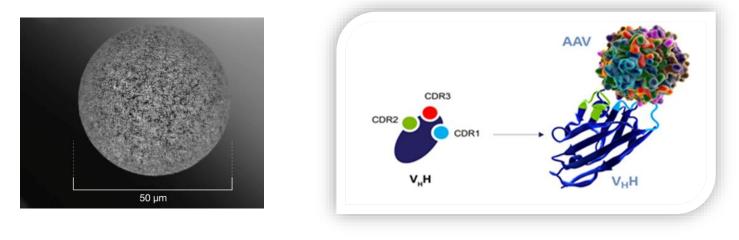
# Productivity optimization and process calculations for AAV affinity chromatography

## INTRODUCTION

The use of recombinant adeno-associated virus (rAAV) as a delivery method for gene therapies continues to be successful with hundreds of ongoing clinical trials and some recent approvals. The diversity of applications for rAAV ranges from rare diseases affecting small patient populations to more prevalent inherited ailments such as hemophilia. The doses required vary widely from ~4E11 vg/eye for subretinal administration to 3.5E14 vg for intrathecal applications [1]. From a manufacturing perspective the field has moved to common approaches for production and purification of rAAV. Upstream approaches typically use transfection of HEK293 cells and titers are routinely in the 1-2E10 vg/mL although higher titers of up to 6E11 vg/mL at a 2000 L scale were recently reported [2]. These high titers will be needed for large dose and/or patient populations to meet the demand of these therapies and reduce costs. For rAAV purification the majority of the field has moved to scalable processes employing an affinity capture chromatography step [3] and commonly utilizing POROS<sup>™</sup> CaptureSelect<sup>™</sup> AAVX resin. In this work, dynamic binding capacity (DBC) data for multiple AAV serotypes were leveraged to estimate an optimal productivity of rAAV using the AAVX resin. An analysis of process conditions and column geometries that would fit maximum processing times and resin utilization was conducted for two case scenarios representing current titers for clinical manufacturing and high titers for commercial manufacturing scales.



POROS<sup>™</sup> base bead technology (polystyrene divinylbenzene, left) and CaptureSelect<sup>™</sup> ligand technology (single-domain antibody, right) are combined in the manufacturing of AAVX resin

## **METHODOLOGY**

### **Dynamic binding capacity:**

AAV2 breakthrough curves were generated using HEK293 clarified lysate to determine DBC at 10% breakthrough. AAV8 and rh10 DBC data were obtained from references 4 and 5, respectively.

Equation I was fitted to the DBC data using a linear regression numerical method. **Productivity:** 

Productivity curves were generated using equations *I* and *II*.

Column volumes and residence time for the non-loading steps were 25 CV and 2 min. Column volumes and residence time for CIP steps were 10 CV and 3 min. Scale-up and process considerations:

GMP pre-packed column pressure limitations were based on literature from multiple vendors.

Pressure drop at 3 bar was based on pressure-flow curves for POROS CaptureSelect AAVX resin (internal pressure-flow data).

