For the 4th Technical Meeting on Divertor Concepts 7-10 Nov 2022



EIRENE modelling with improved CRMs for spectroscopic RT detachment control in EU-DEMO

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Outline

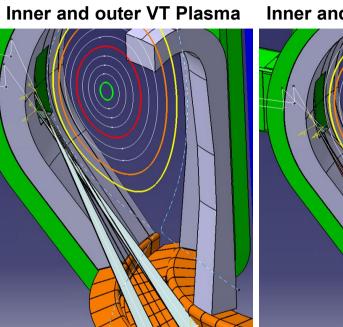


- 1) Collisional-radiative model (CRM) and real time control (RTC) of detachment
- 2) EIRENE code: CRMs and fluid-kinetic hybridisation (FKH)
- 3) EIRENE general development and restructuring: new CRM requirements
- 4) Standalone CRM for EU-DEMO: branching ratios, vibrostates effect.
- 5) HydKin development A&M data pre-processing

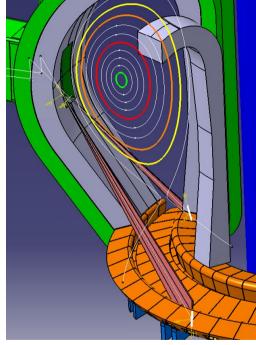


LOS under consideration for EU-DEMO





Inner and outer VT surface



Courtesy of D.Dunai, I.Katona, EUROfusion WP DC

Modelling for Real-time control (RTC) of detachment:

- What spatial resolution and sensitivity is needed?
- Is the current line-of-sight (LOS) choice optimal in general?
- What spectral range and resolution do we need?
- Which line(s) or line ratios do we choose to characterize the degree of detachment?
 - → Lyman + Balmer lines (H)? Fulcher band (H₂)? Impurity lines?..
- What will be effect of opacity (incl. reflection), line blending, etc?

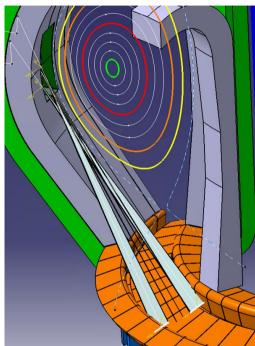


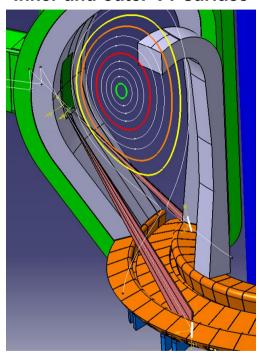
LOS under consideration for EU-DEMO



Inner and outer VT Plasma

Inner and outer VT surface





Modelling for Real-time control (RTC) of detachment:

What are the main show-stoppers from the EIRENE (SOLPS-ITER) modelling side?

- 1) Sufficient code performance on the DEMO scale
- 2) Including of sufficient physics
- Data consistency and reliability

Courtesy of D.Dunai, I.Katona, EUROfusion WP DC



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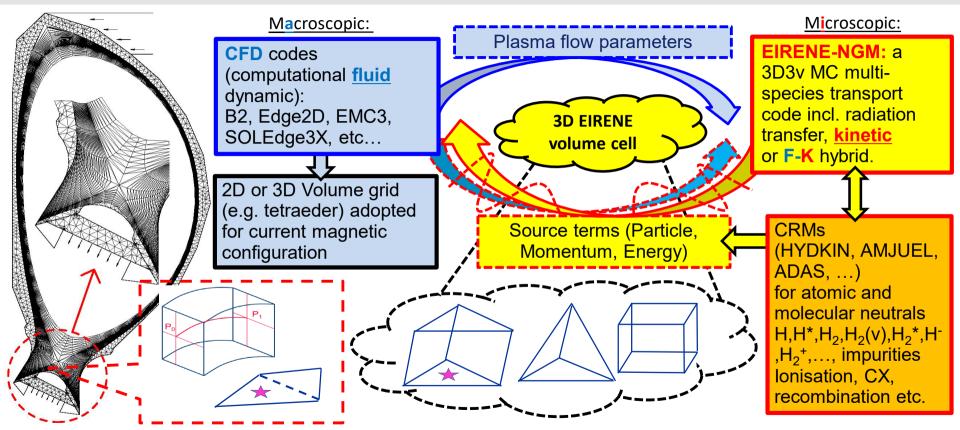


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What is EIRENE in a nutshell (e.g. in SOLPS)?..







