, 54.78, 46.00, 44.44, 43.83, 40.98, 39.50, 39.22, 38.68, 38.26, 37.20, 28.40, 27.95, 24.08, 21.69, 18.84, 15.50, 14.58. HRMS calc for $\text{C}_{32}\text{H}_{44}\text{F}_3\text{N}_3\text{O}_4\text{Na}$ $[\text{M}+\text{Na}]^+$: 614.3182, found 614.3184. HPLC purity: 100.00%, retention time: 6.043 min. ^{19}F NMR (377 MHz, CDCl_3) δ -57.72.

N-(2-(1*H*-Indol-3-yl) ethyl)-4-((*E*)-5-((1*R*,4*aS*,6*R*,8*aS*)-6-hydroxy-5,5,8*a*-trimethyl-2-methylenedecahydronaphthalen-1-yl)-3-methylpent-2-enoyl) piperazine-1-

carboxamide(**15t**)

Colorless oil, yield 84%, ^1H NMR (400 MHz, CDCl_3) δ 8.37 (s, 1H), 7.60 (d, $J = 7.9$ Hz, 1H), 7.37 (d, $J = 8.1$ Hz, 1H), 7.23 – 7.15 (m, 1H), 7.15 – 7.06 (m, 1H), 7.03 (d, $J = 2.2$ Hz, 1H), 5.70 (s, 1H), 4.86 (s, 1H), 4.58 (t, $J = 5.5$ Hz, 1H), 4.51 (s, 1H), 3.63 – 3.52 (m, 4H), 3.50 – 3.40 (m, 2H), 3.39 – 3.30 (m, 2H), 3.27 – 3.16 (m, 3H), 2.99 (t, $J = 6.5$ Hz, 2H), 2.45 – 2.36 (m, 1H), 2.30 – 2.19 (m, 1H), 1.99 – 1.88 (m, 2H), 1.88 (s, 3H), 1.80 – 1.55 (m, 7H), 1.39 (qd, $J = 13.0, 4.3$ Hz, 1H), 1.13 (td, $J = 13.2, 6.6$ Hz, 1H), 1.05 (dd, $J = 12.6, 2.7$ Hz, 1H), 0.99 (s, 3H), 0.77 (s, 3H), 0.69 (s, 3H). ^{13}C NMR (101 MHz, CDCl_3) δ 167.65, 157.60, 151.01, 147.85, 136.58, 127.62, 122.34, 122.16, 119.55, 118.86, 116.95, 113.43, 111.47, 106.96, 78.89, 56.01, 54.82, 46.02, 43.85, 43.59, 41.59, 40.81, 39.53, 39.26, 38.65, 38.31, 37.24, 28.47, 28.01, 25.80, 24.12, 21.68, 18.86, 15.55, 14.64. HRMS calc for $\text{C}_{35}\text{H}_{51}\text{N}_4\text{O}_3$ $[\text{M}+\text{H}]^+$: 575.3961, found 575.3963. HPLC purity: 97.12%, retention time: 13.997 min.

N,N'-(1,4-Phenylene) bis(4-((*E*)-5-((1*R*,4*aS*,6*R*,8*aS*)-6-hydroxy-5,5,8*a*-trimethyl-2-methylenedecahydronaphthalen-1-yl)-3-methylpent-2-enoyl) piperazine-1-carboxamide) (**15u**)

White power solid, melt point: 190 °C, yield 96%, ^1H NMR (400 MHz, $\text{DMSO}-d_6$) δ 8.45 (s, 2H), 7.30 (s, 4H), 5.84 (s, 2H), 4.86 (s, 2H), 4.54 (s, 2H), 4.40 – 4.34 (m, 2H), 3.55 – 3.40 (m, 16H), 3.05 (d, $J = 10.9$ Hz, 2H), 2.42 – 2.34 (m, 2H), 2.22 – 2.11 (m, 2H), 1.98 – 1.86 (m, 4H), 1.82 (s, 6H), 1.77 – 1.45 (m, 14H), 1.32 (qd, $J = 12.8, 4.0$ Hz, 2H), 1.12 (td, $J = 13.0, 4.2$ Hz, 2H), 1.05 (dd, $J = 12.7, 2.5$ Hz, 2H), 0.91 (s, 6H), 0.67 (s, 6H), 0.64 (s, 6H). ^{13}C NMR (101 MHz, $\text{DMSO}-d_6$) δ 166.64, 155.57, 149.50, 148.24, 135.22, 120.63, 118.05, 107.09, 77.17, 55.73, 54.58, 49.44, 45.93, 44.41, 40.40, 39.56, 39.24, 38.43, 38.17, 37.07, 28.86, 28.15, 24.14, 21.64, 18.71, 16.25, 14.80. HRMS calc for $\text{C}_{56}\text{H}_{85}\text{N}_6\text{O}_6$ $[\text{M}+\text{H}]^+$: 937.6531, found 937.6528. HPLC purity: 100.00%, retention time: 17.500 min.

N,N'-(1,3-Phenylene)bis(4-((*E*)-5-((1*R*,4*aS*,6*R*,8*aS*)-6-hydroxy-5,5,8*a*-trimethyl-2-methylenedecahydronaphthalen-1-yl)-3-methylpent-2-enoyl)piperazine-1-carboxamide) (**15v**)

Light yellow oil, yield 91%, ^1H NMR (400 MHz, CDCl_3) δ 7.50 (s, 1H), 7.28 (s, 2H), 7.10 (t, $J = 8.1$ Hz, 1H), 6.95 (d, $J = 8.2$ Hz, 2H), 5.71 (s, 2H), 4.86 (s, 2H), 4.51 (s, 2H), 3.63 – 3.53 (m, 4H), 3.52 – 3.31 (m, 12H), 3.21 (dd, $J = 11.5, 4.4$ Hz, 2H), 2.44 – 2.36 (m, 2H), 2.29 – 2.19 (m, 2H), 1.98 – 1.88 (m, 4H), 1.87 (s, 6H), 1.80 – 1.48 (m, 14H), 1.37 (qd, $J = 12.9, 4.1$ Hz, 2H), 1.12 (td, $J = 13.4, 3.5$ Hz, 2H), 1.04 (dd, $J = 12.5, 2.7$ Hz, 2H), 0.97 (s, 6H), 0.75 (s, 6H), 0.67 (s, 6H). ^{13}C NMR (101 MHz, CDCl_3) δ 167.71, 155.53, 151.23, 147.79, 139.58, 129.01, 116.88, 116.01, 114.01, 106.96, 78.74, 55.93, 54.82, 46.04, 44.22, 43.86, 41.01, 39.49, 39.23, 38.59, 38.29, 37.25, 28.47, 27.94, 24.09, 21.62, 18.91, 15.58, 14.65. HRMS calc for $\text{C}_{56}\text{H}_{85}\text{N}_6\text{O}_6$ $[\text{M}+\text{H}]^+$: 937.6531, found 937.6530. HPLC purity: 99.45%, retention time: 4.390 min.

N,N'-(1,3-Phenylenebis(methylene)) bis(4-((*E*)-5-((1*R*,4*aS*,6*R*,8*aS*)-6-hydroxy-5,5,8a-trimethyl-2-methylenedecahydronaphthalen-1-yl)-3-methylpent-2-enoyl) piperazine-1-carboxamide) (**15w**)

Colorless oil, yield 93%, ¹H NMR (400 MHz, CDCl₃) δ 7.29 – 7.24 (m, 1H), 7.20 – 7.14 (m, 3H), 5.71 (s, 2H), 5.28 (d, *J* = 6.8 Hz, 2H), 4.86 (s, 2H), 4.51 (s, 2H), 4.36 (d, *J* = 5.7 Hz, 4H), 3.66 – 3.55 (m, 4H), 3.52 – 3.31 (m, 12H), 3.22 (dd, *J* = 11.7, 4.4 Hz, 2H), 2.44 – 2.35 (m, 2H), 2.29 – 2.19 (m, 2H), 1.98 – 1.88 (m, 4H), 1.87 (s, 6H), 1.78 – 1.53 (m, 14H), 1.38 (qd, *J* = 12.8, 4.0 Hz, 2H), 1.14 (td, *J* = 12.9, 2.8 Hz, 2H), 1.06 (dd, *J* = 11.6, 1.2 Hz, 2H), 0.98 (s, 6H), 0.76 (s, 6H), 0.68 (s, 6H). ¹³C NMR (101 MHz, CDCl₃) δ 167.64, 157.70, 151.22, 147.85, 139.87, 129.04, 127.00, 126.67, 116.86, 106.95, 78.82, 56.02, 54.83, 46.02, 44.93, 43.99, 43.87, 40.96, 39.53, 39.26, 38.69, 38.31, 37.25, 28.45, 27.99, 24.11, 21.69, 18.86, 15.55, 14.64. HRMS calc for C₅₈H₈₉N₆O₆ [M+H]⁺: 965.6844, found 965.6841. HPLC purity: 98.75%, retention time: 17.390 min.

N,N'-(Naphthalene-1,5-diyl)bis(4-((*E*)-5-((1*R*,4*aS*,6*R*,8*aS*)-6-hydroxy-5,5,8a-trimethyl-2-methylenedecahydronaphthalen-1-yl)-3-methylpent-2-enoyl)piperazine-1-carboxamide) (**15x**)

Colorless oil, yield 96%, ¹H NMR (400 MHz, CDCl₃) δ 7.84 (s, 2H), 7.60 – 7.52 (m, 2H), 7.37 – 7.29 (m, 2H), 7.30 – 7.24 (m, 2H), 5.66 (s, 2H), 4.85 (s, 2H), 4.50 (s, 2H), 3.32 – 3.14 (m, 14H), 3.07 – 2.92 (m, 4H), 2.39 (d, *J* = 13.6 Hz, 2H), 2.26 – 2.18 (m, 2H), 1.97 – 1.86 (m, 4H), 1.85 (s, 6H), 1.78 – 1.50 (m, 14H), 1.37 (qd, *J* = 12.8, 3.9 Hz, 2H), 1.13 (td, *J* = 10.7, 6.9 Hz, 2H), 1.03 (dd, *J* = 12.3, 2.4 Hz, 3H), 0.95 (s, 6H), 0.74 (s, 6H), 0.67 (s, 6H). ¹³C NMR (101 MHz, CDCl₃) δ 167.48, 156.66, 150.63, 147.75, 134.82, 130.84, 125.37, 123.85, 121.05, 116.96, 106.81, 78.53, 55.95, 54.73, 45.89, 43.79, 43.60, 40.80, 39.41, 39.12, 38.53, 38.20, 37.18, 28.36, 27.85, 24.01, 21.63, 18.77, 15.49, 14.56. HRMS calc for C₆₀H₈₆N₆O₆Na [M+Na]⁺: 1009.6507, found 1009.6505. HPLC purity: 97.51%, retention time: 17.890 min.

¹H NMR, ¹³C NMR, HRMS, ¹⁹F NMR and HPLC spectra of synthetic compounds

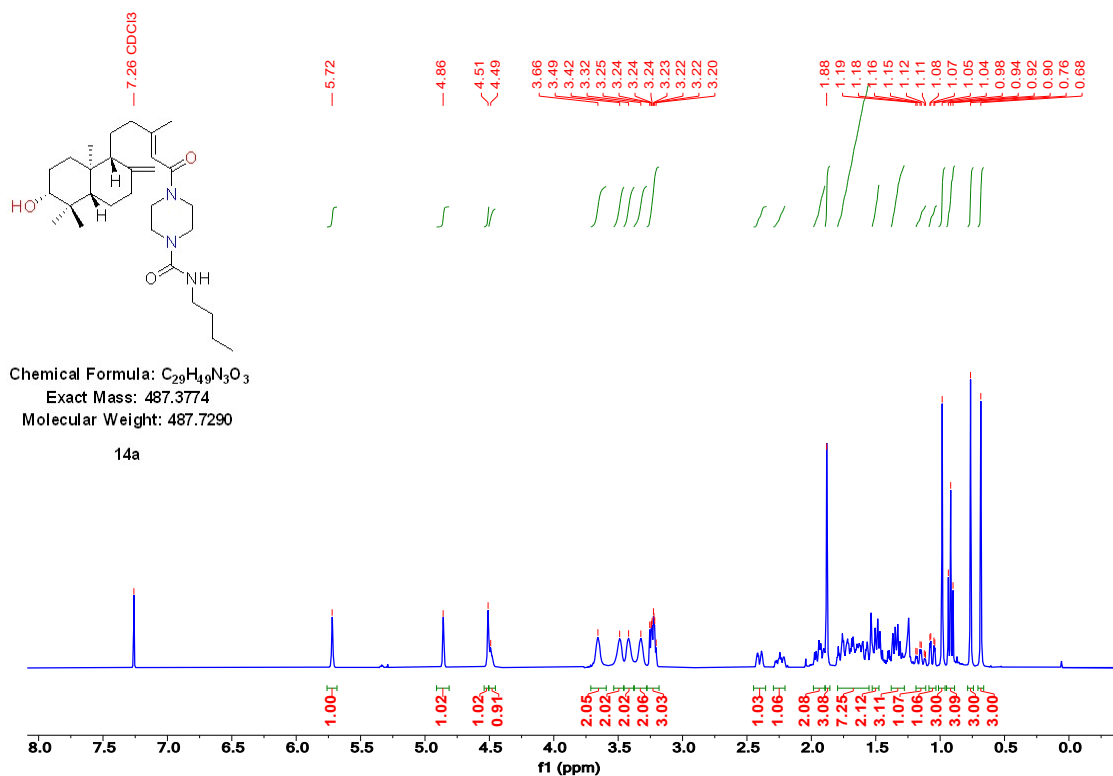


Figure S1. 1H NMR spectrum of 14a

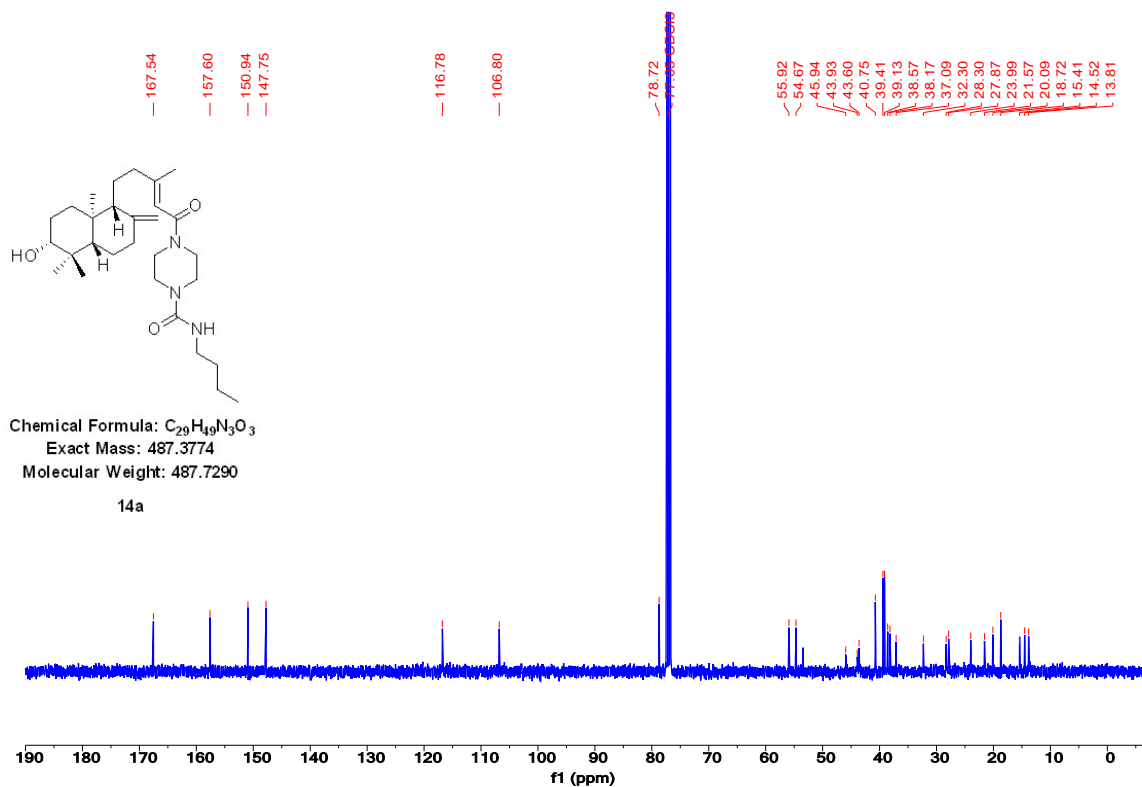


Figure S2. ^{13}C NMR spectrum of 14a

Single Mass Analysis

Tolerance = 5.0 mDa / DBE: min = -1.5, max = 50.0

Element prediction: Off

Number of isotope peaks used for i-FIT = 3

Monoisotopic Mass, Even Electron Ions

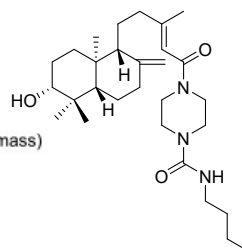
16 formula(e) evaluated with 1 results within limits (up to 50 best isotopic matches for each mass)

Elements Used:

C: 29-56 H: 47-85 N: 2-6 O: 2-6

7

250310-2-Q-21 21 (0.065)



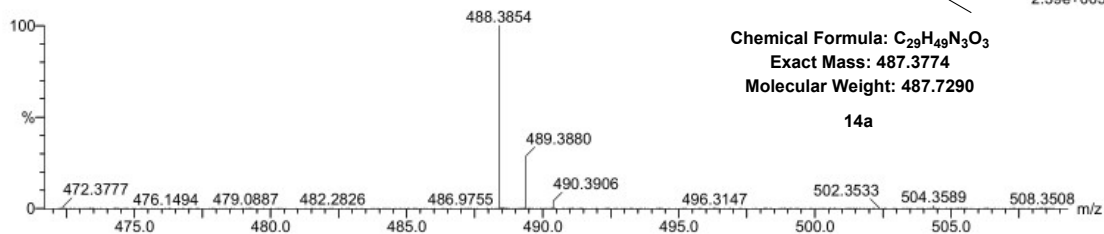
1: TOF MS ES+
2.39e+005

Chemical Formula: C₂₉H₄₉N₃O₃

Exact Mass: 487.3774

Molecular Weight: 487.7290

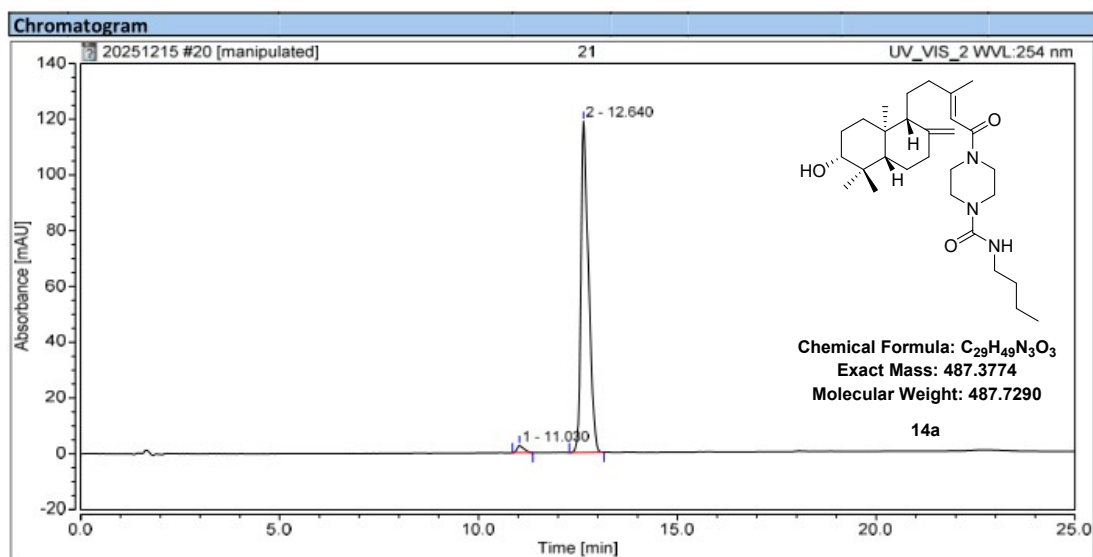
14a



Minimum: -1.5
Maximum: 5.0 10.0 50.0

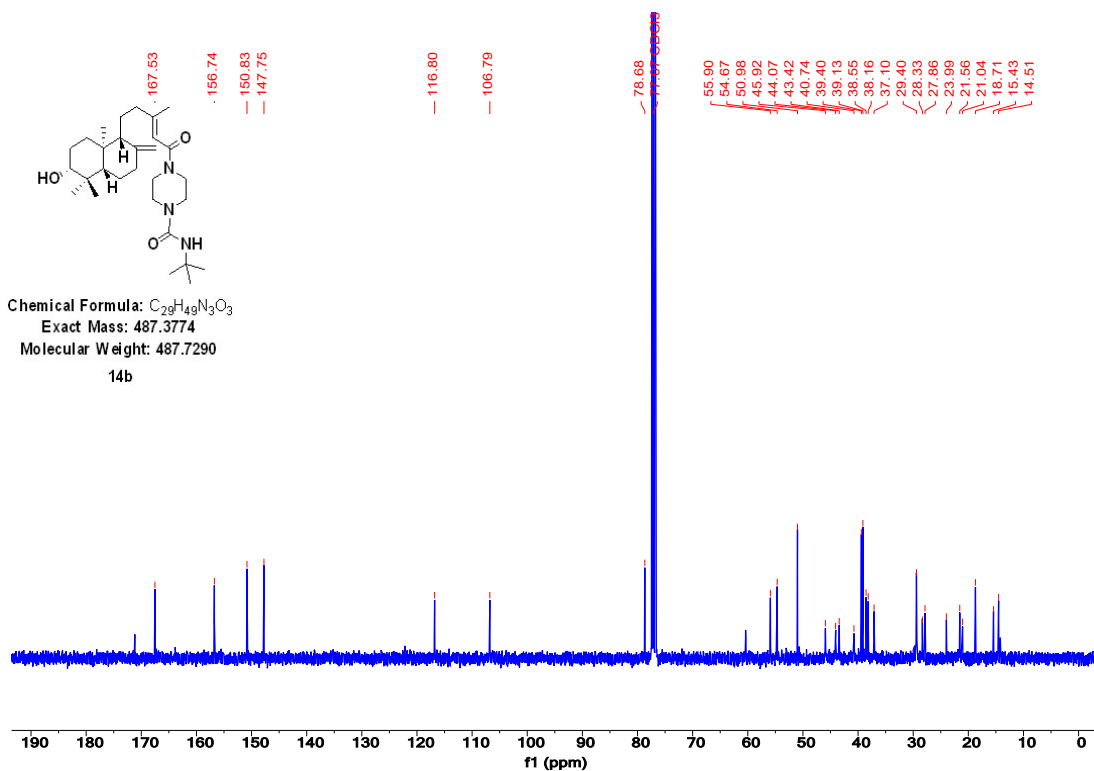
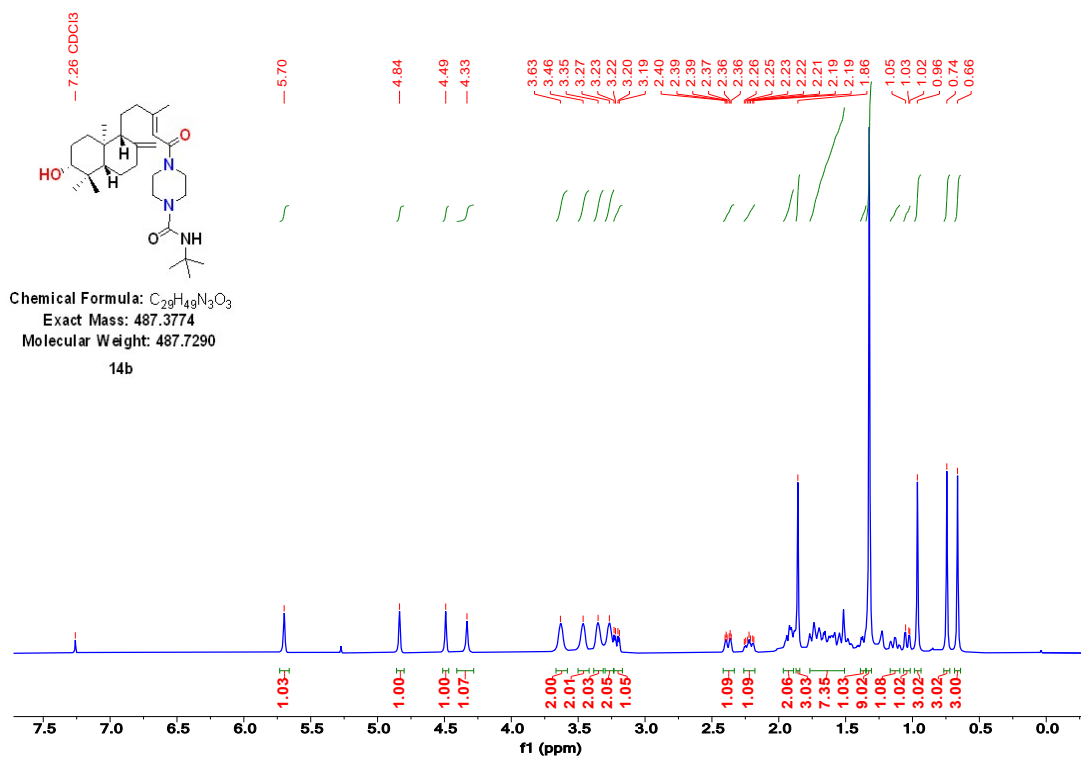
Mass	Calc. Mass	mDa	PPM	DBE	i-FIT	Norm	Conf (%)	Formula
488.3854	488.3852	0.2	0.4	6.5	475.7	n/a	n/a	C ₂₉ H ₅₀ N ₃ O ₃

Figure S3. HRMS spectrum of 14a



Integration Results							
No.	Peak Name	Retention Time min	Area mAU*min	Height mAU	Relative Area %	Relative Height %	Amount
1		11.030	0.525	2.598	1.97	2.14	n.a.
2		12.640	26.063	118.781	98.03	97.86	n.a.
Total:			26.588	121.379	100.00	100.00	

Figure S4. HPLC spectrum of 14a



Single Mass Analysis

Tolerance = 5.0 mDa / DBE: min = -1.5, max = 50.0

Element prediction: Off

Number of isotope peaks used for i-FIT = 3

Monoisotopic Mass, Even Electron Ions

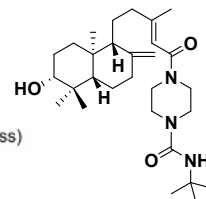
9 formula(e) evaluated with 1 results within limits (up to 50 best isotopic matches for each mass)

Elements Used:

C: 27-33 H: 44-51 N: 2-5 O: 2-5

7

250310-2-Q-14 32 (0.086)



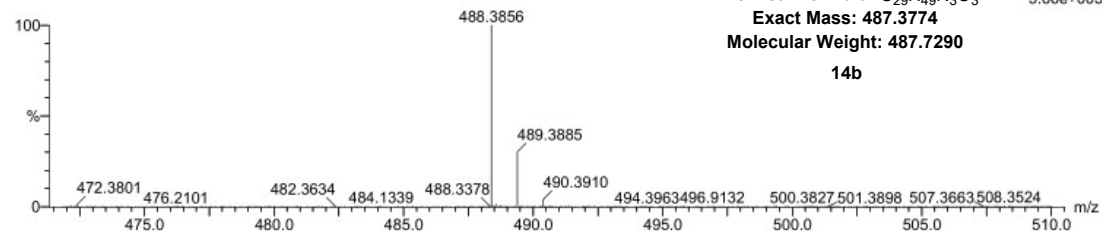
Chemical Formula: $C_{29}H_{49}N_3O_3$

Exact Mass: 487.3774

Molecular Weight: 487.7290

14b

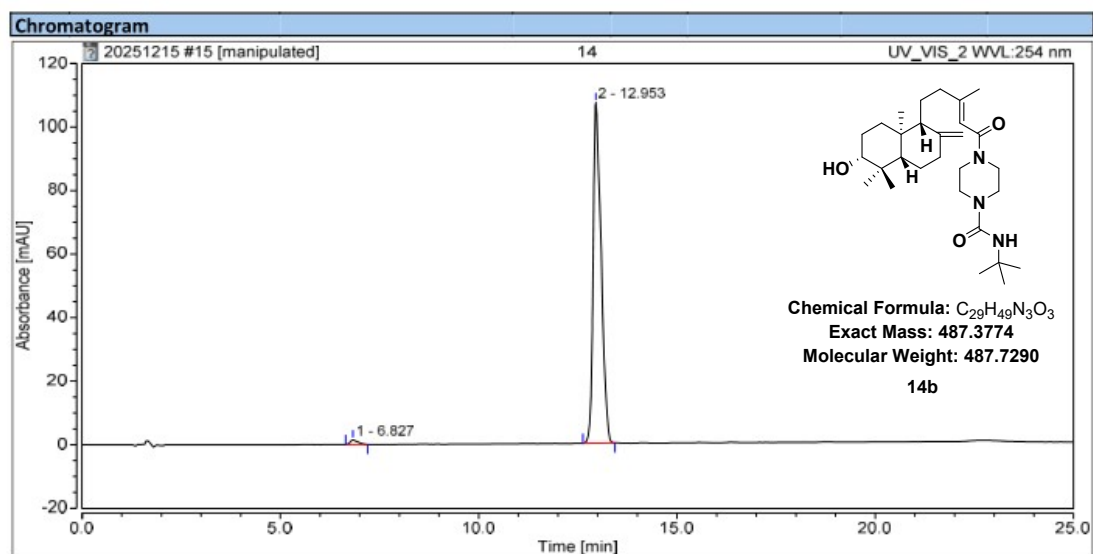
1: TOF MS ES+
5.66e+005



Minimum: -1.5
Maximum: 5.0 10.0 50.0

Mass	Calc. Mass	mDa	PPM	DBE	i-FIT	Norm	Conf (%)	Formula
488.3856	488.3852	0.4	0.8	6.5	612.9	n/a	n/a	C29 H50 N3 O3

Figure S7. HRMS spectrum of 14b



Integration Results							
No.	Peak Name	Retention Time min	Area mAU*min	Height mAU	Relative Area %	Relative Height %	Amount
1		6.827	0.322	1.453	1.29	1.34	n.a.
2		12.953	24.750	107.268	98.71	98.66	n.a.
Total:			25.073	108.721	100.00	100.00	

Figure S8. HPLC spectrum of 14b

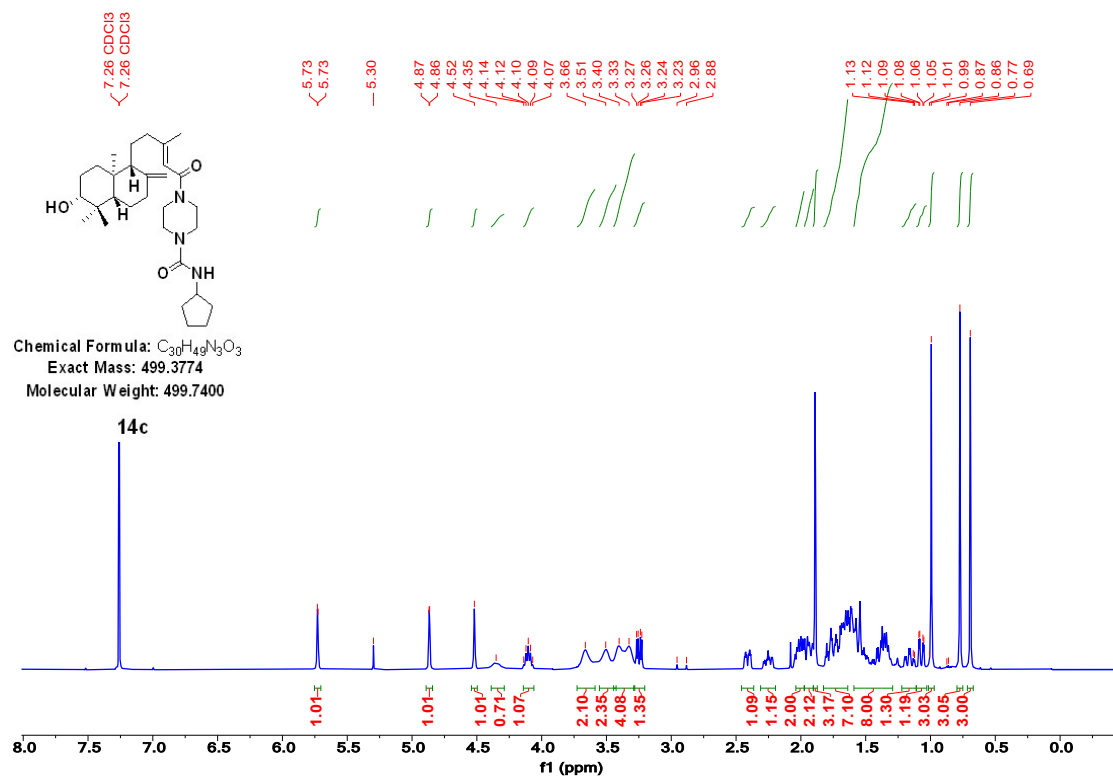


Figure S9. 1H NMR spectrum of **14c**

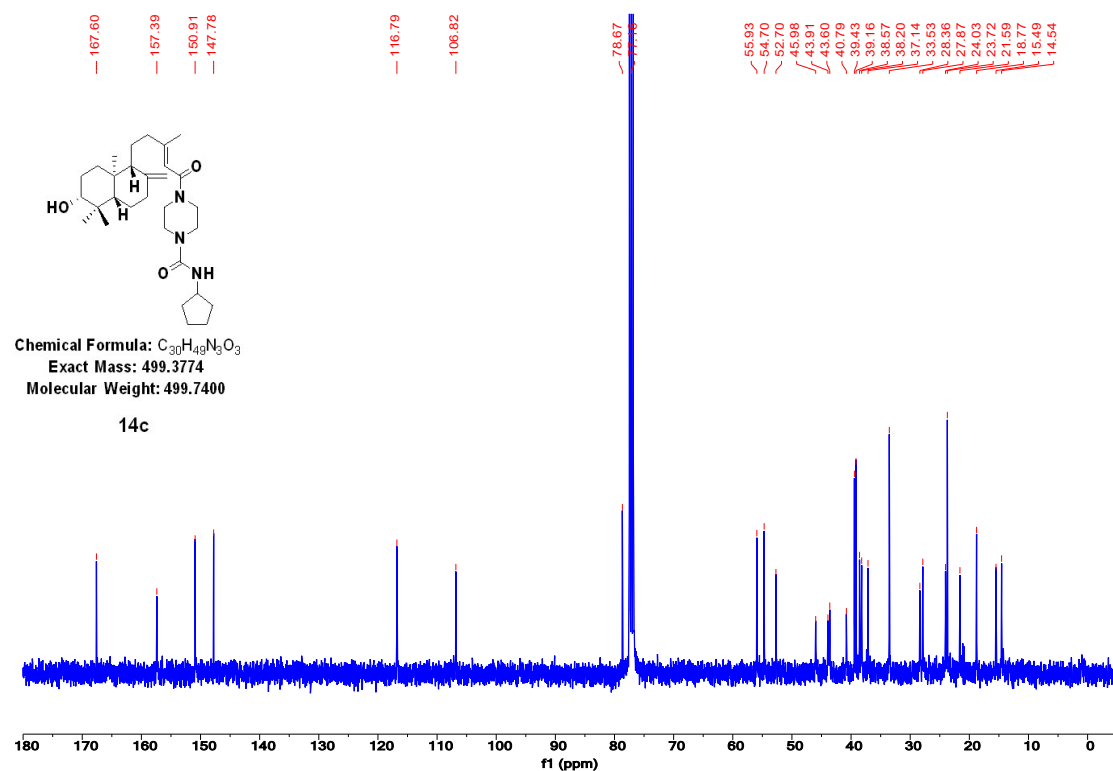


Figure S10. ^{13}C NMR spectrum of **14c**

Single Mass Analysis

Tolerance = 5.0 mDa / DBE: min = -1.5, max = 50.0

Element prediction: Off

Number of isotope peaks used for i-FIT = 3

Monoisotopic Mass, Even Electron Ions

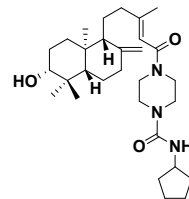
9 formula(e) evaluated with 1 results within limits (up to 50 best isotopic matches for each mass)

Elements Used:

C: 27-33 H: 44-51 N: 2-5 O: 2-5

7

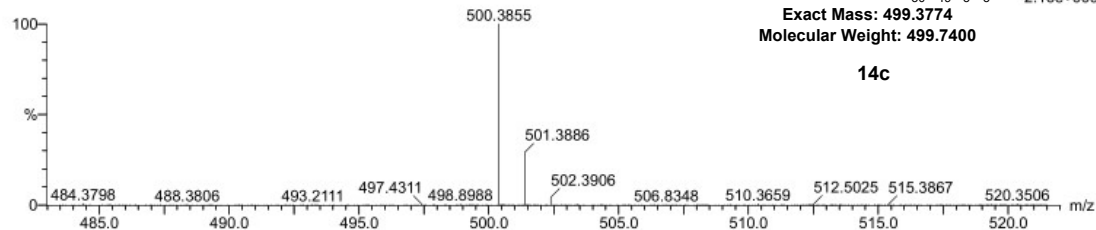
250310-2-Q-15 41 (0.103)



Chemical Formula: $C_{30}H_{49}N_3O_3$
Exact Mass: 499.3774
Molecular Weight: 499.7400

1: TOF MS ES+
2.16e+005

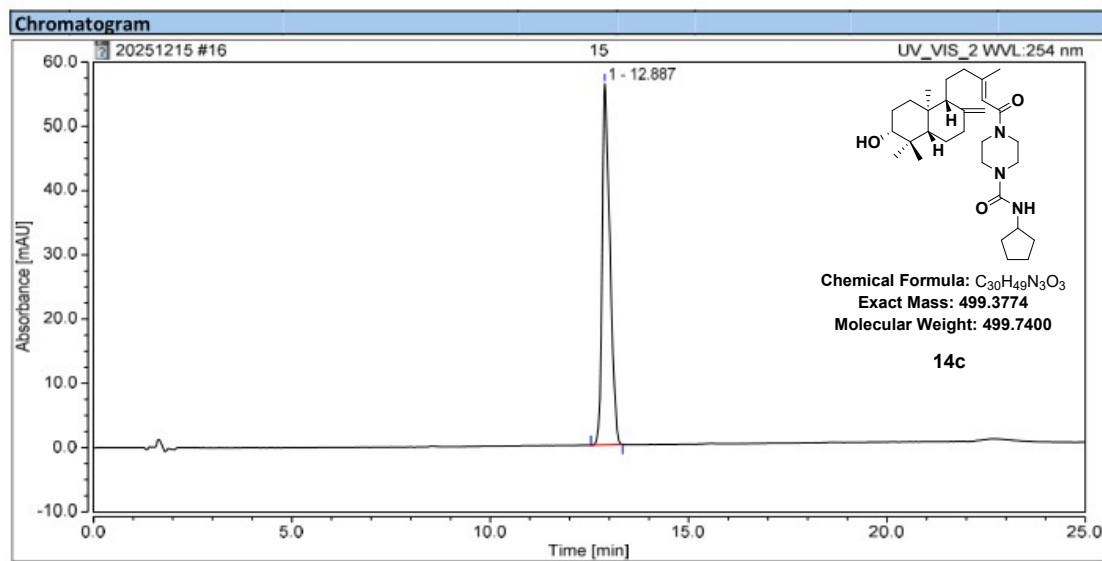
14c



Minimum: -1.5
Maximum: 5.0 10.0 50.0

Mass	Calc. Mass	mDa	PPM	DBE	i-FIT	Norm	Conf(%)	Formula
500.3855	500.3852	0.3	0.6	7.5	406.1	n/a	n/a	C30 H50 N3 O3

