# **Supporting Information**

## Synthesis and Mechanism Studies of 1,3-Benzoazolyl Substitued Pyrrolo[2,3-*b*]pyrazine Derivatives as Non-intercalative Topoisomerase II Catalytic Inhibitors

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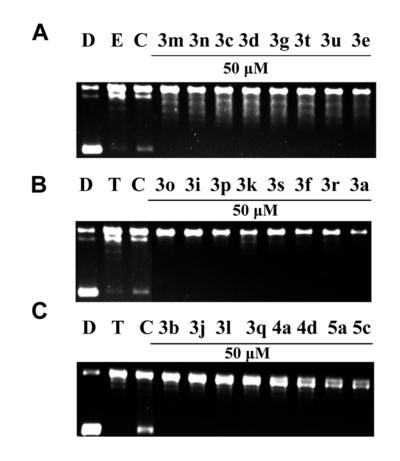
School of Pharmaceutical Sciences, Sun Yat-sen University, Guangzhou, 510006, China

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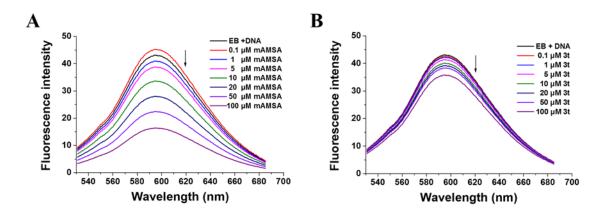
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**Topo I-midated DNA relaxation assay.** Topo I-midated DNA relaxation assay was performed to test whether BPPs act as a dual inhibitor of Topo I and Topo II.<sup>1, 2</sup> The result was presented at Figure S1. All of the synthesized compounds had no Topo I inhibitory activity at 50  $\mu$ M.



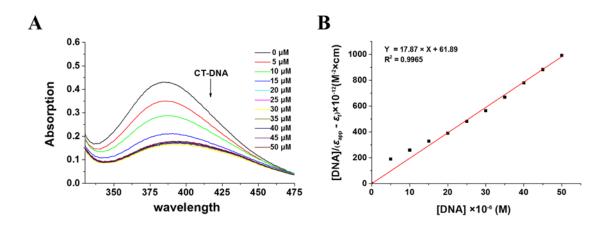
**Figure S1.** Topo I inhibitory activity of BPPs. (A), (B) and (C) lane **D**: pBR322 DNA; lane **T**: pBR322 DNA + Topo I; lane **C**: pBR322 DNA + Topo I + Camptothecin (100 μM) other lanes: pBR322 DNA + Topo I + BPPs (50 μM).

**EB displacement assay.** The ability of **3t** to interact with DNA was also confirmed by fluorescence-based EB displacement assay (Figure S2). The DNA bound form of EB has a stronger fluorescence emission than free EB, so the displacement of EB from DNA could be monitored by a decrease in the fluorescence signal.<sup>6</sup> As expected, the addition of the DNA intercalator mAMSA was accompanied by a concomitant decrease in EB fluorescence (Figure S2A). In contrast, the addition of increasing concentration of **3t** did not alter the intensity of EB fluorescence (Figure S2B), indicating that **3t** was incapable of intercalating into DNA.



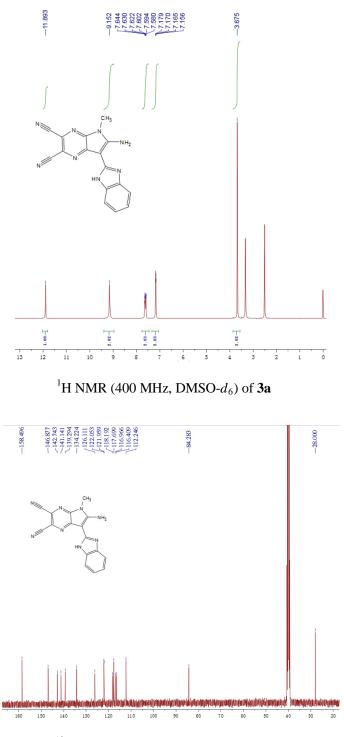
**Figure S2.** (A) and (B) pBR322 DNA (20 nM) was incubated in the presence of EB (2.5  $\mu$ M). Arrow indicates the fluorescence intensity change upon increasing concentrations of mAMSA (A) or **3t** (B).

**UV-vis titration assay**. UV-vis titration was employed to measure the binding affinity of BPPs to CT-DNA. Generally, interaction between compounds and DNA would result in hypochromic effect and red shifts in the absorption spectrum due to the strong stacking interaction between the aromatic chromophore of the compounds and the base pair of DNA, and the electron transfer effect in the process.<sup>3, 4</sup> As showed in Figure S3A, with the increasing addition of CT-DNA to compound **3t** resulted in a hypochromic effect, but without an obvious red shifts, which was different from the classical intercalation interaction. The apparent binding constant ( $K_b$ ) to CT-DNA was determined to be  $2.8 \times 10^5$  M<sup>-1</sup> (Figure S3B), means that BPPs showed weak interaction with CT-DNA. This is consistent with the results obtained from the unwinding assay.

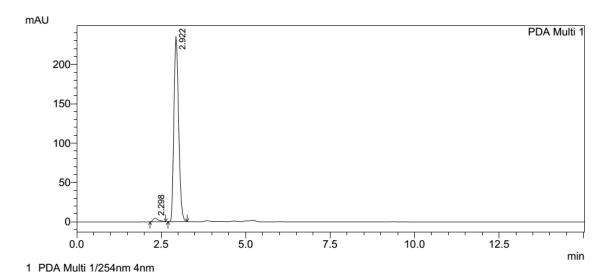


**Figure S3.** (A) Absorption spectra of compound **3t** upon addition of CT-DNA. [Compound **3t**] = 5  $\mu$ M, [DNA] = 0 – 50  $\mu$ M. Arrow indicates the absorption change upon increasing amount of CT-DNA. (B) Graph shows the plot of [DNA]/ ( $\varepsilon_a - \varepsilon_f$ ) versus [DNA].

<sup>1</sup>H NMR, <sup>13</sup>C NMR, HRMS, and HPLC analysis of compounds.



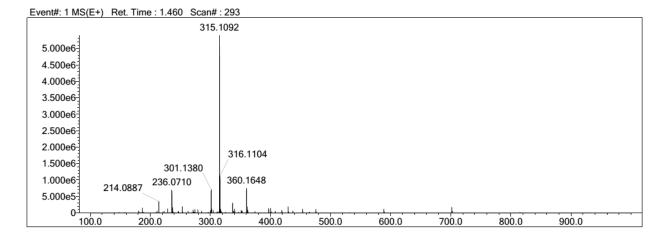
<sup>13</sup>C NMR (100 MHz, DMSO- $d_6$ ) of **3a** 



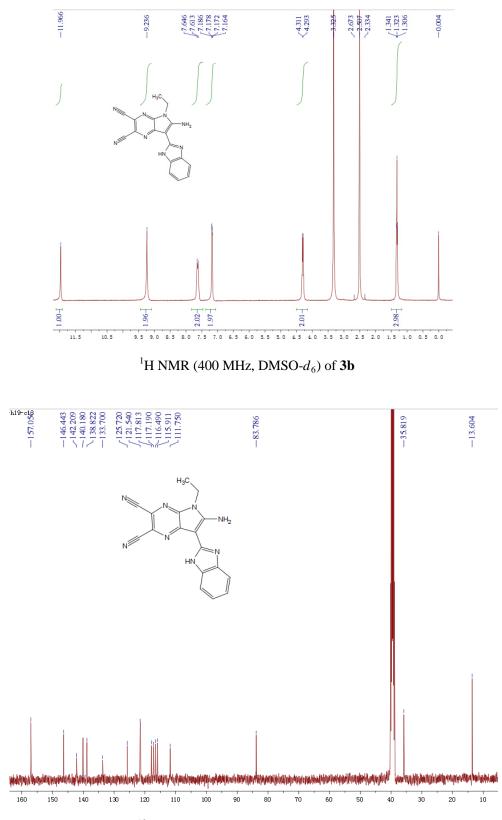
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Peak#	Ret. Time	Area	Height	Area %	Height %				
1	2.298	51581	4682	2.208	1.951				
2	2.922	2284804	235341	97.792	98.049				
Total		2336385	240023	100.000	100.000				

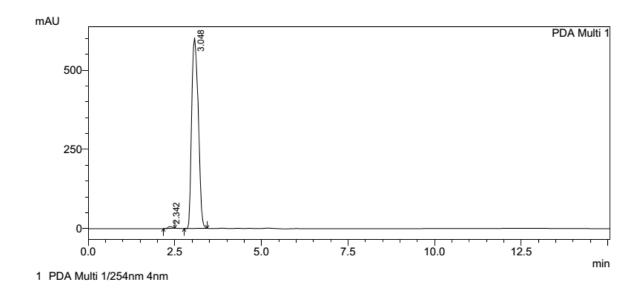
HPLC spectrum of 3a



HRMS spectrum of 3a

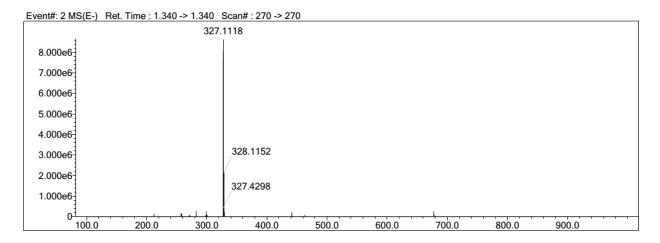


<sup>13</sup>C NMR (100 MHz, DMSO- $d_6$ ) of **3b** 

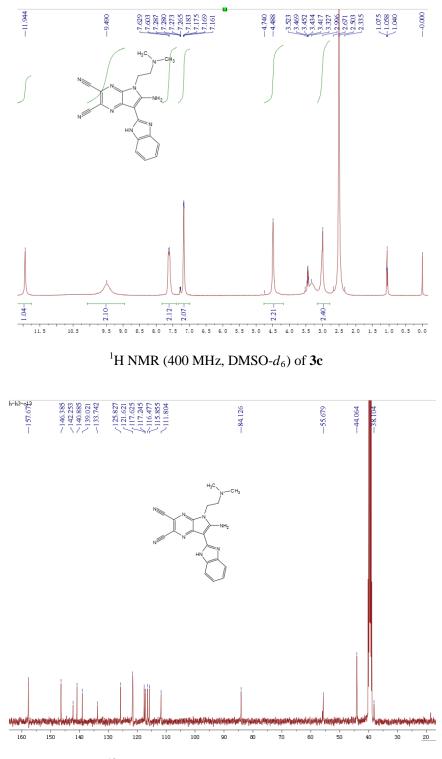


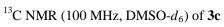
		PeakTable					
PDA Ch1 254nm 4nm							
Peak#	Ret. Time	Area	Height	Area %	Height %		
1	2.342	48688	5864	0.606	0.965		
2	3.048	7982812	601737	99.394	99.035		
Total		8031500	607601	100.000	100.000		

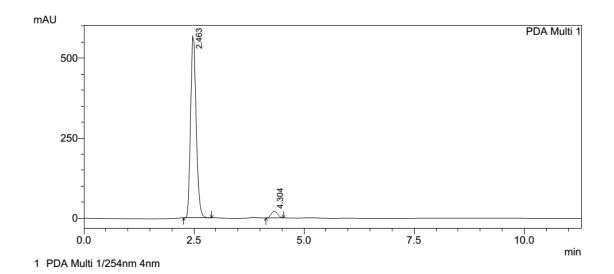
HPLC spectrum of 3b



HRMS spectrum of 3b



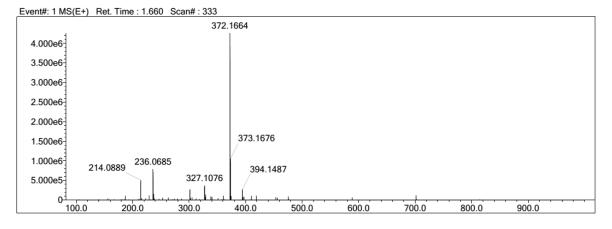




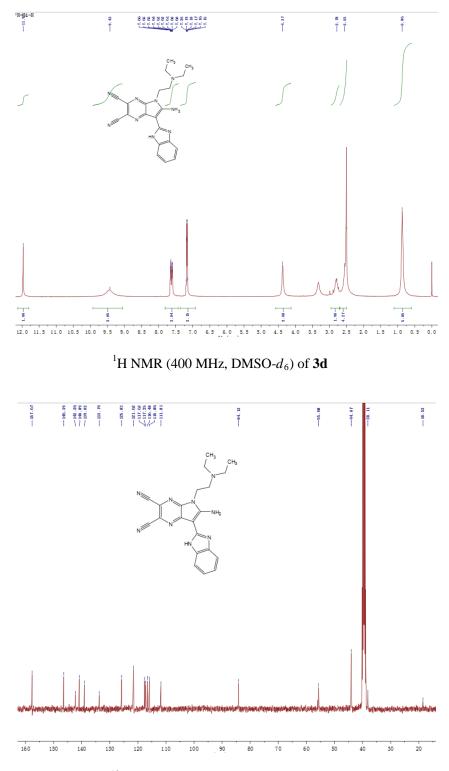
PeakTable

P	PDA Ch1 254nm 4nm								
	Peak#	Ret. Time	Area	Height	Area %	Height %			
	1	2.463	5002713	568967	95.951	96.419			
Γ	2	4.304	211085	21133	4.049	3.581			
	Total		5213799	590100	100.000	100.000			

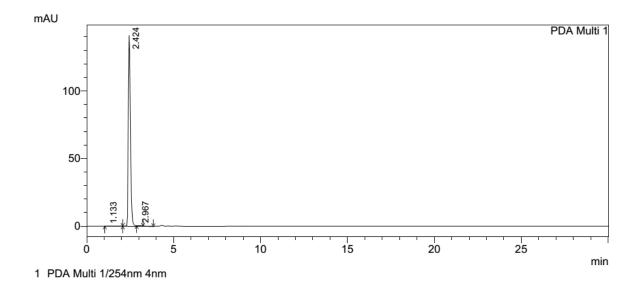
HPLC spectrum of 3c



HRMS spectrum of 3c

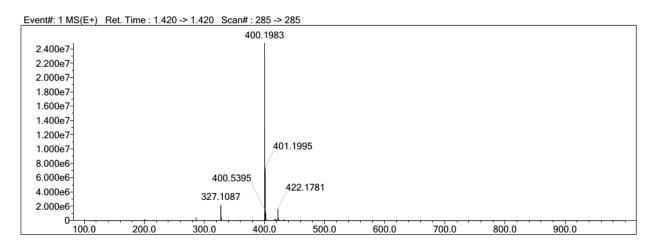


 $^{13}$ C NMR (100 MHz, DMSO- $d_6$ ) of **3d** 

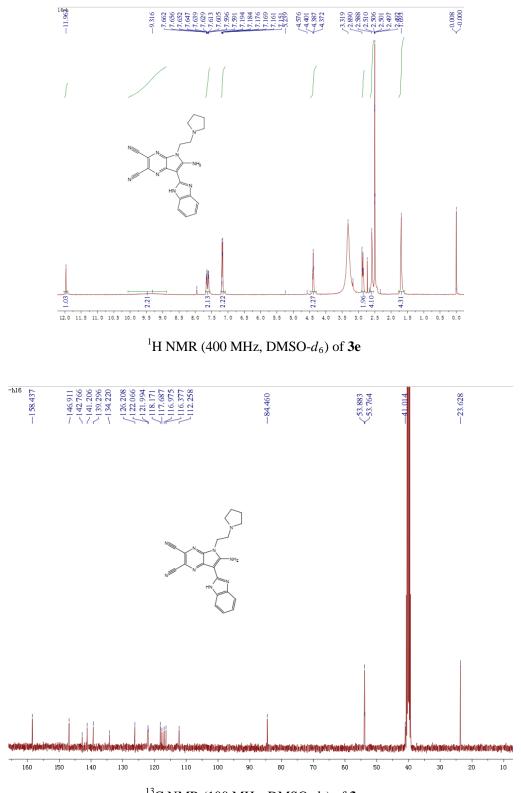


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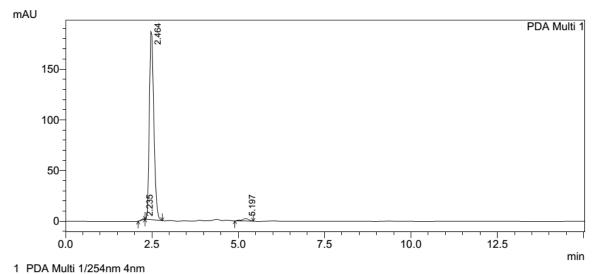
### HPLC spectrum of 3d



