

An Open-source Aerosol Dynamics and Computational Fluid Dynamics Model

aerosolGDEFoam User Guide

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Source

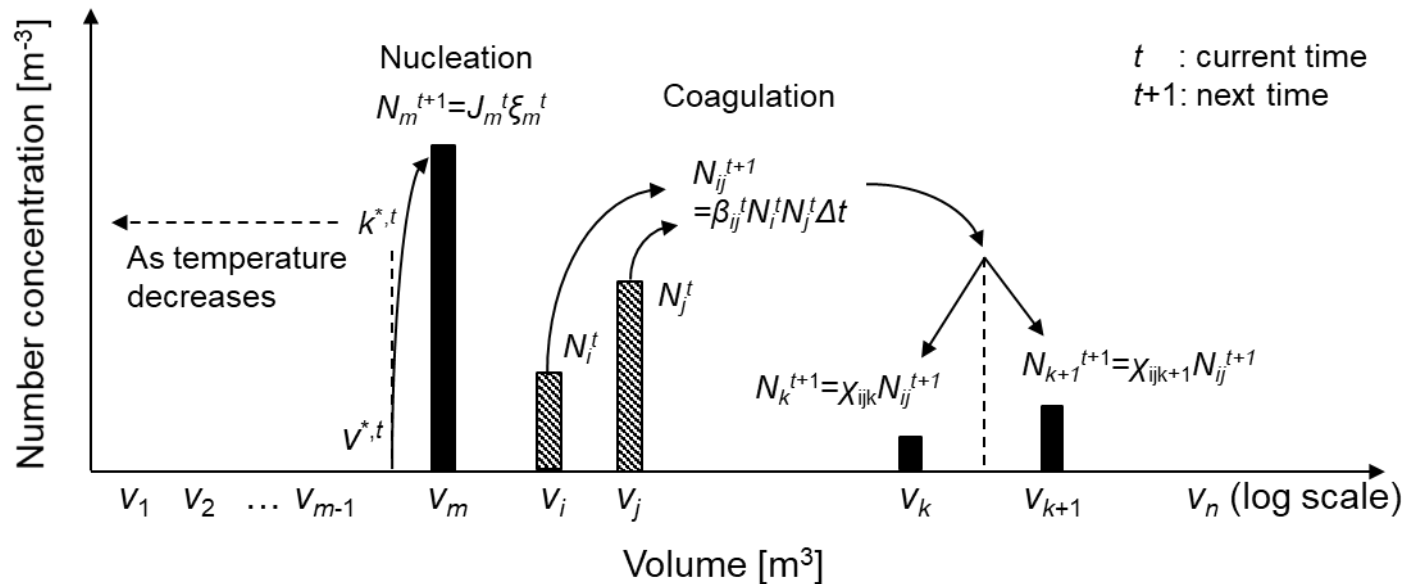
- Woo et al. “Open-source Aerosol Dynamics and Computational Fluid Dynamics 2 Model: Nodal Method for Nucleation, Coagulation, and Surface growth”, [Journal, year, page. doi](#)

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1. Introduction

- Implementation of nodal method for time dependent general dynamic equation including nucleation, surface growth, and coagulation mechanisms



Schematics of nucleation and coagulation algorithms in nodal method

1. Introduction

- Governing equations

Particles

$$\frac{dN_k}{dt} = J_k \xi_k + \underbrace{\frac{1}{2} \sum_{\substack{i=2 \\ j=2}} \chi_{ijk} \beta_{i,j} N_i N_j - N_k \sum_{i=2} \beta_{i,k} N_i}_{\text{coagulation}}$$

$$+ \underbrace{\left\{ \begin{array}{l} \frac{v_1}{v_k - v_{k-1}} \beta_{1,k-1} (N_1 - N_{\text{sat},1,k-1}) N_{k-1} \quad \text{if } N_1 > N_{\text{sat},1,k-1} \\ -\frac{v_1}{v_{k+1} - v_k} \beta_{1,k+1} (N_1 - N_{\text{sat},1,k+1}) N_{k+1} \quad \text{if } N_1 < N_{\text{sat},1,k+1} \\ -\frac{v_1}{v_{k+1} - v_k} \beta_{1,k} (N_1 - N_{\text{sat},1,k}) N_k \quad \text{if } N_1 > N_{\text{sat},1,k} \\ \frac{v_1}{v_k - v_{k-1}} \beta_{1,k} (N_1 - N_{\text{sat},1,k}) N_k \quad \text{if } N_1 < N_{\text{sat},1,k} \end{array} \right.}_{\text{surface growth}}$$

