

Enhanced Cancer Theranostics with Self-Assembled, Multi-labeled siRNAs

Stephen D. Kozuch¹, Christopher N. Cultrara¹, Adah E. Beck¹, Claudia J. Heller¹, Sunil Shah¹,
Mayurbhai R. Patel², Jenny Zilberberg^{3†} and David Sabatino^{1†*}

¹Department of Chemistry and Biochemistry, Seton Hall University, South Orange, NJ 07079

²Nitto Denko AVECIA Inc, 8560 Reading Road, Cincinnati, OH 45215

³Department of Biomedical Research, Hackensack University Medical Center, Hackensack, NJ
07601

*Corresponding Author: David Sabatino: Email ID : david.sabatino@shu.edu

Tel : +1-973-313-6359

† Jenny Zilberberg and David Sabatino contributed equally to this work.

Supplementary Materials

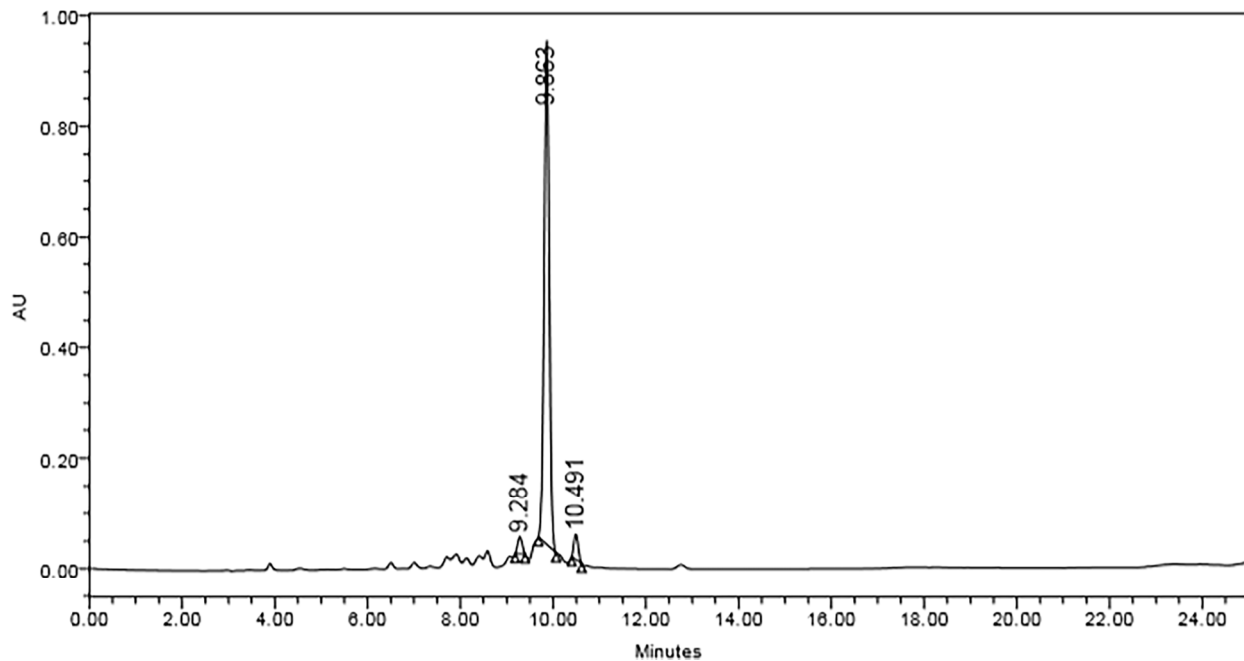
Within this supporting information:

- | | |
|--|-----|
| 1) RNA characterization table | S3 |
| 2) HPLC/ESI-MS traces of RNA sequences | S4 |
| 3) Ultraviolet-Visible spectroscopy of FITC-siRNA hybrids | S23 |
| 4) Fluorescent microscopy images of FITC-V-shape siRNA 4 vs 24 h | S25 |
| 5) Cell viability determined by propidium iodide staining | S26 |

Name	Sequence ^a	Percent Yield (%) ^b	Percent Purity (%) ^c	Mass ^d
GRP78A1	5'-AUC AGA AUC UUC CAA CAC U-3'	91	>99%	5949.6 (5949.9)
GRP78S1	5'-AGU GUU GGA AGA UUC UGA U-3'	92	>99%	6103.7 (6103.9)
GRP94S1	5'-GAA GAA GCA UCU GAU UAC C-3'	90	>99%	6068.8 (6069.0)
GRP75S1	5'-ACU GAC UCG GAG AAU ACA A-3'	91	>99%	6091.8 (6092.2)
V-78A194A1	2'-3'-CUU CUU CGU AGA CUA CUG G-5' rU 3'-5'-AUC AGA AUC UUC CAA CAC U-3'	83	>99%	12929.7 (12929.9)
V-78A178A2	2'-3'-CCU CGC GUA AUC AUG AUC U-5' rU 3'-5'-AUC AGA AUC UUC CAA CAC U-3'	81	>98%	12275.4 (12276.6)
Y-78A194A175A1	2'-3'-CUU CUU CGU AGA CUA CUG G-5' 5'-UUG UA UCU CCG AGU CAG U-3'-5'-rU 3'-5'-AUC AGA AUC UUC CAA CAC U-3'	62	>98%	18335.9 (18357.2)
FITC-78A1	FITC-NH-(CH ₂) ₆ -5'-AUC AGA AUC UUC CAA CAC U-3'	52	>97%	6516.7 (6518.5)
FITC-78S1	FITC-NH-(CH ₂) ₆ -5'-AGU GUU GGA AGA UUC UGA U-3'	48	>98%	6670.8 (6672.5)
FITC-94S1	FITC-NH-(CH ₂) ₆ -5'-GAA GAA GCA UCU GAU UAC C-3'	51	>97%	6635.9 (6636.8)
FITC-75S1*	FITC-NH-(CH ₂) ₆ -5'-ACU GAC UCG GAG AAU ACA A-3'	N/A	>99%	6629.3 (6628.8)
FITC-V-78A194A1	2'-3'-CUU CUU CGU AGA CUA CUG G-5'-(CH ₂) ₆ -NH-FITC rU 3'-5'-AUC AGA AUC UUC CAA CAC U-3'	22	>95%	13013.1 (12884.6)
FITC-V-78A178A2	2'-3'-CCU CGC GUA AUC AUG AUC U-5'-(CH ₂) ₆ -NH-FITC rU 3'-5'-AUC AGA AUC UUC CAA CAC U-3'	20	>95%	12969.1 (12843.9)

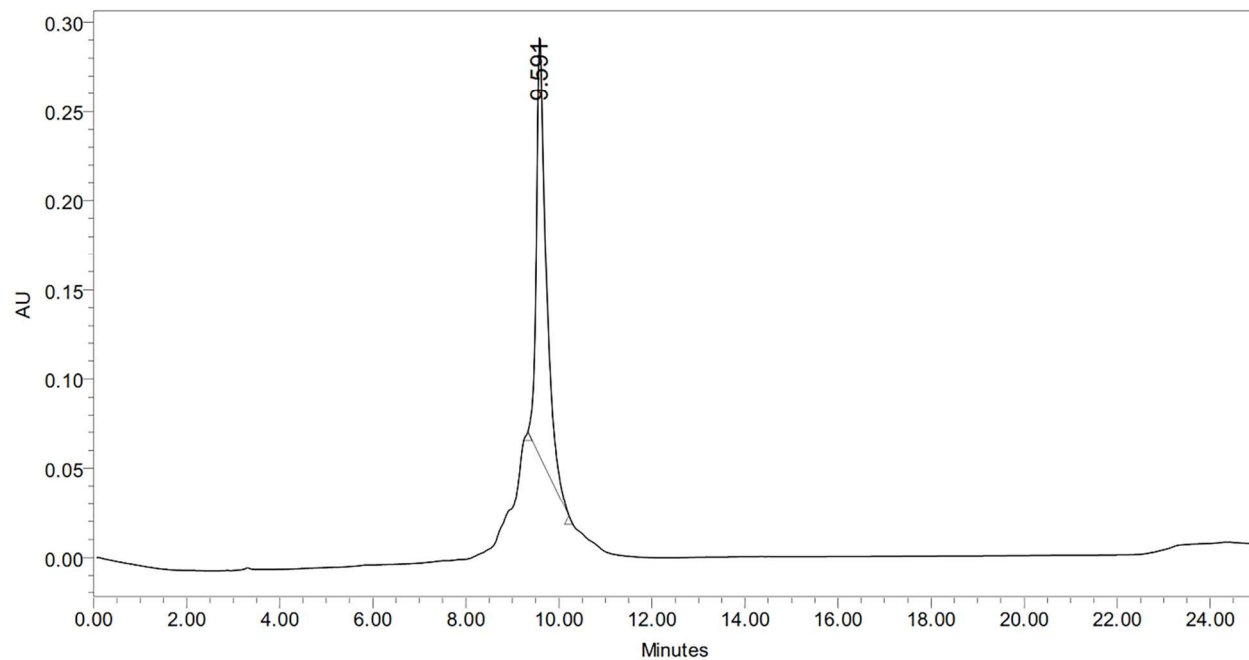
Table S1. ^aLinear, V- and Y-shaped RNA sequences designed to target GRP-78, 94 and 75. ^bDetermined by UV-Vis Spectroscopy. ^cObtained by RP-IP-HPLC using 0.1 mM TEAA in 0-45% MeCN, pH: 7.2 over 20 min. ^dCalculated mass (observed mass) by ESI-MS in negative mode (Novatia LLC, Newton, PA). * Sample purchased from ChemGenes (Wilmington, MA)

