Synthesis, Biological Evaluation and Molecular Modeling of 2-Hydroxyisoquinoline-1,3-dione Analogues as Inhibitors of HIV Reverse Transcriptase Associated Ribonuclease H and Polymerase

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Additional docking to NNRTI binding pocket

PDB	Co-crystallized	Compound 8c		Compound 9c	
Structure	NNRTI				
		Docking	Main Side Chain	Docking	Side Chain
		score	Contacts	score	Contacts
3LP1	Nevirapine	-8.32		-7.44	
3V81	Nevirapine	-8.65		-7.24	
1KLM	Delavirdine	-9.64	Asn103, Tyr181, Tyr188, Trp229	-9.91	Asn103, Tyr181, Tyr188, Trp229
2YNG	GSK560	-9.38		-9.78	

LID for best docked poses at RNase H active site

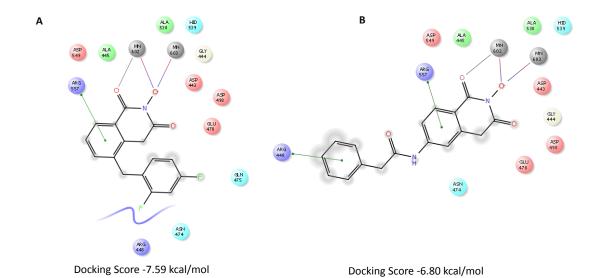


Fig. S1. Ligand Interaction Diagram for best poses of **8c** and **9c** docked into RNase H active center: A) Compound **8c** B) Compound **9c**; LID legend: red – charged (negative) residues; blue – charged (positive) residues; cyan – polar residues; green – hydrophobic residue.

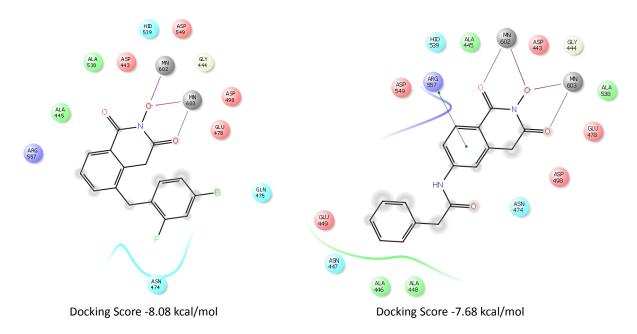


Fig. S2. Ligand Interaction Diagram for best poses of **8c** and **9c** docked into RNase H active center with R448A mutation: A) Compound **8c** B) Compound **9c**; LID legend: red – charged (negative) residues; blue – charged (positive) residues; cyan – polar residues; green – hydrophobic residue.

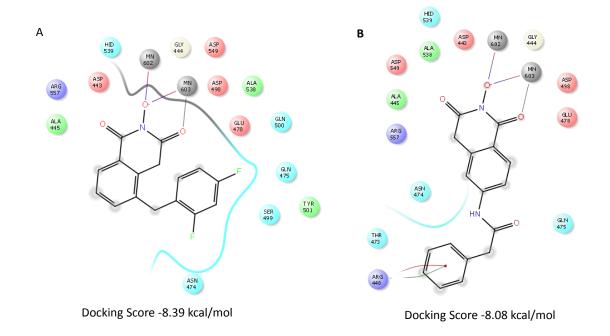


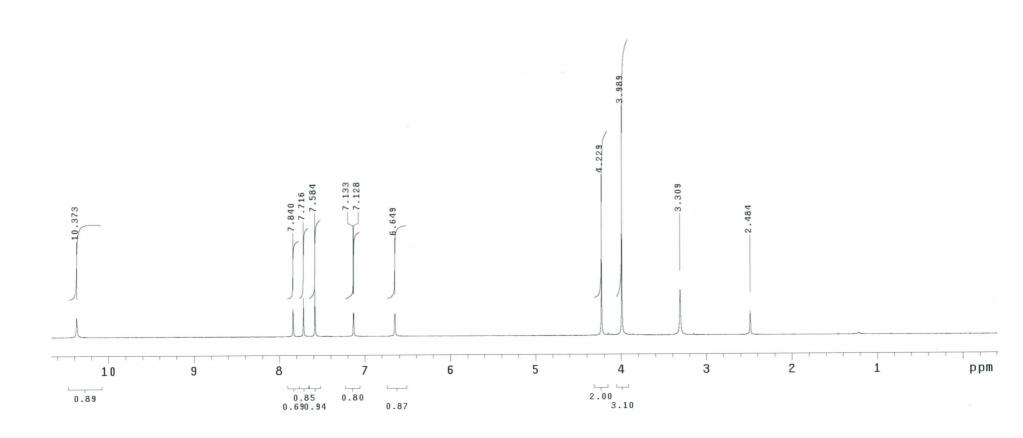
Fig. S3. Ligand Interaction Diagram for best poses of **8c** and **9c** docked into RNase H active center with flexible Arg448 side chain: A) Compound **8c** B) Compound **9c**; LID legend: red – charged (negative) residues; blue – charged (positive) residues; cyan – polar residues; green – hydrophobic residue.

