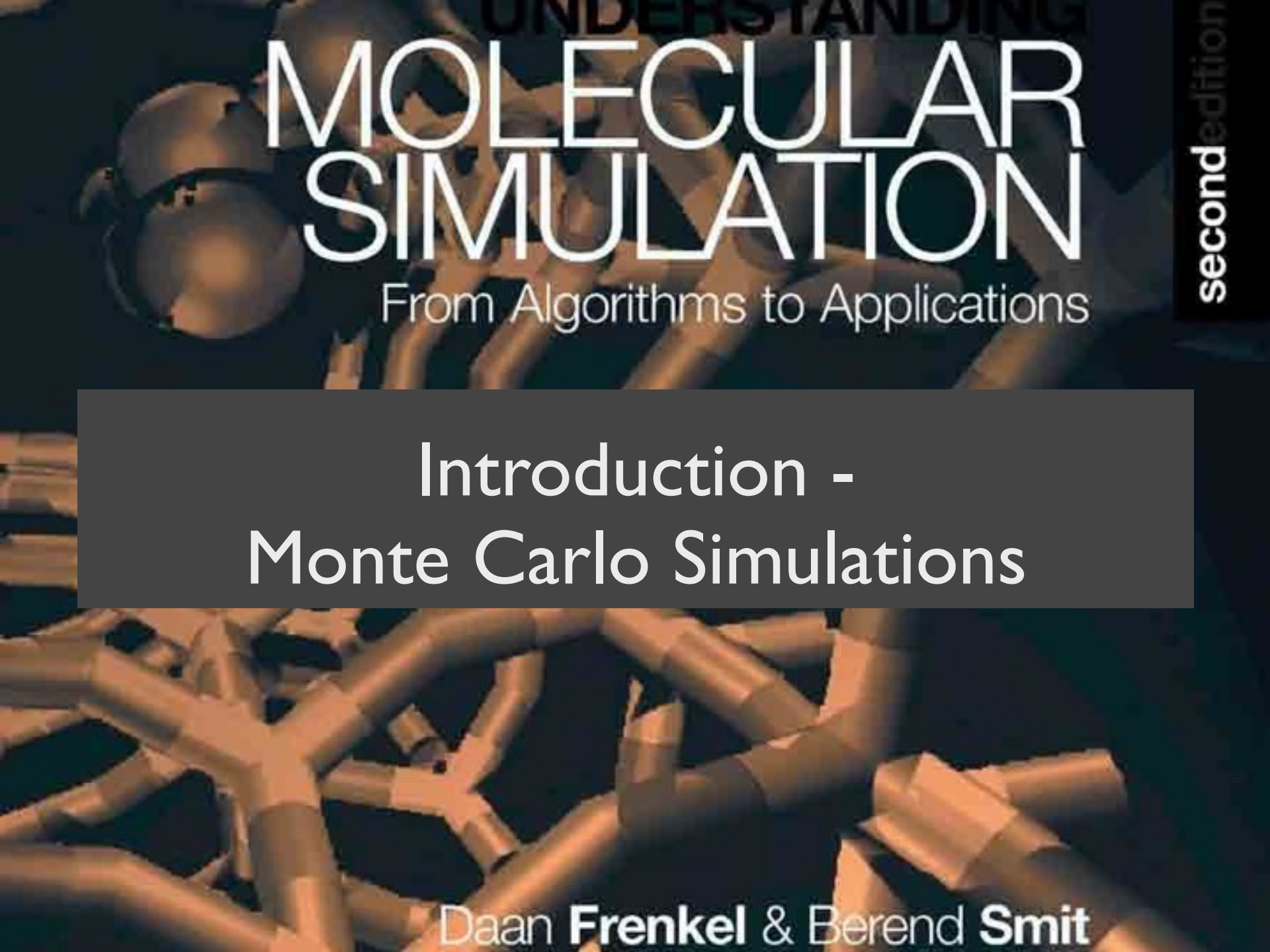


The background of the slide is a 3D molecular simulation. It features a complex network of orange and yellow rods and spheres, representing atoms and bonds in a molecular structure. The lighting is dramatic, with strong highlights and shadows, giving it a sense of depth and three-dimensionality. The overall color palette is dominated by warm tones like orange, yellow, and brown, set against a dark background.