Figure S11. HRMS spectrum of 14c



Integration Results							
No.	Peak Name	Retention Time min	Area mAU*min	Height mAU	Relative Area %	Relative Height %	Amount
1		12.887	12.664	56.239	100.00	100.00	n.a.
Total:			12.664	56.239	100.00	100.00	

Figure S12. HPLC spectrum of 14c

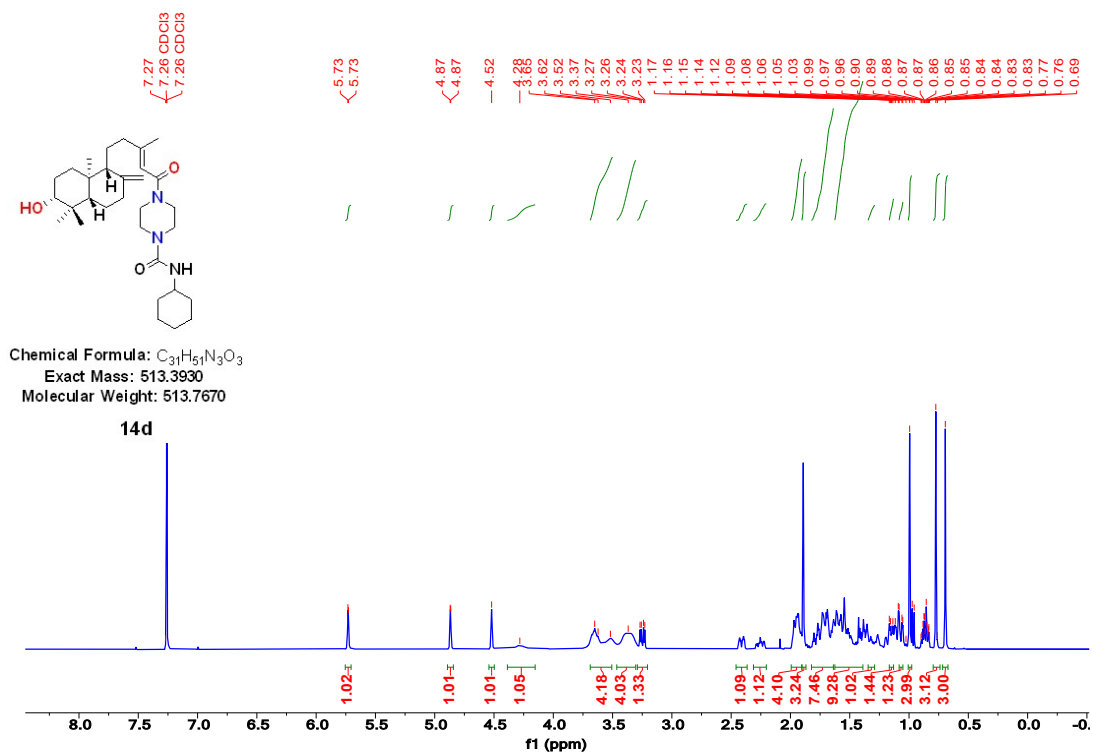


Figure S13. ^1H NMR spectrum of **14d**

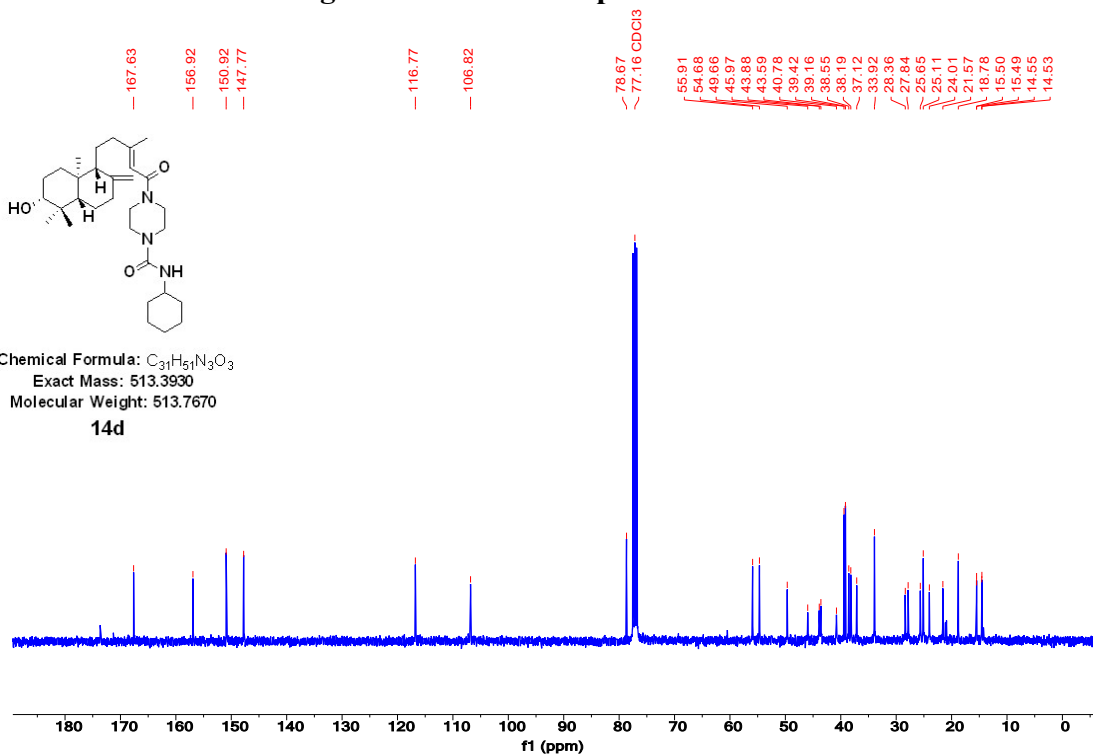


Figure S14. ^{13}C NMR spectrum of **14d**

Single Mass Analysis

Tolerance = 5.0 mDa / DBE: min = -1.5, max = 50.0

Element prediction: Off

Number of isotope peaks used for i-FIT = 3

Monoisotopic Mass, Even Electron Ions

11 formula(e) evaluated with 1 results within limits (up to 50 best isotopic matches for each mass)

Elements Used:

C: 27-33 H: 44-52 N: 2-5 O: 2-5

7

250310-2-Q-17 20 (0.063)

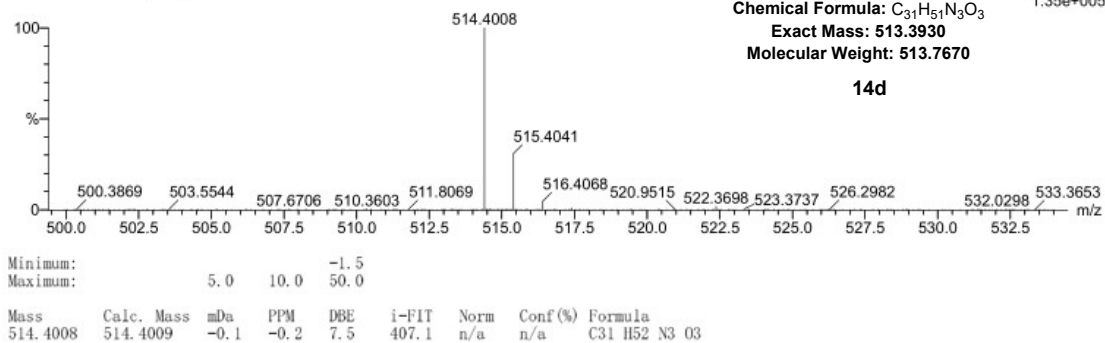


Figure S15. HRMS spectrum of 14d

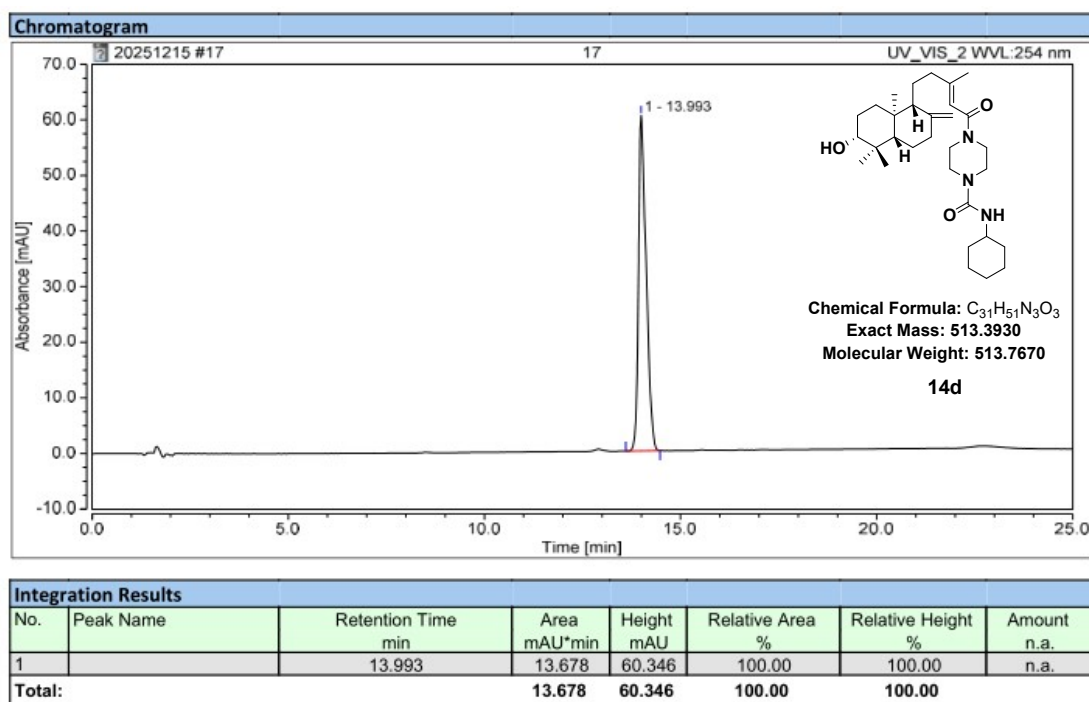


Figure S16. HPLC spectrum of 14d

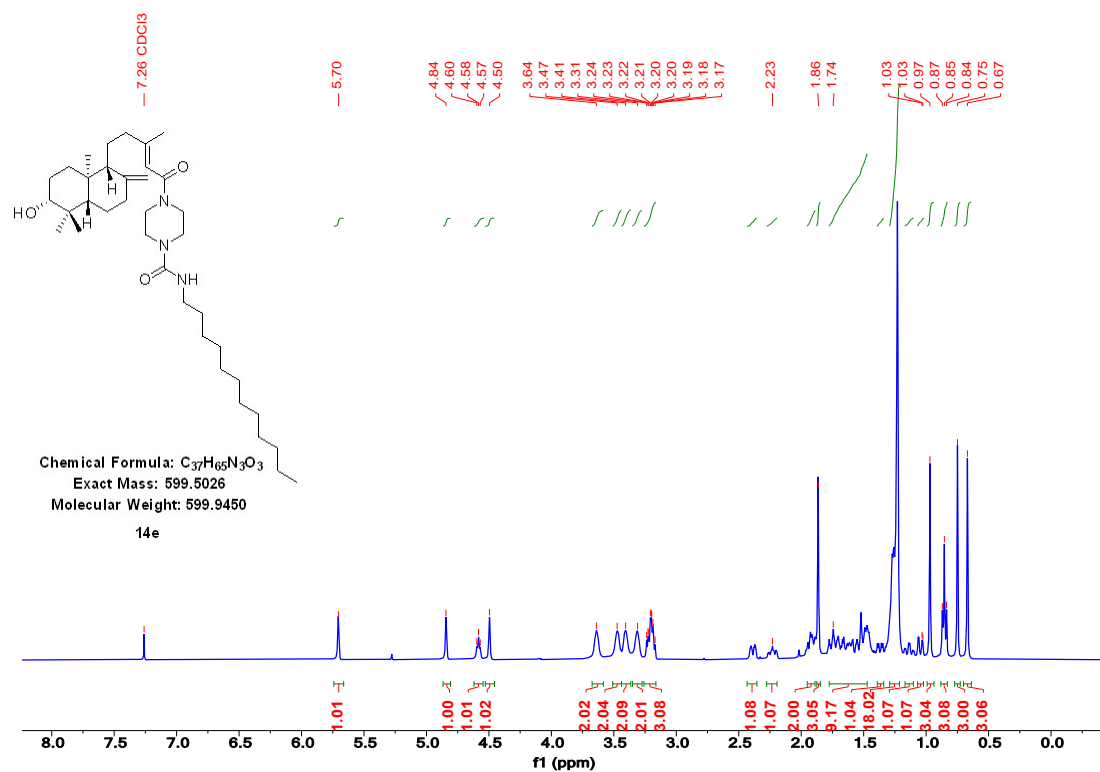


Figure S17. 1H NMR spectrum of 14e

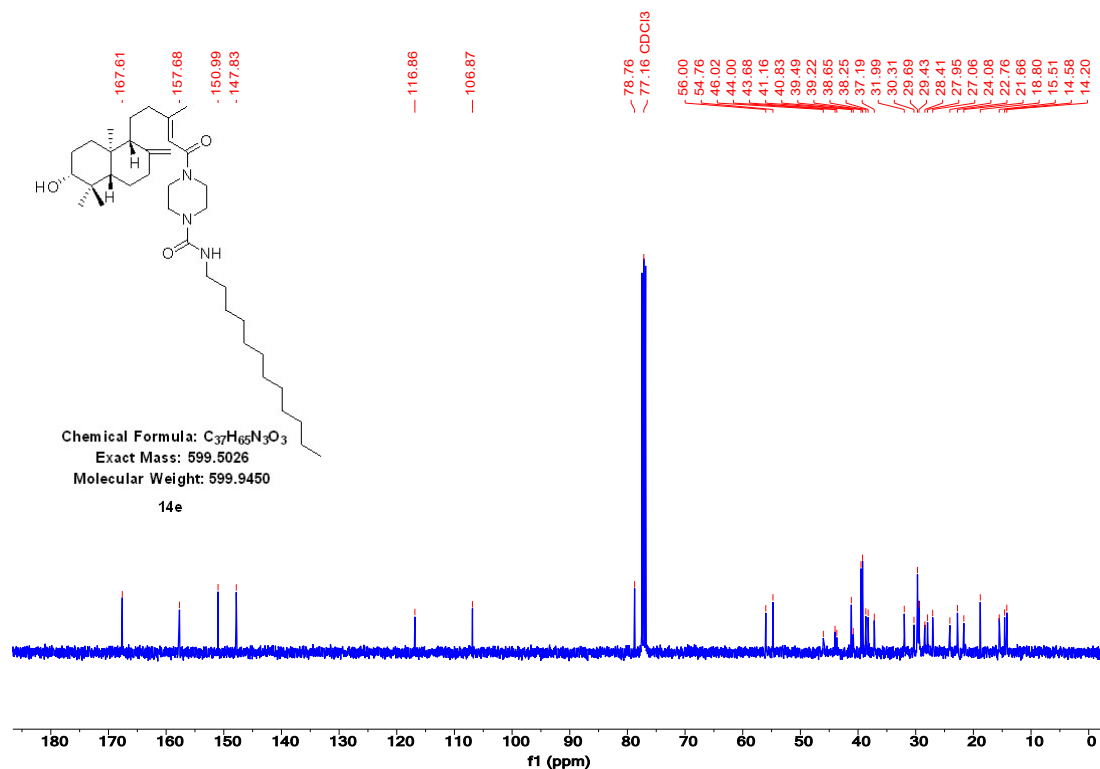


Figure S18. ^{13}C NMR spectrum of 14e

Single Mass Analysis

Tolerance = 5.0 mDa / DBE: min = -1.5, max = 50.0

Element prediction: Off

Number of isotope peaks used for i-FIT = 3

Monoisotopic Mass, Even Electron Ions

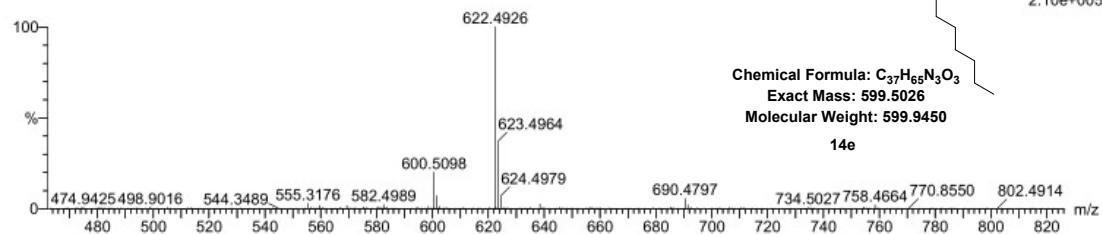
5 formula(e) evaluated with 1 results within limits (up to 50 best isotopic matches for each mass)

Elements Used:

C: 37-37 H: 63-65 N: 2-6 O: 2-6 Na: 0-1

7

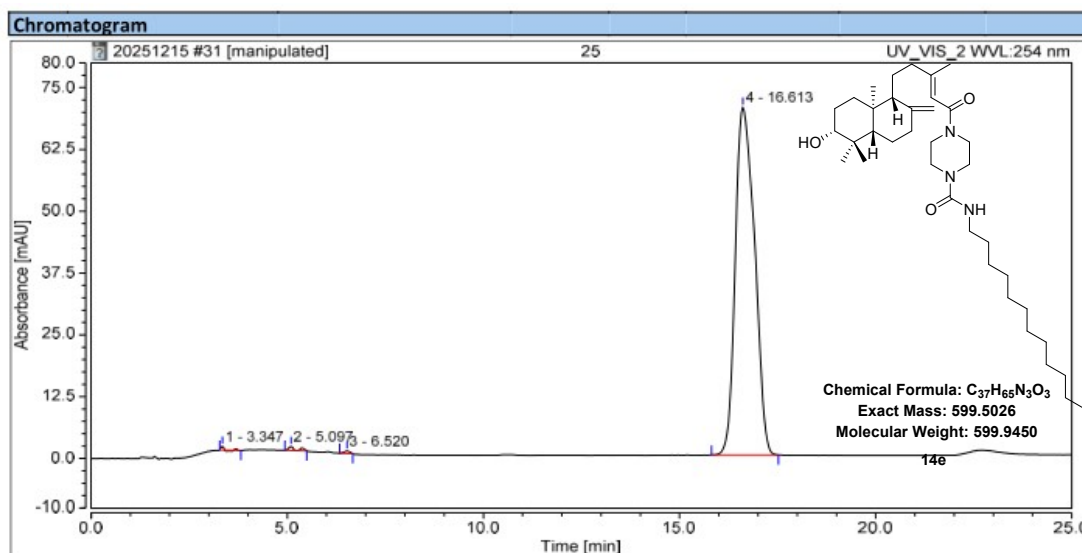
250310-2-Q-25 26 (0.075)



Minimum: -1.5
Maximum: 5.0 10.0 50.0

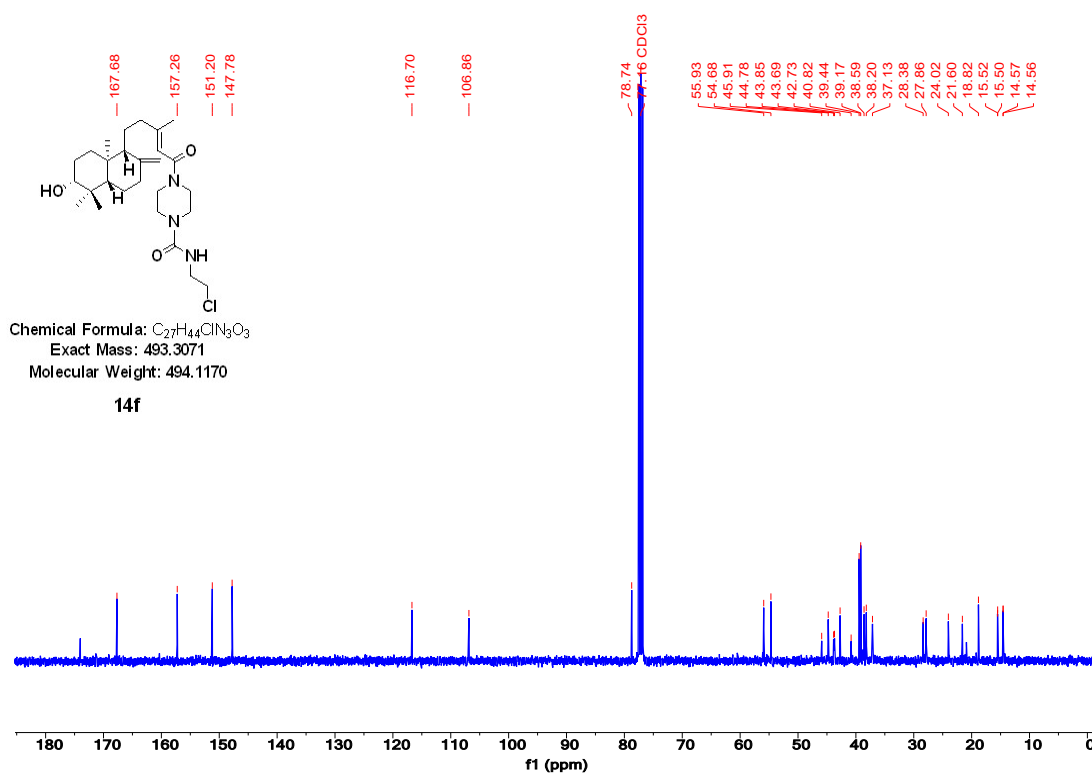
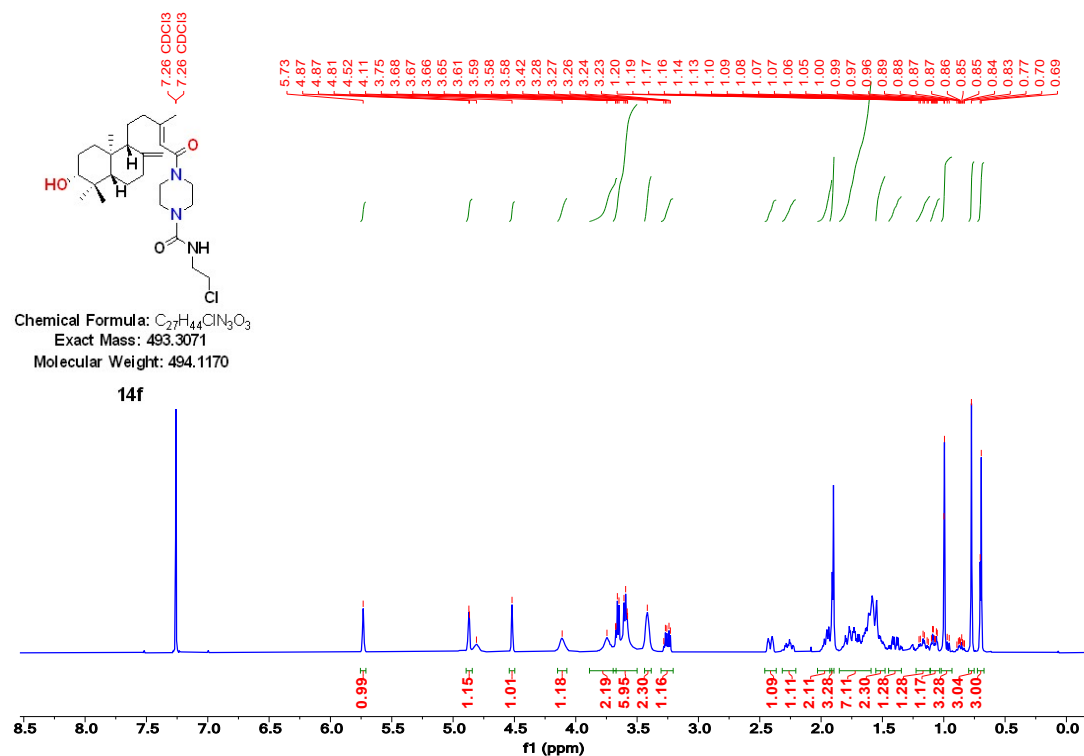
Mass	Calc. Mass	mDa	PPM	DBE	i-FIT	Norm	Conf (%)	Formula
622.4926	622.4924	0.2	0.3	6.5	446.1	n/a	n/a	C37 H65 N3 O3 Na

Figure S19. HRMS spectrum of 14e



No.	Peak Name	Retention Time min	Area mAU*min	Height mAU	Relative Area %	Relative Height %	Amount
1		3.347	0.056	0.817	0.14	1.13	n.a.
2		5.097	0.172	0.800	0.43	1.10	n.a.
3		6.520	0.087	0.516	0.22	0.71	n.a.
4		16.613	40.046	70.367	99.22	97.06	n.a.
Total:			40.361	72.499	100.00	100.00	

Figure S20. HPLC spectrum of 14e



Single Mass Analysis

Tolerance = 5.0 mDa / DBE: min = -1.5, max = 50.0
Element prediction: Off
Number of isotope peaks used for i-FIT = 3

Monoisotopic Mass, Even Electron Ions

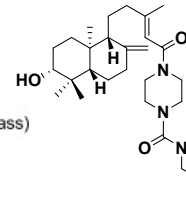
31 formula(e) evaluated with 1 results within limits (up to 50 best isotopic matches for each mass)

Elements Used:

C: 27-33 H: 44-51 N: 2-5 O: 2-5 Na: 0-1 Cl: 0-1

7

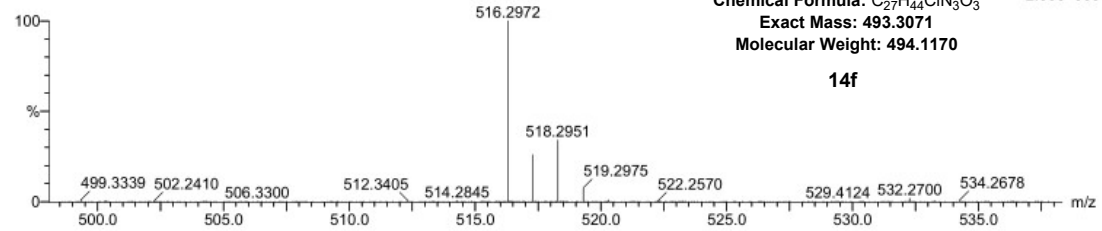
250310-2-Q-13 40 (0.101)



Chemical Formula: $C_{27}H_{44}ClN_3O_3$
Exact Mass: 493.3071
Molecular Weight: 494.1170

14f

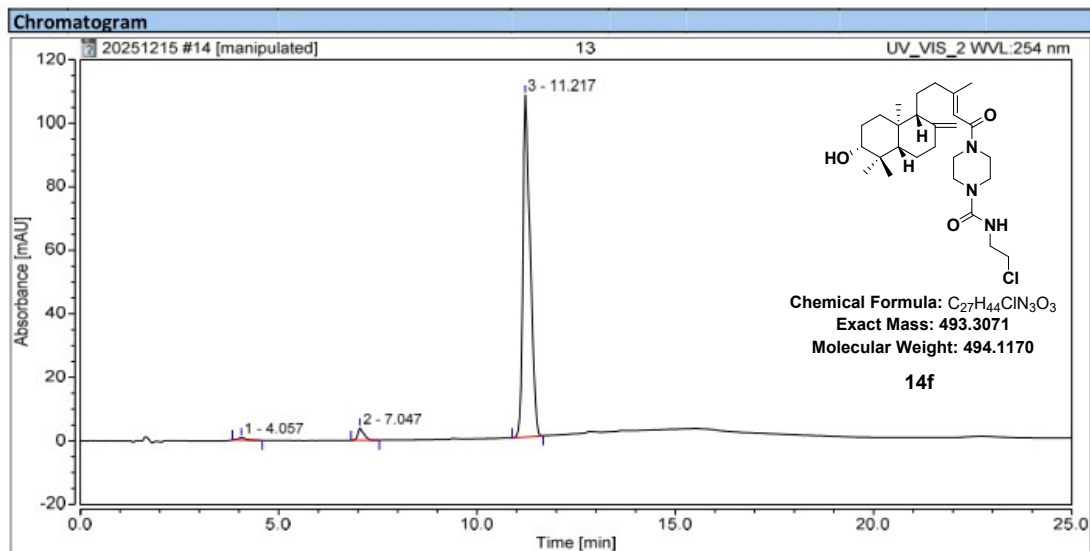
1: TOF MS ES+
2.55e+005



Minimum: -1.5
Maximum: 50.0

Mass	Calc. Mass	mDa	PPM	DBE	i-FIT	Norm	Conf (%)	Formula
516.2972	516.2969	0.3	0.6	6.5	476.1	n/a	n/a	C27 H44 N3 O3 Na Cl

Figure S23. HRMS spectrum of 14f



Chemical Formula: $C_{27}H_{44}ClN_3O_3$
Exact Mass: 493.3071
Molecular Weight: 494.1170

14f

Integration Results							
No.	Peak Name	Retention Time min	Area mAU*min	Height mAU	Relative Area %	Relative Height %	Amount n.a.
1		4.057	0.156	0.743	0.66	0.66	n.a.
2		7.047	0.679	3.871	2.88	3.45	n.a.
3		11.217	22.693	107.674	96.45	95.89	n.a.
Total:			23.527	112.288	100.00	100.00	

Figure S24. HPLC spectrum of 14f

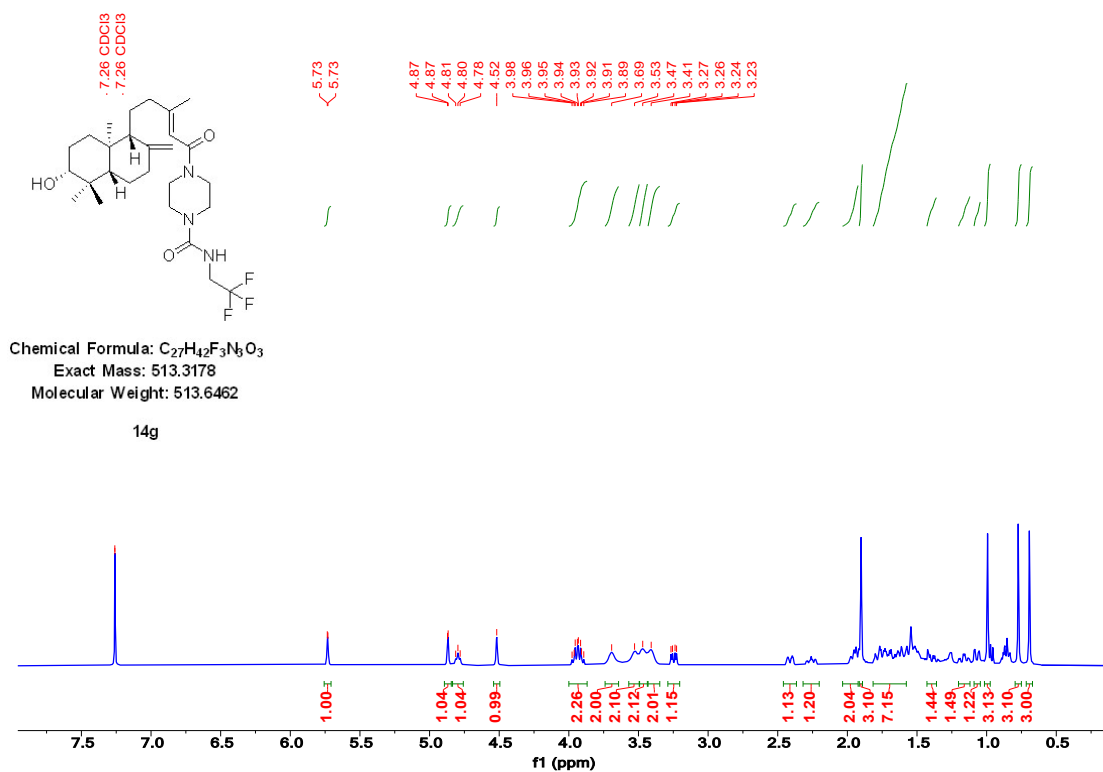


Figure S25. ¹H NMR spectrum of 14g

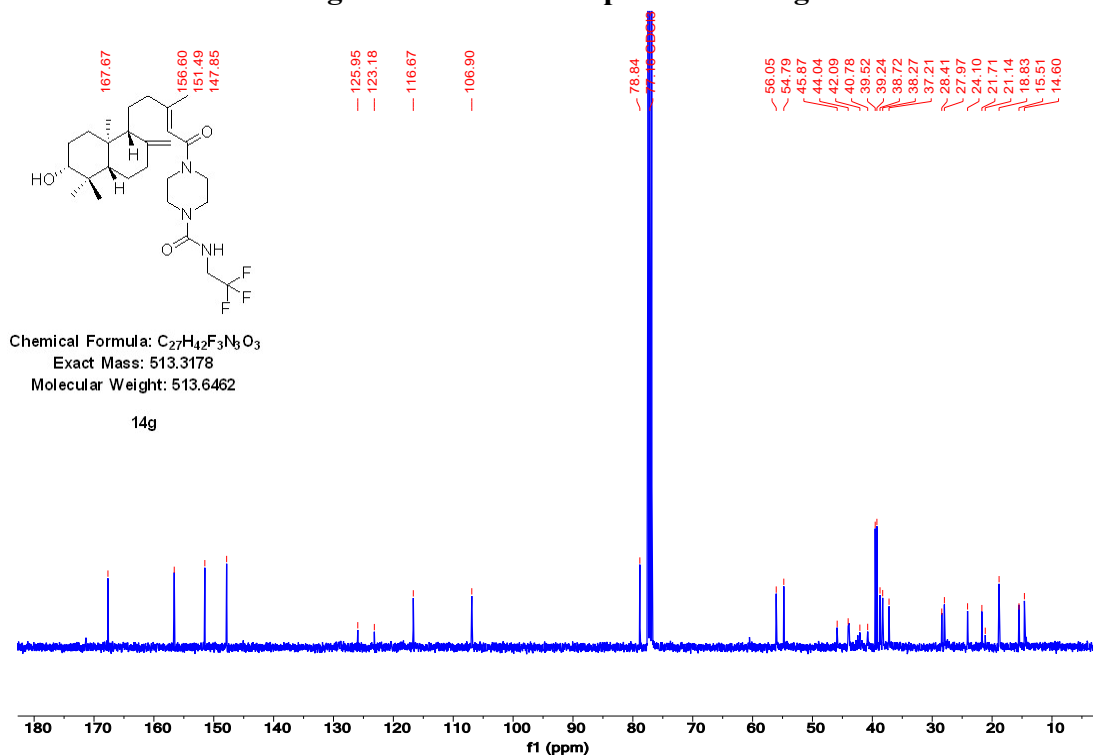


Figure S26. ¹³C NMR spectrum of 14g

Single Mass Analysis

Tolerance = 5.0 mDa / DBE: min = -1.5, max = 50.0

Element prediction: Off

Number of isotope peaks used for i-FIT = 3

Monoisotopic Mass, Even Electron Ions

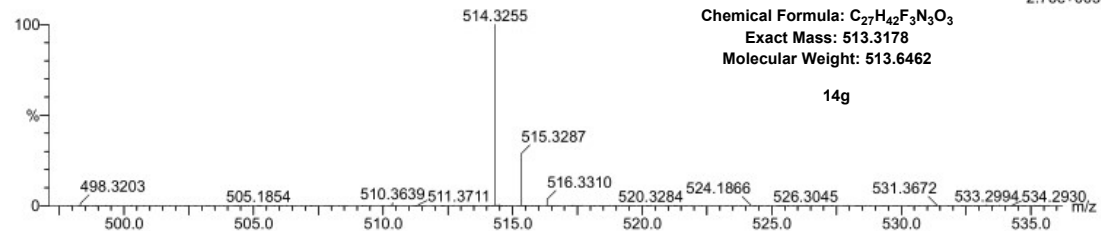
16 formula(e) evaluated with 1 results within limits (up to 50 best isotopic matches for each mass)

Elements Used:

C: 27-27 H: 42-43 N: 2-6 O: 2-6 F: 0-3

7

250310-2-Q-35 20 (0.063)



Chemical Formula: C₂₇H₄₂F₃N₃O₃

Exact Mass: 513.3178

Molecular Weight: 513.6462

14g

1: TOF MS ES+
2.76e+005

Minimum: -1.5
Maximum: 5.0 10.0 50.0

Mass	Calc. Mass	mDa	PPM	DBE	i-FIT	Norm	Conf (%)	Formula
514.3255	514.3257	-0.2	-0.4	6.5	418.6	n/a	n/a	C ₂₇ H ₄₃ N ₃ O ₃ F ₃

Figure S27. HRMS spectrum of 14g

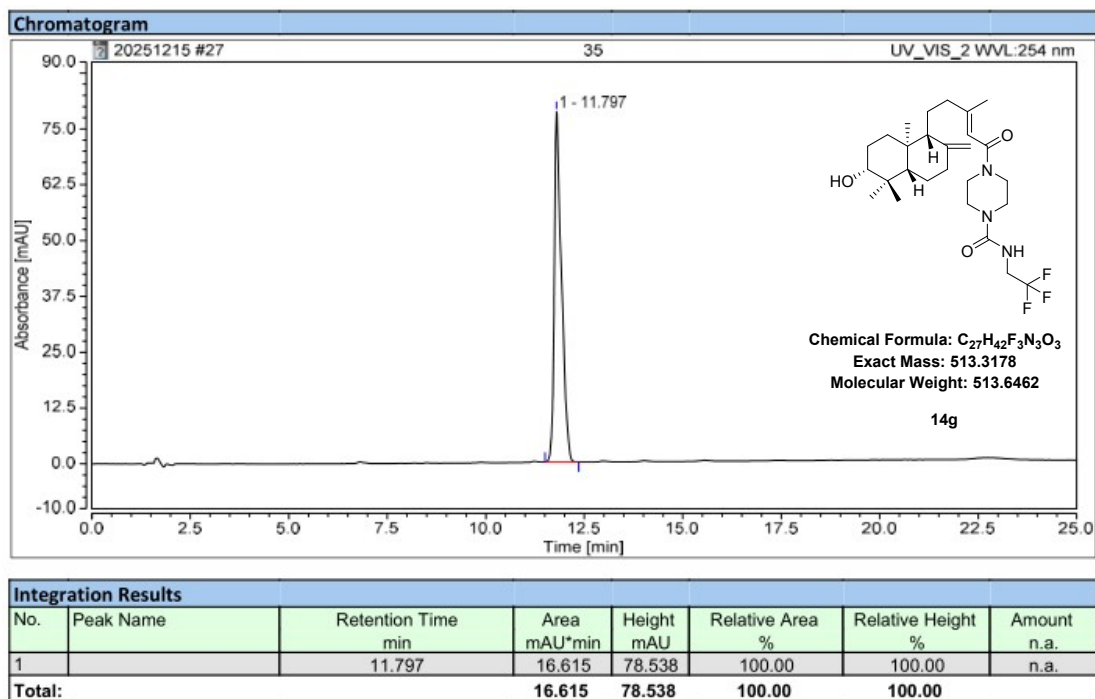
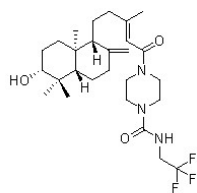


Figure S28. HPLC spectrum of 14g

^{19}F NMR (376 MHz, CDCl_3) -72.99.



