Design and Synthesis of Protein Chemical Crosslinkers: A Modular Approach

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Abstract

The study of protein structure and interactions is pivotal in understanding the function and malfunction of complex biological systems. The structures of some proteins are unable to be determined using traditional high resolution biophysical techniques, requiring the development of amenable low resolution alternatives. Chemical Crosslinking Mass Spectrometry (CXMS) is one technique which can be used to probe protein structure through the formation of covalent linkages between protein residues. The formation of these links is facilitated by chemical crosslinking reagents.

Widespread use of the CXMS technique has been hampered primarily by analytical challenges pertaining to the detection and identification of crosslinked species using Mass Spectrometry (MS). Attempts to mitigate the challenges have been made by modifying the structure of chemical crosslinkers through the addition of functional groups such as affinity tags, isotope labels and cleavable bonds. Crosslinkers combining more than one type of functional group (combination crosslinkers) present the most promising targets for CXMS applications, combining the benefits of each functional group. However, combination crosslinkers are not commercially available, thus necessitating in-house synthesis. Incorporating more than one functionality also results in more complex molecular structures and synthetic processes, making the crosslinkers difficult to adapt to suit a particular experiment. Consequently, the use of combination crosslinkers has been limited to date to a small number of studies.

The research presented in this thesis describes the development of a modular chemical crosslinker design and corresponding synthetic protocol for the synthesis of combination crosslinkers. The modular crosslinker structure can be readily modified to include a range of functional groups using a small number of different reactions, including amide coupling and O-alkylation, and commercially available starting materials such as Boc-serine, from a minimum of five synthetic steps. The utility of the synthetic process was validated through the synthesis of a crosslinker containing an alkyne functional group, which can be used to attach a biotin affinity tag through alkyne-azide Huisgen Cyclisation.

Synthesis of two custom designed combination crosslinkers utilising alkyne tags and cleavable bonds is also described. The function of the cleavable bonds was established using collision induced dissociation processes within the mass spectrometer. Ensuring that a crosslinker is effective in probing quaternary structure and protein-protein interactions is essential as the investigation of these structures is a major goal of CXMS. Therefore, a crosslinking assay using *Staphylococcus aureus* biotin protein ligase, which forms homodimers when substrate bound, was also developed using the commercially available crosslinkers Disuccinimidyl Suberate (DSS) and Dithiobis(succinimidyl) Propionate (DSP), to enable the efficacy of crosslinkers synthesised using the modular synthetic protocol to be determined.

Declaration

I certify that this work contains no material which has been accepted for the award of any other degree or diploma in my name, in any university or other tertiary institution and, to the best of my knowledge and belief, contains no material previously published or written by another person, except where due reference has been made in the text. In addition, I certify that no part of this work will, in the future, be used in a submission in my name, for any other degree or diploma in any university or other tertiary institution without the prior approval of the University of Adelaide and where applicable, any partner institution responsible for the joint-award of this degree.

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Abbreviations

Arom Aromatic Chemical Shift
ATP Adenosine Triphosphate
Boc tert-Butyloxycarbonyl
BPL Biotin Protein Ligase

CID Collision Induced Dissociation

Cryo-EM Cryo-electron Microscopy

CXMS Cross Linking Mass Spectrometry

CXL Chemical Cross Linking

Da Daltons

kDa kilo-Daltons DC Direct Current

DDA Dithiodiglycolic Acid

DEA Diethylamine

DCM Dichloromethane

DIC N'N-diisopropylcarbodiimide

DMF Dimethylformamide
DNA Deoxyribonucleic Acid

DSP Dithiobis(succinimidyl) Propionate

DSS Disuccinimidyl Suberate

EDC-HCl 1-Ethyl-3-(3-dimethylaminopropyl)carbodiimide Hydrochloride

ECD Electron Capture Dissociation

El Electron Ionisation

eq Equivalents

ESI Electrospray Ionisation
FAB Fast Atom Bombardment
Fmoc Fluorenylmethyloxycarbonyl

h hour(s)

HATU 1-[Bis(dimethylamino)methylene]-1H-1,2,3-triazolo[4,5-b]pyridinium

3-oxid hexafluorophosphate

HDX Hydrogen/Deuterium Exchange

HDX-MS Hydrogen/Deuterium Exchange Coupled to Mass Spectrometry

HRMS High Resolution Mass Spectrometry

HTH Helix-Turn-Helix

HOBt 1-Hydroxybenzotriazole

HPLC High Performance Liquid Chromatography

IMMS Ion Mobility Mass Spectrometry

IR Infrared

MALDI Matrix Assisted Laser Desorption Ionisation

MCP Microchannel Plate

MeO Methyl Ester Protecting Group

MeOH Methanol

MS Mass Spectrometry

MS/MS Tandem Mass Spectrometry

MS/MS/MS 3-Stage Tandem Mass Spectrometry

m/z Mass to Charge Ratio

nESI Nano-Electrospray Ionisation

NHS N-hydroxy Succinimide

NMR Nuclear Magnetic Resonance Spectroscopy

PBS Phosphate-Buffered Saline Q-ToF Quadupole-Time of Flight

 $R_{\rm f}$ Retention Factor RF Radio Frequency

Rxn Reaction

SaBPL Staphylococcus aureus Biotin Protein Ligase

SAXS Small Angle X-ray Scattering

SOCl₂ Thionyl Chloride

TFA-NHS N-trifluoroacetoxy Succinimide

TFAA Trifluoroacetic Anhydride

THF Tetrahydrofuran
ToF Time of Flight

TLC Thin Layer Chromatography

WT Wild Type

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