

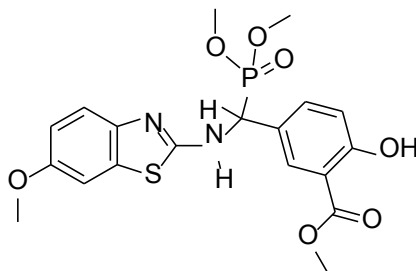
Supplementary Materials

Structure Confirmation and Evaluation of a Nonsteroidal Inhibitor of 17 β -Hydroxysteroid Dehydrogenase Type 10

Content:

- 1) Listing of ^1H and ^{13}C NMR data reported in literature for compound **2**
- 2) Crystal data and structure refinement for CHUL08 (**2**)
- 3) NMR spectra for compound **2** in DMSO- d_6 (^1H NMR, ^{13}C NMR (APT), COSY, HSQC and HMBC)

(1) Listing of ^1H and ^{13}C NMR data reported in literature for compound 2, but without assignation



From the patent (US 2015/0065463 A1):

^1H NMR (500 MHz) in DMSO- d_6 (δ in ppm): 10.48 (s, 1H), 7.91-7.90 (m, 1H), 7.66-7.64 (m, 1H), 7.23-7.22 (m, 1H), 7.06 (d, J = 10 Hz), 6.95 (d, J = 10.0 Hz), 6.89-6.87 (m, 2H), 4.32 (d, J = 15.0 Hz, 1H), 3.90 (s, 3H), 3.72 (s, 3H), 3.55 (d, J = 10.0 Hz, 3H), 3.44 (d, J = 10.0 Hz, 3H).

^{13}C NMR (125 MHz) in DMSO- d_6 (δ in ppm): 169.2, 169.2, 159.1 (d), 157.5, 154.4, 135.7 (d), 134.4, 129.4, 129.2 (d), 122.2, 116.8, 112.6, 112.2 (d), 109.9, 108.1, 66.7, 65.4, 55.6, 52.9 (d), 52.4.

From the article (Chem Biol Drug Des, 2013, 81, 238-249):

^1H NMR (500 MHz) in DMSO- d_6 (δ in ppm): 10.47 (s, 1H), 7.91 (s, 1H), 7.66-7.64 (m, 1H), 7.63-7.60 (m, 1H), 7.25-7.23 (m, 2H), 6.97-6.95 (m, 1H), 6.90-6.88 (m, 1H), 5.35 (dd, J = 20.0, 20.0 Hz, 1H), 3.91 (s, 3H), 3.58 (s, 3H), 3.56-3.46 (m, 6H).

^{13}C NMR (125 MHz) in DMSO- d_6 (δ in ppm): 169.5, 166.2, 158.7 (d, J_{CP} = 20 Hz), 157.8, 155.9, 148.9 (d, J_{CP} = 36.2 Hz), 148.0 (d, J_{CP} = 20.0 Hz), 135.8 (d, J_{CP} = 36.2 Hz), 131.8 (d, J_{CP} = 11.2 Hz), 128.5, 118.0 (d, J_{CP} = 10.0 Hz), 116.3, 112.0 (d, J_{CP} = 22.5 Hz), 111.7, 107.0 (d, J_{CP} = 26.2 Hz), 56.4, 55.3, 52.2.

(2) Crystal data and structure refinement for CHUL08 (2)

checkCIF/PLATON report

Structure factors have been supplied for datablock(s) chul08

THIS REPORT IS FOR GUIDANCE ONLY. IF USED AS PART OF A REVIEW PROCEDURE FOR PUBLICATION, IT SHOULD NOT REPLACE THE EXPERTISE OF AN EXPERIENCED CRYSTALLOGRAPHIC REFEREE.

