## The Development of Small Molecule Inhibitors for Fibrosis Drug Discovery

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### **Supporting Information**

#### Contents

Chapter 1

- S1. <sup>1</sup>H NMR, <sup>13</sup>C NMR, IR, and HRMS Spectra
- S2. Biological Assay Results
  - a. Table S1: Summary of assay results
  - **b.** Bis-*p*NPP assay results
  - c. LPC assay control reactions
  - d. Initial screen in LPC assay: 300 nM 100  $\mu$ M of inhibitor
  - e. LPC assay results:  $30 \text{ nM} 10 \mu \text{M}$  of inhibitor
- **S3.** Physicochemical Data
- S4. Molecular Modelling
- **S5.** Crystallographic Data

### Chapter 2

- S6. <sup>1</sup>H NMR, <sup>13</sup>C NMR, IR, and HRMS Spectra
- S7. References

# S1. <sup>1</sup>H NMR, <sup>13</sup>C NMR, IR and HRMS Spectra















### 3-((6-Chloro-2-methyl-1*H*-indol-3-yl)thio)benzoic acid (1.52).









### 3-((6-Chloro-1,2-dimethyl-1*H*-indol-3-yl)thio)benzoic acid (1.53).





### 3-((6-Chloro-1-cyclopentyl-2-methyl-1*H*-indol-3-yl)thio)benzoic acid (1.54).





### 3-((6-Chloro-2-methyl-1-phenyl-1*H*-indol-3-yl)thio)benzoic acid (1.55).







### 3-((2-Methyl-1-(1-methyl-1*H*-pyrazol-4-yl)-1*H*-indol-3-yl)thio)benzoic acid (1.56).



### 3-((2-Methyl-1*H*-indol-3-yl)thio)benzoic acid (1.57).







### 3-((1,2-Dimethyl-1*H*-indol-3-yl)thio)benzoic acid (1.58).





### 3-((1-Cyclopentyl-2-methyl-1*H*-indol-3-yl)thio)benzoic acid (1.59).





### 3-((2-Methyl-1-phenyl-1*H*-indol-3-yl)thio)benzoic acid (1.60).













EPSRC National Facility Swansea LTQ Orbitrap XL LM406 MW=379? C21H18CIN3O2 (DCM)/MeOH + NH4OAc Lisa Miller 08/04/2015 12:28:46 380.1163 NL: 100<sub>∃</sub> 2.45E6 2:4560 STRWAT463-OR-HNESP#31-34 RT: 0.69-0.77 AV: 4 T: FTMS + p NSI Full ms [140.00-1935.00] 90-80-Observed Data Relative Abundance 70 60 50 40 382.1134 30-381.1197 20-10 383.1167 378.1008 384.2172 0-100 NL: 1.39E4 380.1160 C<sub>21</sub> H<sub>19</sub> ClN<sub>3</sub> O<sub>2</sub>: C<sub>21</sub> H<sub>19</sub> Cl<sub>1</sub> N<sub>3</sub> O<sub>2</sub> p (gss, s /p:40) Chrg 1 R: 100000 Res .Pwr . @FWHM 90 80 Theoretical Isotope Model: [M + H]+ 70-60 50-40 382.1131 30 381.1194 20 383.1165 10-0<sup>Ξ</sup> .... 378 380 382 384 386 388 m/z







### 3-((6-Chloro-2-methyl-1*H*-indol-3-yl)oxy)benzoic acid (1.64).





## 6-((2-Methyl-1-(1-methyl-1*H*-pyrazol-4-yl)-1*H*-indol-3-yl)methyl)picolinic acid (1.65).





## 3-((2-Methyl-1-(1-methyl-1*H*-pyrazol-4-yl)-1*H*-indol-3-yl)methyl)benzoic acid (1.66).





345.1438

345

m/z

346.1468

347.1491

348

349

347

20-10-

ΕO

342

343

344




# 3-((2-Methyl-1-(1-methyl-1*H*-pyrazol-4-yl)-1*H*-indol-3-yl)oxy)benzoic acid (1.67).



## 3-((2-Methyl-1*H*-indol-3-yl)oxy)benzoic acid (1.68).





6-Chloro-2-methyl-1-(1-methyl-1*H*-pyrazol-4-yl)-1*H*-indole (1.72).







