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# Locally refined discrete velocity grids for stationary rarefied flow simulations

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Abstract: Most of deterministic solvers for rarefied gas dynamics use discrete velocity (or discrete ordinate) approximations of the distribution function on a Cartesian grid. This grid must be sufficiently large and fine to describe the distribution functions at every space position in the computational domain. For 3-dimensional hypersonic flows, like in re-entry problems, this induces much too dense velocity grids that cannot be practically used, for memory storage requirements. In this article, we present an approach to generate automatically a locally refined velocity grid adapted to a given simulation. This grid contains much less points than a standard Cartesian grid and allows us to make realistic 3-dimensional simulations at a reduced cost, with a comparable accuracy.

Keywords: rarefied flow simulation, hypersonic flows, re-entry problem, transition regime, BGK model, discrete velocity model

## 1 Introduction

The description of a flow surrounding a flying object at hypersonic speed in the rarefied atmosphere (more than 60 km altitude) is still a challenge in the atmospheric Re-Entry community [2]. When this flow is in a rarefied state, that is to say when the Knudsen number (which is the ratio  $Kn = \frac{\lambda}{L}$  between the mean free path  $\lambda$  of particle and a characteristic macroscopic length  $L$ ) is larger than 0.01, the flow cannot be accurately described by the compressible Navier-Stokes equations of Gas Dynamics. In this case, the kinetic theory has to be used. The evolution of the molecules of the gas is then described by a mass density distribution in phase space, which is a solution of the Boltzmann equation. In the transitional regime, this equation can be replaced by the simpler Bhatnagar-Gross-Krook (BGK) model.

In order to be able to compute parietal heat flux and aerodynamic coefficients in the range of 60-120 km, a kinetic description of the stationary flow is necessary.

The most popular numerical method to simulate rarefied flows is the Direct Simulation Monte Carlo method (DSMC) [9]. However, it is well known that this method is very

expensive in transitional regimes, in particular for flows in the range of altitude we are interested in. The efficiency of DSMC can be improved by using coupling strategies (see [12, 14]) or implicit schemes (see [24, 15]), but these methods are still not very well suited for stationary computations. In contrast, deterministic methods (based on a numerical discretization of the stationary kinetic model) can be more efficient in transitional regimes. Up to our knowledge, there are few deterministic simulation codes specifically designed for steady flows. One of the most advanced ones is the 3D code of Titarev [25] developed for unstructured meshes. Another 3D code has been developed by G. Brook [10]. Other codes exist, but they are rather designed for unsteady problems, see for instance [20, 1] or the recent UGKS scheme developed by K. Xu and his collaborators [27, 18], which is an Asymptotic Preserving scheme for unsteady flows.

In our team, we developed several years ago a code to make 2D plane and axisymmetric simulations of rarefied flows based on the BGK model (see a description in [22, 23, 4]). This code has recently been extended to 3D computations, for polyatomic gases. Due to the physical model (polyatomic gases), the space discretization (block structured mesh), and the parallelization (space domain decomposition with MPI and inner parallelization with openMP), this code is rather different from the other existing 3D codes recently presented in the literature for the same kind of problems (the 3D code of Titarev [25] for example), even if space domain decompositions have already been used for unsteady simulation (see [20]).

All the codes designed for steady flows have a common feature: they are based on a “discrete ordinate” like approach, and use a global velocity grid. This grid is generally a Cartesian grid with a constant step size. The number of points of this grid is roughly proportional to the Mach number of the flow in each direction, and hence can be prohibitively large for hypersonic flows, even with parallel computers. To compute realistic cases (3D configurations with Mach number larger than 20), the velocity space discretization has to be modified in order to decrease CPU time and memory storage requirements. It has already been noticed that a refinement of the grid around small velocities can improve the accuracy and reduce the cost of the computation (especially for large Knudsen numbers in flows close to solid boundaries, see [25]). However, up to our knowledge, there is no general strategy in the literature that helps us to reduce the number of discrete velocities of a velocity grid for any rarefied steady flow, even if some works on adaptive velocity grids have already been presented: the first attempt seems to be [5] for a 1D shock wave problem, and recently, more general adaptive grid techniques designed for unsteady flows have been presented in [21, 13, 19, 11].

The main contribution of this article is to propose an algorithm for an automatic construction of a locally refined velocity grid that allows a dramatic reduction of the number of discrete velocities, with the same accuracy as a standard Cartesian grid. This algorithm uses a compressible Navier-Stokes pre-simulation to obtain a rough estimation of the macroscopic fields of the flow. These fields are used to refine the grid wherever it is necessary by using some indicators of the width of the distribution functions in all the computational domain. This strategy allows us to simulate hypersonic flows that can hardly be simulated by standard methods, since we are indeed able to apply our method to our kinetic code to simulate

a re-entry flow at Mach 25 and for temperature and pressure conditions of an altitude of 90 km. In this example, the CPU time and memory storage can be decreased up to a factor 24, as compared to a method with a standard Cartesian velocity grid. Note that preliminary results have already been presented in [6] and [7].

The outline of this article is the following. In section 2, we briefly present the kinetic description of a rarefied gas. In section 3, we discuss the problems induced by the use of a global velocity grid, and our algorithm is presented. Our approach is illustrated in section 4 with several numerical tests. To simplify the reading of the paper, the presentation of our simulation code is made in the appendix.

## 2 Boltzmann equation and Cartesian velocity grid

### 2.1 Kinetic description of rarefied gases

In kinetic theory, a monoatomic gas is described by the distribution function  $f(t, \mathbf{x}, \mathbf{v})$  defined such that  $f(t, \mathbf{x}, \mathbf{v})d\mathbf{x}d\mathbf{v}$  is the mass of molecules that at time  $t$  are located in an elementary space volume  $d\mathbf{x}$  centered in  $\mathbf{x} = (x, y, z)$  and have a velocity in an elementary volume  $d\mathbf{v}$  centered in  $\mathbf{v} = (v_x, v_y, v_z)$ .

Consequently, the macroscopic quantities as mass density  $\rho$ , momentum  $\rho\mathbf{u}$  and total energy  $E$  are defined as the five first moments of  $f$  with respect to the velocity variable, namely:

$$(\rho(t, \mathbf{x}), \rho\mathbf{u}(t, \mathbf{x}), E(t, \mathbf{x})) = \int_{\mathbb{R}^3} (1, \mathbf{v}, \frac{1}{2}|\mathbf{v}|^2)f(t, \mathbf{x}, \mathbf{v}) d\mathbf{v}. \quad (1)$$

The temperature  $T$  of the gas is defined by the relation  $E = \frac{1}{2}\rho|\mathbf{u}|^2 + \frac{3}{2}\rho RT$ , where  $R$  is the gas constant defined as the ratio between the Boltzmann constant and the molecular mass of the gas.

When the gas is in a thermodynamical equilibrium state, it is well known that the distribution function  $f$  is a Gaussian function  $M[\rho, \mathbf{u}, T]$  of  $\mathbf{v}$ , called Maxwellian distribution, that depends only on the macroscopic quantities :

$$M[\rho, \mathbf{u}, T] = \frac{\rho}{(2\pi RT)^{\frac{3}{2}}} \exp\left(-\frac{|\mathbf{v} - \mathbf{u}|^2}{2RT}\right). \quad (2)$$

It can easily be checked that  $M$  satisfies relations (1).

If the gas is not in a thermodynamical equilibrium state, its evolution is described by the following kinetic equation

$$\partial_t f + \mathbf{v} \cdot \nabla_{\mathbf{x}} f = Q(f), \quad (3)$$

which means that the total variation of  $f$  (described by the left-hand side) is due to the collisions between molecules (described by the right-hand side). The most realistic collision model is the Boltzmann operator but it is still very computationally expensive to use. In this paper, we use the simpler BGK model [8, 26]

$$Q(f) = \frac{1}{\tau}(M[\rho, \mathbf{u}, T] - f) \quad (4)$$

which models the effect of the collisions as a relaxation of  $f$  towards the local equilibrium corresponding to the macroscopic quantities defined by (1). The relaxation time is defined as  $\tau = \frac{\mu}{\rho RT}$ , where  $\mu$  is the viscosity of the gas. This definition allows to match the correct viscosity in the Navier-Stokes equations obtained by the Chapman-Enskog expansion. This viscosity  $\mu$  is usually supposed to fit the law  $\mu = \mu_{ref}(\frac{T}{T_{ref}})^\omega$ , where  $\mu_{ref}$  and  $T_{ref}$  are reference viscosity and temperature determined experimentally for each gas, as well as the exponent  $\omega$  (see a table in [9]).

The interactions of the gas with solid boundaries are described with the diffuse reflection model. Let us suppose that the boundary has a velocity  $\mathbf{u}_w = 0$  and temperature  $T_w$ . In the diffuse reflection model, a molecule that collides with this boundary is supposed to be re-emitted with a temperature equal to  $T_w$ , and with a random velocity normally distributed around 0. This reads

$$f(t, \mathbf{x}, \mathbf{v}) = M[\sigma_w, 0, T_w](\mathbf{v}) \quad (5)$$

if  $\mathbf{v} \cdot \mathbf{n}(\mathbf{x}) > 0$ , where  $\mathbf{n}(\mathbf{x})$  is the normal to the wall at point  $\mathbf{x}$  directed into the gas. The parameter  $\sigma_w$  is defined so that there is no normal mass flux across the boundary (all the molecules are re-emitted). Namely, that is,

$$\sigma_w = - \left( \int_{\mathbf{v} \cdot \mathbf{n}(\mathbf{x}) < 0} f(t, \mathbf{x}, \mathbf{v}) \mathbf{v} \cdot \mathbf{n}(\mathbf{x}) d\mathbf{v} \right) / \left( \int_{\mathbf{v} \cdot \mathbf{n}(\mathbf{x}) > 0} M[1, 0, T_w](\mathbf{v}) \mathbf{v} \cdot \mathbf{n}(\mathbf{x}) d\mathbf{v} \right). \quad (6)$$

There are other reflection models, like the Maxwell model with partial accommodation, but they are not used in this work.

## 2.2 Velocity discretization on Cartesian grid

In most of existing computational codes for the Boltzmann equation that use a deterministic method, the velocity variable is discretized with a *global* Cartesian grid, that is to say the same grid for every point of the space mesh. Consequently, it is necessary to compute a priori a velocity grid in which every distribution function that may appear in the computation can be resolved (see figure 1). This means that the grid must be:

- large enough to contain the main part of the distribution functions *for every position in the computational domain*
- fine enough to capture the core of the distribution functions *for every position in the computational domain*

The corresponding parameters (bounds and step of the grid) can be determined with the following idea: at each point  $\mathbf{x}$  of the computational domain, the macroscopic velocity  $\mathbf{u}(\mathbf{x})$  and temperature  $T(\mathbf{x})$  give information on the support of the distribution function  $f(\mathbf{x}, \cdot)$ . Indeed, if  $f(\mathbf{x}, \cdot)$  is “not too far” from its corresponding Maxwellian, as supposed in the BGK model, its support is centered around  $\mathbf{u}(\mathbf{x})$ , and is essentially localized between bounds that depend on  $\mathbf{u}(\mathbf{x})$  and  $T(\mathbf{x})$ , corresponding to the standard deviation of the Maxwellian ; that

is to say  $f(\mathbf{x}, \cdot)$  is very small outside (see figure 2 where these bounds are  $\mathbf{u}(\mathbf{x}) - c\sqrt{RT(\mathbf{x})}$  and  $\mathbf{u}(\mathbf{x}) + c\sqrt{RT(\mathbf{x})}$  in 1D). When several distribution functions have to be discretized on the same grid, their supports are reasonably well approximated if the bounds are

$$v_{max}^\alpha = \max_{\mathbf{x} \in \Omega} \{u^\alpha(\mathbf{x}) + c\sqrt{RT(\mathbf{x})}\} \quad v_{min}^\alpha = \min_{\mathbf{x} \in \Omega} \{u^\alpha(\mathbf{x}) - c\sqrt{RT(\mathbf{x})}\}, \quad (7)$$

and if the grid step is

$$\Delta v = a \min_{\mathbf{x} \in \Omega} \sqrt{RT(\mathbf{x})}, \quad (8)$$

where  $\Omega$  is the computational domain (in space), and  $\alpha = x, y, z$  (see 3). In (7)-(8), the parameters  $c$  and  $a$  can be chosen as follows. For  $c$ , statistical argument suggest that values between 3 and 4 are needed, and so  $c = 3$  seems to be a good compromise between accuracy and computational cost. The parameter  $a$  allows us to adjust the grid step: it must be at most equal to 1 (which ensures that there are at least six points inside the core of every distribution function), but a smaller value might be necessary for an accurate computation.

To compute these bounds, it is necessary to a priori estimate the values of  $\mathbf{u}$  and  $T$  in the computational domain. A simple way is to use a compressible Navier-Stokes (CNS) pre-simulation (for which the CPU time is negligible as compared to the Boltzmann simulation). This simulation gives the fields  $\mathbf{u}^{CNS}$  and  $T^{CNS}$  in  $\Omega$ , and the bounds and the grid step can be defined by formulas (7)-(8) applied to these fields, that is to say

$$v_{max}^\alpha = \max_{\mathbf{x} \in \Omega} \{u^{\alpha,CNS}(\mathbf{x}) + c\sqrt{RT^{CNS}(\mathbf{x})}\} \quad v_{min}^\alpha = \min_{\mathbf{x} \in \Omega} \{u^{\alpha,CNS}(\mathbf{x}) - c\sqrt{RT^{CNS}(\mathbf{x})}\}, \quad (9)$$

and

$$\Delta v = a \min_{\mathbf{x} \in \Omega} \sqrt{RT^{CNS}(\mathbf{x})}. \quad (10)$$

However, the values of the bounds are mainly determined by the temperatures in the shock zone, which can reach thousands of Kelvin. One must be careful that, at high altitude and high speeds, numerical computations of Navier-Stokes equations can underestimate those temperatures, which would lead to inappropriate bounds.

