

# The Kinetic Monte Carlo method

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- Monte-Carlo refers to a broad class of algorithms that solve problems using random numbers.
- The stochastic method has existed since well before the first computer and was e.g. used by Enrico Fermi in 1930.

 One useful instance is the Metropolis algorithm\* which allows by very simple means to find the equilibrium state of a system.





## Applications range from:

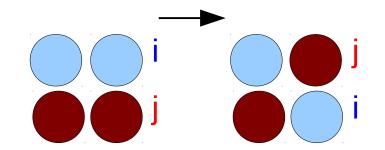
- Graphic ray tracing
- Semiconductor Carrier simulation
- Financial analysis
- Numerical integration
- Neutron transport
- Diffusion in solids
- etc...



## Metropolis Monte-Carlo:

Classically: Transmutation MC, switching of particles

Switch acceptance based on the probability ratio between initial (ij) and final (ji) state:



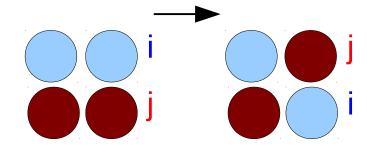
If 
$$P_{ij \to ji} = exp(-\Delta U/k_BT) > 1$$
 then accept,

If not, accept with probability  $P_{ij \rightarrow ji}$  where  $\Delta U$  is the potential energy difference between the states ij and ji.



## Metropolis Monte-Carlo:

Efficient algorithm for finding equilibrium configurations. "Path" to final state without physical meaning.



No information on the time scale.

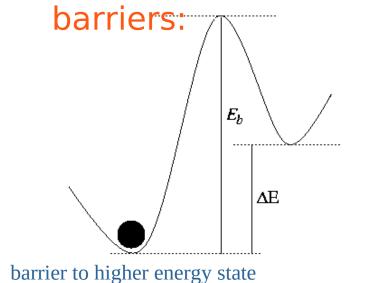
However, coupled with cohesive model, provides path to thermodynamic integration (if atomic relaxation is allowed).

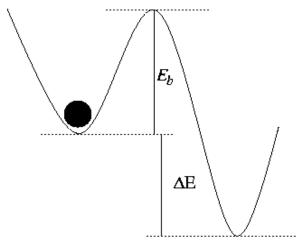


#### **Time for Kinetic Monte-Carlo**

How do we treat the time evolution of Question: systems defined by diffusion processes?

Diffusionerally activated by jumps over energy





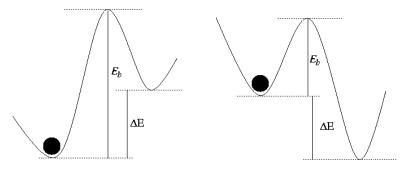
barrier to lower energy state

e.g. thermal activation: 
$$R(T) = R_0 e^{-E_b/k_B T}$$



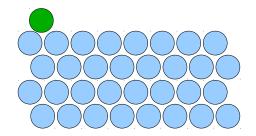
#### **Time for Kinetic Monte-Carlo**

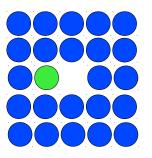
## Surface: Hopping, exchange, adsorption, desorption, etc



$$R(T) = R_0 e^{-E_b/k_B T}$$

Bulk: Vacancy driven, self-interstitial driven, diffusion of interstitial elements.







#### **Time for Kinetic Monte-Carlo**

How to advance the system:  $R^i(T) = R_0^i e^{-E_b^i/k_BT}$ 

1) Use short time step  $\Delta t$ . For each step go trough all particles i and randomly choose according to  $R^i$  if a process is activated.

Limit:  $\Delta t \ll 1/R_{max} \rightarrow very slow algorithm if rates are strongly varying$ 

2) Pick a particle *i* at random, with probability proportional to R<sup>i</sup>. Every step produces a jump; time step association easy.

The KMC method!



#### **Outline**

#### 1. Introduction

- Origin of Monte-Carlo methods
- ☐ Time for Kinetic Monte-Carlo

#### 2. Kinetic Monte-Carlo method

- Physical motivation
- Mathematical formalism
- Numerical recipie
- Range of applicability
- Advanced methods

#### 3. Applications

- Atomistic KMC thermal ageing & microstructure evolution
- ☐ Object KMC resistivity recovery



### KMC – History

Atomistic Monte-Carlo: The Metropolis algorithm (1953) No timescale!

#### Kinetic Monte-Carlo: ?

- 1) Flinn and McManus, Phys. Rev. 124 (1961) 54. Vacancy driven but no explicit time
- 2) Young and Elcock, Proc. Phys. Soc. 89 (1966) 735. Simple time scale
- 3) Bortz, Kalos and Lebowitz, J. Comp. Phys. 17 (1975) 10. Ising sping KMC with rigorous time scale (no reference to Young and Elcock...)



## **KMC** – History

#### Kinetic Monte-Carlo:

The algorithm derived in the BKL paper is usually called the residence time algorithm (you will see why), also called the n-fold way, the BKL method, or more generally the KMC method.

#### Modern reviews:

Fichtorn and Weinberg, J. Chem. Phys. 95 (1991) 1090.

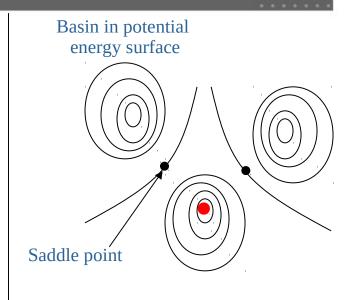
Voter, in Radiation Effects in Solids, (Springer, NATO Publishing Unit, Dordrecht, The Netherlands, 2005)



Many interesting processes consist mainly of diffusive jumps from one state to another.

Skip vibrational dynamics and go for the heart of the problem: The state-tostate transitions.

Normally the transition from one basin to another is an infrequent event, i.e.  $t_{in basin} >> t_{transition}$ 



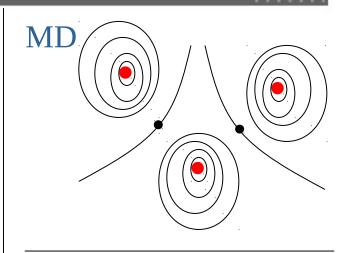
Thus the memory of how each transition transpired is lost. All transitions are independent of the system history.

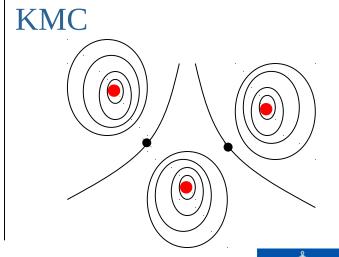
Thus, for each transition from one basin i to another j, there is a rate constant  $r_{ij}$  that characterizes the probability to escape from i to end up in j.

No dependency on state i-1 exists:

→ a Markov chain.

If all  $\{r_{ij}\}$  are known, the system trajectory will be indistinguishable from that of a fully dynamic simulation.







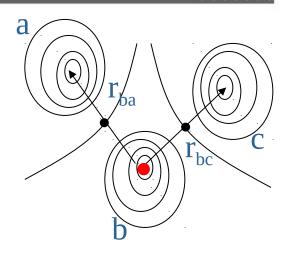
Due to the memory loss of the system, the probability of transition is identical for all equal (and short) time steps t.

⇒ a process with first-order decay statistics (Poisson process), e.g. radioactivity.

The probability of the system staying in a certain basin is:

$$P_{\text{stay}}(t) = \exp(-r_{\text{tot}}t),$$

where  $r_{tot}$  is the total rate for escape from the particular state.