### HRMS spectrum of 3d



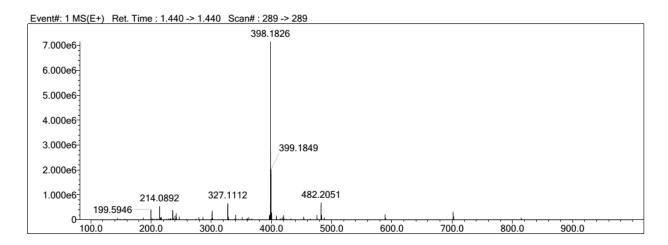
<sup>13</sup>C NMR (100 MHz, DMSO- $d_6$ ) of **3e** 



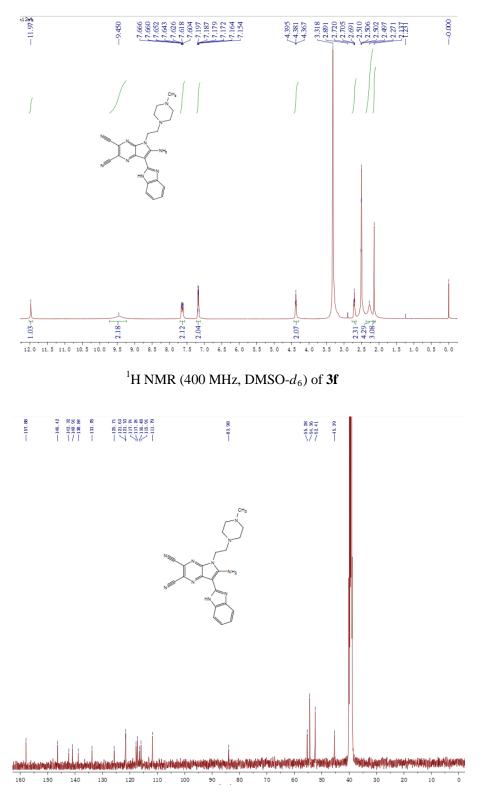
PeakTable

		1 cur i uoic						
ł	PDA Ch1 254nm 4nm							
	Peak#	Ret. Time	Area	Height	Area %	Height %		
	1	2.235	7490	1099	0.438	0.580		
	2	2.464	1670709	185942	97.612	98.145		
Γ	3	5.197	33380	2415	1.950	1.275		
	Total		1711579	189456	100.000	100.000		

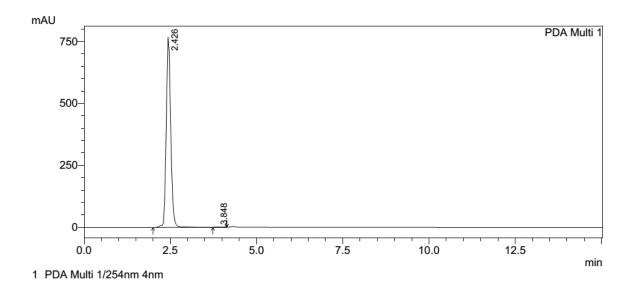
HPLC spectrum of 3e



HRMS spectrum of 3e



13C NMR (100 MHz, DMSO-d6) of 3f



PeakTable

PDA Ch1 254nm 4nm								
Peak#	Ret. Time	Area	Height	Area %	Height %			
1	2.426	6912189	767448	99.816	99.882			
2	3.848	12720	906	0.184	0.118			
Total		6924909	768354	100.000	100.000			

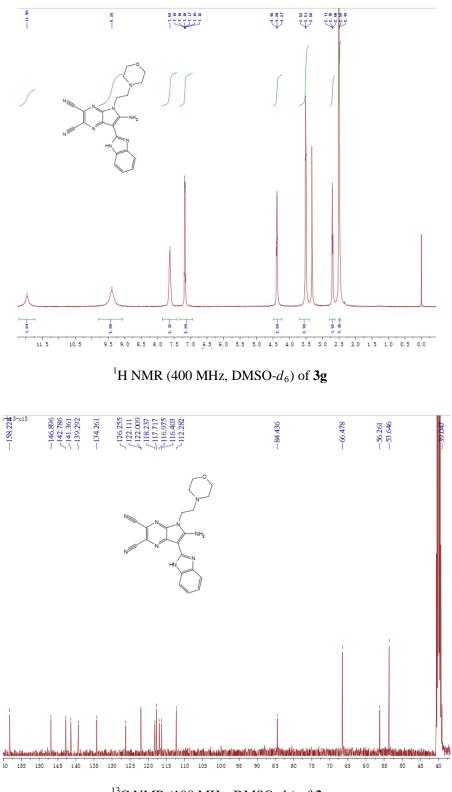
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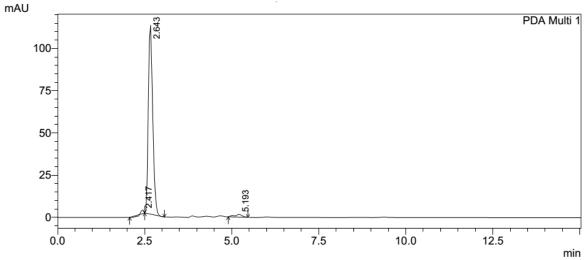
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HPLC spectrum of **3f** 

Event#: 1 MS(E+) Ret. Time :	: 1.470 -> 1.470 Scan# : 295 -> 295
214.	.1075
1.300e7-	
1.200e7-	
1.100e7-	
1.000e7-	
9.000e6-	
8.000e6	
7.000e6-	
6.000e6	214.6072 427.2083
5.000e6-	4212003
4.000e6-	
3.000e6-	428.2087
2.000e6	301.1015
1.000e6-	327.1074
100.0 200.	Lington and the second se

HRMS spectrum of 3f





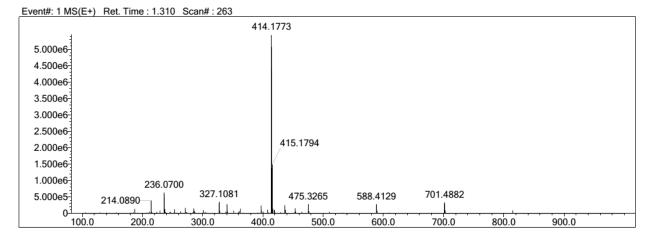
1 PDA Multi 1/254nm 4nm

DDA Ch1 254mm 4mm

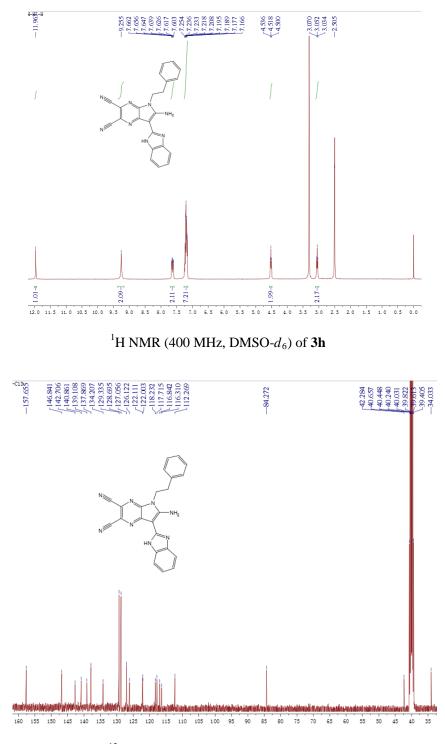
PeakTable

PDA Chi 2	PDA Ch1 254nm 4nm								
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1	2.417	4789	2177	0.473	1.885				
2	2.643	983937	111775	97.094	96.763				
3	5.193	24659	1562	2.433	1.352				
Total		1013385	115515	100.000	100.000				

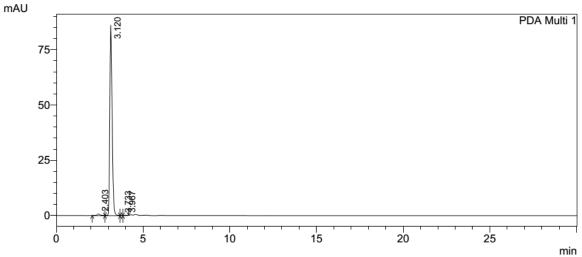
HPLC spectrum of 3g



HRMS spectrum of 3g



<sup>13</sup>C NMR (100 MHz, DMSO- $d_6$ ) of **3h** 

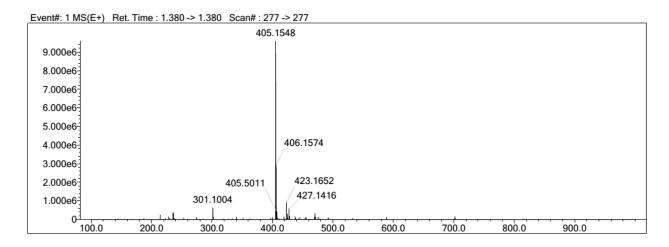


1 PDA Multi 1/254nm 4nm

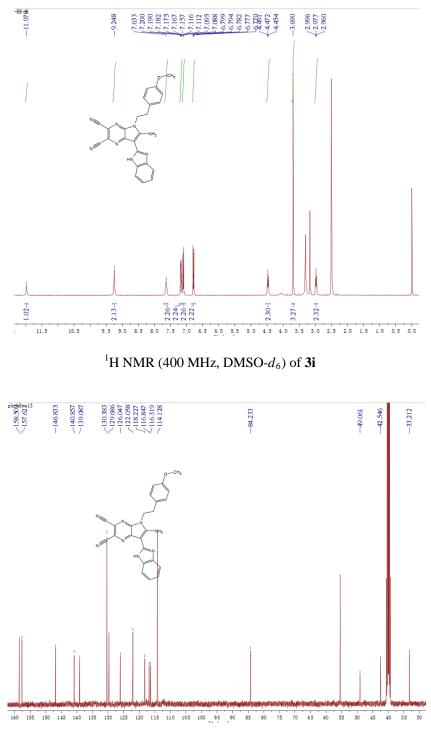
PeakTable

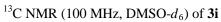
		Peak lable					
Р	DA Ch1 2	54nm 4nm					
	Peak#	Ret. Time	Area	Height	Area %	Height %	
	1	2.403	11289	766	1.369	0.877	
	2	3.120	810652	86229	98.287	98.718	
	3	3.733	1146	211	0.139	0.241	
	4	3.967	1693	143	0.205	0.164	
	Total		824780	87349	100.000	100.000	

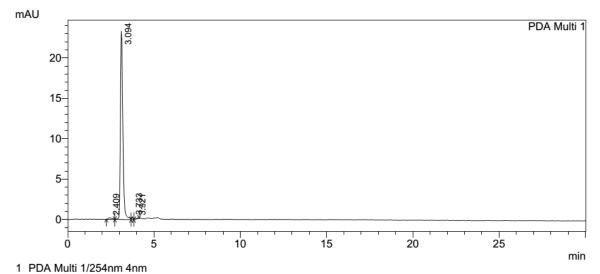
HPLC spectrum of 3h



HRMS spectrum of 3h

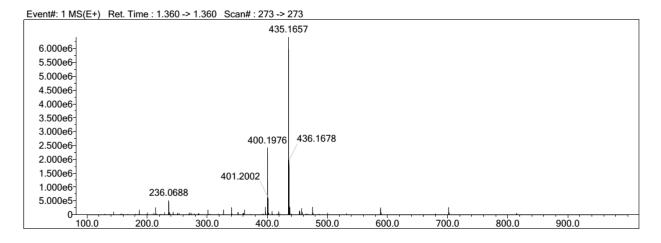




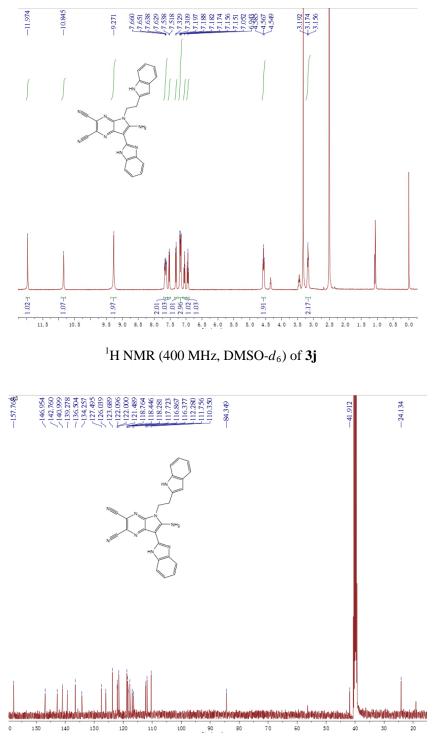


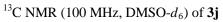
			PeakTable	e	
PDA Ch1 2	54nm 4nm				
Peak#	Ret. Time	Area	Height	Area %	Height %
1	2.409	3074	207	1.301	0.863
2	3.094	230303	23333	97.522	97.411
3	3.733	1428	269	0.605	1.121
4	3.921	1350	145	0.572	0.605
Total		236154	23953	100.000	100.000

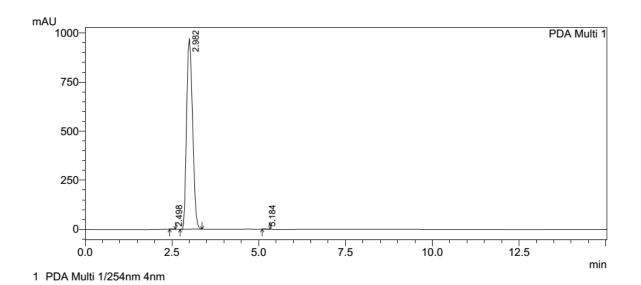
HPLC spectrum of 3i



HRMS spectrum of 3i

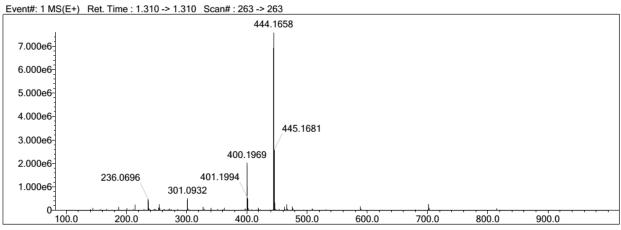




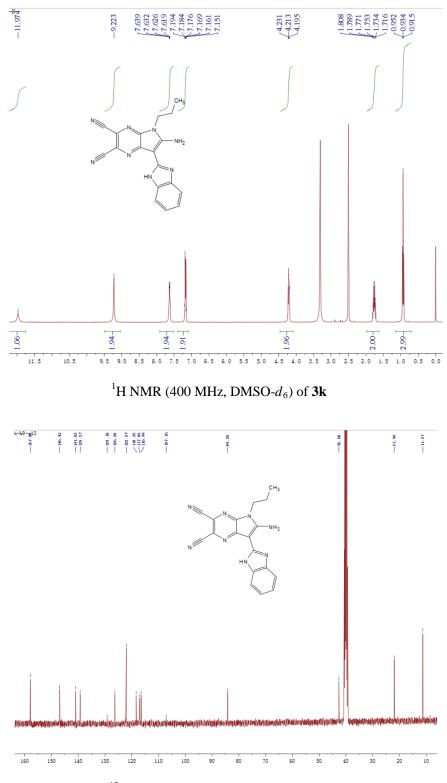


PDA Ch1 2	PeakTable DA Ch1 254nm 4nm						
Peak#	Ret. Time	Area	Height	Area %	Height %		
1	2.498	2473	408	0.021	0.042		
2	2.982	11670675	972320	99.925	99.874		
3	5.184	6254	822	0.054	0.084		
Total		11679402	973550	100.000	100.000		

