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**Köhn, Christoph; Enghoff, Martin Andreas Bødker; Svensmark, Henrik**

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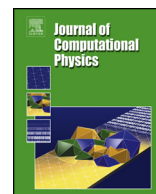
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# A 3D particle Monte Carlo approach to studying nucleation

Christoph Köhn\*, Martin Bødker Enghoff, Henrik Svensmark

Technical University of Denmark, National Space Institute (DTU Space), Elektrovej 328, 2800 Kgs Lyngby, Denmark



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## ABSTRACT

The nucleation of sulphuric acid molecules plays a key role in the formation of aerosols. We here present a three dimensional particle Monte Carlo model to study the growth of sulphuric acid clusters as well as its dependence on the ambient temperature and the initial particle density. We initiate a swarm of sulphuric acid–water clusters with a size of 0.329 nm with densities between  $10^7$  and  $10^8$   $\text{cm}^{-3}$  at temperatures between 200 and 300 K and a relative humidity of 50%. After every time step, we update the position of particles as a function of size-dependent diffusion coefficients. If two particles encounter, we merge them and add their volumes and masses. Inversely, we check after every time step whether a polymer evaporates liberating a molecule. We present the spatial distribution as well as the size distribution calculated from individual clusters. We also calculate the nucleation rate of clusters with a radius of 0.85 nm as a function of time, initial particle density and temperature. The nucleation rates obtained from the presented model agree well with experimentally obtained values and those of a numerical model which serves as a benchmark of our code. In contrast to previous nucleation models, we here present for the first time a code capable of tracing individual particles and thus of capturing the physics related to the discrete nature of particles.

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## 1. Introduction

Nucleation of aerosols is the fundamental process by which gas condenses to form stable clusters. These clusters can potentially grow all the way to sizes where they can serve as cloud condensation nuclei (CCN), typically 50–100 nm. The nucleation phenomenon has been observed all around the globe [23] and is also considered to contribute to the formation of clouds on brown dwarfs and exoplanets [12,25]. About half of all CCN are estimated to originate from nucleated aerosols [29], making nucleation a relevant topic not only for aerosol research but also for its implications for cloud formation. Additionally both aerosols and clouds are relevant for e.g. climate change due to their large forcing effects [3]. The key molecule for aerosol nucleation has long been thought to be sulphuric acid (with water or other stabilizing molecules) due to its ability to form strong bonds [5]. Recently it has been shown that highly oxygenated organic molecules are also able to nucleate at high altitudes [2].

Traditionally nucleation has been described by classical thermodynamic nucleation theory [11], kinetic numerical models [32,27,39], or parametrisations based on either nucleation theory (e.g., [37]) or experimental data (e.g., [7]). The parametrisations and numerical models have the advantage that they can be adapted for use in global modelling [35,41,31] due to being computationally quick. Kazil and Lovejoy [17] used a semi-analytical approach to add aerosols to a global model.

\* Corresponding author.

E-mail address: [koehn@space.dtu.dk](mailto:koehn@space.dtu.dk) (C. Köhn).

More recently an Atmospheric Cluster Dynamics Code (ACDC) has been developed solving the so-called birth–death equations, ordinary differential equations (ODEs) describing the temporal evolution of cluster densities of a given size [28]. The novelty of such a model is the automatic generation of ODEs for a given cluster size and its implementation into the solver whenever needed. This approach aims to reduce typographical errors when implementing ODEs manually. Such traditional numerical models can provide information with regards to particle size distributions and may reflect the physics satisfactorily using actual data for condensation and evaporation. The information provided by these models is, however, focused on developments in time and not in space. If each molecule could be tracked in a three dimensional space it might be possible to achieve new insights into the process of nucleation.

Tracing individual particles in space and time is the main advantage of particle Monte Carlo codes. They are for example widely used to simulate the properties of lightning discharges [4,19,20] by tracing individual electrons and photons or to study the nanostructure growth of atoms on surfaces in electrochemical models [14,15,9]. Recently, molecular dynamic models have also been used to study vapour-to-liquid nucleation [6] or homogeneous water nucleation [1].

Monte Carlo models give the opportunity to include all relevant microphysical processes as well as the interaction amongst all involved particles. In contrast to kinetic models or pure parametrisations which are based on averaged quantities, such as the density or concentration of particles or the mean energy, these models are able to capture rare events initiated by single particles [33,13] as for example the production of gamma rays or positrons in the vicinity of lightning discharges [18].

The disadvantage of Monte Carlo codes is their runtime. Depending on the size of the problem Monte Carlo simulations can take up to several weeks whereas models based on averaged quantities take several hours to days [26]. For small systems, however, the time difference is not significant and Monte Carlo models offer a much better approach to the discrete nature of particles.

