

# Supporting Information

## Chemical Genetic Control of Protein Levels: Selective *in vivo* Targeted Degradation

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Hospital, David Geffen School of Medicine at UCLA, Gwynn Hazen Cherry Memorial  
Laboratories, Molecular Biology Institute, and Jonsson Comprehensive Cancer Center,  
Los Angeles, California 90095-1752.

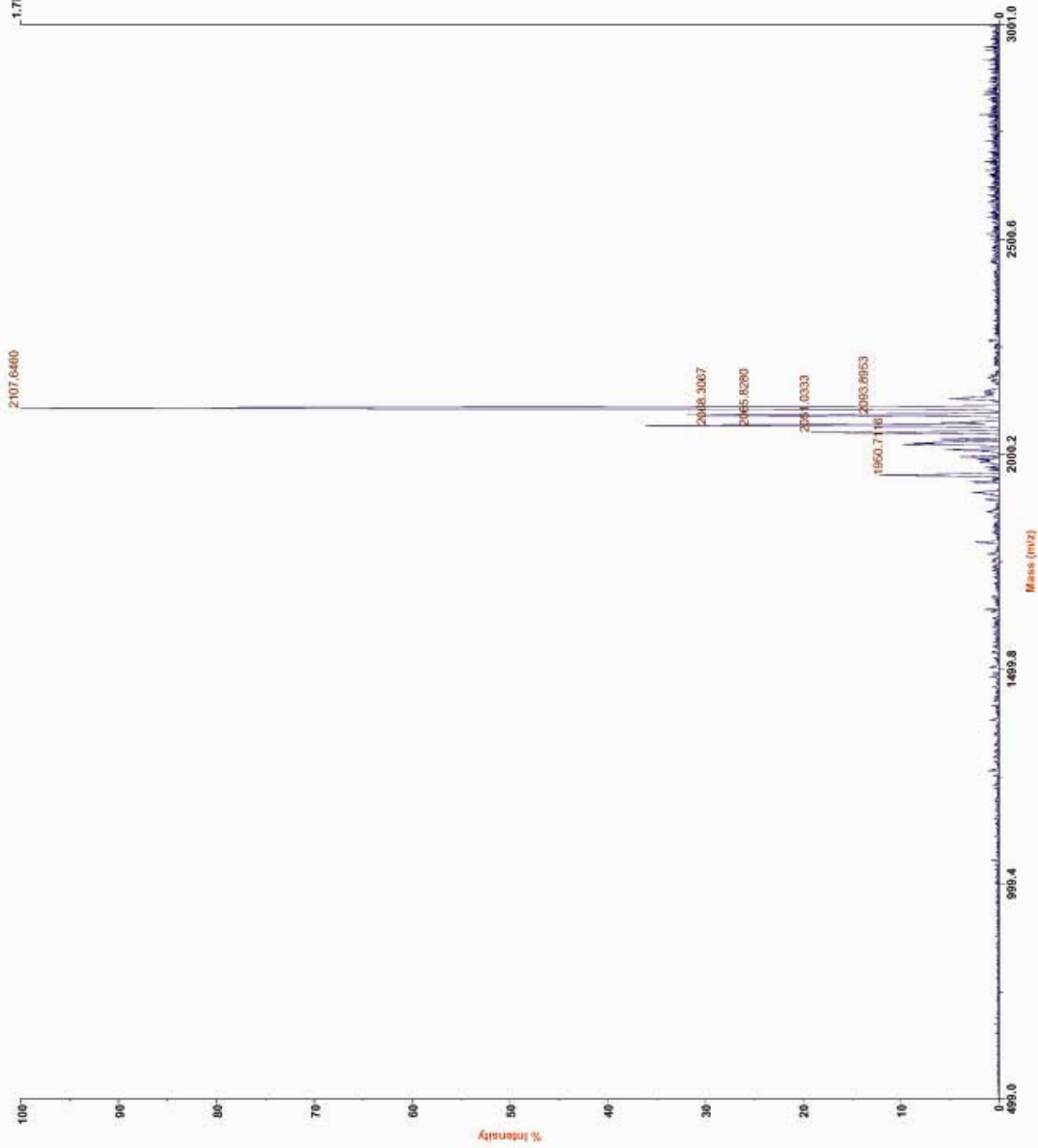
### **Purification of HIF-polyarginine Peptide (HIF1)**

HIF1 (H<sub>2</sub>N-(CH<sub>2</sub>)<sub>5</sub>CO-ALAPYIP-(D-Arg)<sub>8</sub>NH<sub>2</sub>) was purified by RP-HPLC (Rainin  
Dynamax System) comprising two solvent delivery pumps (Model SD200) and variable  
wavelength detector (model UV-1) set at 214nm. The column used was an YMC-Pack