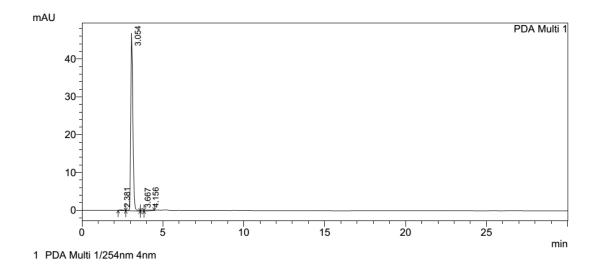
HPLC spectrum of 3j



HRMS spectrum of **3j** 



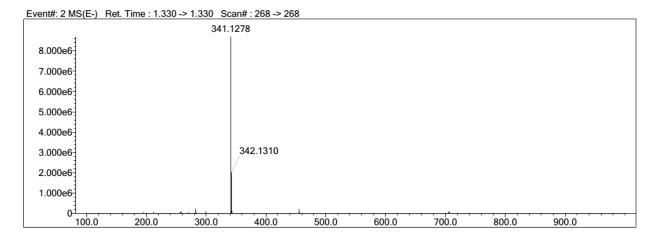
<sup>13</sup>C NMR (100 MHz, DMSO- $d_6$ ) of **3**k



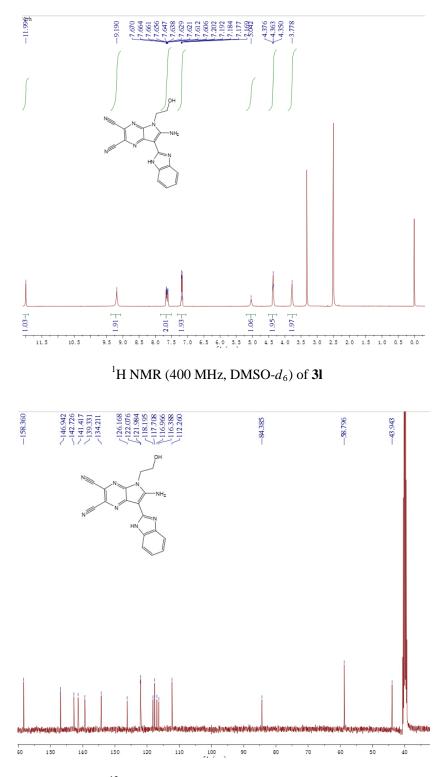
PeakTable

	Peak I able							
PDA Ch1 2	PDA Ch1 254nm 4nm							
Peak#	Ret. Time	Area	Height	Area %	Height %			
1	2.381	3094	168	0.707	0.354			
2	3.054	426908	46944	97.607	98.853			
3	3.667	2292	186	0.524	0.392			
4	4.156	5079	190	1.161	0.401			
Total		437373	47489	100.000	100.000			

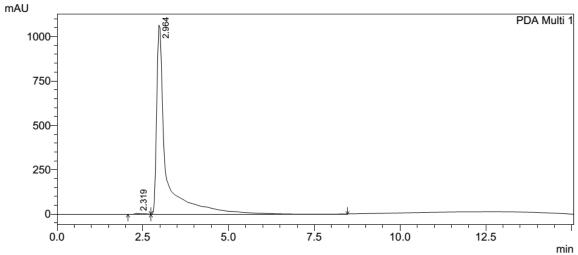
HPLC spectrum of 3k



HRMS spectrum of 3k



<sup>13</sup>C NMR (100 MHz, DMSO- $d_6$ ) of **3**l



1 PDA Multi 1/254nm 4nm

PeakTable

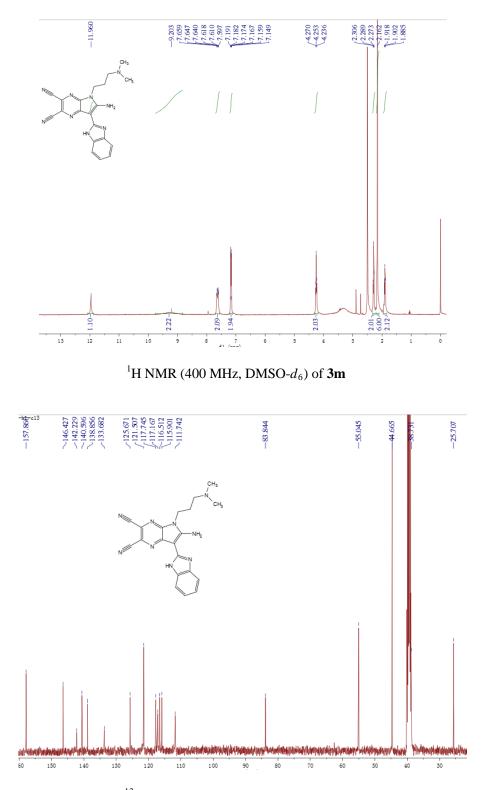
			Peak I able	;			
PDA Ch1 254nm 4nm							
Peak#	Ret. Time	Area	Height	Area %	Height %		
1	2.319	75854	4656	0.368	0.435		
2	2.964	20510840	1065559	99.632	99.565		
Total		20586693	1070215	100.000	100.000		

### HPLC spectrum of 31

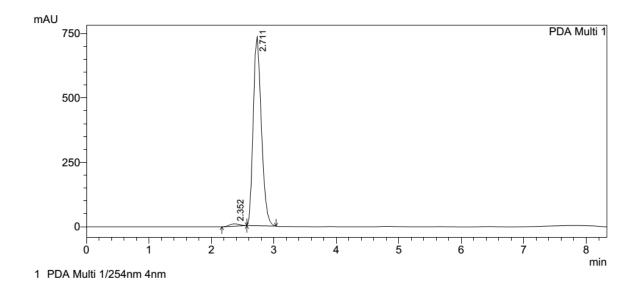
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1.200e7-								
1.100e7-								
1.000e7-								
9.000e6-								
8.000e6-								
7.000e6-								
6.000e6								
5.000e6-			346.1210					
4.000e6-								
3.000e6								
2.000e6-		301.0932	345.4445					
1.000e6-		1	/ .					
0 100.0	200.0		400.0	F00.0	600.0	700.0	800.0	000.0
100.0	200.0	300.0	400.0	500.0	600.0	700.0	800.0	900.0

HRMS spectrum of 31



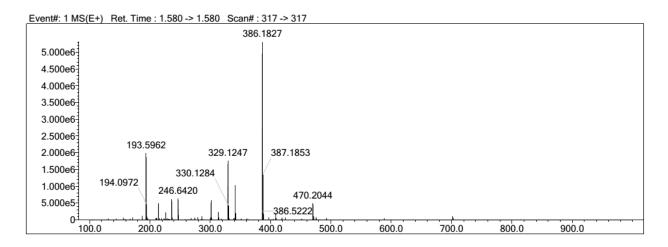
<sup>13</sup>C NMR (100 MHz, DMSO- $d_6$ ) of **3m** 



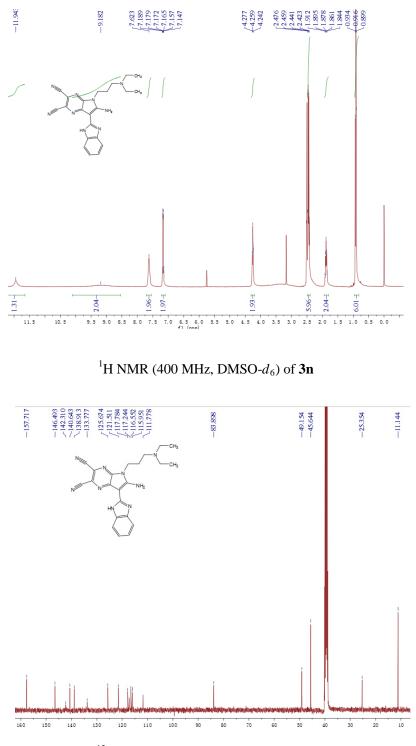
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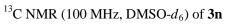
1 van 1 uorv								
PDA Ch1 254nm 4nm								
Ret. Time	Area	Area Height		Height %				
2.352	103081	9181	1.519	1.232				
2.711	6684290	736173	98.481	98.768				
	6787372	745354	100.000	100.000				
	Ret. Time 2.352	Ret. Time         Area           2.352         103081           2.711         6684290	Area         Height           2.352         103081         9181           2.711         6684290         736173	Area         Height         Area %           2.352         103081         9181         1.519           2.711         6684290         736173         98.481				

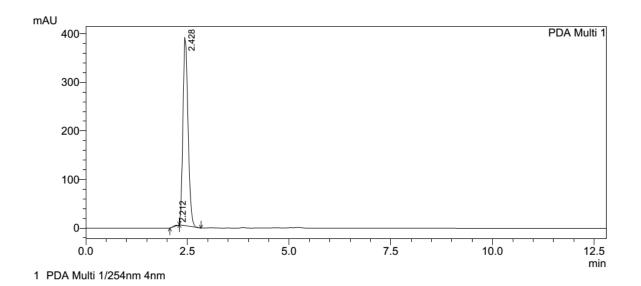
HPLC spectrum of 3m



HRMS spectrum of **3m** 



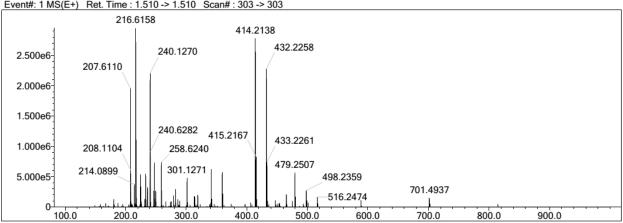




PeakTable

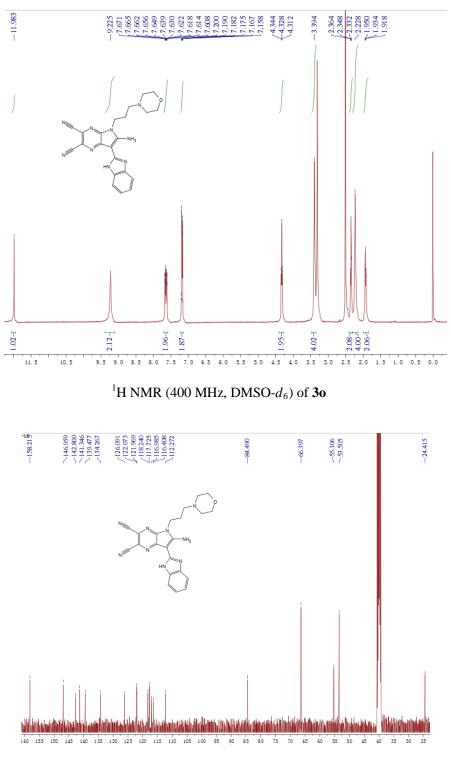
	1 cuit 1 uoiv								
PDA Ch1 254nm 4nm									
Peak# Ret. Time		Area	Height	Area %	Height %				
1	2.212	17235	1812	0.507	0.465				
2	2.428	3381663	387697	99.493	99.535				
Total		3398898	389509	100.000	100.000				

HPLC spectrum of **3n** 

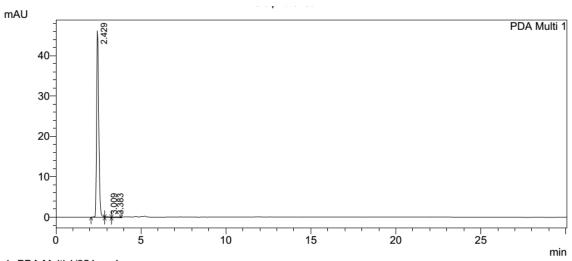


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HRMS spectrum of 3n



<sup>13</sup>C NMR (100 MHz, DMSO- $d_6$ ) of **30** 



1 PDA Multi 1/254nm 4nm

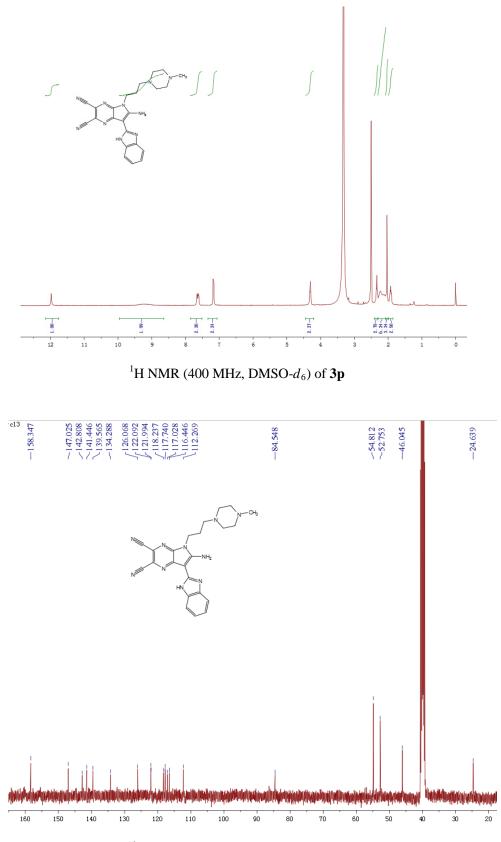
PeakTable

	Peak lable								
PDA Ch1 254nm 4nm									
Peak#	Ret. Time	Area	Height	Area %	Height %				
1	2.429	408159	46214	97.805	99.191				
2	3.009	3630	193	0.870	0.414				
3	3.383	5533	184	1.326	0.395				
Total		417321	46591	100.000	100.000				

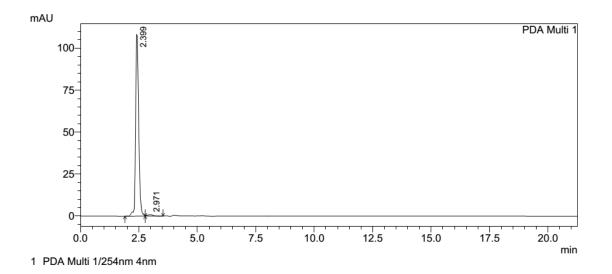
HPLC spectrum of 30

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		428.1929				
9.000e6						
8.000e6						
7.000e6						
6.000e6	214.5993					
5.000e6						
4.000e6		429.1964				
3.000e6	215.1001					
2.000e6	341.1	428.5480				
1.000e6						
0 <sup>4</sup> 100.0	<u>, , , , , , , , , , , , , , , , , , , </u>	400.0 500.0	600.0	700.0	800.0	900.0

HRMS spectrum of 30

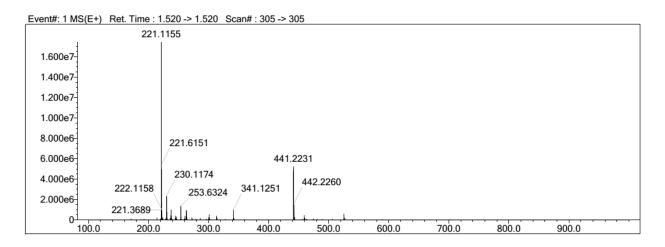


<sup>13</sup>C NMR (100 MHz, DMSO- $d_6$ ) of **3p** 

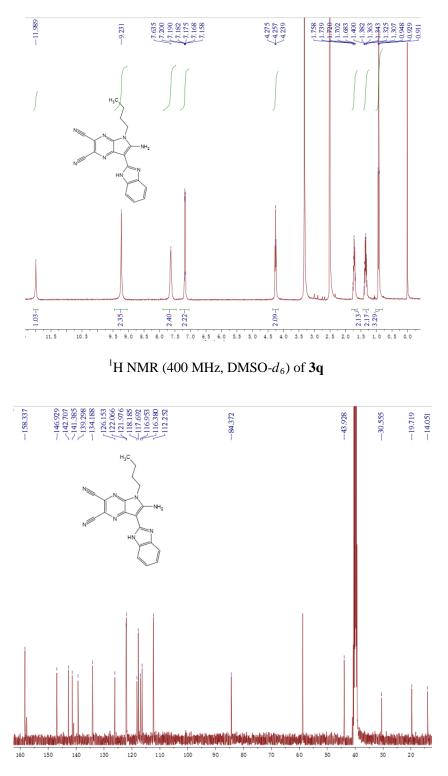


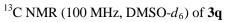
PeakTable PDA Ch1 254nm 4nm Height 108355 Ret. Time 2.399 Area % 98.425 Height % 99.239 Peak# Area 996891 15950 1 2.971 831 109186 1.575 0.761 2 1012841 100.000 100.000 Total

