EFFECT OF IRON IN BACTERIAL GROWTH & FBP PRODUCTION Faculty Member(s) Student(s) Gökşin Liu Gizem Çile İlknur Şafak Demirel Zehra Sayers

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PROGRAM FOR UNDERGRADUATE RESEARCH

ABSTRACT

FBP is an iron binding protein in gram negative bacteria that has the molecular weight of 37 kDa¹. It sequesters iron from transferrin which is an iron binding protein in humans. If the iron-binding of FBP is prevented, the bacteria would die as it can not take iron from transferrin? Since an alternative drug to antibiotics can be found, further investigations of FBP structure is needed. We focused on the optimization of iron concentration to synthesize more recombinant FBP in bacteria to be used in structural studies. After the optimization of iron concentration, we have made the purification of our protein of interest.

METHODS & RESULTS: Purification

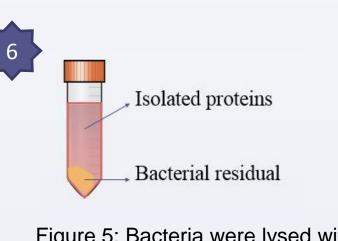


Figure 5: Bacteria were lysed with **BugBuster protocol**

6-Lysis with BugBuster Protocol

The pellet was lysed and centrifuged and isolated proteins containing supernatant was taken.

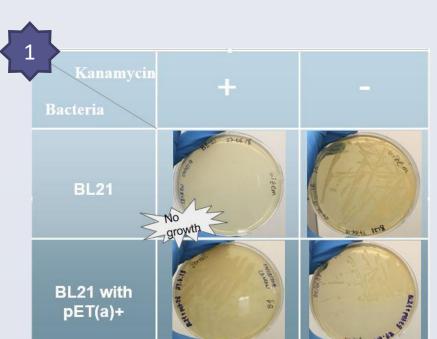
7-FBP Purification

Supernatant were poured into Ni-affinity column, Washing Buffer and Elution Buffer were added respectively. In each steps, samples were taken.

OBJECTIVES

As *E.Coli* do not normally express FBP, the plasmid pET28a (+) containing Kanamycin resistance and FBP gene was inserted into BL21 strain of *E.Coli* which is ideal for protein expression. In order to investigate the structure of FBP, the expression of FBP in BL21 should be increased via optimizations. Therefore in this study, we worked on the concentrations of iron for optimizing expression and performed purification to obtain the expressed FBP and for its quality control.

METHODS & RESULTS: Expression



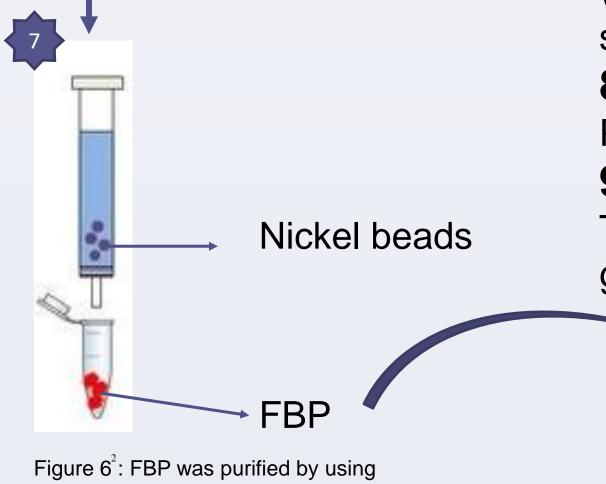
1-Plate Preparation

Bacteria were seeded as seen in Table 1.