Monomer

$$\frac{dN_1}{dt} = J_1 \xi_1 + \left\{ \begin{array}{l} -\beta_{1,k-1} (N_1 - N_{\text{sat},1,k-1}) N_{k-1} \quad \text{if } N_1 > N_{\text{sat},1,k-1} \\ -\beta_{1,k+1} (N_1 - N_{\text{sat},1,k+1}) N_{k+1} \quad \text{if } N_1 < N_{\text{sat},1,k+1} \\ -\beta_{1,k} (N_1 - N_{\text{sat},1,k}) N_k \quad \text{if } N_1 > N_{\text{sat},1,k} \\ -\beta_{1,k} (N_1 - N_{\text{sat},1,k}) N_k \quad \text{if } N_1 < N_{\text{sat},1,k} \end{array} \right.$$

Nucleation rate

$$J_k = N_{\text{sat}}^2 S v_1 \left(\frac{2\sigma}{\pi m_1} \right)^{0.5} \exp \left(\theta - \frac{4\theta^3}{27 \log^2 S} \right)$$

Size operator

$$\xi_k = \begin{cases} \frac{v^*}{v_k}; & \text{if } v_{k-1} \leq v^* \leq v_k, \\ \frac{v^*}{v_2}; & \text{if } v^* \leq v_1, \\ 0; & \text{otherwise} \end{cases}$$

Collision frequency

- Free molecular regime ($\text{Kn} > 10$)

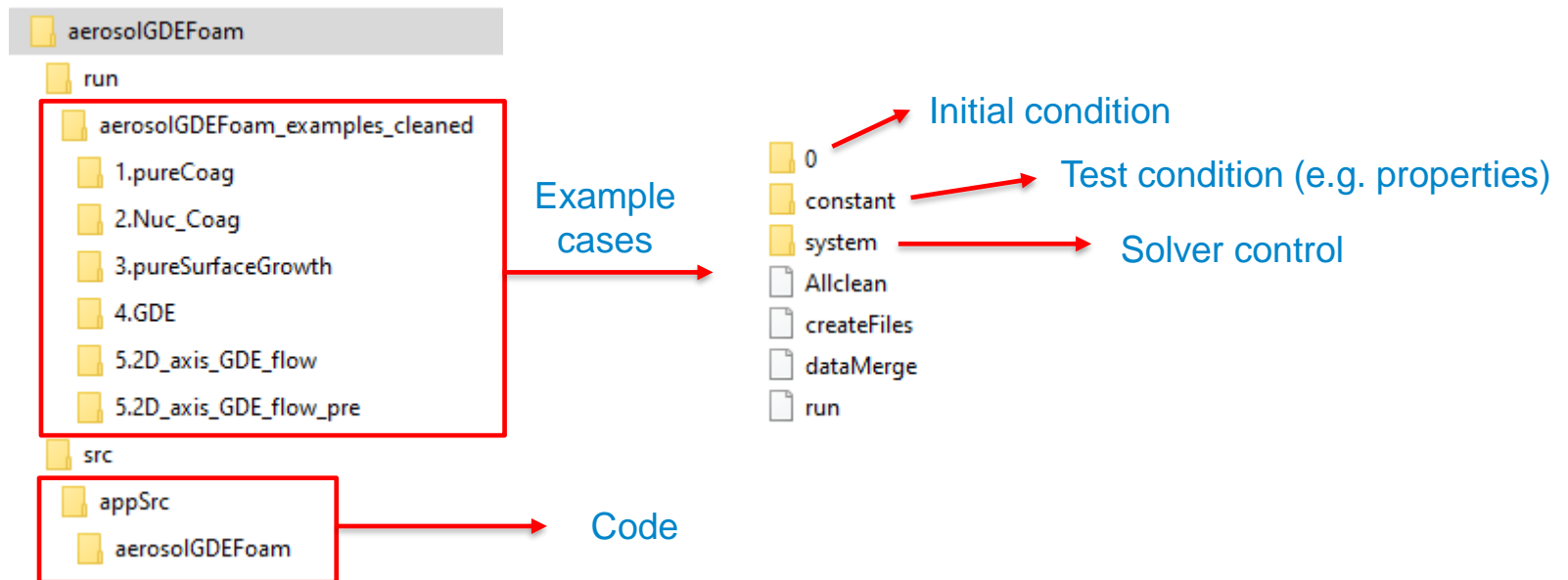
$$\beta_{ij} = \left(\frac{3}{4\pi} \right)^{1/6} \left(\frac{6k_B T}{\rho_p} \right)^{1/2} \left(\frac{1}{v_i} + \frac{1}{v_j} \right)^{1/2} (v_i^{1/3} + v_j^{1/3})^2$$