Chemical Formula: $\text{C}_{27}\text{H}_{42}\text{F}_3\text{N}_3\text{O}_3$

Exact Mass: 513.3178

Molecular Weight: 513.6462

14g

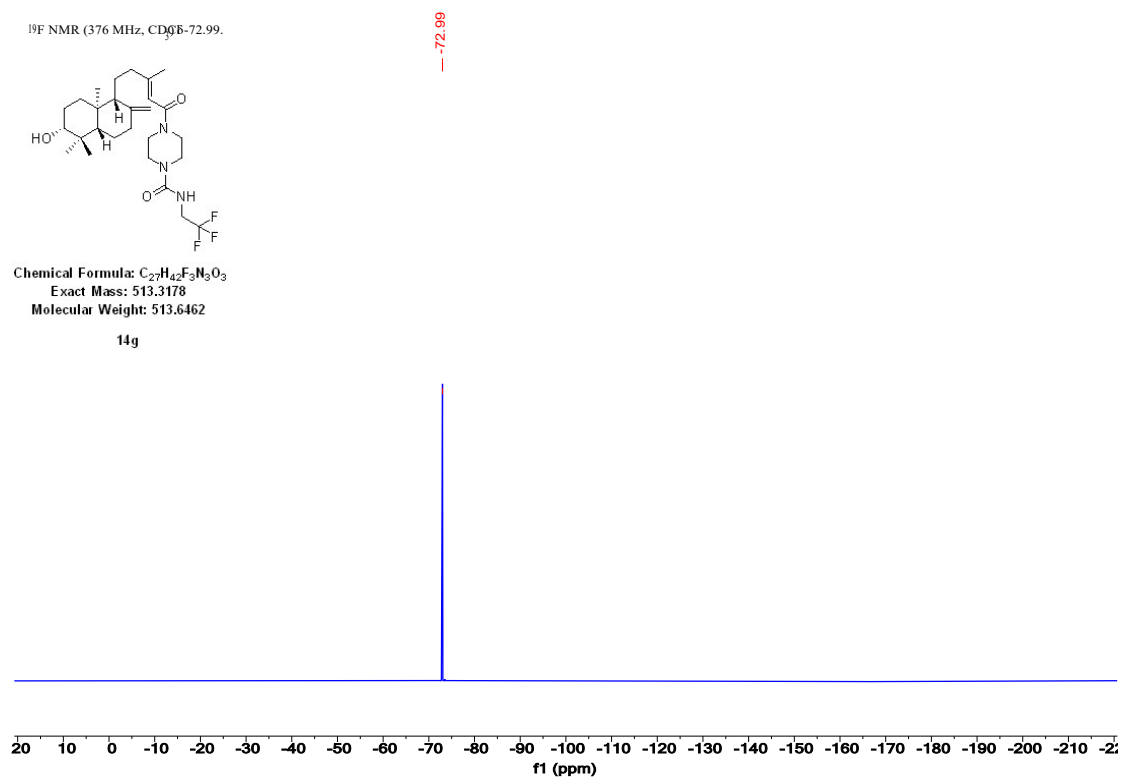


Figure S29. ^{19}F NMR spectrum of 14g

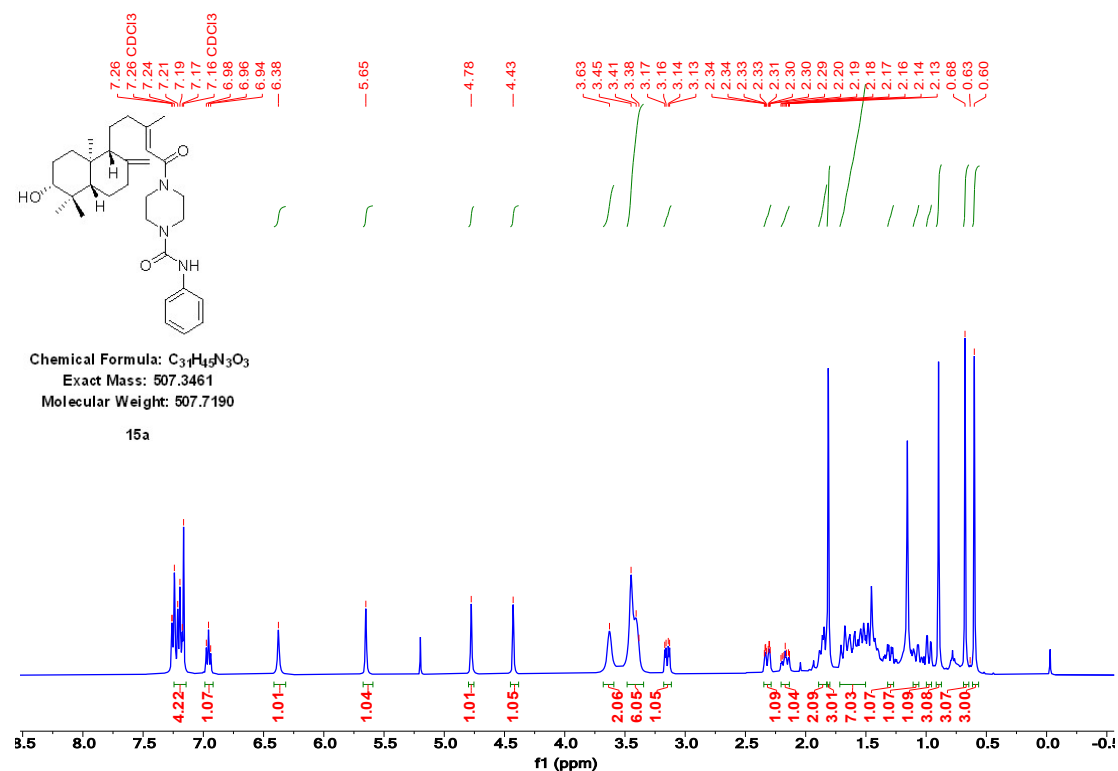


Figure S30. ^1H NMR spectrum of 15a

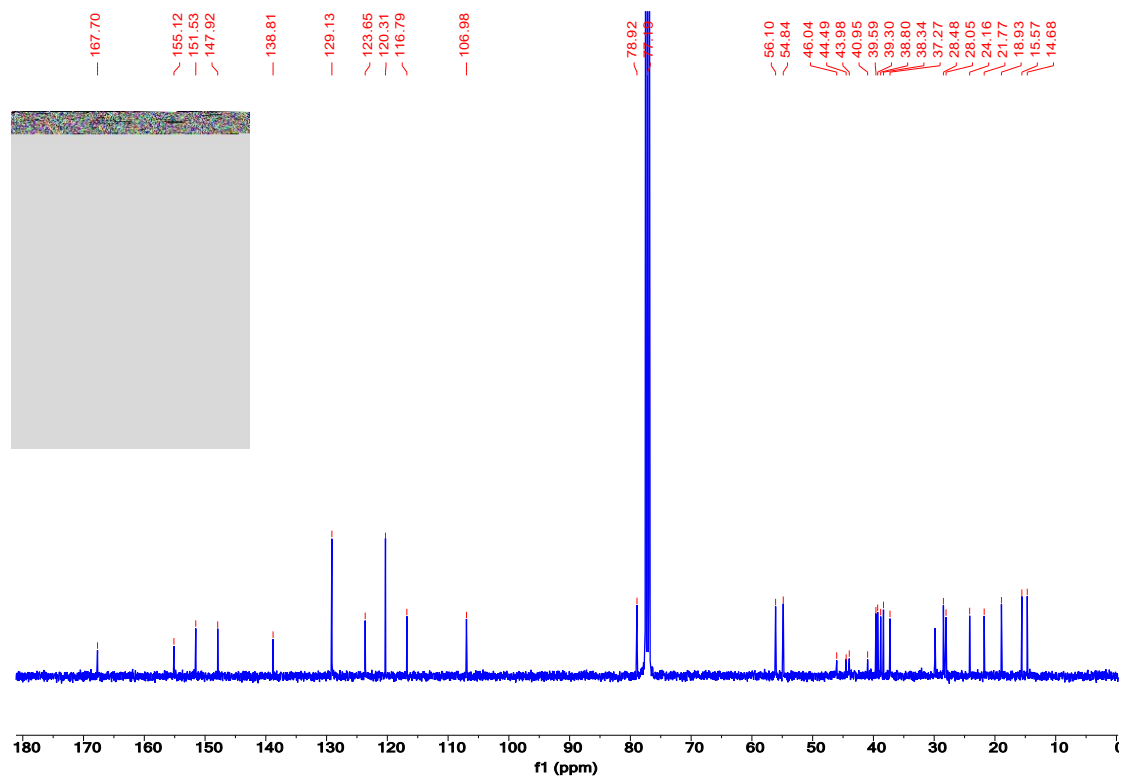


Figure S31. ^{13}C NMR spectrum of 15a

Single Mass Analysis

Tolerance = 5.0 mDa / DBE: min = -1.5, max = 50.0

Element prediction: Off

Number of isotope peaks used for i-FIT = 3

Monoisotopic Mass, Even Electron Ions

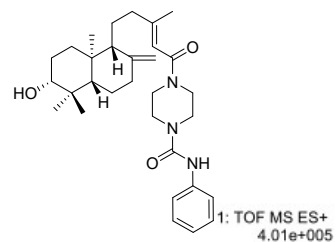
48 formula(e) evaluated with 1 results within limits (up to 50 best isotopic matches for each mass)

Elements Used:

C: 31-32 H: 24-45 N: 2-7 O: 2-7 Na: 0-1

7

250310-2-Q-41 21 (0.065)

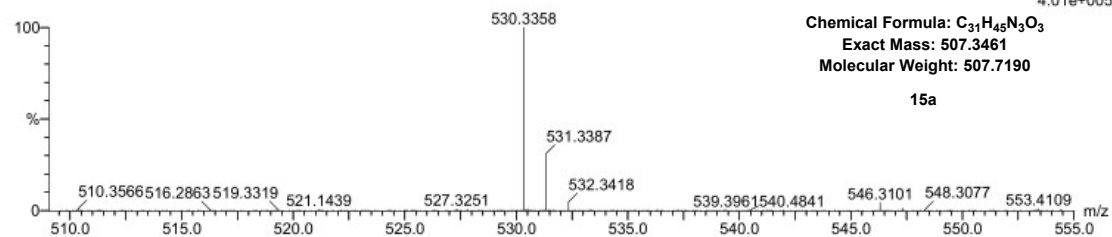


Chemical Formula: $\text{C}_{31}\text{H}_{45}\text{N}_3\text{O}_3$

Exact Mass: 507.3461

Molecular Weight: 507.7190

15a



Minimum: -1.5
Maximum: 5.0 10.0 50.0

Mass	Calc. Mass	mDa	PPM	DBE	i-FIT	Norm	Conf(%)	Formula
530.3358	530.3359	-0.1	-0.2	10.5	472.2	n/a	n/a	C31 H45 N3 O3 Na

Figure S32. HRMS spectrum of 15a

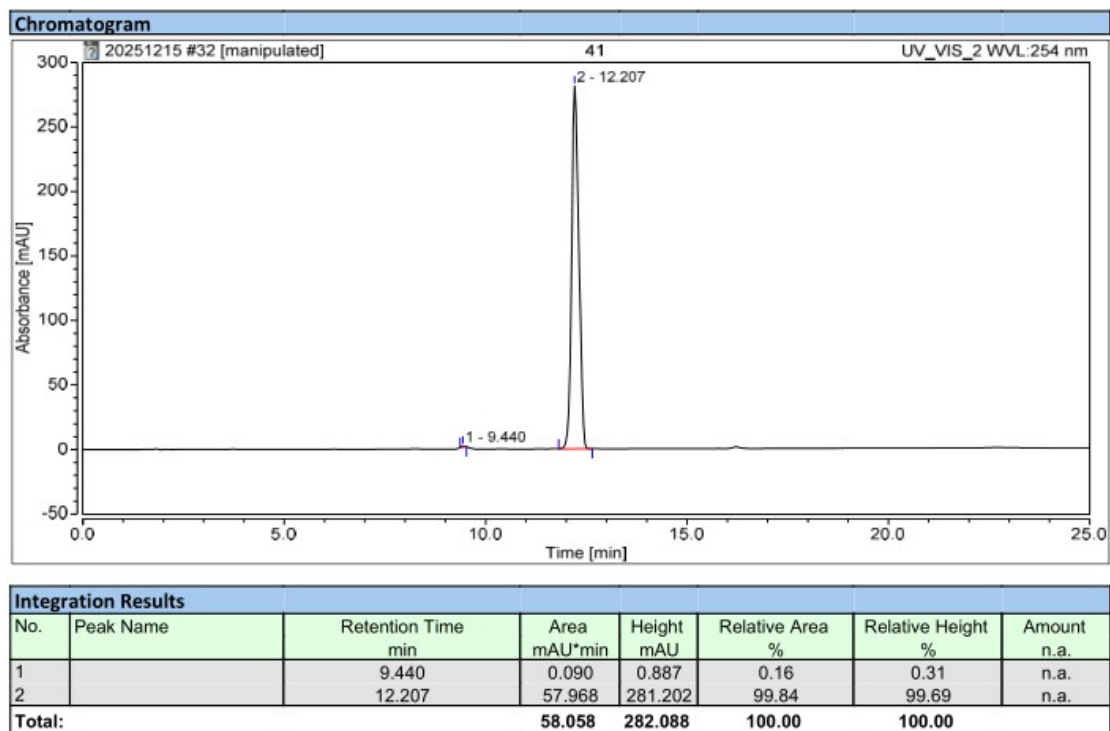


Figure S33. HPLC spectrum of 15a

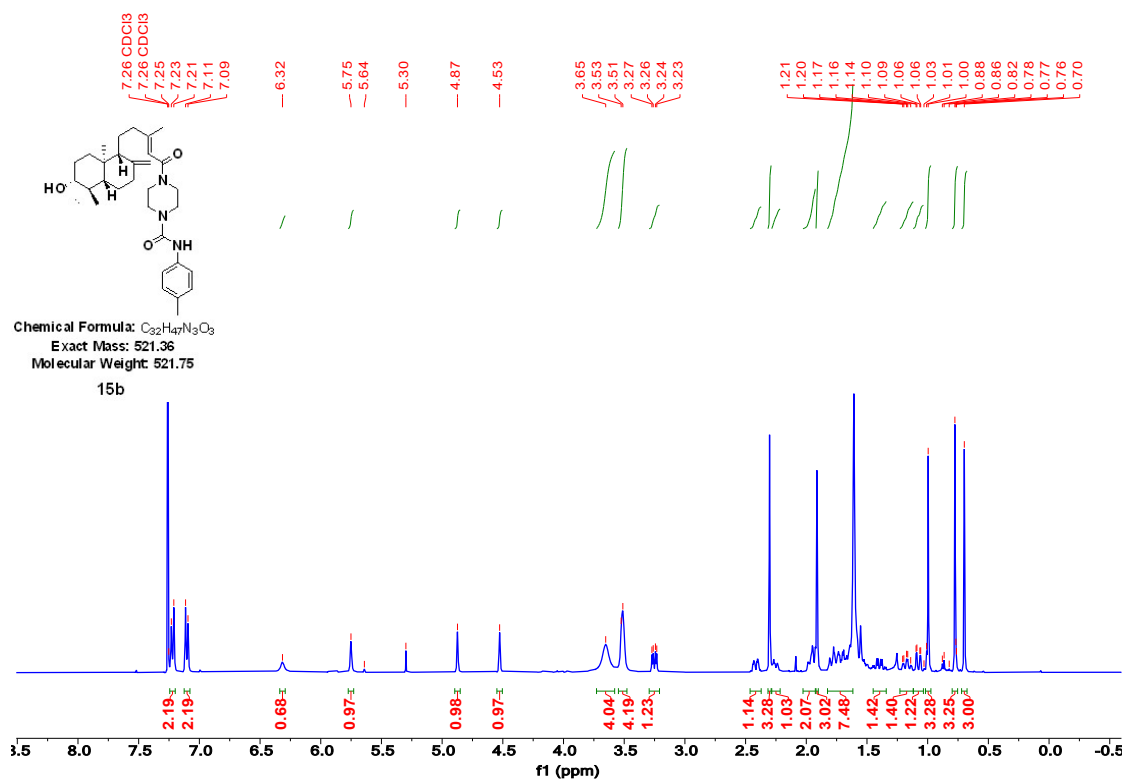


Figure S34. ^1H NMR spectrum of 15b

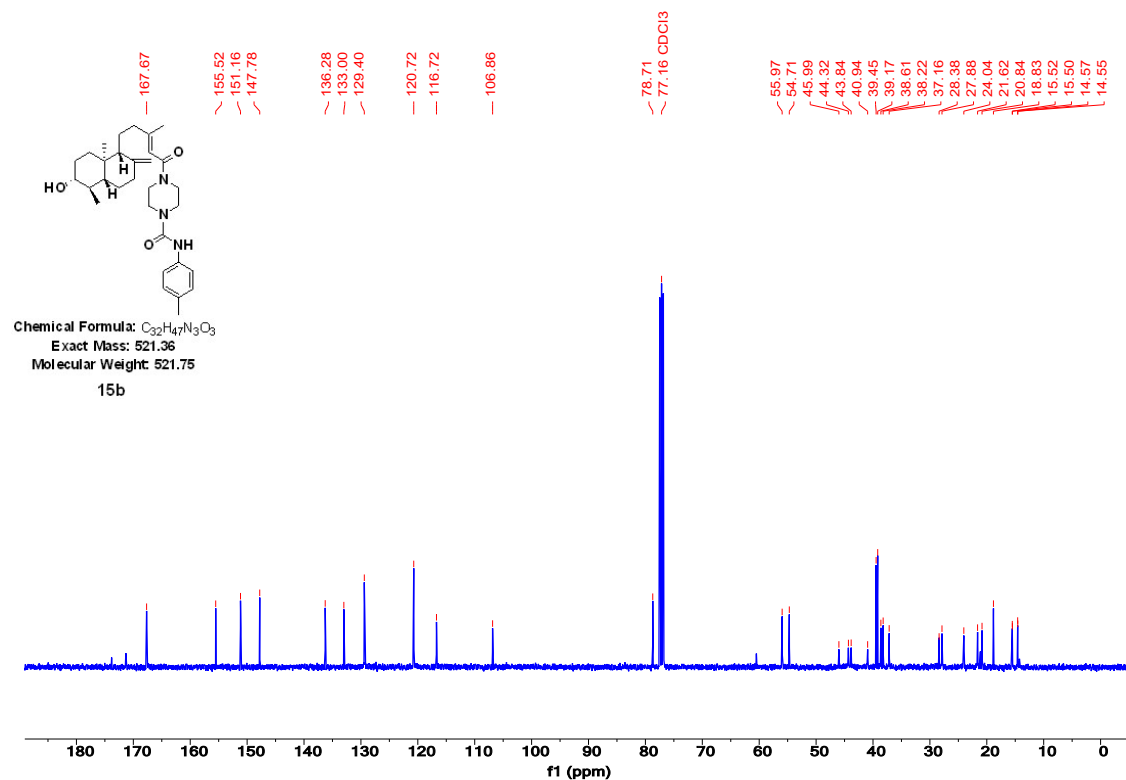


Figure S35. ^{13}C NMR spectrum of 15b

Single Mass Analysis

Tolerance = 5.0 PPM / DBE: min = -1.5, max = 50.0
 Element prediction: Off
 Number of isotope peaks used for i-FIT = 3

Monoisotopic Mass, Even Electron Ions

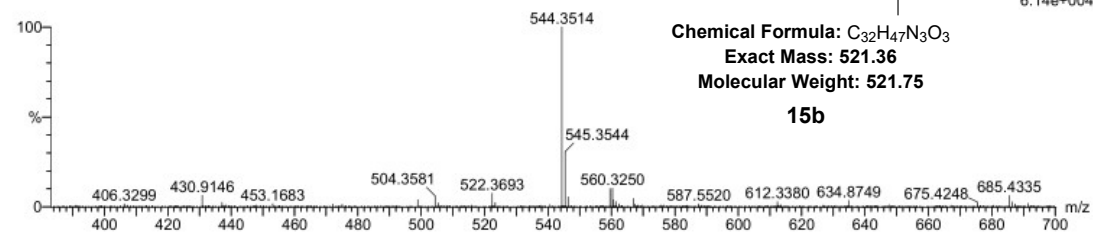
445 formula(e) evaluated with 1 results within limits (all results (up to 1000) for each mass)

Elements Used:

C: 32-32 H: 16-50 N: 1-3 O: 1-20 Na: 1-8

30

250310-2-Q-1 59 (0.137)

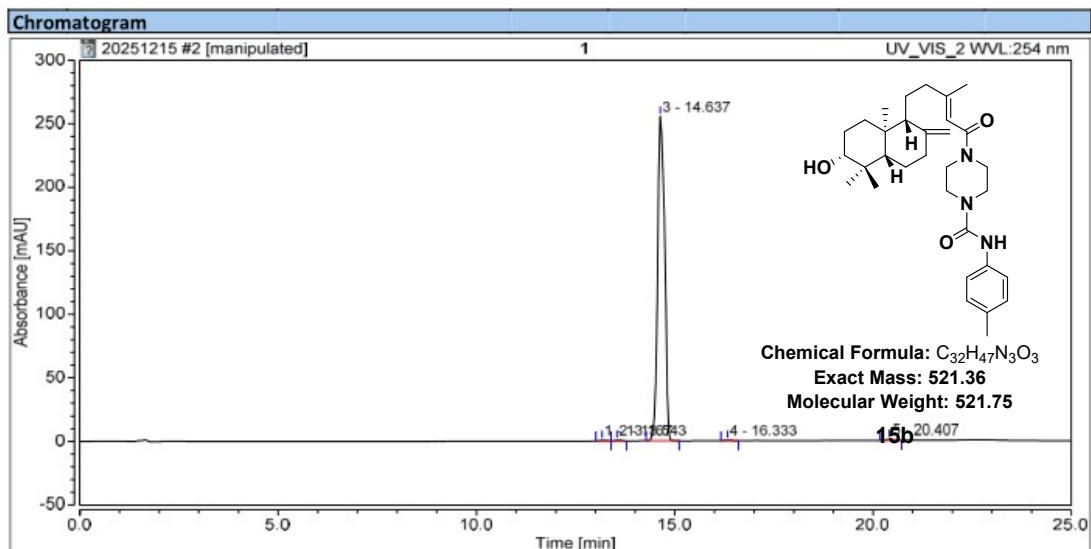


Minimum: -1.5

Maximum: 5.0 5.0 50.0

Mass	Calc. Mass	mDa	PPM	DBE	i-FIT	Norm	Conf(%)	Formula
544.3514	544.3515	-0.1	-0.2	10.5	226.5	n/a	n/a	C ₃₂ H ₄₇ N ₃ O ₃ Na

Figure S36. HRMS spectrum of 15b



Integration Results

No.	Peak Name	Retention Time min	Area mAU*min	Height mAU	Relative Area %	Relative Height %	Amount
1		13.167	0.111	0.557	0.21	0.22	n.a.
2		13.543	0.123	0.636	0.23	0.25	n.a.
3		14.637	52.236	255.633	99.02	98.96	n.a.
4		16.333	0.107	0.533	0.20	0.21	n.a.
5		20.407	0.177	0.964	0.34	0.37	n.a.
Total:			52.754	258.322	100.00	100.00	

Figure S37. HPLC spectrum of 15b

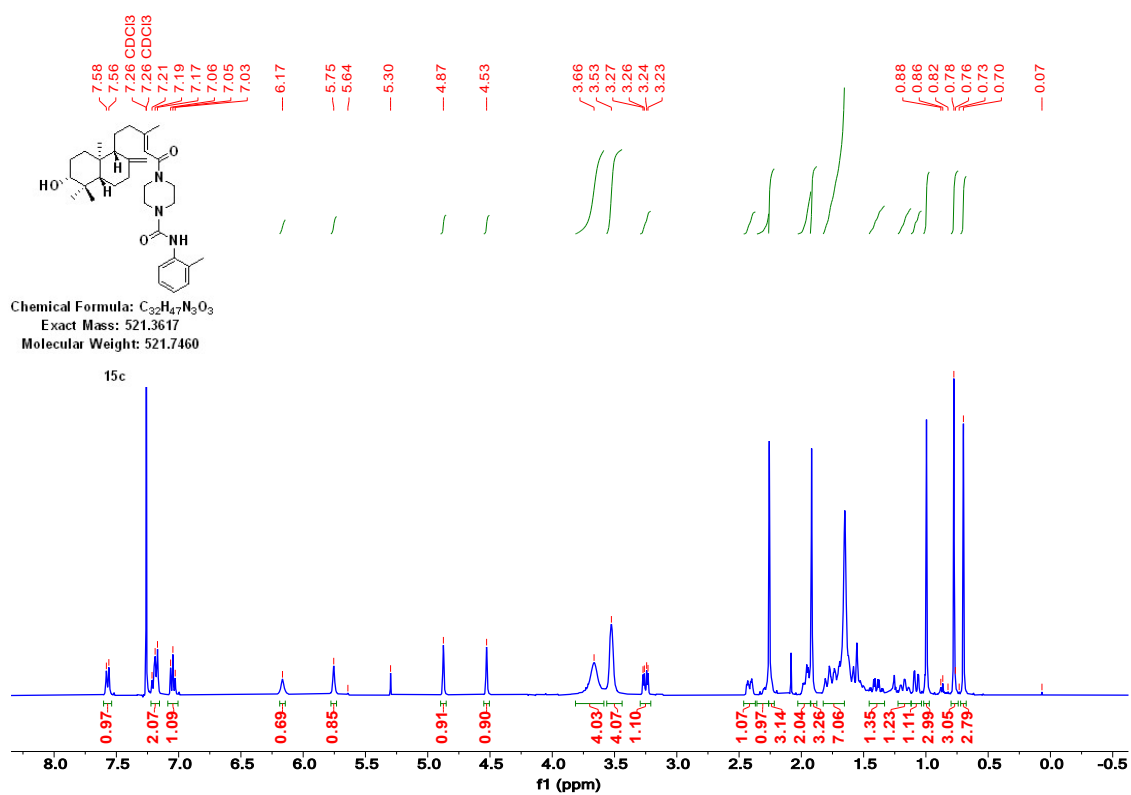


Figure S38. 1H NMR spectrum of 15c

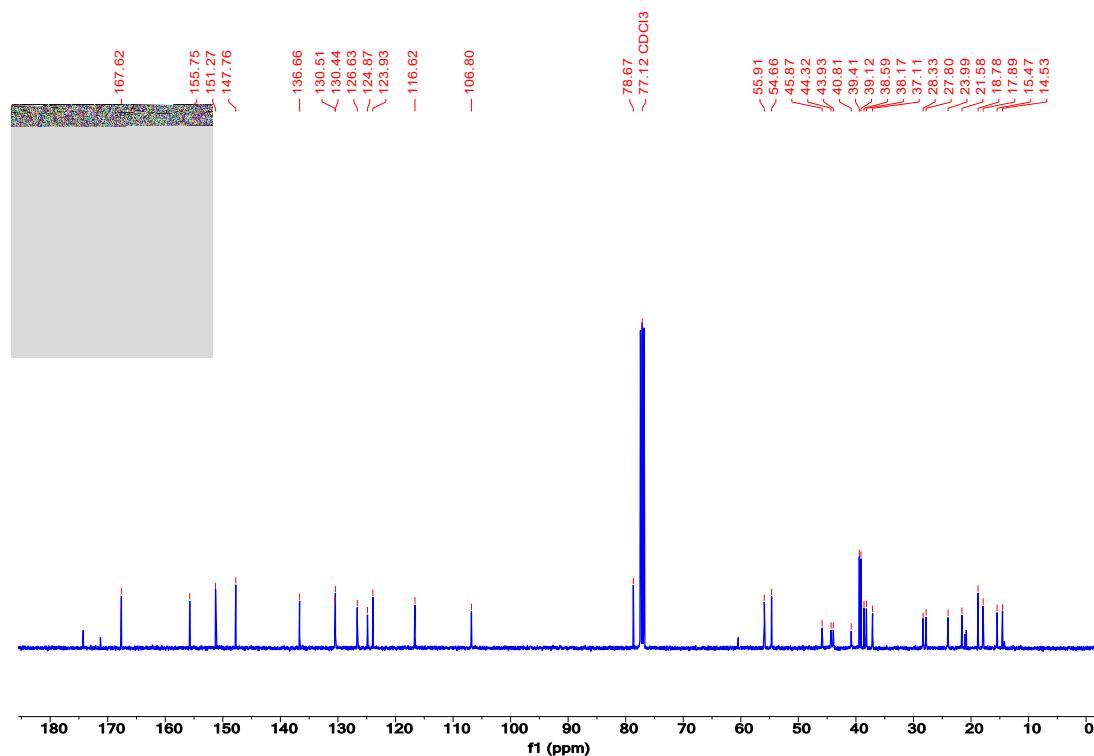


Figure S39. ^{13}C NMR spectrum of 15c

Single Mass Analysis

Tolerance = 3.0 PPM / DBE: min = -1.5, max = 50.0

Element prediction: Off

Number of isotope peaks used for i-FIT = 3

Monoisotopic Mass, Even Electron Ions

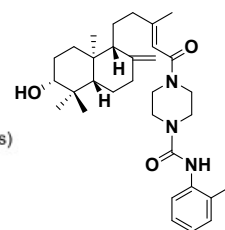
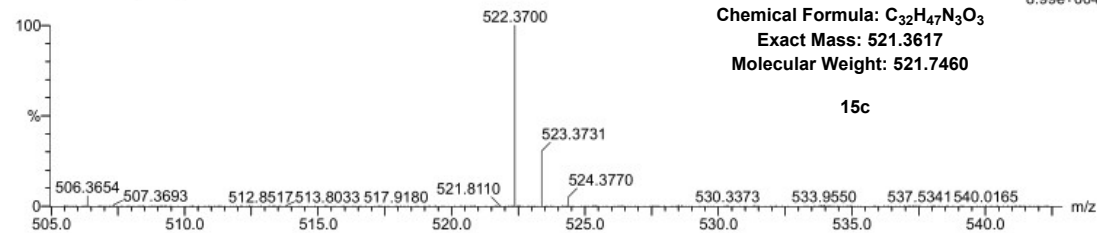
4 formula(e) evaluated with 1 results within limits (up to 50 best isotopic matches for each mass)

Elements Used:

C: 30-34 H: 44-50 N: 3-5 O: 3-6

30

250310-2-Q-2 20 (0.063)



Chemical Formula: $\text{C}_{32}\text{H}_{47}\text{N}_3\text{O}_3$

Exact Mass: 521.3617

Molecular Weight: 521.7460

15c

1: TOF MS ES+
8.99e+004

Minimum: -1.5
Maximum: 50.0

Mass	Calc. Mass	mDa	PPM	DBE	i-FIT	Norm	Conf (%)	Formula
522.3700	522.3696	0.4	0.8	10.5	238.4	n/a	n/a	C32 H48 N3 O3

Figure S40. HRMS spectrum of 15c

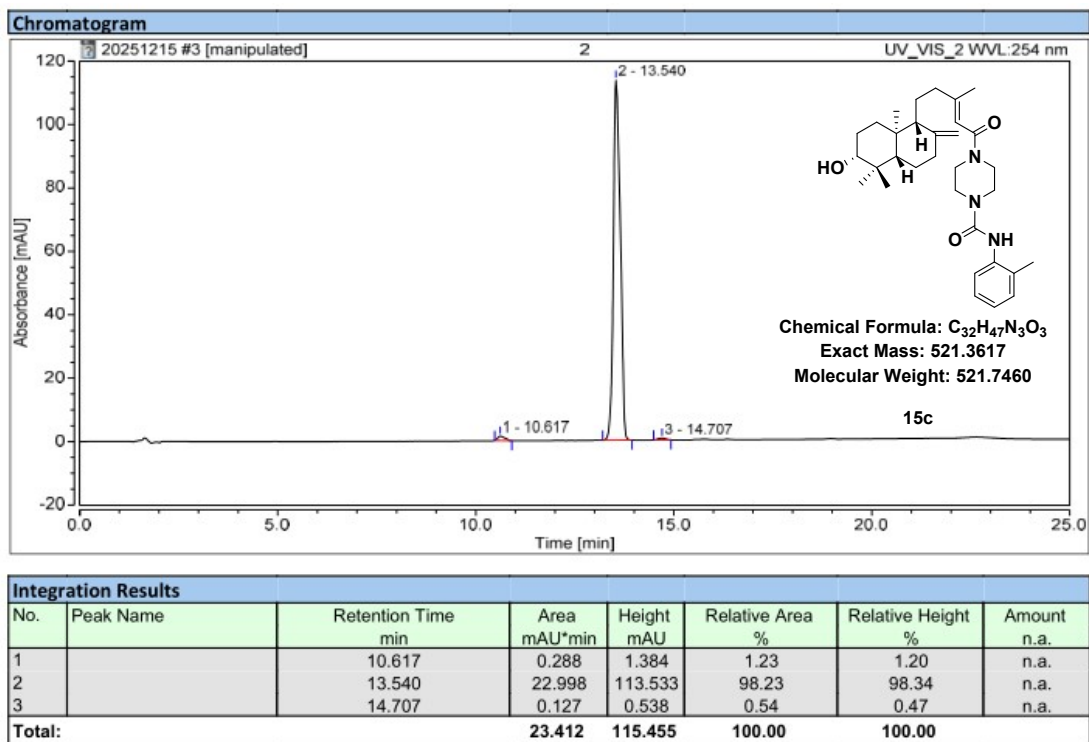


Figure S41. HPLC spectrum of 15c

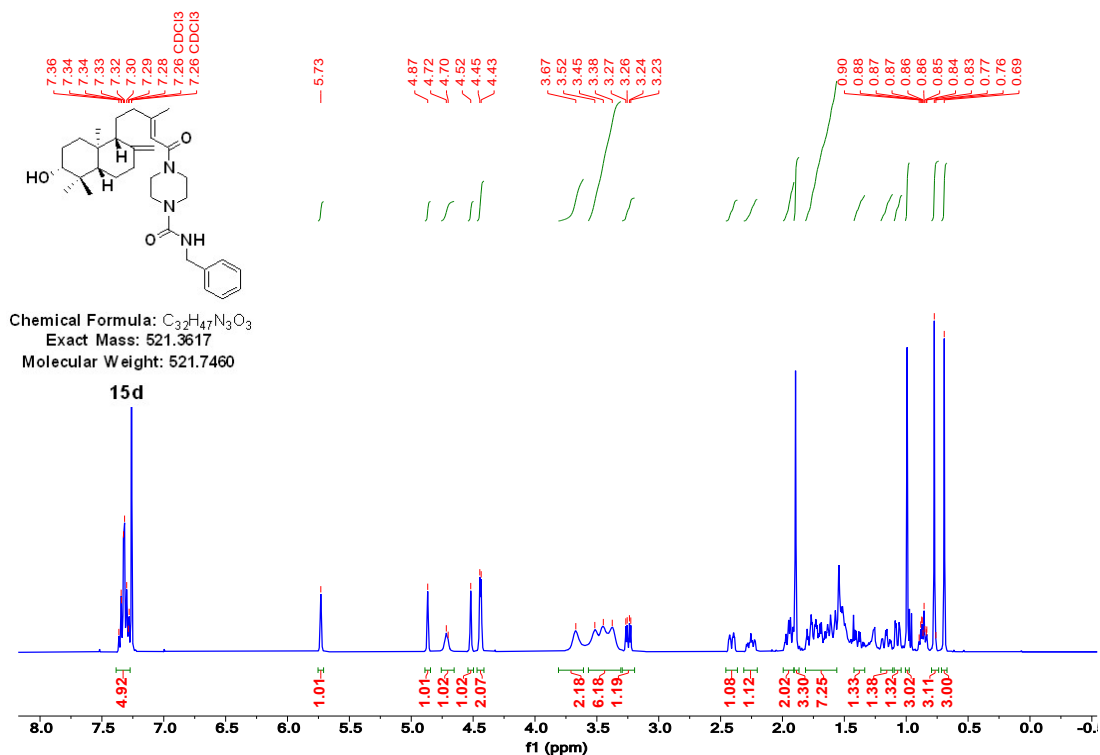


Figure S42. 1H NMR spectrum of 15d

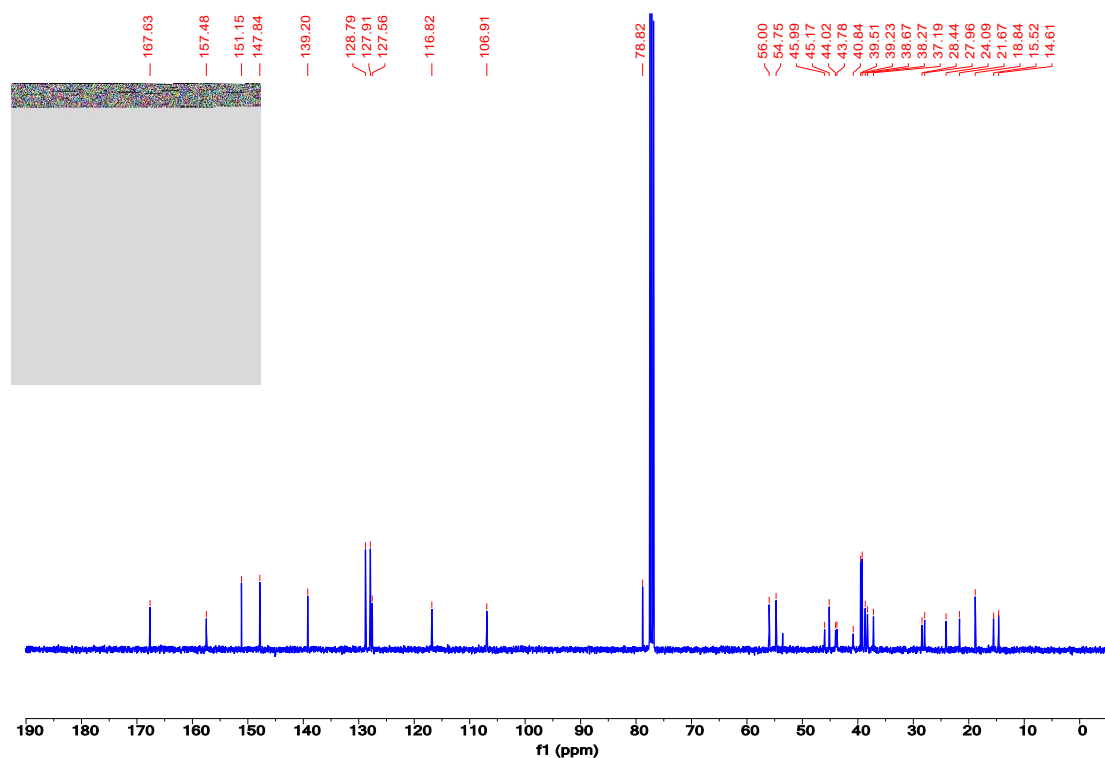


Figure S43. ^{13}C NMR spectrum of 15d

Single Mass Analysis

Tolerance = 5.0 mDa / DBE: min = -1.5, max = 50.0

Element prediction: Off

Number of isotope peaks used for i-FIT = 3

Monoisotopic Mass, Even Electron Ions

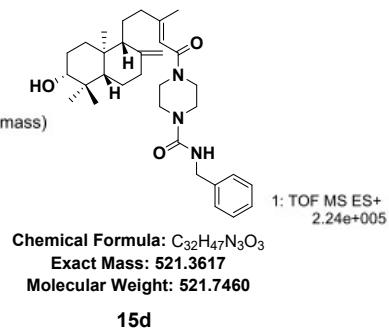
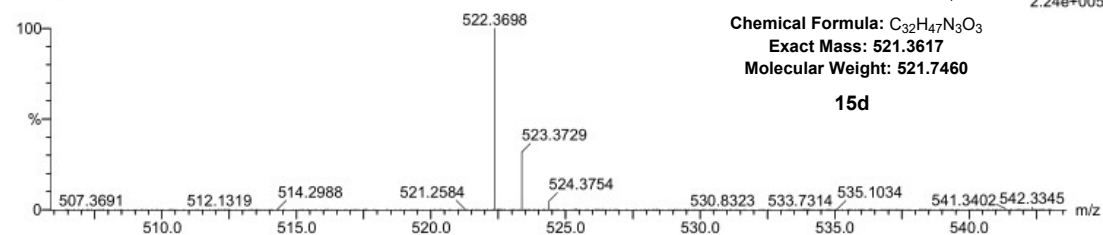
33 formula(e) evaluated with 1 results within limits (up to 50 best isotopic matches for each mass)

