Chempl: a playable package for modeling interstellar chemistry

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Abstract

Astrochemical modeling is needed for understanding the formation and evolution of interstellar molecules, and for extracting physical information from spectroscopic observation of molecular lines. The modeling usually involves handling of a chemical reaction network and solution of a set of coupled nonlinear ordinary differential equations, which is traditionally done using code written in compiled languages such as Fortran or C/C++. While being computationally efficient, there is room for improvement in the ease of use and interactivity for such an approach. In this work we present a new public code named CHEMPL, which emphasizes interactivity in a modern Python environment, while remaining computationally efficient. We demonstrate its usage with a few astrochemically relevant examples.

1 Introduction

The interstellar medium (ISM) is ever-evolving under the effect of electromagnetic radiation, cosmic-rays, and hydrodynamic motion coupled with gravitational and magnetic field. As the physical condition changes, the chemical composition changes accordingly. The buildup of chemical complexity in the ISM, the chemical heritage of forming stellar and planetary systems, and the possible connection between astronomy and the origin of life are fascinating topics. Over the years the chemistry of ISM has proved to be a powerful diagnostic of the physical parameters. For this to work at a quantitative level, it is necessary to model the chemistry in detail. Except for simple cases, a comprehensive chemical network has to be used, because most of the time it is not easy to figure out beforehand which reactions are important and which are not. With over 200 molecules detected in space (not counting isotopologues) (Endres et al. 2016; McGuire 2018), the scale of chemical networks for modeling interstellar molecules keeps growing, with the largest at the moment containing tens of thousands of reactions. The evolution of a large and interconnected chemical network can only be tracked with numerical methods.

In the field of astrochemistry — namely, the study of the chemical side of astronomy — there are already a few codes, many of which are publicly available, including: Nahoon (Wakelam et al. 2012), MAGICKAL (Garrod 2013), KROME (Grassi et al. 2014), Nautilus (Ruaud et al. 2016), ALCHEMIC (Semenov 2017), and UCLCHEM (Holdship et al. 2017). So why create yet another code?

Traditionally, codes for astrochemistry are written in Fortran or C/C++, and the mode of running is usually like this: (1) The user prepares a few files containing the reaction network, the initial chemical abundances, and the physical parameters, (2) then feeds these files to the compiled program to get the evolution of the abundances of different species as a function of time, with the results saved in files. In some codes a preprocessing step to generate the source code describing the differential equations from a reaction network is needed. (3) To analyze the results, separate code has to be written to load the files containing the results. While this whole process can be pipelined and parallelized (at the process level) to work efficiently, it lacks interactivity. The user has to switch between different execution environments back and forth, e.g., a command line terminal to run the chemical code, and a visualization environment (such as Python) for analyzing and making plots. Such a context switching and the associated data loading could be cumbersome. When working with models with time-dependent physical conditions, one usually has to hard-code the time-dependency into the code and recompile the binary executable each time a change is made, which could be distracting to say the least.

Here we present a new code CHEMPL, which is aimed

1CHEMPL is available at https://github.com/fjdu/chempl

A few reaction networks and initial conditions are also included in this repository. All the scripts for the example models in this paper are available at
at improving interactivity without sacrificing efficiency. In writing CHEMPL, one goal we keep in mind is to make the code as “playable” as possible, hence the name. It should be relatively straightforward to make changes to the model configuration, and be fast to see the effects of the changes that just have been made. To ensure performance, under the hood the code is written in C++, and the ODE (ordinary differential equation) solver being used is the time-tested DLSODES solver of the ODEPACK package (Hindmarsh 1983) written in Fortran. The interactivity is achieved through integration with Python, and is designed to be used in the IPython or Jupyter environment. We use Cython (Behnel et al. 2011) to wrap the C++ code into a Python module. All the model data can be saved as human-readable or binary data files, or it may be more beneficial to save the Python data structures using the “pickle” facility of Python (or other structured data format such as the Advanced Scientific Data Format (ASDF)). Note that CHEMPL can also be used in the “traditional” mode (i.e., working with configuration files and launched in the command line), or incorporated into other simulations as a subroutine with a few lines of additional code for setting it up.

This paper focuses on code capabilities and demonstration of use cases for CHEMPL, instead of the scientific implications of the models. In Section 2 we depict the mathematical framework for interstellar chemistry. In Section 3 we show a few example astrochemical models simulated with CHEMPL. We conclude in Section 4.