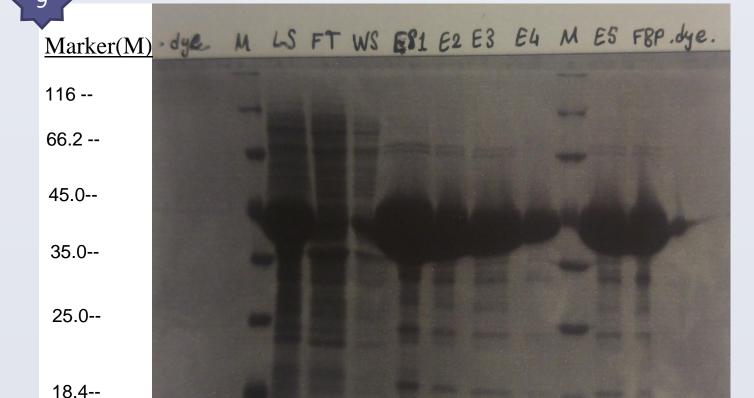
2-Growth Culture

A chosen colony from Kanamycin environment was added to four flasks in 100 ml LB medium and growth at 37 °C was monitored at different iron concentrations. **3-FBP Expression via Induction Culture** When the OD₆₀₀ value reaches to 0.65, IPTG was added

into 100 ml of bacterial culture in four flasks with different



Nickel affinity column. Elution Buffer has Imidazole in it that binds to Nickel to elute FBP.



8-Dialysis

PBS Buffer was used.

9-SDS-PAGE Analysis

The samples were runned in SDS-PAGE gel electrophoresis.





Figure 7: The remaining Imidazole was removed by dialysis with PBS Buffer.

Figure 8: Purified FBP after dialysis.



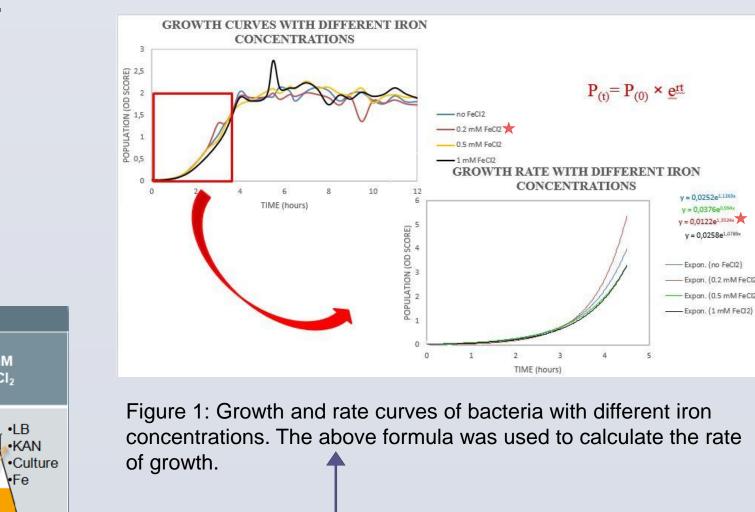


Table 1: Four different plates were prepared with and without pET28a(+) in KAN(+) and KAN(-) environments.

iron concentration to induce FBP gene expression.

4-Data Analysis

Optical densities at 600 nm were recorded every hour and graphs were plotted to check the growth rates of the populations.



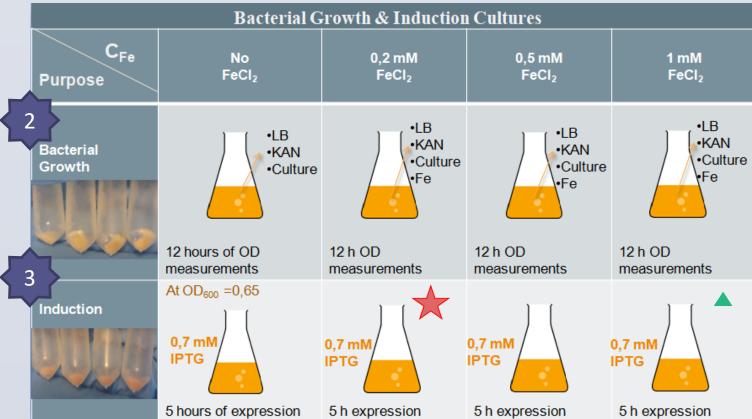


Table 2: Bacterial growth and induction cultures with four different iron concentrations.