It is quite clear that the size of the velocity grid increases with the Mach number. Indeed, a large Mach number implies large upstream velocities and large temperature in the shock wave, which lead to very large bounds (see (7)), while the temperature of the body remains small, thus leading to small grid step (see (8)). For realistic re-entry problems, this can lead to prohibitively large grids. For instance, for the flow around a cylinder of radius 0.1 m at Mach 20 and altitude 90 km, formulas (9) and (10) used with a Compressible Navier-Stokes pre-simulation lead to a velocity grid of  $52 \times 41 \times 41$  points and around 350 GB memory requirements with a coarse 3D mesh in space.

However, this is mainly due to the use of a Cartesian grid with a constant step. In order to decrease the size of the grid, it is attractive to refine it only wherever it is necessary, and to coarsen it elsewhere. In the following section, we show that this can be done automatically with our new algorithm.

This approach may not be well designed for cases where  $f$  is very far from it corresponding Maxwellian (like in shock interactions problem). However, for reentry problems, our assumption is realistic for transitional regime we are interested in.

### 3 An algorithm to define locally refined discrete velocity grids

Since we have to represent many distribution functions on the same grid, it is natural to refine the grid in the cores of these distributions, and to coarsen it in the tails (see figure 4). To achieve this goal, we define the concept of “support function”, and we use it to design an AMR (Adaptive Mesh Refinement) velocity grid.

#### 3.1 The support function

At each point  $\mathbf{x}$  of the computational space domain  $\Omega$ ,  $\mathbf{u}(\mathbf{x})$  and  $T(\mathbf{x})$  give information on the “main” support of the distribution function  $f(\mathbf{x}, \cdot)$  in the velocity space: this support is centered at  $\mathbf{u}(\mathbf{x})$  and of size  $2c\sqrt{RT(\mathbf{x})}$  (see (7)). We define the *support function*  $\phi$  in the velocity space as follows: for every  $\mathbf{v}$  in the velocity space, we set  $\phi(\mathbf{v}) = \sqrt{RT(\mathbf{x})}$  if there exists in  $\Omega$  a  $\mathbf{x}$  such that  $\|\mathbf{v} - \mathbf{u}(\mathbf{x})\| \leq c\sqrt{RT(\mathbf{x})}$ , that is to say that  $\mathbf{v}$  is inside the support of a distribution, considered as a sphere of center  $\mathbf{u}(\mathbf{x})$  and of radius  $c\sqrt{RT(\mathbf{x})}$ . This function gives an incomplete mapping of the velocity domain (see figure 5). Now we want to extend this function to the whole velocity space, so that it can be used as a refinement criterion to design an optimal velocity grid.

Indeed, so far, this function is not defined for every  $\mathbf{v}$ , especially for large ones, as we cannot find for every  $\mathbf{v}$  an appropriate pair  $(\mathbf{u}(\mathbf{x}), T(\mathbf{x}))$  to match the definition of the support function. Moreover, this function might be multivalued, as there might be two different space points  $\mathbf{x}_1$  and  $\mathbf{x}_2$  with same velocity  $\mathbf{u}(\mathbf{x}_1) = \mathbf{u}(\mathbf{x}_2)$ , but with different temperatures  $T(\mathbf{x}_1) \neq T(\mathbf{x}_2)$ . Since our goal is to obtain for every  $\mathbf{v}$  the minimum size of the support of every distribution centered around  $\mathbf{v}$ , these two problems can be avoided as follows :

- (a) we set  $\phi(\mathbf{v}) = \min(\sqrt{RT(\mathbf{x}_1)}, \sqrt{RT(\mathbf{x}_2)})$  if  $\|\mathbf{v} - \mathbf{u}(\mathbf{x}_1)\| \leq c\sqrt{RT(\mathbf{x}_1)}$  and  $\|\mathbf{v} - \mathbf{u}(\mathbf{x}_2)\| \leq c\sqrt{RT(\mathbf{x}_2)}$  with  $(\mathbf{u}(\mathbf{x}_1), T(\mathbf{x}_1)) \neq (\mathbf{u}(\mathbf{x}_2), T(\mathbf{x}_2))$ . Indeed, the size of the grid around  $\mathbf{v}$  is constrained by the distribution centered on  $\mathbf{v}$  that has the smallest support, hence  $\phi$  must have the corresponding smallest possible value.
- (b) if there is no  $\mathbf{x}$  such that  $\|\mathbf{v} - \mathbf{u}(\mathbf{x})\| \leq c\sqrt{RT(\mathbf{x})}$ , then  $\mathbf{v}$  is in the tail of every distribution function, and there is no reason to refine the grid there, so we set  $\phi(\mathbf{v})$  to its largest possible value  $\phi(\mathbf{v}) = \max \sqrt{RT(\mathbf{x})}$ .

These ideas lead to the following algorithm for an automatic construction of  $\phi$ . Note that below, the computational domain in space is supposed to be discretized with a mesh composed of cells numbered with three indices  $(i, j, k)$ . This is not a restriction of our algorithm.

*Algorithm 3.1* (Construction of the support function).

1. CNS velocity and temperature are stored in arrays  $\mathbf{u}(i, j, k)$  and  $T(i, j, k)$  for  $i, j, k = 1 : i_{max}, j_{max}, k_{max}$ .
2. construction of a (fine) Cartesian velocity grid (based on the previous global criterion (9)-(10)): points  $\mathbf{v}(q)$  with  $q = 0 : q_{max}$ .
3. computation of the field  $\sqrt{RT}$ , stored in an descending order in the 1D array  $\psi(I)$ , for  $I = 1 : i_{max} \times j_{max} \times k_{max}$ , so that  $\psi(I) = \sqrt{RT((i, j, k)(I))}$ .
4. initialization of the array  $\phi(0 : q_{max}) = \max(\psi)$  on the velocity grid (one value per discrete velocity)
5. *do*  $I = 1 : i_{max} \times j_{max} \times k_{max}$  (loop on the cells of the space mesh)
  - do*  $q = 0 : q_{max}$  (loop on the nodes of the velocity grid)
    - if  $\mathbf{v}(q)$  is in the sphere of center  $\mathbf{u}((i, j, k)(I))$  and radius  $c\psi(I)$ , then it is in the support of a distribution function, and we set  $\phi(q) := \psi(I)$

This algorithm ensures that the array  $\phi$  satisfies the following property.

**Property 3.1.** *For every  $q$  between 0 and  $q_{max}$ , we have :*

$$\phi(q) = \min \left( \min_{i,j,k} \left\{ \sqrt{RT(i, j, k)} \text{ such that } \|\mathbf{v}(q) - \mathbf{u}(i, j, k)\| \leq c\sqrt{RT(i, j, k)} \right\}, \max_{i,j,k} \sqrt{RT(i, j, k)} \right)$$

This means that if we take a velocity  $\mathbf{v}(q)$  of the fine grid, then among all the distribution functions whose support contains  $\mathbf{v}(q)$ , one of them has a smallest support, and  $2c\phi(q)$  is the size of its support. This support function hence tells us how the fine Cartesian grid should be refined, or coarsened, around  $\mathbf{v}(q)$ .

The generation of the corresponding adapted grid is described in the following section.

**Remark 3.1.** It may happen that with such a strategy, the support function does not take sufficiently into account the support of the wall Maxwellian (for diffuse reflection conditions on a solid obstacle): this Maxwellian is centered around  $\mathbf{u} = 0$  and has a temperature  $T_{wall}$  which is often the smallest temperature of the computational domain. This can be the case when the macroscopic CNS flow is not resolved enough close to the wall, or if the CNS equations are solved with slip boundary conditions. In this case, it is safer to add the element  $\phi(q_{max} + 1) := \sqrt{RT_{wall}}$  in  $\phi$  and to add the corresponding velocity  $\mathbf{v}(q_{max} + 1) := 0$  in the fine grid.

## 3.2 AMR grid generation

The idea is to start with a unique cell defined by the bounds of the fine velocity grid (that is the full velocity domain), and then to apply a recursive algorithm: each cell is cut if its dimensions are larger than the minimum of  $a\phi$  in the cell. The algorithm is the following:

*Algorithm 3.2* (Recursive refinement of a cell  $\mathcal{C}$ ).

1. compute the minimum of the field  $a\phi$  in the cell  $\mathcal{C}$ , that is to say the minimum of the elements of  $a\phi$  that have the same indices as the discrete velocities of the fine grid included in this cell  $\mathcal{C}$ :

$$m := a \min\{\phi(q), \text{ such that } \mathbf{v}(q) \in \text{cell } \mathcal{C}\}$$

2. if one edge of  $\mathcal{C}$  is larger than  $m$ , then the cell is cut into 8 new subcells on which the refinement algorithm is applied, recursively.
3. else, the cell is kept as it is, and the cell and its vertices are numbered.

At the end of the algorithm, the final grid satisfies the following property: every cell has a size smaller than  $a$  times the minimum of the support function in the cell.

## 3.3 An example

We anticipate on the numerical results that will be discussed in section 4 to illustrate the previous algorithms.

The test case is a 2D steady flow around a cylinder at Mach 20 (see the geometry in figure 6 and the physical parameters in section 4.2). A CNS pre-simulation and the use of formula (9)-(10) with parameters  $c = 4$  and  $a = 2$  give us a fine velocity grid with  $52 \times 41$  points, see figure 7. Algorithms 3.1 and 3.2 applied to the CNS fields give the support function and the AMR velocity grid shown in figure 8. The AMR velocity grid has 529 points, which is one fourth as small as the original fine Cartesian grid.

Note that the AMR grid is refined around the zero velocity and the upstream velocity: these velocities correspond to the flow close to the solid boundary and to the upstream boundary, where the temperature is low, thus where the distribution functions are narrow, and hence where, indeed, we need a fine velocity resolution. At the contrary, the grid is coarse close to its boundaries: the discrete velocities are very large there and are all in the tails of all the distribution functions, where, indeed, we do not need a fine resolution.

The accuracy of the computation on this AMR grid is studied in section 4.

## 3.4 Discrete Velocity Approximation on the AMR velocity grid

When one wants to transform a standard discrete velocity method based on a Cartesian grid to a method using our AMR grid, two points have to be treated carefully: the computation

of the moments of  $f$ , and the approximation of the collision operator. In this paper, we only use the BGK collision model of the Boltzmann equation: therefore its approximation (based on the conservative approach of [23]) reduces to the problem of a correct approximation of the moments of  $f$ , as it is described below.

First note that the standard discrete velocity approach consists in replacing the kinetic equation (3) by the following discrete kinetic equation

$$\partial_t f_q + \mathbf{v}_q \cdot \nabla_{\mathbf{x}} f_q = Q_q(f), \quad (11)$$

where  $f_q(t, x)$  is supposed to be an approximation of  $f(t, x, \mathbf{v}_q)$  for every discrete velocity  $\mathbf{v}_q$ , with  $q = 0$  to  $q_{max}$ . The discrete distribution function is  $f = (f_q(t, x))$ , and  $Q_q$  is the discrete collision operator. The moments of the discrete distribution function are obtained by using a quadrature formula to replace (1) by

$$(\rho(t, \mathbf{x}), \rho \mathbf{u}(t, \mathbf{x}), E(t, \mathbf{x})) = \sum_{q=0}^{q_{max}} (1, \mathbf{v}_q, \frac{1}{2} |\mathbf{v}_q|^2) f_q(t, \mathbf{x}) \omega_q,$$

where  $(\omega_q)$  are the weights of the quadrature.

For the BGK model (4), the conservative approximation of [23] gives the following discrete BGK collision operator

$$Q_q(f) = \frac{1}{\tau} (M_q[\rho, \mathbf{u}, T] - f_q),$$

where  $M_q[\rho, \mathbf{u}, T]$  is the discrete Maxwellian that has the same discrete moments as  $f$ . It can be shown that there exists  $\alpha \in \mathbb{R}^5$  such that  $M_q[\rho, \mathbf{u}, T] = \exp(\alpha \cdot m(\mathbf{v}_q))$ , that is to say,  $\alpha$  is the unique solution of the nonlinear system

$$\sum_{q=0}^{q_{max}} m(\mathbf{v}_q) \exp(\alpha \cdot m(\mathbf{v}_q)) \omega_q = \begin{pmatrix} \rho \\ \rho \mathbf{u} \\ E \end{pmatrix},$$

where we note  $m(\mathbf{v}) = (1, \mathbf{v}, \frac{1}{2} |\mathbf{v}|^2)^T$ . This system can be solved by a Newton algorithm (see details in [23] and a more economic version of the algorithm due to Titarev in [25]).

Consequently, to apply this approach to our AMR velocity grid, we just have to chose a quadrature formula. First, we propose the standard  $\mathbb{Q}_1$  bilinear interpolation of  $f$  on each cell of the AMR grid (by using the four corners of the cell). In this case, the quadrature points are the nodes of the grid, and so are the discrete velocities that are used in the transport term. The quadrature weights are

$$\omega_q = \frac{1}{2^d} \sum_{\text{cell } \mathcal{C}_l \ni \mathbf{v}_q} |\mathcal{C}_l|,$$

where  $d = 3$  in 3D or  $2$  in 2D, and  $|\mathcal{C}_l|$  is the volume (or the area in 2D) of  $\mathcal{C}_l$ .