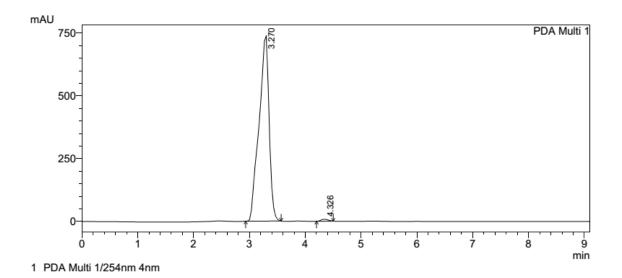
HPLC spectrum of 3p



HRMS spectrum of 3p



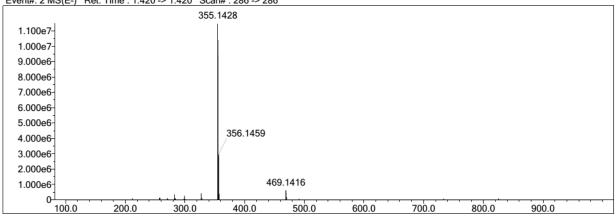




PeakTable

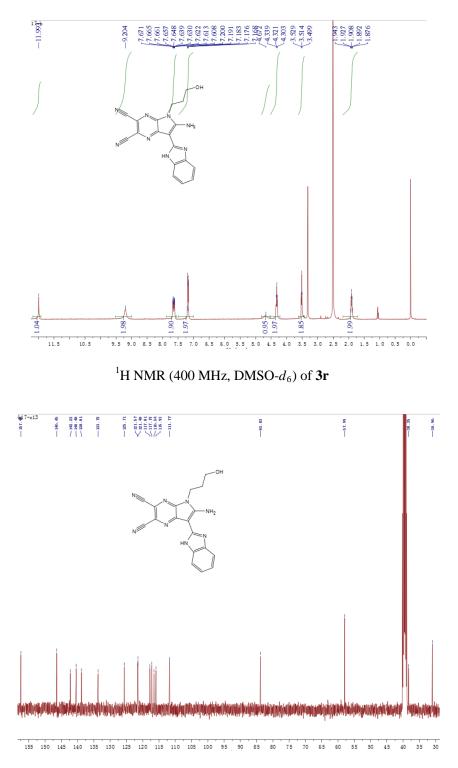
				- cuit - uc		
PDA Ch	1 254nm 41	nm				
Peak#	Ret.	Time	Area	Height	Area %	Height %
	1	3.270	9255149	738770	99.234	98.948
	2	4.326	71470	7855	0.766	1.052
То	tal		9326619	746626	100.000	100.000

HPLC spectrum	of <b>3q</b>
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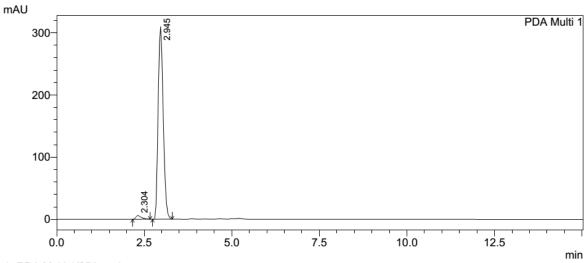


Event#: 2 MS(E-) Ret. Time : 1.420 -> 1.420 Scan# : 286 -> 286

HRMS spectrum of 3q



<sup>13</sup>C NMR (100 MHz, DMSO- $d_6$ ) of **3r** 



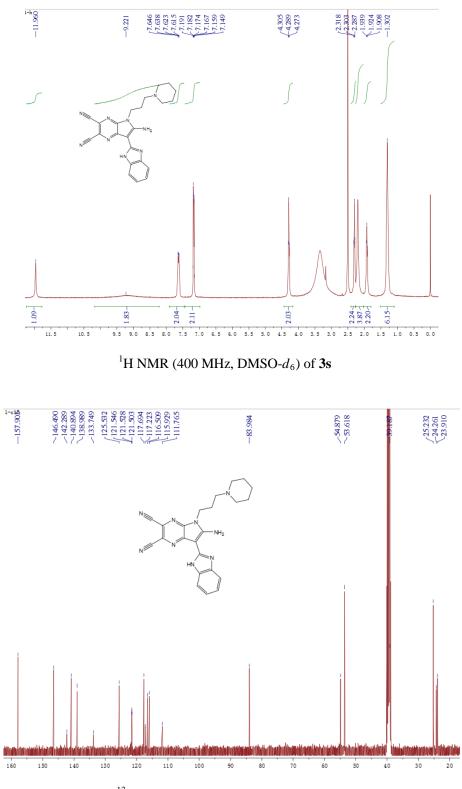
1 PDA Multi 1/254nm 4nm

	PeakTable								
PDA Ch12	PDA Ch1 254nm 4nm								
Peak#	Ret. Time	Area	Height	Area %	Height %				
1	2.304	67001	5820	2.084	1.843				
2	2.945	3147952	309961	97.916	98.157				
Total		3214953	315781	100.000	100.000				

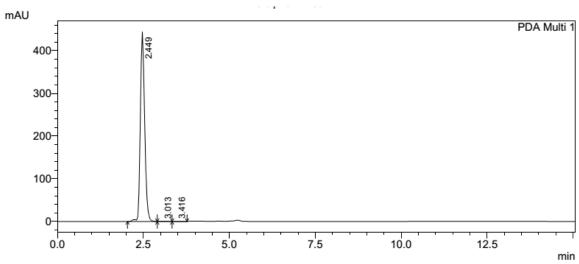
HPLC spectrum of 3r

Event#: 2 MS(E-) F	Ret. Time : 1.350 -> 1.350 Scar	# : 272 -> 27	2					
	357.	199						
9.000e6								
8.000e6								
7.000e6								
6.000e6								
5.000e6								
4.000e6								
3.000e6		358.1211						
2.000e6								
1.000e6	299.0782							
0 <sup>1</sup> 100.0	200.0 300.0	400.0	500.0	600.0	700.0	800.0	900.0	

HRMS spectrum of **3r** 



 $^{13}$ C NMR (100 MHz, DMSO- $d_6$ ) of **3s** 

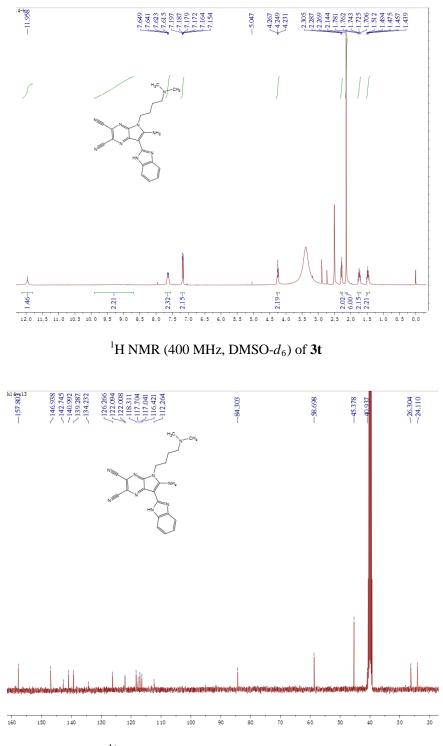


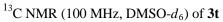
	1 Cak I able						
PDA Ch1 2	254nm 4nm						
Peak#	Ret. Time	Area	Height	Area %	Height %		
1	2.449	3890856	444573	99.647	99.786		
2	3.013	9613	630	0.246	0.141		
3	3.416	4158	322	0.106	0.072		
Total		3904627	445525	100.000	100.000		

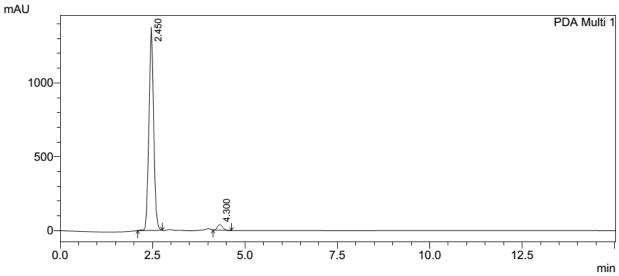
HPLC	spectrum	of	3s

Event#: 1 MS(E+) Ret.	me : 1.390 -> 1.390 Scan# : 279 -> 279
	213.6094
7.000e6	426.2128
6.000e6	
5.000e6	
4.000e6	
3.000e6	214.1056 427.2145
2.000e6	
1.000e6	
0 213.8	<u>, blablan,, b , b , b , b , b , b , b , b , b</u>
100.0	200.0 300.0 400.0 500.0 600.0 700.0 800.0 900.0

HRMS spectrum of 3s





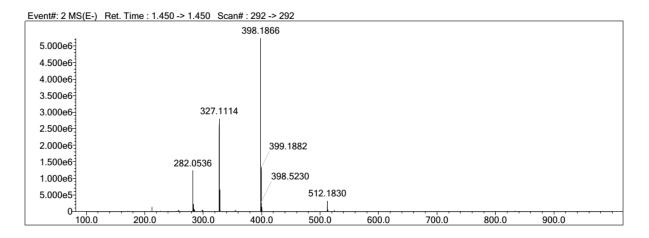


1 PDA Multi 1/254nm 4nm

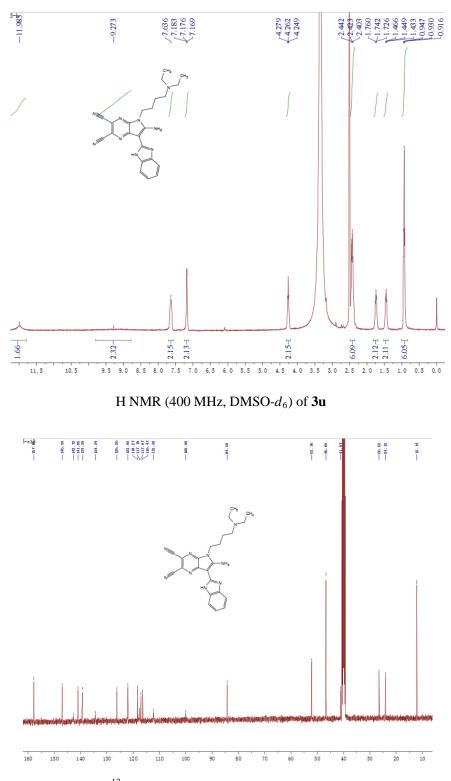
PeakTable

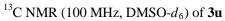
			1 cun i uoic					
PDA Ch1 254nm 4nm								
Peak#	Ret. Time	Area	Height	Area %	Height %			
1	2.450	12089458	1376805	97.099	97.391			
2	4.300	361205	36882	2.901	2.609			
Tota	1	12450663	1413687	100.000	100.000			

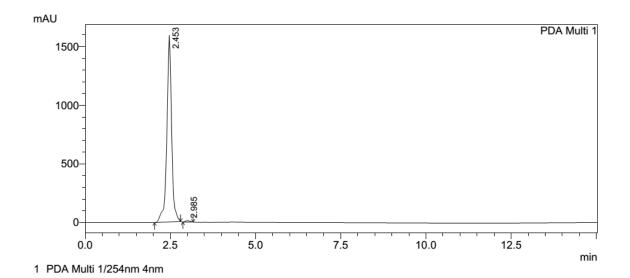
HPLC spectrum of 3t



HRMS spectrum of 3t

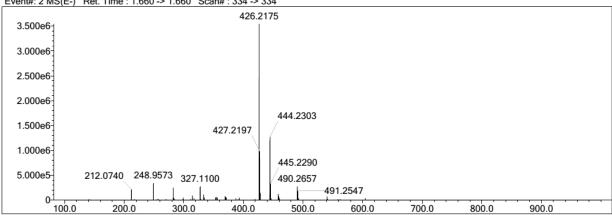






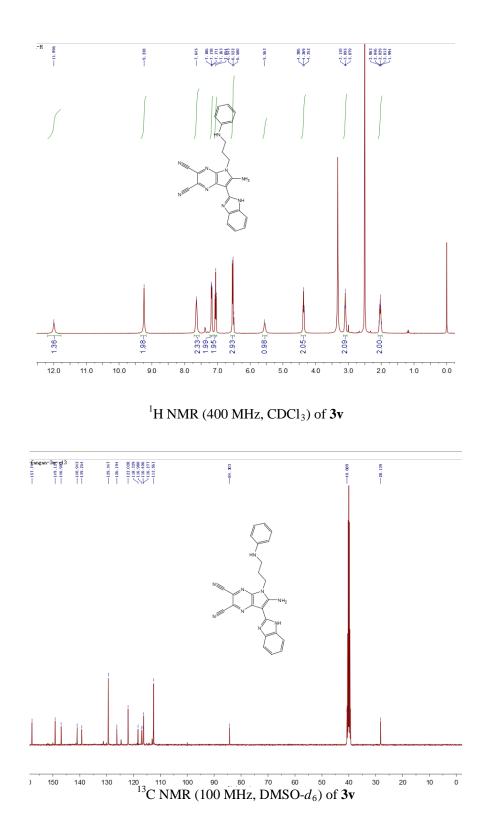
PDA Ch1 2	PDA Ch1 254nm 4nm								
Peak#	Ret. Time	Area	Height	Area %	Height %				
1	2.453	15463068	1592643	99.308	99.280				
2	2.985	107775	11550	0.692	0.720				
Total		15570843	1604193	100.000	100.000				

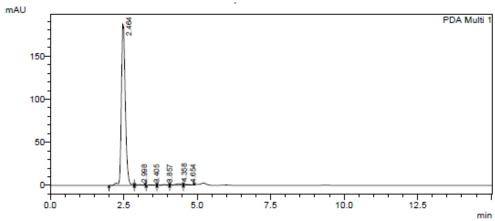
HPLC spectrum of 3u



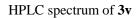
Event#: 2 MS(E-) Ret. Time : 1.660 -> 1.660 Scan# : 334 -> 334

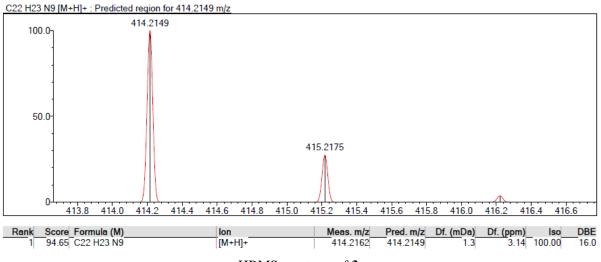
HRMS spectrum of 3u



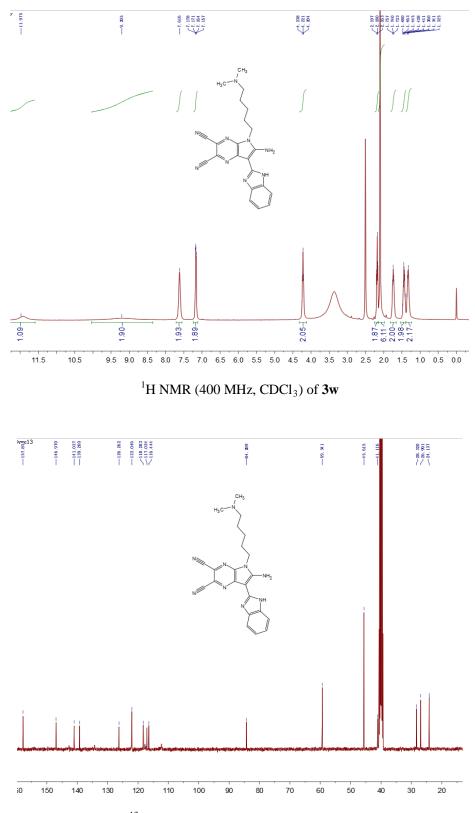


			Peaklable		
PDA Ch1 2	54nm 4nm				
Peak#	Ret. Time	Area	Height	Area %	Height %
1	2.464	1734937	187589	97.007	97.694
2	2.998	9145	706	0.511	0.368
3	3.405	3902	392	0.218	0.204
4	3.857	8535	948	0.477	0.494
5	4.358	23072	1613	1.290	0.840
6	4.654	8876	768	0.496	0.400
Total		1788467	192016	100.000	100.000

