# CRM and photon tracing module in EIRENE

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#### H colrad or the CRM in Eirene

- Based on the collisional radiative model by Sawada (1995)
- Resolve population coefficients for each Eirene cell
- Derive effective rates such as effective ionization and recombination rates to be used in EIRENE
- Provide population densities of excited species as a bulk ion species for the Photon tracing module



#### **CRM for atomic hydrogen**

$$\frac{dn_{H}(p)}{dt} = -\left[\sum_{p>q} \left(A_{(p,q)} + F_{(p,q)}n_{e}\right) + S_{(p)}n_{e} + \sum_{pp} \left[A_{(q,p)} + F_{(q,p)}n_{e}\right] n_{H(q>p)} + \left[\alpha(p)n_{e} + \beta(p)\right] n_{e} n_{H^{+}} + \Gamma_{H(p)}$$

- A = Spontaneous emission rate
- **F** = De-excitation rate coefficient
- S = ionization rate coefficient
- **C** = Excitation rate coefficient
- $\alpha$  = 3-body recombination rate coefficient
- $\beta$  = radiative recombination rate



#### CRM in steady-state, matrix form

 $\begin{bmatrix} -(S_{(1)} + \sum C_{1}) n_{e} & A_{(1,p)} + F_{(1,p)} n_{e} & A_{(1,q)} + F_{(1,q)} n_{e} & (\alpha_{(1)} n_{e} + \beta_{(1)}) n_{e} \\ C_{(1,p)} n_{e} & -A_{(1,p)} - (S_{(p)} + F_{(1,p)} + C_{(p,q)}) n_{e} & A_{(p,q)} + F_{(p,q)} n_{e} & (\alpha_{(p)} n_{e} + \beta_{(p)}) n_{e} \\ C_{(1,q)} n_{e} & C_{(p,q)} n_{e} & -\sum A_{(q)} - (S_{(q)} + \sum F_{(q)}) n_{e} & (\alpha_{(q)} n_{e} + \beta_{(q)}) n_{e} \\ S_{(1)} n_{e} & S_{(p)} n_{e} & S_{(p)} n_{e} & -\sum A_{(q)} - (S_{(q)} + \sum F_{(q)}) n_{e} & (\alpha_{(q)} n_{e} + \beta_{(q)}) n_{e} \\ S_{(q)} n_{e} & -\sum A_{(q)} n_{e} & -\sum A_{(q)} n_{e} & -\sum A_{(q)} n_{e} + \sum \beta n_{e} \end{bmatrix} \begin{bmatrix} n_{H} \\ n_{H(p)} \\ n_{H(q)} \\ n_{H+} \end{bmatrix} = \begin{bmatrix} -\Gamma_{H} \\ -\Gamma_{H(p)} \\ -\Gamma_{H(q)} \\ -\Gamma_{H+} \end{bmatrix}$ 