Sample Name:
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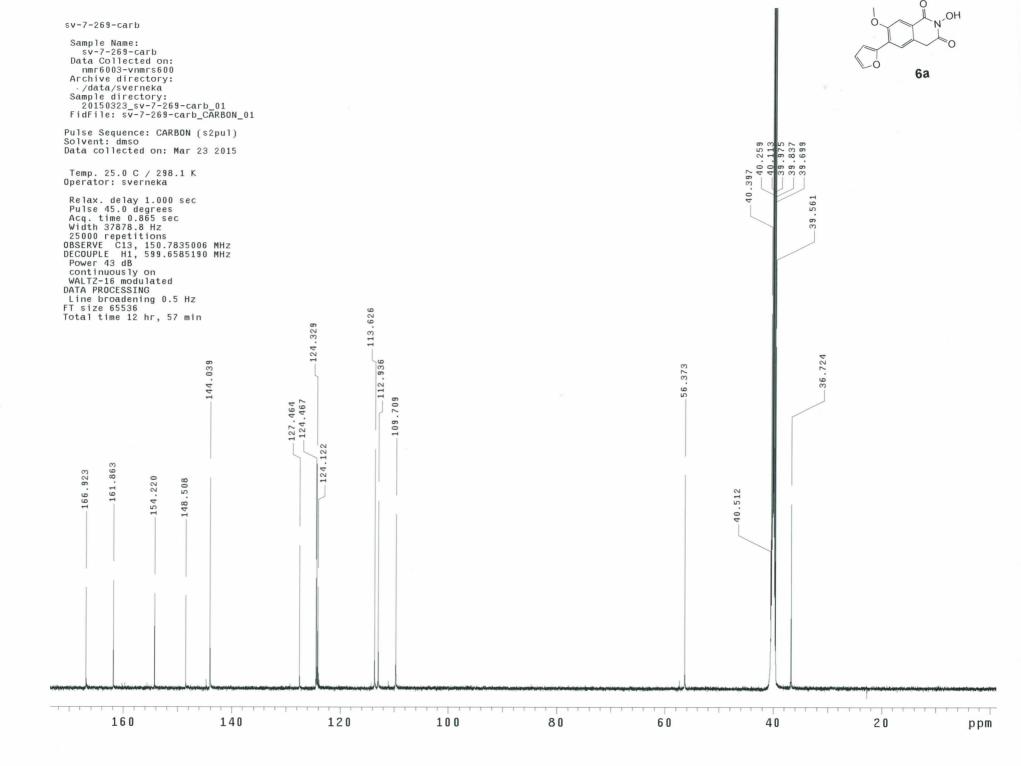
Pulse Sequence: PROTON (s2pul) Solvent: dmso Data collected on: Mar 23 2015

Temp. 25.0 C / 298.1 K Operator: sverneka

Relax. delay 1.000 sec Pulse 45.0 degrees Acq. time 1.704 sec Width 9615.4 Hz 512 repetitions OBSERVE H1, 599.6555207 MHz DATA PROCESSING FT size 32768 Total time 23 min



6a



FT size 32768

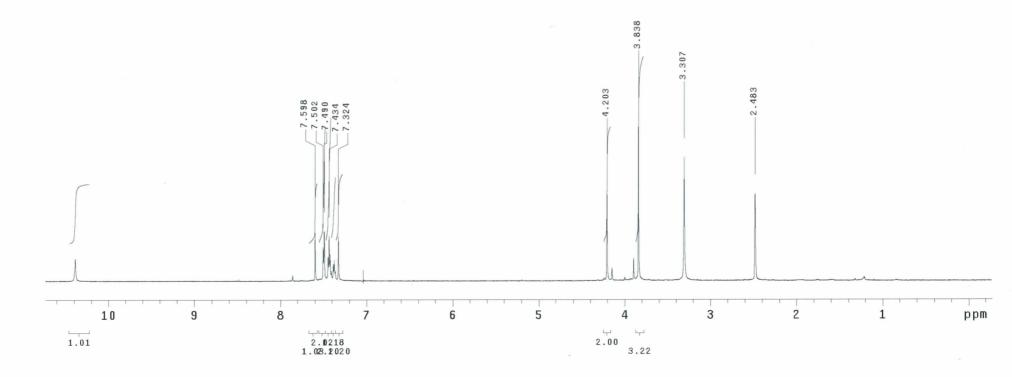
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Temp. 25.0 C / 298.1 K
Operator: sverneka

