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Uncertainty quantification in rarefied dynamics of ² molecular gas: rate effect of thermal relaxation

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The thermal conductivity of a molecular gas consists of the translational and internal parts; 8 although in continuum flows the total thermal conductivity itself is adequate to describe the heat 9 transfer, in rarefied gas flows they need to be modeled separately, according to the relaxation 10 rates of translational and internal heat fluxes in homogeneous system. This paper is dedicated to 11 quantifying how these relaxation rates affect rarefied gas dynamics. The kinetic model of Wu et 12 al. (J. Fluid Mech., vol. 763, 2015, pp. 24-50) is adapted to recover the relaxation of heat fluxes, 13 which is validated by the direct simulation Monte Carlo method. Then the Wu et al. model, 14 having the freedom to adjust the relaxation rates, is used to investigate the rate effects of thermal 15 relaxation in problems such as the normal shock wave, creep flow driven by Maxwell's demon, 16 and thermal transpiration. It is found that the relaxation rates of heat flux affect rarefied gas flows 17 significantly, even when the total thermal conductivity is fixed. 18

19 **1. Introduction**

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When the ratio between the molecular mean free path and the characteristic flow length becomes 20 appreciable, the Navier-Stokes-Fourier equations fail to describe the rarefied gas dynamics and 21 the gas kinetic equation is used instead. For monatomic gas, the Boltzmann equation and the 22 direct simulation Monte Carlo (DSMC) method provide equivalent and successful predictions 23 of rarefied gas dynamics (Bird 1994; Wagner 1992). For molecular gas, however, the internal 24 energy (due to the excitation of rotational, vibrational, or electronic degrees of freedom) other 25 than translational energy exists, making the collision dynamics much more complicated than 26 that of monatomic gas. Wang-Chang & Uhlenbeck (1951) extended the Boltzmann equation by 27 treating the internal degree of freedom quantum mechanically and assigning each internal energy 28 level an individual velocity distribution function. However, it is obvious that the analytical and 29 numerical methods for Wang-Chang & Uhlenbeck equation become difficult and expensive. For 30 example, Tcheremissine & Agarwal (2008) found that in hypersonic flow the computational cost 31 for molecular gas are two orders of magnitude higher than that for monatomic gas. 32

³³ Compared to the dilute monatomic gas, a unique feature of the molecular gas is that it exchange ³⁴ the translational and internal energies during binary collisions. On an averaging sense, in spatial-³⁵ homogeneous systems the relaxation of rotational temperature T_{rot} (for simplicity we assume ³⁶ the molecule has rotational mode excited only, and the rotational degree of freedom is $d_r = 2$ ³⁷ for diatomic and linear molecule, and 3 for all other non-linear molecules) is described by the ³⁸ Jeans-Landau equation

$$\frac{\partial T_{rot}}{\partial t} = \frac{p_{tr}}{\mu} \frac{T - T_{rot}}{Z},\tag{1.1}$$

where t is the time, p_{tr} is the kinetic pressure, μ is the shear viscosity of the gas, T is the total

- $_{40}$ temperature, and Z is the rotational collision number. On the other hand, the relaxation of the
- translational and rotational heat fluxes (q_{tr} and q_{rot} , respectively) are found to satisfy (Mason &
- 42 Monchick 1962; McCormack 1968):

$$\frac{\partial}{\partial t} \begin{bmatrix} \boldsymbol{q}_{tr} \\ \boldsymbol{q}_{rot} \end{bmatrix} = -\frac{p_{tr}}{\mu} \begin{bmatrix} A_{tt} & A_{tr} \\ A_{rt} & A_{rr} \end{bmatrix} \begin{bmatrix} \boldsymbol{q}_{tr} \\ \boldsymbol{q}_{rot} \end{bmatrix}, \qquad (1.2)$$

where the matrix of relaxation rates $\mathbf{A} = [A_{ij}]$ with i, j = t, r determines the translational and internal thermal conductivities, see § 2.2 below. From the physical point of view, the matrix should have two positive eigenvalues.

The DSMC has become the prevailing method to simulate the rarefied dynamics of molecular 46 gases, by using the phenomenological Borgnakke & Larsen (1975) collision model. While the 47 success of DSMC in modeling monatomic gas dynamics lies in its recovery of viscosity and 48 thermal conductivity, and the accurate update of post-collision velocities as per Boltzmann 49 collision operator, the simulation of molecular gas flow in DSMC is not perfect. That is, in DSMC 50 the attention is only paid to realize the correct exchange rate between the translational and internal 51 energies (1.1), which guarantees the exact recovery of bulk viscosity (Boyd 1991; Haas *et al.* 1994; 52 Gimelshein et al. 2002). However, it cannot always recover the thermal conductivity (Wu et al. 53 2020), either the total value or its translational and internal components. So far, the consequence 54 of this overlooked problem remains unknown, as to our knowledge no one has considered (or 55 there is no mechanism to recover) the relaxation of heat fluxes (1.2) in DSMC, which determines 56 the thermal conductivity of gas. 57 The relaxation rates play important roles in the gas dynamics (Candler 2018). Although in 58 DSMC and other kinetic models (Morse 1964; Holway 1966; Rykov 1975; Gorji & Jenny 2013; 59