Elements Used:

C: 29-56 H: 47-85 N: 2-6 O: 2-6

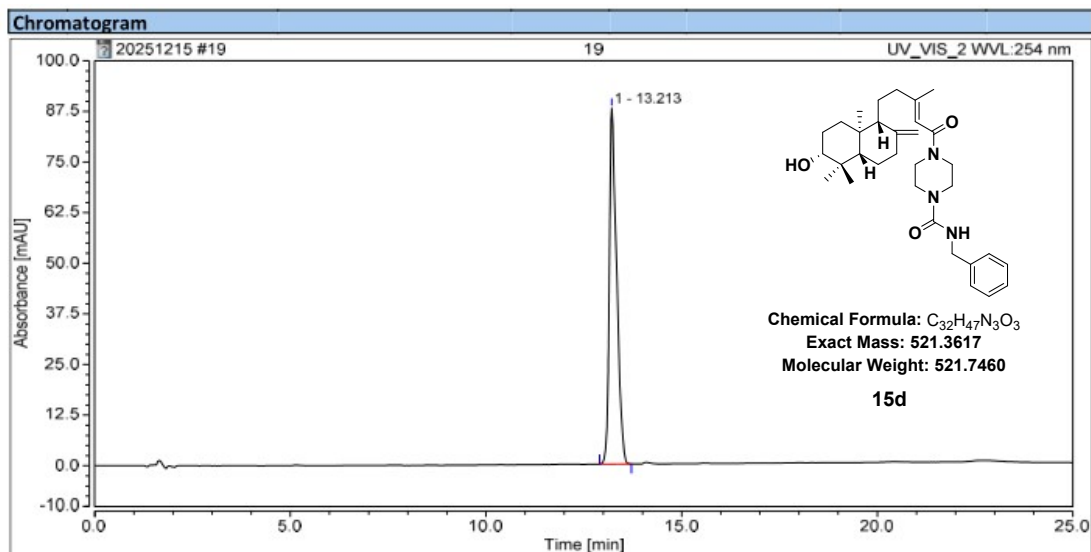
7

250310-2-Q-19 23 (0.069)



Mass	Calc. Mass	mDa	PPM	DBE	i-FIT	Norm	Conf (%)	Formula
522.3698	522.3696	0.2	0.4	10.5	437.7	n/a	n/a	$\text{C}_{32}\text{H}_{48}\text{N}_3\text{O}_3$

Figure S44. HRMS spectrum of 15d



Integration Results

No.	Peak Name	Retention Time min	Area mAU*min	Height mAU	Relative Area %	Relative Height %	Amount n.a.
1		13.213	19.026	87.926	100.00	100.00	n.a.
Total:			19.026	87.926	100.00	100.00	

Figure S45. HPLC spectrum of 15d

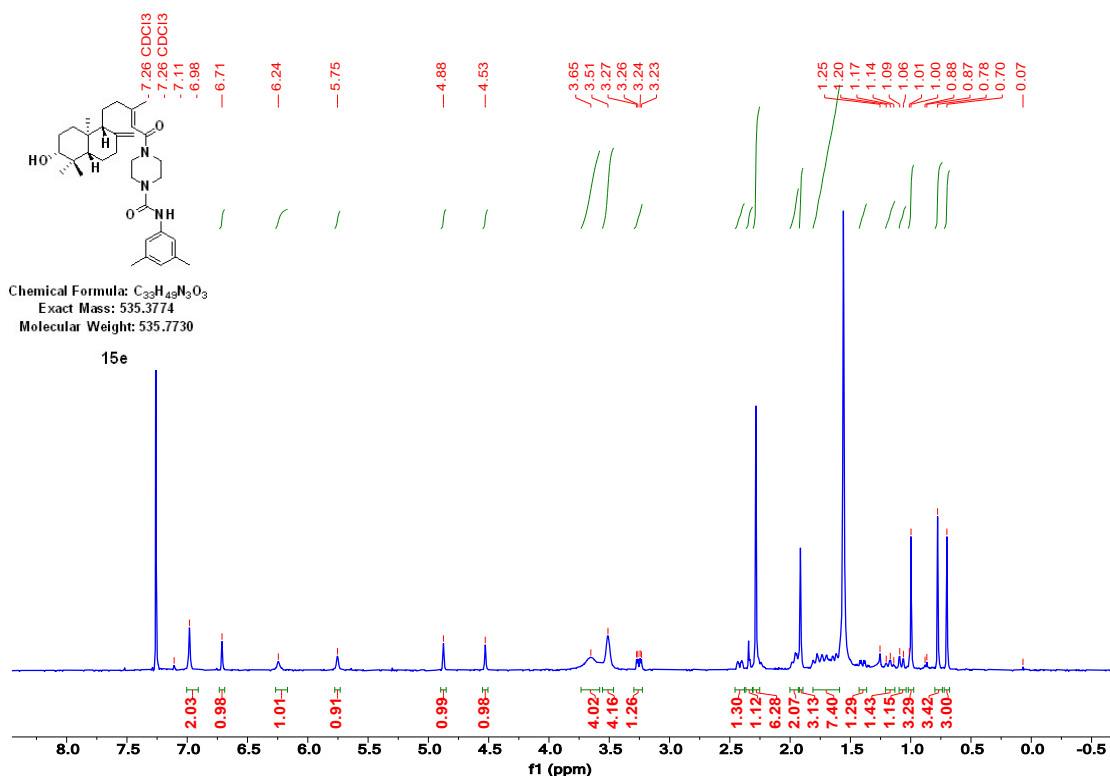


Figure S46. 1H NMR spectrum of 15e

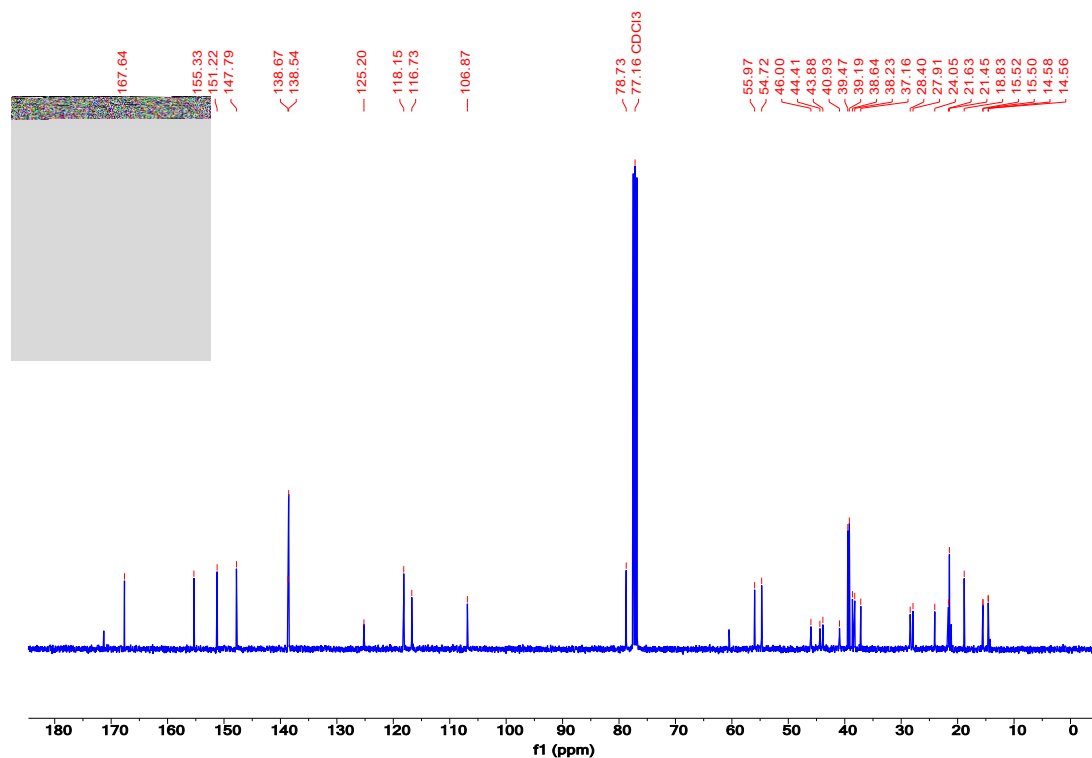


Figure S47. ^{13}C NMR spectrum of 15e

Single Mass Analysis

Tolerance = 3.0 PPM / DBE: min = -1.5, max = 50.0

Element prediction: Off

Number of isotope peaks used for i-FIT = 3

Monoisotopic Mass, Even Electron Ions

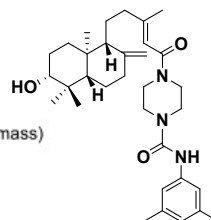
5 formula(e) evaluated with 1 results within limits (up to 50 best isotopic matches for each mass)

Elements Used:

C: 30-34 H: 44-50 N: 3-5 O: 3-6

30

250310-2-Q-3 22 (0.067)



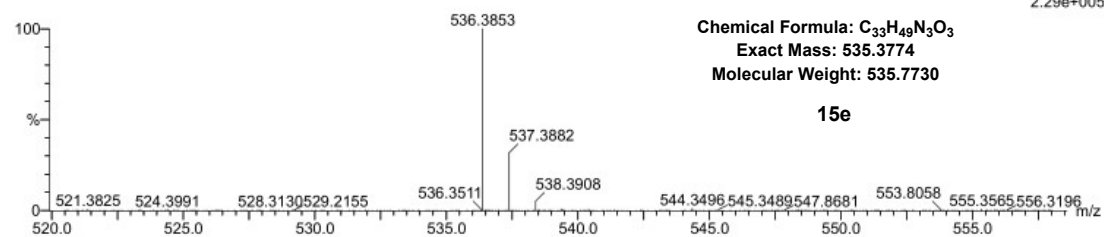
1: TOF MS ES+
2.29e+005

Chemical Formula: $\text{C}_{33}\text{H}_{49}\text{N}_3\text{O}_3$

Exact Mass: 535.3774

Molecular Weight: 535.7730

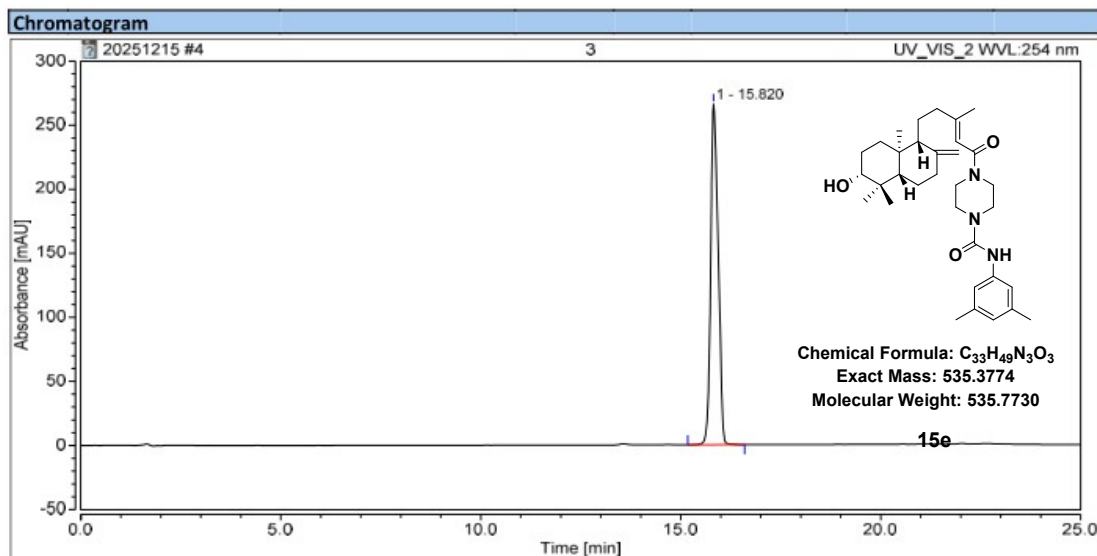
15e



Minimum: -1.5
Maximum: 5.0 3.0 50.0

Mass	Calc. Mass	mDa	PPM	DBE	i-FIT	Norm	Conf (%)	Formula
536.3853	536.3852	0.1	0.2	10.5	421.5	n/a	n/a	$\text{C}_{33}\text{H}_{50}\text{N}_3\text{O}_3$

Figure S48. HRMS spectrum of 15e



Integration Results

No.	Peak Name	Retention Time min	Area mAU*min	Height mAU	Relative Area %	Relative Height %	Amount
1		15.820	57.987	266.136	100.00	100.00	n.a.
Total:			57.987	266.136	100.00	100.00	

Figure S49. HPLC spectrum of 15e

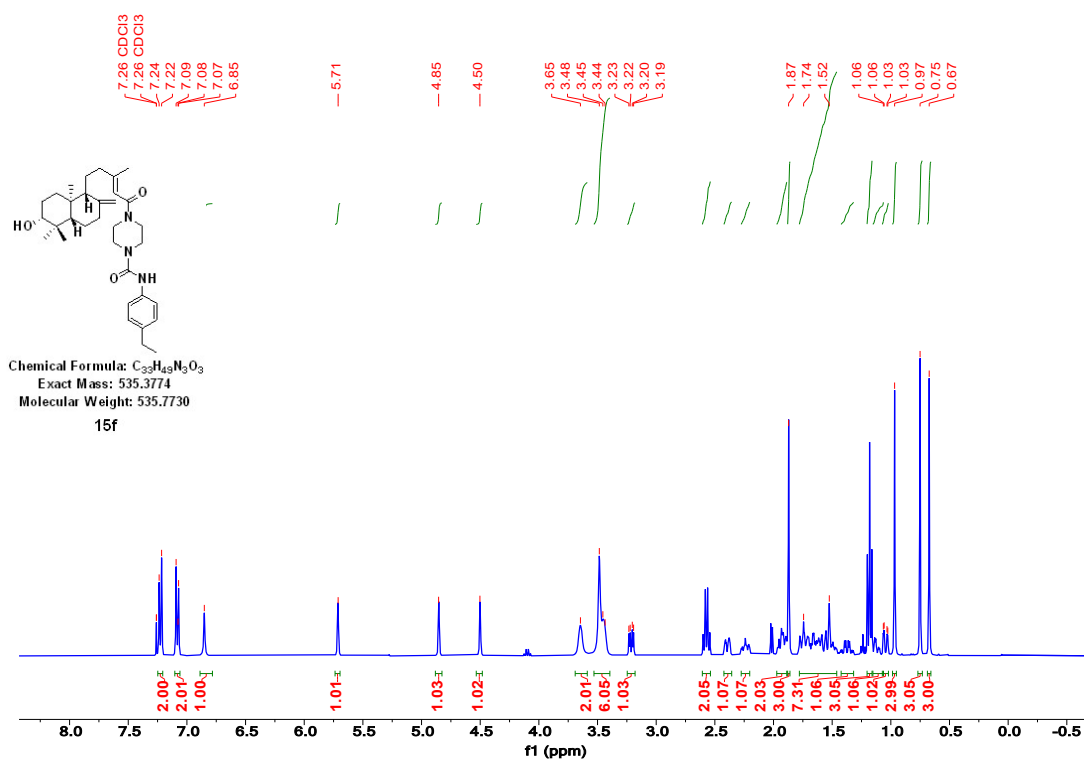


Figure S50. 1H NMR spectrum of 15f

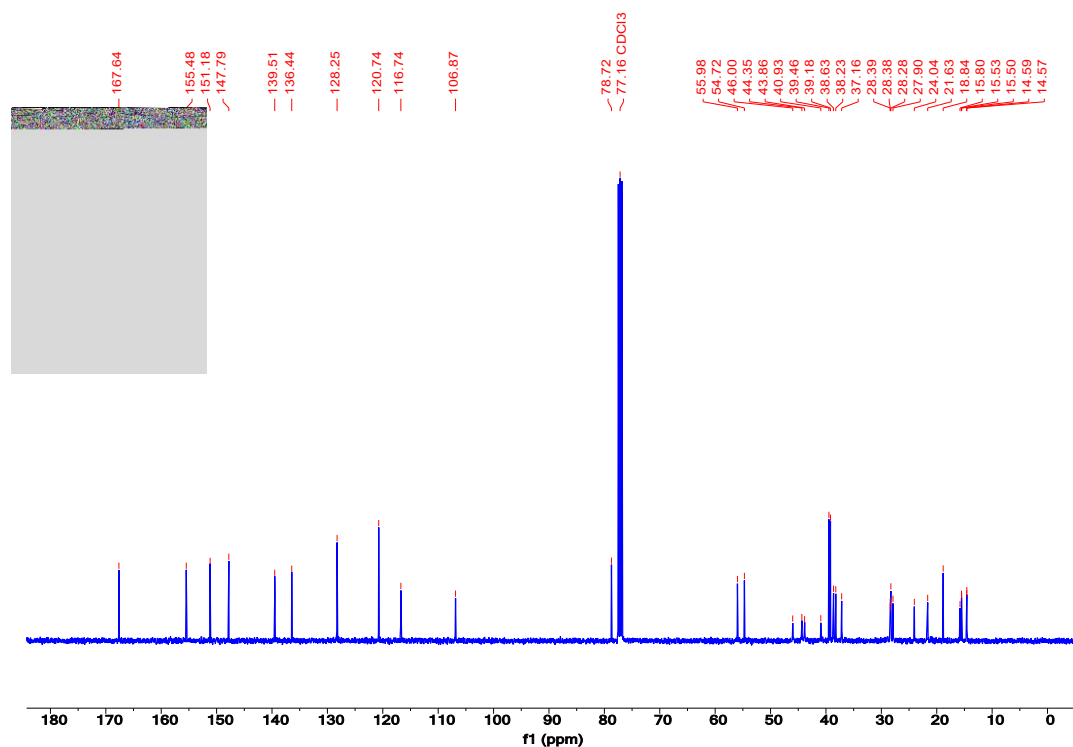


Figure S51. ^{13}C NMR spectrum of 15f

Single Mass Analysis

Tolerance = 3.0 PPM / DBE: min = -1.5, max = 50.0

Element prediction: Off

Number of isotope peaks used for i-FIT = 3

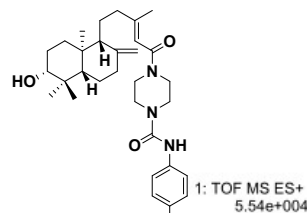
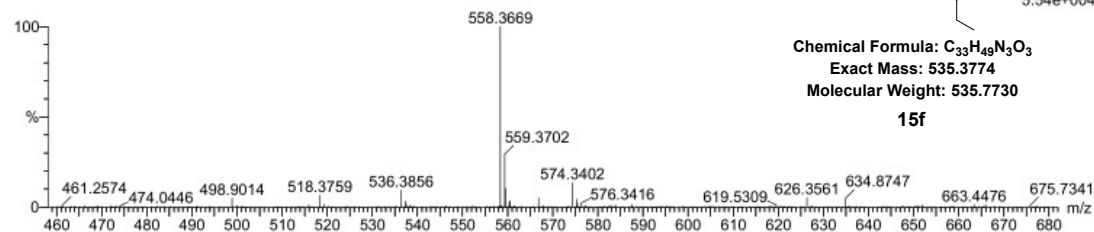
Monoisotopic Mass, Even Electron Ions

7 formula(e) evaluated with 1 results within limits (up to 50 best isotopic matches for each mass)

Elements Used:

C: 33-34 H: 44-50 N: 3-5 O: 3-6 Na: 0-1

30
250310-2-Q-4 18 (0.059)



Chemical Formula: $\text{C}_{33}\text{H}_{49}\text{N}_3\text{O}_3$

Exact Mass: 535.3774

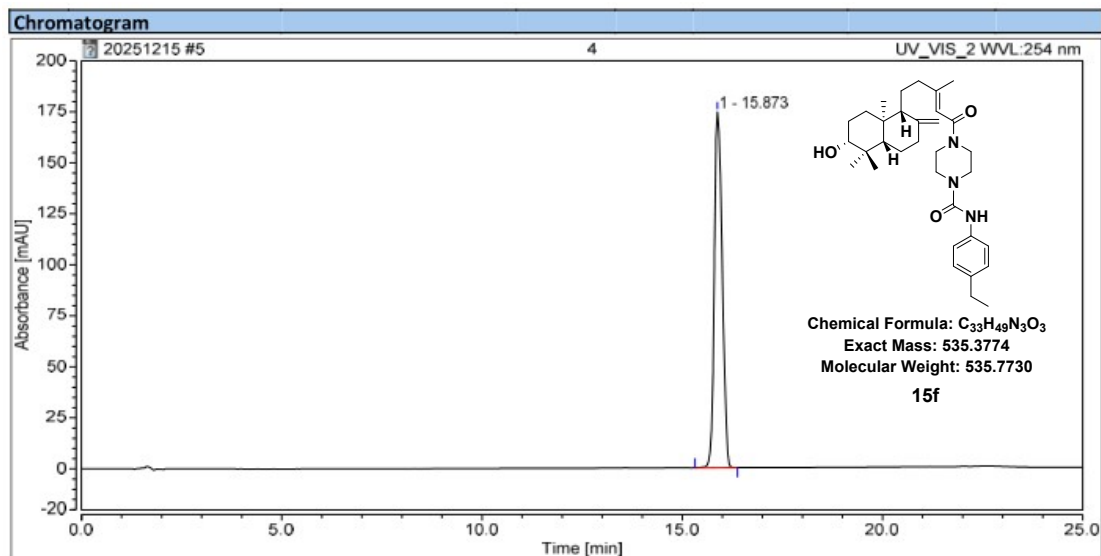
Molecular Weight: 535.7730

15f

Minimum: -1.5
Maximum: 50.0

Mass	Calc. Mass	mDa	PPM	DBE	i-FIT	Norm	Conf (%)	Formula
558.3669	558.3672	-0.3	-0.5	10.5	253.9	n/a	n/a	$\text{C}_{33}\text{H}_{49}\text{N}_3\text{O}_3$ Na

Figure S52. HRMS spectrum of 15f



Integration Results

No.	Peak Name	Retention Time min	Area mAU*min	Height mAU	Relative Area %	Relative Height %	Amount n.a.
1		15.873	40.006	174.420	100.00	100.00	n.a.
Total:			40.006	174.420	100.00	100.00	

Figure S53. HPLC spectrum of 15f

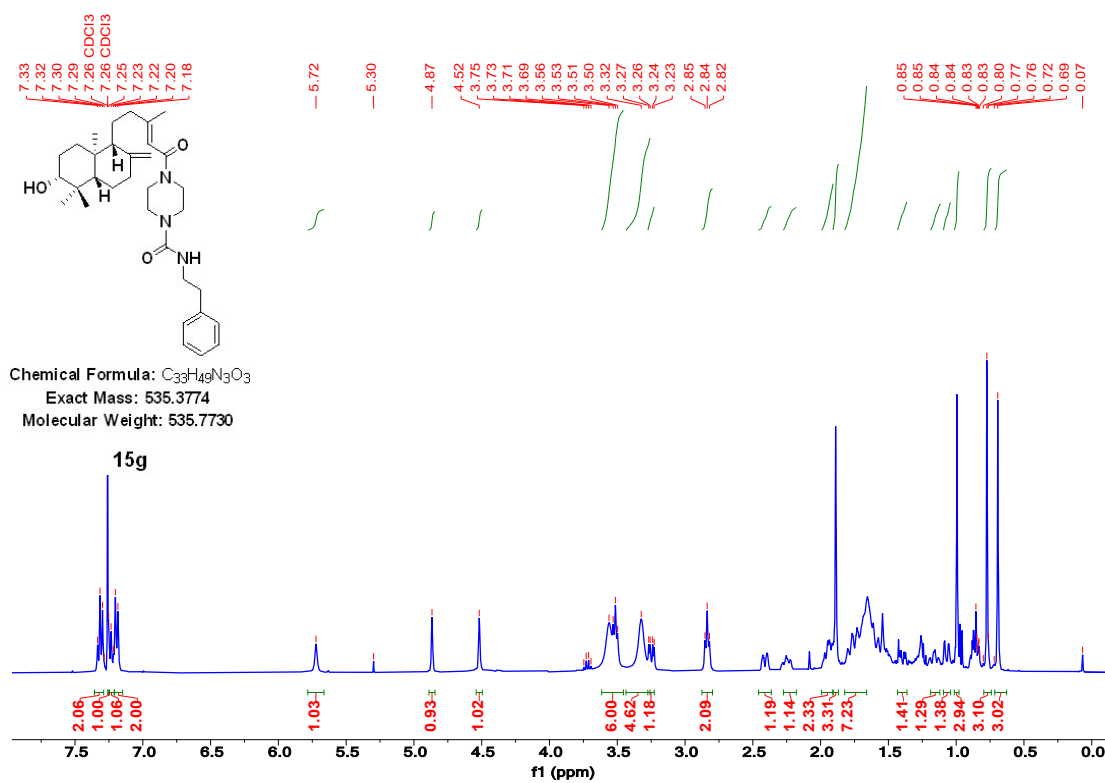


Figure S54. 1H NMR spectrum of 15g

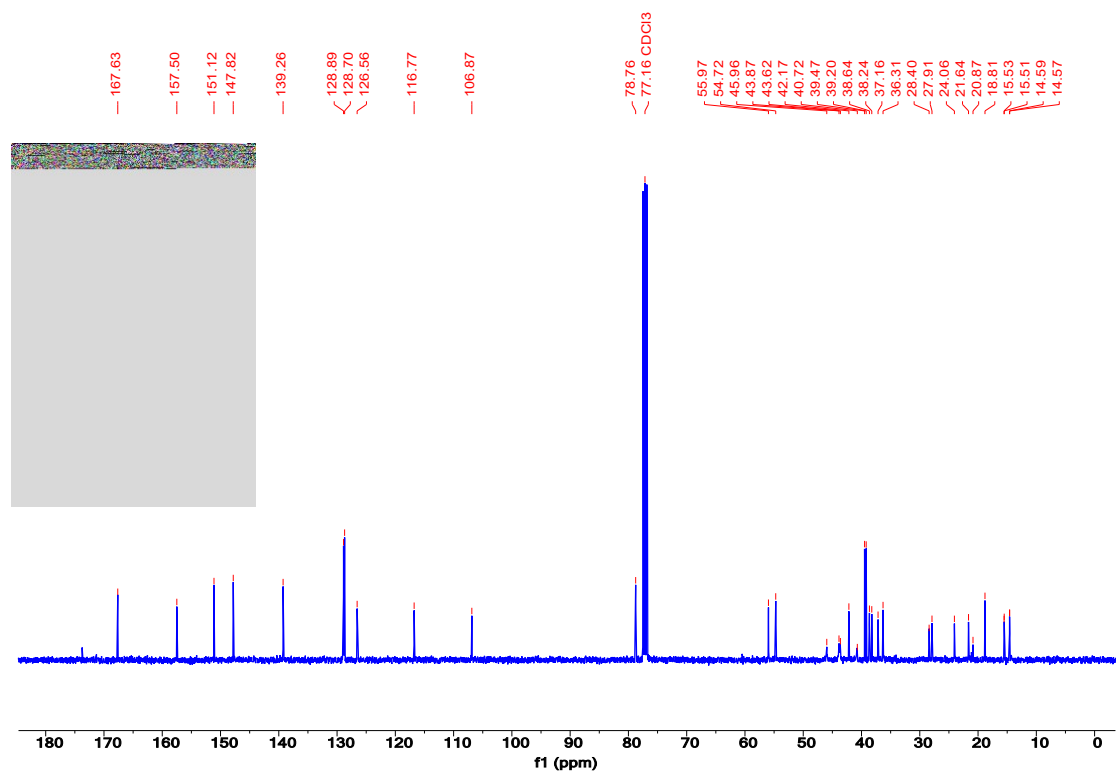


Figure S55. ^{13}C NMR spectrum of 15g

Single Mass Analysis

Tolerance = 5.0 mDa / DBE: min = -1.5, max = 50.0

Element prediction: Off

Number of isotopic peaks used for i-FIT = 3

Monoisotopic Mass, Even Electron Ions

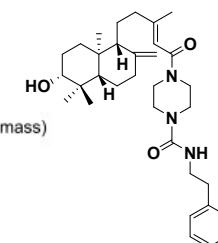
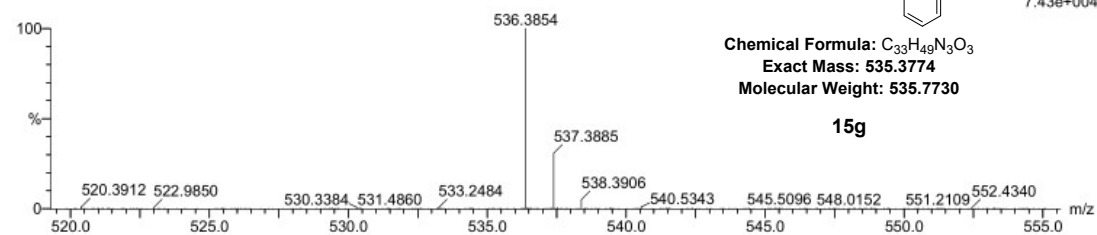
14 formula(e) evaluated with 1 results within limits (up to 50 best isotopic matches for each mass)

Elements Used:

C: 30-35 H: 44-51 N: 2-5 O: 2-5 K: 0-1

30

250310-2-Q-6 46 (0.113)



1: TOF MS ES+
7.43e+004

Chemical Formula: $\text{C}_{33}\text{H}_{48}\text{N}_3\text{O}_3$

Exact Mass: 535.3774

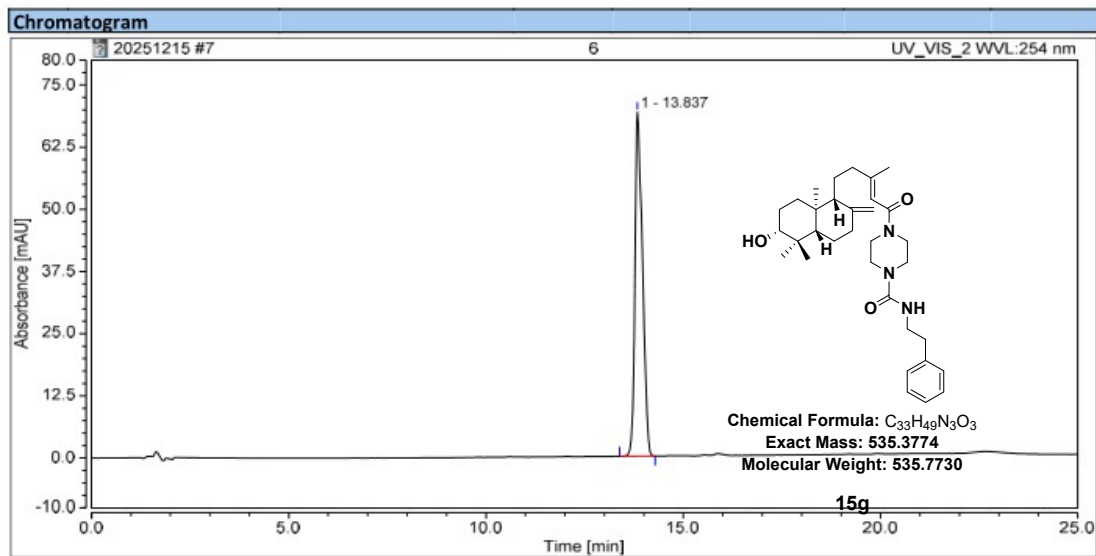
Molecular Weight: 535.7730

15g

Minimum: -1.5
Maximum: 50.0

Mass	Calc. Mass	mDa	PPM	DBE	i-FIT	Norm	Conf (%)	Formula
536.3854	536.3852	0.2	0.4	10.5	296.0	n/a	n/a	C33 H50 N3 O3

Figure S56. HRMS spectrum of 15g



Integration Results

No.	Peak Name	Retention Time min	Area mAU*min	Height mAU	Relative Area %	Relative Height %	Amount
1		13.837	15.263	69.196	100.00	100.00	n.a.
Total:			15.263	69.196	100.00	100.00	

Figure S57. HPLC spectrum of 15g

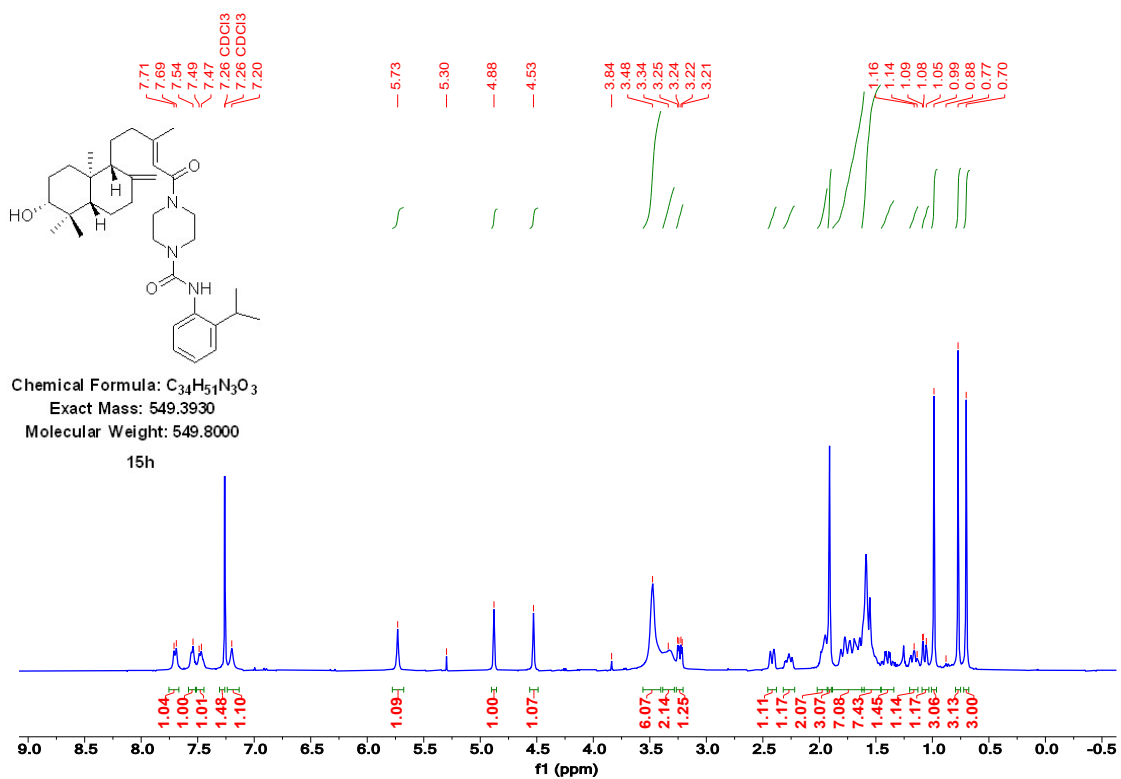


Figure S58. 1H NMR spectrum of 15h

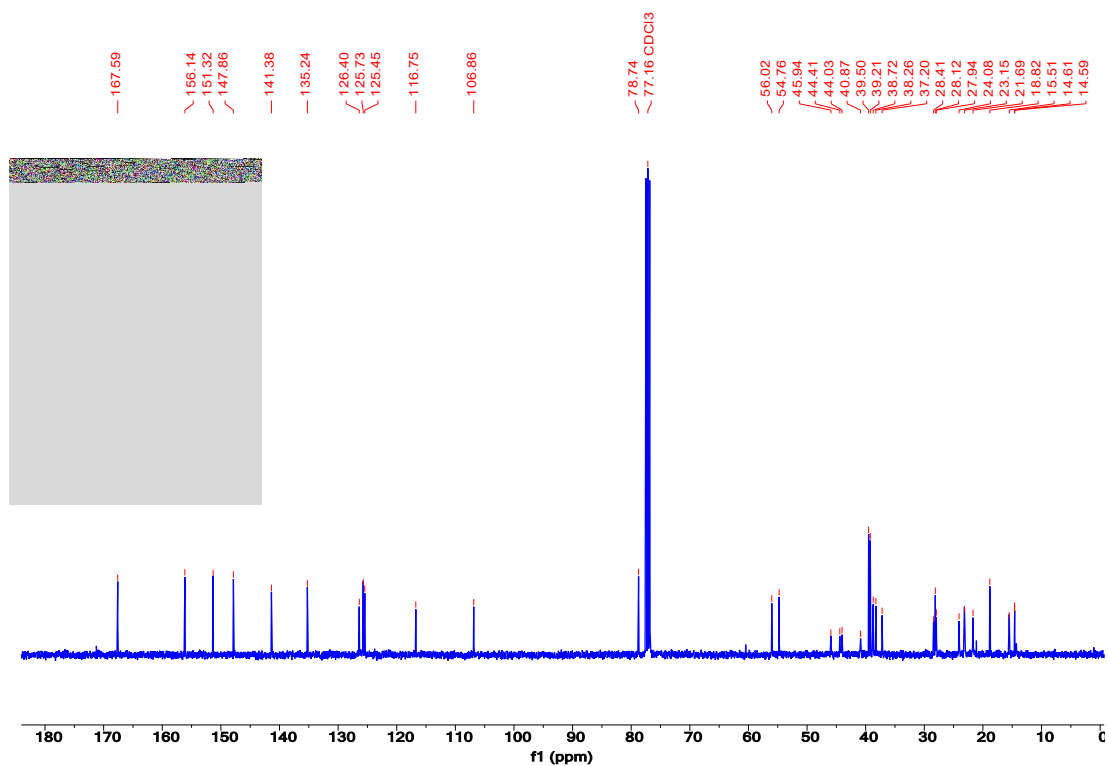


Figure S59. ^{13}C NMR spectrum of 15h

Single Mass Analysis

Tolerance = 5.0 mDa / DBE: min = -1.5, max = 50.0

Element prediction: Off

Number of isotope peaks used for i-FIT = 3

Monoisotopic Mass, Even Electron Ions

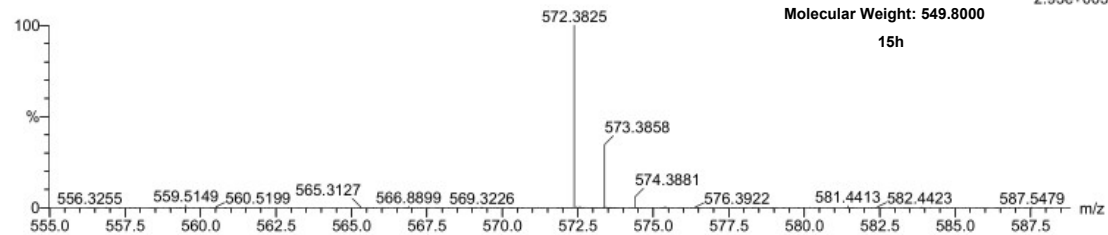
13 formula(e) evaluated with 1 results within limits (up to 50 best isotopic matches for each mass)

Elements Used:

C: 34-34 H: 42-51 N: 2-6 O: 2-6 Na: 0-1

7

250310-2-Q-36 19 (0.061)

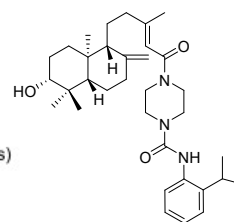


Minimum:

Maximum:

5.0 10.0 -1.5
-0.3 -0.5 10.5

Mass	Calc. Mass	mDa	PPM	DBE	i-FIT	Norm	Conf (%)	Formula
572.3825	572.3828	-0.3	-0.5	10.5	475.6	n/a	n/a	C34 H51 N3 O3 Na



Chemical Formula: $\text{C}_{34}\text{H}_{51}\text{N}_3\text{O}_3$; TOF MS ES+
Exact Mass: 549.3930 2.93e+005
Molecular Weight: 549.8000
15h

