

Chemical networks with KROME

(getting started)



KROME needs a chemical network

```
$ python3 krome
```

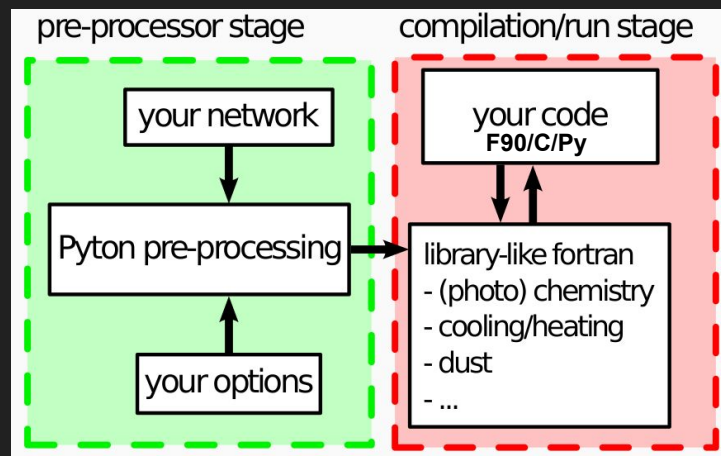
```
*****
```

```
WELCOME TO KROME
```

```
*****
```

```
ERROR: you must define -n FILENAME or -network FILENAME, where  
FILENAME is the reaction file!
```

```
$ python3 krome -n my_network.dat
```



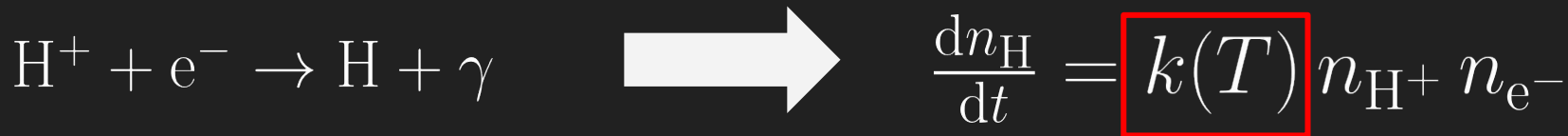
(answer in this presentation)

Choose a chemical network (it's problem-dependent)

GIGO: your model cannot be better than the quality of your data

1. Use one of the pre-built KROME networks (networks/ folder, with caution)
2. Use the KIDA → KROME converter tool to select reactions from standard database (automatic selection is dangerous)
3. Find a chemical network in literature (recommended, updates?)
4. Prepare your own network based on databases, literature, experts, and pre-existing networks (ideal, time consuming)

A simple example



KIDA | KINETIC DATABASE FOR ASTROCHEMISTRY

Home Species Download References Help

H+ + e-

Your query may contain lot of results.
Try some of these tips to filter results:

REACTANTS | PRODUCTS | BOTH

ASTRO | PLANETO | BOTH

ION + NEUTRAL | NEUTRAL

SEARCH

SPECIES 2 | UNIMOLECULAR REACTIONS 3 | **BIMOLECULAR REACTIONS 1** | TERMOLECULAR REACTIONS 0 | SURFACE REACTIONS 0

1 result(s)

Type	Reaction	α	β	γ	T (K)	Formula ⁱ	Evaluation [?]
ER	H+ + e- → H + Photon	3.50E-12	-7.00E-1	0.00E+0	10-280	Modified Arrhenius equation	[?]

Arrhenius

$$k(T) = \alpha \left(\frac{T}{300 \text{ K}} \right)^\beta \exp \left(-\frac{\gamma}{T} \right)$$

A simple example / 2



α	β	γ	$T(K)$
3.50E-12	-7.00E-1	0.00E+0	10-280

@format: idx, R, R, P, Tmin, Tmax, rate

1, H+, E, H, 10, 280, 3.5e-12 * (Tgas / 3e2)**(-0.7)

Service variables

@var: T32=Tgas/3e2

1, H+, E, H, 10, 280, 3.5e-12 * T32**(-0.7)

↑
speed-up

Build by KROME →

$$\frac{dn_{\text{H}}}{dt} = k(T) n_{\text{H}^+} n_{\text{e}^-}$$

$$k(T) = \alpha \left(\frac{T}{300 \text{ K}} \right)^{\beta} \exp \left(-\frac{\gamma}{T} \right)$$

