

Supporting Information for the Paper

Effect of a Bromo Substituent on the Glutathione Peroxidase Activity of a Pyridoxine-like Diselenide

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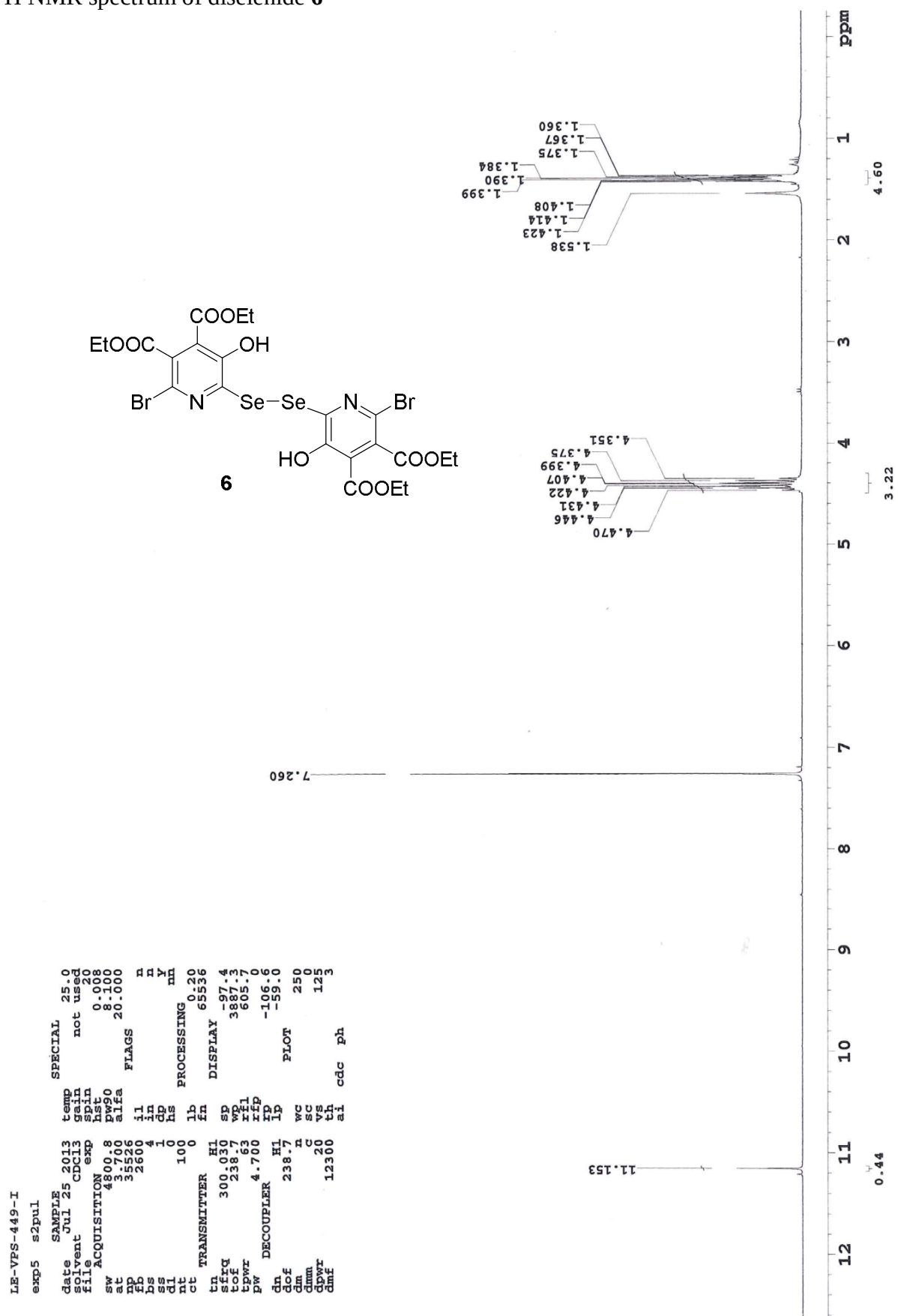
*E-mails: vijaypalchem@gmail.com; lars.engman@kemi.uu.se

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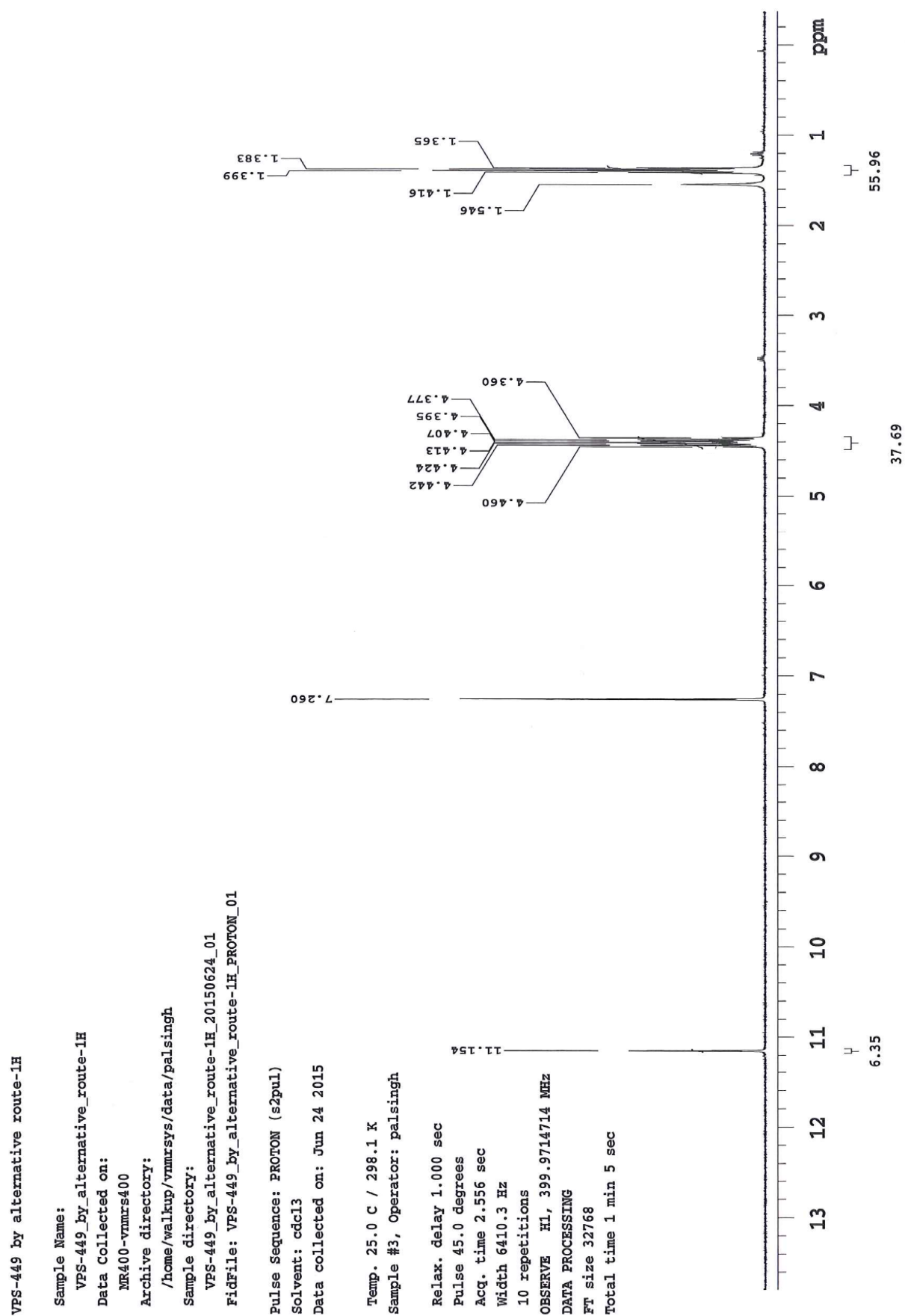
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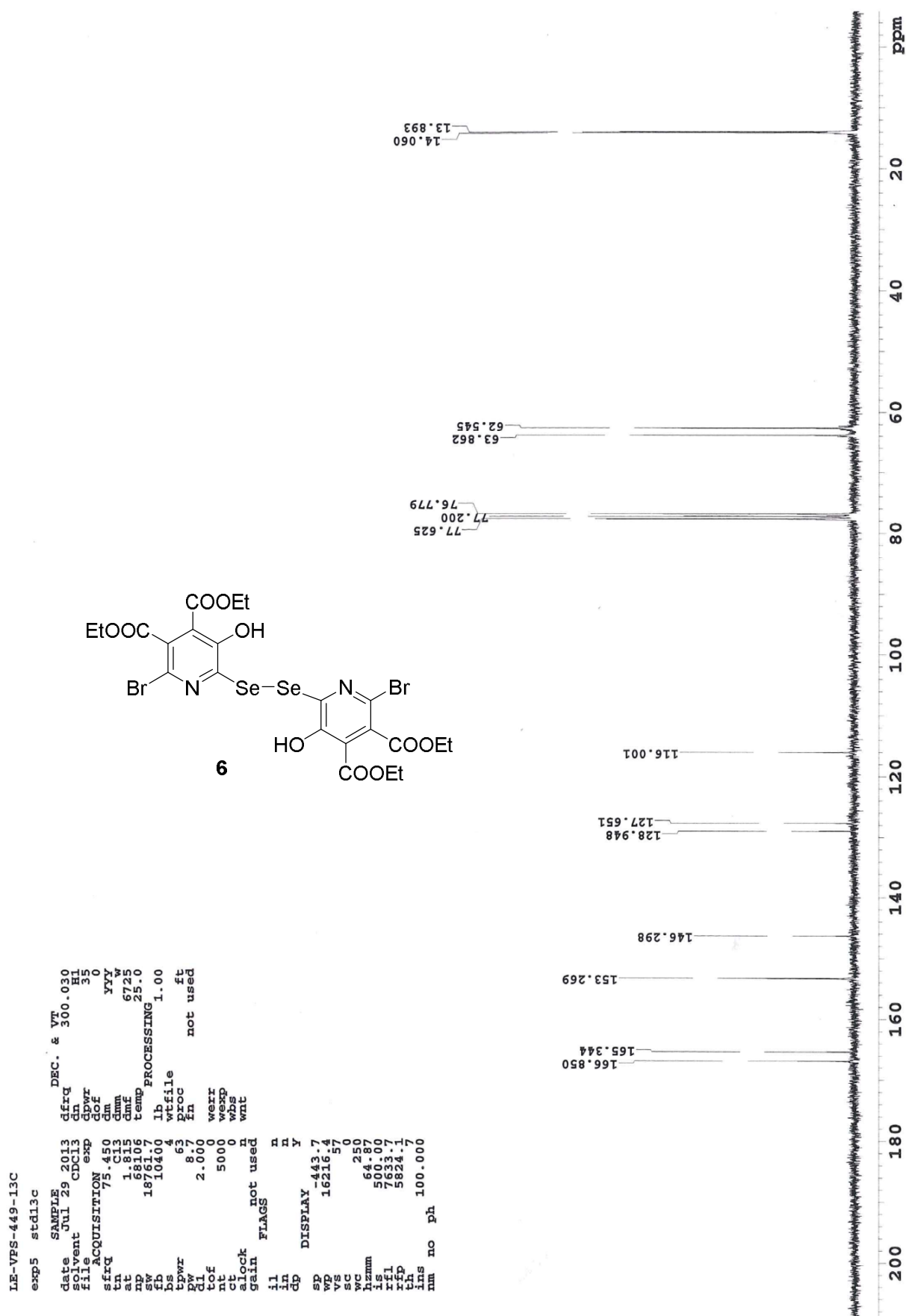
¹H NMR spectrum of diselenide **6**



^1H NMR spectrum of diselenide **6** via oxidation of selenide **8**



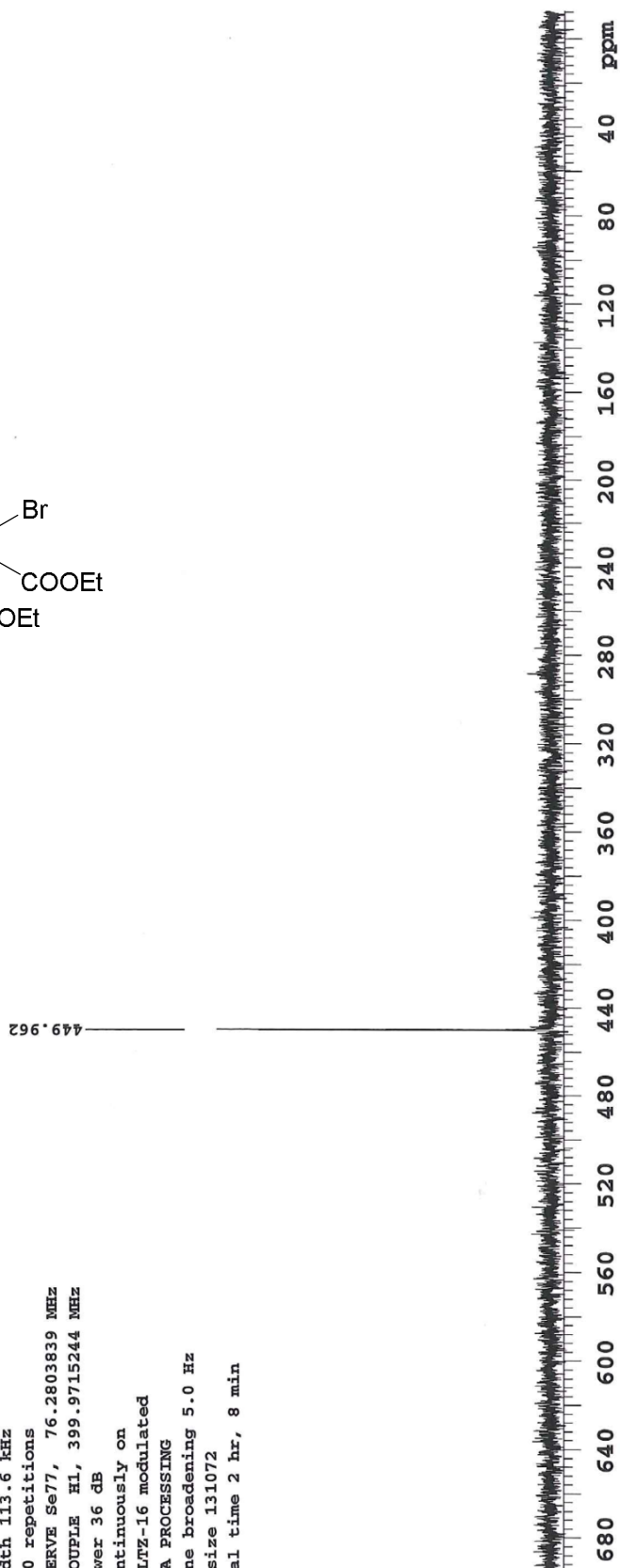
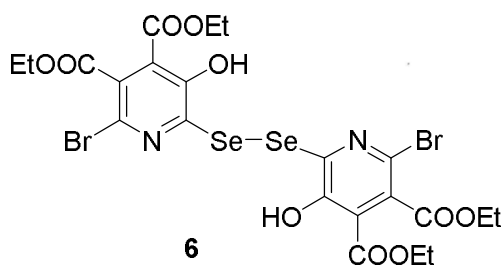
¹³C NMR spectrum of diselenide **6**



⁷⁷Se NMR spectrum of diselenide **6**



Sample Name:
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Data Collected on:
Agilent-NMR-vnmrs400
Archive directory:
Sample directory:
FidFile: SELENIUM77
Pulse Sequence: SELENIUM77 (s2pul)
Solvent: cdcl3
Data collected on: Aug 3 2013
Temp. 25.0 C / 298.1 K
Operator: palsingh
Relax. delay 2.000 sec
Pulse 24.2 degrees
Acq. time 0.577 sec
Width 113.6 kHz
920 repetitions
OBSERVE Se77, 76.2803839 MHz
DECOUPLE H1, 399.9715244 MHz
Power 36 dB
continuously on
WALTZ-16 modulated
DATA PROCESSING
Line broadening 5.0 Hz
FT size 131072
Total time 2 hr, 8 min



¹H NMR spectrum of 7

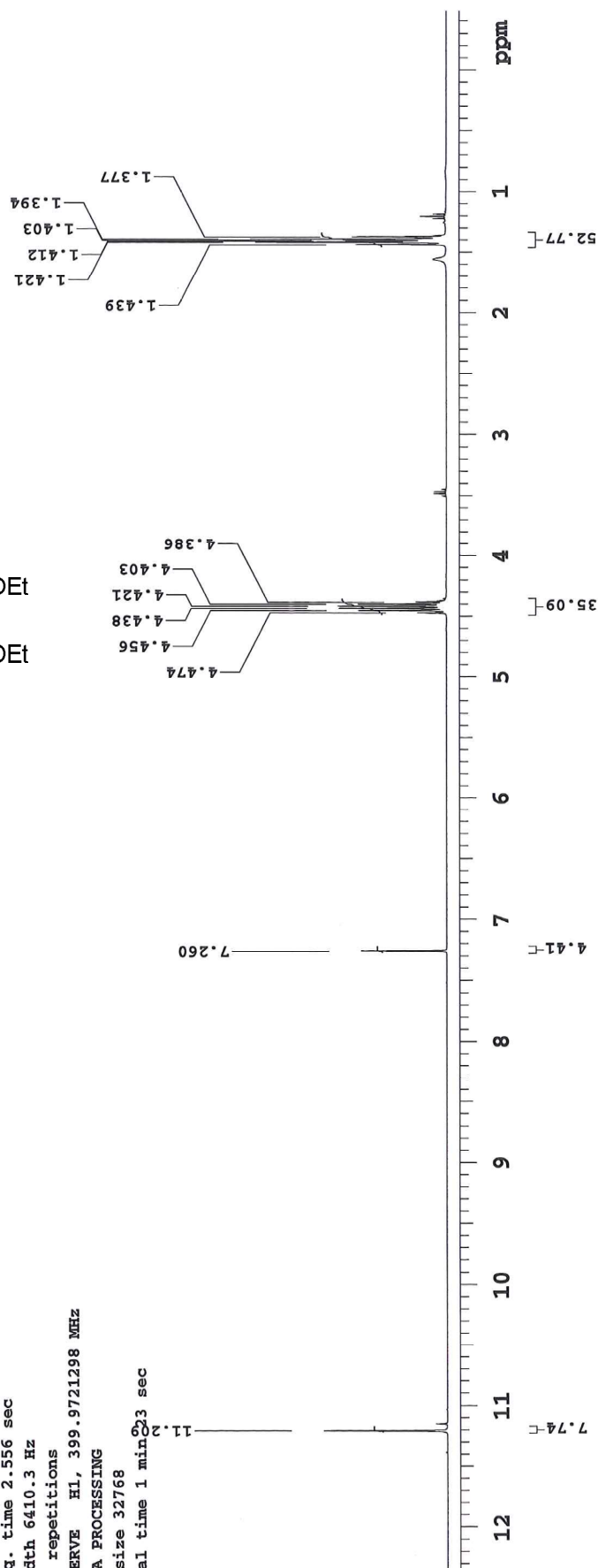
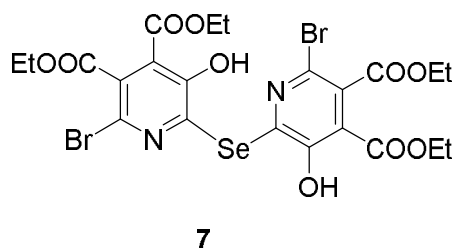
VPS-422-H1

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Archive directory:
/home/walkup/vnmrsys/data/palsingh
Sample directory:
VPS-422-H1_20140120_01
FidFile: VPS-422-H1_PROTON_01

Pulse Sequence: PROTON (s2pul)
Solvent: cdcl3
Data collected on: Jan 20 2014

Temp. 25.0 C / 298.1 K
Sample #2, Operator: palsingh

Relax. delay 1.000 sec
Pulse 45.0 degrees
Acq. time 2.556 sec
Width 6410.3 Hz
15 repetitions
OBSERVE H1, 399.9721298 MHz
DATA PROCESSING
Ft size 32768
Total time 1 min 23 sec



¹³C NMR spectrum of 7

VPS-422-13C

Sample Name:

VPS-422-13C

Data Collected on:

MR400-vnmrs400

Archive directory:

/home/walkup/vnmrsys/data/palsingh

Sample directory:

VPS-422-13C_20140121_01

FidFile: VPS-422-13C_CARBON_01

Pulse Sequence: CARBON (s2pul)

Solvent: cdcl3

Data collected on: Jan 21 2014

Temp. 25.0 C / 298.1 K

Sample #12, Operator: palsingh

Relax. delay 1.000 sec

Pulse 45.0 degrees

Acq. time 1.311 sec

Width 25000.0 Hz

400 repetitions

OBSERVE C13, 100.5730542 MHz

DECOUPLE H1, 399.9741302 MHz

Power 35 dB

continuously on

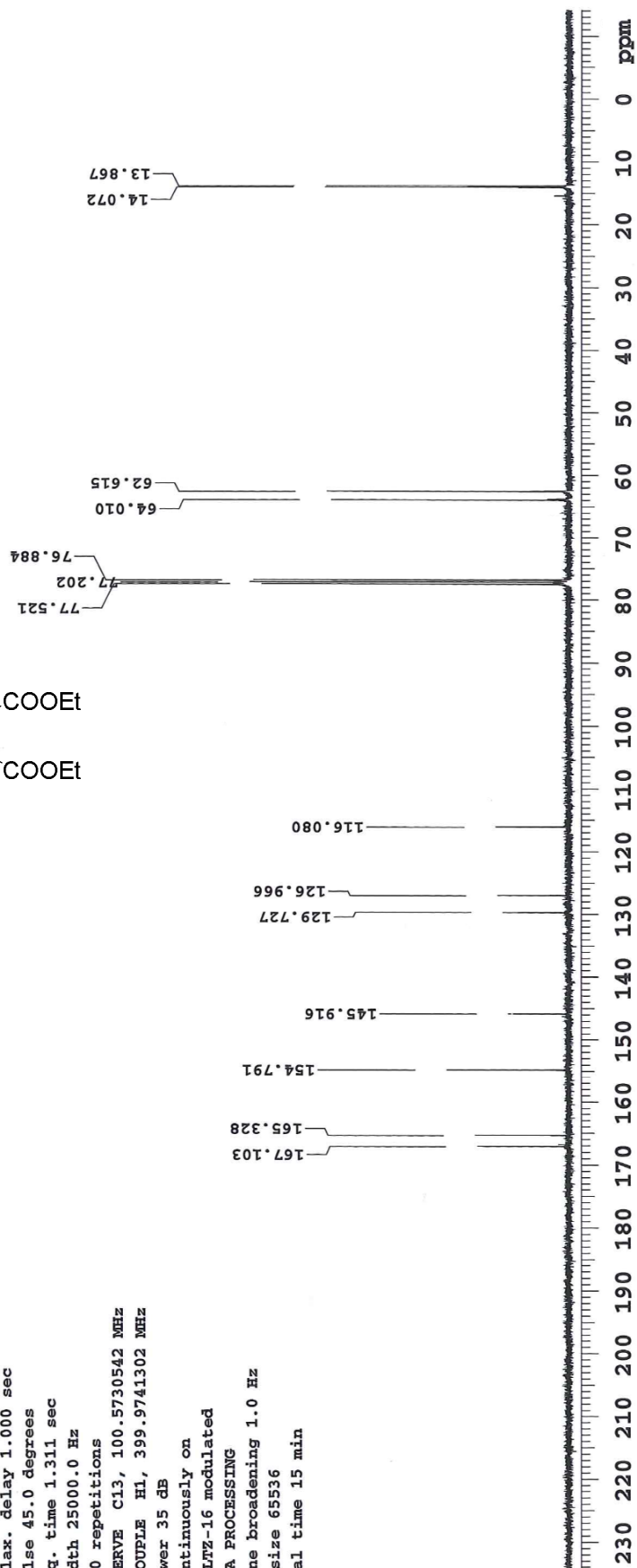
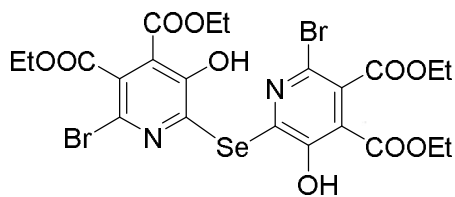
WALTZ-16 modulated

