

Preview of molecular population models

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University of Strathclyde

ADAS-EU course – 26 – 30 Mars 2012

- 1 Objectives
- 2 ADAS 900 formats
- 3 ADAS 900 series overview
 - ADAS902
 - ADAS903
 - ADAS904
 - ADAS901
- 4 Future work

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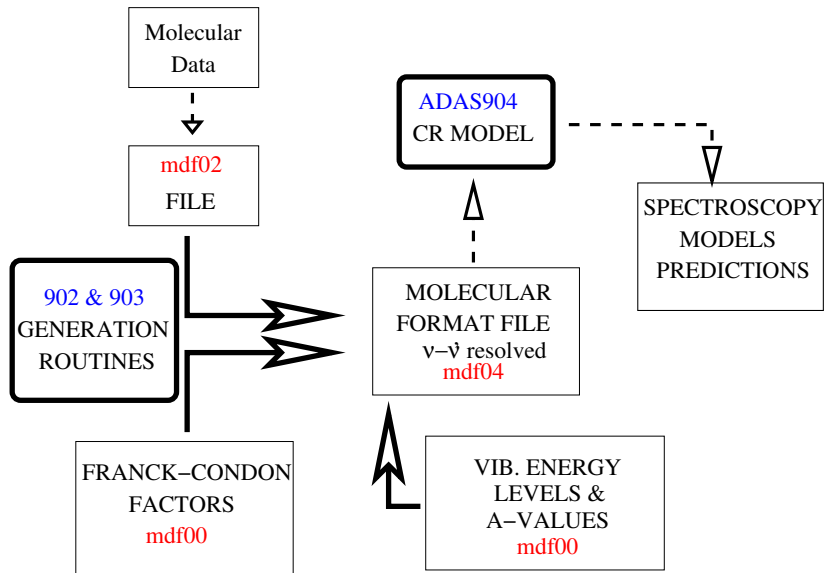
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- ADAS 900 series main objective is to extend ADAS Database to molecular data
- We want to provide molecular data tools to use in plasma edge and divertor calculations and experiments.

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ADAS900



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Two resolution are accounted: vibrational and electronic.

Formats

- mdf00** General information. Vibrational energies, Potential curves. FC factors. A-values.
- mdf02 Cross sections and rates obtained from external sources or calculations.
- mdf33/34 ADAS902 product. Rates (upsilons and $cm^3 s^{-1}$) obtained from maxwellian integratrion of *mdf02*.
- mdf04/14 ADAS903 product. Maxwellian rates completed with EIQIP and EICIP to be used in population models.
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MDF02: Compiling H₂ system

- **Excitation data** The available data have been checked to be in good agreement with the fitting formulas.
- **Ionization data** Checked as well. Only ionization from ground state. Ionization from excited states can be achieved using IP models.
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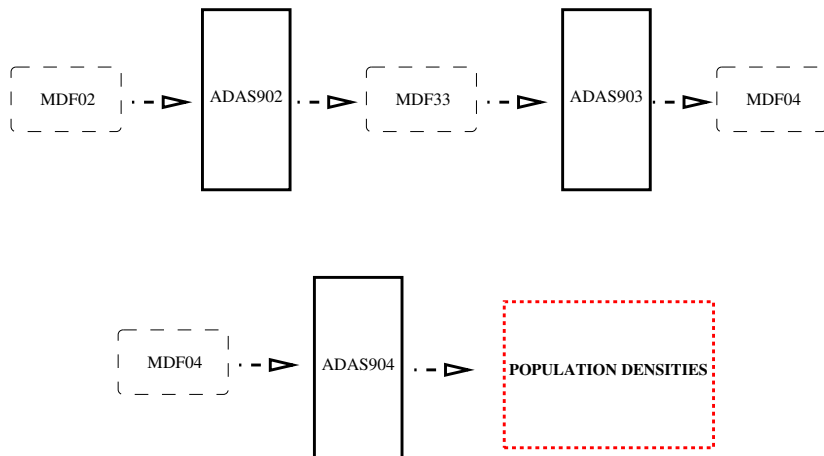
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General Scheme



