



Quasi-State-Specific QCT Method for Calculating the Dissociation Rate of Nitrogen in Thermal Non-Equilibrium

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The dissociation of nitrogen was studied using a quasi-classical trajectory (QCT) analysis in the context of calculating the dissociation rate surface for a dense range of temperatures for use in computational fluid dynamics (CFD) applications. By sampling rovibrational states from a Boltzmann distribution but uniformly sampling the relative speed, the dissociation rate was calculated for translational and rovibrational temperatures between 8000 K and 20000 K. The justification for this approach was verified by analyzing different sampling techniques. It was found that uniformly sampling the relative speed increased the uncertainty of the thermally averaged dissociation rate, but the same QCT results could be used for a large range of temperatures. This is in contrast to Monte Carlo sampling techniques, where a new batch of trajectories must be simulated for each desired temperature. To generate the dissociation rate surface, 500 million trajectories were simulated, and the non-equilibrium rates were compared to other models and experimental data, generally showing good agreement.

Nomenclature

b	Impact parameter, Å
E_{rv}	Quantized rovibrational energy
f_g	Relative speed probability distribution function
f_r	Rotational state probability distribution function
f_{rv}	Rovibrational state probability distribution function
f_v	Vibrational state probability distribution function
g	Relative translational speed, Å/fs
g_s	Spin degeneracy
J	Rotational quantum number
k_B	Boltzmann constant, J/K
k_d	Thermal non-equilibrium dissociation rate, cm ³ /s/mol
k_{eq}	Thermal equilibrium dissociation rate, cm ³ /s/mol
k_{Park}	Park model dissociation rate, cm ³ /s/mol
N	Total number of trajectories
N_d	Number of dissociative trajectories
P_d	Probability of dissociation

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Q_{rv}	Rovibrational partition function
Q_v	Vibrational partition function
T	Temperature, K
T_t	Translational temperature, K
T_r	Rotational temperature, K
T_{rv}	Rovibrational temperature, K
T_v	Vibrational temperature, K
v	Vibrational quantum number
x	target function
xi_r	Reactant rotational state
xi_t	Reactant translational state
xi_v	Reactant vibrational state
ξ	Reactant state vector
δk_d	Standard deviation of the dissociation rate, $\text{cm}^3/\text{s}/\text{mol}$
δP_d	Uncertainty of the probability of dissociation
δx	Uncertainty of the target function
$\delta \sigma$	Uncertainty of the dissociation cross-section, \AA^2
η	Normalization factor
μ	Reduced mass, kg
τ	Reactant orientation vector
φ	Efficiency function
σ	Dissociation cross-section, \AA^2
ζ	Optimized Park model parameter

I. Introduction

Dissociation rates of nitrogen in a bath, i.e. $\text{N}_2 + \text{N}_2 \rightarrow \text{N}_2 + \text{N} + \text{N}$, at high temperatures with thermal non-equilibrium have been studied extensively due to interest in re-entry vehicle design.¹⁻⁴ Computational fluid dynamics (CFD) codes integrate thermal nonequilibrium models into existing platforms through a variety of means, including multiple-temperature and state-specific models. For the re-entry problem, the conventional model is based on two temperatures, wherein the translational and rotational energy modes of the molecules are characterized by a temperature T , and the vibrational energy modes of the molecules are characterized by a vibrational temperature T_v . In each cell, the dissociation rate of nitrogen is calculated based on the local T and T_v , which varies with both spatially and temporally. Thus, access to the full nonequilibrium dissociation rate surface, i.e. the dissociation rate over a range of T and T_v fields, is critical to achieving accurate computational results in a CFD calculation. This work presents a robust methodology for calculating the dissociation rate surface of $\text{N}_2 + \text{N}_2 \rightarrow \text{N}_2 + \text{N} + \text{N}$ and integrating the results alongside modern CFD programs.

For nitrogen dissociation, the most common approach to define the dissociation rate on a continuous field is by using Park's model.⁵ That is, for any T and T_v , the dissociation rate is calculated using the conventional Arrhenius kinetics rate equation at an effective temperature, which is a combination of T and T_v . A single parameter is used to define the effective temperature relation, which is determined based on a subset of directly calculated nonequilibrium rates. Because there is only one parameter to fit, only a small subset of the nonequilibrium rates calculated over wide range of T and T_v need to be directly calculated. The tradeoff for this simple model is global accuracy. Specifically, the relative impact of T_v compared to T on the rate is not constant for all T and T_v .

In contrast to the fitting procedure described above, this work presents a robust approach for calculating the nonequilibrium rate surface using a combination of direct calculation and interpolation. First, the nonequilibrium rate is directly calculated for a coarse-scale grid of T and T_v . Then, the nonequilibrium rate is interpolated from the original grid onto the fine-scale grid of T and T_v . This new set is tabulated to be sufficiently dense so that in a CFD application, accurate nonequilibrium rates can be interpolated given any realized T and T_v . There are three sources of error in this approach: (1) the uncertainty associated with directly calculating the nonequilibrium rate, (2) the interpolation error associated with defining the rate on a fine-scale grid, and (3) the interpolation error associated with defining the spatially and temporally local

rate in a CFD simulation. In this work, the fine-scale grid is assumed to be sufficiently dense so that the error accrued during the CFD application is negligible. The correlation of errors (1) and (2) is nontrivial, but understanding this balance is critical to minimizing the global uncertainty of the nonequilibrium rate surface.

The nonequilibrium dissociation rate was numerically calculated using a quasi-classical trajectory (QCT) analysis.⁶ Such QCT studies have been used in previous studies.^{3,7} In QCT studies, the nuclei of a system are treated as classical particles driven by a potential energy surface (PES), and, in the case of nitrogen dissociation, bimolecular collisions are simulated. The initial conditions of the collision, denoted as a trajectory, are sampled based on specified energy states or from Maxwell-Boltzmann distributions characterized by temperature states, respectively denoted as state-specific and state-averaged sampling techniques. The dissociation rate given this state is then determined from the statistics associated with the sampled trajectories. Typically, many trajectories are required at a given state to ensure a small error in the computed rate. For a new state however, a new batch of trajectories must be simulated. As mentioned before, minimizing the variance (or uncertainty) associated with each rate versus directly calculating more rates (as opposed to using interpolation) is a balance when the full nonequilibrium rate surface is being tabulated.

This work is an extension of the work done by Candler and Truhlar's research groups to account for integration into CFD programs. As a brief summary, the global N_4 potential energy surface was developed by Pauku *et al.* from approximately 17000 *ab initio* data points,² and later improved by Bender *et al.* by introducing mixed exponential-Gaussian variables in the fit.⁷ These surfaces have been used in QCT studies to understand the effect of rovibrational nonequilibrium on the reaction rate. Bender *et al.* validated the thermal equilibrium rate and analyzed the dissociation rate at 13 points $T-T_{rv}$ space. Though not sufficient to define the full rate surface, the results showed that the T_{rv} more dramatically affected the dissociation rate compared to T . More recently, Bender *et al.* decoupled the rotational and vibrational modes to calculate the rate versus T and T_v , as described above.⁷ This work extensively analyzed the effect of the reactant state on different collision paths, including single dissociation, double dissociation, and nitrogen swapping. This work extends this previous work by developing a robust process for integrating QCT-derived rates into CFD applications and understanding the uncertainty associated with these rates.