DATA PROCESSING

Line broadening 1.0 Hz

Ft size 65536

Total time 15 min



⁷⁷Se NMR spectrum of 7

VPS-422-Se77

Sample Name:

VPS-422-Se77

Data Collected on:

MR400-vnmrs400

Archive directory:

/home/walkup/vnmrsys/data/palsingh

Sample directory:

VPS-422-Se77_20140121_01

FidFile: current

Pulse Sequence: SELENIUM77 (s2pul)

Solvent: cdcl3

Data collected on: Jan 21 2014

Temp. 25.0 C / 298.1 K

Sample #12, Operator: palsingh

Relax. delay 1.000 sec

Pulse 45.0 degrees

Acq. time 1.887 sec

Width 138.9 kHz

340 repetitions

OBSERVE Se77, 76.2807368 MHz

DECOUPLE H1, 399.9741302 MHz

Power 35 dB

on during acquisition

off during delay

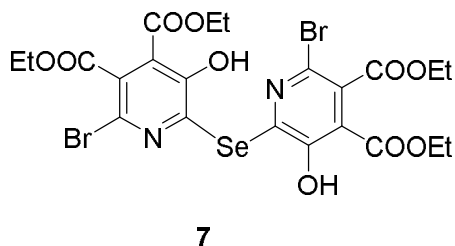
WALTZ-16 modulated

DATA PROCESSING

Line broadening 2.0 Hz

FT size 524288

Total time 24 min

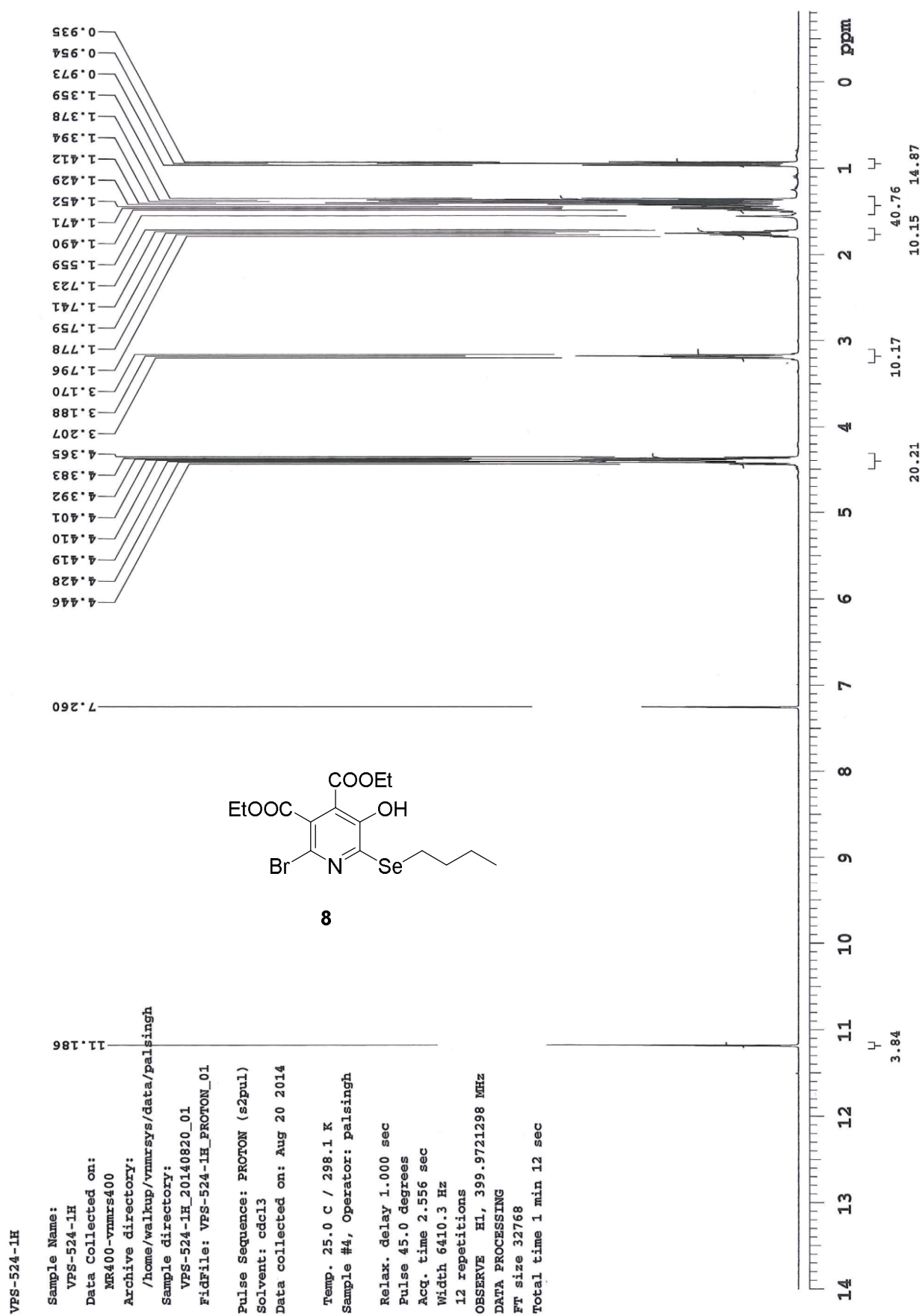


7

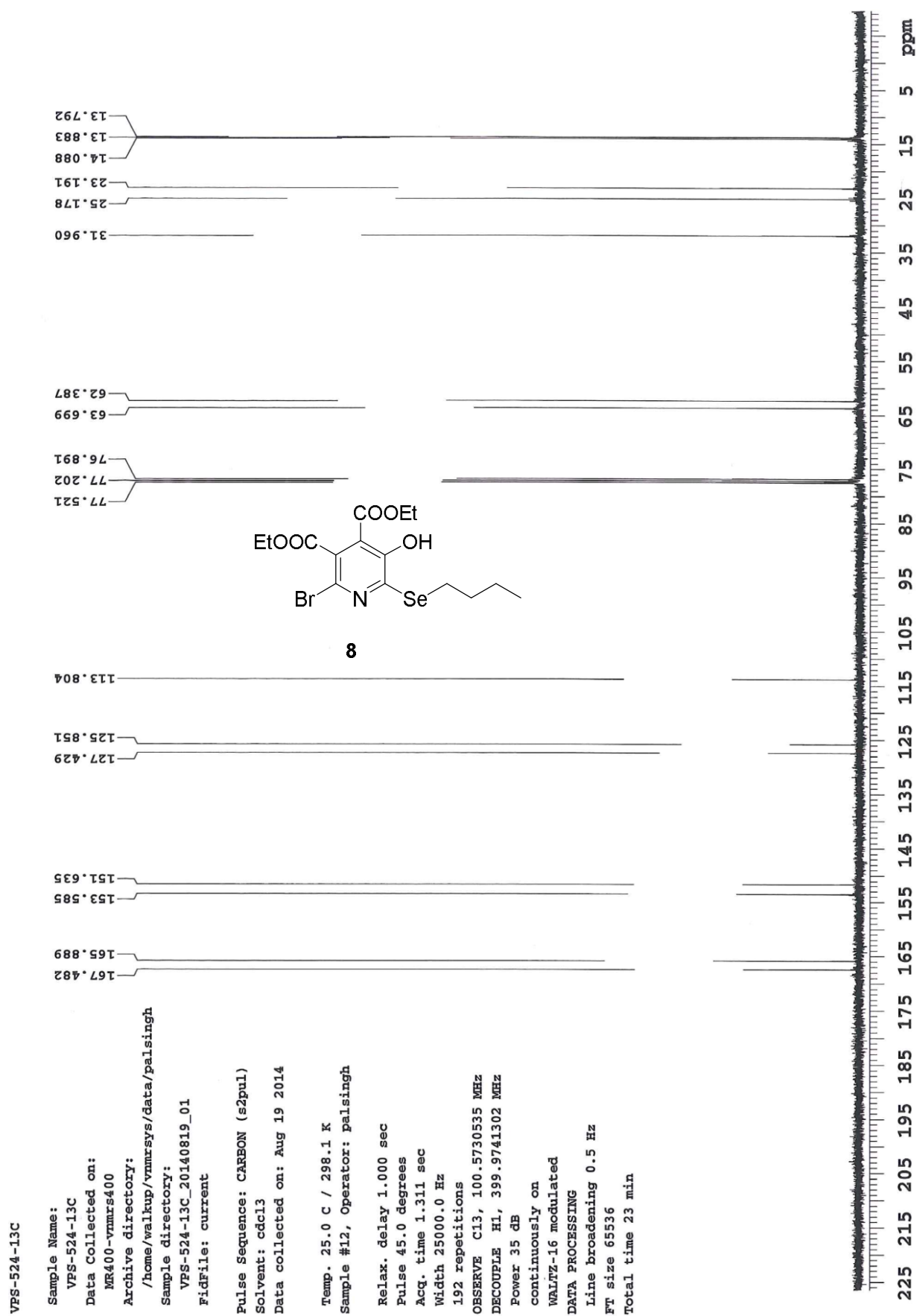
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1150 1050 950 850 750 650 500 400 300 250 150 100 50 ppm

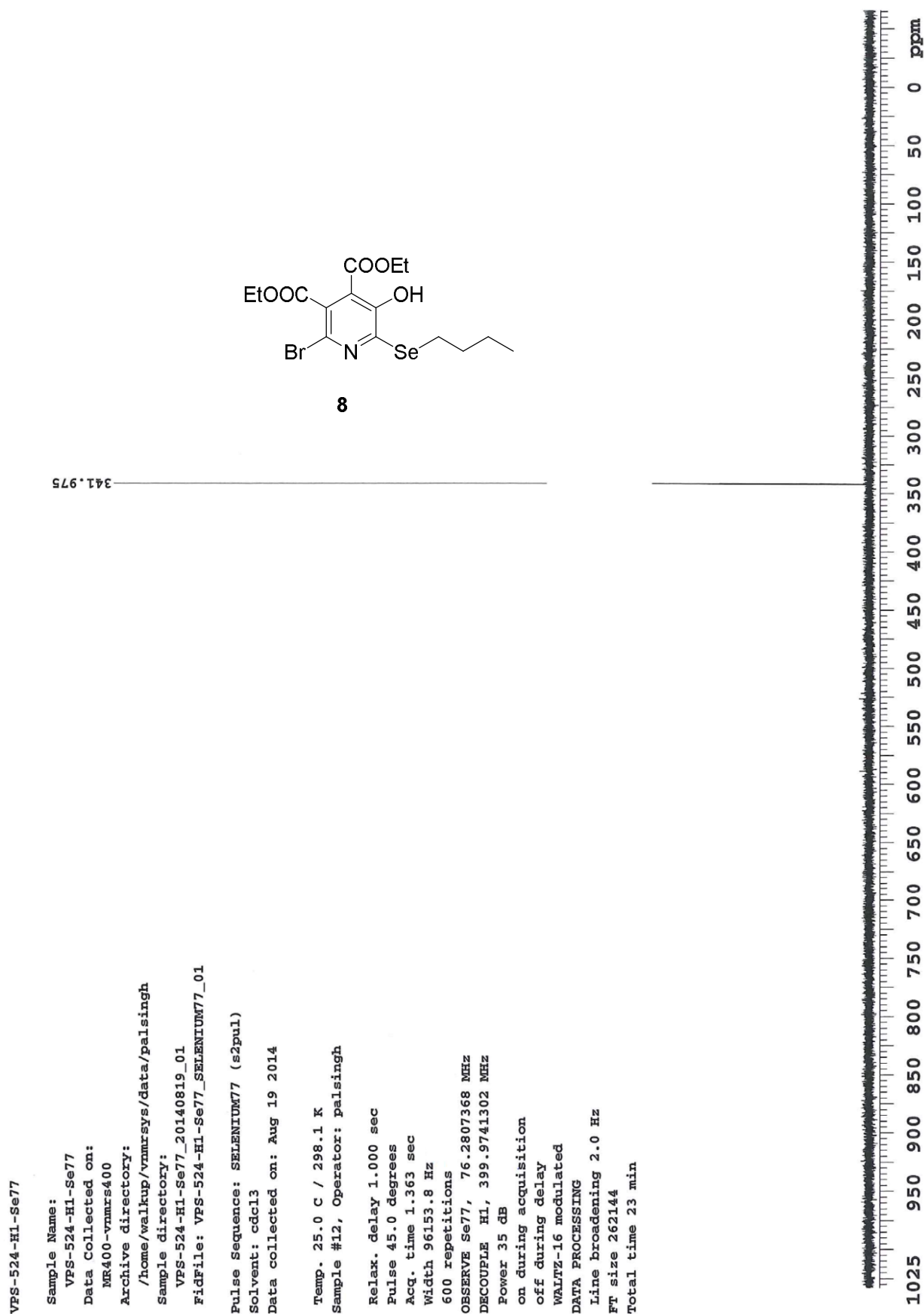
¹H NMR spectrum of **8**



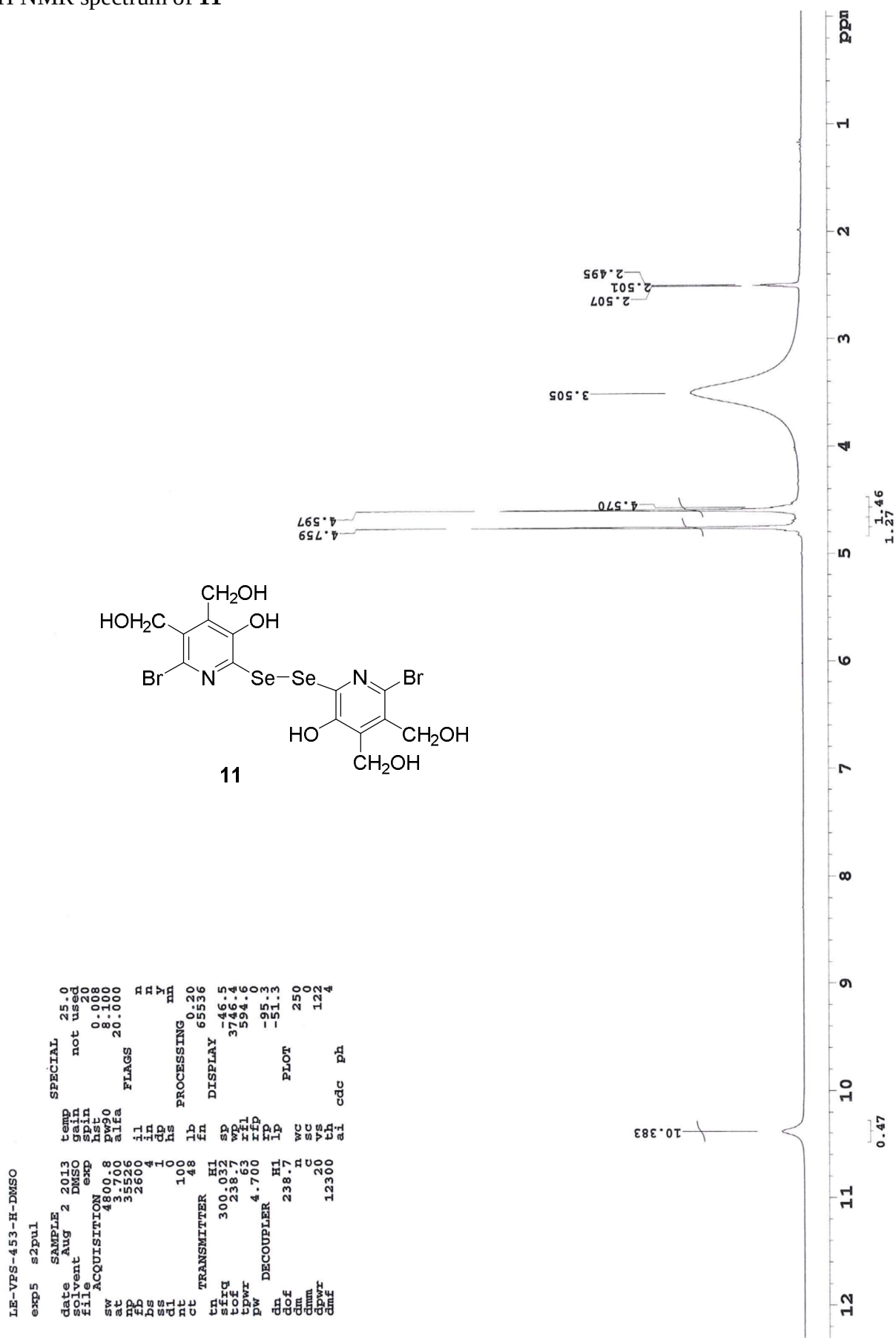
¹³C NMR spectrum of **8**



⁷⁷Se NMR spectrum of **8**



¹H NMR spectrum of **11**



Chemical structure of compound 11:

OCC1=CC=C(C(=O)O)N=C1[Se][Se]C2=CC=C(C(=O)O)N=C2Br

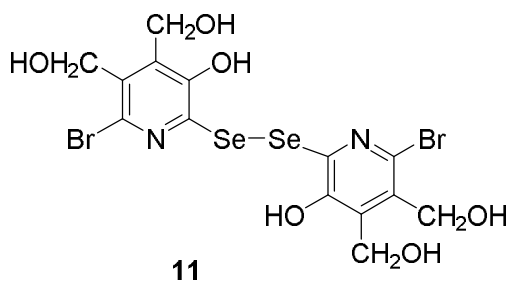
¹³C NMR spectrum (DMSO-d₆) data:

Chemical Shift (ppm)
150.092
141.448
136.580
134.557
133.400
59.646
56.200

⁷⁷Se NMR spectrum of **11**



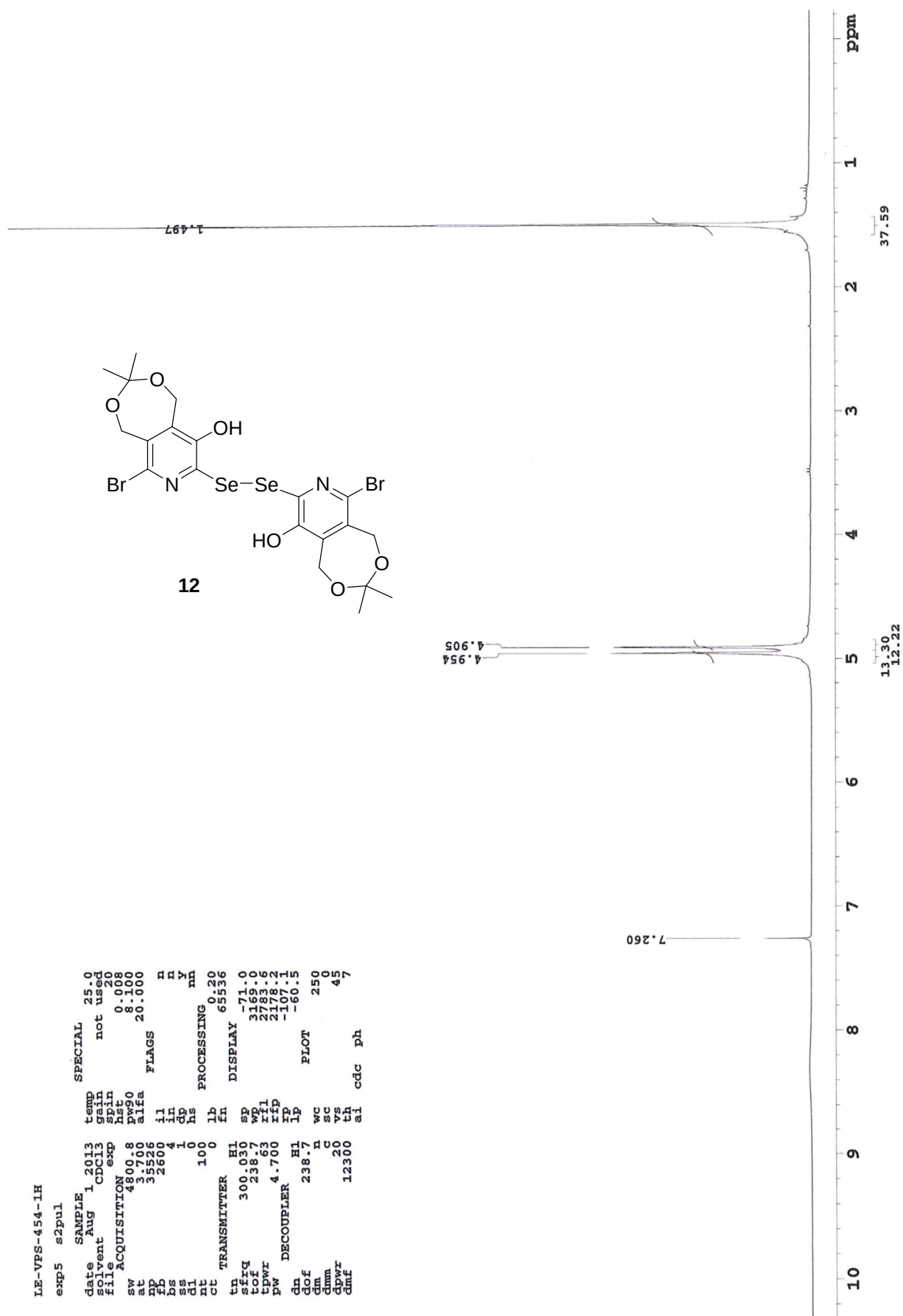
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Archive directory:
Sample directory:
FidFile: SELENIUM77
Pulse Sequence: SELENIUM77 (s2pul)
Solvent: dmsd
Data collected on: Aug 3 2013
Temp. 25.0 C / 298.1 K
Operator: palsingh
Relax. delay 2.000 sec
Pulse 24.2 degrees
Acq. time 0.577 sec
Width 113.6 kHz
4000 repetitions
OBSERVE Se77, 76.2807463 MHz
DECOUPLE H1, 399.9734243 MHz
Power 36 dB
continuously on
WALTZ-16 modulated
DATA PROCESSING
Line broadening 5.0 Hz
FT size 131072
Total time 2 hr, 51 min