- Continuum regime ($\text{Kn} \ll 1$)

$$\beta_{ij} = 2\pi(D_i + D_j)(d_i + d_j) \times \left[\frac{d_i + d_j}{d_i + d_j + 2(g_i^2 + g_j^2)^{1/2}} + \frac{8(D_i + D_j)}{(\bar{c}_i^2 + \bar{c}_j^2)^{1/2} (d_i + d_j)} \right]^{-1}$$

2. Code structure

- Developed on OpenFOAM 6 from The OpenFOAM Foundation
 - For download and install, visit <https://openfoam.org/version/6/>
- File structure



2. Code structure

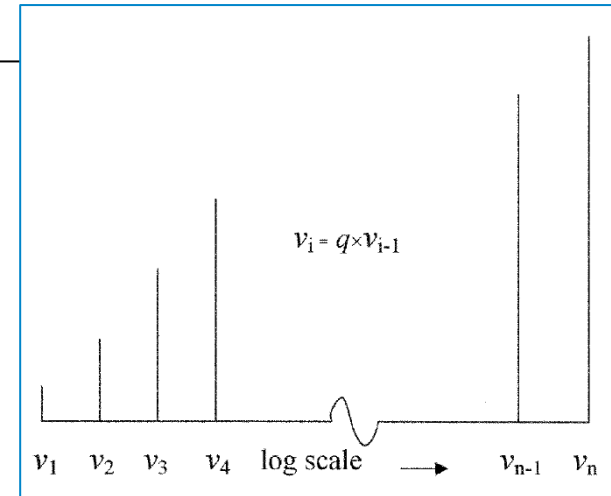
- constant/transportProperties (cont.)

```
nodes nodes [0 0 0 0 0 0 0] 41; // number of nodes
```

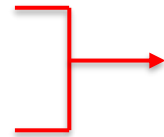
Node spacing according to the Eq. (1) and (2)
in the paper

$$v_k = v_1 q^k \quad \text{and} \quad q = 10^{\frac{12}{k_{\max} - 2}}$$

Figure: Illustration of node spacing on a logarithmic volume space where q is the geometric spacing factor*



```
nucleation true;  
coagulation true;  
growth false;  
advection false;
```



Choose the models of interest

```
// ***** //
```

*Ref.: A simple numerical algorithm and software for solution of nucleation, surface growth, and coagulation problems, Prakash et al., Aerosol Science Technology, 37:892-898, 2003.

2. Code structure

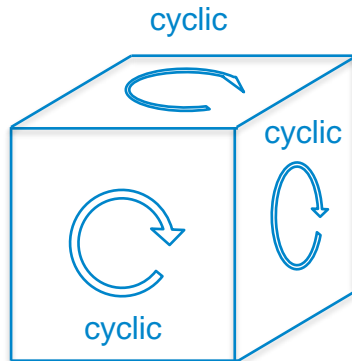
- system/controlDict

```
                                :
startFrom      startTime;//latestTime  Start from initial condition or restart at latest time
startTime      0;
stopAt         endTime;
endTime        1.3745;                Time at which the simulation ends
deltaT         0.0001;                Time step
writeControl    timestep;//adjustableRunTime  Write based on time step or time
writeInterval  2749;//0.2749          Write interval based on time step or time
                                :
```

- For more details on the usage of controlDict in OpenFOAM 6
 - <https://cfd.direct/openfoam/user-guide/v6-controldict/>

3. Verification cases: general dynamic equation (GDE) without flow

- Test problem: time evolution of monomer and particles
 - Zero dimensional problem dependent only on time
 - For pure coagulation (1.pureCoag)
 - Self-preserving distribution from monodisperse particles at the smallest node
 - For the other examples (2.NucCoag, 3.pureSurfaceGrowth, 4.GDE)
 - Aluminum particle formation and growth from a vapor at 1773K as it is passed into a condenser with a cooling rate of 1000K/s*.
- Computational domain
 - Cubic domain with cyclic boundary conditions for each pair of opposite faces