ODS-AM, 250 x 20 mm, 5 $\mu$ m particle size and 120 $\text{\AA}$  pore size (Waters, Milford, MA) coupled to a guard column ODSA (10 x 10mm) with the same stationary phase specifications. The separation was carried out with a linear gradient of solvent A (H<sub>2</sub>O/CH<sub>3</sub>CN 98:2 + 0.06% of TFA) into solvent B (CH<sub>3</sub>CN/H<sub>2</sub>O 80:20 + 0.05% of TFA): 15% of B to 35% of B over 85min, 35% B to 98% B over 15 min, run at 5.0ml/min. HIF1 was eluted at 69 min. MALDI-TOF analysis of peptide samples were performed in a Voyager-DE- PRO 6268 (Applied Biosystems) using a-cyano-4-hydroxycinnamic acid matrices. Analytical HPLC traces of purified samples was carried out in a Waters Separation Module 2795 coupled to a Waters 2795 Photodiode Array Detector (set at 214nm) and to a Micromass ZQ 4000 Electrospray Mass detector (cone voltage = +120V). A linear gradient of Solvent A (composition as described for preparative separations) into Solvent B, 2% B to 37.5% B over 63min, was run at 0.2ml/min. The analytical column used was an XTerra MS C18 (4.6mm x 50mm, 2.5 $\mu$ m particle size, 80 $\text{\AA}$  pore size).

# Applied Biosystems Voyager System 6268

Voyager Spec #1=>NF1,0=>AdvBC(32,0,5,0,1)[BP = 2107.7, 16505]



Mode of operation: Reflector  
 Extraction mode: Delayed  
 Polarity: Positive  
 Acquisition control: Manual

Accelerating voltage: 20000 V  
 Grid voltage: 94%  
 Mirror voltage ratio: 1.12  
 Guide wire 0: 0.0025%  
 Extraction delay time: 185 nsec

Acquisition mass range: 500 - 3000 Da  
 Number of laser shots: 200/spectrum  
 Laser intensity: 2227  
 Laser Rep Rate: 20.0 Hz  
 Calibration type: Default  
 Calibration matrix: a-Cyano-4-hydroxycinnamic acid  
 Low mass gate: 500 Da  
 Timed ion selector: Off

Digitizer start time: 22.699  
 Bin size: 0.5 nsec  
 Number of data points: 65433  
 Vertical scale 0: 500 mV  
 Vertical offset: 1.5%  
 Input bandwidth 0: 500 MHz

Sample well: 15  
 Plate ID: 100 WELL PLATE  
 Serial number: 6268  
 Instrument name: Voyager-DE PRO  
 Plate type filename: C:\VOYAGER\100 well plate.plt  
 Lab name: Yale University

Absolute x-position: 22624.3  
 Absolute y-position: 41422  
 Relative x-position: 716.77  
 Relative y-position: -805.514  
 Shots in spectrum: 200  
 Source pressure: 3.218e-007  
 Mirror pressure: 6.037e-008  
 TC2 pressure: 0.001  
 TIS gate width: 8  
 TIS flight length: 690.5

Figure 1: MALDI-TOF spectrum of purified HIF 1 sample used on the synthesis of (3).

Analytical conditions are described above.