Wu *et al.* 2015; ?), the effect of temperature relaxation (1.1), or equivalently the bulk viscosity, 60 has been extensively studied, e.g. by Frezzotti & Ytrehus (2006), ?, and Kosuge & Aoki (2018), 61 the role of thermal relaxation of heat fluxes (1.2) has seldom been investigated. In experiments, 62 the total thermal conductivity can be measured straightforwardly, and sometimes its translational 63 part (Mason 1963; Gupta & Storvick 1970; Porodnov et al. 1978; Wu et al. 2020) can also be 64 measured; we will show in the following section that, there are still at least two elements in the 65 thermal relaxation rates of heat fluxes A not determined. Therefore, it is the aim of the present 66 work to quantify these uncertainties caused by the variation of A in rarefied gas flows, although 67 they rarely affect the continuum flow described by the Navier-Stokes-Fourier equations when the 68 shear viscosity, bulk viscosity and total thermal conductivity are fixed. 69

To fulfill this goal, a kinetic model which is able to recover the relaxation rates in (1.1) and (1.2)70 is urgently needed. In this paper, the Wu et al. (2015) model is firstly introduced, which is then 71 modified to include the general relaxations for both temperatures and heat fluxes. The modified 72 model is validated by DSMC when both models have the same relaxation rates. Finally, the new 73 kinetic model is used to study the influence of thermal relaxation rates in rarefied gas flows, by 74 keeping other parameters unchanged. Note that here we do not use DSMC because when the shear 75 viscosity, bulk viscosity and Schmidt number (i.e., Sc = $\mu/\rho D$, where ρ is the mass density and 76 D is the diffusion coefficient) are fixed, the matrix A in DSMC is fixed, but the resulting thermal 77 conductivities may not be equal to the experimentally measured values (Wu et al. 2020), not to 78 mention its translational and internal components. 79

2. Thermal relaxation and transport coefficients

The essential difference between monatomic and molecular gases is that molecules exhibit internal relaxation that exchanges the translational and internal energies, which lead to several new transport coefficients including the bulk viscosity and internal thermal conductivity. For simplicity, we consider the case where only rotational modes are activated and treated in the way

⁸⁵ of classical mechanics.

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2.1. Bulk viscosity

In dilute gas, the exchange of translational and internal energy through inelastic collisions leads to a resistance in the compression or expansion of gas, which is quantified by the bulk viscosity μ_b . According to the Chapman & Cowling (1970) expansion, when the relaxation time $Z\mu/p_{tr}$ between the translational and rotational energies is much shorter than the characteristic time of gas flow, the bulk viscosity is expressed as:

$$\mu_b = \frac{2d_r Z}{3(d_r + 3)}\mu.$$
 (2.1)

The most widely used phenomenological model for molecular gas in DSMC is the Borgnakke & Larsen (1975) model, in which the relaxation rate is controlled by making a fraction of collisions inelastic. And this fraction gives the inverse of rotational collision number in DSMC, denoted as Z_{DSMC} . Note that when the variable-soft-sphere model is used in DSMC, Z_{DSMC} is related to the rotational collision number Z in (1.1) as

$$Z = \frac{\alpha(5 - 2\omega)(7 - 2\omega)}{5(\alpha + 1)(\alpha + 2)} Z_{\text{DSMC}},$$
(2.2)

where ω is the viscosity index such that $\mu(T) = \mu(T_0) (T/T_0)^{\omega}$, T_0 is the reference temperature,

 $_{98}$ and α is the parameter that determines the scattering angle after binary collision; it can be chosen

⁹⁹ freely, but in the variable-soft-sphere model it is usually determined by the Schmidt number (in

¹⁰⁰ order to simulate the diffusion process) through the following equation (Bird 1994):

$$Sc = \frac{5(2+\alpha)}{3(7-2\omega)\alpha}.$$
(2.3)

In other words, the bulk viscosity of the molecular gas can be exactly recovered by adjusting the value of Z_{DSMC} in DSMC simulations.

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2.2. Thermal conductivity

¹⁰⁴ Compared to the monatomic gas, the thermal relaxations not only reduce the value of ¹⁰⁵ translational thermal conductivity κ_{tr} , but also result in the rotational thermal conductivity κ_{rot} . ¹⁰⁶ According to the Chapman & Cowling (1970) expansion, the translational and rotational thermal ¹⁰⁷ conductivities satisfy (Mason & Monchick 1962)

$$\begin{bmatrix} \kappa_{tr} \\ \kappa_{rot} \end{bmatrix} = \frac{k_B \mu}{2m} \begin{bmatrix} A_{tt} & A_{tr} \\ A_{rt} & A_{rr} \end{bmatrix}^{-1} \begin{bmatrix} 5 \\ d_r \end{bmatrix},$$
 (2.4)

where k_B is the Boltzmann constant, and *m* is the molecular mass.

It will be convenient to express the thermal conductivity κ of a molecular gas in terms of the dimensionless Eucken (1913) factors:

$$\frac{\kappa m}{\mu k_B} = \frac{3}{2} f_{tr} + \frac{d_r}{2} f_{rot} = \frac{3+d_r}{2} f_u, \qquad (2.5)$$

where f_u is the total Eucken factor, while f_{tr} and f_{rot} are the translational and internal Eucken factors, respectively:

$$f_{tr} = \frac{2}{3} \frac{m\kappa_{tr}}{k_B \mu}, \quad f_{rot} = \frac{2}{d_r} \frac{m\kappa_{rot}}{k_B \mu}.$$
 (2.6)

From (2.4) and (2.6), it is clear that the Eucken factors are determined by the four relaxation

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rates in the matrix **A**. However, the values of these relaxation rates are difficult to be obtained experimentally. For monatomic gas, $A_{tr} = A_{rt} = A_{rr} = 0$ and $A_{tt} = 2/3$, so the translational Eucken factor is 2.5. In molecular gas, the energy exchange between translational and rotational energy makes the off-diagonal components A_{tr} and A_{rt} negative, which leads to a translational Eucken factor f_{tr} lower than 2.5.

