

Identification of Gallic acid based Glycoconjugates as a novel anti-tumor agents targeting tubulin polymerization

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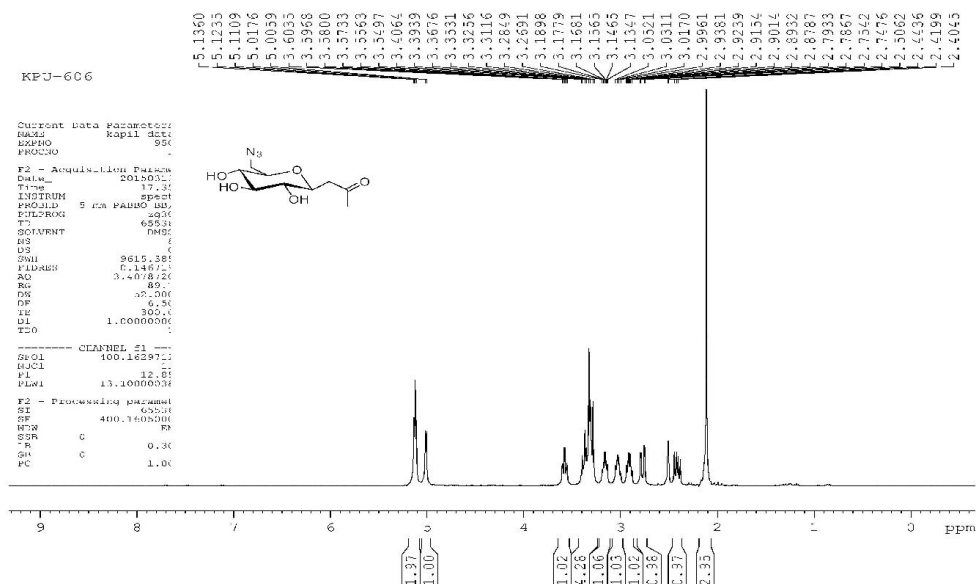
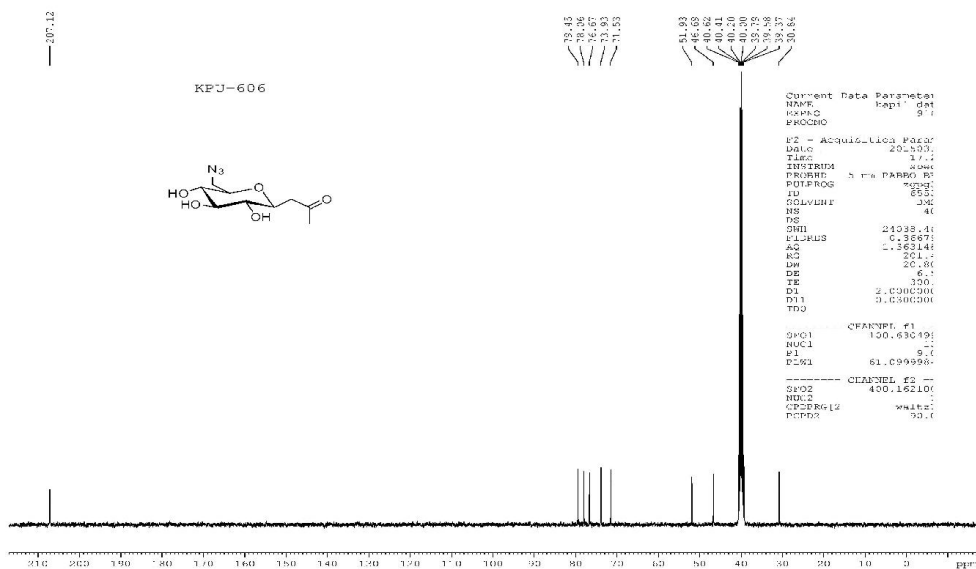
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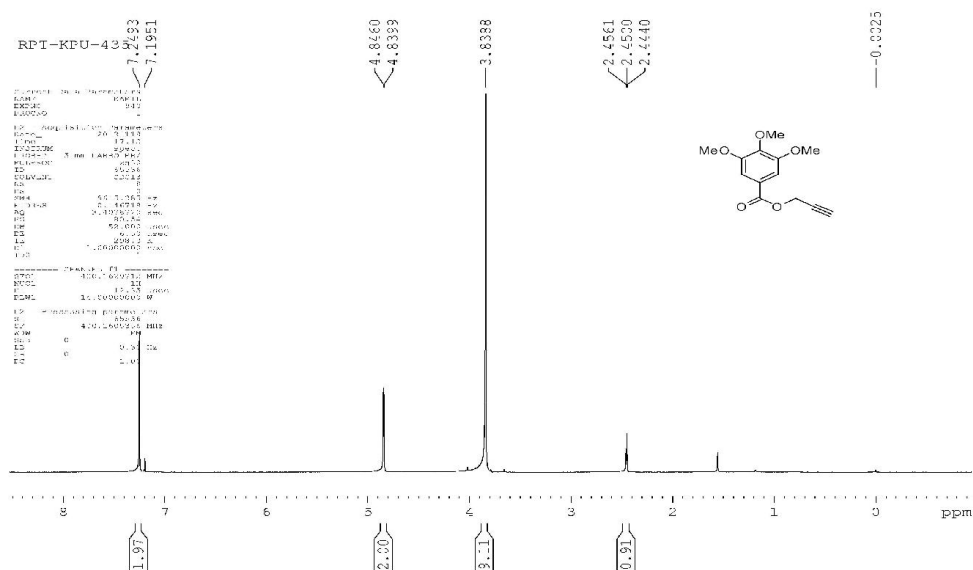
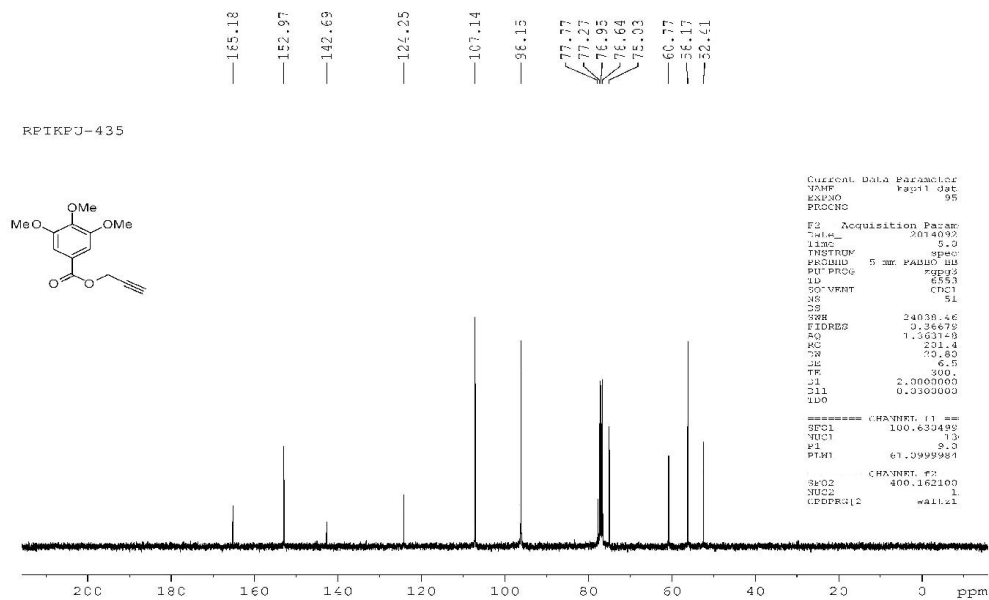
Corresponding author: Tel.: +91 0522 2612411; Fax: +91 522 2623405/ 2623938/ 2629504.

E-mail address: rpt.cdri@gmail.com (Dr.Rama P. Tripathi)

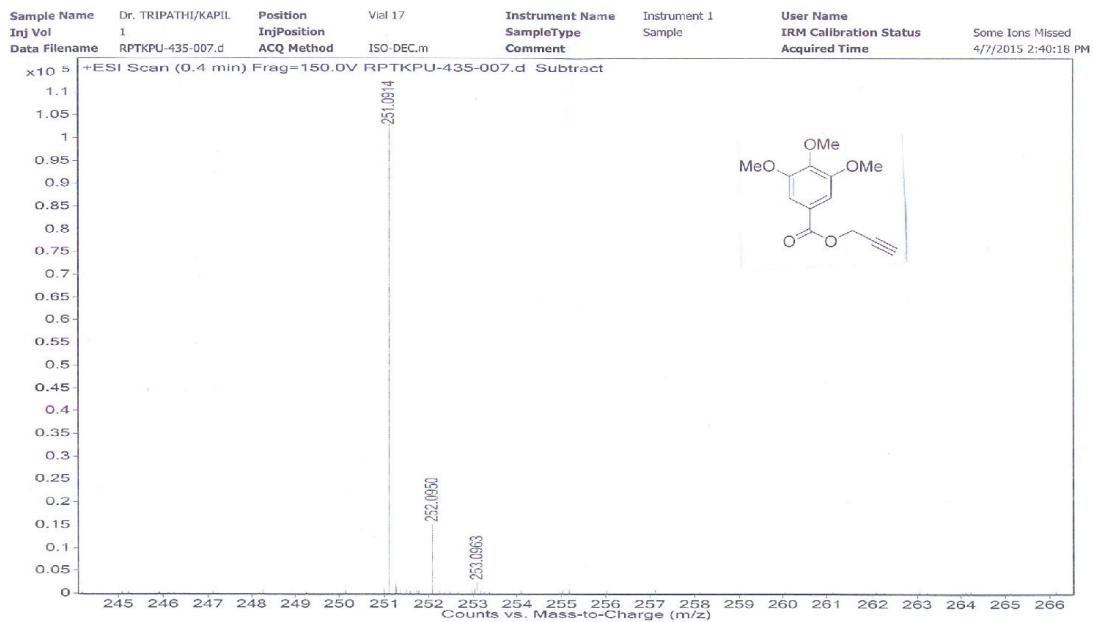
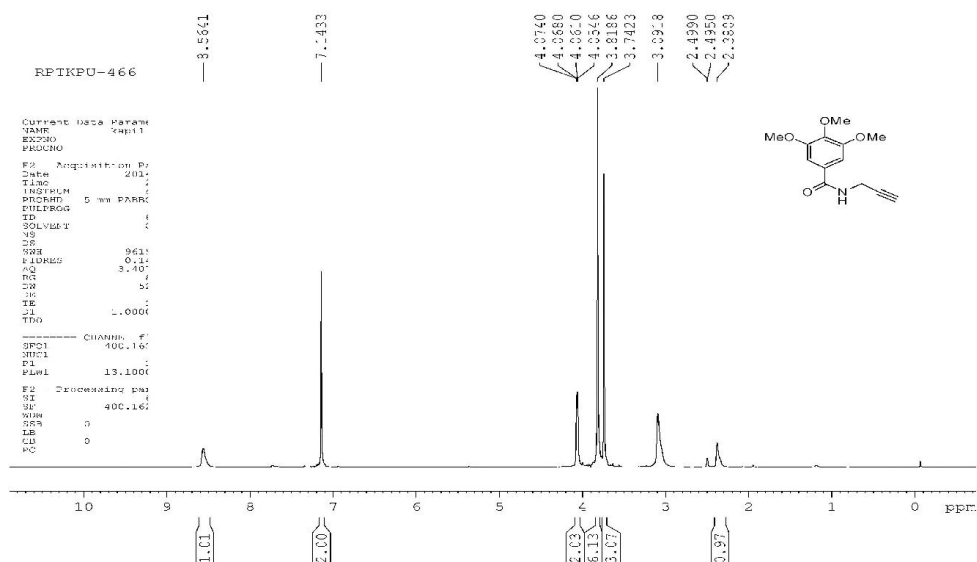
Supporting Information

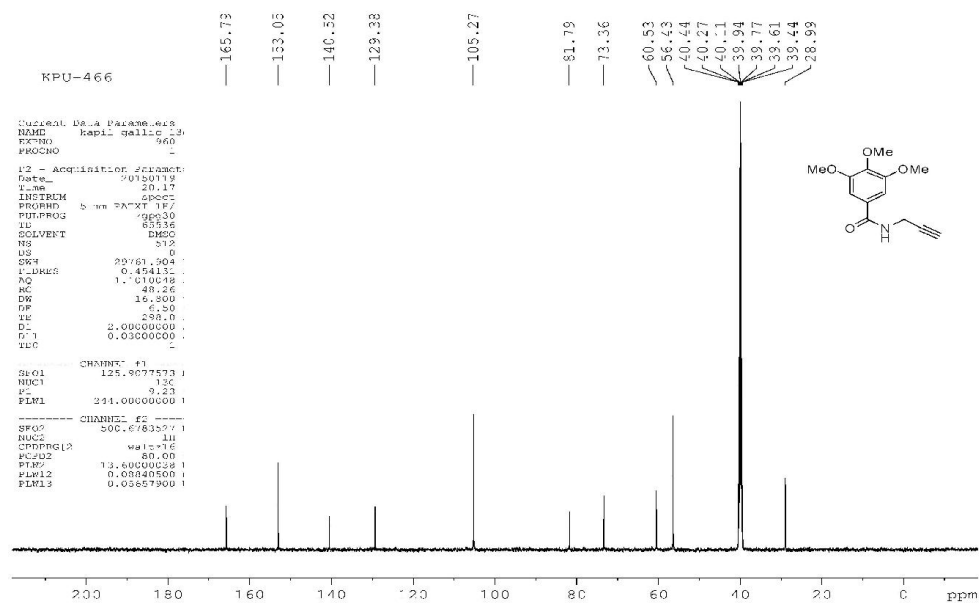
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Copies of HRMS, ^1H NMR and ^{13}C NMR Spectra of Compounds: ^1H NMR spectrum of compound b' ^{13}C NMR spectrum of compound b'

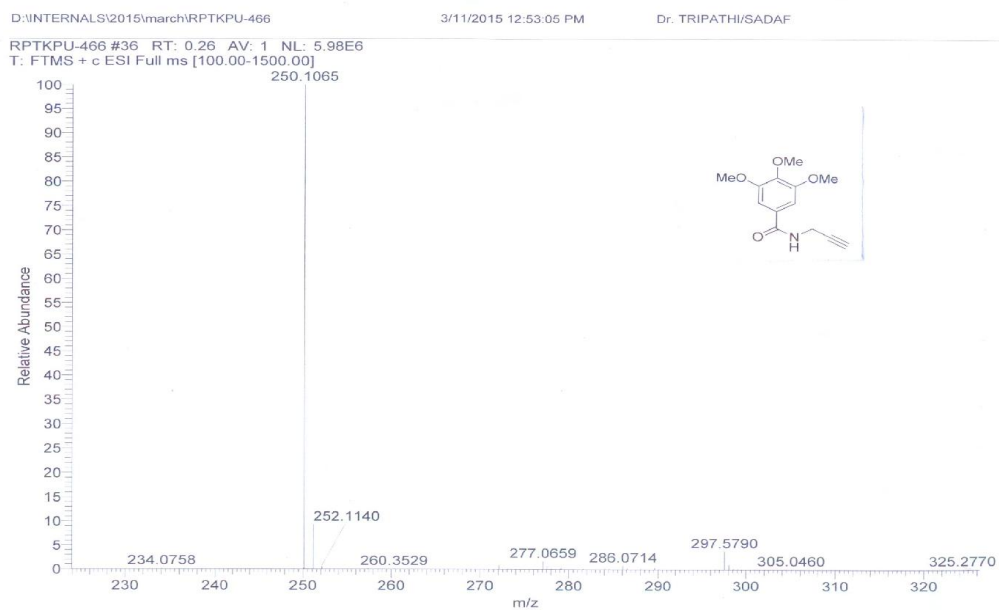
¹H NMR spectrum of compound 2¹³C NMR spectrum of compound 2

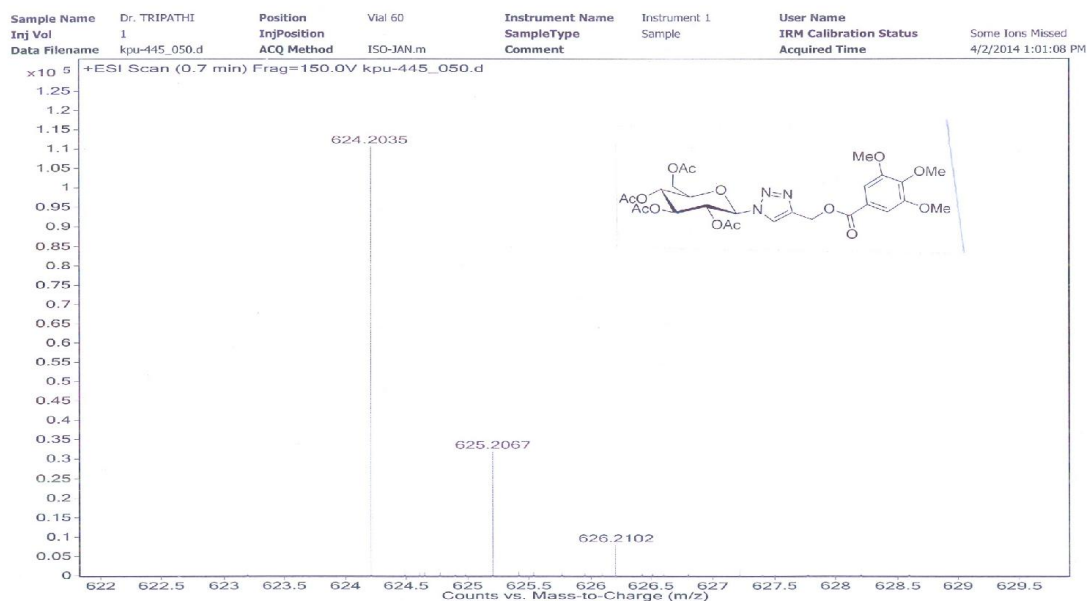
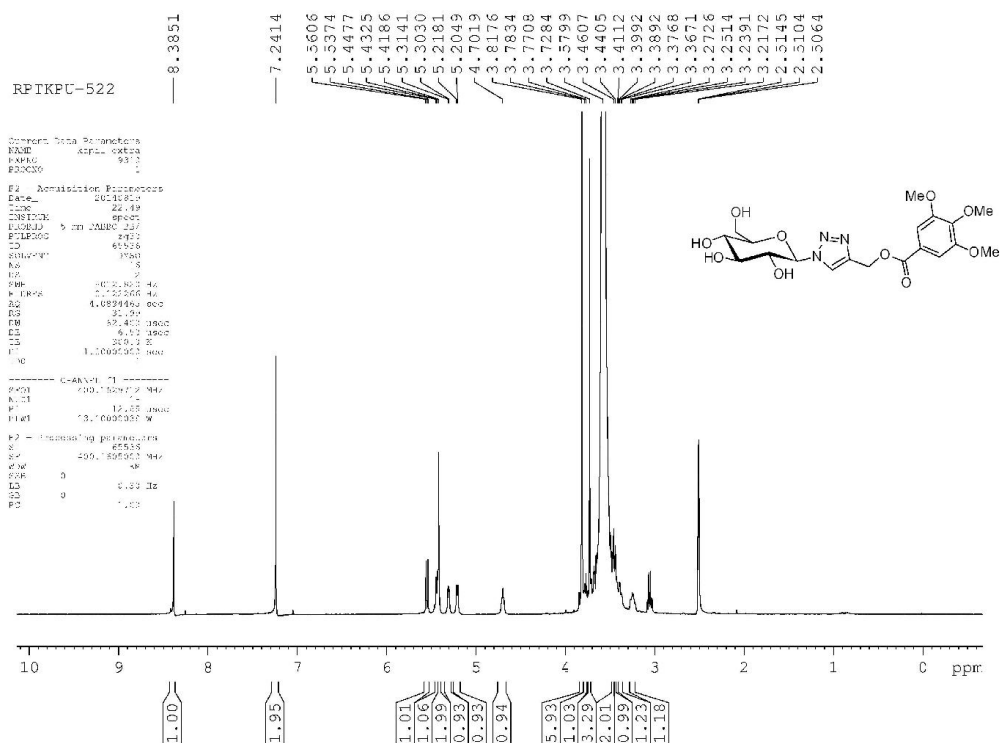
HRMS spectrum of compound 2

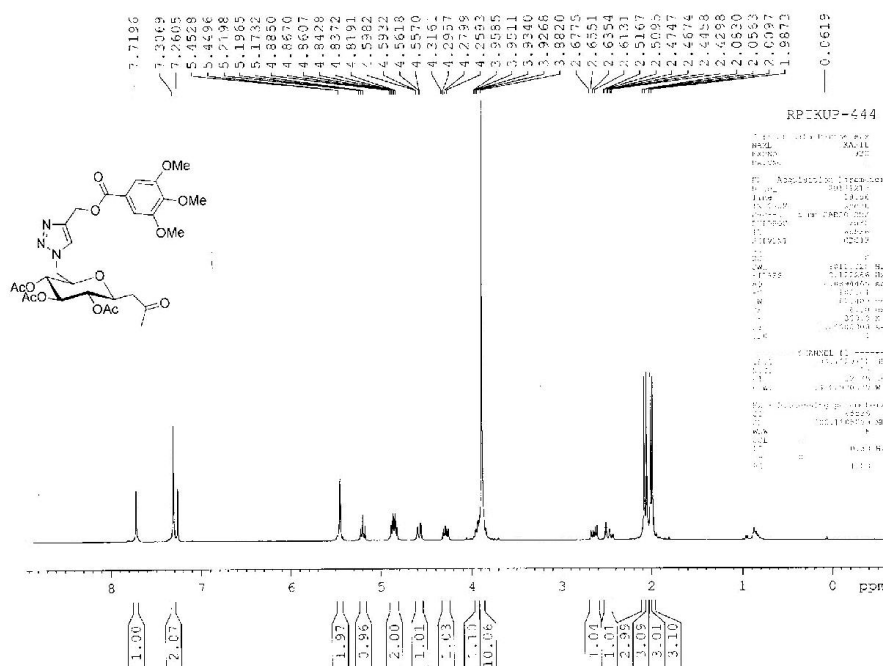
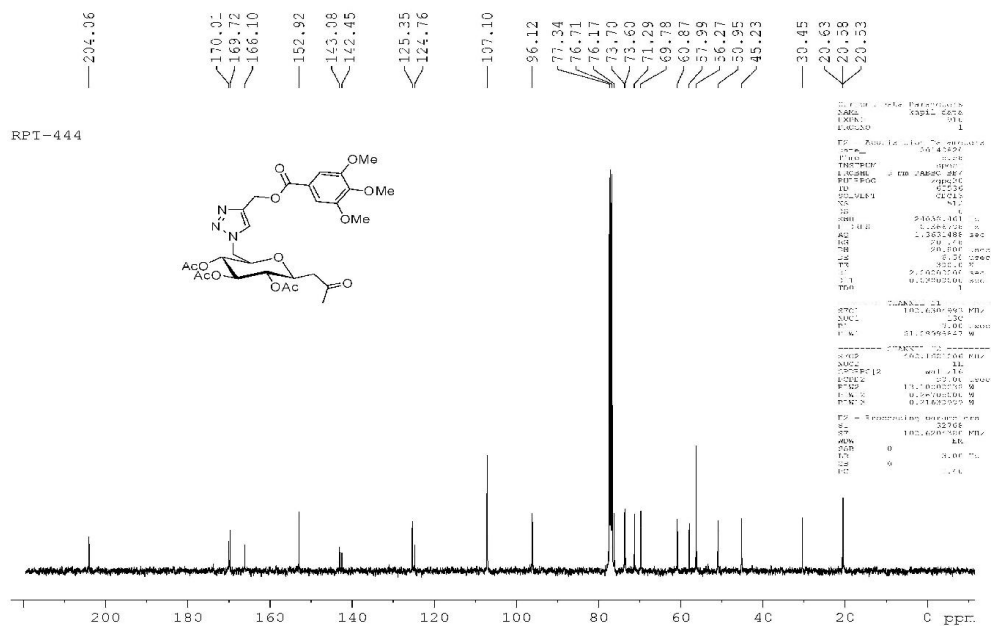
¹H NMR spectrum of compound 4

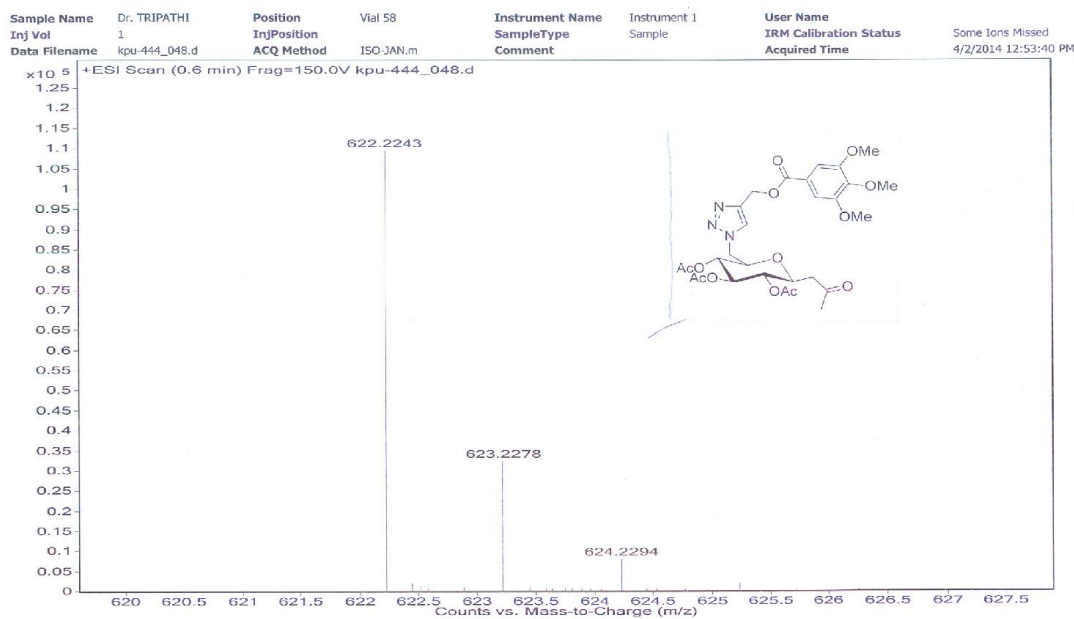
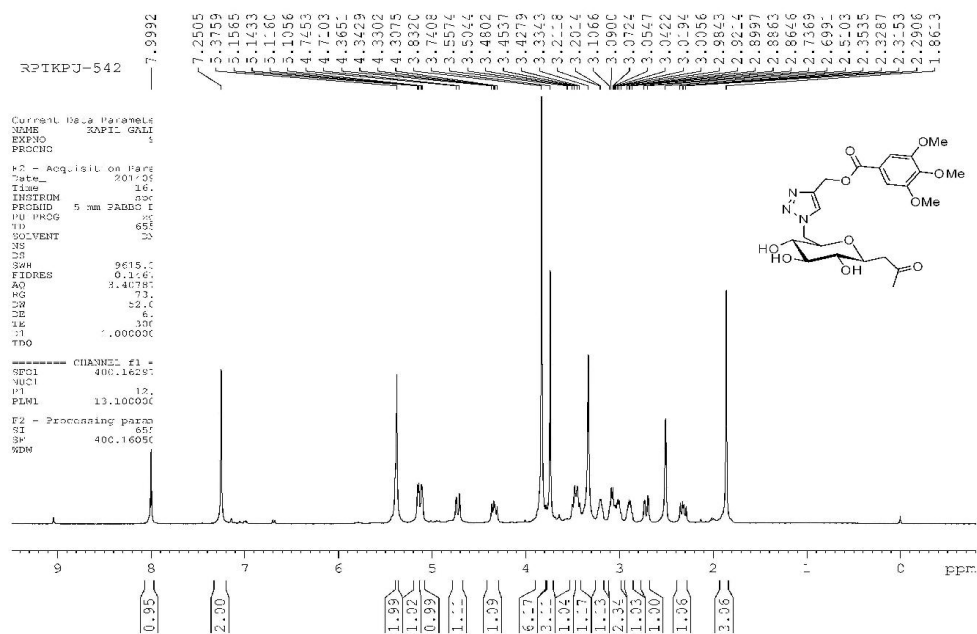
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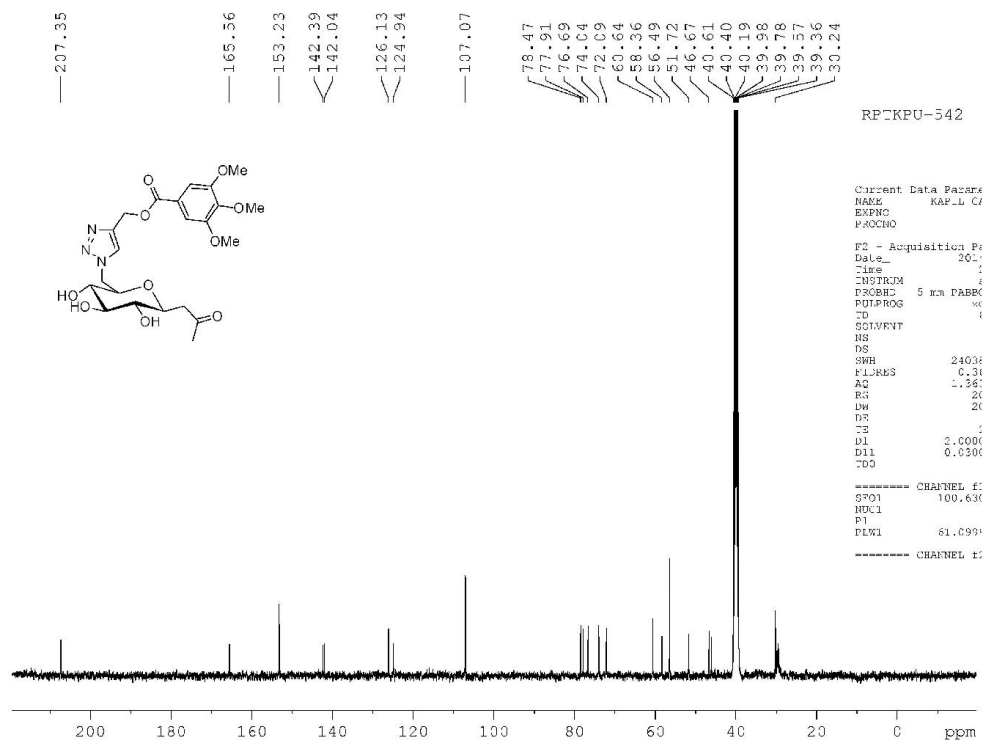
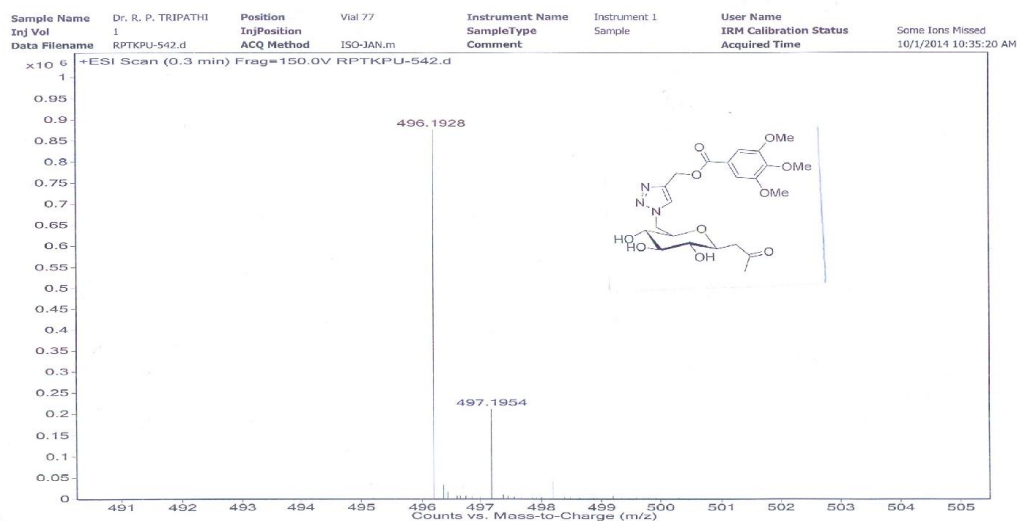
HRMS spectrum of compound 4

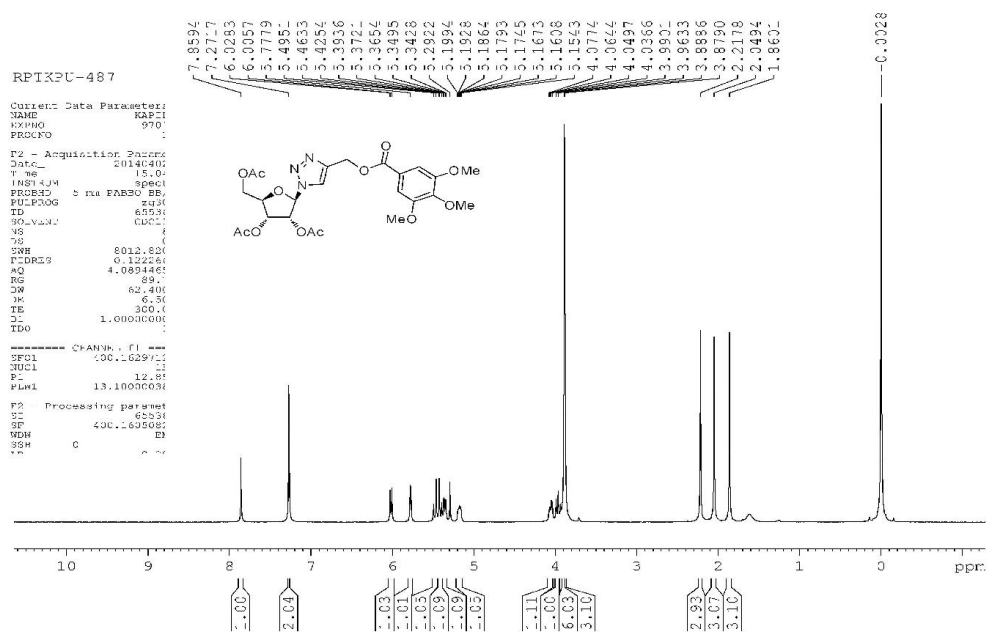
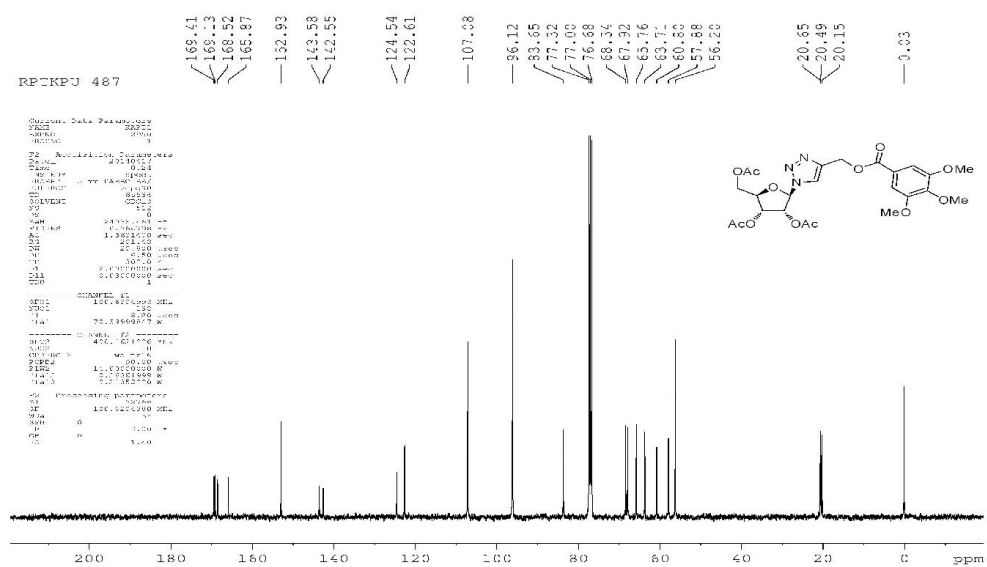


