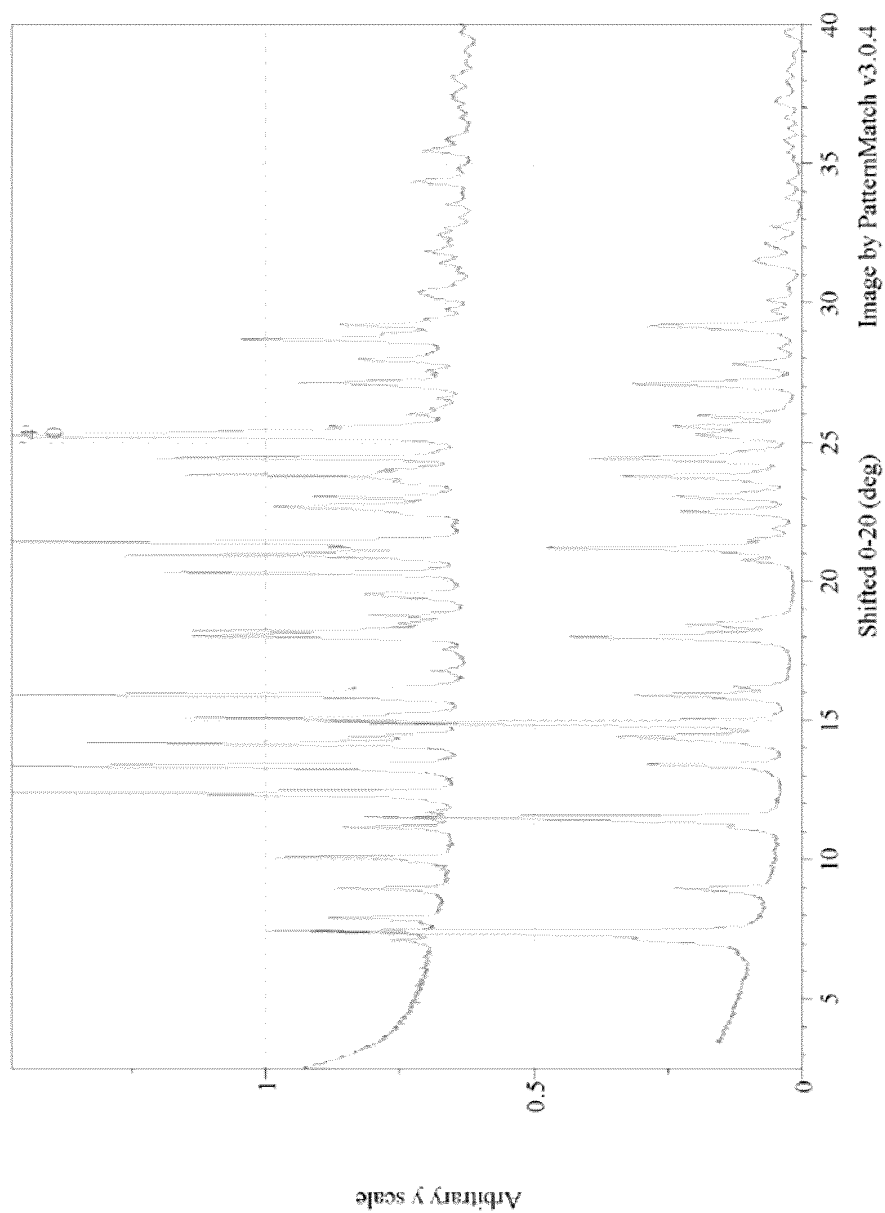
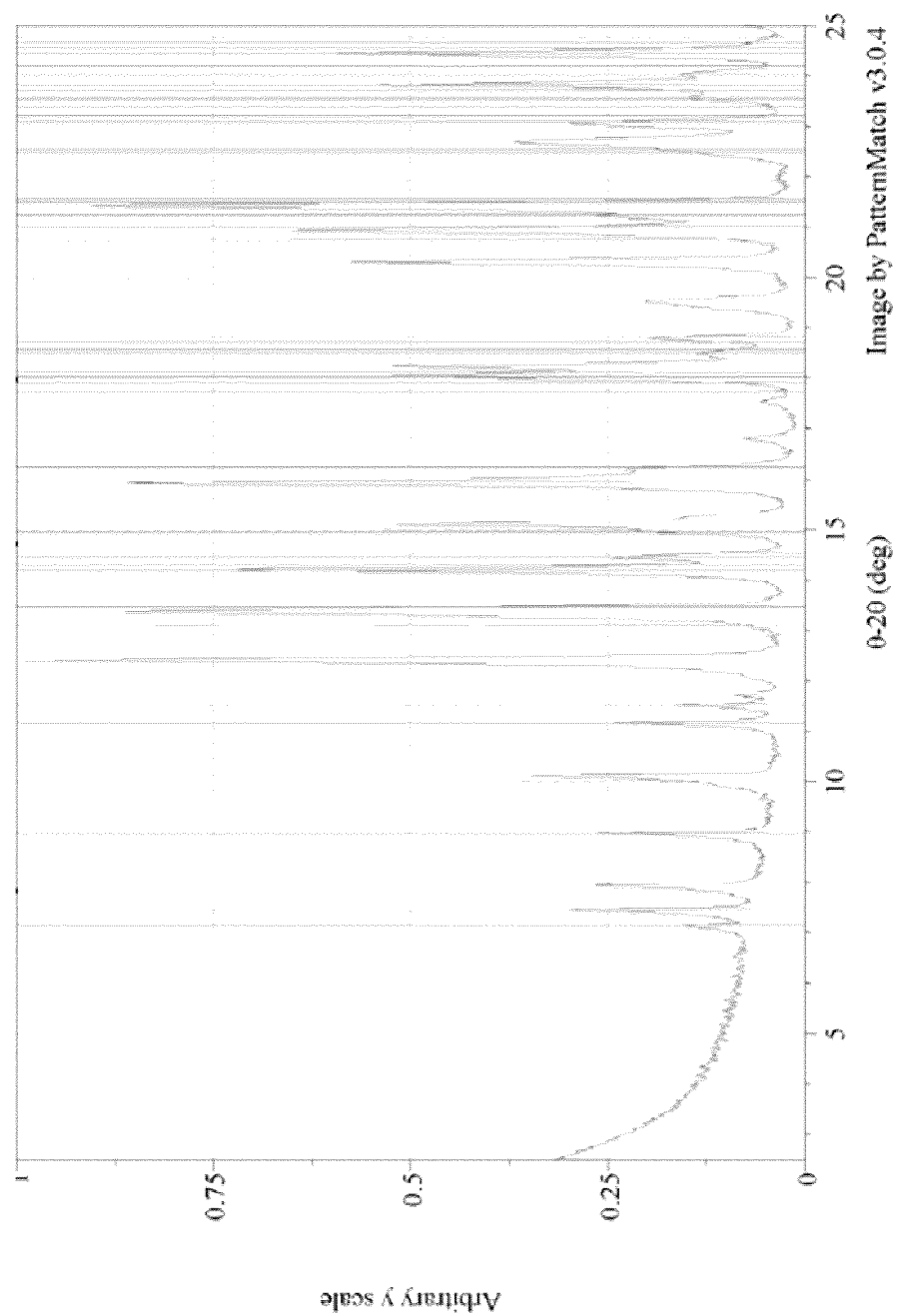


**Figure 1**

**Figure 2**

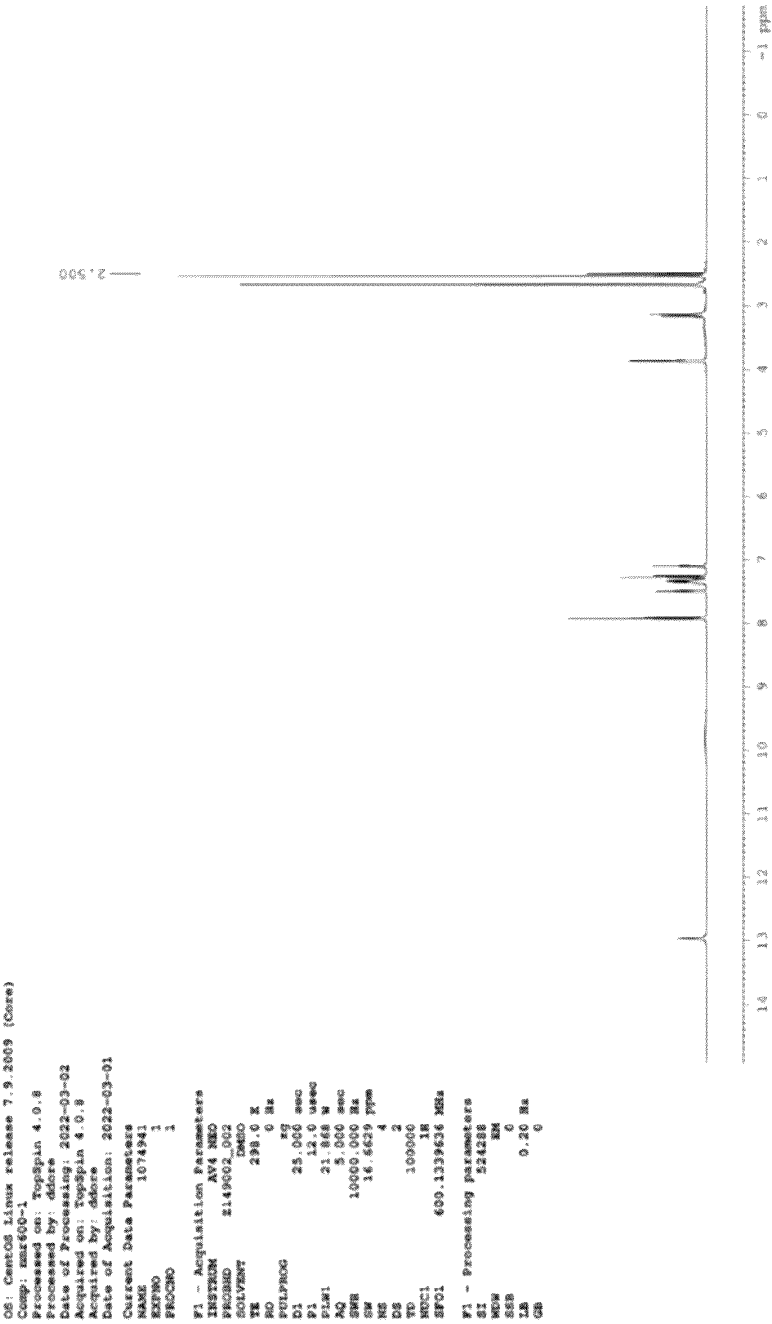


Figure 3

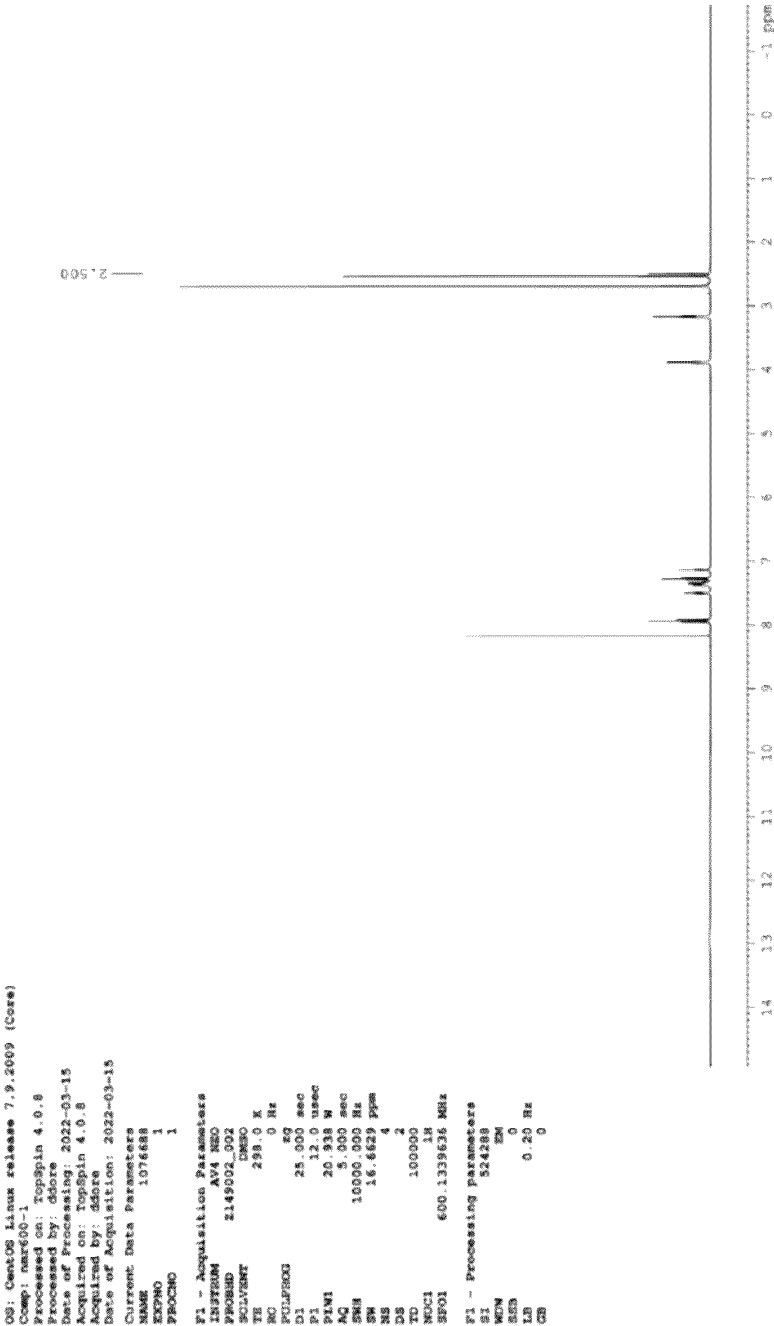
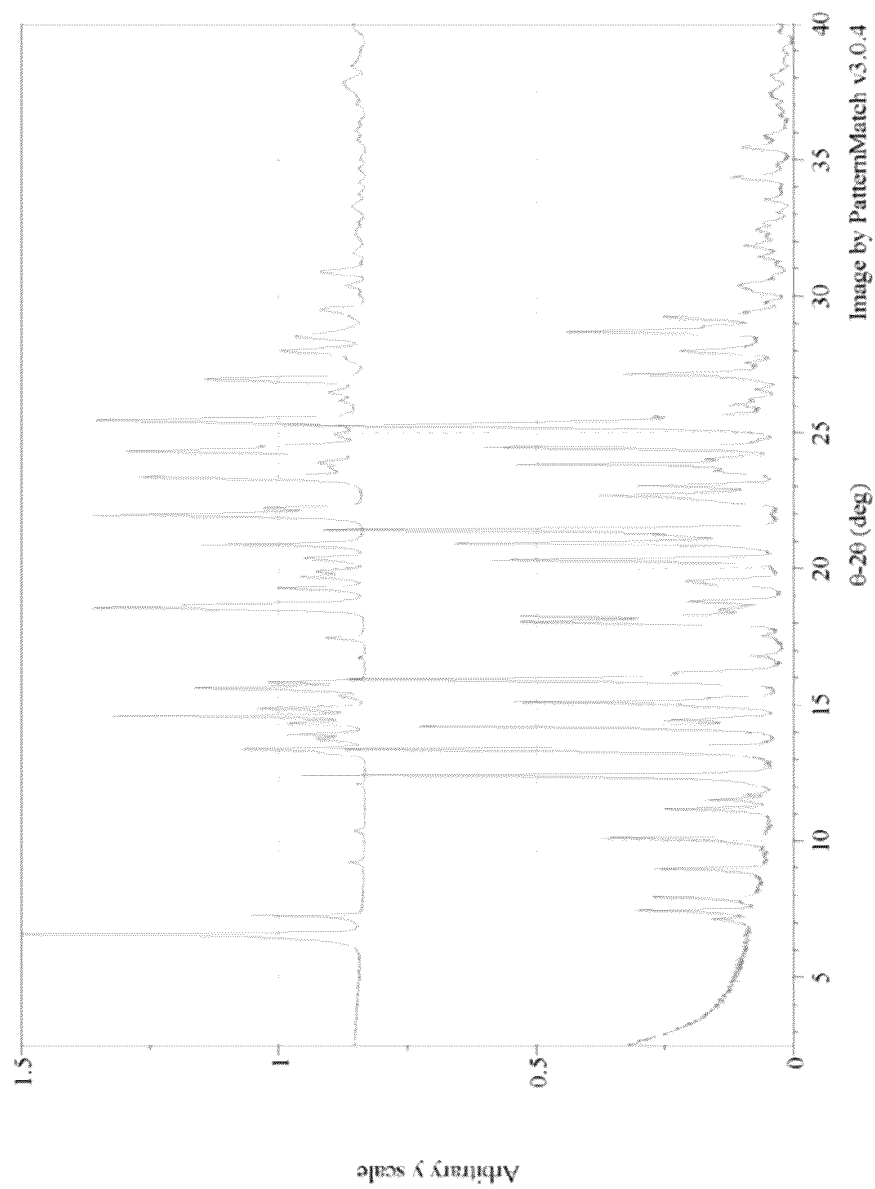