In DSMC, as the only parameter modifying the energy exchange between different energy modes, the collision number Z determines the values of relaxation rates **A** (and hence the thermal conductivities). Considering the discussion in § 2.1, both bulk viscosity and thermal conductivity of molecular gas are determined by Z, so that they cannot be adjusted independently in DSMC. Therefore, generally speaking, these two transport coefficients cannot be matched to the experimental values simultaneously in the conventional DSMC method with Borgnakke-Larsen model.

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2.3. Extraction of thermal relaxation rates in DSMC

Since DSMC does not allow free adjustment of **A** but only the collision number Z_{DSMC} , here we extract the relaxation rates **A** by varying Z_{DSMC} . To this end, we consider both nitrogen and hydrogen chloride, which have only classical rotational motions excited (with $d_r = 2$) at room temperature.

We extract the thermal relaxation rates A in the spatial-homogeneous relaxation problem: 10^{6} 131 simulation particles are generated over a cubic cell of the size (10 nm)³, where periodic condition 132 is employed at all the boundaries. The gas density is $n_0 = 2.69 \times 10^{25} \text{m}^{-3}$ and the temperature is 133 $T_0 = 300$ K. At the beginning of DSMC simulation, simulation particles with positive velocity 134 in the x direction are generated from the Maxwell velocity distribution of T = 200 K, while the 135 rest are generated from Maxwell velocity distribution of T = 400 K, see figure 1(a); similarly, the 136 rotational energy assigned to the particles with $v_x > 0$ is generated from the Maxwell distribution 137 of T = 200 K, while those moving to the opposite direction obey the Maxwell distribution of 138 T = 400 K, see figure 1(b). In this manner we generate an initial velocity and energy distribution 139 which leads to initial non-zero values of translational and rotational heat fluxes. Then, the system 140 with prescribed initial heat fluxes evolves with respect to the time, and both the translational and 141 rotational heat fluxes are monitored until the entire system reaches thermal equilibrium. Both 142 nitrogen and hydrogen chloride are simulated to extract the relaxation rates A with the variable-143 soft-sphere molecular collision model and the corresponding parameters are listed in table 1. 100 144 independent runs were conducted to get smooth results. 145

Parameters	N_2	HC1
Molecular mass: $m (\times 10^{26} \text{kg})$	4.65	6.14
Viscosity index: ω	0.74	1.0
Diameter: $d (\times 10^{10} \text{m})$	4.11	5.59
Schmidt number: Sc	1/1.34	1/1.33
Scattering parameter: α	1.36	1.59
Simulation time step ($\times 10^{12} \text{s}$):	4.74	5.45

Table 1: Parameters in the variable-soft-sphere model of DSMC, for N₂ and HCl, which are collected from tables A1 and A3 in the book of Bird (1994). In this case and all the following cases, the simulation time step is chosen to be one fifth of the minimum cell size divided by the most probable speed $\sqrt{2k_BT_0/m}$.



Figure 1: The initial distribution of (a) molecular velocity and (b) rotational energy of nitrogen molecules in DSMC (the open-source code SPARTA is used), where the abscissas are normalized by $\sqrt{2k_BT_0/m}$ and k_BT_0 , respectively. (c, d) The evolution of heat fluxes and their time derivatives, circles in (d) represent the numerical fitting used to extract the relaxation rates **A** from DSMC. (e – h) Extracted **A** from DSMC for nitrogen (squares) and hydrogen chloride (circles). DSMC simulation parameters are summarized in table 1.

Figure 1(c, d) plots the evolution of the translational and rotational heat fluxes and their time derivatives for nitrogen with $Z_{\text{DSMC}} = 4.0$. It can be seen that the time derivative of the translational heat flux is significantly increased, due to its strong coupling with the rotational heat flux: from (1.2) it can be inferred that A_{tr} is negative. That the time derivative of rotational heat flux decreases monotonically with respect to the time implies that $|A_{rt}|$ is very small if A_{rt} is negative.

We adopt the least square method to solve the linear regression problem (1.2) to extract 152 the relaxation rates \mathbf{A} , and the results in figure 1(e-h) show that these parameters exhibit linear 153 dependence with $1/Z_{\text{DSMC}}$. When the collision number Z_{DSMC} is increased, the energy exchange 154 between translational and internal motions vanishes gradually, hence the relaxation rates A_{tr} and 155 A_{rt} approach zero, while A_{tt} and A_{rr} approach 2/3 and Sc, respectively. According to (2.4), 156 that A_{rr} approaches Sc means that the translational thermal conductivity is proportional to the 157 diffusion coefficient. This is comprehensible because the diffusion of gas molecules transports 158 the heat. 159