- ADAS902 integrates, interpolates and extrapolates (if necessary) the **existing** cross sections and creates the maxwellian rate coefficients. Also create different resolution files.
- ADAS903 look for the gaps in the transitions matrix and calculate them by semi-classical Impact Parameter approach in the required resolution.
- ADAS904 calculates the molecular effective coefficients, write them in the various *mdf11* formats and invert the matrix to obtain the populations.
- ADAS901 package is a set of routines to obtain effective lifetimes of the various processes.
- MDFLIBS are different tools to deal with the data stored.

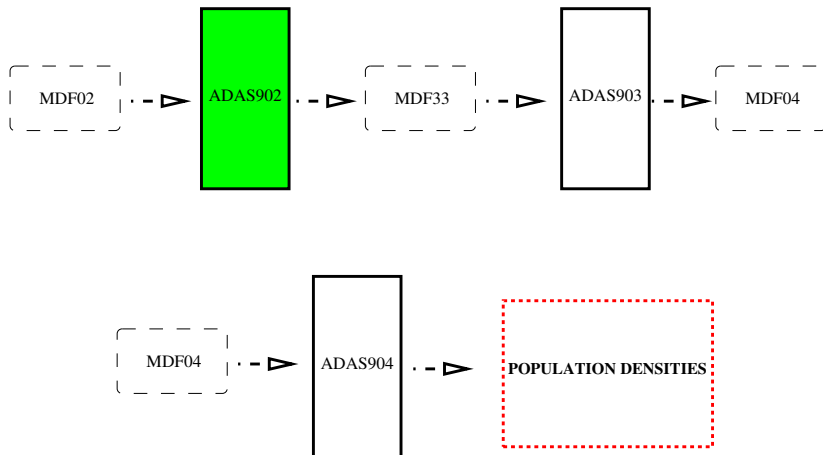
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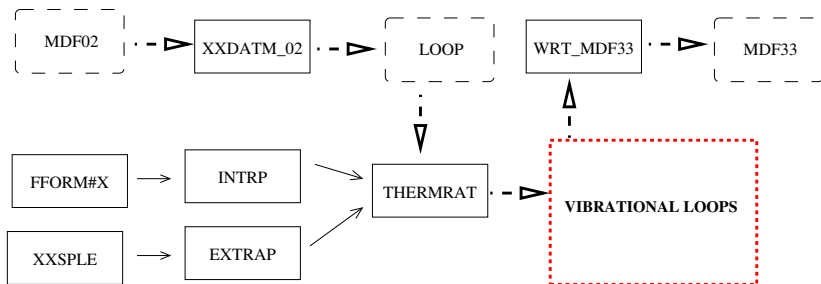
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ADAS902: Interpolation and maxwellians



Interpolation & Extrapolation

If a fitting formula exist, this is chosen by default.

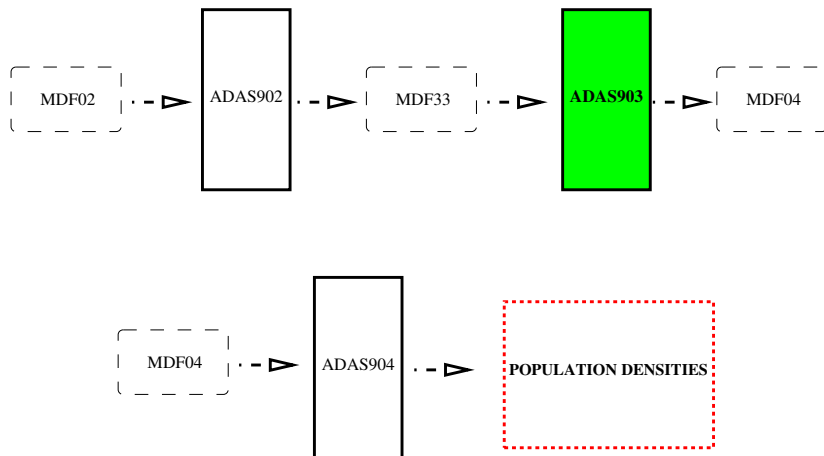
Extrapolation approaches:

- Bethe model for high energy ionization.
- Spline for high and low energies excitation.
- Overbarrier model for high energies CX.

