### Accepted Manuscript

Microbial expression systems for membrane proteins

Marvin V. Dilworth, Mathilde S. Piel, Kim E. Bettaney, Pikyee Ma, Ji Luo, David Sharples, David R. Poyner, Stephane R. Gross, Karine Moncoq, Peter J. F. Henderson, Bruno Miroux, Roslyn M. Bill

PII:S1046-2023(17)30377-8DOI:https://doi.org/10.1016/j.ymeth.2018.04.009Reference:YMETH 4446To appear in:MethodsReceived Date:18 January 2018Revised Date:8 March 2018Accepted Date:10 April 2018



Please cite this article as: M.V. Dilworth, M.S. Piel, K.E. Bettaney, P. Ma, J. Luo, D. Sharples, D.R. Poyner, S.R. Gross, K. Moncoq, P. J. F. Henderson, B. Miroux, R.M. Bill, Microbial expression systems for membrane proteins, *Methods* (2018), doi: https://doi.org/10.1016/j.ymeth.2018.04.009

This is a PDF file of an unedited manuscript that has been accepted for publication. As a service to our customers we are providing this early version of the manuscript. The manuscript will undergo copyediting, typesetting, and review of the resulting proof before it is published in its final form. Please note that during the production process errors may be discovered which could affect the content, and all legal disclaimers that apply to the journal pertain.

## ACCEPTED MANUSCRIPT

#### MICROBIAL EXPRESSION SYSTEMS FOR MEMBRANE PROTEINS

Marvin V. Dilworth<sup>1,2#</sup>, Mathilde S. Piel<sup>3#</sup>, Kim E. Bettaney<sup>4#</sup>, Pikyee Ma<sup>4,5</sup>, Ji Luo<sup>4</sup>, David Sharples<sup>4</sup>, David R. Poyner<sup>1</sup>, Stephane R. Gross<sup>1</sup>, Karine Moncoq<sup>3</sup>, Peter J. F. Henderson<sup>4\*</sup>, Bruno Miroux<sup>3\*</sup> and Roslyn M. Bill<sup>1\*</sup>

<sup>1</sup>School of Life & Health Sciences, Aston University, Aston Triangle, Birmingham, B4 7ET, UK

<sup>2</sup>Current address: Department of Chemistry, King's College London, Britannia House, 7 Trinity Street, London, SE1 1DB, UK

<sup>3</sup>Laboratoire de Biologie Physico-Chimique des Protéines Membranaires, UMR 7099 (CNRS - Université Paris Diderot), Institut de Biologie Physico-Chimique, 13 rue Pierre et Marie Curie, 75005 Paris, France

<sup>4</sup>Astbury Centre for Structural Molecular Biology and School of Biomedical Sciences, University of Leeds, Leeds, LS2 9JT, UK

<sup>5</sup>Current address: Paul Scherrer Institute, CH-5232 Villigen PSI, Switzerland

<sup>#</sup>Equal contributions by: Marvin V. Dilworth, Mathilde Piel and Kim E. Bettaney

\*Correspondence to: Peter J. F. Henderson (+44 113 343 3175; p.j.f.henderson@leeds.ac.uk), Bruno Miroux (+33 1 58 41 52 25; bruno.miroux@ibpc.fr), Roslyn M. Bill (+44 121 204 4274; r.m.bill@aston.ac.uk)

#### Abstract

Despite many high-profile successes, recombinant membrane protein production remains a technical challenge; it is still the case that many fewer membrane protein structures have been published than those of soluble proteins. However, progress is being made because empirical methods have been developed to produce the required quantity and quality of these challenging targets. This review focuses on the microbial expression systems that are a key source of recombinant prokaryotic and eukaryotic membrane proteins for structural studies. We provide an overview of the host strains, tags and promoters that, in our experience, are most likely to yield protein suitable for structural and functional characterization. We also catalogue the detergents used for solubilization and crystallization studies of these proteins. Here, we emphasize a combination of practical methods, not necessarily high-throughput, which can be implemented in any laboratory equipped for recombinant DNA technology and microbial cell culture.

Download English Version:

# https://daneshyari.com/en/article/10156928

Download Persian Version:

https://daneshyari.com/article/10156928

Daneshyari.com