

*Supporting Information*

Comparing hydrazine-derived reactive groups as inhibitors of quinone-dependent amine oxidases

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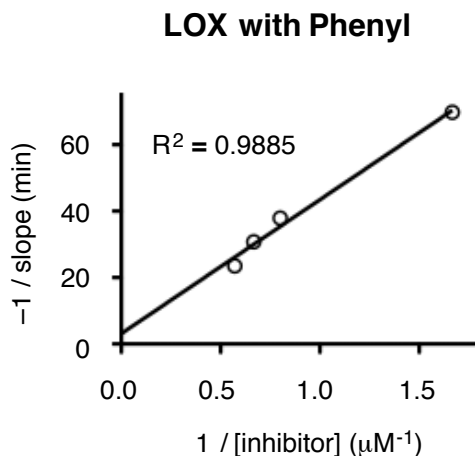
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## 1. Analysis of Previous Data

The kinetics of inhibition of LOX with phenyl hydrazine has been previously studied by Williamson and coworkers (1). Their data (from their Figure 3A) was analyzed using the Kitz–Wilson method that we have used to analyze our own data. (The Y axis was converted to  $\log(\% \text{ activity remaining})$ , the slopes were measured,  $1/\text{slope}$  vs  $1/[\text{inhibitor}]$  was plotted, and a line of best fit was calculated with Prism, Supplementary Figure S1).



Supplementary Figure S1. Kitz–Wilson analysis of data from Williamson and coworkers for phenyl hydrazine inhibiting LOX.

### Kinetics Parameters for Inhibiting LOX with Phenyl Inhibitor 1

Kitz–Wilson Y-Intercept =  $3.1 \pm 3.1$  min

Kitz–Wilson slope =  $40 \pm 3$  min  $\cdot \mu\text{M}$

$k_2 = 0.32 \text{ min}^{-1}$  (error range  $> 0.16 \text{ min}^{-1}$ )

$KI^{-1} = 13 \mu\text{M}$  (error range  $> 6 \mu\text{M}$ )

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<sup>1</sup> Williamson P R, Kittler J M, Thanassi J W, Kagan H M. Reactivity of a functional carbonyl moiety in bovine aortic lysyl oxidase. *Biochem. J.* 1986;235:597–605.

## 2. Results of Inhibition Assay

### Average Percent Activity Remaining for Inhibiting LSDAO with Phenyl Inhibitor 1

Minutes	0.005 $\mu\text{M}$	0.0075 $\mu\text{M}$	0.010 $\mu\text{M}$	0.015 $\mu\text{M}$	0.03 $\mu\text{M}$	0.1 $\mu\text{M}$	0.5 $\mu\text{M}$
10	80	62	50	13	0	0	0

### Average Percent Activity Remaining for Inhibiting LOX with Phenyl Inhibitor 1

Minutes	1 $\mu\text{M}$	2 $\mu\text{M}$	20 $\mu\text{M}$
10	83	73	17

### Average Percent Activity Remaining for Inhibiting LSDAO with Hydrazide Inhibitor 3

Minutes	1 $\mu\text{M}$	2 $\mu\text{M}$	4 $\mu\text{M}$	8 $\mu\text{M}$	14 $\mu\text{M}$	20 $\mu\text{M}$	40 $\mu\text{M}$	500 $\mu\text{M}$
4	90	80	71	54	40	33	4	0.7
10	76	52	33	13	8	4		

### Kinetics Parameters for Inhibiting LSDAO with Hydrazide Inhibitor 3

Kitz–Wilson Y-Intercept = 1.5 +/- 1.2 min  
Kitz–Wilson slope = 81 +/- 3 min  $\cdot$   $\mu\text{M}$   
 $k_2 = 0.67 \text{ min}^{-1}$  (error range = 0.37–3.3  $\text{min}^{-1}$ )  
 $K_i^{-1} = 54 \mu\text{M}$  (error range = 29–280  $\mu\text{M}$ )

### Average Percent Activity Remaining for Inhibiting LOX with Hydrazide Inhibitor 3

Minutes	4 $\mu\text{M}$	7 $\mu\text{M}$	20 $\mu\text{M}$	100 $\mu\text{M}$	500 $\mu\text{M}$
4	93	78	69	27	8
10	84	75	48	3	

### Kinetics Parameters for Inhibiting LOX with Hydrazide Inhibitor 3

Kitz–Wilson Y-Intercept = 2.5 +/- 1.8 min  
Kitz–Wilson slope = 496 +/- 14 min  $\cdot$   $\mu\text{M}$   
 $k_2 = 0.40 \text{ min}^{-1}$  (error range = 0.23–1.4  $\text{min}^{-1}$ )  
 $K_i^{-1} = 200 \mu\text{M}$  (error range = 110–730  $\mu\text{M}$ )

### Average Percent Activity Remaining for Inhibiting LSDAO with Alkyl Inhibitor 4

Minutes	0.03 $\mu\text{M}$	0.08 $\mu\text{M}$	0.1 $\mu\text{M}$	0.2 $\mu\text{M}$	0.5 $\mu\text{M}$	1 $\mu\text{M}$
4	81	60	43	22	1	1
10	45	15	7	2		

### Kinetics Parameters for Inhibiting LSDAO with Alkyl Inhibitor 4

Kitz–Wilson Y-Intercept = 1.0 +/- 0.4 min  
Kitz–Wilson slope = 0.88 +/- 0.02 min  $\cdot$   $\mu\text{M}$   
 $k_2 = 1.0 \text{ min}^{-1}$  (error range = 0.75–1.6  $\text{min}^{-1}$ )  
 $K_i^{-1} = 0.9 \mu\text{M}$  (error range = 0.6–1.4  $\mu\text{M}$ )