Figure 4

**Figure 5a**

Up : Compound 2
Down : Compound 1

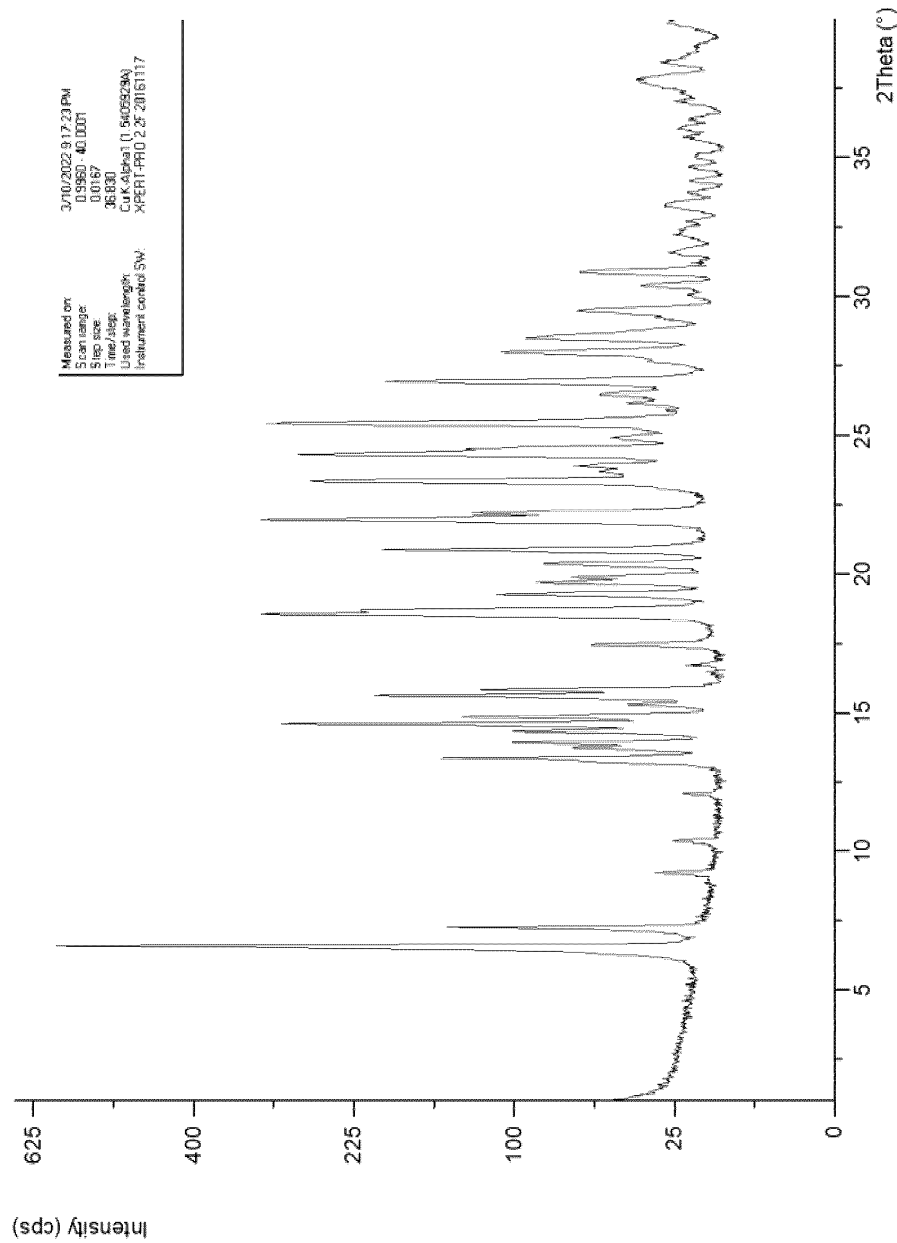
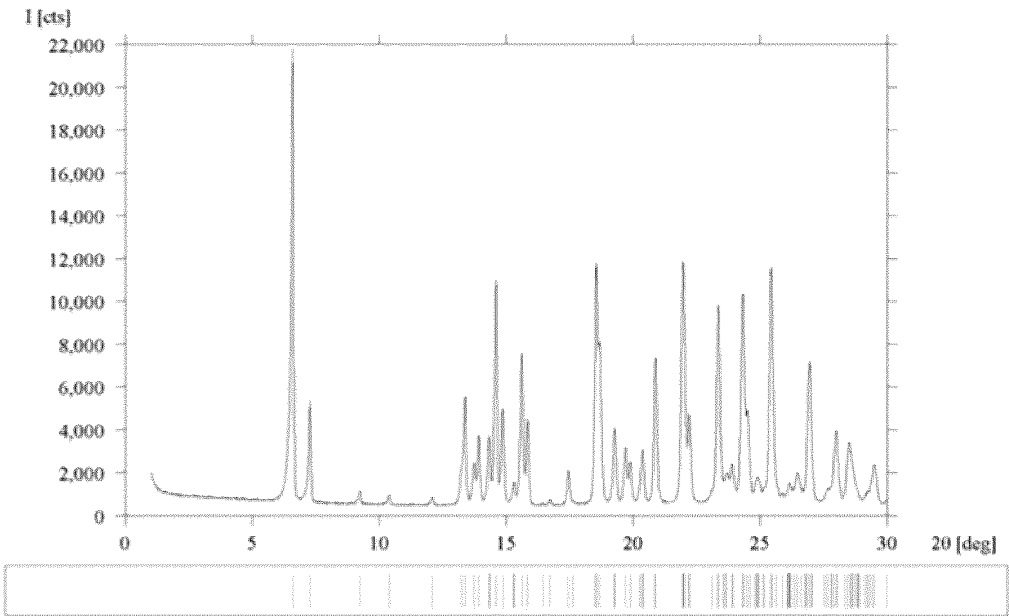


Figure 5b

Indexing results for XRPD file 1075870 collected with Cu-K α radiation.



Bravais Type	Primitive Monoclinic
a [Å]	12.217
b [Å]	7.610
c [Å]	26.945
α [deg]	90
β [deg]	96.66
γ [deg]	90
Volume [Å ³ /cell]	2,488.2
Chiral Contents?	Not Specified
Extinction Symbol	P 1 2 ₁ /c 1
Space Group(s)	P2 ₁ /c (14)
Source	Manual Input

Figure 6

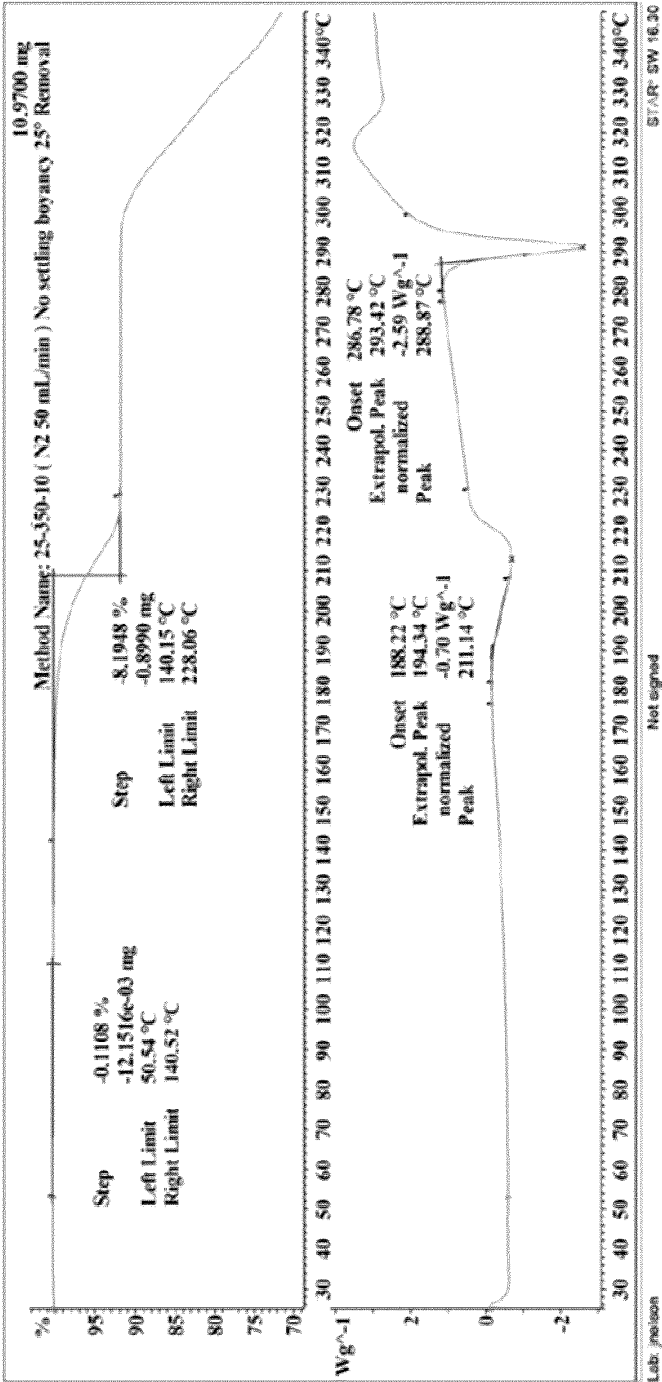


Figure 7

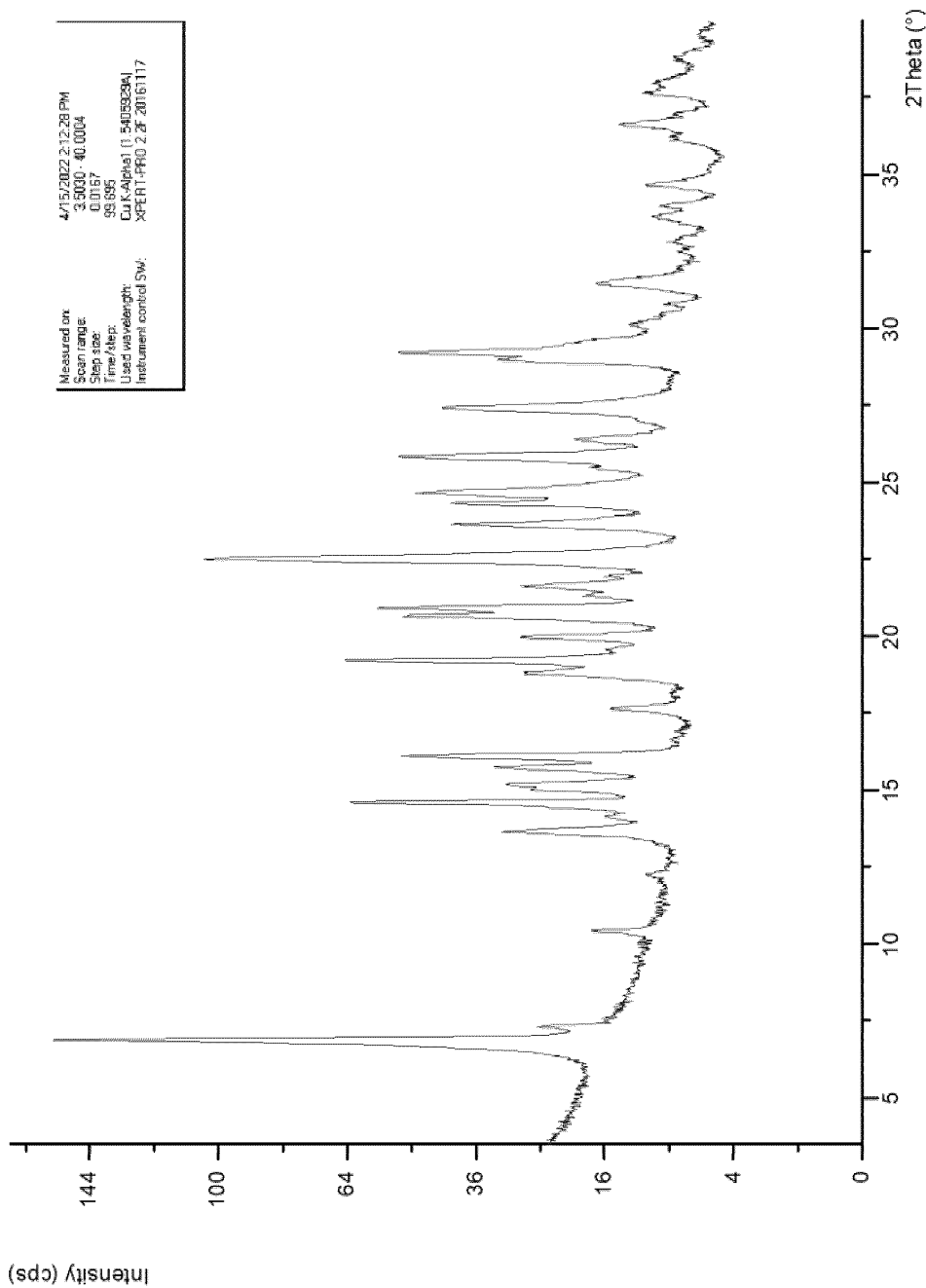


Figure 8

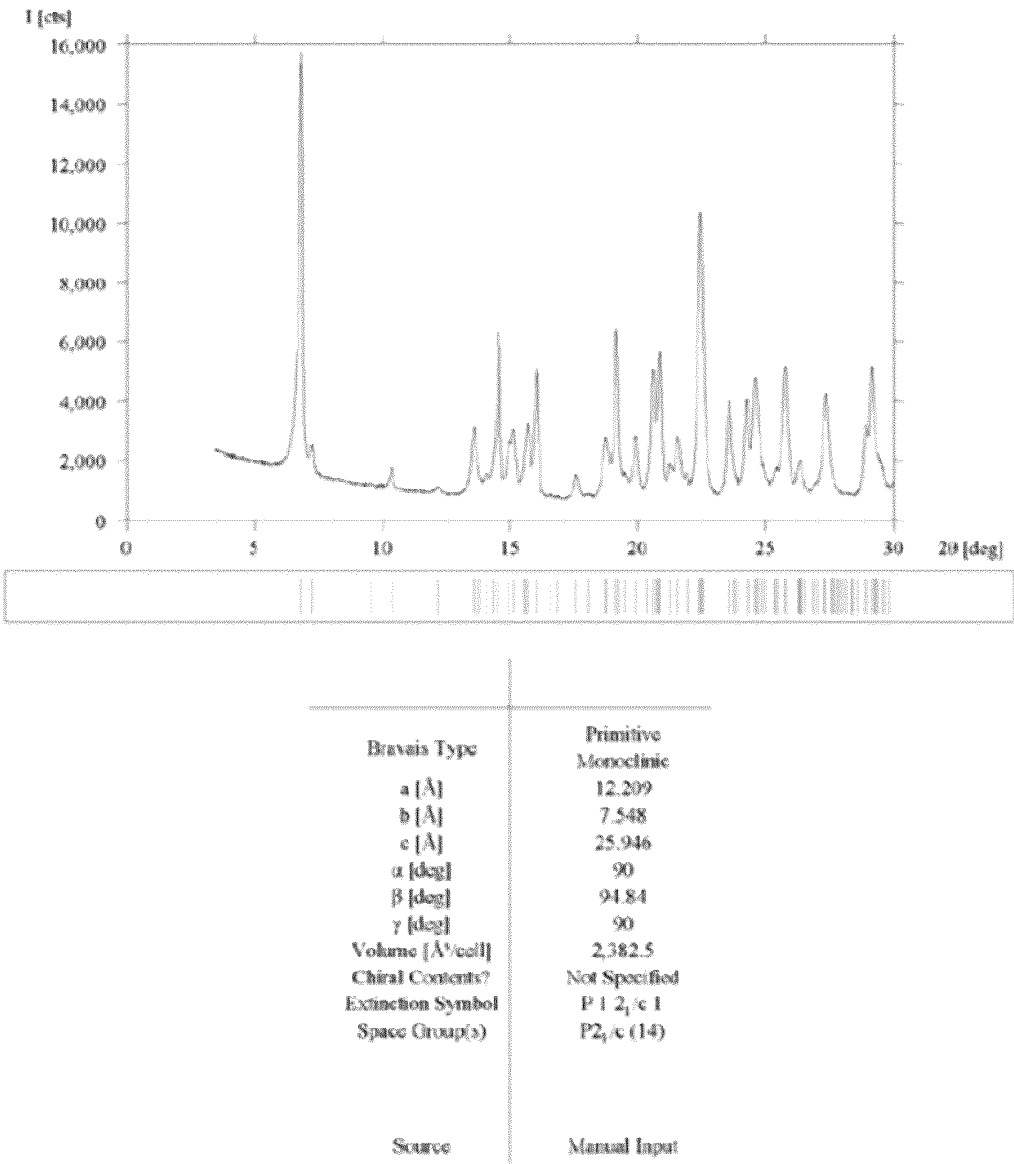
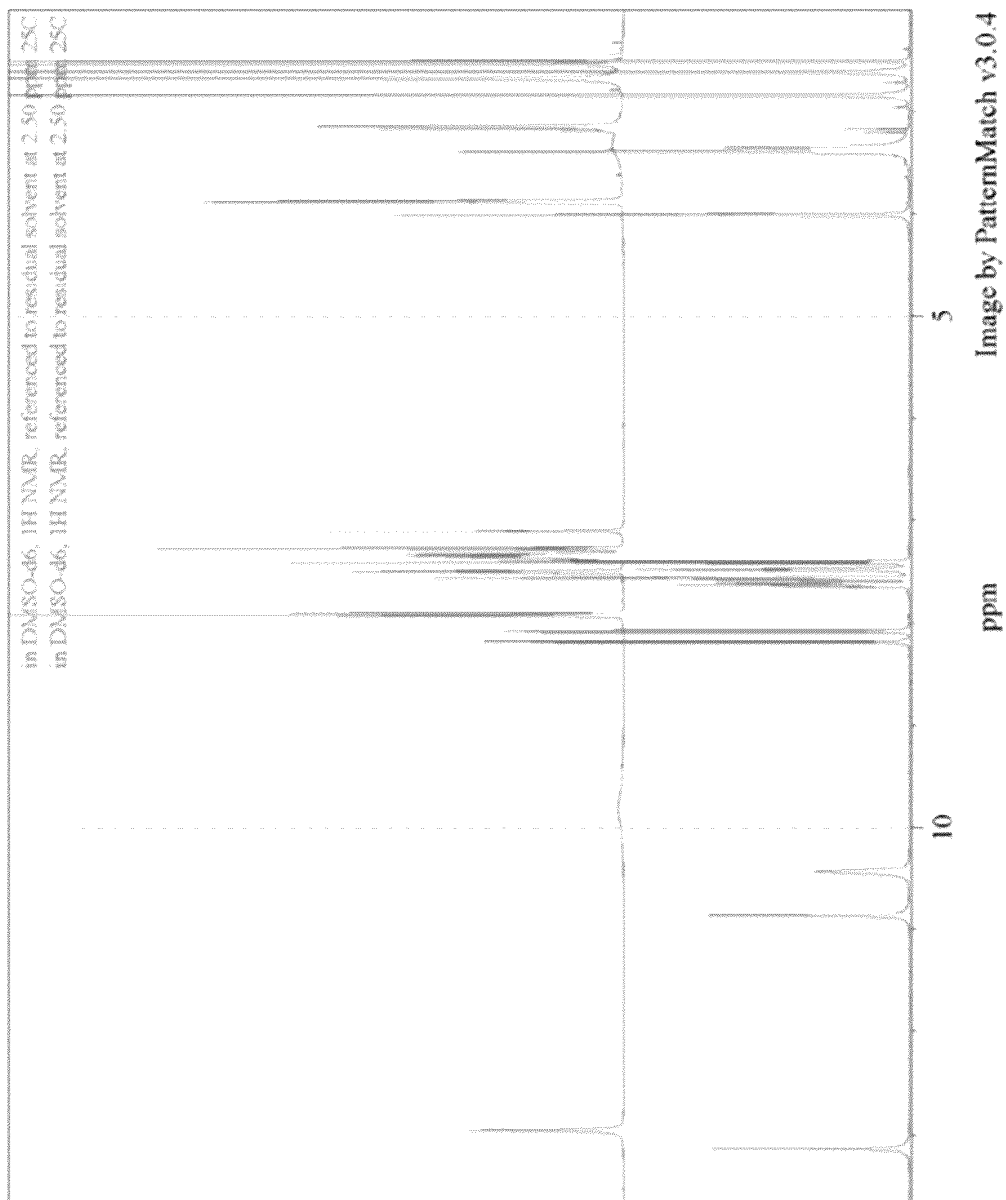