Figure S60. HRMS spectrum of 15h

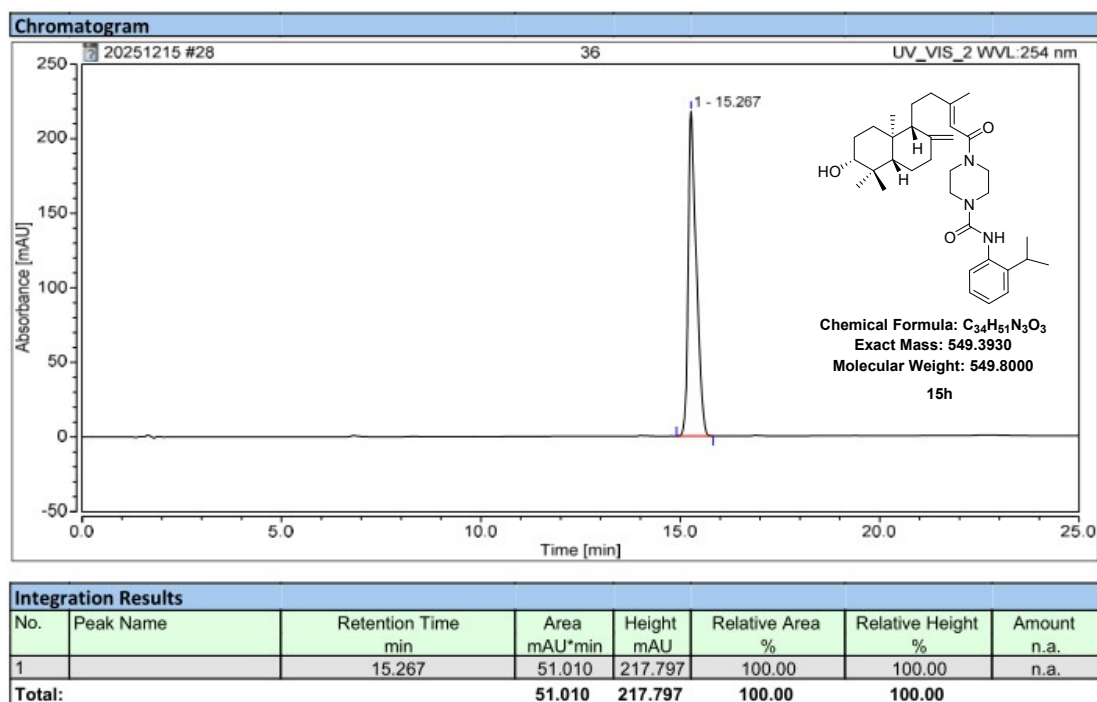


Figure S61. HPLC spectrum of 15h

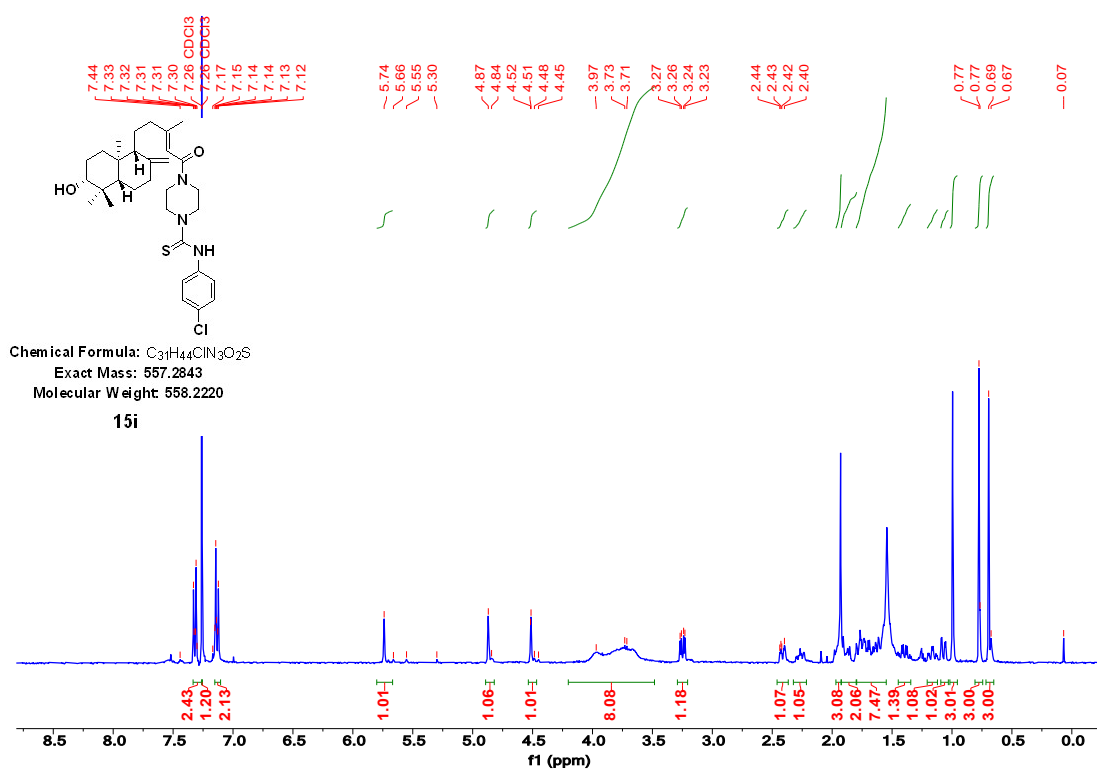


Figure S62. 1H NMR spectrum of 15i

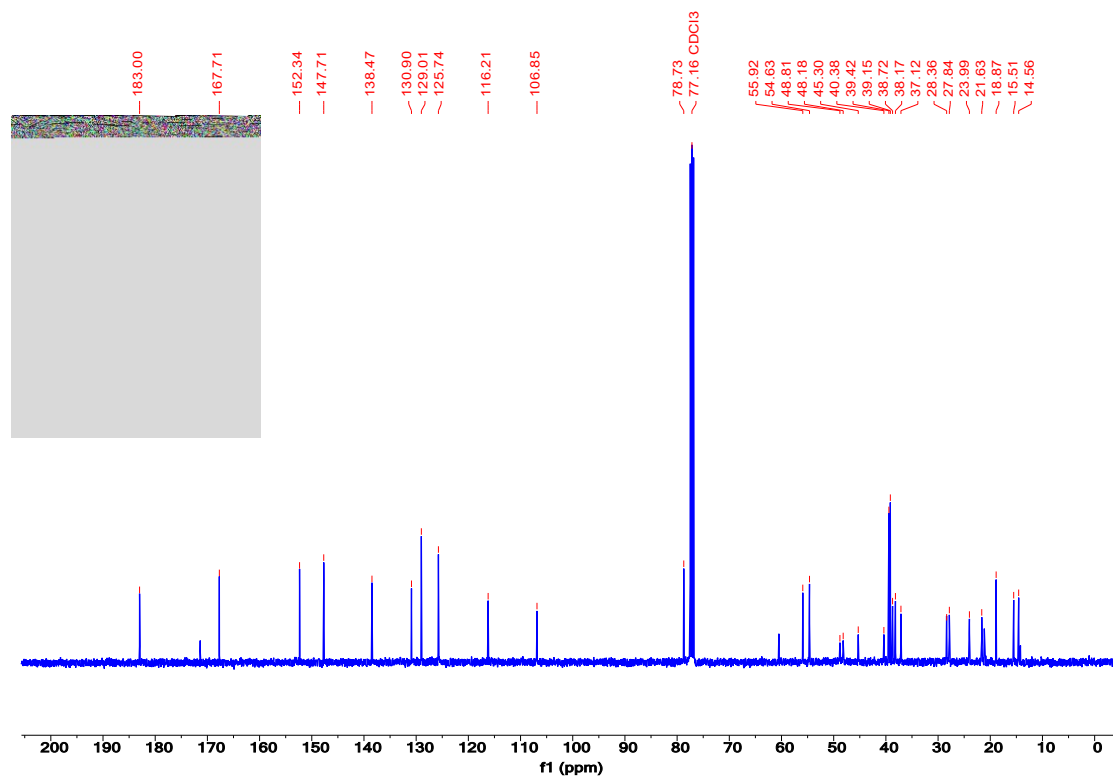


Figure S63. ¹³C NMR spectrum of 15i

Monoisotopic Mass, Even Electron Ions

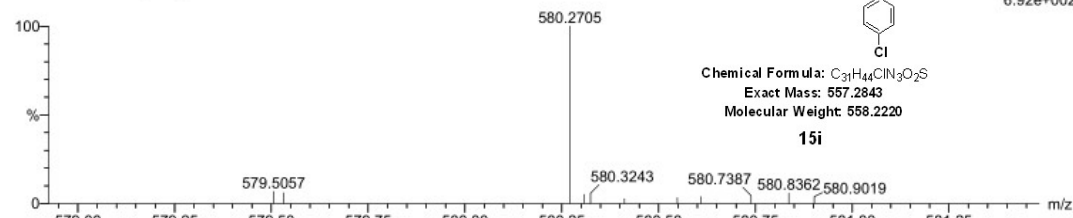
573 formula(e) evaluated with 1 results within limits (all results (up to 1000) for each mass)

Elements Used:

C: 31-31 H: 14-90 N: 1-6 O: 1-20 Cl: 1-5 K: 1-1

30

250310-2-Q-10 72 (0.162)



Chemical Formula: C₃₁H₄₄ClN₃O₂S

Exact Mass: 557.2843

Molecular Weight: 558.2220

15i

1: TOF MS ES+
6.92e+002

Minimum: -1.5
Maximum: 5.0 5.0 50.0

Mass	Calc. Mass	mDa	PPM	DBE	i-FIT	Norm	Conf(%)	Formula
580.2705	580.2708	-0.3	-0.5	10.5	78.0	n/a	n/a	C ₃₁ H ₄₄ N ₃ O ₃ Cl K

Figure S64. HRMS spectrum of 15i

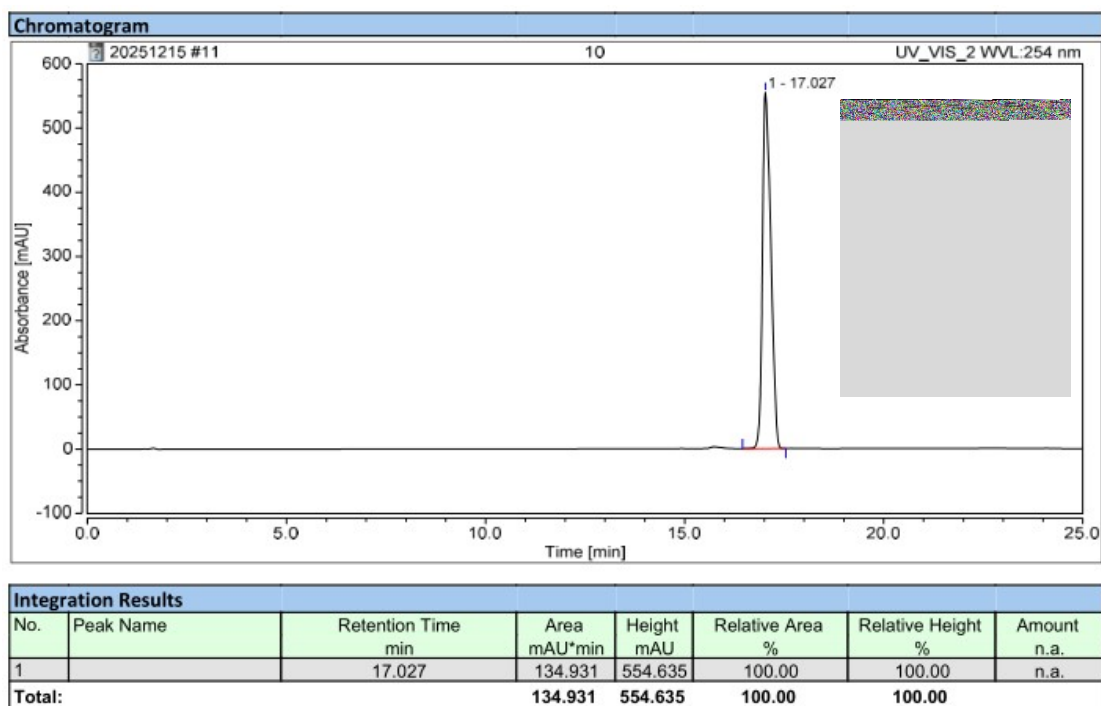
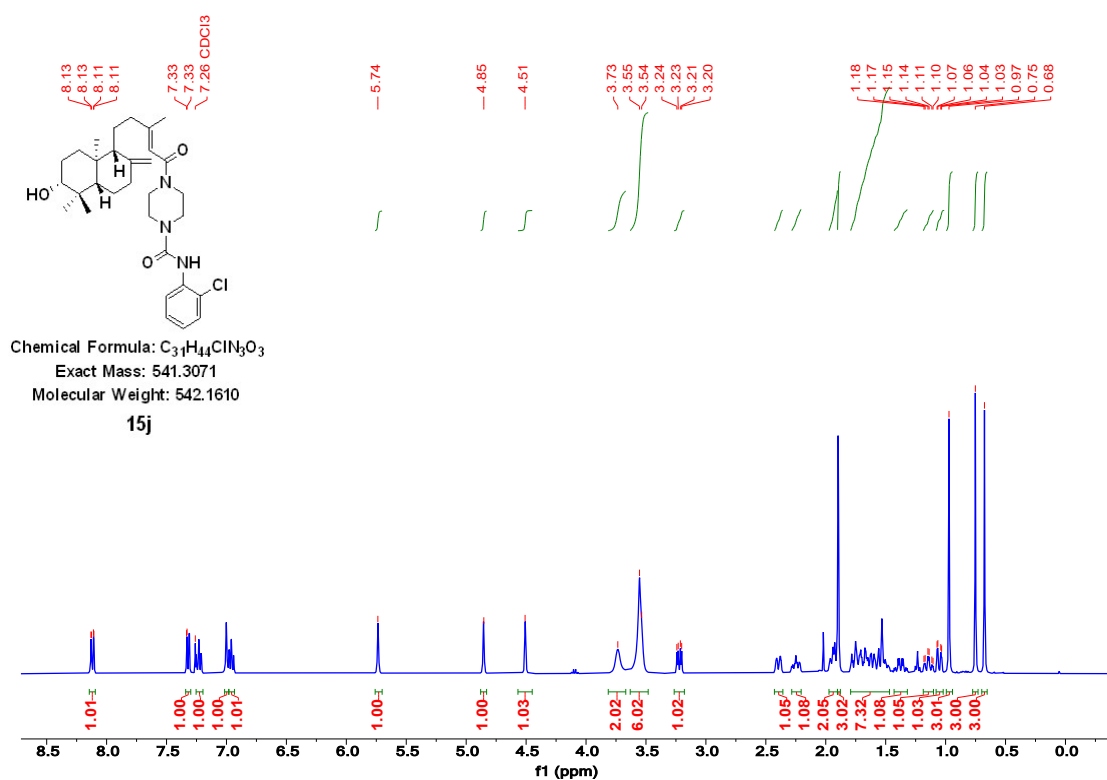


Figure S65. HPLC spectrum of 15i



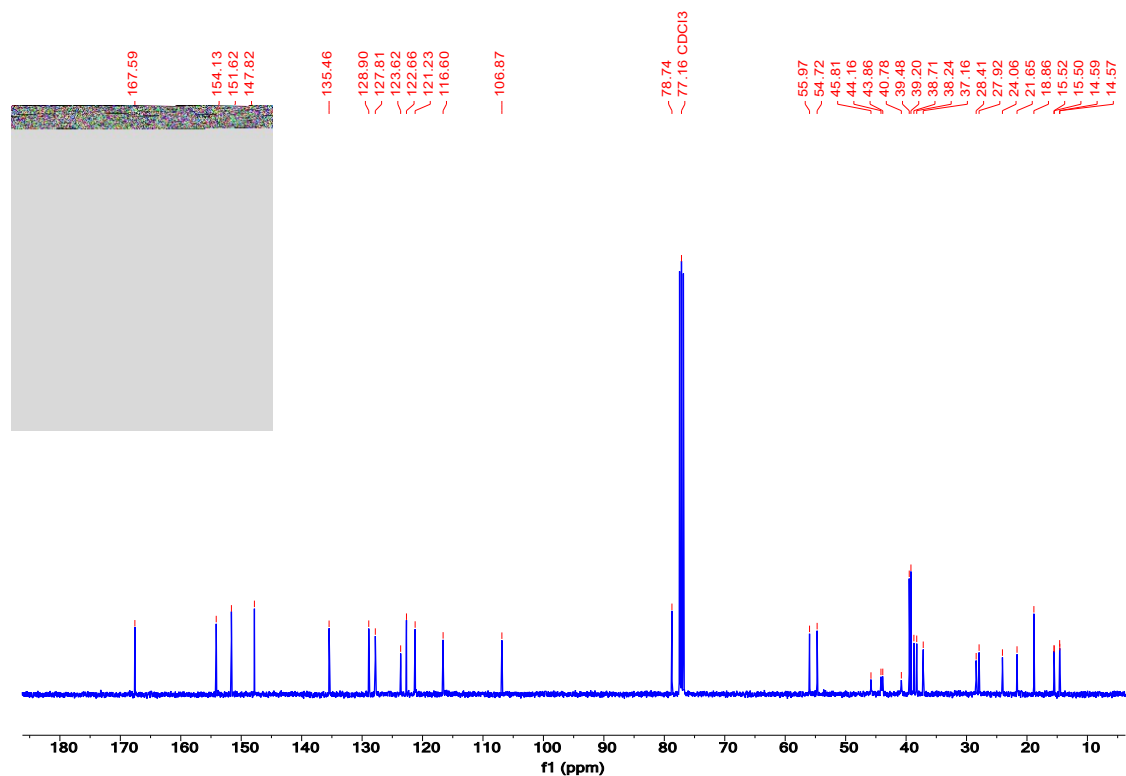


Figure S67. ¹³C NMR spectrum of 15j

Single Mass Analysis

Tolerance = 5.0 mDa / DBE: min = -1.5, max = 50.0

Element prediction: Off

Number of isotope peaks used for i-FIT = 3

Monoisotopic Mass, Even Electron Ions

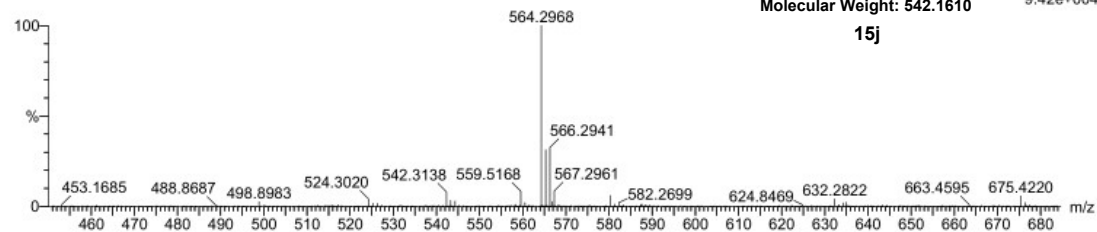
32 formula(e) evaluated with 1 results within limits (up to 50 best isotopic matches for each mass)

Elements Used:

C: 30-35 H: 44-51 N: 2-5 O: 2-5 Cl: 0-1 Na: 0-1

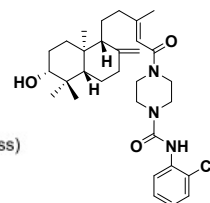
30

250310-2-Q-7 53 (0.126)



Minimum: -1.5
Maximum: 5.0 10.0 50.0

Mass	Calc. Mass	mDa	PPM	DBE	i-FIT	Norm	Conf(%)	Formula
564.2968	564.2969	-0.1	-0.2	10.5	334.7	n/a	n/a	C31 H44 N3 O3 Cl Na



Chemical Formula: C₃₁H₄₄ClN₃O₃
Exact Mass: 541.3071 1: TOF MS ES+
Molecular Weight: 542.1610 9.42e+004
15j

Figure S68. HRMS spectrum of 15j

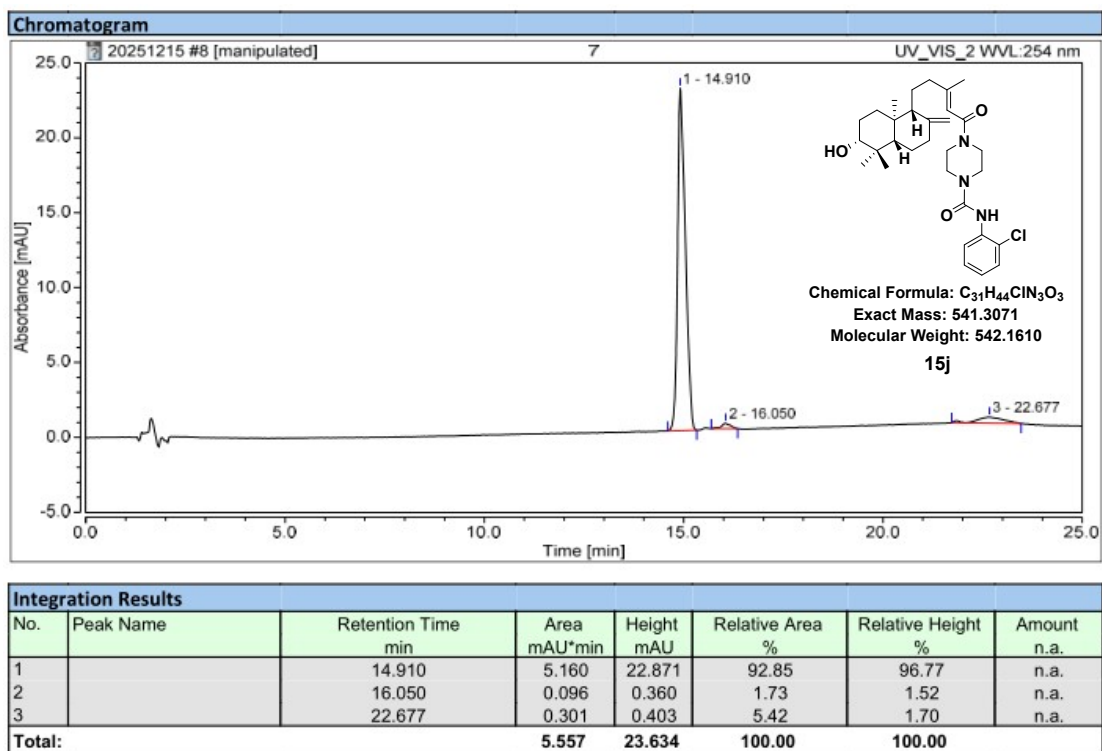


Figure S69. HPLC spectrum of 15j

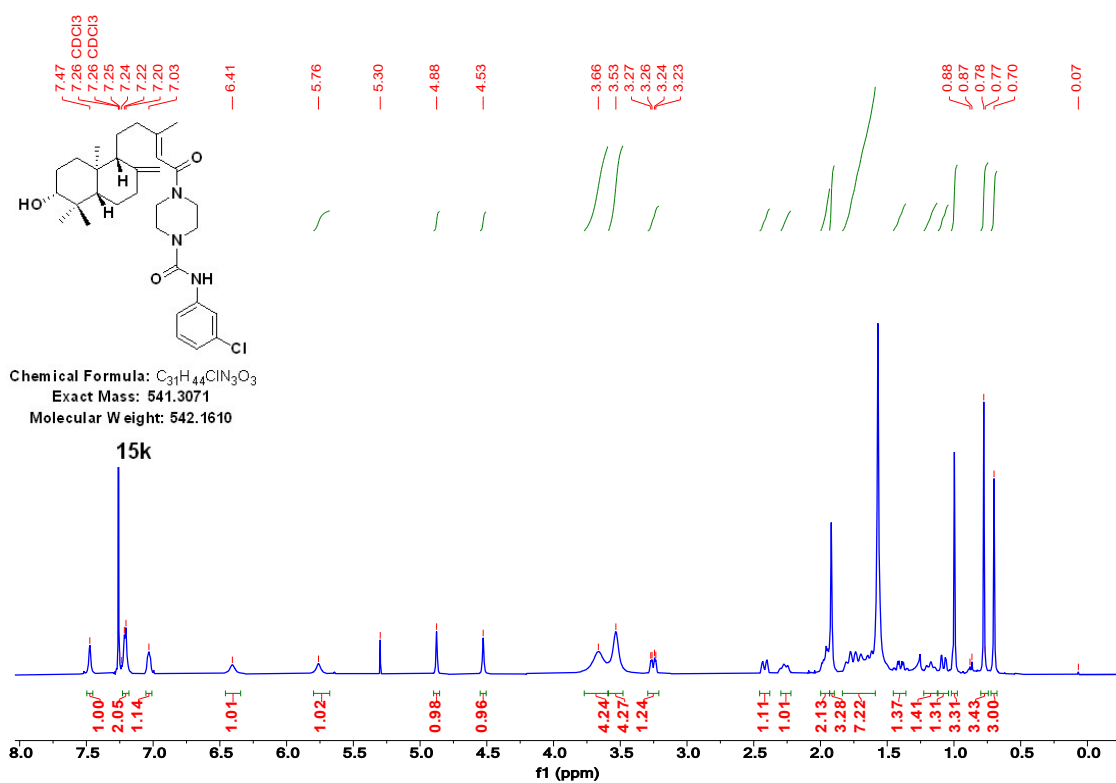


Figure S70. 1H NMR spectrum of 15k

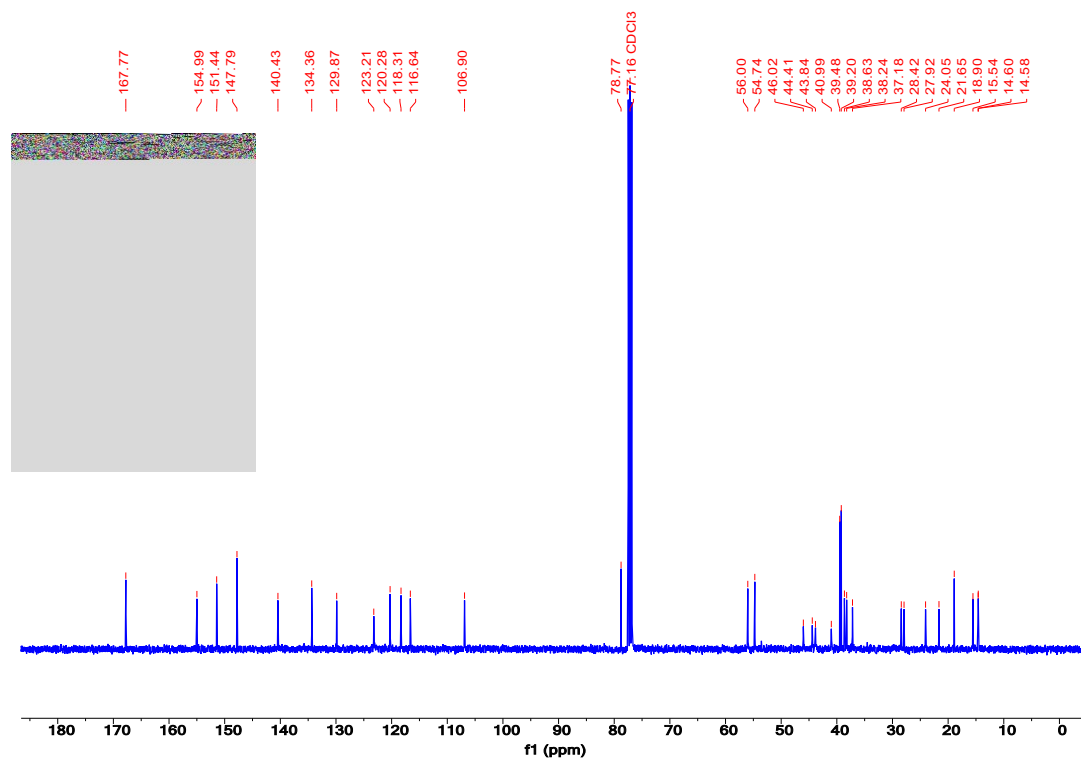


Figure S71. ^{13}C NMR spectrum of 15k

Single Mass Analysis

Tolerance = 5.0 mDa / DBE: min = -1.5, max = 50.0

Element prediction: Off

Number of isotope peaks used for i-FIT = 3

Monoisotopic Mass, Even Electron Ions

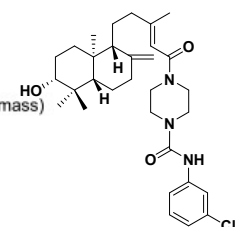
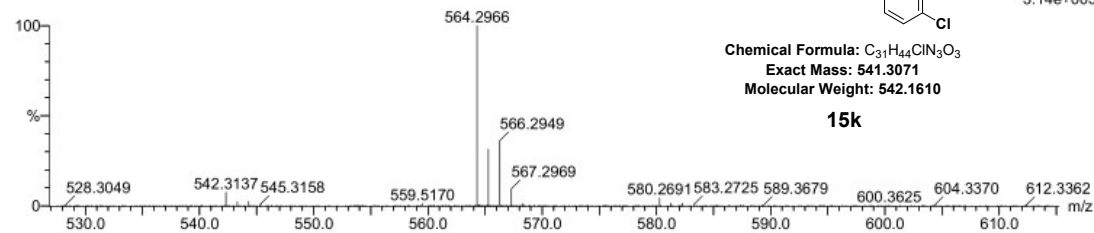
32 formula(e) evaluated with 1 results within limits (up to 50 best isotopic matches for each mass)

Elements Used:

C: 30-35 H: 44-51 N: 2-5 O: 2-5 Na: 0-1 Cl: 0-1

30

250310-2-Q-8 20 (0.063)



1: TOF MS ES+
3.14e+005

Chemical Formula: $\text{C}_{31}\text{H}_{44}\text{ClN}_3\text{O}_3$
Exact Mass: 541.3071
Molecular Weight: 542.1610

15k

Minimum: -1.5
Maximum: 5.0 10.0 50.0

Mass	Calc. Mass	mDa	PPM	DBE	i-FIT	Norm	Conf (%)	Formula
564.2966	564.2969	-0.3	-0.5	10.5	466.1	n/a	n/a	C31 H44 N3 O3 Na Cl

Figure S72. HRMS spectrum of 15k

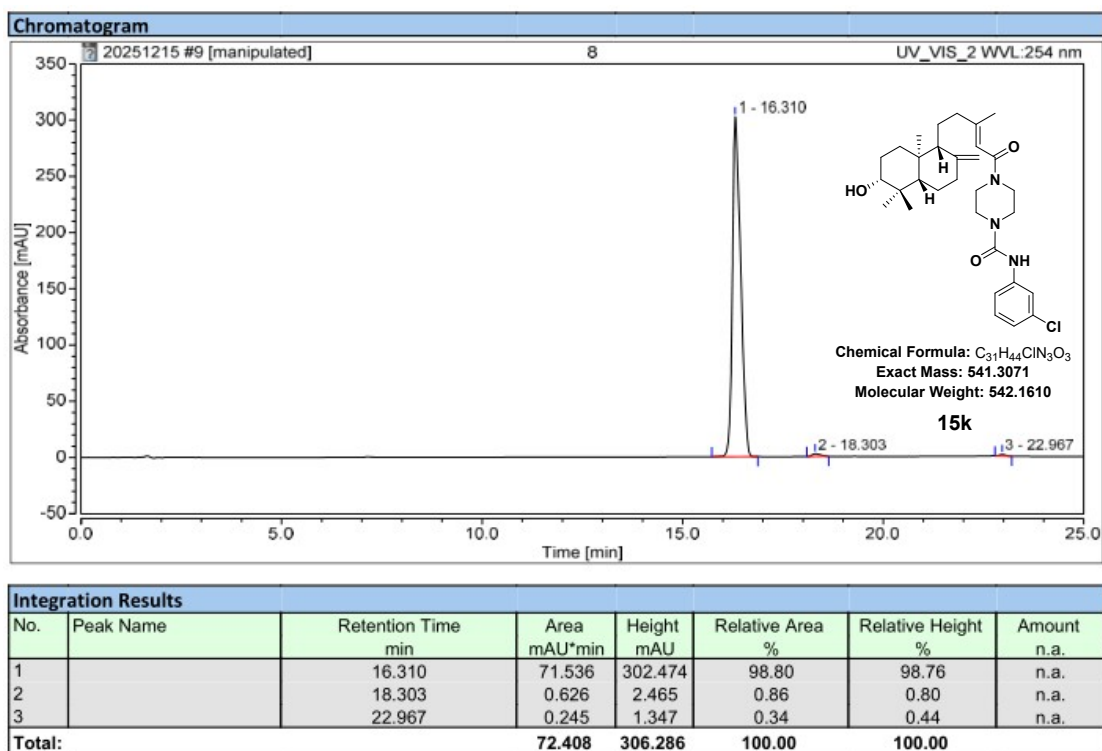


Figure S73. HPLC spectrum of 15k

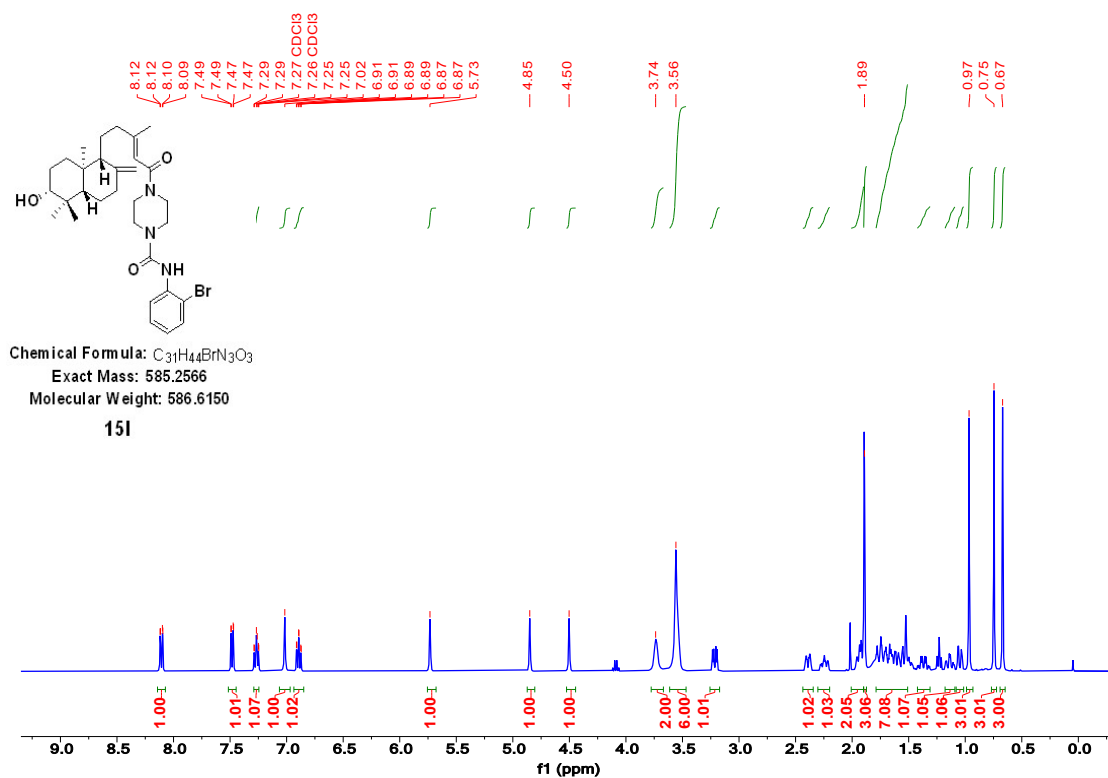


Figure S74. 1H NMR spectrum of 15l

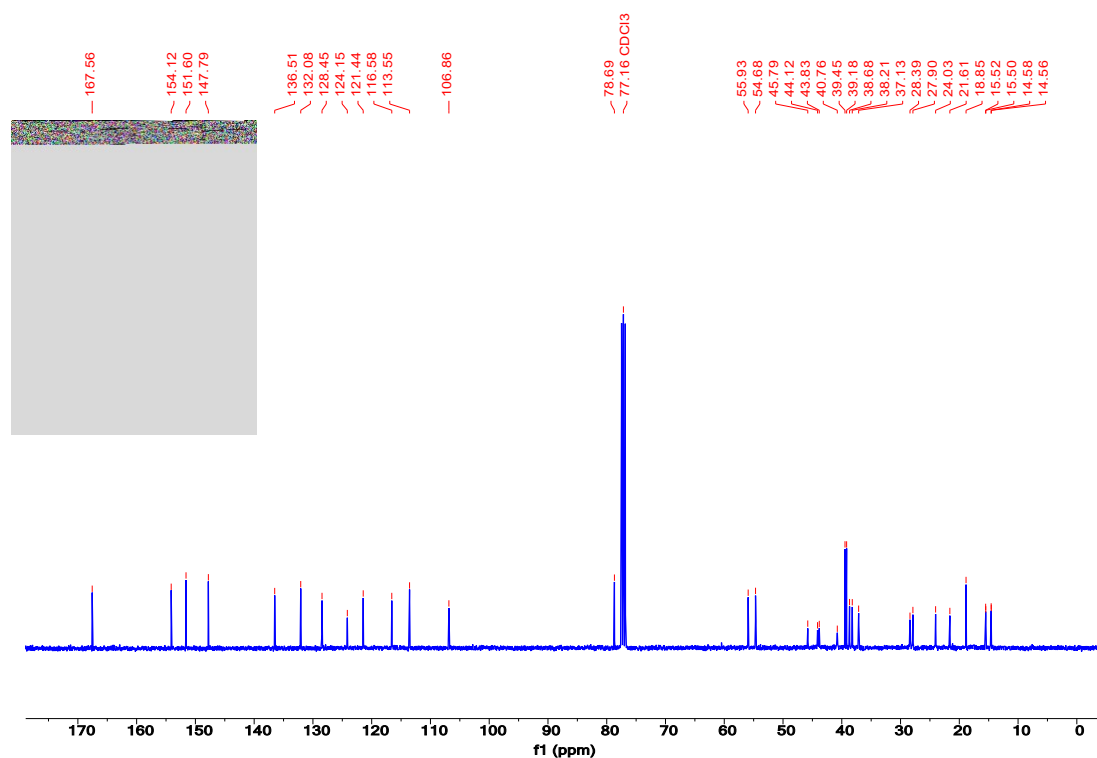


Figure S75. ¹³C NMR spectrum of 15I

Single Mass Analysis

Tolerance = 5.0 PPM / DBE: min = -1.5, max = 50.0

Element prediction: Off

Number of isotope peaks used for i-FIT = 3

Monoisotopic Mass, Even Electron Ions

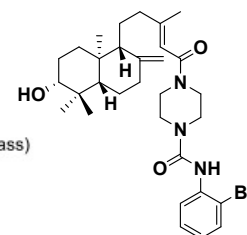
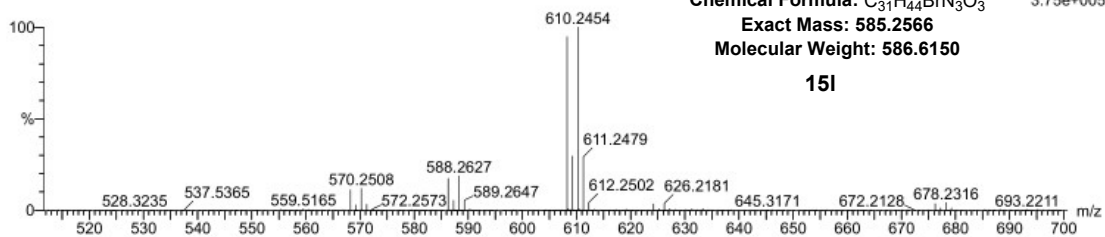
878 formula(e) evaluated with 1 results within limits (all results (up to 1000) for each mass)

Elements Used:

C: 31-31 H: 16-50 N: 1-3 O: 1-20 Na: 1-8 Br: 1-2

7

250310-2-Q-11 22 (0.067)