In conjunction with the previously described objectives, this work also analyzed strategies associated with sampling the initial conditions and monitoring the uncertainty. First, a mode-independent sampling technique was implemented into a new massively-parallel QCT program. That is, the translational, rotational, and vibrational energy are all independently sampled a Maxwell-Boltzmann PDF, a uniform PDF, or from a set of predetermined values. A similar strategy was also developed for the impact parameter, which is another input for a trajectory. This allowed us to assess how different sampling techniques affect the uncertainty of the dissociation cross-section. For instance, the benefit of uniformly sampling the relative translational energy is that the rate can be calculated at any temperature, but, as will be presented below, the tradeoff is higher uncertainty. Also, an algorithm was developed for monitoring and controlling uncertainty in QCT calculations. In general, the dissociation cross-section has a direct correlation to the probability of dissociation for a specific state. But, for a single trajectory, the product state is typically a binary result: either the products were formed or not. For an infinitely large set of trajectories, the probability based on these binary results converges to a fixed value. For finite resources, this fixed value is never reached, but the variance of the present result can be estimated. Using this variance, an automation strategy was developed that dynamically adapts the QCT algorithm to ensure that the overall variance uniformly conforms to a user-defined tolerance.

After implementing these strategies, nonequilibrium dissociation rates for the reaction $N_2+N_2 \rightarrow N_2+N+N$ were calculated using a PES developed by Bender *et al.*⁷ The set of directly calculated rates was used in an interpolation scheme to define the rate surface on a fine-scale grid, tabulated for use in CFD applications. The balance between minimizing the variance of the directly calculated dissociation rates versus sampling more states was analyzed in the context of minimizing the global variance of the averaged dissociation rate surface. The remainder of this abstract is organized as follows. Section II discusses the formulation of the QCT model. Section A defines the the sampled PDFs, Section B describes the process of calculating the dissociation rate and its uncertainty, and C presents the QCT program features. Finally, Section III analyzes the results of this work. These results consist of two studies. The first study analyzes the dissociation rate uncertainty in the context of sampling techniques, and the second study calculates the dissociation rate for translational and rovibrational temperatures ranging from 8000 K to 20000 K using a quasi-state-specific sampling technique. From these results, the rates are validated by thermally averaging the data and comparing the results to

other existing models and experimental data, the dissociation rate and efficiency function are analyzed in context of how dramatically the rovibrational and translational temperature affect the dissociation rate, and Park's two-temperature model is calibrated and compared to the non-equilibrium data.

II. QCT Formulation

The QCT methodology is derived from two main assumptions. First, from quantum mechanics, the Born-Oppenheimer approximation assumes that the nuclear and electron states of the system are decoupled.⁸ From this assumption and by *ab initio* methods, the eigenvalues of the electronic Schrödinger equation are (numerically) calculated for a fixed nuclear position. These eigenvalues correspond to the potential energy of the system, and when defined on the whole space of nuclear configurations, this energy acts as a continuous function, aptly referred to as the potential energy surface (PES). This surface is differentiable almost everywhere, and the only isolated non-differentiable points being where two or more nuclei from the system have the same global position. Second, the classical assumption extends the Born-Oppenheimer approximation and assumes that not only are the nuclear and electron motions decoupled, but that the nuclei can be treated as classical particles,⁶ so the wave function associated with each nucleus is assumed to be a delta function. The driving force between the nuclei is defined as the negative derivative of the aforementioned PES.

Solving the classical equations of motion for a set of nuclei is the main process in a QCT simulation. Computationally, this amounts to solving a set of ordinary differential equations (ODE) representing the evolution of the phase space of the nuclei for a variety of initial conditions and analyzing the outcomes. The numerical ODE solver chosen for this work was the adaptive Runge-Kutta Prince-Dormand (8,9) method as implemented in the open source GNU Scientific Library (GSL).⁹ Though the method is not symplectic, the error tolerance was set sufficiently low so that the total energy of the system did not drift noticeably over a typical trajectory time frame.

Regarding the initial conditions of the ODE system, the reactants' state bounds the initial phase space of the nuclei in the system. For this work, multiple state definitions have been implemented. In short, the translational, rotational, and vibrational energy of the reactants are prescribed based on this state, which in turn restricts the relative position and velocity of the reactants. The orientation of each reactant is sampled from a uniformly distributed sphere and the separation of the reactants was chosen to be greater than 15 Å (the maximum separation recognized for this particular PES).² The final specification is the impact parameter b : the distance component between the reactants perpendicular to the relative motion. In this work, the maximum impact parameter b_{max} was set to 8.0 Å based on the previous analysis by Bender *et al.*³

A. State Probability Distributions

To calculate the thermally averaged dissociation rate, i.e. the rate as a function of one or more temperatures, the PDFs associated with the energy modes of the molecules needs to be defined. The relative translational energy PDF is given by

$$f_g(g; T_t) = \left(\frac{\mu}{2\pi k_B T_t} \right)^{3/2} 4\pi g^2 e^{-\mu g^2 / 2k_B T_t} \quad (1)$$

where g is the relative translational speed, T_t is a translational temperature, μ is the reduced mass of the reactant pair, and k_B is the Boltzmann constant. The rovibrational energy PDF as a function of the rovibrational quantum numbers v and J , and characterized by a rovibrational temperature T_{rv} , is given by

$$f_{rv}(v, J; T_{rv}) = \frac{g_s(J)(2J+1)e^{-(E_{rv}(v,J)-E_{rv}(0,0))/k_B T_{rv}}}{Q_{rv}(T_{rv})}, \quad (2)$$

where E_{rv} is the quantized rovibrational energy, g_s is the spin degeneracy, and Q_{rv} denotes the partition function. In re-entry vehicle simulations, typically, the rovibrational states of the nitrogen in the system after the bow shock are decoupled so that the vibrational and rotational states are characterized by T_v and T_r , respectively. Such a decoupling procedure is not well-defined, and the derivation presented by Bender *et*

al.,⁷ a vibration-prioritized framework, was used. Here, the rovibrational PDF is given by

$$f_{rv}(v, J; T_v, T_r) = \frac{f_v(v; T_v) f_r(J; T_r | v)}{\eta(T_v, T_r)}, \quad (3)$$

where f_v is the vibrational PDF, $f_r(J; T_r | v)$ is the rotational PDF conditioned based on the vibrational state, and η is a normalization factor. In this form, the PDFs are defined by

$$f_v(v; T_v) = \frac{e^{-(E_{rv}(v,0) - E_{rv}(0,0))/k_B T_v}}{Q_v(T_v)} \quad (4)$$

$$f_r(J; T_r | v) = \frac{g_s(J)(2J+1)e^{-(E_{rv}(v,J) - E_{rv}(0,0))/k_B T_r}}{Q_{rv}(T_r)} \frac{1}{f_v(v; T_r)}. \quad (5)$$

Bender *et al.*⁷ have shown that the normalization factor varies between 0.95 and 1.10 for a large range of T_v and T_r .