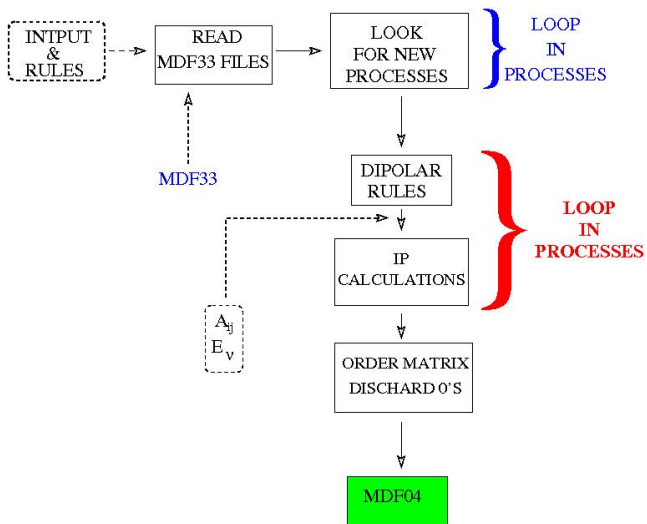
Dissociation can come from excitation or ionization so the approaches above are taken.

The extrapolation models are normalized to the data to avoid peaks and steps.

General Scheme



ADAS903: filling the matrix



ADAS903 input file

- There is the possibility to put a lower limit in the rates to consider the transitions,
- Driver *input903.dat* must be created and dissociative states should be defined with rule “ED” (Excitation to dissociative).
- Temperatures are the same as in mdf33 and mdf34. Interpolation are carried out when reading.

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ADAS903 matrix completion

- EIQIP (bound-bound collisional IP) is used for excitation inside a molecular specie.
- ECIP (bound-free collisional IP) is used for ionization (dissociative and non-dissociative) and dissociative excitation.
- Calculation is performed over vibronic levels and summed.
- EIQIP uses A-values to calculate the oscillator strengths so Franck- Condon factors unitarity is fulfilled.
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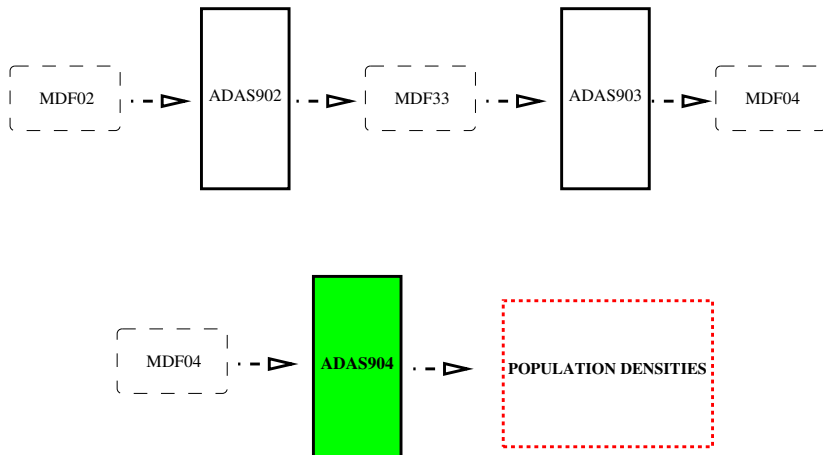
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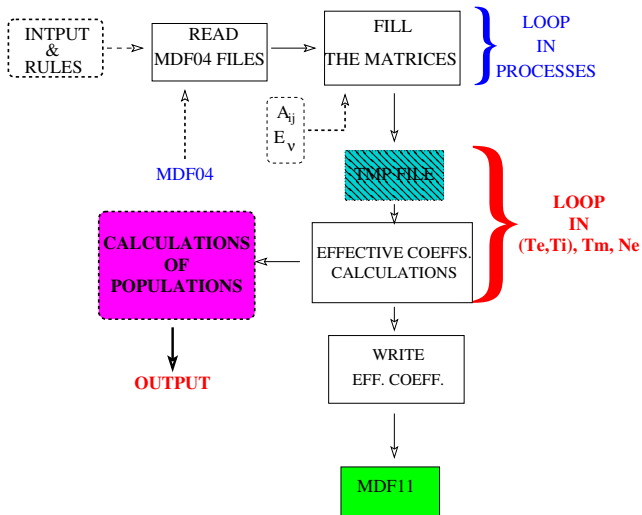
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mdf04 are ascii and human readable. Not so funny though!