Figure 9

**Figure 10**

Up : Compound 1

Down : Compound 3 mono-hydrochloride salt unsolvated, Form A

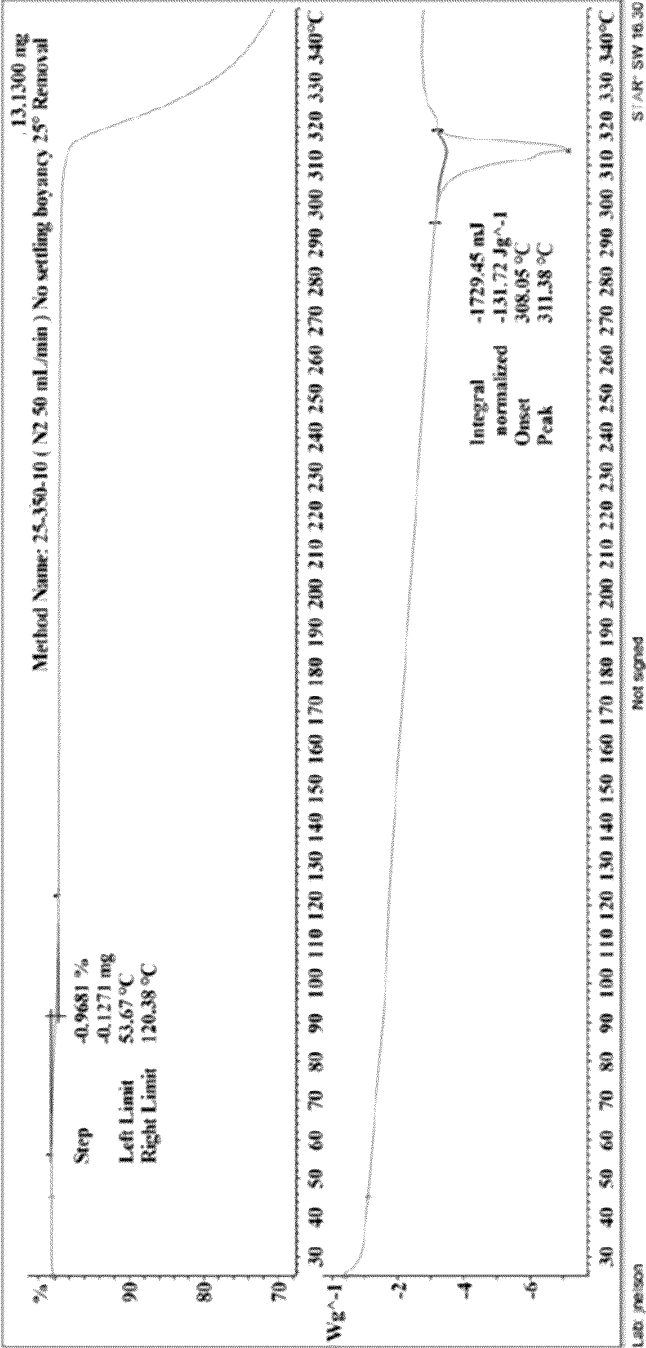
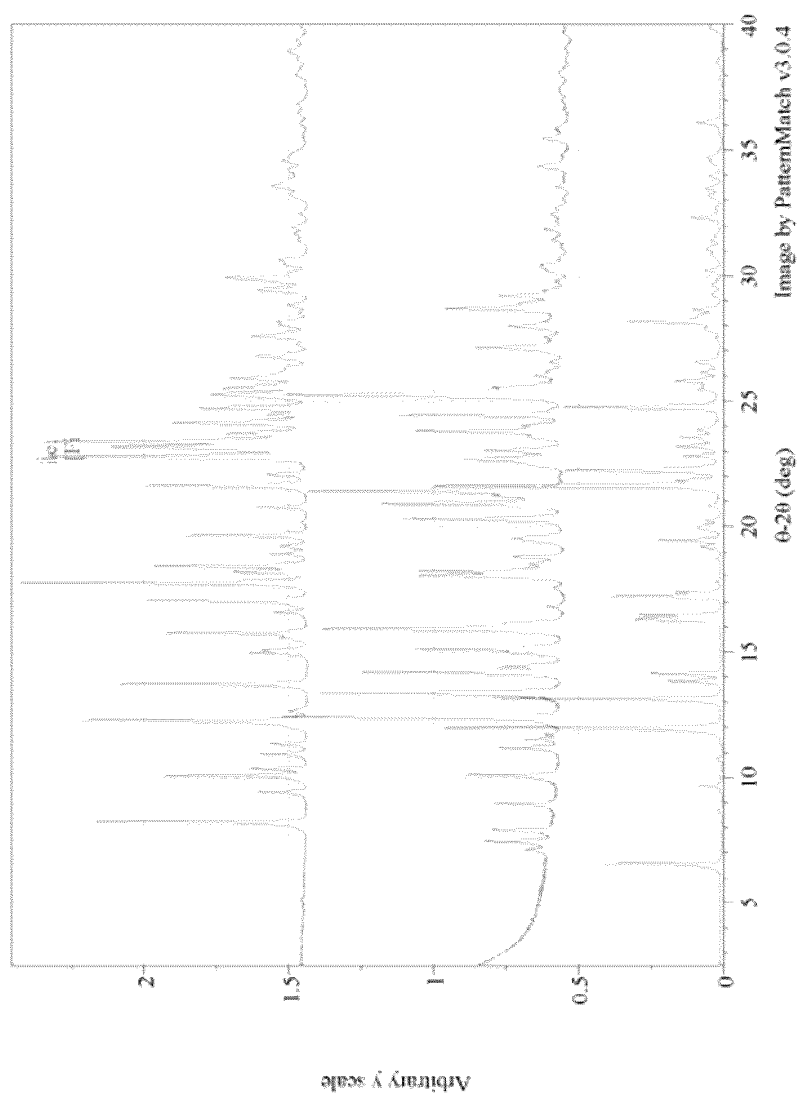


Figure 11

**Figure 12**

Up : Compound 4
Middle : Compound 1
Down : Compound 1 (Form C)

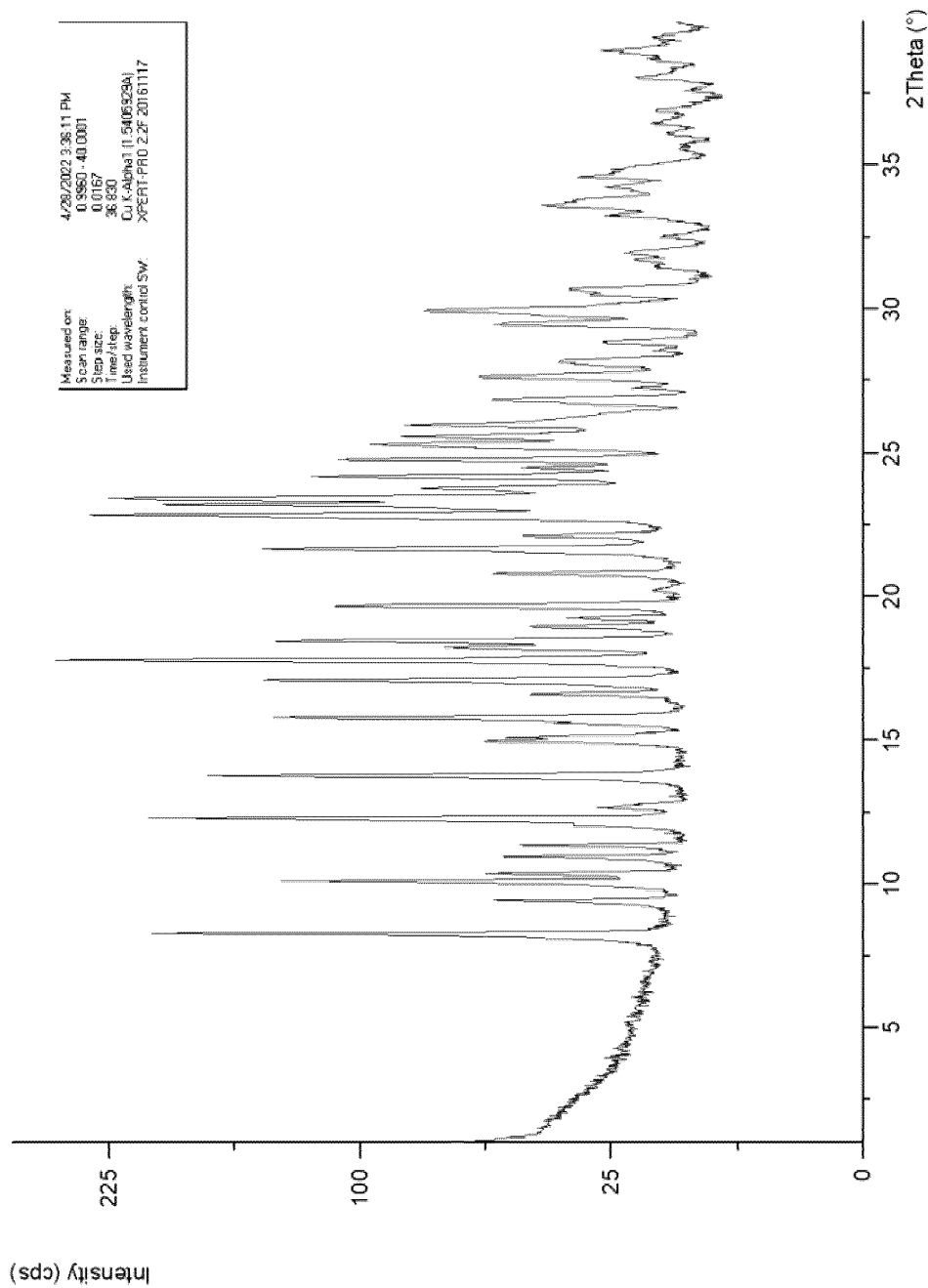
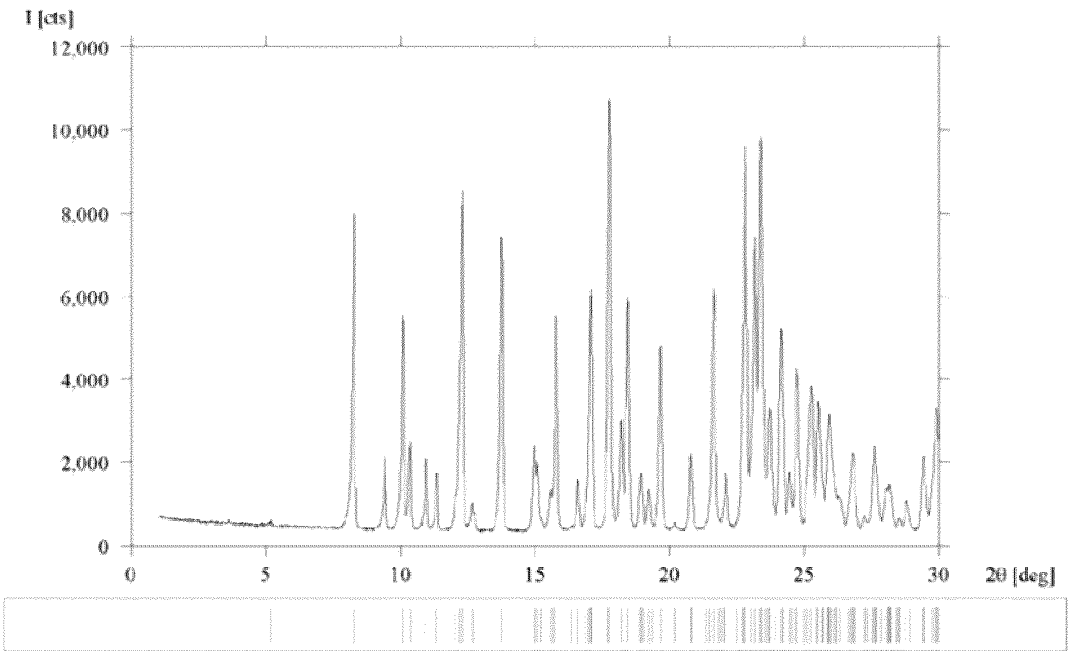


Figure 13



Bravais Type	Triclinic
a [Å]	8.554
b [Å]	11.355
c [Å]	17.120
α [deg]	85.72
β [deg]	89.54
γ [deg]	70.80
Volume [Å ³ /cell]	1,565.9
Chiral Contents?	Not Specified
Extinction Symbol	P -
Space Group(s)	P1 (1), P $\bar{1}$ (2)
Source	Triads® Algorithm