- 1×1×1 unit cell for uniform field (no spatial difference)
- Enable 0D simulation by OpenFOAM based on 3D solver

*Ref.: Panda, S. and S.E. Pratsinis, *Modeling the synthesis of aluminum particles by evaporation-3 condensation in an aerosol flow reactor*. *Nanostructured Materials*, 1995. **5**(7): p. 755-767

3. Verification cases: general steps to setup the test case

- Pre-step 1: Generate mesh
 - Go to run folder (e.g. ./1.pureCoag/40) and ./system/blockMeshDict

```

:
vertices
(
  (0 0 0) //0
  (1 0 0) //1
  (1 1 0) //2
  (0 1 0) //3
  (0 0 1) //4
  (1 0 1) //5
  (1 1 1) //6
  (0 1 1) //7
);
blocks
(
  hex (0 1 2 3 4 5 6 7) block1 (1 1 1)
  simpleGrading (1 1 1)
);
edges
(
);
Boundary
(
  front
  {
    type cyclic;
    neighbourPatch back;
    faces
    (
      (4 5 6 7)
    );
  }
  :
);
```

Domain geometry

Boundary conditions of front
(identical for other boundaries)

Mesh cells and grading

3. Verification cases: general steps to setup the test case

- Pre-step 2: Compile
 - Go to source folder `cd src/appSrc/aerosolGDEFoam`
 - `./wmake`
- Pre-step 3: Data creation before running
 - Go back to run folder and `./createFiles`
 - Create a number of files depending on the number of nodes defined in `constant/tranportProperties`
 - Initial condition should be specified after file creation
- Post-step: data merging for verification cases
 - In run folder `./dataMerge`
 - Take the lines for number concentrations at each node and merge

3.1 Pure coagulation

- Step 1: test condition
 - Go to ./constant/transportProperties
 - Set the number of node, coolrate and models

```
coolrate coolrate [0 0 -1 1 0 0 0] 0;
```

```
mwParticle mwParticle [1 0 0 0 -1 0 0] 0.026981539; // Molecular weight(kg/mol)  
rhoParticle rhoParticle [1 -3 0 0 0 0 0] 2700; // Particle density (kg/m3)  
surTens_A surTens_A [1 0 -2 0 0 0 0] 948; // Surface tension coefficient  
surTens_B surTens_B [1 0 -2 -1 0 0 0] 0.202; // Surface tension coefficient  
satVap_A satVap_A [0 0 0 0 0 0 0] 13.07; // Saturation vapor pressure coefficient  
satVap_B satVap_B [0 0 0 1 0 0 0] 36373; // Saturation vapor pressure coefficient
```

```
nodes nodes [0 0 0 0 0 0 0] 41; // number of nodes
```

```
nucleation false;  
coagulation true;  
growth false;  
advection false;
```

3.1 Pure coagulation

- Step 2: Set initial condition
 - Run ./createFile
 - Go to ./0 and open nPartNode2

```
dimensions      [0 -3 0 0 0 0 0];  
internalField   uniform 1e24;  
boundaryField  
{  
    front  
    {  
        type  
        cyclic;  
    }  
    back  
    {  
        type  
        cyclic;  
    }  
}
```

3.1 Pure coagulation

- Step 3: solver control
 - Go to ./system/controlDict
 - Setup time step and write interval

```
application      scalarTransportFoam;  
  
startFrom        latestTime;  
  
startTime        0;  
  
stopAt           endTime;  
  
endTime          1e-6;  
  
deltaT           1e-11;  
  
writeControl     adjustableRunTime;  
  
writeInterval    1e-7;
```

- Step 4: Restart with increased time step depending on characteristic coagulation time