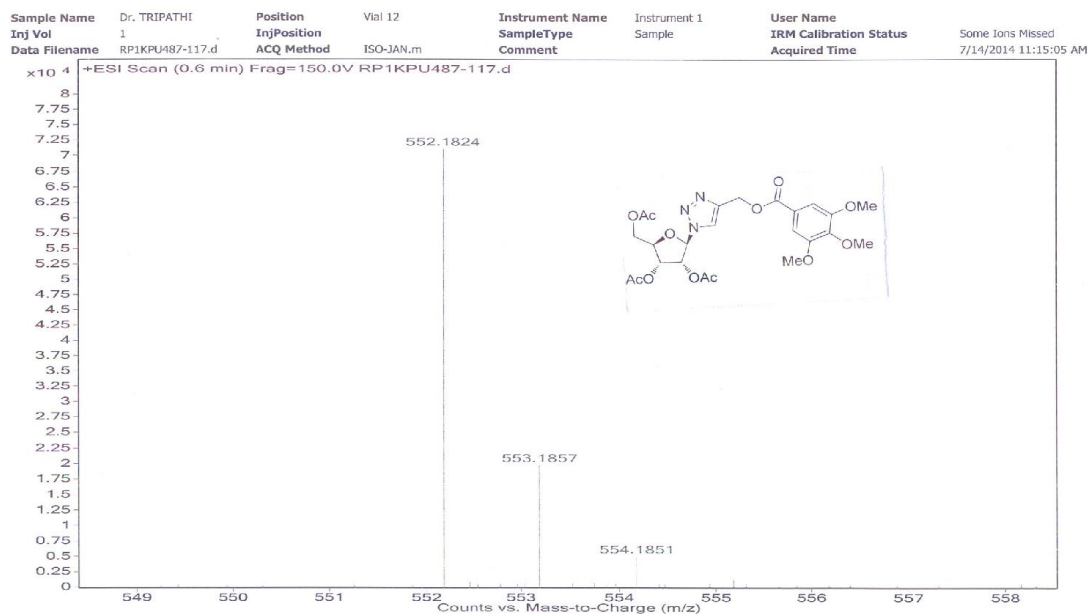
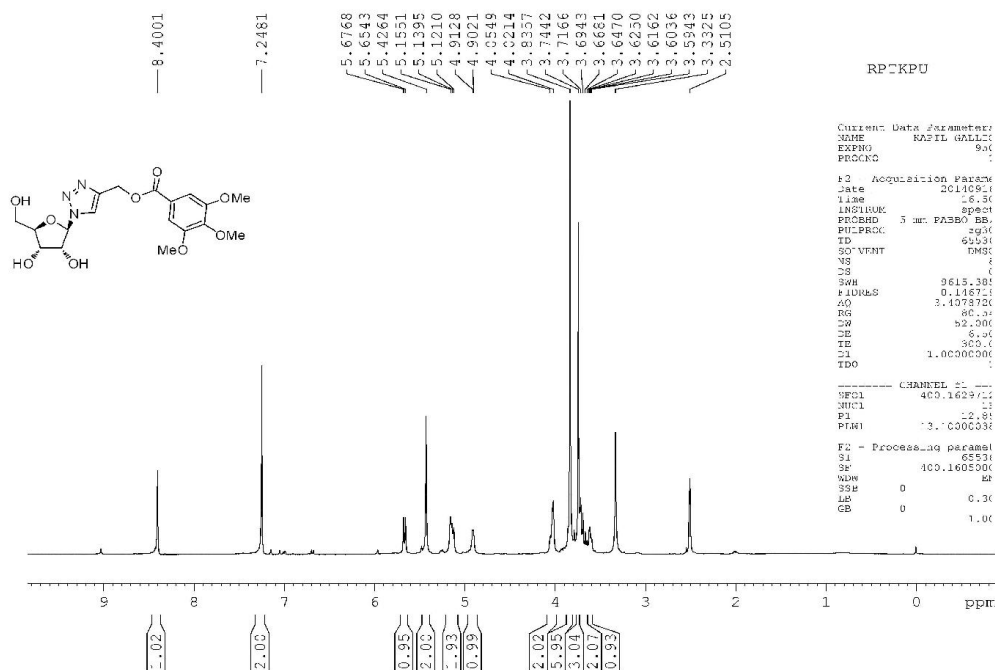
HRMS spectrum of compound **2a** ^1H NMR spectrum of compound **2a'**

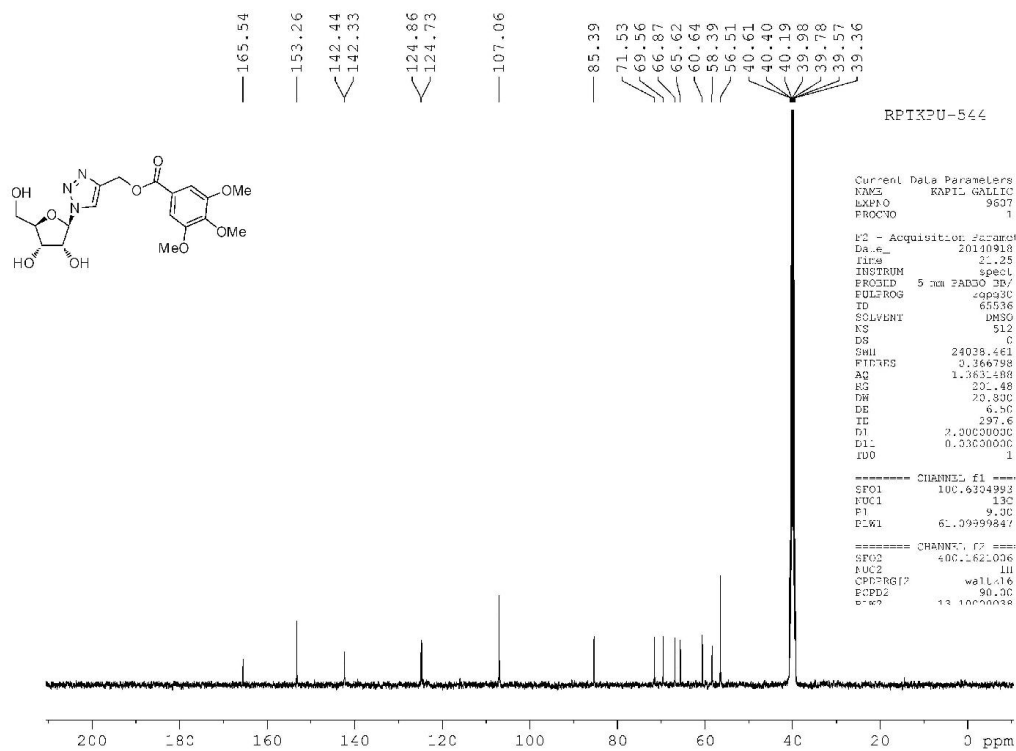
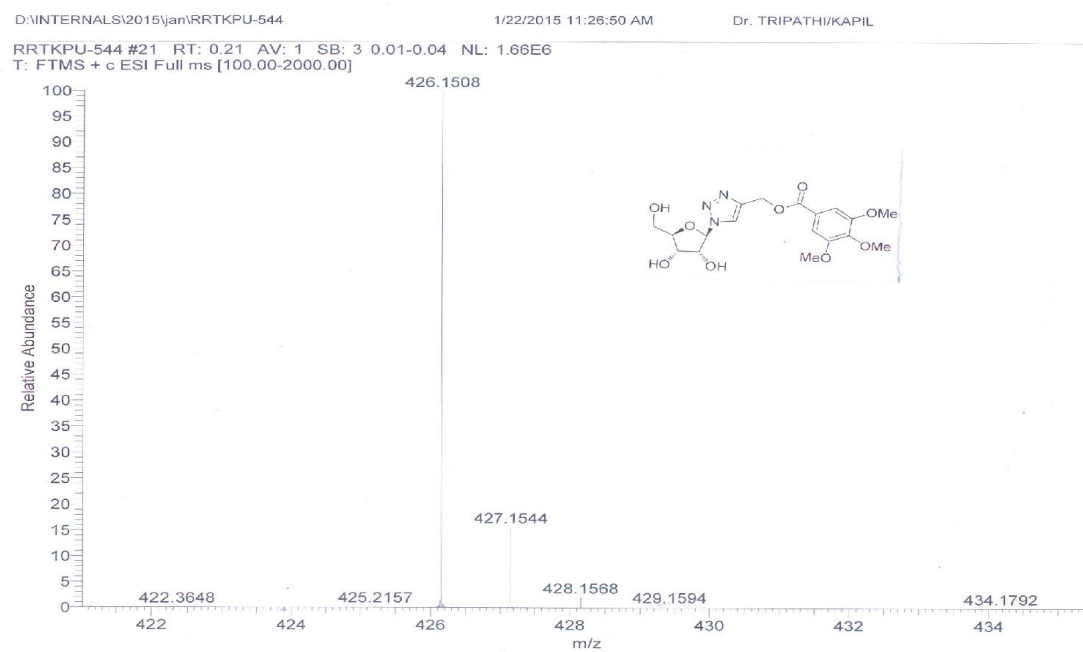
^1H NMR spectrum of compound **2b** ^{13}C NMR spectrum of compound **2b**

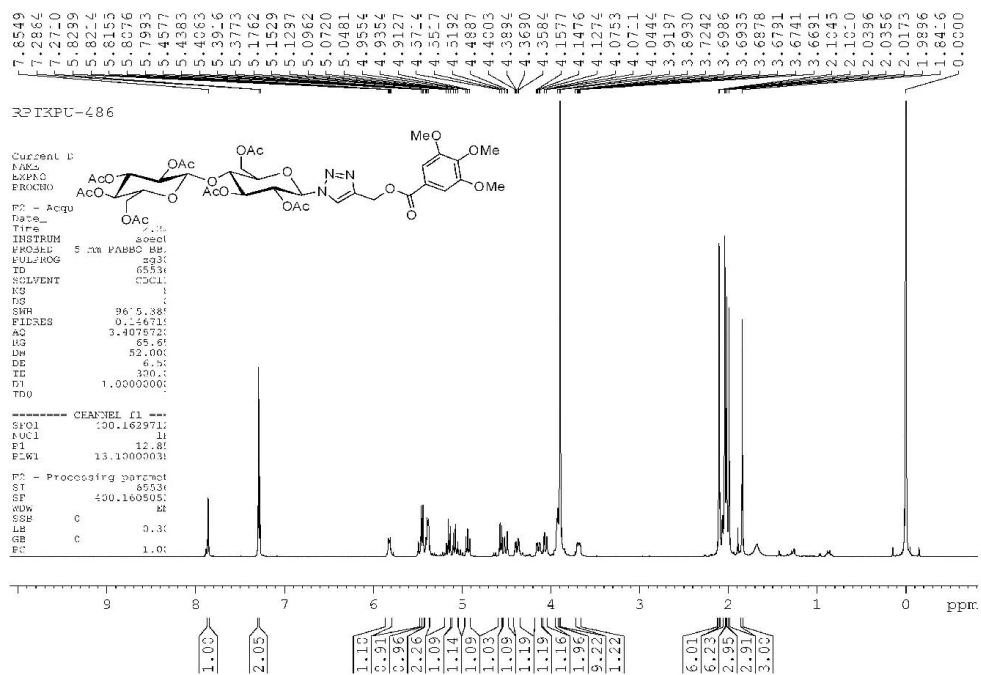
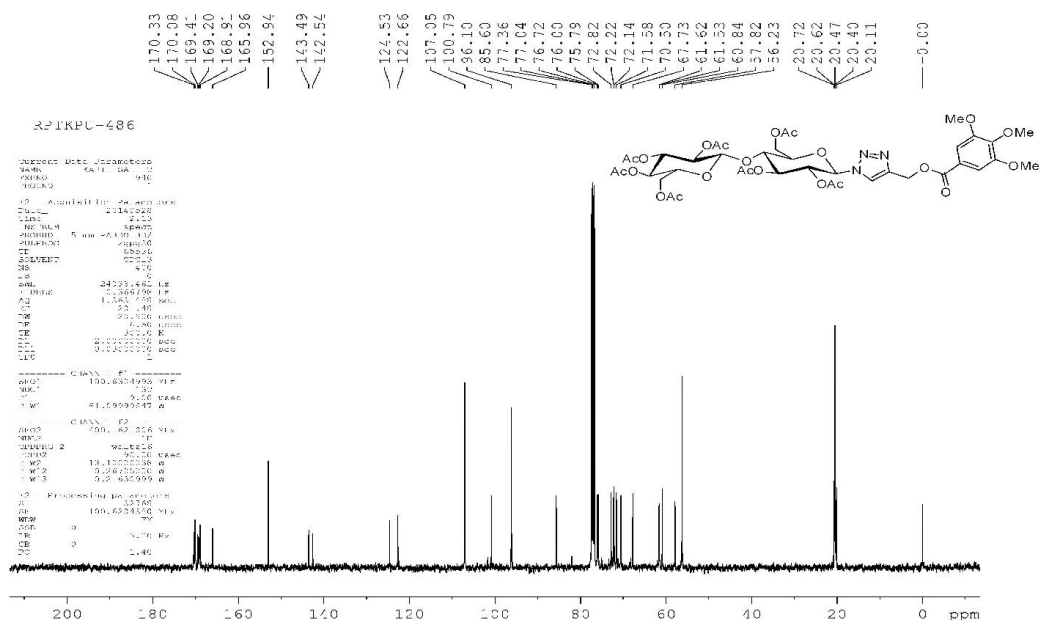
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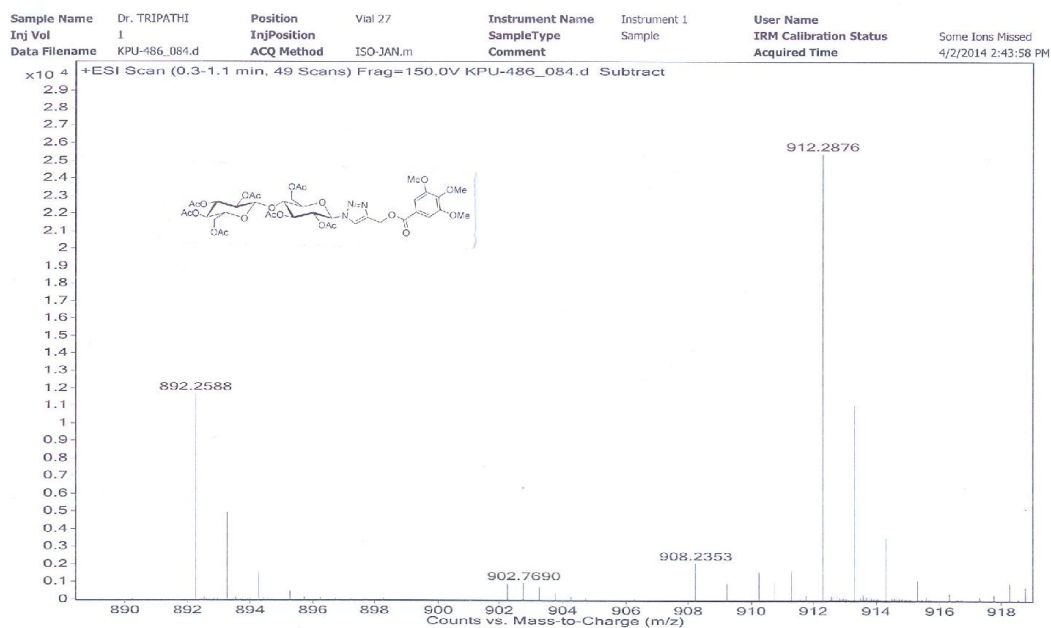
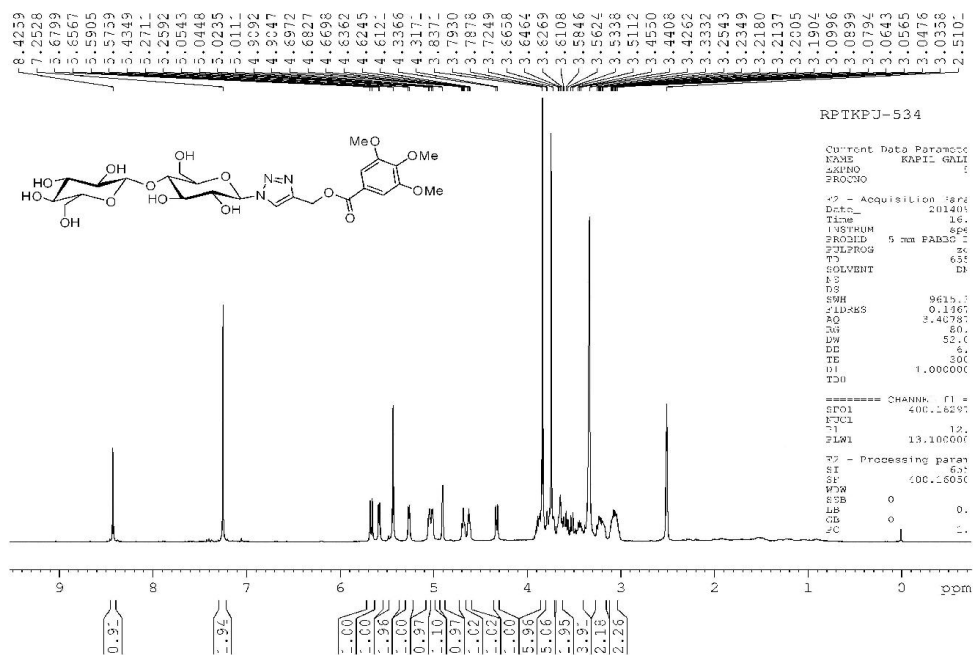
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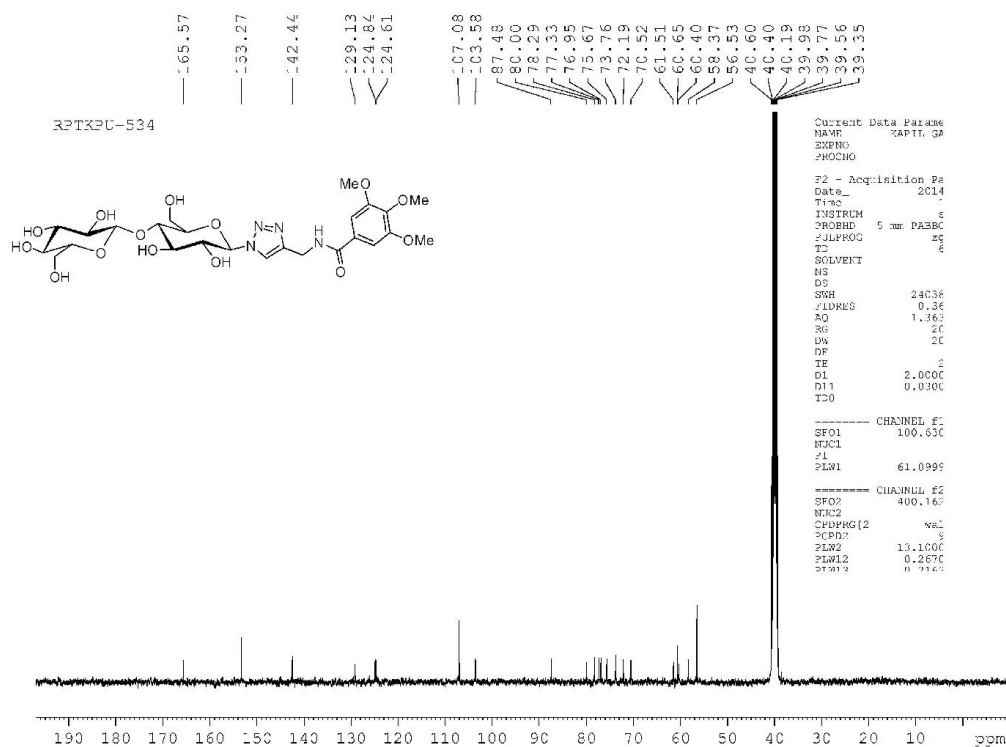
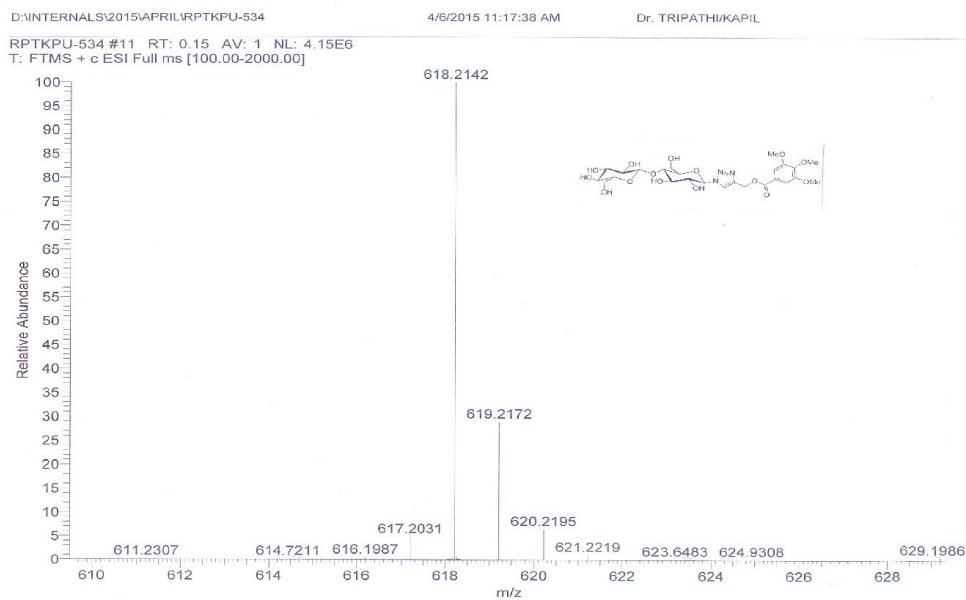
¹H NMR spectrum of compound 2c¹³C NMR spectrum of compound 2c

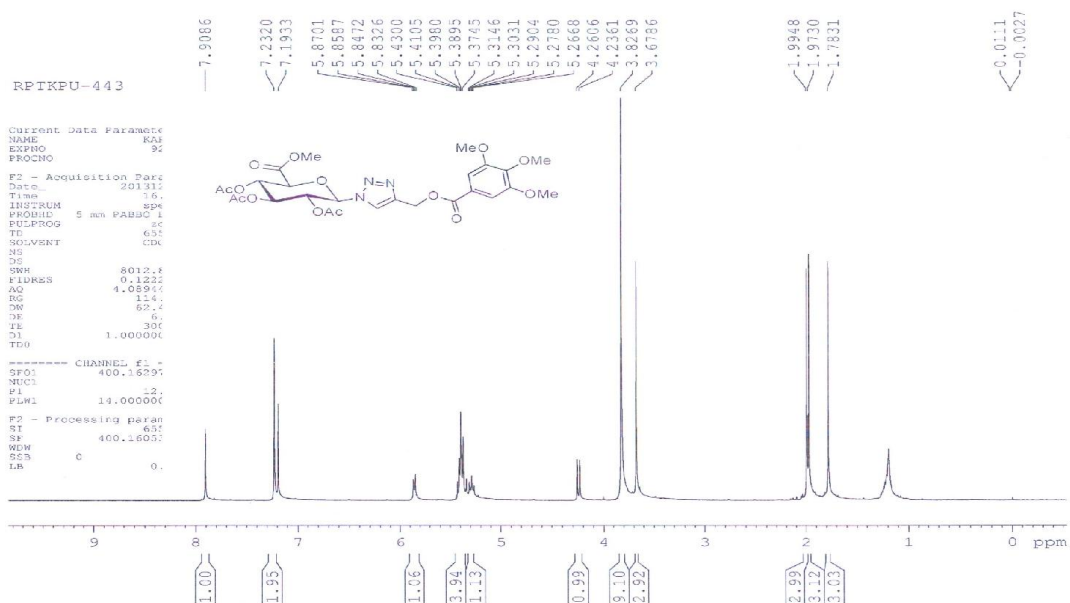
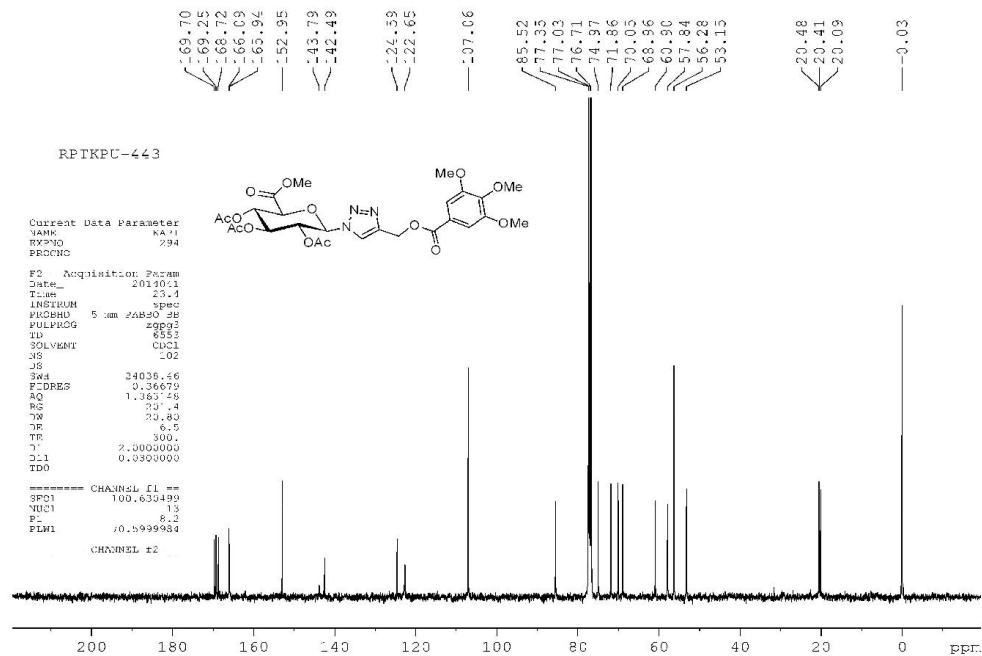
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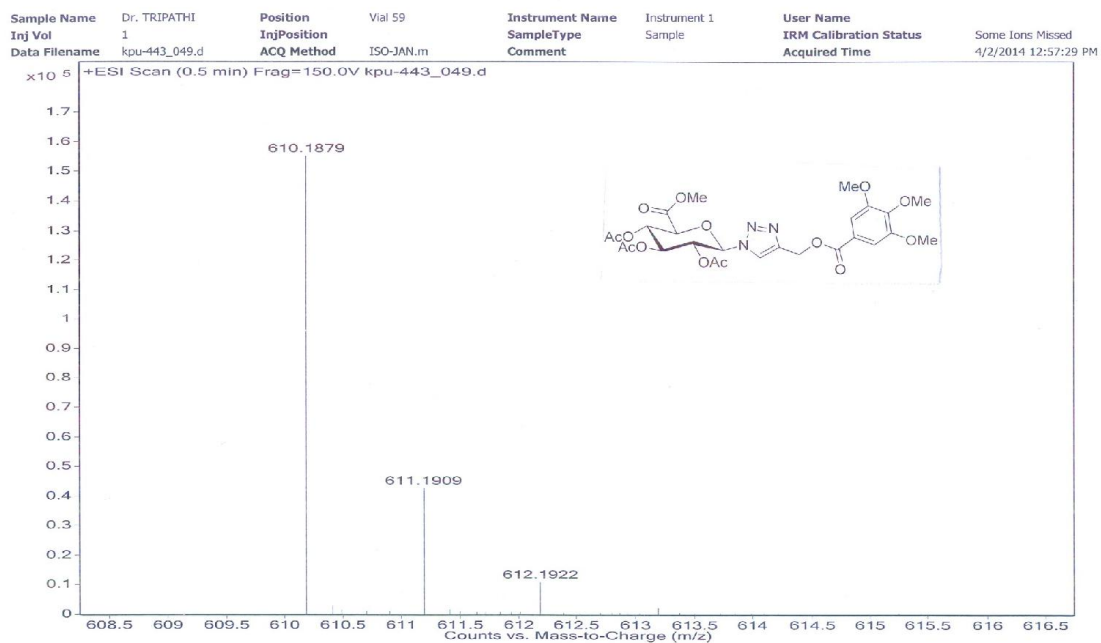
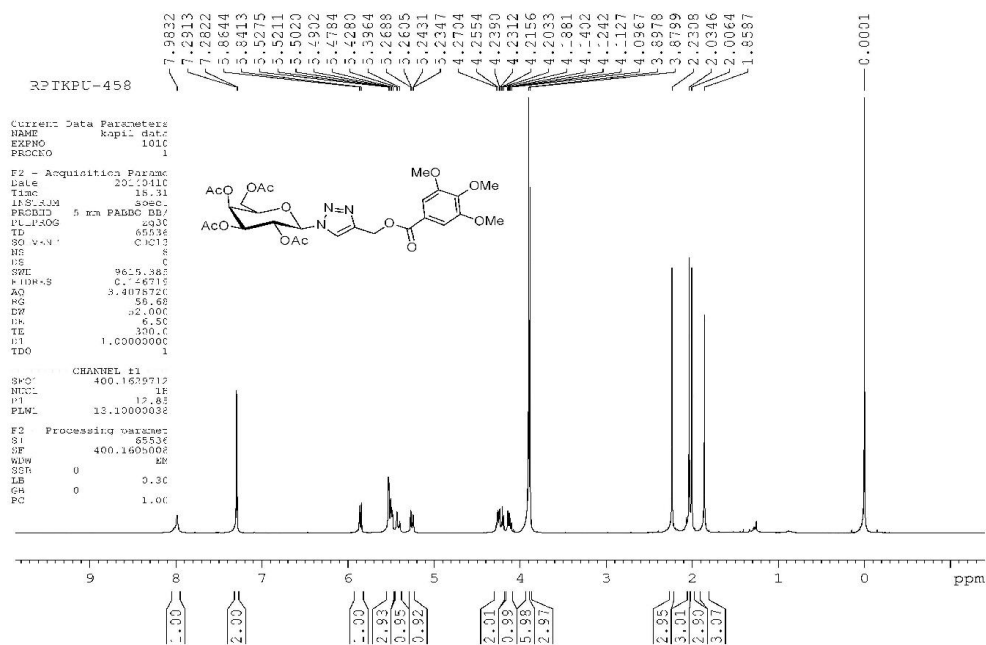
¹³C NMR spectrum of compound **2c'**HRMS spectrum of compound **2c'**

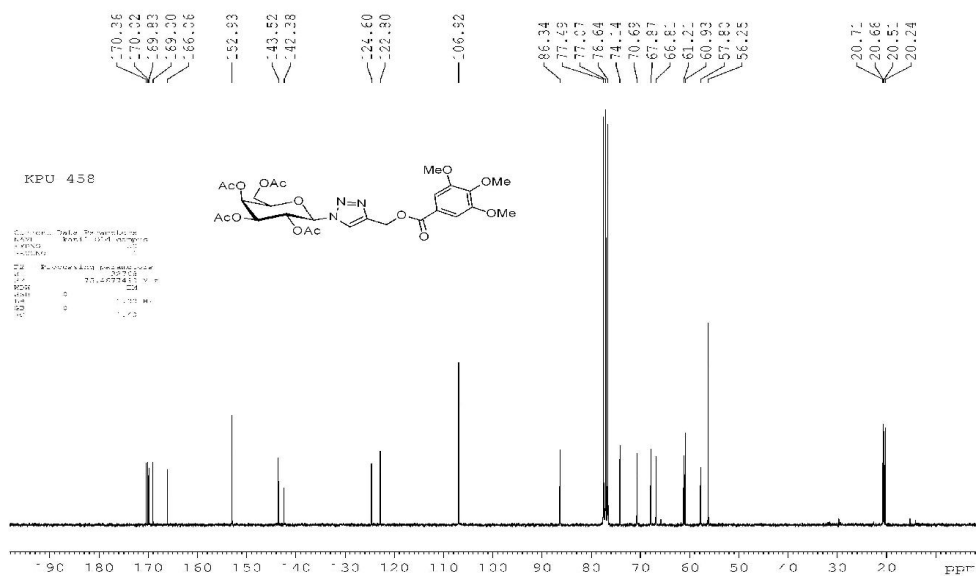
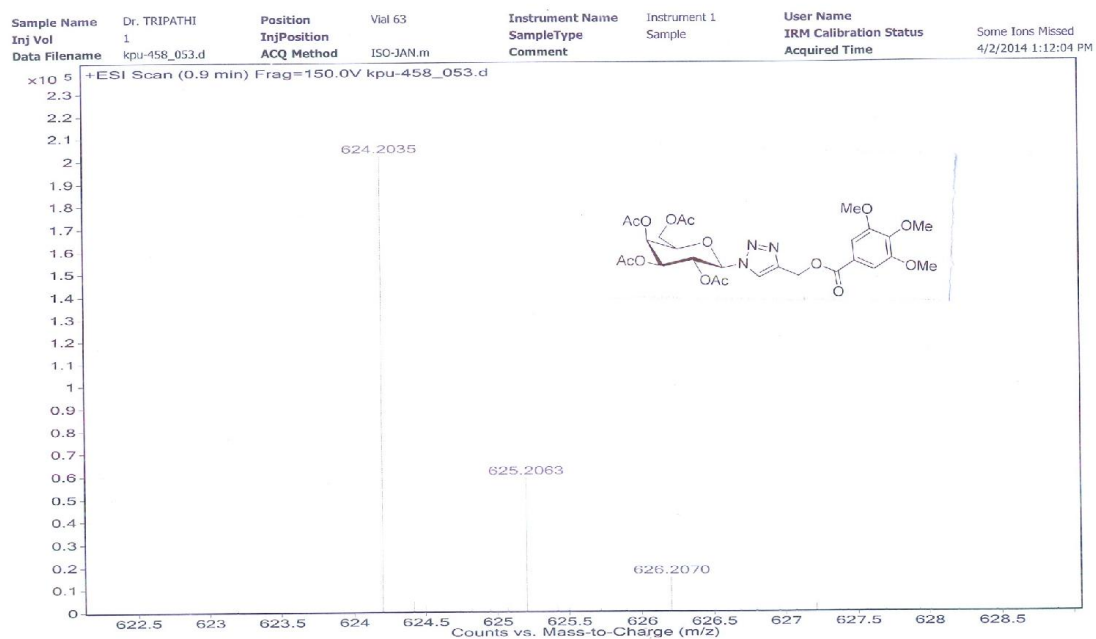
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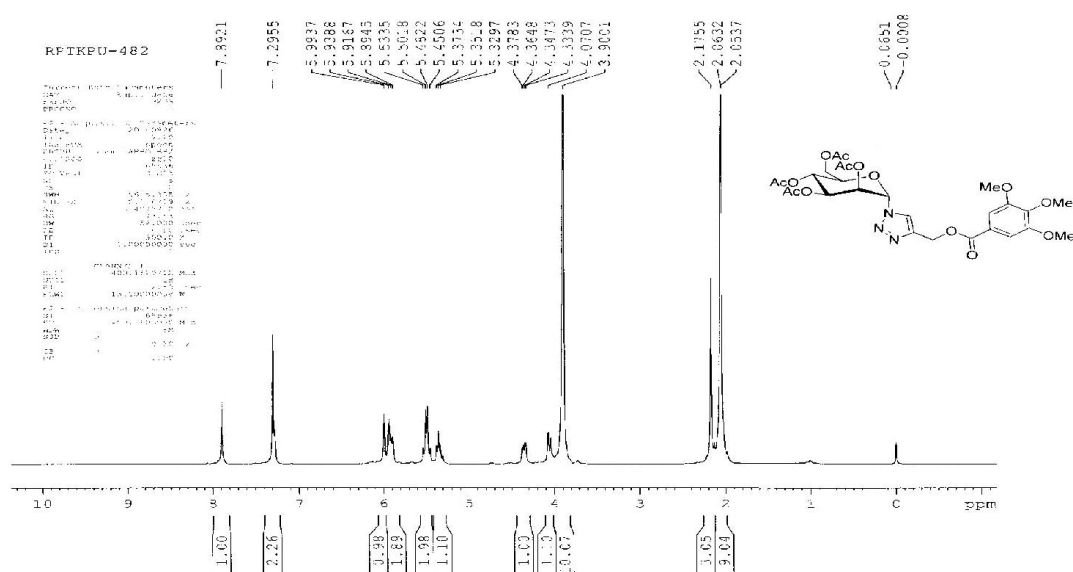
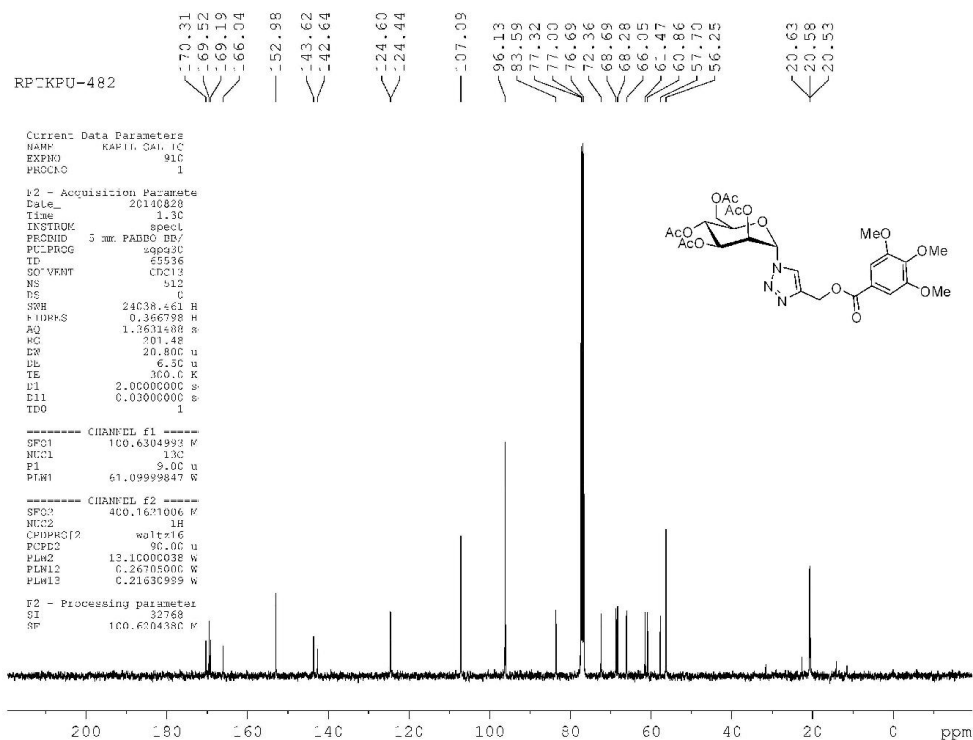
HRMS spectrum of compound **2d** ^1H NMR spectrum of compound **2d'**