Figure S1 RP IP HPLC ANALYSIS OF LINEAR GRP78A1



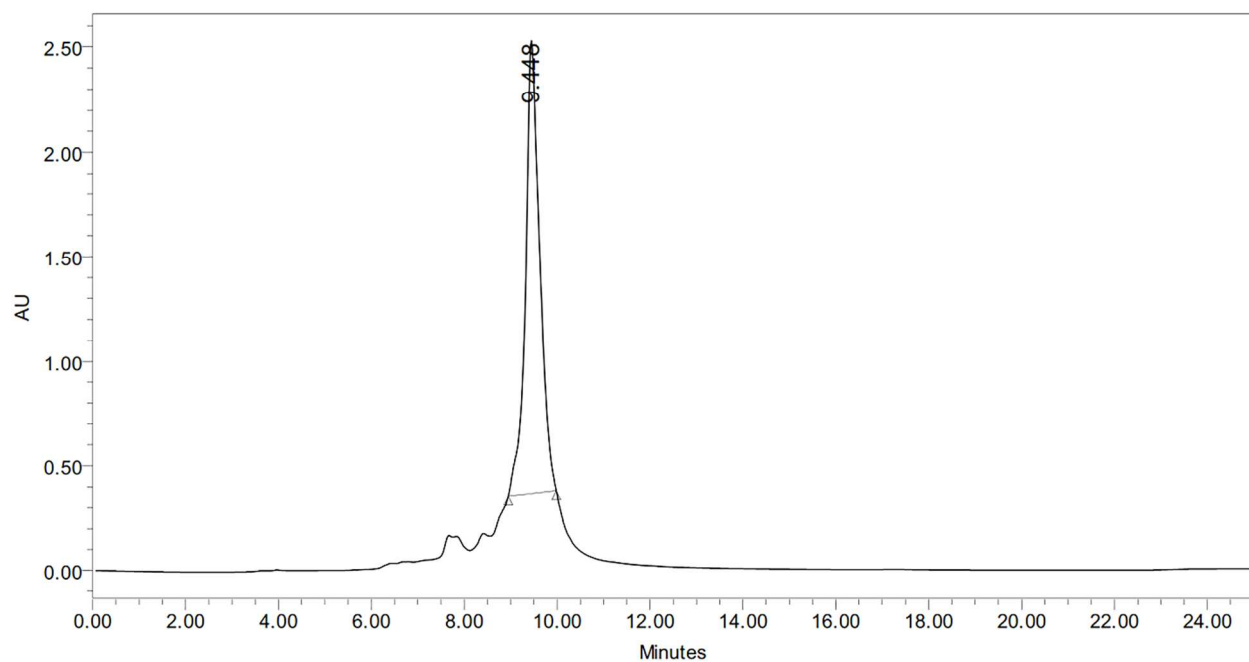
	RT	% Area	Area ($\mu\text{V}\cdot\text{sec}$)
1	9.28	2.78	211473
2	9.86	93.29	7107854
3	10.49	3.93	299732

Figure S2 RP IP HPLC ANALYSIS OF LINEAR GRP78S1



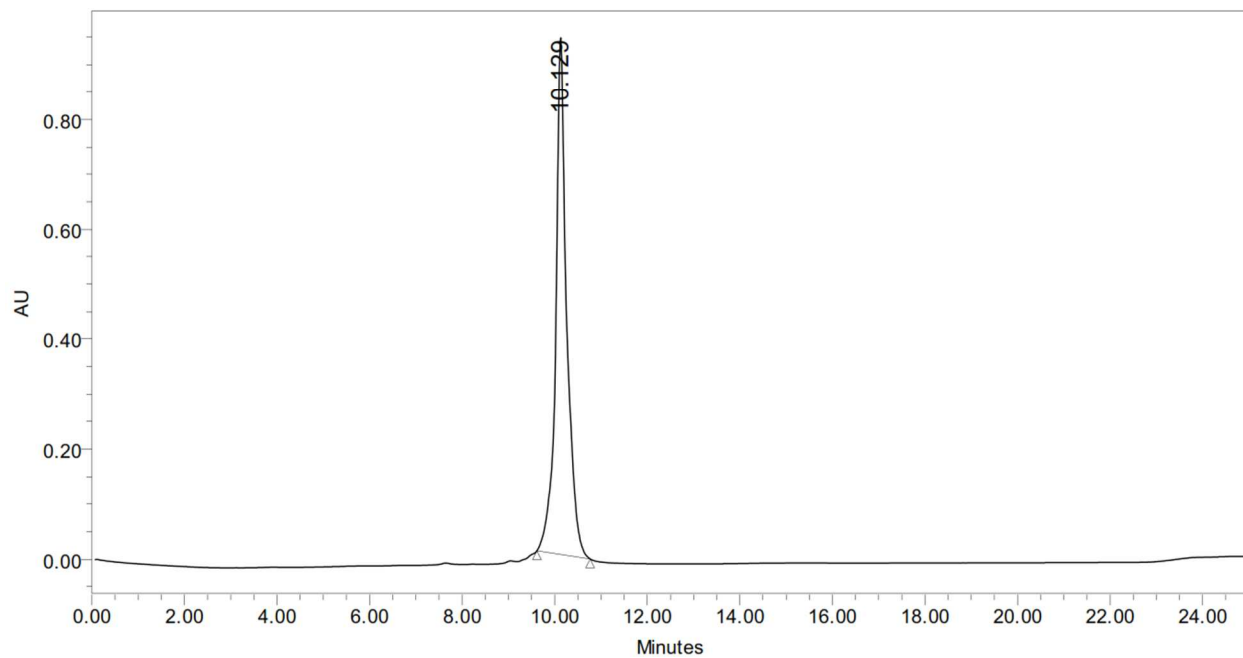
	RT	% Area	Area ($\mu\text{V}\cdot\text{sec}$)
1	9.59	100.00	3449874

Figure S3 RP IP HPLC ANALYSIS OF LINEAR GRP94S1



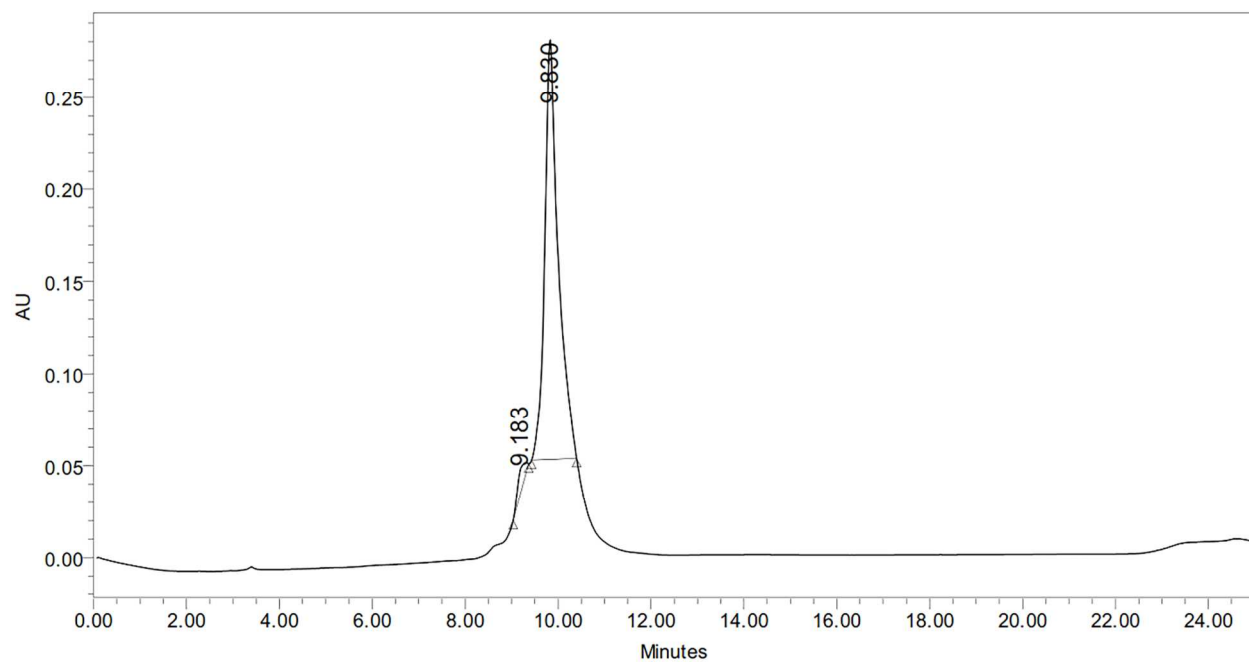
	RT	% Area	Area ($\mu\text{V}\cdot\text{sec}$)
1	9.45	100.00	45867536