#### Average Percent Activity Remaining for Inhibiting LOX with Alkyl Inhibitor 4

Minutes	7 $\mu\text{M}$	20 $\mu\text{M}$	100 $\mu\text{M}$
4	89	82	59
10	83	63	30

#### Kinetics Parameters for Inhibiting LOX with Alkyl Inhibitor 4

Kitz-Wilson Y-Intercept = 12.6 +/- 1.1 min  
Kitz-Wilson slope = 719 +/- 13 min  $\cdot$   $\mu\text{M}$   
 $k_2 = 0.079 \text{ min}^{-1}$  (error range = 0.073–0.087  $\text{min}^{-1}$ )  
 $K_I^{-1} = 57 \mu\text{M}$  (error range = 52–64  $\mu\text{M}$ )

#### Average Percent Activity Remaining for Inhibiting LSDAO with Semicarbazide Inhibitor 5

Minutes	1 $\mu\text{M}$	2 $\mu\text{M}$	4 $\mu\text{M}$	20 $\mu\text{M}$	100 $\mu\text{M}$
4	89	78	69	41	1
10	84	71	50	7	

#### Kinetics Parameters for Inhibiting LSDAO with Semicarbazide Inhibitor 5

Kitz-Wilson Y-Intercept = 1.1 +/- 0.9 min  
Kitz-Wilson slope = 123 +/- 2 min  $\cdot$   $\mu\text{M}$   
 $k_2 = 0.91 \text{ min}^{-1}$  (error range = 0.50–5.0  $\text{min}^{-1}$ )  
 $K_I^{-1} = 112 \mu\text{M}$  (error range = 61–625  $\mu\text{M}$ )

#### Average Percent Activity Remaining for Inhibiting LOX with Semicarbazide Inhibitor 5

Minutes	40 $\mu\text{M}$	100 $\mu\text{M}$	500 $\mu\text{M}$
4	93	76	35
10	78	52	6

#### Kinetics Parameters for Inhibiting LOX with Semicarbazide Inhibitor 5

Kitz-Wilson Y-Intercept = -0.7 +/- 2.7 min  
Kitz-Wilson slope = 3806 +/- 170 min  $\cdot$   $\mu\text{M}$   
 $k_2 = >0.50 \text{ min}^{-1}$   
 $K_I^{-1} = >1000 \mu\text{M}$

### 3. General Synthetic Procedures

#### Abbreviations

AcOH = acetic acid	HOBt = hydroxybenzotriazole hydrate
DCM = dichloromethane	MeOH = methanol
DMF = <i>N,N</i> -dimethylformamide	Quant = quantitative conversion
DMSO = dimethylsulfoxide	TEA = triethylamine
EDC = <i>N</i> -(3-dimethylaminopropyl)- <i>N'</i> -ethylcarbodiimide	TFA = trifluoroacetic acid
Ether = diethyl ether	THF = tetrahydrofuran
EtOAc = ethyl acetate	y = yield

#### General Procedures

**Column chromatography** was performed with 60 Å 40–63 µm silica-P flash silica gel.

**Solvents** for reactions (DMF, DCM, THF, and toluene) were dried using a LC Technology Solutions purification system. Other solvents were used as received unless noted otherwise.

**Chemicals** were purchased from Fisher, VWR, or Sigma-Aldrich and used as received unless noted otherwise.

**NMR Spectra** were measured in CDCl<sub>3</sub> at ambient temperature unless otherwise noted.

**<sup>1</sup>H NMR** spectra were recorded on either a 600 or 200 MHz Varian spectrometer. Chemical shifts are reported in ppm (δ) relative to tetramethylsilane using the solvent as a reference (CDCl<sub>3</sub> = 7.26 ppm, DMSO-*d*<sub>6</sub> = 2.49 ppm, D<sub>2</sub>O = 4.80 ppm, CD<sub>3</sub>OD = 3.30). The following is an example data point: chemical shift (multiplicity [s = singlet, d = doublet, t = triplet, q = quartet, pent = pentet, sext = sextet, sept = septet, oct = octet, m = multiplet, br = broad, and combinations thereof], coupling constants [Hz], integration, assignment [if any]).

**<sup>13</sup>C NMR** spectra were recorded on a 600 or 200 MHz (150 or 50 MHz) Varian spectrometer with complete proton decoupling. Chemical shifts are reported in ppm (δ) relative to tetramethylsilane using the solvent or MeOH as a reference (CDCl<sub>3</sub> = 77.16 ppm, DMSO-*d*<sub>6</sub> = 39.52 ppm, CD<sub>3</sub>OD = 49.00 ppm, MeOH = 49.50).

**IR** spectra were recorded on a Perkin Elmer Spectrum 100 FT-IR spectrometer with Perkin Elmer Spectrum software. Spectra are partially reported ( $\nu_{\max}$ , cm<sup>-1</sup>).

**MS** were obtained either on an Agilent Technologies 6120 quadrupole LC/MS system with an 1260 Infinity liquid chromatography system at Clark University or at The University of Illinois Urbana-Champaign's Mass Spectrometry Center.

**TLC** was performed on 60 Å F<sub>254</sub> pre-coated silica gel plates. Samples were visualized by either ultraviolet irradiation, potassium permanganate staining, or cerium ammonium molybdate staining.