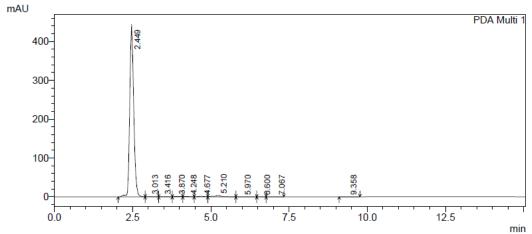






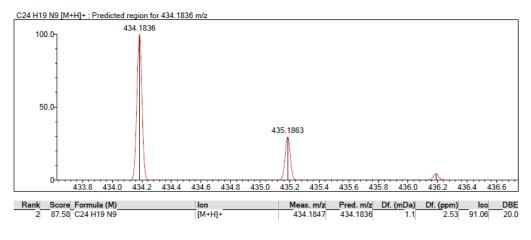


 $^{13}$ C NMR (100 MHz, DMSO- $d_6) of <math display="inline">\mathbf{3w}$ 

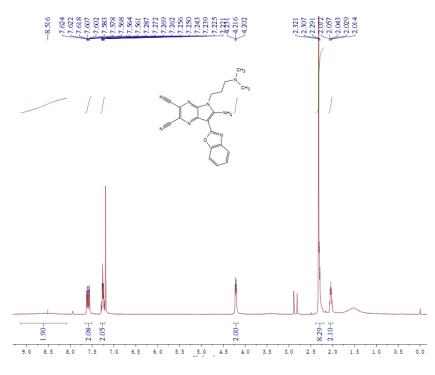


			PeakTable		
PDA Ch1 2	54nm 4nm				
Peak#	Ret. Time	Area	Height	Area %	Height %
1	2.449	3892143	444597	96.900	98.236
2	3.013	11222	686	0.279	0.152
3	3.416	6412	401	0.160	0.089
4	3.870	13162	1085	0.328	0.240
5	4.248	12897	786	0.321	0.174
6	4.677	18664	1156	0.465	0.255
7	5.210	50522	3193	1.258	0.705
8	5.970	4559	234	0.113	0.052
9	6.600	1607	139	0.040	0.031
10	7.067	2102	88	0.052	0.019
11	9.358	3364	217	0.084	0.048
Total		4016654	452581	100.000	100.000

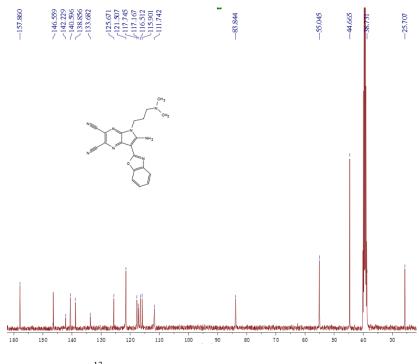
HPLC spectrum of 3w



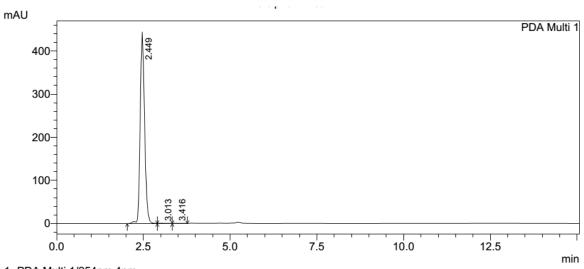
HRMS spectrum of 3w



<sup>1</sup>H NMR (400 MHz, CDCl<sub>3</sub>) of 4m



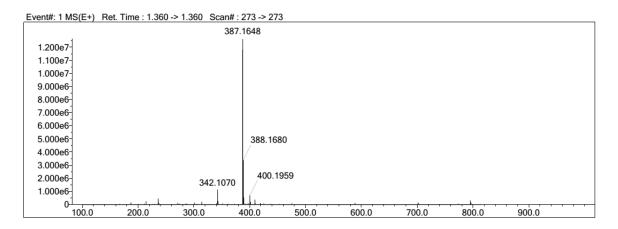
 $^{13}$ C NMR (100 MHz, DMSO- $d_6$ ) of **4m** 



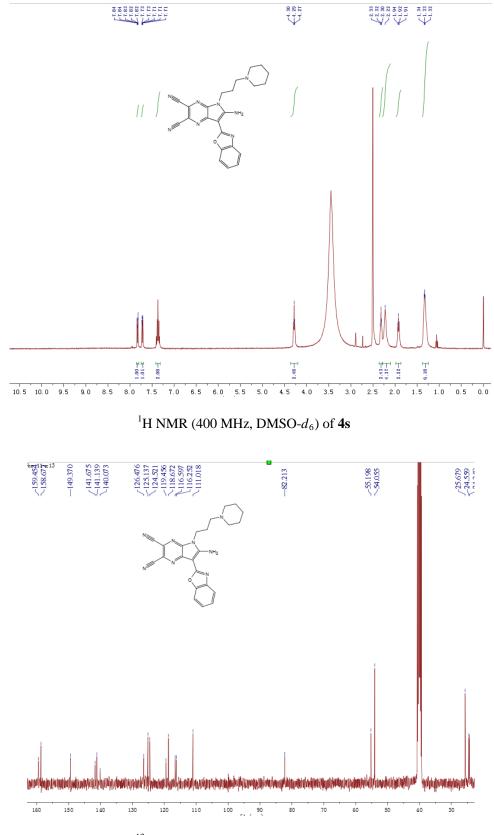
1 PDA Multi 1/254nm 4nm

				1	currituoie	
1	PDA Ch1 2	254nm 4nm				
ſ	Peak#	Ret. Time	Area	Height	Area %	Height %
	1	2.449	3890856	444573	99.647	99.786
[	2	3.013	9613	630	0.246	0.141
	3	3.416	4158	322	0.106	0.072
	Total		3904627	445525	100.000	100.000

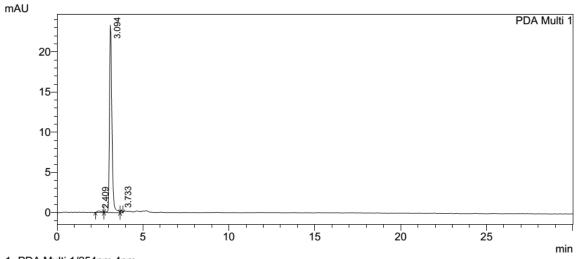
HPLC spectrum of 4m



HRMS spectrum of 4m



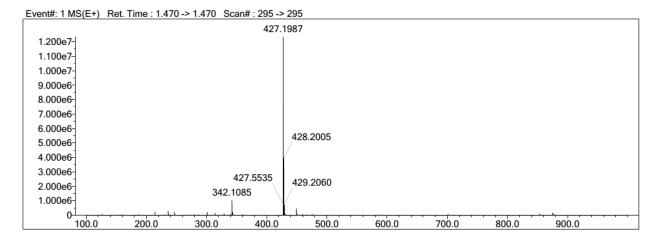
 $^{13}$ C NMR (100 MHz, DMSO- $d_6$ ) of **4s** 



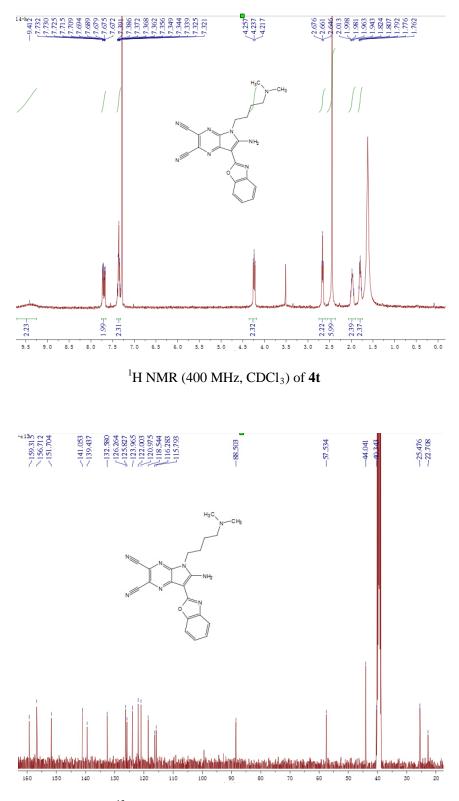
1 PDA Multi 1/254nm 4nm

			PeakTable				
PDA Ch1 2	PDA Ch1 254nm 4nm						
Peak#	Ret. Time	Area	Height	Area %	Height %		
1	2.409	3074	207	1.309	0.868		
2	3.094	230303	23333	98.083	98.004		
3	3.733	1428	269	0.608	1.128		
Total		234804	23808	100.000	100.000		

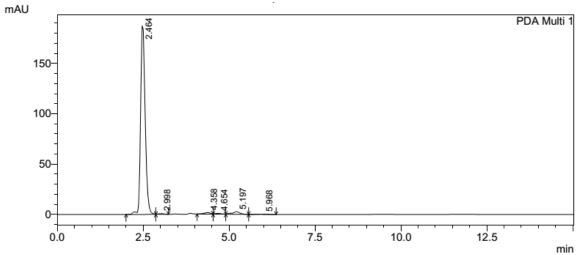
HPLC spectrum of 4s



HRMS spectrum of 4s

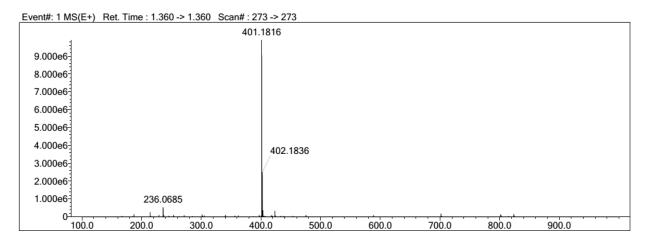


<sup>13</sup>C NMR (100 MHz, DMSO- $d_6$ ) of **4**t

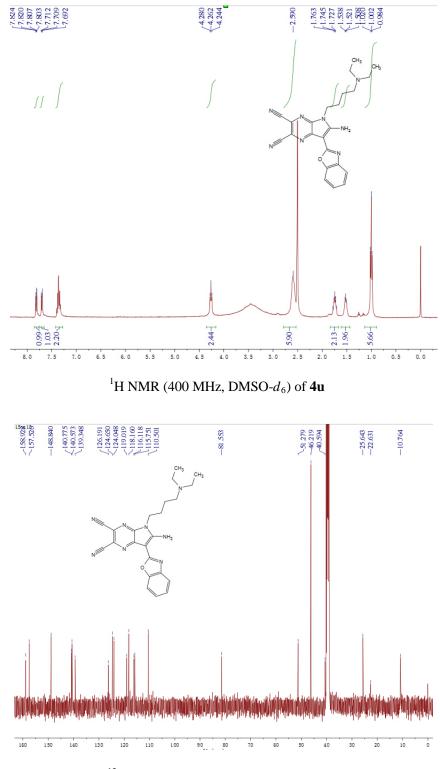


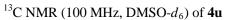
				PeakTa	able	
Р	DA Ch1 2	254nm 4nm				
	Peak#	Ret. Time	Area	Height	Area %	Height %
Γ	1	2.464	1734937	187589	95.334	96.871
Г	2	2.998	9145	706	0.502	0.365
Γ	3	4.358	20626	1550	1.133	0.801
Γ	4	4.654	10929	834	0.601	0.431
Γ	5	5.197	38972	2581	2.142	1.333
	6	5.968	5234	387	0.288	0.200
	Total		1819842	193648	100.000	100.000

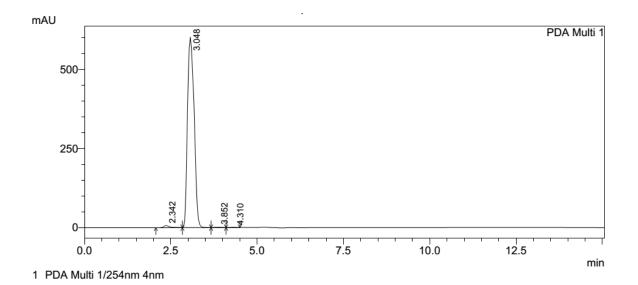
HPLC spectrum of 4t



HRMS spectrum of 4t

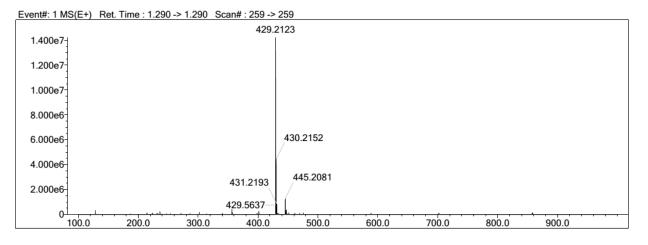




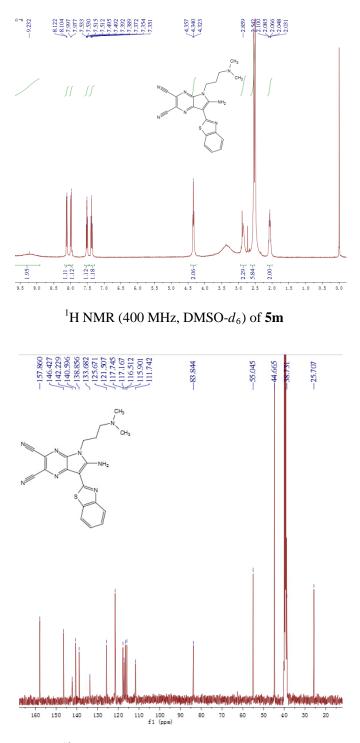


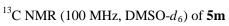
PeakTable PDA Ch1 254nm 4nm nm 41111 Ret. Time 2.342 Height % 1.136 Peak# Area Height Area % 84960 6950 1.045 2 3.048 8018747 602527 98.603 98.452 0.241 3 3.852 16423 1472 0.202 4 4.310 12259 1055 0.151 0.172 8132389 612004 100.000 Total 100.000

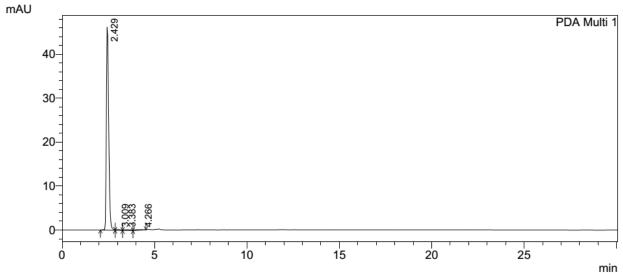
<sup>1</sup>H NMR (400 MHz, DMSO- $d_6$ ) of **4u** 



HRMS spectrum of 4u





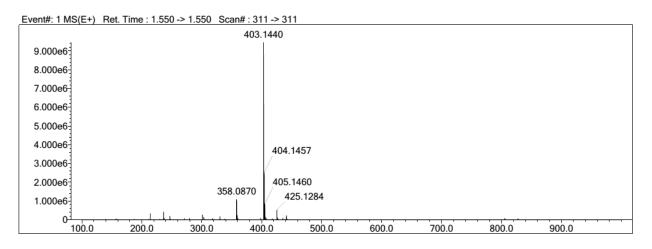


1 PDA Multi 1/254nm 4nm

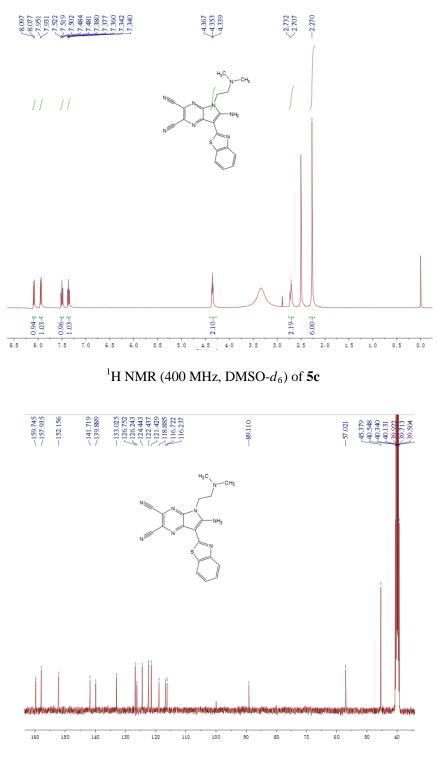
D			. 1	
Pea	κI	a	bl	e

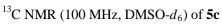
	i cuit i doite					
PDA Ch1 2	254nm 4nm					
Peak#	Ret. Time	Area	Height	Area %	Height %	
1	2.429	408159	46214	96.806	98.926	
2	3.009	3630	193	0.861	0.413	
3	3.383	5533	184	1.312	0.394	
4	4.266	4304	125	1.021	0.267	
Total		421625	46716	100.000	100.000	