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ADAS904: solving CR



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- Dissociative processes are all that go to dissociation (diss. excitation, ionisation, CX, recombination . . .).
- Metastables are always vibrationally resolved \longrightarrow 150×150 maximum dimension of metastable matrix.
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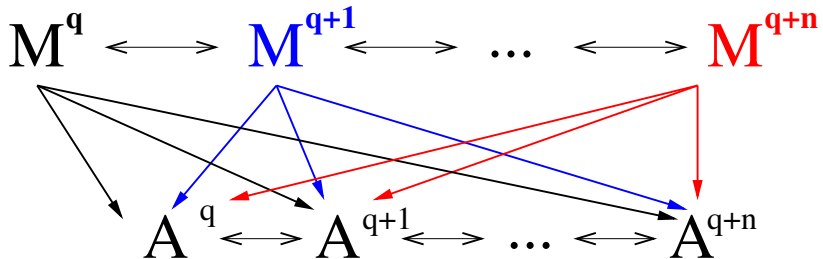
ADAS904: The population model

CR Equation

$$\left(\frac{dN_X}{dt} \right) = \underbrace{\left(\begin{array}{c|c|c} \text{H}_2 & \sim 0 & 0 \\ \hline \neq 0 & \text{H}_2^+ & 0 \\ \hline \neq 0 & \neq 0 & \text{H}(n), \text{H}^+ \end{array} \right)}_{\left(\begin{array}{c} \dots \\ \text{states} \\ \dots \end{array} \right)} \times \left(N_X \right) - \left(\Gamma_{in} \right)$$

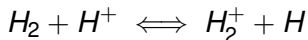
Γ_{in} can correspond to the collisional terms from constant populations.

The population model



Non linear terms

Charge exchange is creating non linear terms to be sum to the background populations of neutrals and protons.

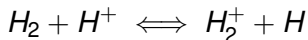


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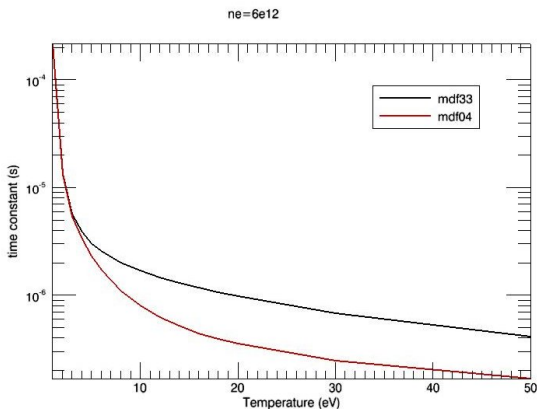
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Steps:

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- 2 Optimization and extension of data base for Hydrogen
- 3 Implementation in ADAS and IDL
- 4 Interaction to cover the experimental needs.
- 5 Extension to other diatomic molecules.
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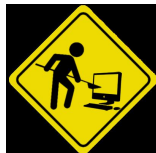
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