Given the thermal relaxation rate A, the Eucken factors can be calculated by (2.4) and (2.6). 160 We find that in order to match the experimental thermal conductivity (or equivalently $f_u = 1.993$) 161 of nitrogen at $T_0 = 300$ K, the collision number has to be chosen as $Z_{\text{DSMC}} = 4.0$, and the 162 corresponding relaxation rates are $A_{tt} = 0.786$, $A_{tr} = -0.201$, $A_{rt} = -0.059$, $A_{rr} = 0.842$; hence 163 we have $f_{tr} = 2.365$ and $f_{rot} = 1.435$. Note that this value of Z_{DSMC} may not lead to the correct 164 value of bulk viscosity. However, for hydrogen chloride, no matter what the value of Z_{DSMC} is, the 165 calculated total thermal conductivity from DSMC can never recover the experimental value (Wu 166 et al. 2020). This is the problem of DSMC which, generally speaking, cannot recover the bulk 167 viscosity and translational/internal thermal conductivity of molecular gas simultaneously in the 168 phenomenological Larsen-Borgnakke collision model. 169

3. The modified Wu model and its validation

¹⁷¹ Due to the limitation of DSMC method, a kinetic model is desired and developed in this section, ¹⁷² which allows free adjustment of relaxation rates (and hence free adjustment of bulk viscosity and 6

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translational/internal thermal conductivities). To this end, we modify the Wu *et al.* (2015) model so that it can reflect the general relaxations for temperature and heat flux. Then, we validate the accuracy of the proposed model by comparing its solutions for the normal shock wave and creep flow driven by Maxwell's demon with the DSMC results. In order to make consistent comparison, the relaxation rates in the modified Wu *et al.* (2020) should be the same as those in the DSMC simulations.

3.1. The modified kinetic model

Like the Wang-Chang & Uhlenbeck (1951) equation, all the kinetic models divide the binary 180 collision into the elastic and inelastic collisions. The elastic collision conserves the translational 181 energy, while the inelastic collision exchanges the translational and rotational energies. The 182 linearized kinetic model for molecular gas is developed by Hanson & Morse (1967), while one 183 of the practical models for nonlinear flows is proposed by Rykov (1975). As an extension of the 184 Rykov model, the kinetic model equation developed by Wu et al. (2015) also treats the elastic 185 and inelastic collision separately. While in order to improve the modeling accuracy, the Wu et al. 186 model replaces the elastic collision operator in the Rykov model with the Boltzmann collision 187 operator for monatomic gas, and thus introduces a more realistic elastic collision relaxation time 188 that is dependent on the molecular velocity (i.e., in the limit without translational-internal energy 189 exchange, it is reduced to the Boltzmann equation for monatomic gas). 190

In the original Wu *et al.* (2015) model, two velocity distribution functions, $G(\mathbf{x}, \mathbf{v}, t)$ and $R(\mathbf{x}, \mathbf{v}, t)$, where \mathbf{x} and \mathbf{v} are respectively the spatial coordinates and molecular velocity, are used to describe the translational and rotational motions of gas molecules; their evolution are governed by the following kinetic equations:

$$\frac{\partial G}{\partial t} + \mathbf{v} \cdot \frac{\partial G}{\partial \mathbf{x}} + \mathbf{a} \cdot \frac{\partial G}{\partial \mathbf{v}} = Q(G) + \frac{G_{rot} - G_{tr}}{Z\tau},$$

$$\frac{\partial R}{\partial t} + \mathbf{v} \cdot \frac{\partial R}{\partial \mathbf{x}} + \mathbf{a} \cdot \frac{\partial R}{\partial \mathbf{v}} = \frac{R_{tr} - R}{\tau} + \frac{R_{rot} - R_{tr}}{Z\tau},$$
(3.1)

where *a* is the external acceleration, $\tau = \mu/p_{tr}$ is the characteristic collision time related to the translational motion of gas molecules, and *Q*(*G*) is the Boltzmann collision operator for monatomic gases (Wu *et al.* 2013, 2014). The four reference distribution functions *G*_{tr}, *G*_{rot}, *R*_{tr} and *R*_{rot} are modeled as

$$G_{tr} = n \left(\frac{m}{2\pi k_B T_{tr}}\right)^{3/2} \exp\left(-\frac{mc^2}{2k_B T_{tr}}\right) \left[1 + \frac{2mq_0 \cdot c}{15k_B T_{tr} p_{tr}} \left(\frac{mc^2}{2k_B T_{tr}} - \frac{5}{2}\right)\right],$$

$$G_{rot} = n \left(\frac{m}{2\pi k_B T}\right)^{3/2} \exp\left(-\frac{mc^2}{2k_B T}\right) \left[1 + \frac{2mq'_0 \cdot c}{15k_B T p} \left(\frac{mc^2}{2k_B T} - \frac{5}{2}\right)\right],$$

$$R_{tr} = \frac{d_r k_B T_{rot}}{2} G_{tr} + \left(\frac{m}{2\pi k_B T_{tr}}\right)^{3/2} \exp\left(-\frac{mc^2}{2k_B T_{tr}}\right) \frac{mq_1 \cdot c}{k_B T_{tr}},$$

$$R_{rot} = \frac{d_r k_B T}{2} G_{rot} + \left(\frac{m}{2\pi k_B T}\right)^{3/2} \exp\left(-\frac{mc^2}{2k_B T}\right) \frac{mq'_1 \cdot c}{k_B T},$$
(3.2)