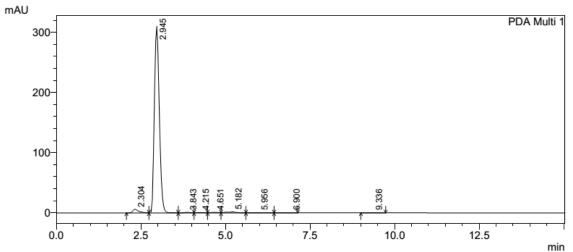
## HPLC spectrum of 5m



HRMS spectrum of 5m



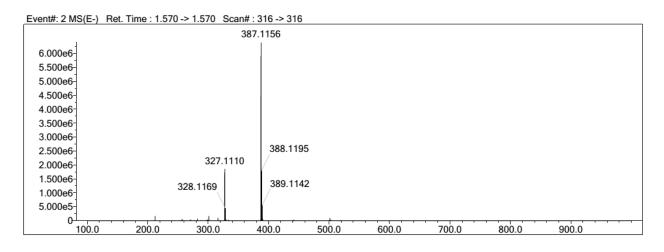




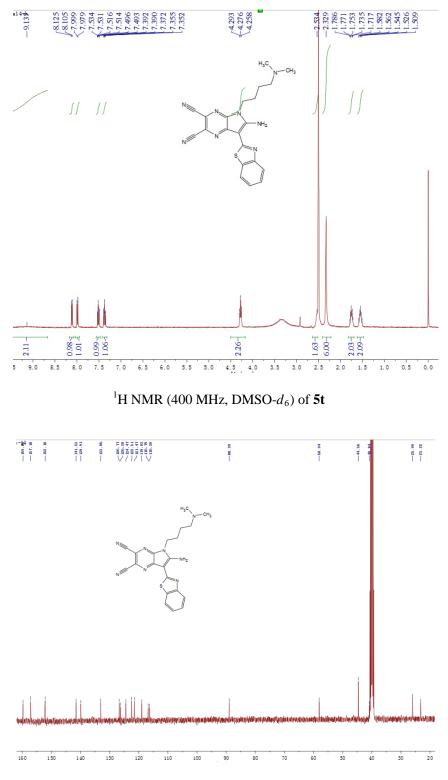
1	PDA	Multi	1/254nm 4nm	
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		PeakTable				
PDA Ch1 2	54nm 4nm					
Peak#	Ret. Time	Area	Height	Area %	Height %	
1	2.304	73831	5963	2.226	1.857	
2	2.945	3164434	310263	95.387	96.639	
3	3.843	12101	1093	0.365	0.340	
4	4.215	8685	512	0.262	0.159	
5	4.651	14401	950	0.434	0.296	
6	5.182	32280	1661	0.973	0.517	
7	5.956	4226	218	0.127	0.068	
8	6.900	1836	62	0.055	0.019	
9	9.336	5685	331	0.171	0.103	
Total		3317480	321054	100.000	100.000	

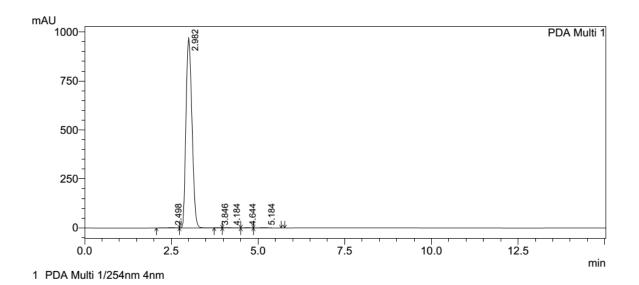
HPLC	spectrum	of	5c
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HRMS spectrum of 5c

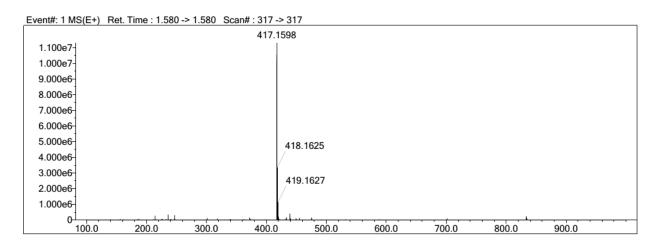


<sup>13</sup>C NMR (100 MHz, DMSO-*d*<sub>6</sub>) of **5**t

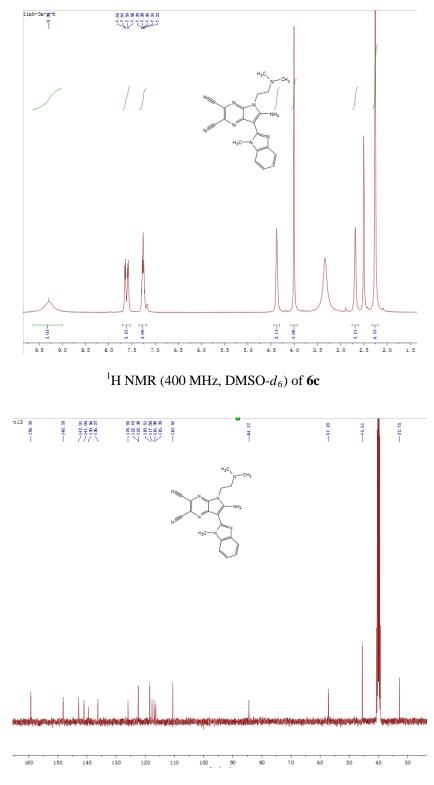


PeakTable PDA Ch1 254nm 4nm Peak# Ret. Time Height Area % Height % Area 2.498 29389 1521 0.248 0.155 1 23 2.982 11737626 973178 99.162 99.408 3.846 0.108 9695 1053 0.082 4 4.184 19081 733 0.161 0.075 4.644 12833 926 0.095 5 0.108 6 5.184 28148 1558 0.238 0.159 Total 11836772 978969 100.000 100.000

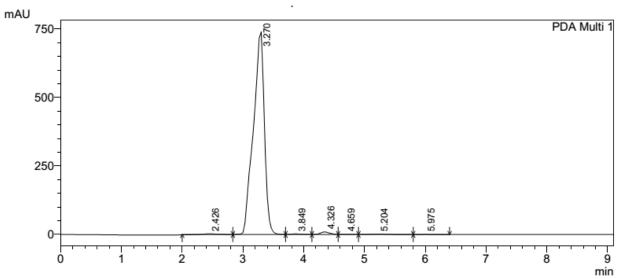
## HPLC spectrum of 5t



HRMS spectrum of 5t



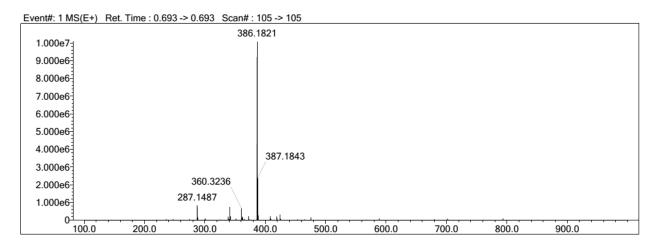
<sup>13</sup>C NMR (100 MHz, DMSO- $d_6$ ) of **6c** 



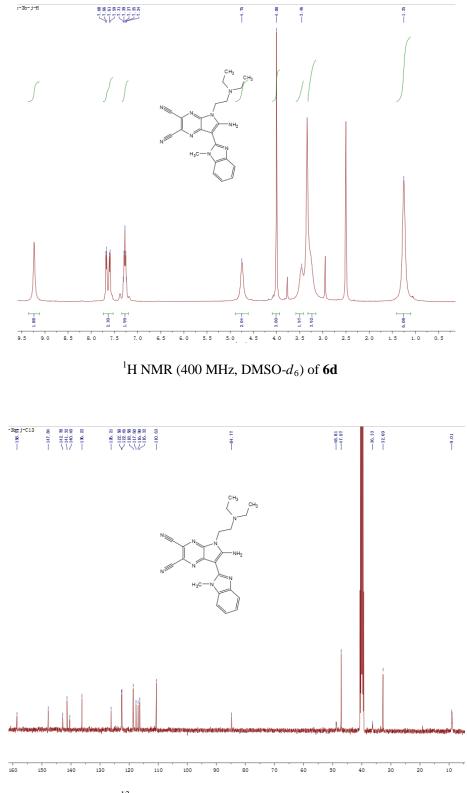
1 PDA Multi 1/254nm 4nm

		PeakTable				
PDA Ch1 2	54nm 4nm					
Peak#	Ret. Time	Area	Height	Area %	Height %	
1	2.426	41024	2562	0.429	0.338	
2	3.270	9328333	740436	97.556	97.702	
3	3.849	33990	2352	0.355	0.310	
4	4.326	101766	9085	1.064	1.199	
5	4.659	15929	1131	0.167	0.149	
6	5.204	33863	1748	0.354	0.231	
7	5.975	7119	539	0.074	0.071	
Total		9562024	757853	100.000	100.000	

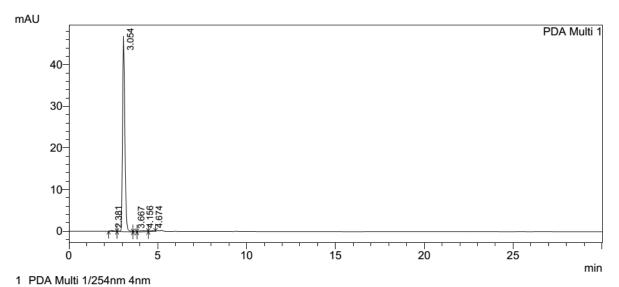
HPLC	spectrum	of	6c
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HRMS spectrum of 6c

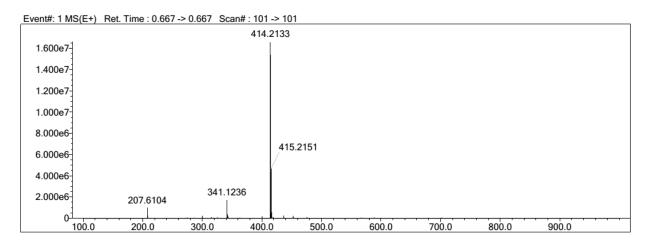


<sup>13</sup>C NMR (100 MHz, DMSO- $d_6$ ) of **6d** 

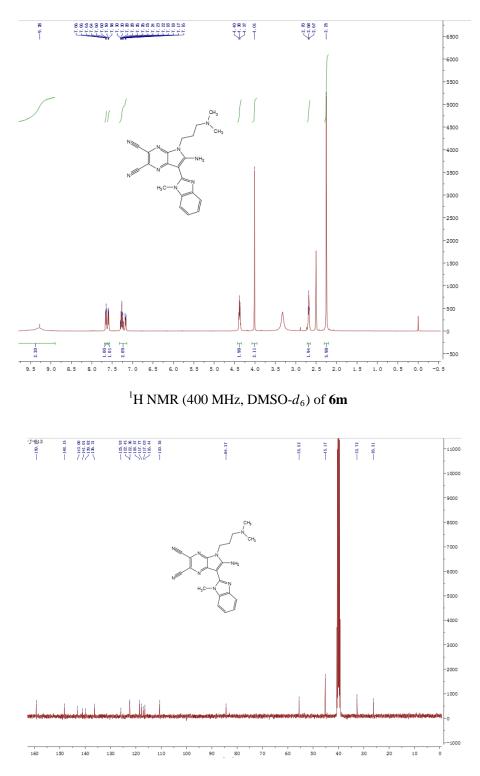


			PeakTable		
PDA Ch1 2	54nm 4nm				
Peak#	Ret. Time	Area	Height	Area %	Height %
1	2.381	3094	168	0.702	0.353
2	3.054	426908	46944	96.880	98.508
3	3.667	2292	186	0.520	0.391
4	4.156	6389	226	1.450	0.473
5	4.674	1973	131	0.448	0.275
Total		440657	47655	100.000	100.000

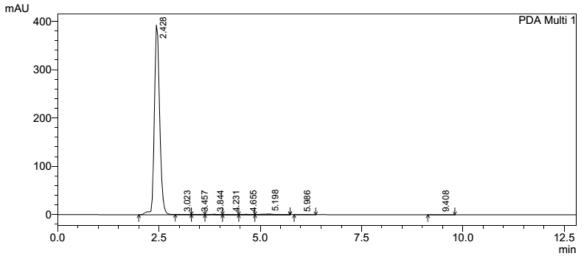
HPLC spectrum of 6d







 $^{13}$ C NMR (100 MHz, DMSO- $d_6$ ) of **6m** 



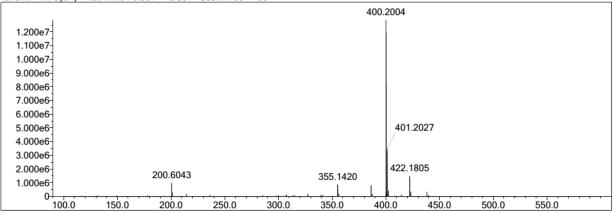
1 PDA Multi 1/254nm 4nm

PDA Ch1 254nm 4nm							
Peak#	Ret. Time	Area	Height	Area %	Height %		
1	2.428	3588432	393141	98.049	98.765		
2	3.023	4612	273	0.126	0.068		
3	3.457	1937	175	0.053	0.044		
4	3.844	9251	1141	0.253	0.287		
5	4.231	6626	414	0.181	0.104		
6	4.655	13461	903	0.368	0.227		
7	5.198	28810	1521	0.787	0.382		
8	5.986	3554	287	0.097	0.072		
9	9.408	3151	204	0.086	0.051		
Total		3659834	398058	100.000	100.000		

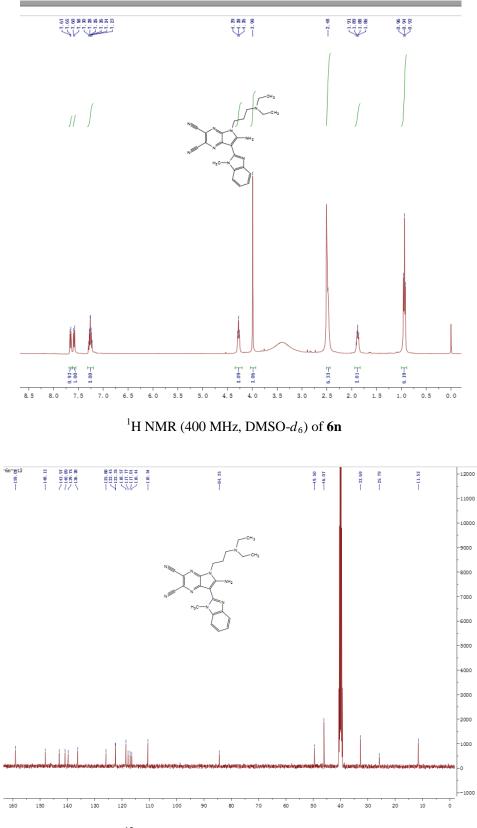
PeakTable

## HPLC spectrum of 6m

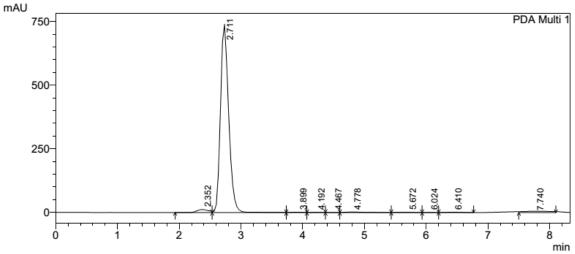
Event#: 1 MS(E+) Ret. Time : 0.587 -> 0.587 Scan# : 89 -> 89