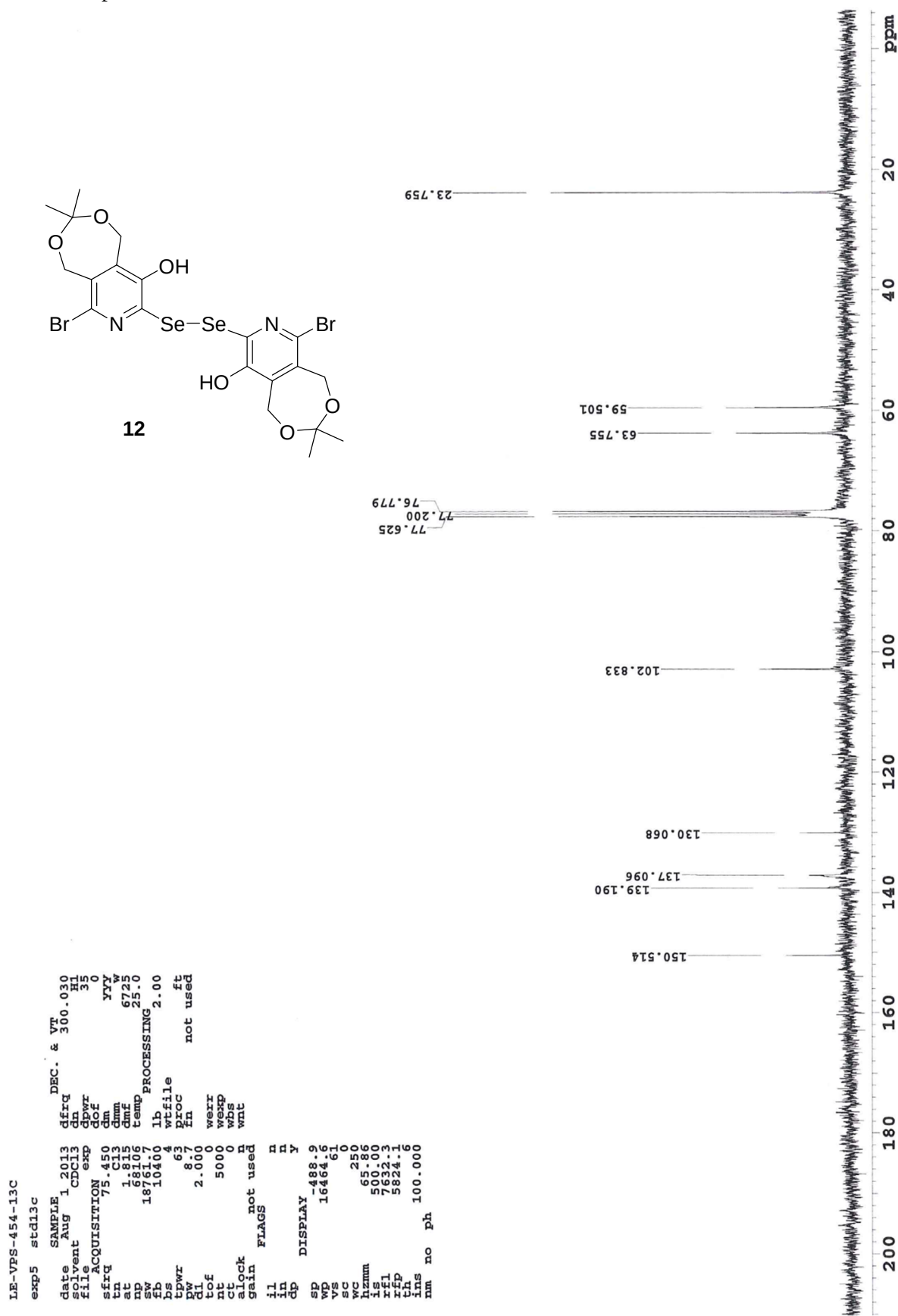
407.886



¹H NMR spectrum of **12**



¹³C NMR spectrum of **12**



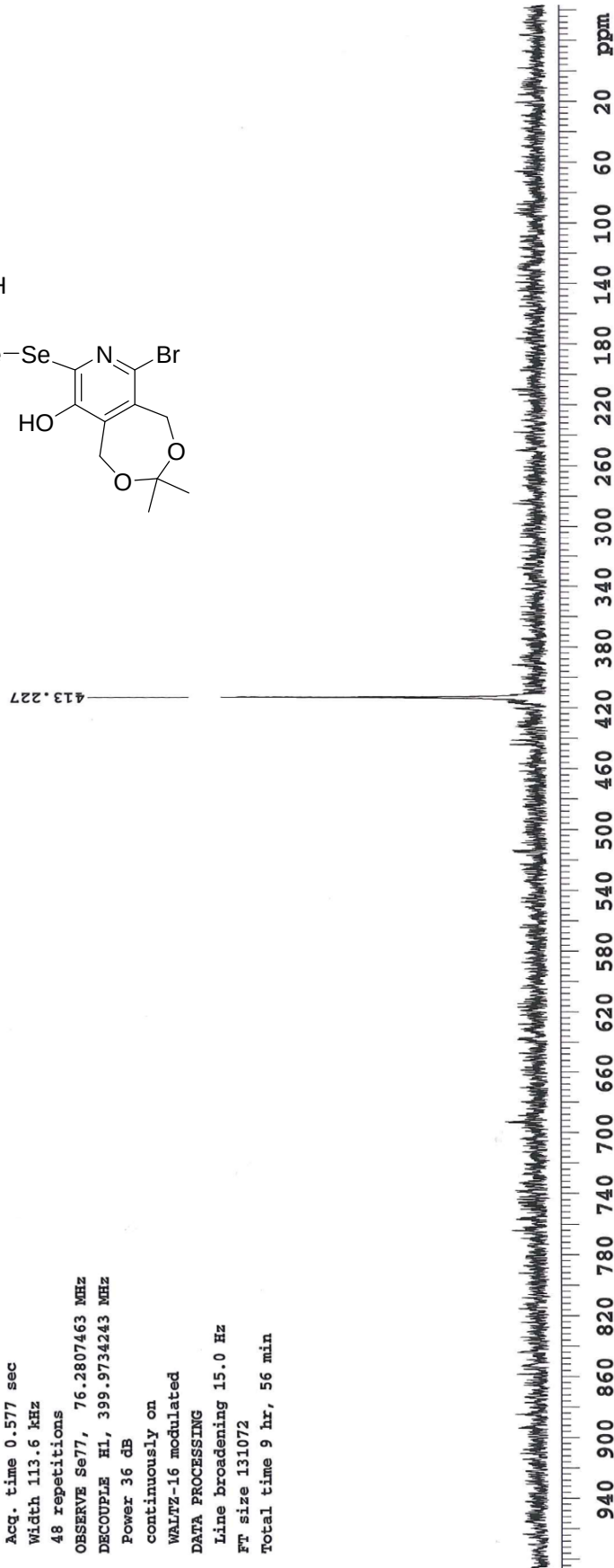
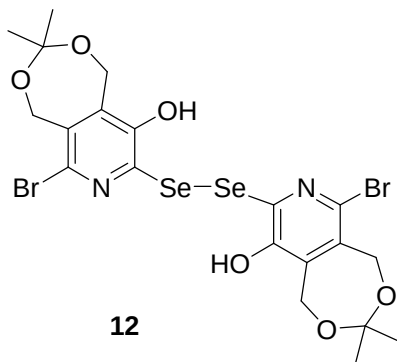
⁷⁷Se NMR spectrum of **12**

Sample Name:
VPS454-Se
Data Collected on:
Agilent-NMR-vmrs400
Archive directory:
/home/walkup/vmrsys/data/palsingh
Sample directory:
VPS454-Se_20130812_01
FidFile: current

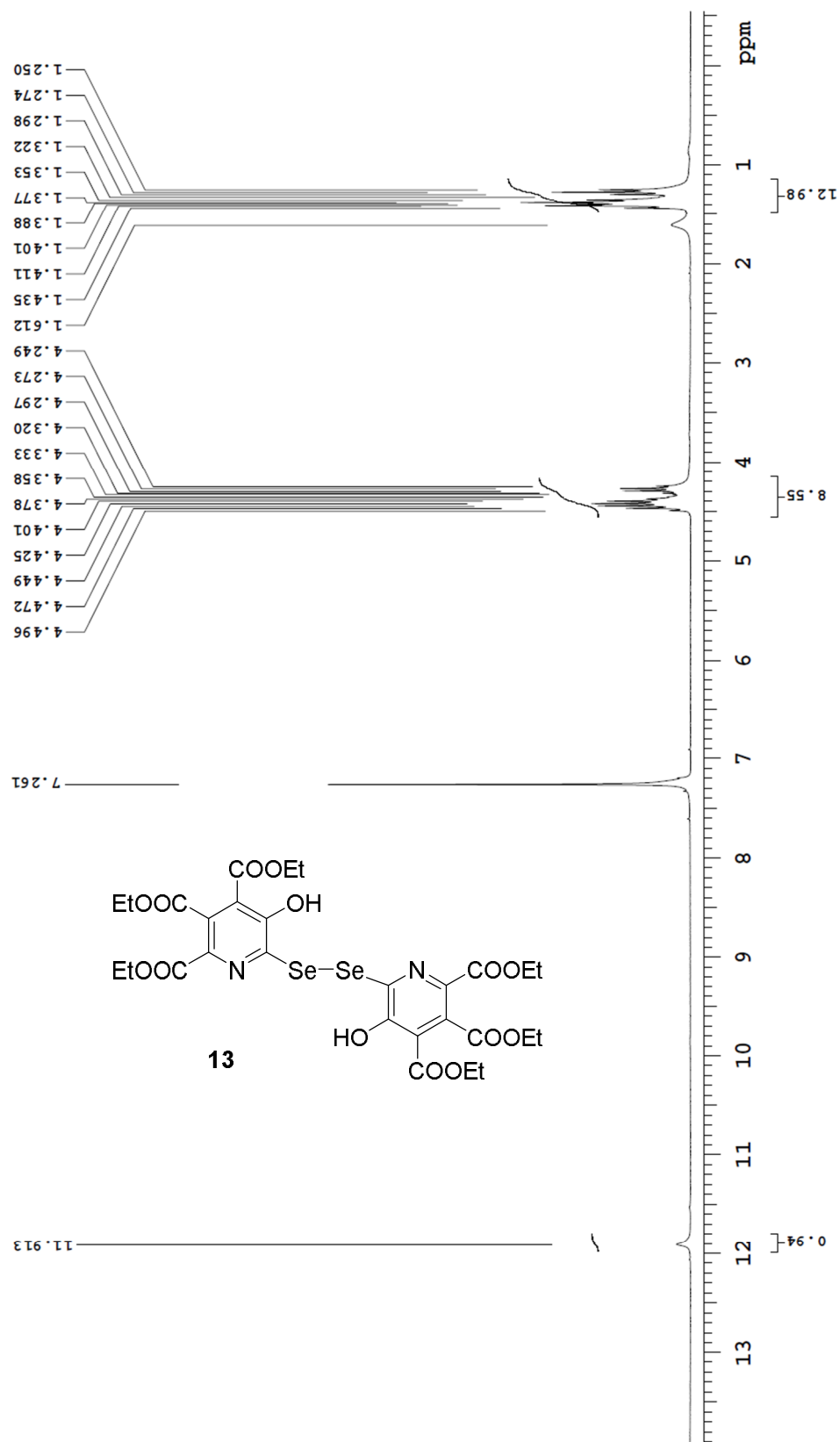
Pulse Sequence: SELENIUM77 (s2pul)
Solvent: dmsd
Data collected on: Aug 12 2013

Temp. 25.0 C / 298.1 K
Sample #11, Operator: palsingh

Relax. delay 3.000 sec
Pulse 24.2 degrees
Acq. time 0.577 sec
Width 113.6 kHz
48 repetitions
OBSERVE Se77, 76.2807463 MHz
DECOUPLE H1, 399.9734243 MHz
Power 36 dB
continuously on
WALTZ-16 modulated
DATA PROCESSING
Line broadening 15.0 Hz
Ft size 131072
Total time 9 hr, 56 min

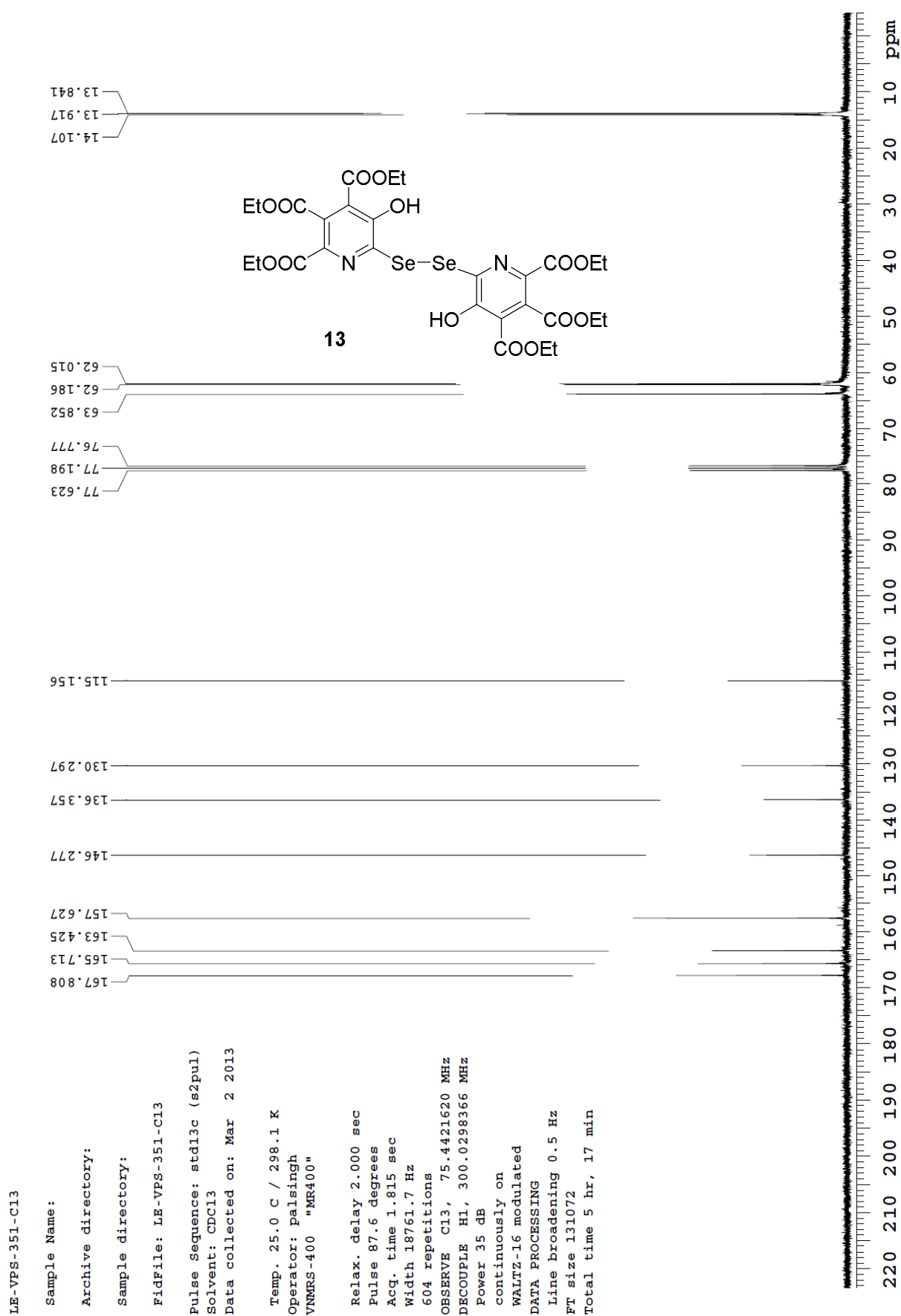


¹H NMR spectrum of **13**

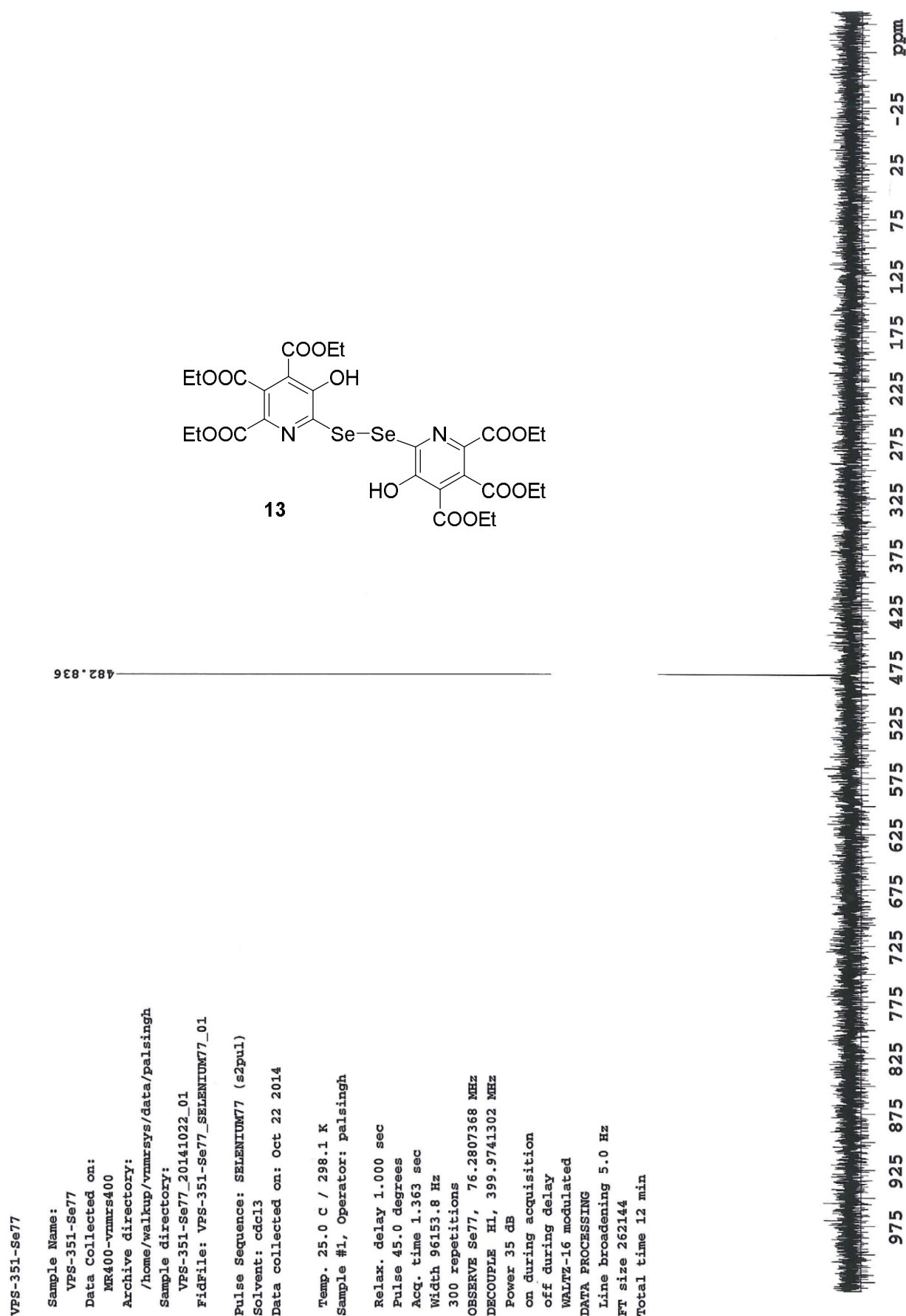


PULSE SEQUENCE	DATA PROCESSING	OBSERVE	
Pulse 52.2 degrees Acq. time 3.700 sec Width 4800.8 Hz 100 repetitions	LE-VPS-351 Solvent: CDCl ₃ Temp. 25.0 C / 298.1 K Operator: palsingh File: LE-VPS-351 UNITYplus-400 "MR400"	H1, 300.0282804 Line broadening 1.0 Hz FT size 65536 Total time 6 minutes	

¹³C NMR spectrum of **13**

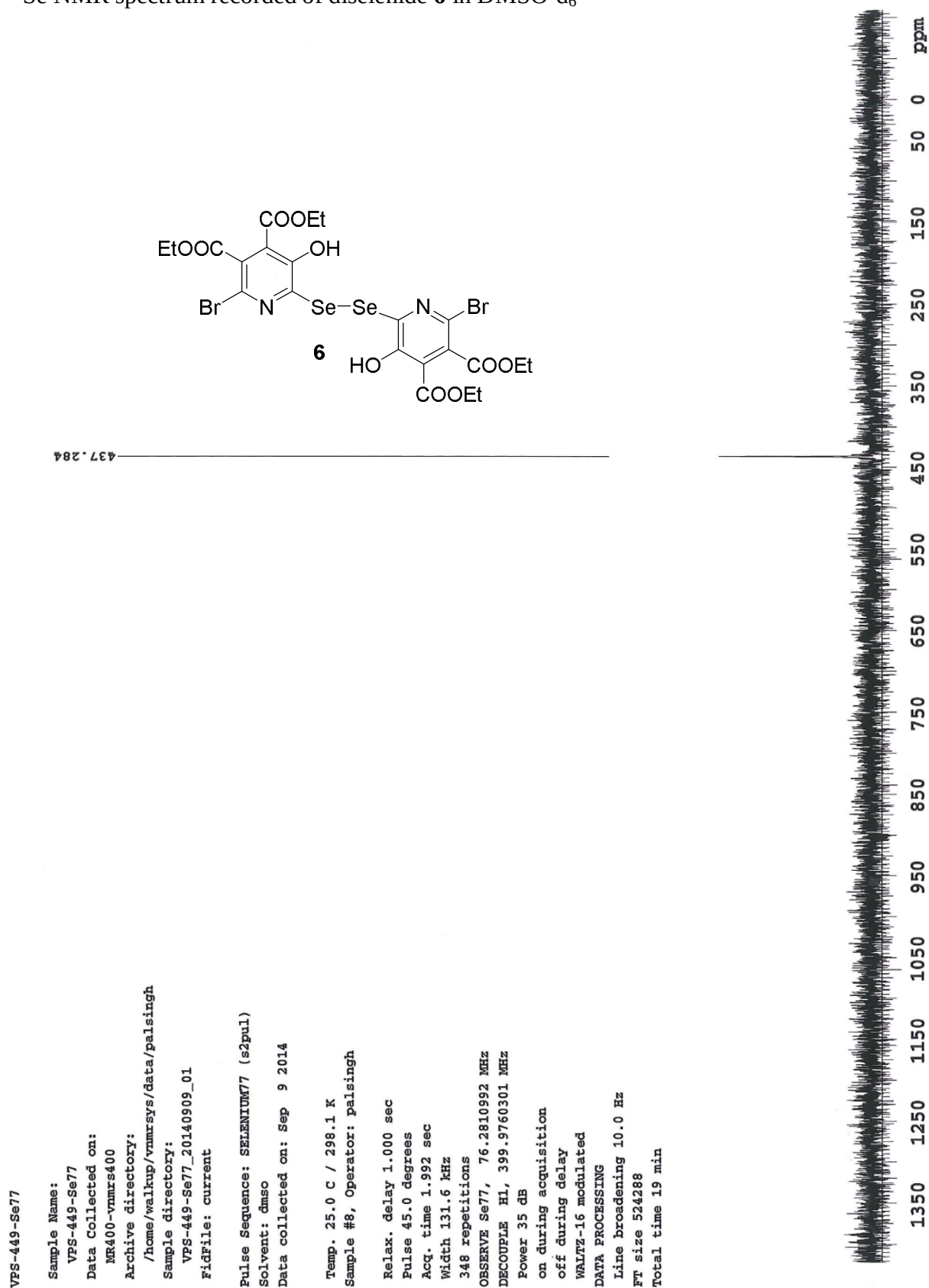


⁷⁷Se NMR spectrum of 13



⁷⁷Se NMR Experimental Data for the Mechanistic Studies with Diselenide 6

⁷⁷Se NMR spectrum recorded of diselenide **6** in DMSO-d₆



⁷⁷Se NMR spectrum recorded after the addition of 1 equiv of GSH (dissolved in 200 μ L H₂O) to a solution of diselenide **6** in DMSO-d₆.