where c = v - U is the peculiar velocity, and

$$q_{0} = q_{tr}, \quad q'_{0} = \omega_{0}q_{tr}, q_{1} = (1 - Sc)q_{rot}, \quad q'_{1} = (1 - Sc)\omega_{1}q_{rot},$$
(3.3)

where ω_0 and ω_1 are the constants to recover both the translational and rotational thermal conductivity coefficients of molecular gases. Further, the macroscopic quantities, number density n, flow velocity U, translational temperature T_{tr} , rotational temperature T_{rot} , translational heat flux q_{tr} , the rotational heat flux q_{rot} , and pressure tensor p_{ij} are calculated from the velocity moments of the two distribution functions *G* and *R*:

$$n = \int G d\mathbf{v}, \quad \mathbf{U} = \frac{1}{n} \int G \mathbf{v} d\mathbf{v},$$

$$T_{tr} = \frac{1}{3nk_B} \int mGc^2 d\mathbf{v}, \quad T_{rot} = \frac{2}{d_r nk_B} \int R d\mathbf{v},$$

$$q_{tr} = \frac{1}{2} \int mGc^2 c d\mathbf{v}, \quad q_{rot} = \int Rc d\mathbf{v}, \quad p_{ij} = \int mGc_i c_j d\mathbf{v}.$$
(3.4)

The total temperature *T*, total pressure *p* and its translational counterpart are $T = (3T_{tr} + d_r T_{rot})/(3+d_r)$, $p = nk_B T$ and $p_{tr} = nk_B T_{tr}$, respectively. It can be verified that (1.1) and (1.2) are satisfied in the kinetic model.

²⁰⁸ Considering the general expression of thermal conductivity coefficients (or Eucken factors ²⁰⁹ equivalently) based on equations (2.4) and (2.6), there are still two unknown values in relaxation ²¹⁰ rates **A** even when both f_{tr} and f_{rot} have been fixed. It implies that the coefficients ω_0 and ω_1 ²¹¹ in the kinetic model above, which is determined by the thermal conductivities, may not able to ²¹² give fully recovery of all the transport information in molecular gases. Therefore, we modify the ²¹³ kinetic model by incorporating the relaxation rates **A** into the reference distribution functions as:

$$q_{0} = q_{tr}, \quad q'_{0} = \left[-3Z(A_{tt} - \frac{2}{3}) + 1\right] q_{tr} - 3ZA_{tr}q_{rot},$$

$$q_{1} = 0, \quad q'_{1} = -Z\left[A_{rt}q_{tr} + (A_{rr} - 1)q_{rot}\right],$$
(3.5)

so that (1.1) and (1.2) are exactly recovered.

3.2. Numerical validation

Now we assess the accuracy of the kinetic model (3.1) with (3.2) and (3.5), by comparing its numerical solutions of normal shock wave and thermal creep flow in nitrogen with the DSMC results. To make fair comparisons, the relaxation rates **A** are equal to those extracted from the DSMC. Therefore, the collision number and relaxation rates take the values determined in §2.3, and the rotational collision number in (3.1) is Z = 2.6671 according to (2.2).

The obtained macroscopic flow quantities will be shown in non-dimensional values: the number density, temperature, spatial coordinate, velocity, pressure, and heat flux are normalized by $n_0 = 2.69 \times 10^{25} \text{ m}^{-3}$, $T_0 = 300 \text{ K}$, the characteristic length L_0 , the most probable speed $v_m = \sqrt{2k_BT_0/m}$, $n_0k_BT_0$, and $n_0k_BT_0v_m$, respectively. The Knudsen number is defined as

$$Kn = \frac{\mu(T_0)}{n_0 L_0} \sqrt{\frac{\pi}{2mk_B T_0}}.$$
(3.6)

3.2.1. Normal shock wave

First, we consider the normal shock wave when the Mach number is Ma = 4 and the upstream 226 mean free path ($L_0 = 59.59$ nm) is chosen as the characteristic length. The simulation domain 227 used in both the kinetic model and DSMC are $30L_0$ in x direction with the wavefront in the 228 center of it, so that the equilibrium states determined by the Rankine-Hugoniot relation can be applied at both ends of the domain. The kinetic model equation is solved by discretize velocity 230 method with fast spectral method dealing with its Boltzmann collision term (Wu et al. 2015). The 231 entire domain is divided into 150 non-uniform cells, with more cells located around the shock 232 center. And $48 \times 32 \times 32$ discrete velocities, which are uniformly distributed within the range 233 $[-7.5v_m, 7.5v_m]$, are used. In the DSMC simulation, 360 uniform spatial cells with size of 5 234 nm are applied, and there are 7.2×10^5 simulation particles in the whole computational domain. 235 When the steady state is reached, by doing time average over 2500 sampling steps, we get the 236

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Figure 2: Comparison between the DSMC (circles) and the modified Wu model (lines) for normal shock wave in nitrogen with Ma = 4. The macroscopic quantity $Q = \rho, u, T$ is normalized by $(Q-Q_u)/(Q_d-Q_u)$, where the subscripts u and d represent the upstream and downstream, respectively. Note that the shock wave is shifted so that the density at x = 0 is $(\rho_d + \rho_u)/2$; and other profiles are shifted accordingly.

final results as the reference for comparison. The accuracy is guaranteed because the cell size is 237 much smaller than the molecular mean free path and there are about 2000 simulation particles 238 per cell.

