

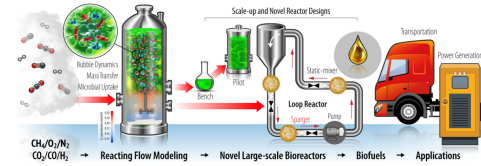
# Multiphysics computational fluid dynamics for design and scale-up of CO<sub>2</sub>/syngas bioreactors

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## Overview

**What are we trying to do:** Develop validated, high fidelity computational models for gas fermentation using CO<sub>2</sub>, CO and H<sub>2</sub>.  
**How is it done today?:** microbiological advances are made at the lab scale in well mixed reactors. Process scale-up remains an open question when considering novel microorganisms/reaction systems.  
**Why is this important?:** lab-to-production scale transition is required to achieve CO<sub>2</sub> conversion at Mton/Gton scale and computational models can de-risk/accelerate this transition.



**Goal:** enable scaled-up bioreactor design and optimization for CO<sub>2</sub> conversion through validated high fidelity Multiphysics simulations

Milestone	Criteria	Date	Status
Survey different methodologies to simulate bubble size distributions (BSD)	Literature on novel approaches to modeling and simulation of bubble breakage, aggregation, and mass transfer using population balances	3/3/2022	Completed
Kinetic mechanisms on bioreaction pathways	Selection/construction of lumped bioreaction kinetic models	6/30/2022	Completed
Influence of BSD on gas composition	Using high-fidelity Eulerian methods, determine differences in BSD and mass transfer for varying gas mixtures	9/30/2022	Completed
Hydrodynamics validation set-up and preliminary simulations towards go/no go decision	Quantity validation parameters and set-up preliminary simulations to compare against experiments reported in literature	12/31/2022	Completed
Meshing and preliminary simulation of sparger geometries	Create high resolution meshes for ring/porous plate sparger designs	03/31/2023	Completed
Novel design of sparging systems	Use multiphase flow models to explore various sparger designs such as ring/porous plates used at scale and explore variations in geometric parameters.	06/30/2023	85% completion

Table 1. Technical approach

## Computational models

$$\alpha_L + \alpha_G = 1$$

**Volume fraction constraint**

$$\frac{\partial}{\partial t}(\alpha_i \rho_i) + \nabla \cdot (\alpha_i \rho_i \mathbf{V}_i) = 0$$

**Mass conservation**

$$\frac{\partial}{\partial t}(\alpha_i \rho_i \mathbf{V}_i) + \nabla \cdot (\alpha_i \rho_i \mathbf{V}_i \mathbf{V}_i)$$

$$= -\alpha_i \nabla P + \alpha_i \rho_i \mathbf{g} + \nabla \cdot (\alpha_i \bar{\mathbf{R}}_i) + \mathbf{F}_i$$

**Momentum conservation**

$$\frac{\partial}{\partial t}(\alpha_i \rho_i Y_{ij}) + \nabla \cdot (\alpha_i \rho_i Y_{ij} \mathbf{V}_i)$$

$$= \nabla \cdot (\alpha_i \rho_i \bar{\mathbf{D}}_{ij} \nabla Y_{ij}) + \dot{R}_{ij}^{MT}$$

**species transport**

$$\frac{\partial n_v}{\partial t} + \nabla \cdot (v_{p,i} n_v) = h_v$$

**Bubble size distribution transport (population balances)**

$$h_v = \frac{1}{2} \int_0^\infty n_{v,coalescence} C_{v,coalescence} d'v - n_v \int_0^\infty n_{v,coalescence} d'v + \int_0^\infty n_{v,breakup} \beta_{v,breakup} d'v - \frac{n_v}{v} \frac{\partial \langle v \rangle}{\partial v} + \dot{n}_{v,translocation}$$

**Bubble dynamics**

## Lumped bioreaction kinetics

$$\frac{d\rho_{H_2}}{dt} = k_L a_{H_2} (\rho_{H_2}^* - \rho_{H_2}) - \left( \frac{\mu_{H_2}}{Y_{X,H_2}} + \frac{\mu_{P,H_2}}{Y_{P,H_2}} \right) \rho_{H_2}$$

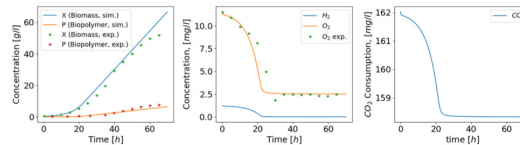
$$\frac{d\rho_{CO_2}}{dt} = k_L a_{CO_2} (\rho_{CO_2}^* - \rho_{CO_2}) - \left( \frac{\mu_{H_2}}{Y_{O_2,H_2}} + \frac{\mu_{P,H_2}}{Y_{P,O_2}} \right) \rho_{CO_2}$$

$$\frac{d\rho_{CO_2}}{dt} = k_L a_{CO_2} (\rho_{CO_2}^* - \rho_{CO_2}) - \left( \frac{\mu_{H_2}}{Y_{X,CO_2}} + \frac{\mu_{P,H_2}}{Y_{P,CO_2}} \right) \rho_{CO_2}$$