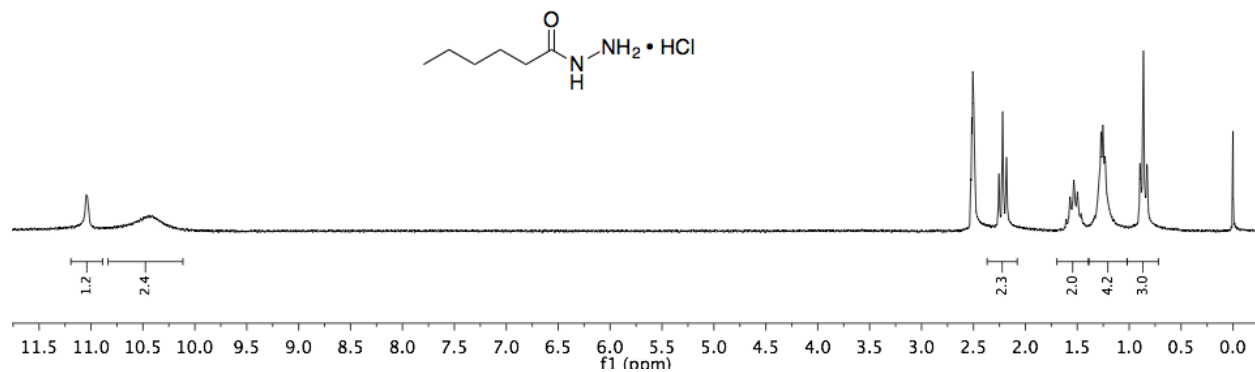
**Optical Rotations** were obtained with a Rudolph Research Autopol II automatic polarimeter.

**Yield** refers to isolated material.

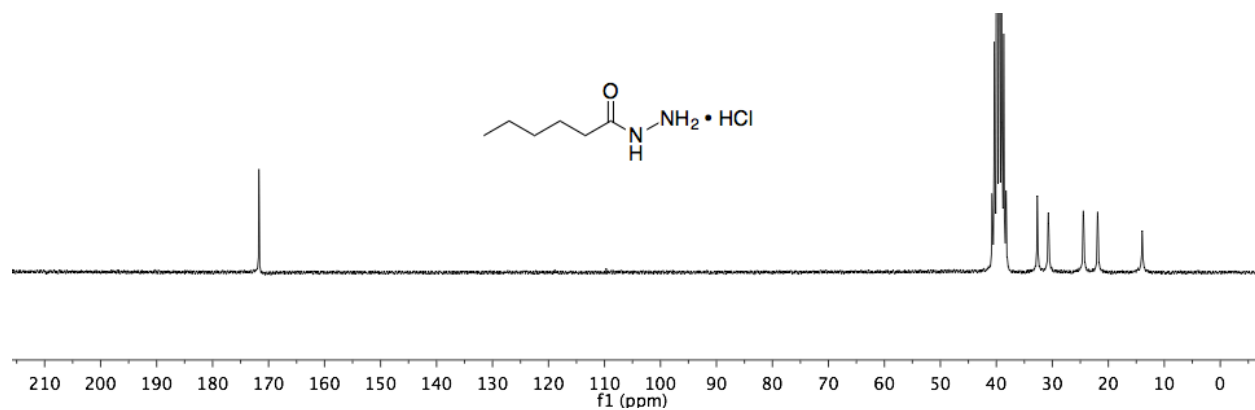
**Quantitative** recovery means that mostly pure material was recovered in approximately the expected mass, and the material was used directly for the next step without purification.

