PLASMAKIN Manual

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# Table of Contents

**Preface** .................................................................................................................................................. i

Abstract ..................................................................................................................................................... i

Intended Audience .................................................................................................................................... i

Other Sources of Information .................................................................................................................... i

Conventions Used in this Manual ............................................................................................................. i

1. **Introduction** ...................................................................................................................................... 1

   What is _PLASMAKIN_ .............................................................................................................................. 1

   Limitations of _PLASMAKIN_ .................................................................................................................... 3

   Units ........................................................................................................................................................ 4

2. **Overview** ........................................................................................................................................ 1

   Usage ....................................................................................................................................................... 1

   Data Model ............................................................................................................................................. 2

3. **Public Parameters and Variables** .................................................................................................... 4

   KTable .................................................................................................................................................. 6

   NCasc ................................................................................................................................................... 7

   NCh ....................................................................................................................................................... 7

   NKrea ................................................................................................................................................... 8

   NKgas .................................................................................................................................................. 9

   NKPhEm .......................................................................................................................................... 9

   NKReac ........................................................................................................................................... 10

   NKRImp .......................................................................................................................................... 11

   NMaxReacSpc .................................................................................................................................. 11

   NnC .................................................................................................................................................. 12

   NnTV ............................................................................................................................................... 13

   NnV .................................................................................................................................................. 13

   NnVX ............................................................................................................................................... 14

   NPhot ............................................................................................................................................. 14

   NReverse ...................................................................................................................................... 15

   NSKea ........................................................................................................................................... 15

   NSpecies ....................................................................................................................................... 16

   NKSurf ......................................................................................................................................... 17

4. **Derived Data Type Definitions** ....................................................................................................... 18

   DATA_COLUMN ................................................................................................................................. 19

   PHYS_PROPERTY ............................................................................................................................... 19

5. **Procedures** .................................................................................................................................... 21

   pkCleanData ..................................................................................................................................... 22
6. Data Input Format ................................................................. 48
   PLASMAKIN_DATA ......................................................... 49
   CHEM_SPECIES ............................................................ 50
   CHEM_REACTION .......................................................... 54
References .................................................................................. 58
A. Error Messages ........................................................................ 59
B. GNU Free Documentation License ........................................ 62
List of Figures

F-1. Relationships between PLASMAKIN and other program units or data files.................................1

List of Examples

E-1. KTable: Using KTable to control the printing sort order of variables........................................6
E-2. pkCleanData: Deallocating PLASMAKIN data........................................................................23
E-3. pkGetParticle: Reading the names of all species into an array..................................................24
E-4. pkGetParticle: Example showing how the returned values depend on the data type of the Value argument..........................................................................................................................................24
E-5. pkGetPhotonEmission: Getting the photon emission distribution in a 1D discharge..............26
E-6. pkGetPowerLosses: Inquiring the power losses using the keyword form for dummy arguments 28
E-7. pkGetReacCoef: Inquiring the electron collision rates.................................................................30
E-8. pkGetReverseCoef: Evaluating the electron superelastic cross sections from the corresponding forward process cross section.........................................................................................................31
E-9. pkGetSources: Computing the source terms in a 1D discharge taking account of gas temperature gradients..........................................................................................................................................33
E-10. pkGetValue: Inquiring the gas temperature and density............................................................35
E-11. pkIsPhotoElec: Computing the photoelectron emission in a 1D discharge model..................36
E-12. pkReadBaseData: Reading the basic data in the input file.......................................................37
E-13. pkReadChemReactions: Reading the chemical reactions in the input file..........................39
E-14. pkReadData: Reading the input file..........................................................................................40
E-15. pkReadSpecies: Reading the basic data in the input file..........................................................41
E-16. pkSetPhoton: Setting the photon density prior to computing the source terms for other species 42
E-17. pkSetReacCoef: Modifying the value of the ‘reverse’ property..................................................44
E-18. pkSetReacCoef: Updating temperature dependent rate coefficients or coefficients defined through an external routine....................................................................................................................44
E-19. pkSetValue: Setting the ratio of the number of molecules in the two first vibrational levels.....46
E-20. PLASMAKIN_DATA: Examples of PLASMAKIN_DATA NAMELIST syntax..........................50
E-21. CHEM_SPECIES: Example of CHEM_SPECIES NAMELIST syntax.....................................53
E-22. CHEM_REACTION: Example of CHEM_REACTION NAMELIST syntax...............................56
Preface

Abstract

*PLASMAKIN* is a software library to handle physical and chemical data used in plasma physics modeling and to compute kinetics data from the reactions taking place in the gas or at the surfaces — particle production and loss rates, photon emission rates and energy exchange rates.

This manual describes all the public variables and routines in *PLASMAKIN* and the data input syntax. It gives all the information needed to use the *PLASMAKIN* library in programs, to write data input files and to understand the errors messages produced by *PLASMAKIN*. However it does not explains the physics and chemistry concepts underlying the library. For that purpose the indicated references should be consulted.

Intended Audience

This manual is intended for programmers with a good background in physics and chemistry or conversely, physicists or chemists with programming skills. In any case a basic understanding of the Fortran 90/95 language is needed.

The reader should be familiar with the physics and chemistry concepts underlying the library and be able to consult the article describing *PLASMAKIN* [Pinhão01].

Other Sources of Information

The main reference for the physics and chemistry processes included in *PLASMAKIN* is an article published in Computer Physics Communications [Pinhão01].

The *PLASMAKIN* library and a sample program and data file can be obtained from the CPC library (http://www.cpc.uk).

The author maintains a mailing list on *PLASMAKIN* news, errors and updates. This mailing list can be subscribed sending an email to the author (mailto:npinhao@itn.pt?subject=Subscribe pk list).
Conventions Used in this Manual

In this manual the following conventions are used:

**expression**

Terms in bold and courier font indicates information supplied by the user

{option1 | option2...}

Braces enclose a list from which you must choose one, but only one, item.

[option] [option1 | option2...]

Square brackets enclose optional items or a list of optional exclusive items.

... 

A horizontal ellipsis means that the preceding item can be repeated

```call pkReadData```

Courier font is used for code samples

value, Ivalue, Lvalue, Rvalue, Cvalue

These words indicate, respectively, a generic type, a Fortran INTEGER, LOGICAL, REAL or CHARACTER value

**IN, OUT, INOUT**

These terms classify arguments that, respectively, are used to provide data to a procedure but are not modified inside it; are used to pass data from the procedure back to the calling program but are undefined on entry; or can both provide data to the procedure and return data to the calling program.
Chapter 1. Introduction

A large number of problems in plasma physics involve the consideration of several chemical species \(^1\) and reactions. The solution of such problems invariably requires the ability to read, classify, sort and manipulate particles and reactions and, frequently, the evaluation of source terms in chemical kinetic equations, the energy transferred by the electrons to the other particles, the gas heating due to collisions or the photon emission spectra.

The handling of these data frequently represents a significant fraction of codes, development and maintenance time and is a source of errors. It would clearly be advantageous to have a package able to deal with these data independently of the number or nature of the species and chemical reactions involved and of the problem being solved or the method used. Such a package could be used as a "black box" moving the description of particles and reactions from code to a data file, isolating the evaluation of terms related to the chemical kinetics from the remaining program, allowing the user to concentrate on the physical problem and on the algorithm to solve it. Once the code developed, it would allow an easy and fast modification and testing of chemical models. PLASMAKIN was developed to fulfill this purpose.

What is PLASMAKIN

PLASMAKIN is a collection of routines to parse an input file describing the species and reactions present in a discharge, handle this data and to compute chemical kinetics data characterizing the discharge. The library supports any number of species and reactions, is independent of the number of dimensions of the problem and can be used both in stationary and time-dependent problems. Although PLASMAKIN has been written for plasma physics problems it can be used or adapted to any field dealing with chemical kinetics.

The user intervention is limited to the preparation of an input file and in making the appropriate calls to PLASMAKIN procedures in a driver program. The input file uses a language close to the physical and chemical notation and the parser is able to detect errors in this file and to deduce the values of several properties, reducing the amount of data that must be written in the file.

A large number of species properties are supported and fully described in CHEM_SPECIES. Vibrational series can be indicated and the corresponding initial relative populations are estimated using a Gordiets-Treanor distribution [Gordiets72]:

\[
\delta_{n,v} = \begin{cases} 
\exp \left[ -v \left( \frac{\epsilon_v}{\varepsilon_n T_e} - (v - 1) \frac{\epsilon_0}{\varepsilon_n T_e} \right) \right], & v \leq v^* \\
\frac{\delta_{n,v^*}}{\exp \left( -\frac{\epsilon_v}{\varepsilon_n T_e} \right)}, & v^* \leq v \leq v^{**} \\
0, & v > v^{**}
\end{cases}
\]
Unimolecular, bimolecular and termolecular reactions can be included.

Optical emission and absorption are examples of unimolecular reaction important in plasma physics. Radiation imprisonment can also be accounted including a constant or space dependent escape factor [Molisch98].

Cascade levels (influencing the population of lower levels by spontaneous emission but whose density is not followed) are supported and the corresponding branching ratios are automatically evaluated.

A large number of rate coefficients have an Arrhenius temperature dependence. However reactions in plasma can have more complex temperature dependencies or, as is the case for electron collision reactions, depend on the electron temperature. To accommodate this, rate coefficients with a power series dependence on temperature in the exponential term can be used in PLASMAKIN:

\[ k_i = \alpha_i^0 T^\alpha_i \exp \left( \sum_{j=1}^{n} \frac{\alpha_j^0}{T^\alpha_j} \right) \]

where T is the electron temperature for electron collision reactions and the gas temperature for other cases. The initial temperature is read in the data file and can be changed during program execution.

As the rate coefficient for forward and reverse reactions are related by the principle of detailed balancing, PLASMAKIN automatically evaluates the reverse rates from the corresponding forward rates.

A large number of reactions with vibrational levels are formally identical, the only difference being the vibrational number(s). To avoid the need of writing all this reactions PLASMAKIN is able to read a single description of a group of reactions involving species from a vibrational series, create the corresponding individual reactions and compute the rate coefficients.

Finally, to compute rate coefficients known as a function of the reduced field E/N or mean energy (as is frequently the case for electron rate coefficients), rate coefficients depending on vibrational quantum numbers or other expressions, a user supplied routine can be used.

Termolecular Reactions - Two types of pressure-dependent reactions are included: recombination reactions and chemically activated bimolecular reactions. Reaction rates are calculated using the [Lindemann22] and [Troe83] formulation. In both cases the 3rd body concentration can be set proportional to the total pressure of to the partial pressure of a selected gas species.

Surface reaction can be included and the indication of surface names allows differentiating the reactions according to the surface.

Energy losses - The energy exchanged in reactions is calculated from the values of the formation enthalpy at the gas temperature. As the gas temperature in cold plasmas is within a few hundred degrees to \( T_0 \), the formation enthalpy at gas temperature is not much higher than the standard value and is approximated by:
Chapter 1. Introduction

\[ H_f(T_g) = H_f^0(T_0) + C_p \Delta T \]

A throughout description of the equations and expressions used in PLASMAKIN can be found in [Pinhão01].

The package is written in standard Fortran 95 as a MODULE unit. However, to allow an easy integration with FORTRAN 77 code only the FORTRAN 77 intrinsic data types are used for public data and PLASMAKIN procedures can be called by FORTRAN 77 code. The usage of Fortran standard naming conventions, error diagnostics and error messages eases the usage of the package.

Limitations of PLASMAKIN

Although PLASMAKIN is prepared to deal with a large diversity of properties and types of chemical reactions and to respond to different problems found in plasma physics, some reaction types are not yet supported or functions found in other chemical kinetic libraries or programs [Kee96, Carver97] have not been included.

The following functions have been intentionally left out:

- It does not include any particle or chemical reaction database. In fact, although there are some reference databases, the number of gases and reactions covered in those databases is still small. Also, different procedures and levels of approximation for the description of species and reactions are common in this field. Consequently the author has decided to leave to the user the selection of data and has instead provided enough flexibility in the data structures in order to accommodate different ways of data representation.

- PLASMAKIN does not provide either any differential equations solver or any other integration procedure. As the number of numerical procedures available is relatively large and in continuous evolution, the selection of the appropriate algorithm is largely problem dependent and many are freely available and well documented the choice of the appropriate algorithm was again left to the user.

- PLASMAKIN does not includes an electron Boltzmann equation solver. Tough the electron and the chemical kinetics are interconnected - the rate coefficients depend on the electron kinetics and the former depends on species concentration - the available methods of solution for the electron Boltzmann equation are valid only in clearly defined conditions and there is no "universal" solver. Again, for problems where both kinetics must be solved together, the choice of the appropriate electron Boltzmann equation solver was left to the user. Nevertheless an electron kinetics library, build on top of PLASMAKIN, and solving the homogeneous electron Boltzmann equation in the classical two term approximation is available from the author (mailto:npinhao@itn.pt?subject=Inquire on Boltzlib) for non-commercial use.
• Finally PLASMAKIN does not use some approximations common in other fields - i.e. the “family approach” used in atmospheric chemistry - as they are seldom used in "gaseous electronics” problems.

The following capabilities are been considered for the next version:

• Support for ionic excited states.
• Account as the 3rd body in three body collisions, of species with different efficiencies in promoting the reaction.
• Support for surface reactions with different adsorbed atoms.
• Evaluation of surface coverage and sticking coefficients.
• Interfaces for C, Pascal and Python mixed language programming.

Units

With a few exceptions the physical properties are handled in PLASMAKIN using S.I. units. The exceptions are the energy, the electron temperature, where in both cases eV is used, and the vibrational frequency where for historical reasons cm$^{-1}$ is used.

However in the input file the values of several physical properties can also be indicated in other units. In this case the unit used must be indicated after the value, according to the syntax:

```
property = value, 'units'
```

The units available for each property are listed in the corresponding reference page.

When these units are found the corresponding value are converted to S.I. units and the conversion factors are saved. When the calling program inquires the value of a given physical property the corresponding conversion factor is used to back-convert to the initial units. Thus the value passed to the calling program is always in the units used in the data file.

The conversion operations and conversion factors are always hidden from the calling program. This allows some flexibility on data input and output and at the same time frees the programmer from elementary but error-prone operations.

Notes

1. In this manual each quantum level is considered as an independent species
2. See, for instance, GAMS or NetLib.

3. In spectroscopy the vibrational frequency is measured in units of $\frac{\nu}{c}$, where $c$ is the velocity of light.
Chapter 2. Overview

Usage

The PLASMAKIN library is a Fortran 90/95 MODULE unit with public and private data and procedures. The library is accessible through a standard USE instruction:

USE PLASMAKIN

The module reads its data from an input file and can call an optional user supplied routine SetRate, to evaluate complex rate reaction expressions, but is otherwise, self-contained.

Figure F-1 represents the relationships between PLASMAKIN and other routines or data files. The arrows indicate the direction of information flow.
Figure F-1. Relationships between PLASMAKIN and other program units or data files.
Data Model

*PLASMAKIN* makes extensive use of derived data types. However only two of them are public and even so they are not intended to be used outside the *PLASMAKIN* MODULE. The only reason why they are public is because they are used in *NAMELIST* reading.

The hiding of data structures guarantees that future modification of the data model has limited, if any, impact in the calling programs. It also considerably simplifies mixed language programming.

Public data include parameters, scalars and arrays variables, the above derived types and procedure interfaces.

With the indicated exception, all the public variables and arguments in *PLASMAKIN* procedures have intrinsic Fortran data types. This allows the use of the library with FORTRAN 77 conforming code with little modifications\(^1\).

Particle or reaction properties are only accessible through access routines allowing a strict control of which data can be inquired or modified by the calling program.

The calling program should not modify the value of any of the public variables as they are extensively used by *PLASMAKIN* to internal processing. The possibility of modification of any of those variables is a potential source of errors. Again this could be avoided by the use of access routines but it would be cumbersome and introduce time penalties as these variables are frequently used in the calling program.

The author has decided to rely on the user experience to avoid this kind of errors.

Notes

1. You need off course a Fortran 95 compiler and the Fortran 90/95 USE instruction to invoke *PLASMAKIN*
3. Public Parameters and Variables

With the exception of the array KTable(:) and the PARAMETER NMaxReacSpc, all the other variables are counters representing the number of species or chemical reactions satisfying some condition.

These counters are interrelated and other useful values can be deduced from the available counters. The following values have not been explicitly defined and can be useful:

- \( N_{\text{of neutral excited species}} = N_{nV} - N_{nVX} \)
- \( N_{\text{of electron collision direct processes}} = N_{Kea} - N_{NSKea} \)
- \( N_{\text{of gas collision processes between atoms and molecules}} = N_{Kgas} - N_{Kea} \)
- \( N_{\text{of reverse gas collision processes between atoms and molecules}} = N_{Reverse} - N_{NSKea} \)

When the input file is read the species and chemical reactions are classified and sorted. The new order follows the classification shown in Figure F-2 and Figure F-3 for, respectively, species and chemical reactions.

**Figure F-2.** Representation of the order of species in PLASMAKIN. The electrons have the highest index of the charged species. The region (a) represents the neutral, excited species.

**Figure F-3.** Representation of the order of reactions in PLASMAKIN. The region (a) and (b) represent respectively, the number of direct and reverse gas collision processes between atoms and
molecules.

**Caution**

The values of these counters are set by *PLASMAKIN* and should not be modified by the user program.
**KTable**

**Name**
KTable — Translation table for reaction coefficient indexes

**Type and Attributes**
*Integer* array with dimension *NKReac*

**Description**

*KTable* is a translation table to map the internal *PLASMAKIN* reaction indexes to the corresponding reaction indexes in the input file.

When the input file is read the reactions are classified and sorted to speed-up further operations. However, the interpretation of several results depends on the relationship with the input file reactions. The array *KTable* holds this information. A typical use of the array *KTable* is in write instructions as in the example below.

**Example**

Example E-1. *KTable*: Using *KTable* to control the printing sort order of variables.

```fortran
REAL(8), ALLOCATABLE :: KRate(:)
...
ALLOCATE( KRate(NKReac) )
...
CALL pkGetReacCoef( 'value', KRate(:), 1 ) ➊
print *, KRate(:) ➋
print *, KRate( KTable(:) ) ➌