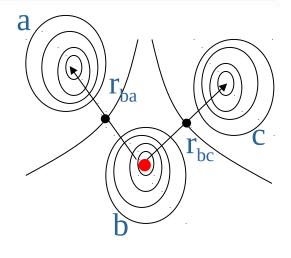


$$r_{tot} = r_{ba} + r_{bc}$$



#### Requirements:

- a) The transitions are Poisson processes
- b) The processes are independent
- c) The time increment is calculated properly



$$r_{tot} = r_{ba} + r_{bc}$$



## KMC - Numerical recipie: meta code

- 0. Set time t=0.
- 1. Form a list of all rates  $\mathbf{r}_i$  of all possible transitions  $\mathbf{W}_i$  in the system.
- 2. Calculate the total rate  $R_i = \sum_j r_j$  for i = 1,...,N where N is the total number of transitions.
- 3. Get a uniform random number  $c \in (0,1]$ .
- 4. Find the event *i* corresponding to c:  $R_{i-1} < cR < R_i$
- 5. Carry out event *i*.
- 6. Find all  $W_i$  and recalculate all  $r_i$  that may have changed.
- 7. Get a new uniform random number  $c \in (0,1]$ .
- 8. Update the time with  $t = t + \Delta t$  where:
- 9. Return to step 1.  $\Delta t = -\frac{ln(c)}{R}$



## KMC – Numerical recipie

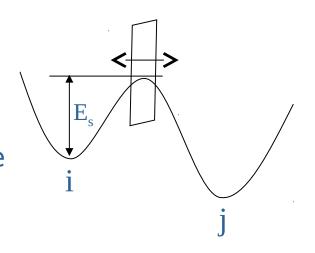
How to calculate the rate constants?

Transition state theory gives us a straightforward scheme:

**r**<sub>ij</sub> is given by the equilibrium flux through the surface separating the two states (the saddle point).

This can be simulated using e.g. the Metropolis algorithm.

However, to further simplify, we can assume a harmonic transition state.



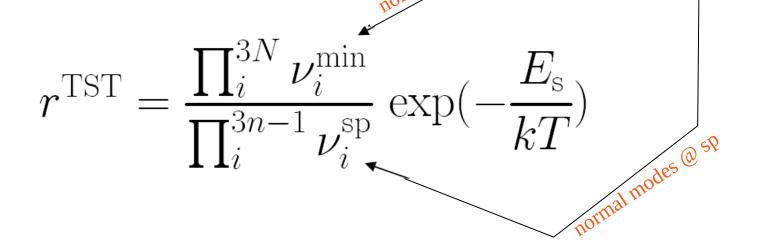


## KMC - Numerical recipie

Harmonic transition state theory (Vineyard theory).

Only allows passage through the saddle point.

Vibration modes around the saddle point are harmonic.





## **Transition rates – drag method**

Choose appropriate reaction coordinate q

Constrain q and relax all other degrees of freedom

Repeat procedure for several q between initial and final state

- highly dependent on good guess for reaction coordinate
- if true reaction coordinate has a large component perpendicular to the initial guess, the method will yield a discontinuous reaction path!

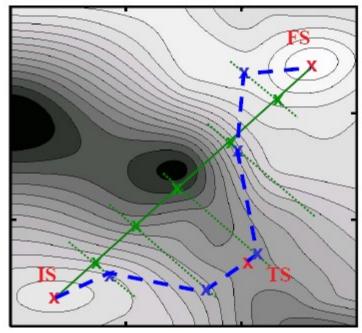


Image from Peter Kratzer



#### **Transition rates – NEB method**

#### Nudged elastic band:

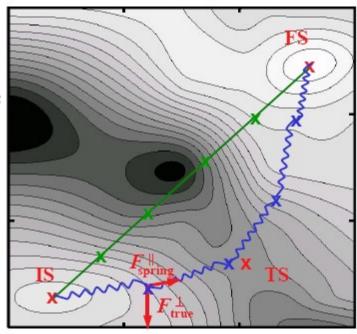
Initialize with several images  $\{R_i\}$  along a straight-line interpolation Minimize

Problem: 
$$S(R_1, ..., R_N) = \sum_i E(R_i) + \sum_i \frac{k}{2} (R_{i+1} - R_i)^2$$

- elastic band cuts corners
- images tend to slide down towards lowenergy IS/FS regions, leaving few images for relevant TS region

#### Solution:

- only spring force component parallel to path (no corner cutting)
- only true force component perpendicular to path (no down-sliding)



G. Mills and H. Jónsson, Phys. Rev. Lett. **72**, 1124 (1994) Image from Peter Kratzer



KMC does give a trajectory identical to a fully dynamic simulation if (and only if) the rate catalogue is *complete*.

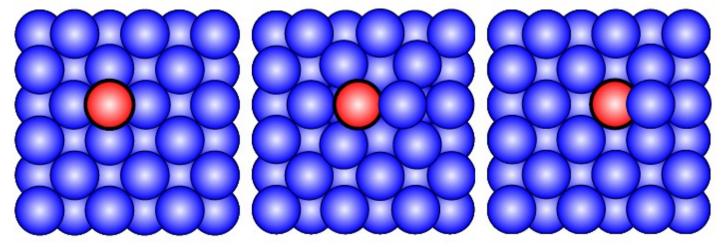
In simple cases (1D, etc) this condition might be fulfilled but in complex systems like 3D solids or surfaces it is *not* trivial!

If only our intuition is allowed to define the rate catalogue we will miss processes.

The question is how important they are...



Example from Voter: Adatom exchange on fcc(100) [Pt, Ir, Al]



Exchange mechanism discovered by DFT (Feibelman 1990)

Previous studies assumed surface hopping: simulations not in agreement with experiments.



#### Computational scaling

Depends on:

the size N of the rate catalogue.

which in turn depends on the system size unless the density of "jumpers" is not constant.

Special cases exist where computation time is independent on N.

Below-N scaling is possible using binning methods and recursive tree searches.

Place all events with rate  $r_i$  into bin j Select events by first choosing bin, then randomly inside the bin.

Scales as log(N)

Maksym, Semicond. Sci. Technol. 3 (1988) 594.



#### Advantages:

Simulated objects can change character: reactions introducing new reaction products annihilation/recombination emission

The  $r_{tot}$  is updated all the time: time scale varies with population

Fast moving population evolving into slow moving population dynamically changes the time scale.

Several orders of magnitude in time can be treated in one simulation!

initially: Δt~fs, finally: Δt~h



## Disadvantages:

All rates and reaction have to be known.

Method cannot predict new reactions! (unlike MD)

Exp/ab initio/MD can give rates and reactions. (complex combinatorics, model simplifications, etc)

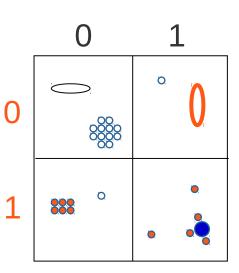


## KMC lab next!



How do we reach longer time scales or larger simulation boxes?

"Standard" solution: parallelize the problem. Decompose the domain and share the load between processors.



One processor per domain, speedup is determined by degree of communication (surface/volume ratio).



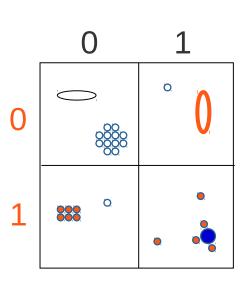
but... KMC is an inherently sequential method.

Take a simulation box with a number of objects and a simple spatial domain composition:

In each domain (00,01,10,11) the time step depends on the characteristics of the simulated event.

e.g: A single vacancy jump differs from an interstitial loop recombination.

