# Spirocycles as Rigidified sp<sup>3</sup>-Rich Scaffolds for a Fragment Collection

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Novel divergent methodology to access  $sp^3$ -rich spirocyclic fragments is reported. Firstly, a robust modular synthesis of bis-alkene amino ester building blocks was developed. Three different carbocycles and six heterocycles were then constructed to assemble eight spirocycles. Importantly, strategic exit vectors were incorporated within each scaffold to aid fragment growth and were elaborated via chemical modifications. Finally, computational methods demonstrate higher levels of rigidity, three-dimensionality and structural diversity of the library compared to a commercial collection.

Since the rise of fragment-based drug discovery (FBDD) over two decades ago this strategy has proven particularly effective, producing numerous clinical candidates and two FDA-approved drugs, Vemurafenib<sup>1</sup> and Venetoclax<sup>2</sup> (Figure 1. A). The success of this approach can be linked to two main benefits. Firstly, due to the considerably fewer number of possible fragment-sized molecules, the chemical space coverage of a relatively small fragment library is exceedingly more efficient than that of a vast high-throughput screening (HTS) library. Secondly, fragment hits possess fewer but nonetheless high-quality binding interactions with the protein target, which can be later elaborated to afford highly potent lead compounds.<sup>3-7</sup>

Within this paradigm the generation of a suitable screening library is paramount. However, despite undoubted success of FBDD, within recent years organic synthesis has been identified as a significant bottleneck within this process, owing to overrepresentation of predominantly the 'flat' (hetero)aromatic fragments lacking three-dimensionality as well as synthetically tractable exit vectors that could be utilised in rapid structure-activity relationship (SAR) studies.<sup>4,5,8</sup> Whilst complexity of fragments remains under debate within the literature,<sup>9,10</sup> importantly, more three dimensional (3D) fragments displaying exit vectors increase the potential for multidirectional fragment growth and the ability to identify leads for challenging targets such as protein-protein interactions.<sup>8,11</sup> Thus, recent efforts from within the synthetic community have focused on the development of novel strategies to access 3D fragments.12-15



**Figure 1.** A) Two FBDD-derived FDA-approved drugs: Venetoclax and Vemurafenib. B) Examples of spirocyclic natural products (Griseofulvin and Histrionicotoxin) and FDA-approved drugs (Spironolactone and Irbesartan).

Spirocyclic motifs remain an important bioactive substructure appearing within several natural products,<sup>16–18</sup> and FDAapproved drugs<sup>19,20</sup> (Figure 1, B). Importantly, as a direct result of their architecture, these small molecules often provide several advantages.<sup>21</sup> Firstly, the spiranic centre generates an inherently 3D structure that gives rise to higher levels of complexity, a feature which has been linked to improved clinical success.<sup>10,22,23</sup> Moreover, the conformationally restricted nature of spirocycles can reduce both the conformational entropy penalty of target binding and the number of possible conformations (distinct 3D shapes) that a molecule can adopt leading to higher potency and selectivity, respectively.<sup>24,25</sup> However, despite their utility, these motifs remain underrepresented in fragment screening collections. Indeed, very few compounds within the ChemBridge spirocycle library meet the size criteria of FBDD,<sup>26,27</sup> whilst only three spirocycles feature within the Maybridge core fragment collection. Thus, there is an urgent need for spirocyclic fragments and calls from within the field have encouraged the development of novel strategies to access such motifs.<sup>25</sup>

Whilst strategies to access spirocycles within sp<sup>3</sup>-rich screening libraries have been reported, they either do not sole-ly seek to construct spirocycles,<sup>28–30</sup> or focus on more complex bis-<sup>31</sup> and bridged-<sup>32</sup> spirocycles. In addition, these compounds often lie outside the requirements for FBDD. Thus, we envisaged that a novel approach to access diverse spirocyclic fragments containing a polar heterocycle and a lipophilic carbocycle could give rise to a valuable library complementing already existing screening collections. To achieve this, our efforts were directed at utilising  $\alpha \alpha$ -disubstituted amino acid derivatives as building blocks, providing the potential to exploit the functional handles to generate fragment-like spirocyclic scaffolds. The incorporation of the two alkene handles enabled the carbocycle formation via ring closing metathesis (RCM), forming an essential alkene exit vector, allowing us to alter the properties of this portion of the fragments. In addition, the amino and ester functionalities were installed to enable diverse heterocycle formations, increasing the possible polar interactions and the overall water solubility of the fragments. Accordingly, rapid access to varied scaffolds and the potential to exploit the exit vectors for fragment growth and merging to aid hit-to-lead development was envisioned.

