

The structure and function of Biotin Protein Ligase:

***A focus on *Staphylococcus aureus*,
Saccharomyces cerevisiae, *Candida albicans*
and *Homo sapiens*.***

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Abbreviations

ACC: Acetyl CoA Carboxylase

AMP: Adenosine monophosphate

Apo: Unliganded enzyme

ATP: Adenosine triphosphate

BCCP: Biotin Carboxyl Carrier Protein

BirA: Biotin inducible repressor

BPL: Biotin Protein Ligase

Bt: Biotin

BtOH-AMP: Biotinol-5'-adenosine monophosphate

CaBPL: *Candida albicans* Biotin Protein Ligase

EcBPL: *Escherichia coli* Biotin Protein Ligase

HCS: Holocarboxylase synthetase

Holo: Ligand bound enzyme

MCD: Multiple carboxylase deficiency

MR: Molecular replacement

MRSA: Methicillin resistant *Staphylococcus aureus*

PC: Pyruvate carboxylase

PDB: Protein Data Bank

PDBID: Protein Data Bank identification code

PhBPL: *Pyrococcus horikoshi* Biotin Protein Ligase

R.M.S.D: Root mean square deviation

SaBPL: *Staphylococcus aureus* Biotin Protein Ligase

ScBPL: *Saccharomyces cerevisiae* Biotin Protein Ligase

SAXS: small angle X-ray scattering

VRSA: Vancomycin resistant *Staphylococcus aureus*

Abstract and summary of thesis for Nicole Renee Pardini

Biotin Protein Ligase (BPL) is an essential enzyme responsible for the covalent attachment of biotin to a specific lysine residue of biotin-dependent carboxylases, transcarboxylases and decarboxylases. Due to the fundamental processes that these enzymes are involved in such as lipogenesis, amino acid catabolism and gluconeogenesis, much research has been conducted on these enzymes. Studies encompassing structural, mutational and catalytic functions of these enzymes have led to novel drug developments for the treatment of obesity, diabetes, metabolic syndrome, bacterial and fungal infections.

As BPL is required for activation of these enzymes by biotinylation, it is believed that it too could be targeted in a similar way to produce novel therapeutics. To date, the most characterised BPLs are from the Gram-negative bacteria *Escherichia coli* and the archaea *Pyrococcus hirokoshii*. However minimal information is known about other forms of clinically important bacterial species or eukaryotic forms of this important enzyme.

Through my candidature I have compiled a thorough literature review summarised as chapter 1: Introduction. Furthering this literature analysis, a human BPL model was generated with aid of BPL structural co-ordinates already deposited in the protein data bank (PDB), thus allowing focus on human BPL mutations that cause multiple carboxylase deficiency (chapter 2). I have solved the structure of BPL from the clinically important pathogenic bacteria *Staphylococcus aureus*. This was performed in several ligand-bound and non-bound states (chapters 3 and 4). A novel high-throughput assay was

developed to test BPL activity. This assay allow testing of compounds that could potentially inhibit the BPL from *Candida albicans* (a species responsible for invasive fungal infections) (chapter 5). Large amounts of highly purified BPL from *Saccharomyces cerevisiae* allowed for the first structural analysis of a eukaryotic BPL (Chapter 6). The work has been summarised by a general discussion and future directions for the project (Chapter 7).

Thesis layout:

The thesis will be presented as a series of manuscripts either published or intention to be submitted for publication. Each manuscript will form a self-contained chapter with its own references. Included will be an overall general introduction and general discussion to link the information from these papers so as to present a uniform body of research conducted during candidature.



Declaration for thesis containing published work and/ or work prepared for publication

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Chapter 2:

Nicole R. Pardini, Lisa M. Bailey, Grant W. Booker, Matthew C. Wilce, John C. Wallace & Steven W. Polyak. (2008) Distant relatives of Biotin Protein Ligase aid in understanding multiple carboxylase deficiency. *Biochemica Biophysica Acta – Proteins and Proteomics*, Jul-Aug 2008, 1784(7-8), 973-82. PMID: 18442489

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Chapter 4:

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Communications and Presentations

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Pendini, N. R., Polyak, S.W., Booker G.W., Wallace J.C. and Wilce, M.C., *Staphylococcus aureus* Biotin Protein Ligase as a novel antibiotic target (2009), 34th Lorne Conference on Protein Structure and Function. Abs#157.

Pendini, N. R., Polyak, S.W., Booker G.W., Wallace J.C. and Wilce, M.C., *Staphylococcus aureus* Biotin Protein Ligase as a novel antibiotic target (2008), University of Adelaide, School of Molecular and Biomedical Sciences Research Symposia. **Awarded** \$200 for best poster presentation.

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Pendini, N. R., Polyak, S.W., Booker G.W., Wallace J.C. and Wilce, M.C., Biotin Protein Ligase as a novel antibacterial target (2008), 33rd Lorne Conference on Protein Structure and Function. Abs#361. **Awarded** student travel award \$100.

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Pendini, N. R., Polyak, S.W., Booker G.W., Wilce, M.C. and Wallace J.C., Determination of the Structure of Yeast Biotin Protein Ligase: Implications as a Novel Antifungal Drug Target (2006), Australian Society for Medical Research SA division Scientific Meeting, invited presenter. **Awarded:** \$400 for Best oral presentation by an ASMR student member.

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Pendini, N. R., Polyak, S.W., and Wallace J.C., 2005-Combio
The purification of Eukaryotic Biotin Protein Ligase.

Pendini, N. R., Polyak, S.W., and Wallace J.C., The purification of yeast Biotin Protein Ligase (2005). The East Coast Protein Structure Conference.

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Invited Presentations at National Conferences

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Ng, B., Pendini N.R., Tieu W., Kuan K., Morona R., Abell A., Wilce M.C.J., Wallace J.C., Polyak S.W. and Booker G.W. (2008) Discovery of biotin protein ligase as a novel class of antibiotic. ComBio2008, Pos WED-024

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Mayende, L., Swift, R. D., Bailey, L. M., Pendini, N. R., Wallace, J. C. & Polyak, S. W. (2008) Domain structure of human holocarboxylase synthetase: evidence of an interaction between the N-terminal and C-terminal halves Proc. Lorne Conference on Protein Structure and Function, Pos 255

Polyak, S. W., Stojkoski, C., Pendini, N. R., Booker, G. W. & Wallace, J. C. (2007) Biotin protein ligase: A novel antibiotic target Proc. Lorne Conference on Protein Structure and Function, Pos 323

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