2 Mathematical formulation of Interstellar Chemistry

In astrochemical modeling the abundances of different chemical species are solved as a function of time. The equation to be solved is of the form

\[ \frac{dy}{dt} = f(t, y), \]  

where \( y \) is a vector with component \( y_i \) being the abundance of the \( i \)th species, usually measured relative to hydrogen nuclei. The vector-valued function \( f \) on the right hand side is a nonlinear function of the components of \( y \), and may be explicitly time-dependent.

Sometimes a steady-state solution may be useful, which may be obtained by solving the algebraic equation \( f(t, y) = 0 \), or by evolving \( y(t) \) for a very long time by solving equation (1).

The system of ODE equations in equation (1) is often stiff, requiring that implicit, rather than explicit methods, be used for efficient solution. Solvers of stiff ODEs often require knowledge of the Jacobian of the right hand side. The Jacobian matrix is defined as

\[ J_{ij} \equiv \frac{\partial f_i}{\partial y_j}, \]

which is often sparse for astrochemistry-relevant networks, meaning that a large fraction \((\gtrsim 90\%)\) of the elements of \( J \) are identically zero. CHEMPL constructs the sparse structure from the reaction network.

2.1 The canonical rate equations

In interstellar or protoplanetary conditions, where the majority of the gas under consideration has a density typically in the range of ten to the power of a few per cm\(^3\), and densities higher than \( \sim 10^{13} \text{ cm}^{-3} \) rarely need to be considered, three-body chemical reactions are not important and are excluded in astrochemical networks.

When only first and second order reactions are included, equation (1) may be written in component form as

\[ \frac{dy_i}{dt} = \sum_{j \in F_{i,1}} k_{ij} y_{j,1} + \sum_{j \in F_{i,2}} k_{ij} y_{j,1} y_{j,2} - \sum_{j \in D_{i,1}} k_{ij} y_{j,1} - \sum_{j \in D_{i,2}} k_{ij} y_{j,1} y_{j,2}, \]

where \( F_{i,1} \) and \( F_{i,2} \) are the set of first and second order reactions that form species \( i \), \( D_{i,1} \) and \( D_{i,2} \) are the set of first and second order reactions that destroy species \( i \), \( k_j \) is the rate coefficient of reaction \( j \), and \( y_{j,1} \) and \( y_{j,2} \) are the abundances of the first and second (when applicable) reactants of reaction \( j \). Note that for \( j \in D_{i,1} \), \( y_{j,1} \) is always simply \( y_j \), and for \( j \in D_{i,2} \), one of \( y_{j,1} \) and \( y_{j,2} \) (or both) must be the same as \( y_j \).

Examples of first order reactions are photodissociation, cosmic-ray ionization, adsorption, and desorption reactions. These reactions are not strictly “uni-molecular”, since when one of these reactions happen there must be two particles interacting, with one being the chemical species we care about and the other being a photon, a cosmic-ray particle, or a dust grain. They are treated as first order because the fluxes of photons and cosmic-rays are external inputs and are independent of local conditions, and in the case of adsorption the dust grains can be “reused” (hence effectively being constant). Examples of second order reactions are neutral-neutral, neutral-ion,
Molecules can be adsorbed on the surface of dust grains. After being adsorbed, these molecules are able to migrate over the dust grain surface and may react with each other when two molecules meet at the same surface site. Since a dust grain has a limited surface area, when many molecules have adsorbed on its surface, newly adsorbed ones will likely sit on top of another. In this way, multiple ice layers will form on the dust grains. In such a description of dust grain structure, usually only molecules on the topmost layer are assumed to be mobile and reactive, while molecules underneath (in the mantle) are “frozen” and nonreactive (however, see Chang & Herbst 2014, 2016). When molecules in the topmost layer evaporate, the underneath layer will be revealed and becomes the new reactive layer. This is the so-called three-phase model of interstellar chemistry, in which the three phases are: gas phase, dust grain surface, and dust grain mantle.