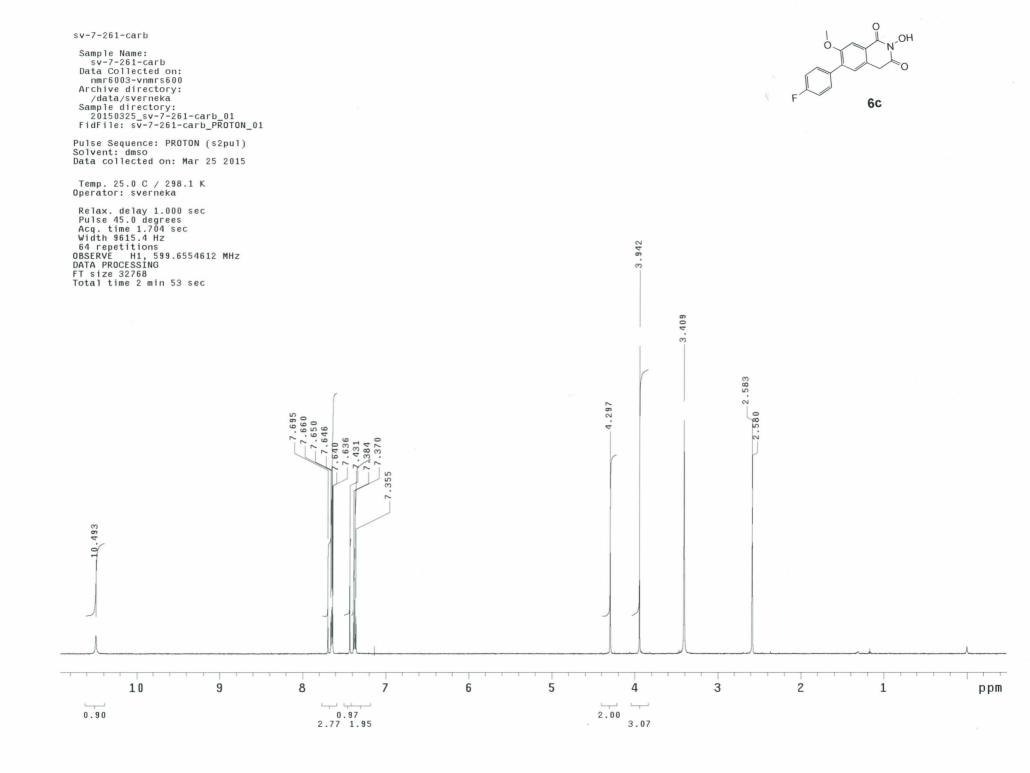
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64 repetitions
OBSERVE H1, 599.6555207 MHz
DATA PROCESSING

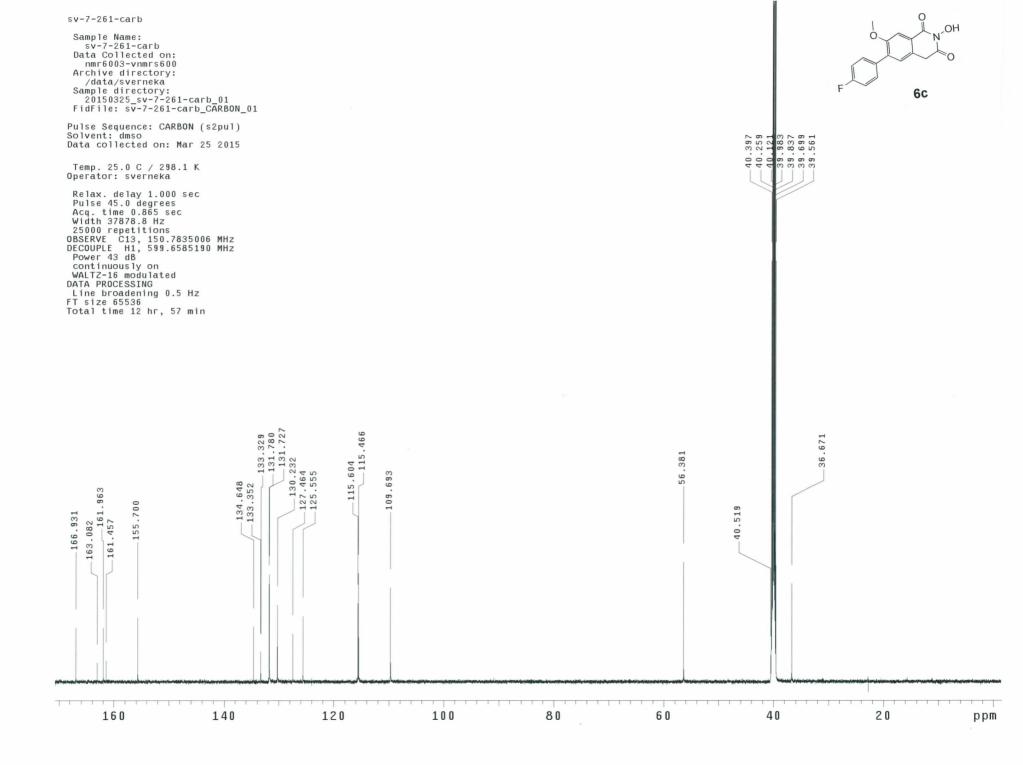
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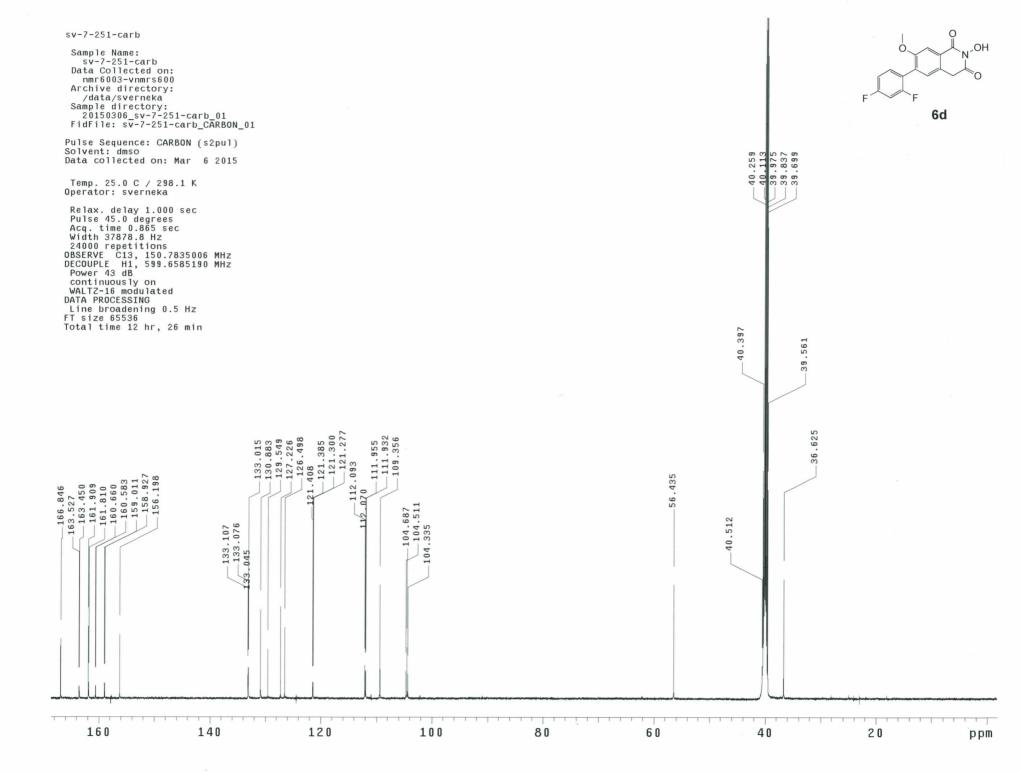