B. Dissociation Rate Calculation

Let the translational state ξ_t denote a specific relative translational speed or a temperature characterizing the distribution of the relative translational speed, and let ξ_r and ξ_v analogously describe the rotational and vibrational state. Then, for N randomly sampled trajectories from the phase space restricted by the reactants' initial state, denoted by $\boldsymbol{\xi} = (\xi_t, \xi_r, \xi_v)$, and an impact parameter b , the mean probability of dissociation is defined by integrating the probability of reaction over the initial orientation of the reactants, i.e.

$$P_d(\boldsymbol{\xi}, b) = \int_{\boldsymbol{\tau}} f_{\boldsymbol{\tau}}(\boldsymbol{\tau}) P_d(\boldsymbol{\xi}, b, \boldsymbol{\tau}) d\boldsymbol{\tau}, \quad (6)$$

where $\boldsymbol{\tau}$ is the initial orientation vector and $f_{\boldsymbol{\tau}}$ is the PDF associated with the initial orientation. In regards to notation throughout this work, please note that if a dependency is dropped, i.e. $P_d(\boldsymbol{\xi}, b)$ instead of $P_d(\boldsymbol{\xi}, b, \boldsymbol{\tau})$, it is implied that the function has been averaged over the dropped variable. Using Monte Carlo integration, Eq. (6) reduces to

$$P_d(\boldsymbol{\xi}, b) = \lim_{N \rightarrow \infty} \frac{N_d(\boldsymbol{\xi}, b)}{N(\boldsymbol{\xi}, b)}, \quad (7)$$

where N_d is the number of trajectories resulting in nitrogen dissociation. The uncertainty of P_d , denoted by δP_d , is defined as two standard standard deviations from the mean, which, once again using Monte Carlo integration theory, reduces to

$$\delta P_d(\boldsymbol{\xi}, b) = 2 \frac{N_d(\boldsymbol{\xi}, b)}{N(\boldsymbol{\xi}, b)} \left(\frac{N(\boldsymbol{\xi}, b) - N_d(\boldsymbol{\xi}, b)}{N_d(\boldsymbol{\xi}, b)(N(\boldsymbol{\xi}, b) - 1)} \right)^{1/2}. \quad (8)$$

As the number of sampled trajectories goes to infinity, the uncertainty goes to zero and the subsequent dissociation rate converges to a fixed value. If no dissociative trajectories are recorded, then the uncertainty is zero, which may occur when the initial state corresponds to a low dissociation probability. Thus, by selecting the number of trajectories to simulate for a given state, the user is also defining the minimum observable probability, i.e. $P = 1/N$, which has a corresponding uncertainty of $\delta P = 2/N$. If a relative level of convergence per state is desired, trajectories are simulated at a particular state until $\delta P/P$ reaches a threshold. Using this approach, the number of trajectories per state will increase significantly for low probability states. Instead, an absolute level of convergence per state was utilized in this work so trajectories are simulated at a particular state until δP reaches a threshold. The threshold is set to a value sufficiently low so that $N_d \gg 1$ for all states, which was determined from preliminary calculations. This ensured that both low and high probability states converge uniformly, thus minimizing the global uncertainty of the probability for all states.

The dissociation cross-section at state $\boldsymbol{\xi}$ is calculated by integrating P along b weighted by $2\pi b$, i.e.

$$\sigma(\boldsymbol{\xi}) = \int_0^{b_{max}} 2\pi b P_d(\boldsymbol{\xi}, b) db. \quad (9)$$

Because the integral is a linear operator, the uncertainty of the cross-section $\delta\sigma$ is defined by replacing P with δP in Eq. (9). For uniform sampling, the impact parameter space is discretized into N_b cells, and the cross-section is the sum of the average values for each cell, i.e.

$$\sigma(\boldsymbol{\xi}) = \sum_{i=1}^{N_b} \left(\int_{b_{i-1/2}}^{b_{i+1/2}} 2\pi b P_d(\boldsymbol{\xi}, b) db \right) \approx \sum_{i=1}^{N_b} 2\pi b_i P_d(\boldsymbol{\xi}, b_i) \Delta b_i, \quad (10)$$

where $P_d(\boldsymbol{\xi}, b_i)$ denotes the probability sampled from all impact parameters in the i^{th} cell, and Δb_i is the cell width. Note that if the impact parameter is sampled based on Monte Carlo integration, then Eq. (9) simplifies to

$$\sigma(\boldsymbol{\xi}) \approx \pi b_{max}^2 P_d(\boldsymbol{\xi}). \quad (11)$$

The dissociation rate is defined by integrating the cross-section over the relative translational speed weighted by the Maxwell speed PDF characterized by a translational temperature T_t , i.e.

$$k_d(T_t, \xi_r, \xi_v) = \int_0^\infty g f_g(g; T_t) \sigma(g, \xi_r, \xi_v) dg. \quad (12)$$

Rather than integrating from zero to infinity, the upper and lower bounds of g are chosen so that $\chi(g)$ is approximately zero beyond these points. As before, the uncertainty of the dissociation rate δk_d is defined by replacing σ with $\delta\sigma$ in Eq. (12). If $\xi_t = g$ and g is uniformly sampled, the relative translational speed is discretized into N_g cells, and the dissociation rate is the sum of the average values for each cell, i.e.

$$k_d(T_t, \xi_r, \xi_v) = \sum_{i=1}^{N_g} \left(\int_{g_{i-1/2}}^{g_{i+1/2}} g f_g(g; T_t) \sigma(g, \xi_r, \xi_v) dg \right) \approx \sum_{i=1}^{N_g} g_i f_g(g_i; T_t) \sigma(g_i, \xi_r, \xi_v) \Delta g_i, \quad (13)$$

where $\sigma(g_i, \xi_r, \xi_v)$ denotes the probability sampled from all relative translational speeds in the i^{th} cell, and Δg_i is the cell width. If $\xi_t = T_t$, then the cross-section is approximated by Monte Carlo integration, i.e.

$$k_d(T_t, \xi_r, \xi_v) \approx \left(\frac{8k_B T_t}{\pi\mu} \right)^{1/2} \sigma(T_t, \xi_r, \xi_v), \quad (14)$$

where k_B is the Boltzmann constant, and μ is the reduced mass of the reactant pair.