The number of discrete velocities can be decreased if we take a simpler  $\mathbb{P}_0$  (constant per cell) interpolation. Here, the quadrature points are the centers of the cells, and so are the

discrete velocities  $\mathbf{v}_q$  that are used in the transport term. The weights are simply the volume (or the area) of the cells:  $\omega_q = |\mathcal{C}_q|$ . Note that the number of discrete velocities that are used with this approach is equal to the number of cells of the AMR grid, which is smaller than the number of nodes, see section 4.2. We advocate the use of this latter approach, since it allows to save a lot of computer memory, and since we observed with our tests that the  $\mathbb{Q}_1$  interpolation does not give more accurate results.

**Remark 3.2.** Our AMR velocity grids are often very coarse at their boundaries (since velocities are very large there, and are in the tails of all the distribution functions). Consequently, passing from  $\mathbb{Q}_1$  to a  $\mathbb{P}_0$  grid may lead to a grid which is not large enough. Indeed, if the  $\mathbb{Q}_1$  grid is of length  $L$  (in the  $x$  direction, for instance) and its outer cells are of length  $l$ , then the length of the  $\mathbb{P}_0$  grid now is  $L - l$ . If  $l$  is large, the  $\mathbb{P}_0$  grid is not large enough. In that case, it is thus safer to modify step 2 of Algorithm 3.1 by increasing the bounds of the fine Cartesian grid:  $v_{max}^\alpha$  and  $v_{min}^\alpha$  are replaced by  $v_{max}^\alpha + \Delta v$  and  $v_{min}^\alpha - \Delta v$  for example, where  $\Delta v$  is the step of the fine grid.

### 3.5 AMR grid generation for axisymmetric flows

Many interesting 3D flows have symmetry axis (like flows around axisymmetric bodies with no incidence). In that case, if we assume that the symmetry axis is aligned with the  $x$  coordinate, it is interesting to write the kinetic equation in the cylindrical coordinate system  $(x, r, \varphi)$ , since the distribution function  $f$  is independent of  $\varphi$ , and we get

$$\partial_t f + v_x \partial_x f + \zeta \cos \omega \partial_r f - \frac{\zeta \sin \omega}{r} \partial_\omega f = Q(f), \quad (12)$$

where  $\zeta$  and  $\omega$  are the cylindrical coordinates of the velocity in the local frame  $(e_r, e_\varphi)$ , that is to say  $v \cdot e_r = \zeta \cos \omega$  and  $v \cdot e_\varphi = \zeta \sin \omega$ .

If the velocities of the upstream flow and of the solid boundaries have no component in the  $\varphi$  direction, then  $f$  is even with respect to  $\omega$ . Indeed, using the Galilean invariance of equation (12), we apply the symmetry with respect to the  $(x, z)$  plane to the equation to get that  $f(t, x, r, v_x, \zeta, -\omega)$  is also a solution.

In this case, the discrete velocity grid must be constructed for variables  $(v_x, \zeta, \omega)$ , where  $\zeta$  is non negative,  $\omega$  is in  $[0, \pi]$ , and  $v_x$  can take any value. The construction of this grid is slightly different from the Cartesian case. We briefly comment on these differences in this section.

First, bounds in  $v_x$  and  $\zeta$  directions can be easily obtained and we set

$$\begin{aligned} v_{x,max} &= \max_{\Omega} \{u_x^{CNS} + c\sqrt{RT}\}, & v_{x,min} &= \min_{\Omega} \{u_x^{CNS} - c\sqrt{RT}\}, \\ \zeta_{max} &= \max_{\Omega} \{u_r^{CNS} + c\sqrt{RT}\}, & \zeta_{min} &= 0, \end{aligned} \quad (13)$$

where  $u_x$  and  $u_r$  are the macroscopic velocities in the axial and radial directions. Since  $\omega$  is a bounded variable, there is nothing else to do at this level.

The steps  $\Delta v_x$  and  $\Delta \zeta$  of the initial grid are obtained by (10) for  $v_x$  and  $\zeta$ . However, the choice of the step in the  $\omega$  direction is not obvious. Indeed, this variable has no link with the width of the distribution. Moreover, the grid in this direction must be fine enough to capture the derivative  $\partial_\omega f$ , while in the other directions, only some moments have to be captured. After several tests (and from the experience made in [23]), it seems that a grid step  $\Delta\omega$  such that there are 30 points uniformly distributed in  $[0, \pi]$  is a good compromise between cost and accuracy.

To refine/coarsen this initial cylindrical grid, we choose to use the procedure described in the steps of sections 3.1 and 3.2 in the plane  $(v_x, \zeta)$  to generate a two-dimensional AMR grid. Indeed, for the same reasons as mentioned above, there is no reason to refine or coarsen the grid in the  $\omega$  direction. Then, the complete grid is obtained by a rotation of the two-dimensional AMR grid in the  $\omega$  direction, with the same step  $\Delta\omega$  as in the initial grid. This procedure is simple and turned out to be very efficient.

Since the AMR grid is uniform in the  $\omega$  direction, the discretization of (12) is made with the approximations derived in [23].

## 4 Numerical results

### 4.1 The 3D kinetic code

Our 3D kinetic code is written in Fortran, and can deal with 2D planar flows, or 2D axisymmetric flows as well as 3D flows on multiblock curvilinear structured grids. A steady state solution of the BGK equation for monoatomic or polyatomic gases is computed with a linearized implicit finite volume scheme, based on a velocity discretization which is conservative and entropic (see [22]). In axisymmetric cases, a specific scheme (named T-UCE) is used in order to ensure a conservative and entropic discretization of the transport operator (see [23]).

In order to deal with high computational costs (in 3D configurations for example), an hybrid parallel implementation is used: a domain decomposition in the position space used with the library routines specified by the Message Passing Interface (MPI) can deal with a large number of mesh cells. Note that this is the opposite of the strategy used in [25] (which uses a domain decomposition in the velocity space). Moreover, the OpenMP library is used with 8 threads for loops computations that are local w.r.t the position (computation of the moments, Maxwellian, etc.), thus large numbers of velocity points can be reached. We also use OpenMP for loops that are local w.r.t the velocity, like the computation of the transport terms, for instance. The communication between different domains are treated as explicit boundary conditions. This code has been run on the super-computer TERA-100 of the CEA (that has around 140 000 cores Intel Xeon 7500), by using at most 1600 cores.

Since a general description of our code is not the main subject of this paper, we do not give more details here. However, for completeness, note that interested readers can find a more detailed description of the code in the appendix.

The distribution is initialized by using a CNS pre-simulation (also used for the AMR

strategy): the number of iterations before reaching convergence is then considerably decreased (see section 4.5).

For the 2D plane examples shown below, we use a simpler version of the code, called CORBIS. This code is based on the same tools but makes use of the reduced distribution technique to reduce the velocity space to 2D only. This code only uses OpenMP directives to work on parallel computers (see [4]). It has been used on a SMP node of 48 cores AMD-Opteron-8439-SE.

For the compressible Navier-Stokes simulations, we use a 3D code developed by the CEA-CESTA for hypersonic flows at moderate to low altitudes. The same techniques as in our 3D kinetic code are used: a finite volume linearized implicit scheme with multiblock structured meshes. On solid walls, no-slip and isothermal boundary condition are used ( $u = 0$  and  $T = T_{wall}$ ).

## 4.2 A 2D plane example

Here, we illustrate our approach with the simulation of a steady flow over a infinite cylinder of radius 0.1m at Mach 20, for density and pressure of the air at an altitude of 90 km. The gas considered here is argon (molecular mass=  $6.663 \times 10^{-24}kg$ ). Namely, we have  $\rho = 3.17 \times 10^{-6}kg/m^3$ ,  $u = 5.81 \times 10^3m/s$ , and  $T = 242.4K$ . The temperature of the wall is 293K. The downstream flow is ignored, and outflow boundary conditions are used at the boundary of the right side of the domain. Since the flow is invariant along  $z$ , the simulation is made with the 2D plane model in the plane  $(x, y, 0)$ . Finally, since the flow is symmetric with respect to the line  $y = 0$ , the computation is made on the upper part on this plane only (see figure 6). See appendix A.2 for the implementation of the boundary conditions in the code.

The space mesh uses  $50 \times 50$  cells, with a refinement such that the first cell at the solid boundary is smaller than one fifth of the mean free path at the boundary.

A CNS pre-simulation and the use of formula (9-10) with parameters  $c = 4$  and  $a = 2$  give us a fine velocity grid with  $52 \times 41$  points (bounds  $\pm 11500$  and  $\pm 8900$  for  $v_x$  and  $v_y$  with a step of 450, in  $m.s^{-1}$ ), see figure 7. Algorithms 3.1 and 3.2 applied to the CNS fields give the support function and the AMR velocity grid shown in figure 8. The AMR velocity grid has 529 points, which is one fourth as small as the original fine Cartesian grid. This gain can be further increased if we define the discrete velocities as the centers of the cells of the AMR grid rather than its vertices. This gives 316 discrete velocities, hence with a gain of 6.7 times instead of 4.

First, we compare the CPU time required by the code CORBIS with the different velocity grids. While the number of iterations to reach steady state is approximately the same with both grids, the CPU time required by the original fine Cartesian velocity grid is around 7 times as large as with the new AMR grid, which is almost the same ratio as the ratio of the number of discrete velocities. The memory required with the uniform grid method is around 170 MB whereas with the use of the AMR grid, only 25 MB of memory storage is required. This shows that the new AMR grid leads to a real gain both in memory storage and CPU

time.

Then, we compare the accuracy of the results with the two grids for the macroscopic quantities. We compute the normal component of the heat flux to the boundary, which is a quantity of paramount importance in aerodynamic simulations: we find a maximum relative difference lower than 5%, which is reasonably small (see the profile of this flux on figure 9). We also compute the differences for the density, temperature and pressure in the whole computational domain:

- the mean quadratic relative difference over all the cells of the computational domain is 5% for the density, 1% for the temperature, 0.6% for the horizontal velocity, and 1.2% for the vertical velocity;
- the maximum relative difference on each cell of the computational domain is 40% for the density, 69% for the temperature, and 157% for the velocity.

The maximum relative differences are observed at the solid wall for the density and the velocity, and in the upstream flow for the temperature.

This difference is quite large, and can be explained as follows. The smallest cells of our AMR velocity grid (that are around small velocities, like velocities at the solid wall) turn out to be smaller than the cells of the Cartesian grid (size 330 instead of 450). This means that our results with the AMR grid are probably *more accurate* than the results of the Cartesian grid. Consequently, the Cartesian grid results should not be considered as a reference for this comparison.

To confirm this analysis, we make a new simulation with a Cartesian grid with a uniform step of 330 (like the smallest step of the AMR grid). We observe that relative differences between the Cartesian grid and the AMR grid are much smaller:

- the mean quadratic relative difference over all the cells of the computational domain is 2% for the density, 0.5% for the temperature, 0.3% for the horizontal velocity, and 0.7% for the vertical velocity;
- the maximum relative difference on each cell of the computational domain is 12% for the density, 13% for the temperature, 80% for the velocity.

The maximum relative difference is still too large for the velocity (80% at the boundary), but this quantity is very small in this zone and probably requires smaller velocity cells (that is a smaller parameter  $a$  for both grids). However, this inaccuracy does not deteriorates the results on the other quantities, in particular for the heat flux at the boundary. Indeed, the comparison of the heat fluxes is even better, since we find a relative difference lower than 2.5%, which is excellent. The comparison in terms of CPU time and memory storage is of course more favorable to the AMR grid here.

Note that we also did the same computations with an finer AMR grid obtained with the parameter  $a = 1$ : the difference with  $a = 2$  is less than 2% for the heat flux at the solid wall, which is quite good. The value  $a = 2$  is then clearly sufficient. However, if one is interested with the flow velocity at the boundary, we advocate to use  $a = 1$ , since the differences here are around 15%.

### 4.3 A 2D axisymmetric example

Here we consider the flow around a sphere of radius 0.1 m, at Mach 20, for density and pressure of the air at an altitude of 90 km (see figure 10). The gas is air (molecular mass =  $4.81 \times 10^{-26}$  kg), considered as a diatomic gas. Namely, the upstream density and pressures are  $\rho = 3.17 \times 10^{-6} \text{ kg/m}^3$ , and  $p = 0.16 \text{ N.m}^{-2}$ , and the temperature of the surface of the sphere is  $T = 280 \text{ K}$ . The downstream flow is ignored, and outflow boundary conditions are used at the boundary of the right side of the domain. Since the flow is rotationally symmetric, the simulation is made with the 2D axisymmetric model in the plane  $(x, y, 0)$ . Finally, since the flow is symmetric with respect to the line  $y = 0$ , the computation is made on the upper part on this plane only.

The space mesh uses  $70 \times 50$  cells, with a refinement such that the first cell at the solid boundary is smaller than one fifth of the mean free path at the boundary. A CNS pre-simulation and the use of formula (13) with parameter  $c = 4$  give us the following velocity bounds  $-8\,700$  and  $11\,000$  for  $v_x$ ,  $0$  and  $7\,500$  for  $\zeta$ , and the bounds for  $\omega$  are always  $0$  and  $\pi$ , as explained in section 3.5.

All the following simulations were made with 140 domains and 4 OpenMP threads per domain.

First, a fine uniform cylindrical velocity grid is obtained with (10) and  $a = 2$ : this gives  $65 \times 33 \times 31 = 66\,495$  discrete velocities. The steady state is reached after 1 828 iterations, in  $5\,476 \text{ s}$ . The corresponding temperature and velocity fields are shown in figure 11.