VPS-Se77-449+GSH(1 eq)

Sample Name:
VPS-Se77-449_GSH_1_eq
Data Collected on:
MR400-vnmrs400
Archive directory:
/home/walkup/vnmrsys/data/palsingh
Sample directory:
VPS-Se77-449_GSH_1_eq_20140731_01
FidFile: VPS-Se77-449_GSH_1_eq_SELENIUM77_01

Pulse Sequence: SELENIUM77 (s2pul)
Solvent: dmso
Data collected on: Jul 31 2014

Temp. 25.0 C / 298.1 K
Sample #5, Operator: palsingh

Relax. delay 1.000 sec
Pulse 45.0 degrees
Acq. time 1.363 sec
Width 96153.8 Hz
640 repetitions

OBSERVE Se77, 76.2810992 MHz
DECOUPLE H1, 399.9760301 MHz
Power 35 dB

on during acquisition
off during delay

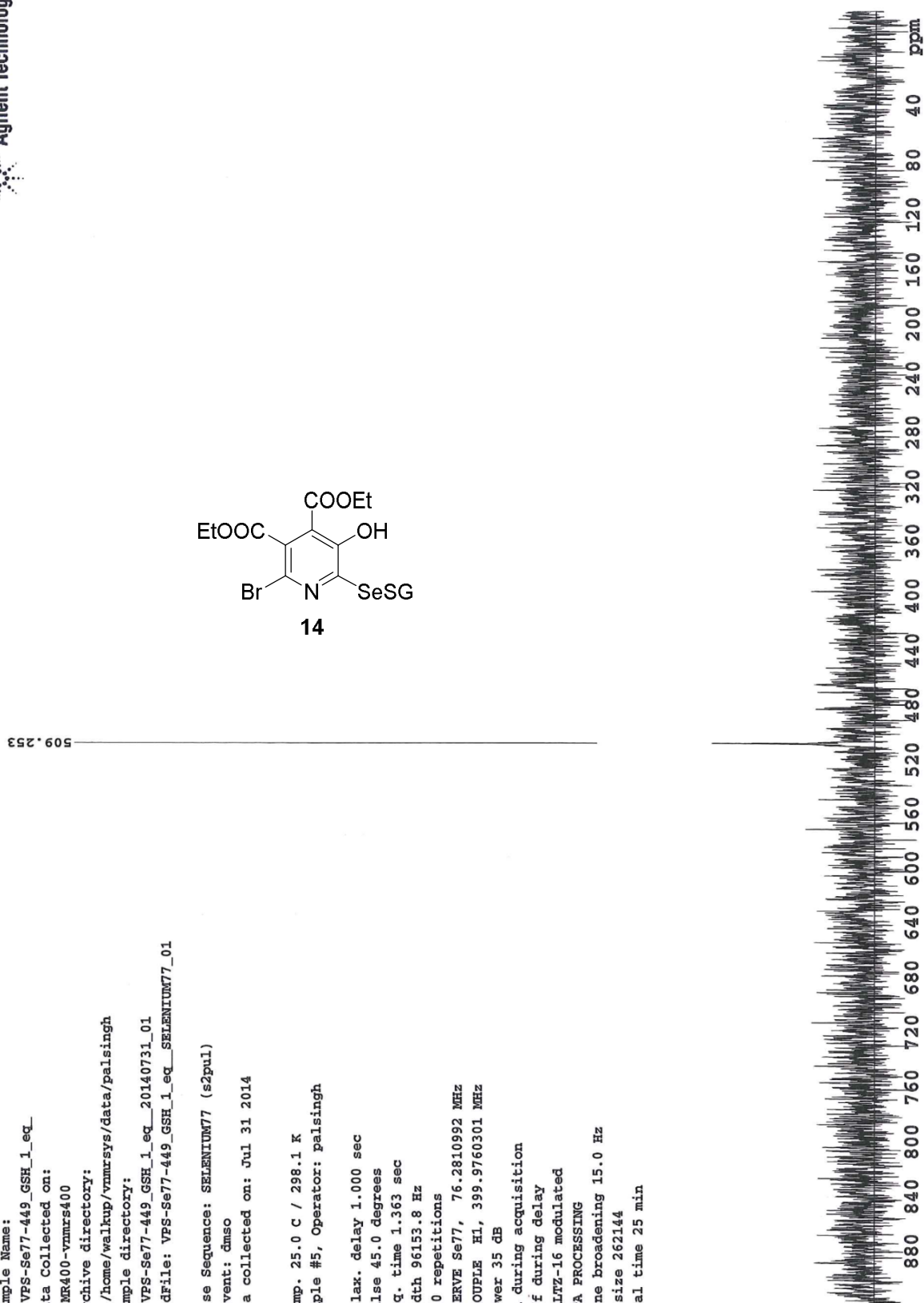
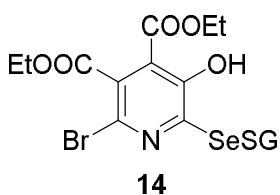
WALTZ-16 modulated

DATA PROCESSING

Line broadening 15.0 Hz

FT size 262144

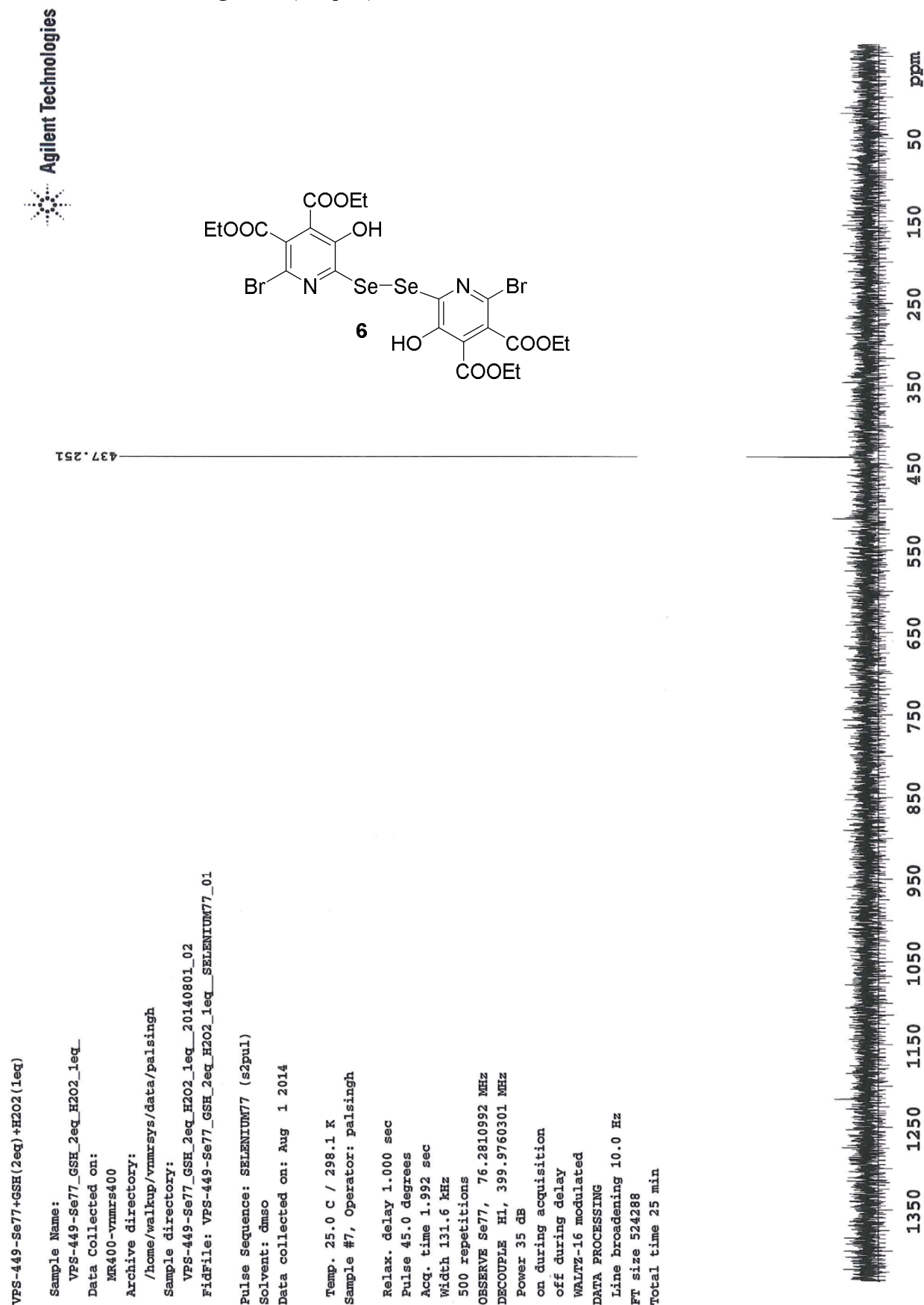
Total time 25 min



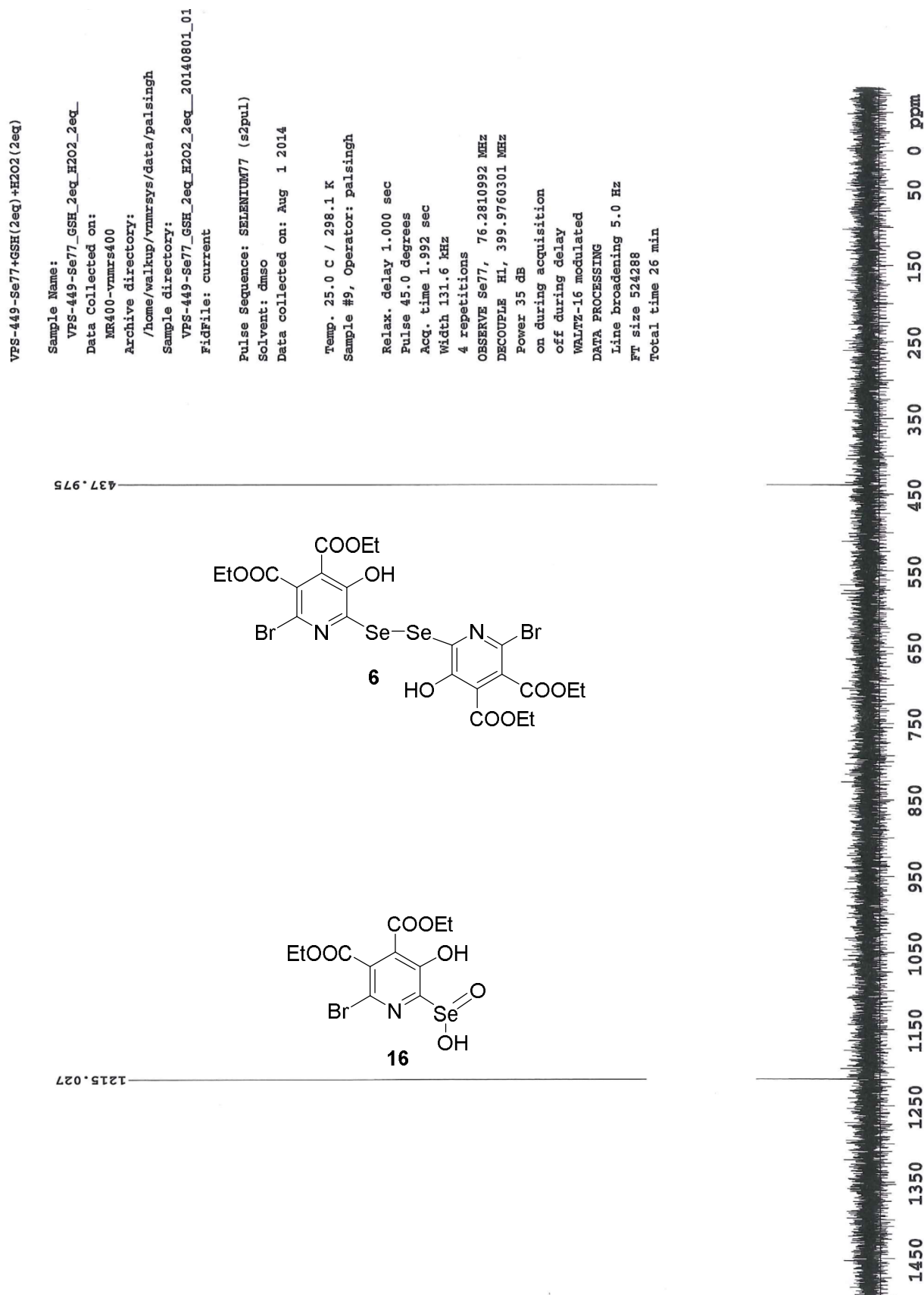
⁷⁷Se NMR spectrum recorded after the addition of 1 more equiv of GSH (2 equiv) to the above mixture in DMSO-d₆.



⁷⁷Se NMR spectrum recorded after the addition of 1 equiv of H₂O₂ to the above mixture of diselenide **6** containing GSH (2 equiv) in DMSO-d₆.



⁷⁷Se NMR spectrum recorded after the addition of 1 more equiv of H₂O₂ (2 equiv) to the above mixture of diselenide **6** in DMSO-d₆.



^{77}Se NMR spectrum recorded after the addition of 2 more equiv of H_2O_2 (4 equiv) to the above mixture of diselenide **6** in DMSO-d_6 .

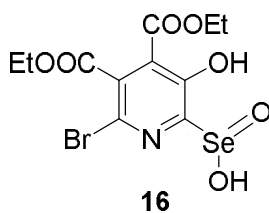
VPS-449-Se77_GSH_2eq_H2O2_4eq_
 Sample Name:
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 Data Collected on:
 MR400-vnmrs400
 Archive directory:
 /home/walkup/vnmrsys/data/palsingh/
 Sample directory:
 VPS-449-Se77_GSH_2eq_H2O2_4eq_20140801_01
 FidFile: current

Pulse Sequence: SELENIUM77 (s2pul)
 Solvent: dmsd
 Data collected on: Aug 1 2014

Temp. 25.0 C / 298.1 K
 Sample #9, Operator: palsingh

Relax. delay 1.000 sec
 Pulse 45.0 degrees
 Acq. time 1.992 sec
 Width 131.6 kHz
 4 repetitions

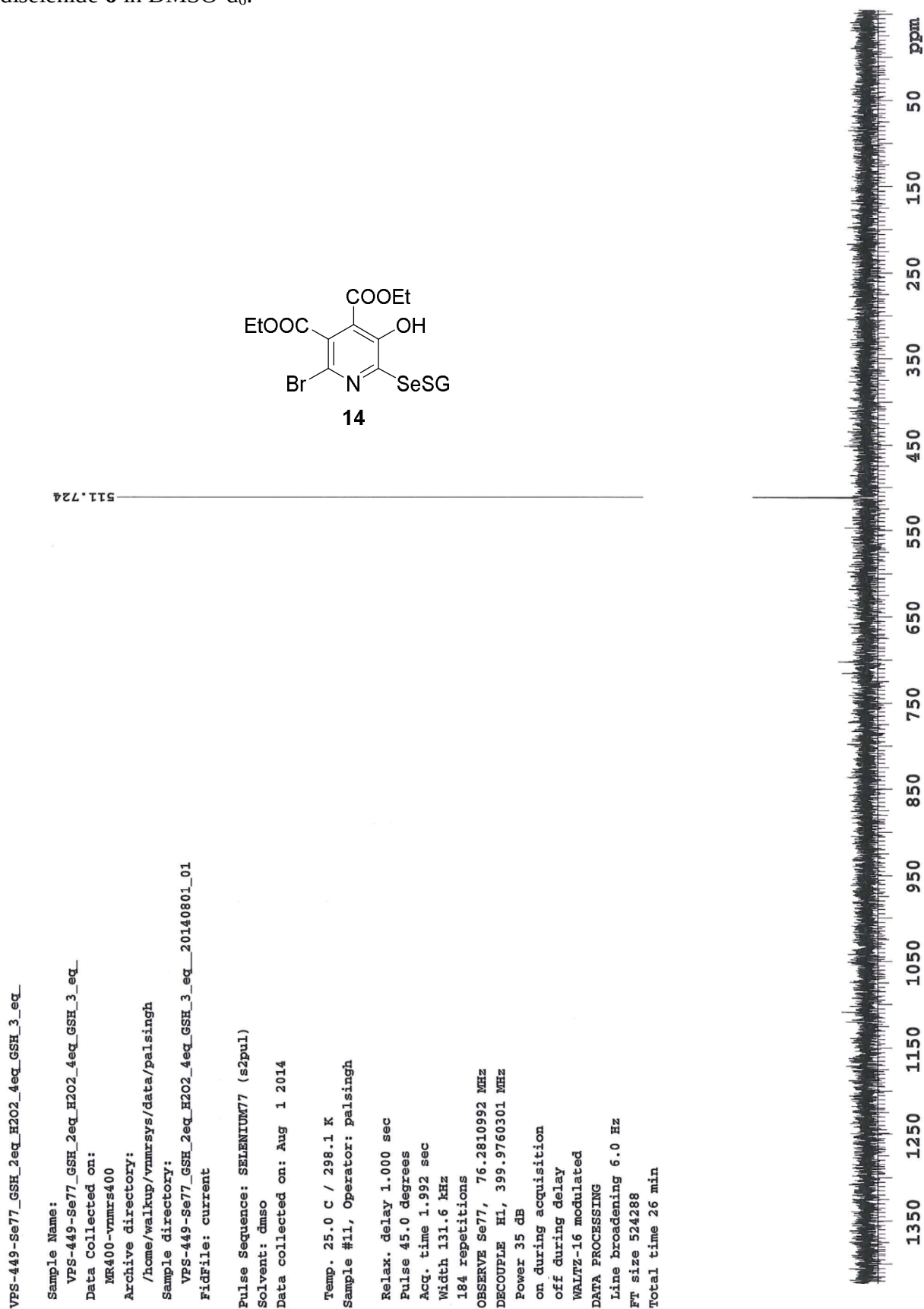
OBSERVE Se77, 76.281092 MHz
 DECOUPLE H1, 399.9760301 MHz
 Power 35 dB
 on during acquisition
 off during delay
 WALTZ-16 modulated
 DATA PROCESSING
 Line broadening 2.0 Hz
 FT size 524288
 Total time 26 min



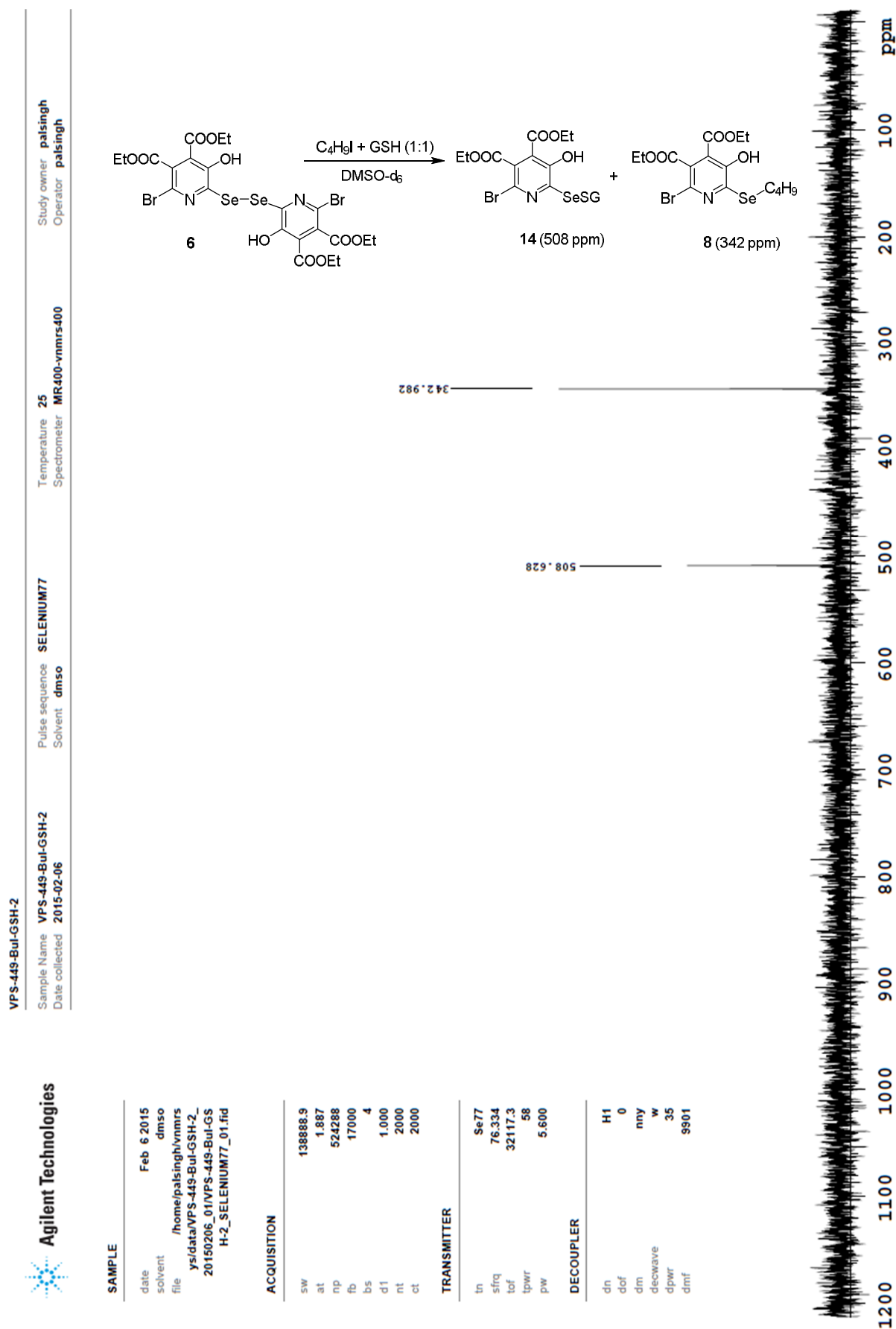
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^{77}Se NMR spectrum recorded after the addition of 3 equiv of GSH to the above mixture of diselenide **6** in DMSO-d_6 .