3.1 Pure coagulation

- Summary of test conditions

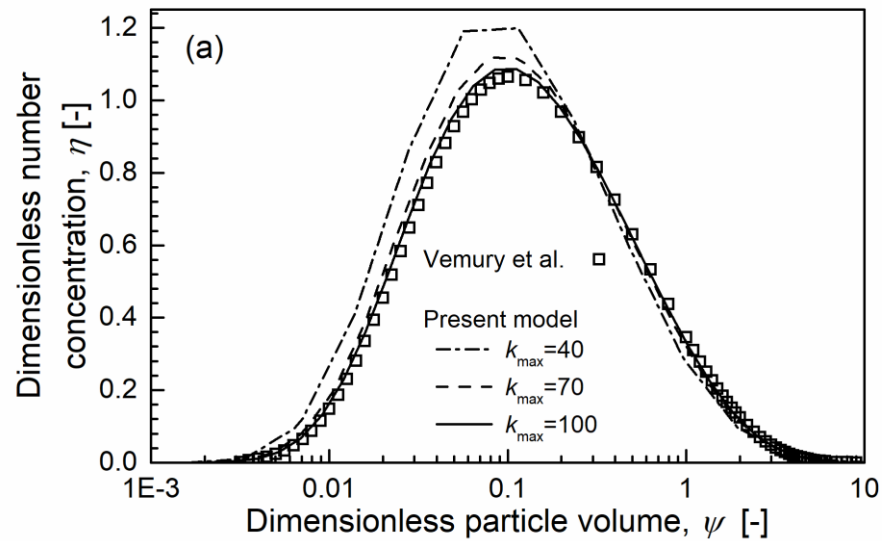
Number of nodes	41, 71 and 101
Initial concentration	nPartNode2: 1e24
Coolrate	0
Time step	Depending on characteristic coagulation time*
Models	nucleation false; coagulation true; growth false; advection false;

*Note:

- Time step is initially given as 1e-11 and increased when restart
- Self-preserving distribution is eventually obtained with time step higher than the criteria but with significant mass defect (valid only for pure coagulation)

3.1 Pure coagulation

- Result



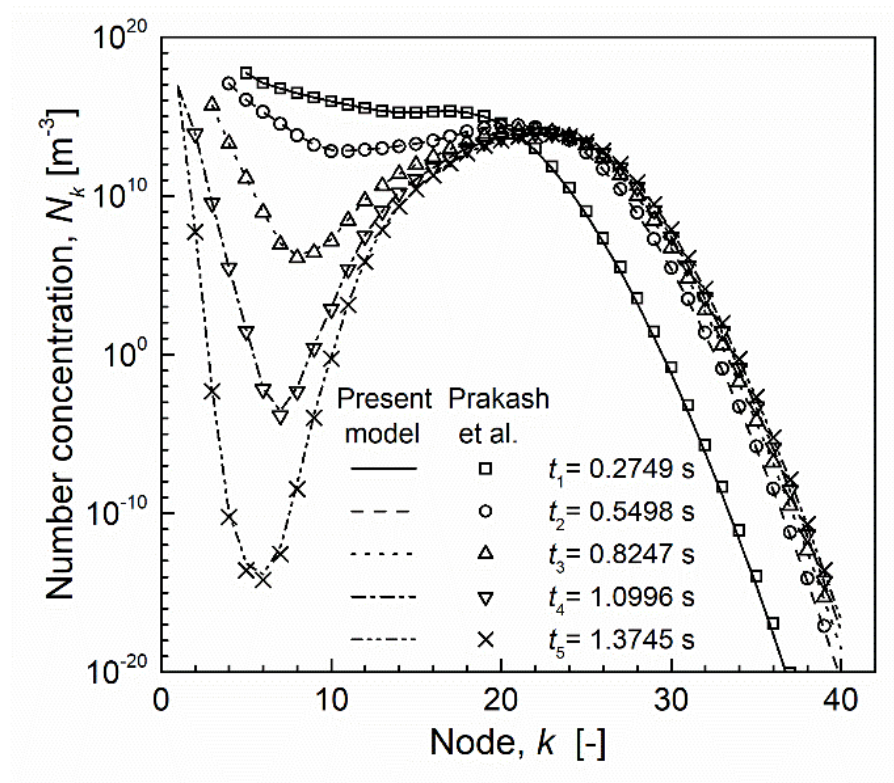
3.2 Nucleation and coagulation

- Test conditions

Number of nodes	41
Initial concentration	nPartNode1: 2.4214e21
Coolrate	1000
Time step	1e-4
Models	nucleation true; coagulation true; growth false; advection false;