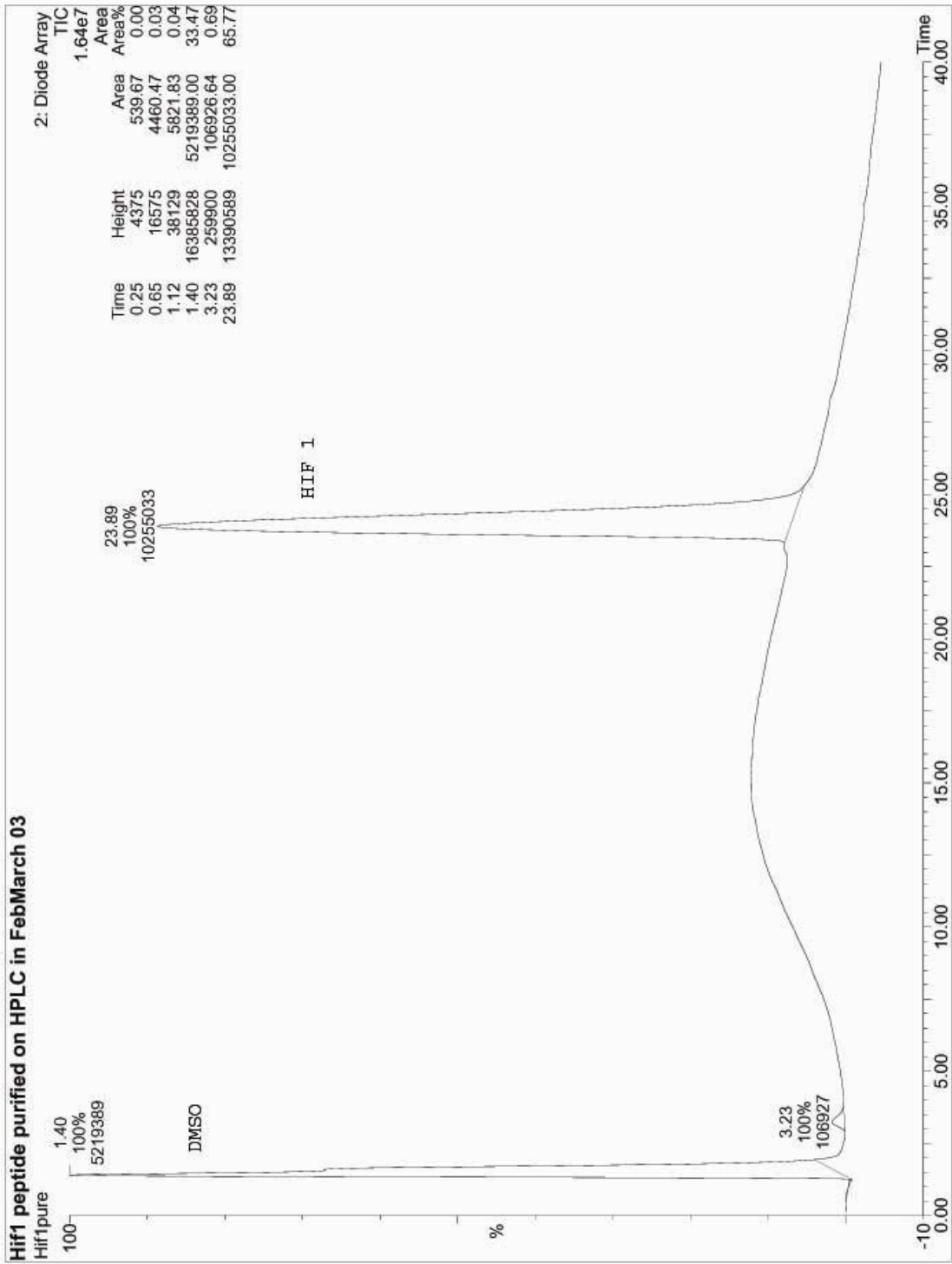


Figure 2: HPLC chromatogram (UV trace) of purified HIF 1 used in the synthesis of Protac-2. Analytical conditions are described above.

### Preparation of PROTAC-4 (3).

AP21998 (**1**) was synthesized as previously described<sup>21,22</sup>, as a 1:1 mixture of diastereomers at C9. To a solution of 15 mg (0.021mmol) of **1** in 1mL DMF was added 6.8mg (0.023mmol) EDCI, 6.4mg (0.053mmol) DMAP, and 4.6mg (0.021mmol) H<sub>2</sub>N-(CH<sub>2</sub>)<sub>5</sub>CO<sub>2</sub>Bn. The reaction was stirred for 24h, and filtered through Celite filter agent. Solvent was removed under vacuum, and the crude product (16mg) was redissolved in 2mL ethanol. To this solution was added 20mg 10% Pd/C, the reaction was placed under an atmosphere of hydrogen (balloon) and stirred for 1h. The reaction was filtered through Celite, after which the solvent was removed *in vacuo* to yield 12mg of crude product (**2**). This material was carried on without further purification.

Crude **2** (12mg, 0.014mmol) was added to a 5mL round bottomed flask along with 3mg (0.0014mmol) HIF1 H<sub>2</sub>N-(CH<sub>2</sub>)<sub>5</sub>CO-ALAPYIP-(D-Arg)<sub>8</sub>NH<sub>2</sub> (W.M. Keck Foundation Biotechnology Resource Laboratory), 1mg PyBrOP, 1μL diisopropylethylamine, and 0.6mL dry DMF. The reaction mixture was stirred for 18 hours, after which solvent was evaporated under a high vacuum.