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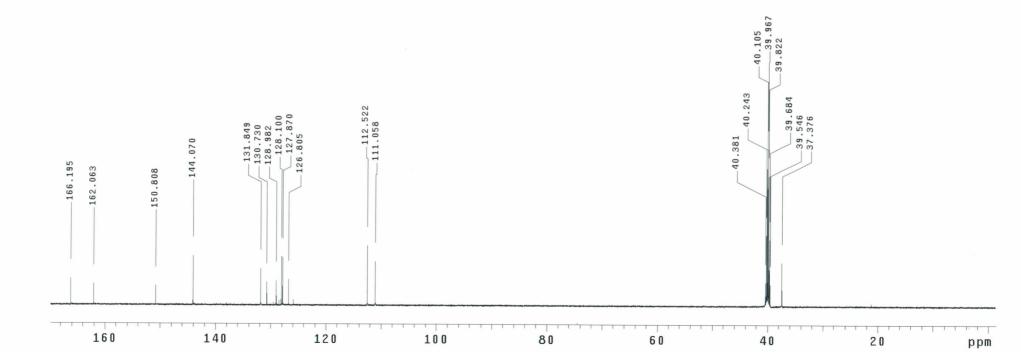
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Archive directory:
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Temp. 25.0 C / 298.1 K Operator: sverneka

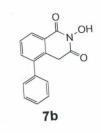
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Power 36 dB
continuously on
WALTZ-16 modulated
DATA PROCESSING
Line broadening 0.5 Hz
FT size 65536
Total time 1 hr, 2 min