Figure S4 RP IP HPLC ANALYSIS OF V-GRP78A194A1



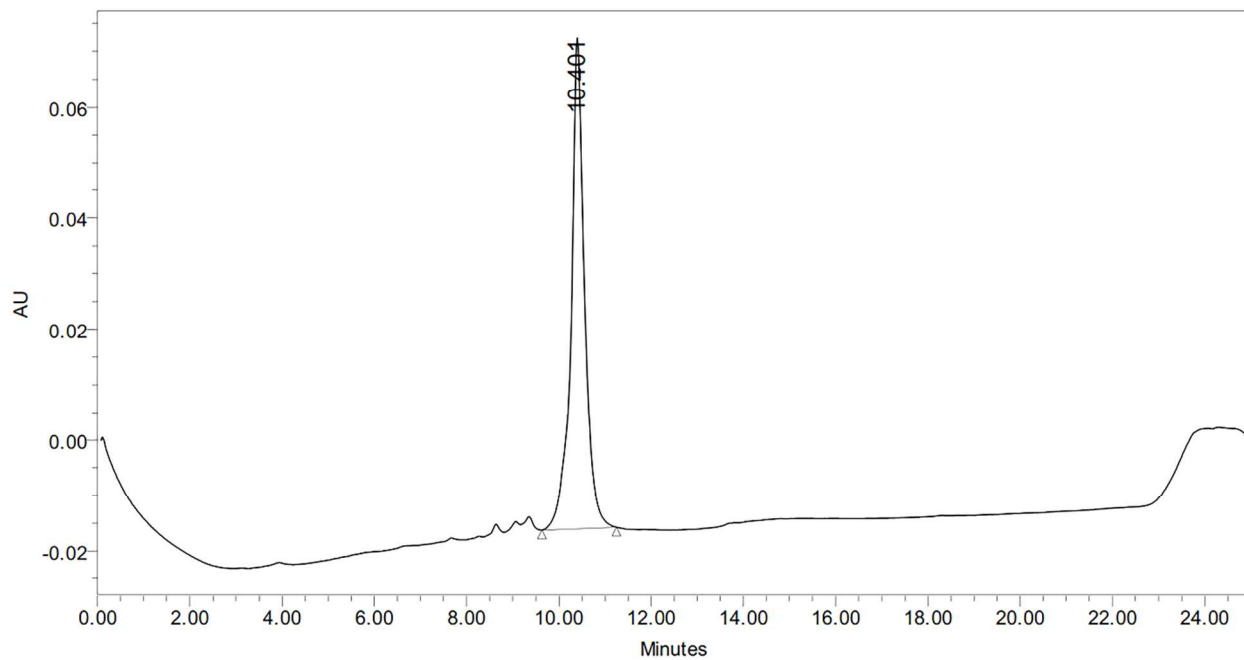
	RT	% Area	Area ($\mu\text{V}\cdot\text{sec}$)
1	10.13	100.00	15461777

Figure S5 RP IP HPLC ANALYSIS OF V-GRP78A178A2



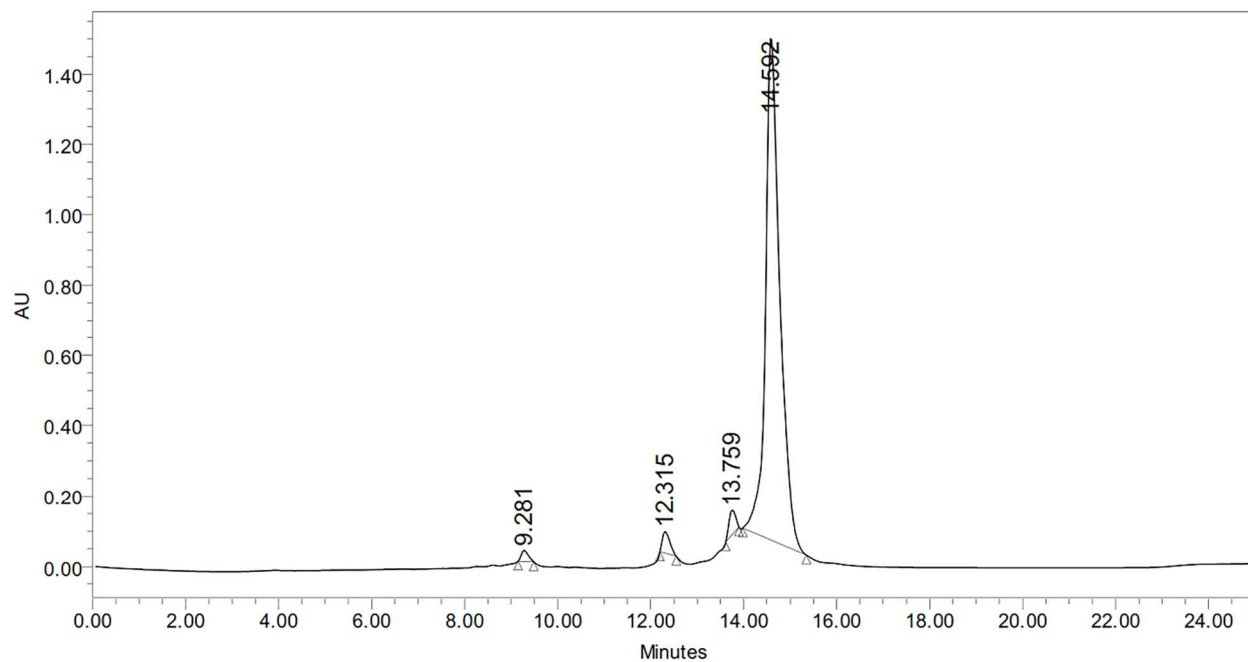
	RT	% Area	Area ($\mu\text{V} \cdot \text{sec}$)
1	9.18	3.07	146735
2	9.83	96.93	4626292

Figure S6 RP IP HPLC ANALYSIS OF Y-GRP78A194A175A1



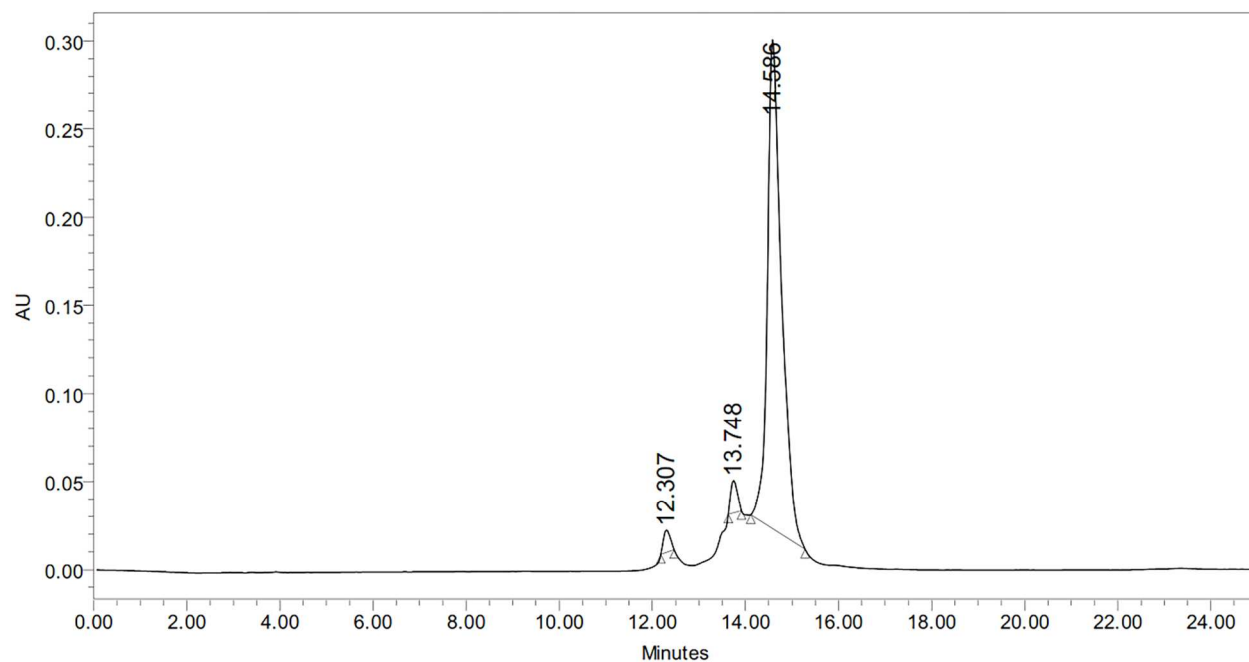
	RT	% Area	Area ($\mu\text{V}\cdot\text{sec}$)
1	10.40	100.00	1797312

Figure S7 RP IP HPLC ANALYSIS OF FL-GRP78A1 (260nm)



	RT	% Area	Area ($\mu\text{V}\cdot\text{sec}$)
1	9.28	1.00	324879
2	12.32	2.10	678728
3	13.76	2.10	680619
4	14.59	94.80	30706756

Figure S8 RP IP HPLC ANALYSIS OF FL-GRP78A1 (488nm)



	RT	% Area	Area ($\mu\text{V}\cdot\text{sec}$)
1	12.31	2.00	123746
2	13.75	2.95	181980
3	14.59	95.05	5866825

Figure S9 ESI-MS ANALYSIS OF FL-GRP78A1

Deconvoluted Mass Spectrum of FITC-GRP78A1 (P1), RT = 0.141 min:

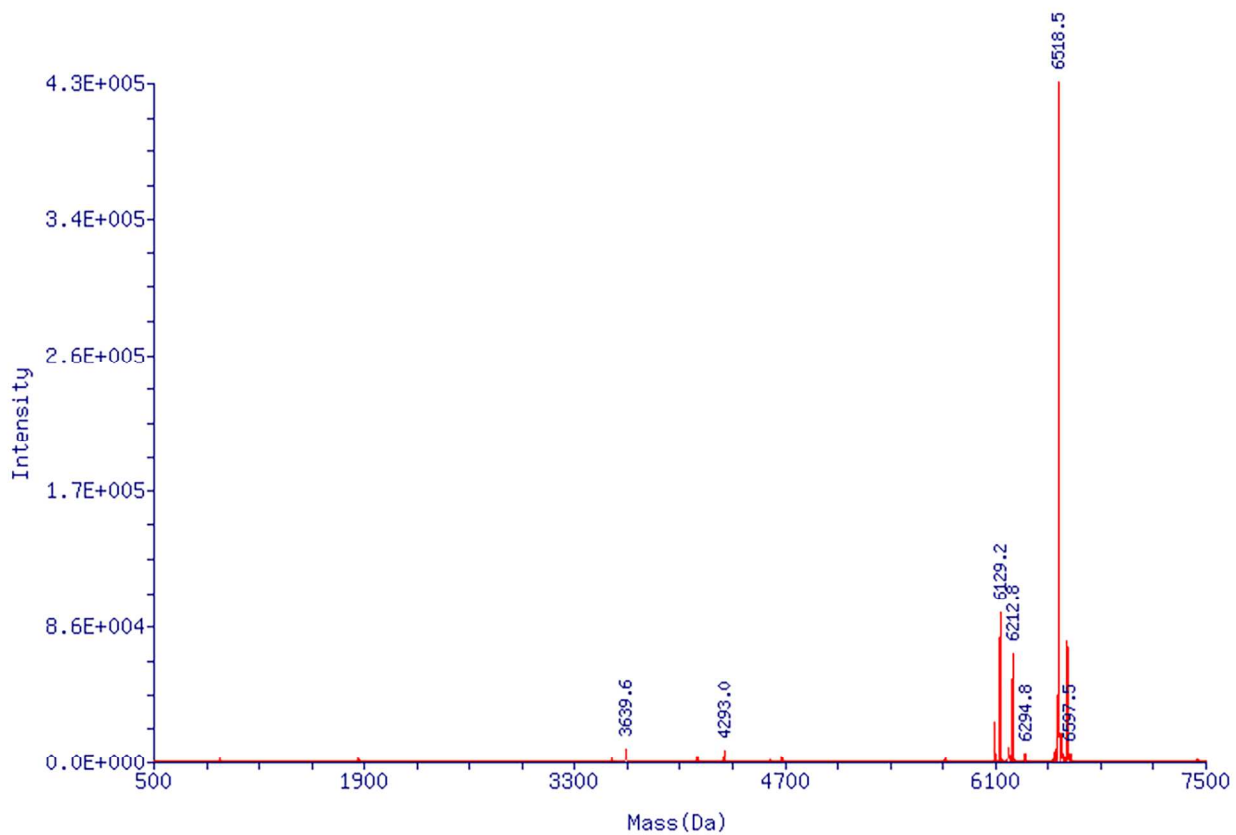
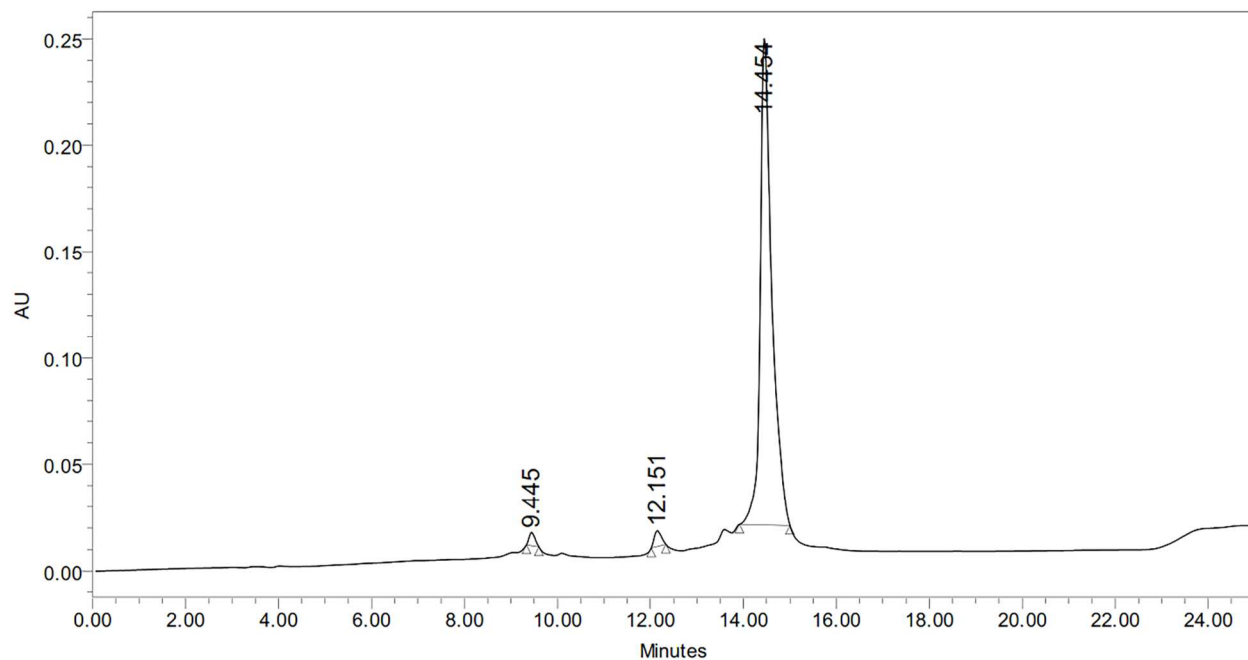
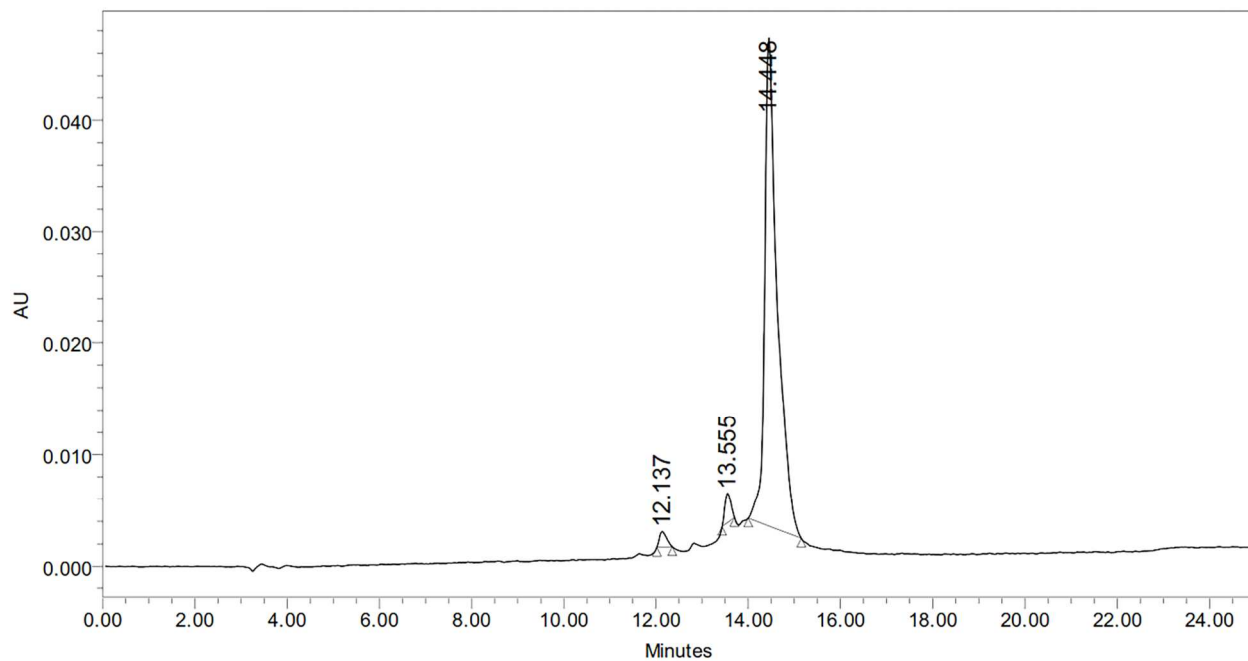


Figure S10 RP IP HPLC ANALYSIS OF FL-GRP78S1 (260nm)



	RT	% Area	Area ($\mu\text{V}\cdot\text{sec}$)
1	9.45	1.29	53177
2	12.15	1.86	76557
3	14.45	96.84	3981950

Figure S11 RP IP HPLC ANALYSIS OF FL-GRP78S1 (488nm)



	RT	% Area	Area ($\mu\text{V}\cdot\text{sec}$)
1	12.14	1.69	14785
2	13.56	2.70	23639
3	14.45	95.61	836278

Figure S12 ESI-MS ANALYSIS OF FL-GRP78S1

Deconvoluted Mass Spectrum of FITC78s1, RT = 0.344 min:

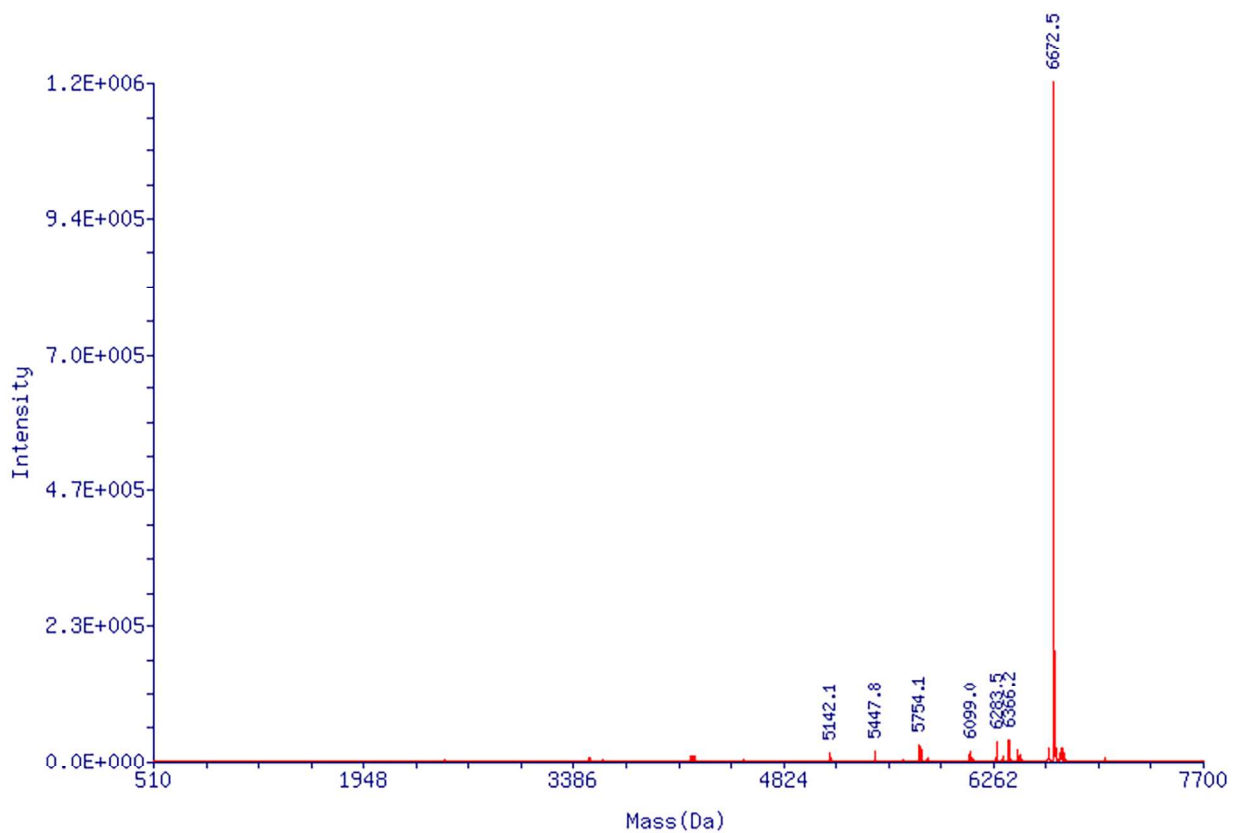
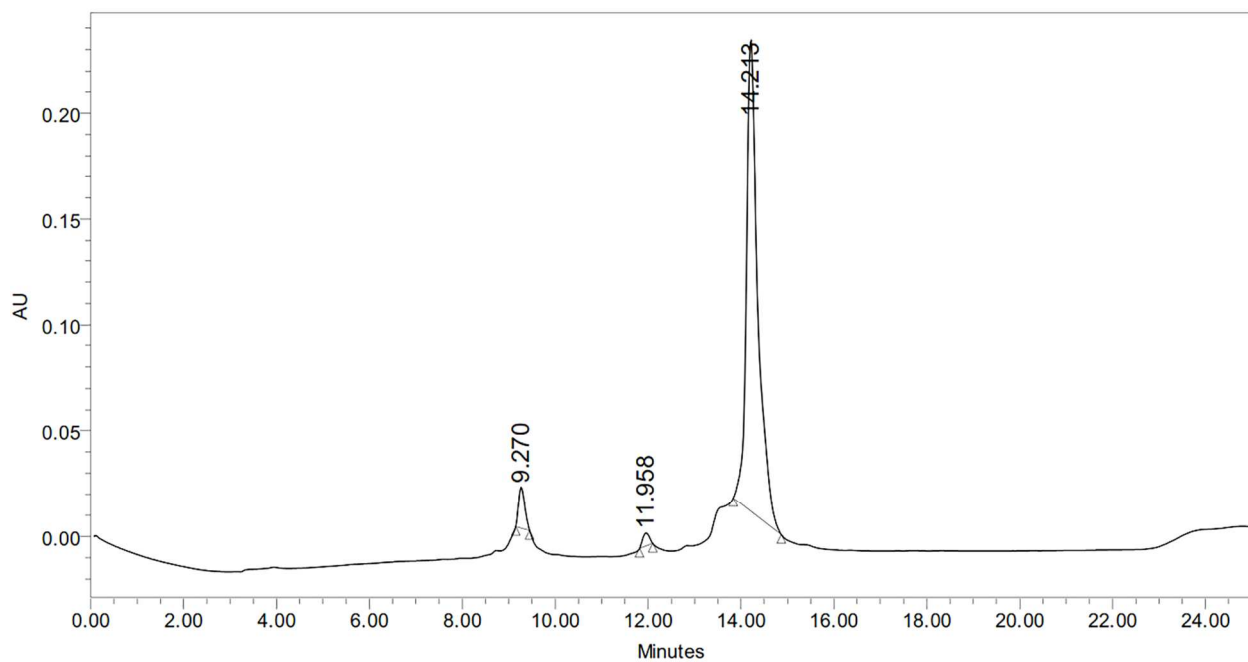


Figure S13 RP IP HPLC ANALYSIS OF FL-GRP94S1 (260nm)



	RT	% Area	Area ($\mu\text{V}\cdot\text{sec}$)
1	9.27	4.44	180347
2	11.96	1.47	59631
3	14.21	94.09	3818389

Figure S14 ESI-MS ANALYSIS OF FL-GRP94S1

Deconvoluted Mass Spectrum of FITC94s1, RT = 0.344 min:

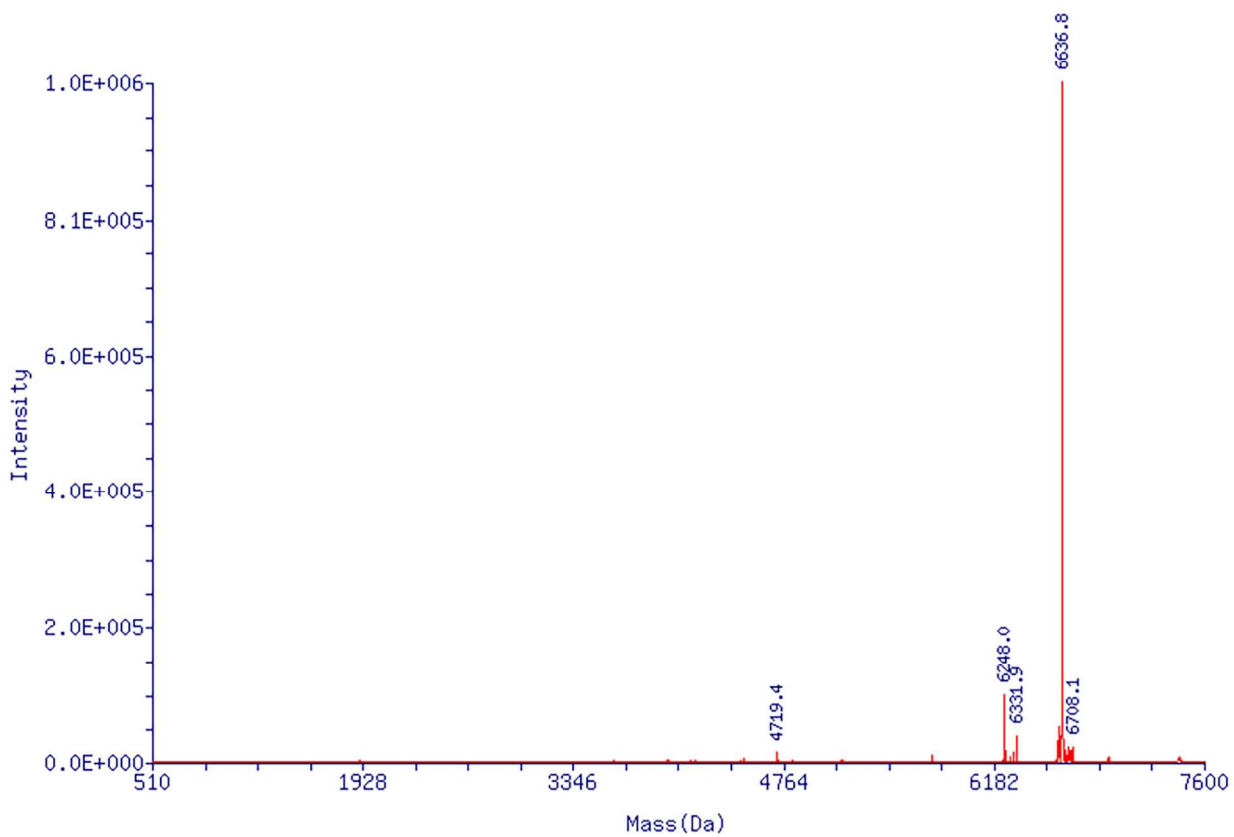
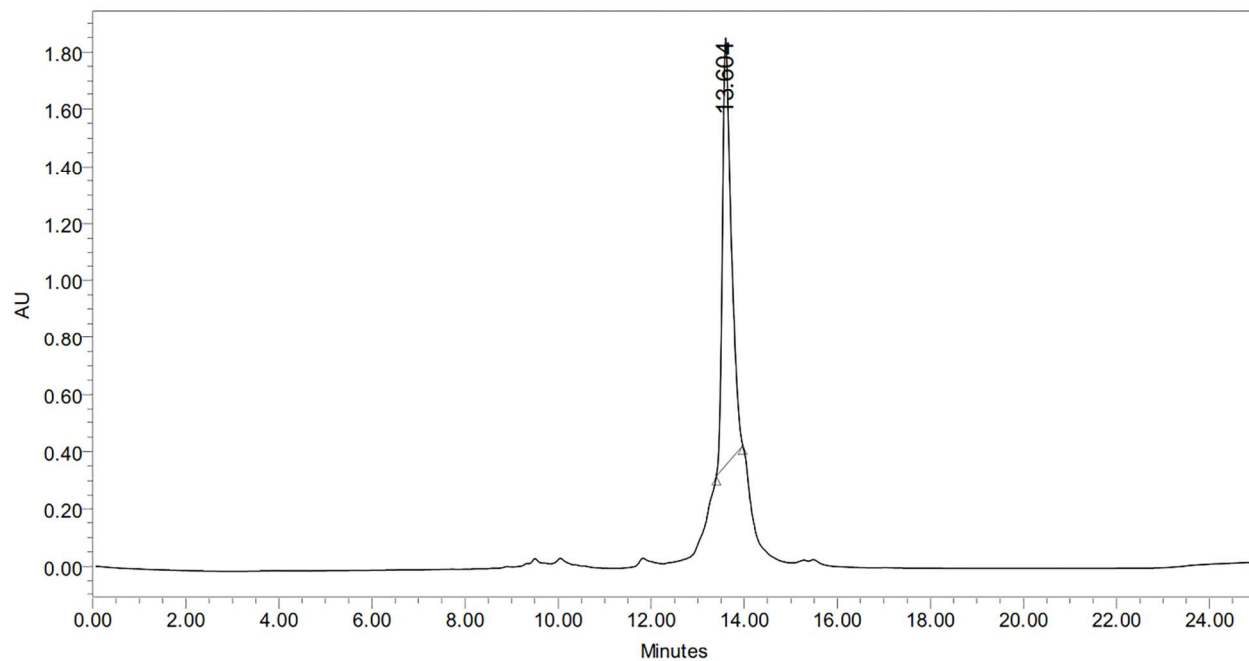
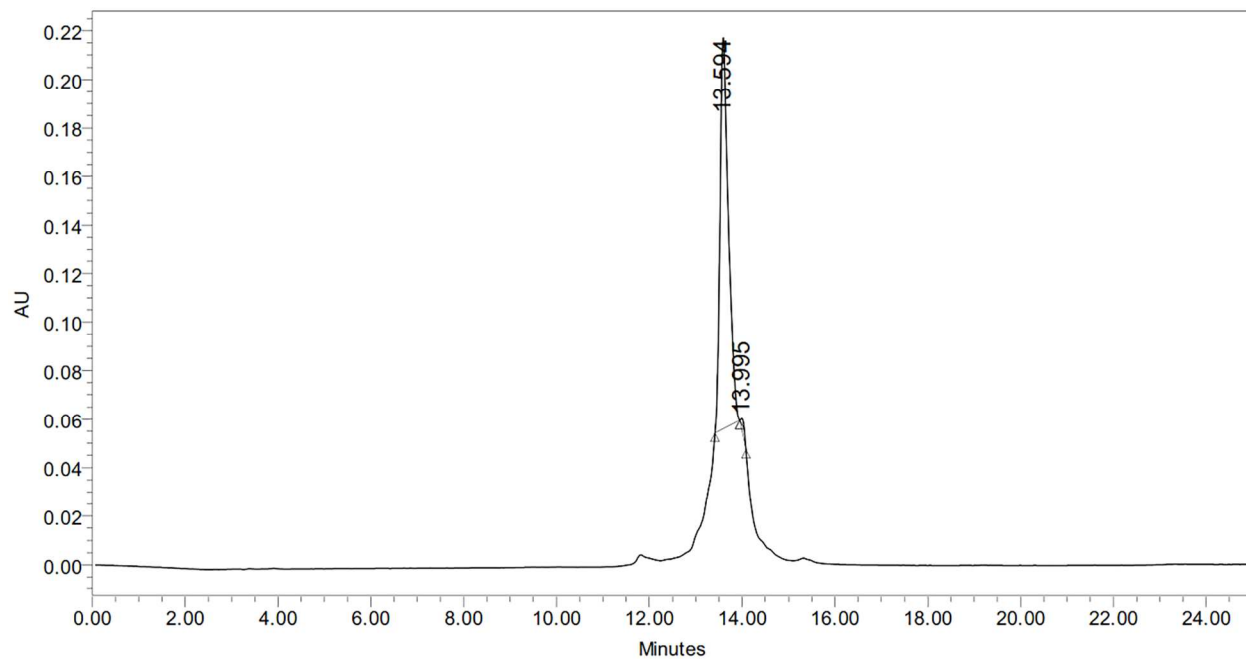


Figure S15 RP IP HPLC ANALYSIS OF FL-V-GRP78A194A1 (260nm)



	RT	% Area	Area ($\mu\text{V}\cdot\text{sec}$)
1	13.60	100.00	19780457

Figure S16 RP IP HPLC ANALYSIS OF FL-V-GRP78A194A1 (488nm)



	RT	% Area	Area ($\mu\text{V}\cdot\text{sec}$)
1	13.59	98.58	2031207
2	14.00	1.42	29335

Figure S17 ESI-MS ANALYSIS OF FL-V-GRP78A194A1

Deconvoluted Mass Spectrum of 7894_amino_FITC, RT = 0.135 min:

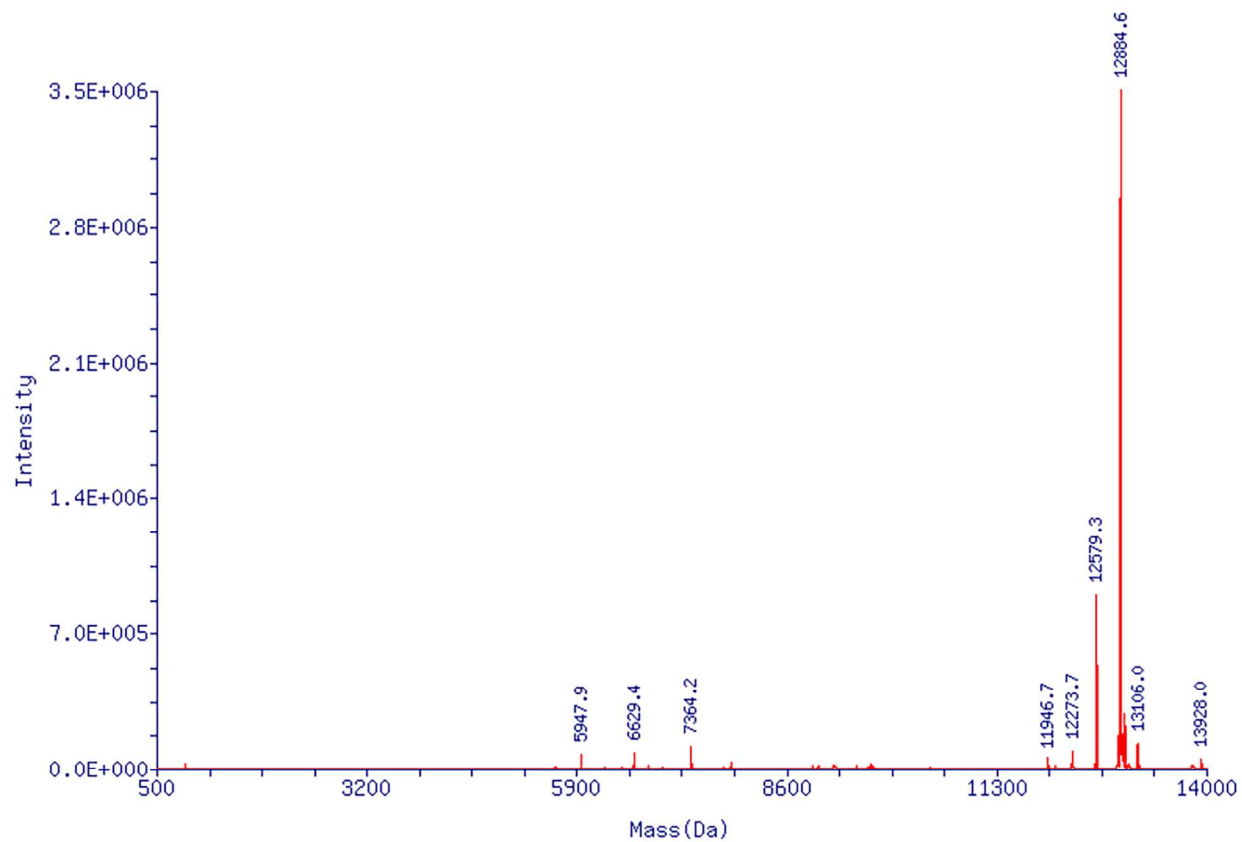
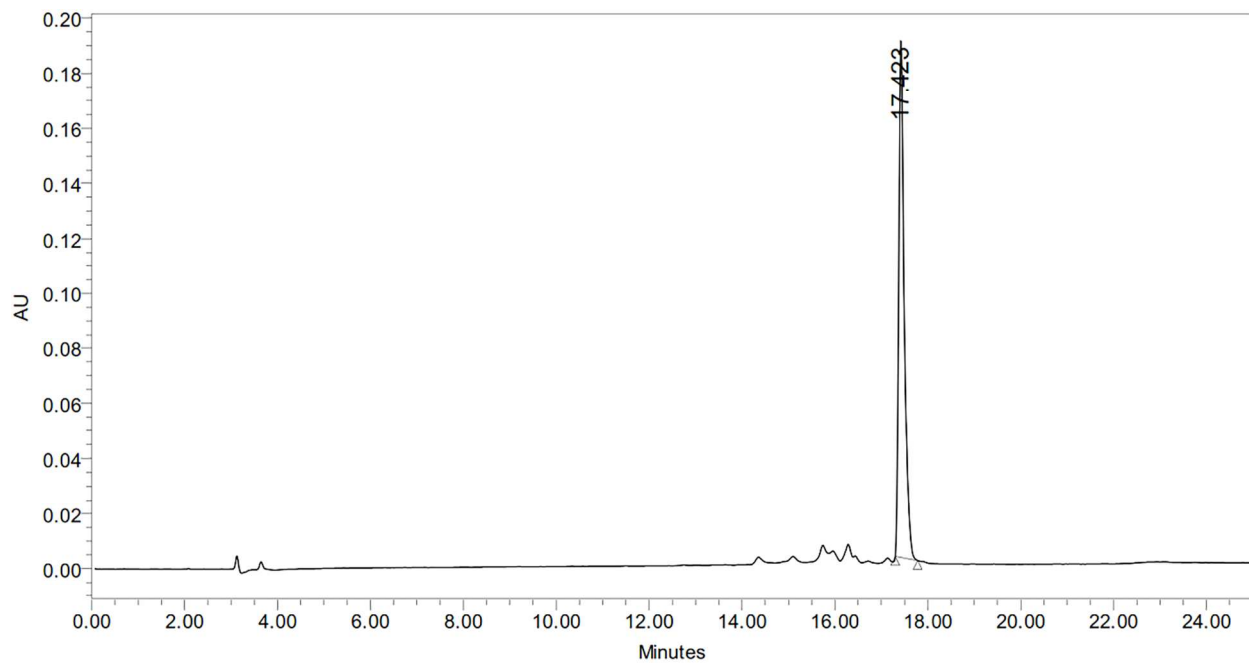