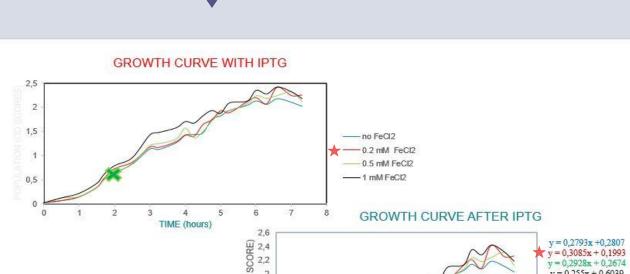
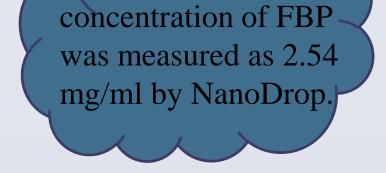




Figure 9: SDS-PAGE gel running.



LS: Supernatant after lysis, FT: Flow through, WS:Sample after washing, ES(1-5): Samples after elution, FBP: Sample after dialysis

CONCLUSION

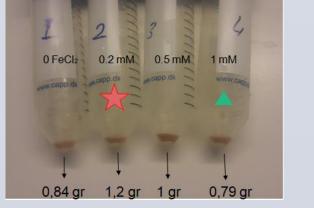
As four different FeCl₂ concentrations of 0 mM, 0.2 mM, 0.5 mM and 1 mM were examined, the samples with 0.2 mM of iron concentration has;

- The highest yield with 1.2 gr,
- The highest growth rate obtained from analysis and modelling of OD₆₀₀ data,
- Thicker bands at 37 kDa in SDS-PAGE gel screening which indicates higher FBP expression.

Therefore, we concluded that the best iron concentration to be used in further experiments is 0.2 mM for both the optimum bacterial growth and **FBP** production.

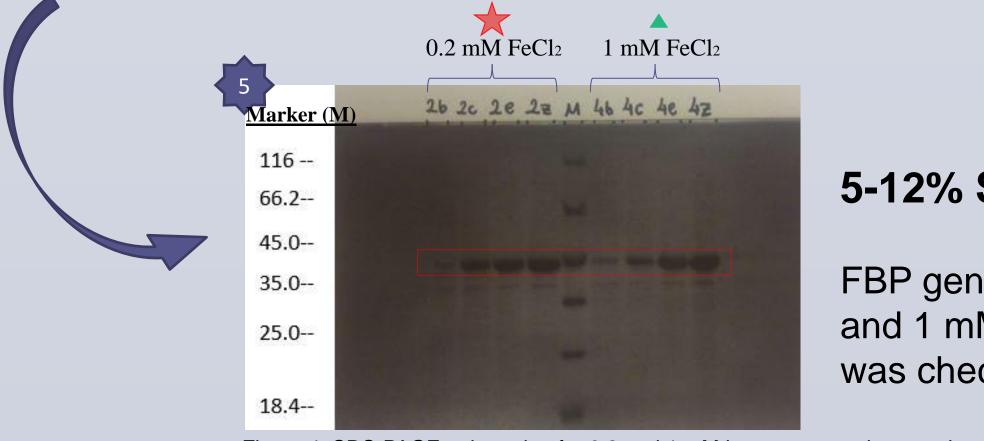
REFERENCES

Liu, G. (2015, December). Design and development of a microfluidic device



r = 0,255x + 0,603- no FeCl - 0.2 mM FeCI2 - 0.5 mM FeCl2 -1 mM FeCl2 Figure 2: Growth curves of bacteria with IPTG addition

Figure 3: Different yields with different iron concentrations. When FBP is bound to iron source, the color turns to pink.



5-12% SDS-PAGE Analysis

FBP gene expression of 0.2 mM and 1 mM iron concentrations was checked.

Figure 4: SDS-PAGE gel running for 0.2 and 1 mM iron concentrations. t= time after IPTG addition (expression). Samples b,c,e,z: t=0,1,3,5 respectively.

- to monitor iron binding dynamics in iron transport proteins.
- 2. Krewulak, K. D., & Vogel, H. J. (2007, August 19). Structural biology of bacterial iron uptake. Retrieved from

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