## 4. NMR Spectra of Synthesized Compounds

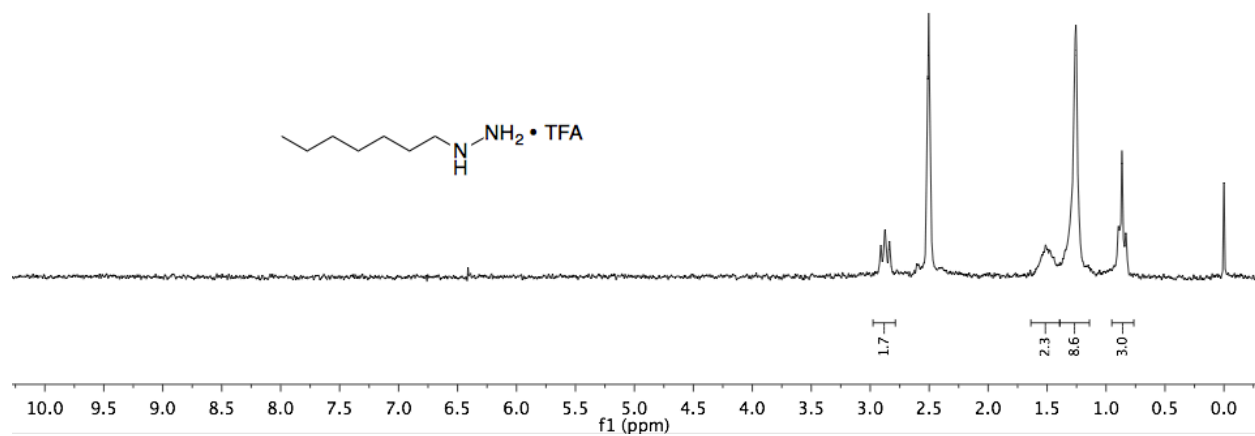
### <sup>1</sup>H NMR Spectrum of 3



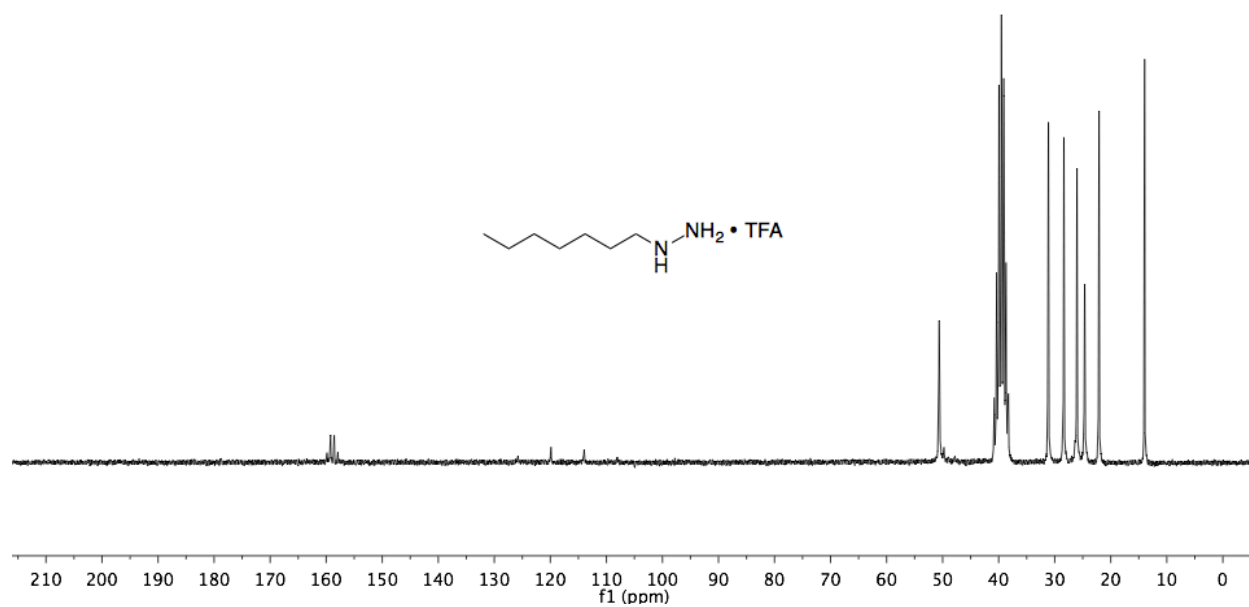