We here present a particle Monte Carlo code to study the nucleation of sulphuric acid clusters in 3D, which to our knowledge has not been done before. We here emphasize, however, that we not only present a new model to simulate the nucleation of sulphuric acid clusters, but also that the implemented physics is sufficient enough to study the nucleation of neutral sulphuric acid clusters in order to benchmark our model. Previously Monte Carlo studies have been used on the sulphuric-acid water system to study cluster parameters such as cluster shape, conformation and dissociation [22] as well as cluster free energies [16]. The paper is organized as follows. In section 2 we introduce the Monte Carlo particle model for the study of nucleation processes and discuss all the ingredients of this model: the implementation of single particles, diffusion coefficients, the collision of particles, the evaporation coefficients as well as the choice of the time step. In section 3 we discuss the spatial and size distribution of single sulphuric acid clusters and compare nucleation rates calculated with the present model with values from literature. We finally conclude in section 4.

## 2. Modelling

For the study of the growth of  $\text{H}_2\text{SO}_4\text{-H}_2\text{O}$  clusters, we introduce a particle Monte Carlo code tracing individual  $\text{H}_2\text{SO}_4\text{-H}_2\text{O}$  clusters. In order to benchmark this model, we perform simulations with initial sulphuric acid molecule densities of  $n = 10^7 \text{ cm}^{-3}$  and of  $10^8 \text{ cm}^{-3}$  at 200 K and 300 K as well as of  $n = 2.5 \cdot 10^7 \text{ cm}^{-3}$  at 238 K with a relative humidity of approximately 50%.

In the simulations we do not distinguish between molecules and clusters, we simply refer to them as particles. Each particle is described as a sphere characterized by its position  $\mathbf{r} = (x, y, z)$  in Cartesian coordinates. The particle's radius  $R$  consisting of  $k$  sulphuric acid molecules is determined through [38]

$$R(k) = \sqrt[3]{\frac{3m(k)}{4\pi\rho(k, T)}} \quad (1)$$

implicitly including water content for a relative humidity of approx. 50%. The mass and the density are given through [37]

$$m(k) = \frac{0.098 \frac{\text{kg}}{\text{mol}}}{N_A} \cdot k + \frac{0.018 \frac{\text{kg}}{\text{mol}}}{N_A} \cdot \frac{k - k \cdot \mathcal{MF}(k)}{\mathcal{MF}(k)} \quad (2)$$

$$\rho(k, T) \left[ \frac{\text{kg}}{\text{cm}^3} \right] = \left[ 0.7681724 + 2.184714 \cdot \mathcal{MF}(k) + 7.163002 \cdot \mathcal{MF}(k)^2 \right. \\ - 44.31447 \cdot \mathcal{MF}(k)^3 + 88.75606 \cdot \mathcal{MF}(k)^4 \\ - 75.73729 \cdot \mathcal{MF}(k)^5 + 23.43228 \cdot \mathcal{MF}(k)^6 + T[\text{K}] \cdot (0.001808255 - 0.009294656 \cdot \mathcal{MF}(k) \\ - 0.03742147 \cdot \mathcal{MF}(k)^2 + 0.2565321 \cdot \mathcal{MF}(k)^3 - 0.5362872 \cdot \mathcal{MF}(k)^4 + 0.4857736 \cdot \mathcal{MF}(k)^5 \\ - 0.1629592 \cdot \mathcal{MF}(k)^6) + T[\text{K}]^2 \cdot (-0.000003478524 + 0.00001335867 \cdot \mathcal{MF}(k) \\ + 0.00005195706 \cdot \mathcal{MF}(k)^2 - 0.0003717636 \cdot \mathcal{MF}(k)^3 + 0.0007990811 \cdot \mathcal{MF}(k)^4 \\ \left. - 0.000745806 \cdot \mathcal{MF}(k)^5 + 0.000258139 \cdot \mathcal{MF}(k)^6) \right] \frac{1}{1000} \quad (3)$$

with the Avogadro constant  $N_A \approx 6.022 \cdot 10^{23} \text{ mol}^{-1}$  and the mole fraction

$$\mathcal{MF}(k) = 0.4505 \cdot k^{-0.2097} \quad (4)$$

which is a powerlaw fit to Fig. 2b of [38] for a temperature of approx. 300 K for a humidity of 50%. According to the same figure, we multiply (4) with a factor of  $\approx 0.8$  for temperatures of 200 K and 238 K.