Chemical Formula: C₃₁H₄₄BrN₃O₃
 Exact Mass: 585.2566
 Molecular Weight: 586.6150

15I

1: TOF MS ES+
 3.75e+005

Minimum: -1.5
 Maximum: 50.0

Mass	Calc. Mass	mDa	PPM	DBE	i-FIT	Norm	Conf (%)	Formula
608.2466	608.2464	0.2	0.3	10.5	643.0	n/a	n/a	C31 H44 N3 O3 Na Br

Figure S76. HRMS spectrum of 15I

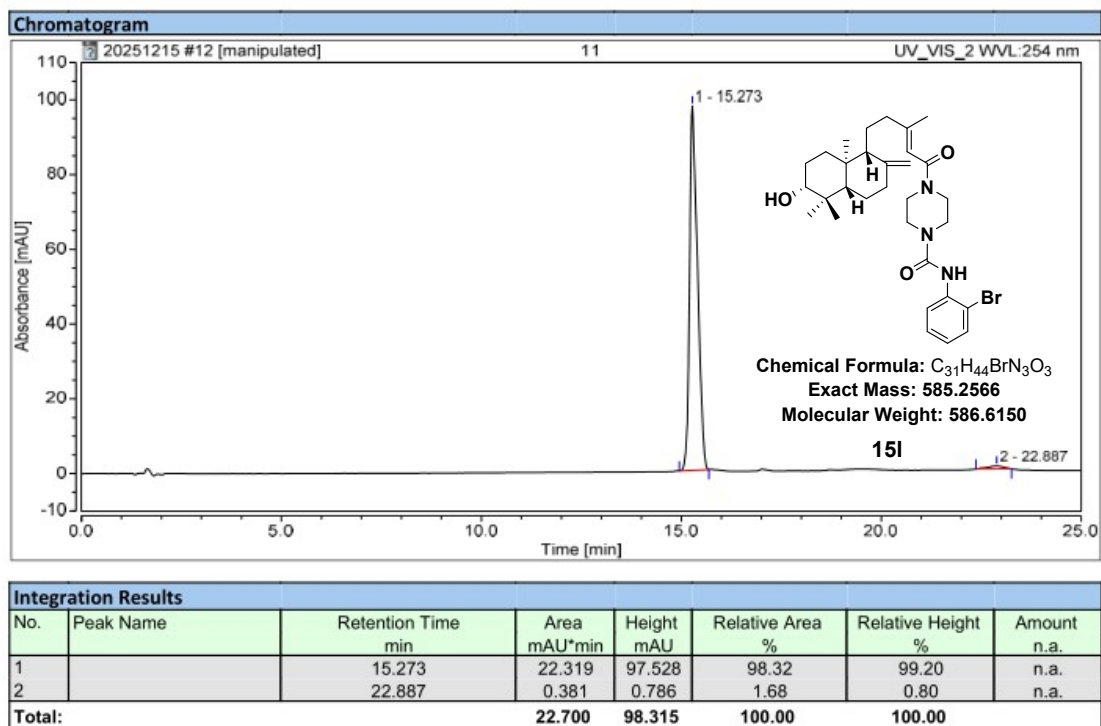


Figure S77. HPLC spectrum of 15I

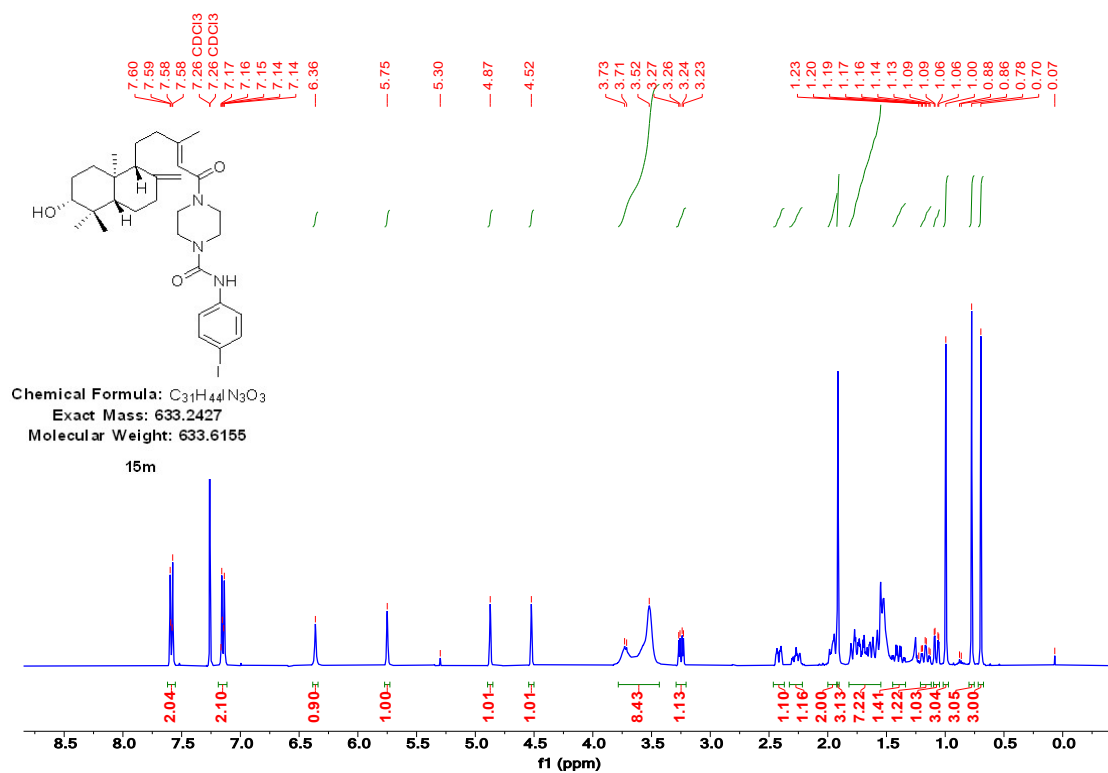


Figure S78. ¹H NMR spectrum of 15m

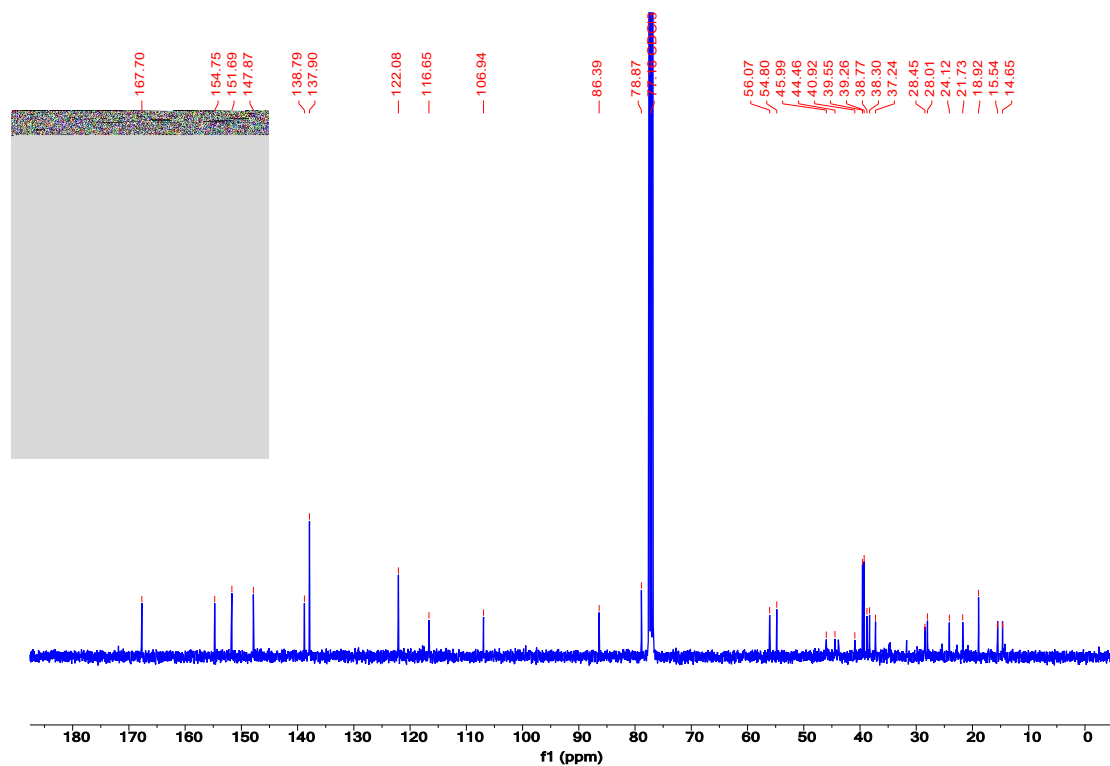


Figure S79. ^{13}C NMR spectrum of 15m

Single Mass Analysis

Tolerance = 5.0 mDa / DBE: min = -1.5, max = 50.0

Element prediction: Off

Number of isotope peaks used for i-FIT = 3

Monoisotopic Mass, Even Electron Ions

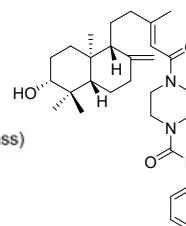
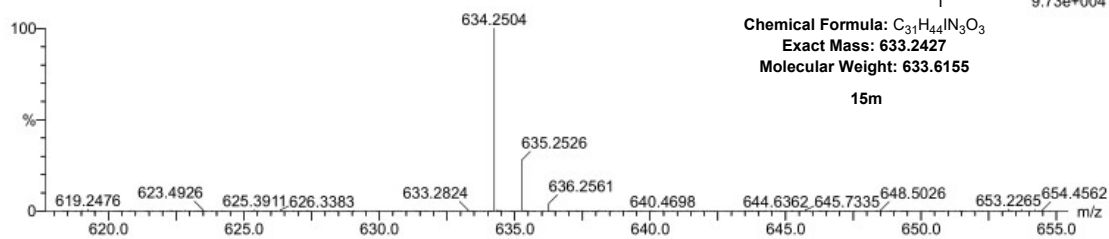
112 formula(e) evaluated with 1 results within limits (up to 50 best isotopic matches for each mass)

Elements Used:

C: 29-56 H: 44-85 N: 2-6 O: 2-6 I: 0-1

7

250310-2-Q-22 27 (0.076)



1: TOF MS ES+
9.73e+004

Chemical Formula: $\text{C}_{31}\text{H}_{44}\text{IN}_3\text{O}_3$

Exact Mass: 633.2427

Molecular Weight: 633.6155

15m

Minimum: -1.5
Maximum: 5.0 10.0 50.0

Mass	Calc. Mass	mDa	PPM	DBE	i-FIT	Norm	Conf (%)	Formula
634.2504	634.2506	-0.2	-0.3	10.5	337.7	n/a	n/a	C31 H45 N3 O3 I

Figure S80. HRMS spectrum of 15m

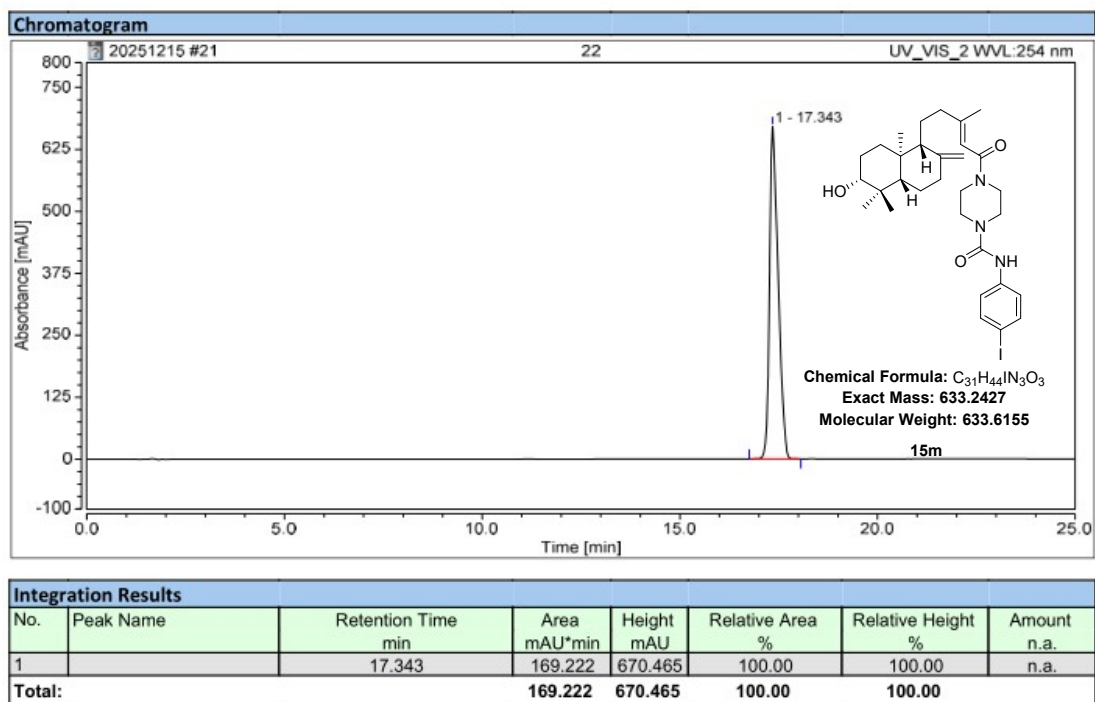


Figure S81. HPLC spectrum of 15m

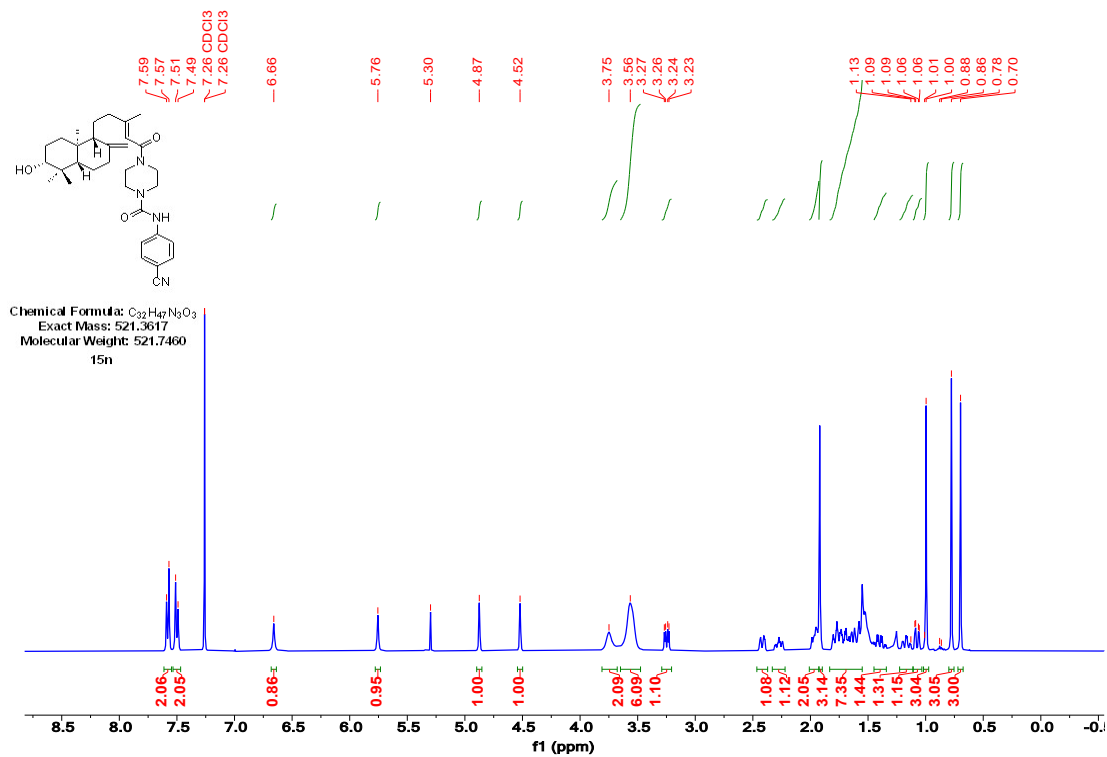


Figure S82. 1H NMR spectrum of 15n

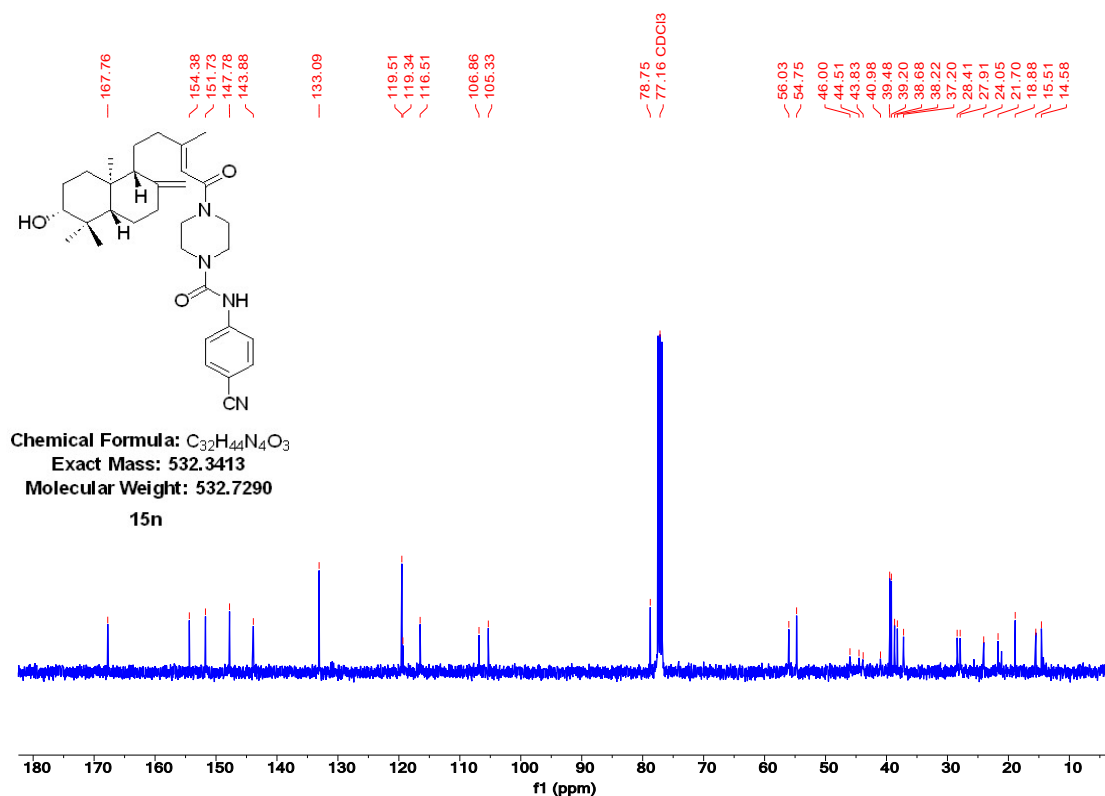


Figure S83. ^{13}C NMR spectrum of 15n

Single Mass Analysis

Tolerance = 5.0 mDa / DBE: min = -1.5, max = 50.0

Element prediction: Off

Number of isotope peaks used for i-FIT = 3

Monoisotopic Mass, Even Electron Ions

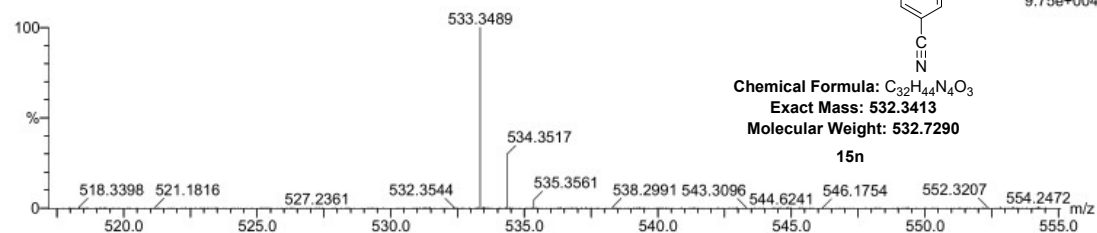
18 formula(e) evaluated with 1 results within limits (up to 50 best isotopic matches for each mass)

Elements Used:

C: 32-37 H: 44-49 N: 2-6 O: 2-6 Na: 0-1

7

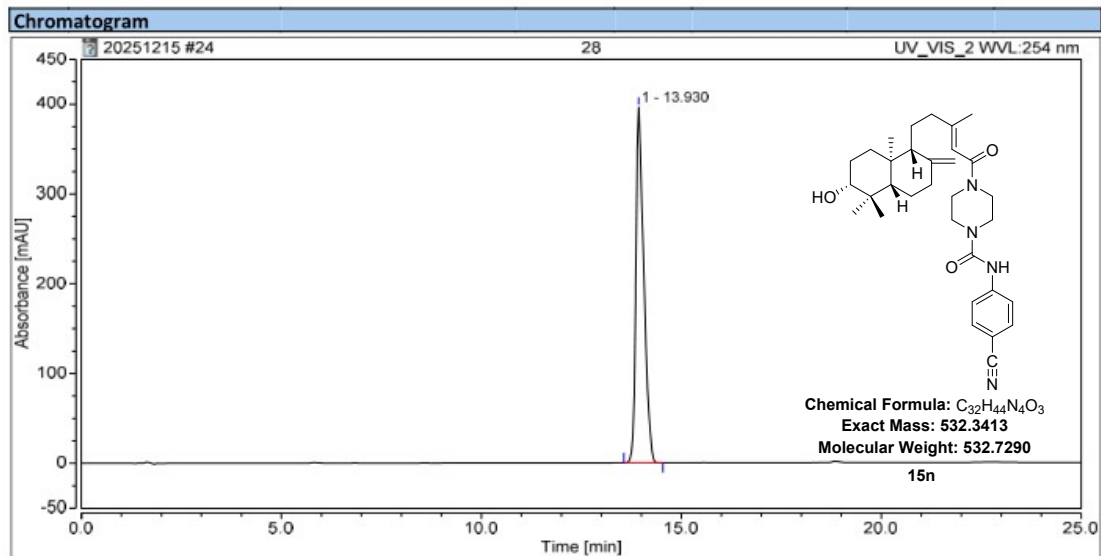
250310-2-Q-28 27 (0.076)



Minimum: -1.5
 Maximum: 50.0

Mass	Calc. Mass	mDa	PPM	DBE	i-FIT	Norm	Conf(%)	Formula
533.3489	533.3492	-0.3	-0.6	12.5	340.0	n/a	n/a	C32 H45 N4 O3

Figure S84. HRMS spectrum of 15n



Integration Results

No.	Peak Name	Retention Time min	Area mAU*min	Height mAU	Relative Area %	Relative Height %	Amount n.a.
1		13.930	90.274	396.385	100.00	100.00	n.a.
Total:			90.274	396.385	100.00	100.00	

Figure S85. HPLC spectrum of 15n

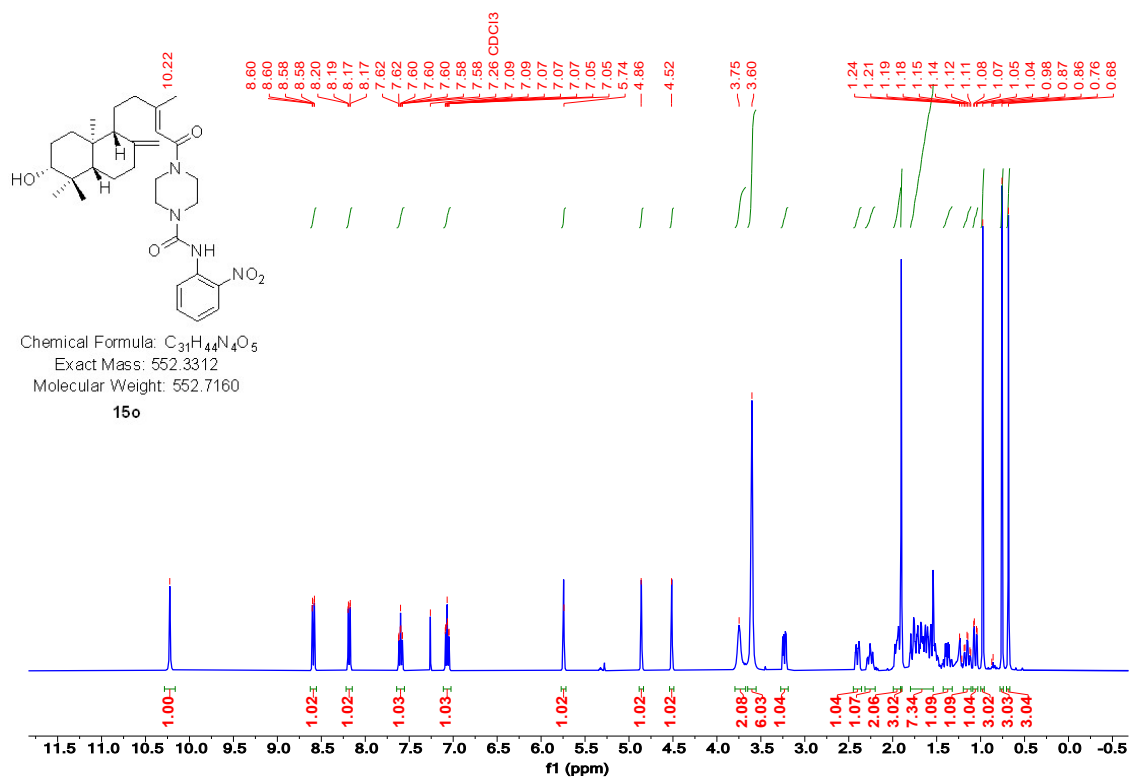


Figure S86. 1H NMR spectrum of 15o

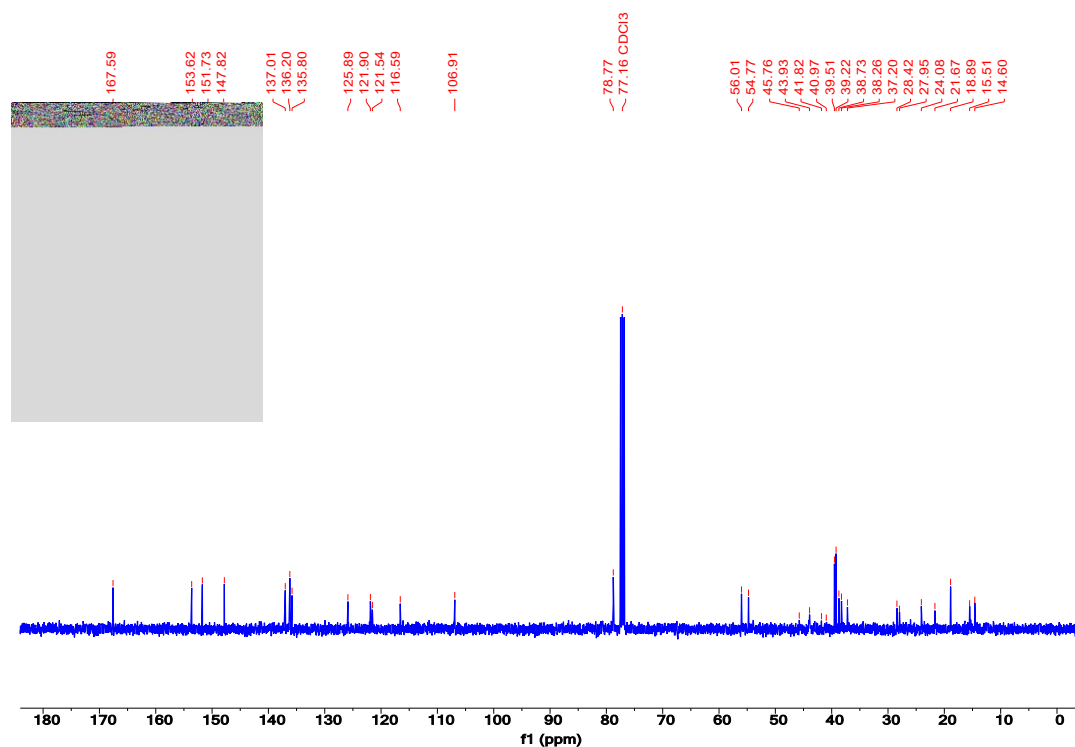


Figure S87. ^{13}C NMR spectrum of **15o**

Single Mass Analysis

Tolerance = 5.0 mDa / DBE: min = -1.5, max = 50.0

Element prediction: Off

Number of isotope peaks used for i-FIT = 3

Monoisotopic Mass, Even Electron Ions

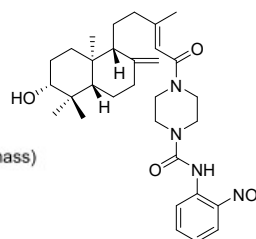
3 formula(e) evaluated with 1 results within limits (up to 50 best isotopic matches for each mass)

Elements Used:

C: 29-31 H: 44-45 N: 2-6 O: 2-6

7

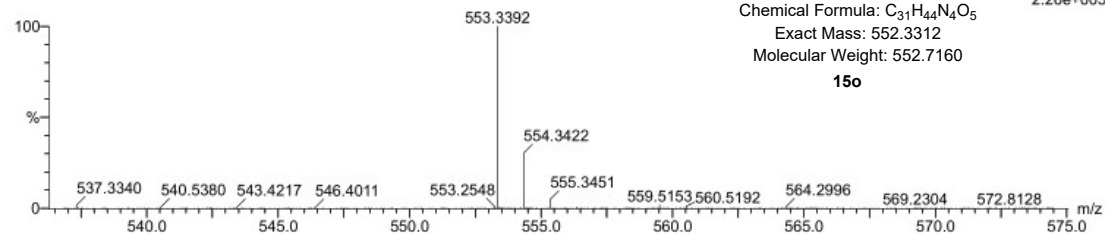
250310-2-Q-24 34 (0.090)



Chemical Formula: $\text{C}_{31}\text{H}_{44}\text{N}_4\text{O}_5$
 Exact Mass: 552.3312
 Molecular Weight: 552.7160

1: TOF MS ES+
2.26e+005

15o



Minimum: -1.5
 Maximum: 5.0 10.0 50.0

Mass	Calc. Mass	mDa	PPM	DBE	i-FIT	Norm	Conf (%)	Formula
553.3392	553.3390	0.2	0.4	11.5	442.7	n/a	n/a	C31 H45 N4 O5

Figure S88. HRMS spectrum of **15o**

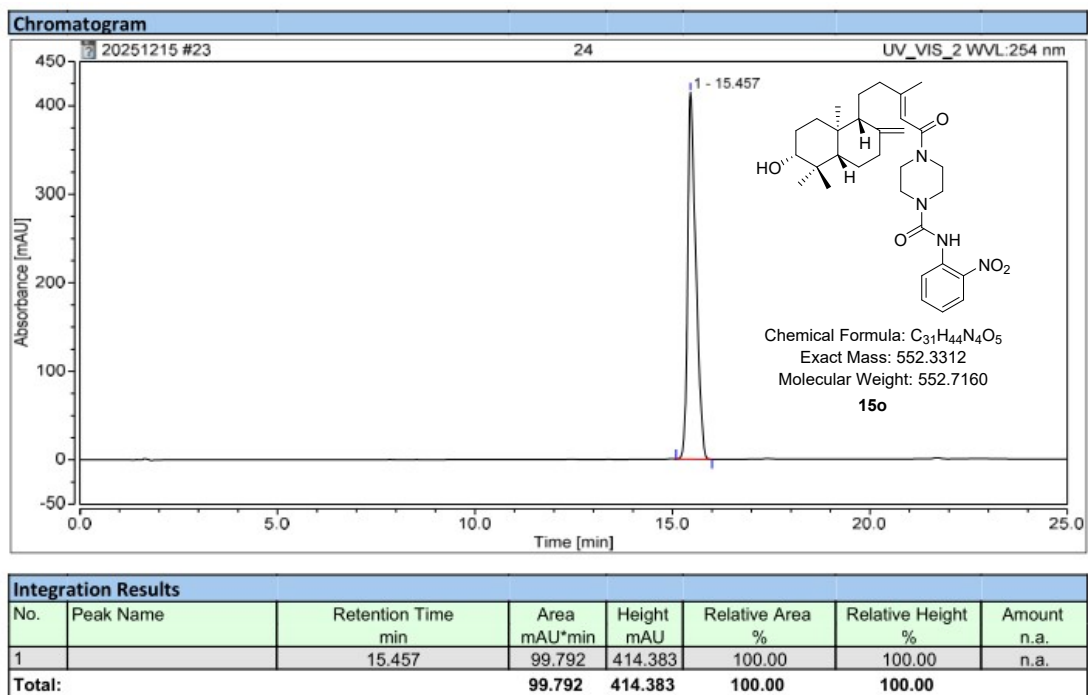


Figure S89. HPLC spectrum of 15o

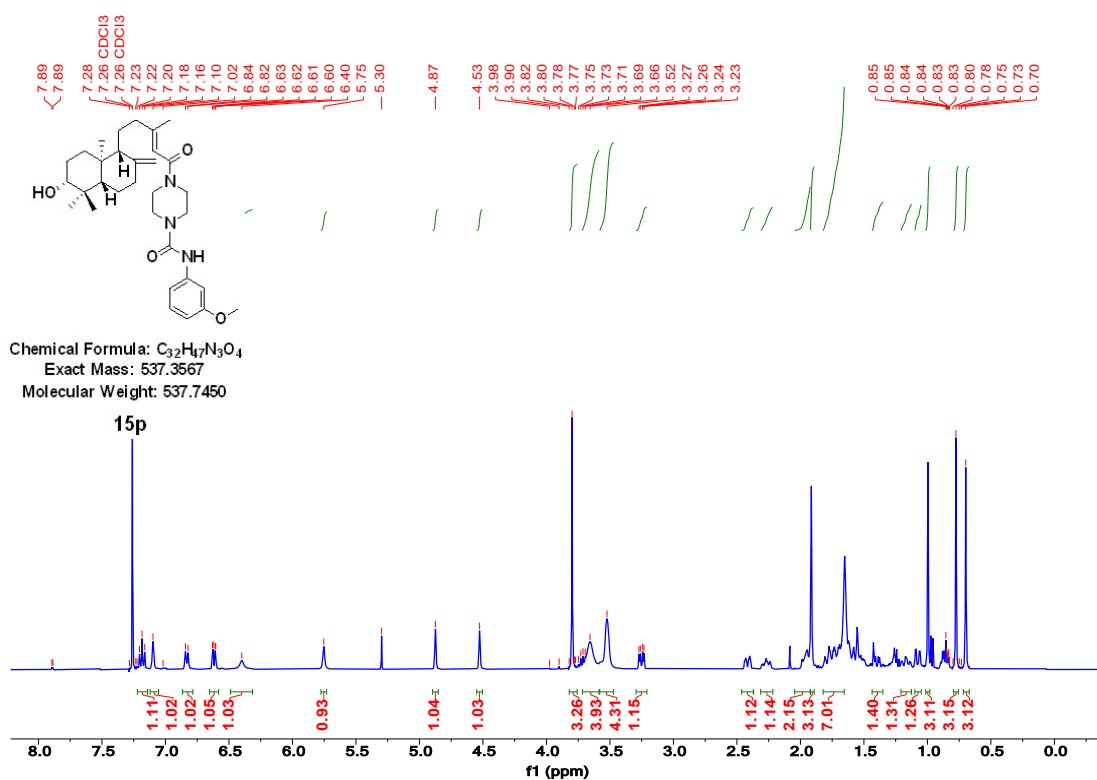


Figure S90. 1H NMR spectrum of 15p

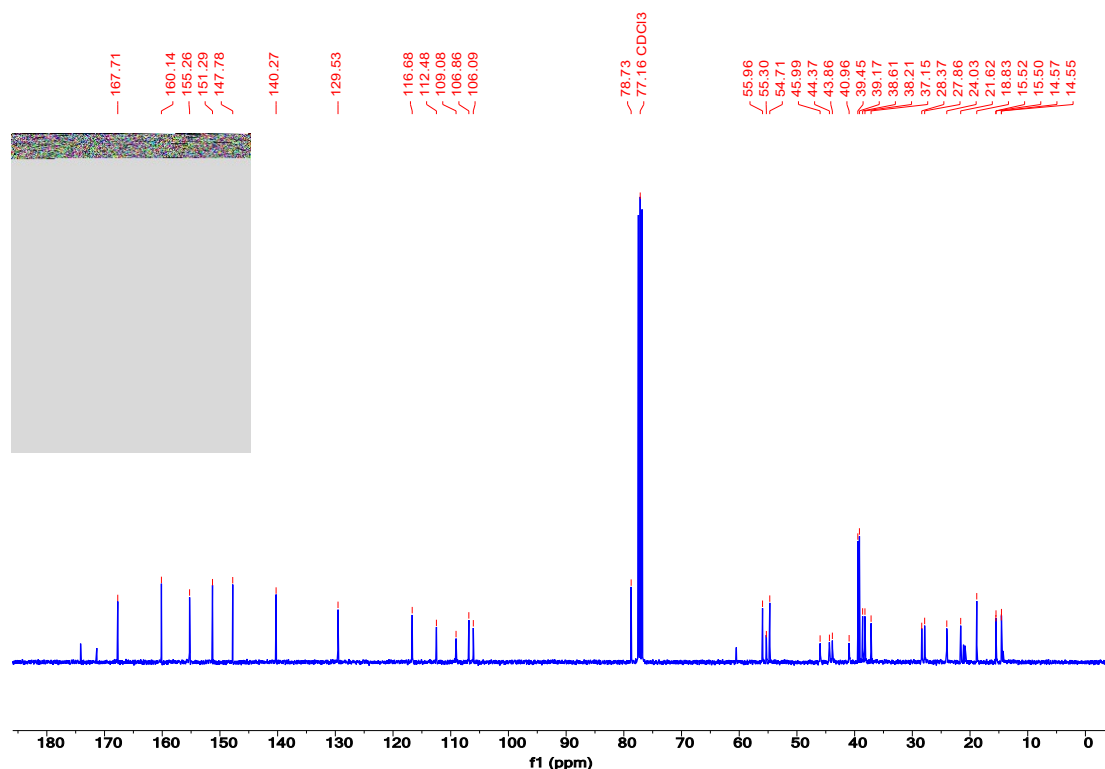


Figure S91. ¹³C NMR spectrum of 15p

Single Mass Analysis

Tolerance = 5.0 mDa / DBE: min = -1.5, max = 50.0

Element prediction: Off

Number of isotope peaks used for i-FIT = 3

Monoisotopic Mass, Odd and Even Electron Ions

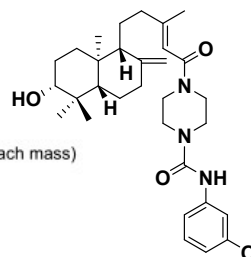
15 formula(e) evaluated with 1 results within limits (up to 50 best isotopic matches for each mass)

Elements Used:

C: 30-35 H: 44-51 N: 2-5 O: 2-5 K: 0-1

30

250310-2-Q-5 39 (0.099)



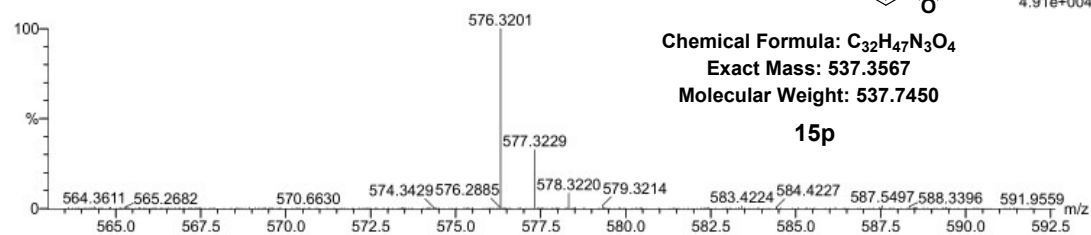
1: TOF MS ES+
4.91e+004

Chemical Formula: C₃₂H₄₇N₃O₄

Exact Mass: 537.3567

Molecular Weight: 537.7450

15p



Minimum: -1.5
Maximum: 5.0 10.0 50.0

Mass	Calc. Mass	mDa	PPM	DBE	i-FIT	Norm	Conf (%)	Formula
576.3201	576.3204	-0.3	-0.5	10.5	361.8	n/a	n/a	C32 H47 N3 O4 K