➊ The array *KRate* holds the values of the reaction rate coefficients sorted by *PLASMAKIN*.
```
This instruction prints the reaction coefficient values in the order used in PLASMAKIN.

This instruction prints the reaction coefficient values in the same order as reactions are listed in the input file.

**NCasc**

**Name**

NCasc — Number of cascade levels

**Type and Attributes**

Integer

**Description**

NCasc is the number of cascading species - excited species decaying only by spontaneous emission and affecting the population of lower levels, but whose population is not followed.

NCasc is calculated counting in the input file the number of species with the *cascade* flag set to T.
**NCh**

**Name**

NCh — Number of charged species

**Type and Attributes**

Integer

**Description**

NCh is the number of different types of charged species, including positively and negatively charged ions and the electrons.

NCh is calculated counting in the input file the number of species with the value of charge different from 0.

**NKea**

**Name**

NKea — Number of coefficients for electron collision reactions

**Type and Attributes**

Integer

**Description**

NKea is the number of different types of reactions with electrons.
Public Parameters and Variables

NKrea is calculated counting the number of reactions indicated in the input file where the name of at least one species in the reactants list is ‘e’.

**NKgas**

**Name**

NKgas — Number of gas phase reactions

**Type and Attributes**

Integer

**Description**

NKgas is the number of different types of reactions taking place in the gas. NKgas is calculated counting in the input file the number of reactions with the value of locus equal to the default value ‘gas’.

**NKPhEm**

**Name**

NKPhEm — Number of photon emitting reactions

**Type and Attributes**

Integer
**Public Parameters and Variables**

**Description**

NKPhEm is the number of different types of reactions emitting photons. NKPhEm is calculated counting the number of reactions with a photon in the *products* list.

**NKReac**

**Name**

NKReac — Total number of reactions

**Type and Attributes**

Integer

**Description**

NKReac is the total number of different types of reactions, including surface reactions and radiation imprisonment reactions.

NKReac is calculated counting the number of reactions defined in CHEM_REACTION NAMELIST groups. The number of reactions counted in each instruction depends on the value of the *opposing* property and on the type of species indicated in the *reactants* and *products* lists:

- If the *opposing* property equals $T$ the number of reactions is doubled;

- If none of the species indicated in the *reactant* or *product* lists represents a vibrational series and *opposing* = $F$, the instruction represents one reaction. However if any of the species represents a vibrational series, all the corresponding reactions are considered and the number of reactions is given by the equation

$$
\prod_{i} (k_{i}^{max} - k_{i}^{min} + 1) \times (v_{i}^{max} - v_{i}^{min} + 1 - \frac{|k_{i}^{max} + k_{i}^{min}|}{2})
$$
where the product is on the number of vibrational species on the side of the equation with the highest number of vibrational species. For clarity let this side be called $S$ and the other side $F$.

$v^i_{\text{min}}$ and $v^i_{\text{max}}$ are, respectively, the minimum and maximum allowed values for the vibrational quantum number for species $i$ in side $S$. If no other limits are imposed they are the minimum and maximum vibrational values for that series. $k^i_{\text{min}}$ and $k^i_{\text{max}}$ are limiting values for the products vibrational quantum number.

**Notes**

1. In most reactions $S$ is the reactants side and $F$ the products side.

However when the number of vibrational species in products is higher (as in the recombination reaction $A^+ + B^- + M \rightarrow AB(v) + M$) the above equation is valid only if side $S$ is the products side.

**NKRImp**

**Name**

NKRImp — Number of radiation imprisonment reactions

**Type and Attributes**

Integer

**Description**

NKRImp is the number of different types of radiation imprisonment reactions.

NKRImp is calculated counting the number of reactions with a photon in the reactants list but none in the products list.
**NMaxReacSpc**

**Name**

*NMaxReacSpc* — Maximum number of particles taking part or produced on a reaction.

**Type and Attributes**

Integer, parameter

**Description**

*NMaxReacSpc* is the maximum number of *reactants or products* in a reaction. In *PLASMAKIN* this number is a parameter constant equal to 4. However if this number is not appropriate, the user only has to modify this value in the declaration of *NMaxReacSpc* in the *PLASMAKIN* source file and recompile the library.

**NnC**

**Name**

*NnC* — Number of neutral species with constant concentration.

**Type and Attributes**

Integer

**Description**

*NnC* is the number of neutral species with constant concentration.
Public Parameters and Variables

\( \text{NnC} \) is calculated counting the number of species with the value \( \textit{constant} = T \).

**NnTV**

**Name**

\( \text{NnTV} \) — Total number of species with variable concentration.

**Type and Attributes**

Integer

**Description**

\( \text{NnTV} \) is the number of species whose concentration changes and must correspond to the number of continuity equations on the model.

\( \text{NnTV} \) is calculated adding the number of charged species \( \text{NnC} \) with the number of neutral species with variable concentration, \( \text{NnVX} \).

**NnV**

**Name**

\( \text{NnV} \) — Number of non-excited neutrals with variable concentration.

**Type and Attributes**

Integer
Description

\( \text{NnV} \) is the number of non-excited neutral species whose concentration changes.
\( \text{NnV} \) is calculated counting the number of species with \( \text{energy} = 0 \) among the \( \text{NnVX} \) species.

\( \text{NnVX} \)

Name

\( \text{NnVX} \) — Total number of neutrals with variable concentration.

Type and Attributes

Integer

Description

\( \text{NnVX} \) is the number of neutral atomic or molecular species with changing concentration.
\( \text{NnVX} \) is calculated counting the number of neutral, non-constant and non-cascade species.

\( \text{NPhot} \)

Name

\( \text{NPhot} \) — Number of different photon species
Public Parameters and Variables

**Type and Attributes**

Integer

**Description**

\( N_{Phot} \) is the number of different photons species. \( N_{Phot} \) is calculated counting the number of species with the word \textsf{photon} in \textit{name}.

**NReverse**

**Name**

\( N_{Reverse} \) — Number of reverse reaction.

**Type and Attributes**

Integer

**Description**

\( N_{Reverse} \) is the number of reverse reaction processes. \( N_{Reverse} \) is calculated counting the number of reverse reactions defined through CHEM_REACTION NAMELIST groups with \textit{opposing} = \texttt{T}.
Public Parameters and Variables

**NSKea**

**Name**
NSKea — Number of super-elastic electron collision reactions.

**Type and Attributes**
Integer

**Description**
NSKea is the number of superelastic collision reactions. NSKea is calculated counting the number of reverse reactions where one of the reactants has the name ‘e’.

**NSpecies**

**Name**
NSpecies — Total number of species

**Type and Attributes**
Integer

**Description**
NSpecies includes all the species defined in the input file, including photons. However the third-body in three-body collisions, identified by the generic name ‘M’ is not counted.
NSpecies is calculated counting the number of species defined in CHEM_SPECIES NAMELIST instructions including of course the vibrational series.

NKSurf

Name

NKSurf — Number of surface reactions.

Type and Attributes

Integer

Description

NKSurf includes all the reactions defined in CHEM_REACTION NAMELIST instructions with the value of locus different from the default value ‘gas’
4. Derived Data Type Definitions

The following two derived data types are public as they are used in data reading. However, the user has no need to use them to communicate with PLASMAKIN as all the data read by PLASMAKIN can be inquired through access routines using only the intrinsic Fortran data types. These derived data types are documented here only to help understanding the data input format used by PLASMAKIN.
DATA_COLUMN

Name

DATA_COLUMN — Identifies the location of input data on an external file.

Structure Members

```
TYPE DATA_COLUMN
  CHARACTER(20) :: name ! Name of datafile
  INTEGER :: index ! Data index (column or block n°)
END TYPE
```

Description

The DATA_COLUMN derived type is used in the input file to address data located in other files. The string represents a file name and the index identifies the location of data in that file.

Frequently the index is used to identify a single column or data block. However if more than one column or data block must be read - as in the case of \((x_{1}, x_{2}, \ldots, y)\) values - the calling program is responsible for selecting the correct columns or data blocks based on this index. Usually the values are organized in consecutive columns or data blocks but the used is free to choose how to organize the data. The drawback is that he is also responsible by reading this data.

Example

In the following example the variable `data_file` has type DATA_COLUMN and is used to input the filename `eTransport.txt` and the index value 1. In this case the file holds the electron transport properties (mobility and diffusion coefficient) as a function of the reduced electrical field.

```
&CHEM_SPECIES name = 'e', charge = -1, data_file = 'eTransport.txt',1 /
```
**PHYS_PROPERTY**

**Name**

PHYS_PROPERTY — Represents a physical property

**Structure Members**

```fortran
TYPE PHYS_PROPERTY
    REAL(double) :: value ! value of physical property
    CHARACTER(10) :: units ! units of physical property
END TYPE PHYS_PROPERTY
```

**Description**

The PHYS_PROPERTY derived type is used in the PLASMAKIN input file to read physical properties. Two fields compose it, one for the numerical value and the other for the unit.

It is used only when a given property can be indicated in more than one unit. Otherwise only the numerical value of the property is used.

**Example**

In the following example all the variables in the NAMELIST have type PHYS_PROPERTY and both the value and the unit can be indicated. In this example although the value of the variable Gas_n, representing the gas density, is not indicated, the unit is needed to force the writing of results in cm$^{-3}$.

```plaintext
&PLASMAKIN_DATA Pressure = 0.15,'mbar', Gas_n = ,'cm-3', Gas_T = 300,'K' /
```

---

Derived Data Type Definitions
5. Procedures
pkCleanData

Name

pkCleanData — Deallocates arrays destroying all the information on chemical data

Description

The subroutine pkCleanData destroys all private arrays of PLASMAKIN. These arrays are used to hold the chemical data and are created when the data file is read.

Syntax

pkCleanData( [STAT] )

<table>
<thead>
<tr>
<th>Name</th>
<th>Description</th>
<th>Data type and attributes</th>
</tr>
</thead>
<tbody>
<tr>
<td>STAT</td>
<td>Execution status indicator</td>
<td>INTEGER, OPTIONAL, INTENT(OUT)</td>
</tr>
</tbody>
</table>

Results

STAT = 0: no errors; STAT = n: error number.

Comments

Why is this subroutine needed?

• Although most modern compilers are able to manage the memory efficiently and avoid memory leaks, it’s a good programming practice to clean all arrays before leaving a routine or program.

• If you wish to build a program that uses more than one input data file (i.e. for batch processing) you must call this routine to clean the internal arrays before reading the next data file.
Examples

Example E-2. pkCleanData: Deallocating PLASMAKIN data.

INTEGER :: ErrorStatus
CHARACTER, PARAMETER :: StopCode = 'pkError: Memory deallocation error'
...
CALL pkCleanData( ErrorStatus )
IF( ErrorStatus /= 0 ) STOP StopCode

pkGetParticle

Name

pkGetParticle — Get the values of chemical species properties

Description

The subroutine pkGetParticle returns the value of a given property for one species or the values of that property for a sequence of species.

Syntax

pkGetParticle( Property, Value, Species[, STAT] )

<table>
<thead>
<tr>
<th>Name</th>
<th>Description</th>
<th>Data type and attributes</th>
</tr>
</thead>
</table>
| Property | Case sensitive name of property | CHARACTER(len=*)
| {Value | Values(s) of property            | (INTEGER |REAL(double))
| Value(:) |                                  | , INTENT(OUT)                           |
**Procedures**

<table>
<thead>
<tr>
<th>Name</th>
<th>Description</th>
<th>Data type and attributes</th>
</tr>
</thead>
<tbody>
<tr>
<td>SpeciesN</td>
<td>{Index of \textit{Species}</td>
<td>starting index of \textit{Species}}</td>
</tr>
<tr>
<td>STAT</td>
<td>Execution status indicator</td>
<td>INTEGER, OPTIONAL, INTENT(OUT)</td>
</tr>
</tbody>
</table>

**Results**

The Value argument can be a scalar or an array. In the first case the value returned is the value of \textit{Property} of \textit{Species}(SpeciesN). In the last case the values returned are the \textit{Property} values from consecutive species from \textit{Species}(SpeciesN) to \textit{Species}(SpeciesN+\text{SIZE(Value)}-1).

When the properties are of a derived-type, the result depends on the intrinsic type of the dummy argument Value and is the derived-type component with the same type as Value.

\texttt{STAT = 0} : no errors; \texttt{STAT = n} : error number.

**Examples**

**Example E-3. pkGetParticle:** Reading the names of all species into an array.

