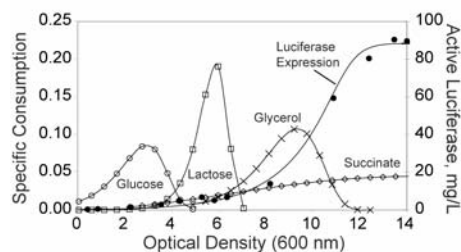


# Center for Eukaryotic Structural Genomics

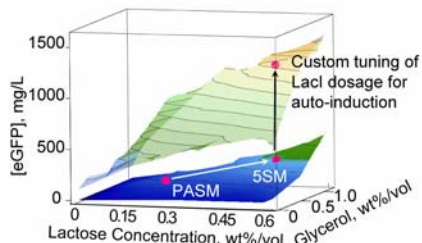
## Technology Dissemination Report

<b>CESG Tech Report No.</b>	006
<b>Title</b>	Factorial Evolved Auto-Induction Medium
<b>Research Unit</b>	Small-Scale Expression Testing and Large-Scale Protein Production
<b>Authors</b>	Blommel, P.G., Becker, K.J., Duvnjak, P. and Fox, B.G.
<b>Primary Contact</b>	<a href="mailto:bgfox@biochem.wisc.edu">bgfox@biochem.wisc.edu</a>

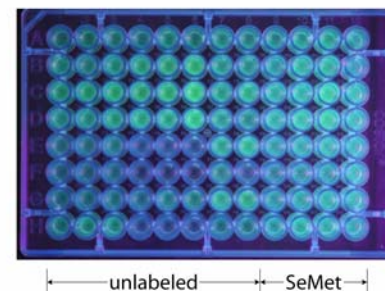
### A. Diauxy during auto-induction



### B. Medium and vector effects



### C. Medium composition array



## Summary

The auto-induction method of protein expression in *E. coli* is based on diauxic growth resulting from dynamic function of *lac* operon regulatory elements (*lacO* and *LacI*) in mixtures of glucose, glycerol, and lactose. Our results show that successful execution of auto-induction is strongly dependent on the plasmid promoter and repressor construction, on the oxygenation state of the culture, and on the composition of the auto-induction medium. Thus expression hosts expressing high levels of *LacI* during aerobic growth exhibit reduced ability to effectively complete the auto-induction process. Manipulation of the promoter to decrease the expression of *LacI* altered the preference for lactose consumption in a manner that led to increased protein expression and partially relieved the sensitivity of the auto-induction process to the oxygenation state of the culture. Factorial design methods were used to optimize the chemically defined growth medium used for expression of two model proteins, *Photinus* luciferase and enhanced green fluorescent protein, including variations for production of both unlabeled and selenomethionine-labeled samples [1]. The optimization included studies of the expression from T7 and T7-*lacI* promoter plasmids and from T5 phage promoter plasmids expressing two levels of *LacI*. Upon the basis of the analysis of over 500 independent expression results, combinations of optimized expression media and expression plasmids that gave protein yields of greater than 1000  $\mu\text{g}/\text{mL}$  of expression culture were identified.

These approaches are incorporated into CESG Technology Dissemination Reports 010, 020, 021, and 022.

Publication:

- [1] Blommel, P.G., Becker, K.J., Duvnjak, P. and Fox, B.G. (2007) Enhanced bacterial protein expression during auto-induction obtained by alteration of *lac* repressor dosage and medium composition. *Biotechnol Prog* 23(3):585-98.

<b>Acquiring the Technology</b>	See publication.
<b>Other Acknowledgements</b>	Also supported by Promega Corporation, Madison, WI (B.G. Fox, PI).
Center for Eukaryotic Structural Genomics (CESG), University of Wisconsin-Madison Biochemistry Department, 433 Babcock Drive, Madison, WI 53706-1549; phone: 608.263.2183; fax: 608.890.1942; email: <a href="mailto:cesginfo@biochem.wisc.edu">cesginfo@biochem.wisc.edu</a> ; website: <a href="http://www.uwstructuralgenomics.org">http://www.uwstructuralgenomics.org</a> . This research funded by NIH / NIGMS Protein Structure Initiative grants U54 GM074901 and P50 GM064598.	