If ξ_v and ξ_r represent specific quantum numbers, i.e. $\xi_v = (v_1, v_2)$ and $\xi_r = (J_1, J_2)$, then the thermally averaged dissociation rate is calculated by averaging over the Boltzmann PDF characterized by vibrational and rotational temperatures T_v and T_r , respectively. For the rovibrationally state-specific reaction



both reactant's initial rovibrational state are independently averaged. So, the thermally averaged dissociation energy is given by

$$k_d(T_t, T_r, T_v) = \sum_{(v_1, J_1)} \sum_{(v_2, J_2)} f_{rv}(v_1, J_1; T_v, T_r) f_{rv}(v_2, J_2; T_v, T_r) k_d(T_t, v_1, v_2, J_1, J_2) \quad (16)$$

where the summations are over the set of valid rovibrational states, and though each reactant state has been independently averaged, T_v and T_r are the same for both reactants.

C. Computational Details

Alongside this work, a new QCT program was developed to take advantage of large-scale computational infrastructure. This program was heavily influenced by the VENUS program¹⁰ and the work of Bender *et al.*³ The parallelization of the QCT algorithm is straightforward because each trajectory is independent of the other trajectories. However, due to the variation in processor speed and trajectory computational cost, a load balancing algorithm was necessary to maintain uniform computational load over all processors for the duration of a QCT simulation. Furthermore, for adaptive algorithms that dynamically adjust the simulation

based on the intermediate uncertainty calculations, a communication infrastructure was necessary so that each processor core knew which cross-sections still needed to be calculated or improved.

These computational objectives were achieved by utilizing a single, master nodes that operates alongside a set of worker nodes. During the QCT initiation processes, the master node determines the necessary states to calculate the desired set of reaction rates. Then, during the online portion of the simulation, each worker node contacts the master node one-by-one for instructions, which is the status of the QCT simulation and the next state to analyze. Once these directions are received by the worker node, it simulates a batch of trajectories sampled from the given state. Afterwards, it recontacts the master node for further instructions, and the cycle continues. Thus, interprocessor communication is minimized because there is a central node in charge of all the directives.

For up to 4,096 cores (the most tested to date), this process computationally efficient. Figure 1 shows a preliminary parallel scaling assessment of the computational time for running five million trajectories on varied numbers of cores.

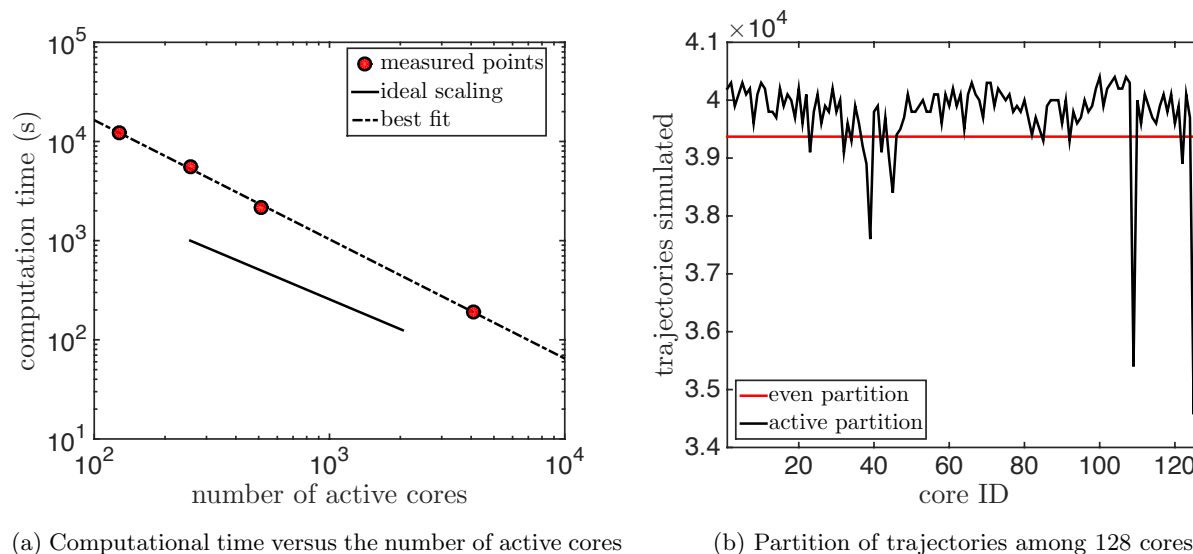


Figure 1: Parallel scaling assessment of QCT program for five million trajectories simulated using a varied number of cores

These results show that the code is both efficient, approximately linearly scaling with cores, and predictable, a small-scale simulation can be used to predict the computational cost of a large-scale simulation. Note that the perception that this scaling is more efficient than ideal is due to the marginalized cost of the master node as the total number of cores increases. The active partition in Figure 1b shows that in general, each cores simulates approximately the same number of trajectories, but a few of the trajectories run significantly fewer trajectories. To minimize the cost, the other cores balance this by simulating more trajectories. If an even partition was used, the slow cores would significantly increase the total simulation time.

Periodically throughout the online portion of the QCT simulation, at a time when the master node is not communicating with the worker nodes, the mater node calculates the set of cross-sections and their associated uncertainties from the results gathered up to that point. If the uncertainty of any cross-section is sufficiently low, below a user-defined absolute or relative tolerance, then the master node marks that state as complete. When all the states are complete, the master node communicates to the worker nodes that the job is complete. Due to the intricacies of post-processing the data for a variety of different sampling techniques and desired data points, this process is left offline and run as a separate program.

The interpolation strategy implemented for computing the nonequilibrium rate surface on a fine-scale grid of T and T_v was the multivariate interpolation and regression (MIR) scheme proposed by Wang *et al.*¹¹ The scheme creates a high-order approximation function constructed from a set of data points on a scattered domain where the estimate of the approximation error at each data point is minimized. The set of data

points, or target functions, used to create the interpolant were given by

$$x_i = \log_{10}(k_d(T_i, T_{v,i})) \quad (17)$$

$$\delta x_i = \log_{10}(k_d(T_i, T_{v,i}) + \delta k_d(T_i, T_{v,i})) - \log_{10}(k_d(T_i, T_{v,i})), \quad (18)$$

for $i = 1, \dots, N_x$ and N_x is the number of rates directly calculated. The target function was chosen as $f = \log_{10}(k)$ because of its expected near-linear relationship with T and T_v . Currently, the interpolation of the rate onto the fine-scale $T - T_v$ grid is done in a post-processing step. However, it is possible to integrate this process alongside the QCT program so that the surface can be dynamically checked throughout the simulation. However, because each directly calculated rate is converged within a user-defined uncertainty tolerance, this step was not implemented for this work.

III. Results

The results of this work are focused on the nonequilibrium dissociation rate of nitrogen in a bath, i.e. $N_2 + N_2 \rightarrow N_2 + N + N$. The analytical PES used in this work was developed by Bender *et al.*,⁷ which was an extension of the surface developed by Paukku *et al.* from a set of approximately 17000 *ab initio* data points.²

A. Study I: Optimal Sampling Techniques

The first study consisted of three test cases designed to analyze the effect of different sampling techniques on the estimated uncertainty. The goal of each test was to calculate the dissociation rate and estimate the corresponding uncertainty for six temperatures ranging from 10000 K to 20000 K. One million trajectories were simulated per temperature, leading to a total of six million trajectories per case. The differences between each case were how the impact parameter and relative translational speed were sampled. The rovibrational states were sampled from a Boltzmann distribution at the corresponding temperatures. Table 1 shows the different PDFs used to sample the initial conditions for each test case.

