

# Hybrid Dynamic Modeling and Identification

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## Park-Ramirez Bioreactor Model

For genetically modified yeast in a fed-batch reactor, predict:

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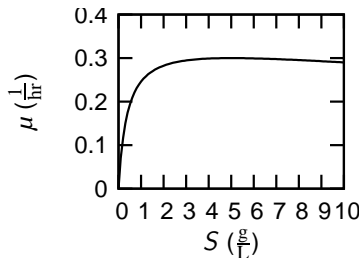
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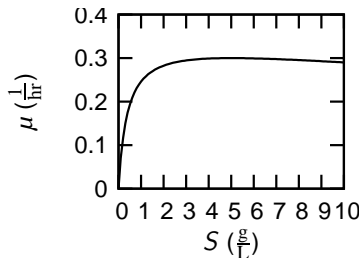


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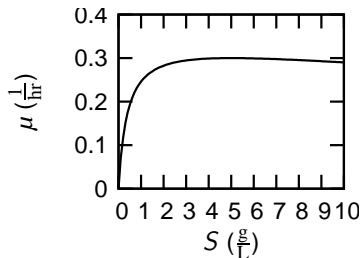


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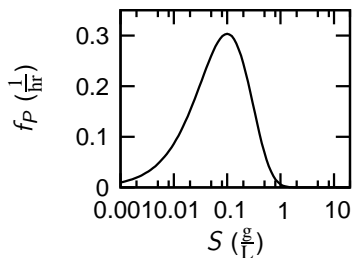
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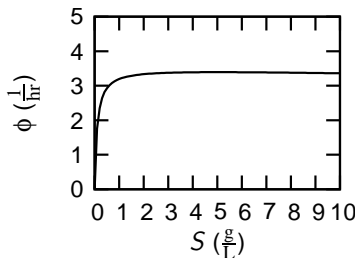
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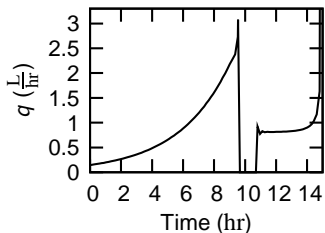
$$\begin{aligned} & \text{MAX}(\Phi) \\ \Phi &= P_M(t_f) \cdot V(t_f) \end{aligned}$$

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How to find  $\omega_1, \omega_2, \omega_3$ ?

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What if not much data available for a state?

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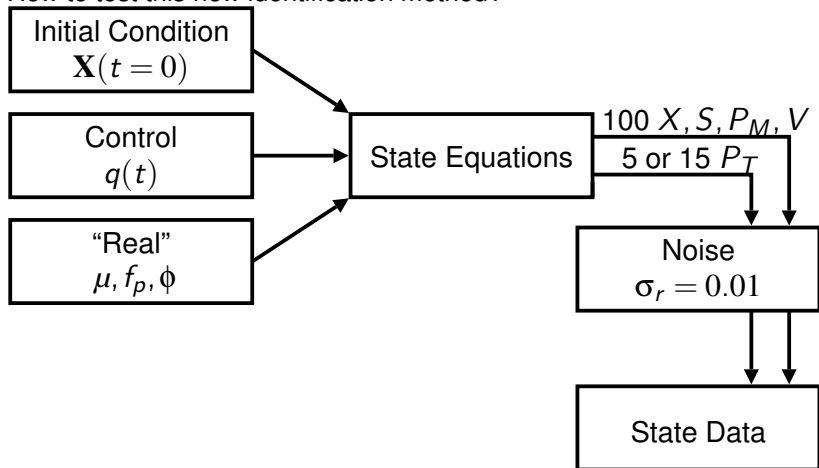
No limitations: if a system can be moved forward in time, this method can be used.

# Testing Optimal Control Identification

How to test this new identification method?

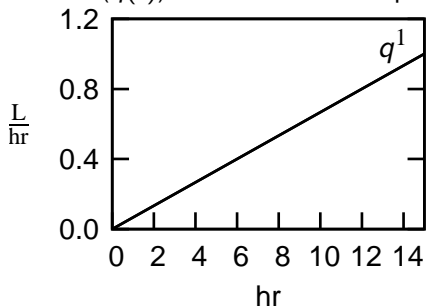
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## Control: $q(t)$

Control ( $q(t)$ ) for first set of examples.



# Identification Problem Using IDP Optimal Control Algorithm

- Animation: Two-step IDP with 25 stages

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(Back to Animation)

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Identify Park-Ramirez hybrid model with 8 artificial data runs:

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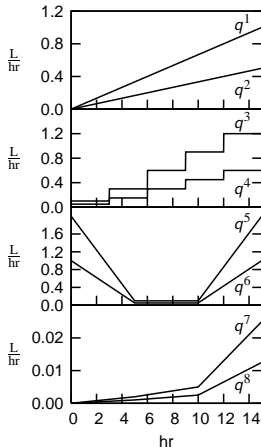
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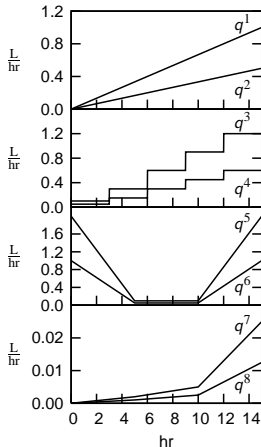
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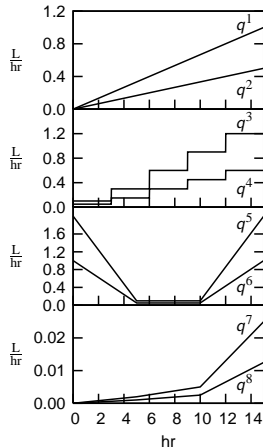
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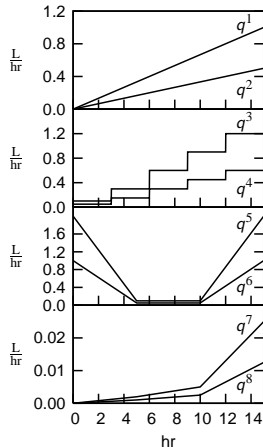
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- 4 Combine input-output pairs from all runs into a single set for each parameter function.

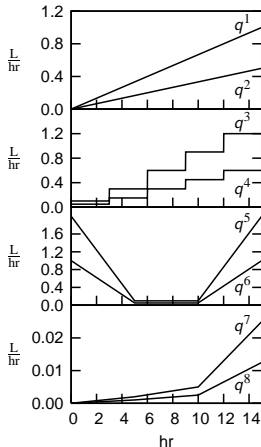




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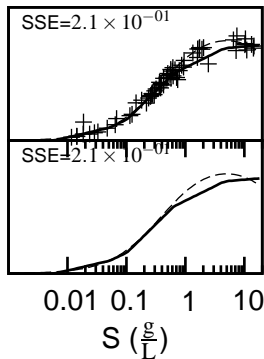
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- 5 Train neural networks on combined input-output set.



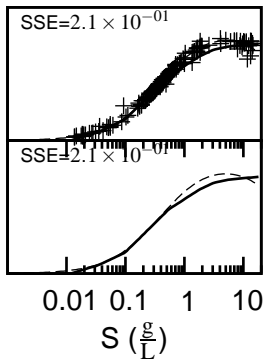
# Growth Function

$\mu$

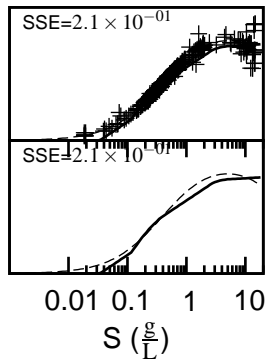
10 stages



25 stages



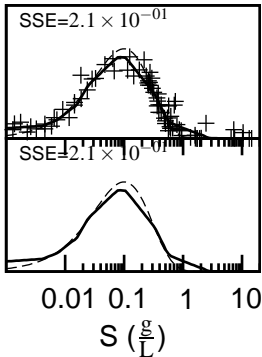
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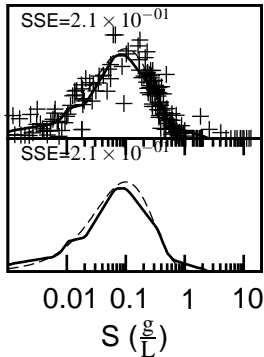
# Protein Production Function

$f_P$

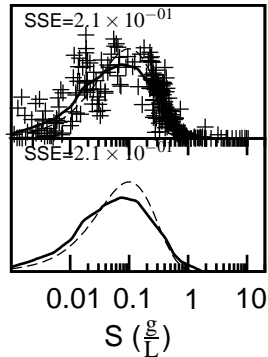
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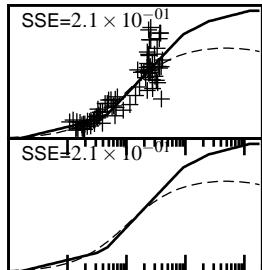
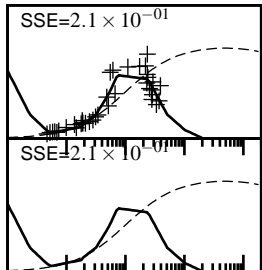
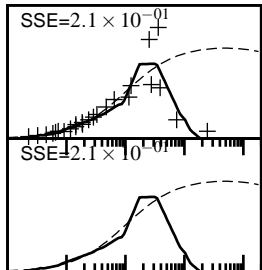
# Protein Secretion Function

$\phi$

10 stages

25 stages

50 stages

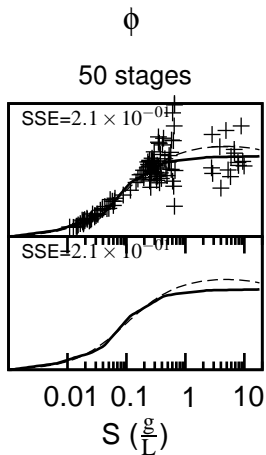


0.01 0.1 1 10  
 $S (\frac{g}{L})$

0.01 0.1 1 10  
 $S (\frac{g}{L})$

0.01 0.1 1 10  
 $S (\frac{g}{L})$

# Protein Secretion Function (glucose bump)



# Conclusions

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  - No sensitivity derivatives needed.
  - Any function estimator can be used (RBF neural nets seem especially promising).



# Acknowledgments

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