Figure 14

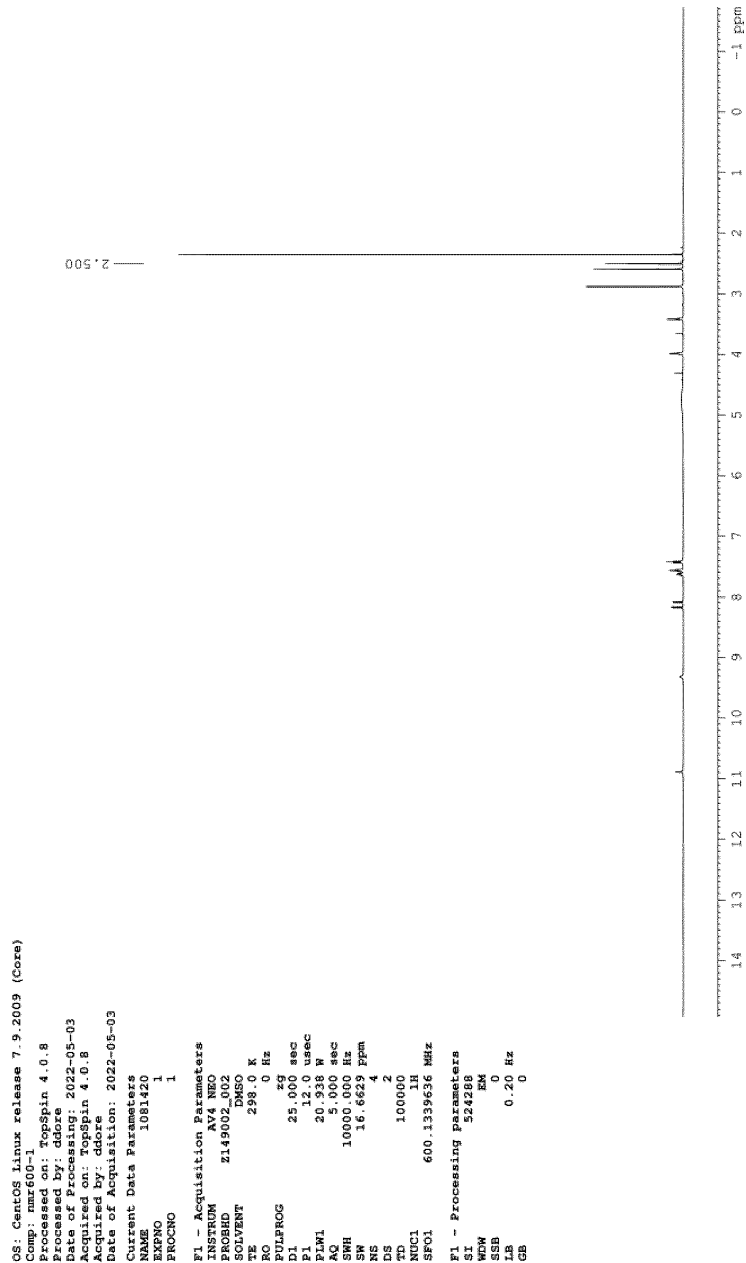


Figure 15

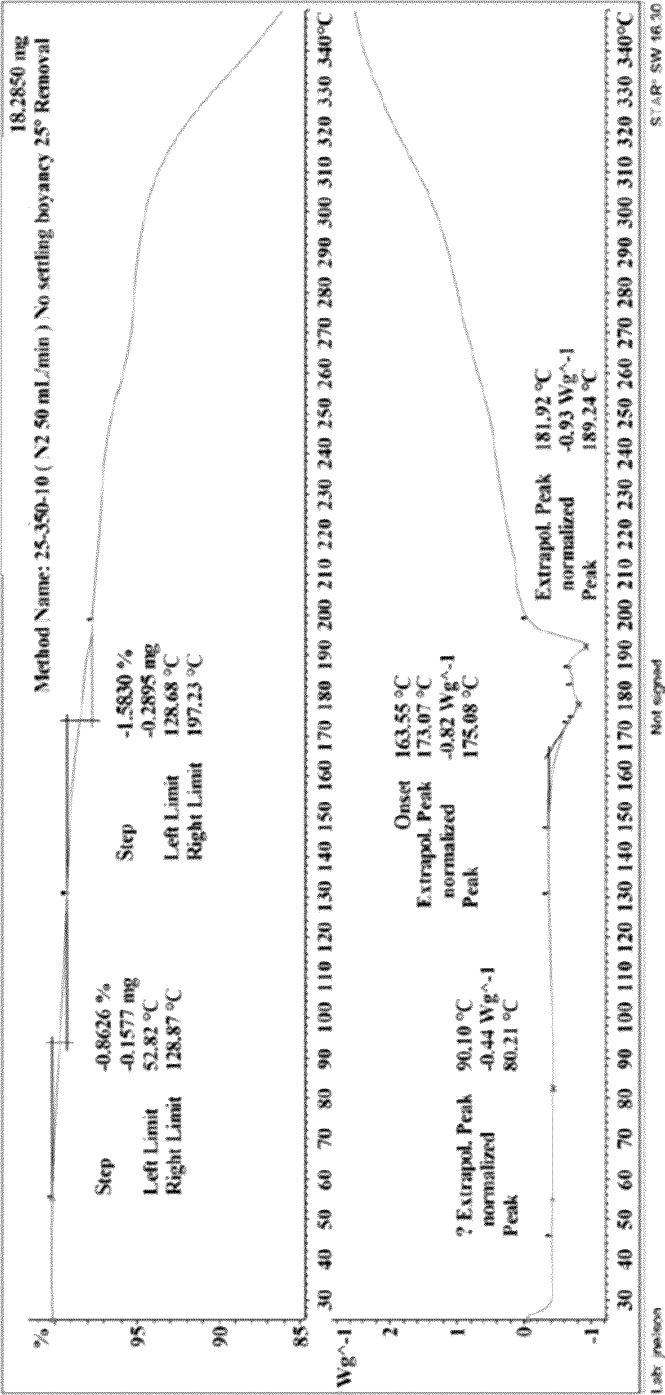
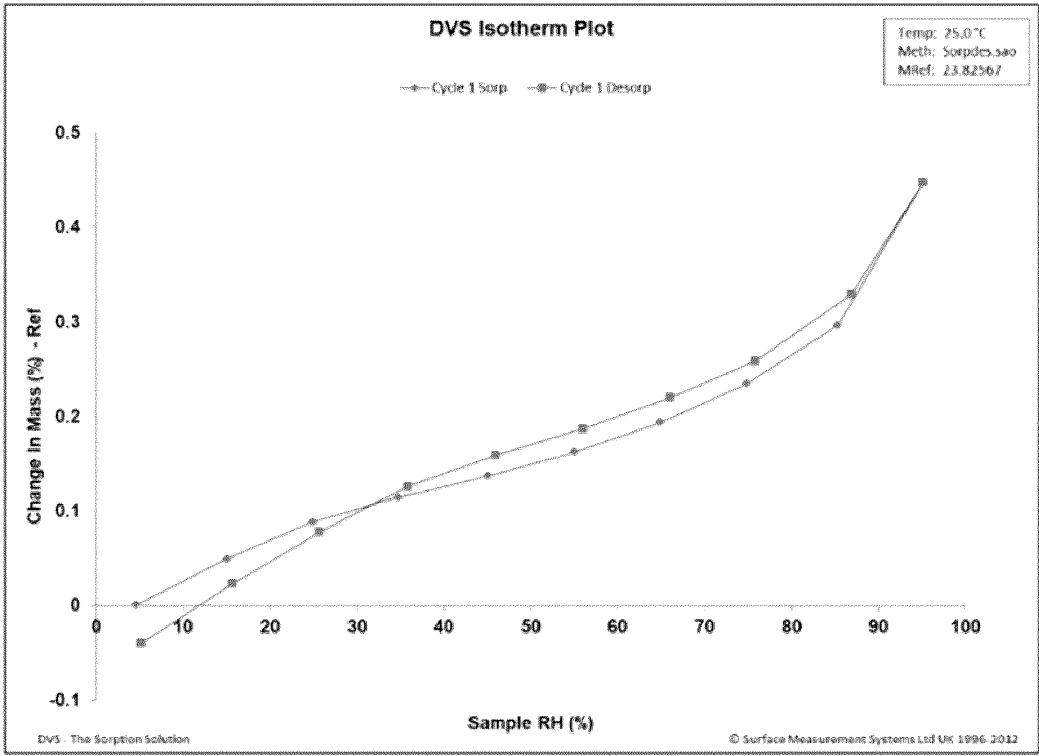
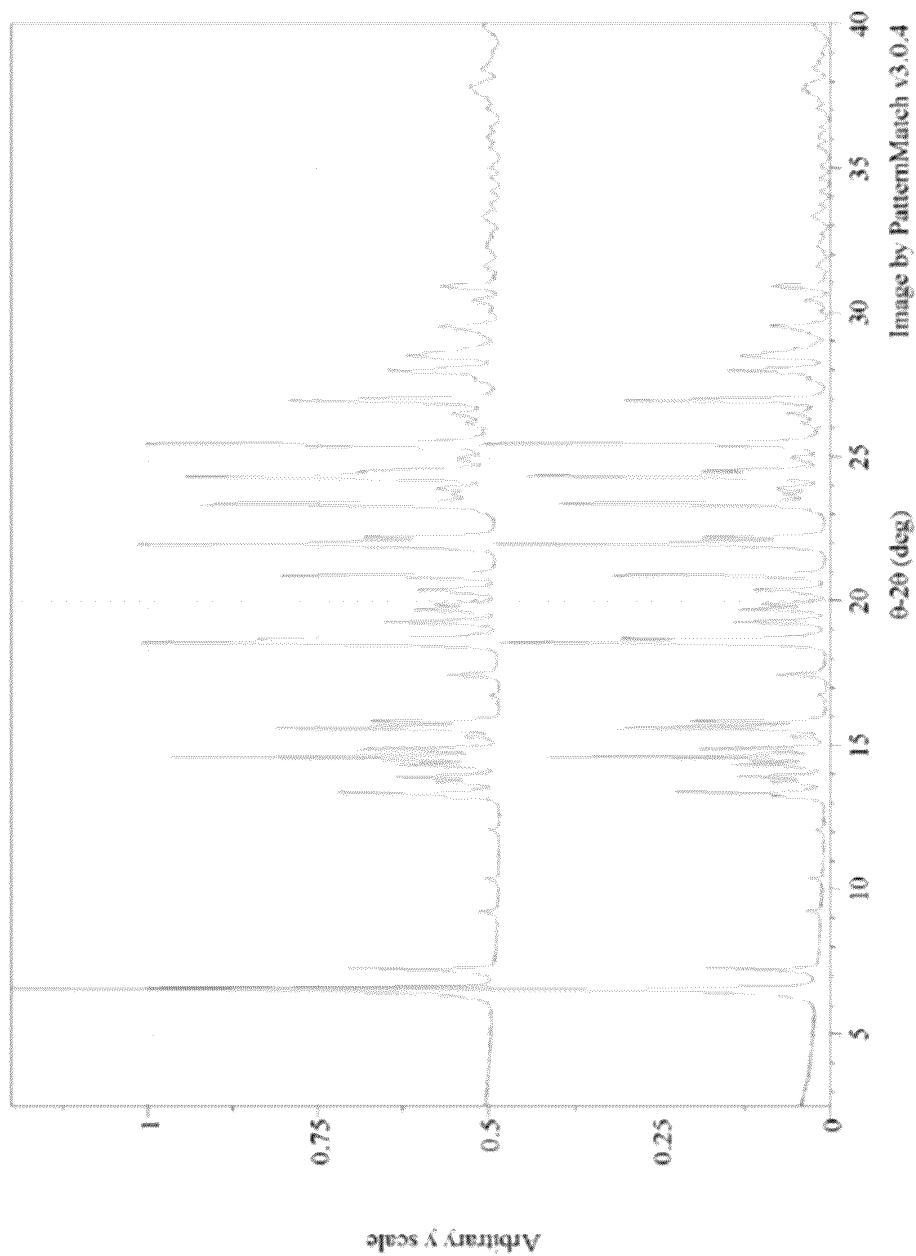


Figure 16

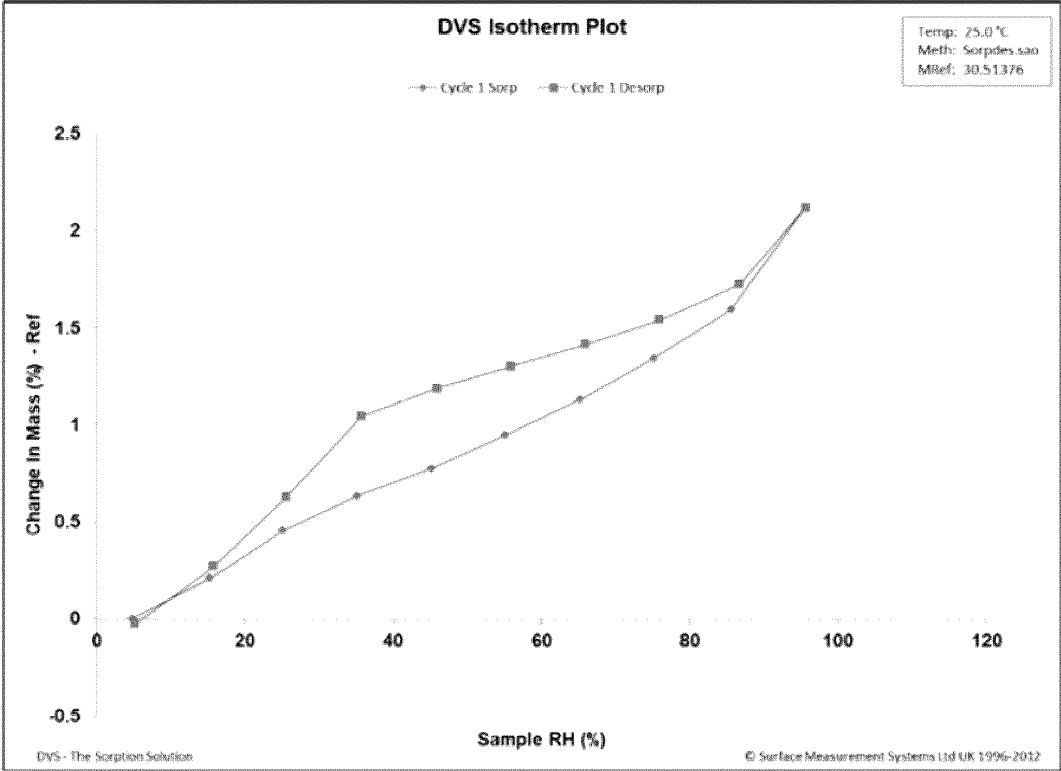


	Target RH (%)	Sample RH (%)	Sorp Mass Change (%)	Sample RH (%)	Desorp Mass Change (%)	Hysteresis
Cycle 1	5.0	4.6	0.0001	5.2	-0.0402	
	15.0	15.0	0.0492	15.7	0.0228	-0.0264
	25.0	24.9	0.0883	25.6	0.0769	-0.0113
	35.0	34.8	0.1143	35.8	0.1256	0.0113
	45.0	45.0	0.1370	46.0	0.1588	0.0218
	55.0	55.0	0.1621	56.0	0.1873	0.0252
	65.0	64.9	0.1936	66.0	0.2196	0.0260
	75.0	74.8	0.2343	75.9	0.2587	0.0243
	85.0	85.3	0.2960	86.8	0.3288	0.0327
	95.0	95.2	0.4475	95.2	0.4475	