Test Case	$f_b(b)$	$f_g(g; T)$	$f_{rv}(v, J; T)$
1	Linear	Maxwell	Boltzmann
2	Uniform	Maxwell	Boltzmann
3	Uniform	Uniform	Boltzmann

Table 1: PDFs used to sample the initial conditions for each test case

Case 1 corresponds to full Monte Carlo sampling, so the cross-section and dissociation rate are given by Eq (11) and Eq (14), respectively. For case 2, the cross-section and dissociation rate are given by Eq (10) and Eq (14), respectively. For case 3, the cross-section and dissociation rate are given by Eq (10) and Eq (13), respectively. For the cases 2 and 3, the impact parameter was discretized into 32 cells ranging from 0 Å to 8 Å. For case 3, the relative translational speed was discretized into 25 cells. The minimum sampled speed was 10^{-4} as likely as the most probable speed based on the Maxwell PDF at 10000 K, and the maximum sampled speed was 10^{-8} as likely as the most probable speed based on the Maxwell PDF at 20000 K.

First, the effect of the impact parameter sampling technique was analyzed by comparing cases 1 and 2. Figure 2 shows the probability of dissociation multiplied by $2\pi b$ versus the impact parameter at 14000 K and 20000 K.

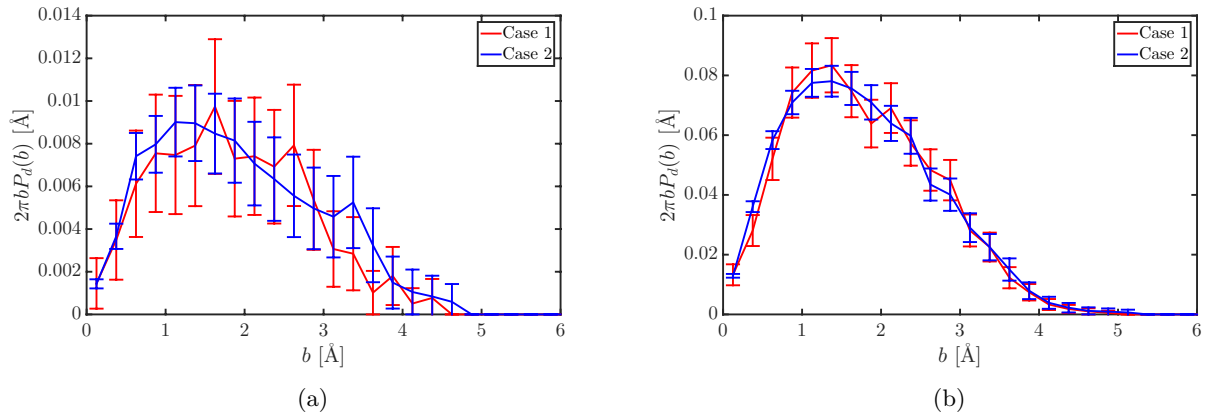


Figure 2: Probability of dissociation multiplied by $2\pi b$ versus impact parameter: (a) $T = 14000$ K, and (b) $T = 20000$ K.

Note that the area under each curve is the corresponding cross-sections. Based on qualitative inspection, it is clear that as the temperature increases, the uncertainty decreases. This trend follows from the uncertainty definition in Eq. (8). The average uncertainty of the dissociation probability $\langle \delta P_d \rangle$ was normalized by the average dissociation probability $\langle P_d \rangle$ to create a quantified measure of comparison. The results are tabulated in Table 2.

Test Case	T [K]	$\langle \delta P_d \rangle / \langle P_d \rangle$
1	14000	0.427
2	14000	0.292
1	20000	0.143
2	20000	0.098

Table 2: Probability distribution functions sampled from for each test case

For both low and high temperatures, the uniform sampling technique (case 2) has a lower average uncertainty compared to the linear sampling technique (case 1). Moving from case 1 to case 2, at 14000 K the relative uncertainty decreased by 13.5%, and at 20000 K the relative uncertainty decreased by 4.5%. For case 1, too few trajectories were sampled for low impact parameters, where dissociation is more likely to occur, resulting in a higher variance. For instance, for one million sampled trajectories at 14000 K, only 348 trajectories resulted in dissociation. However, when uniformly sampling, 1265 trajectories resulted in dissociation at 14000 K. At high impact parameters, case 1 has a lower uncertainty compared to case 2, but the probability of reaction in these regions is already very small. Thus, the difference is not sufficient to improve the average relative uncertainty significantly.

One of the main drawbacks of both cases 1 and 2 are that for each desired translational temperature, a new batch of trajectories needs to be calculated. To account for this, test case 3 uniformly samples both the impact parameter and relative translational speed. With this in mind, Figure 3 shows the computed dissociation rate for each case.

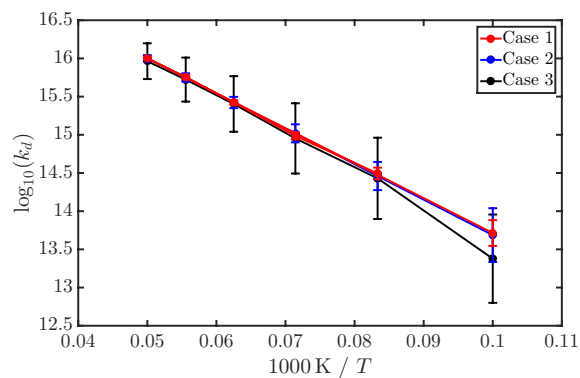


Figure 3: Thermal equilibrium dissociation rate versus the temperature. The units for k_d are $\text{cm}^3/\text{s}/\text{mol}$.

Notice that the dissociation rate's uncertainty for case 1 is now lower than for case 2, which is unexpected because it contradicts the results in Figure 2 and Table 2. This suggests that though Monte Carlo integration does not accurately capture low-level information, e.g. the probability versus the impact parameter, it instead, by assuming the PDF of the state, more accurately captures the high-level information, e.g. the reaction rate. This claim is also supported by the results for case 3, which has the highest uncertainty. However, for case 1 and 2, only 6 cross-sections were calculated, whereas for case 3, 150 cross-sections were calculated. For each cross-section in case 1 and 2, one million trajectories were simulated, but for case 3, only forty thousand trajectories were simulated per cross-section. This is the reason the uncertainty is so large for case 3. However, by uniformly sampling the relative translational speed, the dissociation rate at any translational temperature is calculated by averaging the speed-dependent cross-section weighted by the Maxwell PDF. Thus, because the uncertainty is not too large for case 3, it should be used to calculate the dissociation rate surface because more data points can be generated on a fine-scale grid of T_t to create the interpolant.