Thus, the time advance in each region is asynchronous. When objects cross into neighbouring domains, the times have to match.



$$\Delta t_{00} \neq \Delta t_{01} \neq \Delta t_{10} \neq \Delta t_{11}$$



#### Some ways around the problem:

#### **Rigorous Algorithms**

Conservative asynchronous algorithm

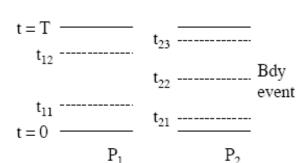
Lubachevsky (1988), Korniss et al (1999), Shim & Amar (2004)

• Synchronous relaxation algorithm

Lubachevsky & Weiss (2001), Shim & Amar (2004)







One Cycle

#### Synchronous relaxation (SR) algorithm

(Lubachevsky & Weiss, 2001)

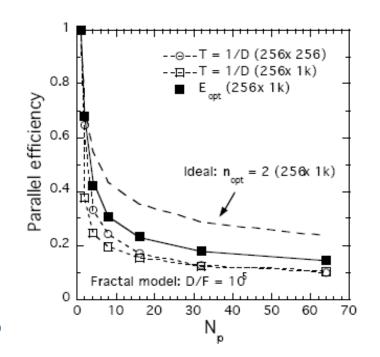
- All processors 'in-synch' at beginning & end of each cycle
- Iterative relaxation at each iteration processors use boundary info. from previous iteration
- Relaxation complete when current iteration identical to previous iteration for all processors

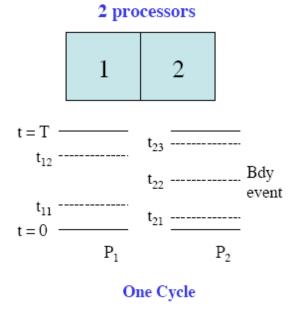


Thin film growth

#### Disadvantages:

- Complex: requires 'keeping list' of all events, random numbers used in each iteration
- Algorithm does not scale: faster than CA algorithm but still slow due to global synchronization and requirement of multiple iterations per cycle







## **Hybrid KMC**

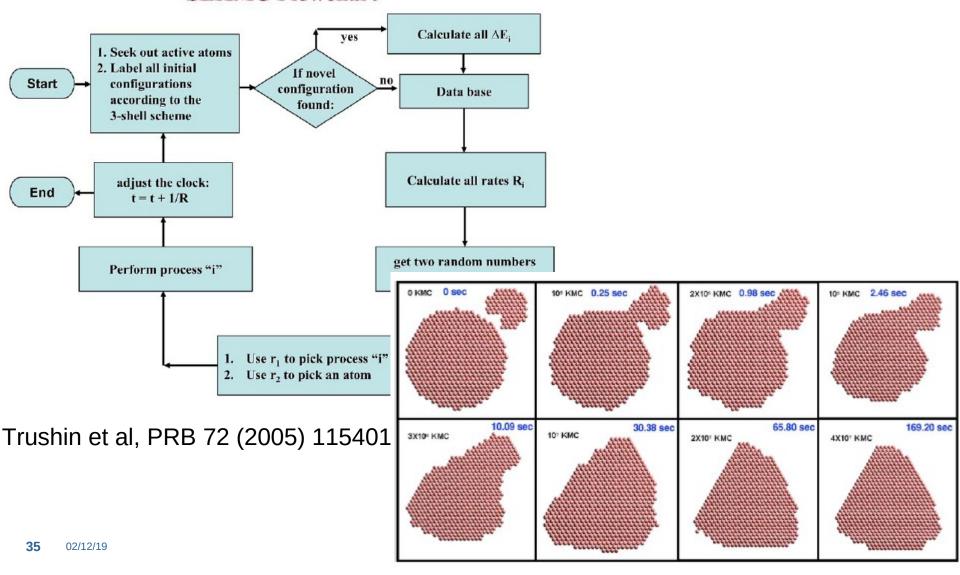
Concurrent multi-scale models (See presentation by P. Geysermans)

## Surface diffusion Continuum Blending Zone Atomistic (KMC) $j_{N-2}$ $j_{N-1}$ $J_2$ Gill, Spencer and Cocks, Materials Science and Engineering A365 (2004) 66–72 Region of crystal strain no (weak) crystal strain



## **Self-learning KMC**

#### **SLKMC Flowchart**



#### **Outline**

#### 1. Introduction

- Origin of Monte-Carlo methods
- ☐ Time for Kinetic Monte-Carlo

#### 2. Kinetic Monte-Carlo method

- Physical motivation
- Mathematical formalism
- Numerical recipie
- Range of applicability
- Advanced methods

#### 3. Applications

- ☐ Atomistic KMC thermal ageing & microstructure evolution
- □ Object KMC resistivity recovery



#### **Atomistic KMC**

- Rigid Lattice
- Atoms
- Vacancy (and interstitial) diffusion
- Recombination
- Binding energies



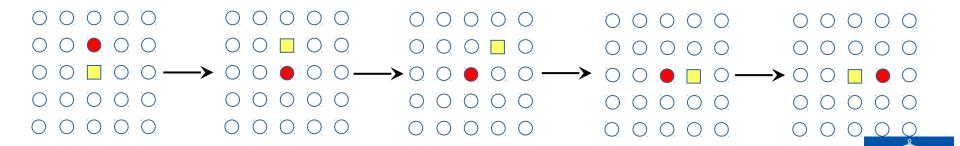
#### **Atomistic KMC**

Matrix + solute elements + vacancies + interstitials

Diffusion of vacancies and interstitials on a fixed lattice.

Vacancies: rigid lattice approximation, migrate to nearest neighbour.

Interstitials: migration depending on lattice type.



#### **Atomistic KMC**

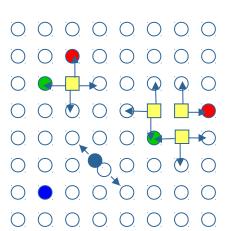
- BCC metal: treatment of multi-component systems:
  - substitutional elements

- vacancies
- interstitials



- Diffusion by first nearest neighbour jumps via:

vacancies interstitials



## Lattice diffusion:

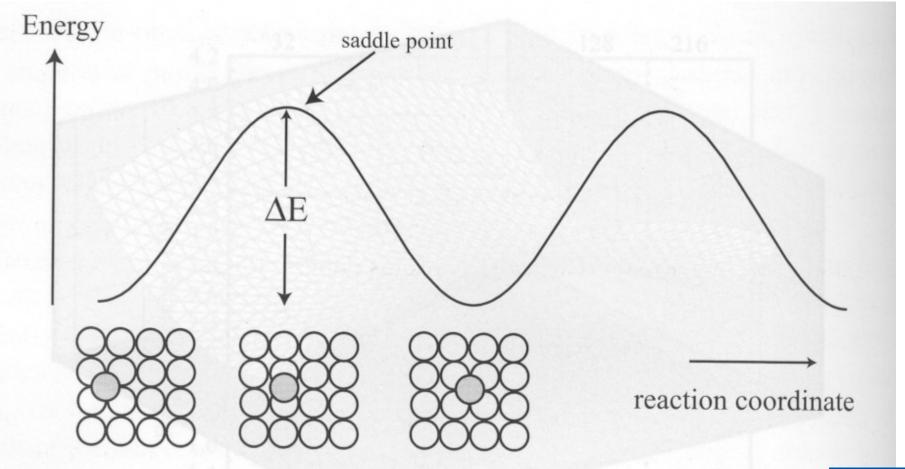
Jump frequency:

$$D = D_0 \exp(-\frac{E_a}{kT})$$

$$\Gamma = \nu \exp(-\frac{E_{\rm a}}{kT})$$
attempt frequency

activation energy



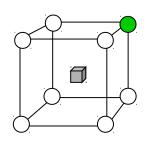


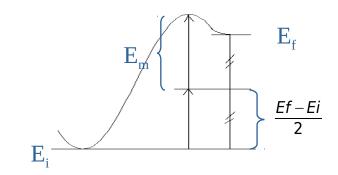


# How to get activation energy $E_a$ ?

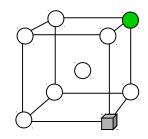
Should obey detailed balance:  $p_i r_{ii} = p_i r_{ii}$ 