Figure 17

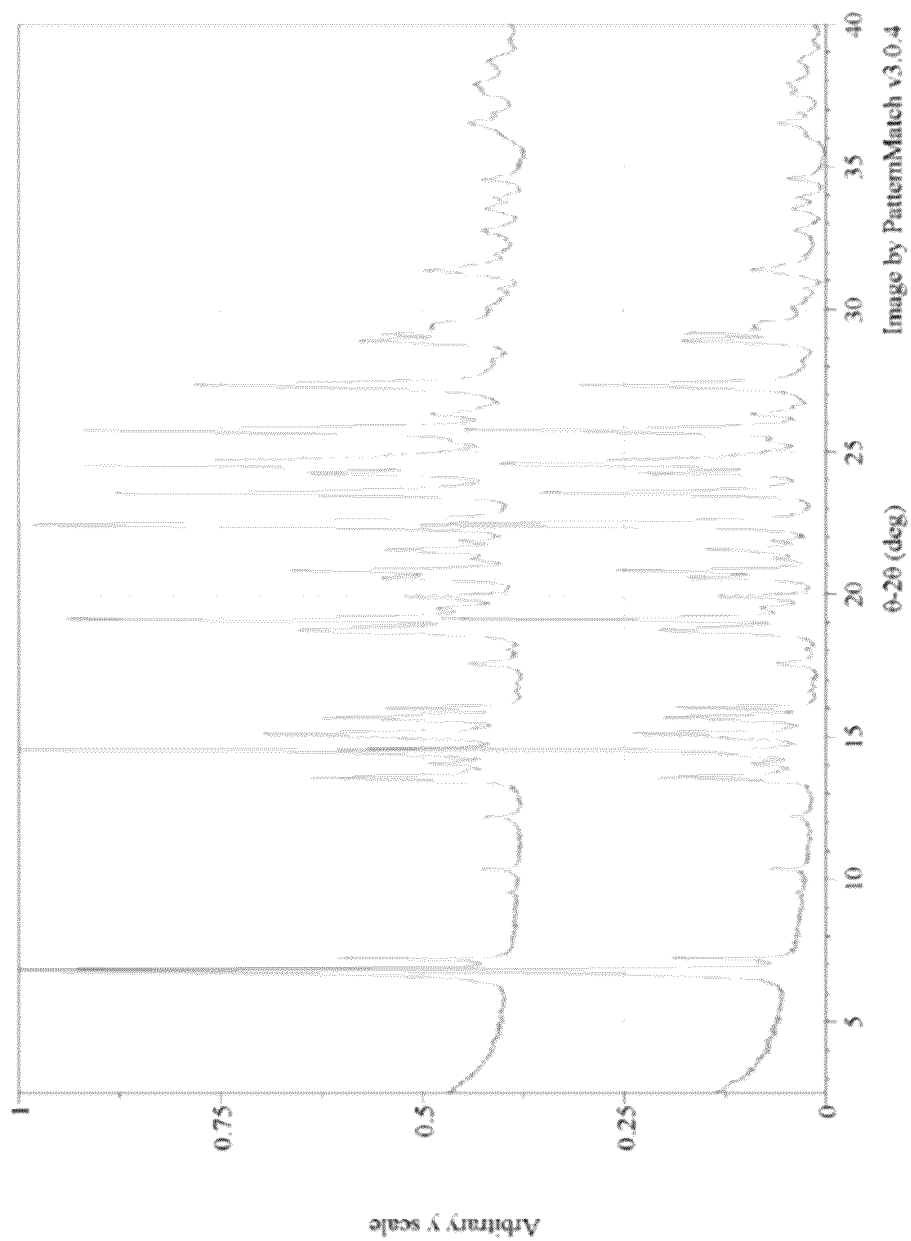
**Figure 18**

Up : Compound 2 pre- DVS
Down : Compound 2 post- DVS



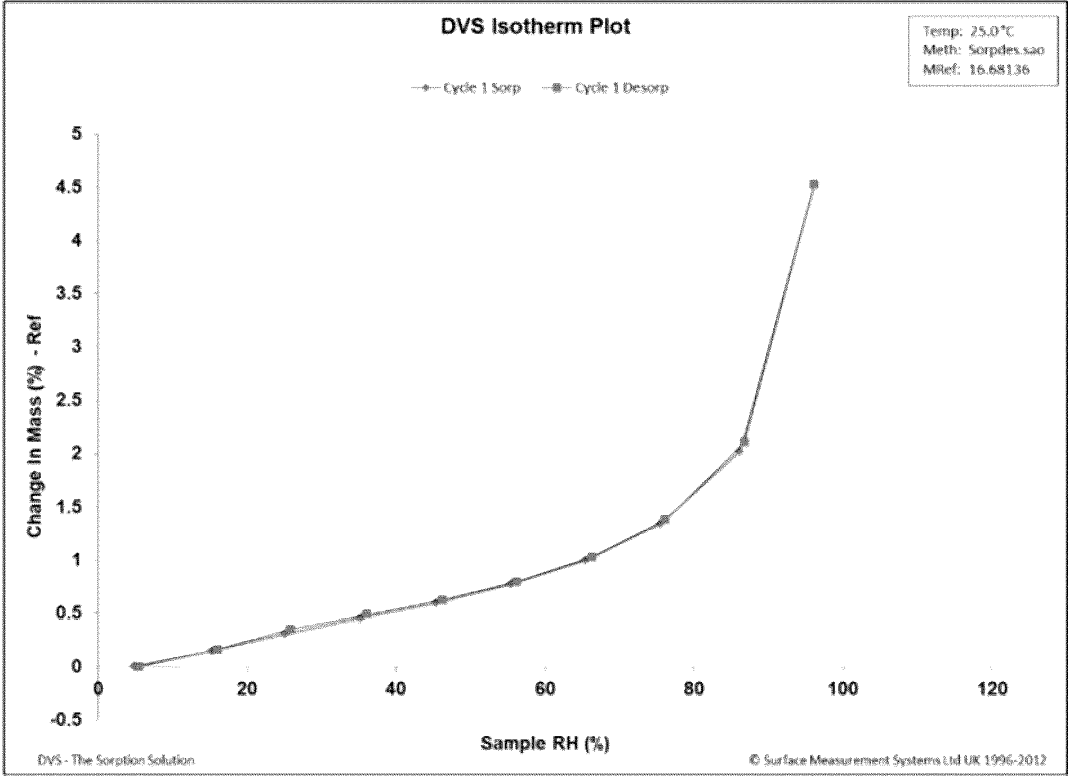
	Target RH (%)	Sample RH (%)	Sorp Mass Change (%)	Sample RH (%)	Desorp Mass Change (%)	Hysteresis
Cycle 1	5.0	4.8	0.000	5.2	-0.029	
	15.0	15.3	0.209	15.7	0.271	0.062
	25.0	25.1	0.454	25.6	0.628	0.174
	35.0	35.1	0.633	35.8	1.044	0.411
	45.0	45.2	0.775	45.9	1.190	0.415
	55.0	55.1	0.948	55.9	1.301	0.353
	65.0	65.2	1.130	65.9	1.413	0.282
	75.0	75.1	1.342	75.9	1.540	0.197
	85.0	85.6	1.594	86.7	1.725	0.131
	95.0	95.6	2.119	95.6	2.119	

Figure 19

**Figure 20**

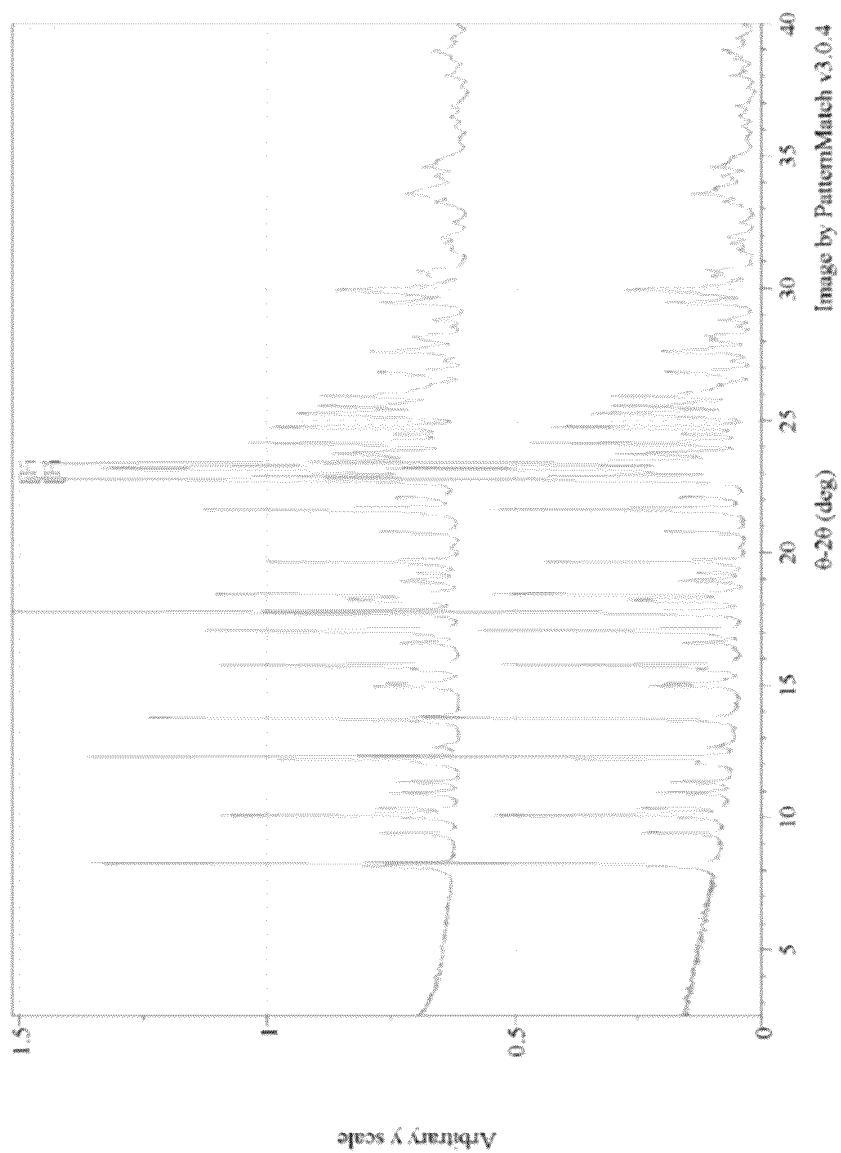
Up : Compound 3 pre- DVS

Down : Compound 3 post- DVS



	Target RH (%)	Sample RH (%)	Sorp Mass Change (%)	Sample RH (%)	Desorp Mass Change (%)	Hysteresis
Cycle 1	5.0	4.8	0.003	5.5	-0.006	
	15.0	15.3	0.142	16.0	0.155	0.013
	25.0	25.1	0.300	25.9	0.346	0.046
	35.0	35.2	0.446	36.1	0.482	0.036
	45.0	45.4	0.590	46.2	0.620	0.030
	55.0	55.4	0.765	56.2	0.790	0.025
	65.0	65.4	0.990	66.3	1.020	0.030
	75.0	75.4	1.335	76.2	1.377	0.041
	85.0	86.1	2.016	86.8	2.100	0.084
	95.0	96.2	4.526	96.2	4.526	

