# **New Peptide-Based Templates**

# Constrained into a $\beta$ -Strand by Huisgen

# Cycloaddition

A thesis submitted in total fulfilment of the requirements for the degree of Doctor of Philosophy

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# **Abstract**

Chapter One introduces the concept of peptide 'secondary structure' with an emphasis on  $\beta$ -strand geometry in macrocycles. This structural design is crucial for targeting different proteases. The significance of the macrocylic  $\beta$ -strand 'bioactive' conformation is discussed in detail. In particular the exploitation of the conformationally constrained peptidomimetic macrocylic backbone, which is constrained by a number of synthetic approaches to lock the 'bioactive' conformation in place.

**Chapter Two** describes simple and scalable methodology for the preparation of *N*-Cbz protected amino acids by reaction with Cbz-Cl which uses a mixture of aqueous sodium carbonate and sodium bicarbonate to maintain the appropriate pH. This method proceeds without the formation of by-products. The method is extended to large scale preparation of an intermediate zofenopril, an ACE inhibitor.

Chapter Three describes new peptidic templates constrained into a  $\beta$ -strand geometry by linking acetylene and azide containing  $P_1$  and  $P_3$  residues of a tripeptide by Huisgen cycloaddition. The conformations of the macrocycles are defined by NMR studies and those that best define a  $\beta$ -strand are shown to be potent inhibitors of the protease calpain. The  $\beta$ -strand templates presented and defined here are prepared under optimized conditions and should be suitable for targeting a range of proteases and other applications requiring such geometry.

Chapter four describes a new approach to non-covalent peptide-based nanotubular or rod-like structures, whereby the monomeric units are preorganised into a  $\beta$ -strand geometry that templates the formation of an extended and unusual parallel  $\beta$ -sheet rod-like structure. The conformational constraint is introduced by Huisgen cycloaddition to give a triazole-based macrocycle, with the resulting self-assembled structures stabilized by a well-defined series of intermolecular hydrogen bonds.

Chapter Five the 26S proteasome has emerged over the past decade as an attractive therapeutic target in the treatment of cancers. Here, we report new tripeptide aldehydes that are highly specific for the chymotrypsin-like catalytic activity of the proteasome. These new CT-L specific proteasome inhibitors demonstrated high potency and specificity for cancer cells, with therapeutic windows superior to those observed for benchmark proteasome inhibitors, MG132 and Bortezomib. Constraining the peptide backbone into the β-strand geometry was associated with decreased activity *in vitro* and reduced anticancer activity, suggesting that the proteasome prefers to bind a conformationally flexible ligand. Using these new proteasome inhibitors, we show that the presence of an intact p53 pathway significantly enhances cytotoxic activity, thus suggesting that this tumor suppressor is a critical downstream mediator of cell death following proteasomal inhibition.

Chapter Six peptide derived protease inhibitors represent an important class of compounds with the potential to treat a wide range of serious medical conditions. Herein we describe the synthesis of a series of triazole containing macrocylic protease inhibitors preorganised in a  $\beta$ -strand conformation and evaluate their selectivity and potency against a panel of protease inhibitors. A series of acyclic azido-alkyne-based aldehydes is also evaluated for comparison. The macrocyclic peptidomimetics showed considerable activity towards Calpain II, Cathepsin L and S and the 26S proteasome chymotrypsin-like activity. Importantly, the first examples of potent and selective inhibitors of Cathepsin S were identified and shown to adopt a well-defined  $\beta$ -strand geometry by NMR, X-ray and molecular docking studies.

**Chapter Seven** describes simple and efficient methodology for the selective acylation and alkylation of biotin at its 3'-nitrogen. This methodology is used to prepare of other biotin derivatives.

# **Declaration**

This work contains no material which has been accepted for the award of any other degree or diploma in any university or other tertiary institution and to the best of my knowledge and belief, contains no material published or written by another person, except where due reference has been made in the text.