Direct migration energy calculation (MD or ab initio)





$$E_{\rm a} = E_{\rm m} + \frac{E_{\rm f} - E_{\rm i}}{2}$$



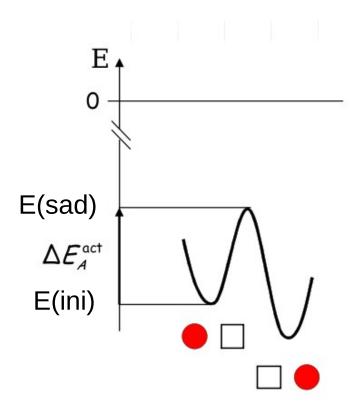


a)

# b)

Jump frequency of point defects

$$\Gamma_{A} = \mathbf{v}_{A} \exp \left\{ -\frac{\Delta E_{A}^{\text{act}}}{k_{b}T} \right\}$$
attempt frequency migration barrier
$$\Delta E_{a} = E(\text{sad}) - E(\text{ini})$$



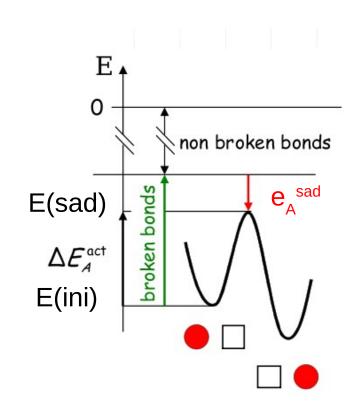
F. Soisson, A. Barbu and G. Martin, Acta Mater. 44 (1996) 3789.



# b)

Jump frequency of point defects

$$\Delta E_A^{act} = E(sad) - E(ini) = e_A^{sad} - \Sigma V_{Aj} - \Sigma V_{iv}$$
  
bond energy for A at saddle



F. Soisson, A. Barbu and G. Martin, Acta Mater. 44 (1996) 3789.

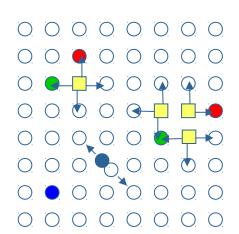


- BCC metal: treatment of multi-component systems:
  - substitutional elements

- vacancies
- interstitials



• Diffusion by first nearest neighbour jumps via: vacancies interstitials

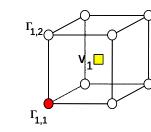


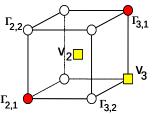
Jump probability 
$$\Gamma_x = \nu_x e^{-\frac{E_{\mathrm{a}}}{kT}}$$
  $\nu_{\mathrm{x}}$  is the attempt frequency

- Residence time algorithm
- Applied to vacancy and interstitial jumps

Average time step:

$$\Delta t = \frac{1}{\sum_{j,k} \Gamma_{jk}}$$



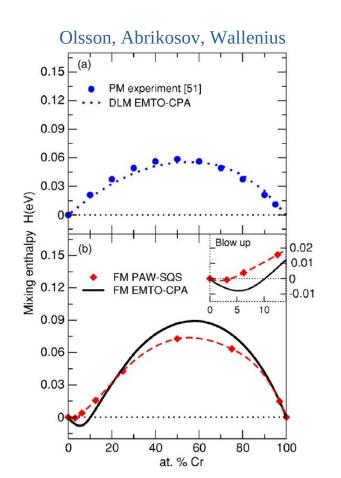


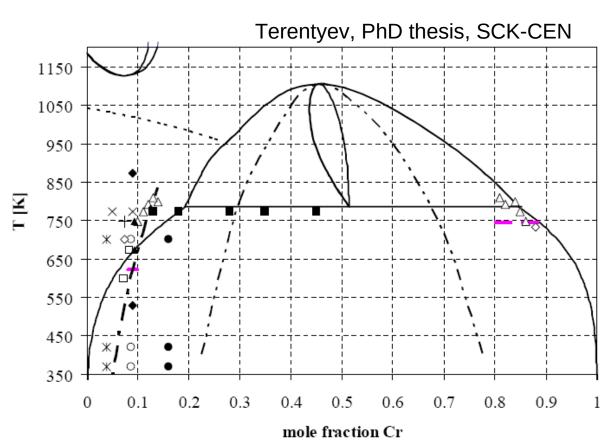




Fe-Cr alloys are very interesting for nuclear applications:

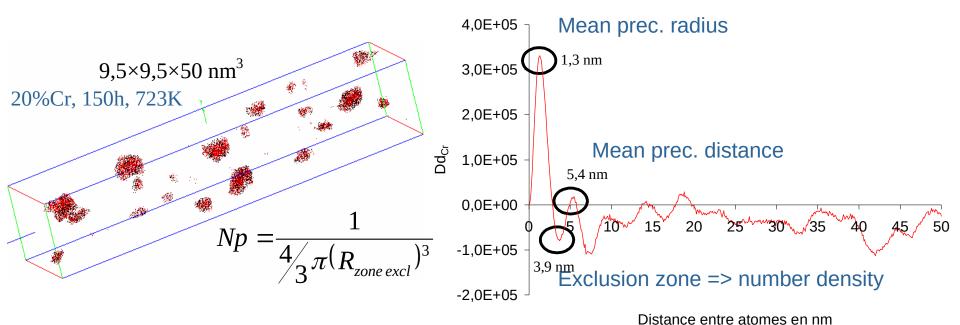
- Swelling resistance
- Low ductile-brittle transition temperature
- Creep resistance
- Corrosion resistance
- etc.





Tomographic Atom Probe (TAP) at Univ. Rouen: ageing study on model Fe-Cr alloys and real steels

$$\mathrm{Dd}_{\mathrm{Cr}} = \mathrm{d}_{\mathrm{Cr-Cr}} - \mathrm{d}_{\mathrm{X-Y}}$$





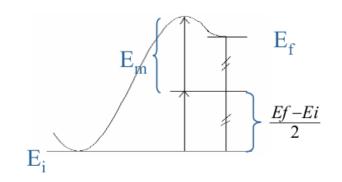
## Vacancy driven atomistic KMC:

only process is the single vacancy jump
only one vacancy (still supersaturated simulation cell)
Olsson et al PRB 72 (2005)
cohesion model: concentration dependent interatomic potential.

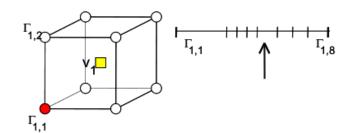
Residence time algorithm:

$$\Gamma_x = \nu_x e^{-\frac{E_a}{kT}}$$

$$E_{\rm a} = E_{\rm m} + \frac{E_{\rm f} - E_{\rm i}}{2}$$

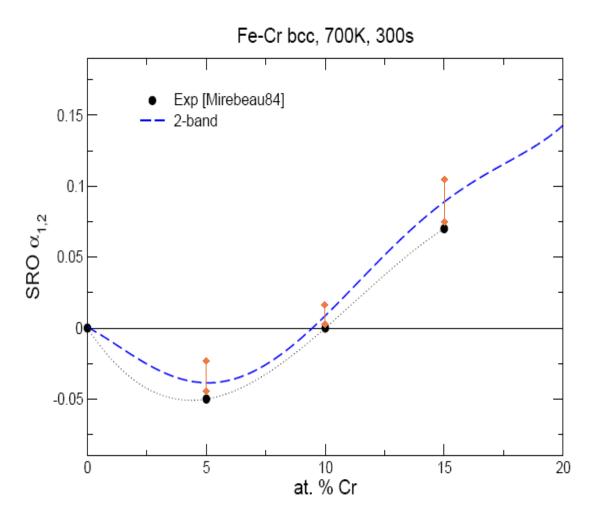


Average time step: 
$$\Delta t = \frac{1}{\sum_i \Gamma_i}$$





## Short range order



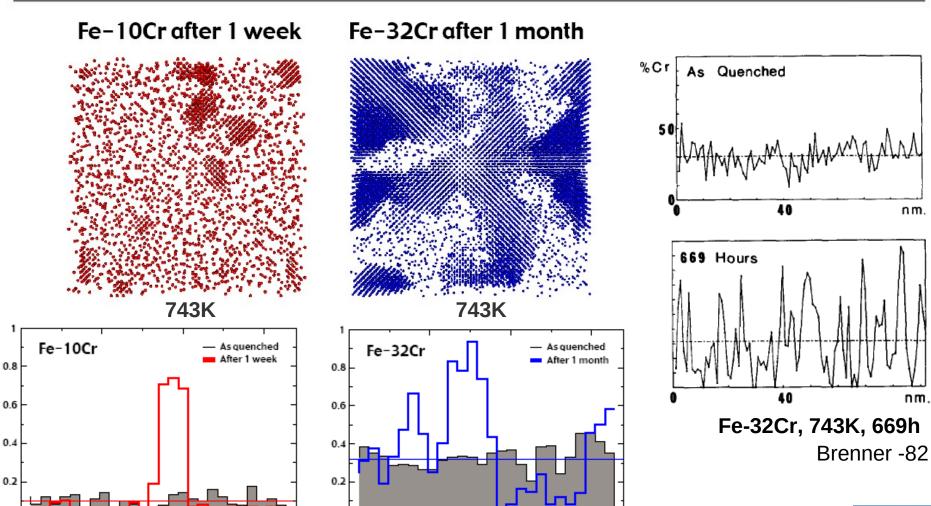
Due to supersaturation of vacancies, the simulated KMC time is not directly comparable to real time.

"Real" time from KMC time scaled to equilibrium vacancy concentration:

$$C_{\text{vac}}^{\text{eq}} = \exp(-\frac{G_{\text{f}}}{kT})$$

$$\Delta t = C_{\rm vac}^{\rm eq} \Delta t_{\rm KMC}$$





Distance along the <110 > direction [nm]



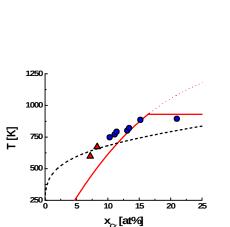
# Homogeneous precipitation in bulk Fe-Cr

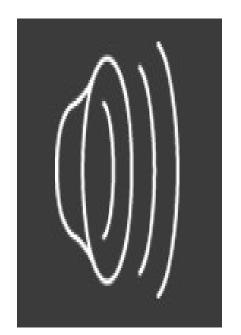
T = 750K ;  $x_{cr} = 0.15$ 



Cr precipitates when fluctuations become larger than a critical size

Precipitates are spherical











## Spinodal decomposition in bulk Fe-Cr

Homogeneous precipitation at  $x_{cr} \sim 0.5$ , within the spinodal

Spinodal decomposition happens by growth of small composition fluctuations

(diffusion against the gradient of x),

in regions of the phase diagram where  $d^2G/dx^2 < 0$ 

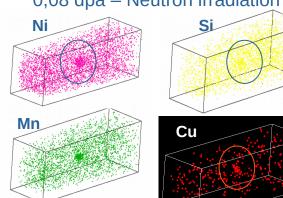
Cr precipitation in Fe.<sub>5</sub>Cr.<sub>5</sub> at 750K

2-D slice of a 3D sample

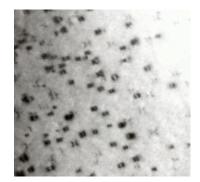




#### 0,08 dpa - Neutron irradiation



15x15x50 nm

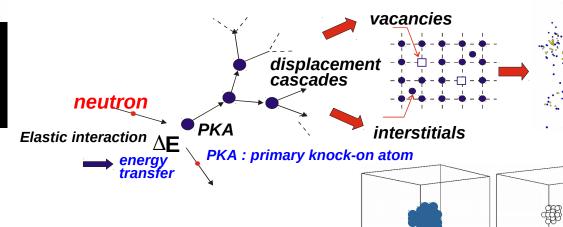


TEM, Barbu, CEA

#### **RPV** steels:

#### Matrix Fe

- + alloying elements: Cu, Ni, Mn, Si, ...
- + carbon, nitrogen
- + dislocations



TAP, Pareige, U. Rouen

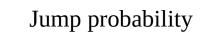
- Multi-component system treated by Atomistic KMC:
  - substitutional elements (Cu, Ni, Mn, Si)
  - interstitials (Fe, Cu, Ni, Mn, Si)

Cohesive model by interatomic potential not feasible!

 Diffusion by first nearest jumps via vacancies interstitials

 $\Gamma_X = v_X \exp\left(-\frac{Ea}{kT}\right)$ 

 $v_x$  attempt frequency

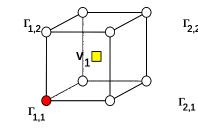


$$\Gamma_X = v_X \exp\left(-\frac{Ea}{kT}\right)$$

Residence time algorithm

Applied to vacancy and interstitial jumps

$$\Delta t = \frac{1}{\sum_{i,k} \Gamma_{jk}}$$



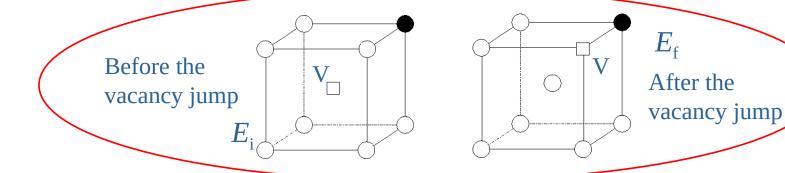




#### Environment dependent form for the activation energy

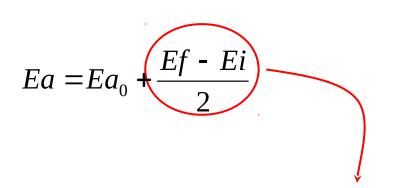
$$Ea = Ea_0 + \frac{Ef - Ei}{2}$$

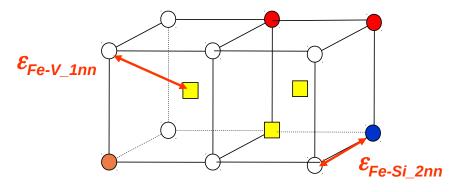
Ea<sub>0</sub> = ab initio migration energy via a vacancy of the migrating atom in an  $\alpha$ -Fe matrix



DFT data	Fe	Cu	Ni	Mn	Si
E <sub>migration</sub> (X) (eV)	0.62	0.54	0.68	1.03	0.42

#### Cohesive model based on pair interactions up to second nearest neighbours





$$E = \sum_{j} \varepsilon_{(Fe\text{-}Fe)}^{(i)} + \sum_{k} \varepsilon_{(V\text{-}V)}^{(i)} + \sum_{l} \varepsilon_{(Fe\text{-}V)}^{(i)} + \sum_{m} \varepsilon_{(Fe\text{-}X)}^{(i)} + \sum_{n} \varepsilon_{(V\text{-}X)}^{(i)} + \sum_{p} \varepsilon_{(X\text{-}Y)}^{(i)}$$

i = 1 or 2, corresponds to first or second nearest neighbour interaction

j = the number of Fe-Fe bonds

m = the number of Fe-X bonds

k =the number of V-V bonds

n = the number of V-X bonds

*l* = the number of Fe-V bonds

p = the number of X-Y bonds

X, Y = solute atoms (Cu, Ni, Mn or Si)



