Engineering Conferences International ECI Digital Archives

Integrated Continuous Biomanufacturing II

Proceedings

Fall 11-3-2015

Design and control of chromatography step in an integrated column sequence

Bernt Nilsson

Dept. of Chemical Engineering, Lund University, bernt.nilsson@chemeng.lth.se

Niklas Andersson

Dept. of Chemical Engineering, Lund University

Anders Holmqvist

Dept. of Chemical Engineering, Lund University

Anton Sellberg

Dept. of Chemical Engineering, Lund University

Peter Tiainen

Novo Norsdisk

See next page for additional authors

Follow this and additional works at: http://dc.engconfintl.org/biomanufact ii



Part of the Biomedical Engineering and Bioengineering Commons

Recommended Citation

Bernt Nilsson, Niklas Andersson, Anders Holmqvist, Anton Sellberg, Peter Tiainen, and Arne Staby, "Design and control of chromatography step in an integrated column sequence" in "Integrated Continuous Biomanufacturing II", Chetan Goudar, Amgen Inc. Suzanne Farid, University College London Christopher Hwang, Genzyme-Sanofi Karol Lacki, Novo Nordisk Eds, ECI Symposium Series, (2015). http://dc.engconfintl.org/biomanufact_ii/77

This Conference Proceeding is brought to you for free and open access by the Proceedings at ECI Digital Archives. It has been accepted for inclusion in Integrated Continuous Biomanufacturing II by an authorized administrator of ECI Digital Archives. For more information, please contact franco@bepress.com.

Authors Bernt Nilsson, Niklas Andersson, Anders Holmqvist, Anton Sellberg, Peter Tiainen, and Arne Staby

DESIGN AND CONTROL OF CHROMATOGRAPHY STEPS IN AN INTEGRATED COLUMN SEQUENCE

Bernt Nilsson, Dept. of Chemical Engineering, Lund University, Sweden bernt.nilsson@chemeng.lth.se

Niklas Andersson, Dept. of Chemical Engineering, Lund University, Sweden Anders Holmqvist, Dept. of Chemical Engineering, Lund University, Sweden Anton Sellberg, Dept. of Chemical Engineering, Lund University, Sweden Peter Tiainen, Novo Nordisk A/S, Måløv, Denmark Arne Staby, Novo Nordisk A/S, Bagsvaerd, Denmark

Key words: Chromatography, Integration, Optimization, Recycle, Robustness

Integration of the purification steps in downstream processing is one way to minimize the number of storage tanks and reduce hold-up time. The result is an integrated column sequence that performs straight through processing of the target protein, with minimal time from expression to formulation. This paper discusses the overall integration design and its expected performance, together with a more detailed analysis of the step behavior based on column simulations. The optimal design of the individual steps in the sequence is presented, both with robustness margins and local control system, as well as with and without local recycle.

An optimization methodology for chromatography steps is used to quantify the performance of the integrated sequence. The propagation of disturbances in connected column sequences forces an open-loop control strategy to be unnecessary conservative. The optimization is modified to also handle disturbances which makes it possible to find the robust optimum. The introduction of local control for disturbance compensation and for guarantee of optimal robust performance is shown for transient control of the dynamic cyclic steady-state.

For some separation problems, it is attractive to introduce local recycles in the process, such as in MCSGP, to enhance the yield of the target protein. These recycles are also introduced in the optimization methodology of separation steps in integrated column sequences, generating optimal robust solutions with recycles. The result shows the advantages and disadvantages of local recycles for the case study analyzed in the work.