Custom functions

$$k(T) = \alpha \left(\frac{T}{300 \text{ K}} \right)^\beta \exp\left(-\frac{\gamma}{T}\right)$$

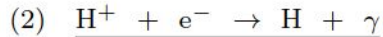
1, H+, E, H, 10, 280, k_arr(3.5d-12, -0.7d0, 0d0, Tgas)

Defined as a F90 function in a dedicated file



Janev et al. 1987 (2.1.5)

$$k_1 = \exp[-32.71396786 + 13.536556 \ln(T) - 5.73932875 \ln(T)^2 + 1.56315498 \ln(T)^3 - 0.2877056 \ln(T)^4 + 3.48255977 \times 10^{-2} \times \ln(T)^5 - 2.63197617 \times 10^{-3} \times \ln(T)^6 + 1.11954395 \times 10^{-4} \ln(T)^7 - 2.03914985 \times 10^{-6} \ln(T)^8] \text{ cm}^3 \text{ s}^{-1}.$$



Our fit to data given by Ferland et al. (1992)

$$k_2 = \exp[-28.6130338 - 0.72411256 \ln(T) - 2.02604473 \times 10^{-2} \ln(T)^2 - 2.38086188 \times 10^{-3} \ln(T)^3 - 3.21260521 \times 10^{-4} \ln(T)^4 - 1.42150291 \times 10^{-5} \ln(T)^5 + 4.98910892 \times 10^{-6} \ln(T)^6 + 5.75561414 \times 10^{-7} \ln(T)^7 - 1.85676704 \times 10^{-8} \ln(T)^8 - 3.07113524 \times 10^{-9} \ln(T)^9] \text{ cm}^3 \text{ s}^{-1}.$$

A simple example / 3



$$\frac{dn_{\text{H}}}{dt} = k_1 n_{\text{H}^+} n_{\text{e}^-} - k_2 n_{\text{H}}$$

$$k_2 = \alpha \zeta$$

Type	Reaction		α	β	γ	UV Field	Formula ?	Evaluation
CR	H + CR	→ H ⁺ + e ⁻	4.60E-1	0.00E+0	0.00E+0		Cosmic-ray ionization	?
Phot	H + Photon	→ H ⁺ + e ⁻	0.00E+0	0.00E+0	3.00E+0	Draine	Photo (Draine)	✓

Preprocessor stage in your network

@format: idx,R,P,P,rate

@common: user_crate

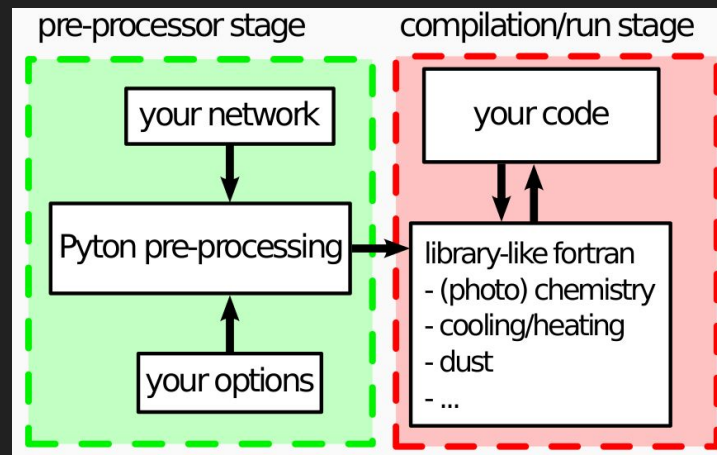
2, H, H+, E, 0.46 * user_crate

model parameter

Change the value at runtime from your code

call krome_set_user_crate(1d-15)

[1/s]



A simple example (network recap)

```
@var: T32=Tgas/3e2
```

```
@common: user_crate
```

```
@format: idx,R,R,P,Tmin,Tmax,rate
```

```
1, H+, E, H, 10, 280, 3.5e-12 * T32**(-0.7)
```

```
@format: idx,R,P,P,rate
```