HRMS spectrum of 6m



<sup>13</sup>C NMR (100 MHz, DMSO- $d_6$ ) of **6n** 

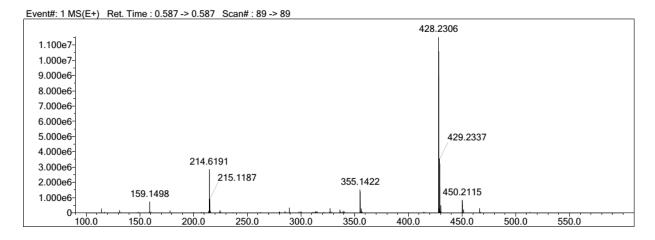


1 PDA Multi 1/254nm 4nm

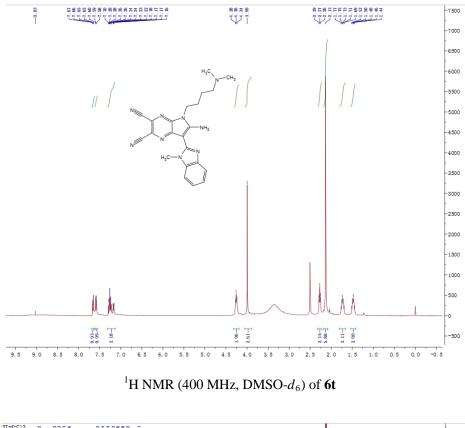
PDA Ch1 254nm 4nm Peak# Height Height % Ret. Time Area Area % 153924 6838316 11518 740524 2.172 96.490 1.518 97.592 2.352 2.711 1 2 3 3.899 15755 1416 0.222 0.187 4 4.192 4827 362 0.068 0.048 5 4.467 3319 295 0.047 0.039 4.778 2411 0.576 6 40815 0.318 7 5.672 9379 708 0.132 0.093 0.022 0.020 8 1571 152 6.024 6.410 7.740 9 17720 1332 0.250 0.176 10 1424 80 0.020 0.011 7087052 758798 100.000 100.000 Total

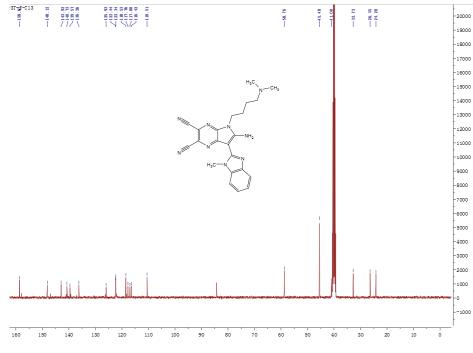
PeakTable

## HPLC spectrum of 6n

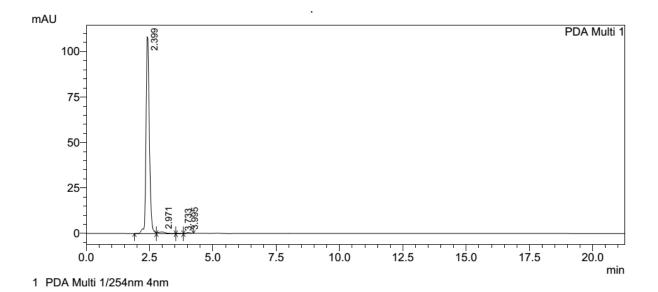


HRMS spectrum of 6n



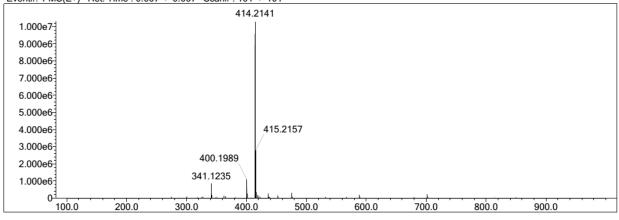


<sup>13</sup>C NMR (100 MHz, DMSO- $d_6$ ) of **6t** 



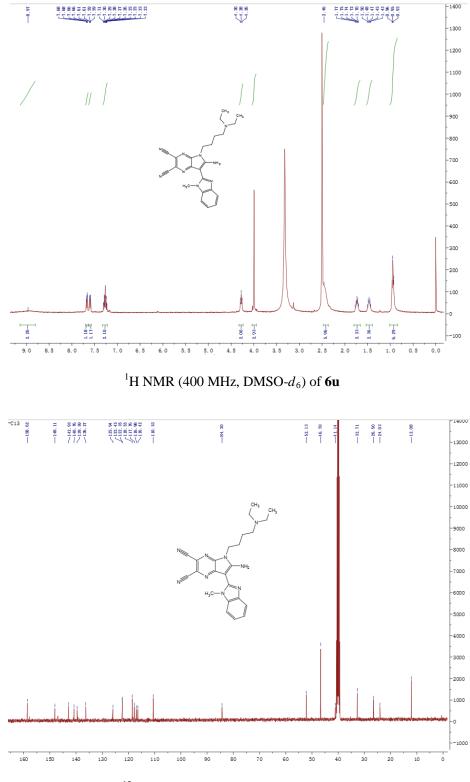
PDA Ch1 2	54nm 4nm		Peak	Table	
Peak#	Ret. Time	Area	Height	Area %	Height %
1	2.399	998863	108398	97.109	98.411
2	2.971	20982	925	2.040	0.840
3	3.733	2509	248	0.244	0.225
4	3.995	6249	577	0.608	0.524
Total		1028603	110149	100.000	100.000

HPLC spectrum of 6t

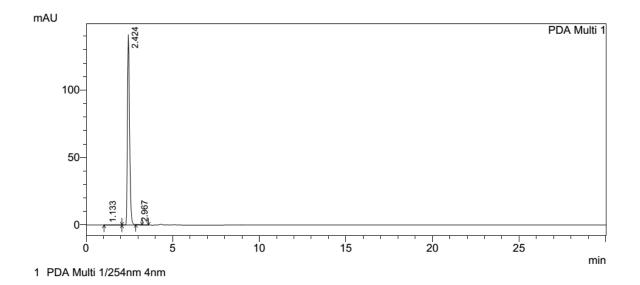


Event#: 1 MS(E+) Ret. Time : 0.667 -> 0.667 Scan# : 101 -> 101

HRMS spectrum of 6t



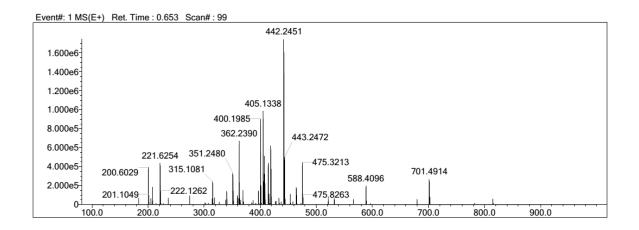
 $^{13}$ C NMR (100 MHz, DMSO- $d_6) of$ **6u** 



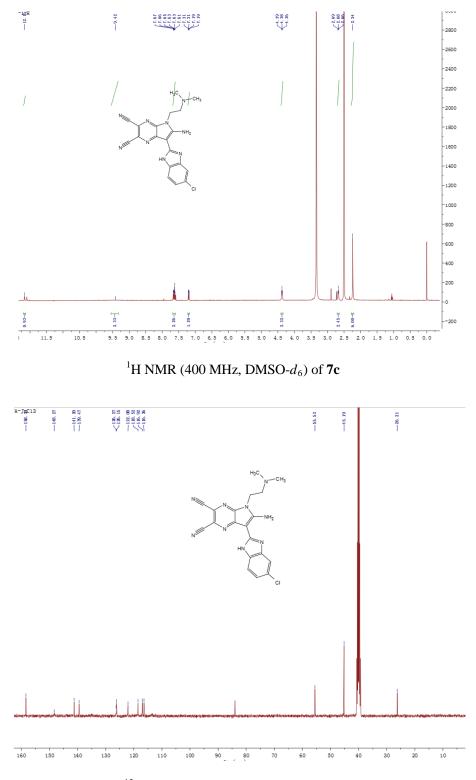
PeakTable

			1 cuit 1 uote						
Pl	DA Ch1 2	254nm 4nm							
	Peak#	Ret. Time	Area	Height	Area %	Height %			
	1	1.133	1063	20	0.087	0.014			
	2	2.424	1216952	141127	99.815	99.891			
Γ	3	2.967	1199	134	0.098	0.095			
	Total		1219213	141282	100.000	100.000			

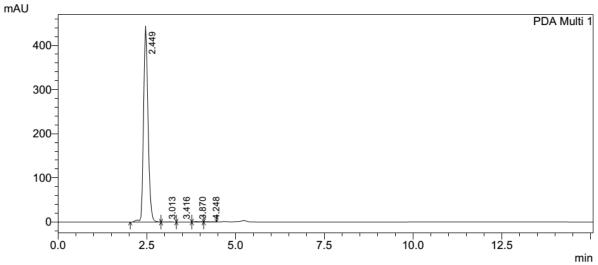
HPLC spectrum of 6u



HRMS spectrum of 6u



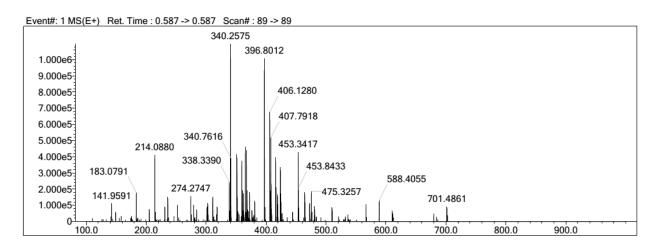
<sup>13</sup>C NMR (100 MHz, DMSO- $d_6$ ) of **7c** 



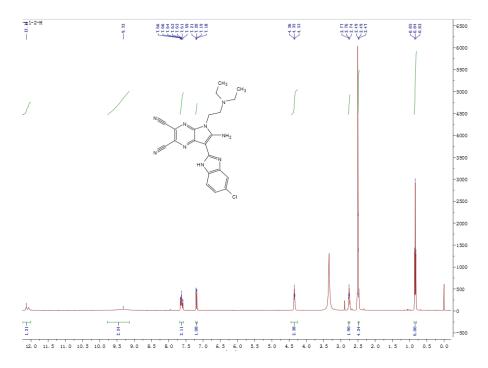
1 PDA Multi 1/254nm 4nm

	PeakTable							
PDA Ch1 2	54nm 4nm							
Peak#	Ret. Time	Area	Height	Area %	Height %			
1	2.449	3890856	444573	99.214	99.456			
2	3.013	9613	630	0.245	0.141			
3	3.416	4158	322	0.106	0.072			
4	3.870	10086	952	0.257	0.213			
5	4.248	6981	529	0.178	0.118			
Total		3921694	447005	100.000	100.000			

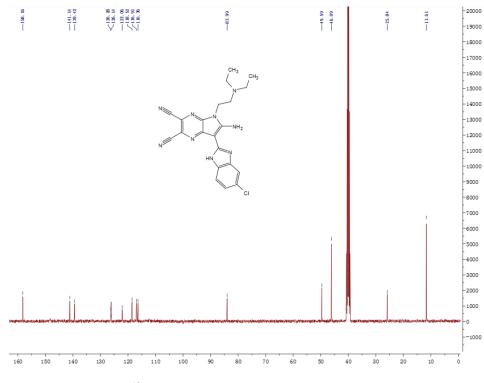
HPLC spectrum of 7c



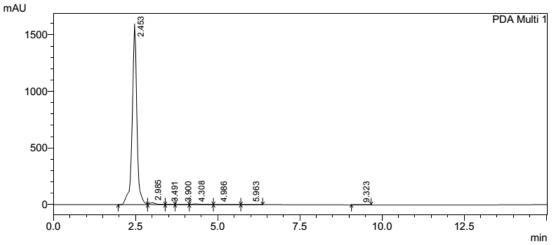
HRMS spectrum of 7c



<sup>1</sup>H NMR (400 MHz, DMSO- $d_6$ ) of **7d** 



 $^{13}$ C NMR (100 MHz, DMSO- $d_6) of 7d$ 

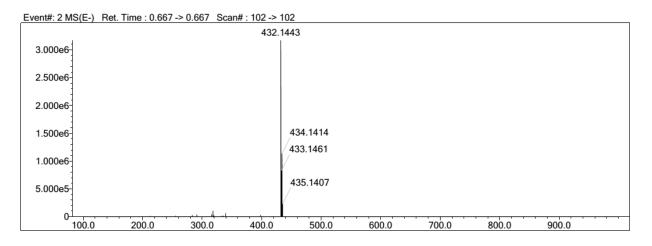


1 PDA Multi 1/254nm 4nm

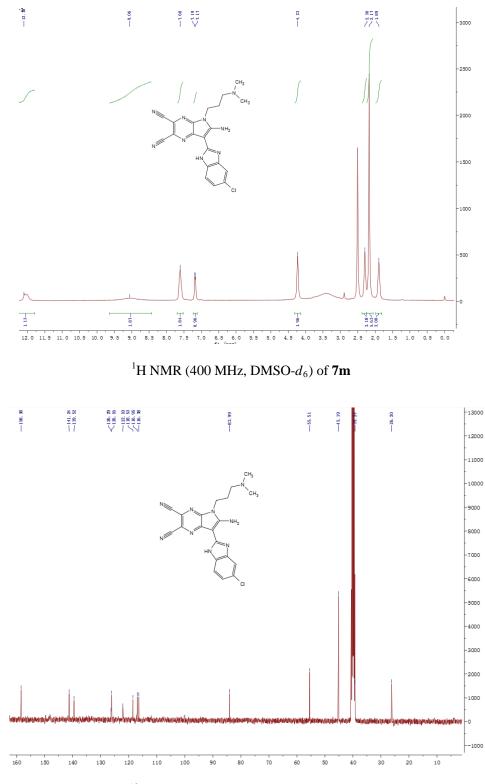
PeakTable

	FCakTable						
PDA Ch1 2	54nm 4nm						
Peak#	Ret. Time	Area	Height	Area %	Height %		
1	2.453	15726222	1598355	97.058	98.155		
2	2.985	228710	16469	1.412	1.011		
3	3.491	33841	3525	0.209	0.216		
4	3.900	48981	2063	0.302	0.127		
5	4.308	129187	6207	0.797	0.381		
6	4.986	28626	1310	0.177	0.080		
7	5.963	4118	265	0.025	0.016		
8	9.323	3205	203	0.020	0.012		
Total		16202891	1628398	100.000	100.000		

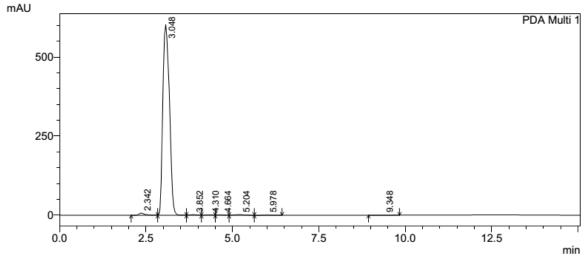
HPLC spectrum of 7d



HRMS spectrum of 7d



<sup>13</sup>C NMR (100 MHz, DMSO- $d_6$ ) of **7m** 

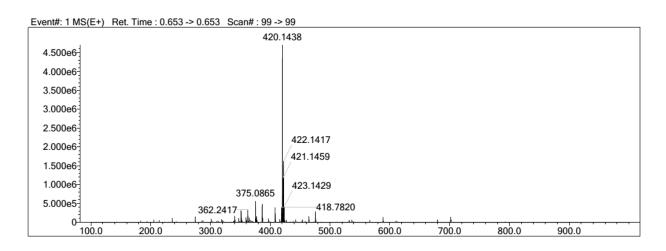


1 PDA Multi 1/254nm 4nm

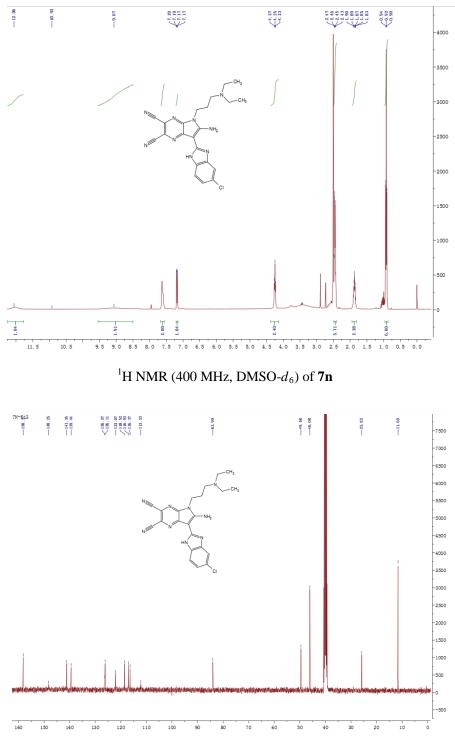
PeakTable

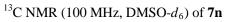
	1 curt i uote							
PDA Ch1 2	54nm 4nm							
Peak#	Ret. Time	Area	Height	Area %	Height %			
1	2.342	87885	6996	1.068	1.134			
2	3.048	8028562	602690	97.587	97.734			
3	3.852	24258	1768	0.295	0.287			
4	4.310	21150	1427	0.257	0.231			
5	4.664	17470	1100	0.212	0.178			
6	5.204	37044	2044	0.450	0.332			
7	5.978	4770	272	0.058	0.044			
8	9.348	5919	367	0.072	0.060			
Total		8227057	616665	100.000	100.000			