```fortran
CHARACTER(20), ALLOCATABLE :: SpcName(:) ➊

...  
ALLOCATE( SpcName(NSpecies) )  
...  
CALL pkGetParticle( ‘name’, SpcName, 1 ) ➋
```

➊ The array \texttt{SpcName} will hold the names of all species  
➋ The names of all species are assigned to the array \texttt{SpcName} following the order used by \textit{PLASMAKIN}
Example E-4. *pkGetParticle*: Example showing how the returned values depend on the data type of the *Value* argument.

```
CHARACTER(20) :: SpcFileName
INTEGER :: SpcFileIndex
...
NAMELIST /TransportData/ Vd, D
...
! Reads transport properties in an external file
DO i = NnC+1, NnC+NnTV
    CALL pkGetParticle( ’data_file’, SpcFileName, i )
    CALL pkGetParticle( ’data_file’, SpcFileIndex, i )
OPEN( IOUnit, SpcFileName, ACTION=’READ’ )
DO j = 1, SpcFileIndex ! Reads file until finding the correct index
    READ( IOUnit, NML=TransportData, IOSTAT )
END DO
CLOSE( IOUnit )
END DO
```

1. The variable *SpcFileName* will hold the name of the external file listing the transport parameters for the inquired *Species*.
2. The variable *SpcFileIndex* indexes the transport data for each *Species* in the external file.
3. In this case the value returned is a character string.
4. In this case the value returned is an integer.

**pkGetPhotonEmission**

**Name**

*pkGetPhotonEmission* — Evaluates photon emission rates
Description

The subroutine `pkGetPhotonEmission` returns the photon energy and the photon emission rate for each photon.

Syntax

`pkGetPhotonEmission( LDensity, PhRate[, STAT] )`

<table>
<thead>
<tr>
<th>Name</th>
<th>Description</th>
<th>Data type and attributes</th>
</tr>
</thead>
<tbody>
<tr>
<td>LDensity(NnTV)</td>
<td>Local density of species with variable concentration</td>
<td>REAL(double), INTENT(IN)</td>
</tr>
<tr>
<td>PhRate(2,NPhot)</td>
<td>Photon energy and local photon emission rate for all photons</td>
<td>REAL(double), INTENT(OUT)</td>
</tr>
<tr>
<td>STAT</td>
<td>Execution status indicator</td>
<td>INTEGER, OPTIONAL, INTENT(OUT)</td>
</tr>
</tbody>
</table>

Results

The array section `PhRate(1,:)` holds the photon energies and the array section `PhRate(2,:)` the photon emission rates.

`STAT = 0`: no errors; `STAT = n`: error number.

Examples

Example E-5. `pkGetPhotonEmission`: Getting the photon emission distribution in a 1D discharge

```fortran
REAL(double), ALLOCATABLE :: Dens(:,:), PhDist(:,:), PhRate(:,:)
...
ALLOCATE( Dens(NnTV,Nx), PhRDist(NPhot,Nx), PhRate(2,NPhot) )
...
DO i = 1, Nx
   CALL pkGetPhotonEmission( Dens(:,1), PhRate )
   PhRDist(:,i) = PhRate(2,:)
END DO
```
**pkGetPowerLosses**

**Name**

pkGetPowerLosses — Evaluates the power losses and the relative contribution of each process

**Description**

The subroutine pkGetPowerLosses computes the electron collisional power losses, the power converted into heat in other collision processes, the power radiated and the relative contribution of each reaction for these losses.

**Syntax**

```
pkGetPowerLosses( LDensity[, eP[, eR][, HeatP][, HeatR][, RadP][, RadR][, STAT] )
```

<table>
<thead>
<tr>
<th>Name</th>
<th>Description</th>
<th>Data type and attributes</th>
</tr>
</thead>
<tbody>
<tr>
<td>LDensity(NnTV)</td>
<td>Local density of species with variable concentration</td>
<td>REAL(double), INTENT(IN)</td>
</tr>
<tr>
<td>eP</td>
<td>Power lost by the electrons by unit volume</td>
<td>REAL(double), OPTIONAL,</td>
</tr>
<tr>
<td></td>
<td>INTENT(OUT)</td>
<td></td>
</tr>
<tr>
<td>eR(NKea)</td>
<td>Relative contribution of each electron collision reaction to the</td>
<td>REAL(double), OPTIONAL,</td>
</tr>
<tr>
<td></td>
<td>electron power losses</td>
<td>INTENT(OUT)</td>
</tr>
<tr>
<td>HeatP</td>
<td>Power converted into heat in other collision processes by unit volume</td>
<td>REAL(double), OPTIONAL,</td>
</tr>
<tr>
<td></td>
<td>INTENT(OUT)</td>
<td></td>
</tr>
<tr>
<td>HeatR(NKgas)</td>
<td>Relative contribution of each reaction to gas heating</td>
<td>REAL(double), OPTIONAL,</td>
</tr>
<tr>
<td></td>
<td>INTENT(OUT)</td>
<td></td>
</tr>
<tr>
<td>RadP</td>
<td>Radiated power by unit volume</td>
<td>REAL(double), OPTIONAL,</td>
</tr>
<tr>
<td></td>
<td>INTENT(OUT)</td>
<td></td>
</tr>
</tbody>
</table>
**Procedures**

<table>
<thead>
<tr>
<th>Name</th>
<th>Description</th>
<th>Data type and attributes</th>
</tr>
</thead>
<tbody>
<tr>
<td>RadR (NPhot)</td>
<td>Relative contribution of each photon emission process to the radiated power</td>
<td>REAL(double), OPTIONAL, INTENT(OUT)</td>
</tr>
<tr>
<td>STAT</td>
<td>Execution status indicator</td>
<td>INTEGER, OPTIONAL, INTENT(OUT)</td>
</tr>
</tbody>
</table>

**Results**

The power/volume losses are in Watt/L^3, where L^3 is the unit used to indicate the gas density in the input file.

STAT = 0: no errors; STAT = n: error number.

**Examples**

**Example E-6. pkGetPowerLosses:** Inquiring the power losses using the keyword form for dummy arguments.

```plaintext
REAL(double) :: ePLosses, HeatPLosses, RadPLosses
...
CALL pkGetPowerLosses( eP=ePLosses, HeatP=HeatPLosses, RadP=RadPLosses )
```

**pkGetReacCoef**

**Name**

pkGetReacCoef — Get the values of reaction coefficients properties
**Procedures**

### Description

The subroutine `pkGetReacCoef` returns the value of a given reaction coefficient property. If `Value` is an array, a list of values is returned.

### Syntax

```fortran
pkGetReacCoef( Property, Value, KReac[, STAT] )
```

<table>
<thead>
<tr>
<th>Name</th>
<th>Description</th>
<th>Data type and attributes</th>
</tr>
</thead>
<tbody>
<tr>
<td>Property</td>
<td>Case sensitive name of property</td>
<td>CHARACTER(len=*) , INTENT(IN)</td>
</tr>
<tr>
<td>`{Value</td>
<td>Value(:)</td>
<td>Value(:,:)}`</td>
</tr>
<tr>
<td>KReacN</td>
<td>Index of reaction inquired if <code>Value</code> is a scalar or starting index of reaction if <code>Value</code> is an array</td>
<td>INTEGER, INTENT(IN)</td>
</tr>
<tr>
<td>STAT</td>
<td>Execution status indicator</td>
<td>INTEGER, OPTIONAL, INTENT(OUT)</td>
</tr>
</tbody>
</table>

### Results

The dummy argument `Value` returns the value of the property inquired and can be a scalar, a one or a two-dimension array. A two-dimension array is only used to inquire the name of the species involved in a reaction as `reactants` or `products`.

If the dummy argument `Value` is a scalar, the result is the value of the property for species `KReacN`. If `Value` is a one dimension array, the returned values are the property values from consecutive `SIZE(Value, DIM=1)` species, starting in species `KReacN`. If `Value` is a two-dimensional array, for each reaction inquired, the returned values are the `SIZE(Value, DIM=2)` first names of the species involved in the reaction as `reactants` or `products`. If `SIZE(Value, DIM=2) > NMaxReacSpc`, the values with index higher than `NMaxReacSpc` are returned empty.

`STAT = 0`: no errors; `STAT = n`: error number.
Examples


```fortran
REAL(double), ALLOCATABLE :: eKRates(:)
...
ALLOCATE( eKRates(NKea) )
...
! The electron collision rates are the first NKea rates
CALL pkGetReacCoef( 'Value', eKRates, 1 )
```

**pkGetReverseCoef**

**Name**

`pkGetReverseCoef` — Get the values of reverse reactions properties

**Description**

The subroutine `pkGetReverseCoef` returns a list of values of a given property for reverse reactions. Two properties are available:

- The ratio, in the forward reaction, between the product of the degeneracy for the *products* species and the product of the degeneracy for the *reactants* species, -`'g_ratio'`
- The energy difference between the products and the reactants in the forward reaction -`'dE'`

**Syntax**

`pkGetReverseCoef( Property, Value, KReacN[, STAT] )`
### Procedures

<table>
<thead>
<tr>
<th>Name</th>
<th>Description</th>
<th>Data type and attributes</th>
</tr>
</thead>
<tbody>
<tr>
<td>Property</td>
<td>Case sensitive name of property</td>
<td>CHARACTER(len=*), INTENT(IN)</td>
</tr>
<tr>
<td>Value(:)</td>
<td>Values of property</td>
<td>REAL(double), INTENT(OUT)</td>
</tr>
<tr>
<td>KReacN</td>
<td>Starting index of reactions inquired</td>
<td>INTEGER, INTENT(IN)</td>
</tr>
<tr>
<td>STAT</td>
<td>Execution status indicator</td>
<td>INTEGER, OPTIONAL, INTENT(OUT)</td>
</tr>
</tbody>
</table>

### Results

The values returned are the values of the property inquired from consecutive `SIZE(Value)` reverse reactions starting in species `KReacN`.

STAT = 0: no errors; STAT = n: error number.

### Examples

**Example E-8. pkGetReverseCoef:** Evaluating the electron superelastic cross sections from the corresponding forward process cross section.