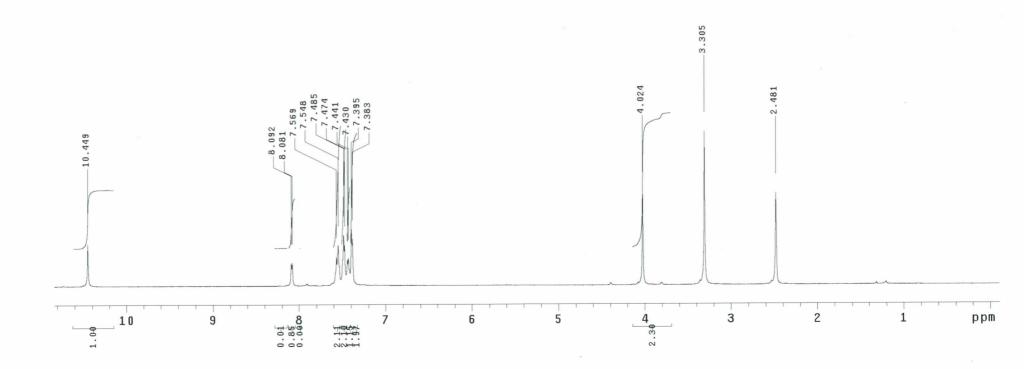


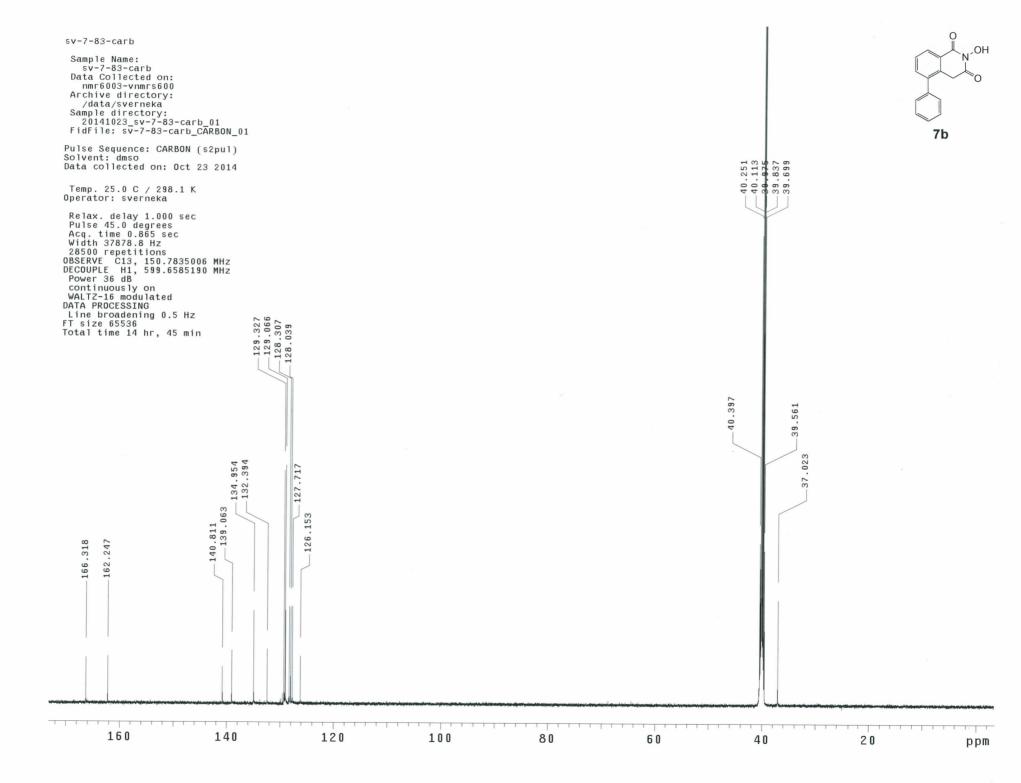
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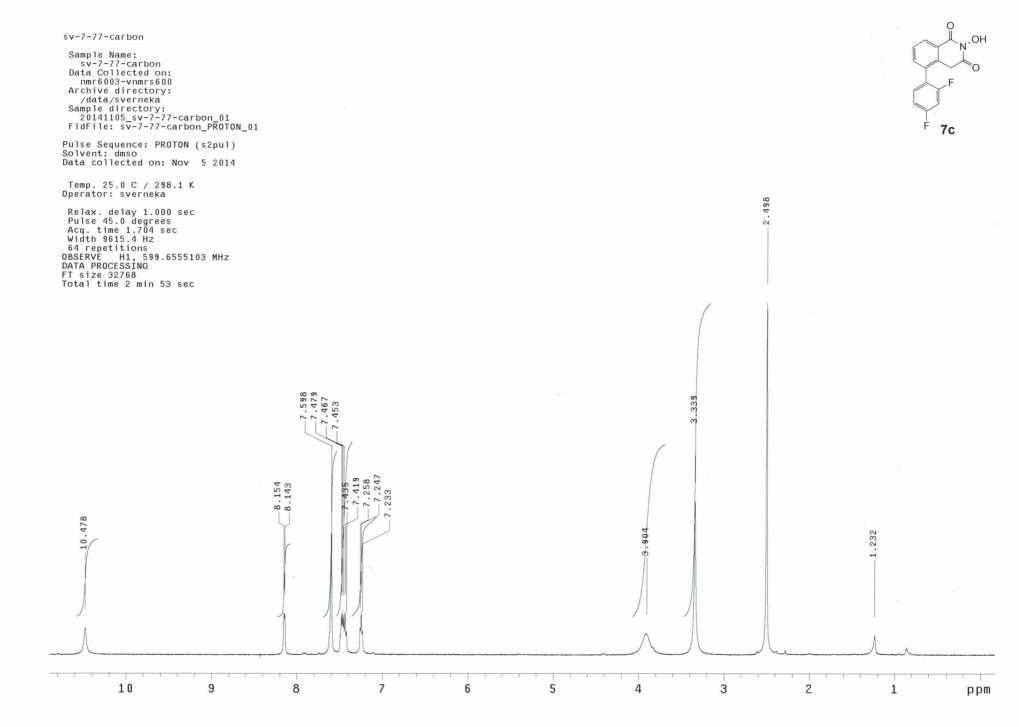
Temp. 25.0 C / 298.1 K Operator: sverneka