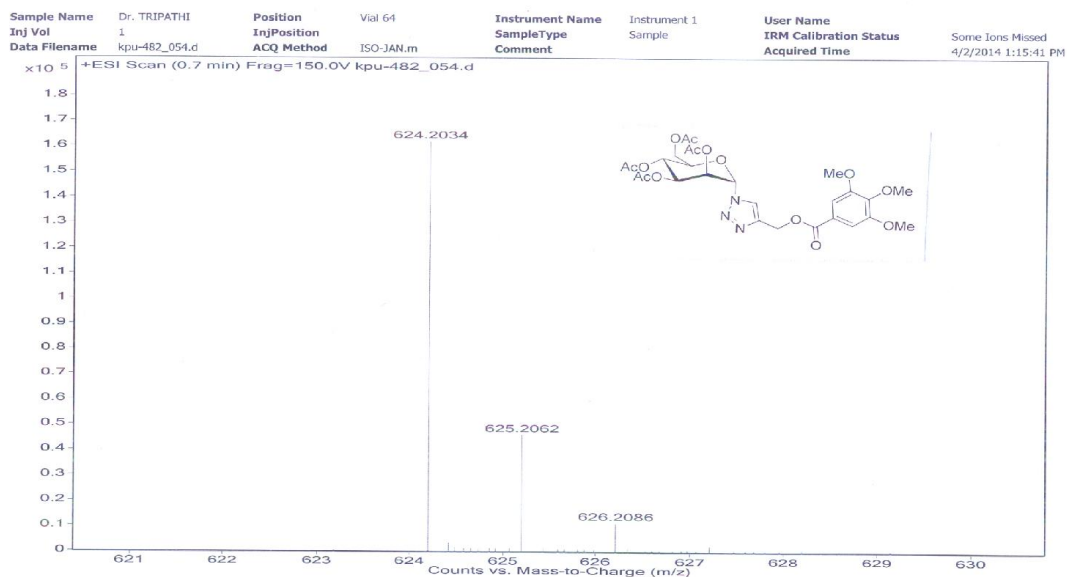
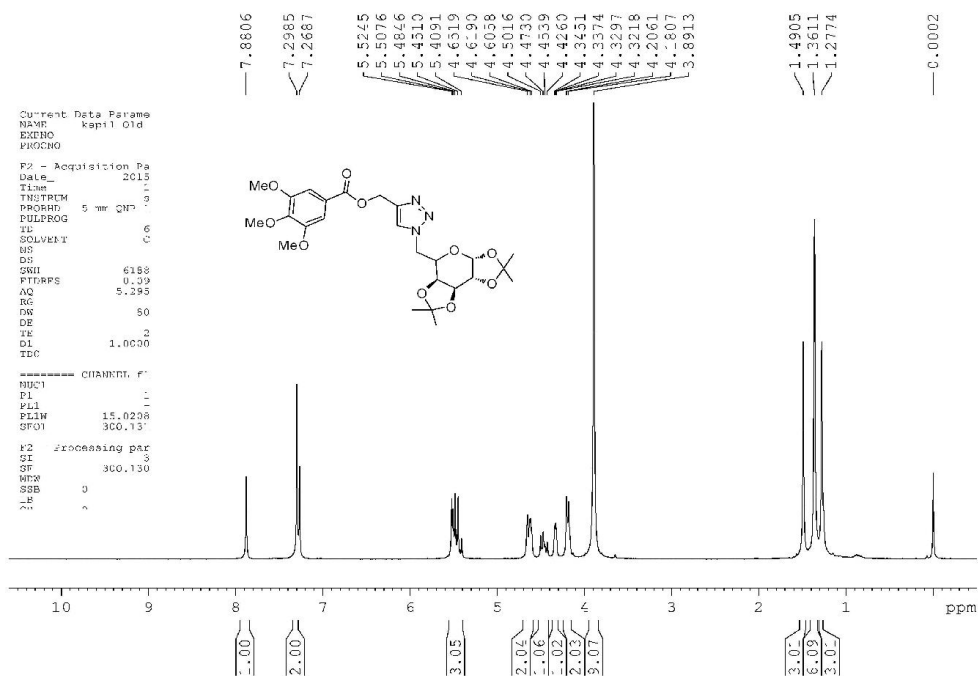
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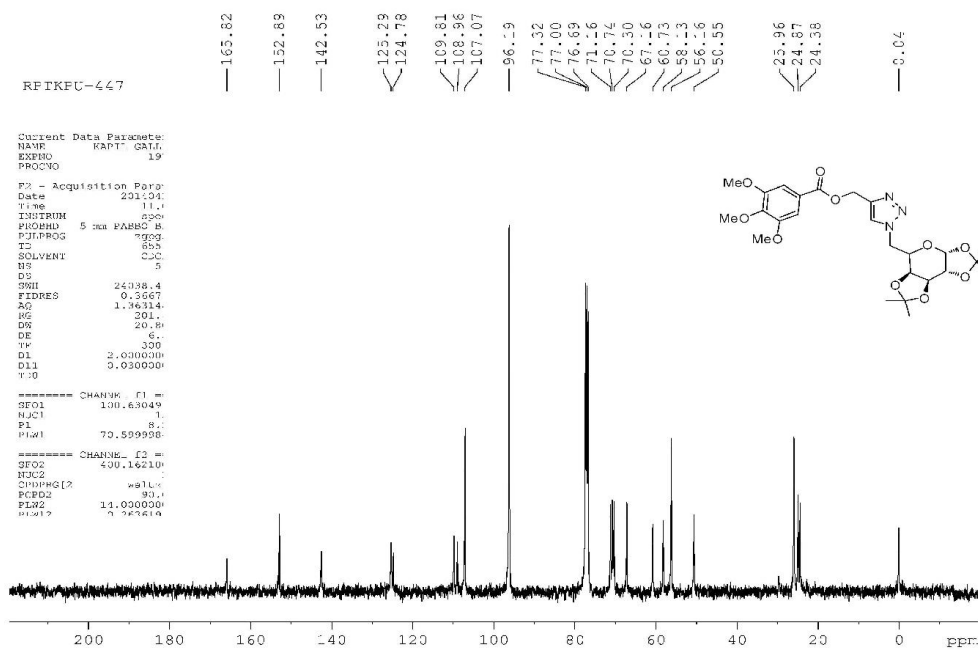
¹H NMR spectrum of compound 2e¹³C NMR spectrum of compound 2e

HRMS spectrum of compound **2e**¹H NMR spectrum of compound **2f**

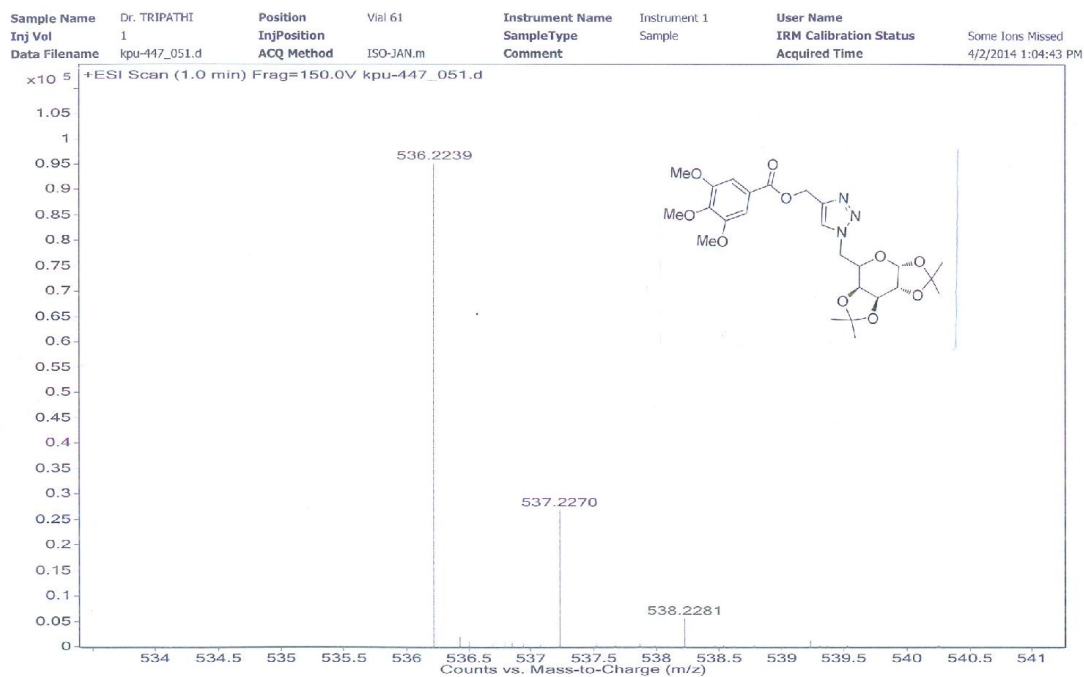
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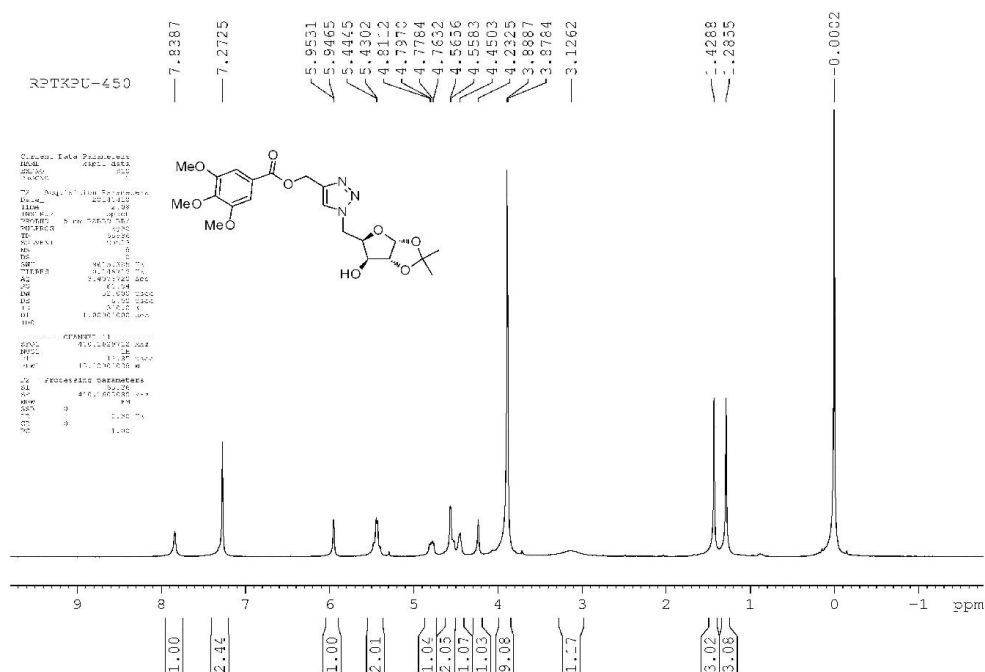
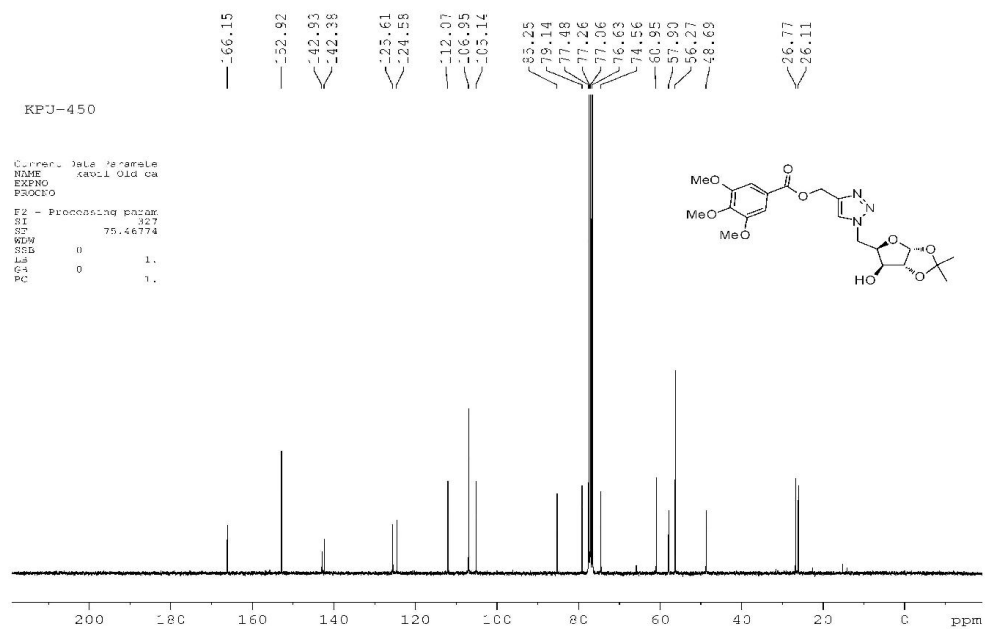
¹H NMR spectrum of compound 2g¹³C NMR spectrum of compound 2g

HRMS spectrum of compound **2g**¹H NMR spectrum of compound **2h**

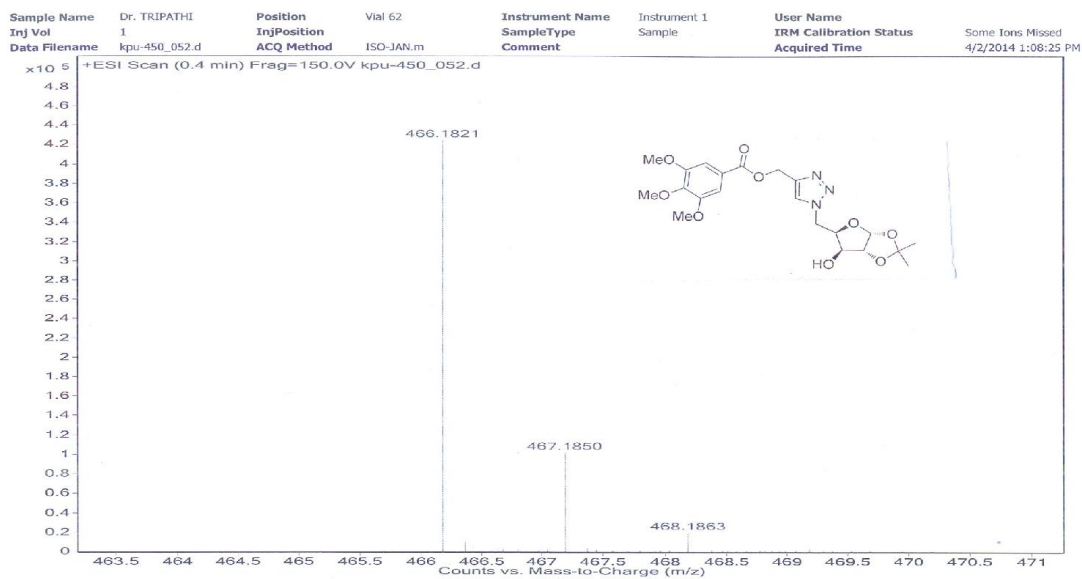
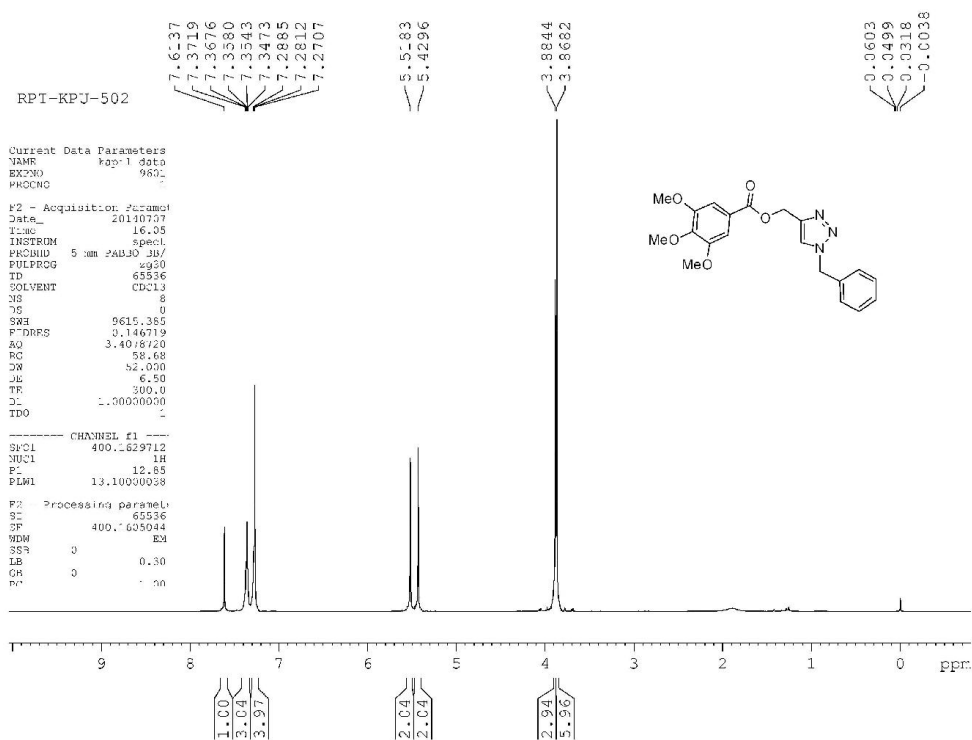
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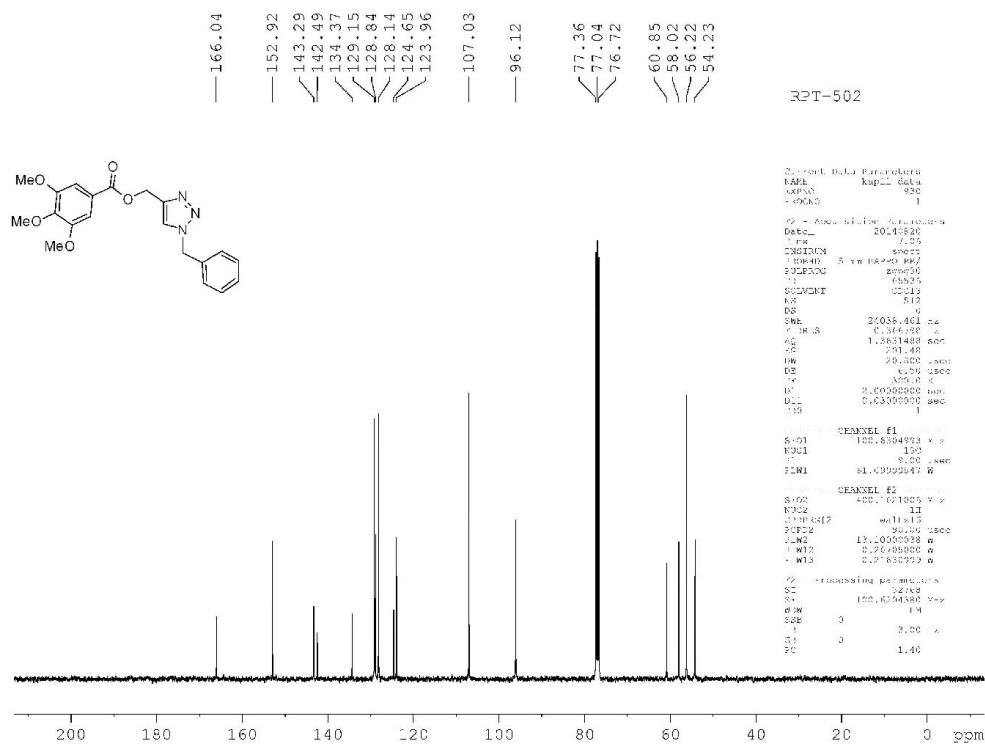
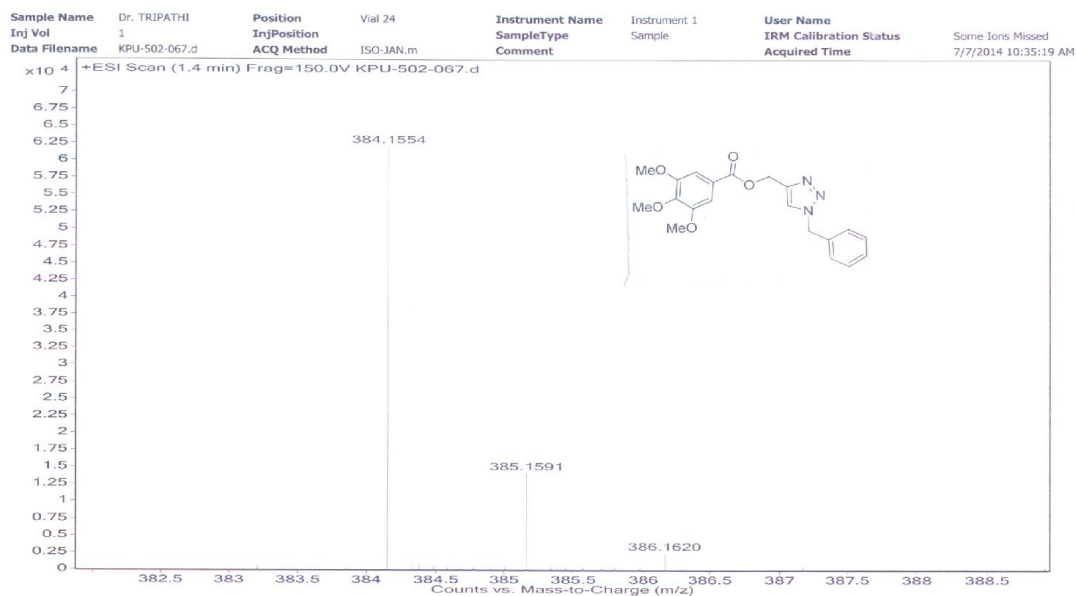
HRMS spectrum of compound 2h

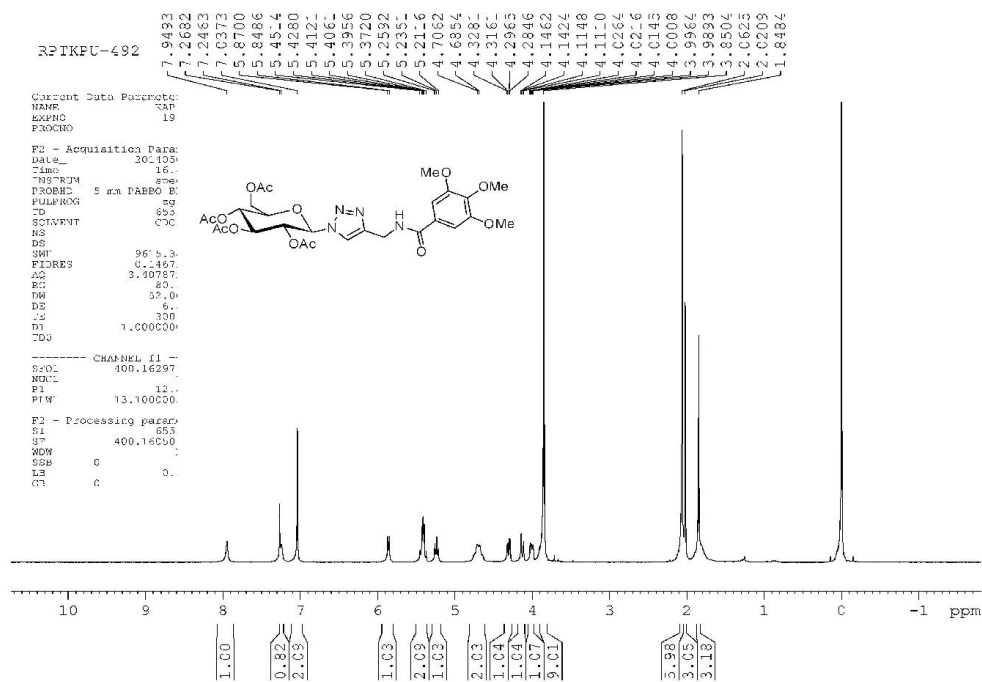
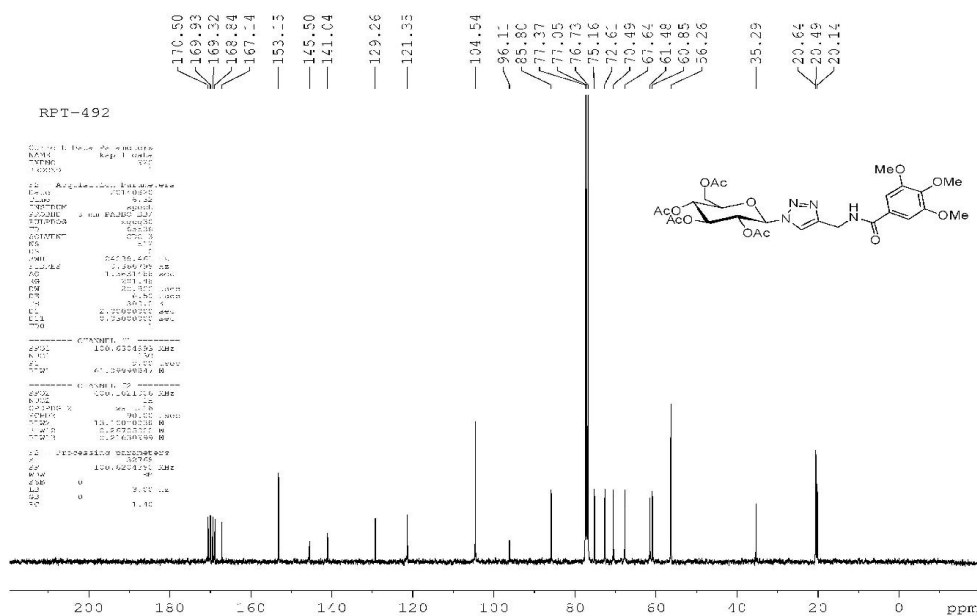


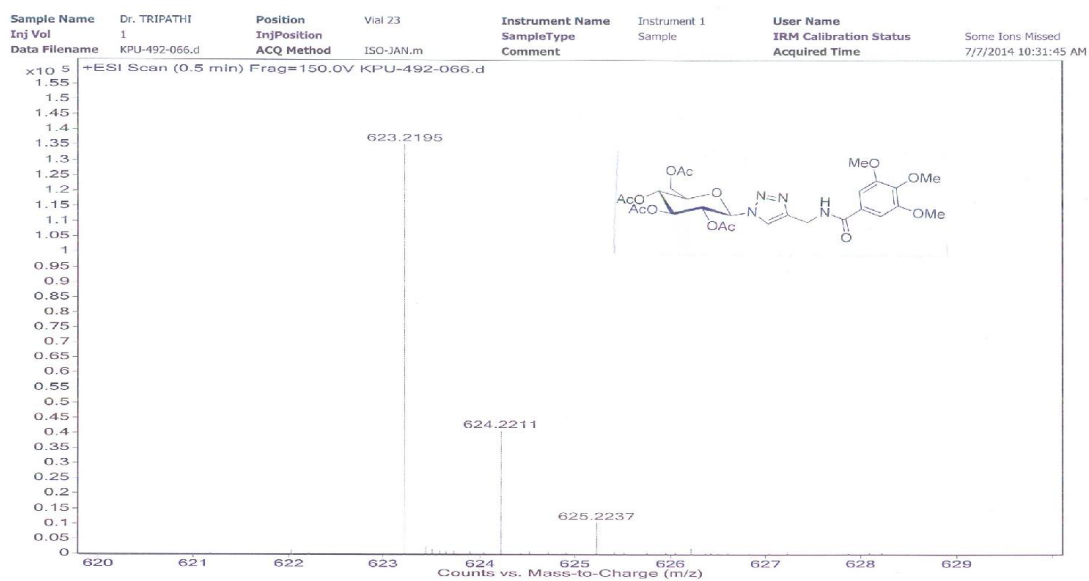
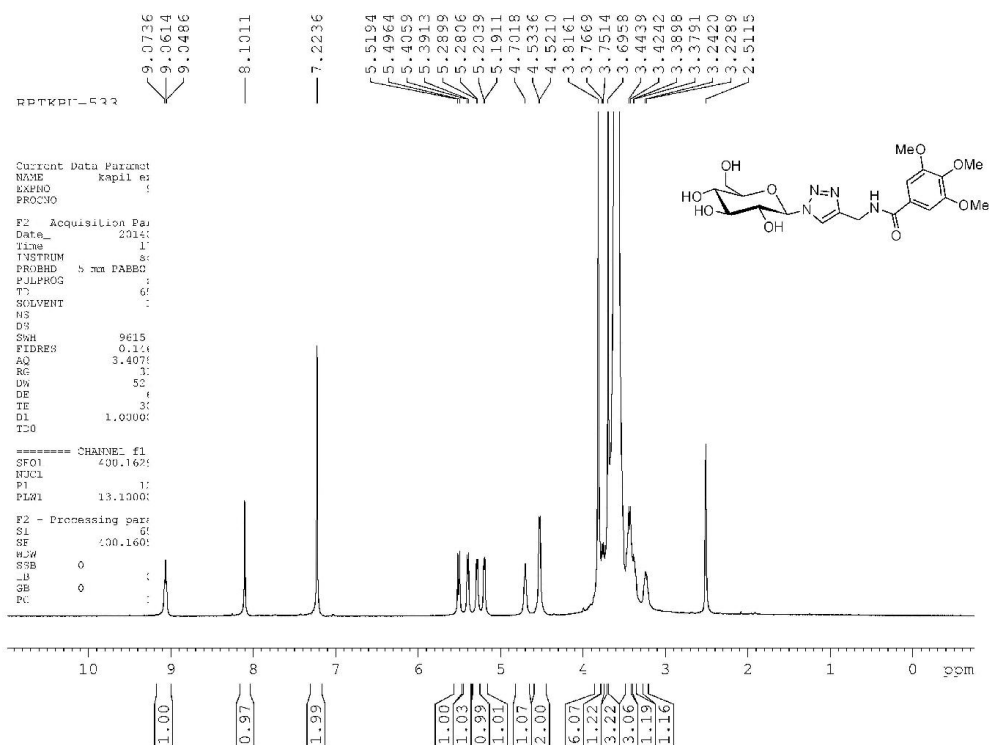
^1H NMR spectrum of compound **2i** ^{13}C NMR spectrum of compound **2i**

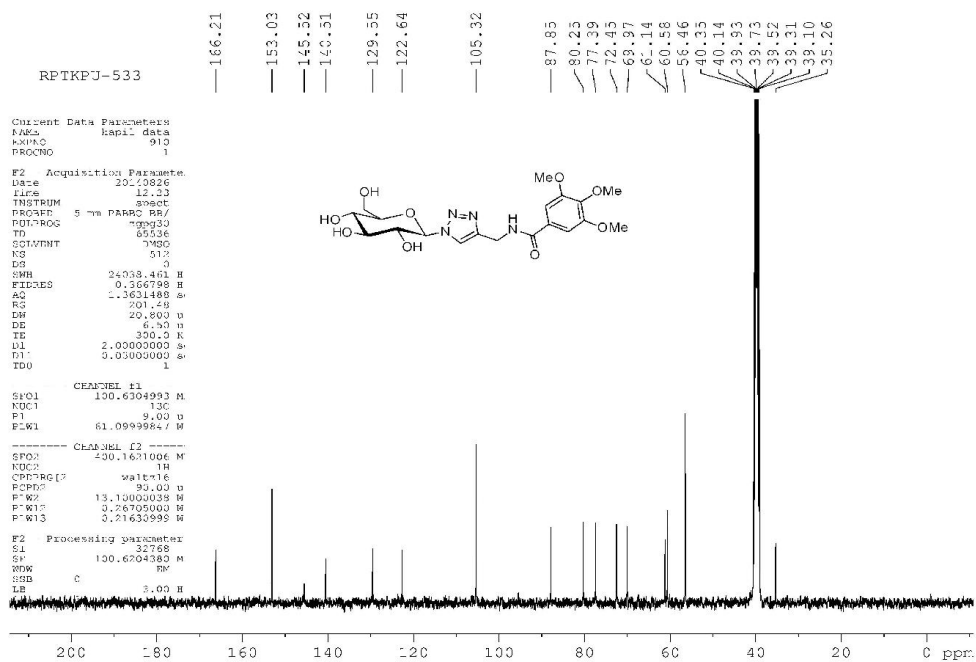
HRMS spectrum of compound 2i

¹H NMR spectrum of compound 2j

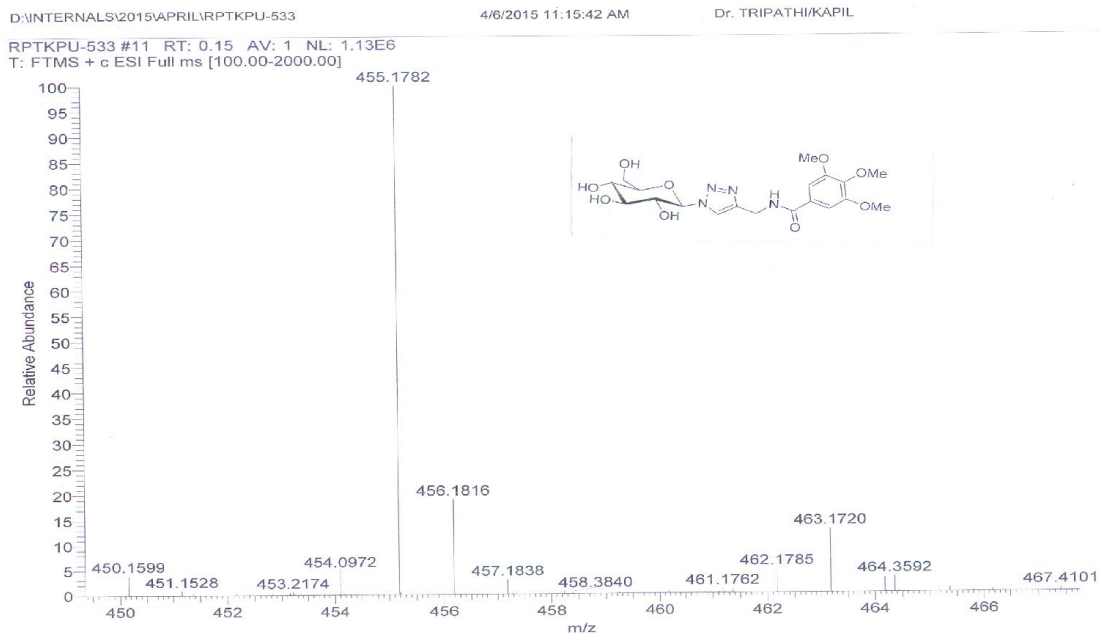
¹³C NMR spectrum of compound **2j**HRMS spectrum of compound **2j**