CHEMPL has implemented the three-phase description of gas-grain chemistry, following the approach of Hasegawa & Herbst (1993) and Garrod & Pauly (2011) (for a detailed description, see also Du 2012). For each surface species (except for very weakly adsorbed species such as H₂) a new mantle species is included in the network. For species $i$ with surface population per grain $n_S(i)$, its evolution equation is

$$\frac{dn_S(i)}{dt} = \sum_{j,k} k_{j \rightarrow k} n_S(j)n_S(k) - \sum_{j} k_{j \rightarrow i} n_S(i)n_S(j) + k_{ads}(i)n_G(i) - k_{ads} n_S(i)/N_S + k_{des} n_M(i)/\max(n_M, n_S) - k_{des}(i)n_S(i),$$

(4)

and the mantle population of $i$ evolves according to

$$\frac{dn_M(i)}{dt} = k_{ads} n_S(i)/N_S - k_{des} n_M(i)/\max(n_M, n_S).$$

(5)

In equation (4) the two summations are over two-body reactions producing and consuming species $i$. $k_{ads}/des(i)$ is the adsorption/desorption rate coefficient of $i$, and $n_G(i)$ is the gas phase population of $i$. $k_{ads}/des$ is the total adsorption/desorption rate of all the species, $n_S$ is the total population of all surface species, $n_M$ is the total population of all mantle species, and $N_S$ is the number of reactive sites per grain ($\sim 10^9$ for a dust grain radius of 0.1 $\mu$m).

The quantity “population-per-grain” $n_S(i)$ is related with abundance with respect to hydrogen nuclei $y_i$ by

$$y_i = n_S(i)/\eta,$$

(6)

where $\eta$ is the dust-to-hydrogen number ratio, with is of the order of $2.8 \times 10^{-12}$ for canonical parameters (dust-to-gas mass ratio of 0.01, dust grain radius of 0.1 $\mu$m, . . .).

With the inclusion of three-phase chemistry, the code will run slower, not just because of the increase in the number of variables due to addition of mantle species, but also because of the increased non-linearity of equation (4) relative to equation (5).

### 2.3 The workflow of CHEMPL

While CHEMPL can be used as a stand-alone program, it is mainly intended to be used as a Python module. Here we describe the basic workflow of CHEMPL when used in this mode. A working example is shown in Table 1.

1. Create the ChemModel object. This is the central Python object to work with in CHEMPL. During the instantiation the user may provide the file names of the reaction network, the initial abundances, and the enthalpies (needed for reaction heat calculation). These files could also be loaded later with the corresponding helper functions. The user could also add reactions one by one using Python code, and this is the way to load reaction network stored in different formats.

2. Set the physical parameters. Similarly, this could be done with an input file, or using a helper function to set the value of each parameter by its name. They could also be provided in the previous step. Optionally, the user could add time-dependency for certain parameters by providing a lookup table. Internally the code linearly interpolate the table to evaluate time-dependent parameters at arbitrary time.

3. Do some “boilerplate” preparation work. These include, e.g., sorting out reactions of different types, getting the elemental composition of each species and calculating their molecular masses and quantum mobilities, and calculating the heat of each reaction. Such boilerplate work usually only need to be done once for each model, unless during some unconventional investigations (e.g., changing the reaction network while the code is running).
4. Setup the ODE solver. This usually amounts to setting the solver ID, which is itself optional (could be useful when running in multi-process mode). Details of the solver setup and preparation are concealed in the C++ code, which include constructing the sparse structure of the ODE system, allocation of the needed working memories, and assignment of flags defining the working modes of the solver.

5. Feed the current time and abundances, and the required output time, to the solver, and wait for the output.

6. Retrieve and store the output time (which could be different from the value provided in the previous step) and abundances.

7. Terminate if the output time is equal to or greater than the targeted final evolution time; otherwise go to step 5.

3 Example models based on CHEMPL

In this section we present a few example models calculated with CHEMPL. The chemical networks we use for these examples are the UMIST network, or extended from it. The latest version of the UMIST network is released in 2012, hence it is called RATE12 [McElroy et al. 2013]. The default reaction network format supported by CHEMPL is similar to the KIDA network [Wakelam et al. 2012], instead of the colon-separated format of RATE12, and can support any format with a simple conversion.

3.1 Dark cloud models

We first run a model with physical conditions relevant for dark clouds. The physical parameters are listed in Table 2. The initial abundances are listed in Table 3. The network for this model is UMIST RATE12, extended by only adding the H2 formation reaction on dust grain surface and the adsorption and evaporation of H and H2, and no other adsorption/desorption processes and surface reactions are included.