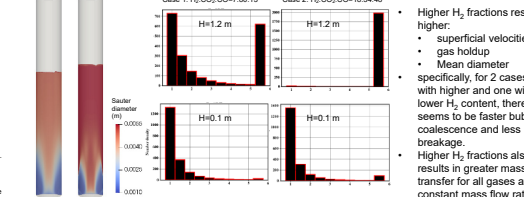
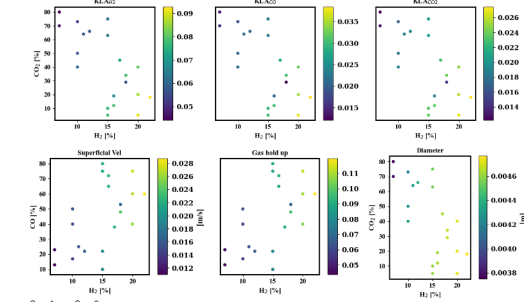
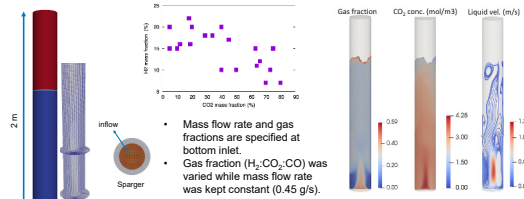
$$\mu_X = \mu_X^{max} \left( \frac{\rho_{H_2}}{K_{X,H_2} + \rho_{H_2}} \right) \times \left( \frac{\rho_{CO_2}}{K_{X,CO_2} + \rho_{CO_2}} \right) \times \left( \frac{\rho_{O_2}}{K_N + \rho_N + \frac{\rho_{CO_2}}{K_{PN}}} \right)$$

$$\mu_P = \mu_P^{max} \left( \frac{\rho_{H_2}}{K_{P,H_2} + \rho_{H_2}} \right) \times \left( \frac{\rho_{CO_2}}{K_{P,CO_2} + \rho_{CO_2}} \right) \times \left( \frac{\rho_{O_2}}{K_{PN} + \rho_N + \frac{\rho_{CO_2}}{K_{PN}}} \right) \times \left[ 1 - \left( \frac{f_P}{f_{P,max}} \right)^\beta \right] \times \frac{K_{IN}}{\rho_N + K_{IN}}$$

- We verify a mixture **Monod model** for a case of PHB production from the fermentation of CO<sub>2</sub>/H<sub>2</sub>O<sub>2</sub> gas mixtures in nitrogenous fermentation medium described by Mczunder et al.
- The kinetic model is general enough because it considers the effect of all gases with possible nutrient limitations as well as the effect of the biopolymer in the rates of growth and production.
- The ODEs represent the term R<sub>i</sub> in the species transport equation.

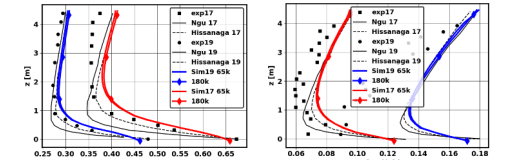


## Influence of bubble size distribution (BSD) and gas composition

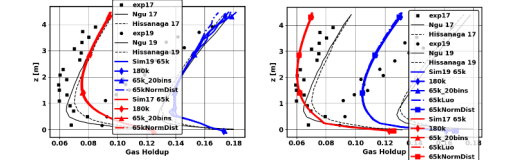


## Validation setup

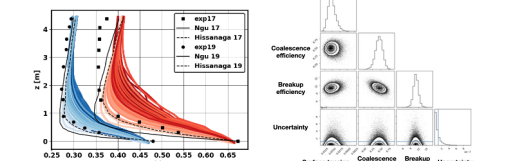
- As first step towards validation, we have compared our CFD modeling against computational/experimental work presented in Ngu et al., Deweer et al., and Hisanaga et al. The work represents dissolution of air/CO<sub>2</sub> mixtures in a bubble column.
- The comparison was carried out assuming **constant bubble size** and modeling the bubble size distribution via **population balances** where we use **breakup** and **binary breakup** models to represent changes in bubble size due to breakup events.
- Additionally, we set up a Bayesian calibration methodology to answer the question: How should one modify hydrodynamics models to capture experiments? This involves hundreds of runs varying different modeling parameters for population balances



Constant diameter (left) and population balances (right) validation



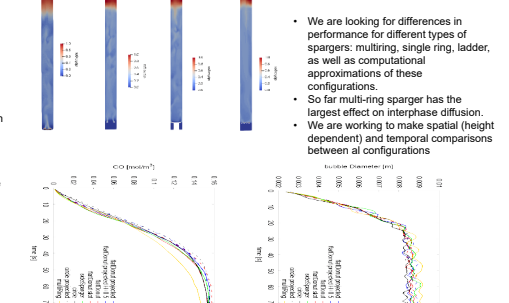
Breakup (left) and binary breakup (right) for population balances



Bayesian calibration for breakup and coalescence efficiencies

## Sparger design

- Sparger design is influential in the control of the bubble size distribution. Additionally, uniformity in gas sparging, affects gas hold-up, interfacial area, mixing time and mass transfer coefficient.
- We have created tools to generate the computational meshes for the spargers:
  - Automated STL generation
  - Automated block-cylindrical meshing
- These tools are publicly available: <https://github.com/NREL/spargerDeSign>



## Bibliography

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- Ngu et al. (2020). Spatio-temporal 1D gas-liquid model for biological methanation in lab scale and industrial bubble column. *Chemical Engineering Science* 251, 117478
- Deweer et al. (1978). A Comprehensive Study on CO-Interphase Mass Transfer in Vertical Co-current and Counter-current Gas-Liquid Flow. *The Canadian Journal of Chemical Engineering*, Vol. 56 43-55
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