Then a cylindrical AMR grid is generated (with  $a = 2$ ): it gives 7 691 discrete velocities (with the  $\mathbb{Q}_1$  points), the steady state is reached after 1 899 iterations, in  $710 \text{ s}$ . The support function and AMR grid are shown in figure 12. Here, the gain factors in memory and in CPU time with respect to the uniform grid are around 9.

Now, we use the same AMR grid, but with the  $\mathbb{P}_0$  points (that is to say the centers of the cells), which gives 6 900 discrete velocities only. The number of iterations is close (1 796), and the CPU time is  $562 \text{ s}$ . Here, the gain factors in memory and CPU time, with respect to the uniform grid, are around 10.

Finally, we compare in figure 13 the normal heat flux at the boundary for the uniform grid and the AMR grid generated by using the CNS fields ( $\mathbb{Q}_1$  version). As in the 2D example, we find that the fluxes are very close (with a relative difference which is less than 1%). We get the same results with the  $\mathbb{P}_0$  version.

Consequently, the AMR grid ( $\mathbb{P}_0$  version) is the most efficient here: it gives the same results as the fine uniform grid, for computational and memory cost which is one ninth as small.

**Remark 4.1.** Before discussing a full 3D case, we briefly mention another experiment we have done here. We have tried to see if the use of a CNS pre-simulation could be avoided by using a direct estimation of the extreme values of the velocity and the temperature given by the Rankine-Hugoniot relations. Indeed, before the computation, we know the upstream values  $\mathbf{u}_{up}$  and  $T_{up}$ , and we can compute the downstream values  $(\mathbf{u}_{down}, T_{down}) = RH(\mathbf{u}_{up}, T_{up})$  by the Rankine-Hugoniot relations for a stationary shock. Then we have three

set of different values: upstream, downstream, and wall values ( $\mathbf{u}_{wall} = 0, T_{wall}$ ), and we can use these three sets in our algorithm 3.1 instead of the CNS fields ( $\mathbf{u}^{CNS}, T^{CNS}$ ).

With the same test case, this strategy gives a grid with 3962 points only (with  $a = 2$ ), which is half as less as with the CNS fields. The support function and AMR grid are shown in figure 14: of course, since we only have three values for the macroscopic fields here, the grid is not as smooth as with the CNS fields.

Moreover, the convergence to steady state is much slower (since we cannot use the CNS solution as an initial state, see section 4.5 for a comment on the initialization). More important, the accuracy observed on the heat flux is not as good as with the CNS fields: the difference with the fine uniform grid is here around 8% instead of 1%.

Consequently, even if strategy is simple and does not require a CNS solver, it seems to be not accurate enough.

## 4.4 A 3D example

Here, we consider the flow around a 3D object composed of cone with a spherical nose, with no incidence (see figure 15). The physical parameters are the same as in section 4.3. The space mesh uses 50000 cells, with 50 in the direction normal to the surface. Again, the downstream flow is ignored, and outflow boundary conditions are used at the boundary of the right side of the domain.

A CNS pre-simulation and the use of formula (13) with parameter  $c = 4$  and  $a = 2$  give us a uniform Cartesian grid with  $65 \times 33 \times 33 = 70785$  discrete velocities. This is much too large to make the simulation (the memory storage itself is huge). Then we only do the simulation with the AMR grid ( $\mathbb{P}_0$  version) generated by our algorithm which has only 2956 points.

The gain factor in memory storage is 22. The steady state is reached in 3424 iterations and 3566s with 400 domains and 4 threads per domain.

To estimate the gain in CPU that would be obtained if the uniform Cartesian grid could be used, we use the following remark: on the previous tests (2D plane and axisymmetric), the number of iterations to reach the steady state is almost the same for the uniform and the AMR velocity grids. Then we make a single iteration with the uniform Cartesian grid (same domain decomposition) and we multiply the CPU time obtained for this iteration by the number of iterations required with the AMR grid. We find a CPU time which is 24 times as large as with the AMR grid, which is quite important. This CPU gain is larger than the gain observed for the memory storage.

In figure 16, we show the uniform Cartesian grid and the AMR grid. In figure 17, we show the pressure field computed by the code with the AMR grid, and the heat flux along the surface is shown in figure 18.

## 4.5 Initialization with the CNS results

Since we use a CNS pre-simulation to build our discrete velocity grid, it is interesting to use it also to initialize the BGK computation, what was done in the previous simulations. It helps to reach the steady state more rapidly, as it is shown in the following example.

We take the same test case as in section 4.3, with the uniform velocity grid. In a first computation, the distribution is initialized with the (Maxwellian) upstream flow (this is the standard initialization), and the scheme converges to the steady state in 3 549 iterations. Then we do the same computation, but now the initial state is the Maxwellian distribution computed with the macroscopic variables that are given by the steady CNS solution. Then the steady state is reached in 1 822 iterations only.

A similar gain (half as many iterations as with the standard initialization) is obtained in all our test cases.

## 5 Conclusion and perspectives

In this paper, we have proposed a method to refine and coarsen a global velocity grid of a discrete velocity approximation of a kinetic equation. It is based on a criterion called “the support function” that links the local size of the velocity grid to the macroscopic temperature and velocity of the flow. Our algorithm uses the macroscopic fields given by a compressible Navier-Stokes pre-simulation to automatically generate an optimal velocity grid.

This approach has been tested in a 3D computational code (and 2D plane and axisymmetric versions) which uses the BGK model of rarefied gas dynamics. Typical test cases in hypersonic re-entry problems (for simplified geometries) have been used. We observed that using our algorithm allows important gains both in CPU time and memory storage, up to a factor 24. This allows to make simulations that are hardly possible with standard grids, even on super computers.

Note that this work might be extended to unsteady flow simulations: we could indeed modify the velocity grid at different time steps according to our refinement algorithm. This might also be used during the iterations of our iterative solver for a steady simulation. However, this would require to interpolate the solution between two successive grids, and this might lead to an important increase of the CPU time. This clearly requires more investigations to get an efficient method.

An obvious, and straightforward, extension of this work will be the use of modified BGK models (like ES or Shakhov models), in order to have a model with a correct Prandtl number.

Moreover, we will try to investigate non isotropic AMR velocity grids by taking into account translational kinetic temperatures in different directions, that can be obtained by the stress tensor computed by the CNS simulation. It could also be interesting to refine the grid so as to capture the discontinuity of the distribution function in the velocity space, but this will require a rather different approach.

Our main short term project is to improve the physical accuracy of our code by implementing a simple BGK-like model for multi-species reactive flows. An application of our

method to the full Boltzmann collision operator is less obvious but might also be explored.

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## A Overview of the 3D kinetic code

### A.1 The linearized implicit scheme

Our code is an extension of the code presented in [23] to 3D polyatomic flows. It is based on the following reduced BGK model

$$\begin{aligned}\partial_t f + \mathbf{v} \cdot \nabla_{\mathbf{x}} f &= \frac{1}{\tau} (M(U) - f) \\ \partial_t g + \mathbf{v} \cdot \nabla_{\mathbf{x}} g &= \frac{1}{\tau} \left( \frac{\delta}{2} RTM(U) - g \right),\end{aligned}$$

where  $U = \langle m f + e^{(5)} g \rangle = (\rho, \rho \mathbf{u}, E = \frac{1}{2} \rho |\mathbf{u}|^2 + \frac{3+\delta}{2} \rho RT)$  is the vector of macroscopic mass, momentum, and energy density. Here, we use the standard notation  $\langle \cdot \rangle = \int_{\mathbb{R}^3} \cdot d\mathbf{v}$  for any vector valued function of  $\mathbf{v}$ ,  $m(v) = (1, \mathbf{v}, \frac{1}{2} |\mathbf{v}|^2)$  is the vector of collisional invariants, and  $e^{(5)} = (0, 0, 0, 0, 1)$ . Moreover,  $M(U)$  is the standard Maxwellian distribution defined through the density, velocity, and temperature corresponding to the vector  $U$  above (see (2)).

This model comes from the reduction of a BGK model for the full distribution function  $F(t, \mathbf{x}, \mathbf{v}, I)$ , where  $I$  is the internal energy variable, and  $\delta$  is the number of internal degrees of freedom (see [17] for the first use of this technique and [3] for an application to polyatomic gases). Consequently, it accounts for any number of internal degrees of freedom. For instance, a diatomic gas can be described with  $\delta = 2$ .

This model is first discretized with respect to the velocity variable. We follow the approximation of [23] and its extension to polyatomic gases [16]. We assume we have a velocity grid  $\{\mathbf{v}_q, q = 0 : q_{max}\}$  (like the grids described in this paper). The continuous distributions  $f$  and  $g$  are then replaced by their approximations at each point  $\mathbf{v}_q$ , and we get the following discrete velocity BGK system

$$\begin{aligned}\partial_t f_q + \mathbf{v}_q \cdot \nabla_{\mathbf{x}} f_q &= \frac{1}{\tau} (M_q(U) - f_q) \\ \partial_t g_q + \mathbf{v}_q \cdot \nabla_{\mathbf{x}} g_q &= \frac{1}{\tau} (N_q(U) - g_q),\end{aligned}$$

where  $(M_q(U), N_q(U))$  is an approximation of  $(M(U), \frac{\delta}{2}RTM(U))$  that has to be defined. As explained in this paper, we assume we have quadrature weights  $\{\omega_q\}$  corresponding to our discrete velocity grid, so that the moment vector of the discrete distributions is

$$U = \sum_{q=0}^{q_{max}} (m(\mathbf{v}_q) f_q + e^{(5)} g_q) \omega_q.$$

As proposed in [16], the discrete equilibrium  $(M_q(U), N_q(U))$  is constructed so that it has the same moments as  $(f, g)$ , that is to say

$$\sum_{q=0}^{q_{max}} (m(\mathbf{v}_q) M_q(U) + e^{(5)} N_q(U)) \omega_q = U. \quad (14)$$

In our code,  $(M_q(U), N_q(U))$  is determined through the entropic variable  $\alpha$  such that  $M_q(U) = \exp(\alpha \cdot m(\mathbf{v}_q))$  and  $N_q(U) = \frac{\delta}{2} \frac{1}{-\alpha_5} \exp(\alpha \cdot m(\mathbf{v}_q))$ , by solving (14) by a Newton algorithm. We mention that the computational cost of this algorithm can be significantly reduced by using a nice idea due to Titarev [25]. This optimization will be used in a future version of our code.

The discrete velocity BGK system is then discretized by a finite volume scheme on a multi-block curvilinear 3D mesh of hexahedral cells  $\Omega_{ijk}$ , with indices  $i, j, k = 1$  to  $i_{max}, j_{max}, k_{max}$ , respectively. Denoting by  $f_{ijkq}^n$  an approximation of the average of  $f$  at time  $t_n$  on a cell  $\Omega_{ijk}$  at the discrete velocity  $\mathbf{v}_q$ , our scheme reads, in its implicit version,

$$\begin{aligned} \frac{f_{ijkq}^{n+1} - f_{ijkq}^n}{\Delta t} + (\mathbf{v}_q \cdot \nabla_{\mathbf{x}} f_q^{n+1})_{ijk} &= \frac{1}{\tau_{ijk}^{n+1}} (M_q(U_{ijk}^{n+1}) - f_{ijkq}^{n+1}), \\ \frac{g_{ijkq}^{n+1} - g_{ijkq}^n}{\Delta t} + (\mathbf{v}_q \cdot \nabla_{\mathbf{x}} g_q^{n+1})_{ijk} &= \frac{1}{\tau_{ijk}^{n+1}} (N_q(U_{ijk}^{n+1}) - g_{ijkq}^{n+1}), \end{aligned}$$

where

$$U_{ijk}^{n+1} = \sum_{q=0}^{q_{max}} (m(\mathbf{v}_q) f_{ijkq}^{n+1} + e^{(5)} g_{ijkq}^{n+1}) \omega_q.$$

The discrete divergence  $(\mathbf{v}_q \cdot \nabla_{\mathbf{x}} f_q)_{ijk}^{n+1}$  is given by the following second order upwind approximation (with the Yee limiter [28]):

$$(\mathbf{v}_q \cdot \nabla_{\mathbf{x}} f_q^{n+1})_{ijk} = \frac{1}{|\Omega_{i,j,k}|} \left( \left( \Phi_{i+\frac{1}{2},j,k} - \Phi_{i-\frac{1}{2},j,k} \right) + \left( \Phi_{i,j+\frac{1}{2},k} - \Phi_{i,j-\frac{1}{2},k} \right) + \left( \Phi_{i,j,k+\frac{1}{2}} - \Phi_{i,j,k-\frac{1}{2}} \right) \right), \quad (15)$$

where

$$\begin{aligned} \Phi_{i+\frac{1}{2},j,k} &= (\mathbf{v}_q \cdot \nu_{i+\frac{1}{2},j,k})^+ f_{ijkq}^{n+1} + (\mathbf{v}_q \cdot \nu_{i+\frac{1}{2},j,k})^- f_{i+1,j,k,q}^{n+1} \\ &\quad + |\mathbf{v}_q \cdot \nu_{i+\frac{1}{2},j,k}| \min\text{mod}(\Delta_{i-\frac{1}{2}}^{n+1}, \Delta_{i+\frac{1}{2}}^{n+1}, \Delta_{i+\frac{3}{2}}^{n+1}) \end{aligned}$$

is the second order numerical flux across the face between  $\Omega_{i,j,k}$  and  $\Omega_{i+1,j,k}$ , and  $\nu_{i+\frac{1}{2},j,k}$  is the normal vector to this face directed from  $\Omega_{i,j,k}$  to  $\Omega_{i+1,j,k}$  while its norm is equal to the area of the face. In the minmod limiter function, we use the notation  $\Delta_{i+\frac{1}{2}}^{n+1} = f_{i+1,j,k,q}^{n+1} - f_{ijkq}^{n+1}$ . Finally, we use the standard notation  $a^\pm = (a \pm |a|)/2$  for every number  $a$ . The numerical fluxes across the other faces are defined accordingly. For the sequel, it is useful to denote by  $(\mathbf{v}_q \cdot \nabla_{\mathbf{x}} f_q^{n+1})_{ijk}^{1st\ order}$  the corresponding linear first order upwind discretization (that is to say, defined by (15) where the minmod term is set to 0).