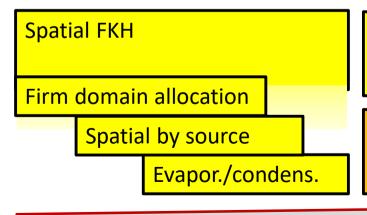
luid-kinetic hybridization (FKH)



A hierarchy of neutral models:

D.V. Borodin et al., FEC-2020, NF (2021)

Code performance Hybrid Fluid-Kinetic approach (FKH) AFN Asymptotic-Preserving MC (APMC) Kinetic, MC



Micro-macro FKH (mMH)

Cancellation error (can be suppressed by enforcing positivity)

Micro-macro kinetic diffusion (KDMC)

- Bias error (can be suppressed via multi-level)
- No coupling with CFD needed

Seamless in volume



Fluid-Kinetic Hybridisation inside EUROfusion TSVV-5



OPEN ACCESS IOP Publishing | International Atomic Energy Agency Nucl. Fusion 62 (2022) 086051 (12pp) https://doi.org/10.1088/1741-4326/ac3fe8

Fluid, kinetic and hybrid approaches for neutral and trace ion edge transport modelling in fusion devices

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W. Van Uytven<sup>2</sup>, M. Baelmans<sup>2</sup>, B. Mortier<sup>3</sup>, G. Samaey<sup>3</sup>
Y. Marandet<sup>4</sup>, P. Genesio<sup>4</sup>, H. Bufferand<sup>5</sup>, E. Westerhof<sup>6</sup>,
J. Gonzalez<sup>6</sup>, M. Groth<sup>7</sup>, A. Holm<sup>7</sup>, N. Horsten<sup>7</sup> and H.J. Leggate<sup>8</sup>
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The development continues according to the statement in the joint paper

- → Combining mMH/SpH/AFN, introducing "evaporation/condensation" (fluid ←→ kinetic consideration)
- → Developing **ML-KDMC** (a sort of "asymptotic preserving" MC) as useful alternative

- Continuing "brute force approach" for the OpenMP/MPI parallelisation as well as more profound parallelisation based on the "toy" model (EIRON).
- General restructuring of the code which includes generalisation of the CRMs



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General code development of EIRENE



- Main goals and decisions:
 - Segregating EIRENE into "starter" and a compact, free of most branching "numeric core"
 - → Optimisation for HPC, transparency and ease of use.
 - Unloading the main loop from branching agreed
 - → Abstractisation of the cell type (removing "geometry" branching)
 - → Abstractisation of A&M reactions + tracking of internal states
 - Version merging scope agreed including synchronisation with SOLPS-ITER
 - Uniform coding style (e.g. variable name conventions, use of OOP, etc.)

With support from the ACHs (Advanced Computing Hubs)

- EIRENE "toy model" EIRON and profiling of EIRENE
 - Test of parallelisation schemes incl. domain decomposition options
 - → Can be used for other purposes e.g. optimisation of ML-KDMC
- EUDAT-based simulation catalogue
 - → Portfolio of simulation cases
- IMASification of the simulation grid (GGD objects) in close contact with ITER (X.Bonnin) and utilizing the positive SOLPS-ITER experience



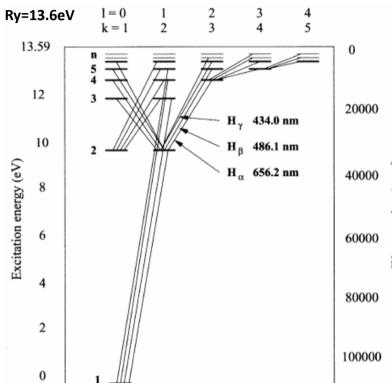


Eirene

Collisional-Radiative Models (CRM)



Grotrian diagram for atomic H (D, T)



