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Radiative transfer and spectroscopic databases: A line-sampling Monte Carlo approach



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ABSTRACT

Dealing with molecular-state transitions for radiative transfer purposes involves two successive steps that both reach the complexity level at which physicists start thinking about statistical approaches: (1) constructing line-shaped absorption spectra as the result of very numerous state-transitions, (2) integrating over optical-path domains. For the first time, we show here how these steps can be addressed simultaneously using the null-collision concept. This opens the door to the design of Monte Carlo codes directly estimating radiative transfer observables from spectroscopic databases. The intermediate step of producing accurate high-resolution absorption spectra is no longer required. A Monte Carlo algorithm is proposed and applied to six one-dimensional test cases. It allows the computation of spectrally integrated intensities (over 25 cm^{-1} bands or the full IR range) in a few seconds, regardless of the retained database and line model. But free parameters need to be selected and they impact the convergence. A first possible selection is provided in full detail. We observe that this selection is highly satisfactory for quite distinct atmospheric and combustion configurations, but a more systematic exploration is still in progress.

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1. Introduction

Recent advances in Monte Carlo methods [1,2] indicate that null-collision algorithms can be used to simulate radiative transfer in semi-transparent media in such a way that grids are no longer required. Even for highly non-

http://dx.doi.org/10.1016/j.jqsrt.2015.10.016 0022-4073/© 2015 Elsevier Ltd. All rights reserved. homogeneous configurations the volume does not need to be discretized and this is achieved while still preserving all the statistical properties of standard Monte Carlo algorithms: convergence toward the exact solution is rigorous (no bias) and for a finite number *M* of samples, the estimate is associated with a faithful statistical uncertainty ($\frac{1}{\sqrt{M}}$ times the standard deviation of the Monte Carlo weights). The initial idea of null-collisions came from the plasma physics and neutron transport communities. It consists in adding a third type of collisional event to the absorption and scattering

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Nomenclature ¹		$\hat{p}_X(x)$	probability density function for absorption/
Ai	Bernoulli random variable		collision events
$\dot{B_{\nu}}$	blackbody intensity	$p_N(\nu)$	probability density function for wavenumber
ď	distance	$p_s(\boldsymbol{\omega} \boldsymbol{\omega}')$	single scattering phase function
Ε	statistical expectation	$Q_{j}(j)$	second component of the proposed transition
$h_{a\nu i}$	contribution of transition i to the absorption		probability expression
u,2 y	coefficient	r	random number sampled uniformly over the
$\hat{h}_{\nu,i}$	contribution of transition j to the extinction		unit interval
- 0	coefficient	S_k	moments of order k
i	index of collision	t	time
i _b	narrowband index	Т	temperature
I_{ν}	monochromatic specific intensity	$t_{1\%}$	computation time require to get a 1% standard
j	index of molecular transition		deviation
$k_{a,\nu}$	absorption coefficient	$T_U(u)$	first component of the proposed transition
$k_{n,\nu}$	null-collision coefficient		probability expression
$k_{s,\nu}$	scattering coefficient	и	index of spectral bands
$\hat{k}_{ u}$	extinction coefficient resulting from null-	VOV	variance of variance
	collision introduction	w	Monte Carlo weight
$ ilde{k}_{ u}$	approximate absorption coefficient	x	location (scalar)
т	index of independent realization	<i>x</i> ₀	observation location
Μ	number of independent realizations	Х	location (vector)
N _b	number of narrowbands	у	absorption/emission location at the end of a
N _t	number of molecular transitions		path involving null-collisions
P_a	probability of absorption	α	multiplicative factor
$P_{a,J}(j)$	probability of absorption for a given transition	YQ	half-width of an arbitrary Lorentz profile
	j	δ	delta-Dirac distribution
$P_b(i_b)$	probability of narrowband i_b	δ_{π}	partial derivative with respect to parameter π
$P_J(j)$	probability of transition <i>j</i>	δν	spectral interval
P_n	probability of null collision	ζ	extinction threshold
$p_X(x)$	probability density function for absorption/	ν	wavenumber
	emission location	π	generic parameter
		τ	transmissivity
		ω	direction (vector)

events. These new collisions change nothing with regard to the propagation of radiation. They are pure-forward scattering events: after the collision, the photon continues its trajectory as if no collision had occurred (the collision is "null" or "virtual"). The amount of such collisions can therefore be tuned to any value without modifying the resulting transfer. This tuning can then be such that the total density of the colliders is uniform: more null-collisions are added where true extinction is weak, fewer where it is strong, so that the total extinction coefficient is uniform (as in a homogeneous semi-transparent medium). The consequence is that collision locations can be sampled within a simple exponential-distribution, according to the Beer extinction law. It is thus only at the sampled location that the question is raised of evaluating the respective amounts of absorbers, true-scatterers and null-colliders, in order to sample one type of events out of the three possible ones (according to their respective probabilities). The photontransport Monte Carlo algorithm is therefore decoupled from the spatial description of radiative properties. It is not required to "know" the true radiative properties when transporting photons until their next collision location.

In preceding null-collision algorithms, the radiative properties were indeed known and the only benefit was a simplification of the procedure for accessing the data (getting rid of the volume-discretization requirement). Here we assume that the monochromatic absorption coefficient $k_{a,\nu}$ is unknown and that the only available data are those of the molecular spectroscopic databases [3–6]. We still have $k_{a,\nu} = \sum_{j=1}^{N_t} h_{a,\nu,j}$, where N_t is the number of molecular transitions and $h_{a,\nu,j}$ is the known contribution of the *j*-th transition to the absorption coefficient at wavenumber ν , but this sum is not precomputed. Null-collisions allow us to initiate the Monte Carlo algorithm without computing $k_{a,\nu}$ and we then observe that the algorithm can be continued without ever making this computation: $\sum_{j=1}^{N_t} h_{a,\nu,j}$ is diluted into the photon-transport algorithm via successive samplings of the transition index.

Why this is possible is quite simple. Let us take only two transitions: $k_{a,\nu} = h_{a,\nu,1} + h_{a,\nu,2}$ and write $k_{a,\nu} = P_1 \frac{h_{a,\nu,1}}{P_1} + P_2 \frac{h_{a,\nu,2}}{P_2}$ with P_1 and P_2 in]0, 1[and $P_1 + P_2 = 1$. This justifies a Monte

¹ We retain here the following standard statistical formalism: random variables are written in upper-case (*e.g. X*), free and bound variables in lower-case (*e.g. x*) and samples of random variables are subscripted (*e.g. x_m*).

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