## 1-((3-Bromophenyl)thio)propan-2-one (1.74).



## 3-((3-Bromophenyl)thio)-6-chloro-2-methyl-1*H*-indole (1.76).





## Methyl 3-((6-chloro-2-methyl-1*H*-indol-3-yl)thio)benzoate (1.77).









(1.78).





## Methyl 6-hydroxypicolinate (1.80).





## Methyl 6-(2-oxopropoxy)picolinate (1.81).





## 1-((6-Bromopyridin-2-yl)oxy)propan-2-one (1.84).





2-Methyl-1-(1-methyl-1*H*-pyrazol-4-yl)-1*H*-indole (1.87).





# 3-(3-Bromobenzyl)-6-chloro-2-methyl-1*H*-indole (1.91).







# 3-((6-Bromopyridin-2-yl)methyl)-6-chloro-2-methyl-1-(1-methyl-1*H*-pyrazol-4-yl)-1*H*-indole

(1.94).







m/z

Methyl 6-((6-chloro-2-methyl-1-(1-methyl-1*H*-pyrazol-4-yl)-1*H*-indol-3-yl)methyl)picolinate

(1.95).







## Methyl 3-((2-methyl-1*H*-indol-3-yl)thio)benzoate (1.100).











# 3-((6-Bromopyridin-2-yl)methyl)-2-methyl-1-(1-methyl-1*H*-pyrazol-4-yl)-1*H*-indole (1.102).





## Ethyl 3-(2-oxopropoxy)benzoate (1.103).










Methyl 6-((2-methyl-1-(1-methyl-1*H*-pyrazol-4-yl)-1*H*-indol-3-yl)methyl)picolinate (1.105).





### Ethyl 3-((2-methyl-1*H*-indol-3-yl)oxy)benzoate (1.106).







### 3-((3-Bromophenyl)thio)-2-methyl-1*H*-indole (1.107).















### S2. Biological Assay Results

Compound	%inhibition at 30 μM inhibitor (bis- <i>p</i> NPP)	IC <sub>50</sub> (bis-pNPP)	IC <sub>50</sub> (LPC)
1.22	66	22 ±6.9 nM	$4 \pm 0.5 \text{ nM}^{a}$
1.24	49	-	258 ±127 nM
1.25	56	-	>30 µM
1.26	100	5700 ±2500 nM	1700 ±400 nM
1.27	38	816 ±564 nM	349 ±125 nM
1.28	48	-	>3500 nM <sup>b</sup>
1.29	52	85 ±22 nM	25 ±12 nM
1.30	44	-	$>10~\mu M^{b}$
1.31	58	75 ±37 nM	124 ±37 nM
1.32	49	-	>30 µM
1.33	31	-	>30 µM
1.34	46	604 ±113 nM	25 ±13 nM
1.35	30	-	>30 µM
1.36	47	-	>30 µM
1.37	33	-	>30 µM
1.38	42	-	${>}20~\mu M$ $^{b}$
1.39	50	-	>30 µM
1.40	11	>30 µM	>30 µM
1.41	42	-	>8000 nM <sup>b</sup>
1.42	21	-	>30 µM
1.43	13	-	>30 µM
1.44	56	>4000 nM <sup>b</sup>	31 ±11 nM
1.45	71	2000 ±785 nM	>2800 nM

**S2a. Table S1.** Summary of assay results

1.46	47	-	_ c
1.47	42	>30 µM	>30 µM
1.48	57	1388 ±746 nM	>30 µM
1.49	50	673 ±237 nM	>30 µM
1.50	63	2000 ±715 nM	>30 µM
1.51	38	-	$11 \pm 7 \text{ nM}^{f}$
1.52	41	290 ±35 nM	747 ±500 nM
1.53	92	100 ±36 nM	$1200 \pm 800 \text{ nM}^{\text{d}}$
1.54	44	>4000 nM <sup>b</sup>	$>20~\mu M^{b}$
1.55	55	5 ±2.5 nM	81 ±28 nM <sup>e</sup>
1.56	67	3000 ±1400 nM	>30 µM
1.57	25	>30 µM	>30 µM
1.58	42	-	>30 µM
1.59	64	>30 µM	>30 µM
1.60	79	>30 µM	>5000 nM <sup>b</sup>
1.61	73	89 ±43 nM	339 ±190 nM
1.62	72	33 ±16 nM	660 ±224 nM
1.63	56	204 ±88 nM	726 ±155 nM
1.64	35	-	>30 µM
1.65	22	>30 µM	>30 µM
1.66	35	>30 µM	>30 µM
1.67	29	>30 µM	>30 µM
1.68	17	>30 µM	>30 µM
1.69	50	>30 µM	>30 µM
1.70	52	-	>30 µM
1.71	18	-	$>10 \ \mu M^{b}$