MOLECULAR SIMULATION

From Algorithms to Applications

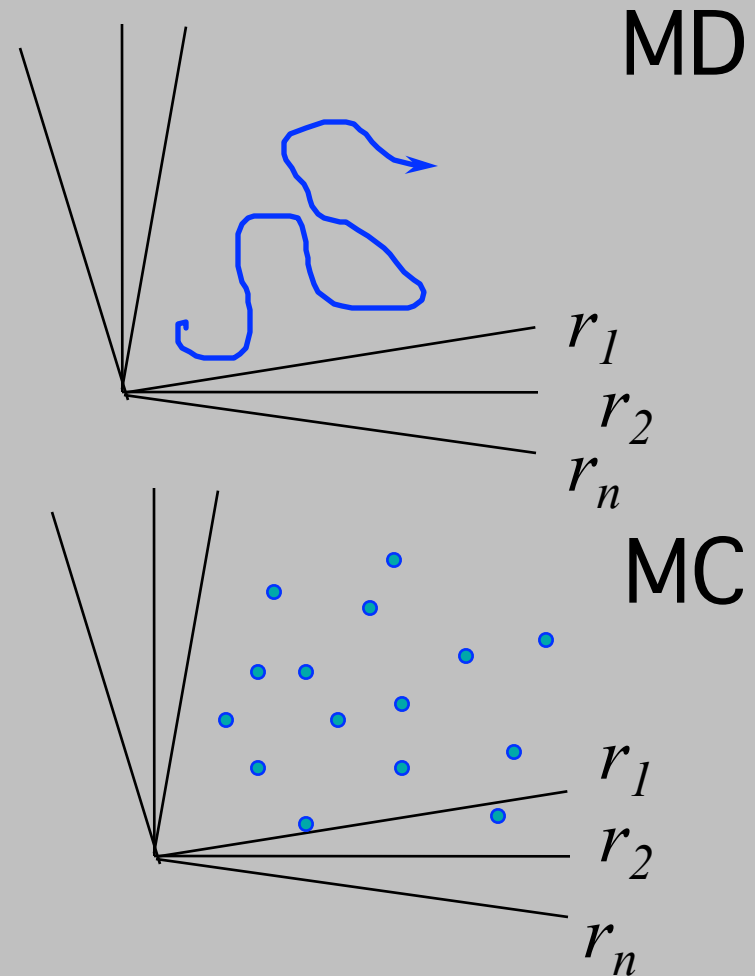
second edition

Introduction - Monte Carlo Simulations

Daan **Frenkel** & Berend **Smit**

Molecular Simulations

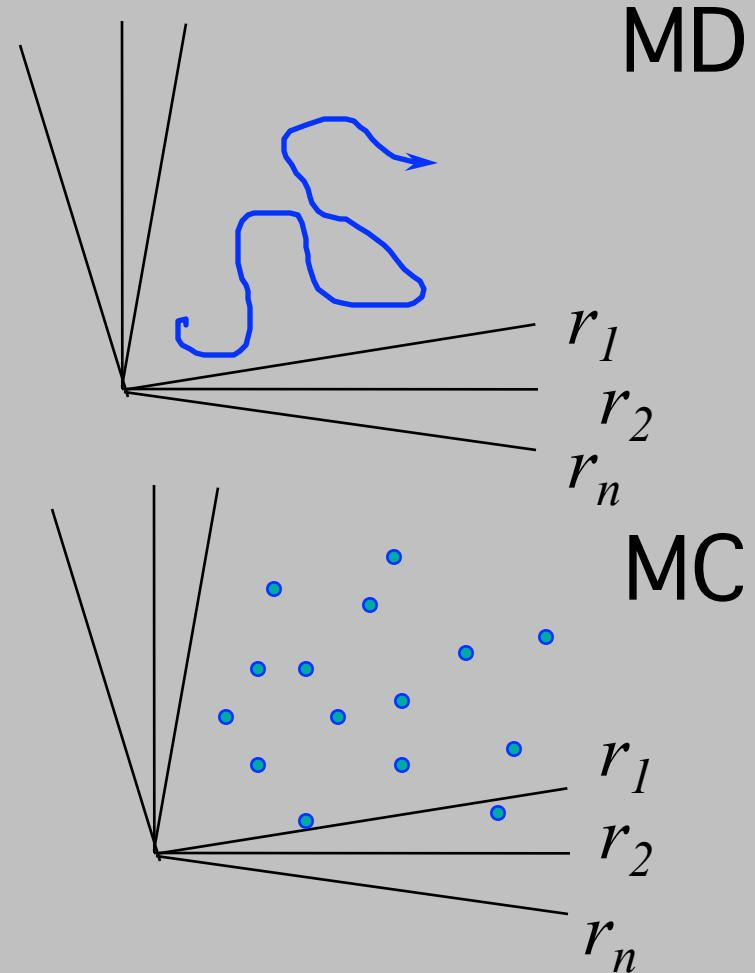
- ◆ Molecular dynamics:
solve equations of motion
- ◆ Monte Carlo:
importance sampling