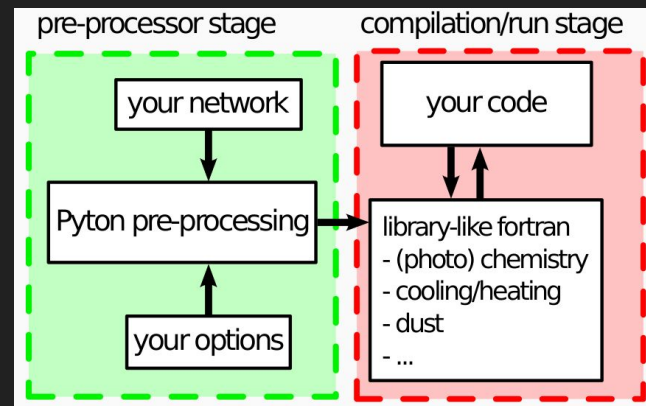
```
@CR_start ← For CR heating
```

```
@storeOnce_start ← To save evaluations
```

```
2, H, H+, E, 0.46 * user_crate
```

```
@storeOnce_stop
```

```
@CR_stop
```



Pre-processing *simple.ntw* (python)

```
tgrassi@multivac:~/repos/krome$ python3 krome -n=simple.ntw
```

```
*****
```

```
WELCOME TO KROME
```

```
*****
```

```
Any key to ignore q to quit...
```

```
Reading from file "simple.ntw"...
```

```
copied H.gfe
```

```
checking sinks/sources...
```

```
checking recombinations...
```

```
done!
```

```
Enthalpy OK!
```

```
ODEs needed: 7
```

```
Reactions found: 2
```

```
Species found: 3
```

```
Species list, reactions, and info saved in build/info.log
```

```
Species index initialization for gnuplot in build/species.gps
```

```
Heating cooling index init for gnuplot in build/heatcool.gps
```

```
Reactions saved in build/reactions.log
```

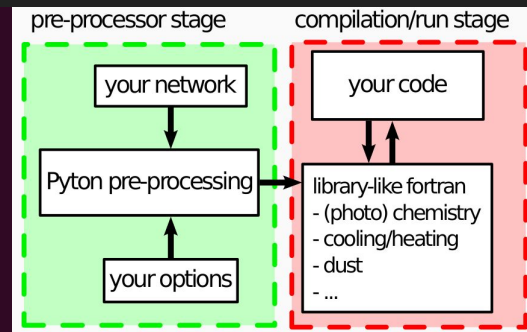
```
ODE partition: [3 atom/mols] + [1 CR] + [1 PHOT] + [1 Tgas] + [1 dummy] = 7 ODEs
```

```
ODEs list: E, H, H+, CR, g, Tgas, dummy
```

```
Max number of reactants: 2
```

```
Max number of products: 2
```

Network checks



Runtime stage (fortran)

```
program test
```

```
use krome_main !use krome (mandatory)  
use krome_user !use utility (for krome_idx_* constants and others)  
implicit none  
integer,parameter::nsp=krome_nmols !number of species (common)  
real*8::Tgas,dt,x(nsp),spy
```

```
spy = 3.65d2 * 2.4d1 * 3.6d3 !seconds per year
```

```
call krome_init() !init krome (mandatory)
```

```
x(:) = 0d0 !default abundances  
x(krome_idx_H) = 1d0 !hydrogen initial abundance
```

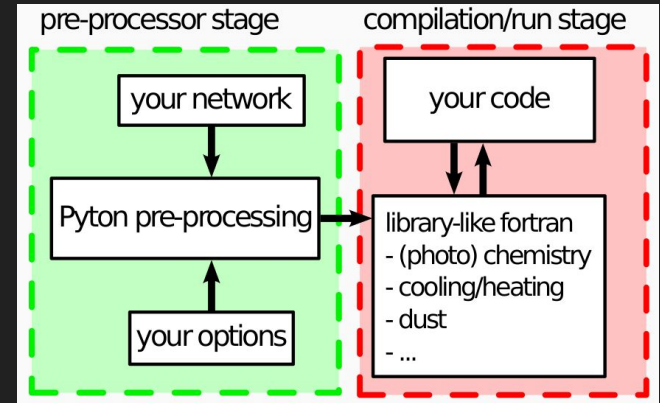
```
Tgas = 270. !gas temperature (K)  
dt = 1d6 * spy !time-step (s)
```

```
call krome_set_user_crate(1d-12)
```

```
!call the solver  
print *, x(krome_idx_H), x(krome_idx_Hj), x(krome_idx_e)  
call krome(x(:), Tgas, dt) !call KROME  
print *, x(krome_idx_H), x(krome_idx_Hj), x(krome_idx_e)
```

```
print *, "Test OK!"
```

```
end program test
```