<b>1.72</b> 9	>30 µM	>30 µM
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<sup>a</sup> Consistent with the value of < 300 nM claimed in the patent, LPC assay. <sup>b</sup> Incomplete curve observed, data extrapolated using GraphPad Prism to determine  $IC_{50}$ . <sup>c</sup> Unable to determine  $IC_{50}$  due to solubility issues in the assay medium. <sup>d</sup> Consistent with the value of >1 uM claimed in the patent, LPC assay. <sup>e</sup> Consistent with the value of < 300 nM claimed in the patent, LPC assay. <sup>f</sup> Consistent with the value of < 300 nM claimed in the patent, LPC assay.





**Figure S1.** Bis-*p*NPP assay results, tested using 30  $\mu$ M inhibitor with a >60% inhibition criterion, n = 2.

Bis-*p*NPP assay results:  $1 \text{ nM} - 30 \mu \text{M}$  of inhibitor.







**S2c.** LPC assay control reactions.

All biochemical studies were performed with hATX. Choline oxidase and horseradish peroxidase (HRP) activity were measured using choline chloride . 20 nM ATX (prepared from HEK 293 Flp-In cells) was incubated with 80  $\mu$ M choline chloride in a final volume of 100  $\mu$ L buffer containing 50 mM Tris pH 7.4, 0.01% Triton X-100, 50 mM CaCl<sub>2</sub>, 1 Unit ml<sup>-1</sup> choline oxidase, 2 Unit ml<sup>-1</sup> HRP, 2 mM homovanilic acid (HVA). The relative amount of released choline was measured by HVA fluorescence in a 96-well plate (Corning). Fluorescent intensity was determined at  $\lambda$ ex/ $\lambda$ em = 320/450 nm every 30 seconds for 90 minutes with a Fluorostar plate reader (BMG Labtech). Four exemplar compounds were tested from 400 nM- 100  $\mu$ M. No inhibition of choline oxidase or HRP was observed. Data analysis was performed using GraphPad Prism version 6.00 for Windows, GraphPad Software, La Jolla California USA, www.graphpad.com.



Figure S2. LPC control reactions ran using 1.29, 1.44, 1.63 and 1.61.











## **S3.** Physicochemical Properties

Compound	logD <sup>a</sup>	HSA <sup>b</sup>	AGP <sup>c</sup>	Sol. <sup>d</sup> (µM)	P <sub>app</sub> <sup>e</sup> (nm s <sup>-1</sup> )
1.22	2.70	98.5	80.3	377	23
1.34	2.19	98.5	78.7	≥395	11
1.44	2.68	98.5	80.4	300	32
1.45	4.05	98.9	93.8	336	290
1.46	4.30	99.4	91.8	485	155
1.51	3.36	98.1	83.2	556	64

 Table S2. Physicochemical properties of select analogues.

<sup>a</sup> logD, Chrom log D at pH 7.4; <sup>b</sup> human serum albumin (HSA) binding; <sup>c</sup> alpha 1 acid glycoprotein (AGP); <sup>d</sup>Chemiluminescent nitrogen detection (CLND) kinetic aqueous solubility assay; <sup>e</sup> P<sub>app</sub> permeability pH 7.4 assay. **S4. Molecular Modeling.** GOLD<sup>1</sup> software version 5.4.1 was used for the docking studies of compound **1.55** with double mutant hATX (PDB 4ZG7). Starting with the energy minimised structure of **1.55** (minimised using the MM2 calculation, Chem3D Pro software version 13.0.2.3021) the GOLD wizard was used to carry out the molecular modelling. The binding site was defined to within 7 Å of the PharmAkea analogue **1.21** and the docking was carried out using the slow method to identify six different solutions. The docking solutions were viewed using Discovery Studio Visualizer<sup>2</sup> with the protein surface colored by hydrophobicity. By comparing the docking solutions to the binding mode of **1.21**, the pose of **1.55** which gave the best overlay was selected for the basis of our docking model.