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Figure 2 compares the structures of the normal shock wave obtained from the kinetic model 240 and the DSMC simulation. Good agreement in macroscopic quantities demonstrates the accuracy 241 of the proposed kinetic model. 242

3.2.2. Creep flow driven by the Maxwell demon 243

Second, we consider the microflow. In the thermal creep along an infinite channel, the gas flow 244 is driven by a temperature gradient at the wall, which is equivalent to applying a small external 245 acceleration. Here, as a thought test, we consider the creep flow driven by the Maxwell demon, 246 where each molecule is subject to an external acceleration based on its kinetic energy: 247

$$a_y = a_0 \left(\frac{v^2}{v_m^2} - \frac{3}{2} \right)$$
(3.7)

see figure 3. It can be seen that the direction of the acceleration is determined by the magnitude 248 of molecular velocity. We solve this creep flow in a one dimensional domain, which is bounded 249 by two parallel walls with fully diffuse boundary condition at the same temperature. Here, 250 the characteristic length L_0 is the distance between the walls and a_0 is a small value set by 251 $2a_0L_0/v_m^2 = 0.0718$ to guarantee that the gas flow deviates slightly from the global equilibrium. 252 The modified Wu model is solved by the general synthetic iterative scheme (Su et al. 2021; 253 ?). There are 100 spatial cells inside the computational domain, with more cells located in the 254 vicinity of solid walls to capture the Knudsen layer structure. And $48 \times 48 \times 48$ non-uniformly 255 distributed discrete velocities within the range $[-6v_m, 6v_m]$ are applied, with dense velocity grids 256 around zero velocity to capture the discontinuity of velocity distribution function therein. In the 257 DSMC simulations, there are 100 uniform cells and 2×10^4 simulation particles between two 258 walls, and both time and ensemble averaging are used which include 10 independent runs with 259 2.5×10^6 sampling times for each one. 260

The results of kinetic model and DSMC are compared in figure 3, for typical Knudsen numbers. 261 It is observed that the flow velocity and heat fluxes obtained from the kinetic model are in good 262 agreement with those from the DSMC. Besides, the rotational heat flux is negligibly small, when 263 compared to the translational heat flux. This implies that the translational thermal conductivity 264 plays the dominated role in the flow velocity in this problem. 265



Figure 3: Comparison between the DSMC (markers) and the modified Wu model (lines) in the creep flow driven by the Maxwell demon. The velocity and heat flux are further normalized by the dimensionless acceleration 0.0718.

4. Uncertainty quantification: rate effect of thermal relaxation

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It can be learned from (2.4) that, even when the translational and rotational thermal conduc-267 tivities (i.e., κ_{tr} and κ_{rot}) are determined, two elements in the matrix A remains unknown; and 268 there will be three undetermined elements if only the total thermal conductivity is known as in 269 many experiments. Here we investigate the effects of these uncertain values based on the modified 270 Wu model, as the DSMC does not have the capability to adjust the thermal relaxation rates once 271 the rotational collision number and Schmidt number are fixed. The uncertainties in rarefied gas 272 flows will be quantified in the following two ways. First, we vary the values of A_{ij} when the 273 translational and rotational Eucken factors (i.e., f_{tr} and f_{rot}) are given. Second, we fix the total 274 Eucken factor f_u , A_{tr} and A_{rt} , but vary the translational and rotational Eucken factors. 275

4.1. Normal shock wave

When f_{tr} and f_{rot} are fixed on top of the fixed shear viscosity and bulk viscosity, the gas 277 dynamics is uniquely determined in the continuum flow. However, different values of A_{ij} could 278 lead to different results in rarefied gas flows. The normal shock wave of nitrogen is firstly studied 279 to demonstrate this uncertainty. Specifically, A_{tr} and A_{rt} are selected to vary within [-5/6Z, 0]280 and [-1/3Z, 0], respectively, while A_{tt} and A_{rr} are determined according to (2.4) and (2.6) to 281 recover the assigned values of $f_{tr} = 2.365$ and $f_{rot} = 1.435$. Given Z = 2.6671, the considered 282 minimum values of A_{rt} and A_{tr} are -0.3124 and -0.1250, respectively, which are about $1 \sim 2$ 283 times larger, in magnitude, than those extracted from the DSMC simulation in §2.3. 284



Figure 4: Influence of the thermal relaxation rates in normal shock wave. The red solid lines are the results of the modified Wu model with **A** extracted from DSMC, while the blue shade regions show the results from the modified Wu model, with $A_{rt} \in [-0.3124, 0.0], A_{tr} \in [-0.1250, 0.0], f_{tr} = 2.365$ and $f_{rot} = 1.435$.



Figure 5: Influence of the translational Eucken factors in normal shock waves. All cases have the same total Eucken factor f_u , but the translational Eucken factor f_{tr} for the green dash-dot, blue dashed, and red solid lines are 1.5, 2, and 2.5, respectively; the rotational Eucken factor is changed accordingly to make f_u fixed. The modified Wu model is used.