Crude mixture of **3** was purified by RP-HPLC (for HPLC system and column specifications see the purification procedure for HIF 1). A linear gradient of Solvent A into B (same solvents composition as described for HIF 1) consisting of 2% B to 37% B over 63 min, 37% B to 75% B over 37 min, 75% B to 98% B over 10 min, 98% B for 20

min, was run at 5.0 mL/min. 3 was eluted at 95 min. MALDI-TOF analyses of purified samples were performed in a Voyager-DE- PRO 6268 (Applied Biosystems) using acyano-4-hydroxycinnamic acid matrices. Purified samples were analyzed by LC-MS (see HIF 1 for equipment, column specifications and solvent composition and analytical parameters) using a linear gradient of B into A: 2% B to 100% B over 40 min, 100% B for 20 min, run at 0.2 mL/min.

**Compound 1.**  $^1\text{H}$  NMR ( $\text{CDCl}_3$ , 400 MHz) (Mixture of rotamers and diastereomers)

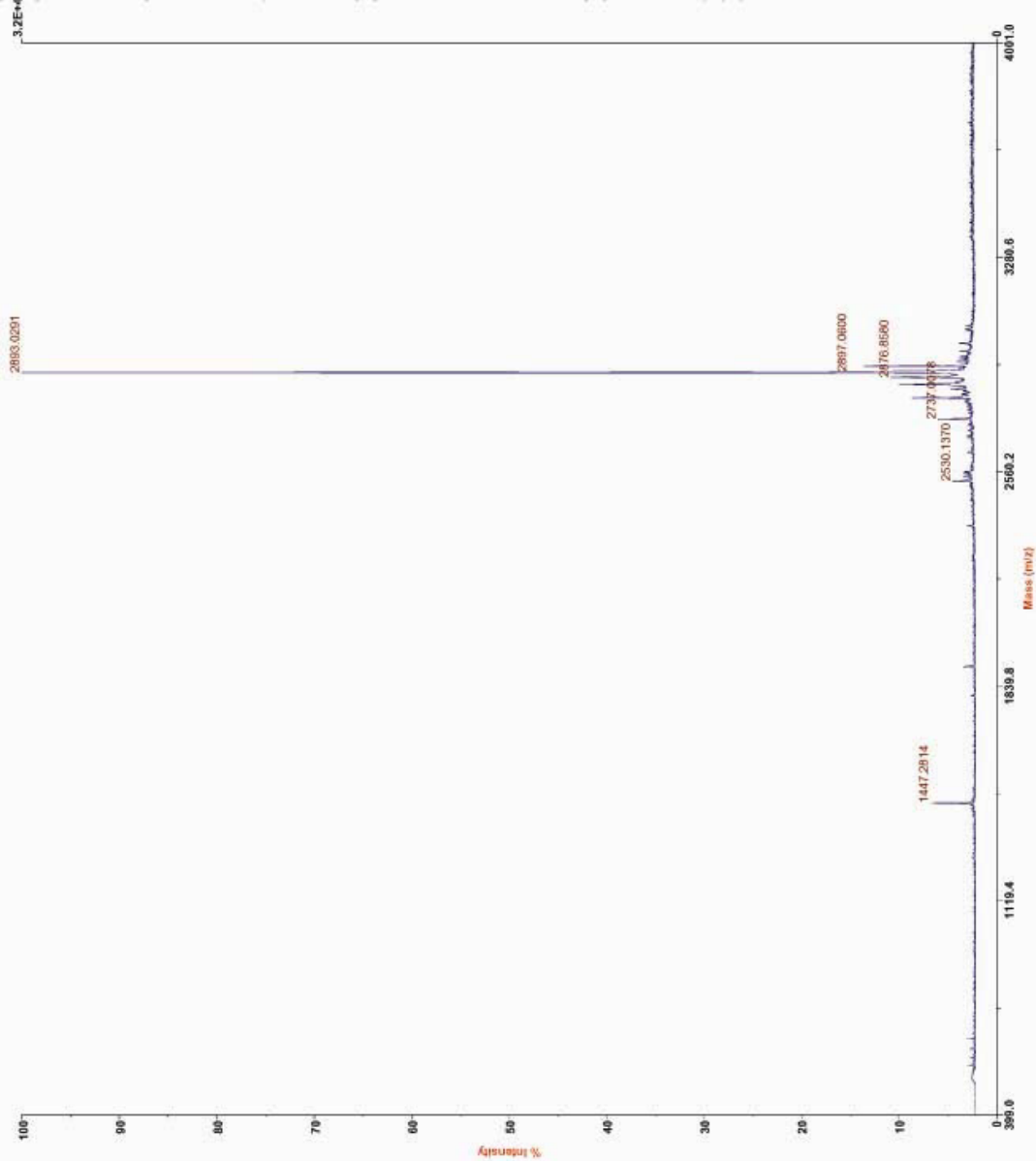
**1(S)** 7.19 (t,  $J=7.9\text{Hz}$ , 1H), 6.67-6.96 (m, 6H) 6.24 (s, 2H), 5.48-5.50 (m, 2H) 4.60-4.74 (m, 2H), 3.86 (s, 3H), 3.85 (s, 3H), 3.78 (s, 3H), 3.61-3.93 (m, 2H), 3.55 (s, 6H), 2.85-2.93 (m, 1H), 2.56-2.68 (m, 2H), 1.98-2.36 (m, 4H), 1.63-1.77 (m, 4H), 1.26-1.46 (m, 2H), 0.90 (t,  $J=7.3\text{Hz}$ , 3H). **1(R)** 7.25 (t,  $J=7.8\text{ Hz}$ ), 6.70-6.90 (m, 6H), 6.45 (s, 2H), 5.66-5.70 (m, 1H), 5.53 (d,  $J=4.5\text{Hz}$ , 1H), 4.59-4.72 (m, 2H), 3.86 (s, 3H), 3.85 (s, 3H), 3.84 (s, 3H), 3.83 (s, 6H), 3.47-3.79 (m, 2H), 3.23-3.32 (m, 1H), 2.52-2.71 (m, 2H), 2.00-2.31 (m, 4H), 1.52-1.75 (m, 4H), 1.28-1.46 (m 2H), 0.83 (t,  $J=7.3\text{Hz}$ , 3H); LRMS (ES) (M + H)<sup>+</sup> calc 694.32, meas. 694.7.