**Figure S3.** Docking studies of **1.55** using PDB 4ZG7. A: **1.55** (yellow) overlayed with **1.21** (colored by atom type), with **1.2** residing in the catalytic site and hydrophobic pocket. B: Potential binding interaction of **1.21**. C: Rotation of the thioether. D: Potential binding interactions of **1.55**. Docked using GOLD<sup>1</sup> and viewed using Discovery Studio Visualizer.<sup>2</sup>



Figure S4. Docking of analogues 1.25 (A) and 1.45 (B).



Figure S5. Overlay of different linker groups: Thioether 1.51 (yellow); ether 1.63 (red); methylene 1.62 (blue).

# **S5.** Crystallographic Data

PDB reference 5LQQ

# Table S3. Crystallography Details

	ATX-Compound 1.55		
Data Collection			
Wavelength (Å)	0.97925		
Resolution (Å)	46-2.49 (8.98-2.4)		
Space Group	P 1 2 <sub>1</sub> 1		
Unit Cell a, b , c (Å)	62.66, 89.07, 77.5		
CC <sub>1/2</sub>	0.999 (0.784)		
R <sub>merge</sub>	0.013 (0.531		
Ι/σΙ	56.6 (1.8)		
Completeness (%)	98.8 (99.4)		
Redundancy	3.1 (3.0)		
Refinement			
No. atoms	6466		
Protein	6252		
Ligand/Metal/Glycan	125		
Water/Iodine	89		
B-factors	62		
Protein	62		
Ligand/Metal/Glycan	67		
Water/Iodine	50		
TLS groups	1		
R <sub>work</sub> /R <sub>free</sub> (%)	21.7/26.2		
Validation			
Rmsd/rmsZ bond lengths (Å)	0.007/0.357		
Rmsd/rmsZ bond angles ( <sup>0</sup> )	1.207/0.535		
Ramachandran (%)	97.1/0.1		
Preferred/ outliers			

Sidechain rotamers	96.0/1.1
Preferred/ outliers	
MolProbity score/ clash score (%-ile)	100/100

High Resolution shell in parentheses





**Figure S6. Experimental density for compound 1.55 bound to ATX.** The 2mFo-Dfc map is shown at 1.0 rms as a blue wireframe model.









2-Amino-N-(3-(2-aminopyridin-4-yl)phenyl)acetamide (2.22).







2-Amino-N-(3-(5-aminopyridin-3-yl)phenyl)acetamide (2.23).









Synthesis of 2-amino-N-(3-(4-aminopyridin-2-yl)phenyl)acetamide (2.24).





## 2-Amino-N-(3-(6-aminopyridin-3-yl)phenyl)acetamide (2.25).






Synthesis of 2-amino-N-(3-(2-aminopyrimidin-4-yl)phenyl)acetamide (2.26).







## 2-Amino-N-(3'-amino-[1,1'-biphenyl]-3-yl)acetamide (2.27).







2-Amino-*N*-(3-(8-aminoimidazo[1,2-α]113yridine-6-yl)phenyl)acetamide (2.28).







2-Amino-N-(3-(2-methylpyridin-4-yl)phenyl)acetamide (2.29).













2-Amino-N-(3-(piperazin-1-yl)phenyl)acetamide (2.31).







2-Amino-N-(3-(5,6,7,8-tetrahydro-1,7-naphthyridin-2-yl)phenyl)acetamide (2.32).



121





N-(3-(1,8-naphthyridin-2-yl)phenyl)-2-aminoacetamide (2.33).







## 2-Amino-N-(3-(2-methyl-1,8-naphthyridin-3-yl)phenyl)acetamide (2.34).







2-Amino-N-(3-(5,6,7,8-tetrahydro-1,8-naphthyridin-2-yl)phenyl)acetamide (2.35).









## 2-Amino-N-(3-(2-methyl-5,6,7,8-tetrahydro-1,8-naphthyridin-3-yl)phenyl) acetamide (2.36).





2-Amino-N-(5'-(aminomethyl)-2'-methyl-[1,1'-biphenyl]-3-yl)acetamide (2.37).







2-Amino-N-(3'-(aminomethyl)-2'-methyl-[1,1'-biphenyl]-3-yl)acetamide (2.38).







2-Amino-N-(3'-(aminomethyl)-[1,1'-biphenyl]-3-yl)acetamide (2.39).







2-Amino-N-(3'-(2-aminoethyl)-[1,1'-biphenyl]-3-yl)acetamide (2.40).