Figure 21

**Figure 22**

Up : Compound 4 pre- DVS

Down : Compound 4 post- DVS

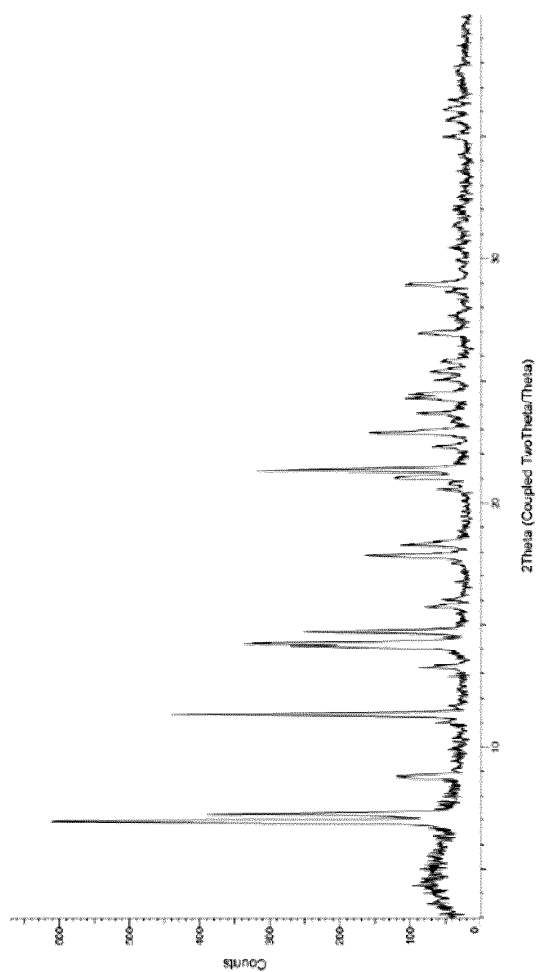


Figure 23

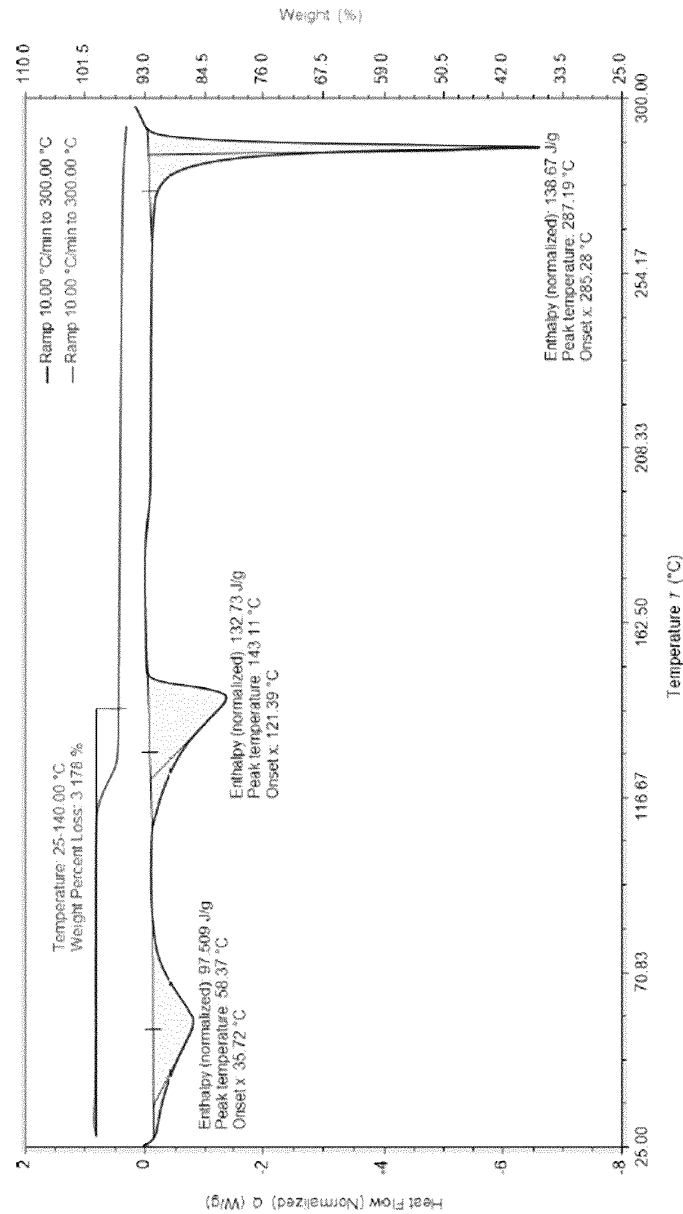


Figure 24

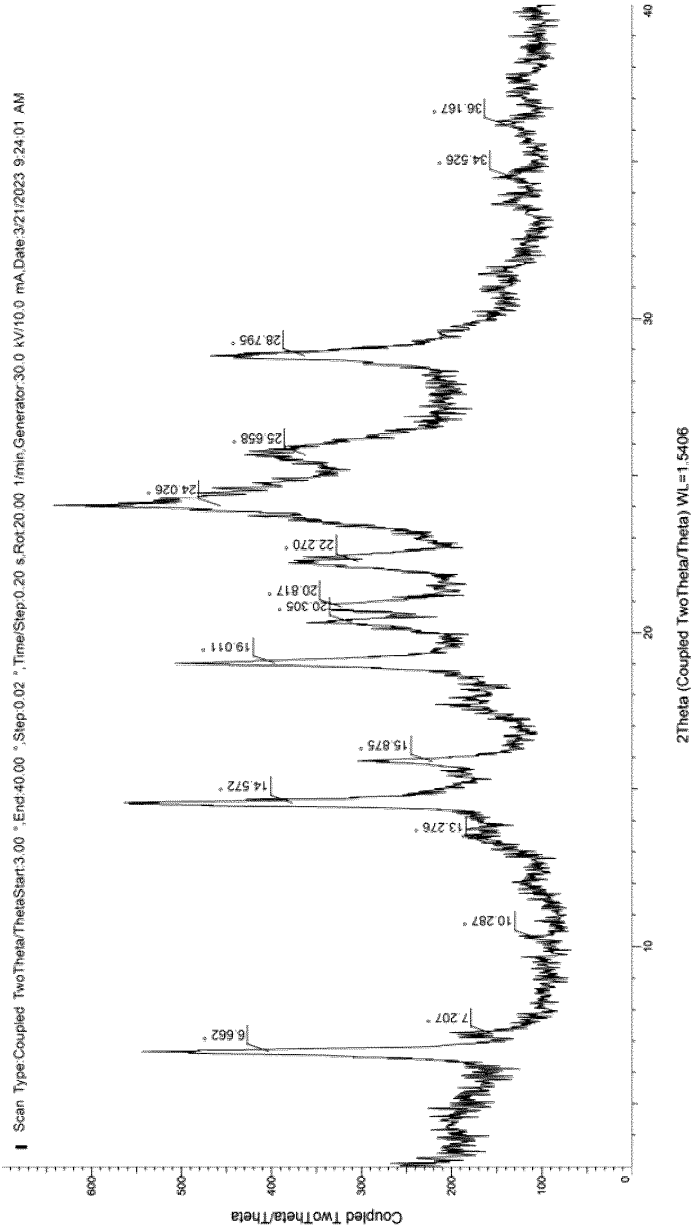


Figure 25

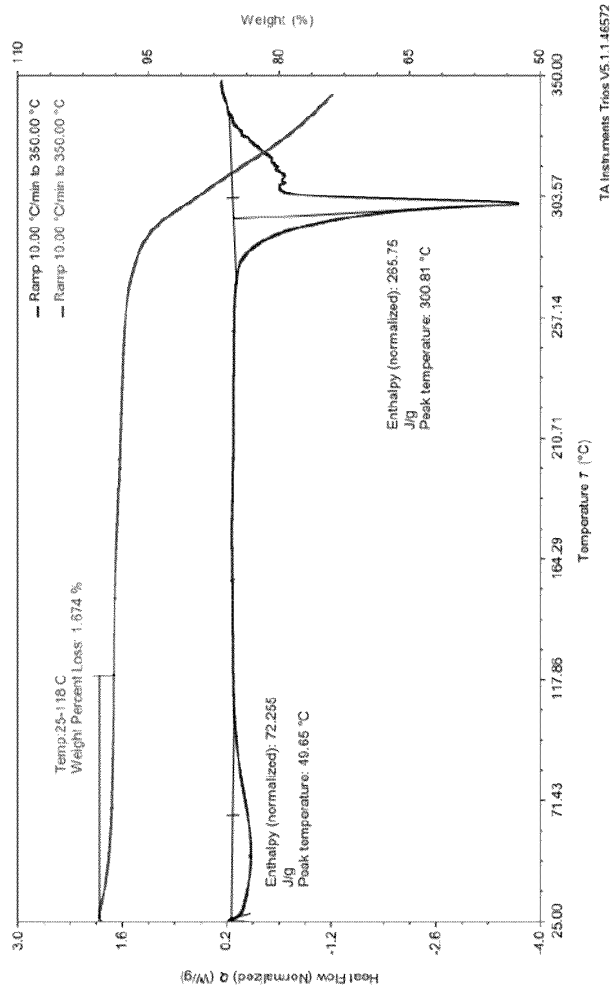


Figure 26

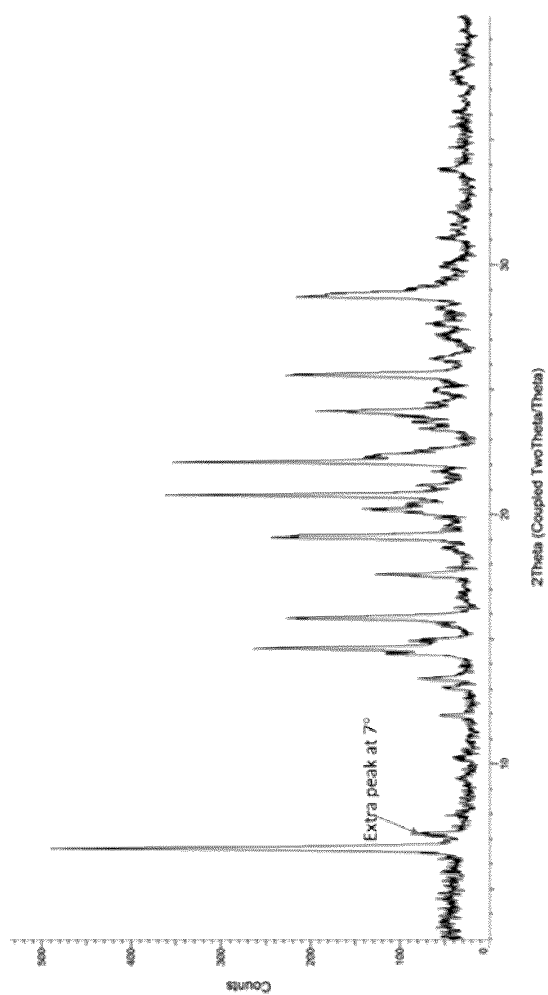


Figure 27

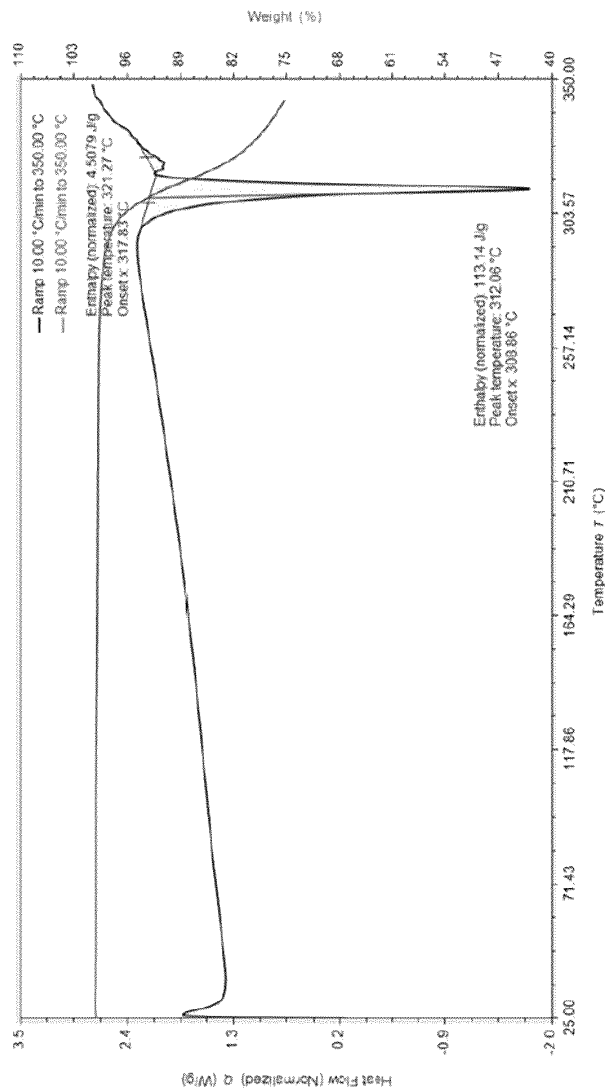


Figure 28

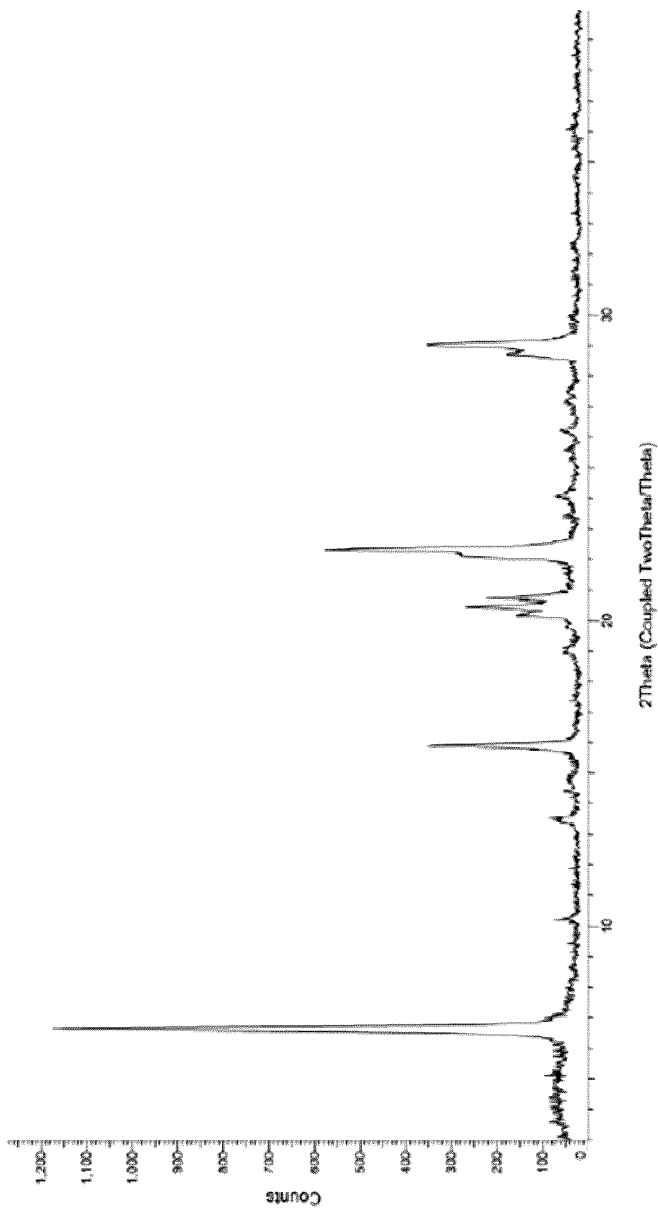


Figure 29

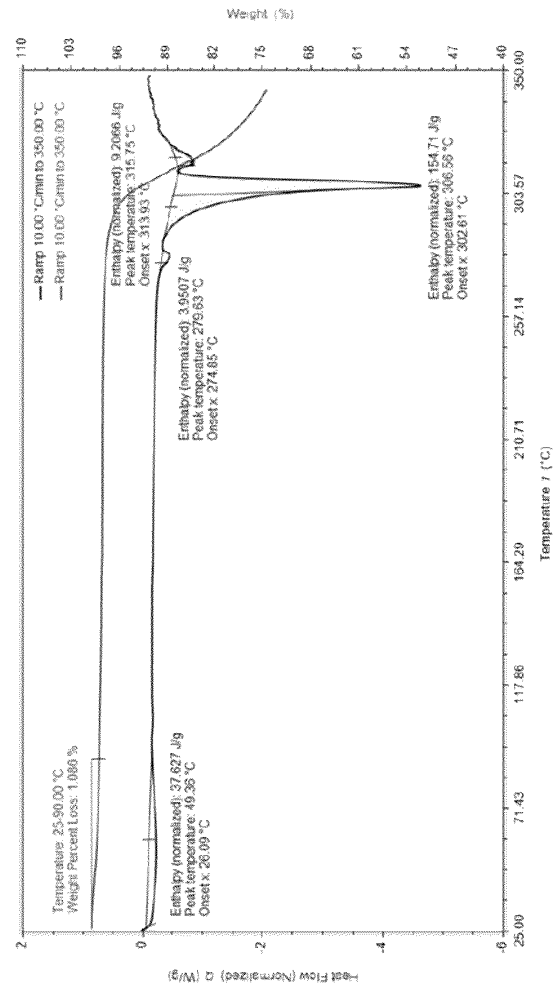