3.2 Nucleation and coagulation

- Result



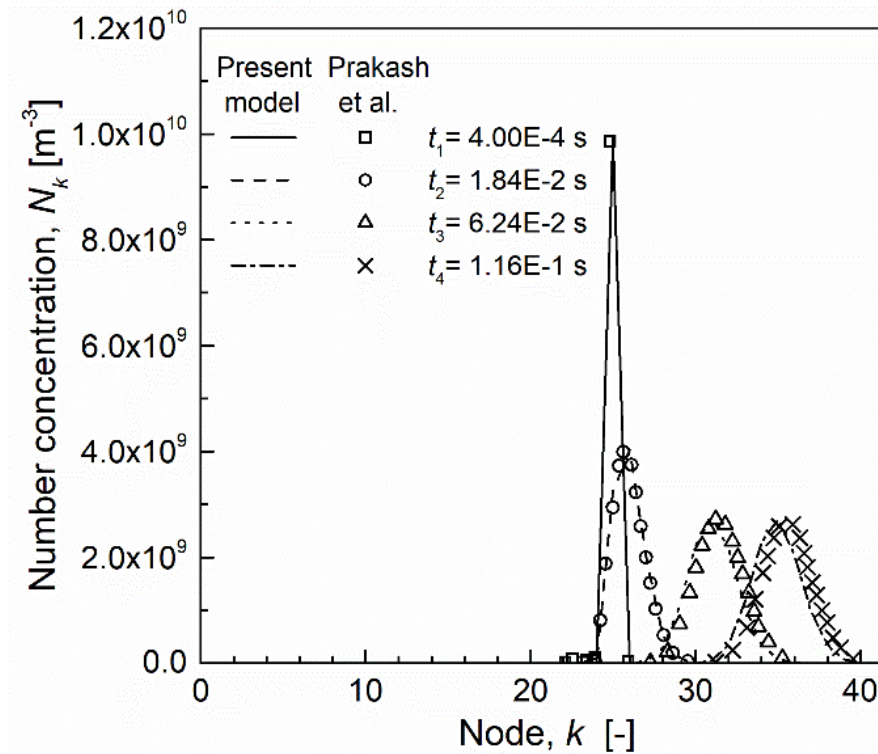
3.3 Pure surface growth

- Test conditions

Number of nodes	41
Initial concentration	nPartNode1: 2.4214e21 nPartNode25: 1e10
Coolrate	1000
Time step	1e-6
Models	nucleation false; coagulation false; growth true; advection false;

3.3 Pure surface growth

- Result



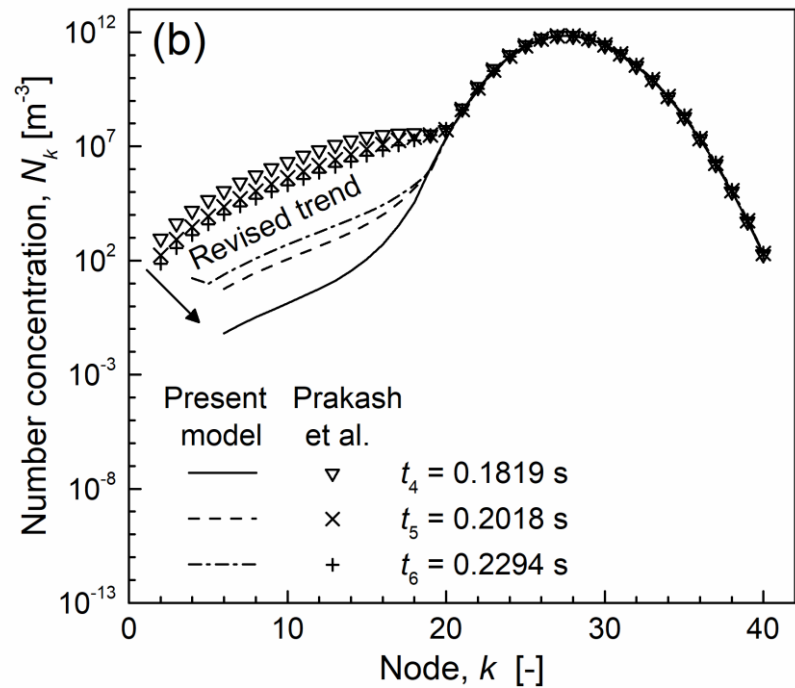
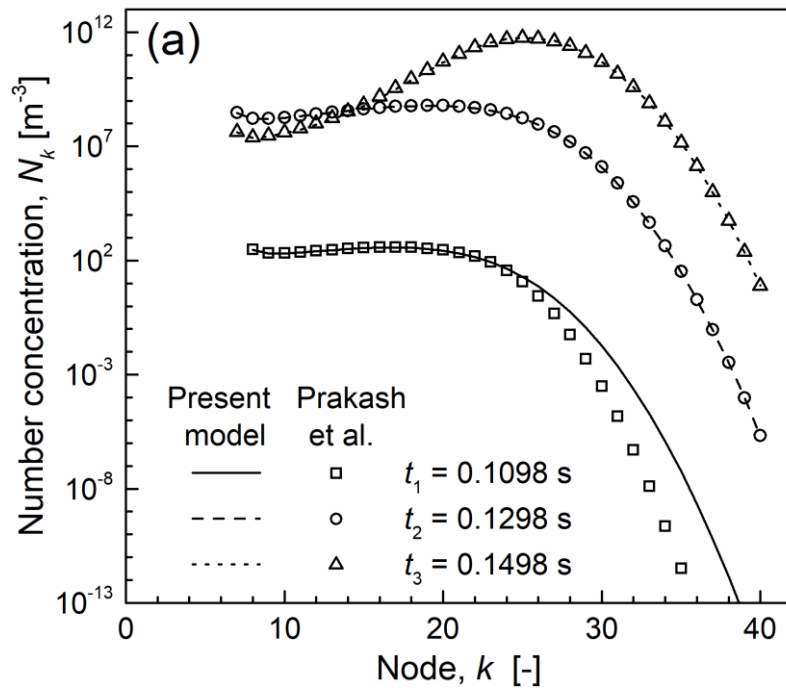
3.4 GDE

