New Molecular Collisional-Radiative Model in ADAS

Francisco Guzmán

ADAS-EU University of Strathclyde IRFM-CEA, Cadarache

ADAS Workshop 24 – 25 September 2012



Outline



2 Molecular ADAS routines: ADAS900

- General view
- Molecular Data
- CR Model







Outline



- 2 Molecular ADAS routines: ADAS900
 - General view
 - Molecular Data
 - CR Model

3 Results





Motivation

- Molecules desorbed from the wall contribute to the neutral density and influence the divertor physics.
- ADAS 900 series main objective is to extend ADAS Database to molecular data



Motivation

- Molecules desorbed from the wall contribute to the neutral density and influence the divertor physics.
- ADAS 900 series main objective is to extend ADAS Database to molecular data



Motivation

- Molecules desorbed from the wall contribute to the neutral density and influence the divertor physics.
- 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.



- Compile a database. Structure it and provide readings tool for it. Do maxwellian integration of cross sections to obtain rates(Boring part).
- There are gaps!. Fill the gaps with estimation of molecular processes. Automatic process is needed due to the big number of states.
- Solve CR model for all molecules and all dissociation products.



- Compile a database. Structure it and provide readings tool for it. Do maxwellian integration of cross sections to obtain rates(Boring part).
- There are gaps!. Fill the gaps with estimation of molecular processes. Automatic process is needed due to the big number of states.
- Solve CR model for all molecules and all dissociation products.



- Compile a database. Structure it and provide readings tool for it. Do maxwellian integration of cross sections to obtain rates(Boring part).
- There are gaps!. Fill the gaps with estimation of molecular processes. Automatic process is needed due to the big number of states.
- Solve CR model for all molecules and all dissociation products.



- Compile a database. Structure it and provide readings tool for it. Do maxwellian integration of cross sections to obtain rates(Boring part).
- There are gaps!. Fill the gaps with estimation of molecular processes. Automatic process is needed due to the big number of states.
- Solve CR model for all molecules and all dissociation products.



Outline



2 Molecular ADAS routines: ADAS900

- General view
- Molecular Data
- CR Model

3 Results





Molecular ADAS routines: ADAS900

Results

ADAS 900





mdf00 General information. Vibrational energies. FC factors. A-values.

- mdf02 Cross sections and rates obtained from external sources or calculations.
- mdf33/34 ADAS902 product. Rates (upsilons and cm^3s^{-1}) obtained from maxwellian integratrion of *mdf02*.
- mdf04/14 ADAS903 product. Maxwellian rates completed with EIQIP and EICIP to be used in population models.
 - mdf11 ADAS904 product. Molecular effective coefficients.
 - mdf15 Molecular PEC, SXB, DXB.
 - mdf25 Mapping Central ADAS adf and mdf formats correlation.
 - mdf38 Autoionization and predissociation rates.



- mdf00 General information. Vibrational energies. FC factors. A-values.
- mdf02 Cross sections and rates obtained from external sources or calculations.
- mdf33/34 ADAS902 product. Rates (upsilons and cm^3s^{-1}) obtained from maxwellian integratrion of *mdf02*.
- mdf04/14 ADAS903 product. Maxwellian rates completed with EIQIP and EICIP to be used in population models.
 - mdf11 ADAS904 product. Molecular effective coefficients.
 - mdf15 Molecular PEC, SXB, DXB.
 - mdf25 Mapping Central ADAS adf and mdf formats correlation.
 - mdf38 Autoionization and predissociation rates.



- mdf00 General information. Vibrational energies. FC factors. A-values.
- mdf02 Cross sections and rates obtained from external sources or calculations.
- mdf33/34 ADAS902 product. Rates (upsilons and cm^3s^{-1}) obtained from maxwellian integratrion of *mdf02*.
- mdf04/14 ADAS903 product. Maxwellian rates completed with EIQIP and EICIP to be used in population models.
 - mdf11 ADAS904 product. Molecular effective coefficients.
 - mdf15 Molecular PEC, SXB, DXB.
 - mdf25 Mapping Central ADAS adf and mdf formats correlation.
 - mdf38 Autoionization and predissociation rates.



- mdf00 General information. Vibrational energies. FC factors. A-values.
- mdf02 Cross sections and rates obtained from external sources or calculations.
- mdf33/34 ADAS902 product. Rates (upsilons and cm^3s^{-1}) obtained from maxwellian integratrion of *mdf02*.
- mdf04/14 ADAS903 product. Maxwellian rates completed with EIQIP and EICIP to be used in population models.
 - mdf11 ADAS904 product. Molecular effective coefficients.
 - mdf15 Molecular PEC, SXB, DXB.
 - mdf25 Mapping Central ADAS adf and mdf formats correlation.
 - mdf38 Autoionization and predissociation rates.