The advantage of the time implicit approximation is that it ensures unconditional stability, which allows us to take large time steps, and hence to get rapid convergence to the steady state. Of course, this scheme is too expensive, since it requires to solve a non linear system at each time iteration. Therefore, we rather use the following linearization. First, the Maxwellian is linearized by a first order Taylor expansion:

$$M_q(U_{ijk}^{n+1}) = M_q(U_{ijk}^n + (U_{ijk}^{n+1} - U_{ijk}^n)) \approx M_q(U_{ijk}^n) + \partial_U M_q(U_{ijk}^n)(U_{ijk}^{n+1} - U_{ijk}^n),$$

and the same for  $N_q(U_{ijk}^{n+1})$  (see section A.4 for explicit expressions of the Jacobian matrices). Then the discrete divergence, which is not differentiable due to the limiter, is linearized by using the corresponding first order upwind approximation. This is sometimes called a ‘‘frozen coefficient technique’’, which gives

$$\begin{aligned} (\mathbf{v}_q \cdot \nabla_{\mathbf{x}} f_q)^{n+1}_{ijk} &= (\mathbf{v}_q \cdot \nabla_{\mathbf{x}} (f_q^n + (f_q^{n+1} - f_q^n)))_{ijk} \\ &\approx (\mathbf{v}_q \cdot \nabla_{\mathbf{x}} f_q^n)_{ijk} + (\mathbf{v}_q \cdot \nabla_{\mathbf{x}} (f_q^{n+1} - f_q^n))_{ijk}^{1st\ order} \end{aligned}$$

Then, denoting by  $\delta f_{i,j,k,q} = f_{ijkq}^{n+1} - f_{ijkq}^n$  (same notation for  $g$ ), and by  $\delta U_{i,j,k}$  the moments of  $(\delta f_{i,j,k}, \delta g_{i,j,k})$ , our scheme reads in the following form:

$$\frac{\delta f_{i,j,k,q}}{\Delta t} + (\mathbf{v}_q \cdot \nabla_{\mathbf{x}} \delta f_q)_{ijk}^{1st\ order} - \frac{1}{\tau_{i,j,k}^n} \partial_U M_q(U_{ijk}^n)(\delta U_{i,j,k}) = RHS f_{i,j,k}^n \quad (16)$$

$$\frac{\delta g_{i,j,k,q}}{\Delta t} + (\mathbf{v}_q \cdot \nabla_{\mathbf{x}} \delta g_q)_{ijk}^{1st\ order} - \frac{1}{\tau_{i,j,k}^n} \partial_U N_q(U_{ijk}^n)(\delta U_{i,j,k}) = RHS g_{i,j,k}^n, \quad (17)$$

where the right-hand sides are given by

$$\begin{aligned} RHS f_{i,j,k}^n &= -(\mathbf{v}_q \cdot \nabla_{\mathbf{x}} f_q^n)_{ijk} + \frac{1}{\tau_{ijk}^n} (M_q(U_{ijk}^n) - f_{ijkq}^n) \\ RHS g_{i,j,k}^n &= -(\mathbf{v}_q \cdot \nabla_{\mathbf{x}} g_q^n)_{ijk} + \frac{1}{\tau_{ijk}^n} (N_q(U_{ijk}^n) - g_{ijkq}^n). \end{aligned}$$

If our scheme converges to steady state, then the right-hand side is zero, and we get a second order discrete steady solution.

## A.2 Numerical boundary conditions

Numerically, the boundary conditions are implemented by the standard ghost cell technique, which is used as follows. When an index  $(i, j, k)$  corresponds to a cell located at the boundaries of the domain, there appear unknown values in the numerical fluxes like  $f_{i,0,k,q}^n$  and  $f_{i,j_{max}+1,k,q}^n$ , for the cells  $(i, 1, k)$  and  $(i, j_{max}, k)$  for instance (see (15)). Corresponding cells  $(i, 0, k)$ ,  $(i, j_{max}+1, k)$ , etc. are called ghost-cells. These values are classically defined according to the boundary conditions that are specified for the problem. Here we use several kinds of boundary conditions: solid wall interactions, inflow and outflow boundary conditions at artificial boundaries, as well as symmetry boundary conditions along symmetry planes and symmetry axes.

For the diffuse reflection, the incident molecules in a boundary cell of index  $(i, 1, k)$  are supposed to be re-emitted by the wall from a ghost cell of index  $(i, 0, k)$ . This cell is the mirror cell of  $(i, 1, k)$  with respect to the wall. The diffuse reflection (5)–(6) is then modeled by

$$(f_{i,0,k,q}^n, g_{i,0,k,q}^n) = \sigma_{i,1,k} (M_q^{wall}, N_q^{wall}), \quad \mathbf{v}_q \cdot \mathbf{n}_{i,1,k} > 0, \quad (18)$$

where  $\sigma_{i,1,k}$  is determined so as to avoid a mass flux across the wall, that is to say between cells  $(i, 0, k)$  and  $(i, 1, k)$ . In this relation,  $(M_q^{wall}, N_q^{wall})$  is a discrete conservative approximation of the wall Maxwellians  $(1, \frac{\delta}{2}RT_{wall}) \frac{1}{(2\pi RT_{wall})^{3/2}} \exp(-|\mathbf{v}|^2/2RT_{wall})$ . Relation (6) gives

$$\sigma_{i,1,k} = - \frac{\sum_{\mathbf{v}_q \cdot \mathbf{n}_{i,1,k} < 0} \mathbf{v}_q \cdot \mathbf{n}_{i,1,k} f_{i,1,k,q}^n \omega_q}{\sum_{\mathbf{v}_q \cdot \mathbf{n}_{i,1,k} > 0} \mathbf{v}_q \cdot \mathbf{n}_{i,1,k} M_q^{wall} \omega_q}.$$

For the inflow boundary condition, for instance at a boundary cell  $(i, j_{max}+1, k)$ , we simply set the ghost cell value to the upstream Maxwellian distributions  $(M_q^{upstream}, N_q^{upstream})$  (defined through the upstream values of  $\rho$ ,  $u$ , and  $T$ ):

$$(f_{i,j_{max}+1,k,q}^n, g_{i,j_{max}+1,k,q}^n) = (M_q^{upstream}, N_q^{upstream}).$$

For the outflow boundary condition, we set the ghost cell value to the value of the corresponding boundary cell:

$$(f_{i,j_{max}+1,k,q}^n, g_{i,j_{max}+1,k,q}^n) = (f_{i,j_{max},k,q}^n, g_{i,j_{max},k,q}^n).$$

Finally, for a cell  $(1, j, k)$  in a symmetry plane (for instance the plane  $(0, y, z)$ ) we use the symmetry of the distribution functions to set

$$(f_{0,j,k,q}^n, g_{0,j,k,q}^n) = (f_{1,j,k,q'}^n, g_{1,j,k,q'}^n),$$

where  $q'$  is such that  $\mathbf{v}_{q'}$  is the symmetric of  $\mathbf{v}_q$  with respect to the symmetry plane. In this case, our AMR velocity grid is constructed such that it is also symmetric with respect to this plane, which gives  $\mathbf{v}_{q'} = (-v_{x,q}, v_{y,q}, v_{z,q})$ . This grid is obtained in two steps: first, a part on one side of the symmetry plane is obtained by using our algorithms (support function and

AMR grid generation), and then this part is symmetrized to obtain the part on the other side.

For the second order numerical flux, we also need the values of a second layer of ghost cells with indices like  $(-1, j, k)$  or  $(i_{max} + 2, j, k)$ , etc. For these values, we simply copy the value of the corresponding ghost cell in the first layer, that is to say  $f_{-1,j,k,q}^n = f_{0,j,k,q}^n$  for instance. This treatment makes the accuracy of our scheme reduce to first order at the boundary (since this makes the flux limiters equal to 0). It may be more relevant to use extrapolation techniques, which is the subject of a work in progress. Only the boundary condition on a symmetry plane is treated differently: here, we use the symmetry of the distributions to set

$$(f_{-1,j,k,q}^n, g_{-1,j,k,q}^n) = (f_{2,j,k,q'}^n, g_{2,j,k,q'}^n),$$

where  $v_{q'}$  has been defined above.

### A.3 Linear solver

At each time iteration, our linearized implicit scheme requires to solve the large linear system (16–17). It is therefore interesting to write it in the following matrix form:

$$\left(\frac{I}{\Delta t} + T + R^n\right)\delta F = RHS^n,$$

where  $\delta F = (\delta f_{i,j,k,q}, \delta g_{i,j,k,q})$  is a large vector that stores all the unknowns,  $I$  is the unit matrix,  $T$  is a matrix such that

$$(T\delta F)_{i,j,k,q} = \left( \left( \mathbf{v}_q \cdot \nabla_{\mathbf{x}} \delta f_q \right)_{ijk}^{1st\ order}, \left( \mathbf{v}_q \cdot \nabla_{\mathbf{x}} \delta g_q \right)_{ijk}^{1st\ order} \right), \quad (19)$$

$R^n$  is the relaxation matrix such that

$$(R^n \delta F)_{i,j,k,q} = \left( -\frac{1}{\tau_{i,j,k}^n} \partial_U M_q(U_{ijk}^n) (\delta U_{i,j,k}), -\frac{1}{\tau_{i,j,k}^n} \partial_U N_q(U_{ijk}^n) (\delta U_{i,j,k}) \right), \quad (20)$$

and  $RHS^n = (RHS f_{i,j,k}^n, RHS g_{i,j,k}^n)$ .

For simplicity, we use explicit boundary conditions, which means  $\delta F_{i,j,k,q} = 0$  in ghost cells. This implies that  $T$  has a simple block structure, which is used in the following.

The algorithm used in our code is based on a coupling between the iterative Jacobi and Gauss-Seidel methods. First, the relaxation matrix  $R^n$  is splitted into its diagonal  $\Delta^n$  and its off-diagonal  $-E^n$ , so that we get the following system (this the Jacobi iteration):

$$\left(\frac{I}{\Delta t} + T + \Delta^n\right)\delta F = RHS^n + E^n \delta F. \quad (21)$$

Note that  $E^n$  is very sparse: the product  $E^n \delta F$  is local in space and can be written  $[E_{i,j,k}^n \delta F_{i,j,k}]_q$  (see section A.4). The left-hand side of this system has a three level tridiagonal block structure. We solve it by using a line Gauss-Seidel iteration which is described below.

With curvilinear grids, in many cases, and in particular for re-entry problems, the flow is aligned with a mesh (generally aligned with solid boundaries), which means that the largest variations occur along the orthogonal direction, say the direction of index  $i$ , for instance. The idea is to use an “exact” inversion of  $T$  along this direction. This is done with a sweeping strategy sometimes called “Gauss-Seidel line iteration”. First, we rewrite system (21) pointwise, as follows:

$$\left( \frac{1}{\Delta t} + A + B + C + \Delta_{i,j,k,q}^n \right) \delta F_{i,j,k,q} + A^- \delta F_{i+1,j,k,q} + A^+ \delta F_{i-1,j,k,q} \\ + B^- \delta F_{i,j+1,k,q} + B^+ \delta F_{i,j-1,k,q} + C^- \delta F_{i,j,k+1,q} + C^+ \delta F_{i,j,k-1,q} = RHS_{i,j,k,q}^n + [E_{i,j,k}^n \delta F_{i,j,k}]_q,$$

where coefficients  $A, B, C, A^\pm, B^\pm, C^\pm$  are standard notations for line-Gauss-Seidel method and can be easily derived from (21), (19), and (15). Then the linear solver reads as shown in Algorithm 1.

- Remark A.1.**
1. We observe that we get a fast convergence to steady state by using a few step of this linear solver (say  $P = 2$  or  $3$ ). This is due to the “exact” inversion along the direction of largest variation of the flow.
  2. In practice, we add one back substitution step right after each forward substitution step.
  3. This solver is a straightforward extension of the linear solver used in [23]. More sophisticated versions of this solver could be tested.