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Date

Publications arising from this thesis:

- "An improved large scale procedure for the preparation of *N*-Cbz amino acids" Pehere, A. D.; Abell, A. D. *Tetrahedron Lett.* **2011**, *52*, 1493-1494.
- 2) "Selective N-acylation and N-alkylation of biotin" Pehere, A. D.; Abell, A. D. *J. Org. Chem.* **2011**, *76*, 9514-9518.
- 3) "New β-Strand Templates Constrained by Huisgen Cycloaddition" Pehere, A. D.; Abell, A. D. *Org. Lett.* **2012**, *14*, 1330-1333.
- 4) "New Cylindrical Peptide Assemblies Defined by Extended Parallel β-Sheets" Pehere, A. D.; Sumby, C. J.; Abell, A. D. (Manuscript is to be submitted to *J. Am. Chem. Soc.*).
- 5) "New 26S-proteasome inhibitors with high selectivity for chymotrypsin-like activity and p53-dependent cytotoxicity". Neilsen , P.M.; Pehere, A. D.; Callen, D. F.; Abell, A. D. (Manuscript is to be submitted to *ACS Chem. Biol.*).
- 6) "Synthesis and extended activity of triazole-containing macrocyclic protease inhibitors". (Manuscript in preparation).

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# **Abbreviations**

aq aqueous

Boc *tert*-butoxycarbonyl br broad (spectroscopic)

calcd calculated

Cbz benzyloxycarbonyl

conc concentrated
Cy cyclohexyl

DCM dichloromethane

DBU 1,8-Diazabicyclo[5.4.0]undec-7-ene

DDQ 2,3-dichloro-5,6-dicyano-1,4-benzoquinone

DIPEA *N,N*-diisopropylethylamine 4-DMAP 4-Dimethylaminopyridine

DMF dimethylformamide

DMP Dess–Martin periodinane

DMSO dimethyl sulphoxide

DMTr 4,4'-dimethoxytrityl group

EDC 1-ethyl-3-(3-(dimethylamino)propyl)carbodiimide

hydrochloride

equiv equivalent

ESI electrospray ionisation

Et ethyl

FTIR Fourier transform infrared

h hour(s)

HATU 2-(7-aza-1*H*-benzotriazol-1-yl)-1,1,3,3-

tetramethyluronium hexafluorophosphate

HIV Human Immunodeficiency Virus

HOAt 1-hydroxy-7-azabenzotriazole

HOBt 1-hydroxybenzotriazole

HPLC high-performance liquid chromatography

HRMS high-resolution mass spectrometry

iPA isopropylalcohol

IR infrared

lit. literature value

Me methyl min minute(s)

mp melting point

Ms methylsulphonyl (mesyl)

MS mass spectrometry m/z mass-to-charge ratio

NMR nuclear magnetic resonance

PDB Protein Data Bank

Ph phenyl

PI protease inhibitor(s)

Ppm part(s) per million

Pr propyl

PTSA *p*-toulenesulphonic acid

Py pyridine quantitative

RCM ring closing metathesis

rt room temperature

SAR structure activity relationship

spec spectrometry

TBAB tetrabutylammonium bromide
TBAI tetrabutylammonium iodide

TCE 1,1,2-trichloroethane

TEA triethylamine temp temperature

TFA trifluoroacetic acid
THF tetrahydrofuran

TLC thin layer chromatography

Ts	para-toluenesulphon	v1	(tosv	1)
13	$\rho a r a$ -toruchicsurphon	yı ı	( tOS y .	IJ

UV ultraviolet

v/v volume per unit volume

w/w weight per unit weight

#### Paper 1

"An improved large scale procedure for the preparation of *N*-Cbz amino acids" Pehere, A. D.; Abell, A. D. *Tetrahedron Lett.* **2011**, *52*, 1493-1494.

#### Mr.Ashok Pehere (candidate)

Performed all the experimental work, interpreted data, and prepared manuscript.

I hereby certify that the statement of contribution is accurate.

Signed .....

date 13/06/2012

#### **Professor Andrew Abell**

Supervised development of work, assisted in data interpretation, and revised the manuscript, and is the corresponding author.

I hereby certify that the statement of contribution is accurate and I give permission for the inclusion of the paper in this thesis.

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date 13/6/2012

#### Paper 2

"New β-Strand Templates Constrained by Huisgen Cycloaddition" Pehere, A. D.; Abell, A. D. *Org. Lett.* **2012**, *14*, 1330-1333.

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#### Paper 4

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Assisted with data interpretation and revision of manuscript.

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#### **Professor Andrew Abell**

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#### Paper 6

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Supervised development of work, assisted in data interpretation, and revised the manuscript, and is the corresponding author.

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