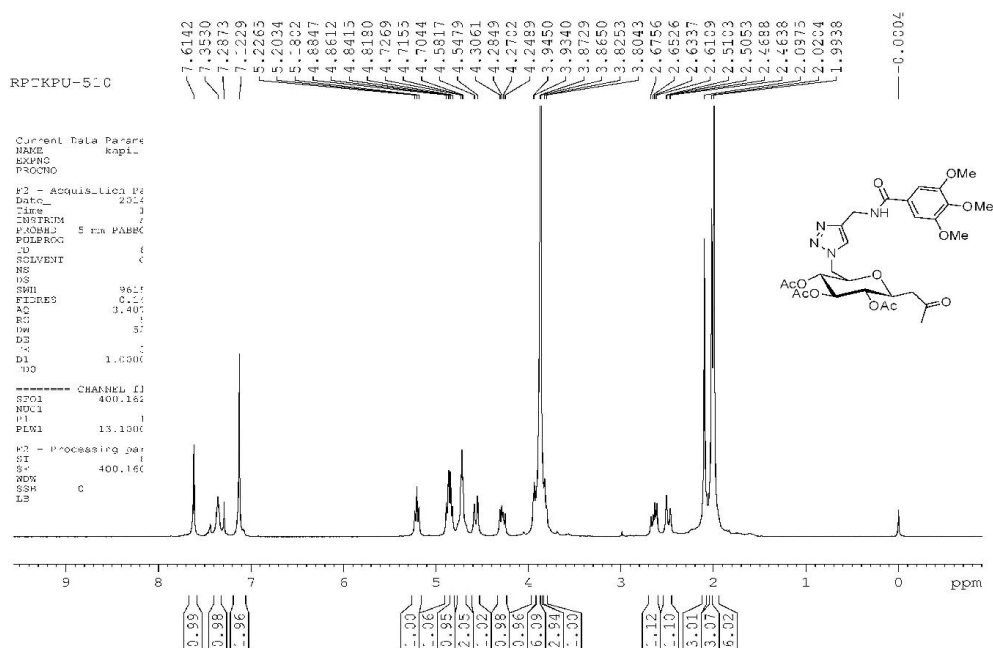
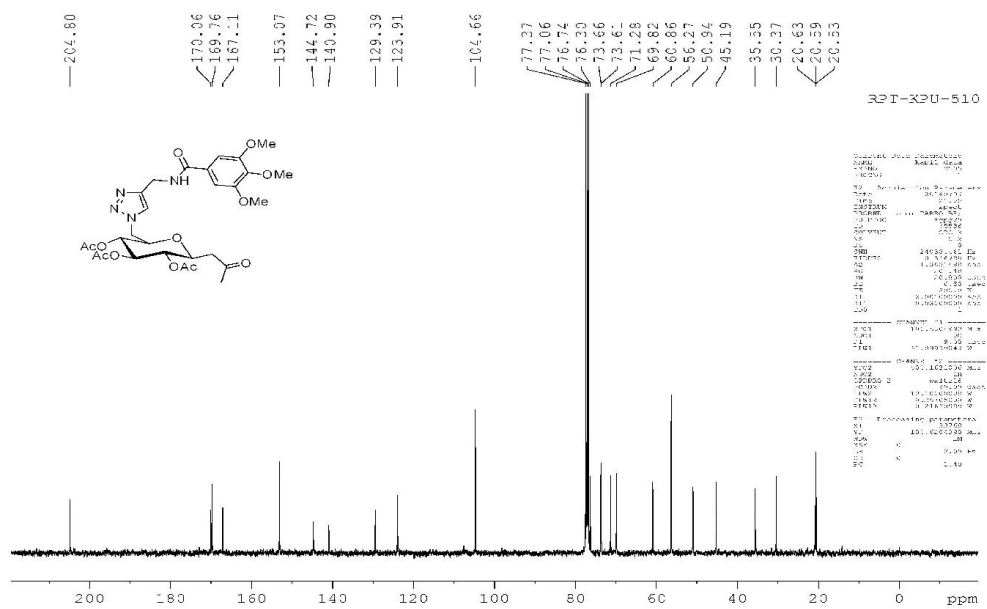
^1H NMR spectrum of compound 4a ^{13}C NMR spectrum of compound 4a

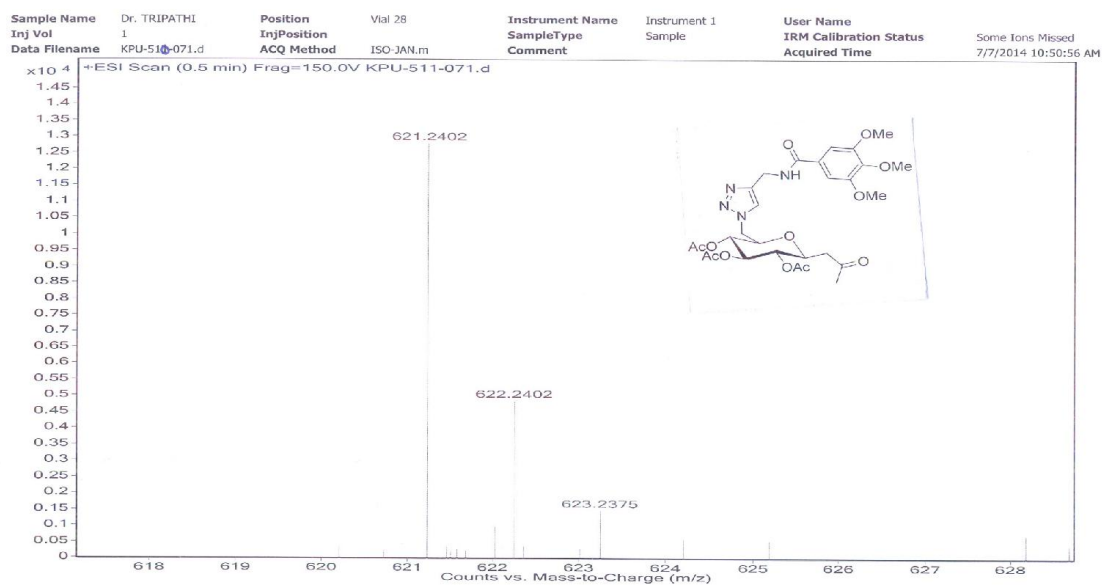
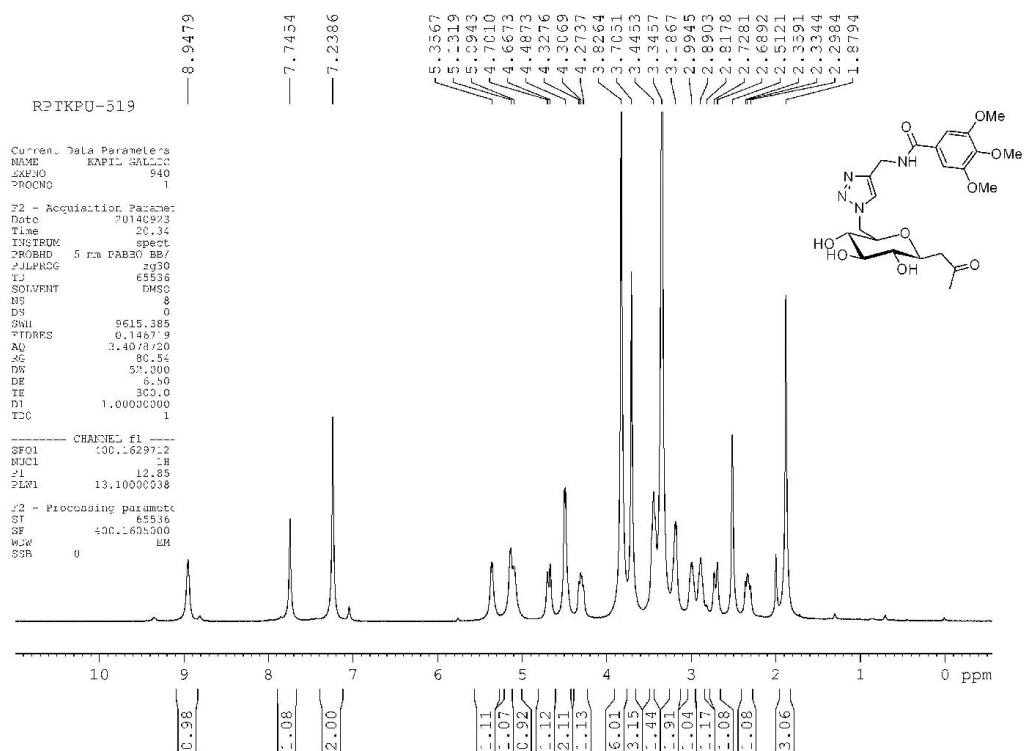
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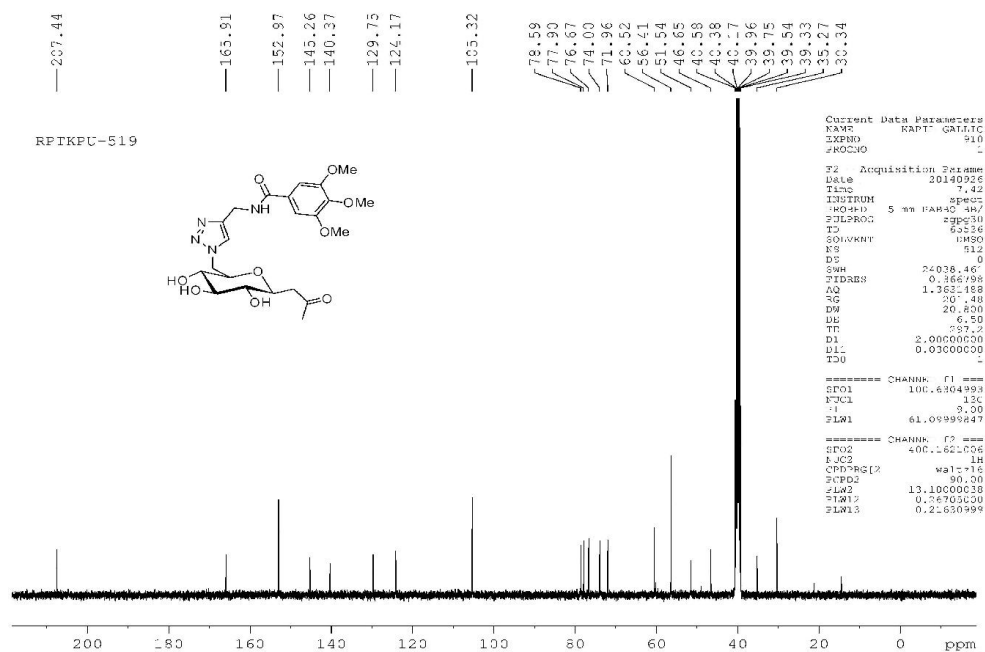
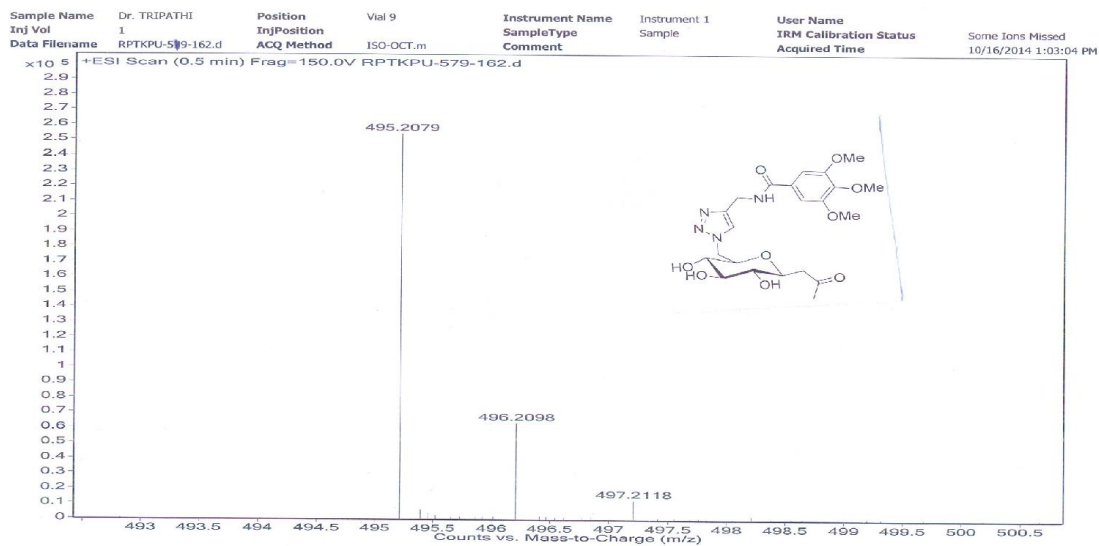
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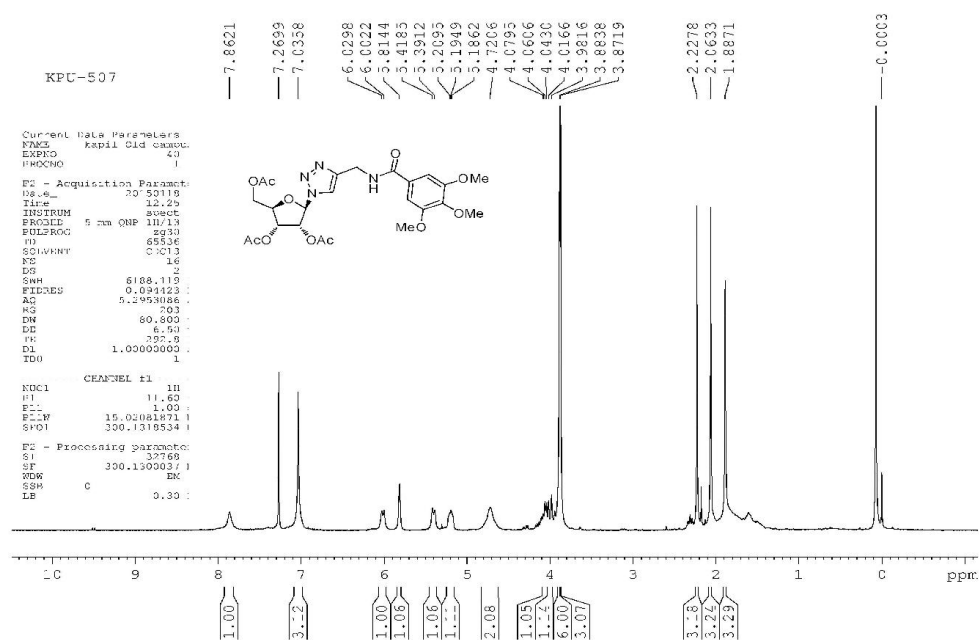
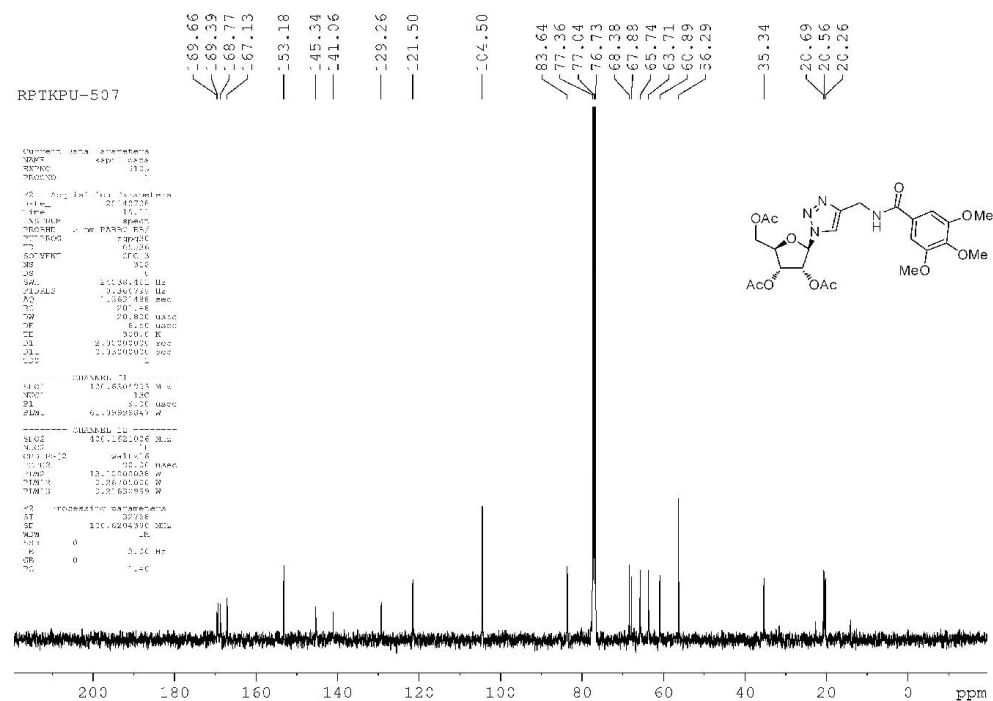
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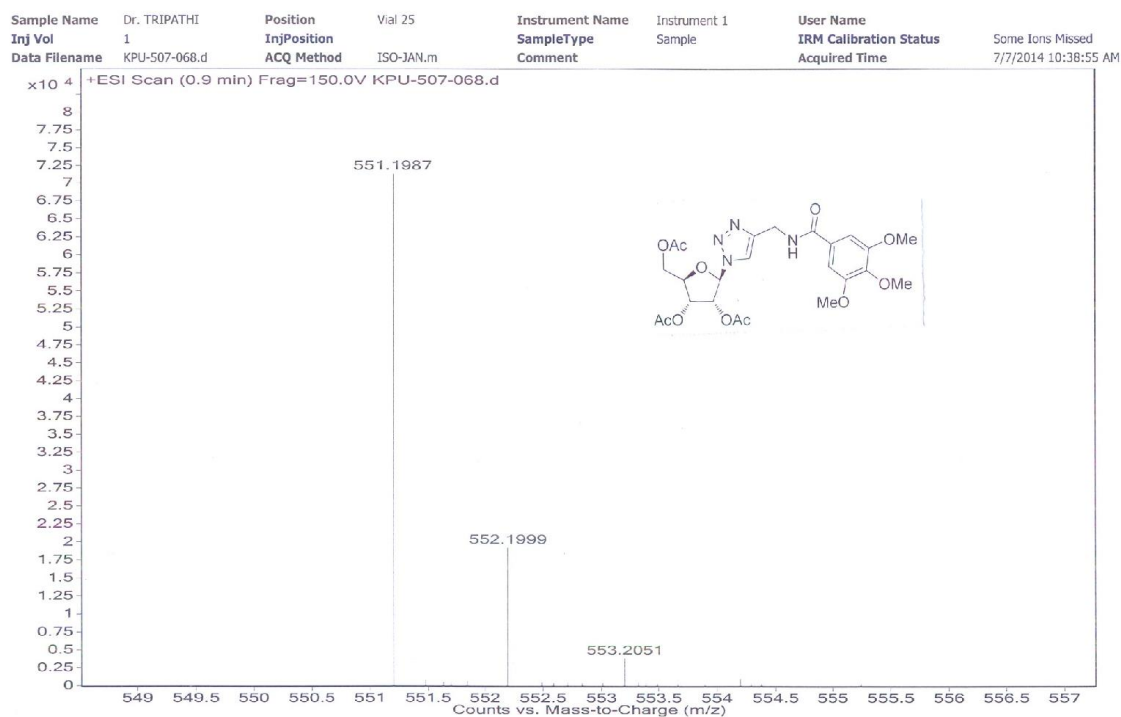
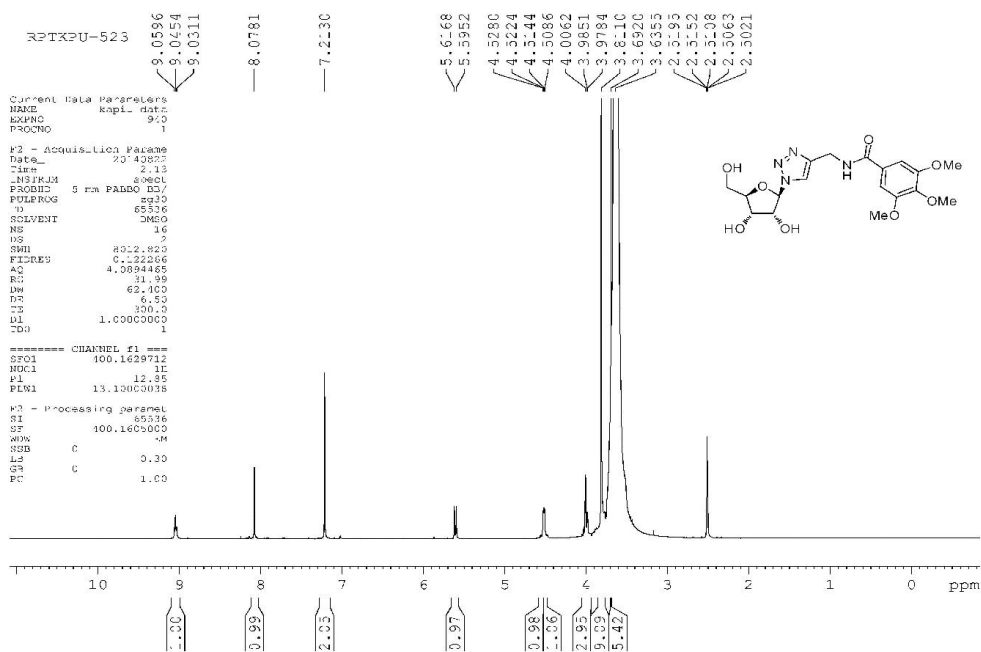


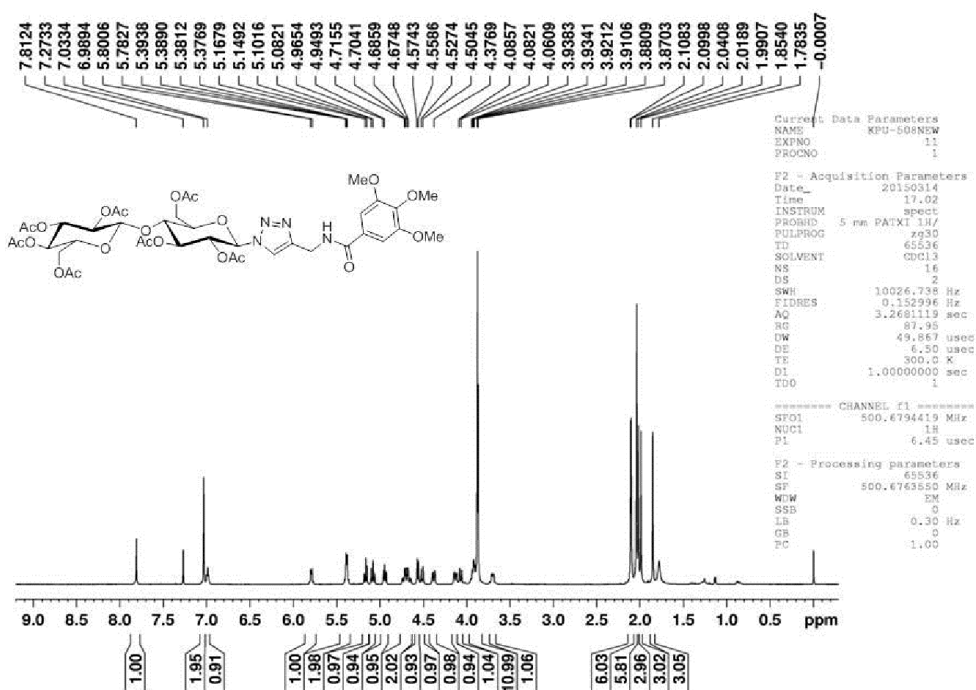
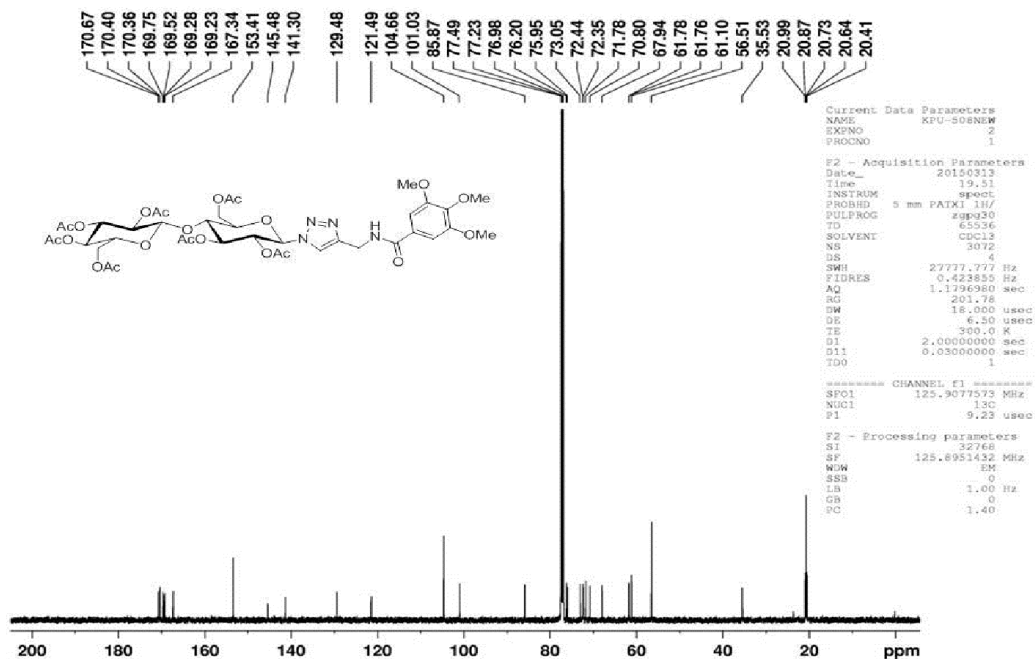
^1H NMR spectrum of compound **4b** ^{13}C NMR spectrum of compound **4b**

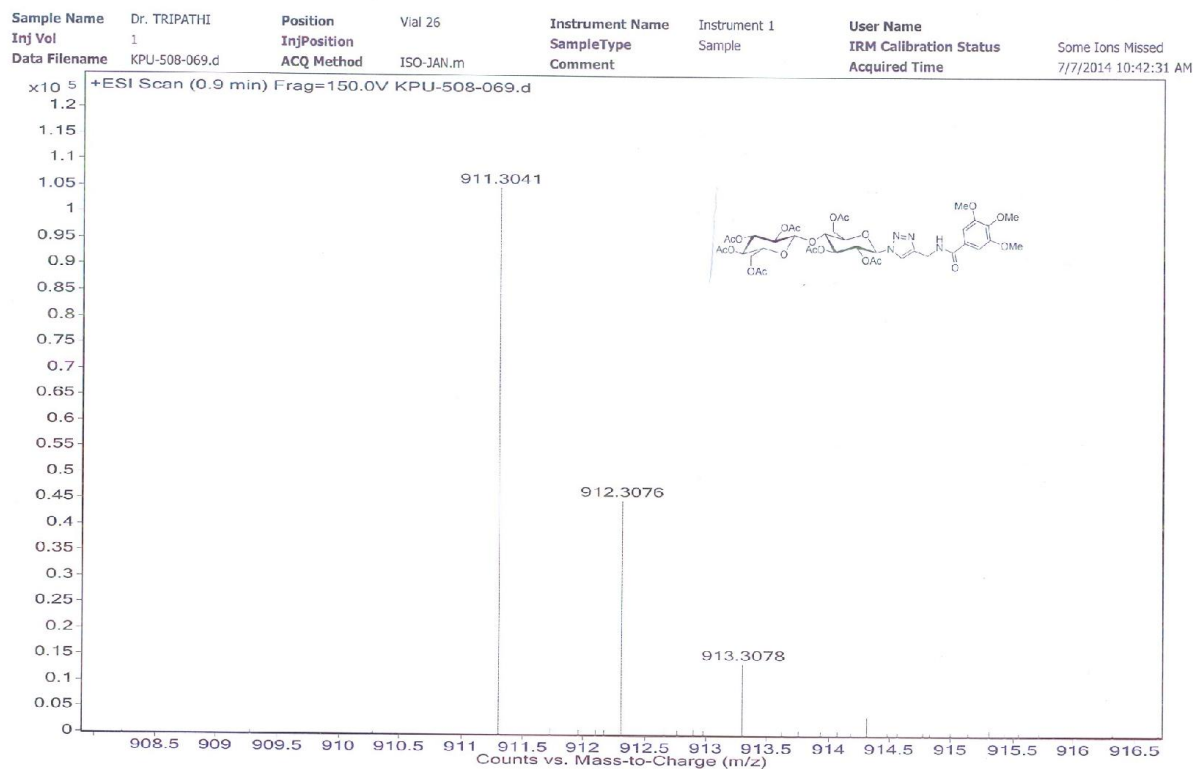
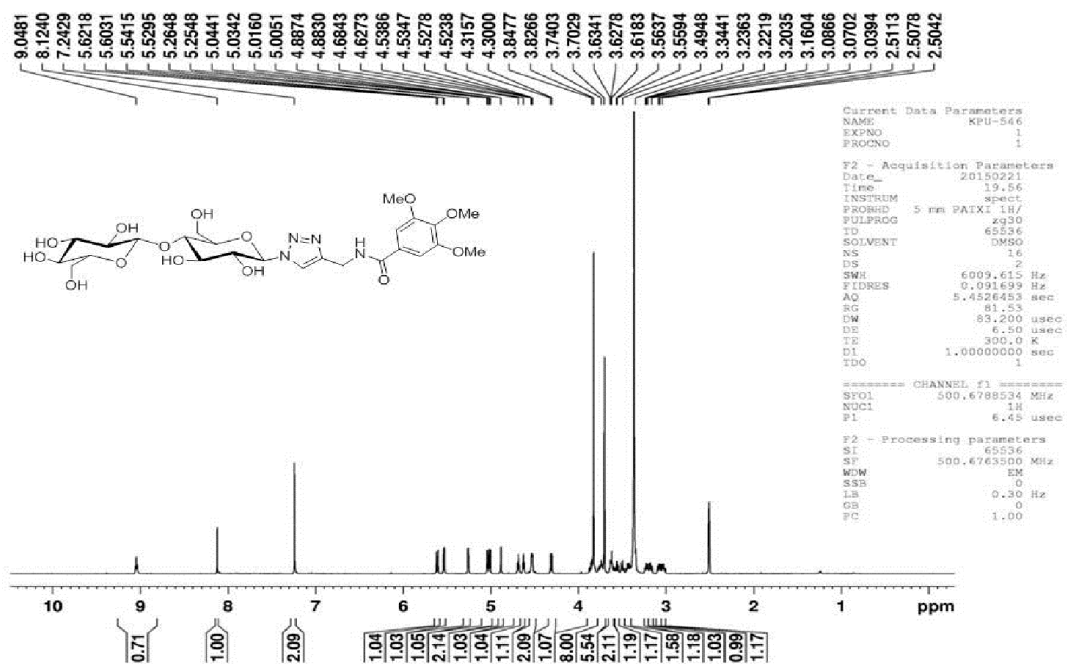
HRMS spectrum of compound **4b**¹H NMR spectrum of compound **4b'**

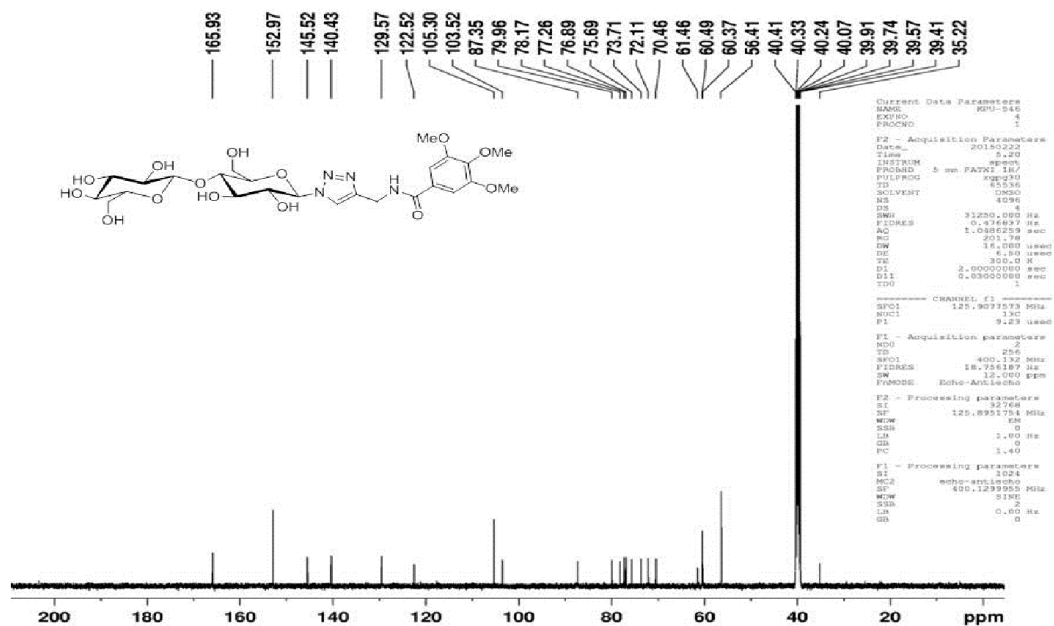
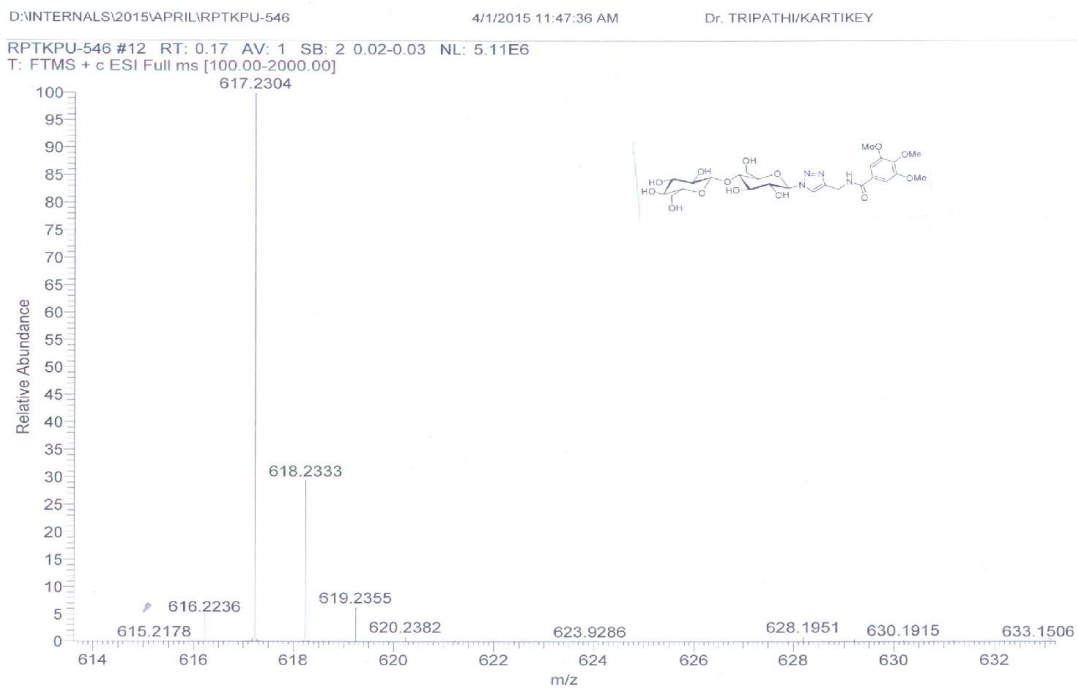
¹³C NMR spectrum of compound **4b'**HRMS spectrum of compound **4b'**