After every time step  $\Delta t$ , the position is updated through

$$\mathbf{r}(t + \Delta t) = \mathbf{r}(t) + \sqrt{2D(R)\Delta t}\mathbf{G} \quad (5)$$

where  $\mathbf{G} = (\varrho \cos \phi \sin \theta, \varrho \sin \phi \sin \theta, \varrho \cos \theta)$  is a Gaussian random number [9] with  $\varrho = \sqrt{-2 \log(r_1)}$ ,  $\phi = 2\pi r_2$ ,  $\theta = \arccos(2r_3 - 1)$ ,  $r_i \in [0, 1)$ .  $D(R) = D_0 \cdot (R(1)/R)^2$  depends on the particle size  $R$  and on the diffusion coefficient [8]

$$D_0 = \frac{2}{3} \sqrt{\frac{k_B^3 T^3}{\pi^3 m(1)} \frac{1}{4PR(1)^2}} \quad (6)$$

for molecules where  $R(1) = 0.329 \text{ nm}$  [21] and  $m(1) = 2.0033 \cdot 10^{-25} \text{ kg}$  are the initial size and the mass of single sulphuric acid molecules without any attached water molecules.  $k_B \approx 1.38 \cdot 10^{-23} \text{ J/K}$  is the Boltzmann constant and  $P = 1 \text{ bar}$  the ambient pressure. For  $T = 300 \text{ K}$ ,  $T = 238 \text{ K}$  and  $T = 200 \text{ K}$  the diffusion coefficients are  $D_0 \approx 1.65 \cdot 10^{-6} \text{ m}^2/\text{s}$ ,  $D_0 \approx 1.16 \cdot 10^{-6} \text{ m}^2/\text{s}$  and  $D_0 \approx 8.96 \cdot 10^{-7} \text{ m}^2/\text{s}$ .

The time step used is related to the time it takes a particle to diffuse the average separation between particles. The average separation between particles is  $\sqrt[3]{1/n}$  and the related diffusion length is  $\sqrt{2D_0\Delta t}$ . Equating the two gives

$$\Delta t \sim \frac{1}{2D_0\sqrt[3]{n^2}}. \quad (7)$$

Hence it is  $\Delta t \sim 0.653 \text{ ms}$  for  $n = 10^7 \text{ cm}^{-3}$  and 300 K,  $\Delta t \sim 0.504 \text{ ms}$  for  $n = 2.5 \cdot 10^7 \text{ cm}^{-3}$  and 238 K,  $\Delta t \sim 1.202 \text{ ms}$  for  $n = 10^7 \text{ cm}^{-3}$  and 200 K as well as  $\Delta t \sim 0.141 \text{ ms}$  for  $n = 10^8 \text{ cm}^{-3}$  and 300 K. Note that the actual time step should be  $\leq \Delta t$  and that  $\Delta t$  limits the maximal distance which a particle travels. Hence, we have chosen  $\Delta t = 100 \mu\text{s}$  for  $n = 10^7 \text{ cm}^{-3}$  and 300 K,  $\Delta t = 250 \mu\text{s}$  for  $n = 2.5 \cdot 10^7 \text{ cm}^{-3}$  and 238 K and  $\Delta t = 1 \text{ ms}$  for  $n = 10^7 \text{ cm}^{-3}$  and 200 K. However, since the diffusion term in (5) depends on a Gaussian random number and since for polymers it is  $D(R) < D_0$ , the actual travelled distance is smaller than the maximally possible value. With  $\Delta t = 0.1 \text{ ms}$  or  $\Delta t = 1 \text{ ms}$ , it is  $\sqrt{2D_0\Delta t} \approx 18 \mu\text{m}$  for 300 K and  $\sqrt{2D_0\Delta t} \approx 42 \mu\text{m}$  for 200 K. Since the mean free path of sulphuric acid molecules in air is approximately 10–60 nm [34], hence significantly smaller than 1  $\mu\text{m}$ , the justification of the diffusion approach in this study is justified.

After every time step we check whether two particles  $i$  and  $j$  with sizes  $R_i$  and  $R_j$  overlap by evaluating the condition

$$|\mathbf{r}_i - \mathbf{r}_j| \leq R_i + R_j \quad (8)$$

where  $\mathbf{r}_i$  are  $\mathbf{r}_j$  are the particles' positions. If this condition is fulfilled, the two particles are merged with new mass and radius according to (1) and (2); the new position is determined through

$$\mathbf{r}_{i+j} = \frac{m_i \mathbf{r}_i + m_j \mathbf{r}_j}{m_i + m_j}. \quad (9)$$

Note that we add a new  $\text{H}_2\text{SO}_4\text{-H}_2\text{O}$  cluster to the simulation domain at a random position after merging two particles if the particle number becomes smaller than the initial particle number; thus, the density of particles does not drop below the initial particle density.