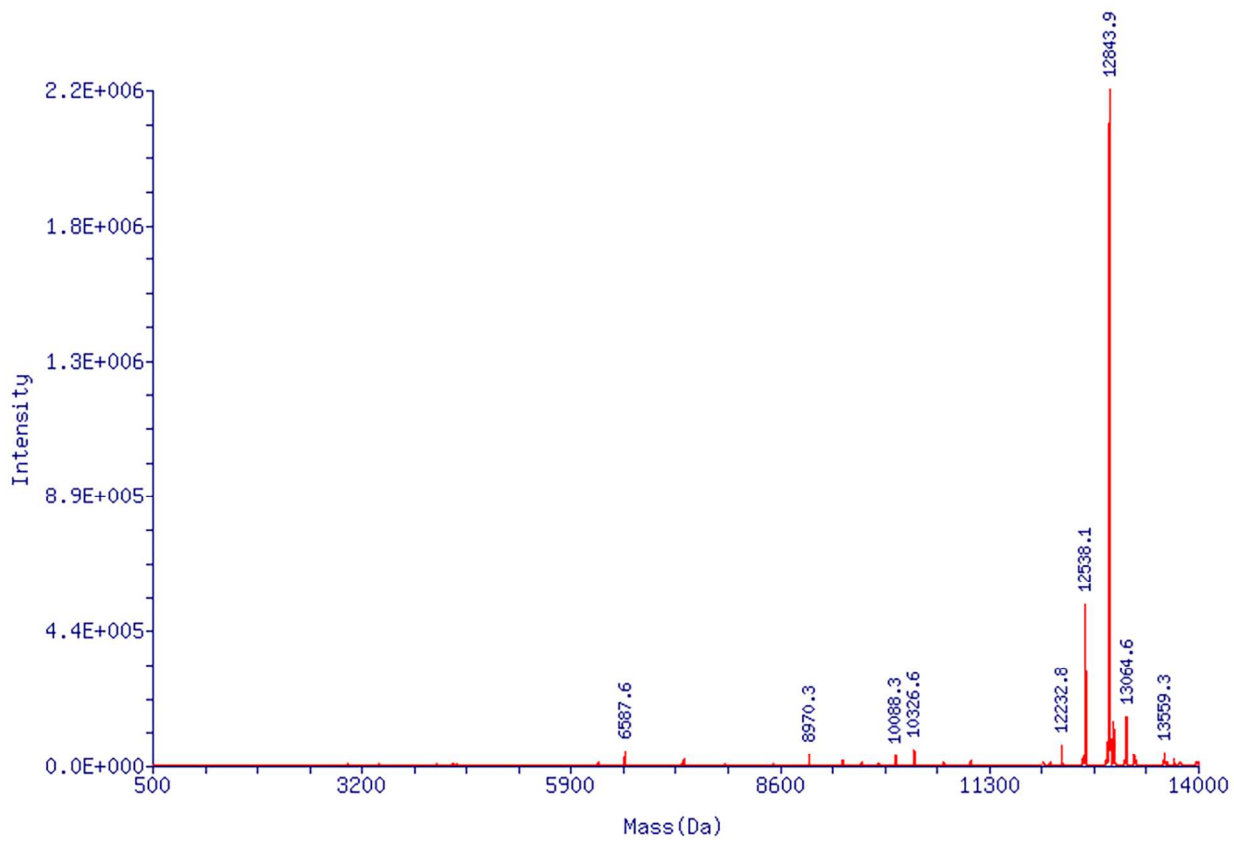
Figure S18 RP IP HPLC ANALYSIS OF FL-V-GRP78A178A2 (488nm)



	RT	% Area	Area ($\mu\text{V}\cdot\text{sec}$)
1	17.42	100.00	1533861

Figure S19 ESI-MS ANALYSIS OF FL-V-GRP78A178A2

Deconvoluted Mass Spectrum of 78a1a2_amino_FITC, RT = 0.125 min:



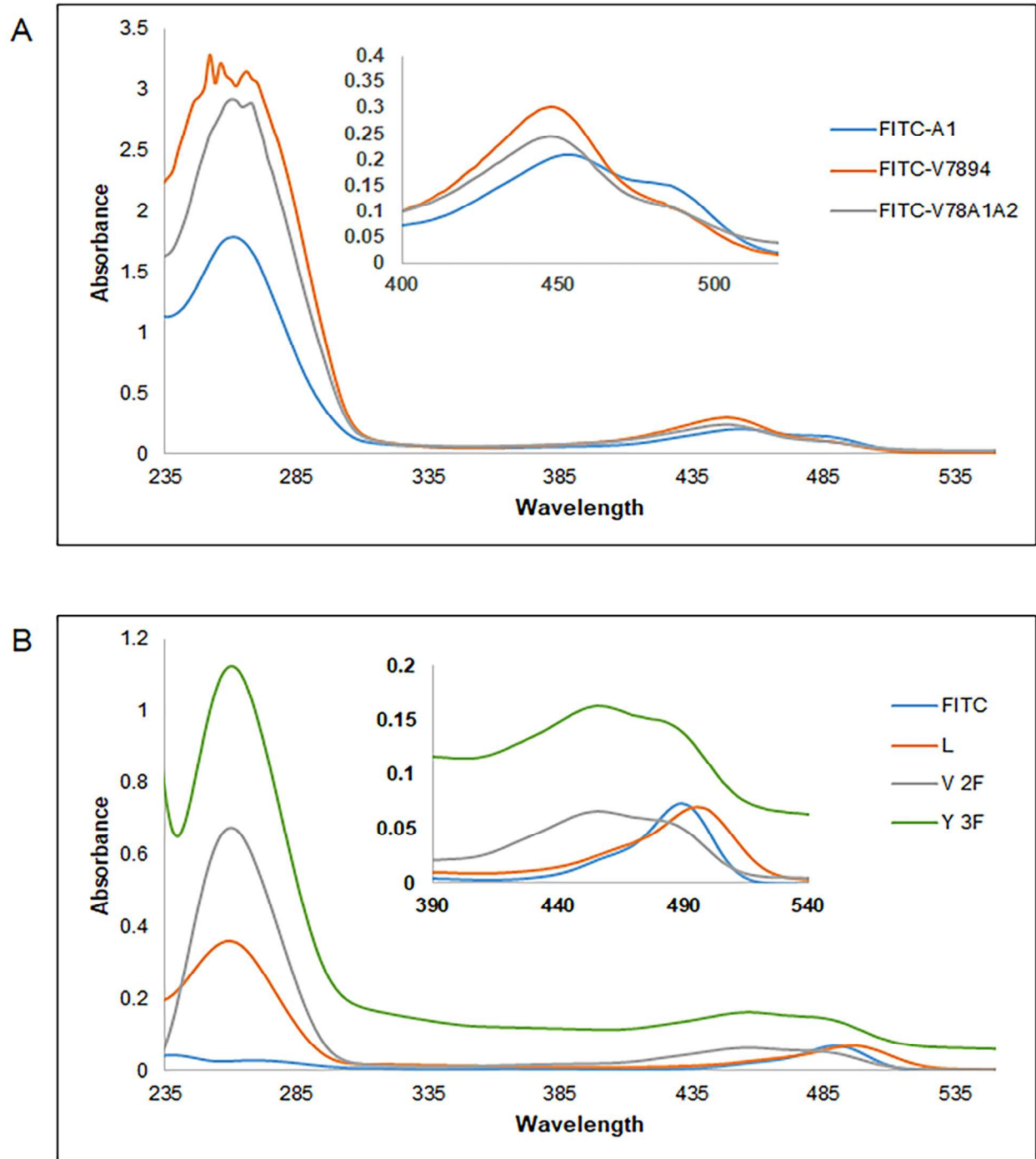


Figure S20 UV absorption spectra of FITC-labeled template siRNAs (**A**) and multi-FITC siRNAs (**B**). All siRNA hybrids were prepared in annealing buffer (1.25 μ M, 10 mM Tris, 50 mM NaCl, 1 mM EDTA, pH 7.5 – 8.0).