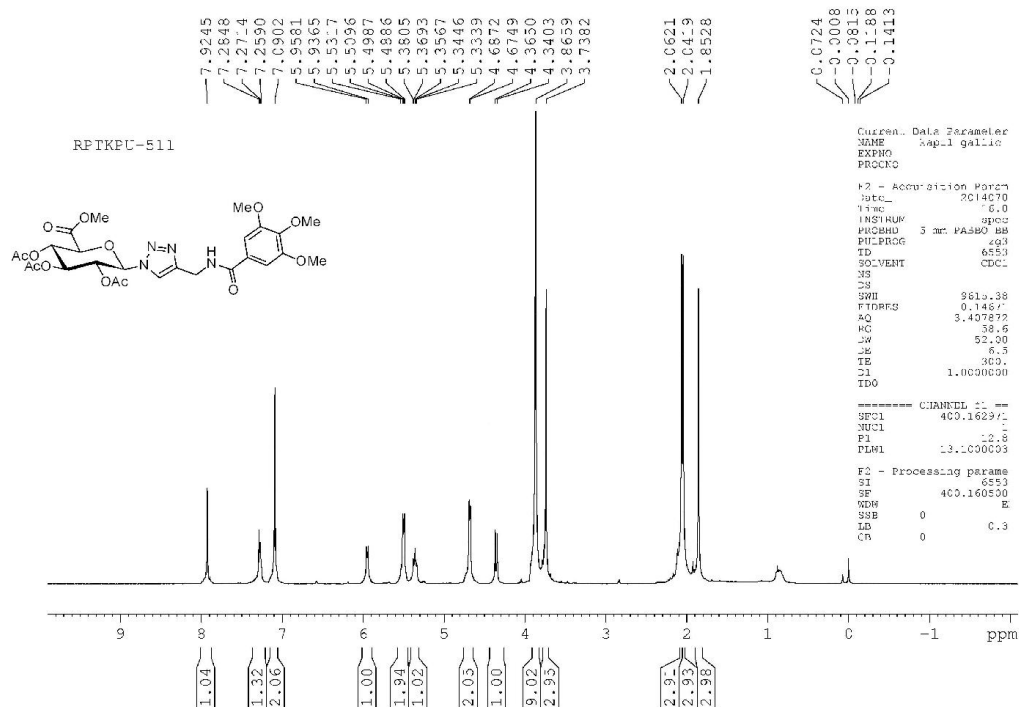
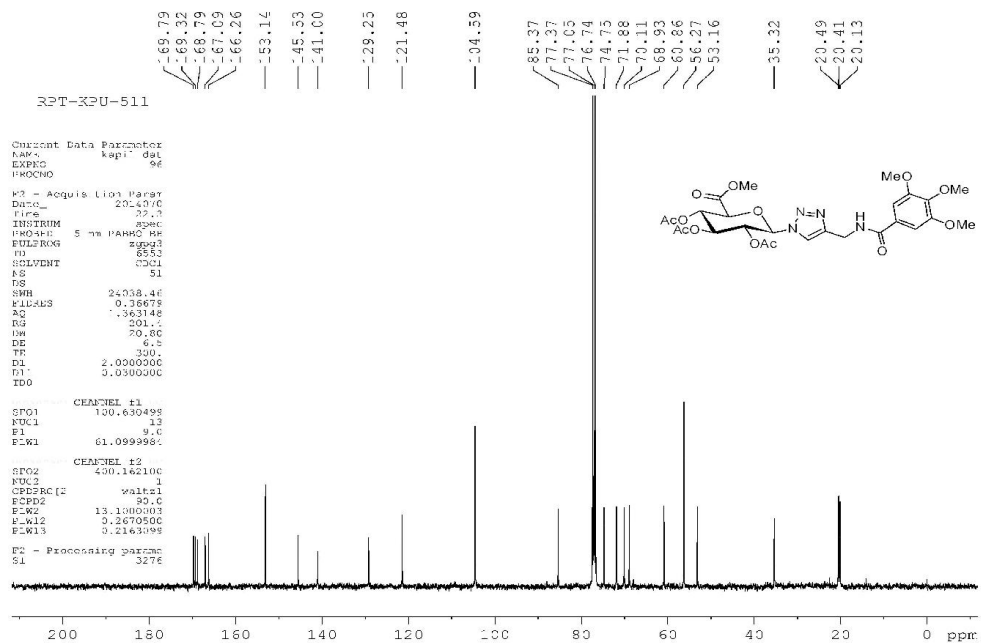
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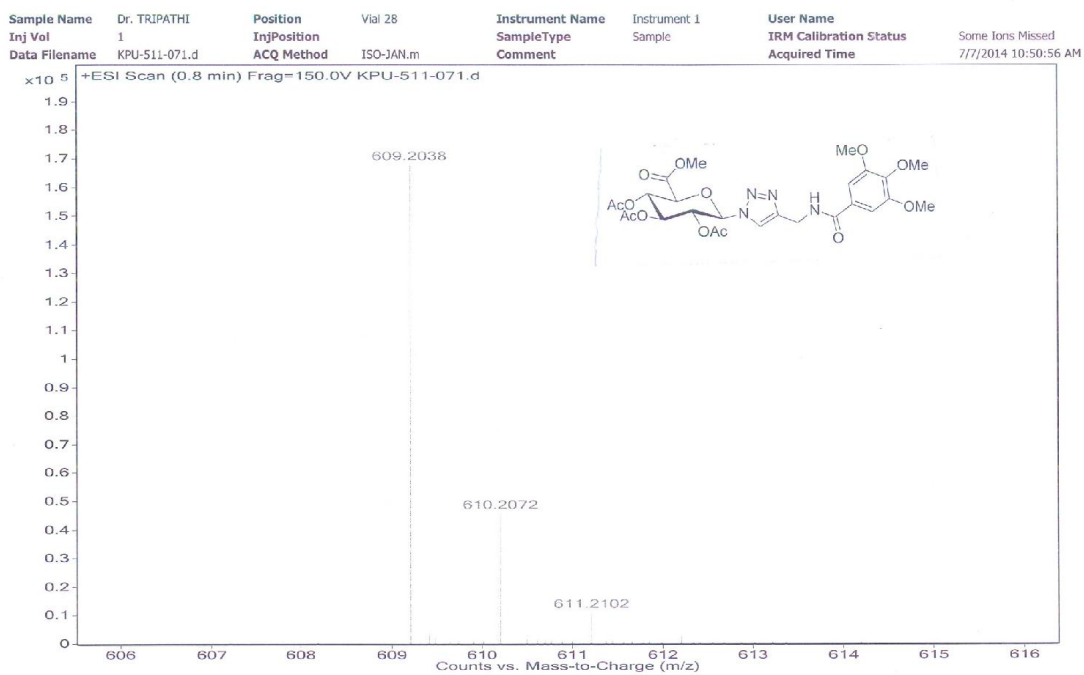
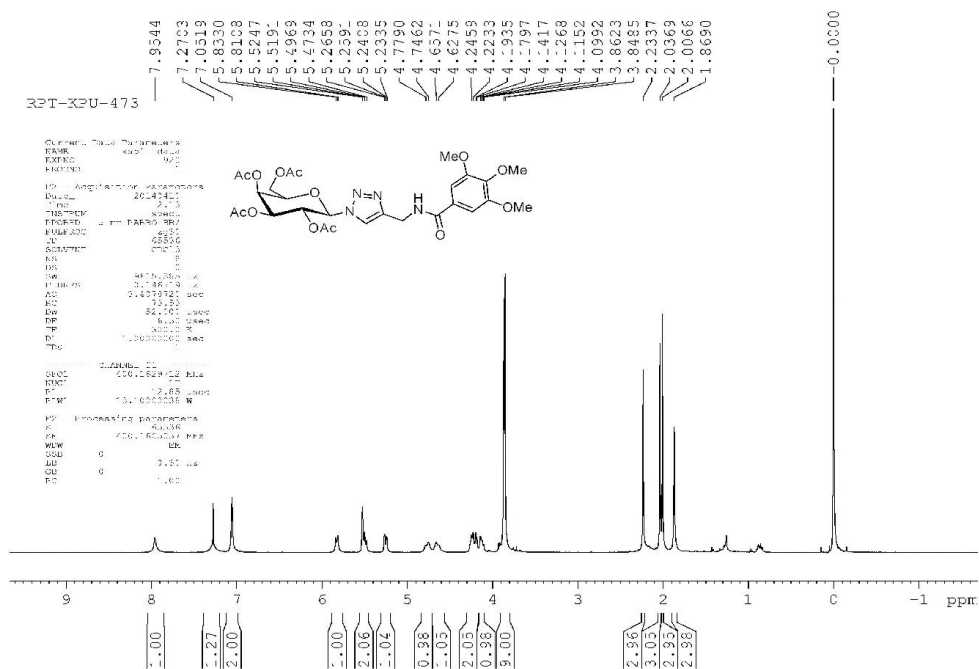
HRMS spectrum of compound **4c** ^1H NMR spectrum of compound **4c'**

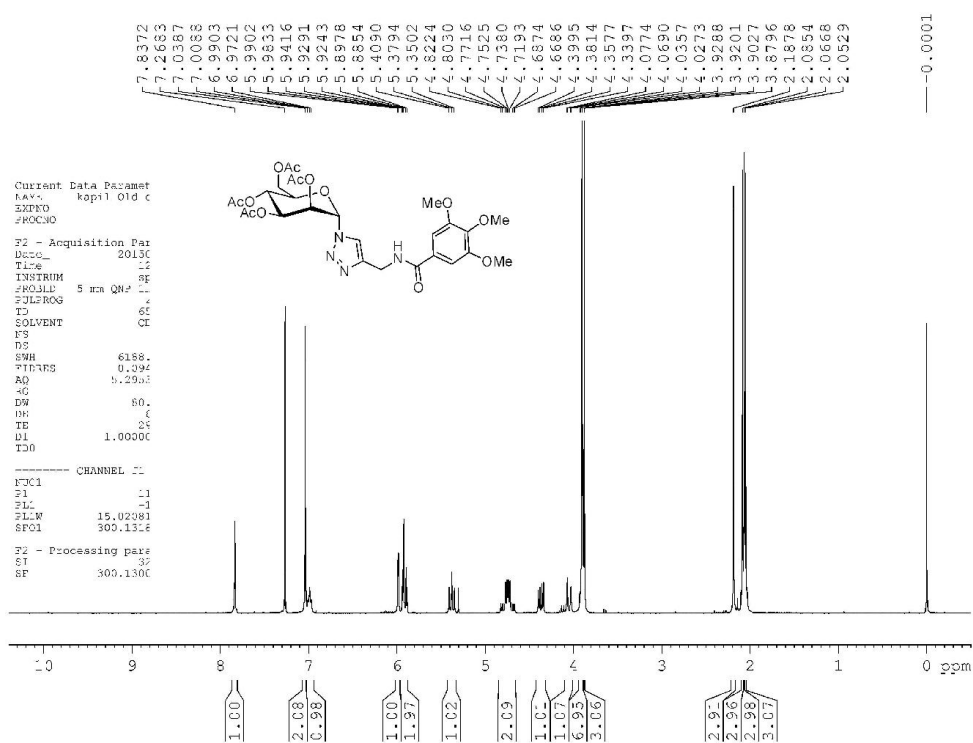
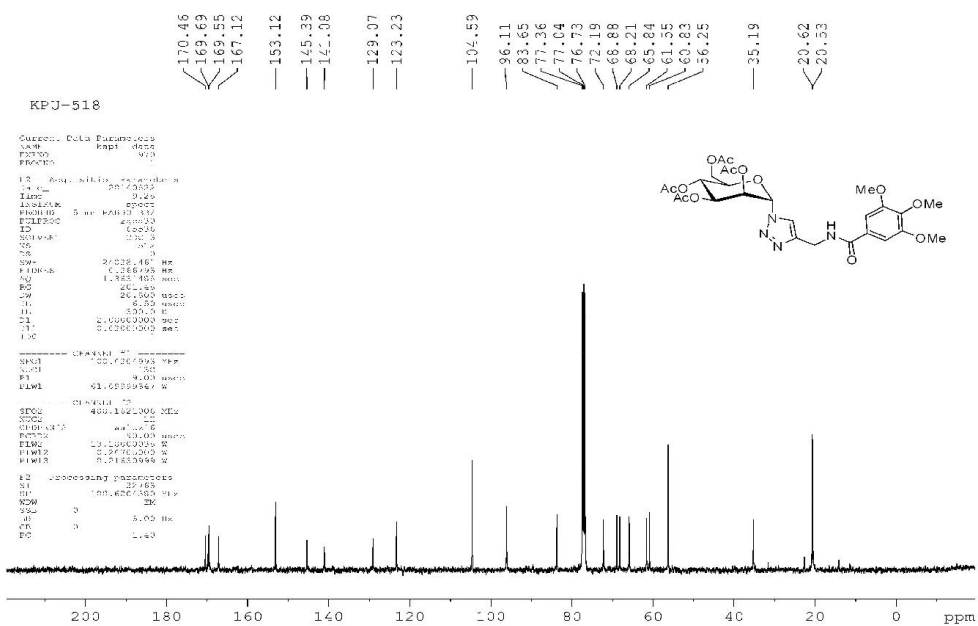
^1H NMR spectrum of compound **4d** ^{13}C NMR spectrum of compound **4d**

HRMS spectrum of compound **4d** ^1H NMR spectrum of compound **4d'**

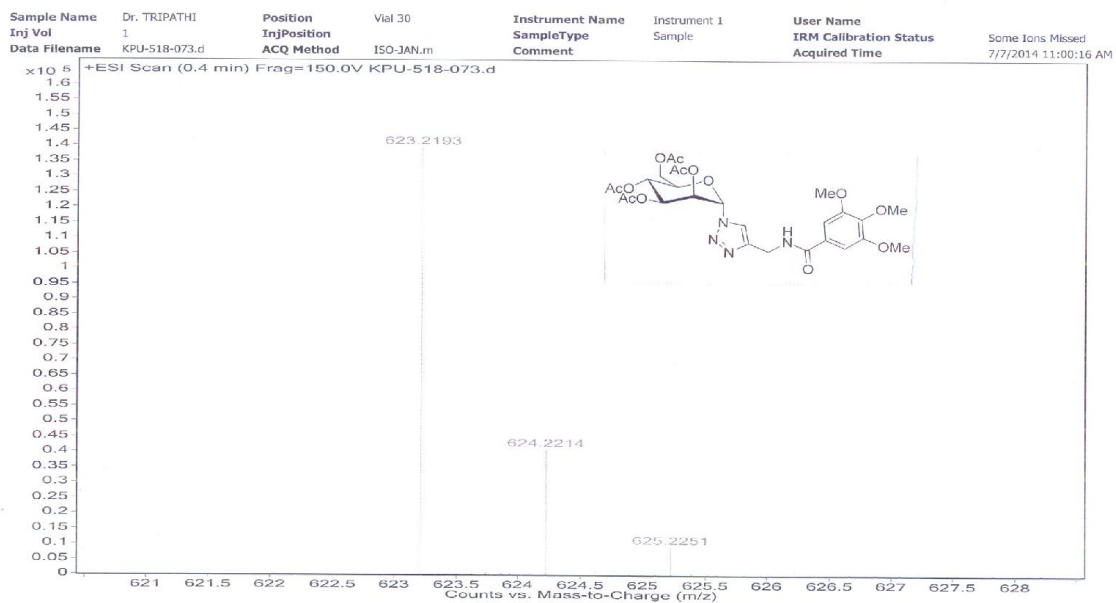
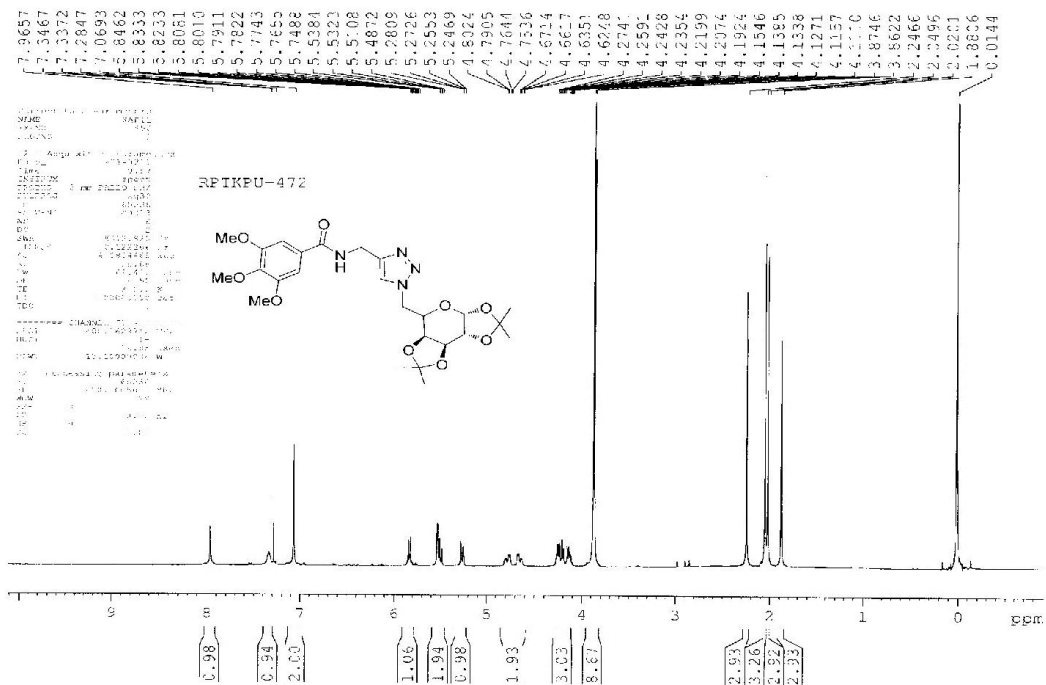
^{13}C NMR spectrum of compound **4d'**HRMS spectrum of compound **4d'**

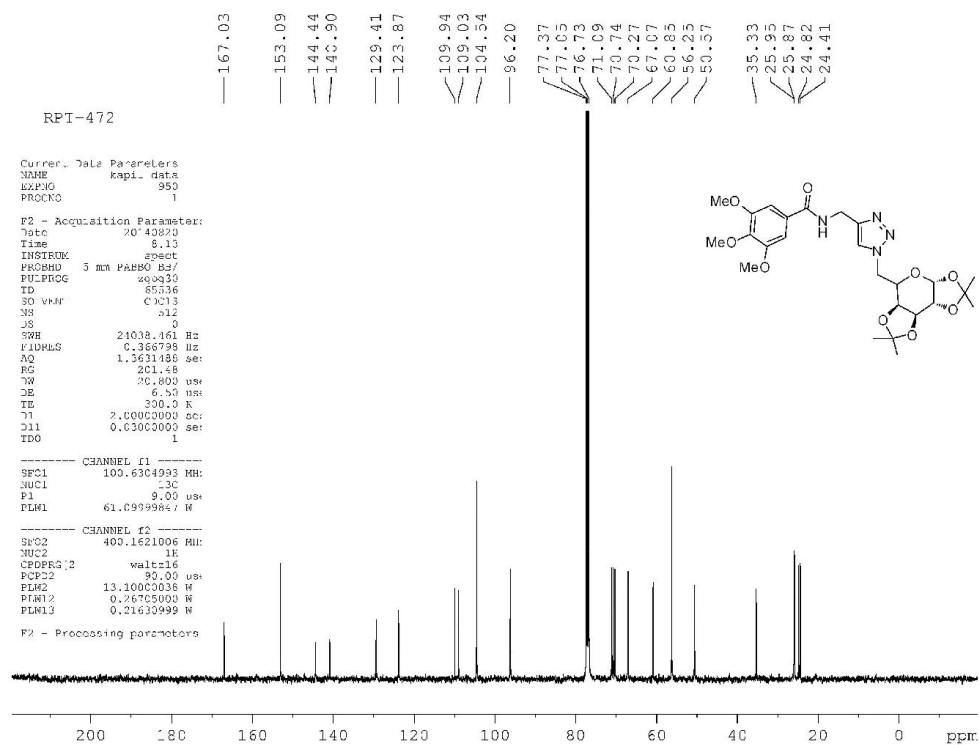
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HRMS spectrum of compound **4e**¹H NMR spectrum of compound **4f**

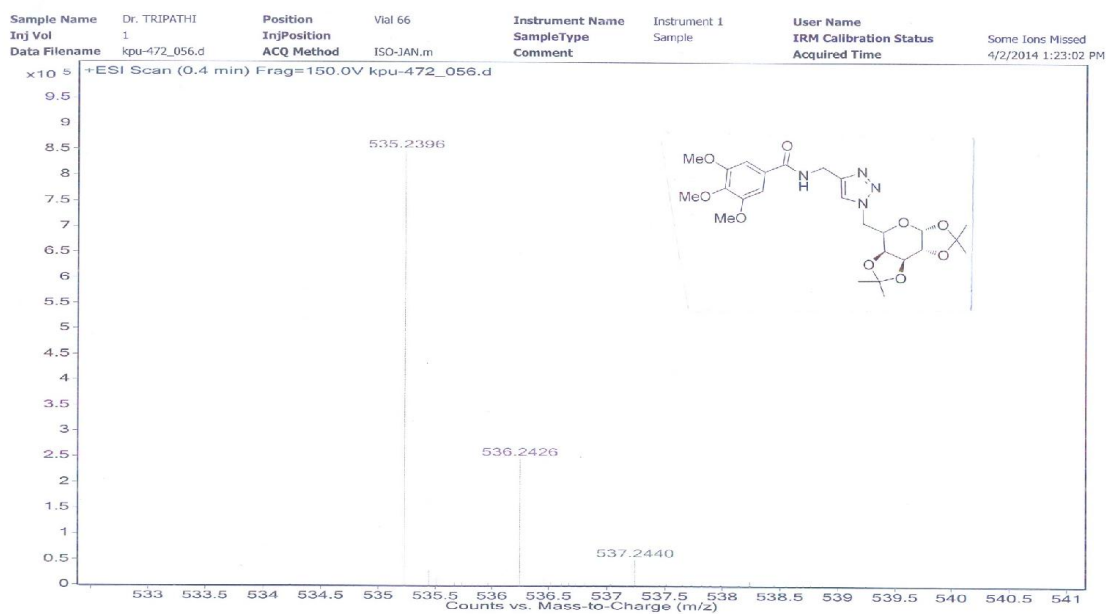
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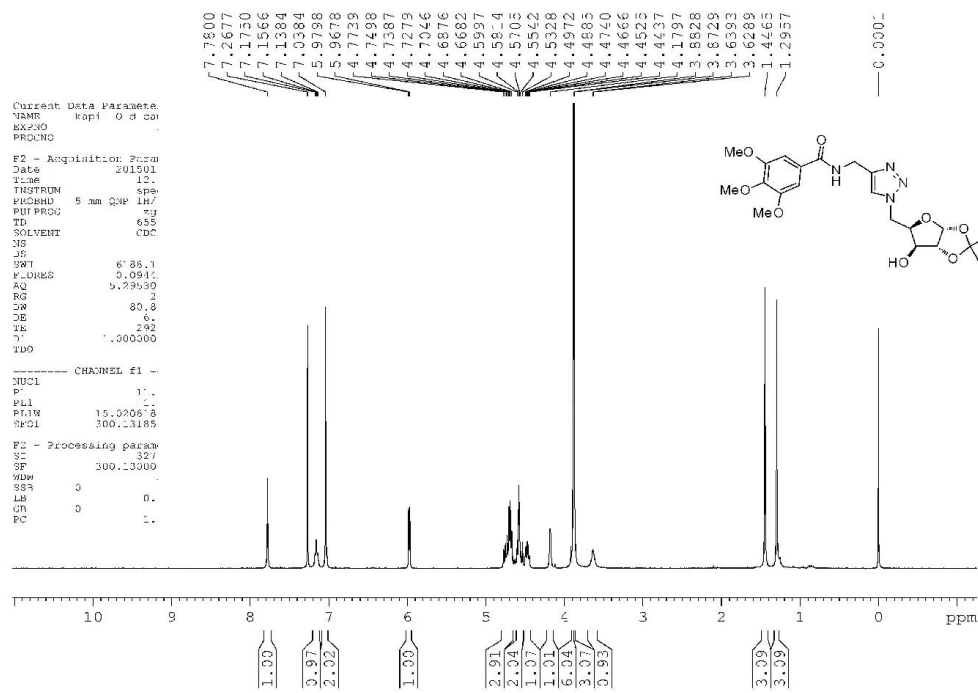
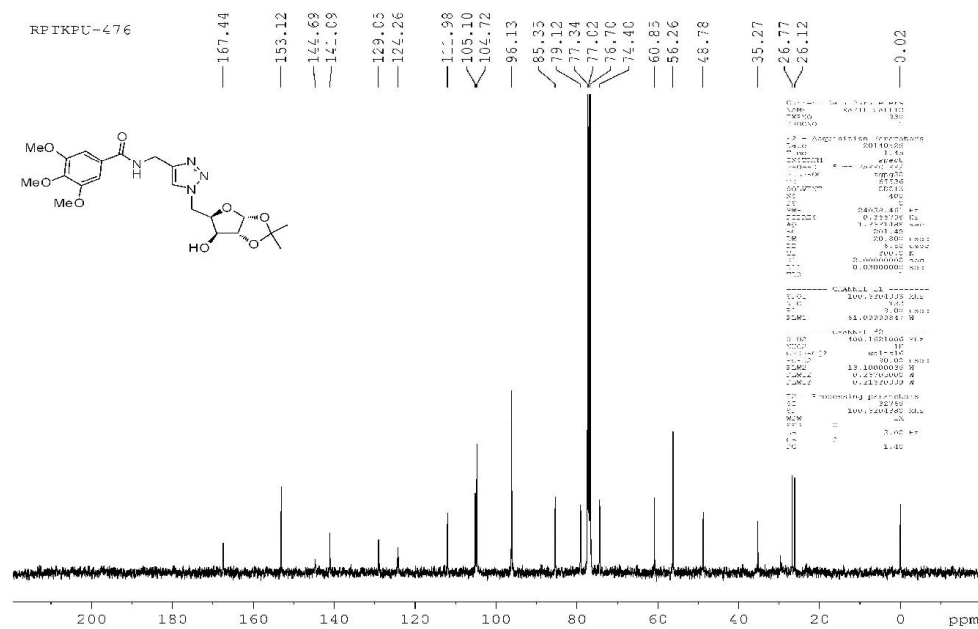
HRMS spectrum of compound 4g

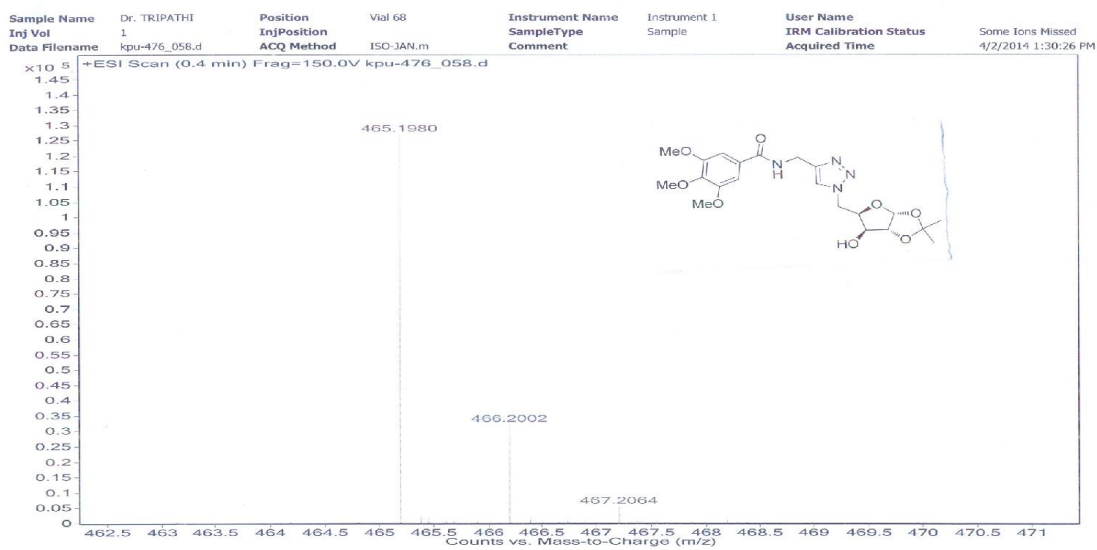
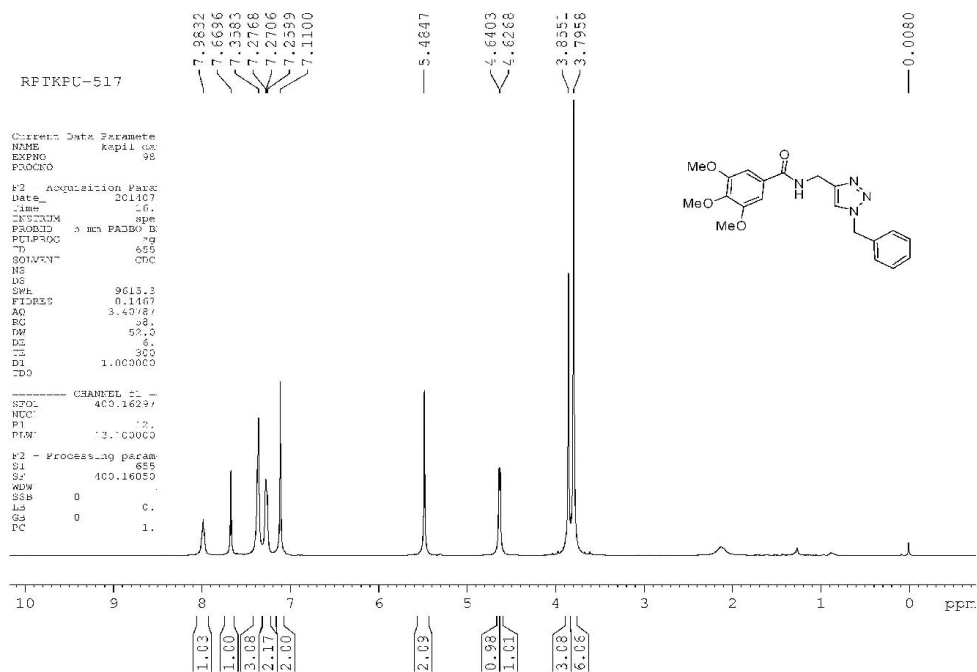
¹H NMR spectrum of compound 4h

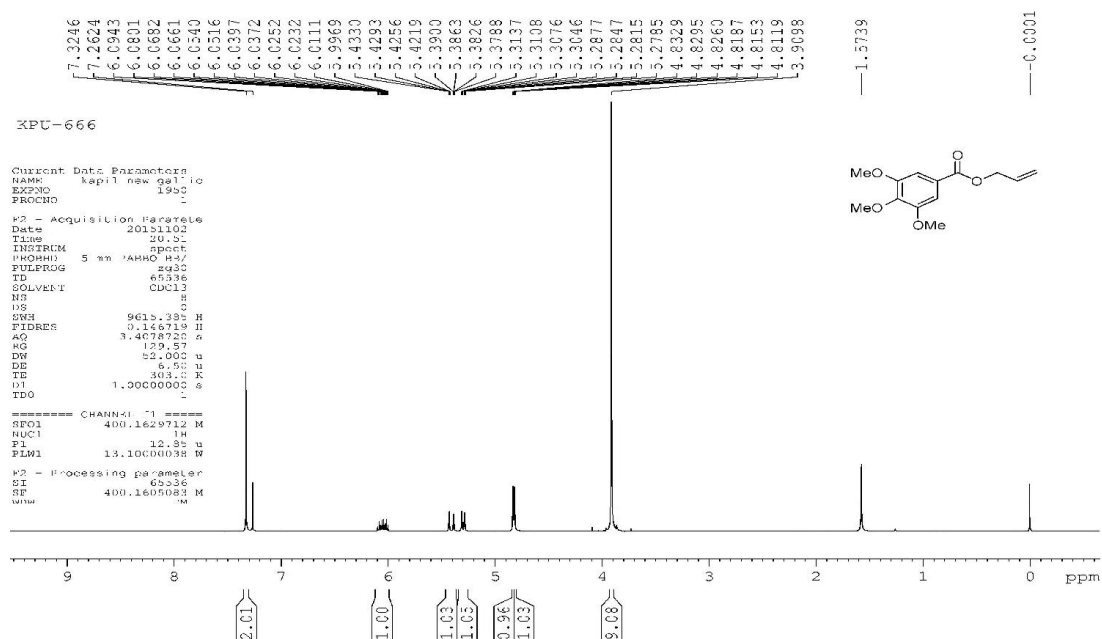
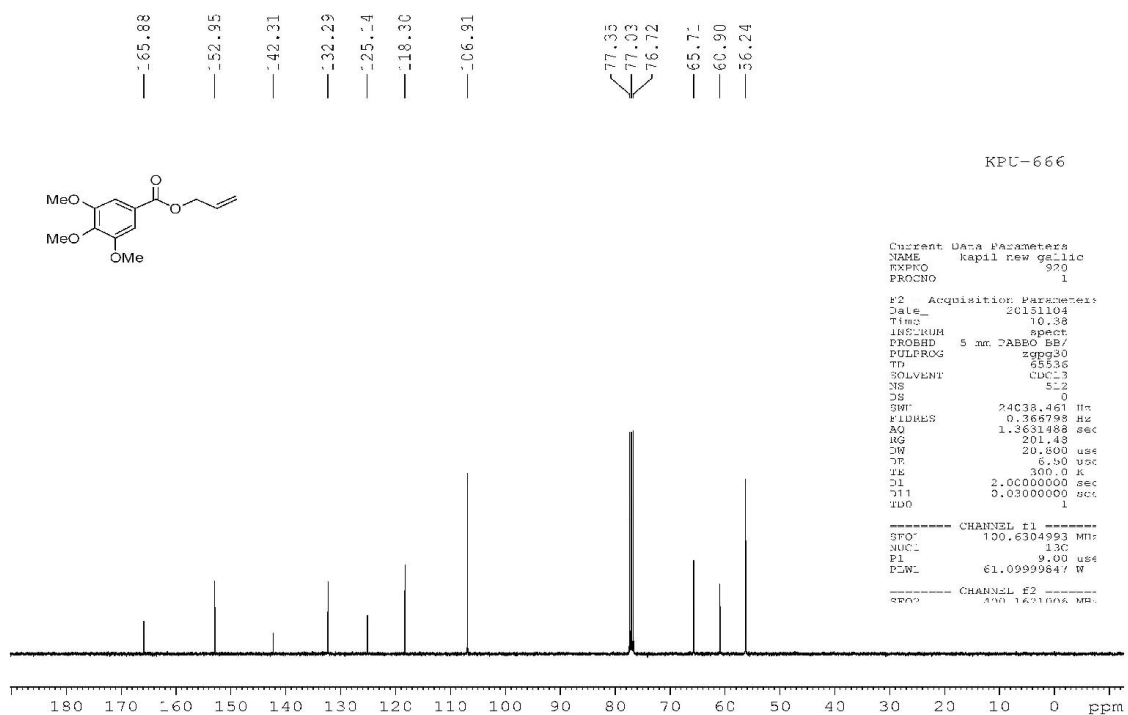
¹³C NMR spectrum of compound 4h