Vice versa we also check after every time step whether a cluster with radius  $R$  and mass  $m$  evaporates by emitting one sulphuric acid molecule added to the simulation domain leaving a cluster with reduced mass and volume behind. For that purpose, we check after every time step and for every cluster whether

$$r \leq P_{eva} \quad (10)$$

for a random number  $r \in [0, 1)$  and for the evaporation probability  $P_{eva} = 1 - \exp(-\gamma \Delta t)$ . Here the evaporation frequency  $\gamma$  for 300 K is adopted from [38]

$$\gamma = \sqrt{\frac{8\pi k_B T (m(1) + m(k))}{m(1)m(k)}} (R(1) + R(k))^2 n_{a,sol}^\infty(k, T) \exp\left(\frac{2M_1\sigma(k, T)}{\varrho(k, T) \cdot \mathcal{R} \cdot TR(k, T)}\right) \quad (11)$$

where the concentration  $n_{a,sol}^\infty$  of  $\text{H}_2\text{SO}_4$  vapour molecules in the equilibrium vapour above a flat source and the surface tension are determined through [30,37]

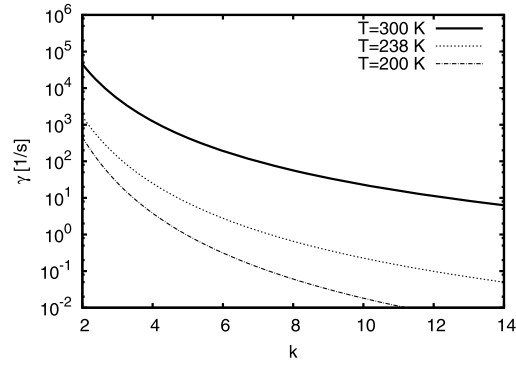


Fig. 1. The evaporation coefficient  $\gamma$  (11) for temperatures of 200 K, 238 K and 300 K as a function of the number  $k$  of sulphuric acid atoms in the cluster.

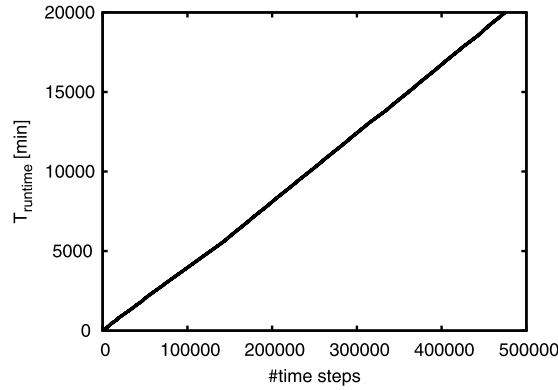


Fig. 2. The runtime of simulations as a function of time steps.

$$n_{a,sol}^{\infty}(k, T)[\text{m}^{-3}] = 0.0017241 \cdot \mathcal{MF}(k)^{10.891} \cdot \frac{133}{T[\text{K}] \cdot \mathcal{R}} \cdot N_A \quad (12)$$

$$\begin{aligned} \sigma[\text{N m}^{-1}] &= 0.11864 - 0.11651 \cdot \mathcal{MF}(k) + 0.76852 \cdot \mathcal{MF}(k)^2 - 2.40909 \cdot \mathcal{MF}(k)^3 + 2.95434 \cdot \mathcal{MF}(k)^4 \\ &\quad - 1.25852 \cdot \mathcal{MF}(k)^5 + T[\text{K}] \cdot (-0.00015709 + 0.00040105 \cdot \mathcal{MF}(k) - 0.0023995 \cdot \mathcal{MF}(k)^2 \\ &\quad + 0.007611235 \cdot \mathcal{MF}(k)^3 - 0.00937386 \cdot \mathcal{MF}(k)^4 + 0.00389722 \cdot \mathcal{MF}(k)^5) \end{aligned} \quad (13)$$