Balance equations:

$$\frac{dN_i}{dt} = \sum_{j \neq i} A_{ji} N_j + n_e \cdot (EXCIT + IZ + CX + REC)$$

$$EXCIT = \sum_{j \neq i} \langle v \sigma_{ji} \rangle N_j$$

$$IZ = \sum_{m} \langle v\sigma_{mi} \rangle N_{m}^{-} + \sum_{z} \langle v\sigma_{zi} \rangle N_{z}^{2-} + \dots$$

j, k, l, m, z, ... states can be fine-superfine

resolved or, opposite, bundled into few quasi-

$$REC = \sum_{k} \langle v \sigma_{ki} \rangle N_{k}^{+} + \sum_{l} \langle v \sigma_{li} \rangle N_{l}^{2+} + \dots$$

Often used: $\langle v\sigma_{ii}\rangle(T_e, n_e)$ - effective Maxwellian averaged rates



D.Borodin | 4th IAEA TM on divertor concepts | 09.11.2022 | Page 11

CRM solvers available



CRM (technically) = reaction data + SOLVER

- \Box Most solvers solve algebraic **stationary** system of equations $\frac{dN_i}{dt} = 0$
- → in such cases the solver is much less valuable as the A&M data collection
- → this statement is underlined even by the developers of the very well established and mostly up-to-date YACORA model

 D.Wünderlich and U.Fantz,

 Atoms 2016, 4(4), 26
- ☐ CRM containing molecular species is even more complex, e.g. one needs to track more processes and more states (vibrational, rotational).
- → Tracking all the states as separate species demands enormous CPU and memory resources.
- □ EIRENE pre-calculates rates for each volume cell. It also allows to pre-calculate values for various T_i/T_e ratios.



New CRM Solver for EIRENE concepted



This CRM is aimed to precompute rate coefficients accounting for all parametric **dependences** (n_a, T_a, but also T_i, ...) in contrast with currently used polynomial fits (AMJUEL, ...) + add a number of levels/processes not accounted for at this time



☐ The **internal states** (e.g. rovibrational states in molecular species) are to be tracked with a flexible a flexible control over this resolution (as separate specie or variable).



The **nonstationary solution** for balance equations should be the default one (with the stationary only as a useful option).



- The solver should be **modular**, thus **usable standalone** or even in other codes.
- The **improved A&M data input** should be readable and structured (for starters JSON, potentially also HDF5). It should be pre-processed mostly automatic and easily exchanged with other codes and tools. We need tools for visualisation and testing.
 - The only way to meet the exploding amounts of data from RMPS and CCC for molecules (with resolution by rovibrational states)
 - ➤ IAEA GNAMPP assists, but also reveals the challenges AMD Unit: GNAMPP

Not only performance and reliability to be improved, however including additional physics can be provided!



Outline

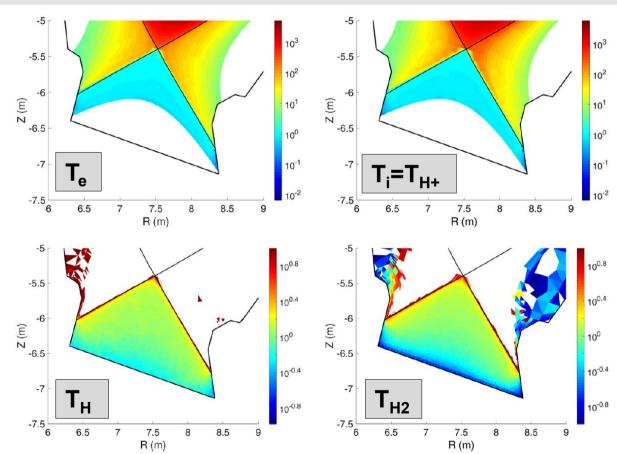


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SOLPS-ITER basic simulation case for EU-DEMO





F.Subba et al., NF 61 (2021) 106013

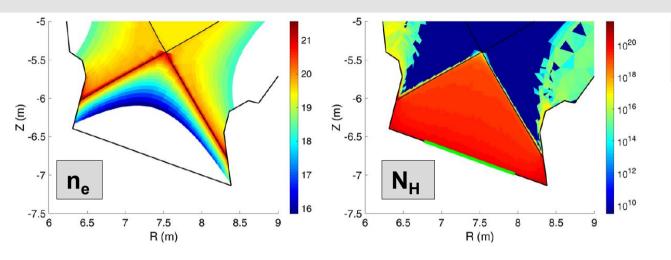
Below X-point:

-) $T_e = T_i = 10eV$
- 2) $T(H)=T(H_2)=10eV$
- 3) $N_e \sim 10e18 \text{ m}^{-3}$
- 4) N(H)=10e20 m⁻³