Relax. delay 1.000 sec Pulse 45.0 degrees Acq. time 1.704 sec Width 9615.4 Hz 64 repetitions OBSERVE H1, 599.6555207 MHz DATA PROCESSING FT size 32768 Total time 2 min 53 sec









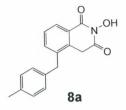
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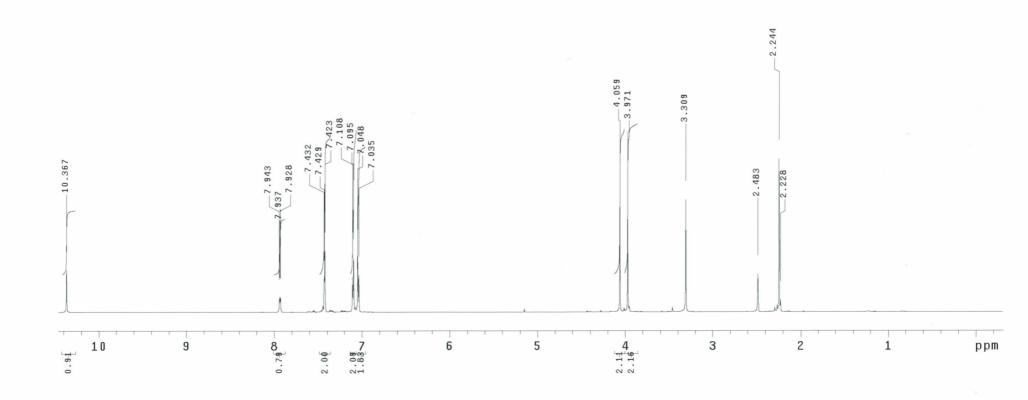
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Pulse Sequence: PROTON (s2pul) Solvent: dmso Data collected on: Sep 24 2014

Temp. 25.0 C / 298.1 K Operator: sverneka

Relax. delay 1.000 sec Pulse 45.0 degrees Acq. time 1.704 sec Width 9615.4 Hz 32 repetitions OBSERVE H1, 599.6555207 MHz DATA PROCESSING FT size 32768 Total time 1 min 27 sec





Sample Name:
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Data Collected on:
nmr6003-vnmrs600
Archive directory:
/data/sverneka
Sample directory:
20140924_sv-7-63-carbon_01
FidFile: sv-7-63-carbon_CARBON_01

Pulse Sequence: CARBON (s2pul) Solvent: dmso Data collected on: Sep 24 2014

Temp. 25.0 C / 298.1 K Operator: sverneka

Relax. delay 1.000 sec
Pulse 45.0 degrees
Acq. time 0.865 sec
Width 37878.8 Hz
24000 repetitions
OBSERVE C13, 150.7835006 MHz
DECOUPLE H1, 599.6585190 MHz
Power 36 dB
continuously on
WALTZ-16 modulated
DATA PROCESSING
Line broadening 0.5 Hz
FT size 65536
Total time 12 hr, 26 min

162.254

160

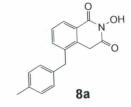
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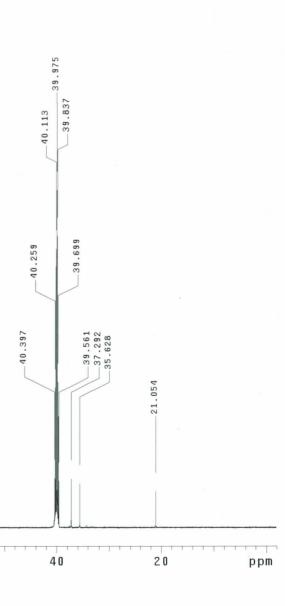
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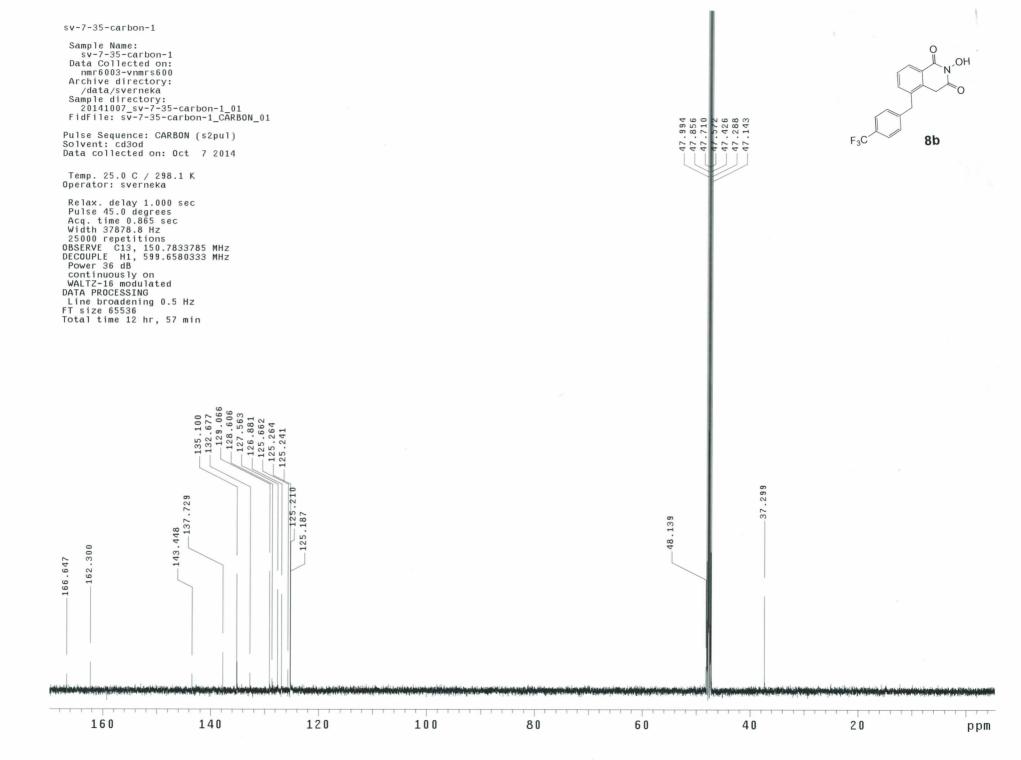
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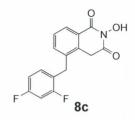


