

行政院原子能委員會
委託研究計畫研究報告

纖維原料水解液高效率發酵菌株之研究及程序開發

**Study of the effective cellulose hydrolysate-fermentative microbe
and development of the microbe-based process**

計畫編號：1022001INER039

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報告日期：102 年 11 月 22 日

目 錄

目 錄.....	I
中文摘要.....	1
英文摘要.....	2
壹、計劃緣起與目的.....	3
貳、研究方法與過程.....	7
一、建構染色體操作工具箱.....	7
二、剷除競爭性的基因.....	8
三、強化關鍵基因的表現.....	9
四、生產菌之醱酵.....	10
參、主要發現與結論.....	10
一、剔除競爭代謝路徑.....	10
二、工程導引三羧酸循環路徑的碳流方向.....	11
三、強化三羧酸循環路徑的關鍵反應步驟.....	12
四、增加前驅物草醋酸的生成.....	13
五、生產菌之初步醱酵檢測.....	13
肆、參考文獻.....	14

中文摘要

為了發展大腸桿菌之琥珀酸發酵的技術平台，在今年度的計劃中我們所提議的主要策略在於改造細胞的代謝途徑，以導引碳流至琥珀酸，目前獲致的成果如下：(1)完成基因剔除工具箱的建構，藉此剔除競爭代謝路徑，以減少能源的消耗和碳流量的損失；(2)完成原位鑲箱啟動子工具箱的建構，藉此強化關鍵基因的表現，以導引碳流匯流至琥珀酸；(3)進行先期發酵條件之探討，結果發現建構菌株可由 30 g/L 葡萄糖轉化生成 10 g/L 琥珀酸，莫爾轉化率達 61%。

關鍵字：代謝工程、染色體工程、琥珀酸

Abstract

This study is aimed at to develop a technique platform for fermentative production of succinic acid in *Escherichia coli*. In this year, the main strategy as proposed is to engineer metabolic pathways of *E. coli* in order to redirect the carbon flux towards succinate. Our current results are as follows. (1) The construction of a gene deletion toolbox was completed. The competing pathways were then removed using this toolbox to curtail the waste of energy and byproduct production. (2) The construction of a promoter-insertion toolbox. The key pathways were enhanced using this toolbox to channel the carbon flux into succinate node. (3) The preliminary study of succinate fermentation was conducted. As a consequence, the engineered strain was able to produce 10 g/L succinate from 30 g/L glucose. This result accounts for a molar conversion yield of 61%.

Keywords: Metabolic engineering, Genomic engineering, succinic acid