The code takes 4–5 seconds of CPU time to reach $10^8$ yr. The resulting evolution of a few common species are plotted as solid curves in Fig. 1. As can be seen, the

```python
import chempl
model = chempl.ChemModel(
    fReactions=b'network_file',
    fInitialAbundances=b'abundance_file',
    phy_params={
        b'Av': 10.0,
        b'G0_UV': 1.0,
        b'Ncol_H2': 1e22,
        b'T_dust': 10.0,
        b'T_gas': 10.0,
        b'chemdesorption_factor': 0.05,
        b'chi_cosmicray': 1.0,
        b'dust_albedo': 0.6,
        b'dust2gas_mass': 1e-2,
        b'dust_albedo': 0.6,
        b'dust_material_density': 2.0,
        b'dust_radius': 1.0e-4,
        b'dust_site_density': 1e15,
        b'dv_km_s': 1.0,
        b'mean_mol_weight': 1.4,
        b'n_gas': 2e4})
model.prepare()
model.set_solver()
t = 0.0
dt = 0.1
tRatio = 1.1
tMax = 3.15e10 # in seconds
nMax = 1000
store = {'t': [], 'y': []}
y = model.abundances
for i in range(nMax):
    t, y = model.update(y, t=t, dt=dt)
    store['t'].append(t)
    store['y'].append(y)
    if t >= tMax:
        break
dt *= tRatio
```

Table 1: A minimal working example of a chemical model written in Python using CHEMPL. Tested with Python 3. The Python bytes objects (in the form of b'***') instead of string objects are needed for interacting with internal C++ code.

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2All the example models reported in this paper are computed with a MacBook Pro (15-inch, 2017), with a processor configuration of “3.1 GHz Quad-Core Intel Core i7”.
system reaches steady state at \( \sim 10^7 \) yr, after which carbon mainly resides in CO, while oxygen mainly resides in CO and O\(_2\). However, this model is not realistic at late times (\( \gtrsim 10^6 \) yr), because at the low temperature for this model, CO should freeze out onto dust grains. So this model should be only viewed as a code benchmark and it does not accurately describe the actual dark cloud chemical processes.

So we run another model with the same initial abundances and physical parameters, but with a full three-phase chemical network. Nonthermal desorption (Garrod et al. 2007) by the heat released in surface reactions is also included. With this added complexity, the code takes \( \sim 15 \) seconds of CPU time to reach \( 10^8 \) yr. The results are plotted as dashed curves in Fig. 1. For species such as CO and C, their early-time (\( \lesssim 10^4 \) yr) evolution from the two models coincide. This is because their abundances are mainly determined by gas-phase reactions at early times. For species such as H\(_2\)O, OH, O\(_2\), and NH\(_3\), their evolution tracks from the two models depart away from each other from the very beginning. This is because they are efficiently formed on the dust grain surface, and the inclusion of surface reactions (together with nonthermal desorption) enhances their abundances from the very beginning.

### 3.2 Time-dependent physical conditions

Chemical models in which physical parameters (most frequently, density and temperature) change with time have been used in the study of warm-up phase of hot cores (Garrod & Herbst 2006). Technically, this time-dependency could be implemented with a series of models with constant physical parameters, and the parameters

| Physical parameter | Value         |
|--------------------|---------------|
| \( A_v \)          | 10            |
| \( G_0 \)          | 1             |
| \( \nu_{\text{H}} \) | \( 10^{22} \) cm\(^{-2} \) |
| \( T_{\text{dust}} \) | 10 K         |
| \( T_{\text{gas}} \) | 10 K         |
| Dust-to-gas mass ratio | 1.3 \times 10\(^{-17} \) s\(^{-1} \) |
| Dust albedo         | 0.01          |
| Dust material density | 2.0 g cm\(^{-3} \) |
| Dust grain radius   | 0.1 \( \mu \)m  |
| Dust site density   | \( 10^{15} \) cm\(^{-2} \) |
| Chemical desorption efficiency | 0.05 |

Table 2: Physical parameters used in the example dark cloud model.

![Figure 1: Example results from a basic dark cloud model. Solid curves are calculated without any dust grain chemistry except \( \text{H}_2 \) formation, and dashed curves with the same color are based on a full three-phase chemical network.](image.png)
only gradually change from one model to another. In the series of models the final abundances of a previous model is used as initial condition of the next. However, such an implementation would be computationally inefficient, because the ODE solver would have to reinitialize itself for each individual model, which is a slow process. So it is necessary to add explicit time-dependency directly in a model, specifically, in the right hand side of equation (1) or equation (3).