- mdf00 General information. Vibrational energies. FC factors. A-values.
- mdf02 Cross sections and rates obtained from external sources or calculations.
- mdf33/34 ADAS902 product. Rates (upsilons and cm^3s^{-1}) obtained from maxwellian integratrion of *mdf02*.
- mdf04/14 ADAS903 product. Maxwellian rates completed with EIQIP and EICIP to be used in population models.
 - mdf11 ADAS904 product. Molecular effective coefficients.
 - mdf15 Molecular PEC, SXB, DXB.
 - mdf25 Mapping Central ADAS adf and mdf formats correlation.
 - mdf38 Autoionization and predissociation rates.



- mdf00 General information. Vibrational energies. FC factors. A-values.
- mdf02 Cross sections and rates obtained from external sources or calculations.
- mdf33/34 ADAS902 product. Rates (upsilons and cm^3s^{-1}) obtained from maxwellian integratrion of *mdf02*.
- mdf04/14 ADAS903 product. Maxwellian rates completed with EIQIP and EICIP to be used in population models.
 - mdf11 ADAS904 product. Molecular effective coefficients.
 - mdf15 Molecular PEC, SXB, DXB.
 - mdf25 Mapping Central ADAS adf and mdf formats correlation.
 - mdf38 Autoionization and predissociation rates.



- mdf00 General information. Vibrational energies. FC factors. A-values.
- mdf02 Cross sections and rates obtained from external sources or calculations.
- mdf33/34 ADAS902 product. Rates (upsilons and cm^3s^{-1}) obtained from maxwellian integratrion of *mdf02*.
- mdf04/14 ADAS903 product. Maxwellian rates completed with EIQIP and EICIP to be used in population models.
 - mdf11 ADAS904 product. Molecular effective coefficients.
 - mdf15 Molecular PEC, SXB, DXB.
 - mdf25 Mapping Central ADAS adf and mdf formats correlation.

ndf38 Autoionization and predissociation rates.



- mdf00 General information. Vibrational energies. FC factors. A-values.
- mdf02 Cross sections and rates obtained from external sources or calculations.
- mdf33/34 ADAS902 product. Rates (upsilons and cm^3s^{-1}) obtained from maxwellian integratrion of *mdf02*.
- mdf04/14 ADAS903 product. Maxwellian rates completed with EIQIP and EICIP to be used in population models.
 - mdf11 ADAS904 product. Molecular effective coefficients.
 - mdf15 Molecular PEC, SXB, DXB.
 - mdf25 Mapping Central ADAS adf and mdf formats correlation.
 - mdf38 Autoionization and predissociation rates.



- Excitation data: There is avalaible data¹ that have been checked to be in good agreement with the fitting formulas from Janev.
- **Ionization data** Only ionization from ground state available.
- Attachment Fitting formulas from Celiberto² for fitting calculations. Vibrational excitation through resonant attachment.

¹R.Janev et al. JUEL 4353 Report, Sept. 2012, ISSN 0944-2952 -EU ^{Conneurope²Celiberto AD&NDT 77, 161 (2001)}

- Excitation data: There is avalaible data¹ that have been checked to be in good agreement with the fitting formulas from Janev.
- **Ionization data** Only ionization from ground state available.
- Attachment Fitting formulas from Celiberto² for fitting calculations. Vibrational excitation through resonant attachment.

¹R.Janev et al. JUEL 4353 Report, Sept. 2012, ISSN 0944-2952 Umm²Celiberto AD&NDT 77, 161 (2001)

- Excitation data: There is avalaible data¹ that have been checked to be in good agreement with the fitting formulas from Janev.
- **Ionization data** Only ionization from ground state available.
- Attachment Fitting formulas from Celiberto² for fitting calculations. Vibrational excitation through resonant attachment.

¹R.Janev et al. JUEL 4353 Report, Sept. 2012, ISSN 0944-2952 U₂₀₀₀²Celiberto AD&NDT 77, 161 (2001)

- Excitation data: There is avalaible data¹ that have been checked to be in good agreement with the fitting formulas from Janev.
- **Ionization data** Only ionization from ground state available.
- Attachment Fitting formulas from Celiberto² for fitting calculations. Vibrational excitation through resonant attachment.

¹R.Janev et al. JUEL 4353 Report, Sept. 2012, ISSN 0944-2952 AS-EU ²Celiberto AD&NDT **77**, 161 (2001)

- **Dissociative Attachment** Available calculations from Atems and Wadehra³ and Celiberto (also compiled by Janev et al.).
- **Double excited** There are not cross sections calculation. No information in cross sections for autoionization.
- **Predissociation** Only rates available. Can be a significant branching ratio. Dependence on rotational states.
- **Metastable** Only excitation and attachment data (no ionization) available.