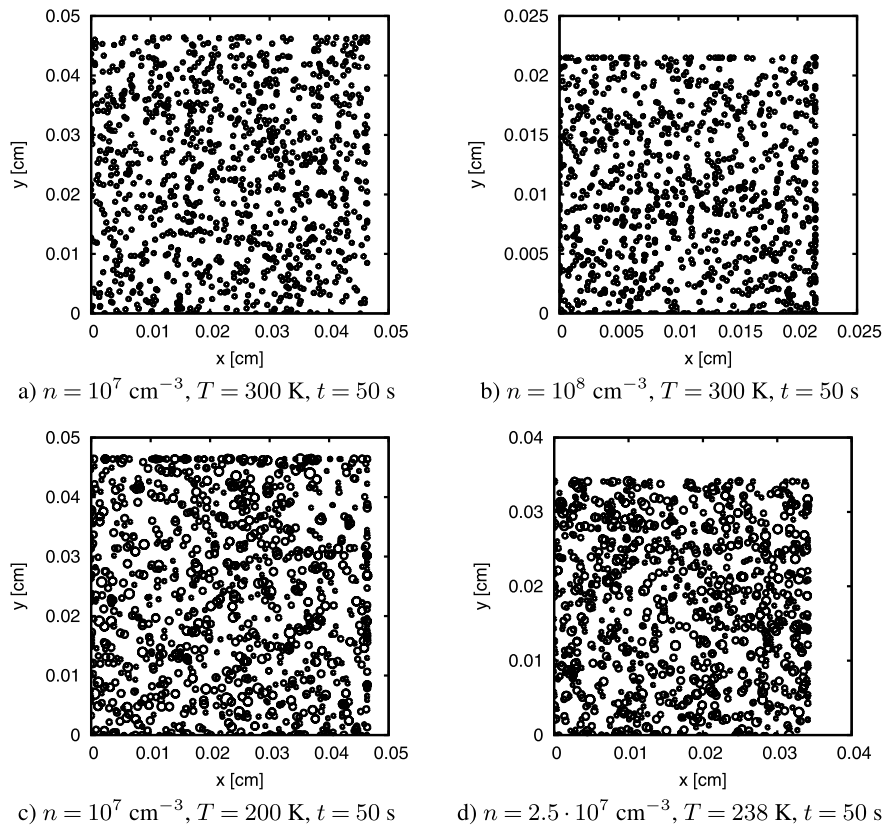
$M_1 = 98 \text{ g mol}^{-1}$  is the molar mass of  $\text{H}_2\text{SO}_4$  [24, pp. 1661–1665] and  $\mathcal{R} \approx 8.31 \text{ J (mol K)}^{-1}$  the universal gas constant. Because of the lack of physical data, e.g. for the equilibrium vapour pressure, we multiply Eq. (11) with a factor of  $\approx 10^{-4}$  for 238 K and of  $\approx 10^{-5}$  for 200 K based on Fig. 4 of [38]. For better accuracy, we use Eq. (11) for  $k > 5$  only; for  $k \leq 5$ , we use distinct values extracted from [38]. Fig. 1 shows the evaporation coefficient  $\gamma$  as a function of  $k$  for 200 K, 238 K and 300 K. For all considered cases evaporation is most efficient for small cluster sizes and becomes less probable with increasing size. For 200 K and 238 K  $\gamma$  is several orders of magnitude smaller than for 300 K, thus compared to 300 K evaporation is much less likely and nearly negligible for 200 K. Since the emitted molecule is added to the simulation domain, the particle density slightly increases; however, we have observed that it does not exceed  $1.002 \cdot 10^7 \text{ cm}^{-3}$  for an initial density of  $10^7 \text{ cm}^{-3}$  and  $1.009 \cdot 10^8 \text{ cm}^{-3}$  for an initial density of  $10^8 \text{ cm}^{-3}$ .

In order to optimize the runtime of our simulations, we have chosen the volume of the simulation domain to be  $10^{-4} \text{ cm}^3$  for a density of  $10^7 \text{ cm}^{-3}$ ,  $4 \cdot 10^{-5} \text{ cm}^3$  for a density of  $2.5 \cdot 10^7 \text{ cm}^{-3}$ , and  $10^{-5} \text{ cm}^3$  for a density of  $10^8 \text{ cm}^{-3}$ . Consequently we initiate all simulations by placing 1000 individual  $\text{H}_2\text{SO}_4$  molecules at random positions into the simulation domain which scales with the concentration.

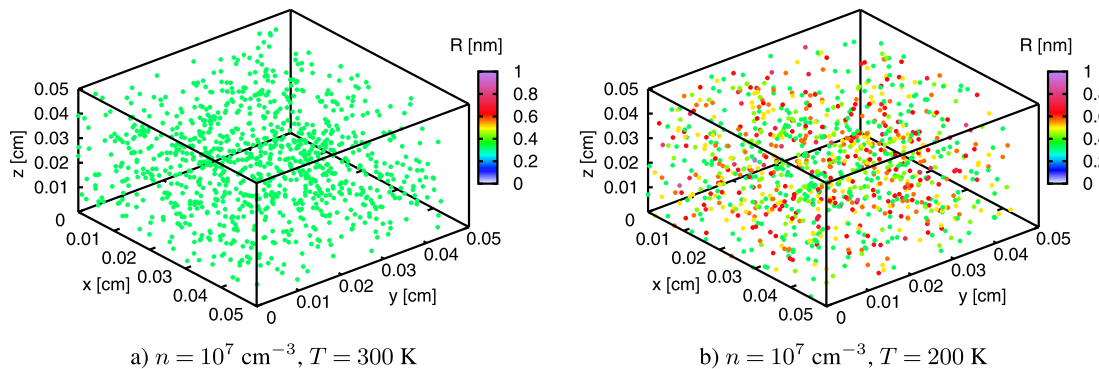
### 3. Results

In the following we discuss the spatial distribution and the size distribution of all particles as well as the nucleation rates of clusters with a radius of 0.85 nm. As supplementary material we have added movies showing the temporal evolution of the particle position and size for  $n = 10^7 \text{ cm}^{-3}$  and 200 K as well as for  $n = 10^7 \text{ cm}^{-3}$  and 300 K.