#### **Binary alloys**

$$E_{mixing} = -4\varepsilon_{(Fe-Fe)}^{(1)} - 3\varepsilon_{(Fe-Fe)}^{(2)} + 8\varepsilon_{(Fe-X)}^{(1)} + 6\varepsilon_{(Fe-X)}^{(2)} - 4\varepsilon_{(X-X)}^{(1)} - 3\varepsilon_{(X-X)}^{(2)}$$

$$E_{\text{interface}(100)} = -2\varepsilon_{(Fe-Fe)}^{(1)} - \varepsilon_{(Fe-Fe)}^{(2)} + 4\varepsilon_{(Fe-X)}^{(1)} + 2\varepsilon_{(Fe-X)}^{(2)} - 2\varepsilon_{(X-X)}^{(1)} - \varepsilon_{(X-X)}^{(2)}$$

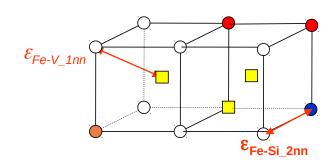
$$E_{coh}(Z) = 4\varepsilon_{(Z-Z)}^{(1)} + 3\varepsilon_{(Z-Z)}^{(2)}$$

$$Z = Fe$$
 or solute atom

$$E_{formation}(V^Z) = 8\varepsilon_{(V-Z)}^{(1)} + 6\varepsilon_{(V-Z)}^{(2)} - 4\varepsilon_{(Z-Z)}^{(1)} - 3\varepsilon_{(Z-Z)}^{(2)}$$

$$E_{binding(V-V)}^{(i)} = 2\varepsilon_{(Fe-V)}^{(i)} - \varepsilon_{(Fe-Fe)}^{(i)} - \varepsilon_{(V-V)}^{(i)}$$

$$E_{binding(V-X)}^{(1)} = \varepsilon_{(Fe-V)}^{(1)} + \varepsilon_{(Fe-X)}^{(1)} - \varepsilon_{(Fe-Fe)}^{(1)} - \varepsilon_{(V-X)}^{(1)}$$



#### **Ternary alloys**

$$E_{\textit{binding}(X-Y)}^{(i)} = \varepsilon_{(\textit{Fe-}X)}^{(i)} + \varepsilon_{(\textit{Fe-}Y)}^{(i)} - \varepsilon_{(\textit{Fe-}Fe)}^{(i)} - \varepsilon_{(X-Y)}^{(i)}$$

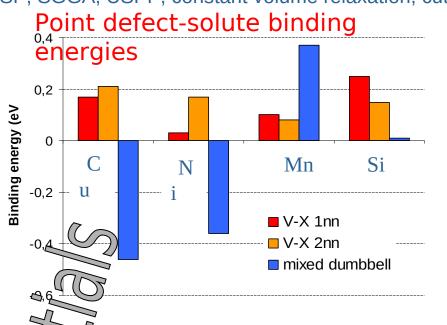
i = 1 or 2

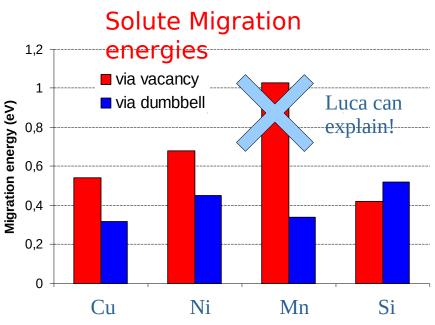
X, Y =solute atoms

Ab initio data

**Parameters** 

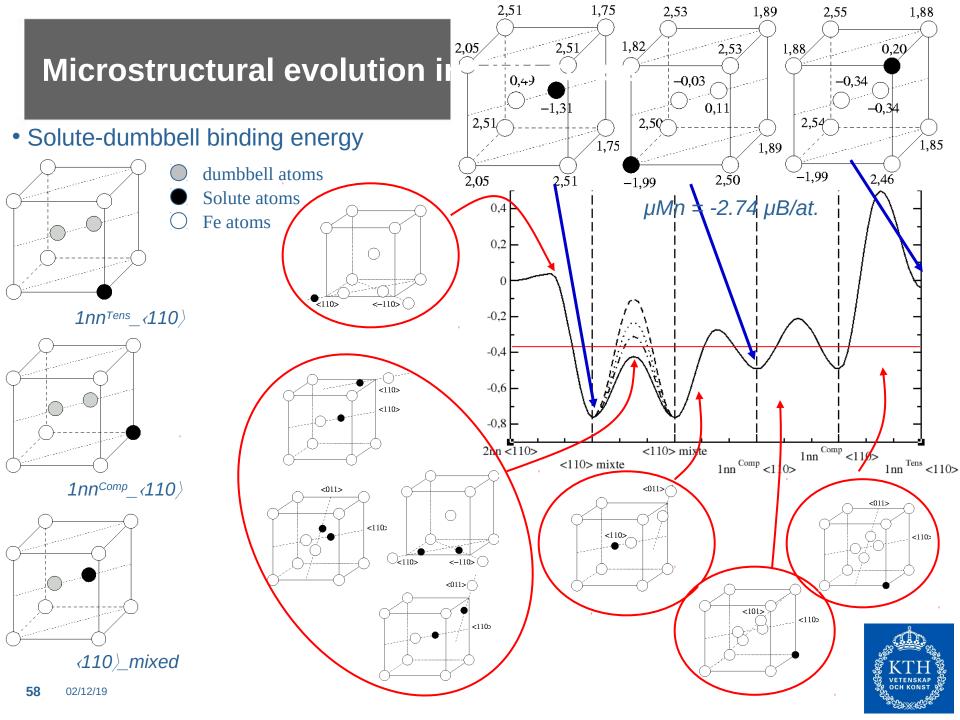
VASP, SGGA, USPP, constant volume relaxation, cut off: 240 eV, 54 or 128 atoms with 125 or 27 k points





- Mn high mixed dumbbell binding energy
  - low migration energy via dumbbells

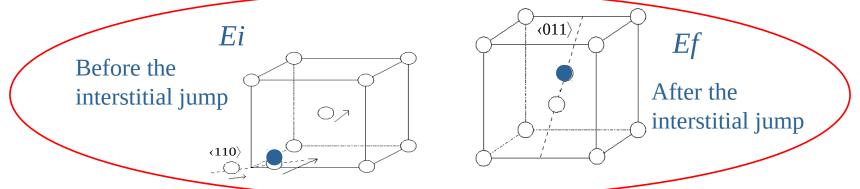
Contrary to the other solutes,
Mn may diffuse preferentially via an interstitial mechanism



# Environment dependent form for the activation energy En - ab initio migration 609

$$Ea = Ea_0 + \frac{Ef - Ei}{2}$$

Ea<sub>0</sub> = ab initio migration-60° rotation energy of the migrating atom in an  $\alpha$ -Fe matrix



	Fe-Fe	Fe-Cu	Fe-Ni	Fe-Mn	Fe-Si
	dumbbell	dumbbell	dumbbell	dumbbell	dumbbell
Migration energy (eV)	0.37	0.32	0.45	0.34	0.52