### <sup>13</sup>C NMR Spectrum of 3



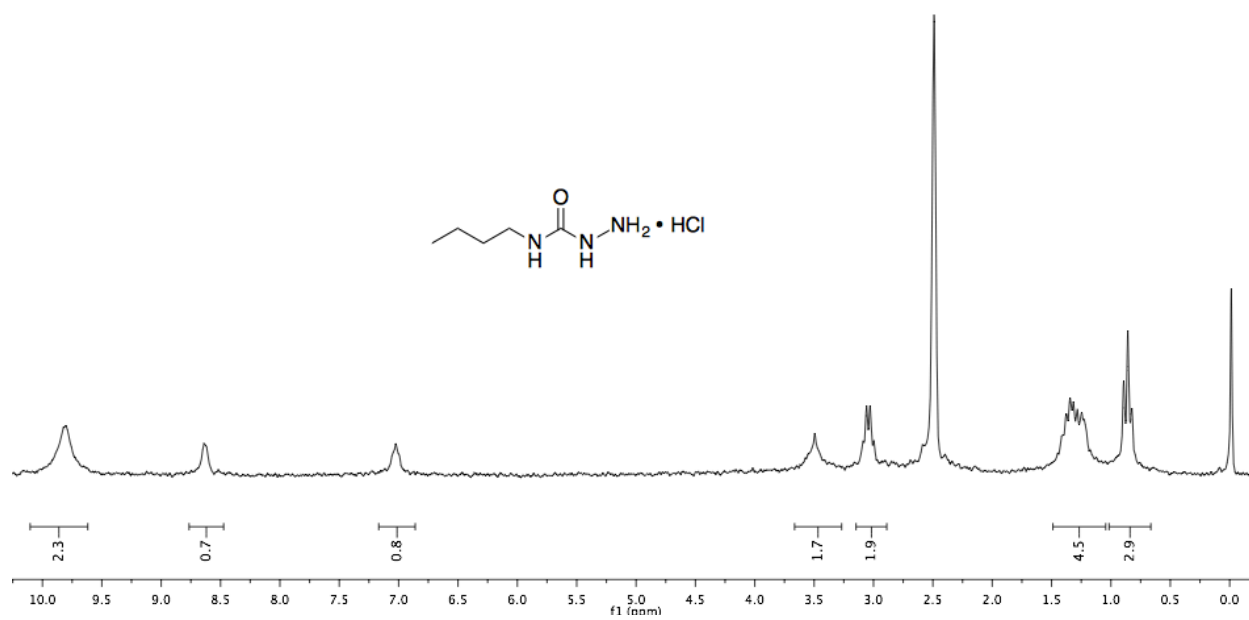
### <sup>1</sup>H NMR Spectrum of 4



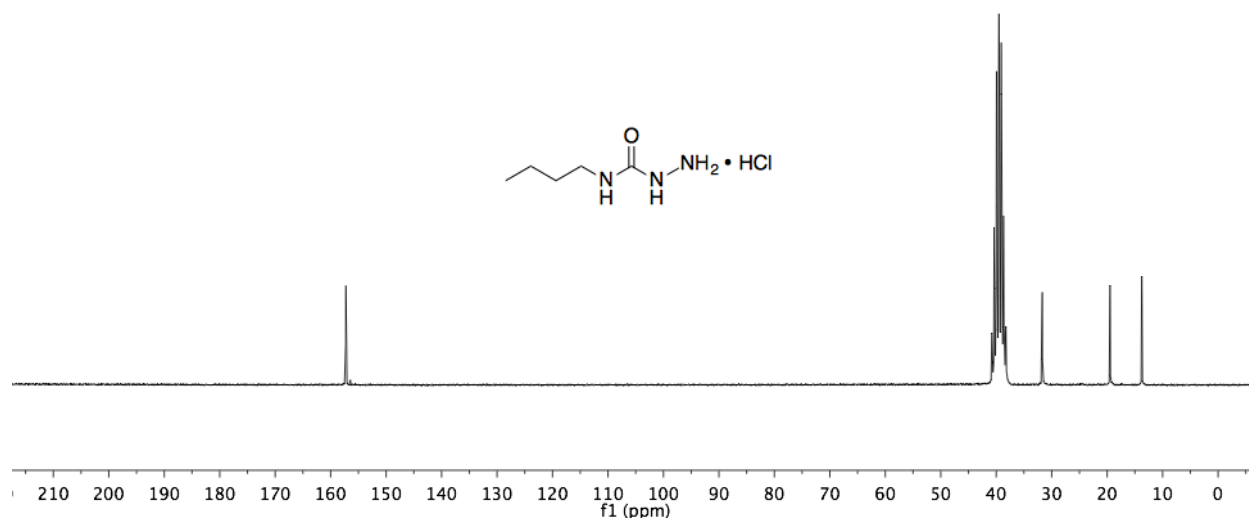
**<sup>13</sup>C NMR Spectrum of 4**



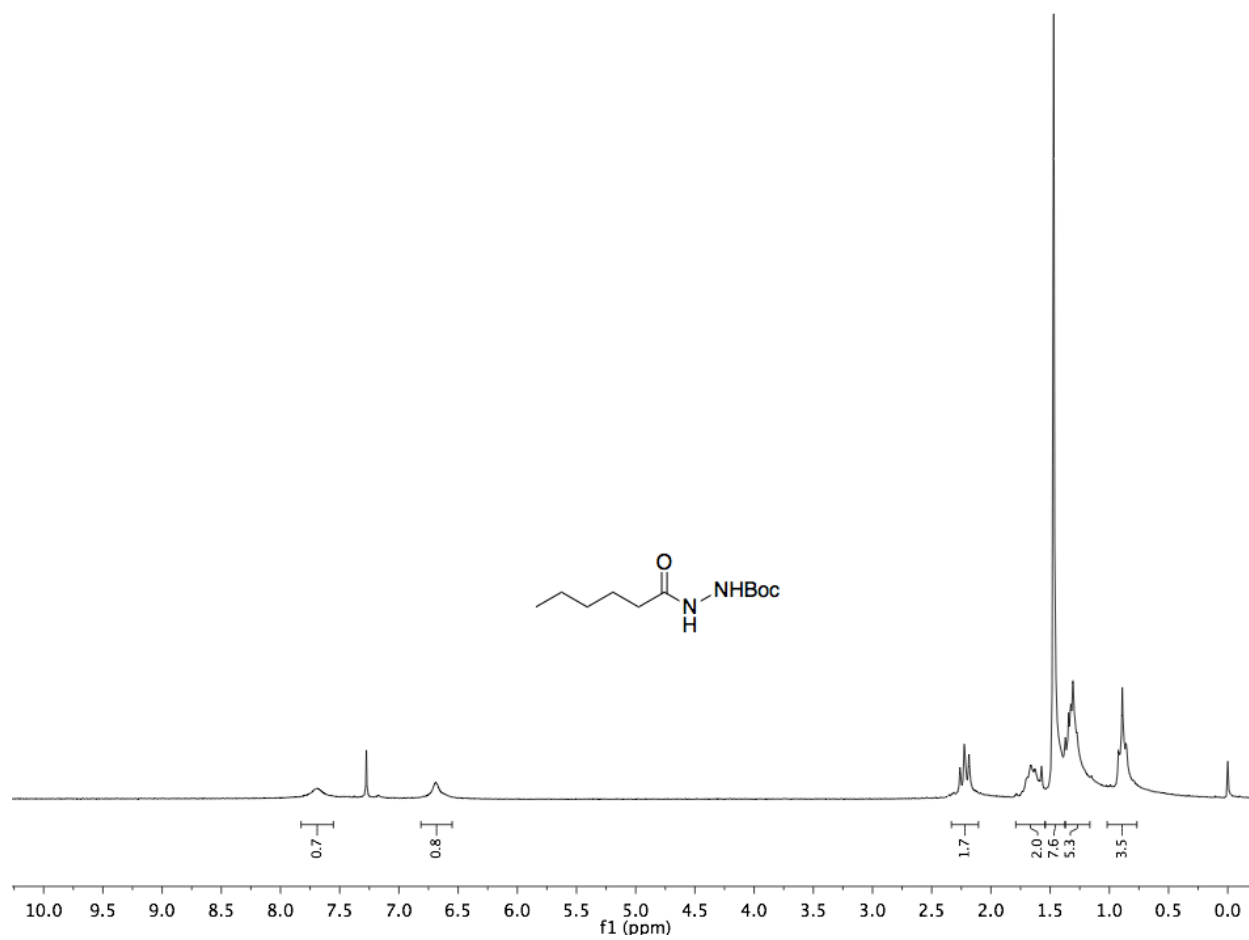