Fig. 2 shows the runtime of all simulations as a function of time steps. The runtime increases linearly because of the approximate constancy of the number of simulated particles. In all considered cases it takes approximately  $10^4$  minutes



**Fig. 3.** The spatial distribution of all particles after 50 s for different densities and different temperatures projected onto the  $xy$  plane. For better visibility the size of each circle is the real size of each particle multiplied by a factor of  $10^6$ .

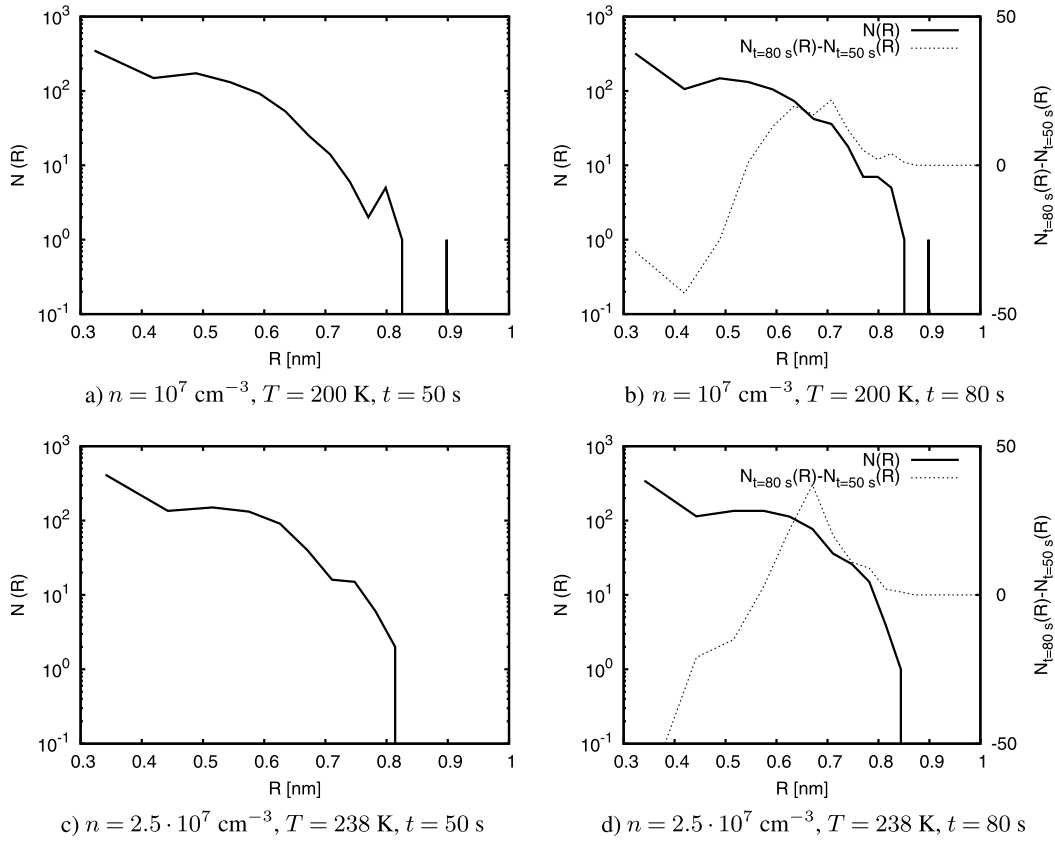


**Fig. 4.** The position of all particles in the three dimensional simulation domain after 50 s for the same densities and temperatures as in Fig. 3 a) and d). The particle size is colour coded. (For interpretation of the colours in the figure(s), the reader is referred to the web version of this article.)

or equivalently 7 days per 250000 time steps which can easily be translated into physical time knowing the size of the computational time step.

### 3.1. Spatial distribution

Fig. 3 shows the spatial distribution of all particles projected onto the  $xy$  plane after 50 s. For better visibility we have multiplied the real size of each particle by a factor of  $10^6$ . Fig. 4 illustrates the capability of the present model for full three dimensional simulations. It shows the spatial distribution after the same time steps and for the same densities and temperatures as in Fig. 3 a) and c). Fig. 3 and 4 demonstrate that in all cases the particles are distributed randomly within the simulation domain. Panels a) and b) of Fig. 3 show that for  $T = 300$  K, there is no growth at all; the average radius of all particles after 50 s is approximately 0.329 nm. In contrast to a temperature of 300 K where the simulation is evaporation



**Fig. 5.** The size distribution of all particles after 50 s (left column) and after 80 s (right column) for the same conditions as in Fig. 3 c) and d). The right y-axis in the right column shows the difference between the particle numbers after 80 s and after 50 s.

dominated, there is a significant growth of clusters for a temperature of 200 K and 238 K where evaporation is less probable to negligible. The average size is 0.48 nm for 200 K and 0.46 nm for 238 K.