No syntax errors found. CIF dictionary Interpreting this report

Datablock: chul08

Bond precision:	C-C = 0.0042 Å	Wavelength=1.34139
Cell:	a=11.7243 (6) alpha=90	b=24.6420 (12) beta=106.034 (2)
		c=7.8468 (4) gamma=90
Temperature:	150 K	
	Calculated	Reported
Volume	2178.83 (19)	2178.83 (19)
Space group	P 21/c	P 1 21/c 1
Hall group	-P 2ybc	-P 2ybc
Moiety formula	2 (C19 H21 N2 O7 P S), C H4 O	C19 H21 N2 O7 P S, 0.5 (C H4 O)
Sum formula Mr	C39 H46 N4 O15 P2 S2	C19.50 H23 N2 O7.50 P S
Dx, g cm ⁻³	1.428	1.428
Z	2	4
Mu (mm ⁻¹)	1.588	1.576
1) F000	980.0	980.0
F000'	984.49	
h,k,lmax	15,31,10	15,31,10
Nref	4941	4900
Tmin,Tmax	0.893,0.95	0.639,0.752
Tmin'	0.789	

Correction method= # Reported T Limits: Tmin=0.639 Tmax=0.752

AbsCorr = MULTI-SCAN

Data completeness= 0.992 Theta(max)= 60.176

R(reflections)= 0.0599 (3862) wR2(reflections)= 0.1718 (4900)

S = 1.065 Npar= 316

The following ALERTS were generated. Each ALERT has the format

test-name_ALERT_alert-type_alert-level.

Click on the hyperlinks for more details of the test.



Alert level C

PLAT340_ALERT_3_C	Low Bond Precision on C-C Bonds	0.00421 Ang.
PLAT413_ALERT_2_C	Short Inter XH3 .. XHn H19C ..H1AC	2.13 Ang.
PLAT906_ALERT_3_C	Large K Value in the Analysis of Variance	3.610 Check



Alert level G

ABSMU01_ALERT_1_G Calculation of _exptl_absorpt_correction_mu not performed for this radiation type.		
PLAT002_ALERT_2_G	Number of Distance or Angle Restraints on AtSite	4 Note
PLAT003_ALERT_2_G	Number of Uiso or Uij Restrained non-H Atoms ...	4 Report
PLAT007_ALERT_5_G	Number of Unrefined Donor-H Atoms	4 Report
PLAT042_ALERT_1_G	Calc. and Reported MoietyFormula Strings Differ	Please Check
PLAT045_ALERT_1_G	Calculated and Reported Z Differ by a Factor ...	0.50 Check
PLAT172_ALERT_4_G	The CIF-Embedded .res File Contains DFIX Records	1 Report
PLAT175_ALERT_4_G	The CIF-Embedded .res File Contains SAME Records	1 Report
PLAT186_ALERT_4_G	The CIF-Embedded .res File Contains ISOR Records	1 Report
PLAT302_ALERT_4_G	Anion/Solvent/Minor-Residue Disorder (Resd 2)	100% Note
PLAT302_ALERT_4_G	Anion/Solvent/Minor-Residue Disorder (Resd 3)	100% Note
PLAT304_ALERT_4_G	Non-Integer Number of Atoms in Resd 2	1.53 Check
PLAT304_ALERT_4_G	Non-Integer Number of Atoms in Resd 3	1.47 Check
PLAT395_ALERT_2_G	Deviating X-O-Y Angle From 120 for O5	120.4 Degree
PLAT395_ALERT_2_G	Deviating X-O-Y Angle From 120 for O6	122.0 Degree
PLAT432_ALERT_2_G	Short Inter X...Y Contact N2 ..C1A	2.95 Ang.
PLAT720_ALERT_4_G	Number of Unusual/Non-Standard Labels	7 Note
PLAT793_ALERT_4_G	Model has Chirality at C1 (Centro SPGR)	R Verify
PLAT860_ALERT_3_G	Number of Least-Squares Restraints	27 Note
PLAT912_ALERT_4_G	Missing # of FCF Reflections Above STh/L= 0.600	41 Note
PLAT933_ALERT_2_G	Number of OMIT Records in Embedded .res File ...	1 Note
PLAT978_ALERT_2_G	Number C-C Bonds with Positive Residual Density.	1 Info