- **Dissociative Attachment** Available calculations from Atems and Wadehra³ and Celiberto (also compiled by Janev et al.).
- **Double excited** There are not cross sections calculation. No information in cross sections for autoionization.
- **Predissociation** Only rates available. Can be a significant branching ratio. Dependence on rotational states.
- **Metastable** Only excitation and attachment data (no ionization) available.

- **Dissociative Attachment** Available calculations from Atems and Wadehra³ and Celiberto (also compiled by Janev et al.).
- **Double excited** There are not cross sections calculation. No information in cross sections for autoionization.
- **Predissociation** Only rates available. Can be a significant branching ratio. Dependence on rotational states.

Metastable Only excitation and attachment data (no ionization) available.

- **Dissociative Attachment** Available calculations from Atems and Wadehra³ and Celiberto (also compiled by Janev et al.).
- **Double excited** There are not cross sections calculation. No information in cross sections for autoionization.
- **Predissociation** Only rates available. Can be a significant branching ratio. Dependence on rotational states.
- Metastable Only excitation and attachment data (no ionization) available.

- EIQIP⁴ (bound-bound collisional IP) is used for excitation inside a molecular specie.
- ECIP⁵ (bound-free collisional IP) has been used for ionization (dissociative and non-dissociative) and dissociative excitation. It is needed a better optimization for the extension to molecules.
- Calculation are performed over vibronic levels and summed in case of electronic resolution.
- EIQIP uses A-values to calculate the oscillator strengths so Franck- Condon factors unitarity is fullfilled.
- Energy gaps are aproximated by the minimun point of the potential energy difference. No storage of full potential is needed.

⁴A. Burgess and H.P. Summers. MNRAS **174**, 345 (1976).

S-EU 5A. Burgess and I. C. Percival. Adv. Atom. Molec. Rhys. 💤 (1968) (199 📱 🧹

- EIQIP⁴ (bound-bound collisional IP) is used for excitation inside a molecular specie.
- ECIP⁵ (bound-free collisional IP) has been used for ionization (dissociative and non-dissociative) and dissociative excitation. It is needed a better optimization for the extension to molecules.
- Calculation are performed over vibronic levels and summed in case of electronic resolution.
- EIQIP uses A-values to calculate the oscillator strengths so Franck- Condon factors unitarity is fullfilled.
- Energy gaps are aproximated by the minimun point of the potential energy difference. No storage of full potential is needed.

⁴A. Burgess and H.P. Summers. MNRAS **174**, 345 (1976).

📙 ⁵A. Burgess and I. C. Percival. Adv. Atom. Molec. 🖓 👘 🖓 (1968) 🕫 🦿

- EIQIP⁴ (bound-bound collisional IP) is used for excitation inside a molecular specie.
- ECIP⁵ (bound-free collisional IP) has been used for ionization (dissociative and non-dissociative) and dissociative excitation. It is needed a better optimization for the extension to molecules.
- Calculation are performed over vibronic levels and summed in case of electronic resolution.
- EIQIP uses A-values to calculate the oscillator strengths so Franck- Condon factors unitarity is fullfilled.
- Energy gaps are aproximated by the minimun point of the potential energy difference. No storage of full potential is needed.

⁴A. Burgess and H.P. Summers. MNRAS **174**, 345 (1976). ^{AS-EU} ⁵A. Burgess and I. C. Percival. Adv. Atom. Molec. Phys. **4** (1968) 109

- EIQIP⁴ (bound-bound collisional IP) is used for excitation inside a molecular specie.
- ECIP⁵ (bound-free collisional IP) has been used for ionization (dissociative and non-dissociative) and dissociative excitation. It is needed a better optimization for the extension to molecules.
- Calculation are performed over vibronic levels and summed in case of electronic resolution.
- EIQIP uses A-values to calculate the oscillator strengths so Franck- Condon factors unitarity is fullfilled.
- Energy gaps are aproximated by the minimun point of the potential energy difference. No storage of full potential is needed.

⁴A. Burgess and H.P. Summers. MNRAS **174**, 345 (1976). AS-EU For base in Europe ⁵A. Burgess and I. C. Percival. Adv. Atom. Molec. Phys. **4** (1968) 109

- EIQIP⁴ (bound-bound collisional IP) is used for excitation inside a molecular specie.
- ECIP⁵ (bound-free collisional IP) has been used for ionization (dissociative and non-dissociative) and dissociative excitation. It is needed a better optimization for the extension to molecules.
- Calculation are performed over vibronic levels and summed in case of electronic resolution.
- EIQIP uses A-values to calculate the oscillator strengths so Franck- Condon factors unitarity is fullfilled.
- Energy gaps are aproximated by the minimun point of the potential energy difference. No storage of full potential is needed.