Structural assignment of **1** based on original synthesis: Yang, W., et. al. *J. Med. Chem.*

**2000**, *43*, 1135-1142.

# Applied Biosystems Voyager System 6268

Voyager Spec #1[BP = 2893.6, 32072]



Mode of operation: Reflector  
Extraction mode: Delayed  
Polarity: Positive  
Acquisition control: Manual

Accelerating voltage: 20000 V  
Grid voltage: 73%  
Mirror voltage ratio: 1.12  
Guide wire 0: 0.0025%  
Extraction delay time: 185 nsec

Acquisition mass range: 400 -- 4000 Da  
Number of laser shots: 200/spectrum  
Laser intensity: 2027  
Laser Rep Rate: 20.0 Hz  
Calibration type: Default  
Calibration matrix: a-Cyano-4-hydroxycinnamic acid  
Low mass gate: 500 Da  
Timed ion selector: Off

Digitizer start time: 20.12  
Bin size: 0.5 nsec  
Number of data points: 86518  
Vertical scale 0: 500 mV  
Vertical offset: 1.5%  
Input bandwidth 0: 500 MHz

Sample well: 34  
Plate ID: 100 WELL PLATE  
Serial number: 6268  
Instrument name: Voyager-DE PRO  
Plate type filename: C:\VOYAGER\100 well plate.plt  
Lab name: Yale University

Absolute x-position: 17066.4  
Absolute y-position: 32985.1  
Relative x-position: 228.874  
Relative y-position: 917.581  
Shots in spectrum: 200  
Source pressure: 4.249e-007  
Mirror pressure: 9.542e-008  
TC2 pressure: 0.001  
TIS gate width: 8  
TIS flight length: 690.5