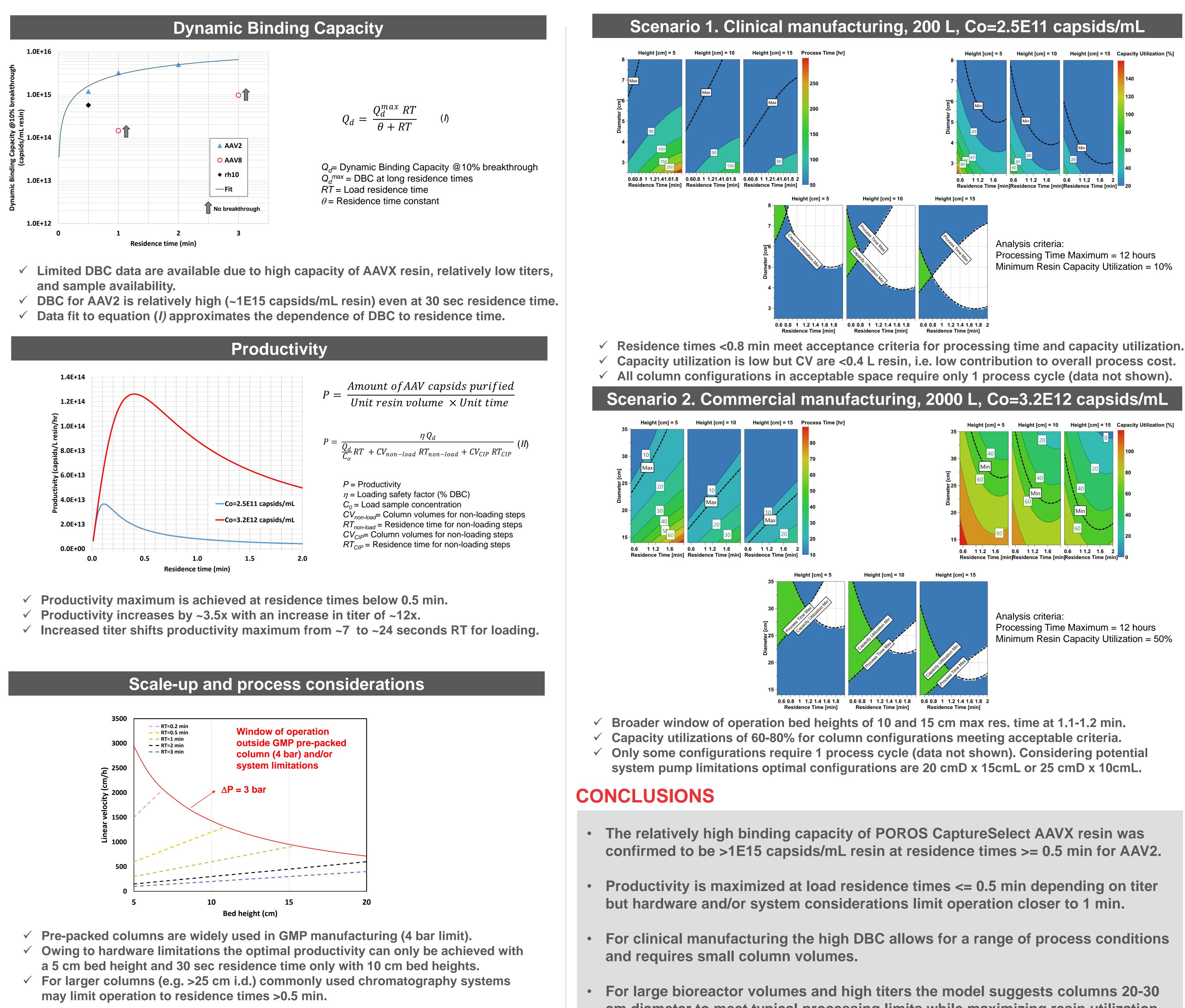
### **Case scenarios**

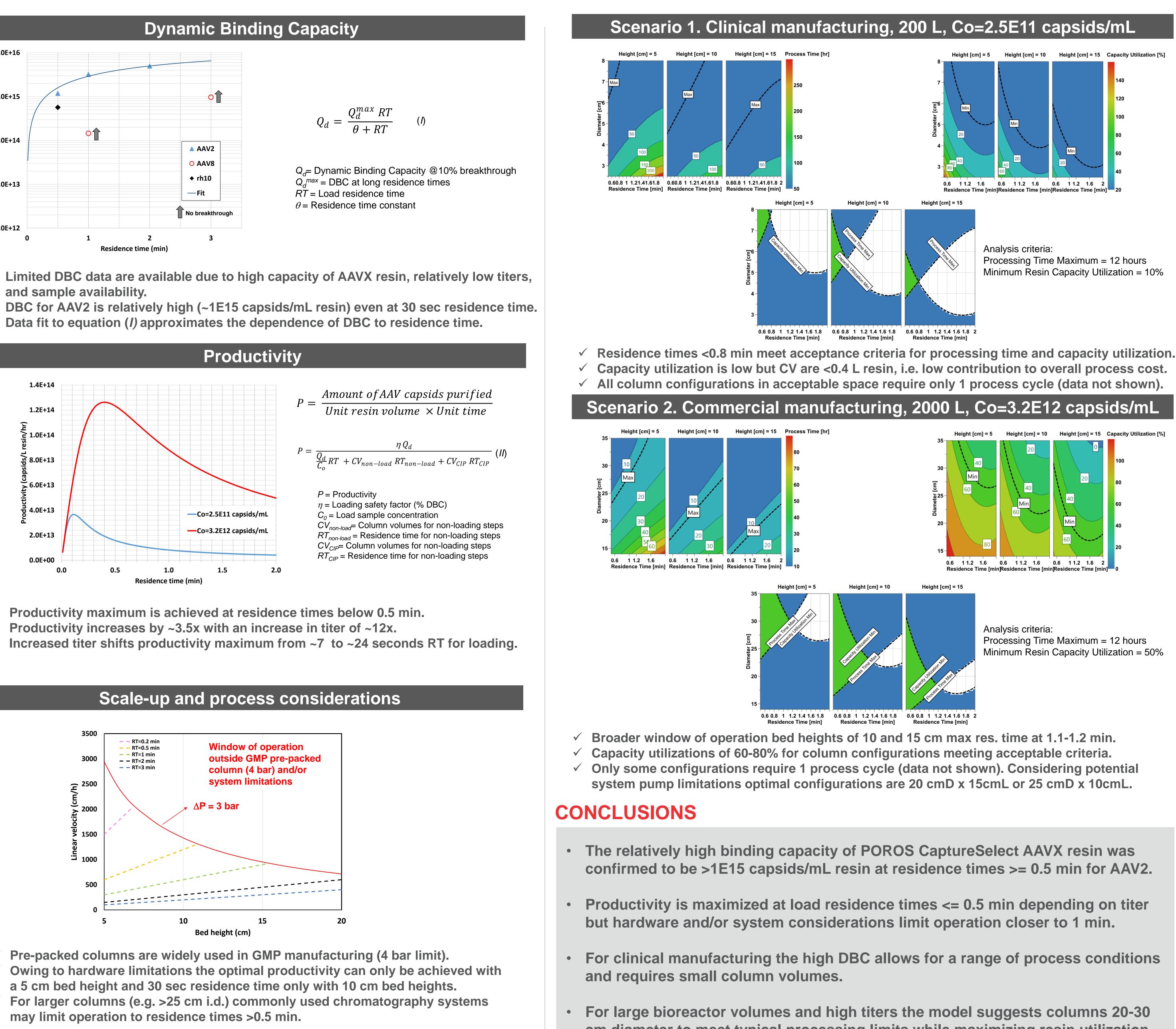
Processing time and resin utilization calculations were performed using Microsoft<sup>®</sup> Excel<sup>®</sup> assuming 20% full capsids and the results were further analyzed and plotted using MODDE<sup>®</sup> software.