HRMS spectrum of compound 4h

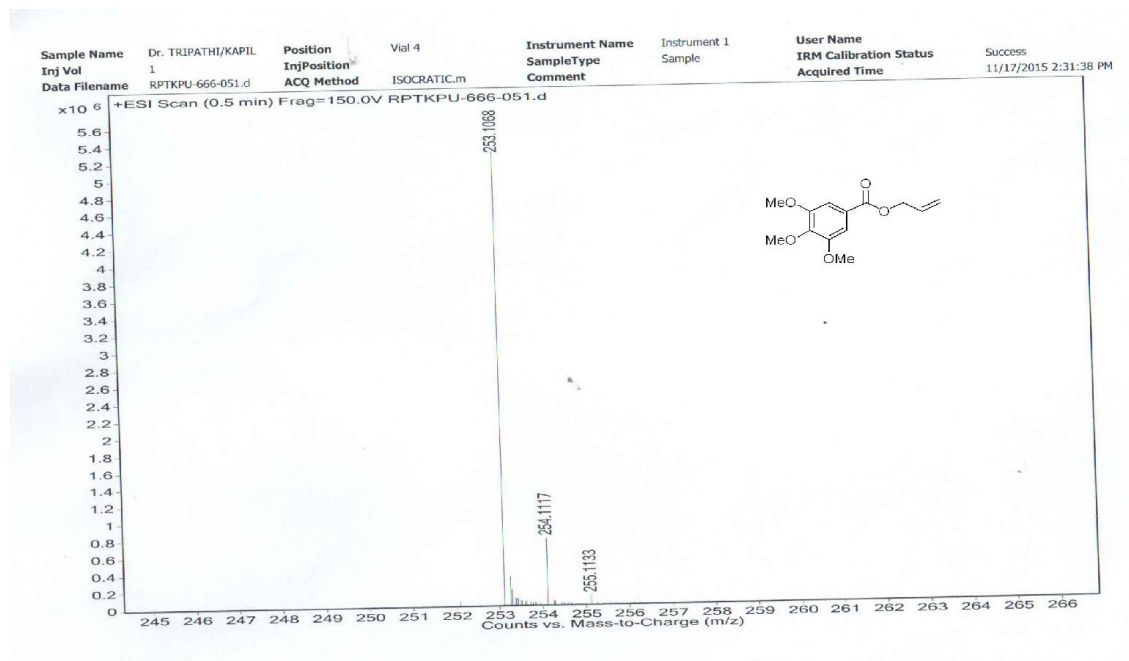
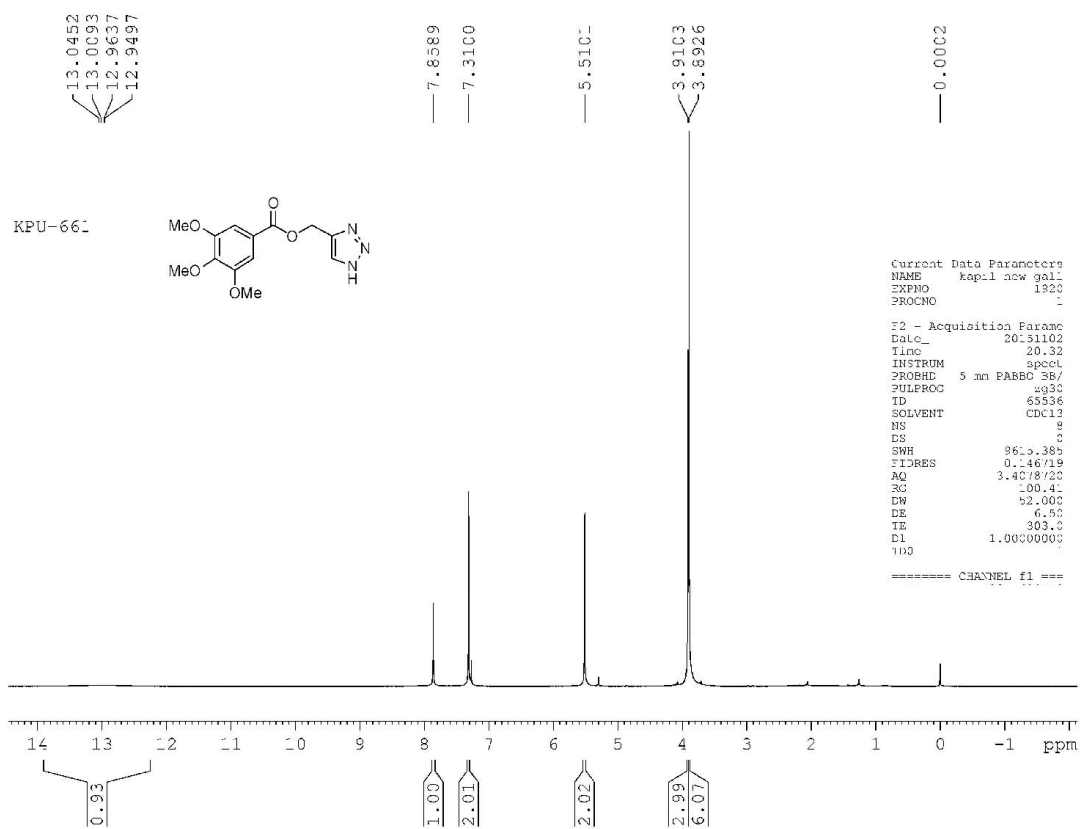


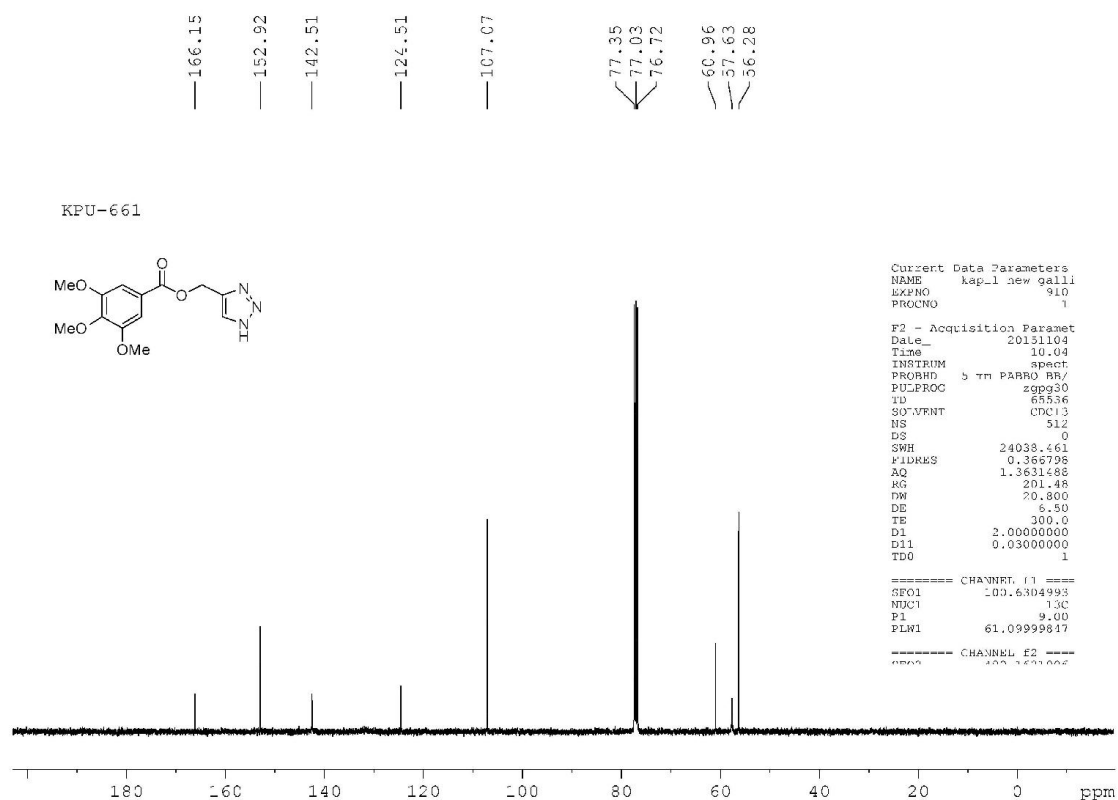
¹H NMR spectrum of compound 4i¹³C NMR spectrum of compound 4i

HRMS spectrum of compound **4i** ^1H NMR spectrum of compound **4j**

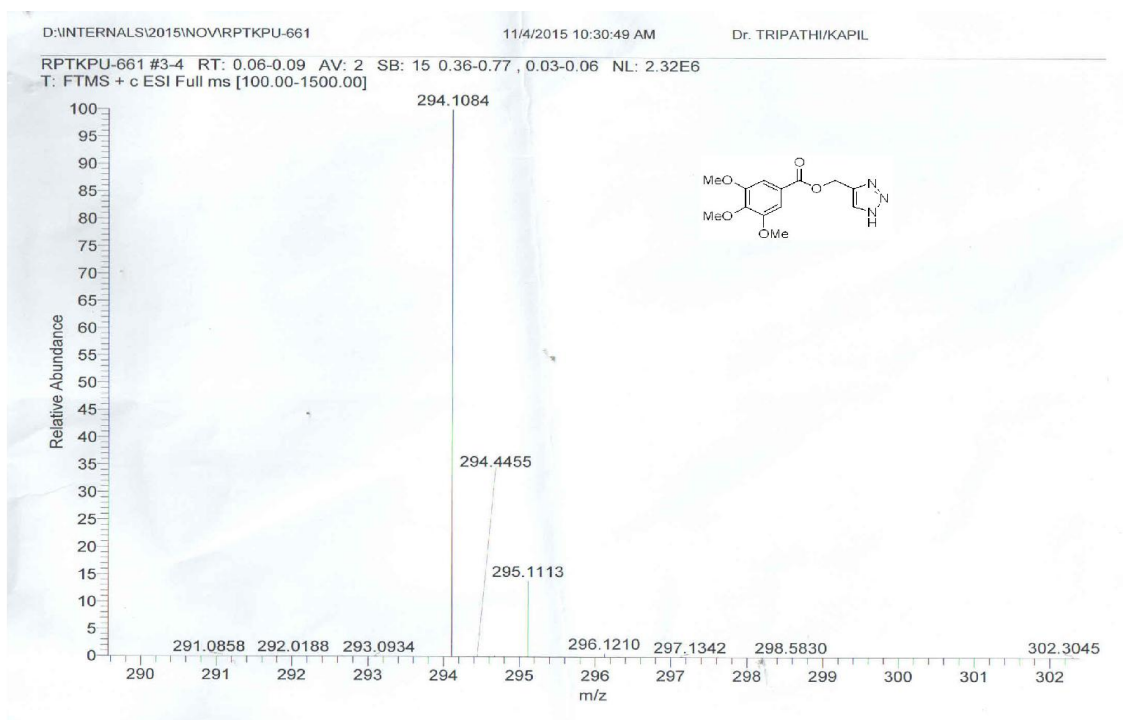
¹H NMR spectrum of compound 5¹³C NMR spectrum of compound 5

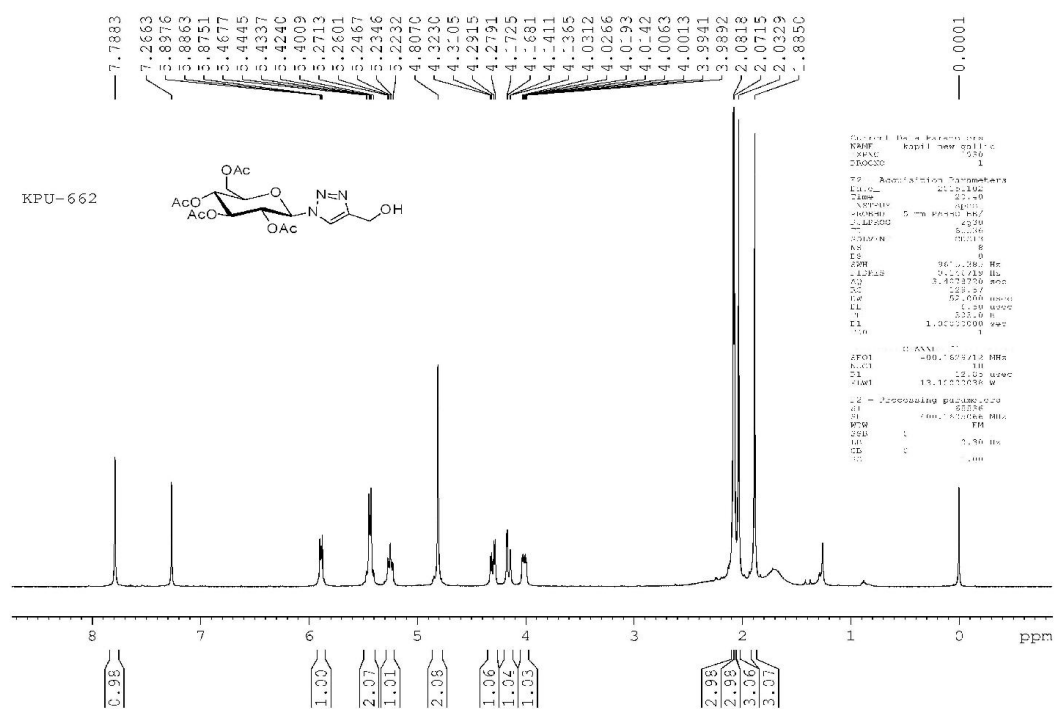
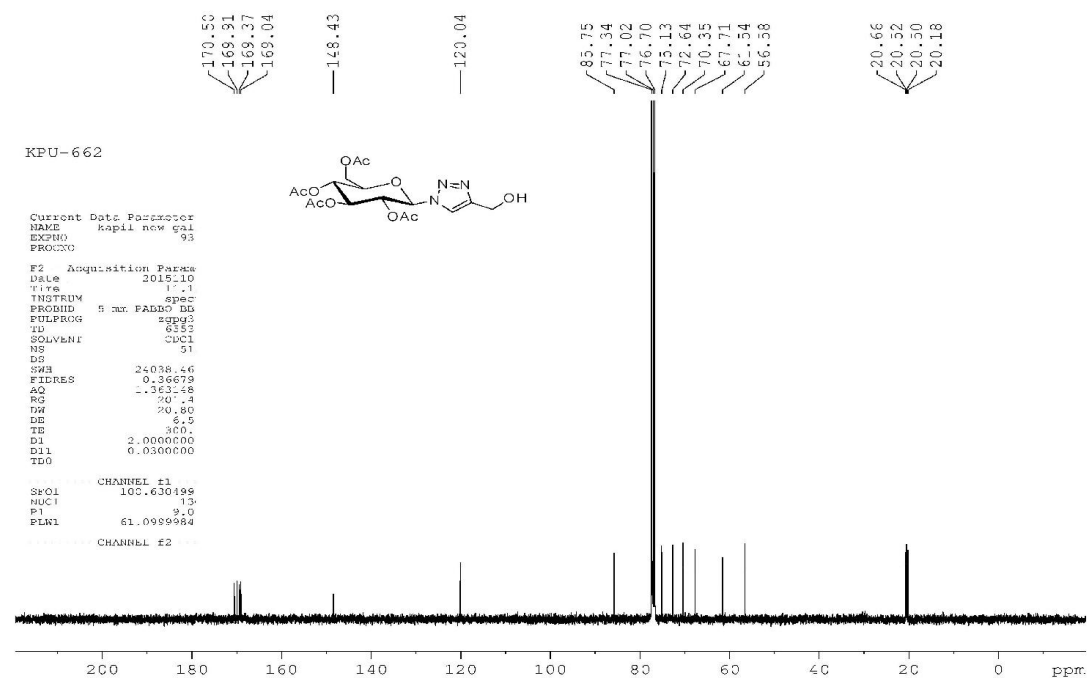
HRMS spectrum of compound 5

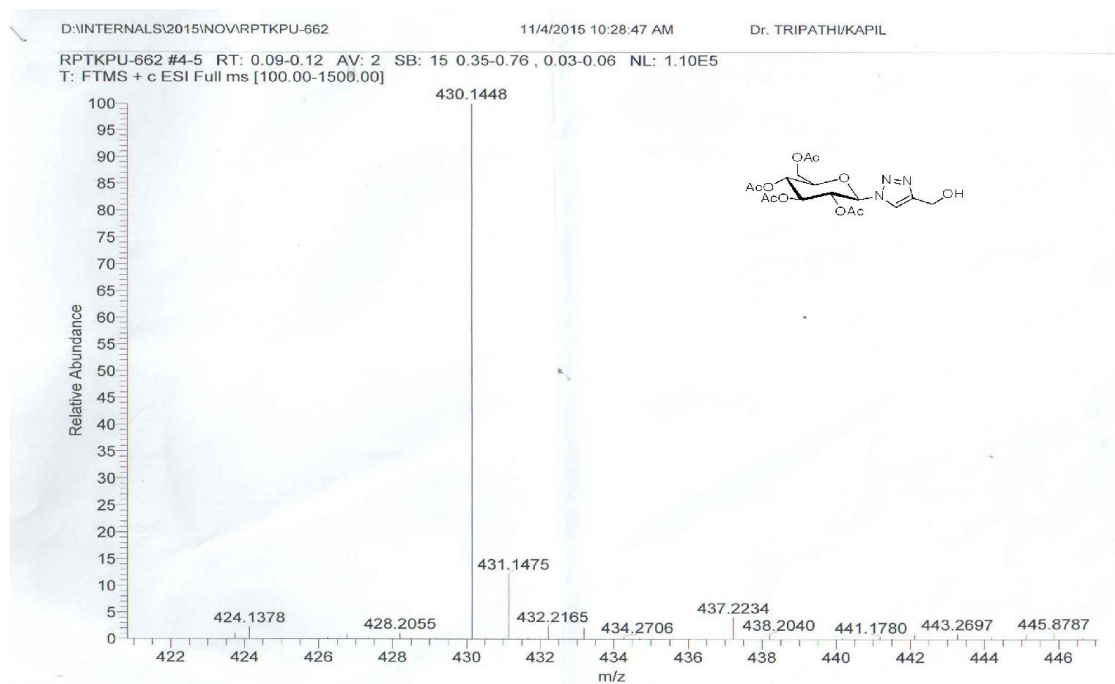
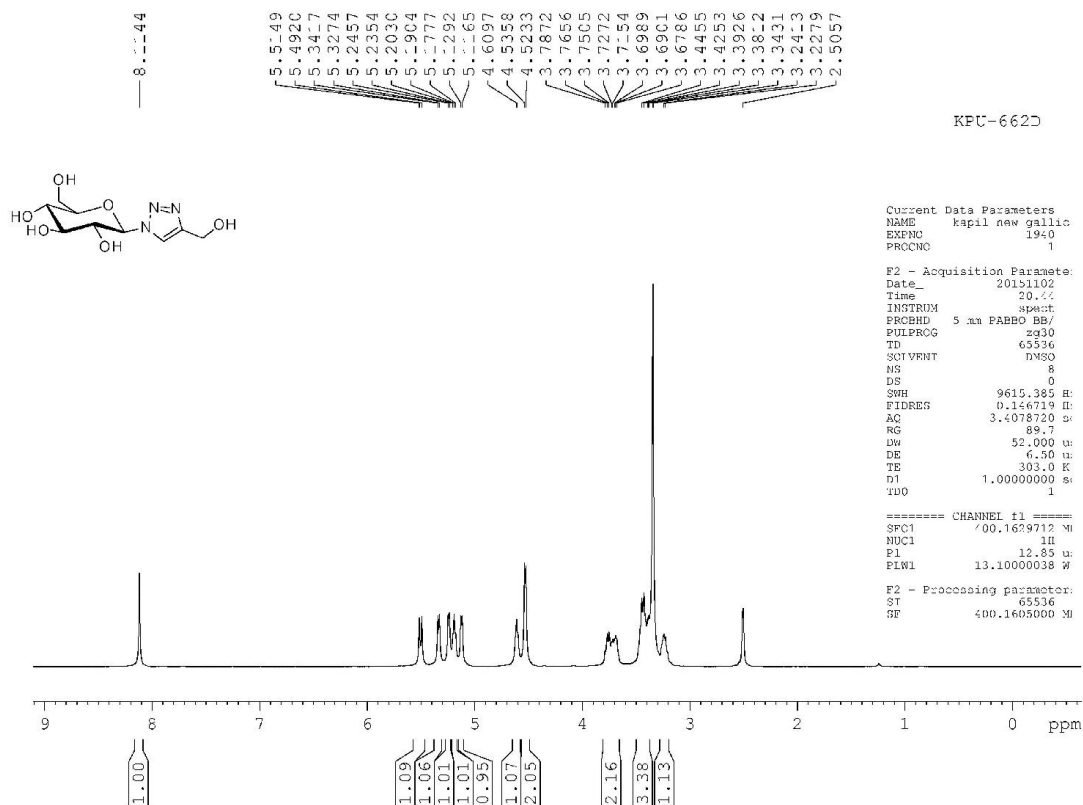
 ^1H NMR spectrum of compound 6

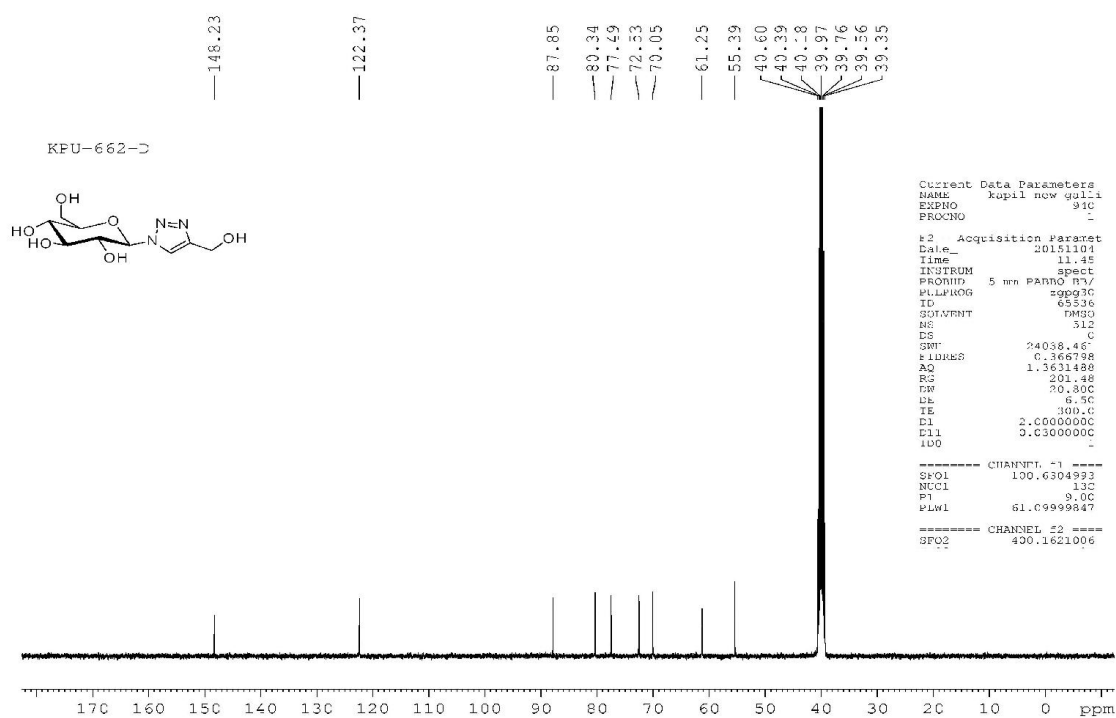
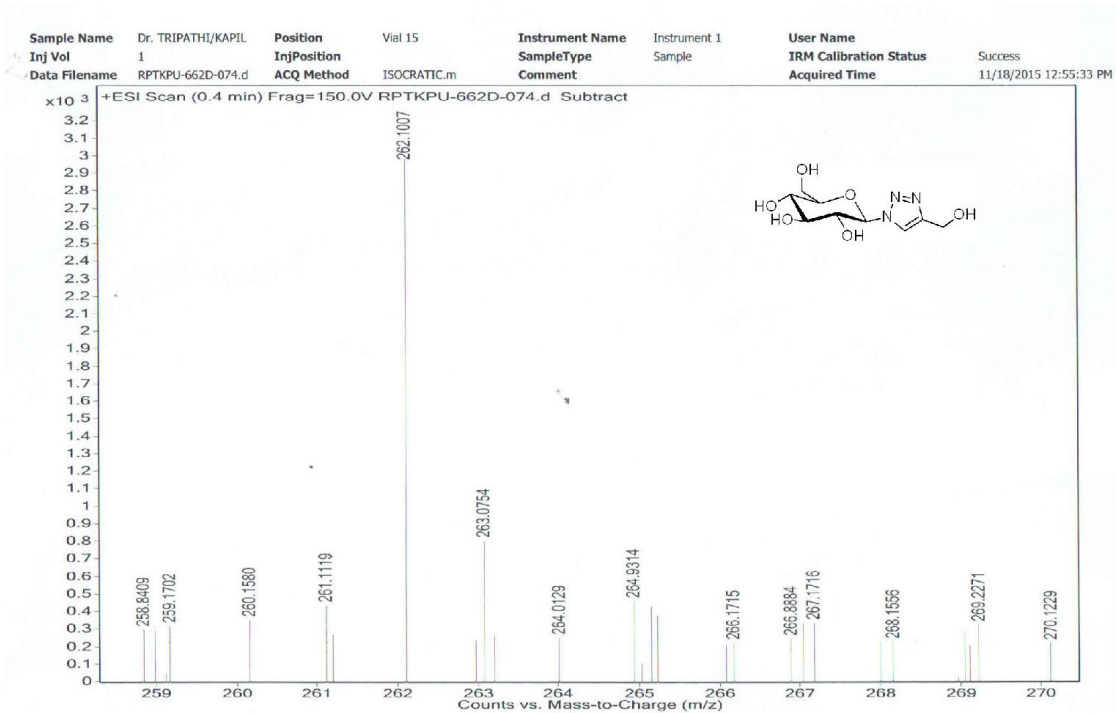
¹³C NMR spectrum of compound 6

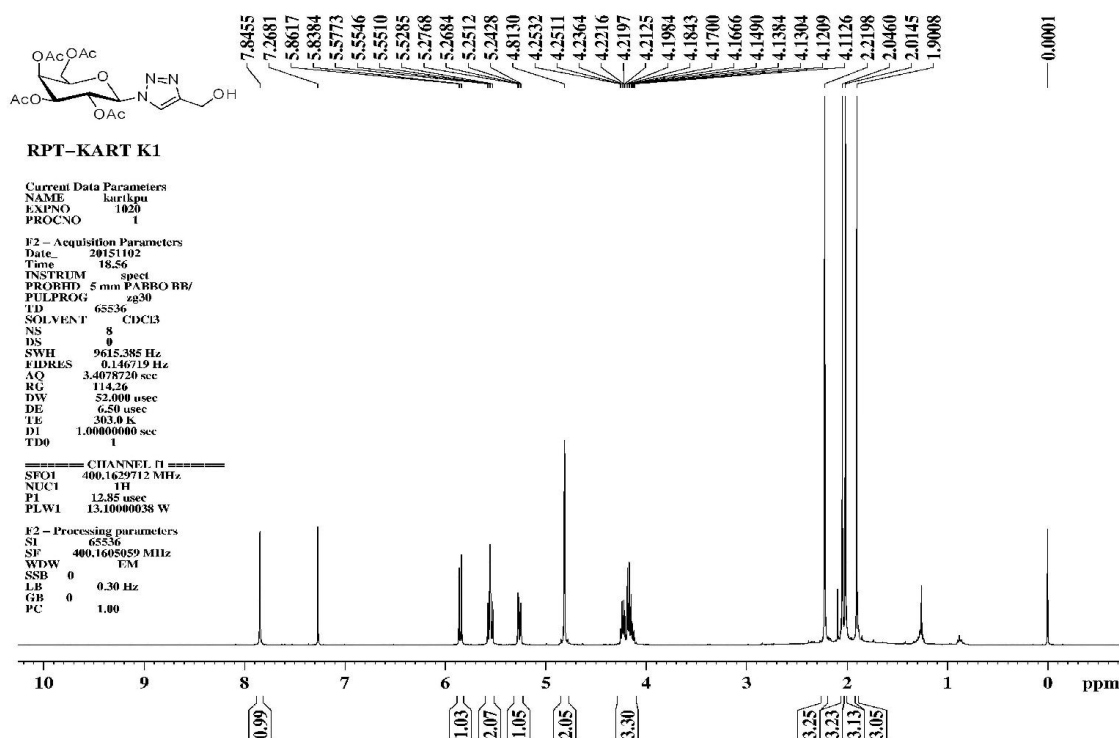
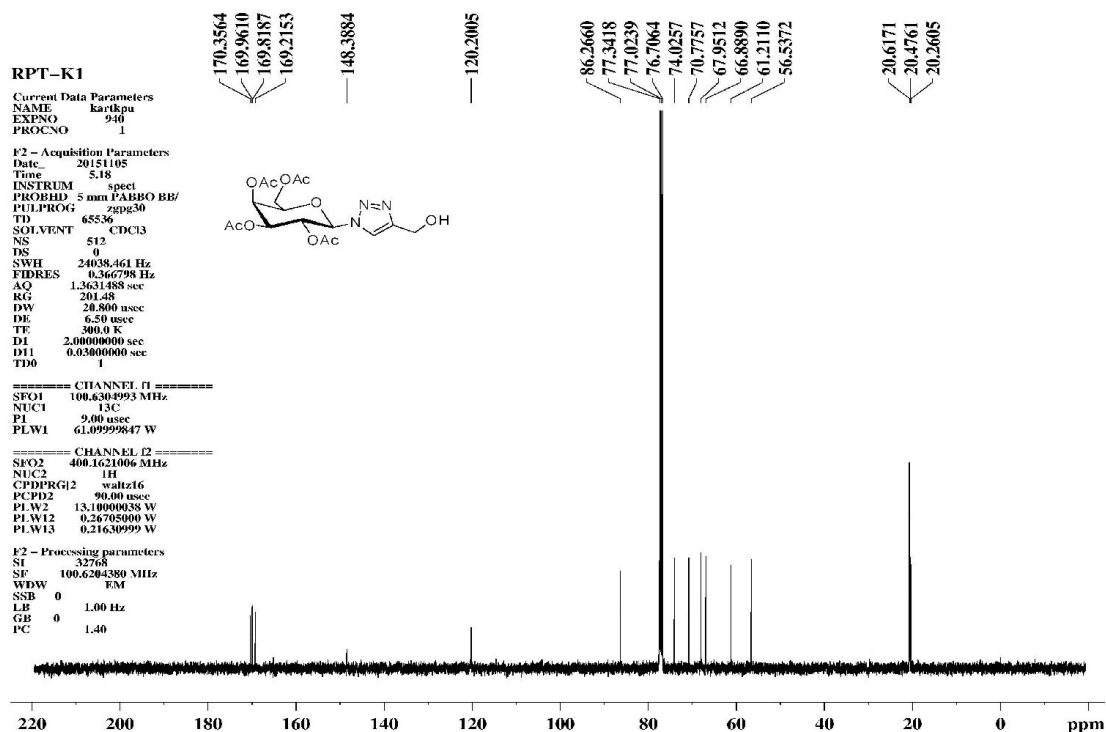
HRMS spectrum of compound 6

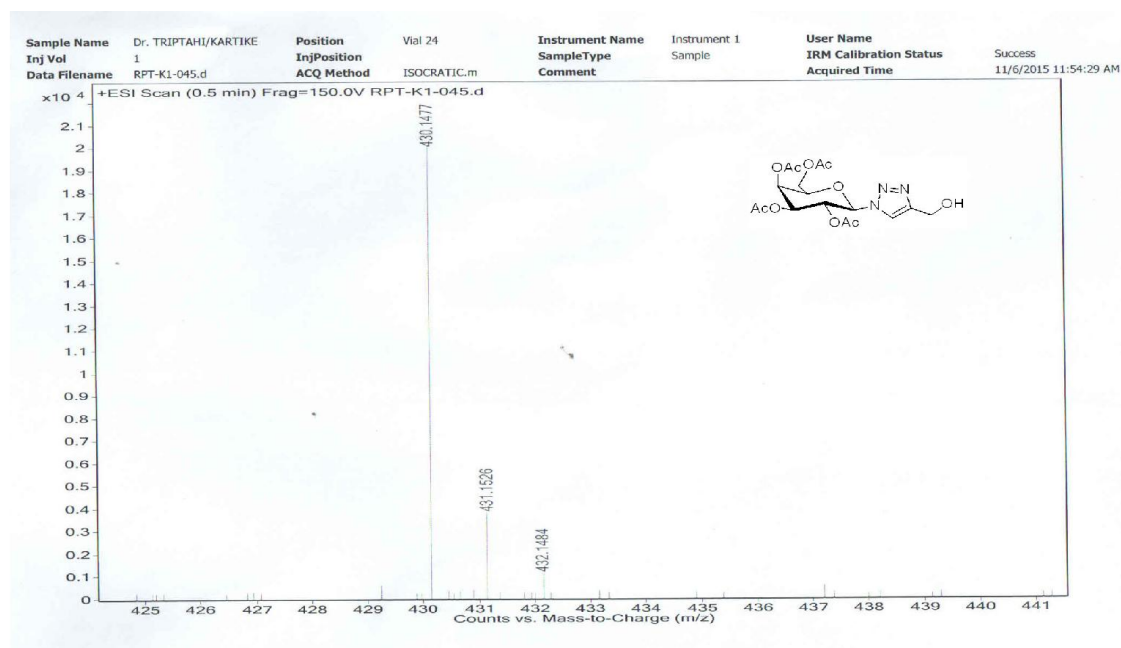
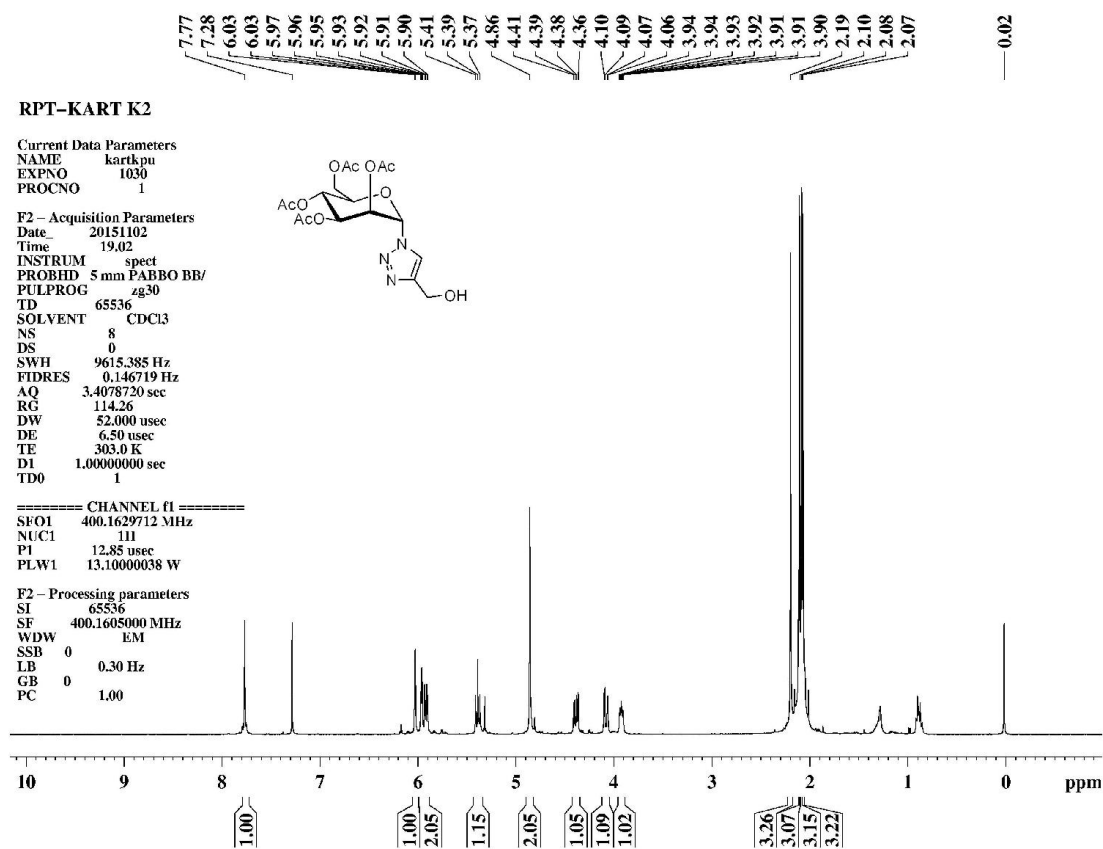


