

**A SYSTEMS BIOLOGY APPROACH FOR
UNDERSTANDING OSMOTIC STRESS IN
ANTIBODY-PRODUCING CELL LINES**

By

Thomas R. Kiehl

An Abstract of a Thesis Submitted to the Graduate
Faculty of Rensselaer Polytechnic Institute
in Partial Fulfillment of the
Requirements for the Degree of
DOCTOR OF PHILOSOPHY

Major Subject: MULTIDISCIPLINARY SCIENCE

The original of the complete thesis is on file
in the Rensselaer Polytechnic Institute Library

Examining Committee:

Susan T. Sharfstein, Thesis Adviser
Lealon L. Martin, Member
Joyce J. Diwan, Member
David Isaacson, Member
Samuel C. Wait, Jr., Member

Rensselaer Polytechnic Institute
Troy, New York

July 2009
(For Graduation August 2009)

ABSTRACT

Growing demand for monoclonal antibodies (mAbs) presents a challenge to the biotechnology industry. Meeting these demands will require improvements in overall viability, proliferation and specific productivity of antibody-producing cell lines. Osmotic stress, when applied to these cultures, has been shown to improve specific productivity. In this work we investigate several aspects of the cellular response to osmotic stress. Well known signaling molecules are probed by western blotting and it is determined that industrially relevant antibody producing Chinese hamster ovary (CHO) cells exhibit constitutive activation of signaling species. It is also demonstrated that these signaling molecules are further activated under hyperosmotic conditions. Osmotic response element (ORE) binding proteins are shown by electrophoretic mobility shift assay to be present in antibody-producing CHO cells. These proteins are shown to increase their ORE binding behavior when these cells experience an increase in extracellular osmolarity. We show that transfecting CHO cells with active and inactive forms of the tonicity element binding protein (TonEBP), an osmotically responsive transcription factor, affects the growth and proliferation of those cells. Hyperosmotic stress is shown to have a dose dependent effect on the regulatory control of cell volume. Also, it is shown that populations of CHO cells under hyperosmotic stress contain a distinct subpopulation with larger cell diameters than those in the bulk of the population. Additionally, we demonstrate the generation of artificial biochemical reaction networks by using a genetic algorithm. We show that networks with specific stochastic behaviors can be generated by the appropriate tuning of the genetic algorithm. These techniques show good potential to be able to integrate broad sets of experimental data into functional models.