Firstly, the building blocks were prepared through the double alkylation of the glycine Schiff base **1** to form building blocks **3a-c** (Scheme 1). Despite similar procedures have been reported before,<sup>33–36</sup> herein we describe the straightforward racemic synthesis of  $\alpha$ -quaternary amino esters with two different alkyl chains bearing terminal alkenes (**3b,c**). A brief optimisation of the related literature procedure<sup>33</sup> allowed us to form **3c** in a simple step-wise process. Further optimisation of the analogous route, however, resulted in the development of a one-pot procedure removing the chromatographic steps to access **3b** on large-scale in an improved 61% yield. This approach also enabled the formation of **3a**.

#### Scheme 1. Synthesis of the Building Blocks



Reaction conditions: (i) *t*BuOK, THF; (ii) *t*BuOK, THF; (iii) HCl, THF/H<sub>2</sub>O; then Na<sub>2</sub>CO<sub>3</sub>. Overall yields: **3a**: 67% (steps i and ii the same, steps i-iii in one-pot); **3b**: 61% (steps i-iii in one-pot); **3c**: 31% (steps i-iii done step-wise).

Subsequently, investigations into the formation of the different heterocycles were pursued using the racemic amino ester building block **3b**. Firstly, the amine was Boc-protected to allow the cyclohexene formation in a RCM, followed by the reduction of the ester to the hydroxymethyl group by LiBH<sub>4</sub> to afford the key intermediate **4** in good yield. Intramolecular base-mediated pairing between the alcohol and Boc groups could then be achieved forming the oxazolidone moiety in **5**. The removal of the Boc protecting group under acidic conditions could be followed by pairing reactions incorporating various reagents. Reaction with cyanogen bromide formed amino oxazoline **6** whereas ethyl benzimidate hydrochloride gave the phenyl-substituted oxazoline **7**. The two morpholinones **8** and **9** were constructed by the chemoselective alkylation/acylation with chloroacetyl chloride and phenyl bromoacetate respectively (Scheme 2).

Building block **3b** could also be acylated with ethyl malonyl chloride, then the base-mediated cyclisation onto the ester group followed by the acid-catalysed decarboxylative hydrolysis yielded the tetramic acid intermediate **10b**. Formation of the carbocycle ring in a RCM gave **12** in a good yield. To exemplify the potential to expand the cyclohexene ring, spirocycles also featuring the 5- and 7-membered carbocycles were also synthesised from **3a**, and **3c** respectively (Scheme 2). Importantly, all the spirocycles were synthesised in no more than five steps from the building blocks.

Scheme 2. Synthesis of Different Core Heterocycles and Carbocycles



Reaction conditions: a) (i)  $Boc_2O$ , THF, 85%; (ii) Grubbs II,  $CH_2Cl_2$ , 81%; (iii) LiBH<sub>4</sub>, THF, 89%; b) *t*BuOK, THF, 90%; c) HCl, dioxane; then BrCN, Et<sub>3</sub>N, EtOH, 58%; d) HCl, dioxane; then ethyl benzimidate hydrochloride, Et<sub>3</sub>N, DCE, 57%; e) (i) HCl, dioxane; then chloroacetyl chloride, Et<sub>3</sub>N, CH<sub>2</sub>Cl<sub>2</sub>, 57%; (ii) *t*BuOK, *t*BuOH, 99%; f) HCl, dioxane; then phenyl bromoacetate, iPr<sub>2</sub>NEt, MeCN, 43%; g) (i) ethyl malonyl chloride, Et<sub>3</sub>N, CH<sub>2</sub>Cl<sub>2</sub>, 73–81%; (ii) *t*BuOK, THF; then aq. HCl, THF, 86–92%; h) Grubbs II, CH<sub>2</sub>Cl<sub>2</sub>, 69–99%.

Although racemic compounds were sought for our fragment library, the ability to produce optically pure isomers e.g. for SAR studies was also crucial. Thus, a second asymmetric route to the desired building blocks was devised utilising the well-precedented stereoselective alkylation of iminolactones derived from phenylglycinol.<sup>37,38</sup> In this case, only one diastereomer of the aminolactone (*R*)-15 was observed, which

was successively deprotected to form the optically pure building block (R)-3d. As proof of concept, the single R-enantiomer of 12, with the spiro[4,5] scaffold, was also synthesised (Scheme 3).

#### Scheme 3. Enantioselective Synthesis of (R)-12



Reaction conditions: a) allyl bromide, Zn, DMF, 68%; b) (i) SOCl<sub>2</sub>, MeOH; (ii) Pb(OAc)<sub>4</sub>, MeOH/CH<sub>2</sub>Cl<sub>2</sub>; then HCl, H<sub>2</sub>O, 92% over 2 steps; c) (i) ethyl malonyl chloride, Et<sub>3</sub>N, CH<sub>2</sub>Cl<sub>2</sub>, 46%; (ii) *t*BuOK, THF; then aq. HCl, THF, 92%; (iii) Grubbs II, CH<sub>2</sub>Cl<sub>2</sub>, 97%.