**<sup>1</sup>H NMR Spectrum of 5**



**<sup>13</sup>C NMR Spectrum of 5**

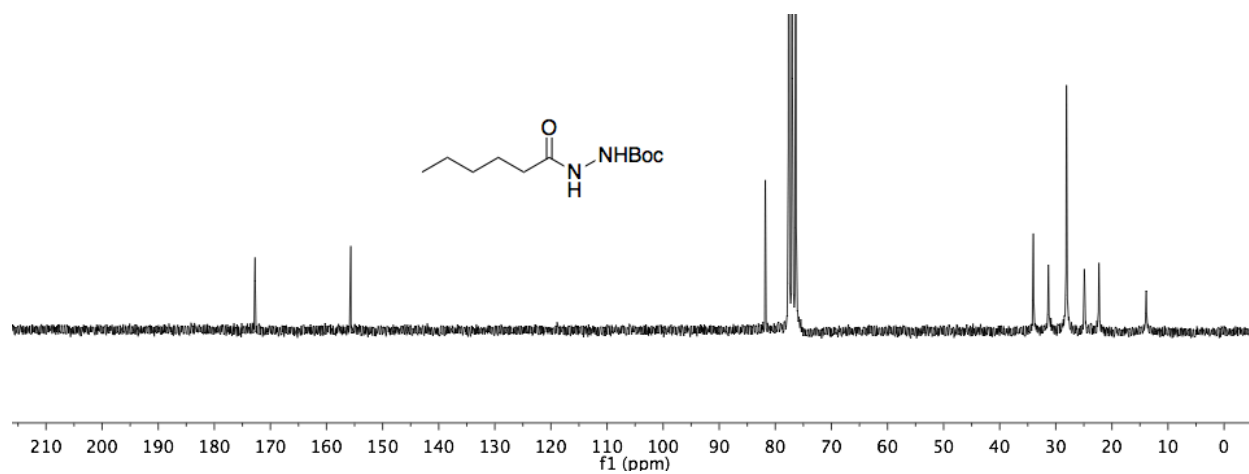


**<sup>1</sup>H NMR Spectrum of 8**