SOLPS-ITER basic simulation case for EU-DEMO





F.Subba et al., NF 61 (2021) 106013

Below X-point:

- $T_e = T_i = 10eV$
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- $N(H)=10e20 \text{ m}^{-3}$

- These are semi-detached conditions (1 point)
 - → Attached vs. detached + 2-3 semidetached cases would be wishful to have!
 - → We use EIRENE or standalone CRM to identify spectral features useful for the degree of detachment control for this plasma conditions
- We can run (and mean to in future!) full EIRENE: SOLPS with a "frozen" fluid side (B2.5)
 - → This option is not mature enough for meaningful physics results
 - → We use standalone CRM and vary consrtant plsma conditions around 🂢



YACORA: inclusion molecular source term



 $H_2(v=0)$ $H_2(v=1)$

 $H_2(v=2)$

 $H_2(v=3)$ $H_2(v=4)$

 $H_2(v=5)$

 $H_2(v=6)$

 $H_2(v=7)$

 $H_2(v=8)$

 $H_2(v=9)$

 $H_2(v=10)$

 $H_2(v=11)$

 $H_2(v=12)$

 $H_2(v=13)$

 $H_2(v=14)$

Stationary run vibrationally resolved:

 p, H_2^+, Z **Particle species**

 $H_2(v=0),\ldots,H_2(v=14)$

YACORA (+ Source term) adjustable to the Edge2D-EIRENE

F.Cianfrani, **EPS-2022**

Reactions

electron impact

proton impact

electron impact

vibrational transitions $H_2 + e \rightarrow H_2(b_{\text{triplet}}) \rightarrow H^+ + H^+ + e + e$

 $H_2 + e \to H_2^+ + e + e$ $H_2 + p \rightarrow H_2^+ + p + e$

 $H_2 + e \rightarrow H^+ + H^+ + e + e$ $H_2(v=0) \rightleftharpoons H_2(v=1)$ $H_2(v=1) \rightleftharpoons H_2(v=2)$

 $H_2(v=13) \rightleftharpoons H_2(v=14)$

 H_2 source term $\langle Z \rightarrow H_2(v=0) \rangle$

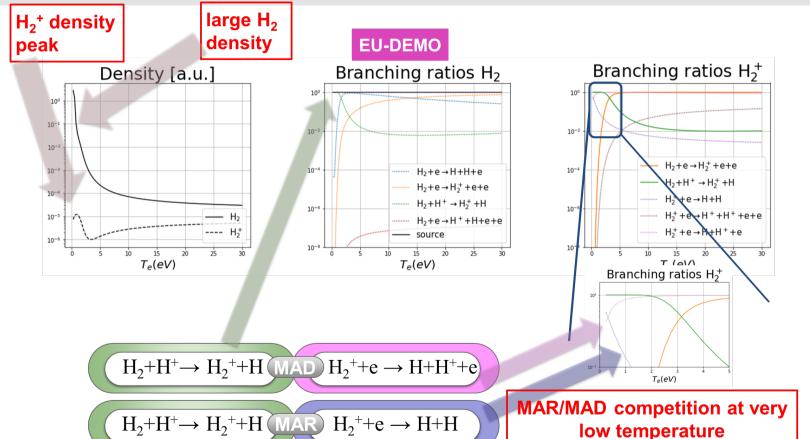
 10^{3} 100 10^{-3} 10-6

 T_e (ev)

 H_2 density (m⁻³)

DEMO-relevant conditions – branching ratios

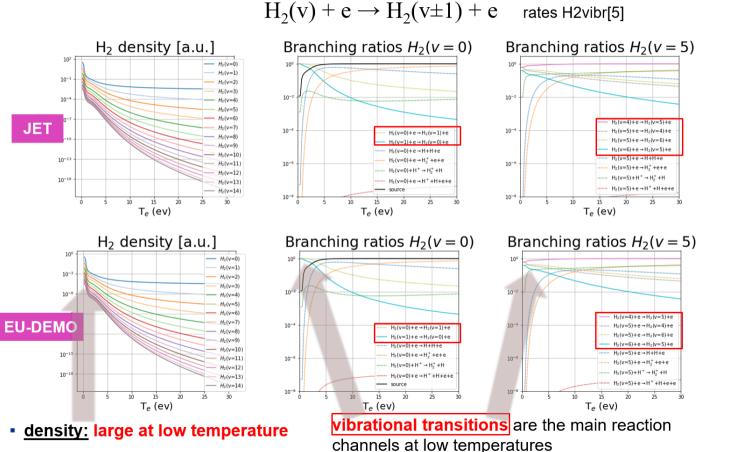






EU-DEMO– effect of resolution by vibrostates





F.Cianfrani et al., EPS-2022

Fully in line with the EIRENE and other CRMs: "Up to 40% reduction in effective dissociation rate due to transport of vibrational states"