With eight fragment scaffolds in hand, it was next crucial to demonstrate the utility of the exit vectors installed within the molecules. Thus, *N*-alkylation of the oxazolidone (16), *O*-alkylation (17) and cross-coupling of the tetramic acid (18) were demonstrated. In addition, a modified heterocycle with an alcohol functionality (19) and two with new double bonds (20 and 21) were synthesised, incorporating new exit vectors to the resultant molecules (Scheme 4).

#### Scheme 4. Heterocycle Modification



Reaction conditions: a) PMBCl, NaH, DMF, 96%; b) KHMDS, EtBr, THF, 54%; c) (i) Tf<sub>2</sub>O, Et<sub>3</sub>N, CH<sub>2</sub>Cl<sub>2</sub>, 59%; (ii) PMPB(OH)<sub>2</sub>, Pd(PPh<sub>3</sub>)<sub>4</sub>, Na<sub>2</sub>CO<sub>3</sub>, H<sub>2</sub>O/THF, 73%; d) NaBH<sub>4</sub>, MeOH, 23%; e) TFAA, Et<sub>3</sub>N, CH<sub>2</sub>Cl<sub>2</sub>; then KHCO<sub>3</sub>, MeOH, 30%; f) Pb(OAc)<sub>4</sub>, MeCN, 92%.

Finally, to exhibit the versatility of the double bond as an exit vector, modifications such as Wacker-oxidation (22a,b), dihydroxylation (23a,b), difluoro-cyclopropanation (24a,b), aziridination (25a,b), epoxidation (26a,b) and dibromination (27a,b) were explored (Scheme 5). Initial attempts of the epoxidation of 5, however proved challenging with respect to the isolation and purification of 26a,b. Thus, a PMB group was installed (16) indeed proving to be compatible with several reaction conditions generating the diversified fragments in good to excellent yields. Removal of the PMB protecting group was also exemplified by the treatment of compound 27a with CAN to generate the unprotected modified spirocycle 28.

Scheme 5. Double Bond Modification and PMB Deprotection



R = PMB unless specified otherwise. All products are racemic. Reaction conditions (combined yields with ratios  $\mathbf{a}/\mathbf{b}$  are given): a) Fe(acac)<sub>2</sub>, *t*BuOH, air, 47%, 1.9:1; b) OsO<sub>4</sub>, NMO, citric acid, H<sub>2</sub>O/THF, 99%, 2.5:1; c) TMSCF<sub>3</sub>, NaI, THF, 62%, 20:1; d) TsNCINa.3H<sub>2</sub>O, PhNMe<sub>3</sub>Br<sub>3</sub>, 4 Å MS, 70%, 1.4:1; e) R = H, mCPBA, NaHCO<sub>3</sub>, CH<sub>2</sub>Cl<sub>2</sub>, 52%, 8:1; f) PhNMe<sub>3</sub>Br<sub>3</sub>, CH<sub>2</sub>Cl<sub>2</sub>, 96%, 18:1; g) CAN, MeCN/H<sub>2</sub>O, 96% (R = H in product).

 Table 1. Physicochemical Properties of Fragment Libraries Compared to the Ideal Range

property <sup>a</sup>	spirocycles <sup>b</sup>	Maybridge <sup>c</sup>	ideal range <sup>d</sup>
MW	$186\pm41$	$182\pm42$	140-230
HBD	$1.3\pm0.7$	$1.0\pm0.8$	≤3
HBA	$1.8\pm0.6$	$1.9\pm0.7$	≤3
SlogP	$0.9\pm0.9$	$1.8\pm0.8$	0–2
RBC	$0.2\pm0.5$	$2.0\pm1.5$	≤3
TPSA	$48 \pm 13$	$39 \pm 14$	≤60

<sup>*a*</sup>MW = molecular weight (Da), HBD = number of hydrogenbond donors, HBA = number of hydrogen-bond acceptors, SlogP = partition coefficient, RBC = rotatable bond count, TPSA = topological polar surface area (Å<sup>2</sup>). <sup>*b*</sup>Protecting groups virtually removed from our library. <sup>*c*</sup>Maybridge core fragment collection of 1000 fragments. <sup>*d*</sup>Guidelines set by Astex Pharmaceuticals.<sup>26,27</sup> See Supporting Information.