- Test conditions

Number of nodes	41
Initial concentration	nPartNode1: 2.4214e21
Coolrate	1000
Time step	1e-5
Models	nucleation true; coagulation true; growth true; advection false;

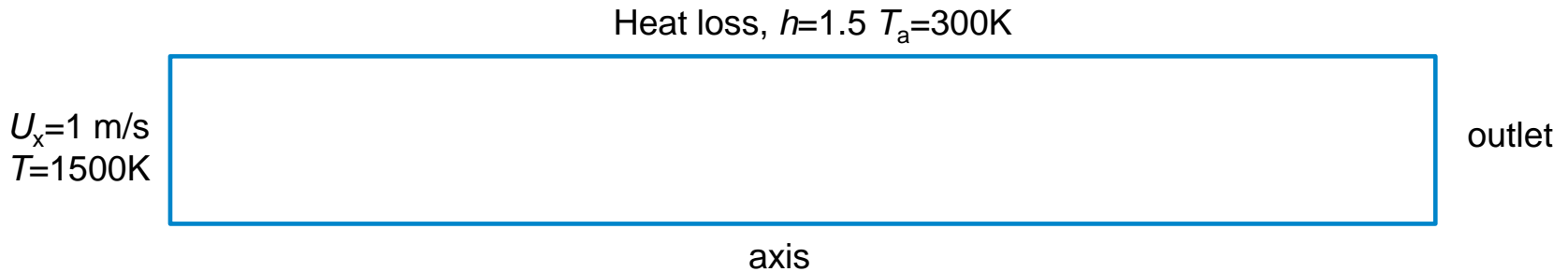
3.4 GDE

- Result



4. Example case with flow

- Pre-calculation
 - 2D axisymmetric computational domain and boundary conditions



- Mesh
 - 200 × 20 × 1 2D axisymmetric mesh

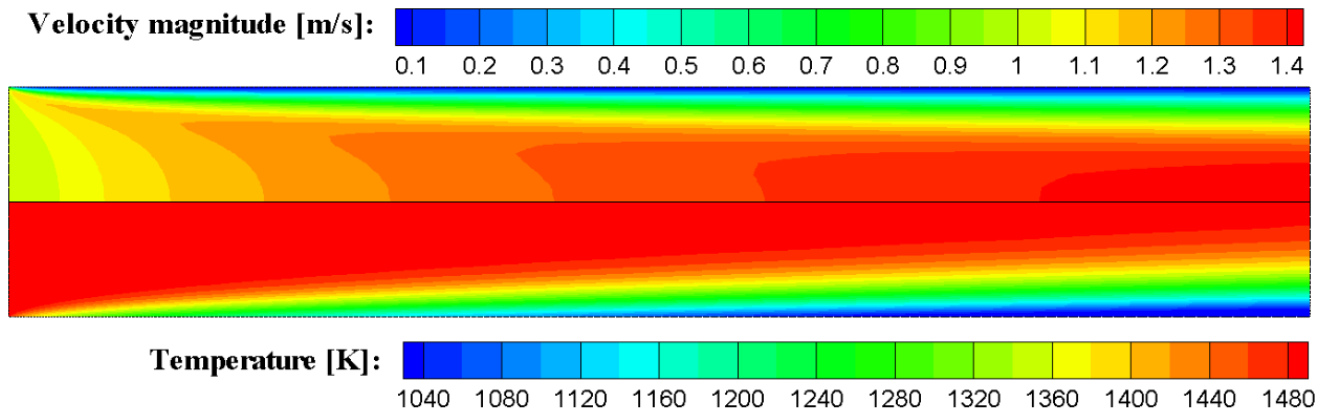
```
blocks(    hex (0 3 4 1 0 3 5 2) block1 (200 20 1)    simpleGrading (5 0.1 1) );
```