^{77}Se NMR spectrum of diselenide **6** with a mixture of iodobutane and GSH in DMSO- d_6



⁷⁷Se NMR spectrum recorded for diselenide **6** with 4 equiv of GSH in DMSO-d₆.

VPS-449-4eqGSH-Se77

Agilent Technologies

Sample Name: VPS-449-4eqGSH-Se77
Date collected: 2015-02-13

Pulse sequence: SELENIUM77
Solvent: dmsd

Temperature: 25
Spectrometer: MR400-vnmrs400

Study owner: palsingh
Operator: palsingh

SAMPLE

date: Feb 13 2015
solvent: dmsd
file: /home/palsingh/vnm
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-Se77_20150213_01\VPS-44
9-4eqGSH-Se77_SELENIUM77
_01.fid

ACQUISITION

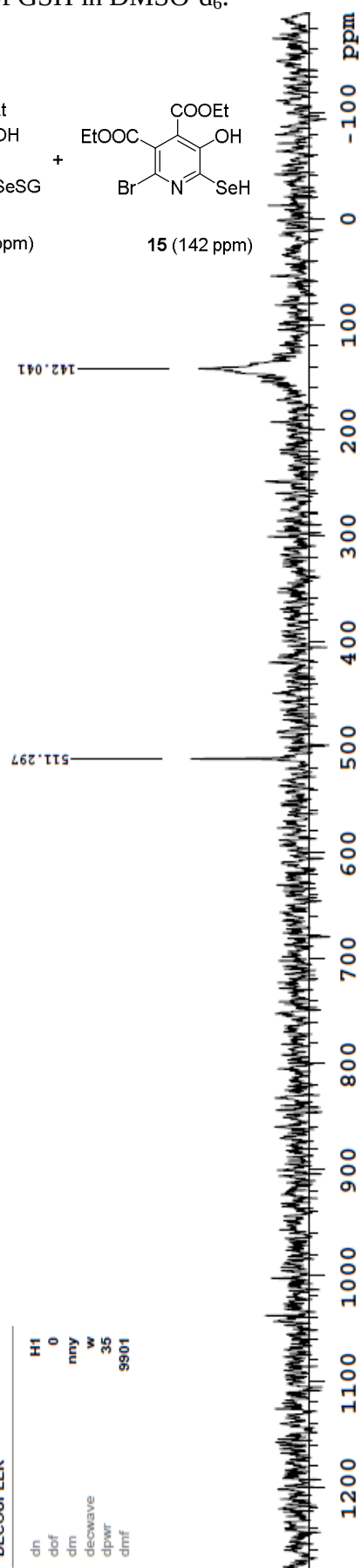
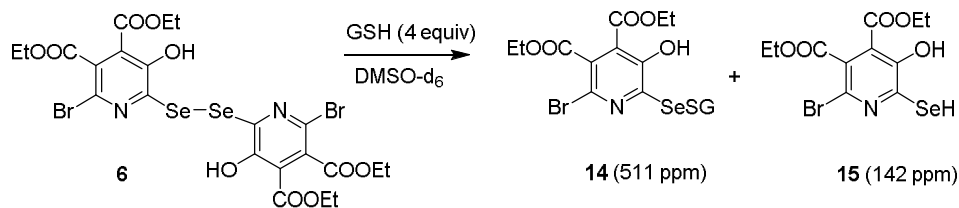
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bs: 4
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nt: 10000
ct: 10000

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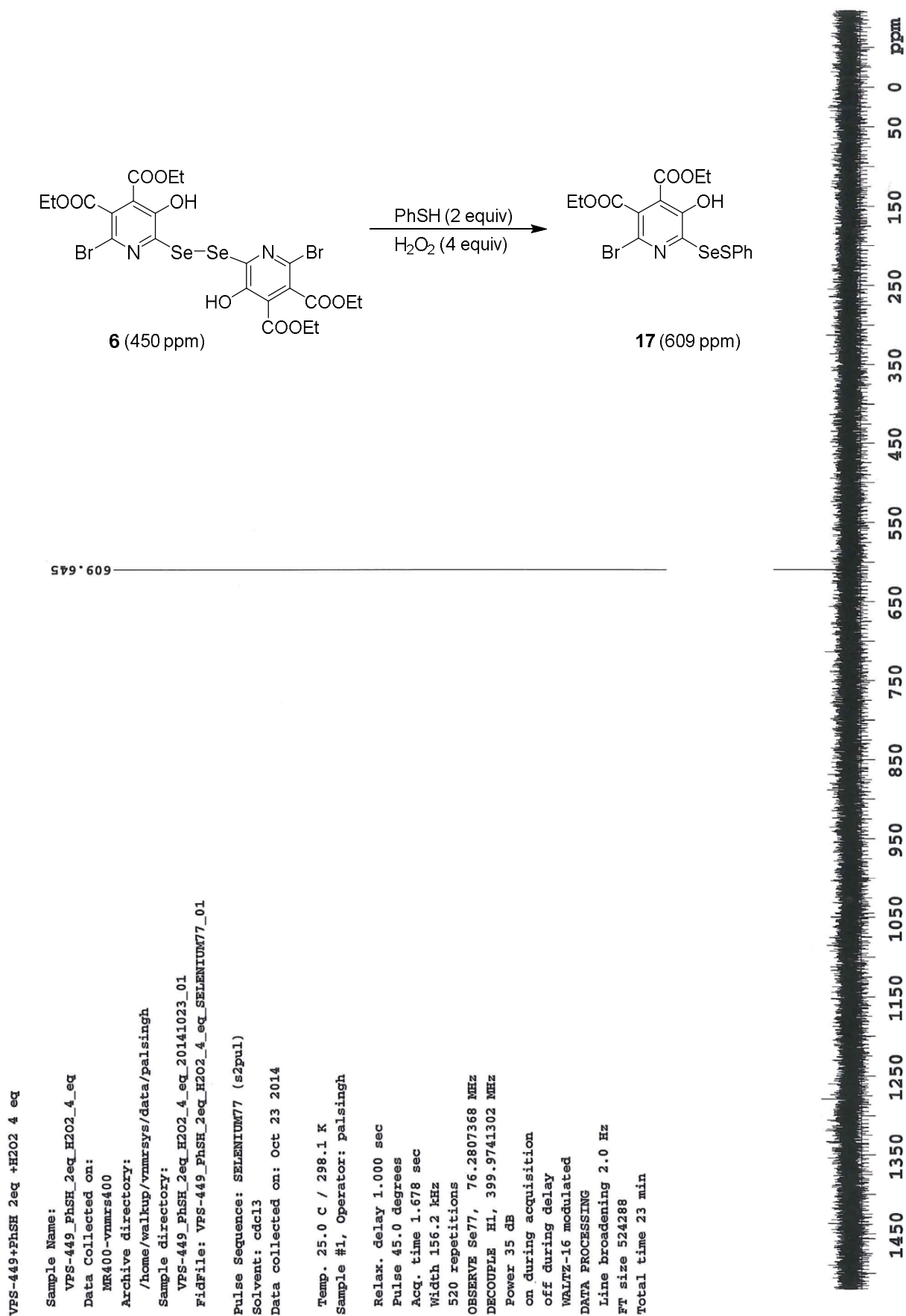
tn: Se77
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tof: 20675.1
tpwr: 58
pw: 5.600

DECOUPLER

dn: H1
dof: 0
dm: nny
decwave: w
dpwr: 35
dmf: 9901



^{77}Se NMR spectrum of diselenide **6** with PhSH (2 equiv) and H_2O_2 (4 equiv) in CDCl_3



VPS-449_2_eq_PhSH_20_eq_H2O2



Study owner
Operator

palsingh
palsingh

Temperature
Spectrometer

25
MR400-vnmrs400

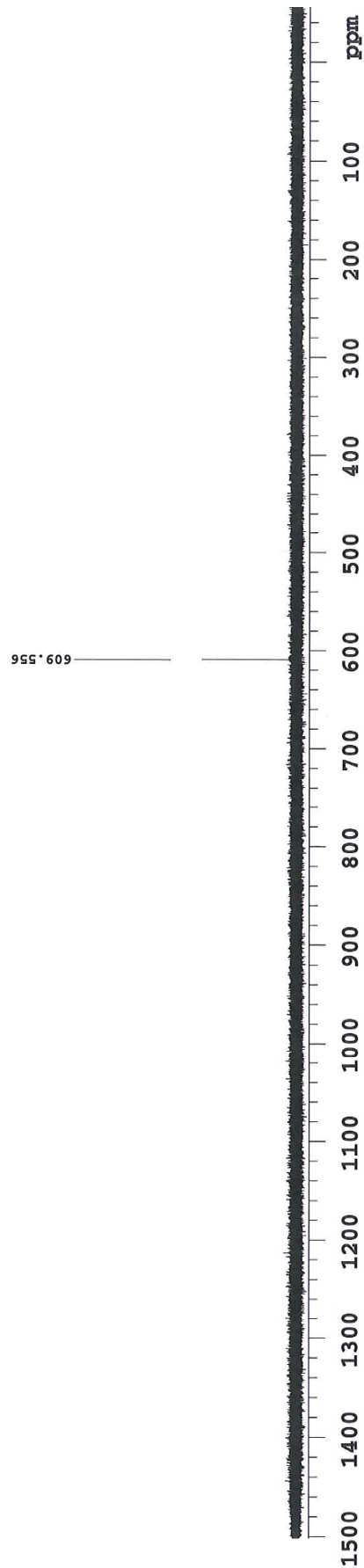
SELENIUM77
Solvent

sequence
cdcl3

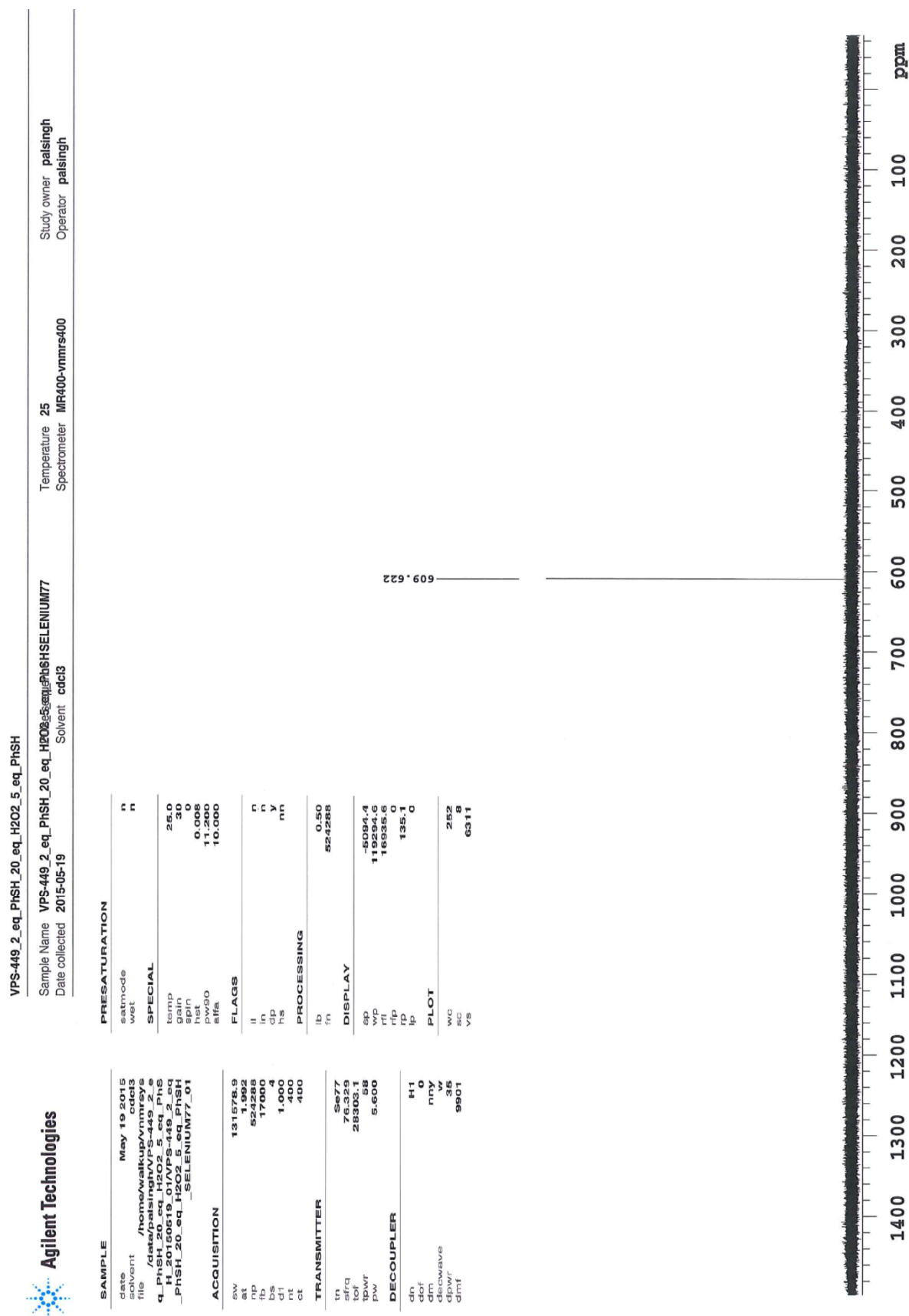
Sample Name
Date collected

VPS-449_2_eq_PhSH_20_eq_H2O2
2015-05-19

SAMPLE		PRESATURATION	
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comment		wet	
file	/home/walkup/vnmrsys	SPECIAL	
q	/data/palsingh/VPS-449_2_e	temp	25.0
q	PhSH_20_eq_H2O2_20150519	spin	30
q	01/VPS-449_2_eq_PhSH_20_e	spit	0
q	q_H2O2_SELENIUM77_01	hst	0.008
		pw90	11.200
		alfa	10.000
ACQUISITION		FLAGS	
sw	131578.9	ll	n
sf	524288	ln	n
np	1.992	ls	y
fb	17000	ns	nm
bs	4		
dt	1.000		
ct	400		
ct	400		
TRANSMITTER		PROCESSING	
tn	Se77	fn	524288
afreq	76329		
tof	28303.1		
tpwr	58		
pw	5.000		
DECOUPLER		DISPLAY	
dn	H1	sp	-4308.9
dof	0	wp	118841.4
dmf	nmr	rf	16355.0
decouwa	w	rp	134.9
cpwr	35	lp	0
dmf	9801		
PLOT		PLOT	
wo	252	wo	252
sc	8	sc	8
ve	2235	ve	2235
gn	11	gn	11



⁷⁷Se NMR spectrum of diselenide **6** after the addition of PhSH (3 equiv) to the above [total PhSH (5 equiv) and H₂O₂ (20 equiv)]



VPS-449_20eq_H2O2_20eq_PhSH

Sample Name VPS-449_20eq_H2O2_20eq_PhSH sequence SELENIUM77
Date collected 2015-05-19 Solvent cdcl3
Temperature 25 Spectrometer MR400-vnmrs400
Study owner palsingh
Operator palsingh

SAMPLE
date May 19 2015
solvent cdcl3
file /home/walkup/vnmrs/sequences/20150519/01/enterQ.macdir/loc8_004/current

ACQUISITION
sw 131575.9
st 1.992
np 524288
ls 17000
d1 1.000
nt 200
ct 4

TRANSMITTER
fr 6077
sfrq 76.329
tofr 283.031
tpwr 58
pwr 5.600

DECOUPLER
dn H1
dof 0
dm nny
dswelve 35
dsw 9901
dmf

PRESATURATION
saltmode n
wet n
SPECIAL

temp 25.0
spin 50
hst 0.008
pw90 11.200
sfrq 16.600

FLAGS
ll n
in n
dp y
hs nn

PROCESSING
lb 1.00
fn 524288

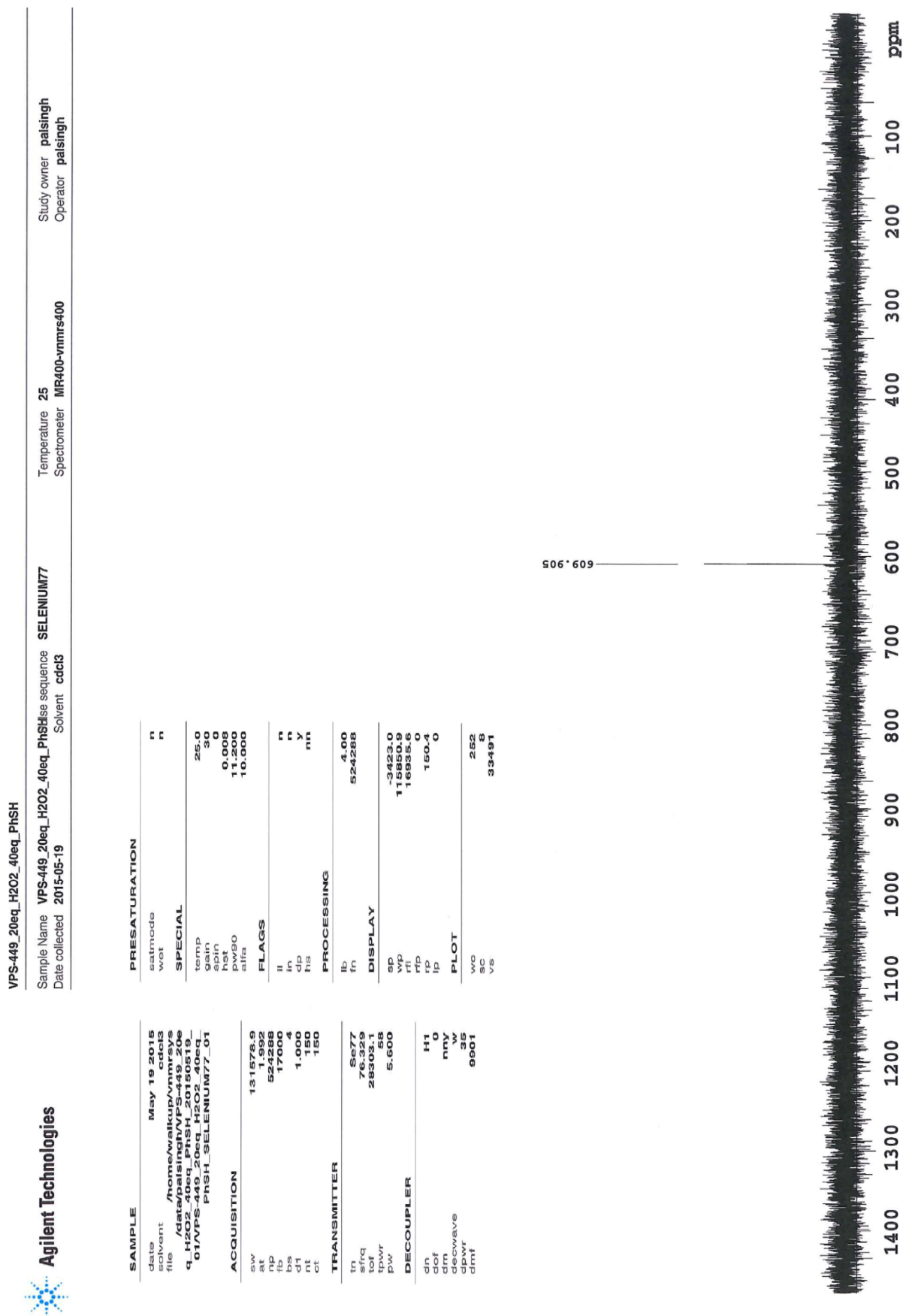
DISPLAY
sp -2991.8
wp 115753.5
rl 16935.6
rp 138.0
lp 0

PLOT
wc 252
sc 5
vs 10740

609.773



⁷⁷Se NMR spectrum of diselenide **6** after the addition of PhSH (20 equiv) to the above [total PhSH (40 equiv) and H₂O₂ (20 equiv)]



⁷⁷Se NMR spectrum of **11** with H₂O₂ (4 equiv)

Sample Name:
VFS453_4H2O2
Data Collected on:
MR400-vnmrs400
Archive directory:
/home/walkup/vnmrsys/data/ruishixa
Sample directory:
xrs_23_87_20131003_01
FidFile: SELENIUM77