Acquired: 15:59:00, May 12, 2003

D:\data\Fabiana\Protac2L3\Purification\_0001.dat

Printed: 16:28, September 26, 2003

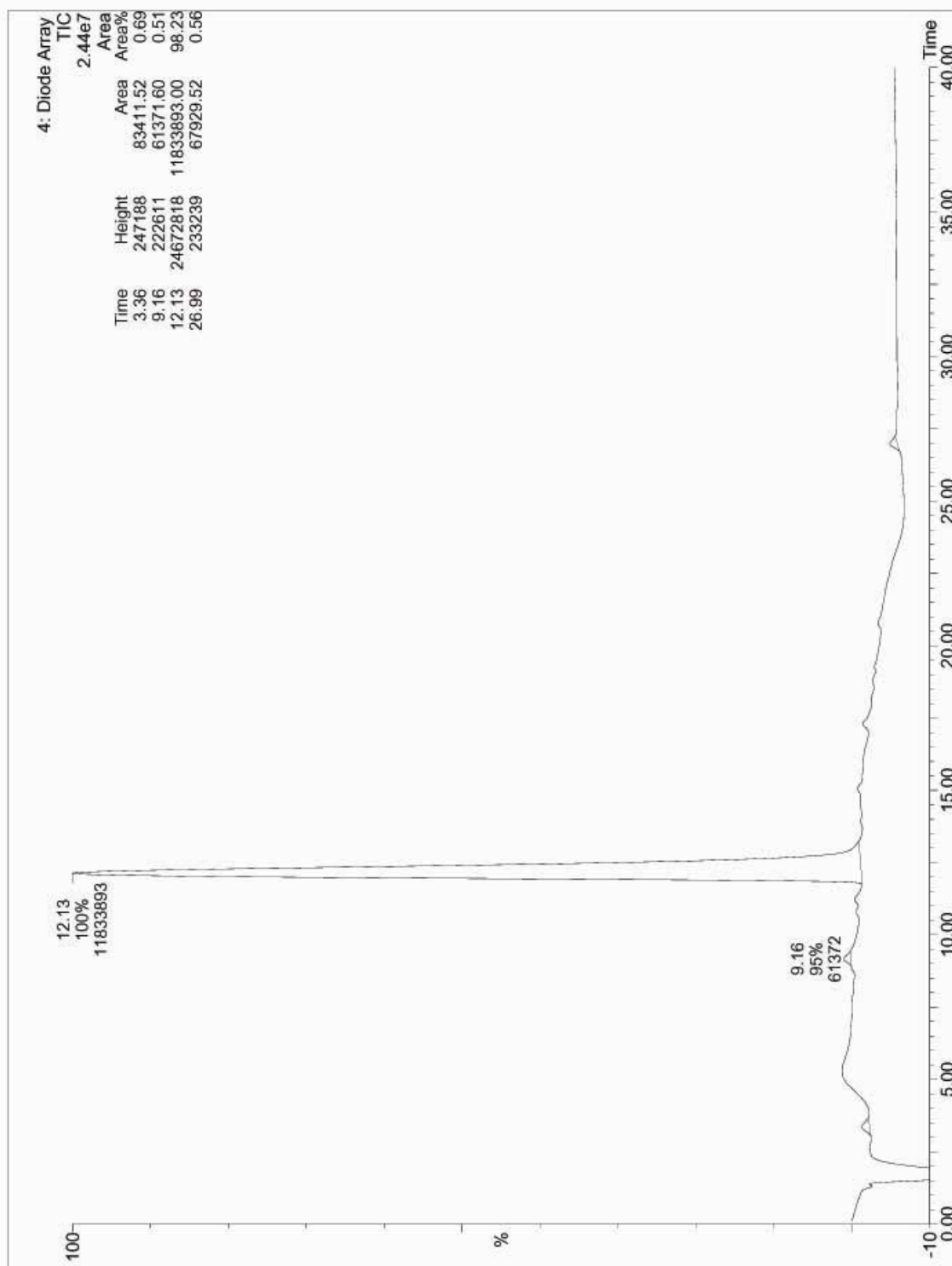


Figure 4: HPLC chromatogram (UV trace) of purified **3**. Analytical conditions described above.