## 2-(3-(2-Aminoacetamido)phenyl)-6-(aminomethyl)pyridine 1-oxide (2.43).












## 2-Acetamido-N-(5'-(acetamidomethyl)-2'-methyl-[1,1'-biphenyl]-3-yl)acetamide (2.45).





2-Amino-N-(5'-(hydroxymethyl)-2'-methyl-[1,1'-biphenyl]-3-yl)acetamide (2.46).







2-Amino-N-(5'-cyano-2'-methyl-[1,1'-biphenyl]-3-yl)acetamide (2.47).







N-(5'-(acetamidomethyl)-2'-methyl-[1,1'-biphenyl]-3-yl)-2-aminoacetamide (2.48).



153







N-(5'-((allylamino)methyl)-2'-methyl-[1,1'-biphenyl]-3-yl)-2-aminoacetamide (2.49).





2-Amino-N-(4'-(aminomethyl)-2'-methyl-[1,1'-biphenyl]-3-yl)acetamide (2.50).













2-Amino-N-(2'-(trifluoromethyl)-[1,1'-biphenyl]-3-yl)acetamide (2.52).









N-([1,1'-biphenyl]-3-yl)-2-aminoacetamide (2.53).







2-Amino-N-(3-(1-amino-2,3-dihydro-1H-inden-4-yl)phenyl)acetamide (2.54).







2-Amino-N-(2'-(hydroxymethyl)-[1,1'-biphenyl]-3-yl)acetamide (2.55).







2-Amino-N-(2'-((methylamino)methyl)-[1,1'-biphenyl]-3-yl)acetamide bis(2,2,2-



trifluoroacetate) (2.56).





2-Amino-N-(2'-(((cyclohexylmethyl)amino)methyl)-[1,1'-biphenyl]-3-yl)acetamide (2.57).







2-Amino-N-(2'-((benzylamino)methyl)-[1,1'-biphenyl]-3-yl)acetamide (2.58).









2-Amino-N-(2'-(((2-methoxyethyl)amino)methyl)-[1,1'-biphenyl]-3-yl)acetamide (2.59).







2-Amino-N-(2'-(morpholinomethyl)-[1,1'-biphenyl]-3-yl)acetamide (2.60).







2-Amino-N-(2'-(((2-aminoethyl)amino)methyl)-[1,1'-biphenyl]-3-yl)acetamide tris(2,2,2-



trifluoroacetate) (2.61).






2-Amino-*N*-(2'-(piperazin-1-ylmethyl)-[1,1'-biphenyl]-3-yl)acetamide bis(2,2,2-trifluoroacetate) (2.62).









*tert*-Butyl (2-oxo-2-((3-(4,4,5,5-tetramethyl-1,3,2-dioxaborolan-2-yl)phenyl) amino)ethyl) carbamate (2.63).









## 3-(4,4,5,5-Tetramethyl-1,3,2-dioxaborolan-2-yl)aniline (2.64).



186





*tert*-Butyl (6-bromoimidazo[1,2-α]pyridin-8-yl)carbamate (2.69).









tert-Butyl 4-(3-(2-((tert-butoxycarbonyl)amino)acetamido)phenyl) piperazine-1-carboxylate

















tert-Butyl (2-((5'-cyano-2'-methyl-[1,1'-biphenyl]-3-yl)amino)-2-oxoethyl) carbamate (2.75).









tert-Butyl (2-((3'-formyl-[1,1'-biphenyl]-3-yl)amino)-2-oxoethyl)carbamate (2.77).









*tert*-Butyl (2-((3'-(((tert-butylsulfinyl)amino)methyl)-[1,1'-biphenyl]-3-yl)amino)-2-oxoethyl) carbamate (2.78).









2-(3-Bromophenethyl)isoindoline-1,3-dione (2.80).





2500 2 Wavenumber (cm-1)



3-Bromo-2,4-dimethylbenzaldehyde (2.83).





## tert-Butyl (2-((3'-(((tert-butylsulfinyl)amino)methyl)-2',6'-dimethyl-[1,1'-biphenyl]-3-

yl)amino)-2-oxoethyl)carbamate (2.85).









