## **Engineering Conferences International ECI Digital Archives**

Cell Culture Engineering XV

**Proceedings** 

Spring 5-10-2016

## Critical process parameter identification using the ambr15(tm) for process characterization

Matthew Zustiak Patheon Biologics, matt.zustiak@patheon.com

Michael Bennett Patheon Biologics

Stephanie Chakravarty Patheon Biologics

Matt Caple Patheon Biologics

Follow this and additional works at: http://dc.engconfintl.org/cellculture xv



Part of the Biomedical Engineering and Bioengineering Commons

## Recommended Citation

Matthew Zustiak, Michael Bennett, Stephanie Chakravarty, and Matt Caple, "Critical process parameter identification using the ambr15(tm) for process characterization" in "Cell Culture Engineering XV", Robert Kiss, Genentech Sarah Harcum, Clemson University Jeff Chalmers, Ohio State University Eds, ECI Symposium Series, (2016). http://dc.engconfintl.org/cellculture\_xv/103

This Abstract is brought to you for free and open access by the Proceedings at ECI Digital Archives. It has been accepted for inclusion in Cell Culture Engineering XV by an authorized administrator of ECI Digital Archives. For more information, please contact franco@bepress.com.

## CRITICAL PROCESS PARAMETER IDENTIFICATION USING THE AMBR15™ FOR PROCESS CHARACTERIZATION

Matthew Zustiak, Patheon Biologics
Matt.zustiak@patheon.com
Michael Bennett, Patheon Biologics
Stephanie Chakravarty, Patheon Biologics
Matt Caple, Patheon Biologics

Key Words: Process characterization, Critical process parameters, DoE.

Process characterization is a critical phase in the development of a commercial process for biotherapeutic production. Knowing the critical quality attributes of your molecule prior to beginning process development and/or characterization is imperative when using a quality by design (QbD) approach. Here we use a (QbD) approach for the characterization of a fed-batch process using an NSO cell line to express an IgG. For this molecule, the glycosylation profile, and in particular, the total fucosylation was identified as a critical quality attribute. After performing a primary hazard analysis, several process inputs were determined to potentially have an impact on this critical quality attribute. These parameters were then studied in a screening DoE using the ambr15<sup>™</sup> to model the first and second order effects for each parameter on both the critical quality attributes and process performance. Of the 9 parameters studied, 5 were determined to have a statistically significant effect on the fucosylation of the molecule. In addition, 6 parameters were identified to have a significant impact on process performance. Through process modeling using JMP, a design space was determined for further studies to determine the proven acceptable range (PAR) for each parameter using the 10L, qualified scale down model. An example of the predicted PAR for pH and the timing of the temp shift can be seen in figure 1. Following the 10L studies, a PAR was determined for each parameter and compared with the predicted PAR from ambr. Here we demonstrate the feasibility to use the ambr15™ as a tool for key and even critical process parameter identification to reduce timelines for process characterization.

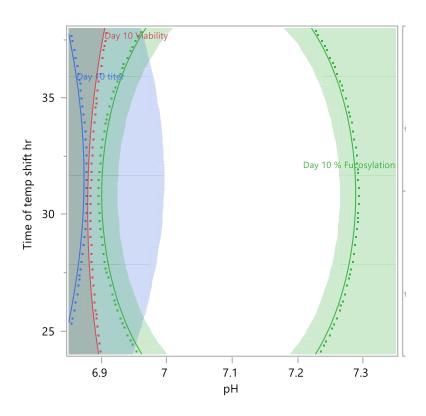


Figure 1 – Design space for critical process parameters satisfying the critical quality attributes of an IgG