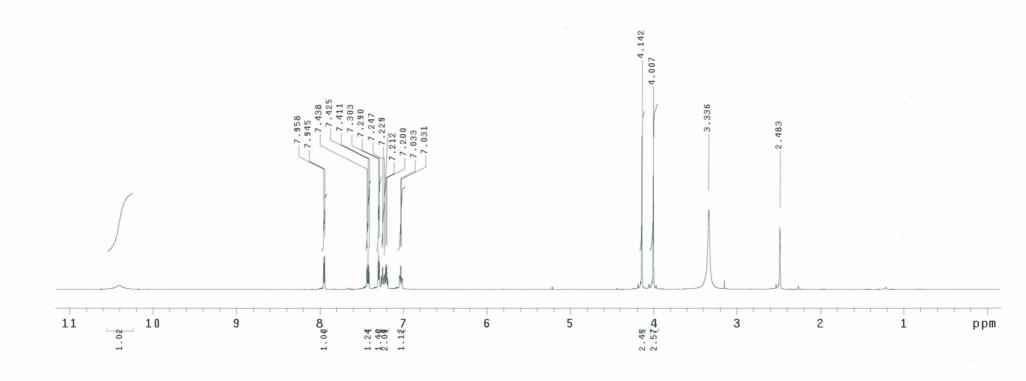
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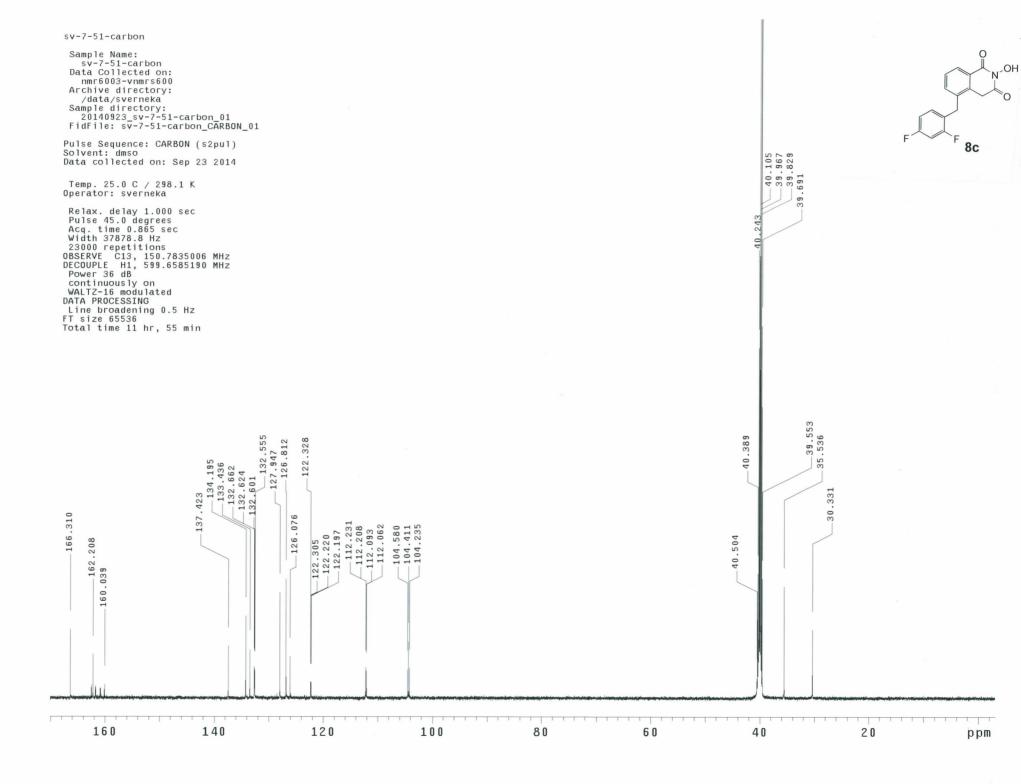
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Temp. 25.0 C / 298.1 K Operator: sverneka