## A.4 Jacobian matrices and the relaxation matrix $R^n$

Elementary calculus gives the following formula:

$$\partial_U M_q(U) = M_q(U) m(\mathbf{v}_q) A(U)^{-1} \\ \partial_U N_q(U) = N_q(U) \left( m(\mathbf{v}_q) - \frac{1}{\alpha_5(U)} e^{(5)} \right) A(U)^{-1},$$

where  $A(U)$  is the following  $5 \times 5$  matrix

$$A(U) = \sum_{q=0}^{q_{max}} \left( m(\mathbf{v}_q)^T m(\mathbf{v}_q) M_q(U) + e^{(5)T} \left( m(\mathbf{v}_q) - \frac{1}{\alpha_5(U)} e^{(5)} \right) N_q(U) \right) \omega_q.$$

Consequently, by using the definition (20) of the relaxation matrix  $R^n$ , a careful algebra shows that its diagonal elements  $\Delta_{i,j,k,q}^n$  can be separated into the following two blocks

---

**Algorithm 1** Jacobi/line-Gauss-Seidel algorithm

---

$\delta F^{(0)} = 0$  % initialization

**for**  $p$  from 0 to  $P$  **do** % iterations of the solver

**for**  $q$  from 0 to  $q_{max}$  **do** % loop over the discrete velocities (Jacobi loop)

        % one sweep in  $j$  direction (forward substitution)

**for**  $k$  from 1 to  $k_{max}$  **do**

**for**  $j$  from 1 to  $j_{max}$  **do**

                    solve exactly the tridiagonal system

$$\begin{aligned} & \left( \frac{1}{\Delta t} + A + B + \Delta_{i,j,k,q}^n \right) \delta F_{i,j,k,q}^{(p+\frac{1}{2})} + A^- \delta F_{i+1,j,k,q}^{(p+\frac{1}{2})} + A^+ \delta F_{i-1,j,k,q}^{(p+\frac{1}{2})} \\ & = RHS_{i,j,k,q}^n + [E_{i,j,k}^n \delta F_{i,j,k,q}^{(p)}] - B^+ \delta F_{i,j-1,k,q}^{(p+\frac{1}{2})} - B^- \delta F_{i,j+1,k,q}^{(p)} \end{aligned}$$

**end for**

**end for**

        % one sweep in  $k$  direction (forward substitution)

**for**  $j$  from 1 to  $j_{max}$  **do**

**for**  $k$  from 1 to  $k_{max}$  **do**

                    solve exactly the tridiagonal system

$$\begin{aligned} & \left( \frac{1}{\Delta t} + A + C + \Delta_{i,j,k,q}^n \right) \delta F_{i,j,k,q}^{(p+1)} + A^- \delta F_{i+1,j,k,q}^{(p+1)} + A^+ \delta F_{i-1,j,k,q}^{(p+1)} \\ & = RHS_{i,j,k,q}^n + [E_{i,j,k}^n \delta F_{i,j,k,q}^{(p)}] - C^+ \delta F_{i,j,k-1,q}^{(p+1)} - C^- \delta F_{i,j,k+1,q}^{(p+\frac{1}{2})} \end{aligned}$$

**end for**

**end for**

**end for**

    compute the moments of  $\delta F^{(p+1)}$  % for the action of  $\Delta_{i,j,k,q}^n$  and  $E_{i,j,k}^n$  onto  $\delta F^{(p+1)}$

**end for**

set  $\delta F = \delta F^{(P+1)}$

---

$(\Delta_{i,j,k,q}^{n,f}, \Delta_{i,j,k,q}^{n,g})$ :

$$\begin{aligned}\Delta_{i,j,k,q}^{n,f} &= \frac{1}{\tau_{i,j,k}^n} (M_q[U_{ijk}^n]m(\mathbf{v}_q)A(U_{ijk}^n)^{-1}m(\mathbf{v}_q)^T\omega_q - 1), \\ \Delta_{i,j,k,q}^{n,g} &= \frac{1}{\tau_{i,j,k}^n} \left( N_q[U_{ijk}^n] \left( m(\mathbf{v}_q) - \frac{1}{\alpha_5(U_{ijk}^n)} e^{(5)} \right) A(U_{ijk}^n)^{-1} e^{(5)T} \omega_q - 1 \right).\end{aligned}$$

Therefore, the product of the off-diagonal part  $-E^n$  of  $R^n$  with  $\delta F$  (as used in algorithm 1) is simply

$$\begin{aligned}- [E_{i,j,k}^n \delta F_{i,j,k}]_q &= \left( -\frac{1}{\tau_{i,j,k}^n} \partial_U M_q(U_{ijk}^n)(\delta U_{i,j,k}) - \Delta_{i,j,k,q}^{n,f} \delta f_{i,j,k,q} \right. \\ &\quad \left. , -\frac{1}{\tau_{i,j,k}^n} \partial_U N_q(U_{ijk}^n)(\delta U_{i,j,k}) - \Delta_{i,j,k,q}^{n,g} \delta g_{i,j,k,q} \right).\end{aligned}$$

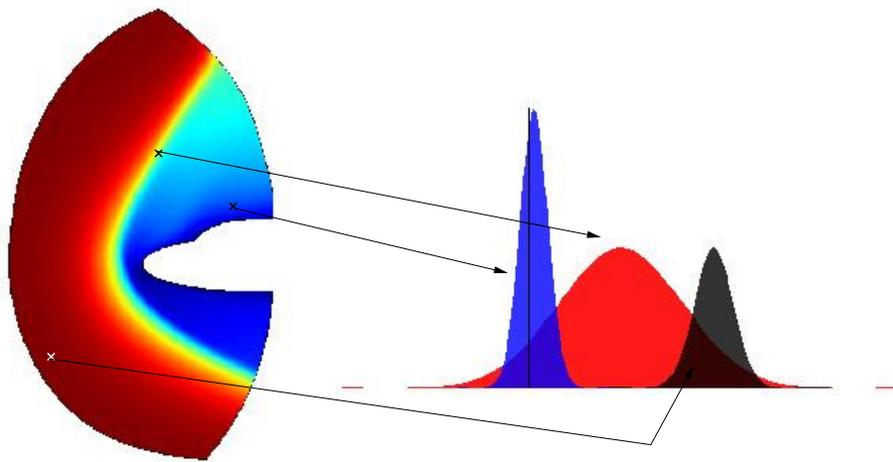


Figure 1: Three distribution functions in different space points of a computational domain for a re-entry problem

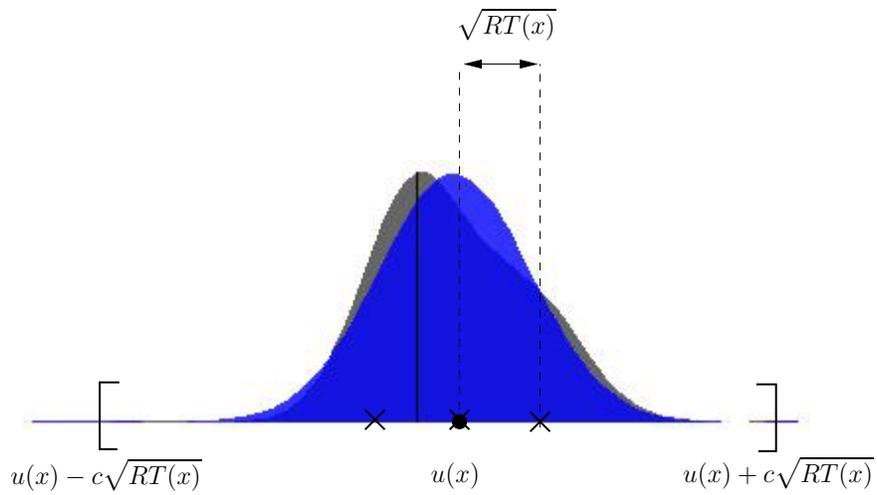


Figure 2: Support of a distribution function  $f(x, \cdot)$ : the distribution (in black), its corresponding Maxwellian (in blue), and the corresponding support based on the bulk velocity  $u$  and the temperature  $T$ .

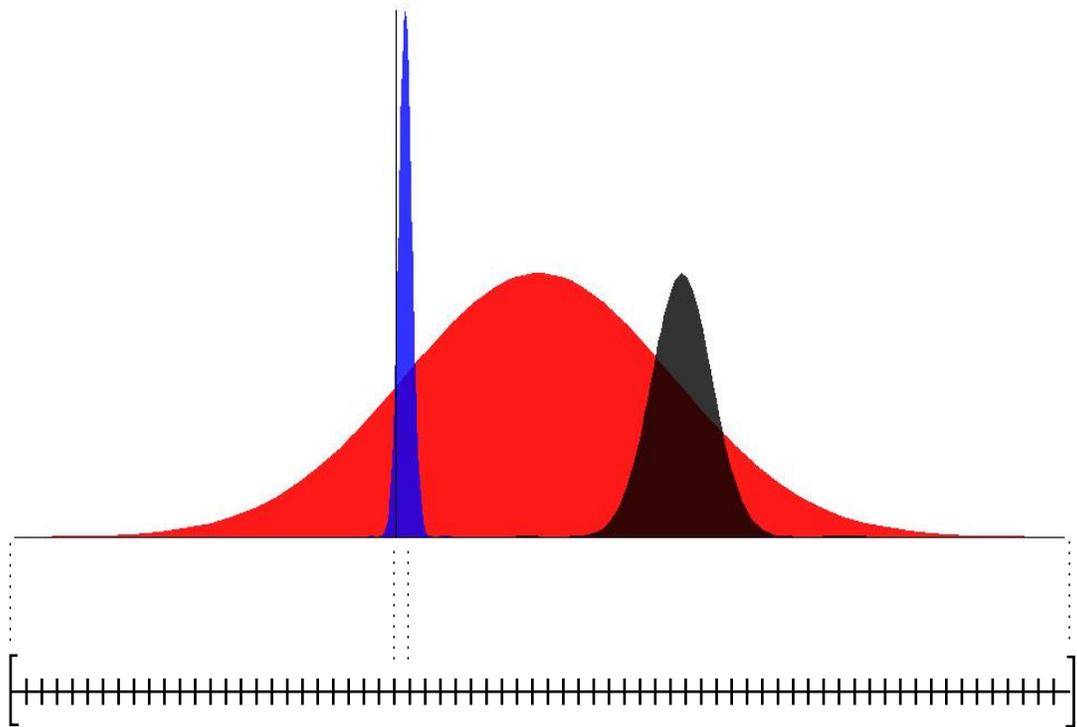


Figure 3: Three different distribution functions and a global grid that satisfies (7)-(8): the bounds are given by the largest distribution, the step is given by the narrowest distribution.

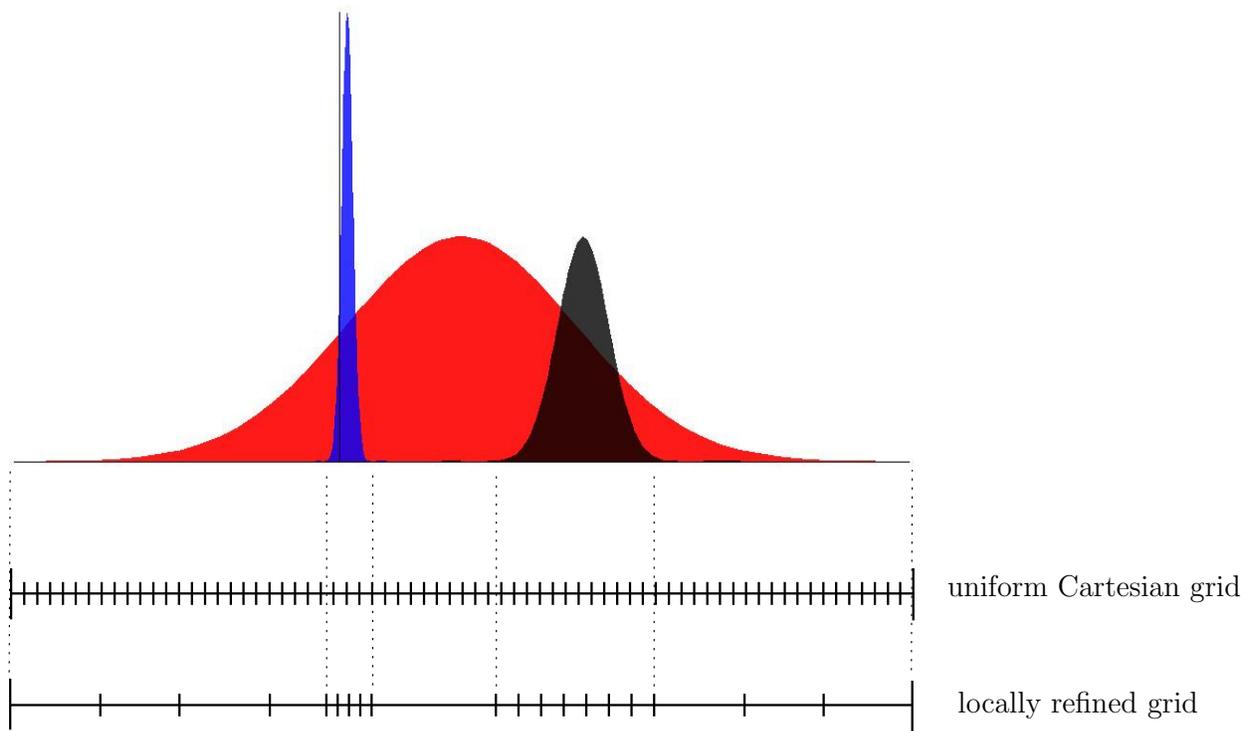


Figure 4: Three different distribution functions (top), the corresponding uniform Cartesian velocity grid (middle), and the grid locally refined in the support of the distributions (coarsened elsewhere).

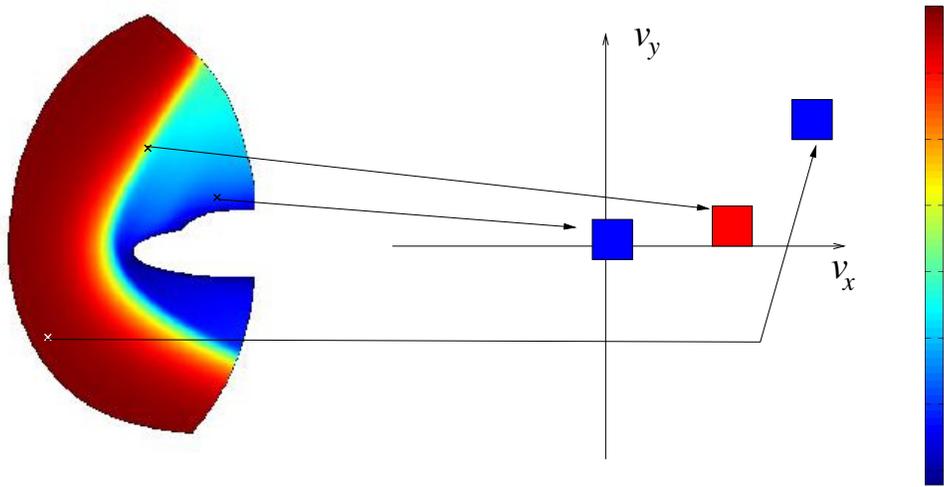


Figure 5: The macroscopic CNS flow (left), and some values of  $\phi$  at three different  $v$  (right).