#### **Direct calculation**

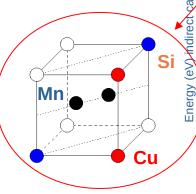
Complex binding energies

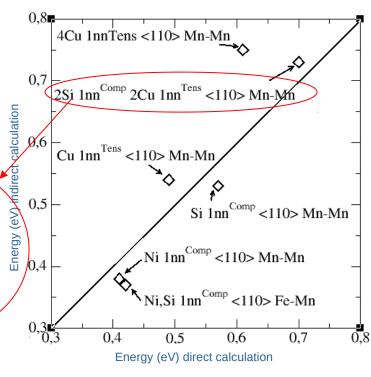
(1 dumbbell + several solute atoms)

$$Ea = Ea_0 + \underbrace{\frac{Ef - Ei}{2}}$$

#### **Indirect calculation**

Sum of elementary binding energies of the complex





#### Complex binding energy addition:

• solute - dumbbell

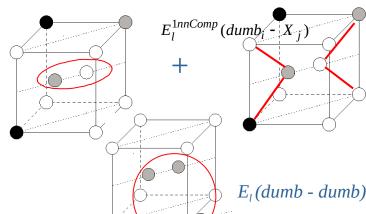
Solute

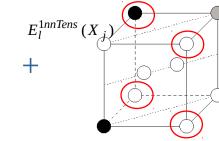
Solute atoms

○ Fe atom

• dumbbell - dumbbell

 $E_l^{mixed}(X_i - X_k)$ 







#### "NEUTRON" IRRADIATION of FeCunimnsi Alloy

Fe-0.2Cu-0.53Ni-1.26Mn-0.63Si (%at.) at 300°C

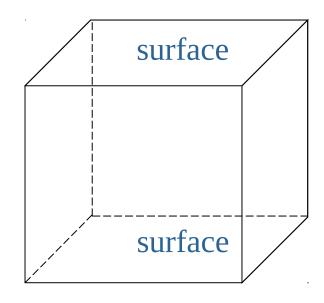
Flux: 6.5 10<sup>-4</sup> dpa.s<sup>-1</sup>, Dose: 2 10<sup>-3</sup> dpa

#### For neutron irradiation:

 flux of 20 keV and 100 keV cascades debris obtained by Molecular Dynamics (R. Stoller, J. Nucl. Mater. 307-311 (2002) 935)

Frenkel pairs

(For electron irradiation: Frenkel pair flux)

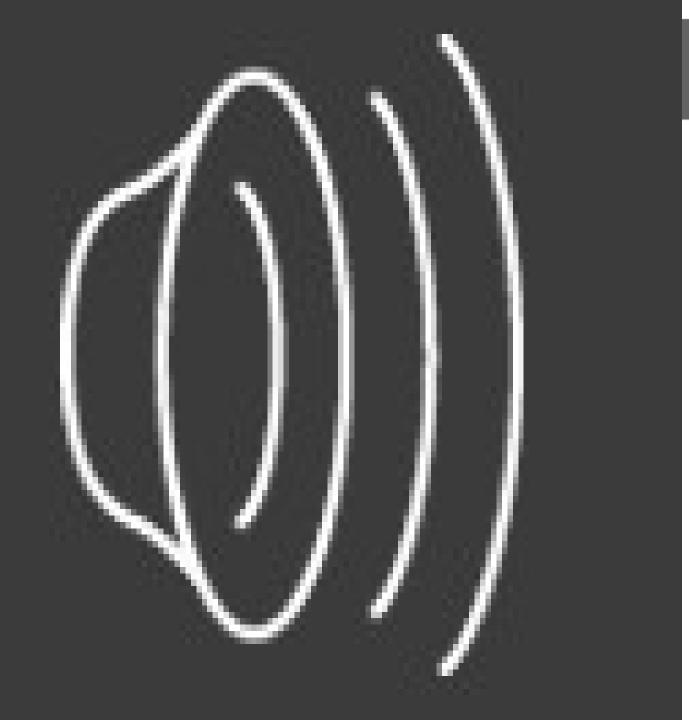


Surfaces along z direction  $100a_0 \times 100a_0 \times 100a_0$  $\sim (30 \text{ nm})^3$ 

- Mn

- c SIA





- $\bullet$ Cu
- Si
- Mn
- Ni
- $\circ$  V
- O SIA



#### Ni effect under electron irradiation

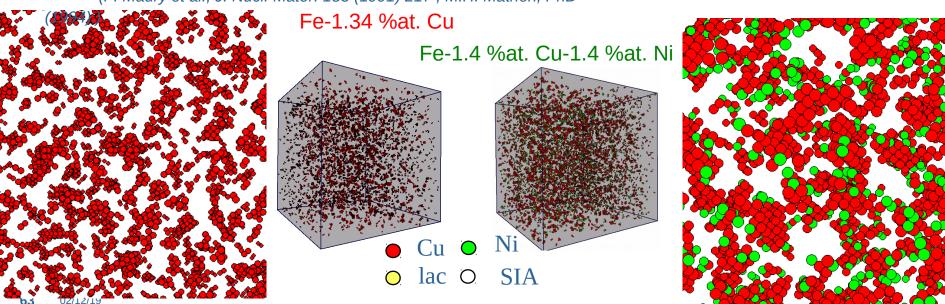
Flux: 5.10-8 dpa.s-1; dose: 1-1.2.10-3 dpa; T: 300 °C

#### Ni effect:

- Similar cluster density
- No modifications of Cu precipitation

#### Good agreement with exp. results

(F. Maury et al., J. Nucl. Mater. 183 (1991) 217; M.H. Mathon, PhD



 $100a_0 \times 100a_0 \times 100a_0$  Surfaces along z axis

# clusters

FeCu

Cluster size (number Cu)

FeCuNi

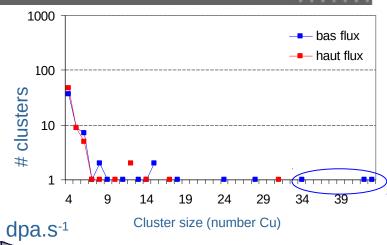
#### Flux effect under neutron irradiation

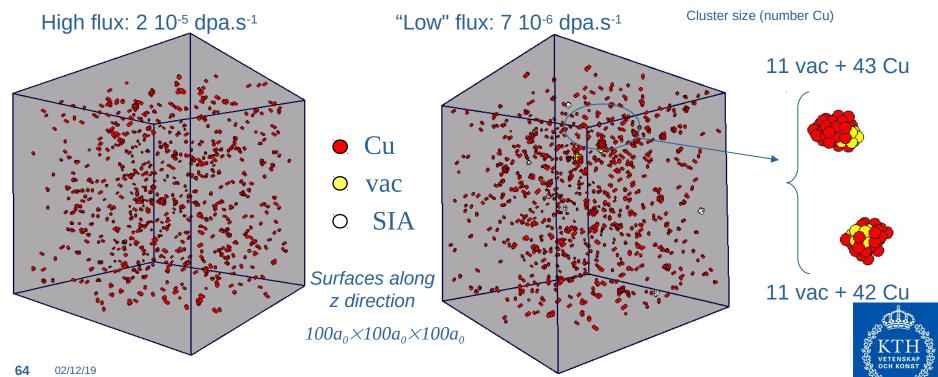
Fe-0.79 %at. Cu at 300°C Dose: 5 10-4 dpa

Bigger Cu and vacancy-solute complex clusters with the lower flux

#### Good agreement with exp. results

(S. Yanagita et al., ASTM STP 1366 (2000) 516; R. Kasada et al., ASTM STP 1405 (2001) 237.)





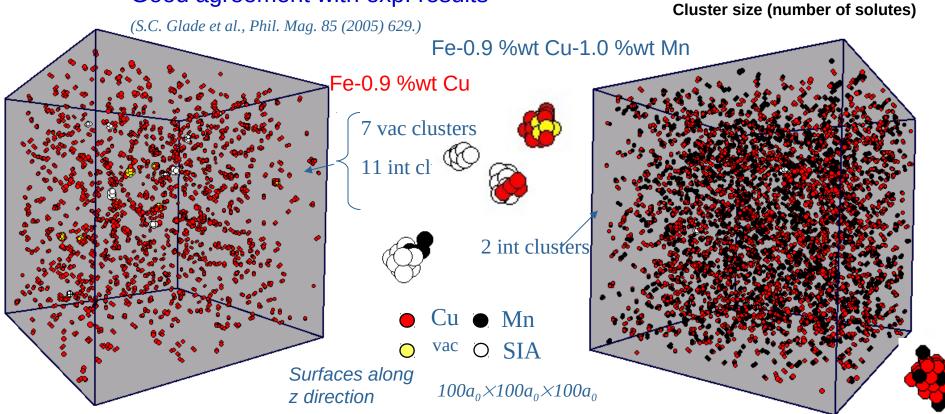
#### Mn effect under neutron irradiation

#### Flux: 0.5-1.0 10<sup>-5</sup> dpa.s<sup>-1</sup> Dose: 1.6-1.7 10<sup>-3</sup> dpa

#### Mn effect:

- higher cluster density, seemingly smaller clusters
- no vacancy-solute clusters

#### Good agreement with exp. results



1000

100

10

- FeCu - FeCuMn

## Objects:

- Vacancies (isolated, clustered, loops)
- Interstitials (isolated, clustered, loops)
- Solute atoms
- Traps, wells (impurities, dislocations,...)