```fortran
INTEGER :: NfKea, & ! N of forward e-atom reactions NEbin, & ! N of energy bins idE, & ! Index for the energy diff. between levels jmax ! Maximum index for superelastic CS REAL(double) :: Ebin ! Energy bin REAL(double), & ALLOCATABLE :: g2_g1(:), & ! Ratio between degeneracy values dE(:), & ! Energy difference between levels CrossSec(:,:) ! Electron collision cross sections ... NfKea = NKea - NSkea ALLOCATE( CrossSec(NEbin,NfKea) ) IF( NSkea > 0 ) THEN ! - Gets g_ratio and dE for superelastic processes CALL pkGetReverseCoef( 'g_ratio', g2_g1, NfKea+1 ) CALL pkGetReverseCoef( 'dE', dE, NfKea+1 ) ! - Computes superelastic cross sections DO i = NfKea + 1, NfKea idE = NINT(dE(i)/Ebin); jmax = NEbin-idE FORALL( j = 1:jmax ) & CrossSec(j,i) = (1.d0+dE(i)/E(j)) / g2_g1(i) & * CrossSec(j+idE,Nindex(i))
```
**pkGetSources**

**Name**

pkGetSources — Get the source terms for the continuity equations and the relative contribution of each process

**Description**

The subroutine *pkGetSources* evaluates the collisional source and loss terms of the continuity equations for each species. The relative contribution of each reaction can also be evaluated.

**Syntax**

```
pkGetSources( LDensity, SrcC[, SrcP][, Ratios][, locus][, STAT] )
```

<table>
<thead>
<tr>
<th>Name</th>
<th>Description</th>
<th>Data type and attributes</th>
</tr>
</thead>
<tbody>
<tr>
<td>LDensity(NnTV)</td>
<td>Local density of species with variable concentration</td>
<td>REAL(double), INTENT(IN)</td>
</tr>
<tr>
<td>SrcC(NnTV)</td>
<td>Collisional gain term of continuity equations</td>
<td>REAL(double), INTENT(OUT)</td>
</tr>
<tr>
<td>SrcP(NnTV)</td>
<td>Collisional loss term of continuity equations divided by LDensity</td>
<td>REAL(double), OPTIONAL, INTENT(OUT)</td>
</tr>
<tr>
<td>Ratios(NKgas,NnTV)</td>
<td>Percent contribution of each reaction to the collisional source and loss terms of each species</td>
<td>REAL(double), OPTIONAL, INTENT(OUT)</td>
</tr>
<tr>
<td>locus</td>
<td>Identifies where the reactions take place</td>
<td>CHARACTER(10), OPTIONAL, INTENT(IN)</td>
</tr>
</tbody>
</table>
**Procedures**

<table>
<thead>
<tr>
<th>Name</th>
<th>Description</th>
<th>Data type and attributes</th>
</tr>
</thead>
<tbody>
<tr>
<td>STAT</td>
<td>Execution status indicator</td>
<td>INTEGER, OPTIONAL, INTENT(OUT)</td>
</tr>
</tbody>
</table>

**Results**

The gain term is in $L^{-3}s^{-1}$ where $L^{-3}$ is the unit used to indicate the gas density in the data file; the loss term is in $s^{-1}$.

$\text{STAT} = 0$: no errors; $\text{STAT} = n$: error number.

**Comments**

If $\text{locus}$ is not present or is equal to ‘gas’ the reactions are in the gas. Otherwise the reactions are on a surface and the value of $\text{locus}$ identifies the surface. This allows having different reactions in different surfaces.

**Examples**

*Example E-9. pkGetSources:* Computing the source terms in a 1D discharge taking account of gas temperature gradients.

```fortran
INTEGER :: i, Nxp
REAL(double) :: Tgas(Nxp)
REAL(double), ALLOCATABLE :: Dens(:,,:), SrcC(:,,:), SrcP(:,,:)
...
ALLOCATE( Dens(NnTV,Nxp), SrcC(nnTV,Nxp), SrcP(NnTV,Nxp) )
...
DO i = 1, Nxp
   CALL pkSetValue( GasTemp=Tgas(i) )
   CALL pkGetSources( Dens(:,i), SrcC(:,i), SrcP(:,i) )
END DO
```
**pkGetValue**

**Name**

pkGetValue — Get the values of global properties

**Description**

The subroutine pkGetValue returns the gas temperature, electron temperature and total gas density.

**Syntax**

pkGetValue( [GasTemp][[,] eTemp][[,] GasN][[,] STAT] )

<table>
<thead>
<tr>
<th>Name</th>
<th>Description</th>
<th>Data type and attributes</th>
</tr>
</thead>
<tbody>
<tr>
<td>GasTemp</td>
<td>The gas temperature</td>
<td>REAL(double), OPTIONAL,</td>
</tr>
<tr>
<td></td>
<td></td>
<td>INTENT(OUT)</td>
</tr>
<tr>
<td>eTemp</td>
<td>The electron temperature</td>
<td>REAL(double), OPTIONAL,</td>
</tr>
<tr>
<td></td>
<td></td>
<td>INTENT(OUT)</td>
</tr>
<tr>
<td>GasN</td>
<td>The total gas density</td>
<td>REAL(double), OPTIONAL,</td>
</tr>
<tr>
<td></td>
<td></td>
<td>INTENT(OUT)</td>
</tr>
<tr>
<td>STAT</td>
<td>Execution status indicator</td>
<td>INTEGER, OPTIONAL,</td>
</tr>
<tr>
<td></td>
<td></td>
<td>INTENT(OUT)</td>
</tr>
</tbody>
</table>

**Results**

Values are returned using the units indicated in the input file for these properties. If for a given property no units were explicitly indicated, the corresponding value is returned in S.I. units.  

$STAT = 0$: no errors; $STAT = n$: error number.
Examples

Example E-10. `pkGetValue`: Inquiring the gas temperature and density.

```fortran
REAL(double) :: gasT, gasDens
...
CALL pkGetValue( gasT, GasN=gasDens )
```

**pkIsPhotoElec**

**Name**

`pkIsPhotoElec` — Tests if a photon produces photoelectric emission on a given surface

**Description**

The function `pkIsPhotoElec` is used to test if a given photon induces photoelectron emission in a given surface. Additionally, it also returns the index of the parent species. The main use of this function is in computing the photoelectron emission from the volume distribution of excited species.

**Syntax**

```fortran
pkIsPhotoElec( PhotN, Surf, NParent, KParent )
```

<table>
<thead>
<tr>
<th>Name</th>
<th>Description</th>
<th>Data type and attributes</th>
</tr>
</thead>
<tbody>
<tr>
<td>PhotN</td>
<td>Photon index</td>
<td>INTEGER(IN)</td>
</tr>
<tr>
<td>Surf</td>
<td>Surface name</td>
<td>CHARACTER(len=*) , INTENT(IN)</td>
</tr>
<tr>
<td>NParent</td>
<td>Index of parent species</td>
<td>INTEGER, INTENT(OUT)</td>
</tr>
<tr>
<td>KParent</td>
<td>Index of photoelectric emission reaction</td>
<td>INTEGER, INTENT(OUT)</td>
</tr>
</tbody>
</table>
**Procedures**

**Results**

The function returns a LOGICAL value.

Using a name to identify the surface allows to compute the photoelectron emission only in selected surfaces and to use different values of the photoelectron emission coefficient in different surfaces.

**Examples**

**Example E-11. pkIsPhotoElec: Computing the photoelectron emission in a 1D discharge model.**

In this example the matrix MPhot represents the probability of transmission of radiation from a plane \( i \), perpendicular to the surface, to a surface "slice" \( j \):

```fortran
INTEGER :: dx, NParent, KParent, iPh, i
REAL(double) :: MPhot(Nxp,Nxp), PhotElectrons(Nxp), KYield
...
PhotElectrons(:) = 0.d0
DO iPh = 1, NPhot
  IF( pkIsPhotoElec( iPh, 'wall', NParent, KParent ) ) THEN
    CALL pkGetReacCoef('Value', KYield, KParent )
    DO i = 1, Nxp
      DO j = 1, Nxp
        dx = ABS(j-i)+1
        PhotElectrons(i) = PhotElectrons(i) + &
        KYield * Dens(j,KParent) * MPhot(i,dx)
      END DO
    END DO
  END IF
END DO
```

**pkReadBaseData**

**Name**

`pkReadBaseData` — Reads the basic data in the input file
**Description**

The subroutine `pkReadBaseData` reads in the input file the **NAMELIST** instructions characterizing base properties as gas temperature, density or pressure and electron temperatures.

**Syntax**

```
pkReadBaseData( UNIT[, IOSTAT] )
```

<table>
<thead>
<tr>
<th>Name</th>
<th>Description</th>
<th>Data type and attributes</th>
</tr>
</thead>
<tbody>
<tr>
<td>UNIT</td>
<td>Unit number for the input file</td>
<td>INTEGER, INTENT(IN)</td>
</tr>
<tr>
<td>IOSTAT</td>
<td>Execution status indicator</td>
<td>INTEGER, OPTIONAL, INTENT(OUT)</td>
</tr>
</tbody>
</table>

**Results**

\[ IOSTAT = 0: \text{no errors}; \quad IOSTAT = n: \text{error number} \]

**Comments**

The calling program is responsible by opening and closing the input file.

**Examples**

**Example E-12. pkReadBaseData:** Reading the basic data in the input file.