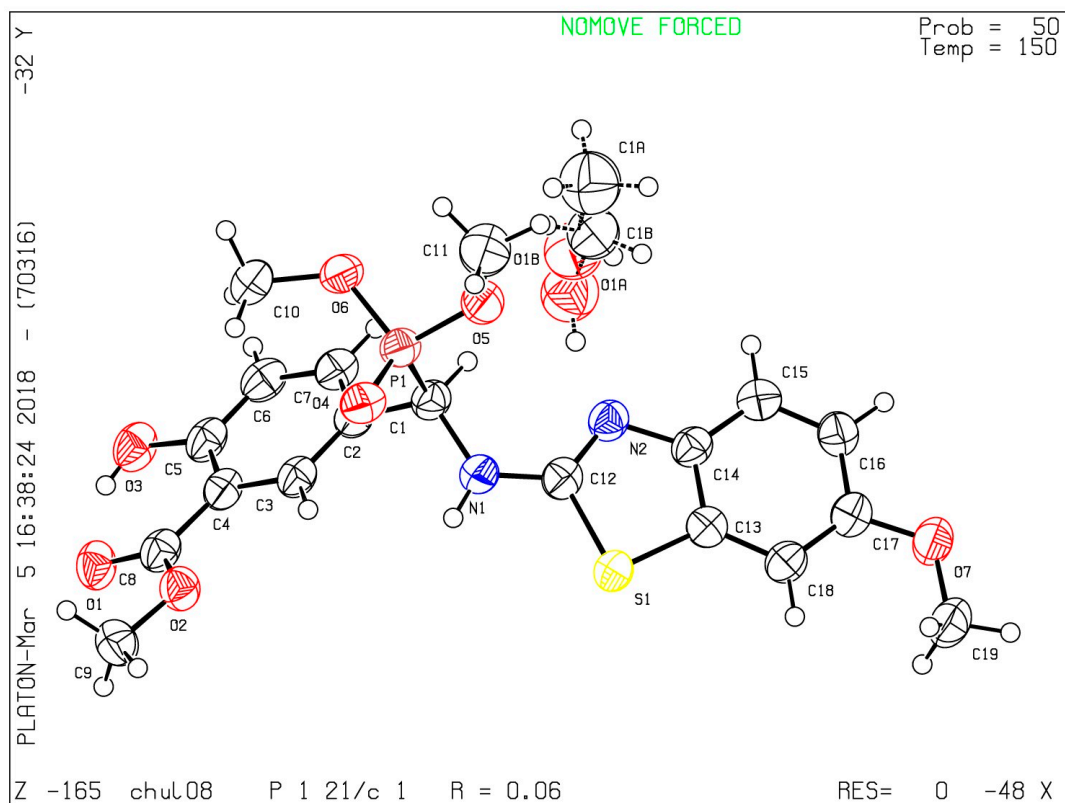
-
- 0 **ALERT level A** = Most likely a serious problem - resolve or explain
 0 **ALERT level B** = A potentially serious problem, consider carefully
 3 **ALERT level C** = Check. Ensure it is not caused by an omission or oversight
 22 **ALERT level G** = General information/check it is not something unexpected
- 3 ALERT type 1 CIF construction/syntax error, inconsistent or missing data
 8 ALERT type 2 Indicator that the structure model may be wrong or deficient
 3 ALERT type 3 Indicator that the structure quality may be low
 10 ALERT type 4 Improvement, methodology, query or suggestion
 1 ALERT type 5 Informative message, check
-

It is advisable to attempt to resolve as many as possible of the alerts in all categories. Often the minor alerts point to easily fixed oversights, errors and omissions in your CIF or refinement strategy, so attention to these fine details can be worthwhile. In order to resolve some of the more serious problems it may be necessary to carry out additional measurements or structure refinements. However, the purpose of your study may justify the reported deviations and the more serious of these should normally be commented upon in the discussion or experimental section of a paper or in the "special_details" fields of the CIF. checkCIF was carefully designed to identify outliers and unusual parameters, but every test has its limitations and alerts that are not important in a particular case may appear. Conversely, the absence of alerts does not guarantee there are no aspects of the results needing attention. It is up to the individual to critically assess their own results and, if necessary, seek expert advice.

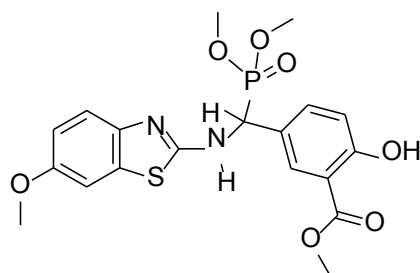
Publication of your CIF in other journals

Please refer to *the Notes for Authors* of the relevant journal for any special instructions relating to CIF submission.