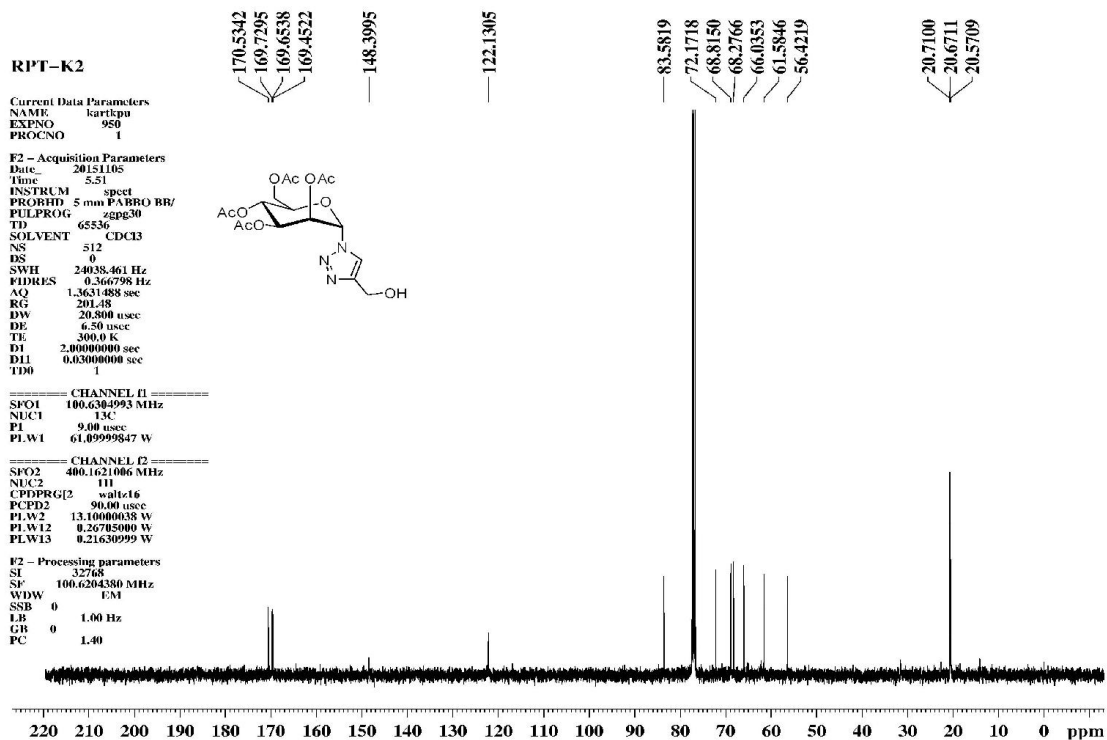
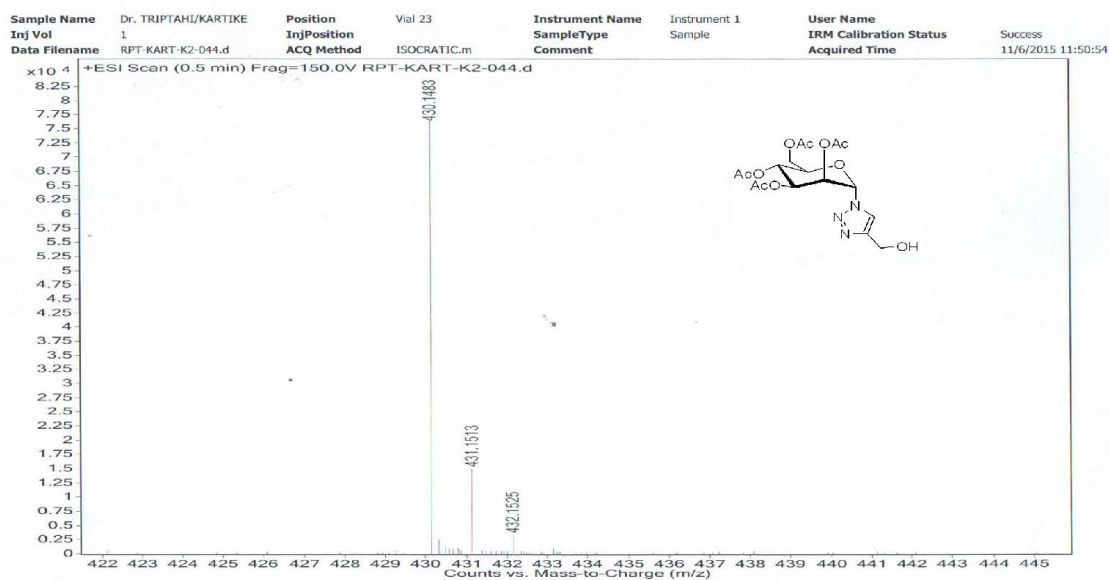
¹H NMR spectrum of compound 7a¹³C NMR spectrum of compound 7a

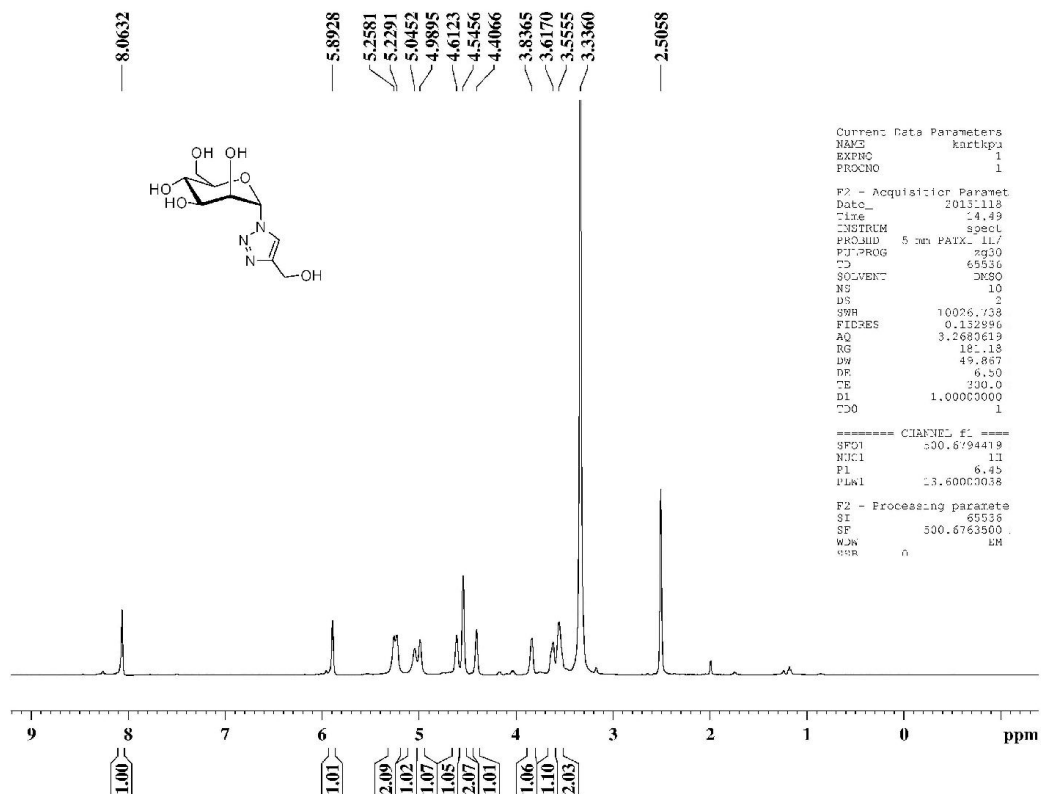
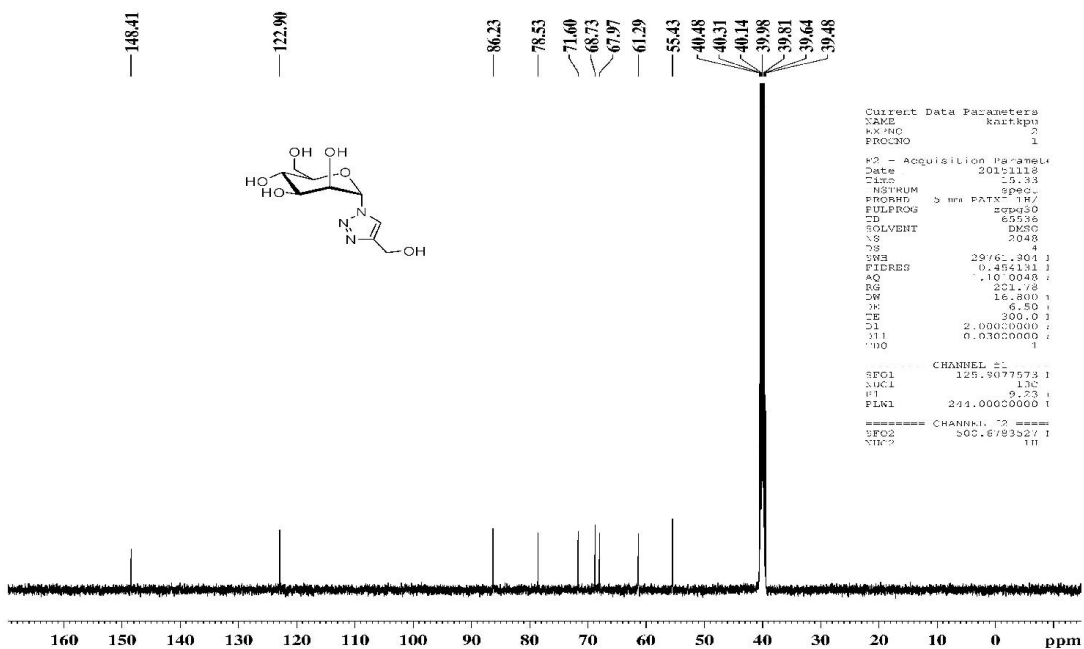
HRMS spectrum of compound **7a** ^1H NMR spectrum of compound **7a'**

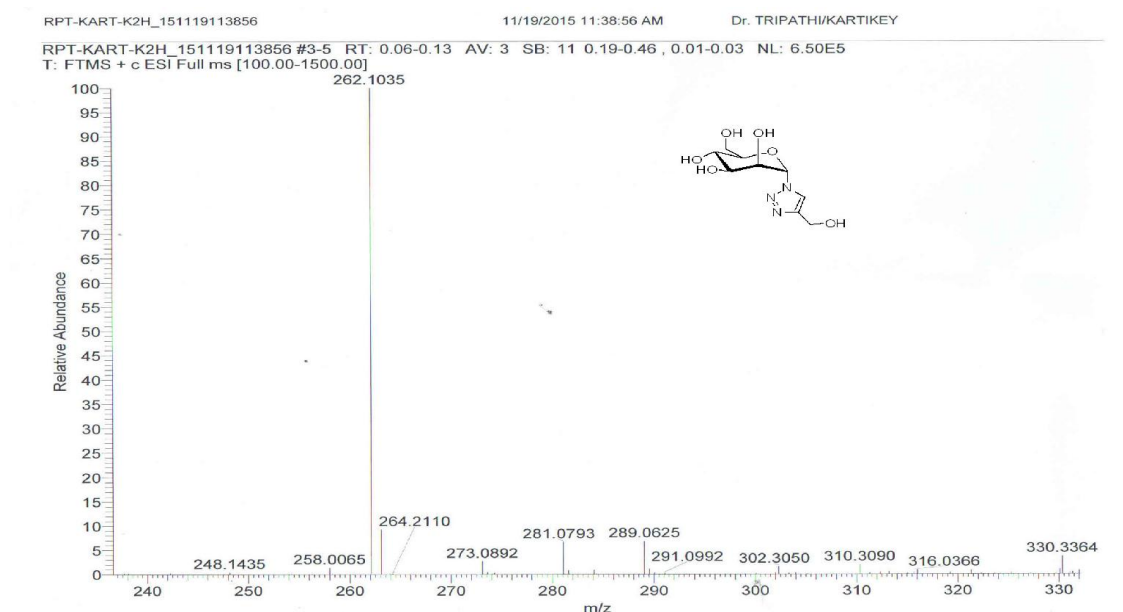
¹³C NMR spectrum of compound **7a'**HRMS spectrum of compound **7a'**

¹H NMR spectrum of compound 7f¹³C NMR spectrum of compound 7f

HRMS spectrum of compound **7f**¹H NMR spectrum of compound **7g**

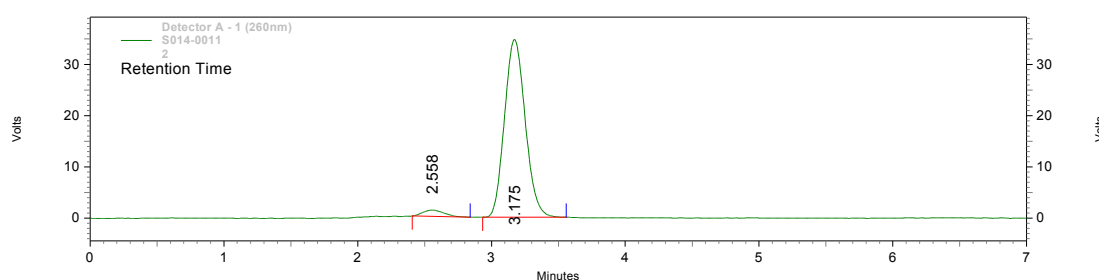
¹³C NMR spectrum of compound **7g**HRMS spectrum of compound **7g**

¹H NMR spectrum of compound **7g'**¹³C NMR spectrum of compound **7g'**

HRMS spectrum of compound **7g'**

HPLC Purity Determination Assay

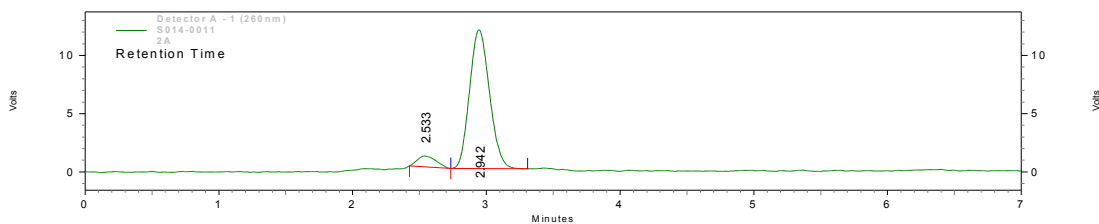
The purity of the compounds was determined by using reversed phase HPLC method on a Discovery HS C-18 (Supelco, Bellefonte, USA) column (5 μm , 100 x 4.6 mm id) preceded with a C-18 guard column (5 μm , 20 x 4.0 mm, id) packed with the same material under isocratic condition at ambient temperature using mobile phase [acetonitrile:ammonium acetate buffer (10 mM, pH 3.5), 70:30, %v/v] at a flow rate of 0.9 mL/min. The HPLC system consisted of a pump (LC-10AT VP with FCV-10AL VP), degasser (DGu-14A) and auto-injector (SIL-HTc, fixed with a 100 μl loop) (Shimadzu, Japan). Eluents were monitored at 260 nm with UV-Vis multiple wavelength detector and chromatograms were integrated using Class-VP (version 6.12 SP5) software (Shimadzu, Japan).

HPLC chromatogram of compound **2**

Detector A - 1 (260nm)

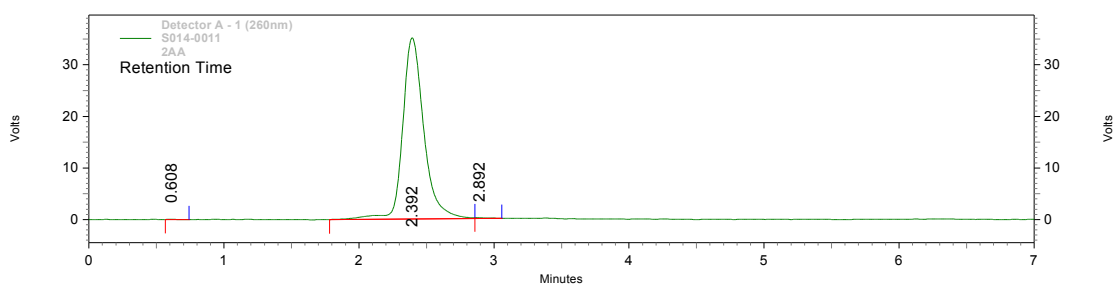
Pk #	Retention Time	Area	Area %	Height	Height %
1	2.558	13762	3.549	1253	3.487
2	3.175	373988	96.451	34680	96.513

Totals		387750	100.000	35933	100.000
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HPLC chromatogram of compound **2a****Detector A - 1 (260nm)**

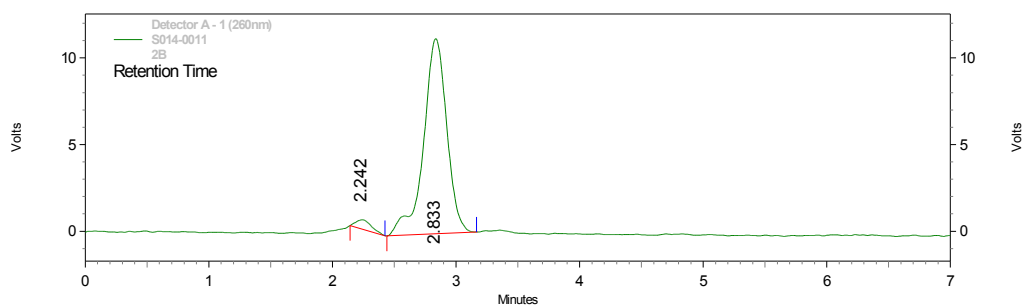
Pk #	Retention Time	Area	Area %	Height	Height %
1	2.533	3576	2.924	508	4.143
2	2.942	118738	97.076	11754	95.857

Totals		122314	100.000	12262	100.000
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HPLC chromatogram of compound **2a'****Detector A - 1 (260nm)**

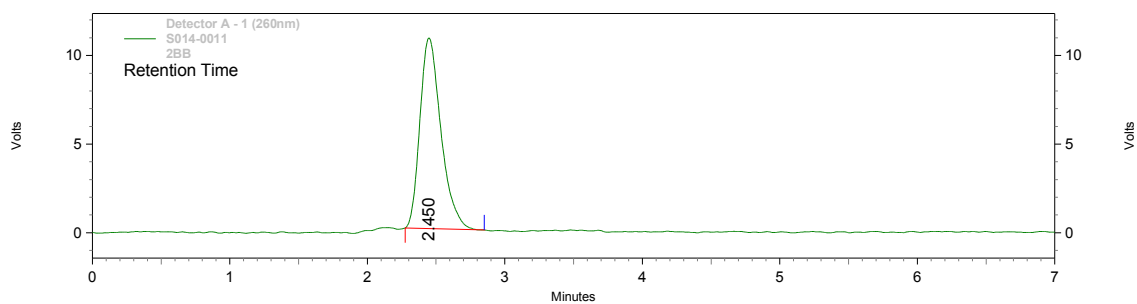
Pk #	Retention Time	Area	Area %	Height	Height %
1	0.608	256	0.067	46	0.130
2	2.392	381332	99.671	35105	99.450
3	2.892	1001	0.262	148	0.419

Totals		382589	100.000	35299	100.000
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HPLC chromatogram of compound **2b****Detector A - 1 (260nm)**

Pk #	Retention Time	Area	Area %	Height	Height %
1	2.242	4781	3.144	545	4.615
2	2.833	147278	96.856	11264	95.385

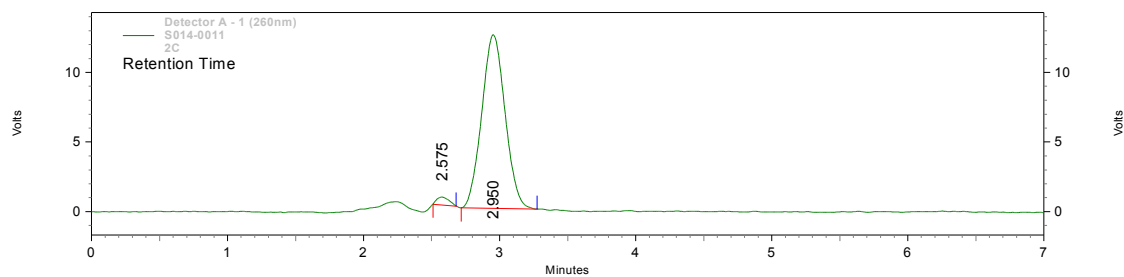
Totals		152059	100.000	11809	100.000
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HPLC chromatogram of compound **2b'****Detector A - 1 (260nm)**

Pk #	Retention Time	Area	Area %	Height	Height %
1	2.450	114346	100.000	10753	100.000

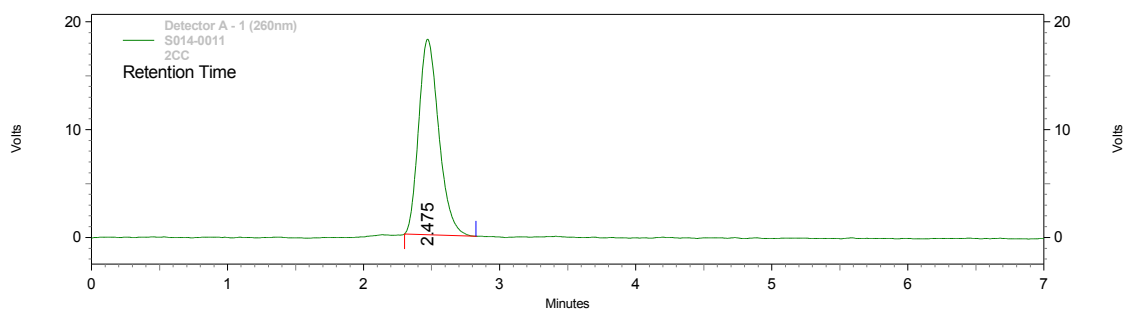
Totals		114346	100.000	10753	100.000
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HPLC chromatogram of compound **2c**


Detector A - 1 (260nm)

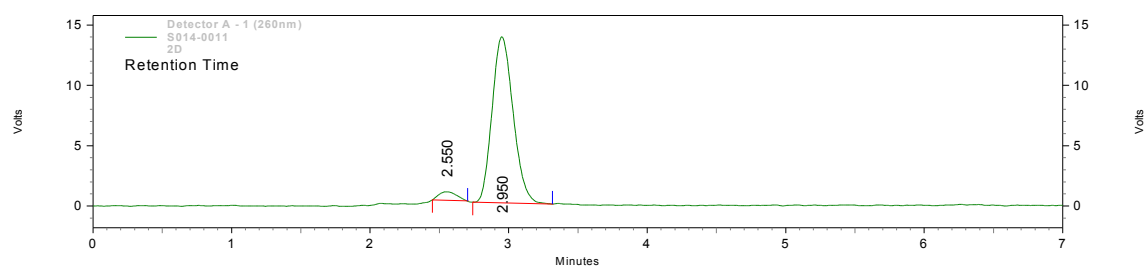
Pk #	Retention Time	Area	Area %	Height	Height %
1	2.575	3577	2.350	567	4.349
2	2.950	148618	97.650	12471	95.651

Totals		152195	100.000	13038	100.000

HPLC chromatogram of compound 2c'

Detector A - 1 (260nm)

Pk #	Retention Time	Area	Area %	Height	Height %
1	2.475	186721	100.000	18113	100.000

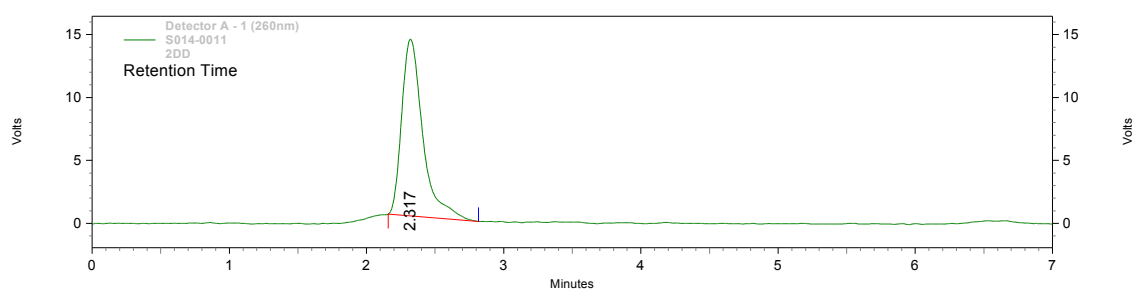
Totals		186721	100.000	18113	100.000

HPLC chromatogram of compound 2d


Detector A - 1 (260nm)

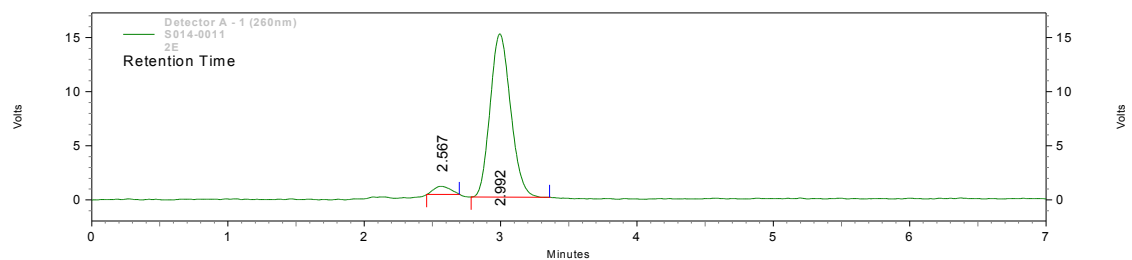
Pk #	Retention Time	Area	Area %	Height	Height %
1	2.550	6326	4.169	705	4.867
2	2.950	145421	95.831	13781	95.133

Totals		151747	100.000	14486	100.000

HPLC chromatogram of compound **2d'****Detector A - 1 (260nm)**

Pk #	Retention Time	Area	Area %	Height	Height %
1	2.317	150094	100.000	14035	100.000

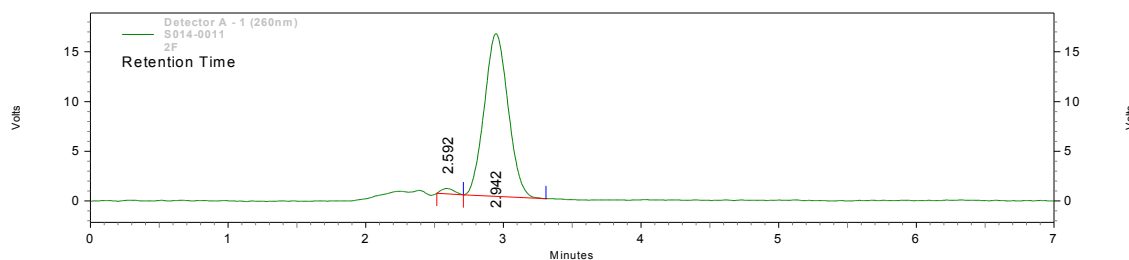
Totals		150094	100.000	14035	100.000

HPLC chromatogram of compound **2e****Detector A - 1 (260nm)**

Pk #	Retention Time	Area	Area %	Height	Height %
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1	2.567	6386	3.882	755	4.760
2	2.992	158114	96.118	15105	95.240

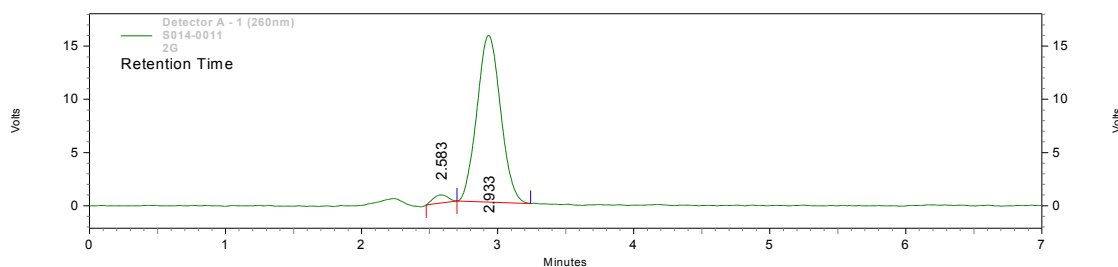
Totals		164500	100.000	15860	100.000
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HPLC chromatogram of compound **2f**

Detector A - 1 (260nm)

Pk #	Retention Time	Area	Area %	Height	Height %
1	2.592	3417	1.713	531	3.147
2	2.942	196044	98.287	16340	96.853

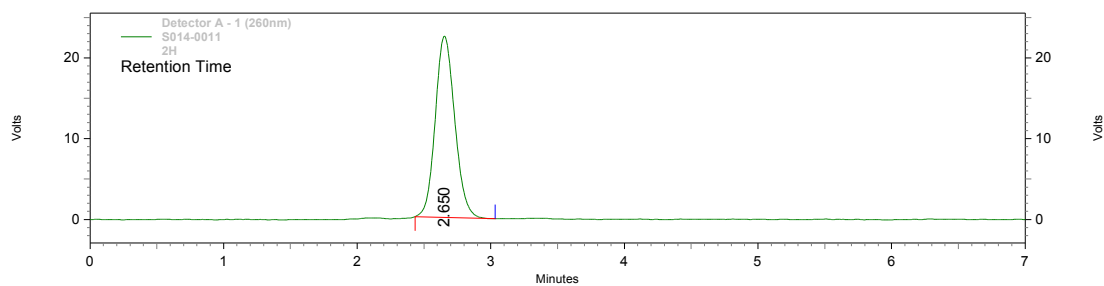
Totals		199461	100.000	16871	100.000
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HPLC chromatogram of compound **2g**

Detector A - 1 (260nm)

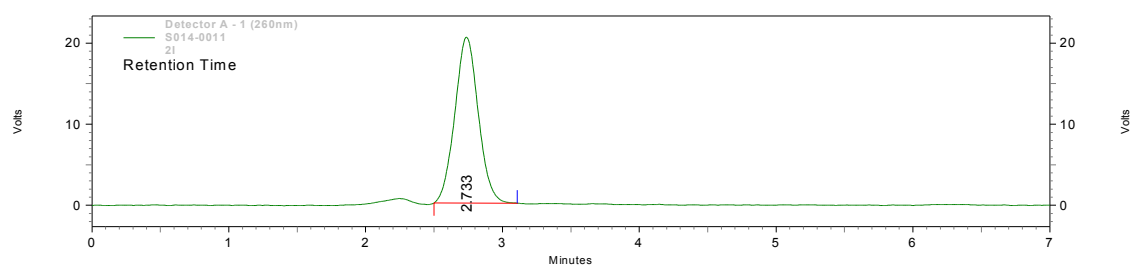
Pk #	Retention Time	Area	Area %	Height	Height %
1	2.583	5892	3.021	779	4.724
2	2.933	189160	96.979	15711	95.276

Totals		195052	100.000	16490	100.000
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HPLC chromatogram of compound **2h****Detector A - 1 (260nm)**

Pk #	Retention Time	Area	Area %	Height	Height %
1	2.650	228136	100.000	22451	100.000

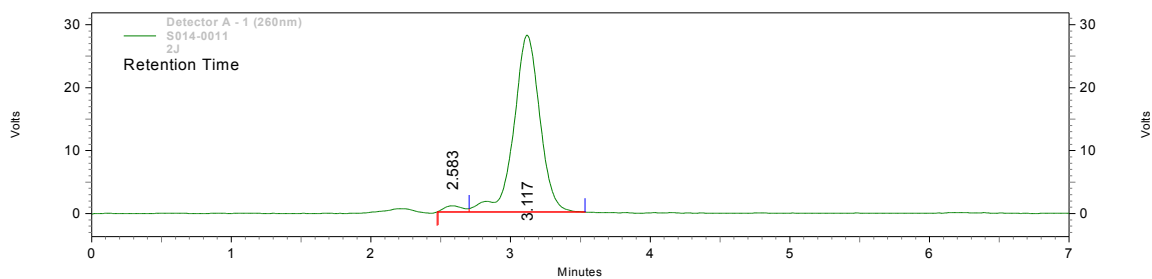
Totals		228136	100.000	22451	100.000
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HPLC chromatogram of compound **2i****Detector A - 1 (260nm)**

Pk #	Retention Time	Area	Area %	Height	Height %
1	2.733	243767	100.000	20451	100.000

Totals		243767	100.000	20451	100.000
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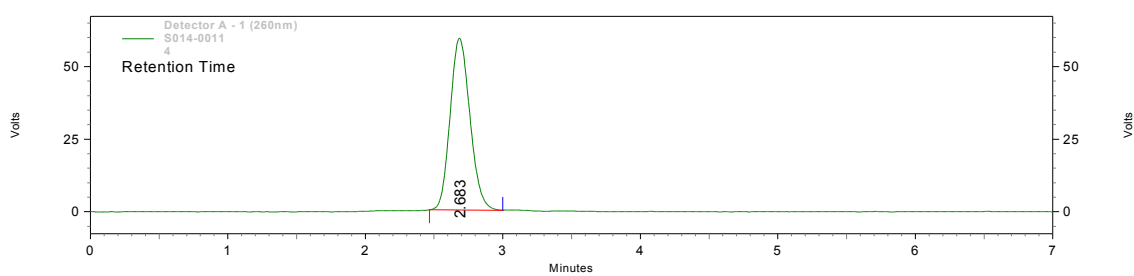
HPLC chromatogram of compound **2j**



Detector A - 1 (260nm)					
Pk #	Retention Time	Area	Area %	Height	Height %
1	2.583	8971	2.340	983	3.380
2	3.117	374454	97.660	28102	96.620

Totals		383425	100.000	29085	100.000
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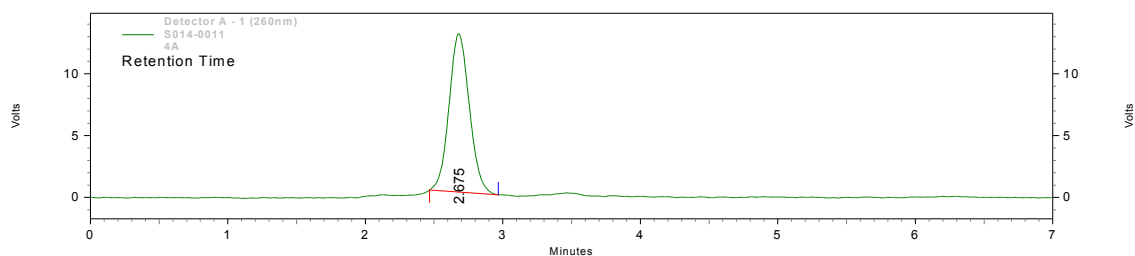
HPLC chromatogram of compound 4



Detector A - 1 (260nm)					
Pk #	Retention Time	Area	Area %	Height	Height %
1	2.683	591966	100.000	59251	100.000

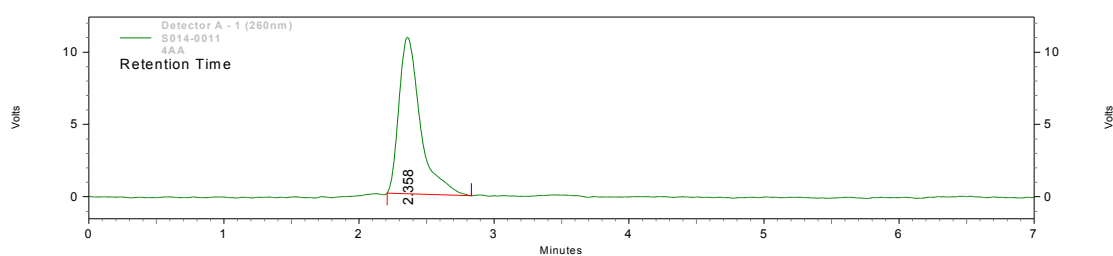
Totals		591966	100.000	59251	100.000
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HPLC chromatogram of compound 4a


Detector A - 1 (260nm)

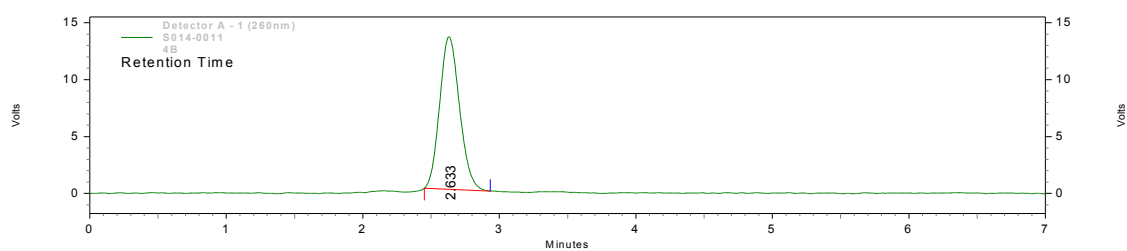
Pk #	Retention Time	Area	Area %	Height	Height %
1	2.675	131474	100.000	12781	100.000

Totals		131474	100.000	12781	100.000
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HPLC chromatogram of compound 4a'