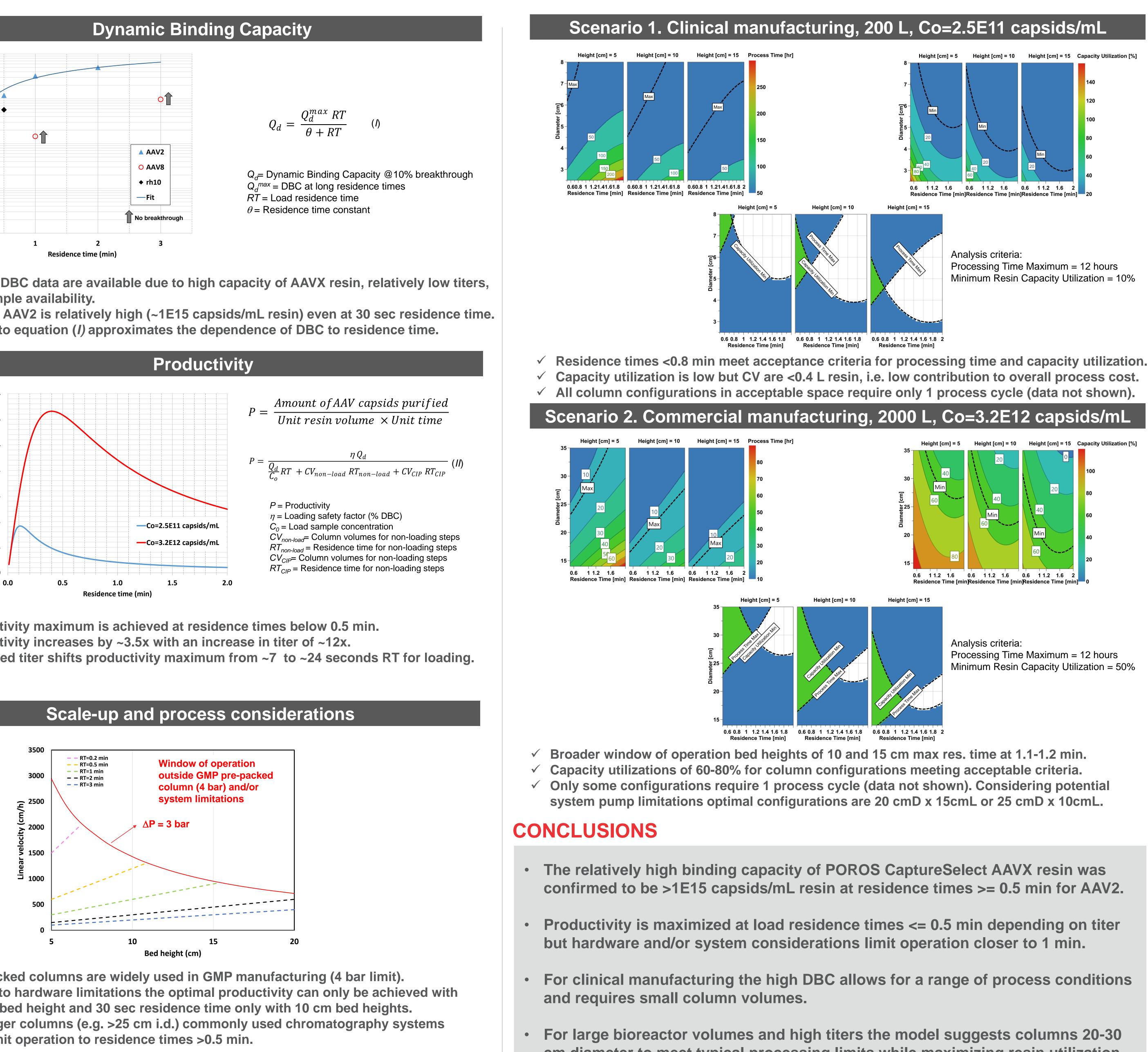
### REFERENCES

- . Burdett and Nuseibeh, Gene Therapy (2023)
- 2. Van Lieshout, et al. Molecular Therapy-Methods in Clinical Development (2023)
- 3. Adams et al., Biotechnology and Bioengineering (2020)
- 4. Ravault and Laroudie, Cell & Gene Therapy Insights (2022)
- 5. Hurwit and Morrison, ASGCT Meeting (2018)

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## Bioprocessing

cm diameter to meet typical processing limits while maximizing resin utilization.

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