Figure 30

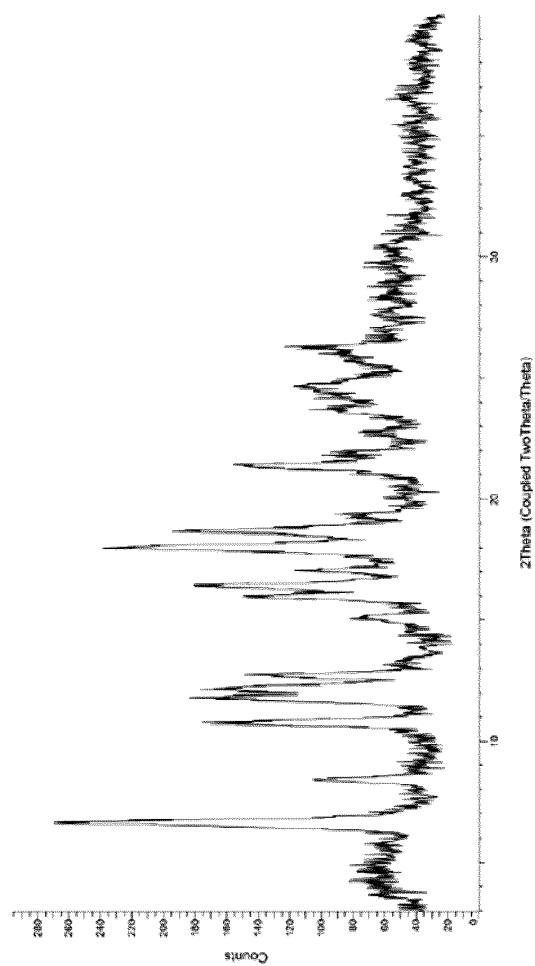


Figure 31

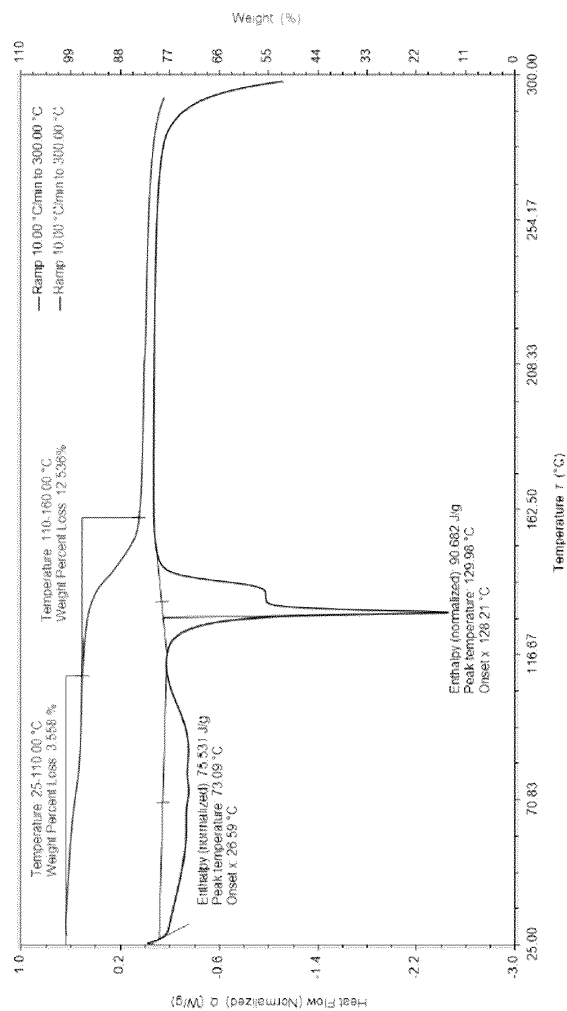


Figure 32

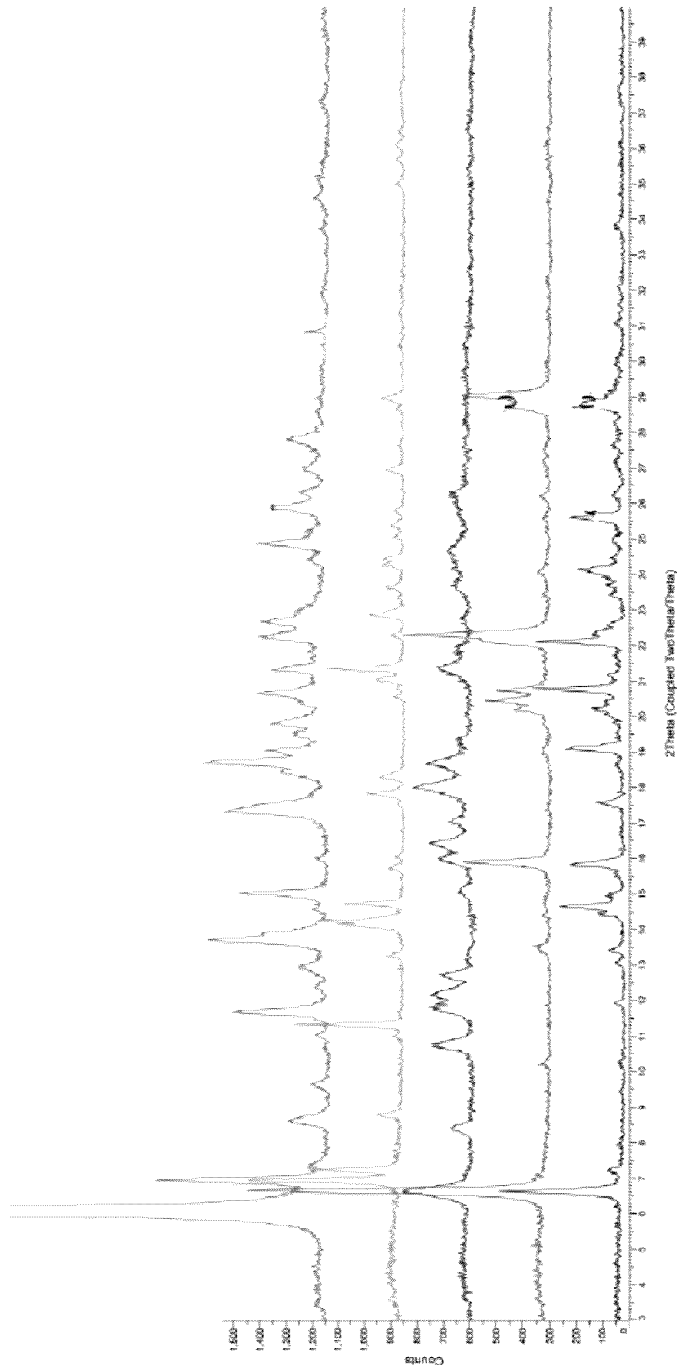


Figure 33

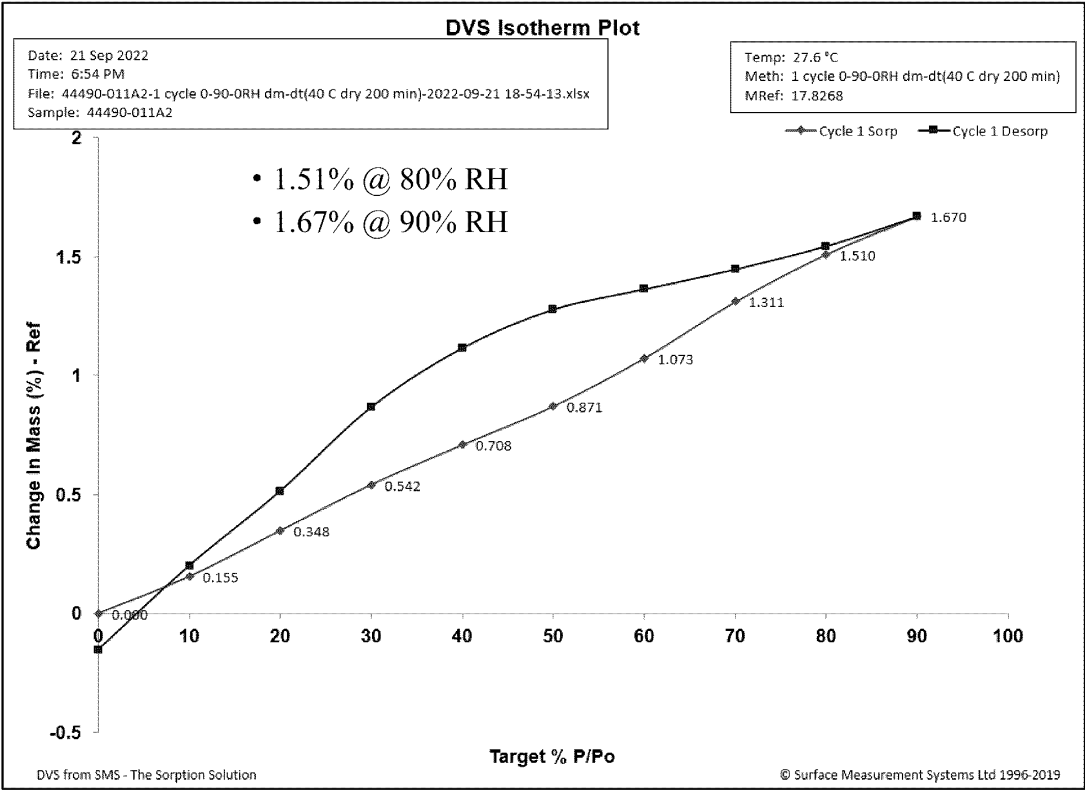


Figure 34

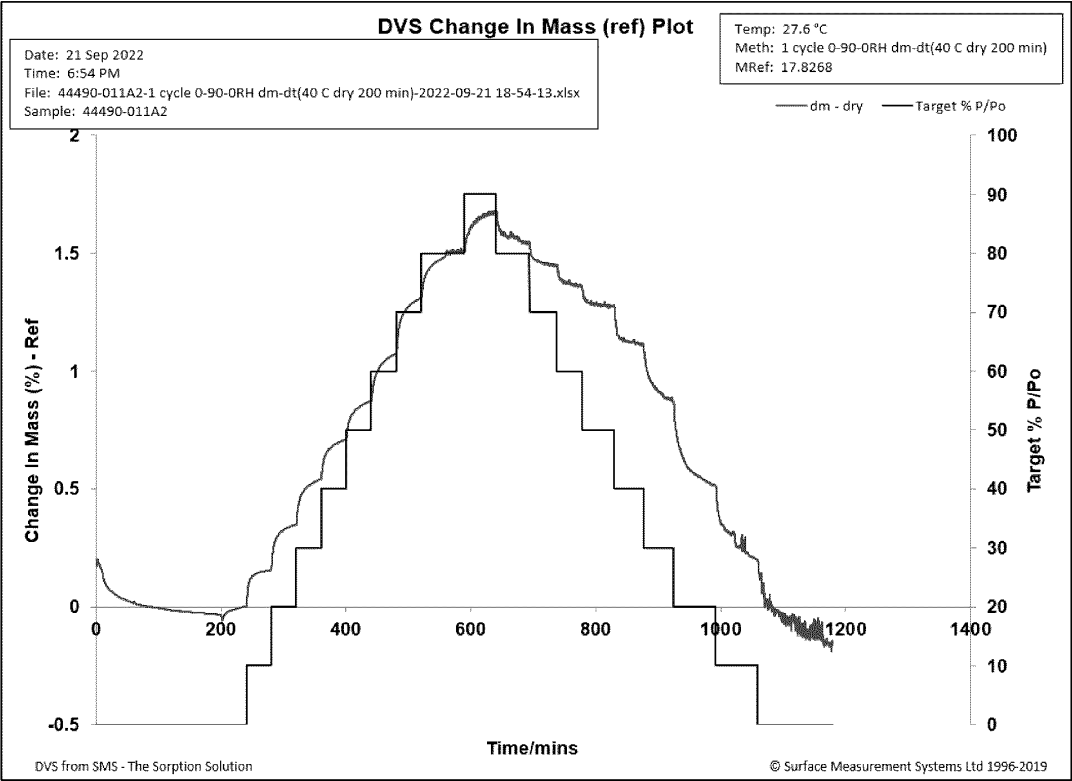


Figure 35

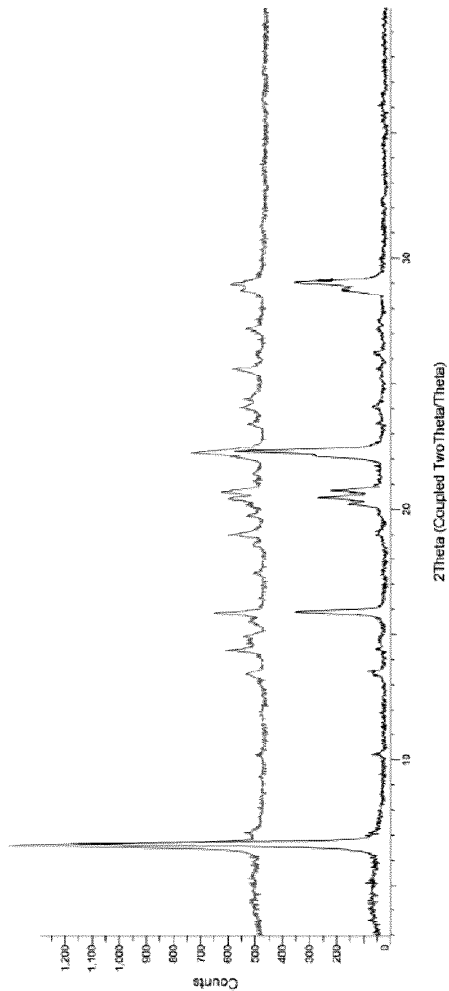


Figure 36

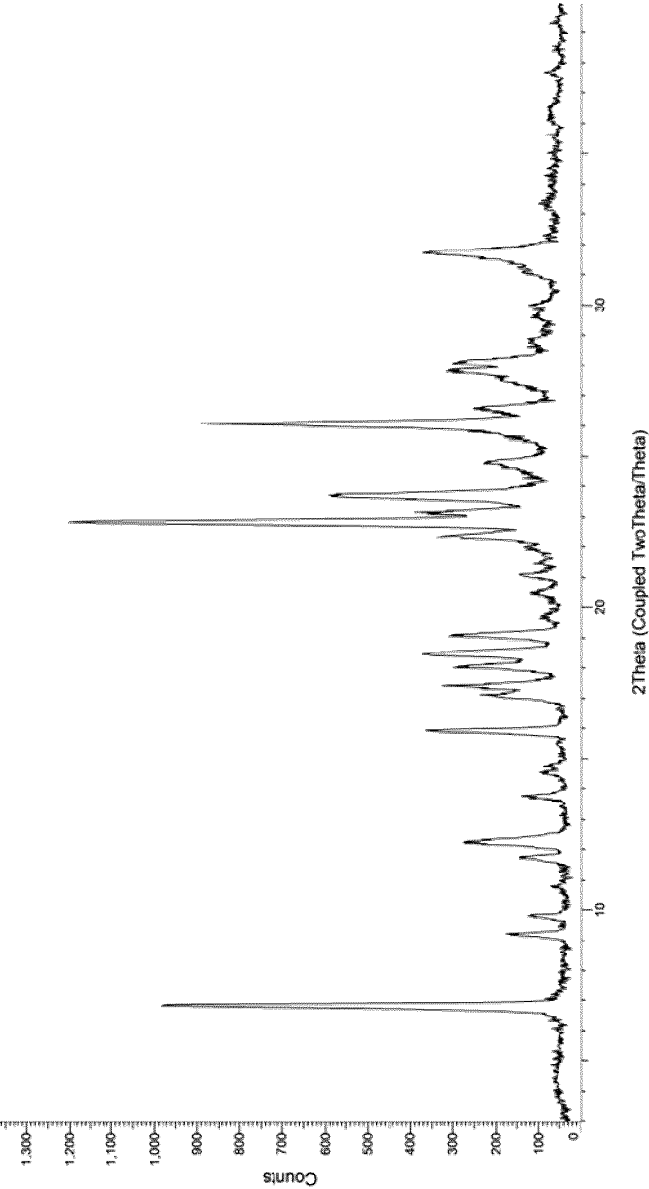


Figure 37

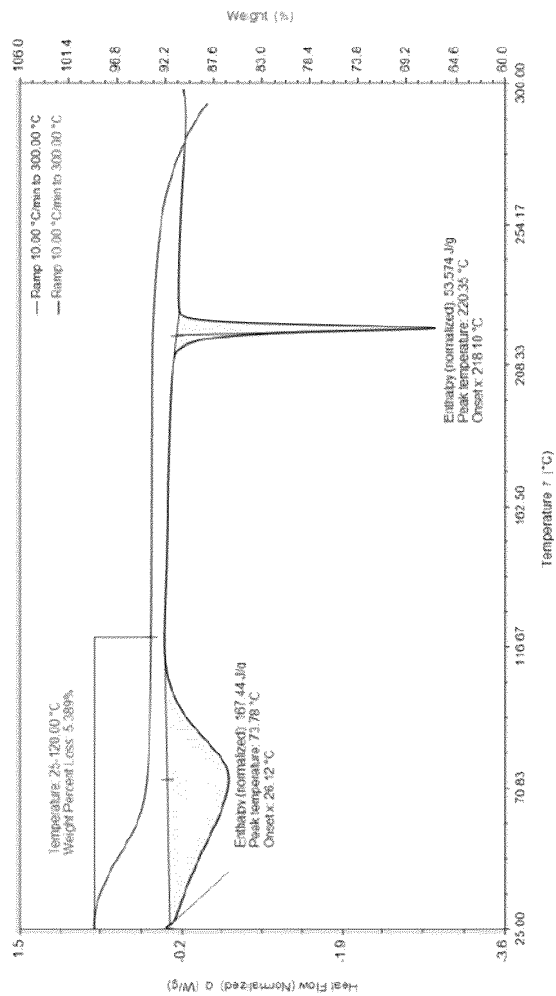
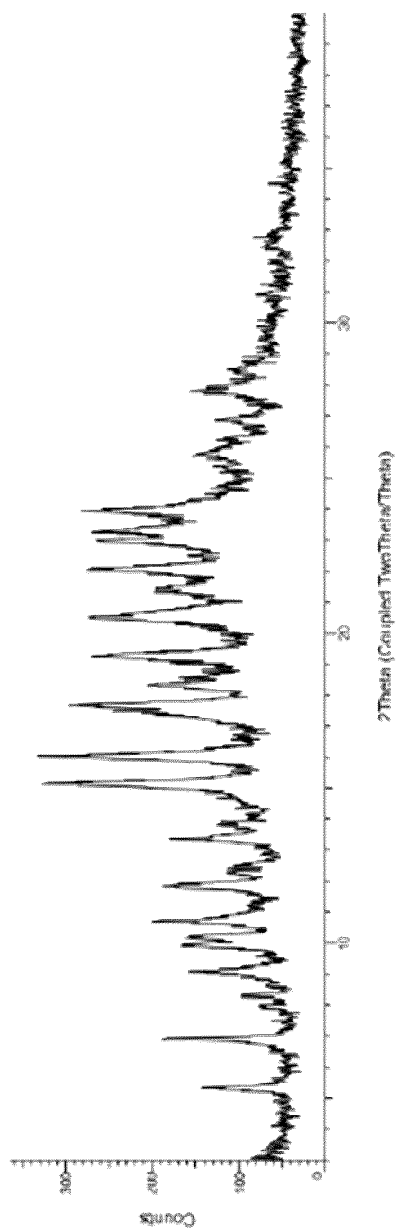


