

A modelling study of the impact of bacteria on cloud development

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1. Introduction

The INSU-DFG bilateral BIOCLOUDS project aimed at investigating and quantifying the role of bioaerosols in tropospheric clouds, focussing in the studies on microorganisms, mainly bacteria. To reach our objectives, we:

- 1. isolated and identified ice nuclei active (INA) bacterial strains in cloud water sampled on the Puy de Dôme mountain station,
- 2. studied in more details ice nuclei (IN) and cloud condensation nuclei (CCN) properties of bacteria isolated from cloud water in the laboratory and in cloud chambers

3. used the new data as input to cloud models.

Two posters present the work, one on the experimental part and this one dedicated to the modelling studies.

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1E-008

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0.1

Dia Fig. 1a : Detailed description of the different

Fig. 1b: Schematic model configuration

hydrometeor class in DESCAM (Planche et

100 1000

2. The DESCAM Model

DESCAM (Detailed SCAvenging Model, Flossmann et al, 1985) is a detailed microphysics model with the specificity of following explicitly the evolution of the mass of the aerosol population (Fig. 1a). This allows to study separately background and bioaerosol.

In this study DESCAM is coupled with a 1D cylindrical dynamical model with lateral exchange (Asai and Kasahara, 1967, Fig. 1b)

The dynamics were initiated from the CCOPE experiment (Dye et al, 1986), a strong storm well studied since the 1990s, regularly used as reference case for convective model studies.

3. Parameterization

The first measurement campaigns in the AIDA Chamber (IMK-AAF, FZK) showed good that bacteria act as cloud condensation nuclei, partly because of their size. Thus, they initiate ice acting as condensation freezing nuclei.

The ice nucleation activity parameterized as following: $N_{bact,ice} = J_{bact} * N_{bact,tot}$

To consider INA bacteria in the model, we used the IN properties measured for the most efficient bacterial strain (Andraud) and those measured by the ICCF (Joly et al, 2012), Fig. 2a, and we first assumed that they represent all of the bioaerosol population present.

We then used in another study results from Fig. 2b : Concentration of INA bioaerosol at the ICCF (Joly et al, 2013) to consider a realistic amount of INA bioaerosol (Fig.2b).





4. Impact of INA bacteria

A recent study (Hiron et al., 2013) demonstrated that condensation freezing in general plays a very important role in the development of the cloud as it forms ice at warm temperatures, having therefore a major dynamical feedback, even though it forms only small amounts of ice. The impact of the INA bacteria is due

to their activity at these warm temperatures as can be seen in Fig. 3b

Time (min) Time (min ud at 0.01 cm⁻¹) 012 at 0.01 cm⁻¹ Time (min)

Fig. 3b : Liquid and ice water content in the ing. so : Liquid and ice water content in the atmospheric column as a function of time with (top) and without (middle) bacteria. Bottom graph gives the resulting rain flux on the graund ulting rain flux on the ground.

5. Impact of INA bacteria in realistic concentrations

- То improve the estimates on the impact of INA bacteria we also used the results from Jolv et al. (2013) where the concentration of INA bacteria was directly measured as a function of temperature (Fig.2b).
- The impact on the then simulated cloud is quite small which is in agreement with the concentration estimations sensitivity study (over one order of magnitude below the optimum: ≈ 0.001 cm⁻³).



6. Conclusions and perspectives

INA bacteria can potentially have a large impact on the development of a storm if present in high concentrations although it seems that often their actual concentration is too low.

Further development of this study will need to consider a 3D dynamical framework and stratiform clouds where the time scales are longer.

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