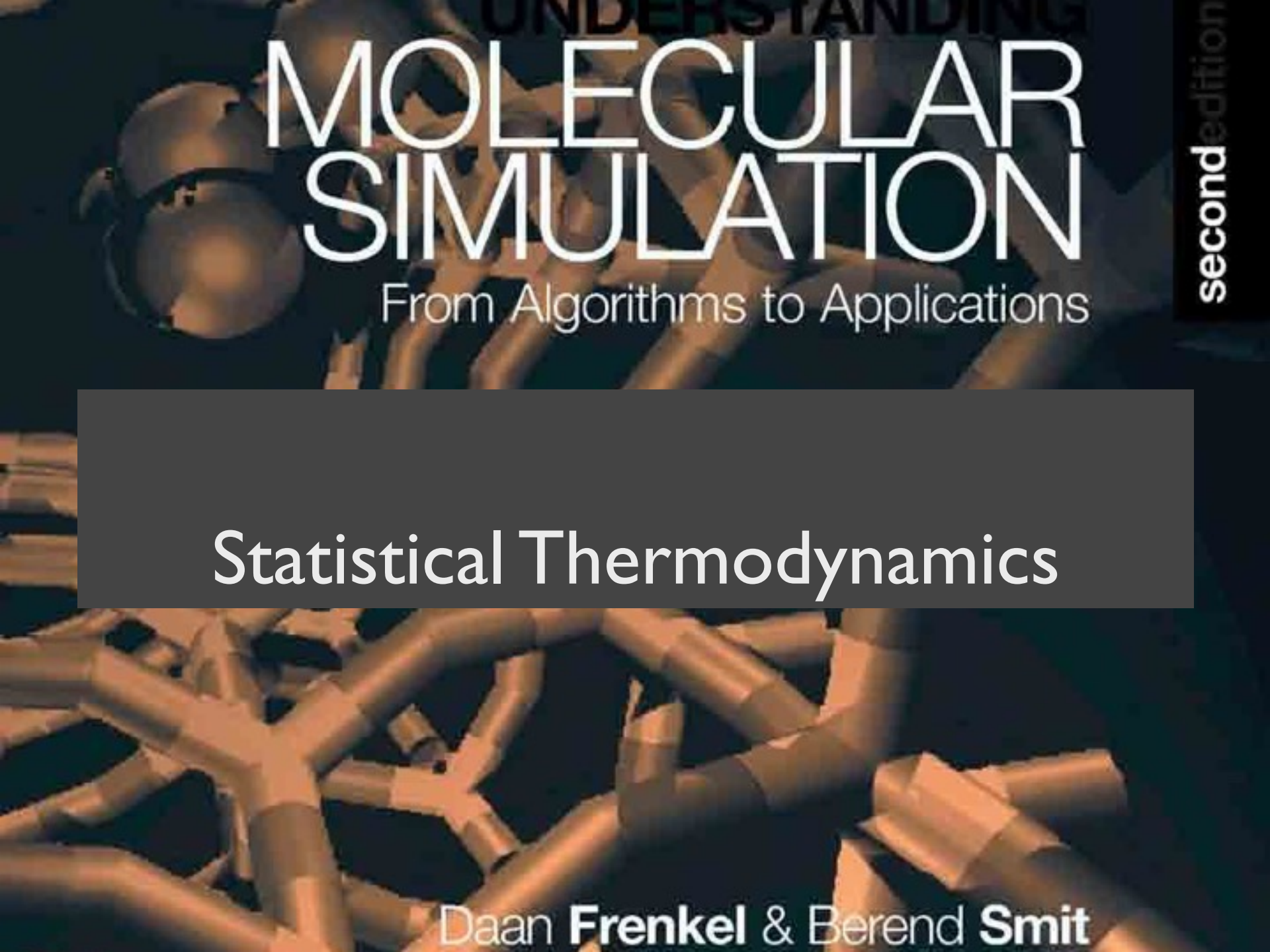


Molecular Simulations

- ◆ Molecular dynamics:
solve equations of
motion

- ◆ Monte Carlo:
importance sampling



The background of the slide is a 3D molecular simulation. It features a complex network of orange and yellow rods representing atoms and bonds, set against a dark blue background. In the upper left, there is a cluster of spheres, some colored red and others blue, representing different types of atoms. The overall image conveys a sense of scientific research and computational chemistry.