Relax. delay 1.000 sec Pulse 45.0 degrees Acq. time 1.704 sec Width 9615.4 Hz 32 repetitions OBSERVE H1, 599.6555207 MHz DATA PROCESSING FT size 32768 Total time 1 min 27 sec





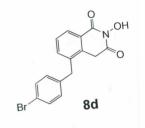


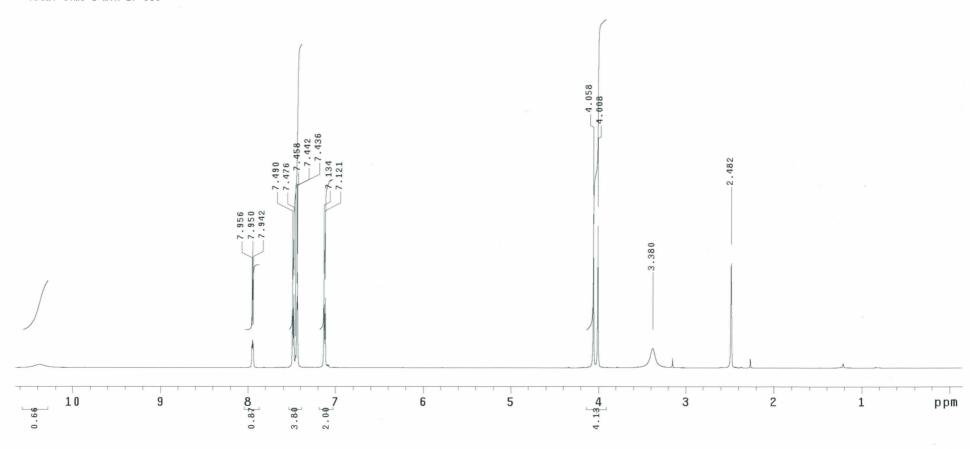
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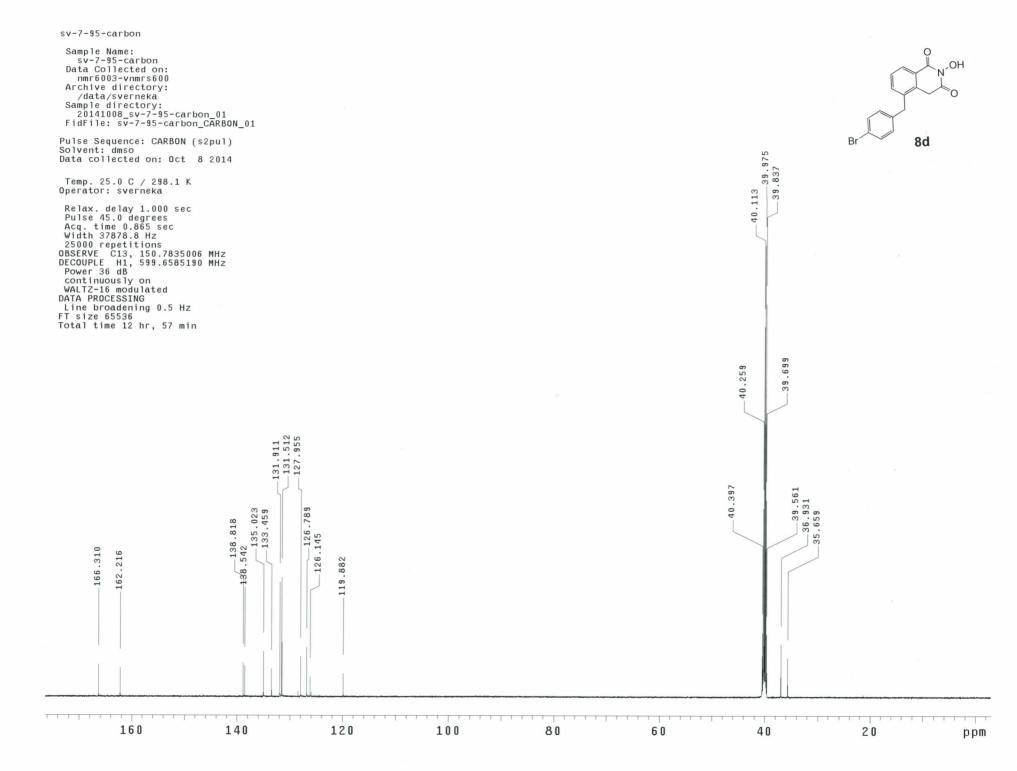
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Temp. 25.0 C / 298.1 K Operator: sverneka

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1.13

2.03

1.06 1.99

1.00 2.00