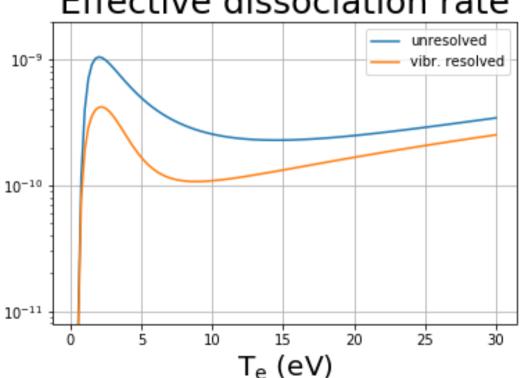
A.Holm, M.Groth, et al., PET, CPP 2021



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HYDKIN (web-based A&M data tool):



- toolbox for plotting and manipulation of A&M data with a user-friendly graphical interface. D.Reiter et al.,
- includes EIRENE databases: AMJUEL, HYDHEL, H2VIBR, METHANE, ...



available at http://www.eirene.de/hydkin/ Password-protected (for access contact the authors).

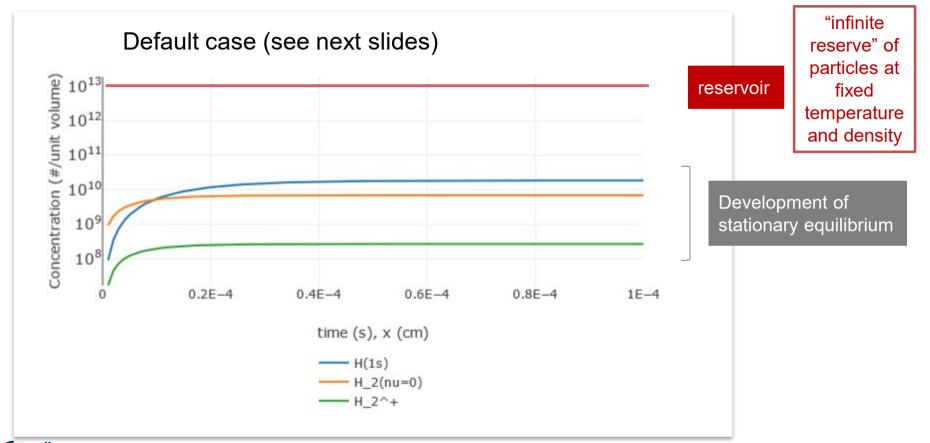
F.Cianfrani et al., ICAMDATA-2022

Phys. Scr. 2009



HYDKIN: particle solver (non-stationary)





HYDKIN: what is inside?..



- CRM solver (transport excluded, pure A&M side of the problem) with extensive features
 - → Stationary and **non-stationary** solution (assuming velocity and plasma pars)
 - → Any specie can be treated as "Reservoir"
 - → "Spectral analysis" based on eigenvalue approach (which reactions are most important?)
 - → Sensitivity analysis vary the plasma parameters, solver settings, reactions, etc.
- Plotting, solver results visualisation
 - → Useful for fast **consistency checks** and trivial analysis while constructing the CRMs
 - → Good for physics analysis in case one follows **established routines**
- Flexible reaction tables interconnected with the solver and plotting
 - → Useful for fast **consistency checks** and trivial analysis while constructing the CRMs
 - → Good for physics analysis in case one follows **established routines**
 - → Input/output in various formats (new: JSON) including directly for EIRENE



HYDKIN: using the power of the web-interface



For H₂ case (example):

Start with own configuration

Start with default-Cases:

Unresolved

Vibrationally resolved 7

Vibrationally resolved 14

- ☐ The basis table is highly customisable:
 - Deselected reaction can be excluded from the view
 - 2) User easily selects the level of info for each reaction to be shown
 - 3) Basic, solver, plotting etc. tables are interlinked
- All is optimised for checking high amounts of similar data
 - 1) Critical things are highlighted
 - 2) Data is grouped by the reactions (even if comes from various sources)
 - 3) One can start with initial configuration (standard or custom) and do only essential (e.g. from physics) changes



HYDKIN – new mode (H₂ case)

Version 2022:

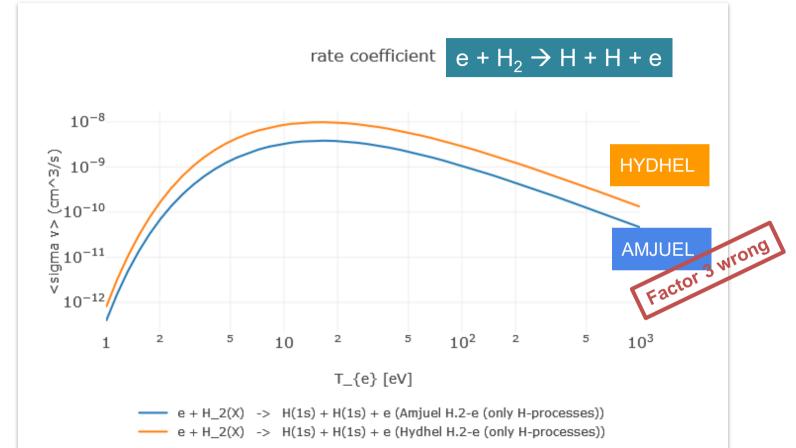
AMJUEL/HYDHEL sorted by reactions

Collision with e selected not included plot Number reaction reference lrange Visibility columns: **Buttons** unselect unselect all solver reaction: switch to off to adjust E min: range: switch to off 1.08e+01 e + H(1s) → \checkmark 2.1.1 JanevEtAl(1987) table H(2p) + eF max: reference: switch to off 2.00e+04 content T min: data type: switch to off 1.26e+00 e + H(1s) → **V** JanevEtAl(1987) H(2p) + eT max: Peculiar properties: switch to on 2.00e+04 generation: switch to on E min: e + H(1s) → 1.08e+01 2.1.2 \checkmark JanevEtAl(1987) data origin: switch to on H(2s)+ e E max: 2.00e+04 File/chapter: switch to off T min: 1.26e+00 e + H(1s) → \checkmark JanevEtAl(1987) H(2s)+ e - T max: 2 00e+04 show only selected reactions (Groups) min: 1.00e-01 T max: \Box n=1 -> n'=2 $e + H(n) \rightarrow$ 1.00e+03 SawadaFujimoto(199 H(n') + en min: 1.00e+08 show only selected reactions (rows) n max: 1.00e+16



HYDKIN: comparison dissociation rates







HYDKIN can be used to



- to import/export data (JSON, tabular and other formats)
- to produce input data for EIRENE and for other codes
- to check data for consistency, abnormal features, etc.
- to check and improve the results of the simulation
- to understand A&M side of the problem and identify the most significant processes (among the selected ones)
- to load/improve/save the developed configuration (selected reactions and parameters) including starting from the standard pre-sets



Summary and outlook



- The development of EIRENE as "neutral gas module" for the integrated HPC-oriented simulations is ongoing
 - → 2 lines of Fluid-Kinetic hybridisation are exploited
 - → Parallelisation is ongoing including testing the parallelisation and domain decomposition schemes with the EIRON "toy"-model and support from the "advanced computing hub".
 - → Code restructuring is aimed to remove branching and segregate compact core
 - → CRM improvement is very significant part of the improvement
- New CRM is concepted and under development
 - → It should be used both in EIRENE and standalone
 - → We use YACORA (with additional source introduced) to gain useful experience
- HydKin is largely updated and foreseen to be the main A&M pre-processing tool
 - → All the development is ongoing in a view of **future massive data** extension
 - → It is meant to provide flexible adjustment of CRMs and auto generation of EIRENE input
- Application to the EU-DEMO case (P.Subba) shows that
 - → Even **standalone model is useful** to understand the critical branching ratios
 - → Tracking of **vibrational states** is essential
 - → Full scale SOLPS-ITER runs, thus further development, is needed for tasks like RTC in DEMO





Thanks for the attention!