MOLECULAR SIMULATION

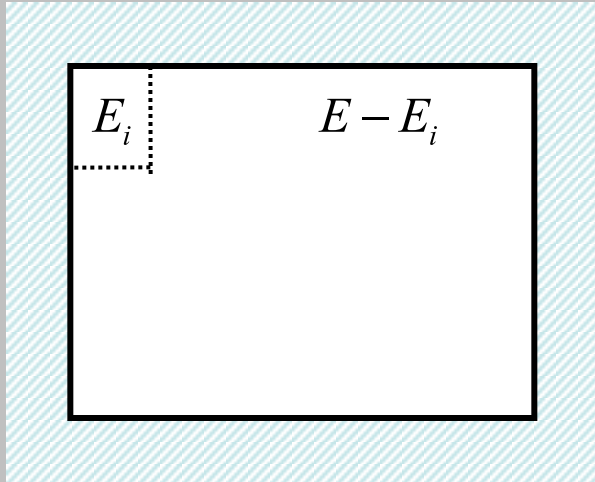
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Statistical Thermodynamics

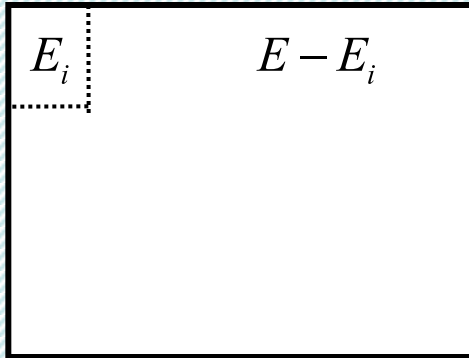
Daan **Frenkel** & Berend **Smit**

Canonical ensemble



Consider a small system that can exchange heat with a big reservoir

Canonical ensemble

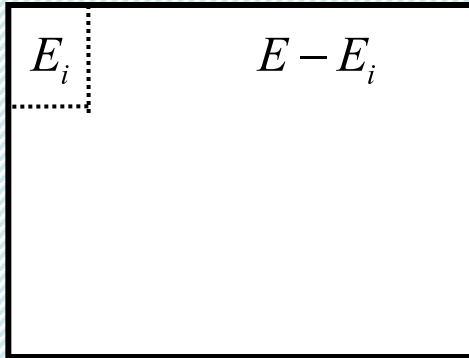


Consider a small system that can exchange heat with a big reservoir

$$\ln \Omega(E - E_i) = \ln \Omega(E) - \frac{\partial \ln \Omega}{\partial E} E_i + \dots$$

Canonical ensemble

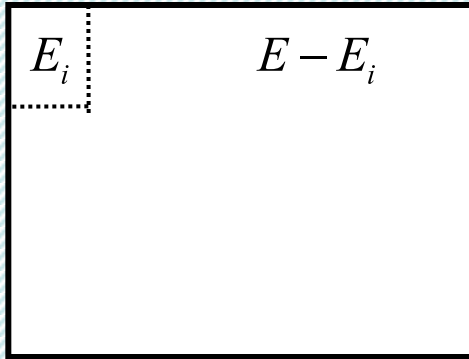
$$1/k_B T$$



Consider a small system that can exchange heat with a big reservoir

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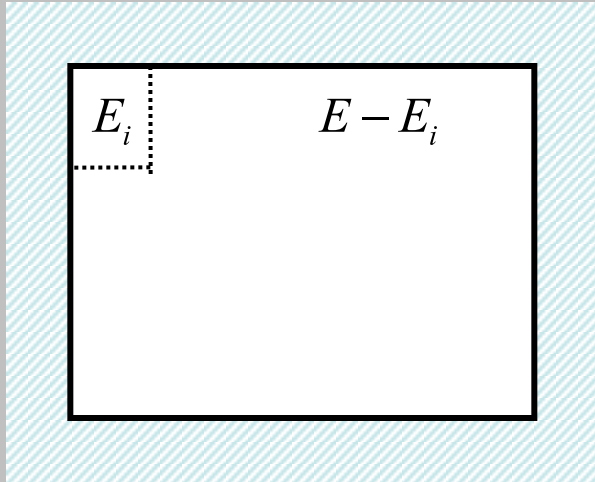
Canonical ensemble