- Solver
 - rhoSimpleFoam

4. Example case with flow

- Results

(a)



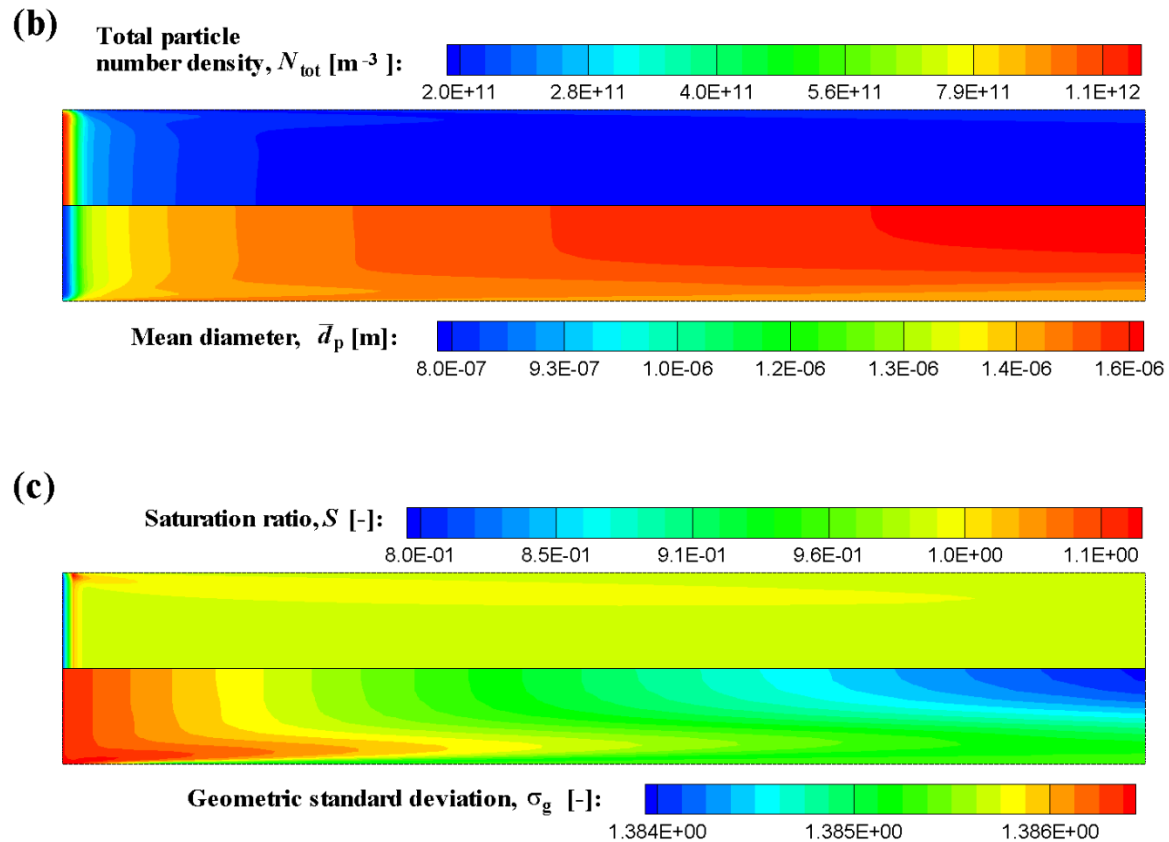
4. Example case with flow

- Copy
 - U,T,p from precalculation
- Test conditions

Number of nodes	41
Initial concentration	_nPart: 1e10 (before createFiles)
Coolrate	0
Time step	1e-5
Models	nucleation true; coagulation true; growth true; advection true;

4. Example case with flow

- Results



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Thank you