$$\begin{bmatrix} -A_{(1,p)} - (S_{(p)} + F_{(1,p)} + C_{(p,q)})n_e & A_{(p,q)} + F_{(p,q)}n_e \\ C_{(p,q)} n_e & -\sum A_{(q)} - (S_{(q)} + \sum F_{(q)})n_e \end{bmatrix} \begin{bmatrix} n_{H(p)} \\ n_{H(q)} \end{bmatrix} = \begin{bmatrix} C_{(1,p)} n_e \\ C_{(1,q)} n_e \end{bmatrix} n_H + \begin{bmatrix} (\alpha_{(p)}n_e + \beta_{(p)})n_e \\ (\alpha_{(p)}n_e + \beta_{(p)})n_e \end{bmatrix} n_{H^+} + \begin{bmatrix} -\Gamma_{H(p)} \\ -\Gamma_{H(q)} \end{bmatrix} = \begin{bmatrix} C_{(1,p)} n_e \\ C_{(1,q)} n_e \end{bmatrix} n_H + \begin{bmatrix} \alpha_{(p)}n_e + \beta_{(p)}n_e \\ \alpha_{(p)}n_e + \beta_{(p)}n_e \end{bmatrix} n_{H^+} + \begin{bmatrix} -\Gamma_{H(p)} \\ -\Gamma_{H(q)} \end{bmatrix} = \begin{bmatrix} C_{(1,p)} n_e \\ \alpha_{(p)}n_e + \beta_{(p)}n_e \end{bmatrix} n_H + \begin{bmatrix} 0 \\ \alpha_{(p)}n_e + \beta_{(p)}n_e \end{bmatrix} n_H + \begin{bmatrix} 0 \\ \alpha_{(p)}n_e + \beta_{(p)}n_e \end{bmatrix} n_H + \begin{bmatrix} 0 \\ \alpha_{(p)}n_e + \beta_{(p)}n_e \end{bmatrix} n_H + \begin{bmatrix} 0 \\ \alpha_{(p)}n_e + \beta_{(p)}n_e \end{bmatrix} n_H + \begin{bmatrix} 0 \\ \alpha_{(p)}n_e + \beta_{(p)}n_e \end{bmatrix} n_H + \begin{bmatrix} 0 \\ \alpha_{(p)}n_e + \beta_{(p)}n_e \end{bmatrix} n_H + \begin{bmatrix} 0 \\ \alpha_{(p)}n_e + \beta_{(p)}n_e \end{bmatrix} n_H + \begin{bmatrix} 0 \\ \alpha_{(p)}n_e + \beta_{(p)}n_e \end{bmatrix} n_H + \begin{bmatrix} 0 \\ \alpha_{(p)}n_e + \beta_{(p)}n_e \end{bmatrix} n_H + \begin{bmatrix} 0 \\ \alpha_{(p)}n_e + \beta_{(p)}n_e \end{bmatrix} n_H + \begin{bmatrix} 0 \\ \alpha_{(p)}n_e + \beta_{(p)}n_e \end{bmatrix} n_H + \begin{bmatrix} 0 \\ \alpha_{(p)}n_e + \beta_{(p)}n_e \end{bmatrix} n_H + \begin{bmatrix} 0 \\ \alpha_{(p)}n_e + \beta_{(p)}n_e \end{bmatrix} n_H + \begin{bmatrix} 0 \\ \alpha_{(p)}n_e + \beta_{(p)}n_e \end{bmatrix} n_H + \begin{bmatrix} 0 \\ \alpha_{(p)}n_e + \beta_{(p)}n_e \end{bmatrix} n_H + \begin{bmatrix} 0 \\ \alpha_{(p)}n_e + \beta_{(p)}n_e \end{bmatrix} n_H + \begin{bmatrix} 0 \\ \alpha_{(p)}n_e + \beta_{(p)}n_e \end{bmatrix} n_H + \begin{bmatrix} 0 \\ \alpha_{(p)}n_e + \beta_{(p)}n_e \end{bmatrix} n_H + \begin{bmatrix} 0 \\ \alpha_{(p)}n_e + \beta_{(p)}n_e \end{bmatrix} n_H + \begin{bmatrix} 0 \\ \alpha_{(p)}n_e + \beta_{(p)}n_e \end{bmatrix} n_H + \begin{bmatrix} 0 \\ \alpha_{(p)}n_e + \beta_{(p)}n_e \end{bmatrix} n_H + \begin{bmatrix} 0 \\ \alpha_{(p)}n_e + \beta_{(p)}n_e \end{bmatrix} n_H + \begin{bmatrix} 0 \\ \alpha_{(p)}n_e + \beta_{(p)}n_e \end{bmatrix} n_H + \begin{bmatrix} 0 \\ \alpha_{(p)}n_e + \beta_{(p)}n_e \end{bmatrix} n_H + \begin{bmatrix} 0 \\ \alpha_{(p)}n_e + \beta_{(p)}n_e + \beta_{(p)}n_e \end{bmatrix} n_H + \begin{bmatrix} 0 \\ \alpha_{(p)}n_e + \beta_{(p)}n_e + \beta_{(p)}n_e \end{bmatrix} n_H + \begin{bmatrix} 0 \\ \alpha_{(p)}n_e + \beta_{(p)}n_e + \beta_{(p)}n_e \end{bmatrix} n_H + \begin{bmatrix} 0 \\ \alpha_{(p)}n_e + \beta_{(p)}n_e + \beta_{(p)}n_e$$

$$\begin{bmatrix} n_{H(p)} \\ n_{H(q)} \end{bmatrix} = R_1 n_H + R_0 n_{H^+} + R_{ext}$$

 $R_1$ ,  $R_0$  and  $R_{ext}$  are population coefficients



#### **General EIRENE test case for testing rate coefficients**

- EIRENE 2D grid with ne and Te varied along x and y dimensions
- Ne, 20 points, ~1e8 1e16 cm3
- Te, 200 points, ~0.5 1e4 eV