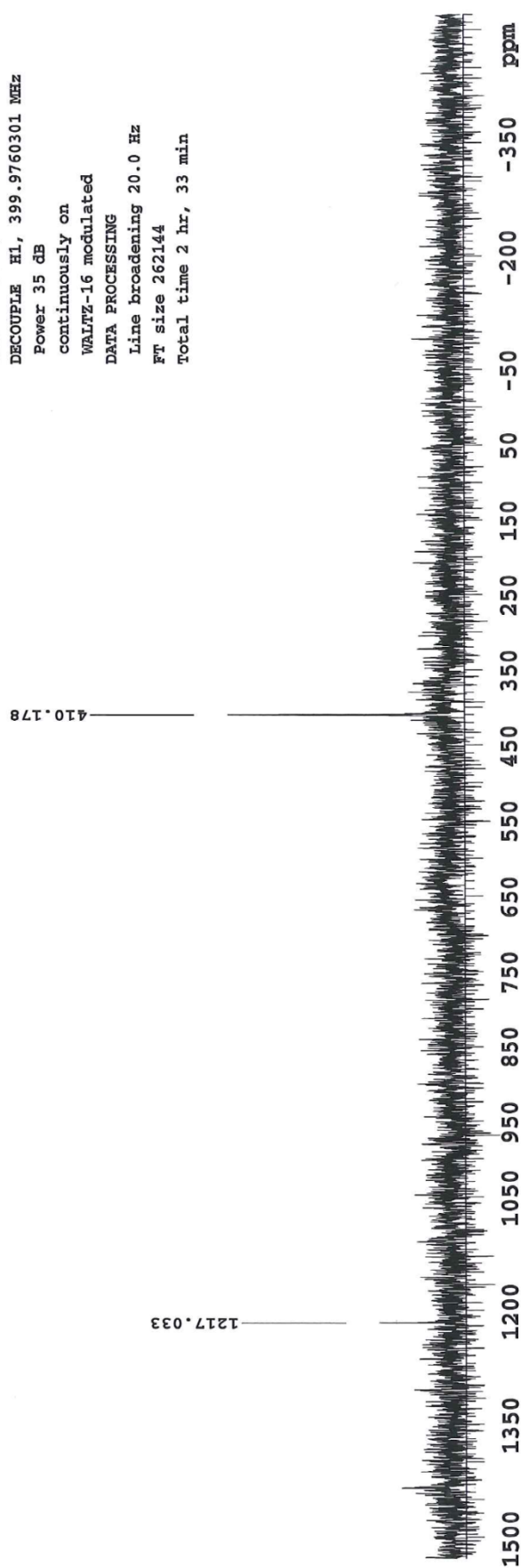
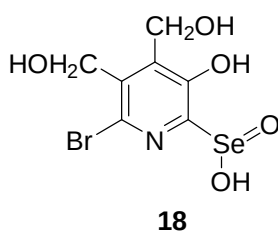
Pulse Sequence: SELENIUM77 (s2pul)
Solvent: dmsd
Data collected on: Oct 5 2013

Temp. 25.0 C / 298.1 K
Operator: jiafei

Relax. delay 1.000 sec
Pulse 24.2 degrees
Acq. time 0.839 sec
Width 156.2 kHz
792 repetitions

OBSERVE Se77, 76.2810992 MHz
DECOUPLE H1, 399.9760301 MHz

Power 35 dB
continuously on
WALTZ-16 modulated
DATA PROCESSING
Line broadening 20.0 Hz
FT size 262144
Total time 2 hr, 33 min



⁷⁷Se NMR spectrum of **11** with H₂O₂ (10 equiv)

Sample Name:
VPS453_4H2O2_6H2O2
Data Collected on:
MR400-vnmrs400
Archive directory:
/home/walkup/vnmrsys/data/ruishix
Sample directory:
xrs_23_87_20131003_01
FidFile: SELENIUM77

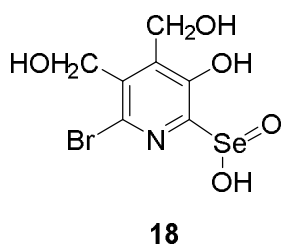
Pulse Sequence: SELENIUM77 (s2pul)
Solvent: dmsd
Data collected on: Oct 5 2013

Temp. 25.0 C / 298.1 K
Operator: jiafei

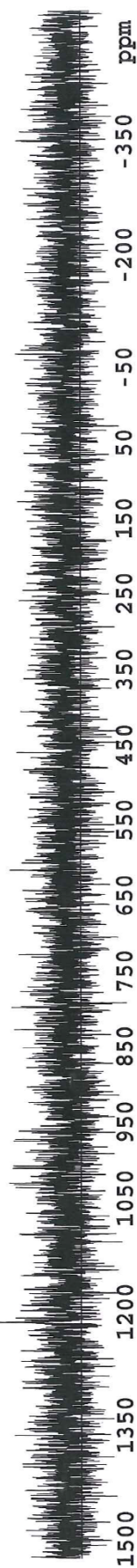
Relax. delay 1.000 sec
Pulse 24.2 degrees
Acq. time 0.839 sec
Width 156.2 kHz
192 repetitions

OBSERVE Se77, 76.2810992 MHz
DECOUPLE H1, 399.9760301 MHz
Power 35 dB

continuously on
WALTZ-16 modulated
DATA PROCESSING
Line broadening 15.0 Hz
FT size 262144
Total time 2 hr, 33 min



1216.314



⁷⁷Se NMR spectrum of **11** with H₂O₂ (10 equiv) and PhSH (20 equiv)

Sample Name:
VPS453_4H2O2_6H2O2_20PhSH
Data Collected on:
MR400-vnmrs400
Archive directory:
/home/walkup/vnmrsys/data/ruishixia
Sample directory:
xrs_23_87_20131003_01
FidFile: SELENIUM77

Pulse Sequence: SELENIUM77 (s2pul)
Solvent: dmsd
Data collected on: Oct 5 2013

Temp. 25.0 C / 298.1 K
Operator: jiafei

Relax. delay 1.000 sec
Pulse 24.2 degrees
Acq. time 0.839 sec
Width 156.2 kHz
184 repetitions

OBSERVE Se77, 76.281092 MHz
DECOUPLE H1, 399.9760301 MHz
Power 35 dB

continuously on

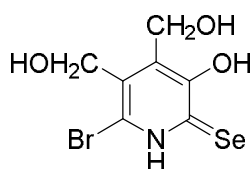
WALTZ-16 modulated

DATA PROCESSING

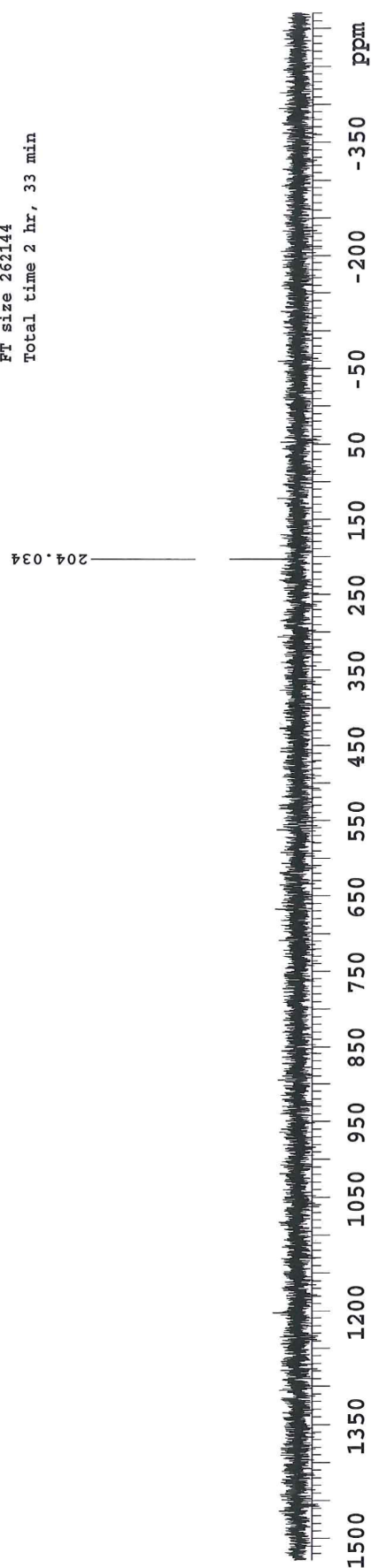
Line broadening 10.0 Hz

FT size 262144

Total time 2 hr, 33 min



Selone



⁷⁷Se NMR spectrum of **12** with H₂O₂ (8 equiv)

Sample Name:
VPS454_8H2O2
Data Collected on:
MR400-vnmrs400
Archive directory:
/home/walkup/vnmrsys/data/ruishixa
Sample directory:
xrs_23_87_20131003_01
FidFile: SELENIUM77

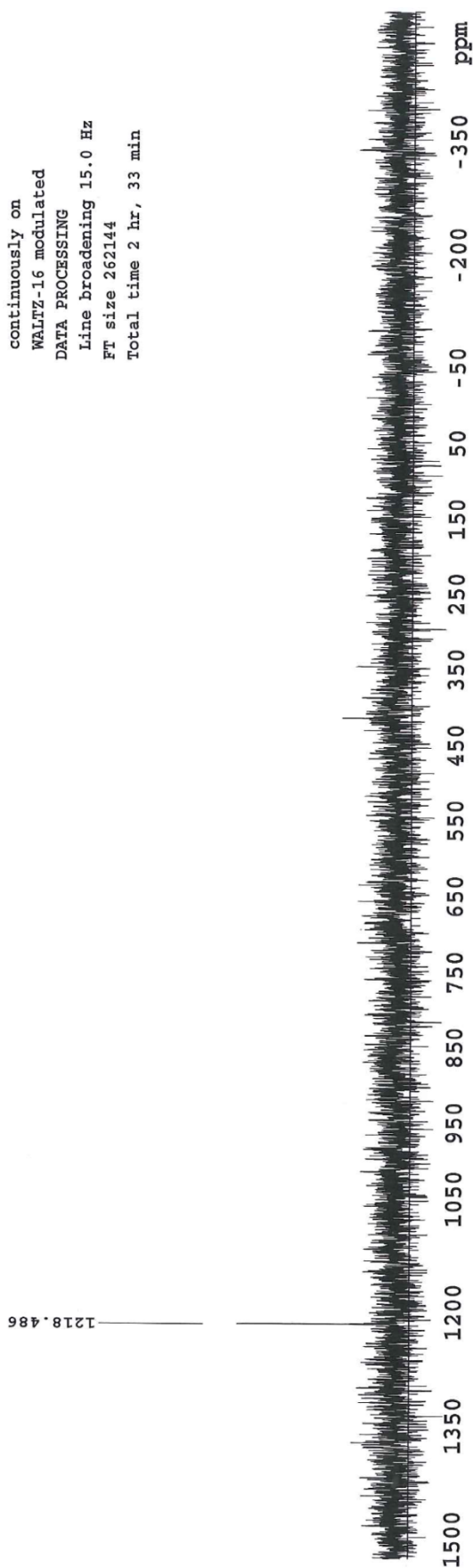
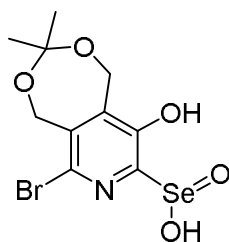
Pulse Sequence: SELENIUM77 (s2pul)
Solvent: dmsd
Data collected on: Oct 5 2013

Temp. 25.0 C / 298.1 K
Operator: jiafei

Relax. delay 1.000 sec
Pulse 24.2 degrees
Acq. time 0.839 sec
Width 156.2 kHz
360 repetitions

OBSERVE Se77, 76.2810992 MHz
DECOUPLE H1, 399.9760301 MHz

Power 35 dB
continuously on
WALTZ-16 modulated
DATA PROCESSING
Line broadening 15.0 Hz
Ft size 262144
Total time 2 hr, 33 min



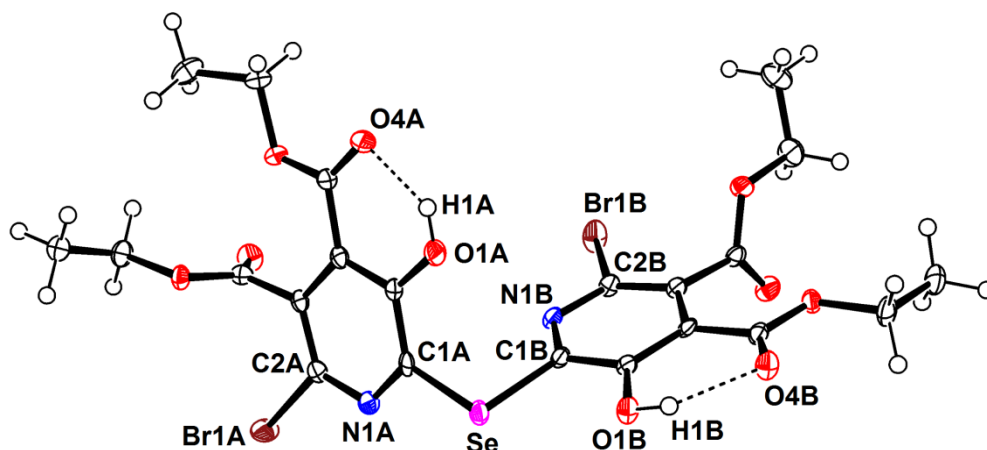


Figure S1. Molecular structure of momoselenide 7. Thermal ellipsoids are set at 50% probability. Significant bond lengths [Å] Se-C1A 1.922(4), Se-C1B 1.920(4), H1A...O4A 1.813, H1B...O4B 1.897, C1-N1 1.352(17), C1-C2 1.420(18), and angles [°] C1A-Se-C1B 97.46(15).

The molecular structure of monoselenide 7 showed a bent geometry around the Se atom with a C1A-Se-C1B bond angle of 97.46(15)°. The pyridinolic OH groups were intramolecularly H-bonded to the oxygen atom neighbouring ester groups. The H...O distances (H1A...O4A 1.813 Å, H1B...O4B 1.897 Å) are significantly shorter than the sum of the van der Waals radii of these two atoms.

Table S1. Optimized geometries and coordinates of **3a**

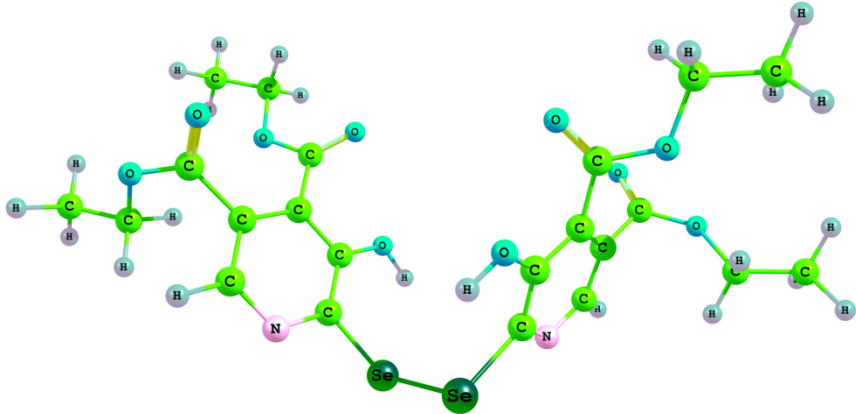
				3a			
7	2.939046	-2.076377	-0.900895	8	-4.352272	2.116894	2.013477
6	2.128193	-1.745912	0.092284	8	-6.078002	0.856391	1.438069
6	2.132218	-0.475884	0.709536	6	-8.139348	-0.061233	0.657924
6	3.068047	0.471058	0.255234	6	-6.631298	-0.163582	0.564889
6	3.938229	0.097999	-0.772104	1	4.457307	-1.484731	-2.156860
6	3.822249	-1.179866	-1.330560	1	0.725636	-0.903390	1.912428
8	1.310944	-0.149062	1.718289	1	-4.089013	-1.309233	2.377331
6	-2.036497	-1.699335	-0.109782	1	-0.854112	-0.951853	-2.122949
6	-2.168789	-0.492108	-0.829401	1	6.182316	4.896426	2.017742
6	-3.091716	0.456841	-0.353524	1	6.449133	3.202053	2.445674
6	-3.818329	0.150613	0.799858	1	6.551149	3.719744	0.750023
6	-3.572937	-1.061413	1.454577	1	4.090481	4.273775	0.771589
7	-2.706756	-1.962776	1.000674	1	3.982075	3.750668	2.455376
8	-1.482616	-0.228010	-1.951907	1	8.621467	-0.529931	0.667273
6	3.091178	1.856324	0.840391	1	8.715751	-0.021417	-1.026634
8	2.118766	2.528894	1.046976	1	8.419615	1.178239	0.241352
8	4.360231	2.243964	1.081552	1	6.196547	0.126112	0.731265
6	6.018207	3.865754	1.691970	1	6.499130	-1.096348	-0.521073
6	4.535018	3.619288	1.523393	1	-6.599295	4.585307	-2.096862
6	4.911489	1.072877	-1.393941	1	-6.828169	2.849973	-2.344808
8	4.534007	1.979237	-2.087135	1	-6.765769	3.514413	-0.699260
8	6.228100	0.850088	-1.218897	1	-4.353335	4.157339	-1.048878
6	8.206925	0.155750	-0.076853	1	-4.407659	3.497099	-2.686735
6	6.715564	-0.072038	-0.208495	1	-8.601441	-0.8191960	0.019470
6	-3.248950	1.779855	-1.051302	1	-8.480295	-0.222718	1.682682
8	-2.342010	2.463610	-1.439161	1	-8.486248	0.921178	0.331530
8	-4.554882	2.100053	-1.161082	1	-6.279591	0.016568	-0.452991
6	-6.349117	3.595599	-1.705424	1	-6.282290	-1.146829	0.888571
6	-4.846000	3.423880	-1.689697	34	0.914660	-3.110281	0.780970
6	-4.761400	1.140995	1.443707	34	-0.851283	-3.079084	-0.823065

Table S2. Optimized geometries and coordinates of **6**

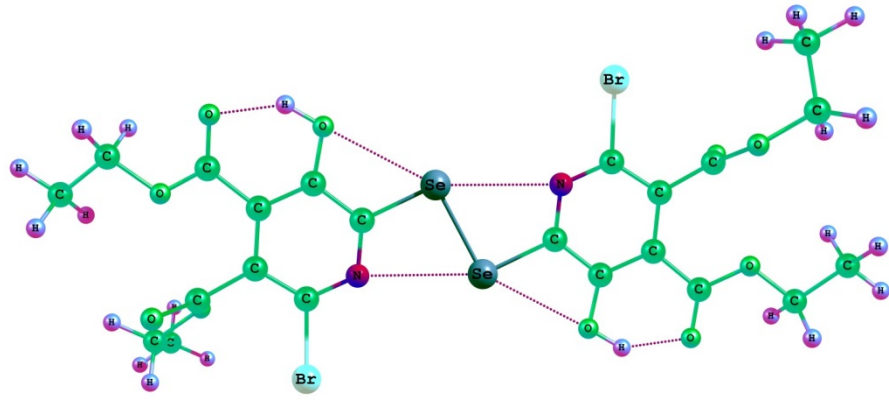
				6			
35	-4.175135	2.901887	0.336984	1	-8.572064	1.229940	-1.810897
35	4.152203	-2.925661	-0.436161	1	-8.770641	0.969375	-0.062836
34	-0.394868	-1.140485	-0.067940	6	-8.195062	2.991895	-0.601602
34	0.387132	1.127966	-0.004564	1	-9.219489	3.367593	-0.683673
8	-2.745045	-2.883890	-0.030479	1	-7.584646	3.501947	-1.349768
8	-5.238870	-3.554940	0.082333	1	-7.818333	3.246823	0.389861
8	-6.827232	-1.978081	0.043215	6	2.227044	0.585064	-0.116731
8	-6.866203	0.700224	1.426177	6	3.202876	1.609157	-0.112133
8	-6.838958	0.951521	-0.822740	6	4.558319	1.241676	-0.176038
8	5.251574	3.521528	-0.158472	6	4.885473	-0.135215	-0.271431
8	6.821454	1.929575	-0.027921	6	3.829254	-1.036870	-0.276802
8	6.808511	-1.082324	0.693605	6	5.562760	2.336679	-0.126338
8	6.850970	-0.679378	-1.528238	6	7.851565	2.960609	-0.023475
7	-2.566489	0.673157	0.1199320	1	7.671309	3.615392	0.830784
7	2.552161	-0.691241	-0.195925	1	7.739580	3.552183	-0.933392
6	-2.236039	-0.601561	0.044227	6	9.194430	2.269220	0.050483
6	-3.207345	-1.629911	0.042106	1	9.988867	3.020249	0.047046
6	-4.564826	-1.269265	0.108600	1	9.346286	1.609228	-0.805761
6	-4.897994	0.107664	0.195965	1	9.288895	1.683280	0.967411
6	-3.845554	1.013417	0.198126	6	6.295345	-0.640729	-0.460332
6	-5.561470	-2.372443	0.079770	6	8.113441	-1.713262	0.622827
6	-7.845155	-3.021276	0.062522	1	8.828075	-0.984391	0.234787
1	-7.679912	-3.636357	0.948359	1	8.054535	-2.538037	-0.090295
1	-7.701660	-3.650121	-0.817826	6	8.470483	-2.183594	2.016363
6	-9.197765	-2.345329	0.076565	1	9.450924	-2.667553	2.002568
1	-9.981977	-3.106623	0.102704	1	7.739707	-2.905180	2.387371
1	-9.347756	-1.737691	-0.818397	1	8.513525	-1.347598	2.718303
1	-9.311876	-1.708404	0.955780	1	-3.504181	-3.507018	0.004603
6	-6.314695	0.605461	0.359851	8	2.747965	2.865926	-0.037151
6	-8.189433	1.490455	-0.823810	1	3.509513	3.484938	-0.083367

Table S3. Optimized geometries and coordinates of **11**

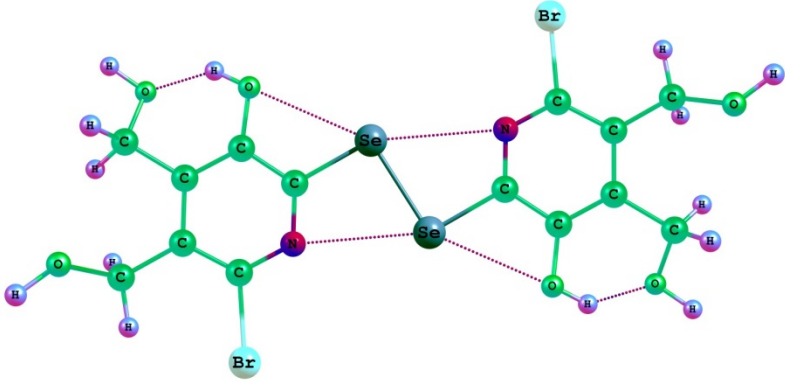
				11			
35	-0.274694	-0.227880	-0.009656	6	4.952724	0.138314	1.528016
35	0.280670	-0.111412	10.151738	1	5.167513	-0.281898	0.545419
34	1.172632	-0.112976	5.338690	1	5.210204	1.202791	1.494005
34	-1.166622	-0.226978	4.803368	6	2.926103	-0.040980	-0.593207
8	3.902468	0.0450060	4.278896	1	2.047414	-0.109180	-1.233102
7	0.786364	-0.150421	2.526077	1	3.571045	-0.902815	-0.819409
7	-0.780924	-0.178035	7.615900	8	5.758135	-0.533202	2.520374
6	1.696204	-0.100816	3.487469	1	6.687973	-0.329711	2.370857
6	3.073245	-0.028970	3.209657	8	3.640000	1.180775	-0.839496
6	3.488714	-0.014758	1.876180	1	3.763650	1.287361	-1.790407
6	2.512527	-0.072347	0.852962	6	-4.956902	-0.259873	8.611969
6	1.181196	-0.134275	1.268050	1	-5.132675	-0.679817	9.602376
6	-1.690639	-0.230426	6.654506	1	-5.312397	0.776457	8.626191
6	-3.068595	-0.281932	6.932163	6	-2.924053	-0.211949	10.734558
6	-3.484674	-0.282464	8.265555	1	-2.043313	-0.186984	11.374787
6	-2.508241	-0.230793	9.288819	1	-3.486606	-1.125578	10.976812
6	-1.176567	-0.176186	8.873743	8	-5.695328	-1.021815	7.632824
1	4.811696	-0.157790	3.989122	1	-6.640227	-0.903968	7.779936
8	-3.900263	-0.304564	5.862521	8	-3.748100	0.942774	10.959482
1	-4.786841	-0.586083	6.156650	1	-3.883114	1.053620	11.908369