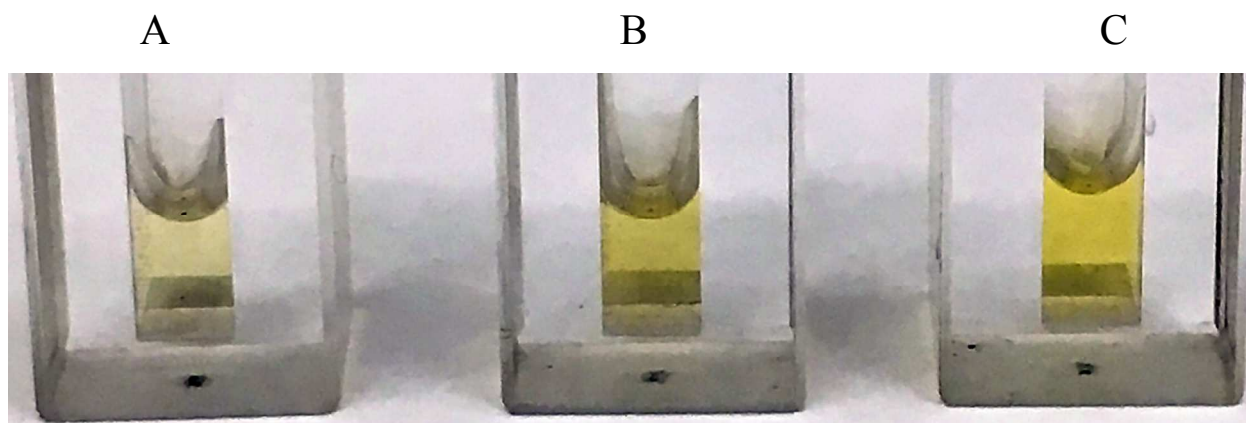
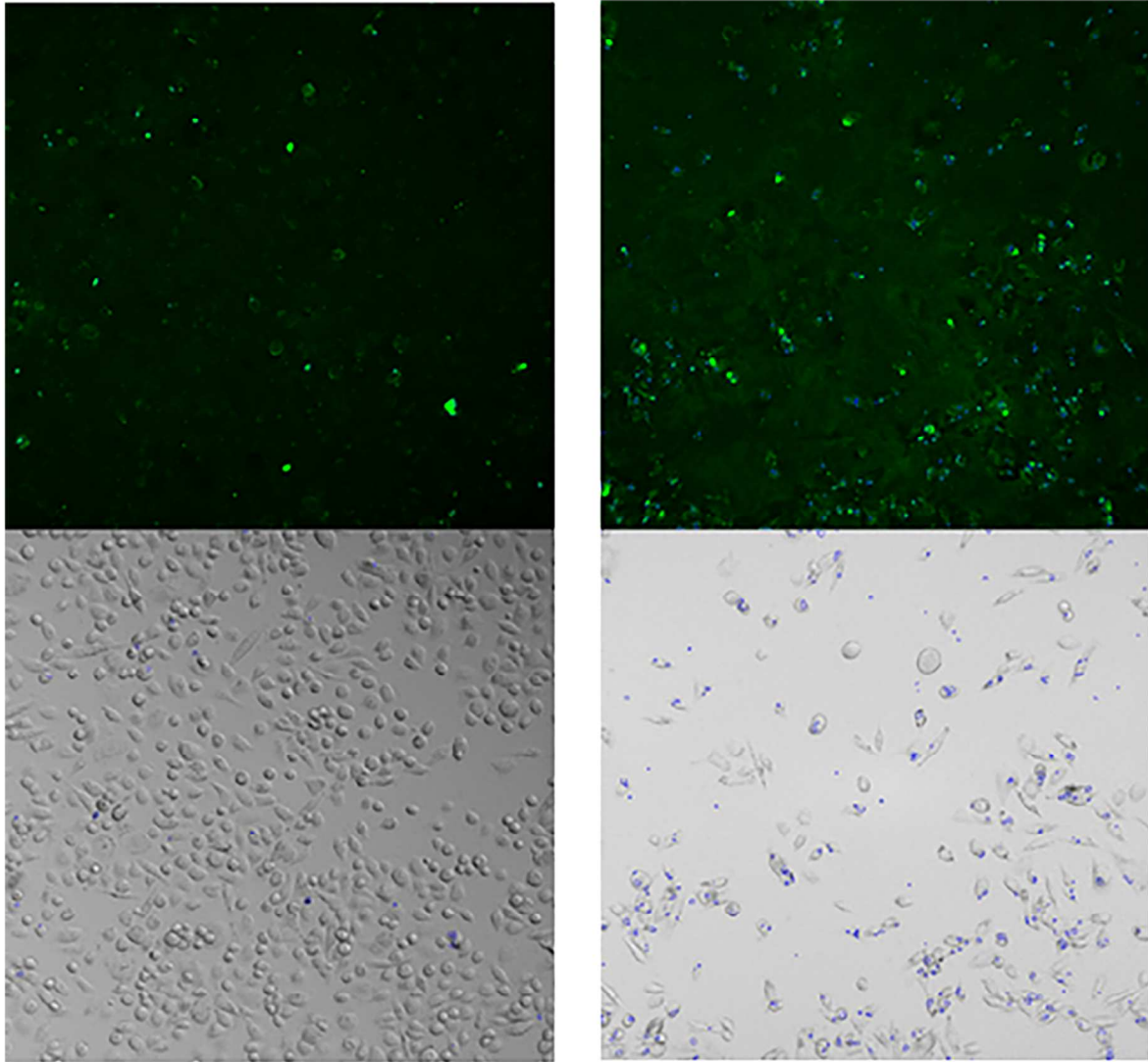


Figure S21 Multi-FITC siRNA samples pre-dilution for fluorescence spectroscopy analysis. Linear 78A1:FL-78S1 (A), V-78A194A1:FL-78S1FL-94S1 (B) and Y78A194A175A1:FL-78S1FL-94S1FL-75S1 (C). siRNA hybrids were prepared in annealing buffer (1.25 μ M, 10 mM Tris, 50 mM NaCl, 1 mM EDTA, pH 7.5 – 8.0).



V-FL-siRNA

4 hours post transfection

24 hours post transfection

Figure S22 Fluorescent microscope images of PC-3 cells treated with V-shape FITC-siRNA containing 2 FITC probes 4 and 24 hours post transfection.

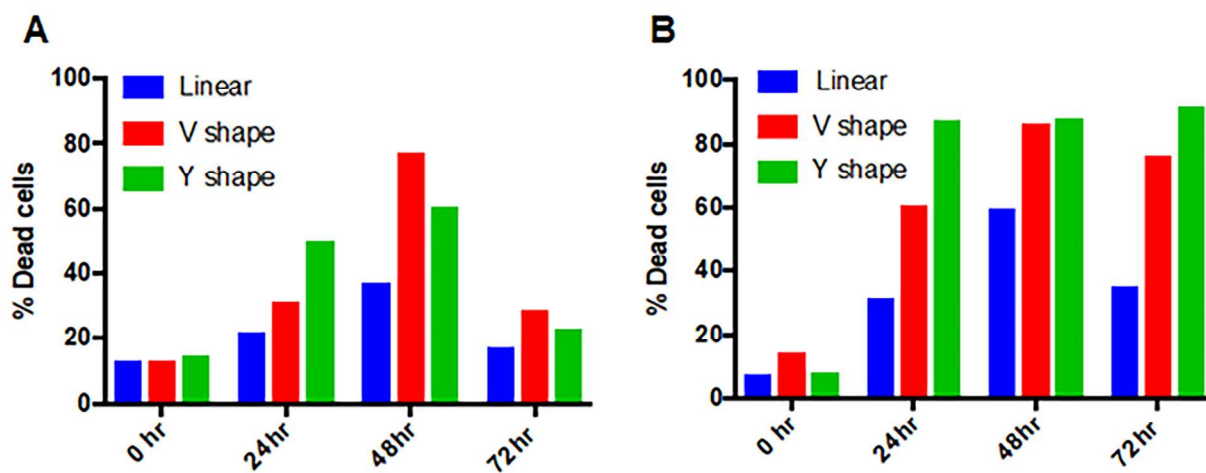


Figure S23 Cell viability determined using PI on flow cytometry, adhered cells (A) and supernatant cells (B). After 24, 48 and 72 h, the supernatant was collected, and the remaining cells were removed using trypsin. Cell samples were pelleted and re-suspended in 1% BSA in PBS (1 mL) and stained using PI according the manufacturer's recommendation.