#### The effective ionization rate of H colrad line perfectly with AMJUEL

Effective ionization rate  $S_{\text{eff}}$  :

$$S_{eff} = S_{(1)} + \sum_{p} \left( C_{(1,p)} - R_{1(p)} (F_{(p,1)} + \frac{A_1}{n_e}) \right)$$

Eirene CR -> H colrad Eirene AMJUEL -> using AMJUEL rates within Eirene

AMJUEL – last update entry: May 18





#### The H colrad recombination rate starts to diverge at Te > 400 eV

Effective recombination rate  $\alpha_{\text{eff}}$  :

$$\alpha_{eff} = \alpha_1 n_e + \beta_1 + \sum_p R_{0(p)} (F_{(p,1)} + \frac{A_1}{n_e})$$

• Discrepancy can be due to the different expressions of the recombination rate coefficient





#### The effective ionization cooling rate of H colrad line perfectly with AMJUEL





#### The H colrad recombination cooling rate starts to diverge at Te > 400 eV

Effective recombination cooling rate  $\alpha E_{eff}$ 

$$\begin{aligned} \alpha E_{eff} &= \beta_1 \bar{E}_1 - \alpha_{(1)} \, n_e E_\alpha \\ &+ \sum_p R_{0(p)} \left( S_p E_{p-\alpha} + \beta_p \bar{E}_p - \alpha_p n_e E_{p-\alpha} \right) \\ &+ \sum_{q>p} \left( R_{0(p)} C_{p,q} E_{p-q} - R_{0(q)} F_{(p,q)} E_{p-q} \right) \end{aligned}$$

• Directly derived from recombination rate, so same discrepancy occurs





#### **Proposed CRM structure in Eirene (H Colrad)**



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#### Proposed H2 Colrad structure in Eirene (currently non-existent)





# The previous (2002) photon tracing model (Reiter et al., PPCF 2002) was revisted

The photon tracing routine in EIRENE is analogous to neutral particle tracing, with v=c and E=hv, differences:

- Bulk ions are now excited species (photon sources)
- Rate coefficients for sources and photon-background interaction must take into account line shapes (natural, Doppler, Zeeman, etc)

Determine opacity with local population escape factors



#### **Determining the population escape factor**

$$\Theta_p = \frac{E - G}{E} = 1 - \frac{G}{E} \xrightarrow{\longrightarrow} \text{absorption}$$

$$\Theta_p = 1 - \frac{\int_{\Omega} \int_{line} \alpha(x,\lambda) L_{\lambda}(x,\lambda,\Omega) d\lambda d\Omega}{\int_{\Omega} \int_{line} \epsilon(x,\lambda) d\lambda d\Omega}$$

Behringer K 1998 Escape factors for line emission and population calculations MPI-Garching Report, IPP 10/11



#### $\Theta_p$ in Eirene

$$\Theta_p = \frac{E - G}{E} = 1 - \frac{G}{E} \longrightarrow \text{absorption}$$

G is simply the number of absorbed photons i.e volume photon sink tallies E is the volume photon source

Thus  $\Theta_p$  can be evaluated per cell of Eirene



#### Photon tracing test case

Cylindrical test case, 20 radial points

Homogeneous plasma and atomic density

 $T_{\rm H} = 1 \text{ eV}, n_{\rm H} = 10^{14} \text{ cm}^{-3}$  , b = 5 cm

Simulated Ly-a and Ly-b photons (2e6) with volumetric sources (H(n=2,3) as bulk ions)

Line shape only doppler broadening





## Population escape factor aligns with analytical function for Ly- $\alpha$ and Ly- $\beta$



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 $\tau = \alpha(\lambda)b$ is the optical depth or 'thickness'

Hollow points with ~10<sup>6</sup> photons Solid points with ~10<sup>9</sup> photons

Agreement for Ly- $\alpha$  at better statistics

## <u>2D (or 1D) profiles of the population escape factor: Ly- $\alpha$ and Ly- $\beta$ opaque at the center</u>





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#### Current project state

Whats on hand:

- H colrad, He colrad
- Photon module
- A&M and photon cylinder test cases

Whats planned:

- CRM-photon coupling (for Planck test)
- Application to JET 81472
- H colrad data update
- He colrad testing
- H<sub>2</sub> colrad creation