### 3.2. Size distribution

Fig. 5 shows the size distributions after 50 s and after 80 s for 200 K and 238 K. The right y-axis in the right column displays the difference between the particle numbers after 80 s and after 50 s.

For all considered cases, the number of monomers with a size of 0.329 nm is dominant after 50 s whilst there are only a few clusters with larger sizes. For  $T = 300$  K, there are no particles larger than 0.329 nm; for  $T = 238$  K (c) and 200 K (a), approximately half of all particles are larger than 0.329 nm. For 200 K and 238 K, the large number of polymers is an effect of the improbable or negligible evaporation.

After 80 s, the size distributions at 300 K have not changed at all. Panel a, however, shows that for a temperature of 200 K, the number of monomers has decreased enormously since there is only nucleation, but no evaporation. The right y-axis of panels b) and d) show that there is an increase of polymers with sizes above 0.8 nm; this effect is more significant for 200 K than for 238 K since evaporation is more likely for the latter case.

### 3.3. Nucleation rate

For 300 K, the simulation is evaporation dominated. Hence we do not observe any polymers and conclude that the nucleation rate is 0.0. In the case of 238 K and 200 K, there is an interplay between the nucleation through the approach of particles and the evaporation. As Fig. 3 and 5 show, there is a significant growth of the particle sizes. After every time step  $t$  we therefore count the number  $N(R, t)$  of particles with size  $R$ . Subsequently the nucleation rate of particles with size  $R$  as a function of time  $t$  is given through

$$\nu(R, t) = \frac{N(R, t)}{V \cdot t} \quad (14)$$

where  $V$  is the volume of the simulation domain.

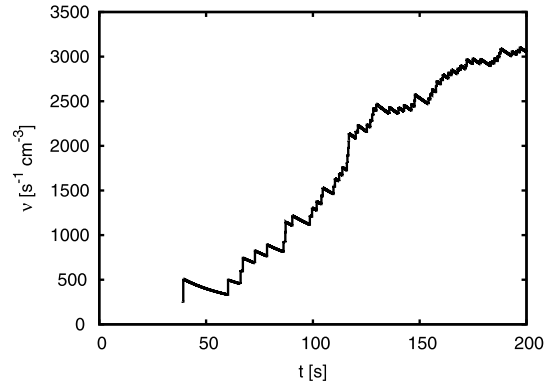


Fig. 6. The nucleation rate  $\nu(t)$  as a function of time for  $n_0 = 10^7 \text{ cm}^{-3}$  for 200 K.

Table 1

The nucleation rates  $\nu_{sim}$  calculated from our simulations, the nucleation rates  $\nu_{par}$  [7] and the nucleation rates by [43].

	$\nu_{sim} [\text{cm}^{-3} \text{s}^{-1}]$	$\nu_{par} [\text{cm}^{-3} \text{s}^{-1}]$	$\nu_{yu} [\text{cm}^{-3} \text{s}^{-1}]$
$n = 10^7 \text{ cm}^{-3}, T = 300 \text{ K}$	0.0	$10^{-8.19}$	$10^{-46.92}$
$n = 10^7 \text{ cm}^{-3}, T = 200 \text{ K}$	$\approx 10^{3.42 \pm 1.77}$	$10^{2.45}$	$10^{3.11}$
$n = 10^8 \text{ cm}^{-3}, T = 300 \text{ K}$	0.0	$10^{-4.57}$	$10^{-35.40}$
$n = 2.5 \cdot 10^7 \text{ cm}^{-3}, T = 238 \text{ K}$	$10^{2.05 \pm 2.38}$	$10^{-0.16}$	$10^{0.04}$

As an example, Fig. 6 shows the nucleation rate  $\nu(R = 0.85 \text{ nm}, t)$  of particles with a radius of 0.85 nm as a function of time for  $n = 10^7 \text{ cm}^{-3}$  and  $T = 200 \text{ K}$ . For comparison with literature, e.g. with [7], we have chosen a radius of 0.85 nm; however, this critical size is rather arbitrary. One of the advantages of the introduced model here is the possibility to determine the nucleation rate of particles of any size. For 200 K, evaporation is negligible and thus particles keep growing and consequently the nucleation rate increases as a function of time. In the beginning, the nucleation rate grows slowly since it takes time for the first clusters to reach the nucleation size. However, the slope of the nucleation rate flattens because of the decreased diffusion coefficient for larger clusters which diffuse more slowly than small clusters and single molecules and consequently collide less with surrounding particles.