Figure 4 shows the density, temperature, and heat flux in the normal shock wave of Mach 285 number Ma = 4, where the red solid lines illustrate the reference solutions with A extracted 286 from the DSMC, while blue shade regions show the divergences caused by the variations of A. It 287 can be seen that the variation of thermal relaxation rates slightly shifts the profiles of rotational 288 temperature and heat fluxes, mainly in the regions $x \in [-2, -1]$ and $x \in [0.5, 2]$. However, the 289 thermal relaxation rates has almost no influence on the profiles of density (hence velocity due to 290 mass conservation) and normal pressure (not shown here). Therefore, there is also little change 291 in the thickness of shock wave. 292

Now we consider different values of f_{tr} and f_{rot} , but fixed value of total thermal conductivity. 293 Figure 5 summaries the numerical results from the modified Wu model with $f_{tr} = 1.5, 2.0, 2.5,$ 294 while A_{tr} and A_{rt} take the values of -5/6Z and -1/3Z, respectively. Note that small values of f_{tr} 295 are possible, especially in polar gases where the translational Eucken factor can be much smaller 296 than 2.5, e.g., $f_{tr} = 1.78$ for water and $f_{tr} = 0.41$ for CH₃OH (Mason & Monchick 1962). 297 Significant discrepancies in macroscopic quantities with different values of f_{tr} are observed, 298 especially in the profiles of temperature. First, larger f_{tr} makes the translational temperature rise 299 earlier to its maximum value, and then decrease faster to the equilibrium value in downstream; 300 the same trend is also observed in the deviation pressure 301

$$P_{xx} = \frac{m}{2} \int \left(c_x^2 - \frac{c^2}{3}\right) G d\boldsymbol{\nu},\tag{4.1}$$



Figure 6: Influence of the thermal relaxation rates in the creep flow driven by the Maxwell demon. Red solid lines are the results with A obtained from the DSMC, the blue shade region shows the results from the modified Wu model, with $A_{rt} \in [-0.3124, 0.0]$ and $A_{tr} \in [-0.1250, 0.0]$. Other parameters are Kn = 0.2, $f_{tr} = 2.365$ and $f_{rot} = 1.435$.



Figure 7: Influence of the translational Eucken factor in the creep flow driven by the Maxwell demon. All cases have the same total Eucken factor f_u and Kn = 0.2, while the translational Eucken factor f_{tr} for the green dash-dot, blue dashed, and red solid lines are 1.5, 2.0, and 2.5, respectively. The modified Wu model is used.

and the magnitude of total heat flux. Second, the influence of Eucken factors on the rotational temperature, however, concentrates around the center of shock structure: lower f_{tr} and hence higher f_{rot} results in larger rotational temperature. Third, larger f_{tr} results in faster rise of density.

4.2. Creep flow driven by the Maxwell demon

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The same sets of values of **A** in normal shock wave cases are used here to study the influence on the velocity and heat flux in the creep flow driven by the Maxwell demon, and the results with Kn = 0.2 are shown in figure 6 when f_{tr} and f_{rot} are fixed. Contrary to the situations in normal shock wave, significant variation in the results with different relaxation rates **A** is observed: the maximum relative uncertainty is 16.7% and 17.6% for the velocity and translational heat flux, respectively. Meanwhile, it is seen that the uncertainty occurs in the middle part of the creep flow, while the velocity slip and heat flux in the vicinity of the wall rarely change.

To further investigate the influence of the translational Eucken factor, $f_{tr} = 1.5, 2.0, 2.5$ are considered in the modified Wu model with Kn = 0.2 and $f_u = 1.993$, while A_{tr} and A_{rt} take the values of -5/6Z and -1/3Z, respectively. As shown in figure 7, both the velocity and translational heat flux vary significantly with f_{tr} : the values of velocity and translational heat flux of $f_{tr} = 2.5$ are 68% larger than those of $f_{tr} = 1.5$. Contrast to the results in figure 6 where the velocity slip



Figure 8: Comparison between the DSMC and the modified Wu model in the thermal transpiration inside a closed cavity. (a) Horizontal velocity. Solid lines are results from the kinetic model, while dots are from the DSMC. (b) Normal pressure $P_{xx} = \frac{m}{2} \int c_x^2 G dv$ along y = 0.5. (c) Flow field in the lower half of the cavity; from top to bottom, the translational Eucken factors are $f_{tr} = 2.37, 2.0, 1.75$, respectively.

and heat flux around the solid wall do not change with fixed f_{tr} , figure 7 shows a significant dependence of the velocity and heat flux on f_{tr} , i.e., both velocity and heat flux on the walls increase with f_{tr} . Thus, it can be concluded that the translational Eucken factor f_{tr} plays a dominant role in this problem.

The importance of the translational Eucken factor in this problem can be understood as follows. 323 It can be seen from (1.2) that the elements A_{tr} and A_{rt} are related to the energy exchange between 324 the translational and rotational motions. In other words, when A_{tr} (or A_{rt}) is zero, the relaxation 325 of translational (or rotational) heat flux will not be affected by the other one. For instance, by 326 varying A, it is found that when $A_{rt} = 0$ the rotational heat flux is always zero. The reason is 327 that in the creep flow driven by the Maxwell demon, only translational energy is changed directly 328 by the external driving force, thus the rotational energy and flux are only affected via the energy 329 exchange, which are determined by A_{tr} , A_{rt} and Z. Since A_{rt} is very small compared to the other 330 three relaxation rates in the matrix **A**, $q_{rot} \approx 0$ and q_{tr} (or f_{tr}) is dominant. 331

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4.3. Thermal transpiration in a cavity