PLATON version of 30/01/2018; check.def file version of 30/01/2018



(3) NMR spectra for compound 2 in DMSO-d₆ (¹H NMR, ¹³C NMR (APT), COSY, HSQC and HMBC)



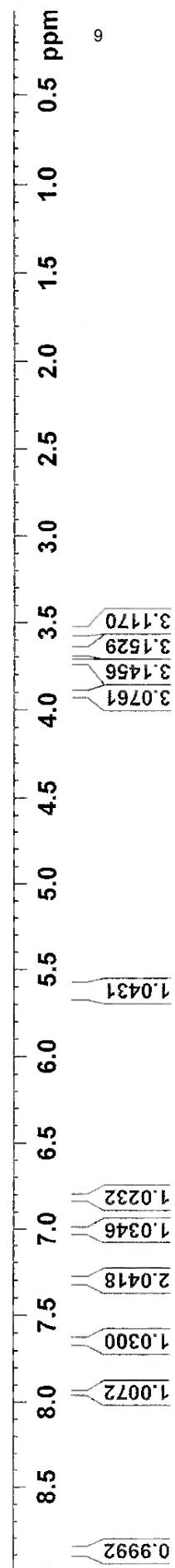
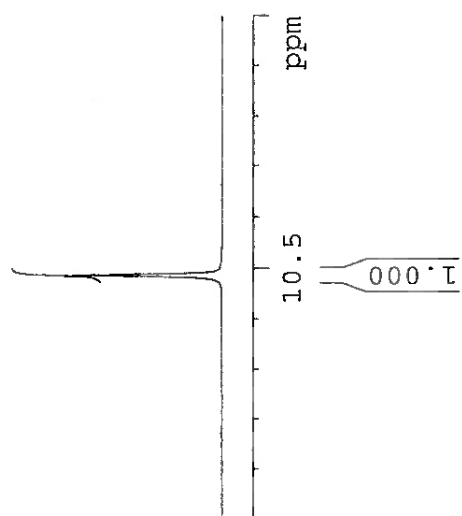
Compound 2

SB-591-38 (INH-USA)

DMSO

1e 21 mars 2018

RMN 1H 400 MHz



Thu Mar 22 15:26:23 EDT 2018

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2	12083.0	3558.246	8.8927	0.10	
3	12097.0	3554.709	8.8839	0.11	
4	12120.6	3548.738	8.8690	0.11	
5	12134.7	3545.190	8.8601	0.11	
6	13575.6	3181.167	7.9503	0.17	
7	13584.3	3178.978	7.9449	0.32	
8	13592.8	3176.821	7.9395	0.18	
9	14030.2	3066.312	7.6633	0.09	
10	14037.6	3064.449	7.6586	0.15	
11	14044.7	3062.654	7.6541	0.10	
12	14064.4	3057.690	7.6417	0.10	
13	14071.8	3055.813	7.6371	0.16	
14	14078.9	3054.027	7.6326	0.10	
15	14573.7	2929.026	7.3202	0.38	
16	14585.0	2926.155	7.3130	0.46	
17	14587.4	2925.560	7.3115	0.47	
18	14623.1	2916.535	7.2890	0.37	
19	15050.0	2808.697	7.0195	0.30	
20	15084.0	2800.094	6.9980	0.29	
21	15346.4	2733.821	6.8323	0.22	
22	15356.9	2731.161	6.8257	0.21	
23	15381.1	2725.040	6.8104	0.20	
24	15391.6	2722.395	6.8038	0.20	
25	17200.9	2265.302	5.6614	0.11	
26	17238.6	2255.785	5.6376	0.11	
27	17285.3	2244.000	5.6082	0.11	
28	17322.9	2234.484	5.5844	0.11	
29	19974.3	1564.680	3.9104	2.09	**
30	20266.7	1490.810	3.7258	2.33	**
31	20339.9	1472.308	3.6796	1.15	*
32	20381.8	1461.714	3.6531	1.15	*
33	20526.8	1425.083	3.5615	1.14	*
34	20568.8	1414.487	3.5351	1.14	*