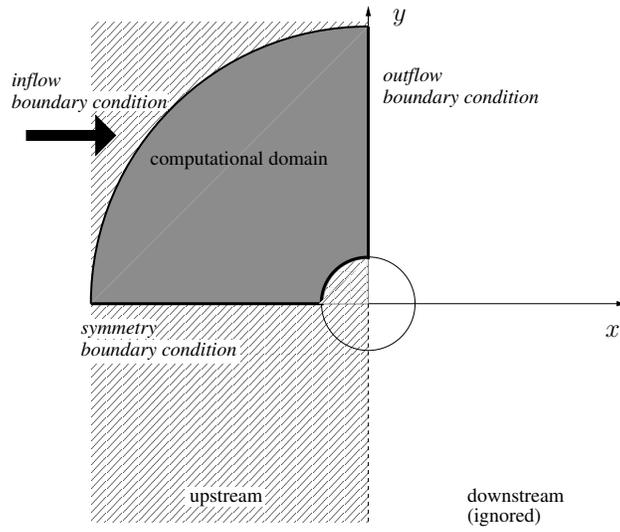


Figure 6: Plane flow around a cylinder: geometry and computational domain. By symmetry with respect to the axis  $x$ , the computational domain is defined for  $y > 0$  only. The downstream flow is not simulated.

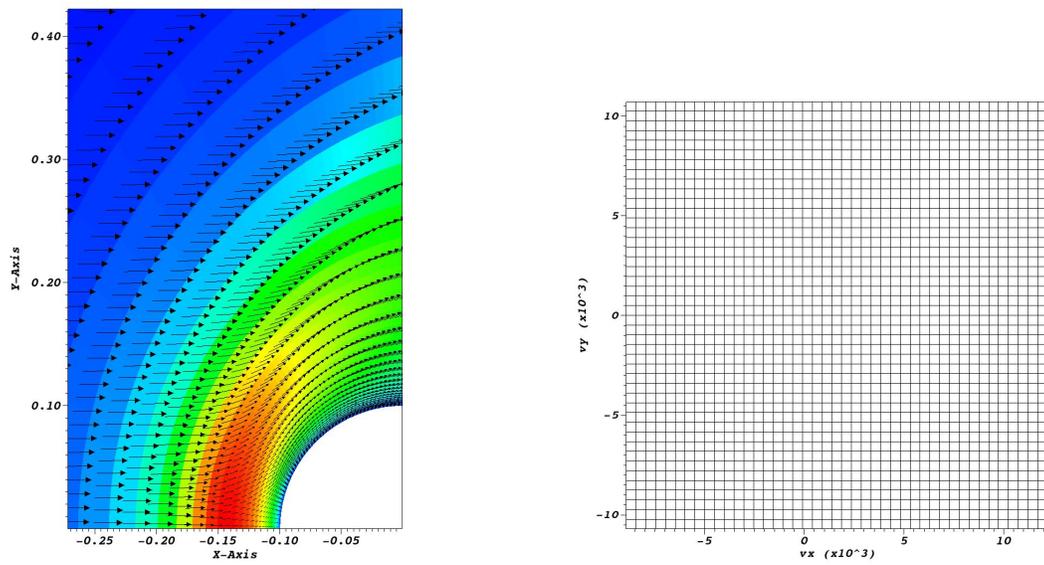


Figure 7: Plane flow around a cylinder: CNS velocity and temperature fields (left), corresponding fine Cartesian velocity grid (right).

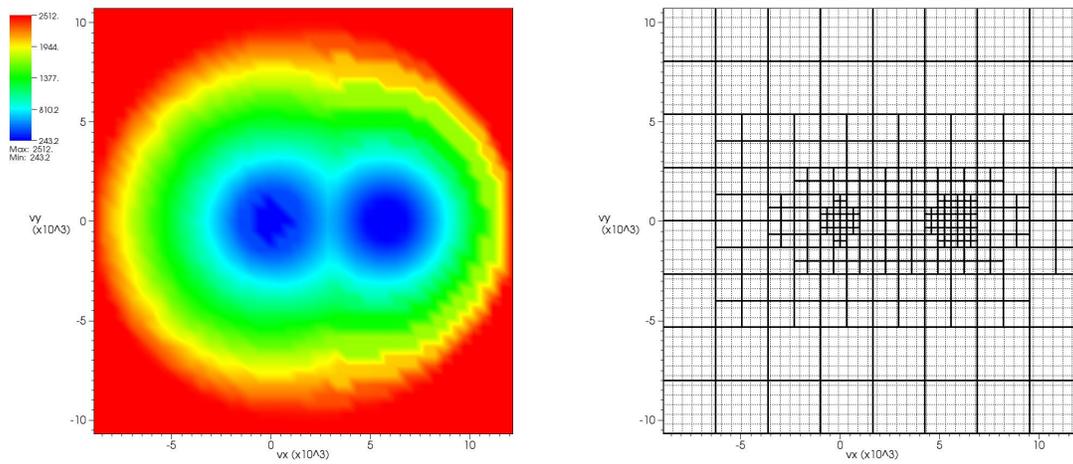


Figure 8: Plane flow around a cylinder. Left: support function, Right: velocity grids (solid line: induced AMR velocity grid, dotted line: initial fine Cartesian grid).

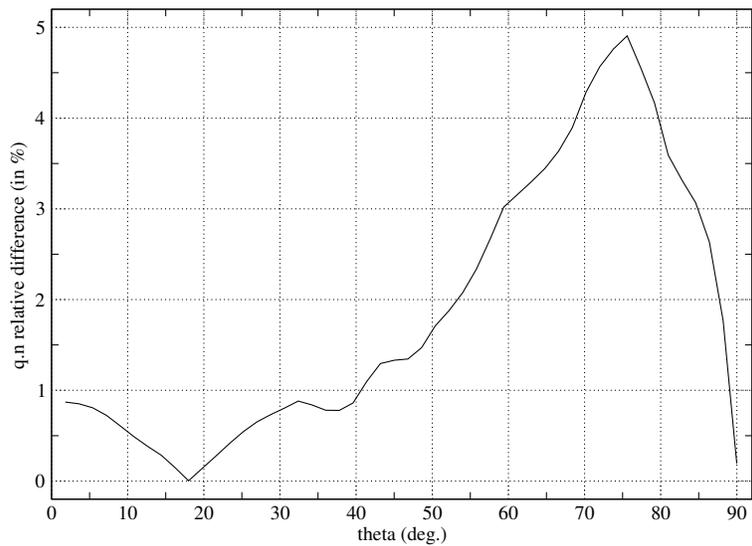
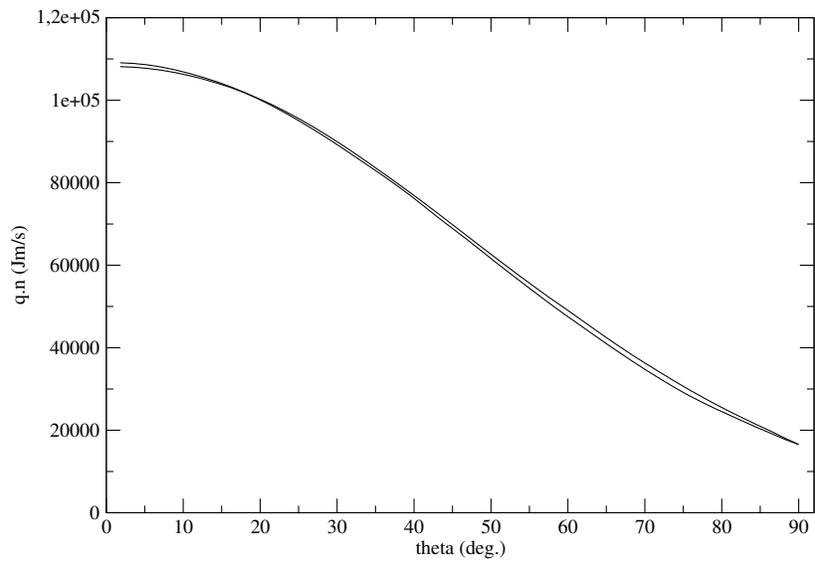


Figure 9: Plane flow around a cylinder. Comparison of the component of the heat flux normal to the solid wall: fluxes obtained with the fine Cartesian grid and the AMR grid (top), relative difference in percent (bottom). The angle  $\theta$  defines the position of the point along the solid boundary, such that  $\theta=0$  at the stagnation point and  $\theta=90$  at the end of the wall.

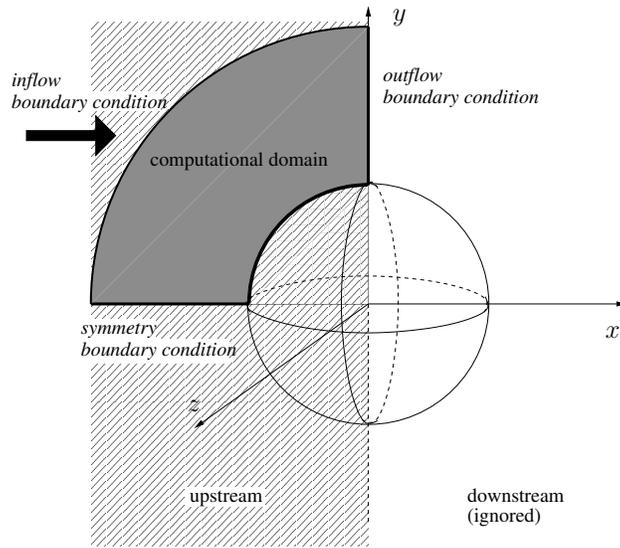


Figure 10: Axisymmetric flow around a sphere: geometry and computational domain. By rotational symmetry and by symmetry with respect to the axis  $x$ , the computational domain is included in the plane  $(x, y, 0)$ . The downstream flow is not simulated.

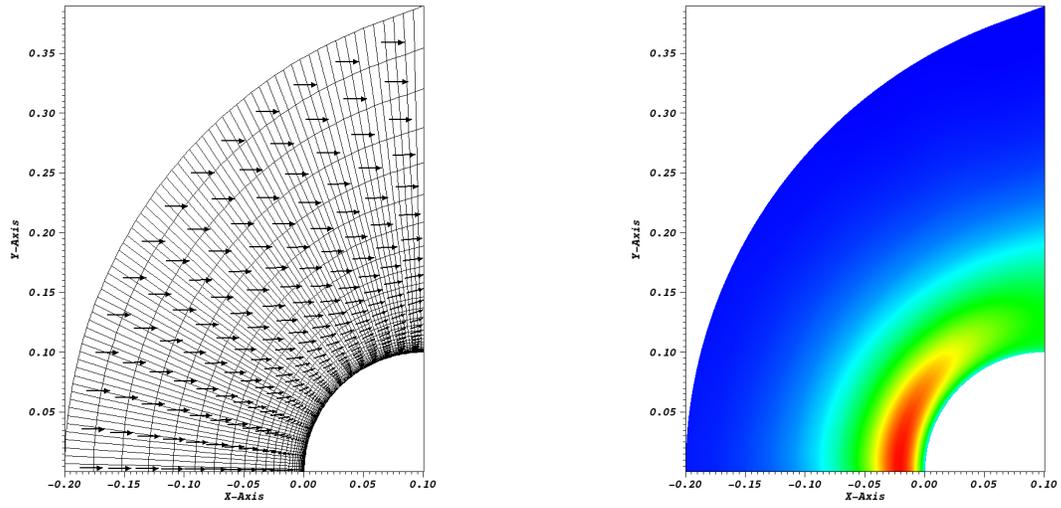


Figure 11: Axisymmetric flow around a sphere: velocity field, spatial mesh and temperature field.