3.2.1 Continuity of reaction rates

In the RATE12 network each reaction has associated a temperature range in which the rate parameters are most reliable. Some reactions have two temperature ranges, which means different set of rate parameters must be chosen at different temperatures. This could cause problems when temperature varies as a function of time: at the boundary of two temperature ranges, the reaction rate calculated from two versions of rate parameters may not necessarily be continuous, which may confuse the ODE solver and the solution will become unreliable (or the solver will stop working and no solution can be obtained after the boundary of a temperature range is reached).

CHEMPL ensures continuity (smoothness actually) by joining the rates calculated with two sets of parameters using a weighting function of the form

$$g(x) = \frac{1}{1 + e^{\pm(x-x_0)/w}}$$

where $x_0$ is the interval edge, and $w > 0$ is the separation from one interval to another. The $\pm$ sign takes “+” when $x_0$ is the right edge, and takes “−” when $x_0$ is the left edge. The function $g(x)$ has the property that for $x < x_0$, it quickly approaches unity, while for $x > x_0$, it quickly approaches zero. The choice of weighting function is arbitrary, in the sense that a weighting scheme is allowed as far as the transition across the boundary is smooth and without any artificial bumps.

Fig. 2 shows the rate coefficients of two reactions as a function of temperature. As can be seen from the left panel, without a treatment such as the one described here, the rate coefficient would jump from one value to another when the temperature only slightly varies at the boundary (from 100 to 101 K). This may halt the ODE solver since it will have to use exceedingly small time steps (since temperature is a function of time in time-dependent models). With a continuous function, the ODE solver will be able to integrate much easier.

Fig. 2 and Fig. 3 show two example models with time-dependent temperature and density. The reaction network used for them are RATE12 with the addition of adsorption and desorption reactions, while no surface reactions beyond the formation of H$_2$ are included. The time evolution of temperature and density in the two models are arbitrarily designed, while keeping in mind that they may mimic realistic processes occurring in certain interstellar environments. The first model simulates a compressing and heating event, which takes about 40 CPU seconds to reach 10$^7$ yr, while the second one simulates a “wavy” oscillating temperature and density profile, and takes about 16 CPU minutes to reach 10$^7$ yr. The abrupt oscillation of temperature and density, especially at late times, limits the step size used internally in the ODE solver and significantly slows down the integration speed.

Interestingly, the evolution curves plotted in the bottom panel of Fig. 4 demonstrate two types of behaviors: those which follow temperature (CO, O$_2$, and NH$_3$), and those which follow density (H$_2$O, OH). For species of the first type, their abundances are mostly controlled by desorption, hence their abundances jumps when the dust temperature reaches their desorption temperature, and stays flat after that. For species of the second type, either they are not able to evaporate at dust temperatures $\lesssim$40 K (the highest dust temperature in the model), and/or their abundances are mostly determined by gas phase reactions, and their abundances tend to follow the evolution of density. The behavior of C and C$^+$ are more complex. They seem to follow both density and the rise and drop of temperature, which is because their abundances are affected by gas phase reactions which involves species that can evaporate at dust temperature less than 40 K.

3.3 Circumstellar envelope

Since we largely use the UMIST RATE12 network as a starting point for CHEMPL, we also run a model similar to the "carbon-rich circumstellar envelope model" presented in the RATE12 paper (McElroy et al. 2013). This may be viewed as a non-strict benchmark, because a strict one is possible only if the exact parameters (including physical constants) used in their model are adopted. Here we try to use the same input as them, though an exact match is not intended for. The initial abundances and density profile are taken from Table 6 of McElroy et al. (2013), and the temperature profile is taken from Cordiner & Millar (2009). The self-shielding of CO from photodissociation is treated using the approach of Morris & Jura (1983), which is also the same as in McElroy et al. (2013). While this model calculates the spatial structure of a circumstellar envelope around a carbon-rich AGB star, however, since the gas in the envelope are launched from very close
Figure 2: Examples showing how the rate coefficients for reactions with two temperature-range-of-applicability in the RATE12 database are calculated to ensure continuity and smoothness with respect to temperature. The left panel corresponds to reaction $H^- + H \rightarrow H_2 + E^-$, and the right panel corresponds to $NH + O \rightarrow NO + H$. Blue and orange curves are the rate coefficients calculated with expressions for each temperature range as shown in the legends (dashed parts are outside the range of applicability). The red curves are the smoothed rates joining the two used in the code.