⁴A. Burgess and H.P. Summers. MNRAS **174**, 345 (1976).

ADAS-EU ⁵A. Burgess and I. C. Percival. Adv. Atom. Molec. Phys., 4 (1968) 109

- EIQIP⁴ (bound-bound collisional IP) is used for excitation inside a molecular specie.
- ECIP⁵ (bound-free collisional IP) has been used for ionization (dissociative and non-dissociative) and dissociative excitation. It is needed a better optimization for the extension to molecules.
- Calculation are performed over vibronic levels and summed in case of electronic resolution.
- EIQIP uses A-values to calculate the oscillator strengths so Franck- Condon factors unitarity is fullfilled.
- Energy gaps are aproximated by the minimun point of the potential energy difference. No storage of full potential is needed.

⁴A. Burgess and H.P. Summers. MNRAS **174**, 345 (1976).

ADAS-EU ⁵A. Burgess and I. C. Percival. Adv. Atom. Molec. Phys. **4** (1968) 109

Molecular ADAS routines: ADAS900

Results

ADAS903: filling the matrix





Molecular ADAS routines: ADAS900

Results

Why filling the matrix?

Maxwellian time constant: $\tau = \frac{1}{n_e \langle \sigma v \rangle}$

$$H_2(X, \nu = 0) \rightarrow H + H$$

$$H_2(X, \sum_{\nu} \nu) \rightarrow H + H$$



Processes from excited vibrational levels are needed.



- Dissociative processes are all that go to dissociation (diss. excitiation, ionisation, CX, recombination ...).
- Metastables are allways vibrationally resolved
 → 150 × 150 maximum dimension of metastable matrix.
- Autoionisation and predissociation are included at this stage from *mdf38* format data.
- Atomic dissociation products equilibrium ionisation balance is included in the CR model.



- Dissociative processes are all that go to dissociation (diss. excitiation, ionisation, CX, recombination ...).
- Metastables are allways vibrationally resolved
 → 150 × 150 maximum dimension of metastable matrix.
- Autoionisation and predissociation are included at this stage from *mdf38* format data.
- Atomic dissociation products equilibrium ionisation balance is included in the CR model.



- Dissociative processes are all that go to dissociation (diss. excitiation, ionisation, CX, recombination ...).
- Metastables are allways vibrationally resolved
 → 150 × 150 maximum dimension of metastable matrix.
- Autoionisation and predissociation are included at this stage from *mdf38* format data.
- Atomic dissociation products equilibrium ionisation balance is included in the CR model.



- Dissociative processes are all that go to dissociation (diss. excitiation, ionisation, CX, recombination ...).
- Metastables are allways vibrationally resolved
 → 150 × 150 maximum dimension of metastable matrix.
- Autoionisation and predissociation are included at this stage from *mdf38* format data.
- Atomic dissociation products equilibrium ionisation balance is included in the CR model.



- Dissociative processes are all that go to dissociation (diss. excitiation, ionisation, CX, recombination ...).
- Metastables are allways vibrationally resolved
 → 150 × 150 maximum dimension of metastable matrix.
- Autoionisation and predissociation are included at this stage from *mdf38* format data.
- Atomic dissociation products equilibrium ionisation balance is included in the CR model.



Molecular ADAS routines: ADAS900

Results

Molecules CR model





ADAS904: The population model

CR Equation

$$\begin{pmatrix} \frac{\mathrm{d}N_{X}}{\mathrm{d}t} \end{pmatrix} = \underbrace{\begin{pmatrix} \frac{\mathrm{H}_{2} & \sim 0 & 0 \\ \neq 0 & \mathrm{H}_{2}^{+} & 0 \\ \neq 0 & \neq 0 & \mathrm{H}(n), \mathrm{H}^{+} \end{pmatrix}}_{\begin{pmatrix} \dots \\ \mathrm{states} \\ \dots \end{pmatrix}} \times \begin{pmatrix} N_{X} \end{pmatrix} - \begin{pmatrix} \Gamma_{in} \end{pmatrix}$$

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



Molecular ADAS routines: ADAS900

Results

ADAS904: CR model





< □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □

Outline



- 2 Molecular ADAS routines: ADAS900
 - General view
 - Molecular Data
 - CR Model







Molecular ADAS routines: ADAS900

Results

Effective coefficients

CR dissociation from ground state



CR ionization from ground state to ground state





Molecular ADAS routines: ADAS900

Results

Population





Outline



- 2 Molecular ADAS routines: ADAS900
 - General view
 - Molecular Data
 - CR Model







Summary

- New molecular collisional-radiative tools have been developped and will be soon available.
- The tools can be extended to other diatomic molecules other than H₂.
- Molecular data are the main problem in building such a model.
- Checking and developing of method of calculations under way.
- Easy implementation of results in models (ADAS structure).

