



AIMS:

- The cloning, expression and characterisation of prokaryotic chitin-binding proteins from *Serratia marcescens*, *Pseudomonas aeruginosa*, *Photobacterium luminescens* and *Photobacterium asymbiotica*
- Development of an assay to assess the activity of chitin-binding proteins
- Mutagenesis of chitin-binding proteins to alter glycan recognition patterns

1. Cloning of Chitin-Binding Proteins

Chitin-binding proteins were amplified from genomic DNA using PCR and subsequently cloned into protein expression vectors.

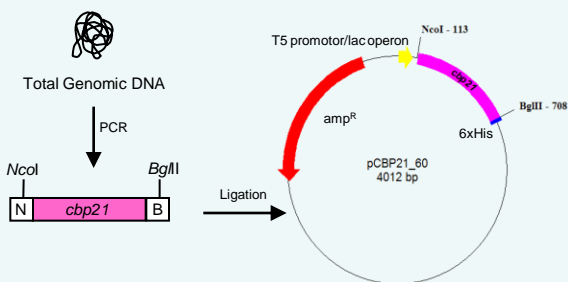


Figure 1: Overview of the cloning of *cbp21*. The *cbp21* gene was cloned into the pQE60 vector from Qiagen. The MSC is located before the (His)₆ amino acid sequence (blue) which allows for the expression of C-terminally (His)₆ tagged proteins. This is under control of the T5 promoter/lac operon (yellow). The *bla* gene encodes beta-lactamase which confers ampicillin resistance to the bacteria (red).

2. Expression of Chitin-Binding Proteins

Recombinant prokaryotic chitin-binding proteins were expressed in *E. coli* under the control of the lac operon with a C-terminal poly histidine tag, to facilitate downstream purification using Immobilised metal affinity chromatography (IMAC).

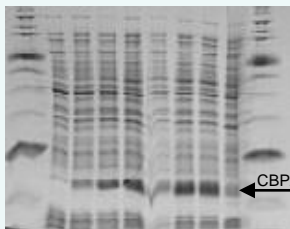


Figure 2: Time course expression analysis of CBP21 in *E. coli* KRX. Optimisation of expression of CBP21 resulted in up expression yields of up to 3.1 mg/g of cell paste

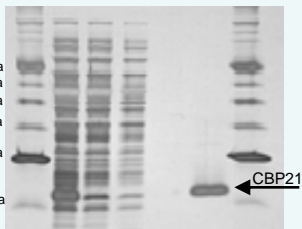
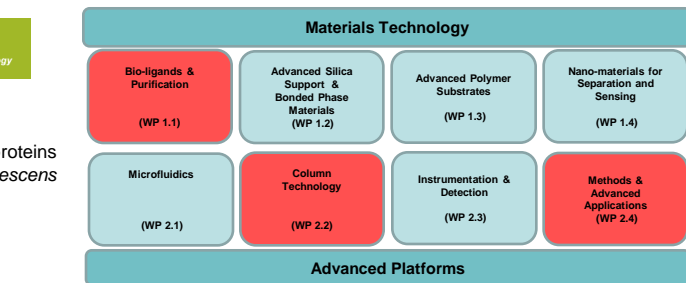


Figure 3: IMAC purification of CBP21. Analysis of purification fractions resulting from IMAC purification of CBP21 reveals that CBP21 purifies to homogeneity.



3. Assessing the activity of Chitin-binding proteins

Enzyme linked lectin assay (ELLA) analysis revealed that CBP21 was not capable of interacting with protein attached glycans in its wild type state (Figure 4). A novel assay to assess chitin-binding activity was developed using PAA-linked (GlcNAc)_N polymers (Figure 5).

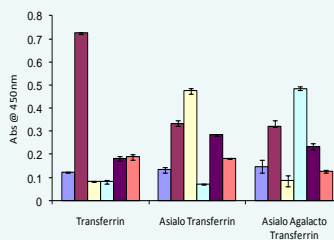


Figure 4: ELLA analysis of CBP21.

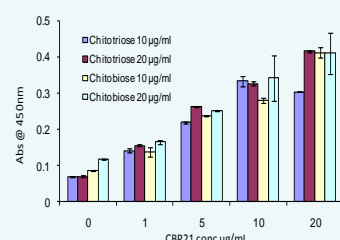


Figure 5: CBP21-(GlcNAc)_N activity assay.

4. Mutagenesis of Chitin-binding proteins

Site directed mutagenesis of amino acids thought to be involved in chitin-binding was undertaken (Figure 6). Mutagenesis did not improve the affinity for protein attached glycans, although some differences in affinities for the insoluble substrates; chitin, chitosan and cellulose were observed (Table 1).

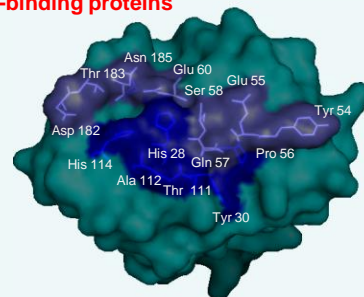


Figure 6: Residues involved in CBP21 binding to chitin.

Table 1: Overview of changes in affinities of CBP21 mutants.

Protein	β-chitin	α-chitin	Chitosan	Cellulose
WT	✓	✓	✓	✓
Y54A	--	--	✓	X
E55A	--	✓	✓	✓
P56A	--	--	✓	✓
Q57A	--	--	++	✓
S58A	--	--	✓	✓
E60A	--	--	✓	--
T111A	--	✓	+	++
H114A	X	--	+	--
D182A	--	✓	✓	++

✓ Binding comparable to WT, slight increase in affinity compared to WT (+), larger increase in affinity compared to WT (++), slightly decrease in affinity compared to WT (-), larger decrease in affinity compared to WT (--), no binding (X).

5. Project Outputs

Poster Presentations

- ISSC Review Meeting, DCU, June 2010.
- DCU School of Biotechnology Research Day, DCU, January 2010.
- UNCSR 10th Anniversary Symposium, The Helix, Dublin City University, October 2009.

• 9th Jenner Medicine and Glycobiology Conference, Royal Academy of Medicine of Belgium, Brussels, September 2009.

This research has been funded by Science Foundation Ireland under grant number 08/SRC/B1412 and the IRCSET Post Graduate Scholarship Scheme