Figure S92. HRMS spectrum of 15p

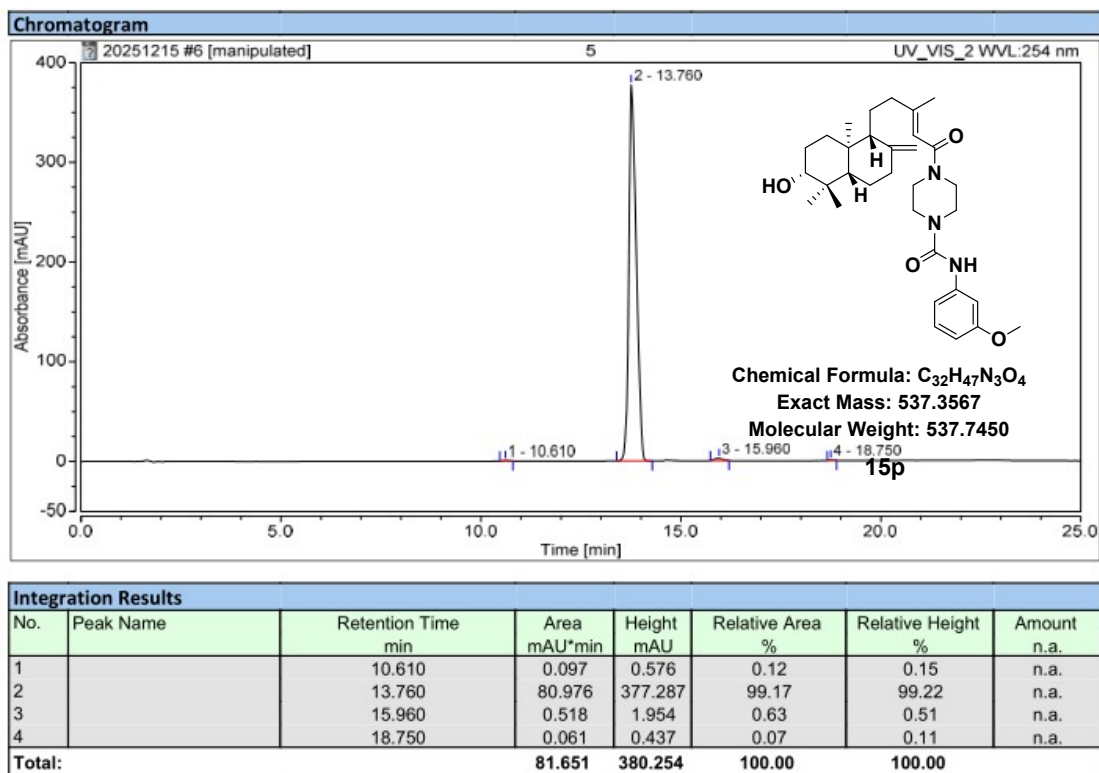


Figure S93. HPLC spectrum of 15p

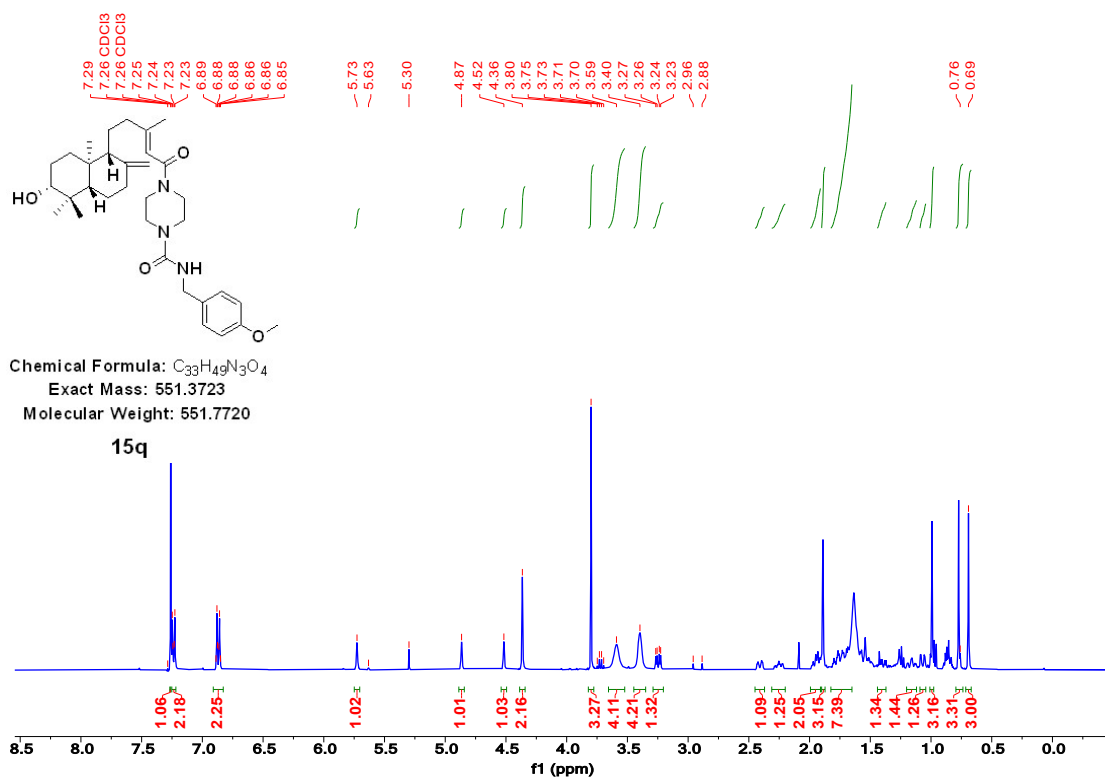


Figure S94. 1H NMR spectrum of 15q

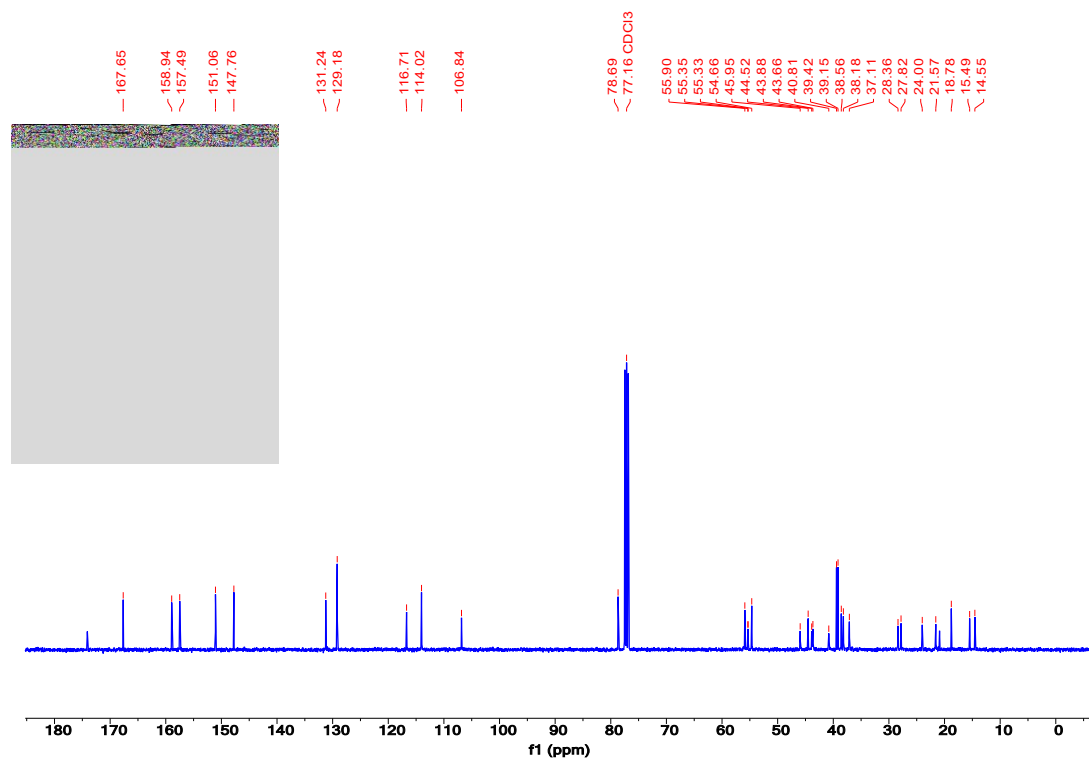


Figure S95. ^{13}C NMR spectrum of 15q

Single Mass Analysis

Tolerance = 5.0 mDa / DBE: min = -1.5, max = 50.0

Element prediction: Off

Number of isotope peaks used for i-FIT = 3

Monoisotopic Mass, Even Electron Ions

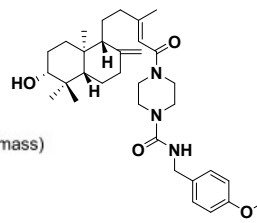
60 formula(e) evaluated with 1 results within limits (up to 50 best isotopic matches for each mass)

Elements Used:

C: 29-56 H: 47-85 N: 2-6 O: 2-6

30

250310-2-Q-9 27 (0.076)



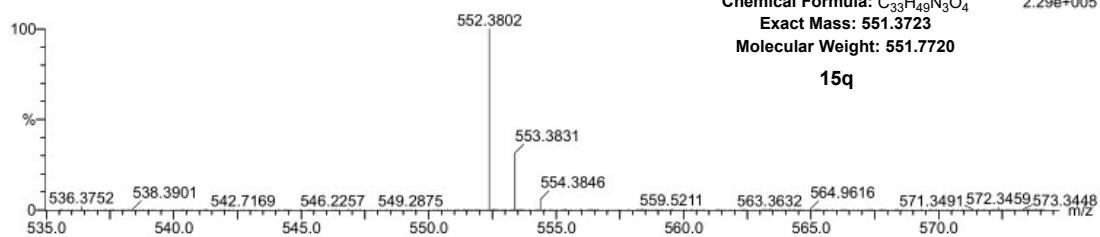
Chemical Formula: $\text{C}_{33}\text{H}_{49}\text{N}_3\text{O}_4$

Exact Mass: 551.3723

Molecular Weight: 551.7720

15q

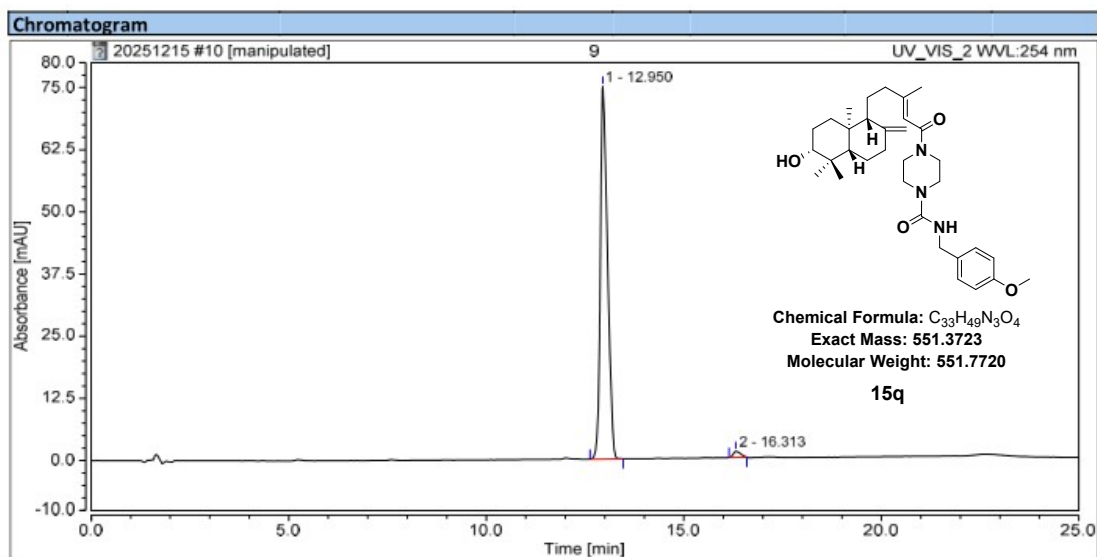
1: TOF MS ES+
2.29e+005



Minimum: -1.5
Maximum: 5.0 10.0 50.0

Mass	Calc. Mass	mDa	PPM	DBE	i-FIT	Norm	Conf(%)	Formula
552.3802	552.3801	0.1	0.2	10.5	450.3	n/a	n/a	C33 H50 N3 O4

Figure S96. HRMS spectrum of 15q



Integration Results

No.	Peak Name	Retention Time min	Area mAU*min	Height mAU	Relative Area %	Relative Height %	Amount
1		12.950	15.670	74.964	98.33	98.39	n.a.
2		16.313	0.266	1.230	1.67	1.61	n.a.
Total:			15.936	76.193	100.00	100.00	

Figure S97. HPLC spectrum of 15q

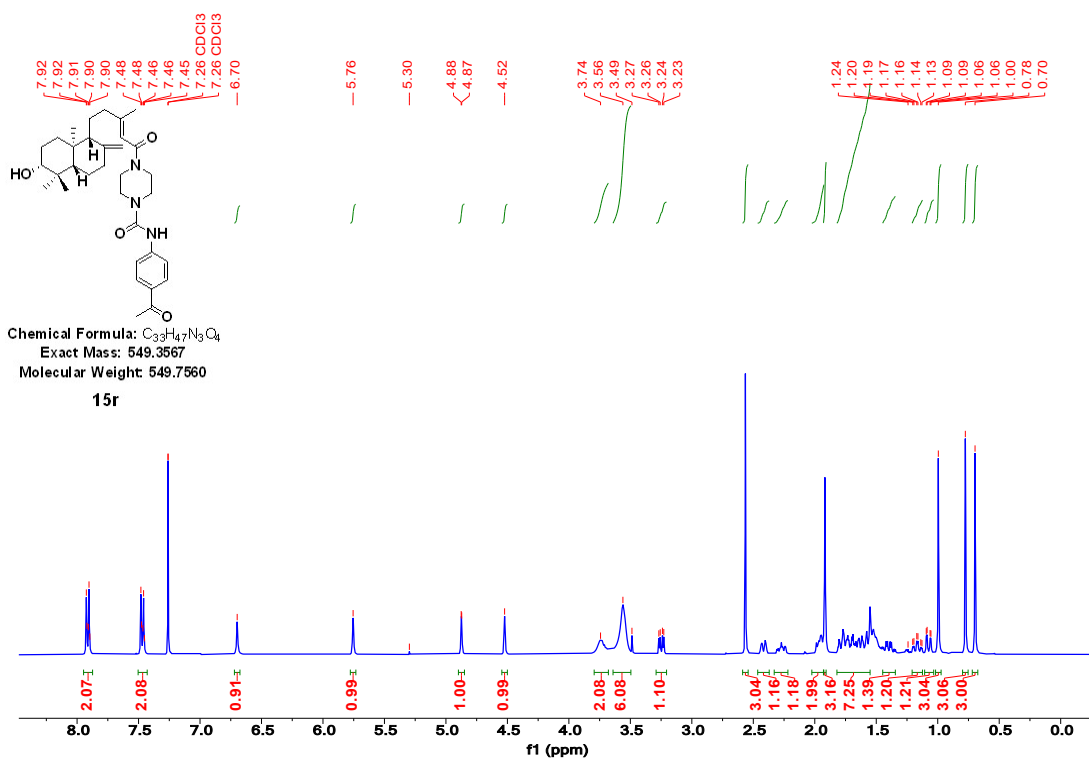


Figure S98. 1H NMR spectrum of 15r

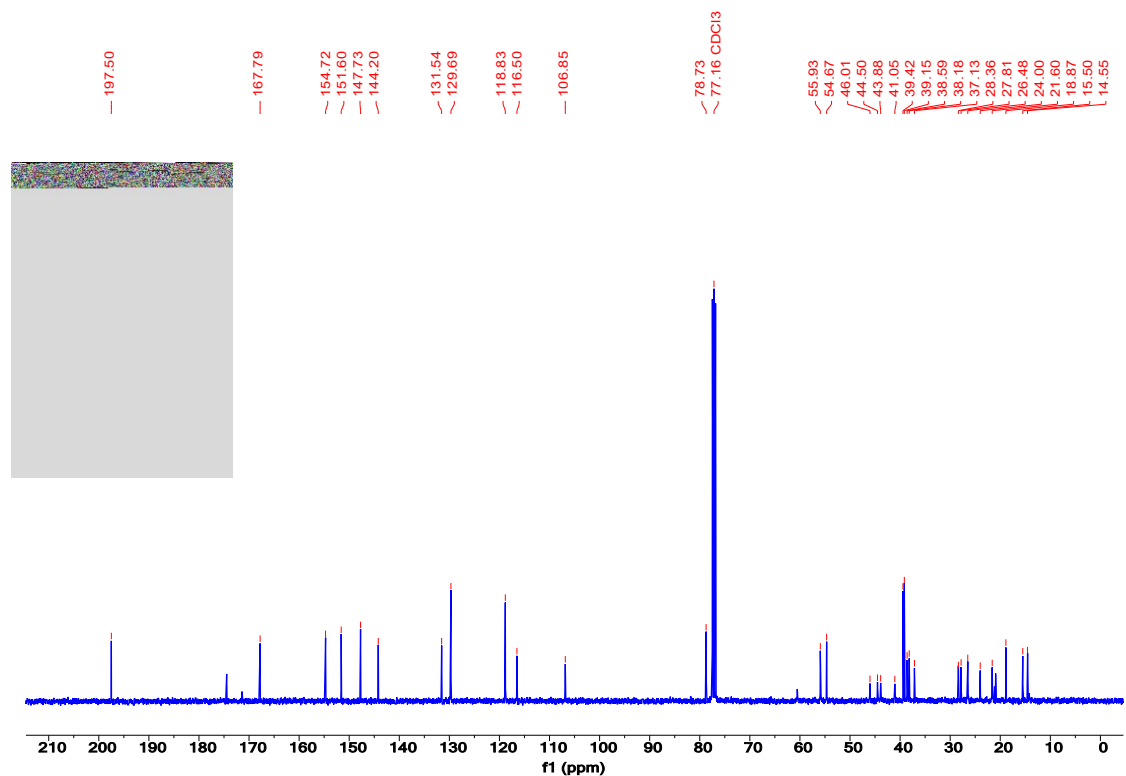


Figure S99. ^{13}C NMR spectrum of 15r

Single Mass Analysis

Tolerance = 5.0 mDa / DBE: min = -1.5, max = 50.0

Element prediction: Off

Number of isotope peaks used for i-FIT = 3

Monoisotopic Mass, Even Electron Ions

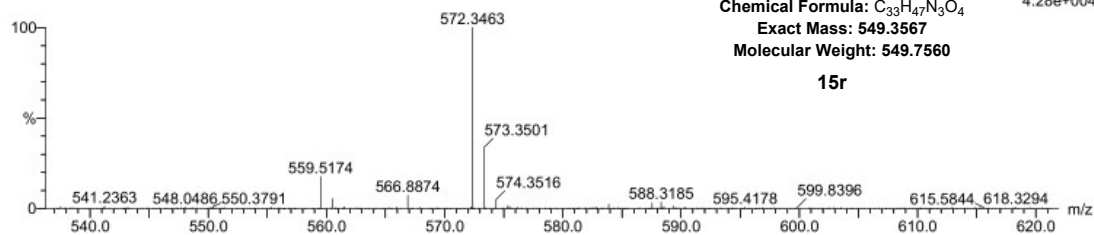
7 formula(e) evaluated with 1 results within limits (up to 50 best isotopic matches for each mass)

Elements Used:

C: 31-33 H: 44-51 N: 2-5 O: 2-5 Na: 0-1

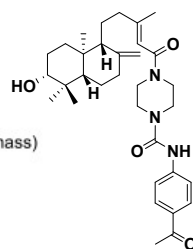
7

250310-2-Q-12 18 (0.059)



Minimum: -1.5
Maximum: 50.0

Mass	Calc. Mass	mDa	PPM	DBE	i-FIT	Norm	Conf(%)	Formula
572.3463	572.3464	-0.1	-0.2	11.5	143.3	n/a	n/a	C ₃₃ H ₄₇ N ₃ O ₄ Na



Chemical Formula: C₃₃H₄₇N₃O₄

Exact Mass: 549.3567

Molecular Weight: 549.7560

15r

1: TOF MS ES+
4.28e+004

Figure S100. HRMS spectrum of 15r

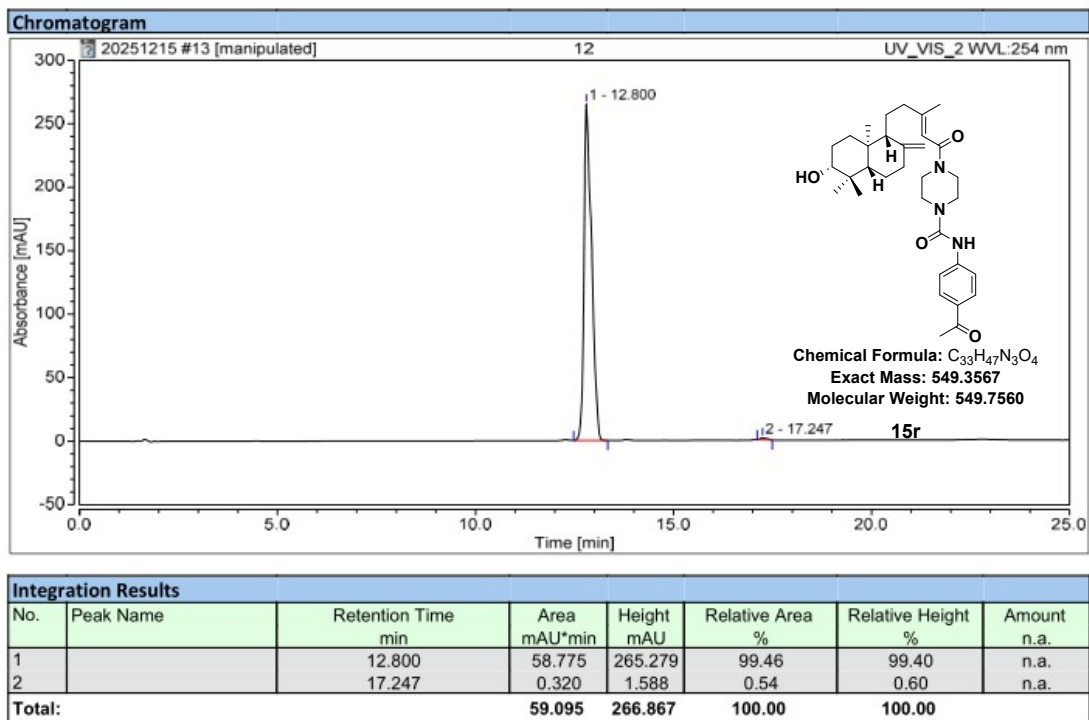


Figure S101. HPLC spectrum of 15r

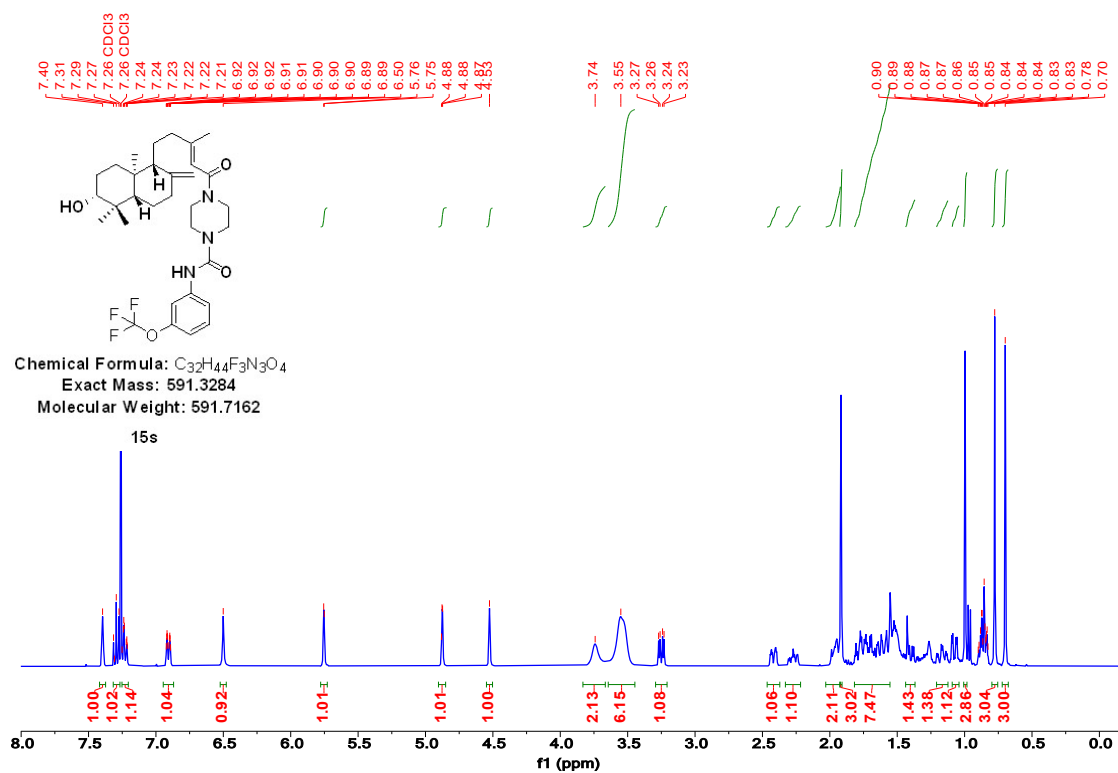


Figure S102. ¹H NMR spectrum of 15s

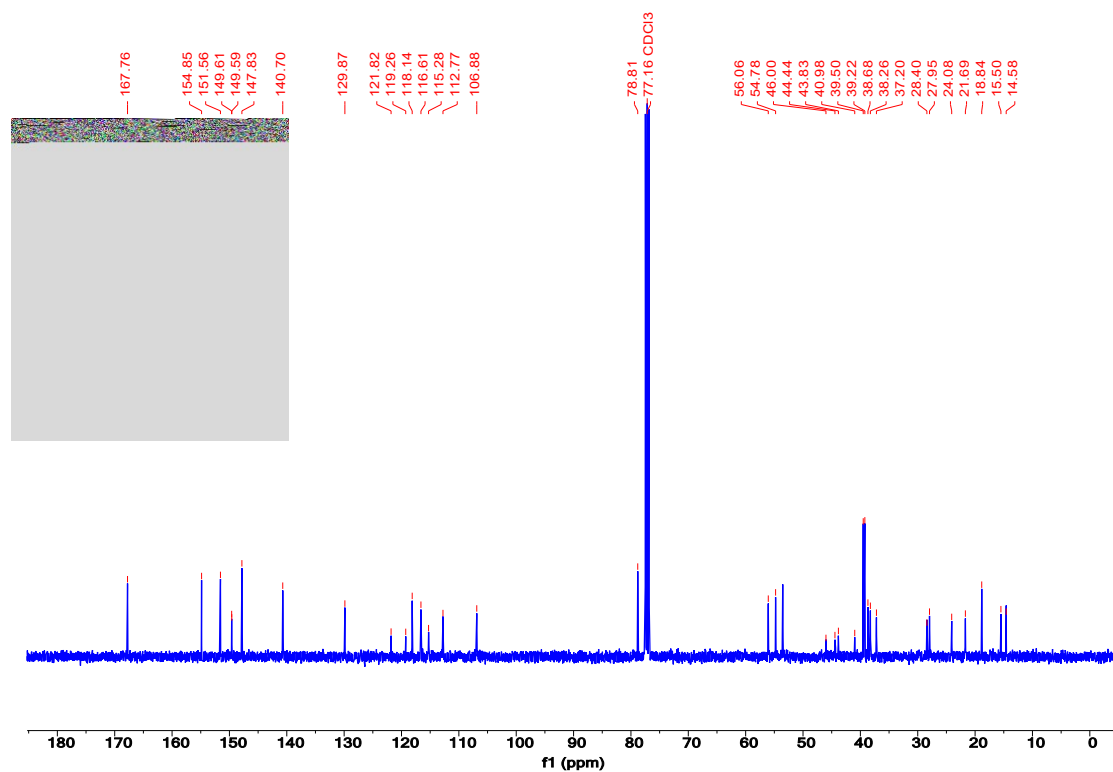


Figure S103. ^{13}C NMR spectrum of 15s

Single Mass Analysis

Tolerance = 5.0 mDa / DBE: min = -1.5, max = 50.0

Element prediction: Off

Number of isotope peaks used for i-FIT = 3

Monoisotopic Mass, Even Electron Ions

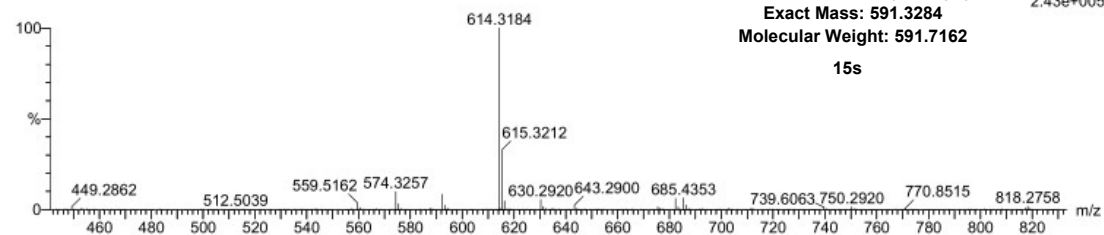
38 formula(e) evaluated with 1 results within limits (up to 50 best isotopic matches for each mass)

Elements Used:

C: 27-33 H: 44-51 N: 2-5 O: 2-5 F: 0-3 Na: 0-1

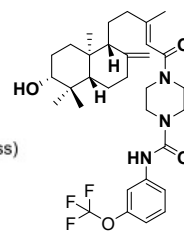
7

250310-2-Q-16 47 (0.115)



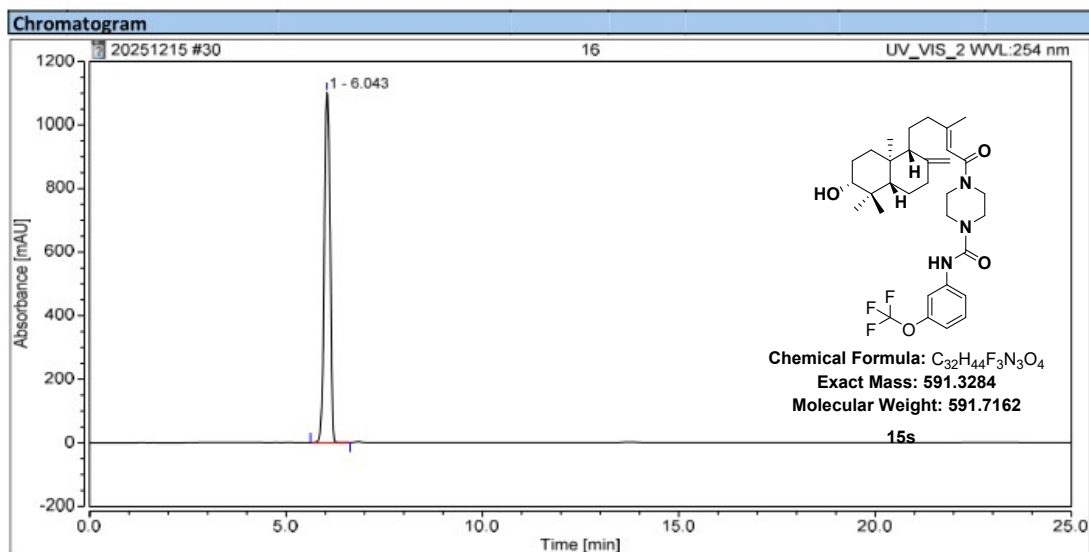
Minimum: -1.5
Maximum: 5.0 10.0 50.0

Mass	Calc. Mass	mDa	PPM	DBE	i-FIT	Norm	Conf (%)	Formula
614.3184	614.3182	0.2	0.3	10.5	425.6	n/a	n/a	C32 H44 N3 O4 F3 Na



Chemical Formula: $\text{C}_{32}\text{H}_{44}\text{F}_3\text{N}_3\text{O}_4$ ¹: TOF MS ES+
Exact Mass: 591.3284 2.43e+005
Molecular Weight: 591.7162
15s

Figure S104. HRMS spectrum of 15s



Integration Results

No.	Peak Name	Retention Time min	Area mAU*min	Height mAU	Relative Area %	Relative Height %	Amount n.a.
1		6.043	189.064	1102.248	100.00	100.00	n.a.
Total:			189.064	#####	100.00	100.00	

Figure S105. HPLC spectrum of 15s

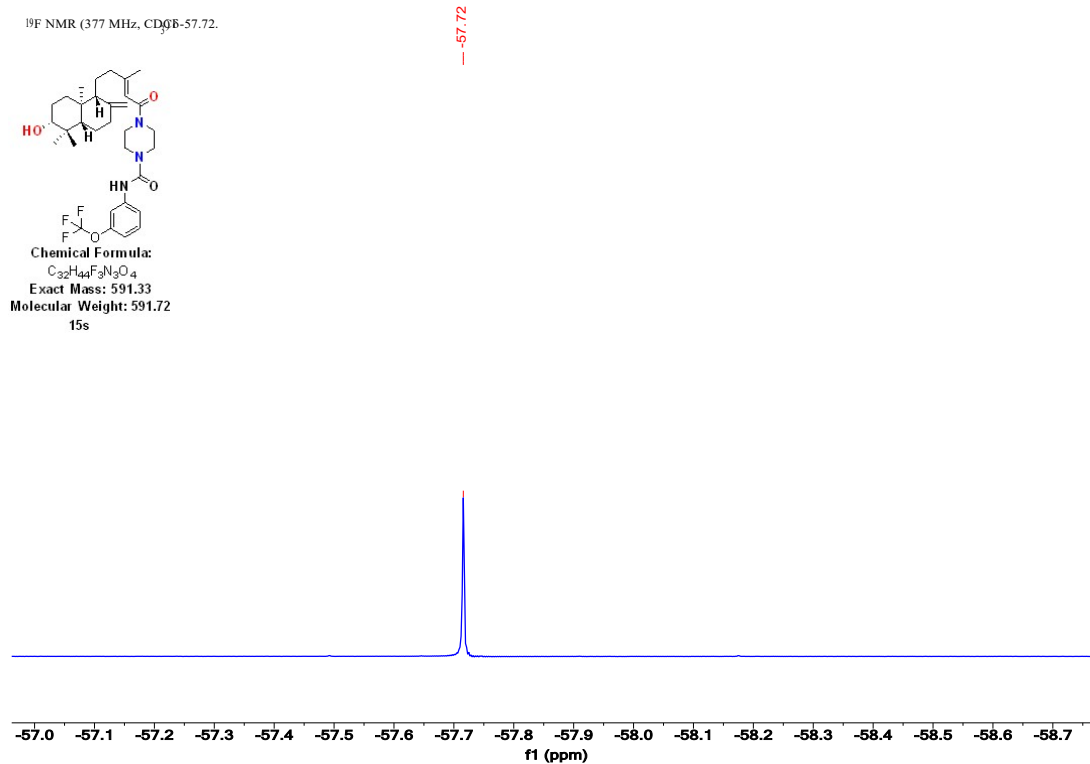


Figure S106. ^{19}F NMR spectrum of 15s

Single Mass Analysis

Tolerance = 5.0 mDa / DBE: min = -1.5, max = 50.0

Element prediction: Off

Number of isotope peaks used for i-FIT = 3

Monoisotopic Mass, Even Electron Ions

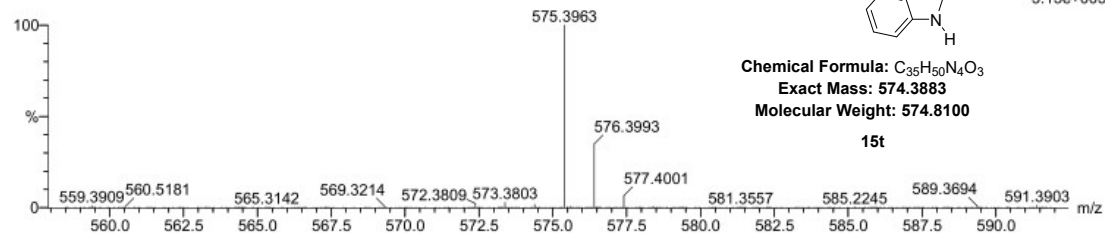
54 formula(e) evaluated with 1 results within limits (up to 50 best isotopic matches for each mass)

Elements Used:

C: 32-56 H: 44-85 N: 2-6 O: 2-6

7

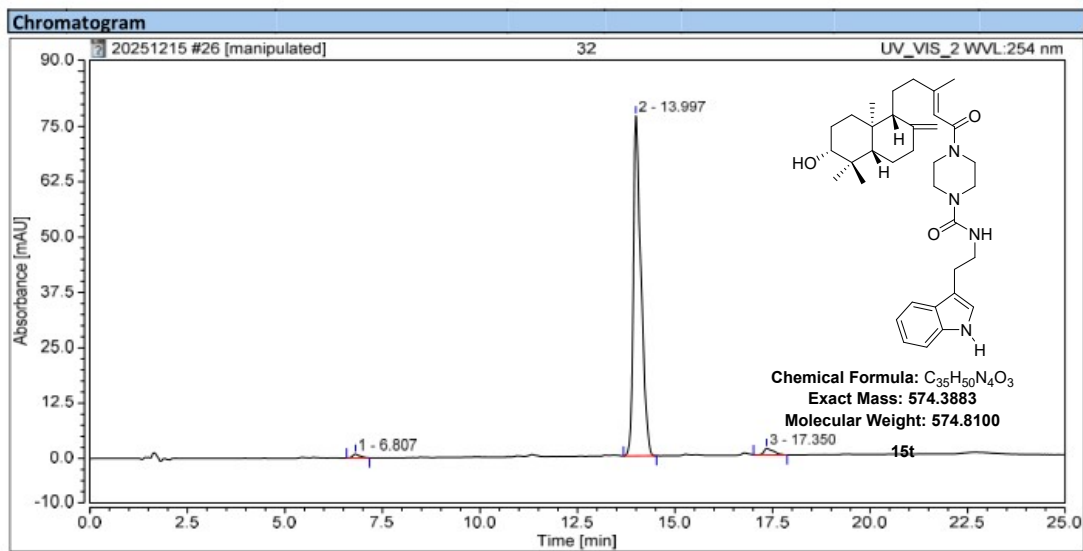
250310-2-Q-32 29 (0.080)



Minimum: -1.5
Maximum: 50.0

Mass	Calc. Mass	mDa	PPM	DBE	i-FIT	Norm	Conf (%)	Formula
575.3963	575.3961	0.2	0.3	12.5	461.6	n/a	n/a	C ₃₅ H ₅₁ N ₄ O ₃

Figure S109. HRMS spectrum of 15t



No.	Peak Name	Retention Time min	Area mAU*min	Height mAU	Relative Area %	Relative Height %	Amount n.a.
1		6.807	0.183	0.846	1.02	1.07	n.a.
2		13.997	17.451	76.864	96.75	97.12	n.a.
3		17.350	0.403	1.430	2.24	1.81	n.a.
Total:			18.038	79.140	100.00	100.00	