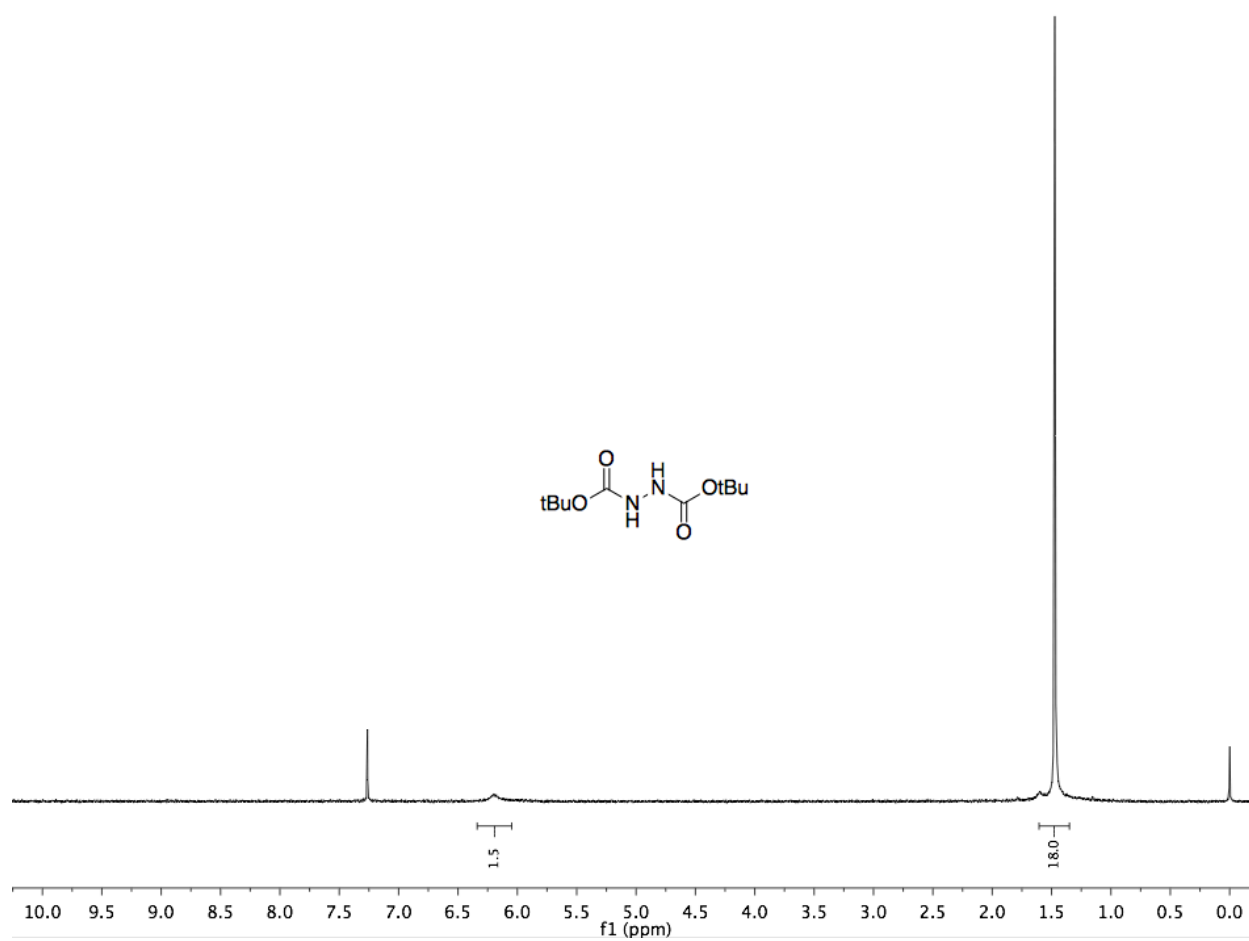




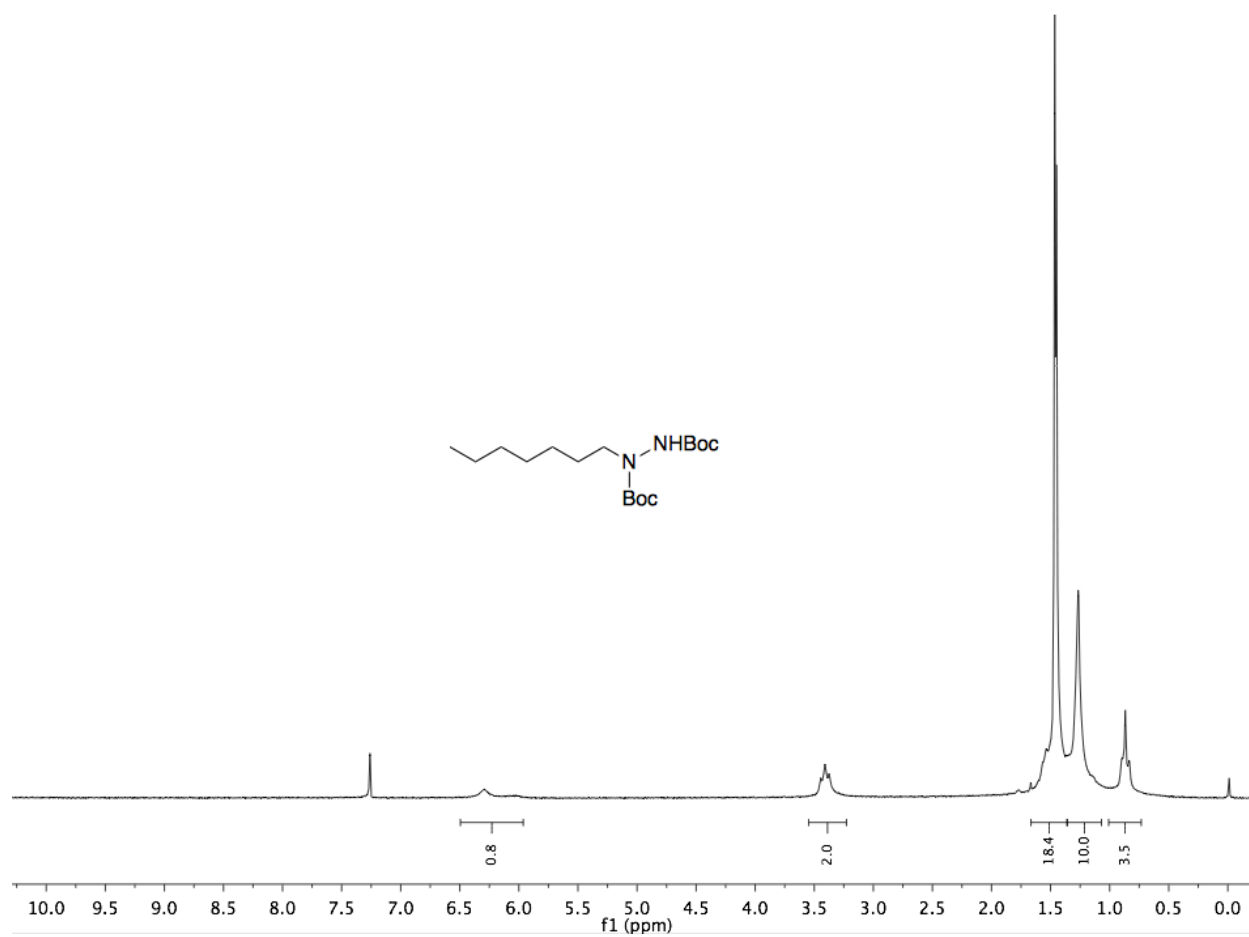
**<sup>13</sup>C NMR Spectrum of 8**



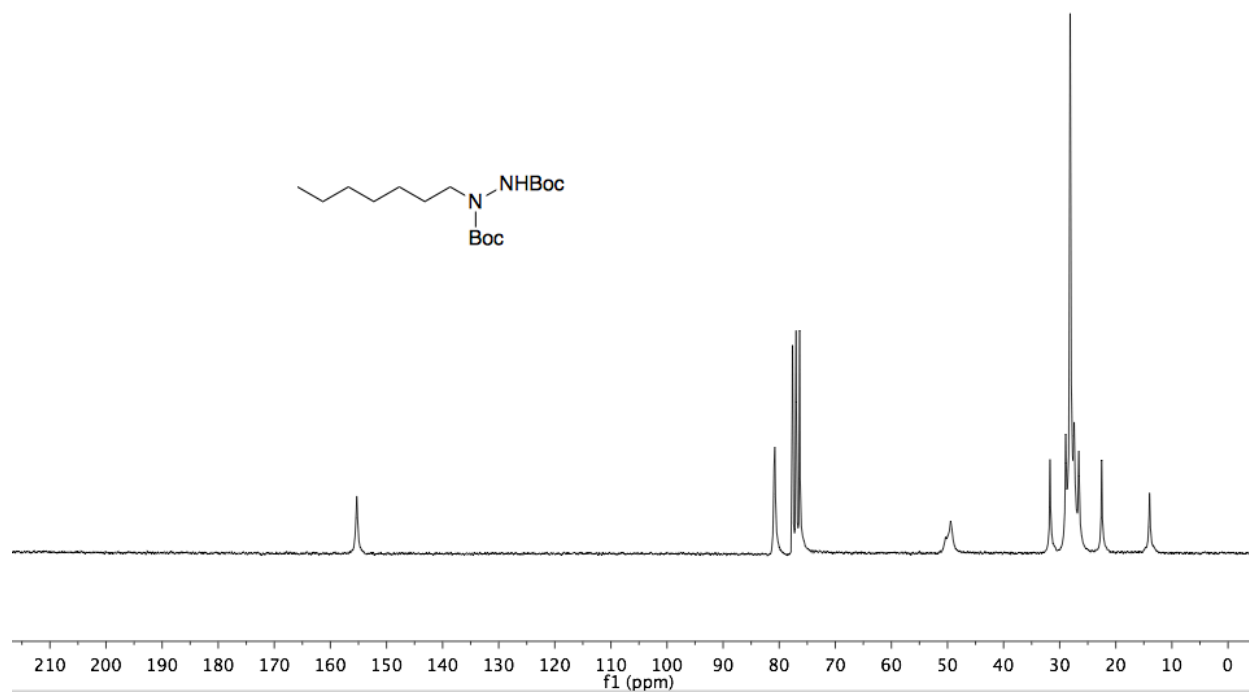
**<sup>1</sup>H NMR Spectrum of 10**