```
OPEN( UNIT=10, FILE='TestData.txt', ACTION='READ' )
CALL pkReadBaseData( 10 )
```
**pkReadChemReactions**

**Name**

pkReadChemReactions — Reads only the chemical reactions in the input file

**Description**

The subroutine pkReadChemReactions reads in the input file the NAMELIST instructions characterizing the chemical reactions and process this information. The reactions are counted, classified, reactions involving vibrational series are split into individual reactions; the reactions are sorted according to the classification of Figure F-3; temperature dependent and reverse reactions rates are evaluated; branching ratios for cascade reactions are computed; etc.

**Syntax**

```
pkReadChemReactions( UNIT[, IOSTAT] )
```

<table>
<thead>
<tr>
<th>Name</th>
<th>Description</th>
<th>Data type and attributes</th>
</tr>
</thead>
<tbody>
<tr>
<td>UNIT</td>
<td>Unit number for the input file</td>
<td>INTEGER, INTENT(IN)</td>
</tr>
<tr>
<td>IOSTAT</td>
<td>Execution status indicator</td>
<td>INTEGER, OPTIONAL, INTENT(OUT)</td>
</tr>
</tbody>
</table>

**Results**

IOSTAT = 0: no errors; IOSTAT = n: error number.

**Comments**

The calling program is responsible by opening and closing the input file.
Examples

Example E-13. pkReadChemReactions: Reading the chemical reactions in the input file.

OPEN( UNIT=10, FILE='TestData.txt', ACTION='READ' )
CALL pkReadChemReactions( 10 )

pkReadData

Name

pkReadData — Reads all data in the input file

Description

The subroutine pkReadData reads in the input file all the NAMELIST instructions characterizing the data handled by PLASMAKIN and process these informations. pkReadData is a wrapper routine calling the other data reading routines.

Syntax

pkReadData( UNIT[, IOSTAT] )

<table>
<thead>
<tr>
<th>Name</th>
<th>Description</th>
<th>Data type and attributes</th>
</tr>
</thead>
<tbody>
<tr>
<td>UNIT</td>
<td>Unit number for the input file</td>
<td>INTEGER, INTENT(IN)</td>
</tr>
<tr>
<td>IOSTAT</td>
<td>Execution status indicator</td>
<td>INTEGER, OPTIONAL, INTENT(OUT)</td>
</tr>
</tbody>
</table>
**Results**

$\text{IOSTAT} = 0$: no errors; $\text{IOSTAT} = n$: error number.

**Comments**

The calling program is responsible by opening and closing the input file.

**Examples**

Example E-14. **pkReadData**: Reading the input file.

```
OPEN( UNIT=10, FILE='TestData.txt', ACTION='READ' )
CALL pkReadData( 10 )
```

**pkReadSpecies**

**Name**

pkReadSpecies — Reads only the species in the input file

**Description**

The subroutine pkReadSpecies reads in the input file the **NAMELIST** instructions characterizing the chemical species and process this information. The species are counted; classified; vibrational series are split into individual species and species concentration calculated; the species are sorted according to the classification of Figure F-2; etc.
**Syntax**

```fortran
pkReadSpecies( UNIT[, IOSTAT] )
```

<table>
<thead>
<tr>
<th>Name</th>
<th>Description</th>
<th>Data type and attributes</th>
</tr>
</thead>
<tbody>
<tr>
<td>UNIT</td>
<td>Unit number for the input file</td>
<td>INTEGER, INTENT(IN)</td>
</tr>
<tr>
<td>IOSTAT</td>
<td>Execution status indicator</td>
<td>INTEGER, OPTIONAL, INTENT(OUT)</td>
</tr>
</tbody>
</table>

**Results**

IOSTAT = 0: no errors; IOSTAT = n: error number.

**Comments**

The calling program is responsible by opening and closing the input file.

**Examples**

**Example E-15. pkReadSpecies: Reading the basic data in the input file.**

```fortran
OPEN( UNIT=10, FILE='TestData.txt', ACTION='READ' )
CALL pkReadSpecies( 10 )
```

---

**pkSetPhoton**

**Name**

pkSetPhoton — Sets local photon density
Description

PLASMAKIN does not evaluate photon transport. However, to compute reactions involving photons such as the photoelectric emission from surfaces of photoabsorption, the local photon density must be known. This routine allows setting the photon density values.

Syntax

pkSetPhoton( PhotDensity, NPhSrc[, STAT] )

<table>
<thead>
<tr>
<th>Name</th>
<th>Description</th>
<th>Data type and attributes</th>
</tr>
</thead>
<tbody>
<tr>
<td>PhotDensity(NPhSrc)</td>
<td>Photon density values</td>
<td>REAL(double), INTENT(IN)</td>
</tr>
<tr>
<td>NPhSrc</td>
<td>Size of PhotDensity</td>
<td>INTEGER, INTENT(IN)</td>
</tr>
<tr>
<td>STAT</td>
<td>Execution status indicator</td>
<td>INTEGER, OPTIONAL, INTENT(OUT)</td>
</tr>
</tbody>
</table>

Results

STAT = 0: no errors; STAT = n: error number.

Examples

Example E-16. pkSetPhoton: Setting the photon density prior to computing the source terms for other species.

```fortran
INTEGER :: Nxp, i
REAL(double), ALLOCATABLE :: Dens(:,,:), SrcC(:,,:), SrcP(:,,:), PhDens(:,,:)
...
ALLOCATE( Dens(NnTV,Nxp), SrcC(NnTV,Nxp), &
          SrcP(NnTV, Nxp), PhDens(NPhSrc,Nxp) )
...
CALL COMPUTE_PH_TRANSPORT( PhDens)  ! User supplied routine
DO i = 1, Nxp
   CALL pkSetPhoton( PhDens(:,i), NPhSrc )
   CALL pkGetSources( Dens(:,i), SrcC(:,i), SrcP(:,i) )
END DO
```
**pkSetReacCoef**

**Name**

pkSetReacCoef — Sets the values of reaction properties

**Description**

The subroutine pkSetReacCoef allows setting the values of selected reaction properties. The properties that can be set are ‘reverse’ and ‘value’. This allows to use user defined expressions for the reaction coefficients and is particularly useful for reactions involving vibrational levels.

**Syntax**

pkSetReacCoef( Property, Value, KReacN[, STAT] )

<table>
<thead>
<tr>
<th>Name</th>
<th>Description</th>
<th>Data type and attributes</th>
</tr>
</thead>
<tbody>
<tr>
<td>Property</td>
<td>Case sensitive name of property</td>
<td>CHARACTER(len=*) , INTENT(IN)</td>
</tr>
<tr>
<td>{Value</td>
<td>Value(:)}</td>
<td>Value(s) of property</td>
</tr>
<tr>
<td>KReacN</td>
<td>Starting index of reaction list</td>
<td>INTEGER, INTENT(IN)</td>
</tr>
<tr>
<td>STAT</td>
<td>Execution status indicator</td>
<td>INTEGER, OPTIONAL, INTENT(OUT)</td>
</tr>
</tbody>
</table>

**Results**

STAT = 0: no errors; STAT = n: error number.
Comments

The value argument sets the value of the property inquired and can be a scalar or an array. In the last case the values set are the \text{SIZE(value)} consecutive values of property.

Examples

Example E-17. \texttt{pkSetReacCoef}: Modifying the value of the 'reverse' property.

The following example is a continuation of Example E-8. In that example the superelastic electron collision cross sections were evaluated from the corresponding cross sections for the forward processes. Consequently we no longer need to consider those reactions as a reverse process and the reverse property is set to \texttt{FALSE}:

\begin{verbatim}
IF( NSKea > 0 ) THEN
  ! See code in pkGetReacCoef routine
  CALL pkSetReacCoef( 'reverse', (/(.FALSE.,i=1,NSKea)/), NdKea+1 )
END IF
\end{verbatim}

Example E-18. \texttt{pkSetReacCoef}: Updating temperature dependent rate coefficients or coefficients defined through an external routine:

\begin{verbatim}
INTEGER :: NKaa
REAL(double), ALLOCATABLE :: Kvalue(:)
...
NKaa = NKgas - NKea
ALLOCATE( Kvalue(NKaa) )
...
CALL pkGetReacCoef( 'value', Kvalue, NKaa )
CALL pkSetValue( GasTemp=GasT )
CALL pkSetReacCoef( 'value' Kvalue, NKea+1 )
\end{verbatim}
Procedures

pkSetValue

Name

pkSetValue — Sets the values of global properties

Description

The subroutine pkSetValue sets the values of several properties: gas or electron temperature, total gas density and vibrational temperature for molecular species.

Syntax

pkSetValue( [GasTemp][[, eTemp][[, GasN][[, N1byN0]]][[,] SpeciesN][[, STAT]] ])

<table>
<thead>
<tr>
<th>Name</th>
<th>Description</th>
<th>Data type and attributes</th>
</tr>
</thead>
<tbody>
<tr>
<td>GasTemp</td>
<td>The gas temperature</td>
<td>REAL(double), OPTIONAL, INTENT(IN)</td>
</tr>
<tr>
<td>eTemp</td>
<td>The electron temperature</td>
<td>REAL(double), OPTIONAL, INTENT(IN)</td>
</tr>
<tr>
<td>GasN</td>
<td>The total gas density</td>
<td>REAL(double), OPTIONAL, INTENT(IN)</td>
</tr>
<tr>
<td>N1byN0(:)</td>
<td>The density ratio for the first two vibrational levels</td>
<td>REAL(double), OPTIONAL, INTENT(IN)</td>
</tr>
<tr>
<td>SpeciesN</td>
<td>Starting index of species sequence</td>
<td>INTEGER, OPTIONAL, INTENT(IN)</td>
</tr>
<tr>
<td>STAT</td>
<td>Execution status indicator</td>
<td>INTEGER, OPTIONAL, INTENT(OUT)</td>
</tr>
</tbody>
</table>

Results

STAT = 0: no errors; STAT = n: error number.
Examples

Example E-19. **pkSetValue**: Setting the ratio of the number of molecules in the two first vibrational levels.