Additional examples

```
#PHOTOREACTIONS THIN WITH Av
@format:idx,R,P,P,rate


@storeOnce_start
# FROM GLOVER+2009, https://arxiv.org/abs/0907.4081
224,H-,H,E,7.1d-7*exp(-.5*user_Av)
225,H2+,H,H+,1.1d-9*exp(-1.9*user_Av)
227,H3+,H2,H+,4.9d-13*exp(-1.8*user_Av)
228,H3+,H2+,H,4.9d-13*exp(-2.3*user_Av)
229,C,C+,E,3.1d-10*exp(-3.*user_Av)
230,C-,C,E,2.4d-7*exp(-.9*user_Av)
231,CH,C,H,8.7d-10*exp(-1.2*user_Av)
232,CH,CH+,E,7.7d-10*exp(-2.8*user_Av)
233,CH+,C,H+,2.6d-10*exp(-2.5*user_Av)
234,CH2,CH,H,7.1d-10*exp(-1.7*user_Av)
235,CH2,CH2+,E,5.9d-10*exp(-2.3*user_Av)
236,CH2+,CH+,H,4.6d-10*exp(-1.7*user_Av)
237,CH3+,CH2+,H,1d-9*exp(-1.7*user_Av)
238,CH3+,CH+,H2,1d-9*exp(-1.7*user_Av)
239,C2,C,C,1.5d-10*exp(-2.1*user_Av)
240,O-,O,E,2.4d-7*exp(-.5*user_Av)
241,OH,O,H,3.7d-10*exp(-1.7*user_Av)
242,OH,OH+,E,1.6d-12*exp(-3.1*user_Av)
243,OH+,O,H+,1d-12*exp(-1.8*user_Av)
244,H2O,OH,H,6d-10*exp(-1.7*user_Av)
245,H2O,H2O+,E,3.2d-11*exp(-3.9*user_Av)
254,O2,O2+,E,5.6d-11*exp(-3.7*user_Av)
255,O2,O,O,7d-10*exp(-1.8*user_Av)
256,CO,C,O,2.d-10*exp(-3.53*user_Av)
@storeOnce_stop
```

```
@noTab_start
#DUST FREEZE-OUT/EVAPORATION
@var:tabTdust = get_table_Tdust(n(:))
@ice CO,freezeout,krate_stickSi(n(:),idx_CO,tabTdust)
@ice CO,evaporation,krate_evaporation(n(:),idx_CO,tabTdust)
@ice H2O,freezeout,krate_stickSi(n(:),idx_H2O,tabTdust)
@ice H2O,evaporation,krate_evaporation(n(:),idx_H2O,tabTdust)

#NON-THERMAL EVAPORATION, Yield=1e-3, Gnot=1.78 is Draine Flux
@ice:CO,evaporation,krate_nonthermal_evaporation(idx_CO,1.78d0, user_Av, user_crate, 1d-3)

@noTab_stop
```

```
@format:idx,R,R,P,P,P,rate
@var:invTe=1d0/Te
@var:logT=log10(Tgas)
@var:invsqrT=1d0/sqrt(Tgas)
@var:kl11 = 1d1**(-27.029d0+3.801d0*logT-29487d0*invT)
@var:kh11 = 1d1**(-2.729d0-1.75d0*logT-23474d0*invT)
@var:ncr11 = 1d1**(5.0792d0*(1d0-1.23d-5*(Tgas-2d3)))
@var:ax11=1.d0/(1.d0+(Hnuclei/(ncr11+1d-40)))
@noTabNext
11,H2,He,H,H,He,kh11**(1.-ax11)*kl11**ax11
```



Recap

- How to prepare a simple chemical network
- Search into a database / literature
- Pre-process your chemical network
- Evolve a chemical network in time