## tert-Butyl (2-((3-(6-formylpyridin-2-yl)phenyl)amino)-2-oxoethyl)carbamate (2.87).





*tert*-Butyl

- 8.87 84.76 44.73 44.73 44.45 44.45 44.45 44.45 3.95 3.95 3.95 - 28000 `S || 0 - 26000 24000 НN `Ń H Å - 22000 s s f f Jin 20000 - 18000 16000 14000 12000 - 10000 - 8000 - 6000 4000 - 2000 AL M - 0 0.84 D-98-I 1.02H 2.11H 2.03 9.12-= 9.11-= 1.00H -2000 6.0 5.5 f1 (ppm) 4.5 4.0 11.0 10.5 10.0 9.5 9.0 8.5 8.0 7.5 7.0 6.5 5.0 3.5 3.0 2.5 2.0 1.5 1.0 0.5 0.0 139.7 137.6 137.6 137.6 129.4 122.7 120.8 1120.8 118.5 ✓ 157.4
✓ 156.2 --- 80.5 --- 28.5 - 22.9 --- 56.4 - 10000 9000 8000 7000 - 6000 - 5000 4000 - 3000 - 2000 1000 - 0 100 f1 (ppm) 200 190 170 130 120 110 90 80 70 60 50 40 30 20 10 0 180 160 150 140

oxoethyl)carbamate (2.88).

Intensity

Intensity





## 3-Bromo-4-methylbenzaldehyde (2.91).





tert-Butyl (2-((5'-formyl-2'-methyl-[1,1'-biphenyl]-3-yl)amino)-2-oxoethyl)carbamate (2.92).









*tert*-Butyl (2-((5'-((allylamino)methyl)-2'-methyl-[1,1'-biphenyl]-3-yl)amino)-2-oxoethyl)carbamate (2.93).





3000 2500 2000 1500 Wavenumber (cm-1)



tert-Butyl (2-((5'-(aminomethyl)-2'-methyl-[1,1'-biphenyl]-3-yl)amino)-2-oxoethyl)carbamate

(2.94).








tert-Butyl (4-bromo-2,3-dihydro-1H-inden-1-yl)carbamate (2.97).







tert-Butyl (2-((2'-formyl-[1,1'-biphenyl]-3-yl)amino)-2-oxoethyl)carbamate (2.99).









t*ert*-Butyl *N*-({[3-(4-{[(*tert*-butoxy)carbonyl]amino}pyridin-2-yl)phenyl] carbamoyl}



methyl)carbamate (2.102).











carbamate (2.106).





tert-Butyl

*N*-({[3-(5-{[(*tert*-butoxy)

carbonyl]amino}pyridin-3-yl)phenyl]

carbamoyl}methyl)carbamate (2.109).









6-Bromoimidazo[1,2-α]pyridin-8-amine (2.110)







tert-Butyl (2-((3-(2-methylpyridin-4-yl)phenyl)amino)-2-oxoethyl) carbamate (2.111).



















tert-Butyl (2-((3'-cyano-2'-methyl-[1,1'-biphenyl]-3-yl)amino)-2-oxoethyl)carbamate (2.113).







tert-Butyl (2-((5'-(hydroxymethyl)-2'-methyl-[1,1'-biphenyl]-3-yl)amino)-2-



oxoethyl)carbamate (2.114).







tert-Butyl (2-((3-bromophenyl)amino)-2-oxoethyl)carbamate (2.115).









*tert*-Butyl (2-(((cyclohexylmethyl)amino)methyl)-[1,1'-biphenyl]-3-yl)amino)-2-

oxoethyl)carbamate (2.116).









*tert*-Butyl (2-((2'-((benzylamino)methyl)-[1,1'-biphenyl]-3-yl)amino)-2-oxoethyl) carbamate (2.117).









tert-Butyl (2-(((2'-(((2-methoxyethyl)amino)methyl)-[1,1'-biphenyl]-3-yl)amino)-2-

oxoethyl)carbamate (2.118).





2500 2 Wavenumber (cm-1)





*tert*-Butyl (2-((2'-(morpholinomethyl)-[1,1'-biphenyl]-3-yl)amino)-2-oxoethyl)carbamate (2.119).









## **S7. References**

[1] Cole, J. C.; Nissink, J. W. M.; Taylor, R. Protein-Ligand Docking and Virtual Screening with GOLD.Virtual Screening in Drug Discovery (Eds. B. Shoichet, J. Alvarez), Taylor & Francis CRC Press, BocaRaton, Florida, USA (2005).

[2] Dassault Systèmes BIOVIA, Discovery Studio Modeling Environment, Release 4.5, San Diego: Dassault Systèmes, **2015**.