Table S4. Optimized geometries and coordinates of **9** and **10**

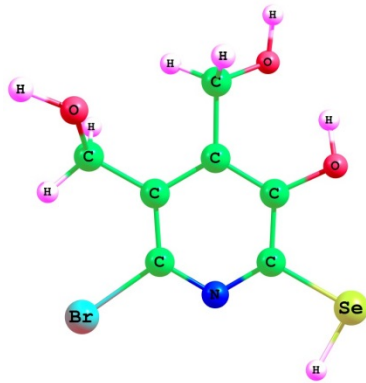
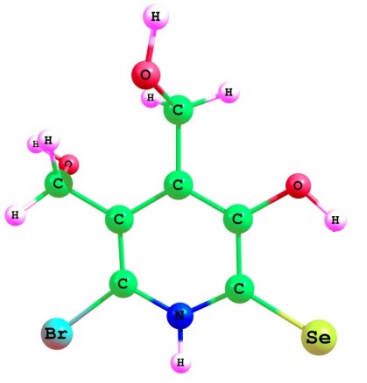
 Selenol 9				 Selone 10			
34	-0.071748000	-0.199011000	0.122877000	6	-0.027562000	-0.005076000	-0.009218000
8	0.156760000	-0.034512000	3.059001000	6	-0.013450000	-0.029745000	1.359995000
7	2.720989000	-0.250531000	0.569261000	7	1.164767000	-0.026190000	2.047763000
6	1.544212000	-0.192657000	1.169719000	6	2.404655000	-0.008466000	1.474408000
6	1.415787000	-0.115479000	2.565947000	6	2.412168000	0.009331000	0.041892000
6	2.578194000	-0.102764000	3.344384000	6	1.235074000	0.003049000	-0.682551000
6	3.834392000	-0.168159000	2.700731000	8	3.599014000	0.026334000	-0.585352000
6	3.813302000	-0.237270000	1.304884000	6	1.292158000	0.018652000	-2.189989000
1	0.178301000	-0.234358000	4.013892000	6	-1.307999000	-0.027823000	-0.800343000
6	2.493834000	0.056885000	4.846302000	1	1.143087000	-0.040218000	3.060071000
1	3.374520000	-0.359990000	5.334576000	1	4.296186000	0.023149000	0.108710000
1	2.451625000	1.122563000	5.097456000	1	0.624602000	-0.752693000	-2.588743000
6	5.104536000	-0.137608000	3.507502000	1	2.311254000	-0.195839000	-2.513275000
1	5.136742000	-0.998948000	4.190809000	1	-1.208434000	0.634194000	-1.663981000
1	5.967530000	-0.208210000	2.846828000	1	-2.137082000	0.331825000	-0.185741000
8	5.139505000	1.085134000	4.259348000	8	0.879174000	1.319575000	-2.642510000
1	6.016120000	1.191659000	4.648022000	8	-1.543807000	-1.380770000	-1.219608000
8	1.313037000	-0.613411000	5.336341000	1	1.107755000	1.412392000	-3.575053000
1	1.194034000	-0.413639000	6.271407000	1	-2.350127000	-1.407047000	-1.749715000
1	0.702268000	-0.281688000	-1.137076000	34	3.949360000	-0.003390000	2.475107000
35	5.452420000	-0.345411000	0.272927000	35	-1.556401000	-0.078866000	2.471924000

Table S5. Second Order Perturbation Theory Analysis of Fock Matrix in NBO Basis for **6**

Donor NBO (i)	Acceptor NBO (j)	E(2)	E(j)-E(i)	F(i,j)
		kcal/mol	a.u.	a.u.
172. LP (1) O5	/996. BD*(1) Se3-Se4	1.11	0.58	0.023
172. LP (1) O5	/997. BD*(1) Se3-C16	0.61	0.72	0.019
190. LP (1) N14	/996. BD*(1) Se3-Se4	0.69	0.34	0.014
190. LP (1) N14	/997. BD*(1) Se3-C16	3.87	0.47	0.039
190. LP (1) N14	/998. BD*(1) Se4-C37	2.50	0.47	0.031
191. LP (1) N15	/997. BD*(1) Se3-C16	2.49	0.63	0.036
191. LP (1) N15	/998. BD*(1) Se4-C37	2.48	0.63	0.036
174. LP (1) O6	/*. BD*(1) O5 - H58	3.74	1.12	0.058

Table S6. Second Order Perturbation Theory Analysis of Fock Matrix in NBO Basis for **11**

Donor NBO (i)	Acceptor NBO (j)	E(2)	E(j)-E(i)	F(i,j)
		kcal/mol	a.u.	a.u.
135. LP (1) O5	/717. BD*(1) Se3-Se4	1.11	0.58	0.023
135. LP (1) O5	/718. BD*(1) Se3-C8	0.61	0.72	0.019
141. LP (1) O19	/717. BD*(1) Se3-Se4	1.11	0.58	0.023
141. LP (1) O19	/719. BD*(1) Se4-C13	0.61	0.72	0.019
137. LP (1) N6	/717. BD*(1) Se3-Se4	0.61	0.33	0.013
137. LP (1) N6	/718. BD*(1) Se3-C8	3.50	0.47	0.037
137. LP (1) N6	/719. BD*(1) Se4-C13	2.49	0.47	0.031
138. LP (1) N7	/717. BD*(1) Se3-Se4	0.61	0.33	0.013
138. LP (1) N7	/718. BD*(1) Se3-C8	2.49	0.47	0.031
138. LP (1) N7	/719. BD*(1) Se4-C13	3.50	0.47	0.037

Figure S2. Molecular graphs of **6**, **10** and **11**

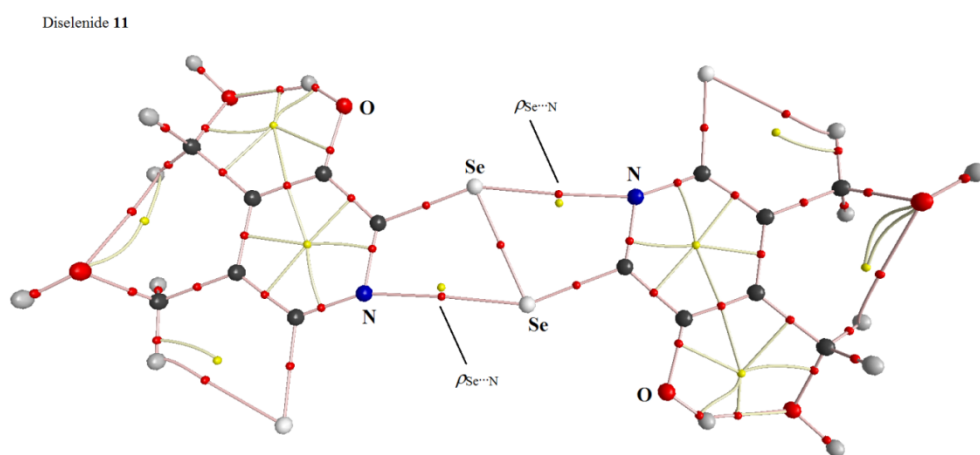
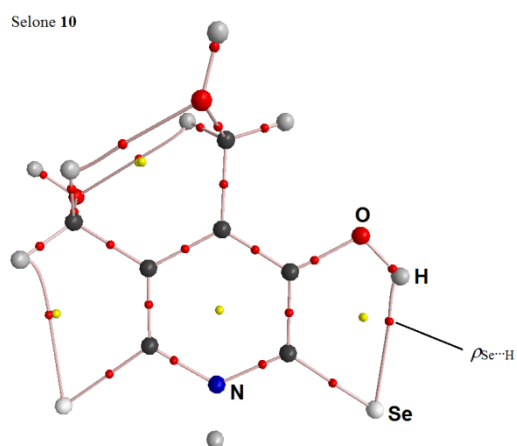
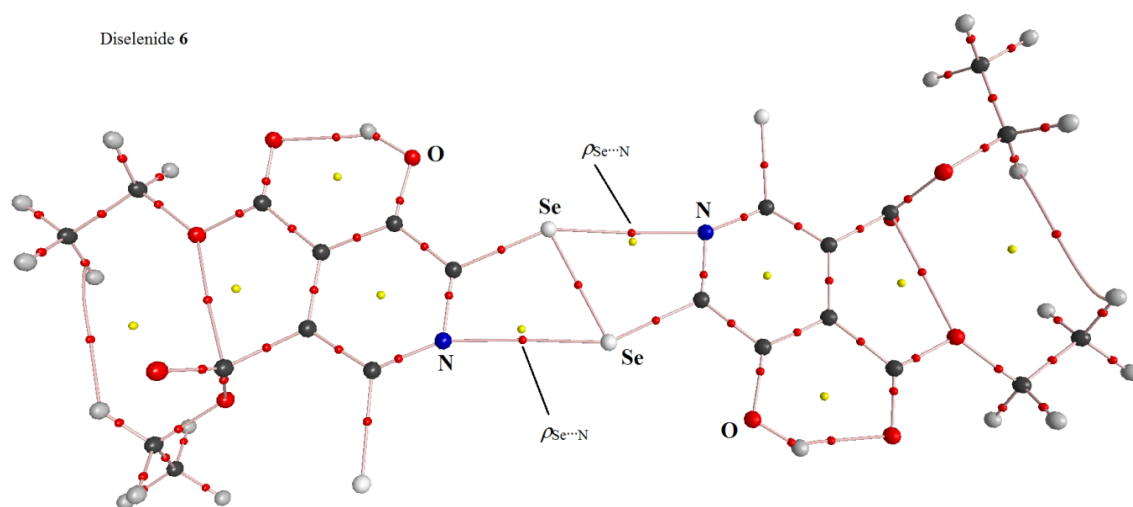


Table S7. Summary of properties of electron density at the bond critical point. AIM calculations were carried out by using the wave functions obtained at the B3LYP/6-311+G(d)//B3LYP/6-311+G(d,p).

Compound	$\rho_{\text{Se}\cdots\text{N}}$ (ea_0^{-3}) ^[a]	$\nabla^2\rho_{\text{Se}\cdots\text{N}}$ (ea_0^{-5}) ^[b]	$H_{\text{Se}\cdots\text{N}}$ (ea_0^{-4}) ^[c]	$\rho_{\text{Se}\cdots\text{H}}$ (ea_0^{-3}) ^[a]	$\nabla^2\rho_{\text{Se}\cdots\text{H}}$ (ea_0^{-5}) ^[b]	$H_{\text{Se}\cdots\text{H}}$ (ea_0^{-4}) ^[c]
6	0.0146	0.0526	+0.00176			
11	0.0144	0.0514	+0.00175			
10				0.0247	0.0525	-0.00098

[a] The electron density at the bond critical point. [b] The Laplacian of the electron density at the bond critical point. [c] The total energy density at the bond critical point.

Table S8. Effect of substrate (GSH) concentration on the initial rate of NADPH-consumption for catalyst **6**.

[GSH] (mM)	v_0 ($\mu\text{M} \cdot \text{min}^{-1}$)	Mean v_0 ($\mu\text{M} \cdot \text{min}^{-1}$)	1/ [GSH] (mM^{-1})	1/ v_0 ($\mu\text{M}^{-1} \cdot \text{min}$)
0.125	209.40 216.27 217.52	214.4 ± 4.4	8.0	0.00466
0.25	221.19 217.04 216.46	218.2 ± 2.6	4.0	0.00458
0.50	221.29 225.62 229.10	225.3 ± 3.9	2.0	0.00443
1.0	242.60 245.11 238.98	242.2 ± 3.1	1.0	0.00413
2.0	244.92 247.23 245.11	245.8 ± 1.3	0.5	0.00407
3.0	263.34 264.40 265.47	264.4 ± 1.1	0.33	0.00378

Assay conditions: phosphate buffer (100 mM), pH 7.5, with EDTA (1 mM), GSH (variable), NADPH (0.20 mM), glutathione reductase (1.3 unit/mL), and H_2O_2 (0.8 mM) catalyst **6** (20 μM).

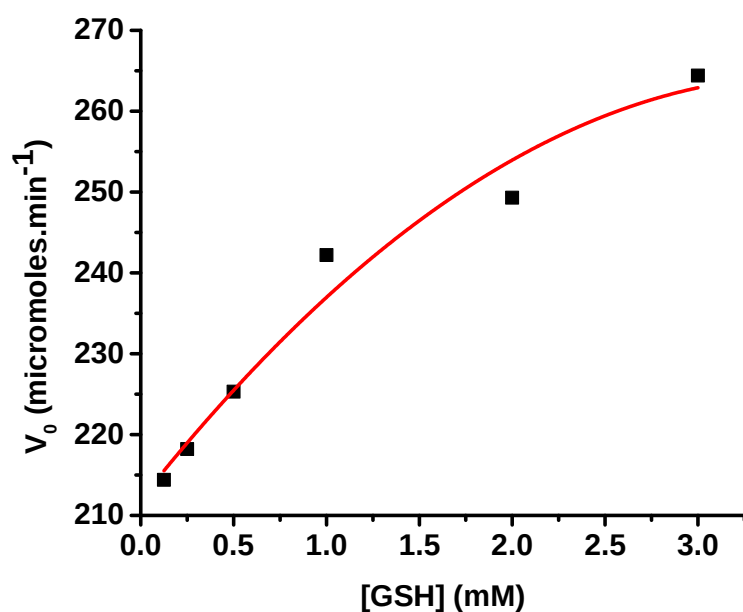


Figure S3. Effect of thiol concentration on initial rate of NADPH consumption for reduction of H_2O_2 in the presence of catalyst **6**.

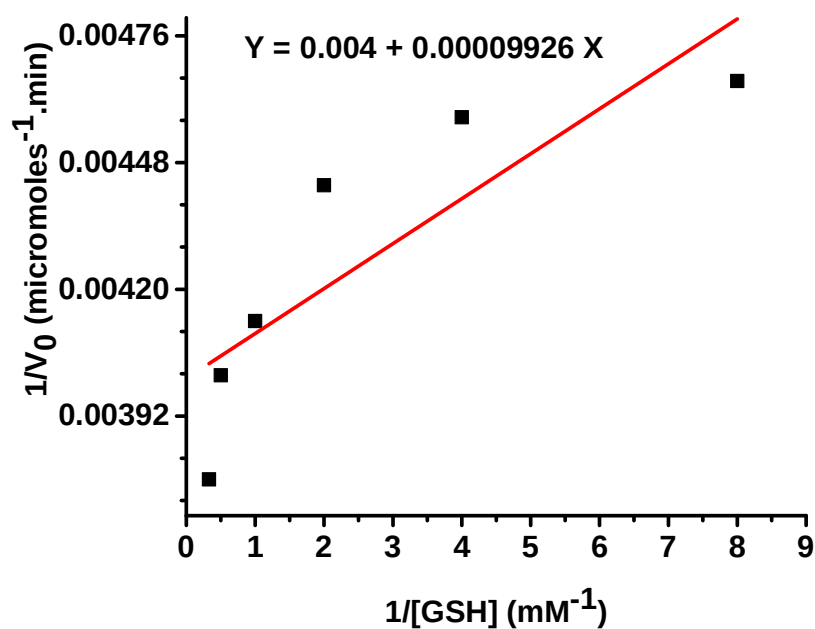


Figure S4. Lineweaver-Burk (double-reciprocal) plot obtained for catalyst **6** with GSH (variable) at a fixed concentration of H_2O_2 .

Table S9. Effect of substrate (H_2O_2) concentration on the initial rate of NADPH-consumption for catalyst **6**.

$[\text{H}_2\text{O}_2]$ (mM)	v_0 ($\mu\text{M}\cdot\text{min}^{-1}$)	Mean v_0 ($\mu\text{M}\cdot\text{min}^{-1}$)	$1/[\text{H}_2\text{O}_2]$ (mM^{-1})	$1/v_0$ ($\mu\text{M}^{-1}\cdot\text{min}$)
0.20	74.28 78.14 77.94	76.8 ± 2.2	5.0	0.01302
0.40	144.41 141.90 136.78	141.0 ± 3.9	2.5	0.00709
0.60	194.86 194.28 195.14	194.8 ± 0.4	1.66	0.00513
0.80	242.60 245.11 238.98	242.2 ± 3.1	1.0	0.00413
1.0	285.28 279.83 285.63	283.6 ± 3.3	1.0	0.00353
1.2	310.99 309.55 306.95	309.2 ± 2.0	0.833	0.00323

Assay conditions: phosphate buffer (100 mM), pH 7.5, with EDTA (1 mM), H_2O_2 (variable), NADPH (0.20 mM), glutathione reductase ($1.3 \text{ unit}\cdot\text{mL}^{-1}$), and GSH (1 mM) catalyst **6** (20 μM).

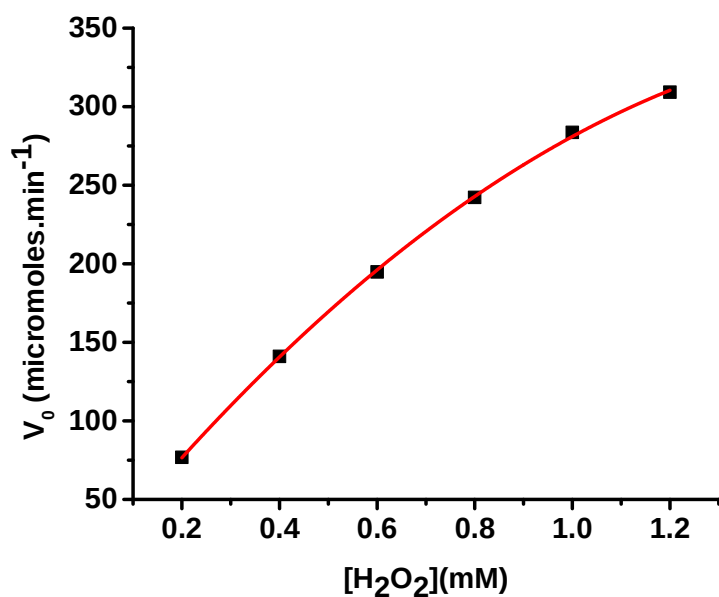


Figure S5. Effect of hydrogen peroxide concentration on initial rate of NADPH consumption for reduction of H₂O₂ in the presence of catalyst **6**.

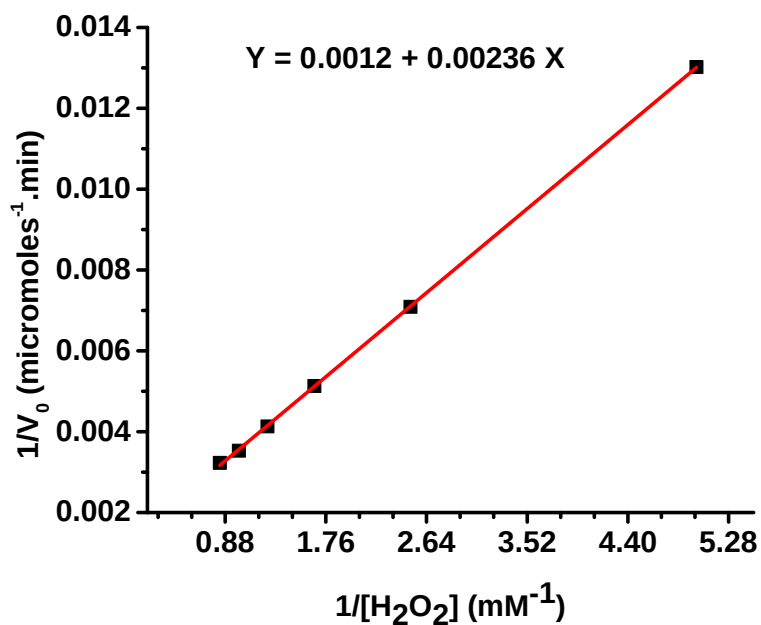


Figure S6. Lineweaver-Burk (double-reciprocal) plot obtained for catalyst **6** with hydrogen peroxide (variable) at a fixed concentration of GSH.

Table S10. Effect of substrate (GSH) concentration on the initial rate of NADPH-consumption for catalyst **11**.

[GSH] (mM)	v_0 ($\mu\text{M}\cdot\text{min}^{-1}$)	Mean v_0 ($\mu\text{M}\cdot\text{min}^{-1}$)	1/ [GSH] (mM^{-1})	1/ v_0 ($\mu\text{M}^{-1}\cdot\text{min}$)
0.125	95.02 92.12 97.52	94.9 ± 2.7	8.0	0.01054
0.25	102.06 106.40 106.50	105.0 ± 2.5	4.0	0.00952
0.50	122.12 126.75 123.57	124.1 ± 2.4	2.0	0.00806
1.0	171.80 166.39 167.27	168.5 ± 2.9	1.0	0.00593
2.0	215.11 211.93 212.60	213.2 ± 1.7	0.5	0.00469
3.0	245.11 243.67 247.33	245.4 ± 1.8	0.33	0.00407

Assay conditions: phosphate buffer (100 mM), pH 7.5, with EDTA (1 mM), GSH (variable), NADPH (0.20 mM), glutathione reductase ($1.3 \text{ unit}\cdot\text{mL}^{-1}$), and H_2O_2 (0.8 mM) catalyst **11** (20 μM).