Thu Mar 22 15:26:23 EDT 2018

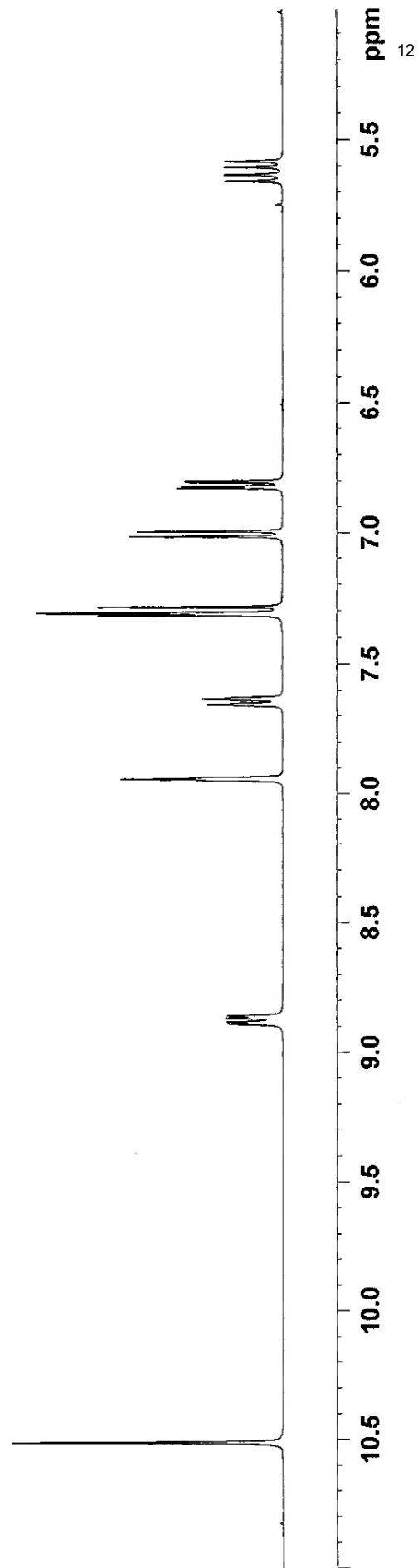
35	20837.0	1346.717	3.3657	0.07	
36	20915.7	1326.840	3.3160	12.50	*****
37	22194.2	1003.863	2.5088	0.20	
38	22201.0	1002.147	2.5046	0.42	
39	22208.1	1000.357	2.5001	0.59	*
40	22215.1	998.579	2.4956	0.44	
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SB-591-38 (INH-USA)

DMSO

1e 21 mars 2018

RMN 1H 400 MHz

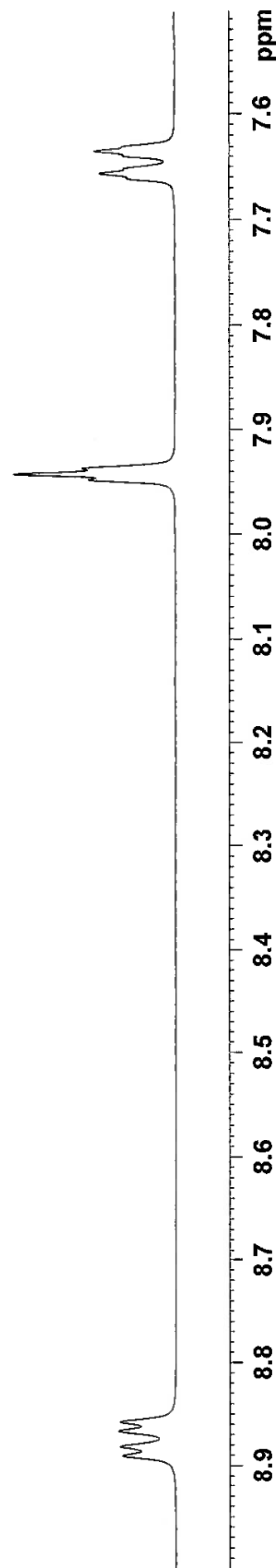


SB-591-38 (INH-USA)