A sampling concern with case 3 is that the rotational and vibrational energy sampling techniques are still state-averaged based on T_r and T_v , respectively. But, for $\text{N}_2 + \text{N}_2$, each reactant has approximately 9,200 bound and quasi-bound rovibrational states, and for both reactants together, there are almost 85 million total rovibrational state combinations. Thus, systematically sampling the rovibrational quantum state makes this problem computationally intractable. Even for a sparse selection of states, interpolation between the states adds a large amount of uncertainty. So, rather than sampling rovibrational quantum numbers, we sample a much smaller selection of rovibrational temperatures and interpolate the values in between. This feasibility of this process was confirmed in the second study.

Finally, the convergence of the three test cases was analyzed. Figure 4 shows the convergence of dissociation rate by plotting the uncertainty in the dissociation rate versus the number of trajectories sampled.

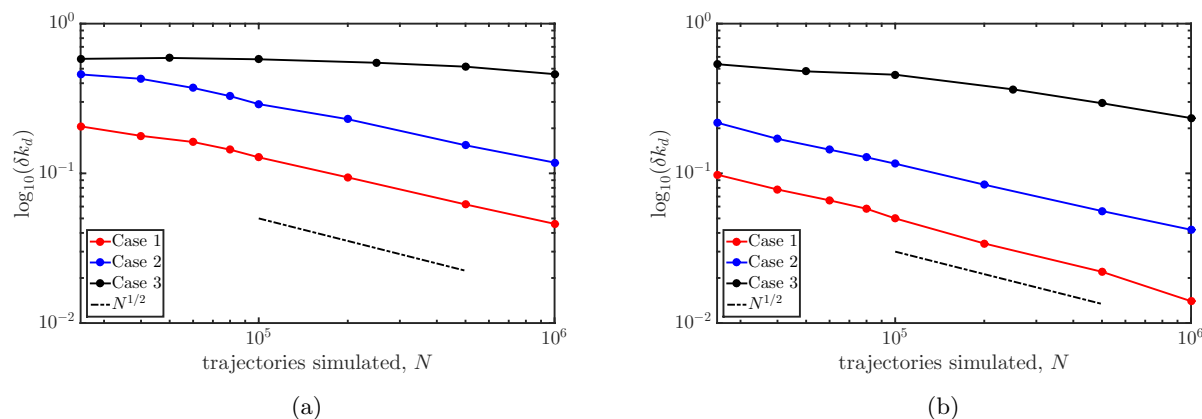


Figure 4: Convergence of dissociation rate based on estimated uncertainty for varied number of total trajectories: (a) $T = 14000$ K, and (b) $T = 20000$ K.

Cases 1 and 2 converge at a rate proportional to $N^{1/2}$, which corresponds to the expected rate for Monte Carlo integration. For case 3, the convergence rate is slower, especially at lower temperatures. Too few trajectories are simulated for each cross-section. Once a sufficient number of trajectories that result in reaction are simulated, especially for low-speed cross-sections, the convergence rate will increase. This is seen in Figure 4b, wherein case 3 converges at approximately the same rate as the other cases because a sufficient number of reactive trajectories have been simulated.

B. Study II: Non-Equilibrium Dissociation Rate Surface Calculations

The second study was a single run designed to calculate 400 unique cross-sections. That is, 100 evenly spaced relative translational speeds were studied for four rovibrational temperatures: 8000 K, 10000 K, 13000 K, and 20000 K. A total of 500 million trajectories were simulated on 2,048 cores over 15 hours. After the simulation, the dissociation rate was calculated at 66 translational temperatures evenly spaced between 8,000 K and 25,000 K for each of the rovibrational temperatures. These dissociation rates were used in the open source MIR library developed by Wang *et al.*¹¹ to calculate the rate surface, as shown in Figure 5a.

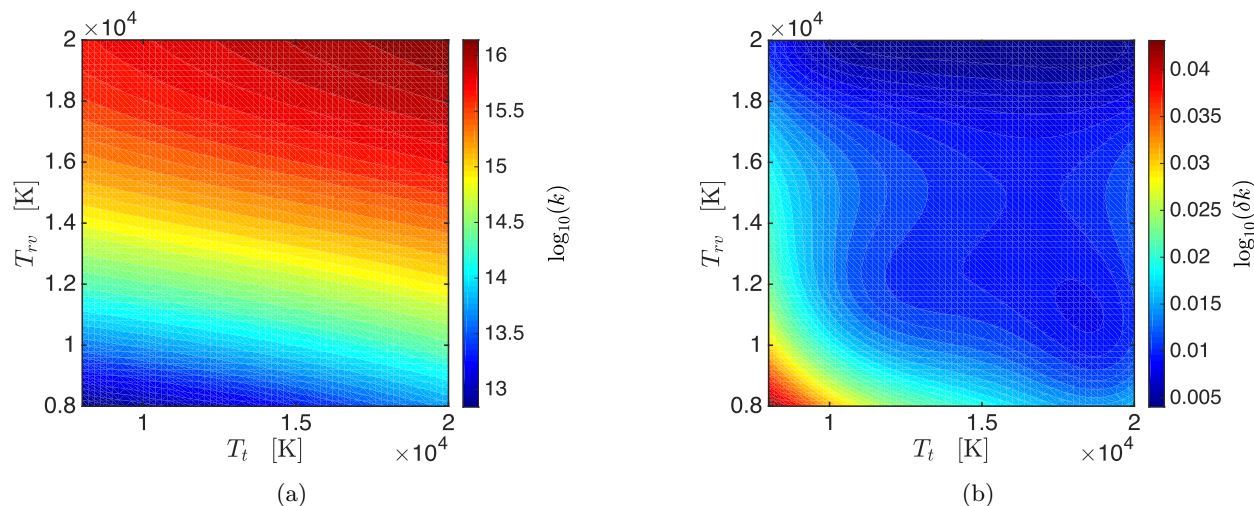


Figure 5: Dissociation rate versus translational and rovibrational temperature from the quasi-state-specific QCT method: (a) the dissociation rate surface, and (b) the standard deviation of the dissociation rate. The units for k and δk are $\text{cm}^3/\text{s}/\text{mol}$.

From this figure, it is clear that the rovibrational temperature has a significantly stronger effect on the dissociation rate than the translational temperature. The results by Bender *et al.* also follow this trend.³ The uncertainty associated with the rate surface, which is predicted by the MIR scheme, is shown in Figure 5b. Looking at the the values between $T_{rv} = 8000$ K, 10000 K, 13000 K and 20000 K, we observe that the uncertainty of the interpolated dissociation rates is most affected by the calculated rate's local uncertainty. The largest uncertainty is for low temperatures, specifically near 8000 K for both T_t and T_{rv} . As with the previous study, this information can be dynamically analyzed during the QCT simulation to decide where more trajectories should be run to improve the global accuracy of the surface.