Figure 3: Results from a toy model with time-dependent temperature and density. Dashed curves in the bottom panel are results calculated with temperature and density kept constant at their initial values.

Figure 4: Results from a toy model with “wavy” and spiky time-dependent temperature and density.
Figure 5: A simple circumstellar envelope model. The solid curves in the lower two panels are calculated with CHEMPL, while the dashed curves are taken from Fig. 2 of McElroy et al. (2013) using data file generated with their code. The top panel shows the adopted temperature (blue) and density (orange) profile.

to the central star with a fully-specified velocity profile, there is an one-to-one mapping between time and radius. Thus the model can be viewed as one in which physical parameters vary with time, which is how we implement it.

The results of the calculation with CHEMPL are shown as solid curves in Fig. 5 overlaid with data from McElroy et al. (2013) shown as dashed curves. It is clear that the two calculations match very well: two versions of most of the curves are almost identical within numerical errors.

3.4 Spatially dependent physical conditions: a toy PDR chemical model

As a final example, we create a model that simulates a plane-parallel photodissociation region. The density is taken to be uniformly $10^4 \text{ cm}^{-3}$, while the dust temperature profile is calculated using equation (9.18) of Tielens (2005), and the gas temperature is assumed to be uniformly 100 K.

The calculation has to proceed from the surface layer ($A_V \sim 0$) to the deeper layers to take into account the self-shielding of H$_2$ and CO. Here the CO self-shielding is approximated using the same approach as in Section 3.3, while the H$_2$ self-shielding is implemented following Draine & Bertoldi (1996). More accurate treatment of self and mutual shielding can also be implemented (Visser et al., 2009; Li et al., 2013; Heays et al., 2017).

The resulting abundances of different molecules are shown as a function of depth ($A_V$) and column density are shown in Fig. 6. As is well-known, the H/H$_2$ transition occurs at $A_V \sim 1$, while the C$^+$/C/CO transition occurs at $A_V \sim 4$.

3.5 Model-level parallelism

As shown before, when the physical parameters are kept constant, CHEMPL runs relatively fast. So it is not necessary to parallelize each individual model. However, when doing a parameter study, in which many models with different configurations need to be computed, parallelization becomes helpful. In the case of CHEMPL, the current parallelization approach is to make use of the Python built-in “multiprocessing” module. With this module, it is possible to run multiple independent models at the same time, which can accelerate the computational speed by a factor depending on the number of CPU cores of the system. We call this approach of parallelization “model-level” parallelism.

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Since there is no interdependence between the models, this type of parallelism is considered to be “embarrassingly parallel.”
When the physical parameters explicitly change with time, especially when the time variation is oscillatory and spiky, as shown in the example model in Fig. 4, the computational speed will be significantly reduced. We don’t expect an easy solution to alleviate the computation burden here, because the problem is inherently difficult. In principle, one could parallelize the matrix inversion (which is frequently carried out in the solver) in the ODE solver to speed it up, however the effectiveness of doing this remains to be tested.

4 Conclusions

In this paper we introduce a new code named CHEMPL for astrochemical study. In comparison with other codes in the field, CHEMPL emphasize interactivity and “playability”. While CHEMPL can be used as a traditional standalone command line program or called as a subroutine within a bigger simulation, it is mainly meant to be used as a Python module. In the recent years the Python programming language and its ecosystem have become prevalent in astronomical community, and CHEMPL is joining this trend.

CHEMPL is not a completed project. It is still evolving. We expect in the future it will keep improving both in performance and in ease-of-use, as well as in incorporating more chemical and physical mechanisms (for example solving the temperature in tandem with chemical abundances, as done in [Du & Bergin 2014]). One potential feature that might be handy is for the user to be able to choose a different ODE solver. It may also be useful for the chemical solver to be able to switch between the rate equation approach and the Monte Carlo ([Vasyunin et al. 2009] [Chang & Herbst 2016]), the moment equation ([Du & Parise 2011]), or the modified-rate-equation approaches ([Caselli et al. 1998] [Garrod 2008]). This will facilitate the study of the fundamental issue of the stochasticity of grain surface reactions.

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