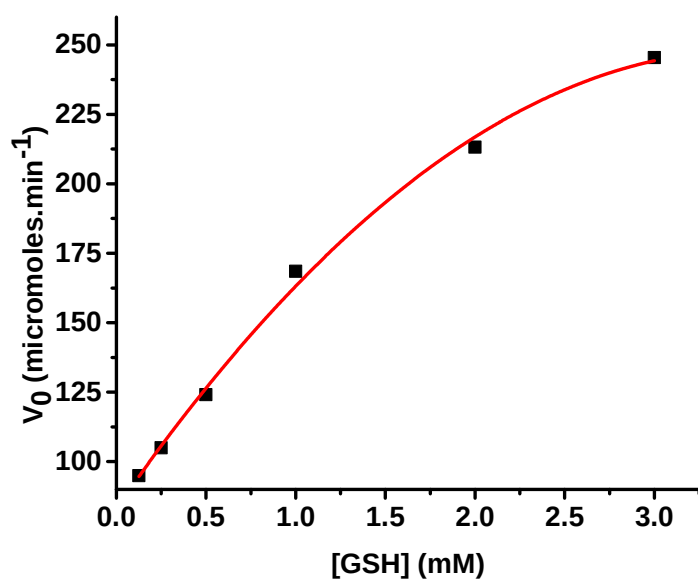


Figure S7. Effect of thiol concentration on initial rate of NADPH consumption for reduction of H_2O_2 in the presence of catalyst **11**.

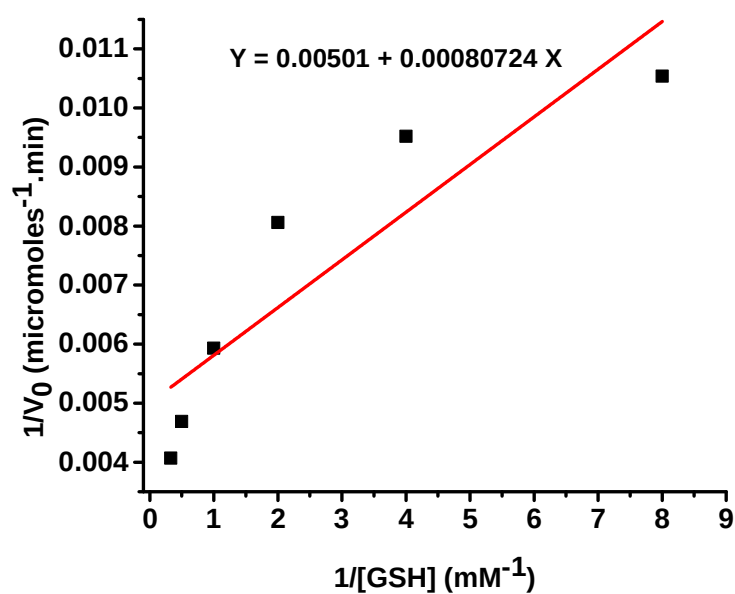


Figure S8. Lineweaver-Burk (double-reciprocal) plot obtained for catalyst **11** with GSH (variable) at a fixed concentration of H_2O_2 .

Table S11. Effect of substrate (H_2O_2) concentration on the initial rate of NADPH-consumption for catalyst **11**.

$[\text{H}_2\text{O}_2]$ (mM)	v_0 ($\mu\text{M}\cdot\text{min}^{-1}$)	Mean v_0 ($\mu\text{M}\cdot\text{min}^{-1}$)	$1/[\text{H}_2\text{O}_2]$ (mM^{-1})	$1/v_0$ ($\mu\text{M}^{-1}\cdot\text{min}$)
0.20	59.04 65.69 63.76	62.8 ± 3.4	5.0	0.01591
0.40	108.14 102.93 103.70	104.9 ± 2.8	2.5	0.00953
0.60	145.76 144.02 139.00	142.9 ± 3.5	1.66	0.00699
0.80	171.80 166.39 167.27	168.5 ± 2.9	1.25	0.00593
1.0	194.76 195.43 197.46	195.9 ± 1.4	1.0	0.00510
1.2	238.17 234.89 231.03	234.7 ± 3.6	0.833	0.00426

Assay conditions: phosphate buffer (100 mM), pH 7.5, with EDTA (1 mM), H_2O_2 (variable), NADPH (0.20 mM), glutathione reductase ($1.3 \text{ unit}\cdot\text{mL}^{-1}$), and GSH (1 mM) catalyst **11** (20 μM).

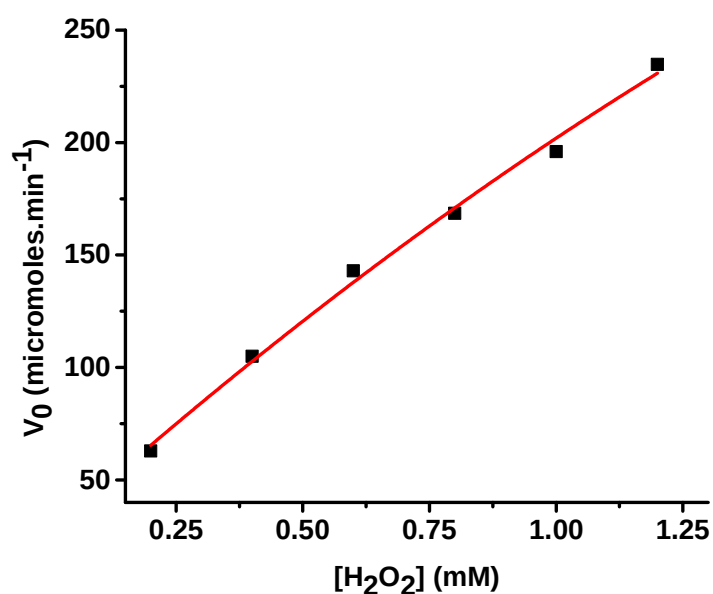


Figure S9. Effect of hydrogen peroxide concentration on initial rate of NADPH consumption for reduction of H₂O₂ in the presence of catalyst **11**.

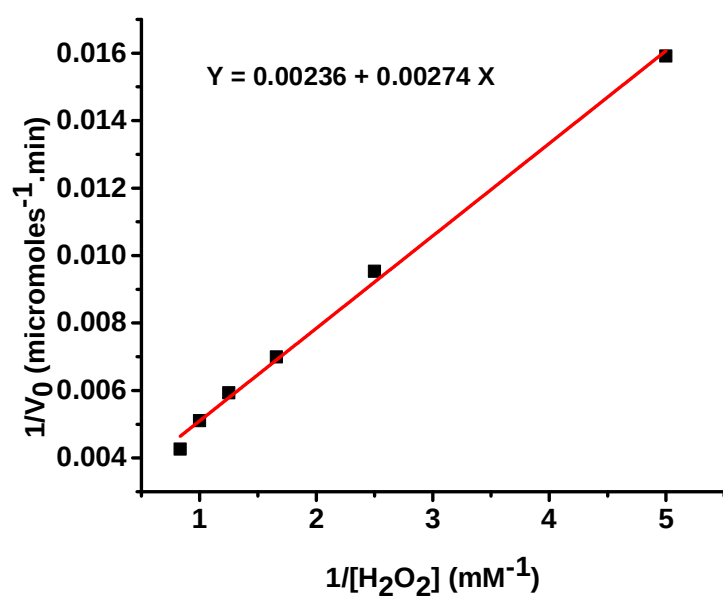


Figure S10. Lineweaver-Burk (double-reciprocal) plot obtained for catalyst **11** with hydrogen peroxide (variable) at a fixed concentration of GSH.

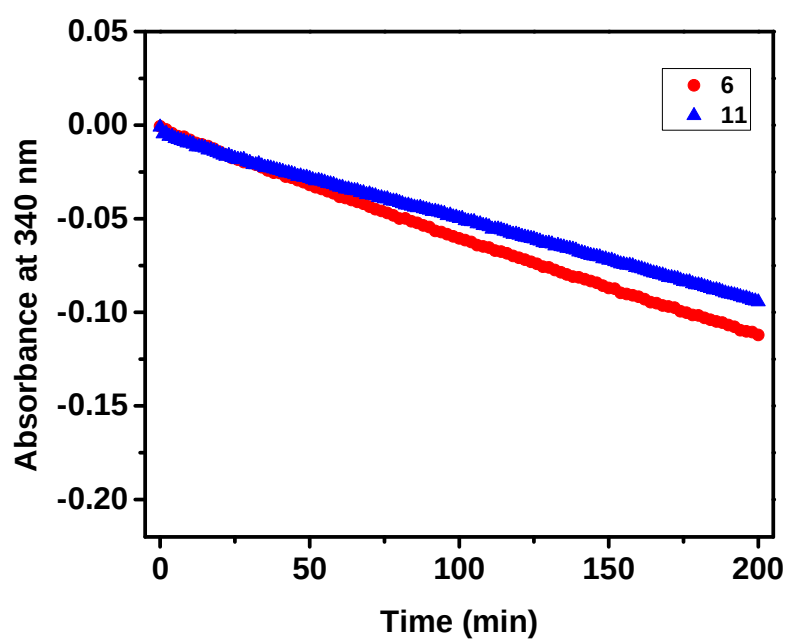


Figure S11. Consumption of NADPH with time in the absence of H_2O_2 (catalyst and GSH) for catalysts **6** and **11**. Assay conditions: reactions were carried out with phosphate buffer (100 mM), pH 7.5, with EDTA (1 mM), GSH (0.10 mM), NADPH (0.20 mM), GR (1.3 $\text{unit}\cdot\text{mL}^{-1}$), and selenium catalyst (2 μM).

Table S12. Crystal Data and Structure Refinement for compounds **6** and **7**.

Compound	6	7
Empirical formula	C ₂₂ H ₂₂ Br ₂ N ₂ O ₁₀ Se ₂	C ₂₂ H ₂₂ Br ₂ N ₂ O ₁₀ Se
Formula weight	792.15	713.19
Crystal system	Monoclinic	Monoclinic
Space group	P <i>c</i>	P 2 ₁ / <i>c</i>
a(Å)	20.039(15)	18.527(6)
b(Å)	5.315(4)	8.443(4)
c(Å)	25.268(2)	16.977(7)
α(deg)	90	90
β(deg)	90.05(3)	102.39(2) ^o
γ(deg)	90	90
V(Å ³)	2691.2(4)	2593.85(18)
Z	4	4
T	100(2)K	120(2)
ρ _{calcd} (Mg/m ³)	1.955	1.806
Abs coeff(mm ⁻¹)	5.774	4.586
Reflections collected	48954	15892
Final R(F) indices [I < 2σ(I)] ^a	R1 = 0.0492 wR2 = 0.1261	R1 = 0.0489 wR2 = 0.1056
wR(F ²) indices	R1 = 0.0536 wR2 = 0.1309 (all data)	R1 = 0.0759 wR2 = 0.01163 (all data)
Data/restrain/parameters	13011 / 896 / 700	7379 / 0 / 340
Goodness of fit on F ²	1.085	1.038

^aDefinitions: $R(F_o) = \frac{\sum ||F_o| - |F_c||}{\sum |F_o|}$ and $wR(F_o^2) = \{\sum [w(F_o^2 - F_c^2)^2] / \sum [w(F_c^2)^2]\}^{1/2}$.

CheckCIF/PLATON (standard) report for compound 6

Datablock: shelx

Bond precision:	C-C = 0.0253 Å	Wavelength=0.71073
Cell:	a=20.0390(15) b=5.3151(4) c=25.268(2)	
	alpha=90 beta=90.051(3) gamma=90	
Temperature:	100 K	

	Calculated	Reported
Volume	2691.3(4)	2691.2(4)
Space group	P c	P c
Hall group	P -2yc	P -2yc
Moiety formula	C22 H22 Br2 N2 O10 Se2	C22 H22 Br2 N2 O10 Se2
Sum formula	C22 H22 Br2 N2 O10 Se2	C22 H22 Br2 N2 O10 Se2
Mr	792.14	792.15
Dx, g cm ⁻³	1.955	1.955
Z	4	4
Mu (mm ⁻¹)	5.774	5.774
F000	1544.0	1544.0
F000'	1541.88	
h, k, lmax	26, 7, 33	26, 7, 33
Nref	13381[6694]	13011
Tmin, Tmax	0.299, 0.630	0.445, 0.746
Tmin'	0.227	

Correction method= # Reported T Limits: Tmin=0.445
Tmax=0.746 AbsCorr = MULTI-SCAN

Data completeness= 1.94/0.97 Theta(max)= 28.333

R(reflections)= 0.0492(12389) wR2(reflections)= 0.1309(13011)

S = 1.085 Npar= 700

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 B

[PLAT341_ALERT_3_B](#) Low Bond Precision on C-C Bonds 0.0253 Ang.
[PLAT369_ALERT_2_B](#) Long C(sp2)-C(sp2) Bond C3A - C6A ... 1.58 Ang.

🟢Alert level C

[DIFMX01_ALERT_2_C](#) The maximum difference density is > 0.1*ZMAX*0.75
_refine_diff_density_max given = 2.925
Test value = 2.625

[DIFMX02_ALERT_1_C](#) The maximum difference density is > 0.1*ZMAX*0.75
The relevant atom site should be identified.

[PLAT034_ALERT_1_C](#) No Flack Parameter Given. Z > Si, NonCentro Please Do !
[PLAT090_ALERT_3_C](#) Poor Data / Parameter Ratio (Zmax > 18) 9.50 Note
[PLAT094_ALERT_2_C](#) Ratio of Maximum / Minimum Residual Density 2.57 Report
[PLAT097_ALERT_2_C](#) Large Reported Max. (Positive) Residual Density 2.92 eA-3
[PLAT213_ALERT_2_C](#) Atom Br2A has ADP max/min Ratio 3.7 oblate
[PLAT220_ALERT_2_C](#) Large Non-Solvent C Ueq(max)/Ueq(min) Range 4.0 Ratio

And 3 other PLAT220 Alerts

More ...

PLAT222_ALERT_3_C Large Non-Solvent H Uiso(max)/Uiso(min) ... 4.3 Ratio
PLAT360_ALERT_2_C Short C(sp3)-C(sp3) Bond C10B - C11B ... 1.42 Ang.
PLAT369_ALERT_2_C Long C(sp2)-C(sp2) Bond C3B - C6B ... 1.53 Ang.
PLAT480_ALERT_4_C Long H...A H-Bond Reported H11A .. BR1B .. 2.98 Ang.

And 6 other PLAT480 Alerts

More ...

PLAT910_ALERT_3_C Missing # of FCF Reflection(s) Below Th(Min) ... 5 Report
PLAT911_ALERT_3_C Missing # FCF Refl Between THmin & STh/L= 0.600 26 Report

Alert level G

PLAT003_ALERT_2_G Number of Uiso or Uij Restrained non-H Atoms ... 71 Report
PLAT007_ALERT_5_G Number of Unrefined Donor-H Atoms 4 Report
PLAT083_ALERT_2_G SHELXL Second Parameter in WGHT Unusually Large. 16.57 Why ?
PLAT112_ALERT_2_G ADDSYM Detects Additional (Pseudo) Symm. Elem... 21 Check
PLAT112_ALERT_2_G ADDSYM Detects Additional (Pseudo) Symm. Elem... 21 Check
PLAT720_ALERT_4_G Number of Unusual/Non-Standard Labels 10 Note
PLAT860_ALERT_3_G Number of Least-Squares Restraints 896 Note
PLAT870_ALERT_4_G ALERTS Related to Twinning Effects Suppressed .. ! Info
PLAT912_ALERT_4_G Missing # of FCF Reflections Above STh/L= 0.600 15 Note
PLAT931_ALERT_5_G Found Twin Law (0 0 1)[] Estimated BASF 0.52 Check

0 **ALERT level A** = Most likely a serious problem - resolve or explain
2 **ALERT level B** = A potentially serious problem, consider carefully
23 **ALERT level C** = Check. Ensure it is not caused by an omission or oversight
10 **ALERT level G** = General information/check it is not something unexpected

2 ALERT type 1 CIF construction/syntax error, inconsistent or missing data
15 ALERT type 2 Indicator that the structure model may be wrong or deficient
6 ALERT type 3 Indicator that the structure quality may be low
10 ALERT type 4 Improvement, methodology, query or suggestion
2 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 IUCr journals

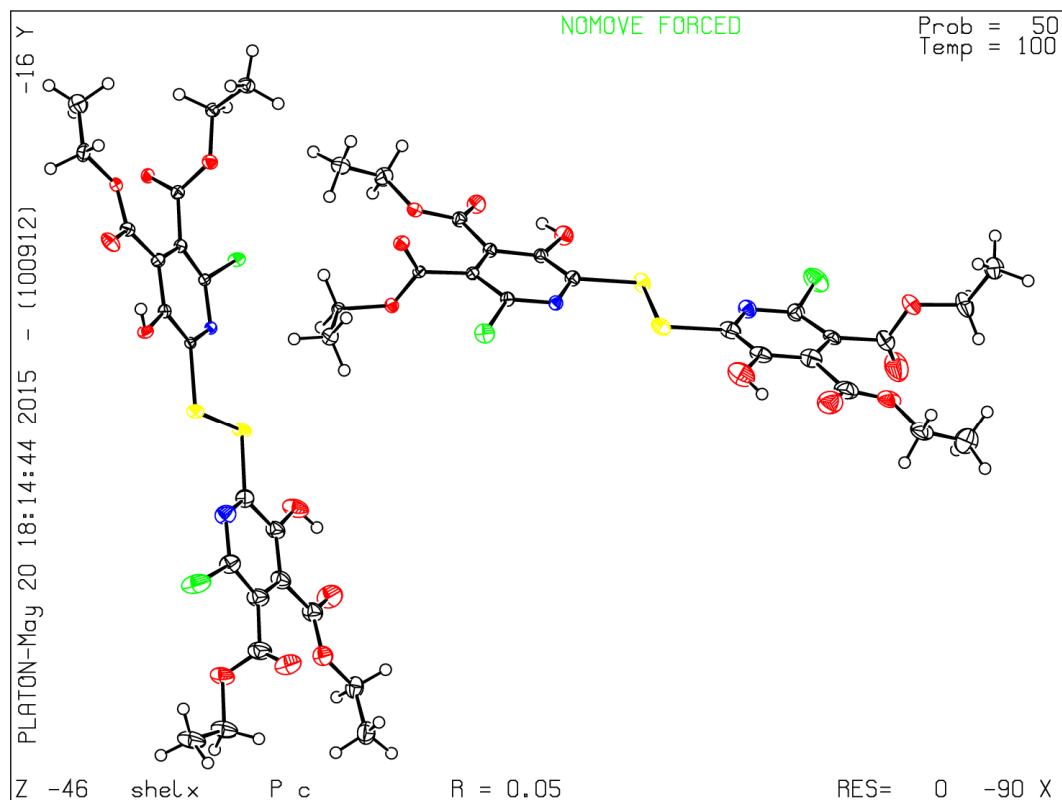
A basic structural check has been run on your CIF. These basic checks will be run on all CIFs submitted for publication in IUCr journals (*Acta Crystallographica*, *Journal of Applied Crystallography*, *Journal of Synchrotron Radiation*); however, if you intend to submit to *Acta Crystallographica Section C* or *E*, you should make sure that [full publication checks](#) are run on the final version of your CIF prior to submission.

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 21/04/2015; check.def file version of 09/03/2015

Datablock shelx - ellipsoid plot



CheckCIF/PLATON (standard) report for compound 7

Datablock: shelx

Bond precision:	C-C = 0.0052 Å	Wavelength=0.71073
Cell:	a=18.5268(6) b=8.4433(4) c=16.9773(7)	
	alpha=90 beta=102.391(2) gamma=90	
Temperature:	120 K	

	Calculated	Reported
Volume	2593.85(18)	2593.85(18)
Space group	P 21/c	P 21/c
Hall group	-P 2ybc	-P 2ybc
Moiety formula	C22 H22 Br2 N2 O10 Se	C22 H22 Br2 N2 O10 Se
Sum formula	C22 H22 Br2 N2 O10 Se	C22 H22 Br2 N2 O10 Se
Mr	713.18	713.19
Dx, g cm ⁻³	1.826	1.826
Z	4	4
Mu (mm ⁻¹)	4.586	4.586
F000	1408.0	1408.0
F000'	1406.13	
h, k, lmax	26, 12, 24	24, 11, 24
Nref	8291	7379
Tmin, Tmax	0.397, 0.577	0.513, 0.746
Tmin'	0.367	

Correction method= # Reported T Limits: Tmin=0.513
Tmax=0.746 AbsCorr = MULTI-SCAN

Data completeness= 0.890 Theta(max)= 31.017

R(reflections)= 0.0489(5533) wR2(reflections)= 0.1163(7379)

S = 1.038 Npar= 340

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

[PLAT094_ALERT_2_C](#) Ratio of Maximum / Minimum Residual Density 2.45 Report

[PLAT480_ALERT_4_C](#) Long H...A H-Bond Reported H1B .. O4B .. 2.62 Ång.

And 2 other PLAT480 Alerts

More ...

[PLAT906_ALERT_3_C](#) Large K value in the Analysis of Variance 3.253 Check

[PLAT911_ALERT_3_C](#) Missing # FCF Refl Between THmin & STh/L= 0.600 79 Report

[PLAT971_ALERT_2_C](#) Check Calcd Residual Density 0.90Å From Se1 2.05 eÅ⁻³

[PLAT971_ALERT_2_C](#) Check Calcd Residual Density 0.69Å From Br1B 1.66 eÅ⁻³

● Alert level G

[PLAT007_ALERT_5_G](#) Number of Unrefined Donor-H Atoms 2 Report

[PLAT083_ALERT_2_G](#) SHELXL Second Parameter in WGHT Unusually Large. 7.28 Why ?

[PLAT432_ALERT_2_G](#) Short Inter X...Y Contact O2B .. C9B .. 2.75 Ång.

[PLAT720_ALERT_4_G](#) Number of Unusual/Non-Standard Labels 10 Note

PLAT910_ALERT_3_G Missing # of FCF Reflection(s) Below Th(Min) ...	1 Report
PLAT912_ALERT_4_G Missing # of FCF Reflections Above STh/L= 0.600	703 Note
PLAT950_ALERT_5_G Calculated (ThMax) and CIF-Reported Hmax Differ	2 Units
PLAT956_ALERT_1_G Calculated (ThMax) and Actual (FCF) Hmax Differ	2 Units

-
- 0 **ALERT level A** = Most likely a serious problem - resolve or explain
0 **ALERT level B** = A potentially serious problem, consider carefully
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- 1 ALERT type 1 CIF construction/syntax error, inconsistent or missing data
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3 ALERT type 3 Indicator that the structure quality may be low
5 ALERT type 4 Improvement, methodology, query or suggestion
2 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 IUCr journals

A basic structural check has been run on your CIF. These basic checks will be run on all CIFs submitted for publication in IUCr journals (*Acta Crystallographica*, *Journal of Applied Crystallography*, *Journal of Synchrotron Radiation*); however, if you intend to submit to *Acta Crystallographica Section C* or *E*, you should make sure that [full publication checks](#) are run on the final version of your CIF prior to submission.

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.

Datablock shelx - ellipsoid plot