DMSO

le 21 mars 2018

RMN 1H 400 MHz

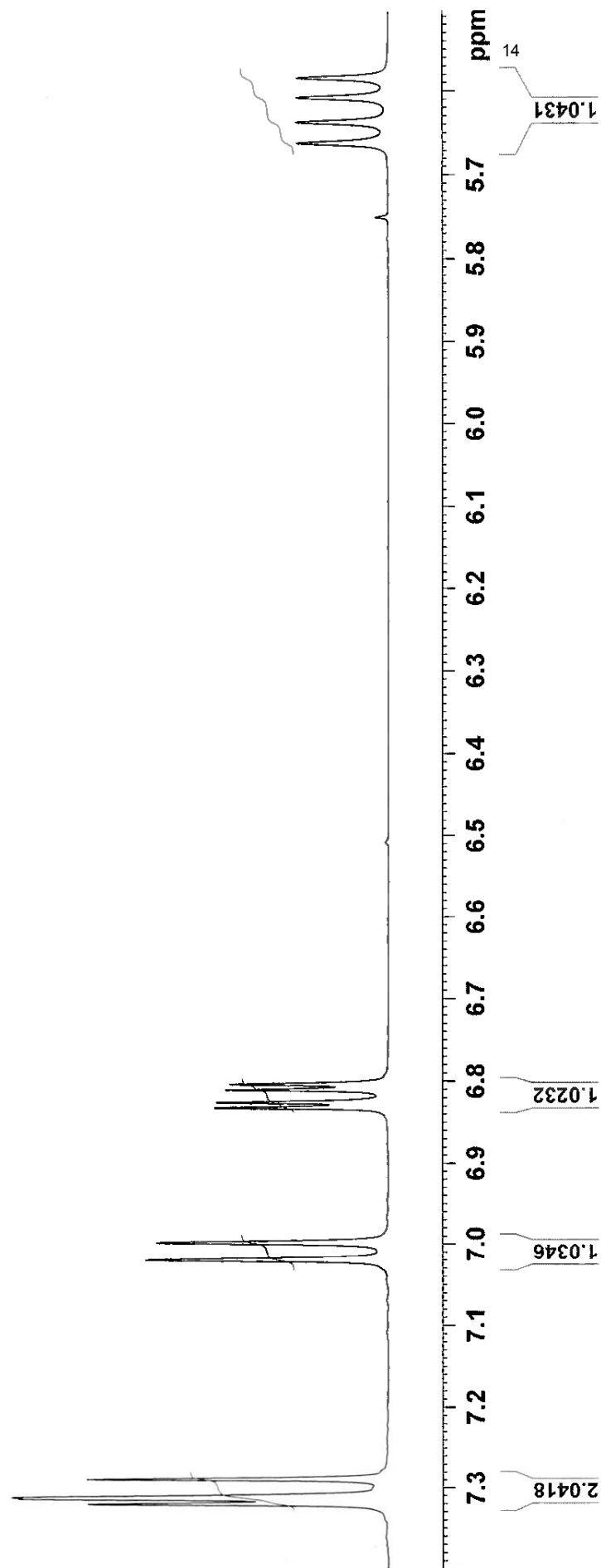


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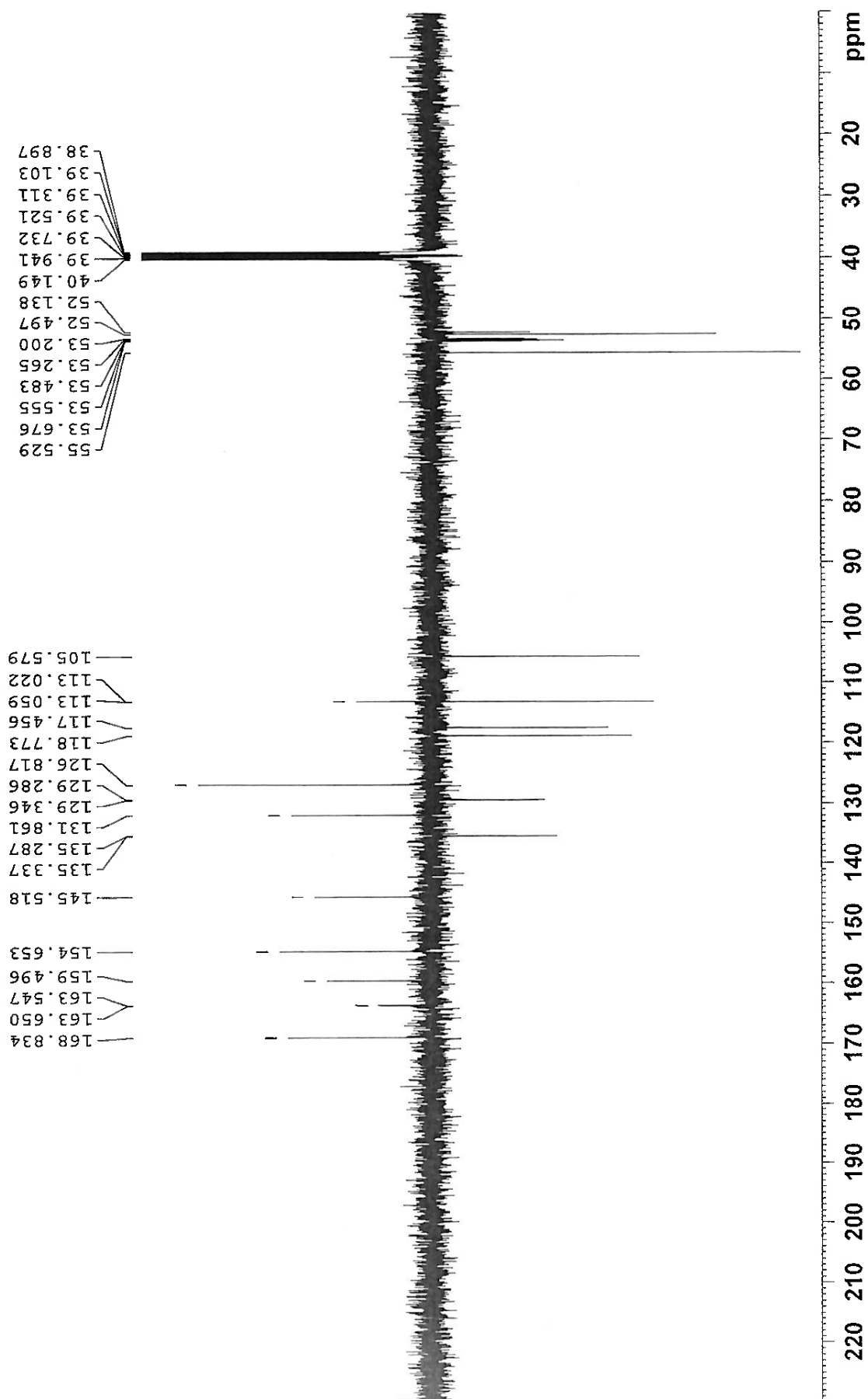
DMSO

1e 21 mars 2018

RMN 1H 400 MHz



SB-591-38-INH-US
Dans le DMSO
22 mars 2018
APT



Fri Mar 23 09:34:54 EDT 2018

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2	9364.4	16465.342	163.6505	0.15	
3	9375.8	16454.885	163.5466	0.15	
4	9819.2	16047.345	159.4960	0.29	
5	10349.2	15560.099	154.6532	0.42	
6	11349.2	14640.929	145.5175	0.32	
7	12463.5	13616.615	135.3368	-0.30	
8	12469.0	13611.583	135.2868	-0.34	
9	12844.0	13266.922	131.8612	0.38	
10	13119.3	13013.844	129.3458	-0.30	
11	13125.8	13007.837	129.2861	-0.31	
12	13396.1	12759.437	126.8172	0.63	*
13	14276.6	11950.070	118.7728	-0.56	
14	14420.7	11817.561	117.4558	-0.49	
15	14902.0	11375.158	113.0587	-0.60	
16	14906.0	11371.505	113.0224	0.20	
17	15720.7	10622.630	105.5793	-0.58	
18	21199.0	5586.979	55.5295	-1.00	
19	21401.8	5400.509	53.6761	-0.25	
20	21415.1	5388.306	53.5549	-0.30	
21	21423.0	5381.067	53.4829	-0.36	
22	21446.9	5359.116	53.2647	-0.30	
23	21454.0	5352.577	53.1998	-0.26	
24	21530.9	5281.874	52.4970	-0.78	
25	21570.2	5245.745	52.1379	-0.28	
26	22882.5	4039.471	40.1487	1.19	*
27	22905.2	4018.584	39.9411	3.57	****
28	22928.2	3997.503	39.7316	6.43	*****
29	22951.2	3976.299	39.5208	6.50	*****
30	22974.2	3955.153	39.3106	5.53	*****
31	22997.0	3934.218	39.1026	3.09	***
32	23019.5	3913.572	38.8974	1.14	*

(1/1)

(wdd) f1