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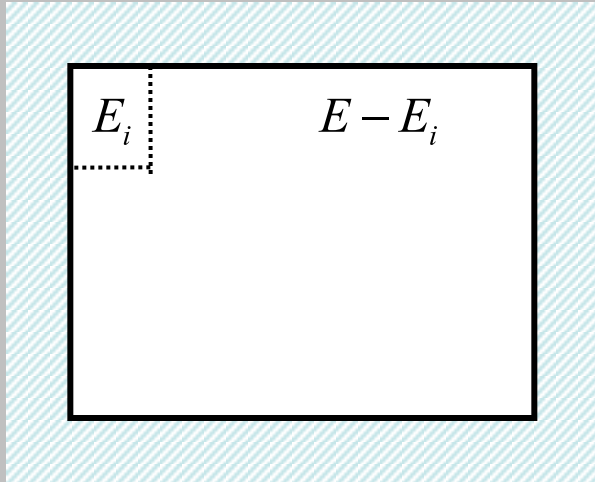


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$$\ln \Omega(E - E_i) = \ln \Omega(E) - \frac{\partial \ln \Omega}{\partial E} E_i + \dots$$

$$\ln \frac{\Omega(E - E_i)}{\Omega(E)} = -\frac{E_i}{k_B T}$$

Canonical ensemble



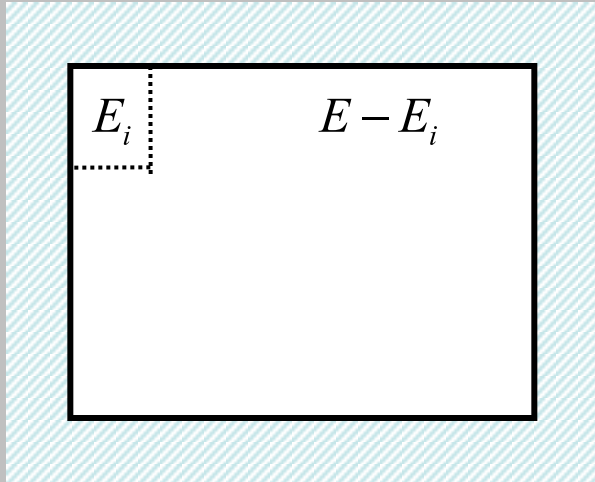
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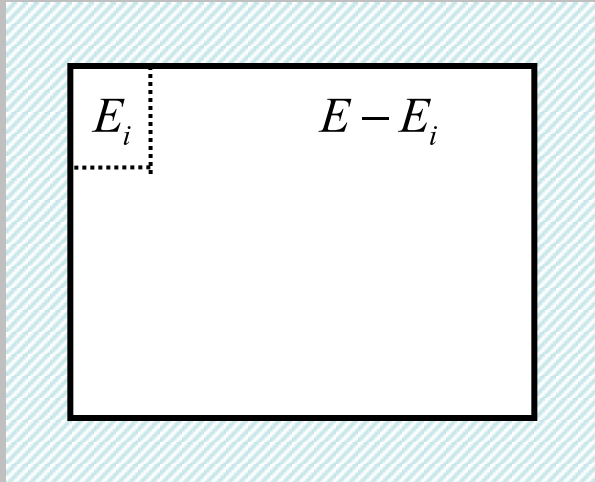
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$$P(E_i) = \frac{\Omega(E - E_i)}{\sum_j \Omega(E - E_j)} = \frac{\exp(-E_i/k_B T)}{\sum_j \exp(-E_j/k_B T)}$$