Thermal transpiration is a classical phenomenon that has many applications, such as the Knudsen pump (Vargo *et al.* 1999), where the mass flow and pressure difference are the quantities of interest. To study this problem, a two-dimensional cavity with an aspect ratio of 5 is considered. The temperature of the left and right walls are 200°C and 400°C, respectively, while the temperature of the horizontal walls increases linearly from 200°C to 400°C. Due to symmetry, only the lower half of the cavity is simulated, and the results of Kn = 0.5959 are shown in figure 8. At the initial stage, due to the thermal transpiration, the gas molecules move



Figure 9: Influence of the Knudsen number Kn in the creep flow driven by Maxwell's demon. All cases have the same f_u , while f_{tr} for green dash-dot, blue dashed, and red solid lines are 1.5, 2.0 and 2.5, respectively. And the Knudsen number are 0.001, 0.1 and 10 from the left column to right column, respectively.

towards the hot ends by the temperature gradient along the solid surfaces, which increases the pressure there. As a consequence, the pressure driven flow is formed in the opposite direction and several vortices are eventually generated in the steady state. Comparisons in the flow velocity and normal pressure in figure 8(a, b) support that the modified Wu model can give good agreement with DSMC simulations.

Similar to the one-dimensional creep flow, the flow fields are expected to be determined by f_{tr} other than f_{tt} in thermal transpiration, where both the normal pressure and the velocity magnitude increases with f_{tr} , see figure 8(b, c). Therefore, the mass flow rate follows the same trend. For the situations with small f_{tr} which may happens for some polar molecular gases, the flow pattern and flow rate could be very different from those of non-polar molecular and monatomic gases.

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4.4. Uncertainty in different flow regimes

In the above cases, the gas flows are in the transition regime, for example, Kn = 0.2 in the creep flow driven by the Maxwell demon. In this section we investigate the uncertainties of thermal relaxation rates when the gas flow is in the near continuum and free molecular regimes. To this end, Kn = 0.001, 0.1, 10 are considered for the case of creep flow driven by Maxwell's demon, and the translational Eucken factors are $f_{tr} = 1.5, 2.0, 2.5$ with $f_u = 1.993$. The thermal relaxation rates **A** are chosen in the same way as that in § 4.2.

Both the velocity and heat flux distribution are examined in figure 9. In the near continuum 357 regime with Kn = 0.001, the thickness of Knudsen layers becomes negligible and the velocity 358 and heat flux are uniformly distributed in the bulk regime. However, the difference caused by 359 different values of f_{tr} are still significant. Specifically, the magnitude of velocity and translational 360 heat flux increases 77.3% and 73.6% when f_{tr} is changed from 1.5 to 2.5. However, it should 361 be noted that, although the relative error is large, the variation of thermal relaxation rates in A is 362 not so important since the flow velocity and heat flux approaches zero along with the Knudsen 363 number. 364

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³⁶⁵ When Kn = 0.1, the variation of velocity and heat flux caused by different f_{tr} are approximately ³⁶⁶ the same as those of Kn = 0.001, which are 76.1% and 72.3% when f_{tr} changes from 1.5 to ³⁶⁷ 2.5. However, the magnitudes of these macroscopic quantities increase by 100 times, compared ³⁶⁸ to those when Kn = 0.001. This implies a roughly linear dependence of the Knudsen number. ³⁶⁹ It can be concluded that, the rarefaction effects disappear gradually when the system approaches ³⁷⁰ the continuum limit, while the relative uncertainty becomes even larger instead.

On the other hand, at large Knudsen number (e.g. Kn = 10), the magnitude of velocity and heat flux become even larger, but the relative uncertainty caused by the changing of f_{tr} reduces to 7.4% and 7.3% for the velocity and translational heat flux, respectively. This is comprehensible, since the effect from collisions between gas molecules is weaken when Kn approaches infinity. Therefore, the uncertainty caused by the thermal relaxation rates of collision becomes negligible at large Kn, though the rarefaction effect is more significant at this regime.

Based on these results, we conclude that the uncertainties in thermal relation rates are only important in the transition flow regime, where, roughly speaking, $0.01 \leq Kn \leq 10$.

379 **5.** Conclusions

In summary, the relaxation rates of translational and rotational heat fluxes play an important 380 role in rarefied flows of molecular gas. Since in experiment only the translational and rotational 381 thermal conductivities are measured (in most cases only the total thermal conductivity is known), 382 there are two (three) underdetermined coefficients. For the first time these uncertainties are 383 properly quantified in this paper. First, a kinetic model which is able to describe the relaxations of 384 energy and heat fluxes are designed. Second, the kinetic model is validated by the DSMC method 385 with the Borgnakke-Larsen collision rule, which can only reflect some fixed values of relaxation. 386 Finally, by varying the thermal relaxation rates in the modified Wu model, we have studied the 387 influence of thermal relaxation rates on the normal shock wave structures, the creep flow driven 388 by Maxwell's demon, and the thermal transpiration in a cavity. 389 This work demonstrates the importance to obtain exact values of thermal relaxation rates used

This work demonstrates the importance to obtain exact values of thermal relaxation rates used in the kinetic model for rarefied gas flow simulations, and to develop a better collision model in DSMC that is able to recover realistic relaxation rates. Research in this direction will help to build correct models for thermal conductivity of molecular gas, especially for molecular gas mixtures and non-equilibrium chemical reactions. In the future work, we plan to investigate whether the molecular dynamics simulation can be used to reduce or remove the uncertainties or not.

Declaration of interests

³⁹⁷ The authors report no conflict of interest.

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