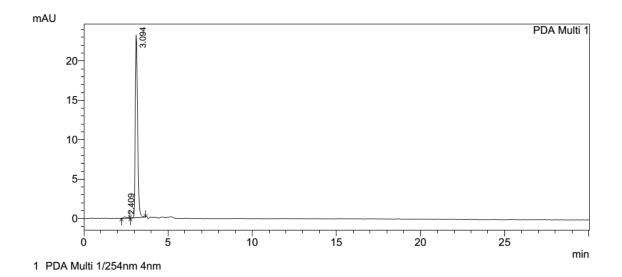
HPLC spectrum of 7m



HRMS spectrum of **7m** 





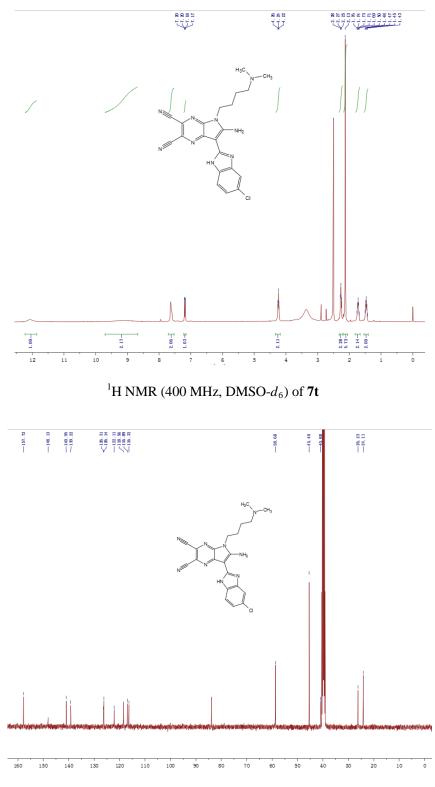


	PeakTable						
PDA Ch1 2	54nm 4nm						
Peak#	Ret. Time	Area	Height	Area %	Height %		
1	2.409	2480	193	1.096	0.823		
2	3.094	223739	23234	98.904	99.177		
Total		226220	23427	100.000	100.000		

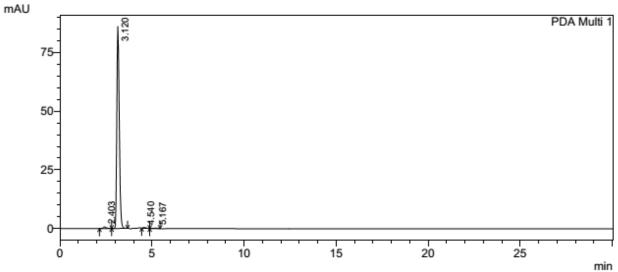
HPLC spectrum of 7n

Event#: 1 MS(E+) F	Ret. Time : 0.680 -> 0.680 Sc	an# : 103 ->	103				
		44	8.1744				
6.000e6							
5.500e6-							
5.000e6							
4.500e6-							
4.000e6-							
3.500e6			450 1715				
3.000e6			450.1715				
2.500e6			449.1754				
2.000e6-			V				
1.500e6	256.6161		451.1716				
1.000e6-	257.6154 351.2494	375.0858	431.1710				
5.000e5	257.1174	al de	Υ				
o <sup>1</sup> ,,	اهم به	المواسات بالل	, <b>b</b> ,b,				
100.0	200.0 300.0	400.0	500.0	600.0	700.0	800.0	900.0

HRMS spectrum of 7n



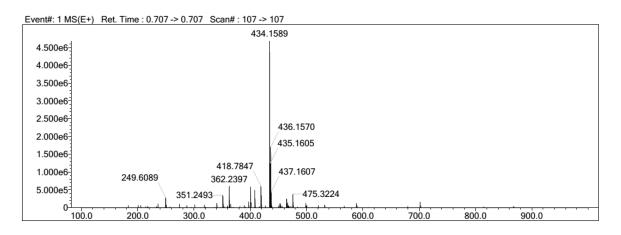
 $^{13}$ C NMR (100 MHz, DMSO- $d_6) of 7t$ 

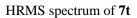


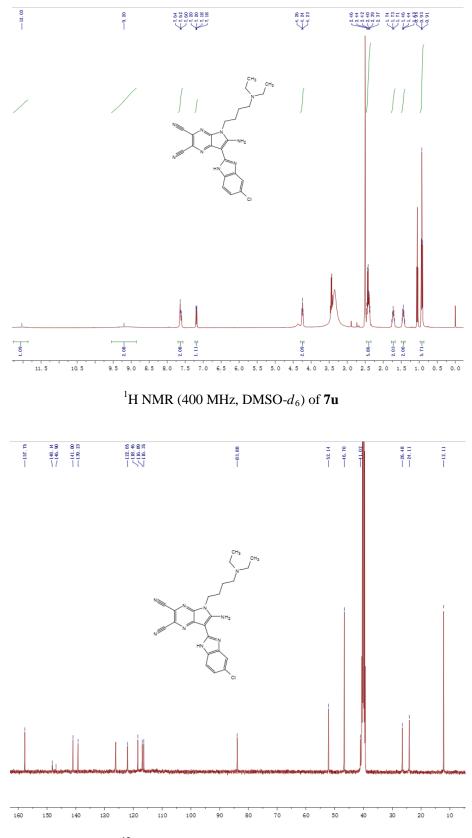
1 PDA Multi 1/254nm 4nm

PDA Ch1 2	254nm 4nm		PeakTa	ble	
Peak#	Ret. Time	Area	Height	Area %	Height %
1	2.403	9695	734	1.182	0.840
2	3.120	804021	86114	97.992	98.557
3	4.540	3854	349	0.470	0.399
4	5.167	2923	178	0.356	0.204
Total		820493	87375	100.000	100.000

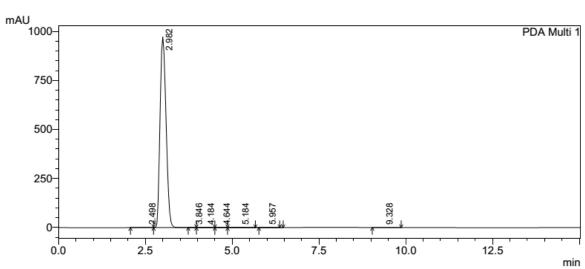
## HPLC spectrum of 7t







 $^{13}$ C NMR (100 MHz, DMSO- $d_6) of 7u$ 

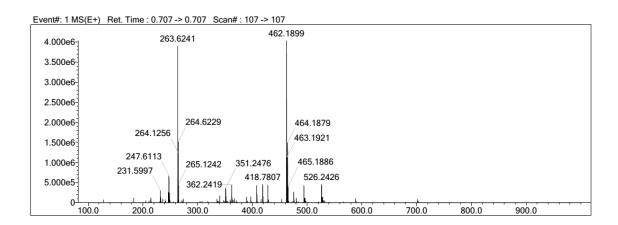


1 PDA Multi 1/254nm 4nm

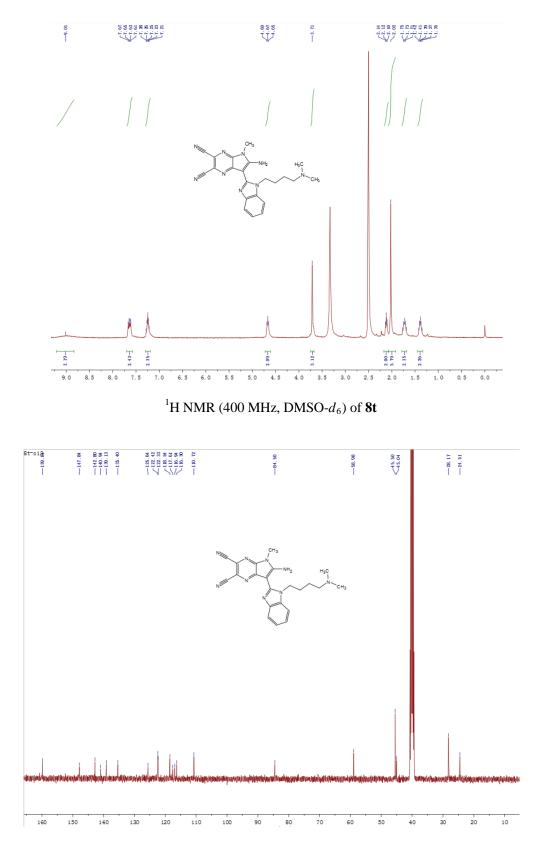
PeakTable

	1 cak I able							
PDA Ch1 2	54nm 4nm							
Peak#	Ret. Time	Area	Height	Area %	Height %			
1	2.498	29509	1525	0.249	0.156			
2	2.982	11742145	973187	99.017	99.282			
3	3.846	9695	1053	0.082	0.107			
4	4.184	19081	733	0.161	0.075			
5	4.644	12833	926	0.108	0.094			
6	5.184	28148	1558	0.237	0.159			
7	5.957	8240	660	0.069	0.067			
8	9.328	9055	582	0.076	0.059			
Total		11858706	980224	100.000	100.000			

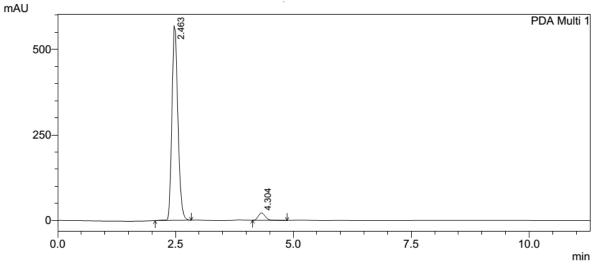
## HPLC spectrum of 7u



HRMS spectrum of 7u



 $^{13}$ C NMR (100 MHz, DMSO- $d_6$ ) of 8t

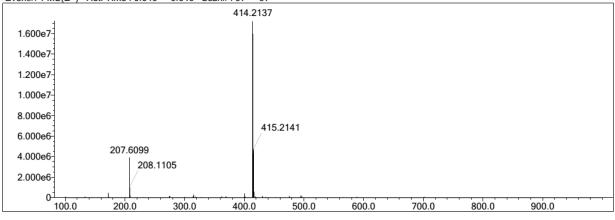


1 PDA Multi 1/254nm 4nm

PeakTable

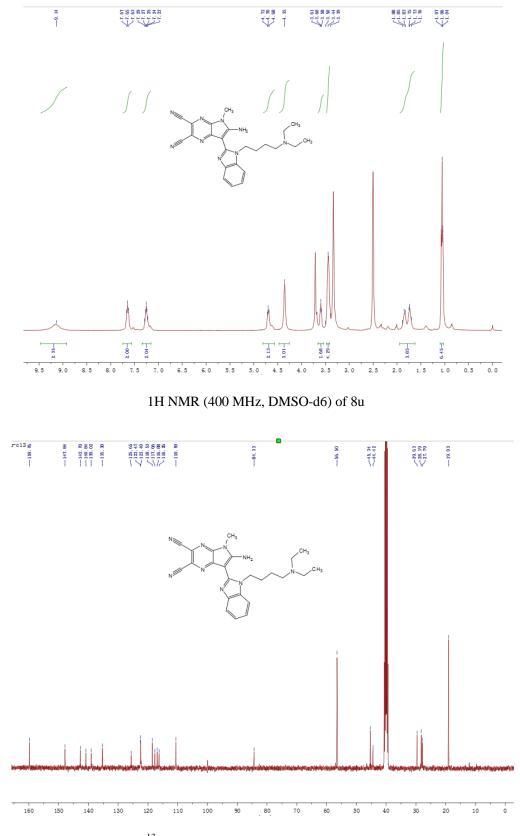
	1 cux 1 uote						
PDA Ch1 2	254nm 4nm						
Peak#	Ret. Time	Area	Height	Area %	Height %		
1	2.463	5034926	569781	95.688	96.370		
2	4.304	226885	21461	4.312	3.630		
Total		5261811	591242	100.000	100.000		

HPLC spectrum of 8t

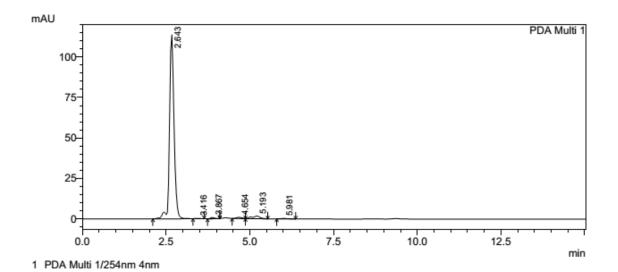


Event#: 1 MS(E+) Ret. Time : 0.640 -> 0.640 Scan# : 97 -> 97

HRMS spectrum of 8t

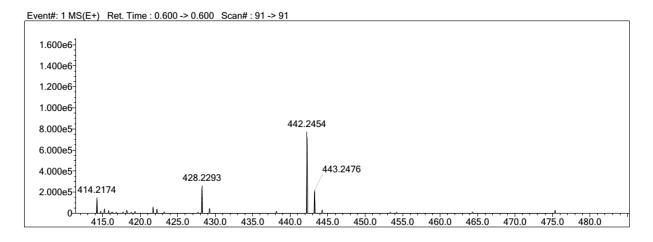


 $^{13}$ C NMR (100 MHz, DMSO- $d_6$ ) of 8u



PeakTable PDA Ch1 254nm 4nm Height 113793 Peak# Ret. Time Area % Height % Area 1068728 95.374 2.643 96.753 23 1679 0.168 3.416 198 0.150 7330 867 0.654 0.737 3.867 4.654 9359 751 0.835 4 0.639 1.424 0.279 5 5.193 28940 1675 2.583 6 5.981 4526 328 117612 0.404 1120562 100.000 Total 100.000

## HPLC spectrum of 8u



HRMS spectrum of 8u

## References

- [1] Liu, L. DNA toisomerase poisons as antitumor drugs. Annu. Rev. Biochem. 1989, 58, 351-375.
- [2] Zhuo, S. T.; Li, C. Y.; Hu, M. H.; Chen, S. B.; Yao, P. F.; Huang, S. L.; Ou, T. M.; Tan, J. H.; An, L. K.; Li, D.; Gu, L. Q.; Huang, Z. S. Synthesis and biological evaluation of benzo[a]-phenazine derivatives as a dual inhibitor of topoisomerase I and II. *Org. Biomol. Chem.* 2013, *11*, 3989–4005
- [3] Sheng, C. Q.; Miao, Z.Y.; Zhang, W. N. New strategies in the discovery of novel non-camptothecin topoisomerase I Inhibitors. *Curr. Med. Chem.* 2011, 18, 4389–4409.
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- [5] Janson T. B., Kang, Z.L., Austin, C.A., Kurz, E.U. Salicylate, a catalytic inhibitor of Topoisomerase II, inhibitors DNA cleavage and selective for the α isoform. *Mol Parmacol*, 2014, 85, 198-207.