```
INTEGER :: Minvib, Nvib, dummy(NSpecies)
CHARACTER(20) :: SpcNames(NSpecies)
...
CALL pkGetParticle( 'name', SpcNames, 1 )
dummy(:) = INDEX( SpcNames(:), 'N2,v' ) ! Indexes of N2,v species
Minvib = MINLOC( dummy, dummy>0 ) - Nnc
Nvib = COUNT( dummy /= 0 )
CALL pkSetValue( N1byN0=SPREAD( Dens(Minvib+1)/Dens( Minvib ), 1, Nvib), &
                 Species=Minvib )
```

### SetRate

#### Name

*SetRate* — User supplied routine to set the values of rate coefficients

#### Description

The subroutine *SetRate* sets the values of rate coefficients when the rate coefficient equation included in *PLASMAKIN* and indicated in chapter one is not appropriate. The user is free to write this routine but the routine must have the interface described below.

#### Syntax

```
SetRate( Idx, rIdx, pIdx, Temp, RateValue )
```

<table>
<thead>
<tr>
<th>Name</th>
<th>Description</th>
<th>Data type and attributes</th>
</tr>
</thead>
<tbody>
<tr>
<td>Idx</td>
<td></td>
<td></td>
</tr>
<tr>
<td>rIdx</td>
<td></td>
<td></td>
</tr>
<tr>
<td>pIdx</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Temp</td>
<td></td>
<td></td>
</tr>
<tr>
<td>RateValue</td>
<td></td>
<td></td>
</tr>
</tbody>
</table>
### Procedures

<table>
<thead>
<tr>
<th>Name</th>
<th>Description</th>
<th>Data type and attributes</th>
</tr>
</thead>
<tbody>
<tr>
<td>Idx</td>
<td>Index of desired rate coefficient equation in the routine</td>
<td>INTEGER, INTENT(IN)</td>
</tr>
<tr>
<td>rIdx(NMaxReacSpc)</td>
<td>Array with the vibrational quantum numbers of reactant species</td>
<td>INTEGER, INTENT(IN)</td>
</tr>
<tr>
<td>pIdx(NMaxReacSpc)</td>
<td>Array with the vibrational quantum numbers of product species</td>
<td>INTEGER, INTENT(IN)</td>
</tr>
<tr>
<td>Temp</td>
<td>The electron or gas temperature used in the rate coefficient equation</td>
<td>REAL(double), INTENT(IN)</td>
</tr>
<tr>
<td>RateValue</td>
<td>The rate coefficient value</td>
<td>REAL(double), INTENT(OUT)</td>
</tr>
</tbody>
</table>
6. Data Input Format

The data organization and the input file syntax were designed to provide as much freedom as possible to the user, to reduce to a minimum the amount of information needed in particular the information not directly related to the physical or chemical nature of the problem. At the same time the input file syntax was designed to enforce correction.

The following strategies were used:

- Exclusive use of **NAMELIST** groups for data input.
- Species and reactions are always initialized with default values. This means that only when the value of a variable is different from the default it must be explicitly included on the input file.
- Values as the total number of species, number of species per type, number of chemical reactions etc, are automatically evaluated by **PLASMAKIN**
- The syntax and data input routines allow the reading and processing of vibrational series and reactions involving vibrational series.
- All CHARACTER values are case sensitive. This facilitates the detection of spelling errors in chemical elements and units.
PLASMAKIN_DATA

Name

PLASMAKIN_DATA — Holds initial values of data applying to the entire system

Syntax

&PLASMAKIN_DATA [Cte_p = Lvalue [,]] [ pressure = Rvalue [, ‘Cvalue’] [,]
gas_T = Rvalue [, ‘Cvalue’] [,] gas_n = Rvalue [, ‘Cvalue’] ]
[ [,] electron_T = Rvalue [, ‘Cvalue’] ] /

<table>
<thead>
<tr>
<th>Name</th>
<th>Description</th>
<th>Data type and attributes</th>
<th>Default value</th>
<th>Units or values supported</th>
</tr>
</thead>
<tbody>
<tr>
<td>Cte_p</td>
<td>Constant pressure system</td>
<td>LOGICAL</td>
<td>.TRUE.</td>
<td>.T, .t, .T, .t, .F, .f, .f, .f</td>
</tr>
<tr>
<td>Pressure</td>
<td>Total pressure</td>
<td>PHYSPROPERTY</td>
<td>0,’Pa’</td>
<td>’Pa’, ‘Nm-2’, ‘bar’, ‘mbar’, ‘torr’, ‘atm’</td>
</tr>
<tr>
<td>Gas_T</td>
<td>Gas temperature</td>
<td>PHYSPROPERTY</td>
<td>300,’K’</td>
<td>’K’, ‘C’</td>
</tr>
<tr>
<td>Gas_n</td>
<td>Total gas density</td>
<td>PHYSPROPERTY</td>
<td>0,’m-3’</td>
<td>‘m-3’, ‘cm-3’</td>
</tr>
<tr>
<td>electron_T</td>
<td>Electron temperature</td>
<td>PHYSPROPERTY</td>
<td>0,’eV’</td>
<td>‘eV’, ‘K’, ‘C’</td>
</tr>
</tbody>
</table>

Comments

This NAMELIST is usually used only once in the data input file. The Cte_p value is used to distinguish between constant pressure and constant volume conditions. The initial gas density can be indicated explicitly - specifying its value - or implicitly indicating the pressure and gas temperature. However even in the later case the user can be interested in indicate the gas density units. The electron temperature is only necessary if electron temperature dependent rate coefficients are used. Note that ‘C is just written C.
Examples

Example E-20. PLASMAKIN_DATA: Examples of PLASMAKIN_DATA NAMELIST syntax.

&PLASMAKIN_DATA Gas_n = 1.e+22 / ➊
&PLASMAKIN_DATA
     Gas_n = ,'cm-3'
     Pressure = 1,'mbar'
     Gas_T = 25,'C'
/
&PLASMAKIN_DATA Cte_p = F, Pressure = 1,'torr', electron_T = 3,'eV' / ➋

➊ Explicit indication of gas density using the default units (m⁻³). Gas temperature has also the default value (300 K).
➋ The density is calculated by PLASMAKIN from the pressure and temperature values but the value of the total density passed to the calling program is in cm⁻³.
➌ Constant volume conditions. Electron rate coefficients are evaluated with an initial electron temperature of 3 eV.

CHEM_SPECIES

Name
CHEM_SPECIES — Read chemical species and species properties
## Syntax


<table>
<thead>
<tr>
<th>Name</th>
<th>Description</th>
<th>Data type and attributes</th>
<th>Default value</th>
<th>Units or values supported</th>
</tr>
</thead>
<tbody>
<tr>
<td>name</td>
<td>Name of species</td>
<td>CHARACTER(20)</td>
<td>&quot;</td>
<td></td>
</tr>
<tr>
<td>constant</td>
<td>Is the species concentration constant?</td>
<td>LOGICAL</td>
<td>.FALSE.</td>
<td>.F., .F., .F., .T., .T.</td>
</tr>
<tr>
<td>charge</td>
<td>Electrical charge in elementary charge units</td>
<td>INTEGER</td>
<td>0</td>
<td></td>
</tr>
<tr>
<td>mass</td>
<td>Mass in a.m.u.</td>
<td>REAL(8)</td>
<td>0</td>
<td></td>
</tr>
<tr>
<td>g</td>
<td>Level multiplicity</td>
<td>INTEGER</td>
<td>1</td>
<td></td>
</tr>
<tr>
<td>energy</td>
<td>Level energy or standard molar enthalpy</td>
<td>PHYSPROPERTY</td>
<td>0, ‘eV’</td>
<td>‘.eV’, ‘kJ/mol’, ‘kcal/mol’</td>
</tr>
<tr>
<td>heatC</td>
<td>Heat capacity (c_p or c_v) in kJ/mol/k</td>
<td>REAL(8)</td>
<td>0</td>
<td></td>
</tr>
<tr>
<td>K_T</td>
<td>Thermal conductivity</td>
<td>REAL(8)</td>
<td>0</td>
<td></td>
</tr>
<tr>
<td>v</td>
<td>Range of quantum vibrational numbers</td>
<td>INTEGER</td>
<td>0, 0</td>
<td></td>
</tr>
<tr>
<td>omega</td>
<td>Vibrational frequency</td>
<td>PHYSPROPERTY</td>
<td>0, ‘cm-1’</td>
<td>‘.cm-1’, ‘.s-1’</td>
</tr>
<tr>
<td>vib_T</td>
<td>Vibrational temperature</td>
<td>PHYSPROPERTY</td>
<td>0, ‘K’</td>
<td>‘K’, ‘C’</td>
</tr>
<tr>
<td>anharmonicity</td>
<td>Anharmonicity parameter</td>
<td>REAL(8)</td>
<td>0</td>
<td></td>
</tr>
</tbody>
</table>
Data Input Format

<table>
<thead>
<tr>
<th>Name</th>
<th>Description</th>
<th>Data type and attributes</th>
<th>Default value</th>
<th>Units or values supported</th>
</tr>
</thead>
<tbody>
<tr>
<td>rotational_cte</td>
<td>Rotational constant</td>
<td>REAL(8)</td>
<td>0</td>
<td></td>
</tr>
<tr>
<td>mpolar</td>
<td>Dipolar or quadrupolar momentum</td>
<td>PHYSPROPERTY</td>
<td>0, &quot;&quot;</td>
<td>&quot;&quot;, ‘ea_o’, ‘ea_o2’</td>
</tr>
<tr>
<td>initial_conc</td>
<td>Species initial concentration</td>
<td>PHYSPROPERTY</td>
<td>0, &quot;&quot;</td>
<td>&quot;&quot;, ‘%’</td>
</tr>
<tr>
<td>num_scheme</td>
<td>Index of an users’ defined algorithm to evaluate species properties</td>
<td>INTEGER</td>
<td>0</td>
<td></td>
</tr>
<tr>
<td>cascade</td>
<td>Is species a cascade level?</td>
<td>LOGICAL</td>
<td>.FALSE.</td>
<td>.F, .f, .T, .t, .t, t</td>
</tr>
<tr>
<td>data_file</td>
<td>Data file with further informations</td>
<td>DATA_COLUMN</td>
<td>&quot;&quot;, 0</td>
<td></td>
</tr>
</tbody>
</table>

Comments

The following rules must be obeyed:

- All species must have a name. This name is used to identify the species in chemical reactions. Although nothing hinders the use of custom names, it is advisable to follow the chemical nomenclature with the limitation that under- or superscript numbers must be written inline. Names are case sensitive.
- The name for electrons must be the letter ‘e’.
- The names of photons must contain the word ‘photon’.
- A vibrational series is identified by a value of $v(2) > 0$. The name of each species in the series is formed by replacing the last closing brace in the series name by suffix ‘=n)’ where $n$ is the corresponding vibrational number. The range of vibrational levels in the series is defined by the values of the array $v$. The values of $v$ must follow the order $v(1) \leq v(2)$. For a single vibrational level only the first value of $v$ must be indicated.
Examples


```plaintext
&CHEM_SPECIES
name = 'Ne', constant = T, mass = 20.18, initial_conc = 100,'%'/