As validation, the slice from Figure 5a where $T_{rv} = T_t$, i.e. the thermal equilibrium rate, is compared with other published rates for nitrogen dissociation. The uncertainty bars correspond to two standard deviations from the mean, or a 95 % confidence interval.

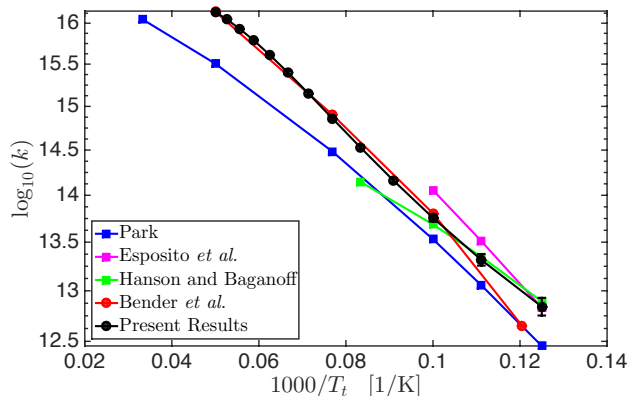


Figure 6: Thermal equilibrium dissociation rate versus the translational temperature ($T_{rv} = T_t$). The units for k are $\text{cm}^3/\text{s}/\text{mol}$.

At low temperatures, the present results match, within the uncertainty bars, both the model from Esposito *et al.* calculated using the Park factor¹ and shock-tube experimental studies of Hanson and Baganoff.¹² From 8000 K to 10000 K, the rate is very close to the results of Hanson and Baganoff. Above 10000 K the two diverge, but from 10000 K to 20000 K, the present results closely match the rate calculated by Bender *et al.*³ The rate never exactly matches the rate from Park's model,⁵ but the two are close from 9000 K to 12000 K. It should be noted that Park's model explicitly assumes $T_r = T_t \neq T_v$, whereas we have used $T_r = T_v \neq T_t$. We made this choice to compare our results directly with the results of Bender *et al.*³

From the dissociation rate surface in Figure 5a, the efficiency function is defined as the relative effectiveness of T_{rv} in increasing or decreasing the dissociation rate,¹³ given as

$$\varphi = \frac{k_d(T_t, T_{rv})}{k_{eq}(T_t)}, \quad (19)$$

where $k_{eq}(T_t) = k_d(T_t, T_{rv} = T_t)$. The logarithm of the efficiency function for T_t and T_{rv} ranging from 8000 K to 20000 K is shown in Figure 7.

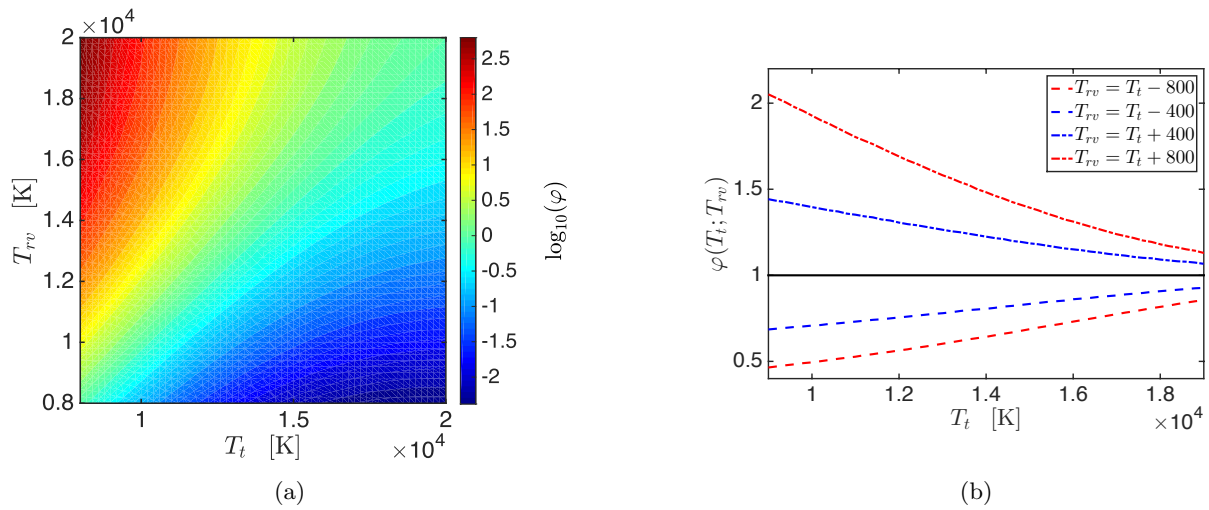


Figure 7: Efficiency function versus translational and rovibrational temperature from the quasi-state-specific QCT method: (a) the efficiency function surface, and (b) slices along $T_{rv} = T_t + \Delta T$.

The surface shows that for T_{rv} either greater or less than T_t , the dissociation rate will be noticeably affected for even moderate differences. At the most extreme $T_t - T_{rv}$ values, the non-equilibrium rate is approximately 300 times greater or less than the equilibrium rate, but these conditions rarely occur in physically realizable flows. At more moderate temperature differences, slices along the surface are insightful, where at 8000 K, if T_{rv} is within 5 % of T_t (i.e. $|T_{rv} - T_t| < 400$ K), then the dissociation rate ranges from 75 % to 150 % of the thermal rate. This effect is amplified even further for larger differences. For the temperature range studied, we observe that increasing T_{rv} relative to T_t had a more substantial effect on the rate compared to decreasing T_{rv} relative to T_t by the same amount. As the temperature increases though, these effects are diminished because the likelihood of dissociation is already high.

Lastly, from these simulations, the present non-equilibrium results are compared to Park's two-temperature model⁵ (with the exception that $T_r = T_v$ as opposed to T_t), whereby the dissociation rate is given by:

$$k_{Park}(T_t, T_{rv}) = k_{eq}((T_t T_{rv}^\zeta)^{1/(1+\zeta)}), \quad (20)$$

where ζ is a constant. Figure 8 shows the rate surface derived using Park's model for a conventional model where $\zeta = 1$ and for when $\zeta = 4$.