Canonical ensemble



Consider a small system that can exchange heat with a big reservoir

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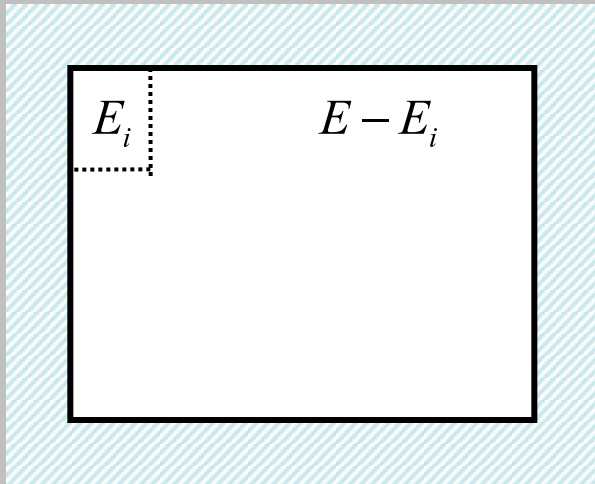
$$\ln \frac{\Omega(E - E_i)}{\Omega(E)} = -\frac{E_i}{k_B T}$$

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$$P(E_i) \propto \exp(-E_i/k_B T)$$

Canonical ensemble



Consider a small system that can exchange heat with a big reservoir

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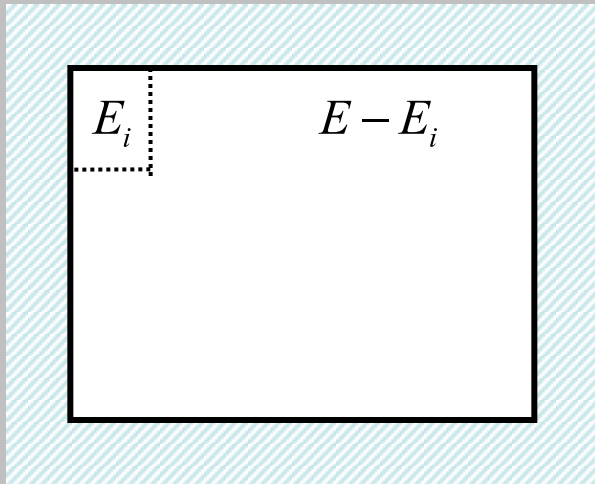
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Boltzmann distribution

Canonical ensemble



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Boltzmann distribution

Summary: Canonical ensemble (N, V, T)

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Partition function:

$$Q(N, V, T) = \frac{1}{\Lambda^{3N} N!} \int d\mathbf{r}^N \exp[-\beta U(\mathbf{r}^N)]$$

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Probability to find a particular configuration:

$$P(\Gamma) \propto \exp[-\beta U(\Gamma)]$$

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Ensemble average:

$$\langle A \rangle_{NVT} = \frac{1}{Q_{NVT}} \frac{1}{\Lambda^{3N} N!} \int d\mathbf{r}^N A(\mathbf{r}^N) \exp[-\beta U(\mathbf{r}^N)]$$