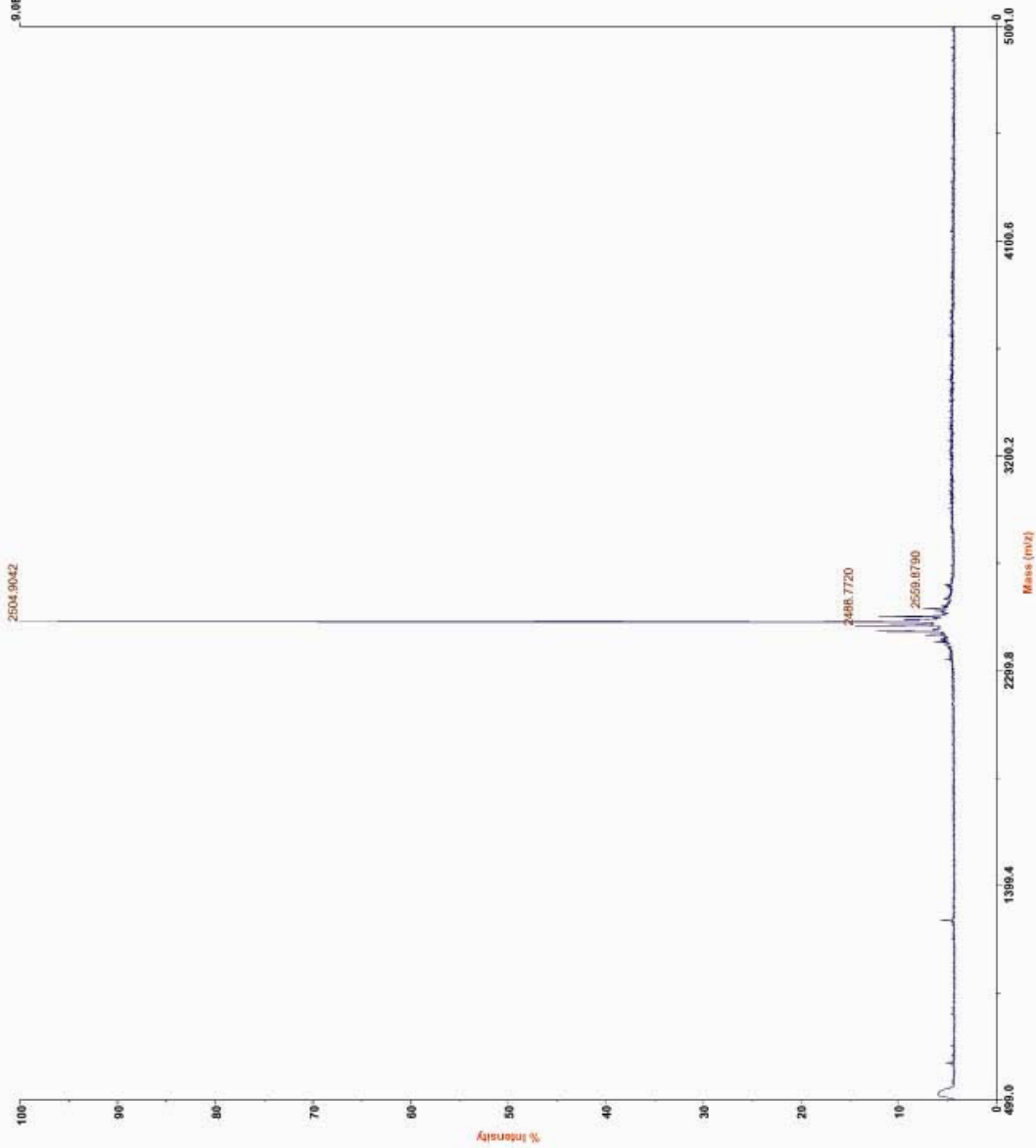
### **Preparation of PROTAC-5 (5).**

Compound **4** was synthesized as previously described<sup>33</sup>. To a solution of 1.2mg (0.003mmol) **4** in 1mL DMF was added 2mg  $\text{H}_2\text{N}-(\text{CH}_2)_5\text{CO-ALAPYIP}-(D\text{-Arg})_8\text{NH}_2$ , 0.3mg (0.014mmol) EDCI, and 0.2mg (0.015mmol) DMAP. The reaction mixture was stirred for 24h, after which solvent was evaporated under high vacuum. The product was purified by HPLC to yield 1.3mg **5**.

The purification procedure was carried out using same equipment, column and Solvents A and B composition as described for HIF 1 and PROTAC-4 (**3**). A linear gradient was done by adding Solvent B into Solvent A: 40%B to 70%B over 100min, 70%B to 98%B over 20min, run at 5.0mL/min. Compound **5** eluted at 30min. Purified samples were analyzed by MALDI-TOF using the same parameters as described earlier. Purified samples of **5** were analyzed by LC-MS using a linear gradient consisting of 2%B to 40%B over 40min. Analytical conditions, equipment and parameters were the same used for HIF 1 and **3**.

# Applied Biosystems Voyager System 6268

Voyager Spec #1[BP = 2504.9, 9019]



Mode of operation: Reflector  
Extraction mode: Delayed  
Polarity: Positive  
Acquisition control: Manual

Accelerating voltage: 20000 V  
Grid voltage: 73%  
Mirror voltage ratio: 1.12  
Guide wire 0: 0.0025%  
Extraction delay time: 180 nsec

Acquisition mass range: 500 – 5000 Da  
Number of laser shots: 600/spectrum  
Laser intensity: 2027  
Laser Rep Rate: 20.0 Hz  
Calibration type: Default  
Calibration matrix: *o*-Cyano-4-hydroxycinnamic acid  
Low mass gate: 500 Da  
Timed ion selector: Off

Digitizer start time: 22.4815  
Bin size: 0.5 nsec  
Number of data points: 96729  
Vertical scale 0: 500 mV  
Vertical offset: 1.5%  
Input bandwidth 0: 500 MHz

Sample well: 12  
Plate ID: 100 WELL PLATE  
Serial number: 6268  
Instrument name: Voyager-DE PRO  
Plate type filename: C:\VOYAGER\100 well plate.plt  
Lab name: Yale University

Absolute x-position: 6660.96  
Absolute y-position: 42228.7  
Relative x-position: -6.53802  
Relative y-position: 1.20874  
Shots in spectrum: 600  
Source pressure: 8.968e-007  
Mirror pressure: 4.637e-007  
TC2 pressure: 0.001  
TIS gate width: 8  
TIS flight length: 690.5

Acquired: 11:21:00, July 24, 2003

D:\data\Fabiana\Protac3L1\Protac3L3\_0001.dat

Printed: 16:29, September 26, 2003

Figure 5: MALDI-TOF spectrum of purified **5**. Analytical conditions are described above.