Figure 38

**Figure 39**

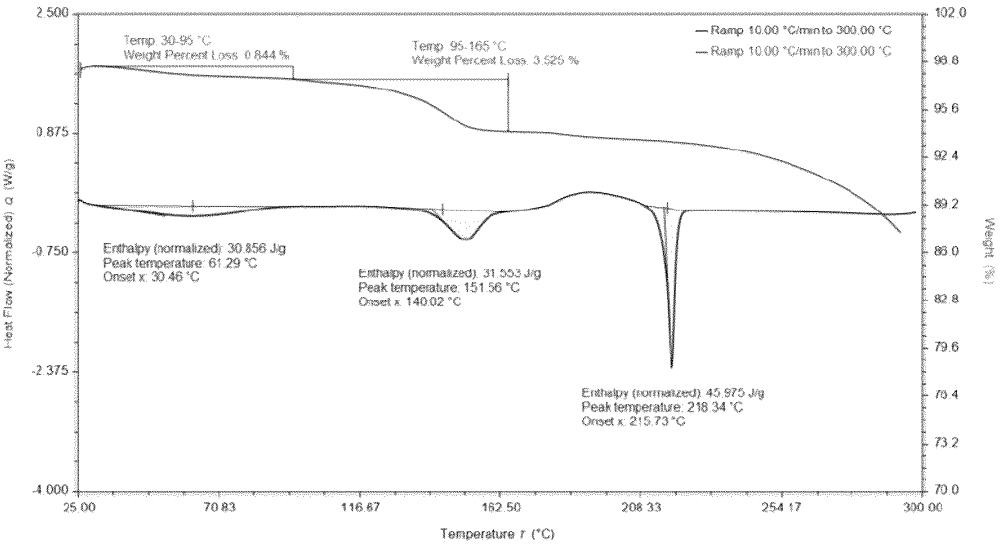


Figure 40

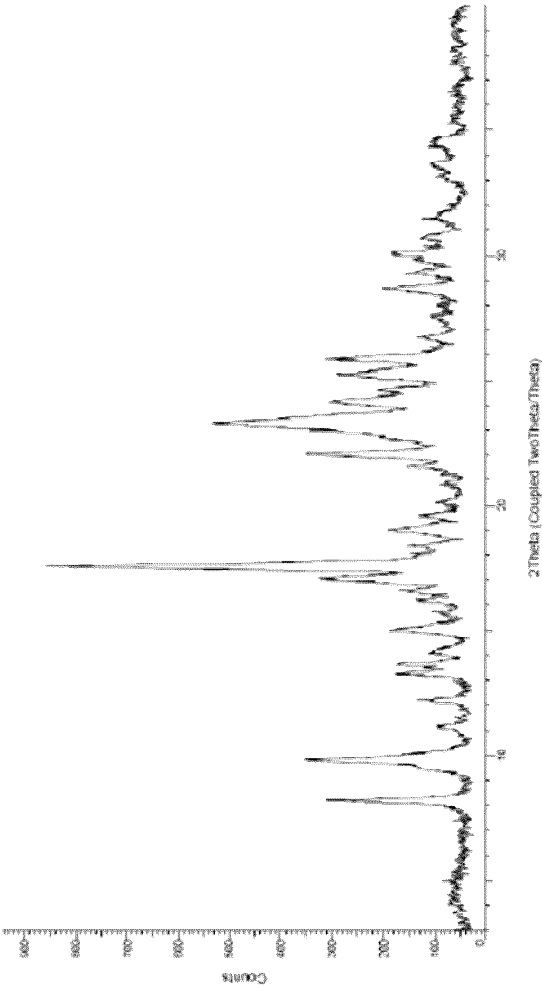


Figure 41

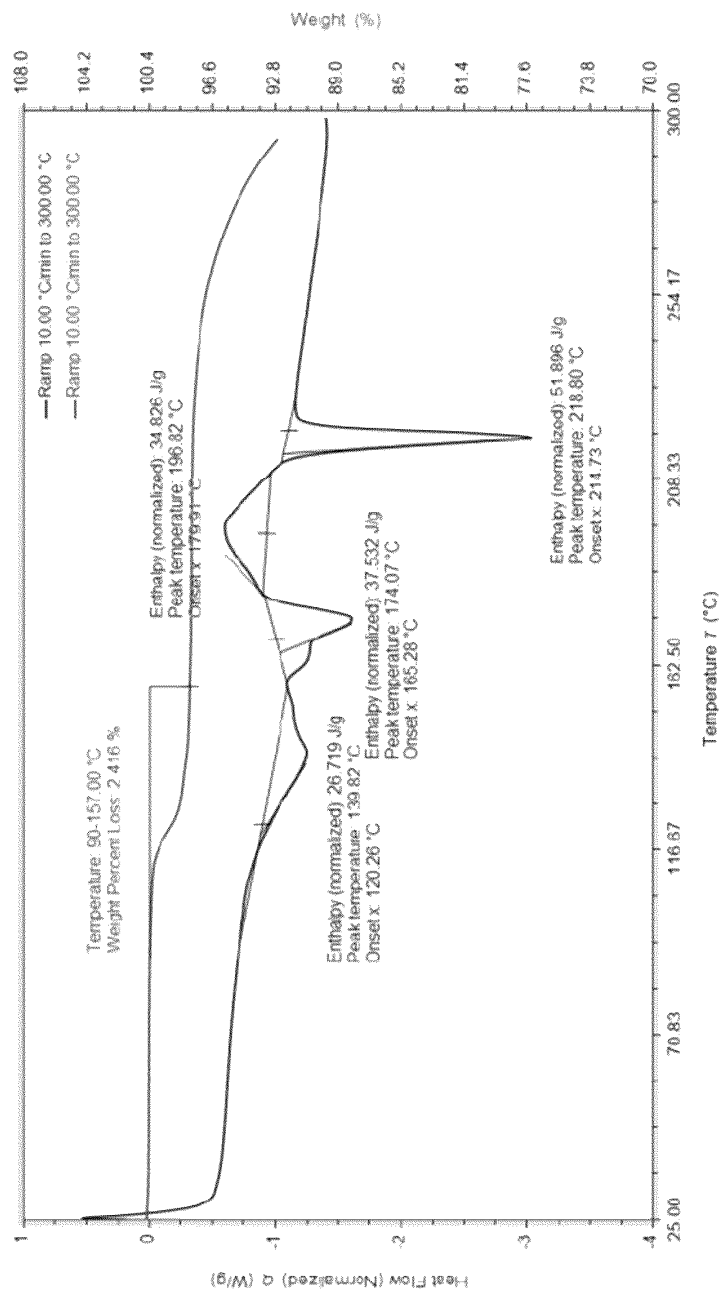


Figure 42

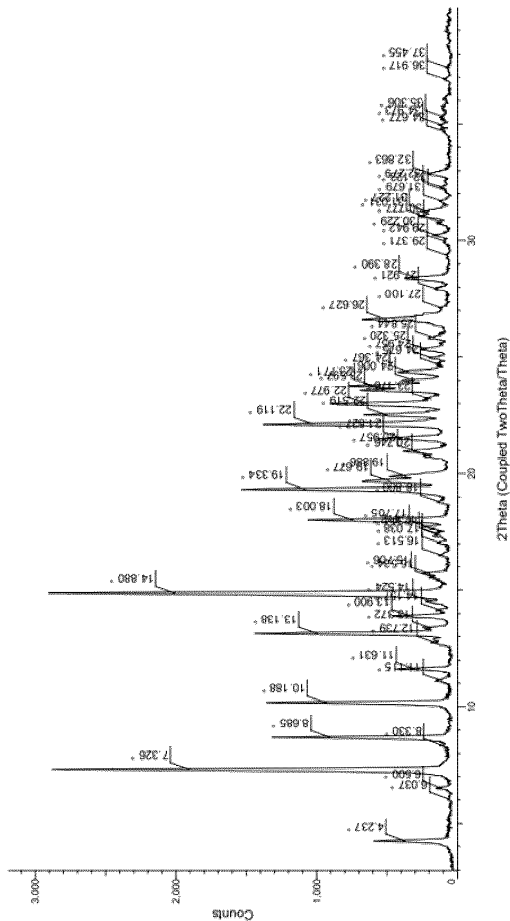


Figure 43

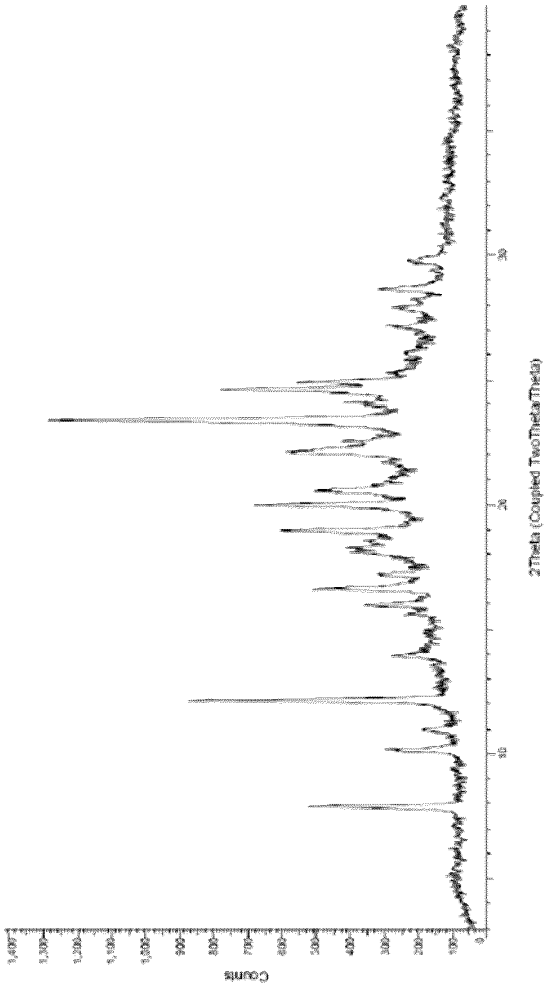


Figure 44

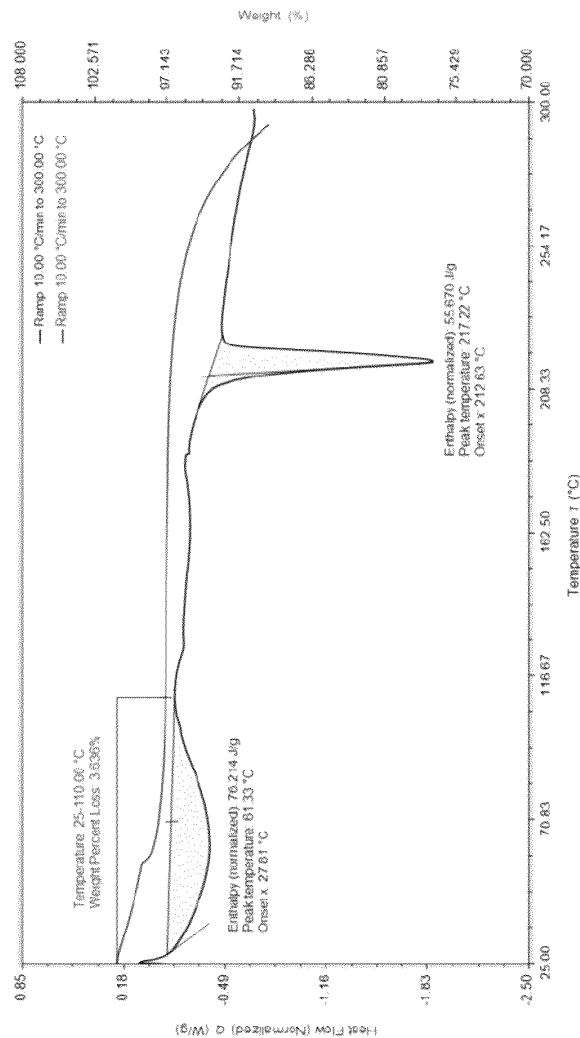


Figure 45

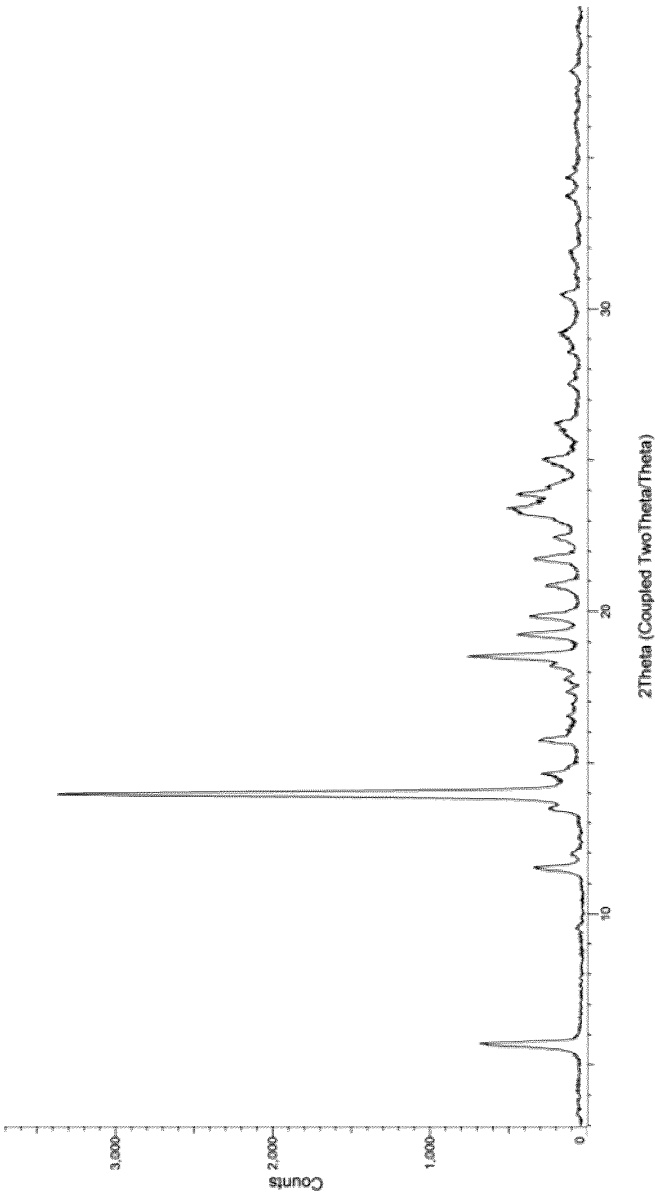


Figure 46

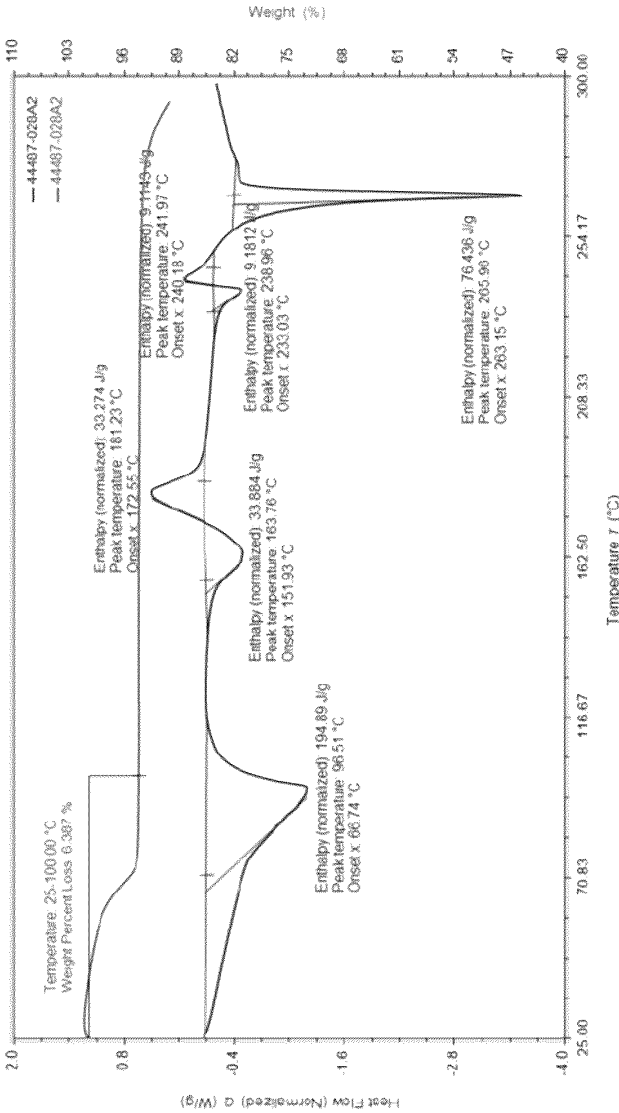


Figure 47

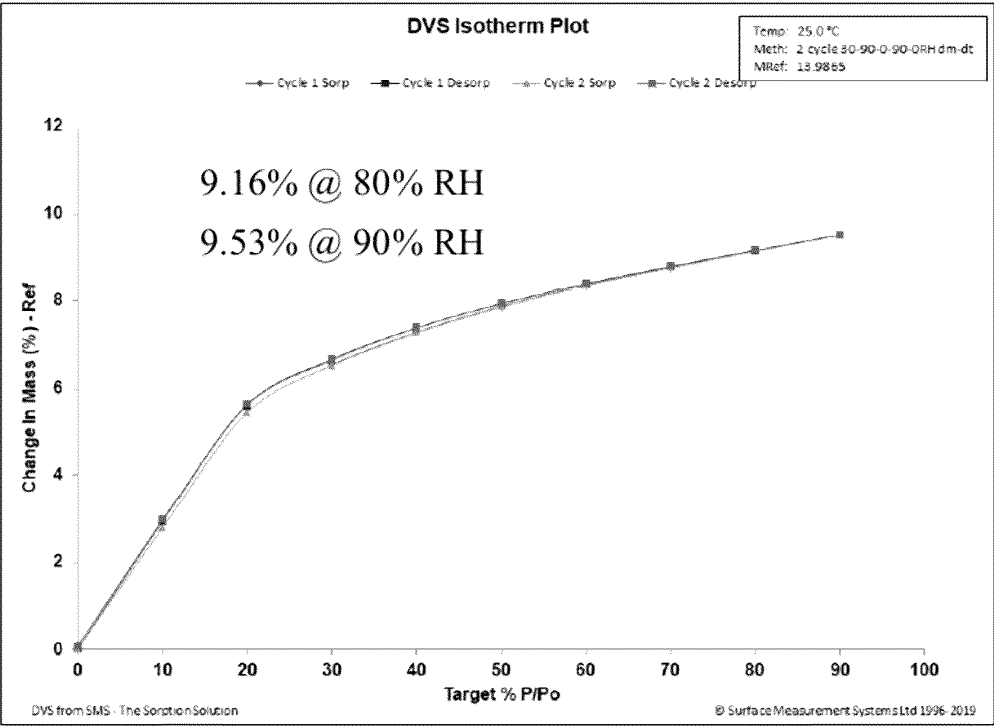
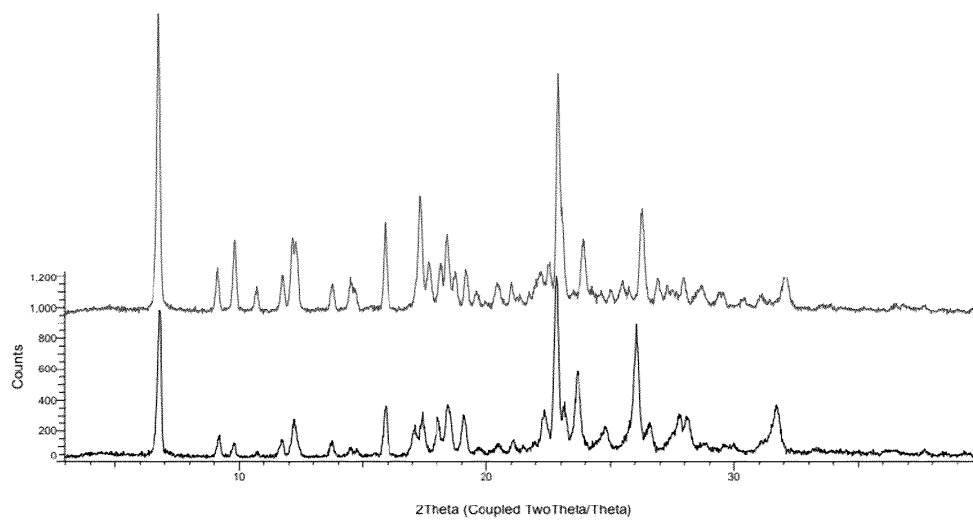


Figure 48

**Figure 49**

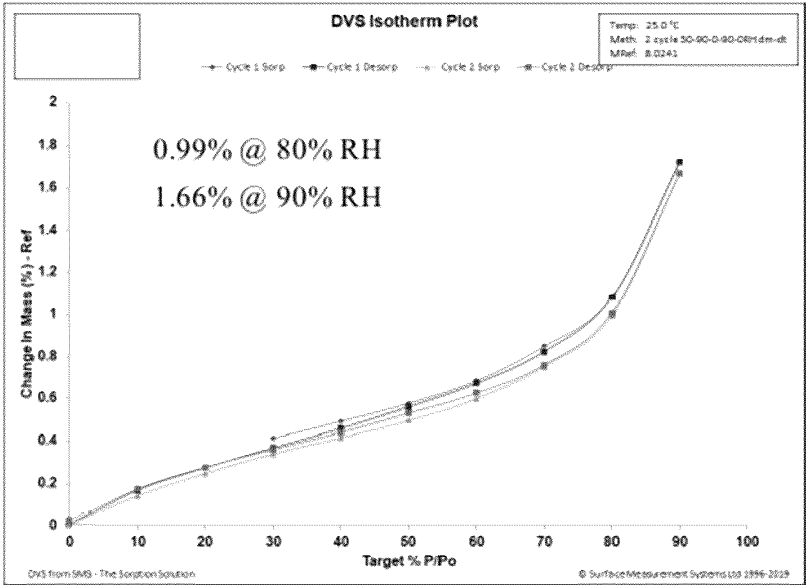
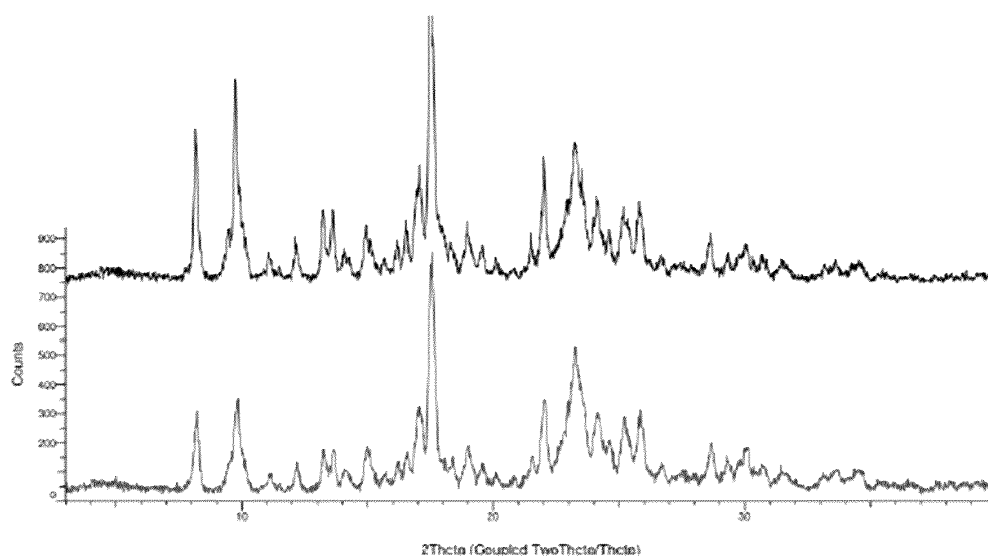


Figure 50

**Figure 51**