Figure S110. HPLC spectrum of 15t

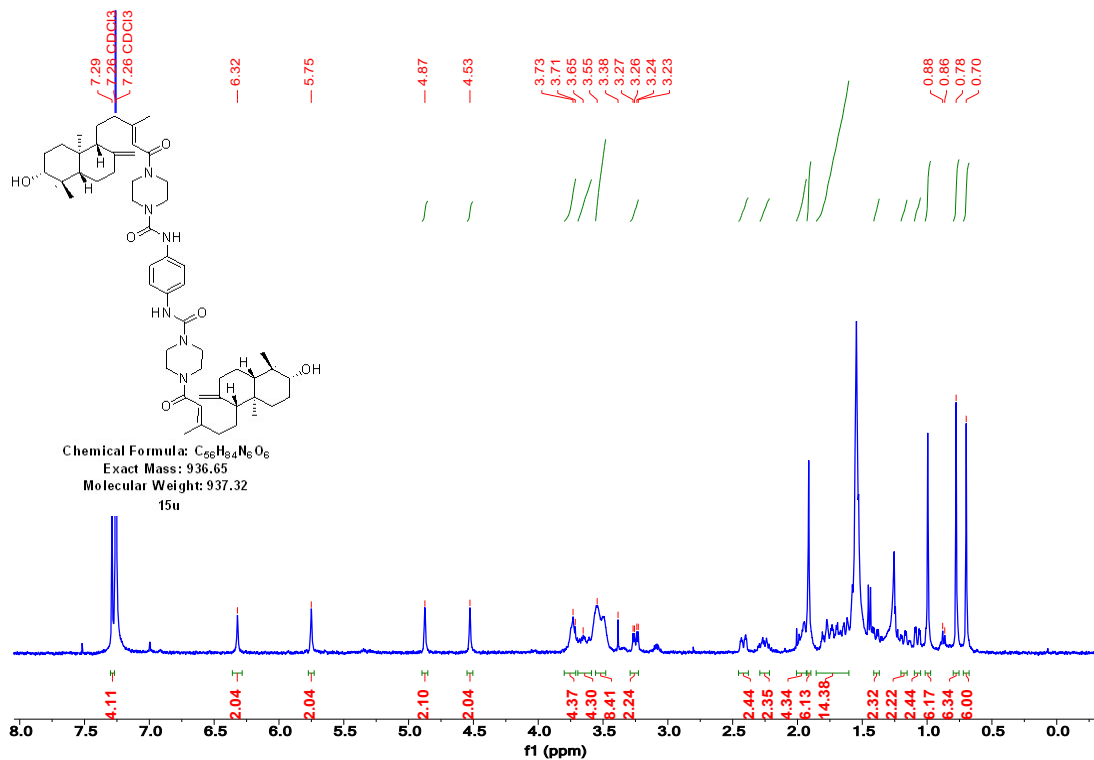


Figure S111. 1H NMR spectrum of 15u

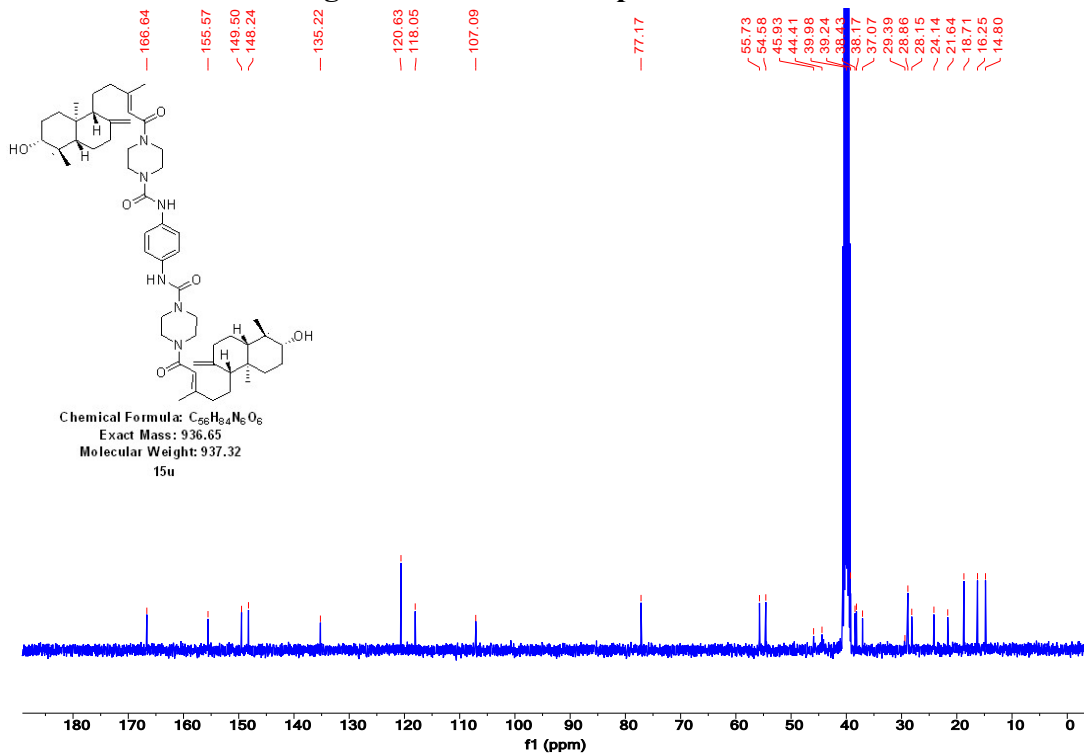


Figure S112. ^{13}C NMR spectrum of 15u

Single Mass Analysis

Tolerance = 5.0 mDa / DBE: min = -1.5, max = 50.0

Element prediction: Off

Number of isotope peaks used for i-FIT = 3

Monoisotopic Mass, Even Electron Ions

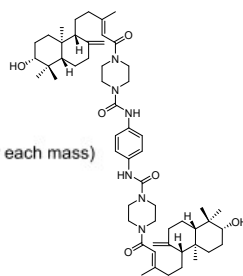
7 formula(e) evaluated with 1 results within limits (up to 50 best isotopic matches for each mass)

Elements Used:

C: 32-56 H: 44-85 N: 2-6 O: 2-6

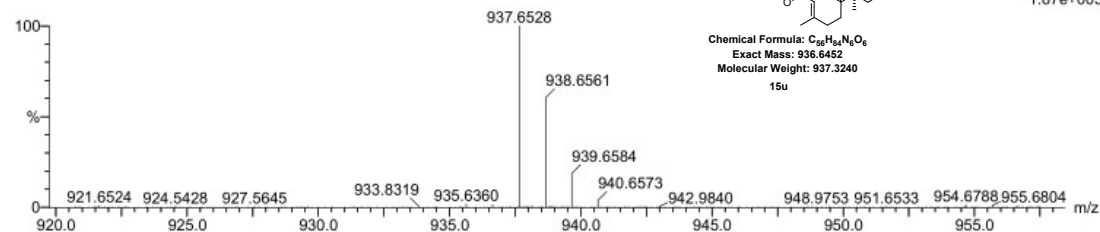
7

250310-2-Q-29 31 (0.084)



Chemical Formula: $C_{26}H_{32}N_2O_6$
Exact Mass: 936.6452
Molecular Weight: 937.3240
15u

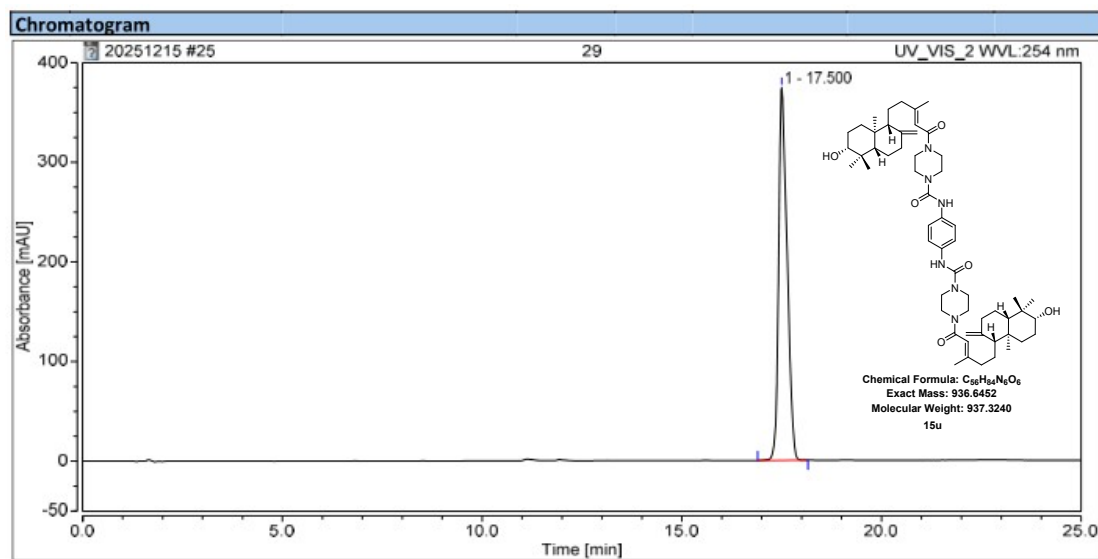
1: TOF MS ES+
1.87e+005



Minimum: -1.5
Maximum: 5.0 10.0 50.0

Mass	Calc. Mass	mDa	PPM	DBE	i-FIT	Norm	Conf (%)	Formula
937.6528	937.6531	-0.3	-0.3	17.5	416.3	n/a	n/a	C56 H85 N6 O6

Figure S113. HRMS spectrum of 15u



Integration Results							
No.	Peak Name	Retention Time min	Area mAU*min	Height mAU	Relative Area %	Relative Height %	Amount n.a.
1		17.500	91.730	374.563	100.00	100.00	n.a.
Total:			91.730	374.563	100.00	100.00	

Figure S114. HPLC spectrum of 15u

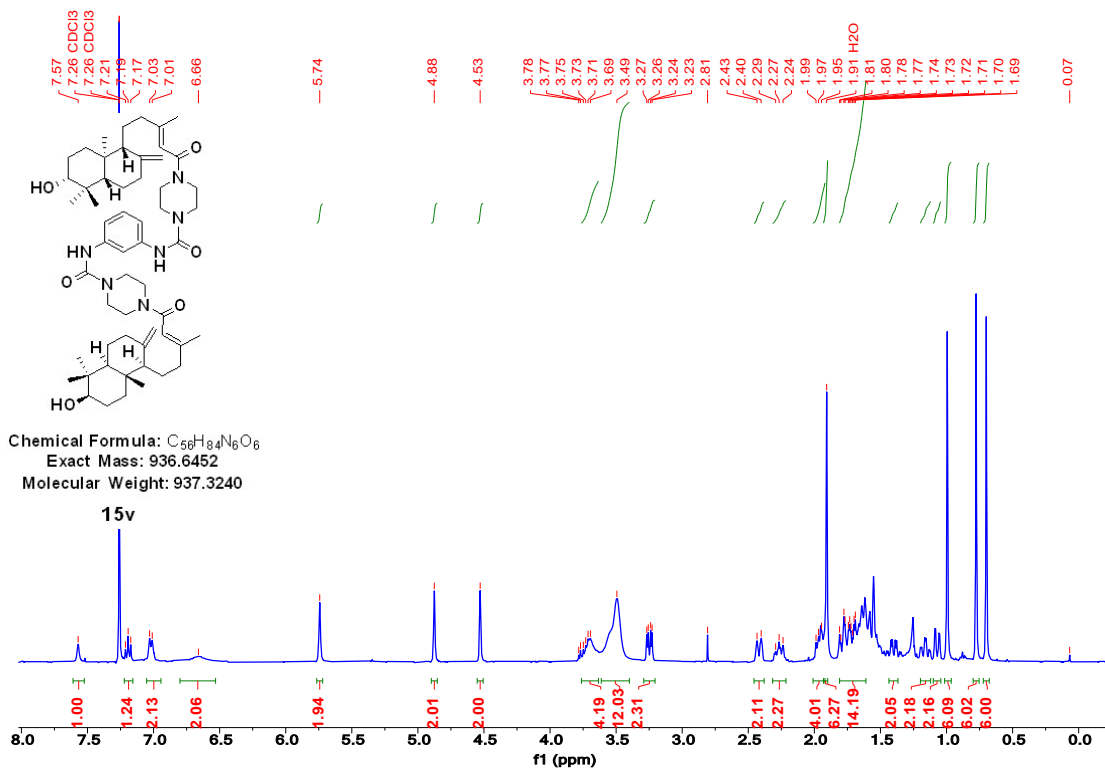


Figure S115. ^1H NMR spectrum of 15v

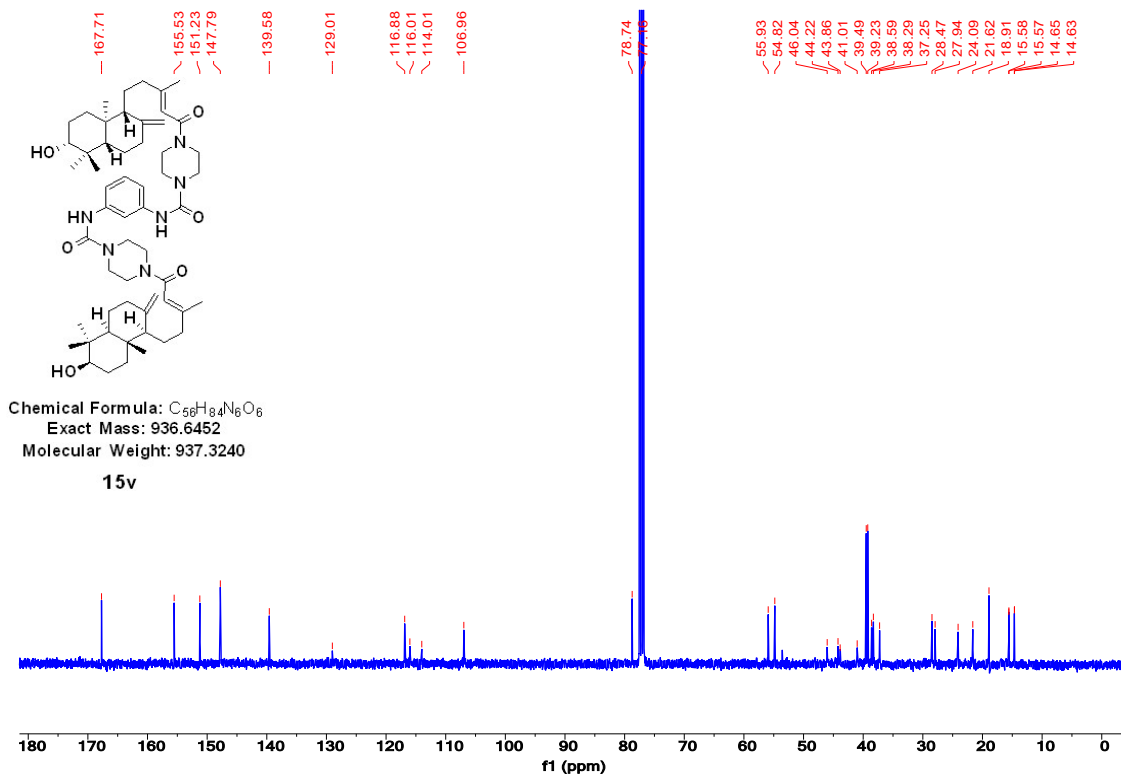


Figure S116. ^{13}C NMR spectrum of 15v

Single Mass Analysis

Tolerance = 5.0 mDa / DBE: min = -1.5, max = 50.0

Element prediction: Off

Number of isotope peaks used for i-FIT = 3

Monoisotopic Mass, Even Electron Ions

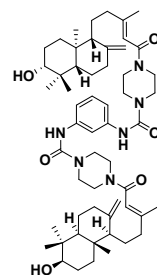
7 formula(e) evaluated with 1 results within limits (up to 50 best isotopic matches for each mass)

Elements Used:

C: 29-56 H: 47-85 N: 2-6 O: 2-6

7

250310-2-Q-18 36 (0.094)



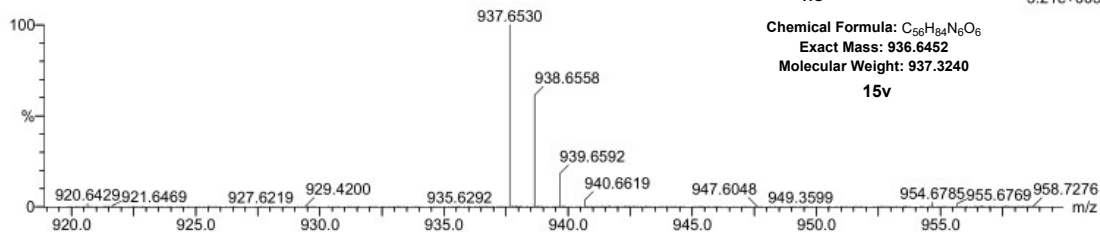
1: TOF MS ES+
3.21e+005

Chemical Formula: C₅₆H₈₄N₆O₆

Exact Mass: 936.6452

Molecular Weight: 937.3240

15v



Minimum: -1.5
Maximum: 5.0 10.0 50.0

Mass	Calc. Mass	mDa	PPM	DBE	i-FIT	Norm	Conf (%)	Formula
937.6530	937.6531	-0.1	-0.1	17.5	405.7	n/a	n/a	C ₅₆ H ₈₅ N ₆ O ₆

Figure S117. HRMS spectrum of 15v

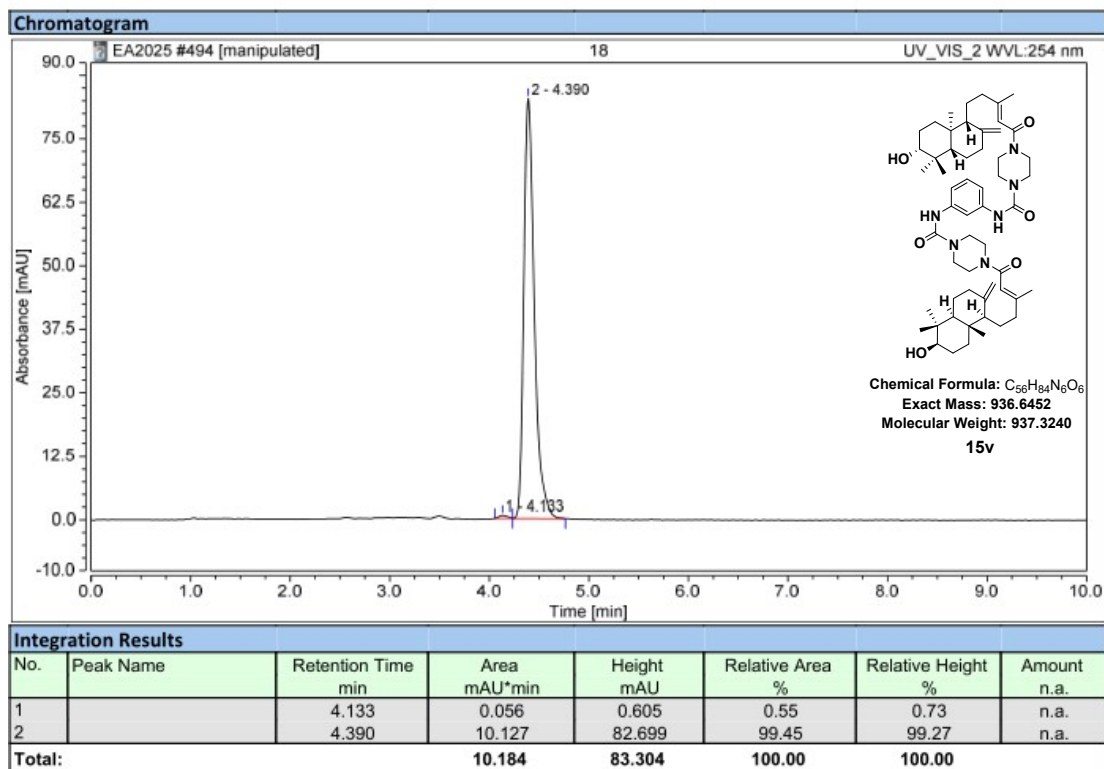


Figure S118. HPLC spectrum of 15v

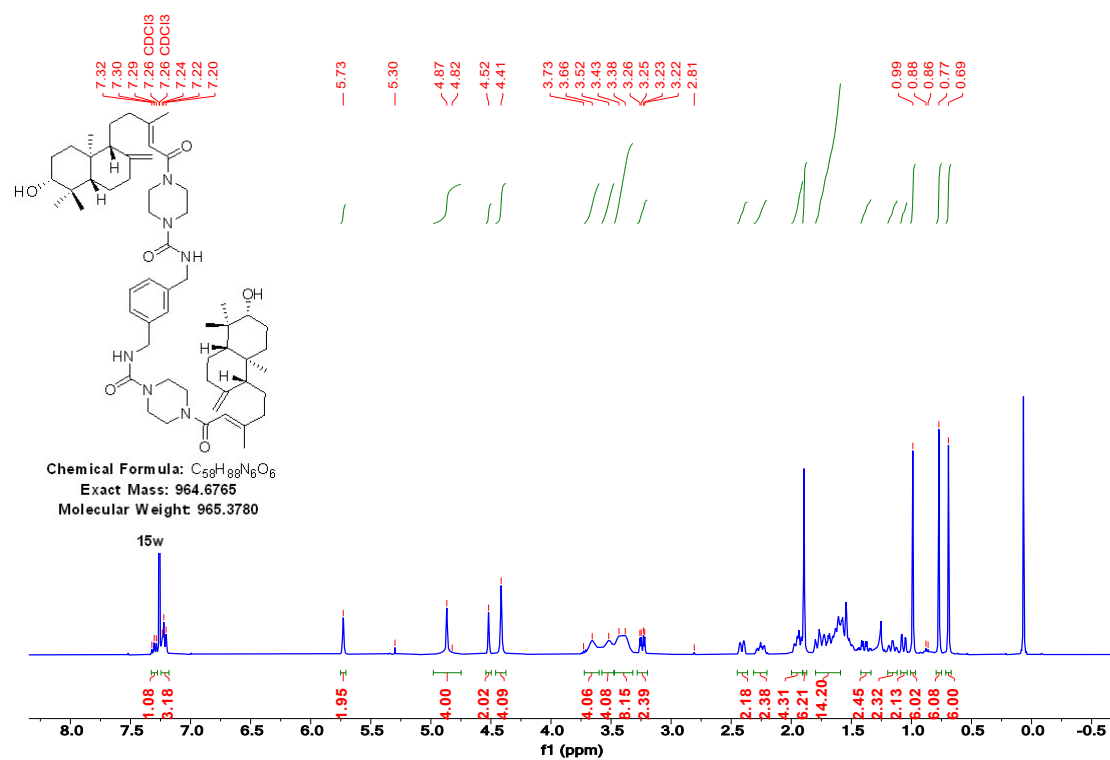


Figure S119. ¹H NMR spectrum of 15w

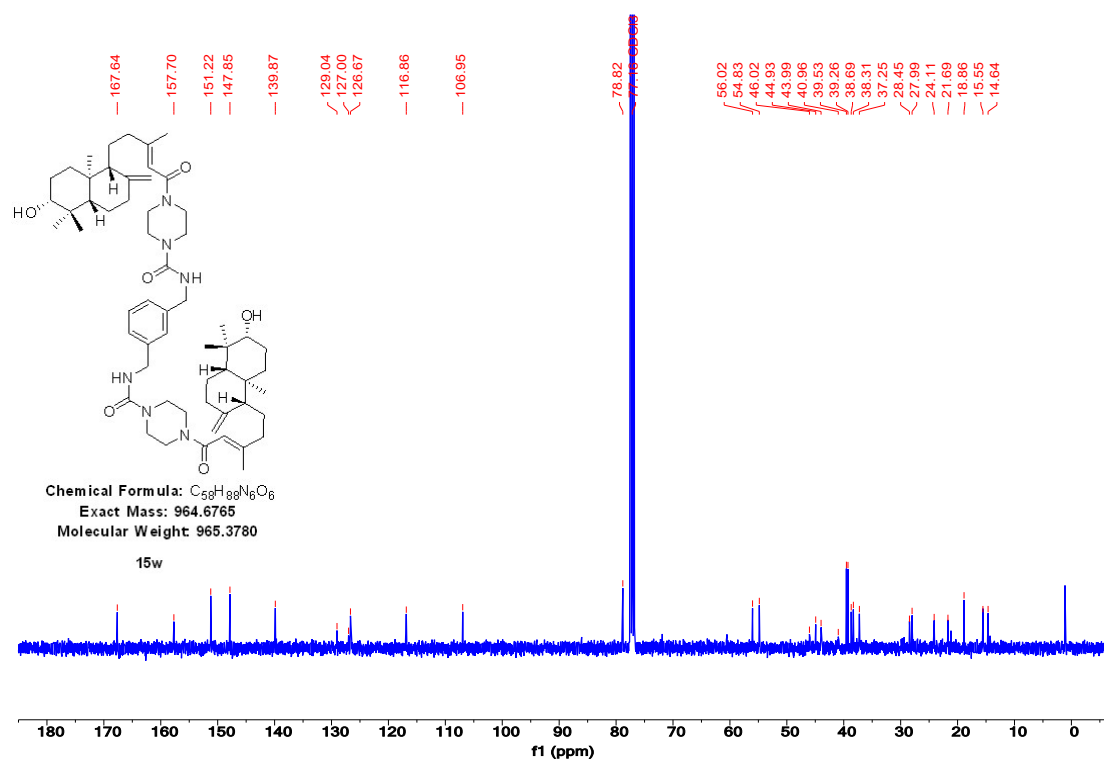


Figure S120. ¹³C NMR spectrum of 15w

Single Mass Analysis

Tolerance = 5.0 mDa / DBE: min = -1.5, max = 50.0

Element prediction: Off

Number of isotope peaks used for i-FIT = 3

Monoisotopic Mass, Even Electron Ions

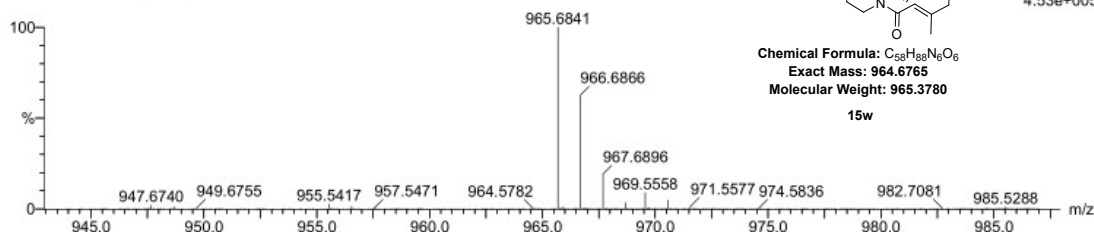
7 formula(e) evaluated with 1 results within limits (up to 50 best isotopic matches for each mass)

Elements Used:

C: 29-58 H: 44-89 N: 2-6 O: 2-6

7

250310-2-Q-23 29 (0.080)



Minimum: -1.5
Maximum: 50.0

Mass	Calc. Mass	mDa	PPM	DBE	i-FIT	Norm	Conf(%)	Formula
965.6841	965.6844	-0.3	-0.3	17.5	477.1	n/a	n/a	C58 H89 N6 O6

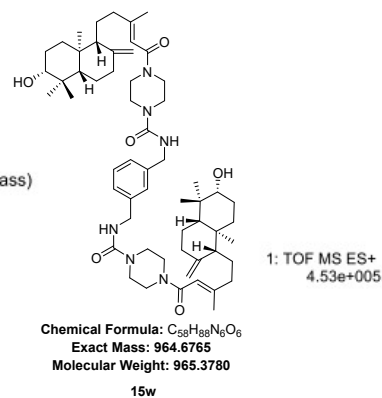
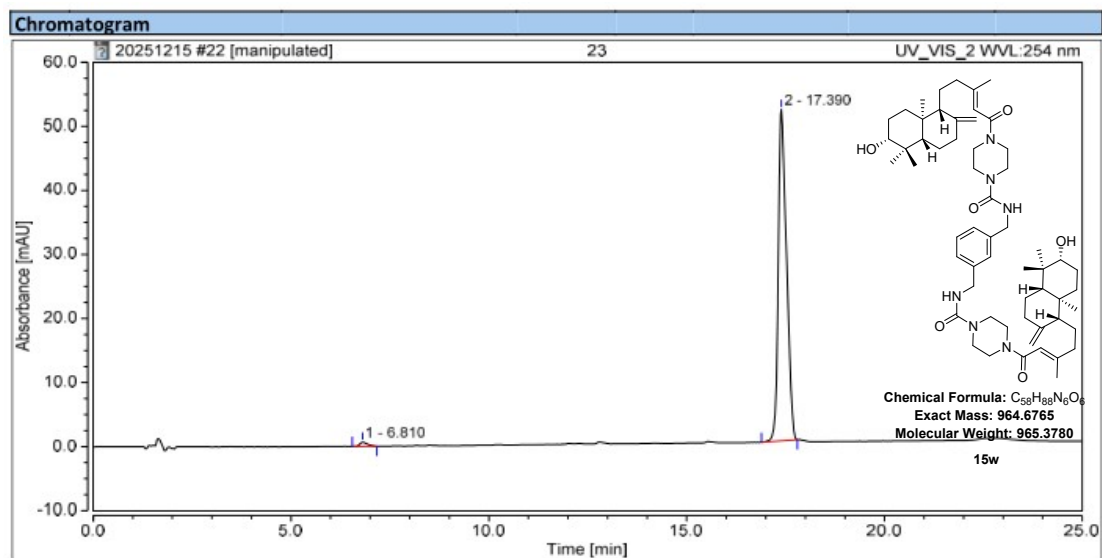


Figure S121. HRMS spectrum of 15w



Integration Results							
No.	Peak Name	Retention Time min	Area mAU*min	Height mAU	Relative Area %	Relative Height %	Amount
1		6.810	0.144	0.658	1.15	1.25	n.a.
2		17.390	12.384	51.772	98.85	98.75	n.a.
Total:			12.528	52.429	100.00	100.00	

Figure S122. HPLC spectrum of 15w

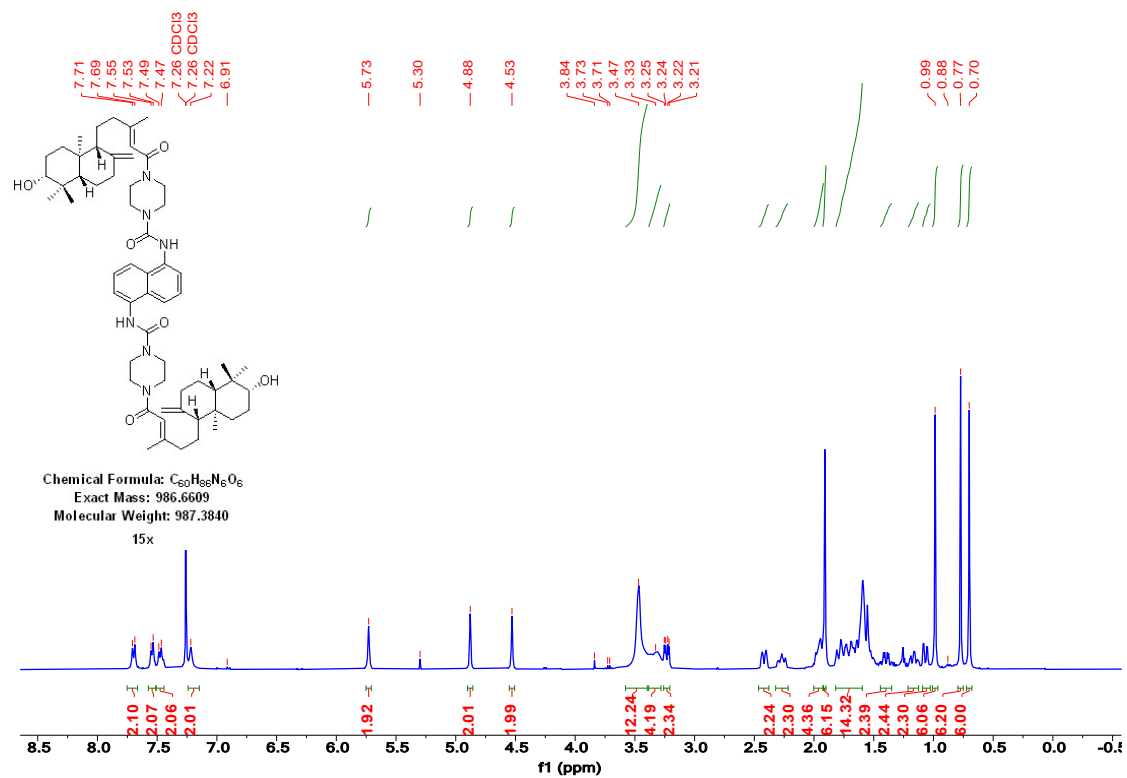


Figure S123. 1H NMR spectrum of 15x

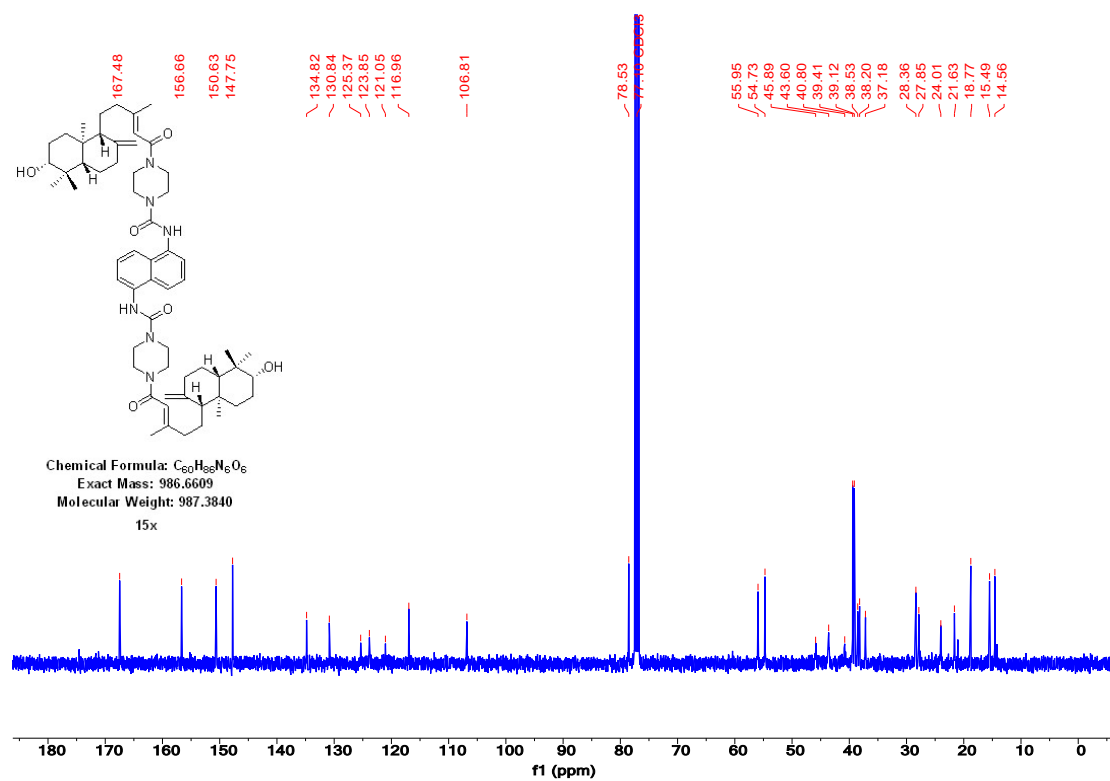


Figure S124. ^{13}C NMR spectrum of 15x

Single Mass Analysis

Tolerance = 5.0 PPM / DBE: min = -1.5, max = 50.0

Element prediction: Off

Number of isotope peaks used for i-FIT = 3

Monoisotopic Mass, Even Electron Ions

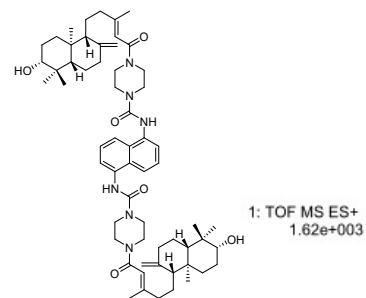
808 formula(e) evaluated with 1 results within limits (all results (up to 1000) for each mass)

Elements Used:

C: 60-60 H: 16-90 N: 1-6 O: 1-20 Na: 1-8

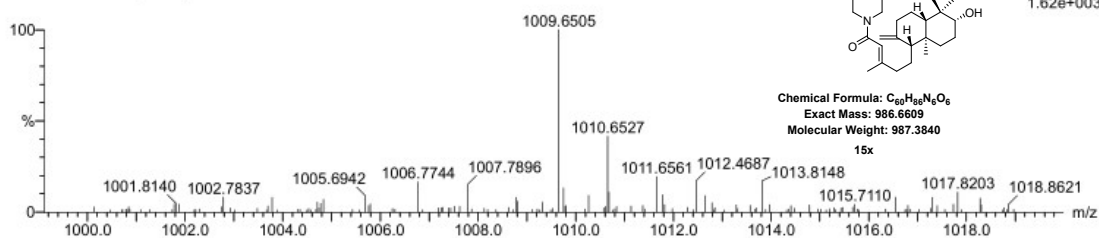
7

250310-2-Q-37 89 (0.203)



Chemical Formula: C₆₀H₈₆N₆O₆
Exact Mass: 986.6609
Molecular Weight: 987.3840

15x

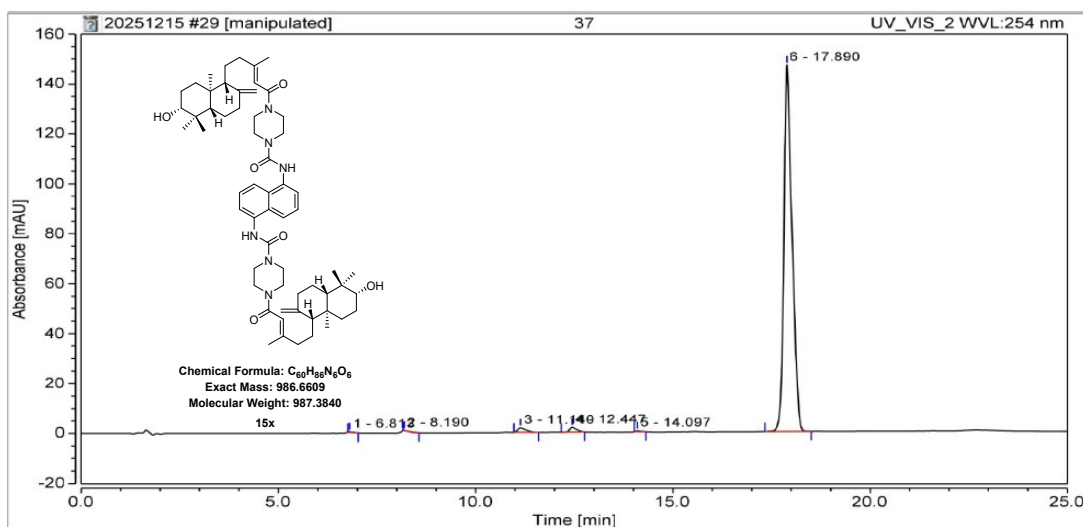


Minimum: -1.5
Maximum: 50.0

Mass	Calc. Mass	mDa	PPM	DBE	i-FIT	Norm	Conf (%)	Formula
1009.6505	1009.6507	-0.2	-0.2	20.5	151.0	n/a	n/a	C60 H86 N6 O6 Na

Figure S125. HRMS spectrum of 15x

Chromatogram



Integration Results

No.	Peak Name	Retention Time min	Area mAU*min	Height mAU	Relative Area %	Relative Height %	Amount
1		6.813	0.020	0.166	0.05	0.11	n.a.
2		8.190	0.017	0.146	0.05	0.10	n.a.
3		11.140	0.429	1.797	1.18	1.19	n.a.
4		12.447	0.394	1.956	1.09	1.29	n.a.
5		14.097	0.054	0.347	0.15	0.23	n.a.
6		17.890	35.374	146.782	97.48	97.08	n.a.
Total:			36.288	151.194	100.00	100.00	

Figure S126. HPLC spectrum of 15x