Detector A - 1 (260nm)

Pk #	Retention Time	Area	Area %	Height	Height %
1	2.358	118706	100.000	10831	100.000

Totals		118706	100.000	10831	100.000
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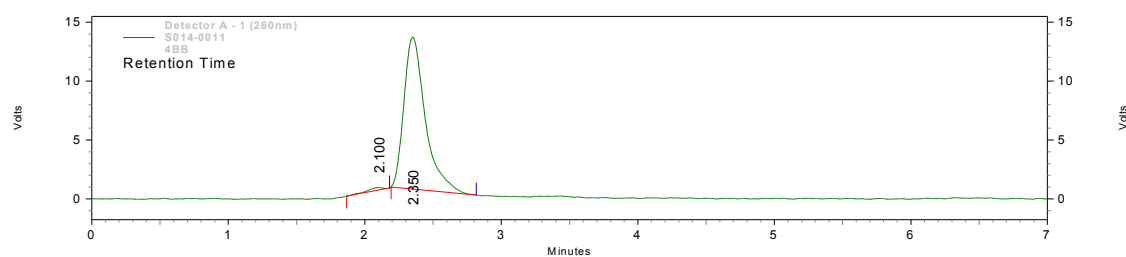
HPLC chromatogram of compound 4b


Detector A - 1 (260nm)

Pk #	Retention Time	Area	Area %	Height	Height %
1	2.633	135982	100.000	13414	100.000

Totals		135982	100.000	13414	100.000
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HPLC chromatogram of compound 4b'

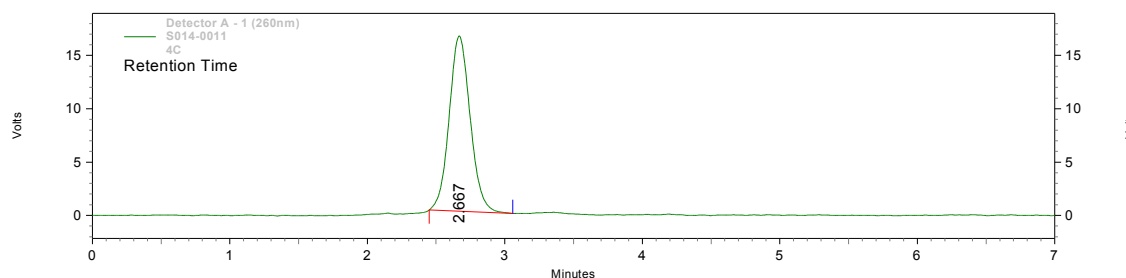


Detector A - 1 (260nm)

Pk #	Retention Time	Area	Area %	Height	Height %
1	2.100	1709	1.205	214	1.629
2	2.350	140173	98.795	12923	98.371

Totals		141882	100.000	13137	100.000
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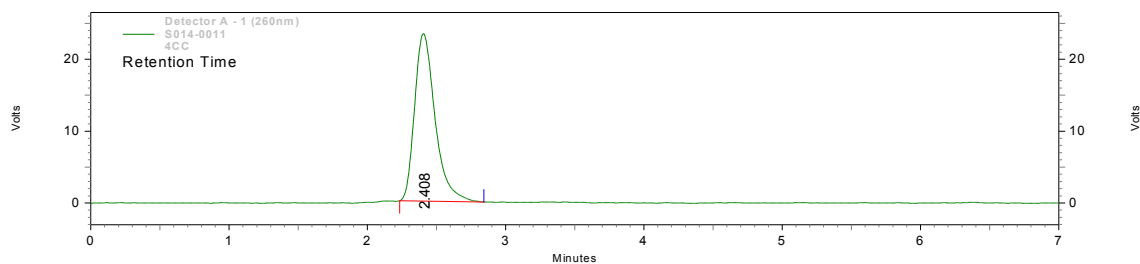
HPLC chromatogram of compound 4c



Detector A - 1 (260nm)

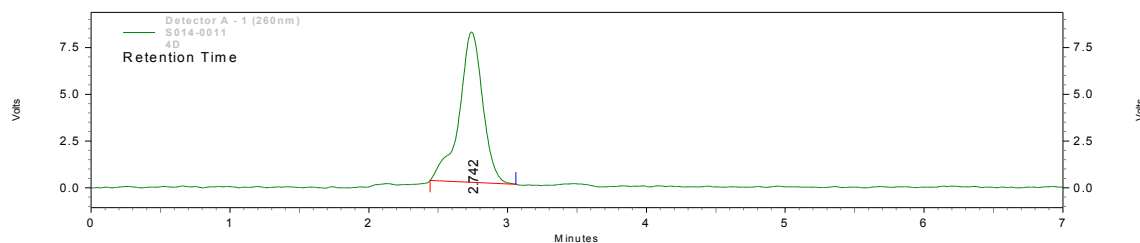
Pk #	Retention Time	Area	Area %	Height	Height %
1	2.667	174489	100.000	16442	100.000

Totals		174489	100.000	16442	100.000
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HPLC chromatogram of compound **4c'****Detector A - 1 (260nm)**

Pk #	Retention Time	Area	Area %	Height	Height %
1	2.408	239832	100.000	23299	100.000

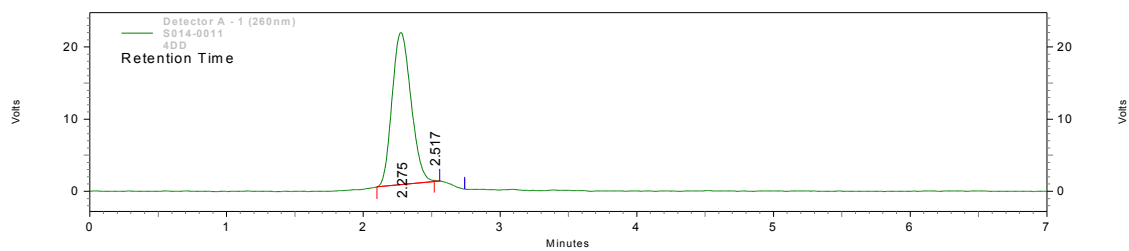
Totals		239832	100.000	23299	100.000
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HPLC chromatogram of compound **4d****Detector A - 1 (260nm)**

Pk #	Retention Time	Area	Area %	Height	Height %
1	2.742	94773	100.000	8041	100.000

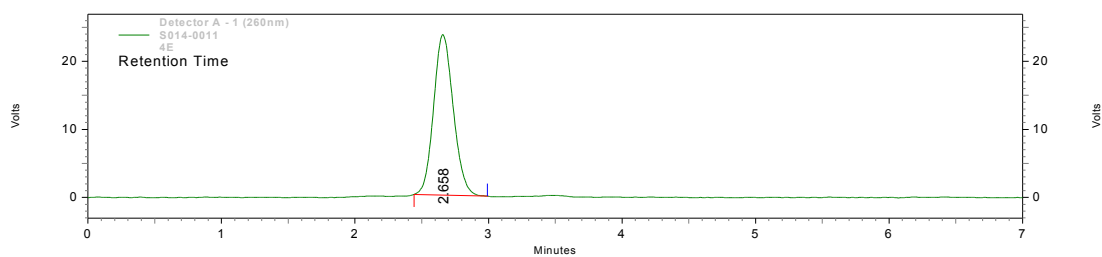
Totals		94773	100.000	8041	100.000
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HPLC chromatogram of compound **4d'**


Detector A - 1 (260nm)

Pk #	Retention Time	Area	Area %	Height	Height %
1	2.275	203233	96.277	21102	99.327
2	2.517	7860	3.723	143	0.673

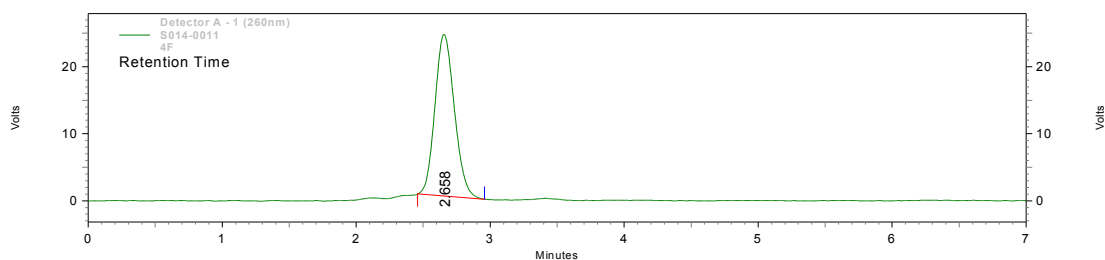
Totals		211093	100.000	21245	100.000
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HPLC chromatogram of compound 4e

Detector A - 1 (260nm)

Pk #	Retention Time	Area	Area %	Height	Height %
1	2.658	242093	100.000	23600	100.000

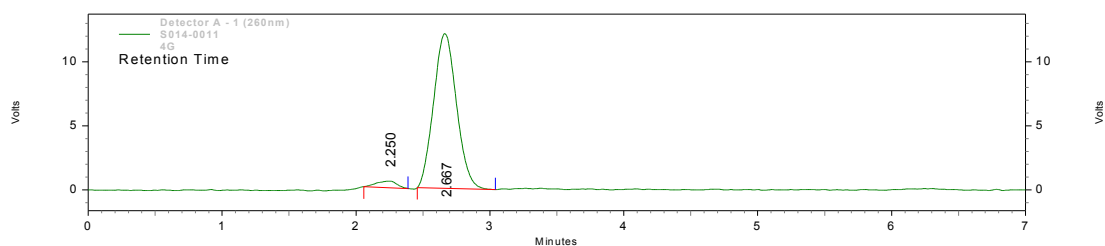
Totals		242093	100.000	23600	100.000
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HPLC chromatogram of compound 4f


Detector A - 1 (260nm)

Pk #	Retention Time	Area	Area %	Height	Height %
1	2.658	242992	100.000	24122	100.000

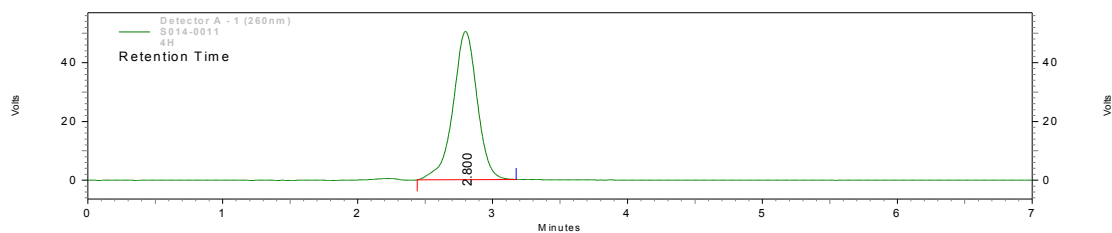
Totals		242992	100.000	24122	100.000
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HPLC chromatogram of compound 4g

Detector A - 1 (260nm)

Pk #	Retention Time	Area	Area %	Height	Height %
1	2.250	5363	3.599	515	4.091
2	2.667	143661	96.401	12074	95.909

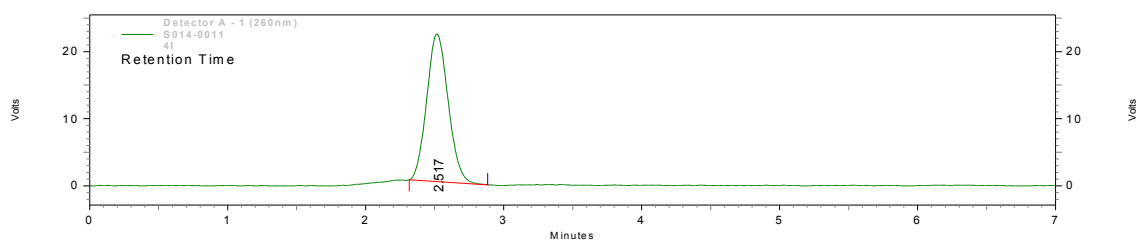
Totals		149024	100.000	12589	100.000
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HPLC chromatogram of compound 4h


Detector A - 1 (260nm)

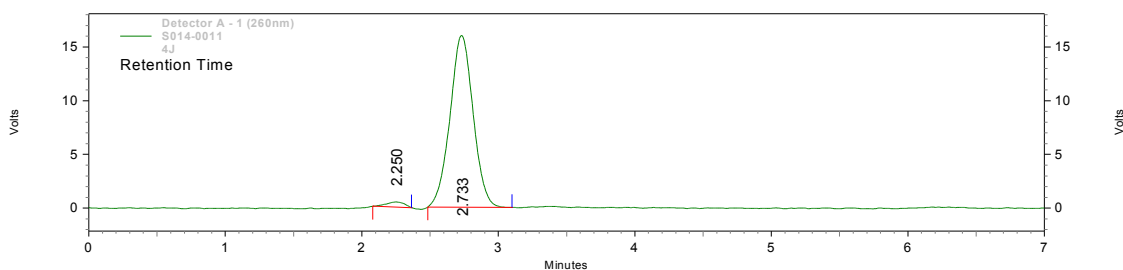
Pk #	Retention Time	Area	Area %	Height	Height %
1	2.800	625507	100.000	50418	100.000

Totals		625507	100.000	50418	100.000
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HPLC chromatogram of compound 4i

Detector A - 1 (260nm)

Pk #	Retention Time	Area	Area %	Height	Height %
1	2.517	237636	100.000	21976	100.000

Totals		237636	100.000	21976	100.000
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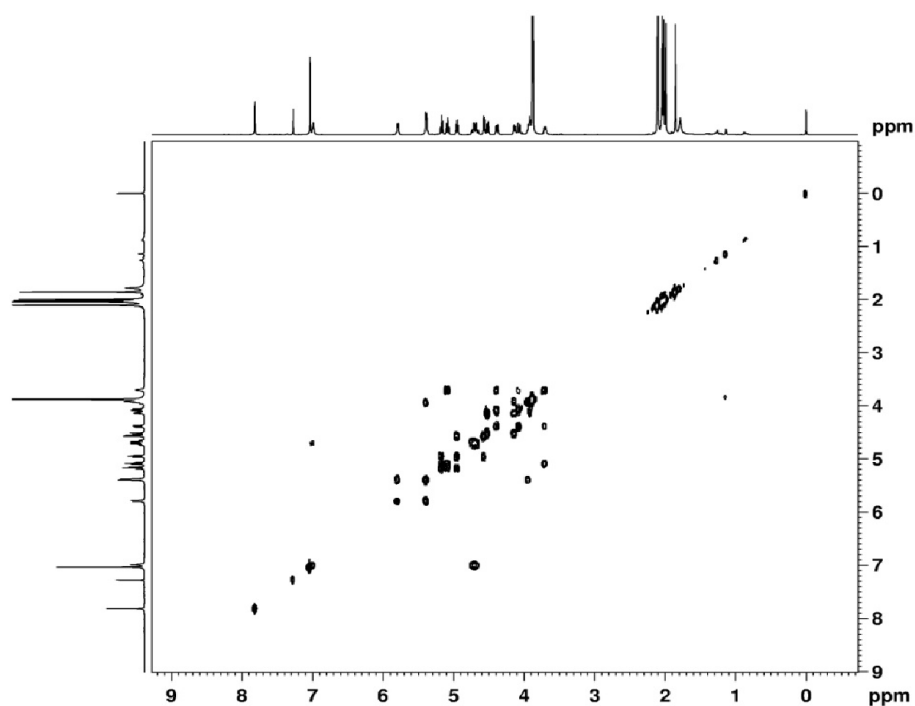
HPLC chromatogram of compound 4j

Detector A - 1 (260nm)

Pk #	Retention Time	Area	Area %	Height	Height %
1	2.250	4486	2.326	485	2.946

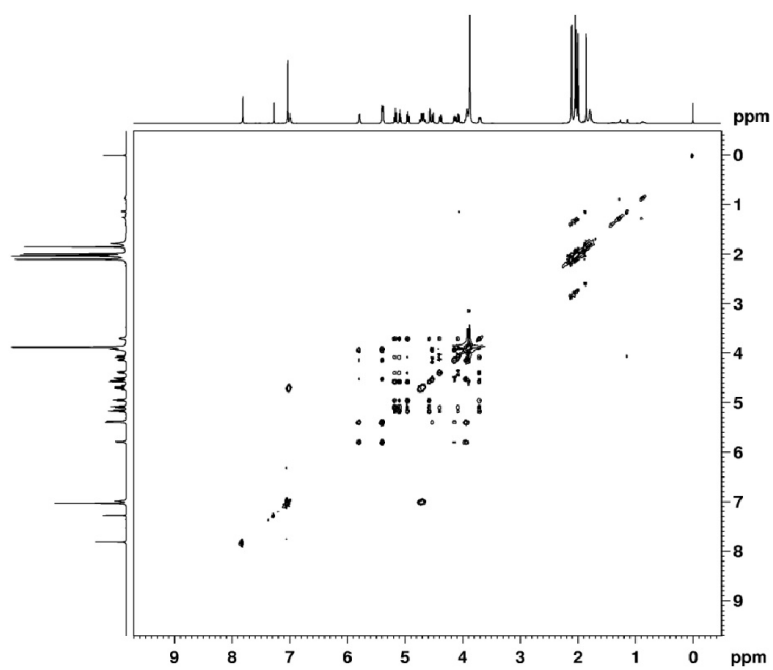
2 2.733 188350 97.674 15979 97.054

Totals		192836	100.000	16464	100.000
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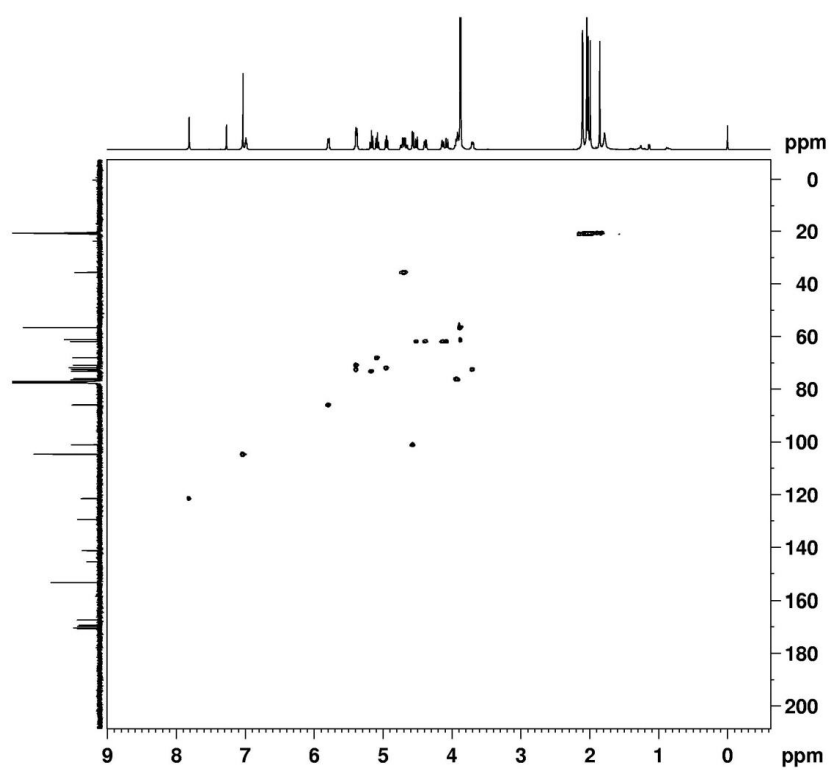
COSY spectrum of **4d**

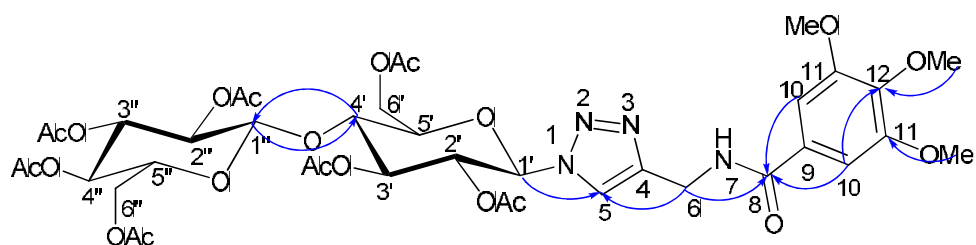
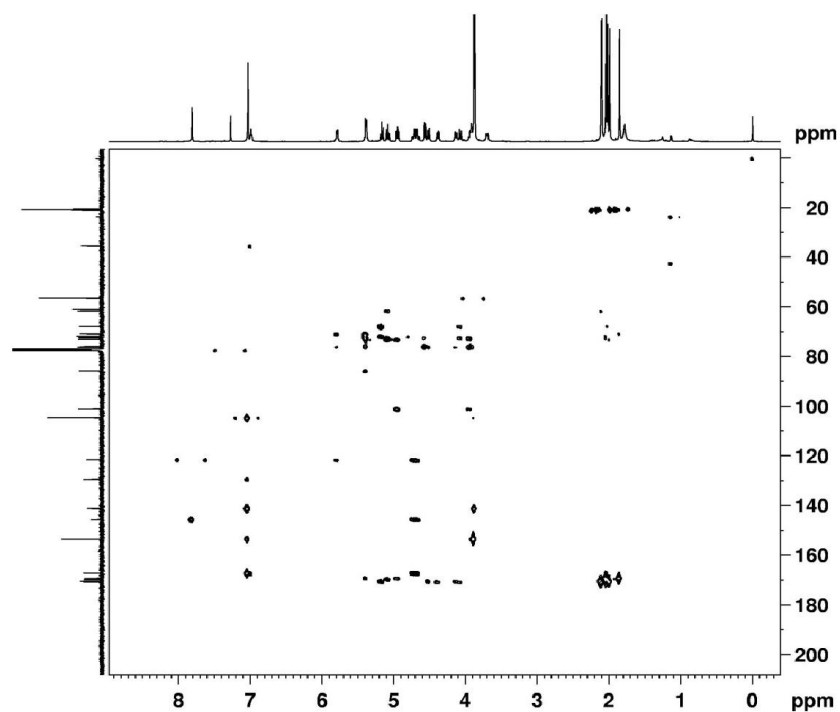


TOCSY spectrum of **4d**



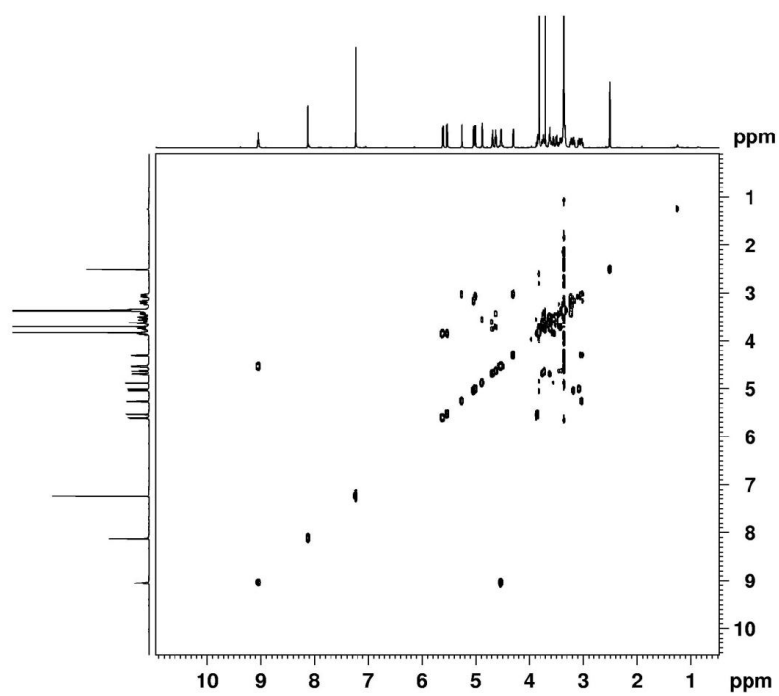
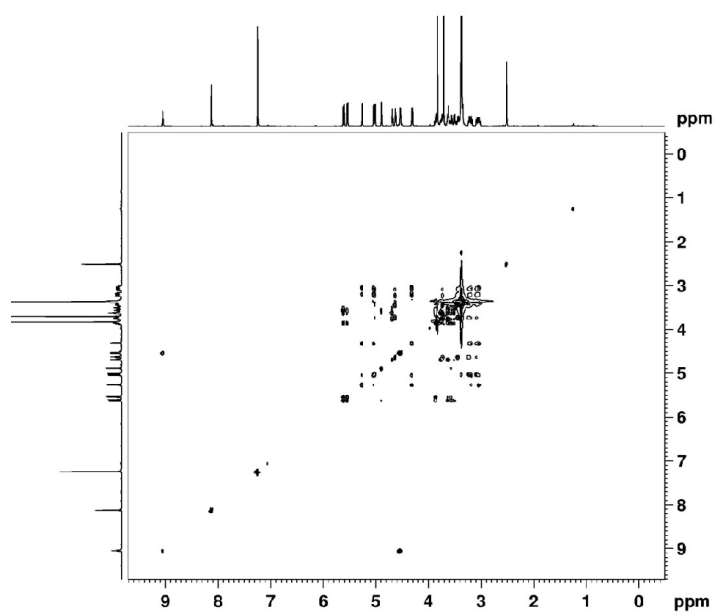
HSQC spectrum of 4d

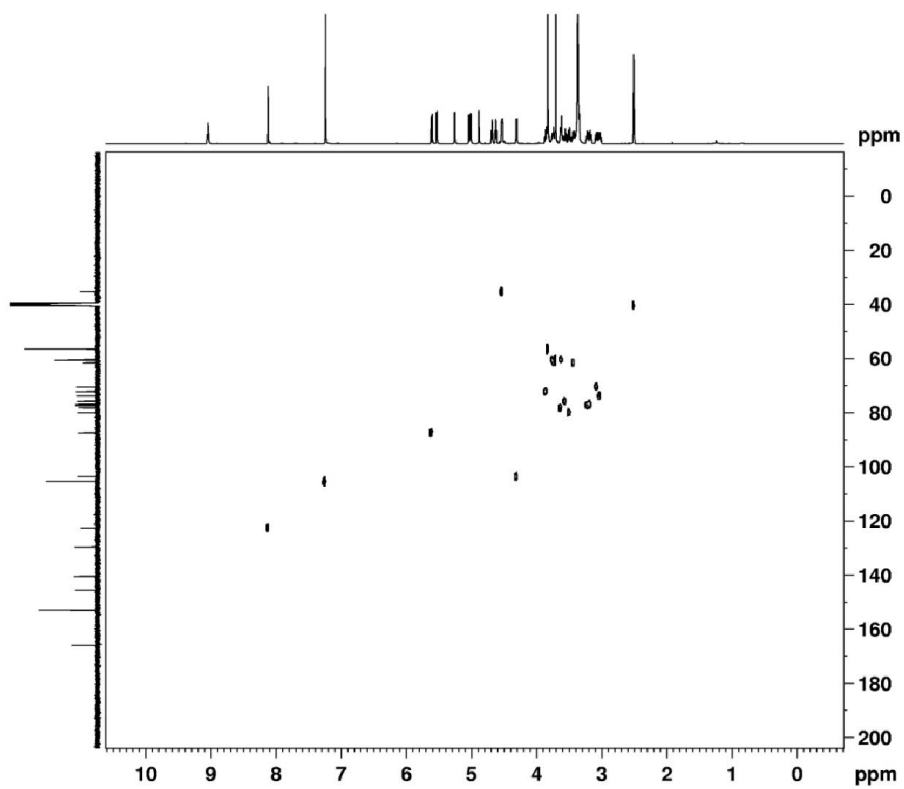


HMBC spectrum of **4d****Figure 1.** Important HMBC correlation of compound **4d**

Complete ^1H and ^{13}C signal assignments of compound **4d**

^1H NMR (500 MHz, CDCl_3) δ 7.81 (s, 1H, H-5), 7.03 (s, 2H, H-10), 6.98 (t, $J = 5.52$ Hz, 1H, H-7), 5.79 (dd, $J_1 = 6.15$ Hz, $J_2 = 2.60$ Hz, 1H, H-1'), 5.39 (m, 2H, H-2', H-3'), 5.14 (t, $J = 9.80$ Hz, 1H, H-3''), 5.08 (t, $J = 9.58$ Hz, 1H, H-4''), 4.96 (t, $J = 8.20$ Hz, 1H, H-2''), 4.70 (m, 2H, H-6), 4.55 (d, $J = 8.42$ Hz, 1H, H-1''), 4.50 (d, $J = 12.33$ Hz, 1H, H_b -6'), 4.36 (dd, $J_1 = 12.6$ Hz, $J_2 = 4.28$ Hz, 1H, H_b -6''), 4.12 (dd, $J_1 = 12.33$ Hz, $J_2 = 4.78$ Hz, 1H, H_a -6'), 4.07 (dd, $J_1 = 12.60$ Hz, $J_2 = 2.00$ Hz, 1H, H_a -6''), 3.93 (m, 2H, H-4', H-5'), 3.88 (s, 6H, OMe-11), 3.87 (s, 3H, OMe-12), 3.69 (m, 1H, H-5'), 2.10 (s, 3H, OAc-6'), 2.09 (s, 3H, OAc-6''), 2.04 (s, 6H, OAc-3', OAc-2''), 2.01 (s, 3H, OAc-4''), 1.99 (s, 3H, OAc-3''), 1.85 (s, 3H, OAc-2'); ^{13}C NMR (125 MHz) δ 170.6 (C-6''-C=O), 170.4 & 170.3 (C-6'-C=O & C-3''-C=O), 169.7 (C-4''-C=O), 169.5 (C-2''-C=O), 169.2 (2C, C-2'-C=O & C-3'-C=O), 167.3 (C-8), 153.4 (2C, C-11), 145.4 (C-4), 141.3 (C-12), 129.4 (C-9), 121.4 (C-5), 104.6 (2C, C-10), 101.0 (C-1'), 85.8 (C-1'), 76.2 & 75.9 (C-4' & C-5'), 73.0 (C-3'), 72.4 & 72.3 (C-3' & C-5''), 71.7 (C-2''), 70.8 (C-2'), 67.9 (C-4''), 61.7 (2C, C-6', C-6''), 61.1 (C-12-OMe), 56.5 (2C, C-11-OMe), 35.5 (C-6), [20.9, 20.8, 20.7, 20.6 & 20.4 (total 7C for 2', 3', 6', 2'', 3'', 4'' & 6''-OAc)]

COSY spectrum of **4d'**TOCSY spectrum of **4d'**

HSQC spectrum of **4d'**HMBC spectrum of **4d'**

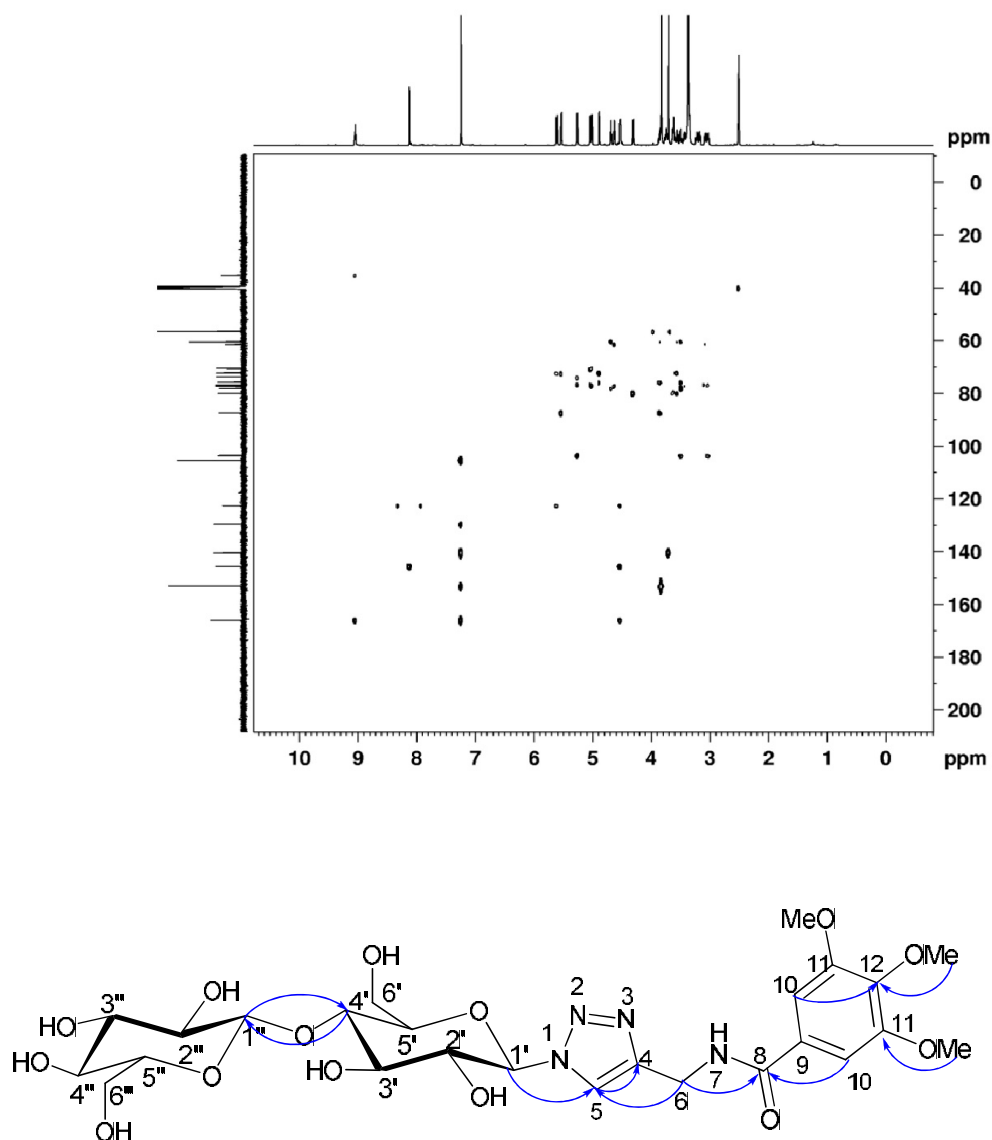


Figure 2. Important HMBC correlation of compound **4d'**

Complete ^1H and ^{13}C signal assignments of compound **4d'**

^1H NMR (500 MHz, CDCl_3) δ 9.04 (t, $J = 5.75$ Hz, 1H, H-7), 8.12 (s, 1H, H-5), 7.24 (s, 2H, H-10), 5.60 (d, $J = 9.55$ Hz, 1H, H-1'), 5.52 (d, $J = 6.15$ Hz, OH-2'), 5.25 (d, $J = 5.01$ Hz, OH-2''), 5.03 (d, $J = 5.10$ Hz, OH-3'), 5.00 (d, $J = 5.47$ Hz, OH-4'), 4.88 (d, $J = 2.30$ Hz, OH-3'), 4.68 (t, $J = 4.90$ Hz, OH-6'), 4.62 (t, $J = 5.36$ Hz, OH-6''), 4.53 (dd, $J_1 = 5.75$ Hz, $J_2 = 2.60$ Hz, 2H, H-6),

4.30 (d, $J = 7.80$ Hz, 1H, H-1''), 3.85 (m, 1H, H-2'), 3.82 (s, 6H, OMe-11), 3.74 (m, 1H, H_b-6'), 3.71 (m, 1H, H_b-6''), 3.70 (s, 3H, OMe-12), 3.63 (m, 1H, H-5'), 3.60 (m, 1H, H_a-6'), 3.55 (m, 1H, H-3'), 3.50 (t, $J = 7.80$ Hz, 1H, H-4'), 3.40 (m, 1H, H_a-6''), 3.23 (m, 1H, H-5''), 3.16 (m, 1H, H-3''), 3.07 (m, 1H, H-4''), 3.01 (m, 1H, H-2''); ¹³C NMR (125 MHz) δ 165.9 (C-8), 152.9 (2C, C-11), 145.5 (C-4), 140.4 (C-12), 129.5 (C-9), 122.5 (C-5), 105.3 (2C, C-10), 103.5 (C-1''), 87.3 (C-1'), 79.9 (C-4'), 78.1 (C-5'), 77.2 (C-5''), 76.8 (C-3''), 75.6 (C-3'), 73.7 (C-2''), 72.1 (C-2'), 70.4 (C-4'), 61.4 (C-6''), 60.4 (OMe-12), 60.3 (C-6'), 56.4 (2C, OMe-11), 35.2 (C-6).