We here define the mean nucleation rate of particles with a radius of 0.85 nm and above within time interval  $\mathcal{T}$  as

$$\nu_{sim} = \langle \nu \rangle = \frac{1}{\mathcal{T}} \sum_t \nu(R \geq 0.85 \text{ nm}, t). \quad (15)$$

Dunne et al. [7] simulated the formation of atmospheric aerosol particles in extensive laboratory experiments. They determined the nucleation rates for aerosols at different temperatures and for different compounds and derived the parametrization

$$\nu_{par}(n[10^6/\text{cm}^3], T[\text{K}]) = n[10^6/\text{cm}^3]^{3.62} \cdot e^{46.3 - \frac{245.07|K|}{1000.0}} \quad (16)$$

for the neutral nucleation rate of sulphuric acid–water clusters with a radius of 0.85 nm as a function of the density  $n$  in units of  $10^6$  molecules/ $\text{cm}^3$  and of the temperature  $T$  in Kelvin. Additionally, Yu [43] set up a website which calculates nucleation rates  $\nu_{yu}$  based on [40] and [42].

Table 1 compares the nucleation rates  $\nu_{sim}$  calculated from our simulations with the nucleation rates  $\nu_{par}$   $\nu_{yu}$ . In all cases  $\nu_{sim}$  is comparable to the values obtained by Dunne et al. [7] or [43], hence there is a good agreement within the error bars. This might be due to underestimating the evaporation coefficient. However, from the available data, it is difficult to get the exact values for the evaporation coefficients. However, we here emphasize that this paper is not about finding the best input data, but to introduce a code that – provided sufficiently accurate input data – models nucleation on a particle level.

#### 4. Conclusions

We have presented a particle Monte Carlo model tracing individual  $\text{H}_2\text{SO}_4$ – $\text{H}_2\text{O}$  clusters for different initial densities,  $n$ , of sulphuric acid molecules and for different temperatures  $T$  taking the growth by collision of particles and evaporation by single  $\text{H}_2\text{SO}_4$  molecules into account.

Four cases were considered,  $n = 10^7 \text{ cm}^{-3}$  with temperature either  $T = 300 \text{ K}$  or  $T = 200 \text{ K}$ , one case with  $n = 2.5 \cdot 10^7 \text{ cm}^{-3}$  and  $T = 238 \text{ K}$  and one case with  $n = 10^8 \text{ cm}^{-3}$  and  $T = 300 \text{ K}$ . In the two cases at 300 K, there is no significant



growth between 50 s and 80 s, in accordance with a low nucleation rate. However, for 200 K and 238 K, growth of clusters between 50 s and 80 s is observed.

The simulations make it possible to calculate the size distribution based on individual particle sizes. For 300 K, all particles are single molecules. This behaviour is rather independent of the initial density. In contrast at 200 K and 238 K, evaporation becomes improbable or negligible and thus the size distribution consists of fewer monomers compared to the case with  $T = 300$  K which increases the number of clusters above 0.329 nm.

Finally we have calculated the nucleation rates as a function of time and the mean nucleation rates averaged over time. Since for 300 K there is an interplay between nucleation and evaporation, the nucleation rate is 0. For lower temperatures, evaporation is less likely and as such the nucleation rate increases with time. However, it tends to an upper limit since the probability of two particles colliding with each other is reduced as an effect of the decreased diffusion coefficient.

We compared the nucleation rates with values experimentally obtained by Dunne et al. [7] and F. Yu [43]. Within the given error bars we see a good agreement between our simulation results and experimental values which serves as a benchmark for our Monte Carlo code. We therefore conclude that the physics implemented in the present Monte Carlo model is appropriate to simulate the growth of sulphuric acid clusters. Its main advantage is that it traces individual particles and therefore reflects their distinct nature and as such reality much better than for example kinetic numerical models or pure parametrisations.

Subsequently, the 3D particle approach allows us to study new phenomena. In future work, we intend to investigate how the temporal evolution of the nucleation rate changes for fixed particle densities and different volumes, thus different particle numbers. Such a task is ideal for a model tracing individual particles and cannot be studied with kinetic numerical models or pure parametrisations.

The model presented in this paper can easily be adjusted to study nucleation phenomena of other species in different temperatures and pressures. Lee et al. [25] studied the formation of  $\text{TiO}_2$  and  $\text{SiO}$  clusters on brown dwarfs and extra-solar planets. They use a classical nucleation theory where they allow only monomers to attach to other monomers or clusters. In future work, we will adjust our model to include cluster–cluster collisions and to study their effects on the nucleation of clusters in the atmospheres of exoplanets.

In a forthcoming paper, we will finally present a more sophisticated model where we include  $\text{HSO}_4^-$  ions and investigate their influence on the nucleation rate. Not only is this of relevance for aerosol formation [10] and growth [36] in Earth's atmosphere, but this model can be adjusted to study the influence of ions in exoplanets' atmospheres which has not been considered before.

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## Appendix A. Supplementary material

Supplementary material related to this article can be found online at <https://doi.org/10.1016/j.jcp.2018.02.032>.

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