The physicochemical properties of our library consisting of 28 different non-protected spirocyclic molecules were then calculated and compared to the commercially available Maybridge core fragment library using the widely-accepted guidelines from within the field (Table 1).<sup>26,27</sup> This revealed the spirocyclic library adheres well to the guidelines, and was additionally predicted to be significantly less lipophilic (SlogP of 0.9 versus 1.8) and more rigid (rotatable bond count of 0.2 versus 2.0) than the Maybridge core fragment collection. Strikingly, our library also displayed a far superior sp<sup>3</sup>-content with an average fraction of sp<sup>3</sup> atoms (Fsp<sup>3</sup>) of 0.52, which also translates into a much lower fraction of aromatic atoms (Faro) of 0.16. Moreover, the average number of chiral centres (1.6) is also considerably higher, resulting in greater stereochemical diversity achieved by the spirocyclic fragments.

In order to qualitatively assess the shape diversity in our library, a principal moments of inertia (PMI) analysis was carried out. Our library was then compared to the whole Maybridge core fragment collection of 1000 fragments and a representative subset consisting of the 147 best-matched compounds based on heavy atoms (Figure 2). Both plots show that the Maybridge fragments tend to aggregate to the left-hand edge (rod- and disc-like features) or the 'flat land', whereas our spirocyclic fragments are more evenly distributed. Analysis showed that more than 70% of the whole Maybridge collection and 75% of the best formula match subset falls within 'flat land' (defined as npr1 + npr2  $\leq 1.1$ ).<sup>11</sup> On the other hand, no spirocyclic fragment in our library was found below the 'flat land' criteria, suggesting more 3D molecules.

In conclusion, we have developed a robust, scalable and modular route to racemic  $\alpha,\alpha$ -disubstituted amino ester building blocks including an adapted stereoselective alkylation protocol to access the optically pure intermediate. These were utilised in the efficient construction of eight novel spirocyclic scaffolds comprising of pharmacophore heterocycles and variable carbocycles. All core scaffolds display an array of 3D exit vectors demonstrated by a number of chemical modifications to both the hetero- and carbocycles. Together with the enantioselective synthesis, rapid SAR studies and binding pocket exploration by fragment growth could therefore be envisioned. Finally, the computational predictions revealed optimal physicochemical properties, higher rigidity increased 3D properties, shape and stereochemical diversity compared to a commercial fragment library.



**Figure 2.** PMI plots for the visual representation of shape diversity. Each corner of the plot represents a unique shape (rod-, disk- and sphere-like features). The dashed line represents the boundary of 'flat land' <sup>11</sup>. Our virtual deprotected library of 28 spirocycles (blue) is compared to 1000 fragments (red, left) and the 147 best-matched fragments based on heavy and heteroatom count (red, right) from the Maybridge collection. The table in the right summarises the physicochemical properties used to describe the 3D properties of the two libraries. "Fsp<sup>3</sup> = fraction of sp<sup>3</sup> atoms, Far = fraction of aromatic atoms, chiral = number of chiral centres, npr = normalised PMI ratio, Fflat = fraction of molecules lying below the 'flat land line'<sup>11</sup>. <sup>b</sup>Virtual deprotected library. <sup>c</sup>147 best-matched fragments from the Maybridge collection. See supporting information.

## ASSOCIATED CONTENT

### **Supporting Information**

The Supporting Information is available free of charge on the ACS Publications website at DOI: 10.1021/acs.orglett.9b01499.

General Remarks, Procedures and Analytical Data, Computational Analysis, Crystallographic Data and NMR Spectra (PDF)

#### **Accession Codes**

CCDC 1912266-1912268 and 1912283-1912289 contains the supplementary crystallographic data for this paper. These data can be obtained free of charge via www.ccdc.cam.ac.uk/data\_request/cif, or by emailing data\_request@ccdc.cam.ac.uk, or by contacting The Cambridge Crystallographic Data Centre, 12 Union Road, Cambridge CB2 1EZ, UK; fax: +44 1223 336033.

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All authors have given approval to the final version of the manuscript.

#### Notes

The authors declare no competing financial interest.

# ACKNOWLEDGMENT

A.S. acknowledges support from the Walters family and Selwyn College, University of Cambridge. A.J.P.N., H.F.S. and D.R.S. acknowledge support from the Engineering and Physical Sciences Research Council (EP/P020291/1) N.M.S. thanks the EU for a Marie Curie Fellowship (2013-IEF-626191). S.L.K. thanks AstraZeneca for funding. D.R.S. acknowledges support from the Royal Society (Wolfson Research Merit Award). The Spring lab acknowledges general lab support from ERC, EPSRC, BBSRC, MRC and Royal Society. The authors would like to thanks Dr. Andrew Bond (Department of Chemistry, University of Cambridge) for carrying out the single-crystal crystallography for the project.

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