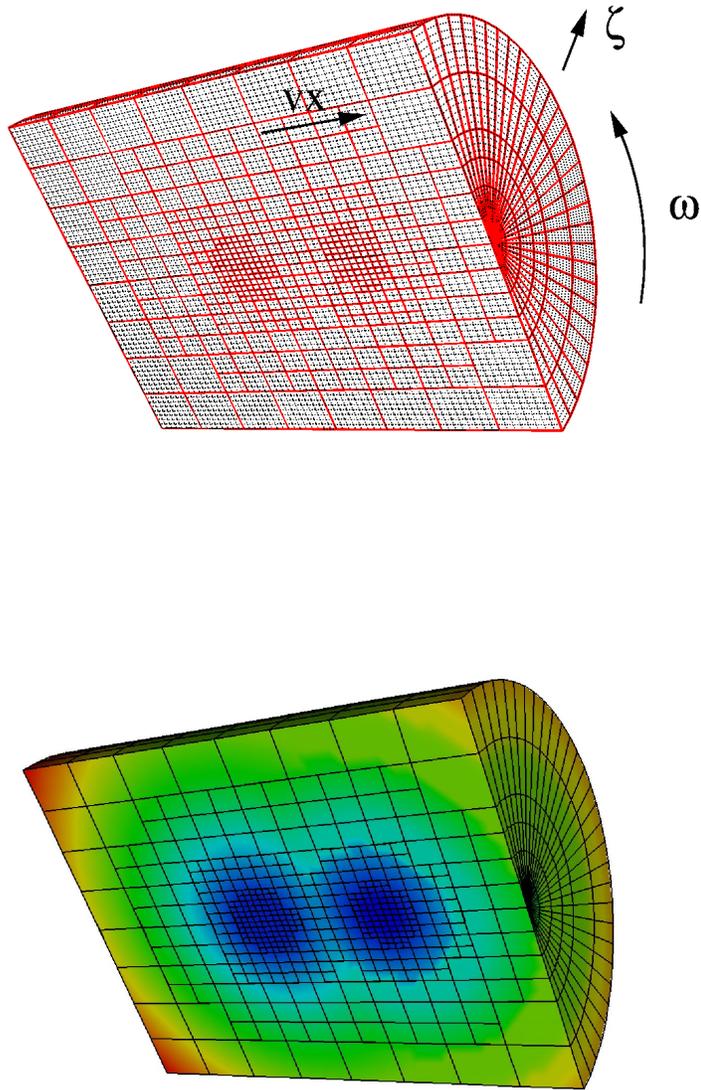


Figure 12: 2D axisymmetric flow around a sphere. Top: uniform cylindrical velocity grid (dashed lines) and AMR velocity grid (solid) with the cylindrical velocity coordinate system  $(v_x, \zeta, \omega)$ . Bottom: The same cylindrical AMR velocity grid with the contours of the support function based on the CNS pre-simulation.

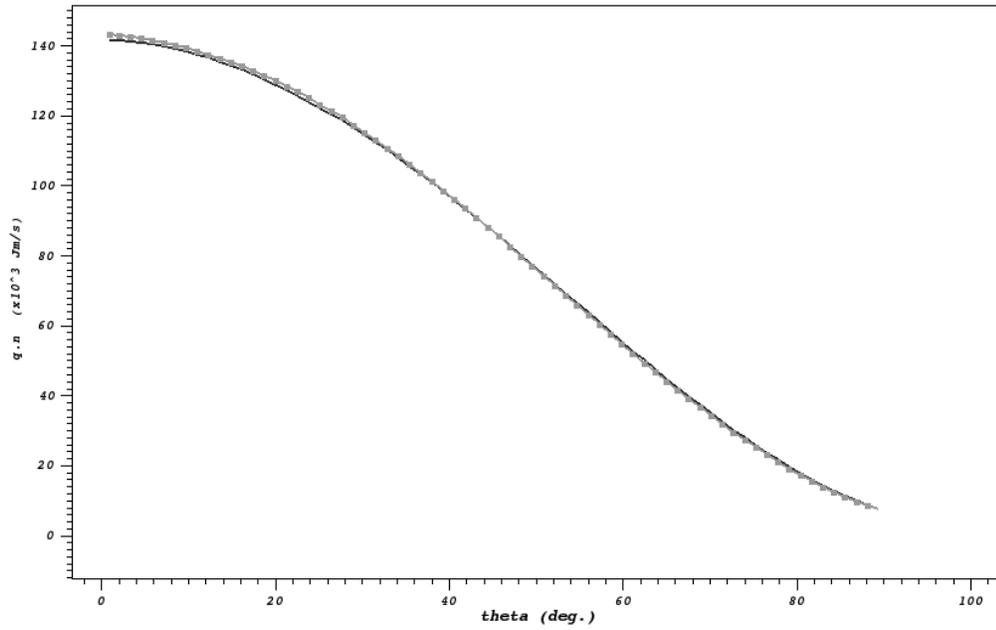


Figure 13: 2D axisymmetric flow around a sphere: comparison of the component of the heat flux normal to the surface: fluxes obtained with the fine uniform velocity grid (dot) and the AMR velocity grid (solid)

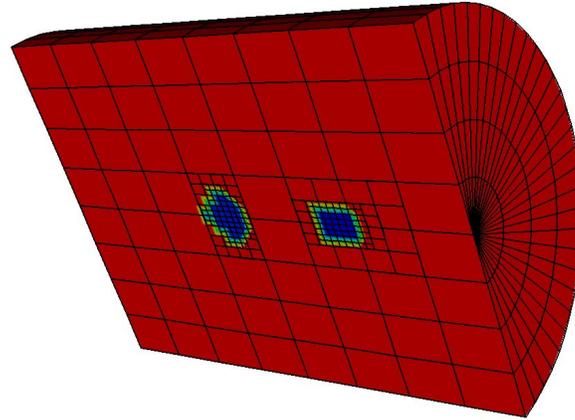


Figure 14: 2D axisymmetric flow around a sphere: cylindrical AMR velocity grid computed with Rankine-Hugoniot relations rather than with the CNS pre-simulation, with contours of the corresponding support function. Note that the refined zones are much smaller than in the previous grid, and that the contours of the support function are different (see figure 12)

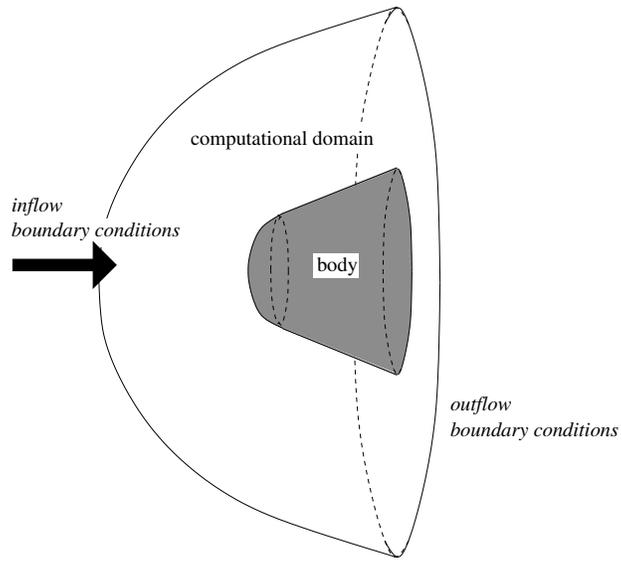


Figure 15: 3D flow around a cone: geometry and computational domain. The downstream flow is not simulated.

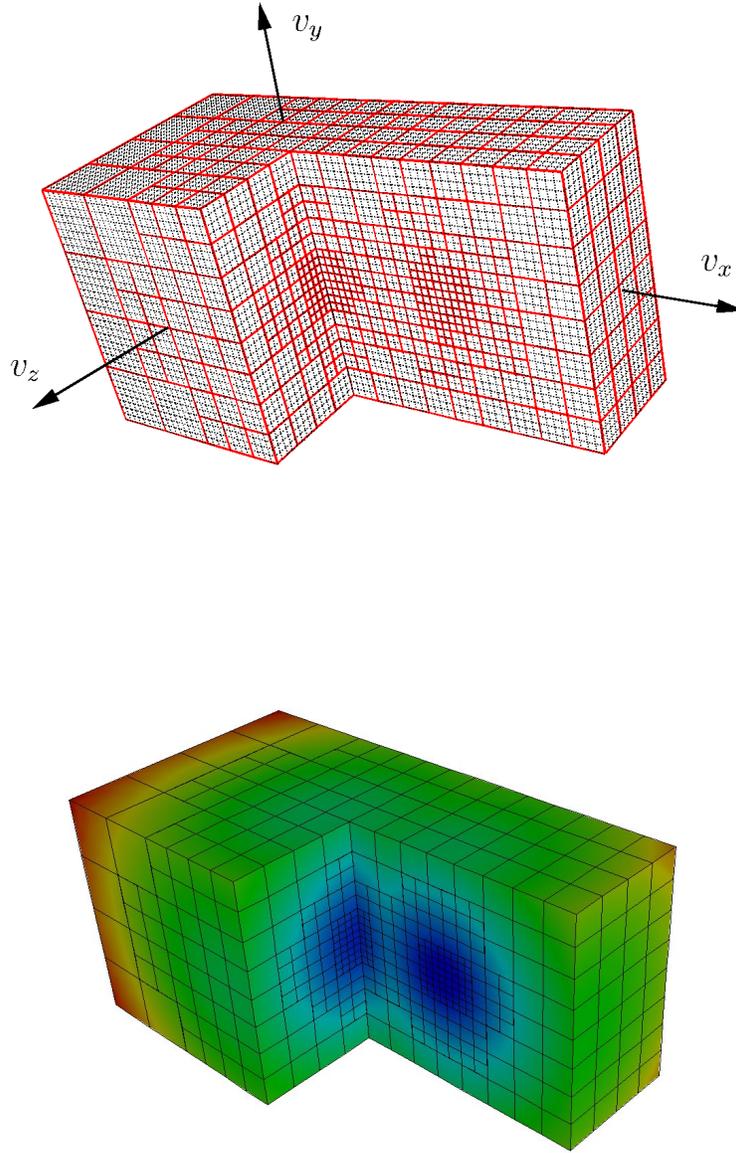


Figure 16: 3D flow around a cone. Top: the 3D Cartesian velocity grid (dashed line) and AMR velocity grid (solid) with the Cartesian coordinate system  $(v_x, v_y, v_z)$ . In order to see the refined zones around  $(0, 0, 0)$  and  $(u_{upstream}, 0, 0)$  that are inside the grid, a part of the grid has been hidden. Bottom: The same AMR velocity grid with the contours of the corresponding support function based on the CNS pre-simulation.

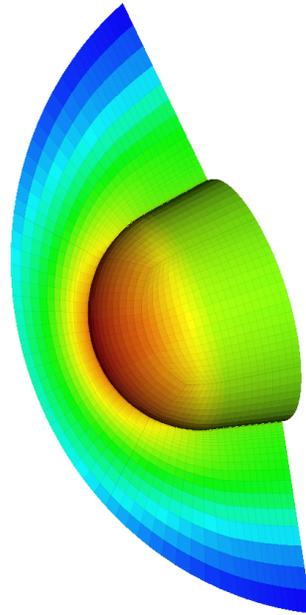


Figure 17: 3D flow around a cone: pressure field computed with the kinetic code and the AMR velocity grid. The pressure is shown on the surface of the solid, as well as in a vertical plane  $(x, y, 0)$  that crosses the cone along its middle part.

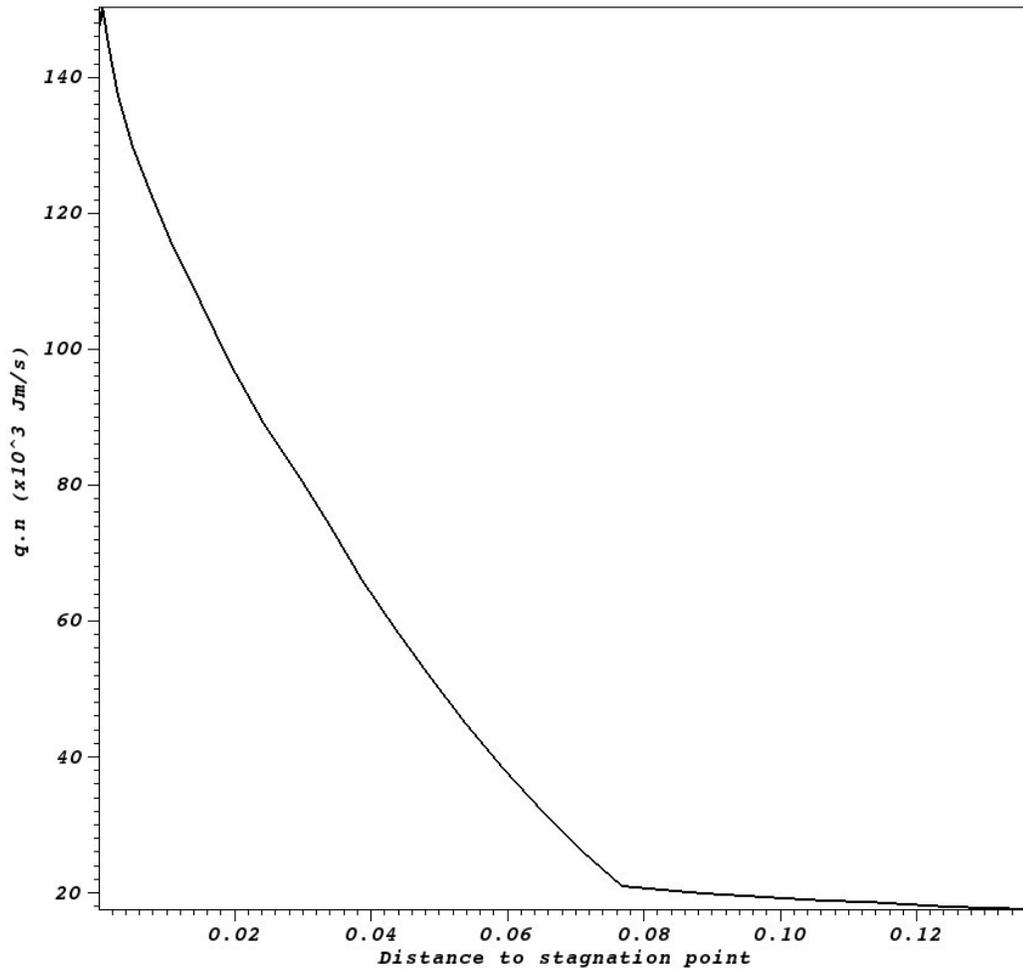


Figure 18: 3D flow around a cone: normal heat flux along the boundary (computed with the kinetic code and the AMR velocity grid).