#### Internal events:

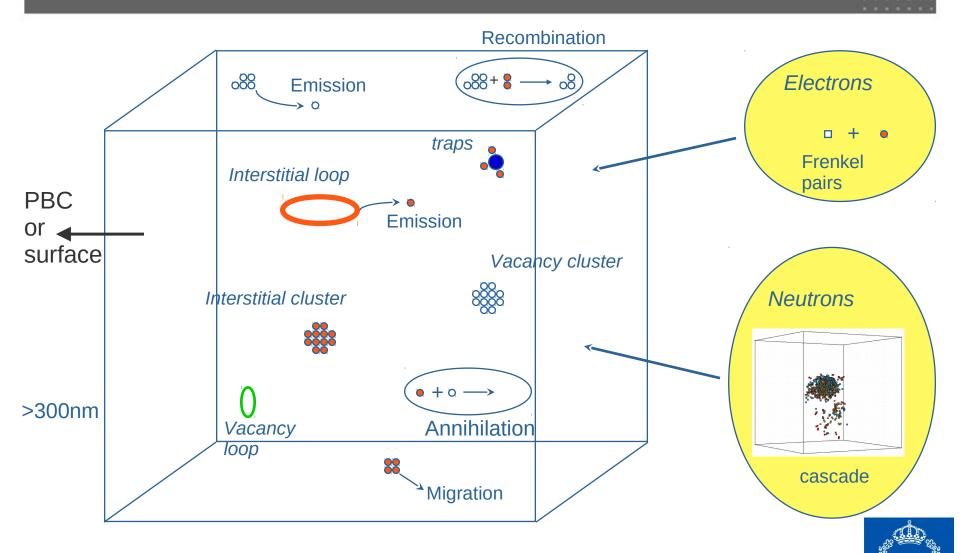
- Diffusion
- Dissociation or recombination

## External events (flux):

Arrival of neutron (or ion or electron)

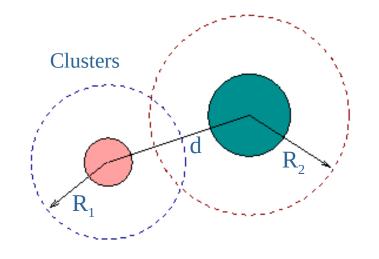
Follow the evolution of radiation damage: diffusion, dissociation, trapping, ...

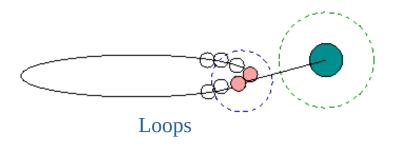




## Interaction between two objects:







$$v+i \underset{n}{\longleftrightarrow} i_{n-1}$$





$$v + i \rightarrow 0$$

$$\mathbf{i} + \mathbf{i}_{n} \rightarrow \mathbf{i}_{n+1}$$



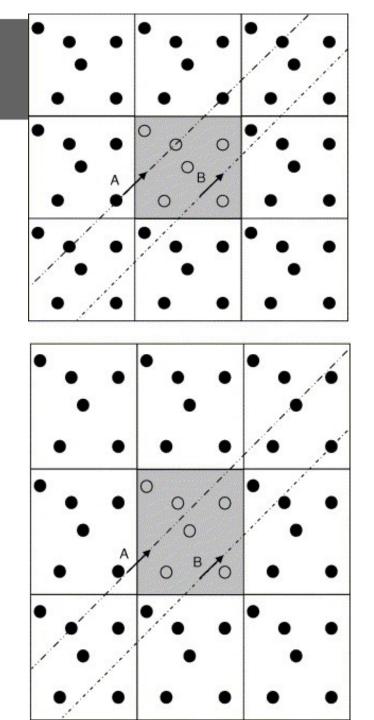
$$\mathbf{v} + \mathbf{v}_{\mathbf{n}} \longleftrightarrow \mathbf{v}_{\mathbf{n}+1}$$





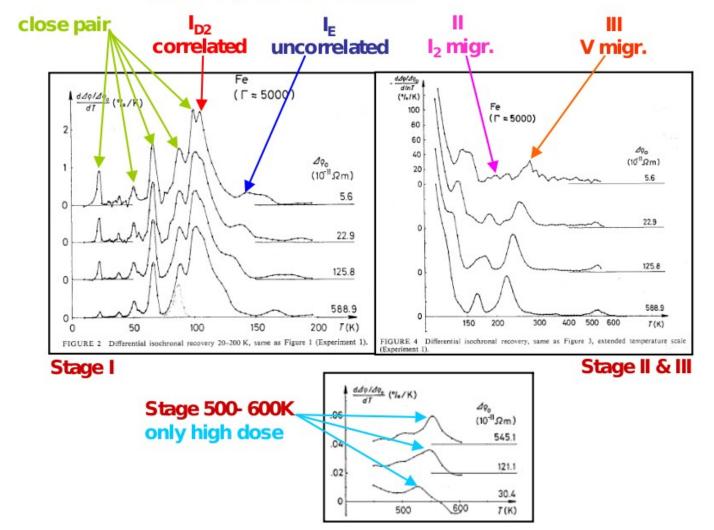
The choice of simulation box can significantly affect the interaction patterns.

In non-cubic boxes the interaction distance can overcome the usual cubic limitations (max one box length or infinite).



#### **Evolution of microstructure after irradiation**

Recovery of pure iron after irradiation with 3 MeV electron Resistivity measurements, Takaki et al. 1983





### **Evolution of microstructure after irradiation**

Interstitial

migration

Correlated

pairs

Recovery of pure iron after irradiation with 3 MeV electron

Modeling : EKMC (JERK) coupled with Ab Initio calculations

J. Dalla Torre, Chu Chun Fu, F. Willaime 2003, Nature Mater. 0,01 Resistivity **Experimental** Ш Ш ŀ  $I_{D2}$ peaks 0,005 107.5K 144K 185K 278K Calculated 170 230 250 110 130 150 190 210 270 290 Defect concentration 10000 1000 100 10 V2 130 170 90 **1**10 150 190 210 230 250 270 290 T (K)

Di-interstitial

migration



Vacancy

migration

#### **Evolution of microstructure after irradiation**

## Recovery of pure iron after irradiation with 3 MeV electron Conclusions - Coupling Ab Initio-EKMC (Jerk)

#### Monte Carlo simulations in excellent agreement with measurements

- temperature peaks reproduced within 10 K
- dose effects as well :
  - ✓ IE, II, III stages shift towards lower temperatures
  - √ 500-600K stage appears at high doses only

#### Identification/ validation of associated mechanisms

- confirms identification of recovery stages
  - ✓ stage 500-600K: associated with vacancy clusters dissociation
  - ✓ stage III: migration of di-vacancies improves agreement / experiment
- confirms E<sub>f</sub> calculations for vacancies :
  - ✓ ab initio values agree with high experimental values (2.1–2.4 eV);
    lower experimental values (1.6 eV) are incompatible: due to C-V binding E



### Conclusions

We have described the basics of the Kinetic Monte-Carlo method, its implementation, difficulties and potential.

Building the rate catalouge!

#### Three application were discussed:

Simlations of thermal ageing of Fe-Cr alloys.

Nanostructure evolution in RPV steels.

Isochronal annealing of iron.