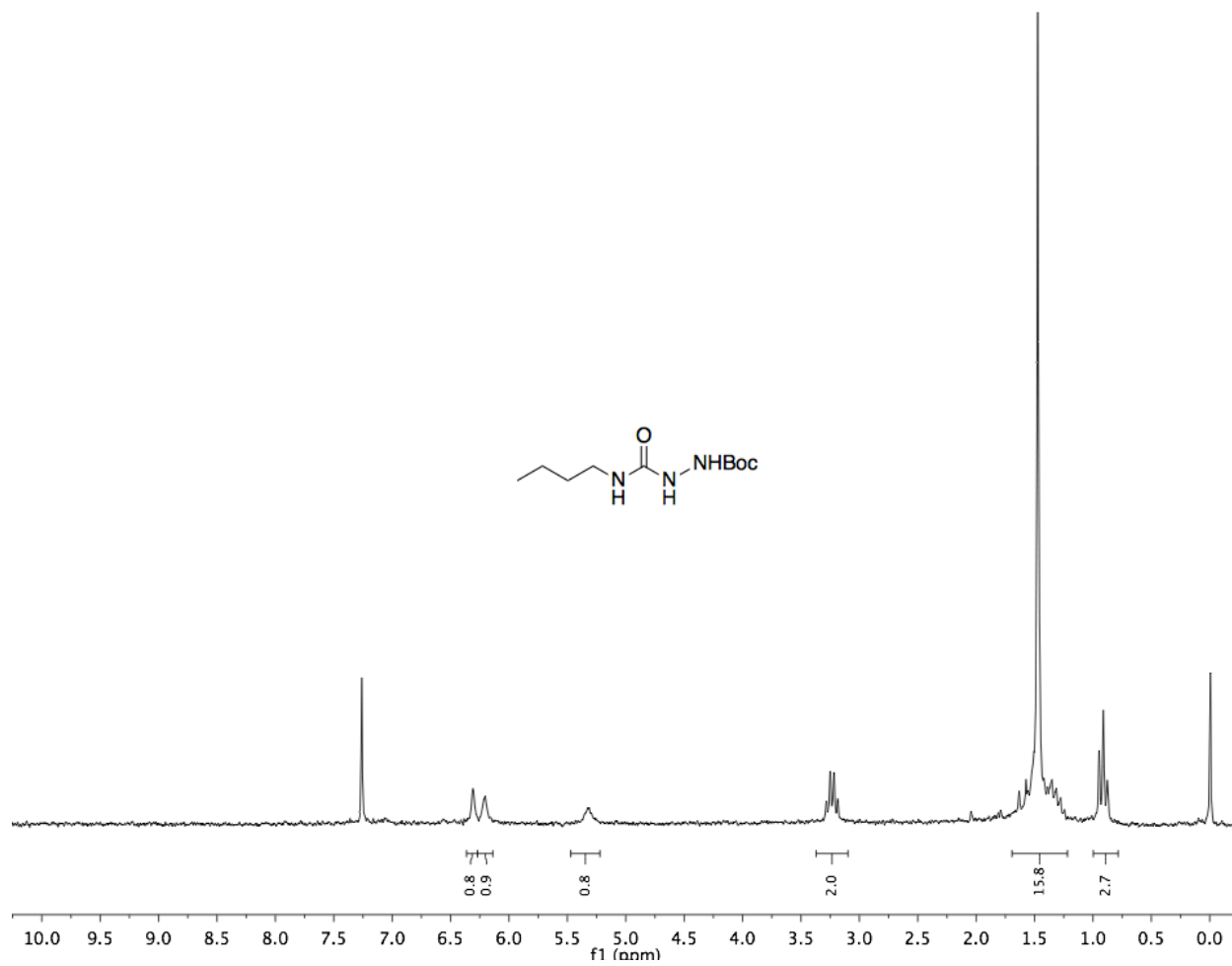
<sup>1</sup>H NMR Spectrum of 11



<sup>13</sup>C NMR Spectrum of 11



### <sup>1</sup>H NMR Spectrum of 14



### <sup>13</sup>C NMR Spectrum of 14