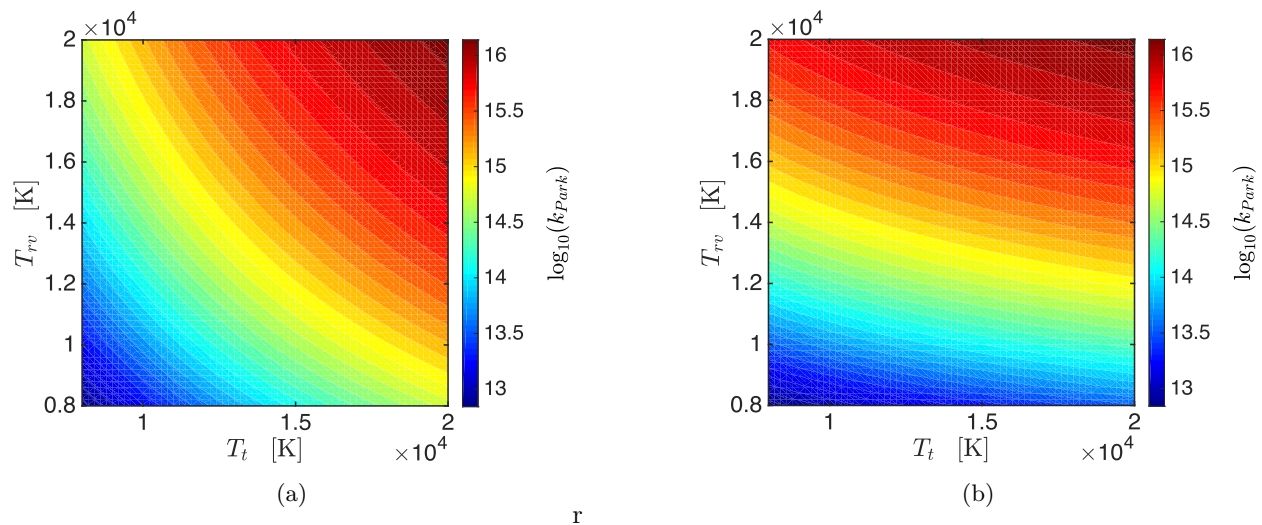


Figure 8: Dissociation rate versus translational and rovibrational temperature based on Park’s two-temperature model: (a) $\zeta = 4$, and (b) $\zeta = 4$. The units for k_d are $\text{cm}^3/\text{s}/\text{mol}$.

For $\zeta = 1$, by construction the k_{Park} surface is symmetric across the $T_t = T_{rv}$ line, which indicates that an increase in either T_t or T_{rv} will have an identical effect on the dissociation rate. This is different from the true non-equilibrium rate as shown in Figure 5a, in which T_{rv} has a larger effect on the rate than T_t . With this in mind, ζ was increased to 4 to more closely match the non-equilibrium rate surface. This implies that T_{rv} is approximately four times more influential than T_t on the non-equilibrium rate. In the final work, an optimized m and n will be calculated. The efficiency function was also calculated for the Park model when $\zeta = 4$. Slices compared the non-equilibrium rate are shown in Figure 9. Even for the non-optimized selection of ζ , this figure validates Park’s model for calculating the efficiency function of N_2 dissociation. Only for relatively low T_t and $T_{rv} \gg T_t$, are significant differences between Park’s model and the non-equilibrium model observed. It is not yet known if there is an optimal selection of ζ that minimizes these differences further.

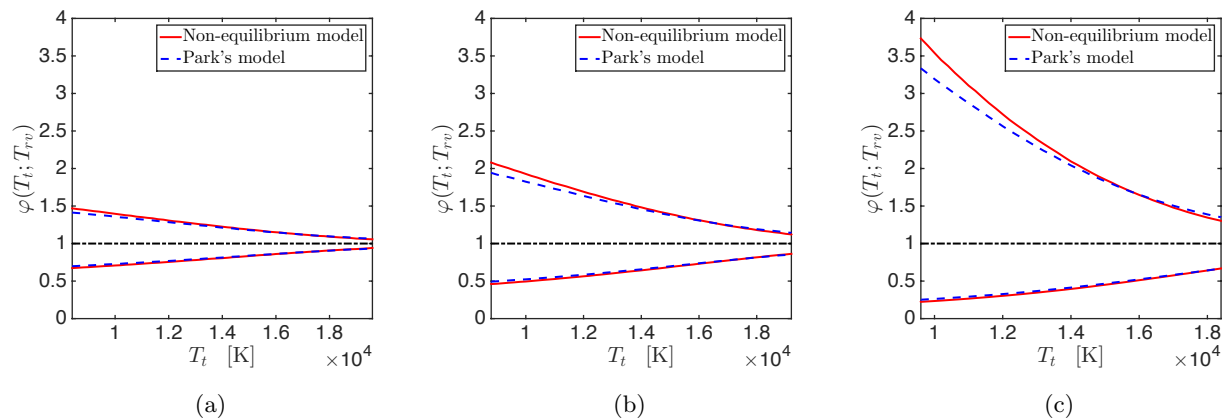


Figure 9: Slices along the efficiency function surface at various $T_{rv} = T_t + \Delta T$ for both the non-equilibrium and Park’s model: (a) $T_{rv} = T_t \pm 400$, (b) $T_{rv} = T_t \pm 800$, and (c) $T_{rv} = T_t \pm 1600$

IV. Conclusions

The focus of this work was accurate and efficient calculation of dissociation rates of nitrogen at high temperatures with thermal non-equilibrium. The strategy for accomplishing these goals was to calculate quasi-state-specific dissociation cross-sections and utilize the results to calculate the dissociation rate surface for a large range of translational and rovibrational temperatures. Minimizing and understanding the uncertainty of these calculations was also a critical goal. Specifically, for a fixed total number of trajectories, this work analyzed the consequences of different sampling techniques.

A new massively-parallel QCT program was used to this effect. The features of this code include its ability to scale to many processors while maintaining a high level of efficiency. Using a master/worker core partition strategy, the uncertainty of the directly calculated rates is estimated online throughout the simulation. This allows the number of trajectories for a given state to be truncated after a user-defined tolerance has been achieved. Also, a mode-independent sampling technique was developed so that the impact parameter, relative translational speed, rotational state, and vibrational state of the reactants are all independently sampled.

This first study examined the role of sampling techniques on the dissociation rate uncertainty. It was found that though Monte Carlo sampling poorly estimates low-level information such as the probability of dissociation, it accurately calculates the high-level information such as the dissociation rate. However, the tradeoff for Monte Carlo sampling is that if the state changes, a new set of trajectories must be simulated to calculate the rate. In contrast, uniformly sampling the state leads to a higher uncertainty, but the results can be averaged offline and weighted against a PDF characterized by any temperature. Thus, for CFD applications, uniform sampling is preferable because the reaction rate can be tabulated on a fine-scale grid of temperatures without using interpolation. However, for the rovibrational states, state-specific sampling is computationally intractable to date, so interpolation must be used.

In the second study, a batch of 500 million trajectories was simulated to verify and validate both the newly developed QCT program and the sampling technique's viability. The non-equilibrium rate surface was built using an interpolation strategy that accounted for the uncertainty of the rates. In general, the uncertainty of the surface was small except for low T_t and T_{rv} . The thermal rate was compared with other models and experimental data, showing strong agreement. The non-equilibrium rate surface showed that T_{rv} has a stronger effect on the dissociation rate than T_t . This was also observed from the efficiency function of the dissociation rate, where relatively small changes of T_{rv} compared to T_t could have a dramatic effect on the rate. Finally, Park's two temperature model was examined and compared to the true non-equilibrium rate. Before calibration, Park's model under-estimated the T_{rv} effect on the dissociation rate compared to T_t , but after an approximate calibration, the efficiency function of Park's model closely matched the non-equilibrium model.

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