&CHEM_SPECIES
name = 'Ne(3P2)', energy = 16.61, g = 5,
data_file = 'NeTransp.txt', /

&CHEM_SPECIES name = 'Ar(3P)', cascade = T /

&CHEM_SPECIES name = 'Ne+', charge = 1, num_scheme = 2 /

&CHEM_SPECIES name = 'photon 1' /

&CHEM_SPECIES name = 'e', data_file = 'eTransp.txt' /

&CHEM_SPECIES
name = 'O2(X,v)'
mass = 32. 
v = 0.15
omega = 1580.19,'cm-1'
vib_T = 2000
anharmonicity = 7.58e-3
initial_conc = 5,'%'/
```

1. Dominant species. The value of constant must be indicated; mass must be indicated if the chemical kinetics model includes reverse reactions.
2. Neon excited level; the energy (in eV) and g values will be used in evaluating superelastic rates. The external data file is used to hold the value of the diffusion coefficient.
3. This level will be handled as a cascade level.
4. As species is an ion the charge value must be indicated. The index of an user defined algorithm is also indicated.
5. Photons produced on radiative transitions and involved on radiation trapping or any other process must be listed and identified by a name containing the word ‘photon’.
Electrons have the mandatory name ‘e’. The data_file can contain the values for the drift velocity and diffusion coefficient.

A vibrational series (from \(v=0\) to \(v=15\)). The names of the corresponding species are ‘\(O_2(X,v=0)\)’, ‘\(O_2(X,v=1)\)’, ‘\(O_2(X,v=15)\)’. The initial populations are computed from the values of the vibrational frequency, vibrational temperature and anharmonicity.

**CHEM_REACTION**

**Name**

CHEM_REACTION — Read a chemical reaction or reactions involving vibrational series

**Syntax**

&CHEM_REACTION [locus = ‘Cvalue’] [reactants = ‘Cvalue’ [[,]...]] [.]
[products = ‘Cvalue’ [[,]...]] [.] [comment = ‘Cvalue’] [.] [opposing = Lvalue] [.]
[ee_loss = Rvalue] [.] [value = Rvalue [[,]...]] [data_file = ‘Cvalue’,Ivalue] [.]
[units = ‘Cvalue’]/

<table>
<thead>
<tr>
<th>Name</th>
<th>Description</th>
<th>Data type and attributes</th>
<th>Default value</th>
<th>Units or values supported</th>
</tr>
</thead>
<tbody>
<tr>
<td>locus</td>
<td>Where the reaction takes place</td>
<td>CHARACTER(10)</td>
<td>‘gas’</td>
<td>‘gas’, ‘user_string’</td>
</tr>
<tr>
<td>reactants</td>
<td>Reactant species</td>
<td>CHARACTER(20)</td>
<td>NMaxReacSpc&quot;</td>
<td></td>
</tr>
<tr>
<td>products</td>
<td>Product species</td>
<td>CHARACTER(20)</td>
<td>NMaxReacSpc&quot;</td>
<td></td>
</tr>
<tr>
<td>comment</td>
<td>Relation between vibrational quantum numbers</td>
<td>CHARACTER(20)</td>
<td>‘'</td>
<td></td>
</tr>
</tbody>
</table>
### Data Input Format

<table>
<thead>
<tr>
<th>Name</th>
<th>Description</th>
<th>Data type and attributes</th>
<th>Default value</th>
<th>Units or values supported</th>
</tr>
</thead>
<tbody>
<tr>
<td>opposing</td>
<td>The <strong>NAMELIST</strong> describes both forward and reverse reactions</td>
<td>LOGICAL</td>
<td>.FALSE</td>
<td>.F, .F, .f, .F, .T, .t</td>
</tr>
<tr>
<td>ee_loss</td>
<td>Energy lost in the reaction by the electron</td>
<td>REAL(8)</td>
<td>0</td>
<td></td>
</tr>
<tr>
<td>units</td>
<td>Units used for reaction coefficient</td>
<td>CHARACTER(10)</td>
<td>&quot;</td>
<td>&quot;, 's-1', 'm3s-1', 'cm3s-1', 'm6s-1', 'cm6s-1'</td>
</tr>
<tr>
<td>value</td>
<td>Reaction rate equation coefficients</td>
<td>REAL(8)(12) (0, i=1,12)</td>
<td>(0, i=1,12)</td>
<td></td>
</tr>
<tr>
<td>data_file</td>
<td>Index of function to compute the reaction rate in the user supplied routine</td>
<td>DATA_COLUMN</td>
<td>&quot;, 0</td>
<td></td>
</tr>
</tbody>
</table>

### Comments

The following rules must be followed:

- If the `locus` of a reaction in the gas is indicated, this value must be `gas`. For surface reactions any name can be used and reactions in different surfaces can be identified by different `locus` values.
- All species indicated in the `reactants` and `products` lists must be described in a `CHEM_SPECIES NAMELIST`.
- Reactions involving vibrational species can be collectively represented indicating the generic name representing the vibrational series. The vibrational species involved in the reaction can be selected through relational expressions involving the vibrational quantum numbers. These expressions can be included in the species name in the `reactants` or `products` fields or, if they involve two different vibrational quantum numbers, in the `comment` field. The vibrational quantum numbers in these expressions can only be represented by the letters `v` or `w`. 

55
The following classes of relational expressions can be used:

a. If the reaction corresponds to a fixed variation of the vibrational quantum number, the relation between the initial and the final values can be translated into the names of the vibrational species in the reactants and products. In this case the arithmetic operators + and - can be used in the species names.

b. If the range of vibrational levels is defined by a relation involving only one vibrational number, this relation can be included in the vibrational series name.

c. If the expression used to select the vibrational levels involves two vibrational numbers, this relation must be written in the comment field.

In case these two last cases the relations can be written using the logical operators $\lt$, $\leq$, $\gt$, $\geq$ and $\equiv$. In the last case the modulo operator can also be used.

- value is an array of 12 elements allowing representing the 6 pair of $\alpha$ and $\beta$ values needed to represent the rate coefficient using the equation indicated in chapter one.

- Finally, if the vibrational rate coefficients must be computed from a table or analytically but the equation in chapter one is not adequate, the user can supply the external routine SetRate with user defined rate equations. The index value in the data_file field is used to selection of the appropriate rate equation. In this case the name value is not used.

Caution: Indistinguishable reactions

For a V-V reaction involving two vibrational levels of the same molecule the direct and reverse reactions are indistinguishable. In this case the opposing property should not be set to T. Otherwise the number of reactions is counted twice.

Examples

Example E-22. CHEM_REACTION: Example of CHEM_REACTION NAMELIST syntax.

```
&CHEM_REACTION
  reactants = 'Ne(3P1)', products = 'Ne','photon 1',
  value = 0.486e8 /
&CHEM_REACTION
  reactants = 'Ne(3P2)','Ne(3P1)', products = 'Ne','Ne+','e',
  value = 3.2e-10, units = 'cm3s-1' /
```

Radiation emission; the value of \( \text{value} \) is the transition probability in \( s^{-1} \).

Penning ionisation with constant rate coefficient.

Electron excitation. The values of the rate coefficient are read in the external file \texttt{Ne_data.txt} and identified by the index 3.

Both the direct and reverse reactions are taken into account. The rate coefficient for the direct reaction has an Arrhenius temperature dependence with the above coefficients. The rate coefficient for the reverse reaction are automatically computed.

Group of reactions on two vibrational series. The syntax for the relations between the vibrational quantum numbers corresponds to case a. above.

Group of reactions involving a vibrational series. The syntax corresponds to case b. above.

Group of reactions involving a vibrational series. The syntax corresponds to case c. above.
References


Appendix A. Error Messages

The number of error messages issued by \textit{PLASMAKIN} is relatively small - twenty-seven - and in the present version no attempt has been made to cover all possible error situations. Most of the messages are related with the reading and processing of the input file where it is easy to make typographical mistakes. This is an area where improvement can be expected in next versions of the library.

The error messages have the structure \textit{PLASMAKIN:}\textless\text{classification and number}\textgreater:\textless\text{description}\textgreater

\textbf{Error messages and description}

\textit{PLASMAKIN}: severe(1): NAMELIST "PLASMAKIN\_DATA" not found

\textit{PLASMAKIN}: severe(2): NAMELIST "CHEM\_SPECIES" not found

\textit{PLASMAKIN}: severe(3): NAMELIST "CHEM\_REACTION" not found

\textit{PLASMAKIN}: severe(4): Syntax error in "CHEM\_SPECIES" NAMELIST #

\textit{PLASMAKIN}: severe(5): Syntax error in "CHEM\_REACTION" NAMELIST #

\textit{PLASMAKIN}: severe(6): Too many values for NAMELIST variable

\textit{PLASMAKIN}: severe(7): Invalid reference to variable in NAMELIST input

\textit{PLASMAKIN}: severe(8): NAMELIST I/O not consistent with OPEN options

\textit{PLASMAKIN}: severe(9): Invalid NAMELIST input format
Appendix A. Error Messages

PLASMAKIN: severe(10): REWIND error

PLASMAKIN: error(11): Undetermined I/O error

PLASMAKIN: severe(12): Error in physical units or units not supported

PLASMAKIN: error(13): Total sum of species initial concentration is not 1

PLASMAKIN: severe(14): Attempt to re-read chemical species list

PLASMAKIN: severe(15): Attempt to re-read reaction list

PLASMAKIN: severe(16): Unknown species in reaction #

PLASMAKIN: severe(17): Violation of charge conservation in reaction #

PLASMAKIN: severe(151): Allocatable array is already allocated

PLASMAKIN: severe(727): Cannot ALLOCATE allocatable array - out of memory

PLASMAKIN: error(20): Undetermined memory error

PLASMAKIN: error(21): Syntax error in call to routine "pkSetValue"

PLASMAKIN: error(22): Syntax error in call to routine "pkSetPhoton"
Appendix A. Error Messages

*PLASMAKIN*: error(23): Syntax error in call to routine "pkGetSources"

*PLASMAKIN*: error(24): Syntax error in call to routine "pkGetReacCoef"

*PLASMAKIN*: warning(25): Interrogation on unknown property

*PLASMAKIN*: warning(26): Unable to set property of rate coefficient

*PLASMAKIN*: warning(27): Attempt to set the value of T-dependent rate coefficient
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