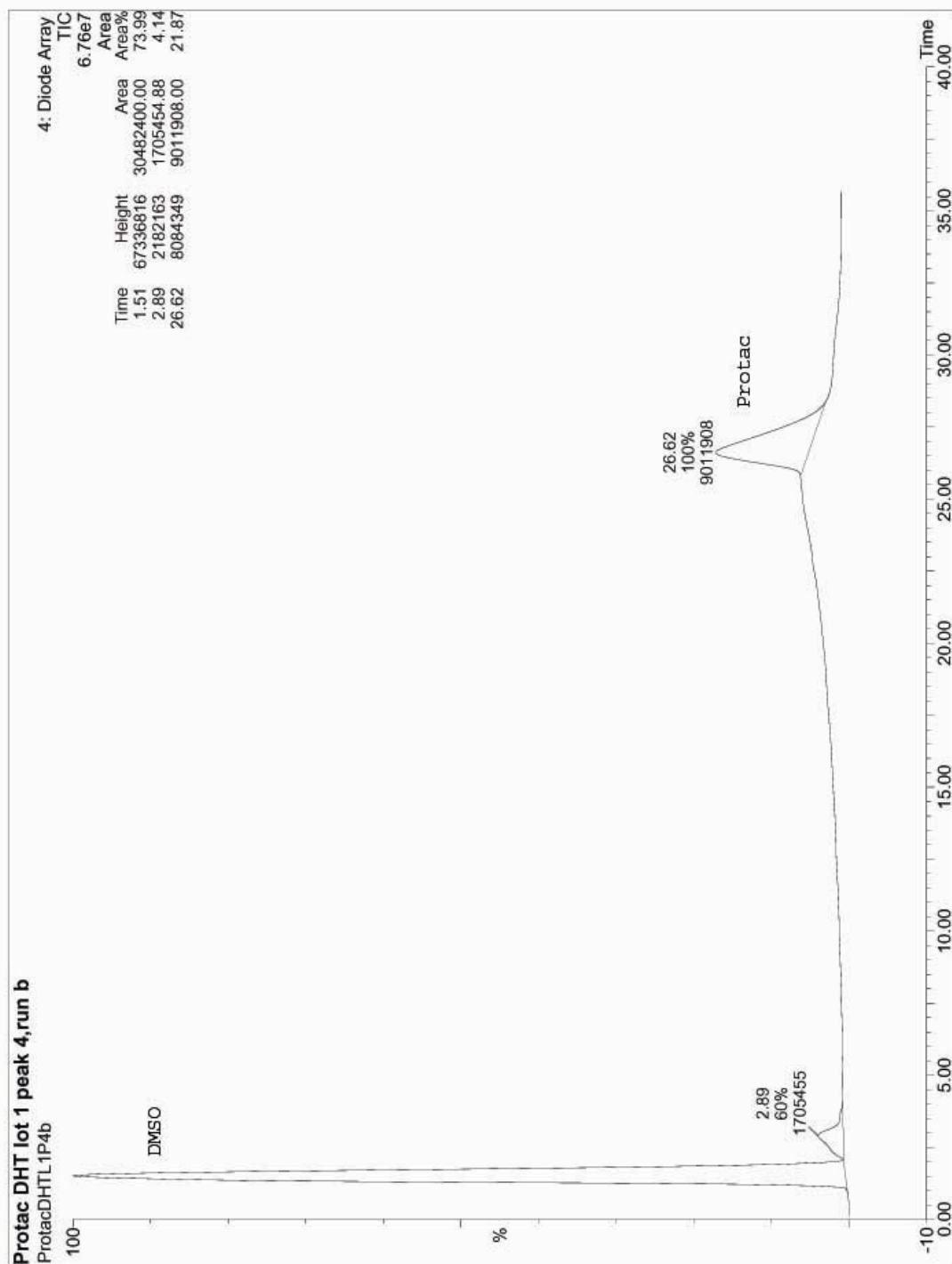


Figure 6: HPLC chromatogram (UV trace) of purified **5**. Analytical conditions are described above.