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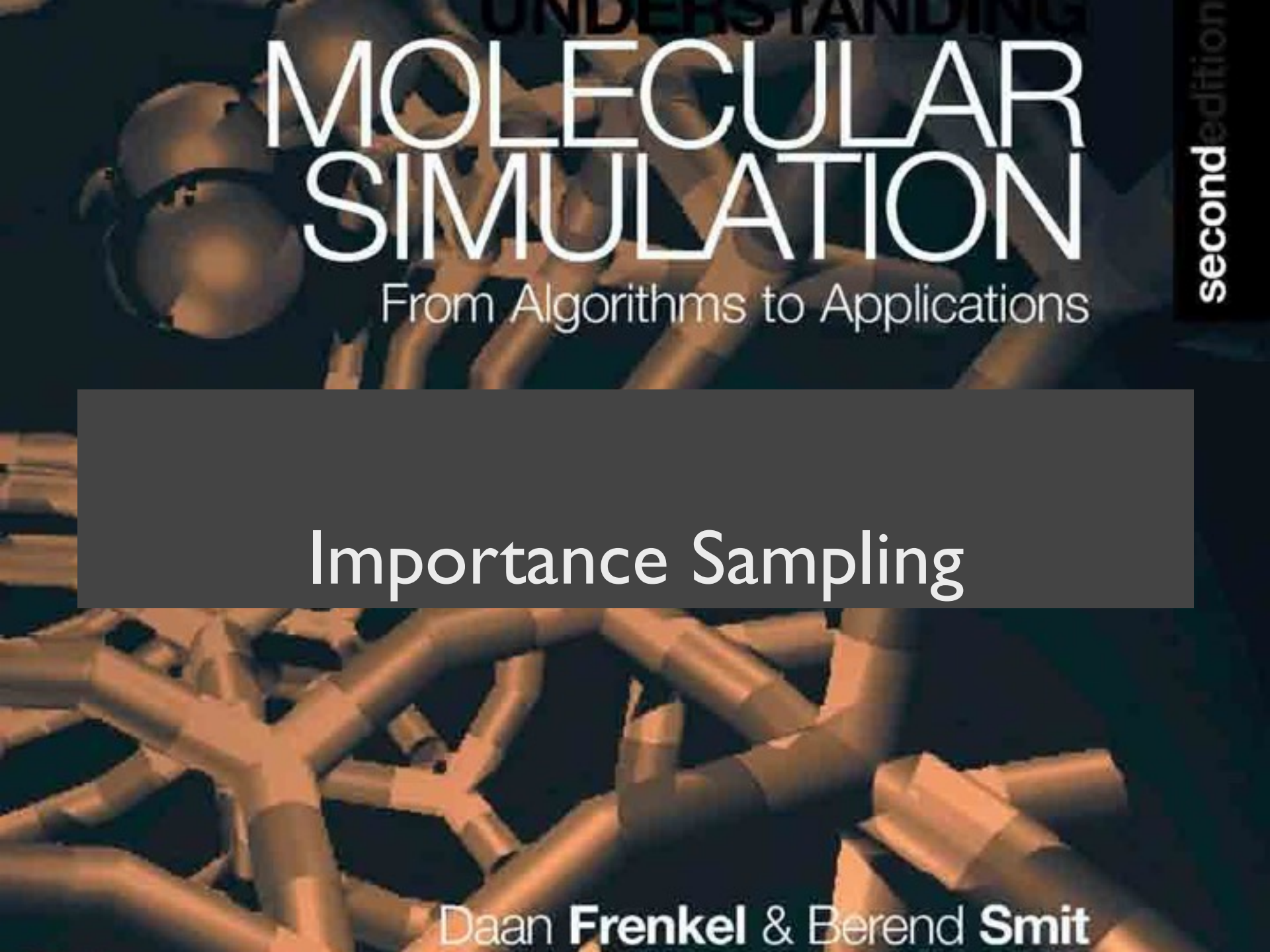
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Free energy:

$$\beta F = -\ln Q_{N,V,T}$$

The background of the slide is a 3D molecular simulation. It features a complex network of orange and yellow rods representing atoms or molecules, with some spheres visible. The overall color scheme is dark with warm, glowing highlights from the simulation.

MOLECULAR SIMULATION

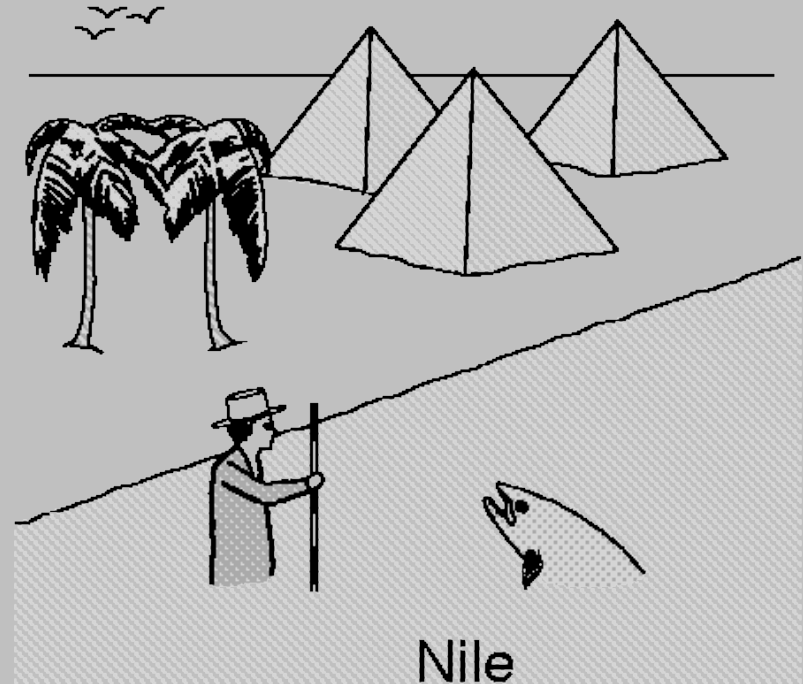
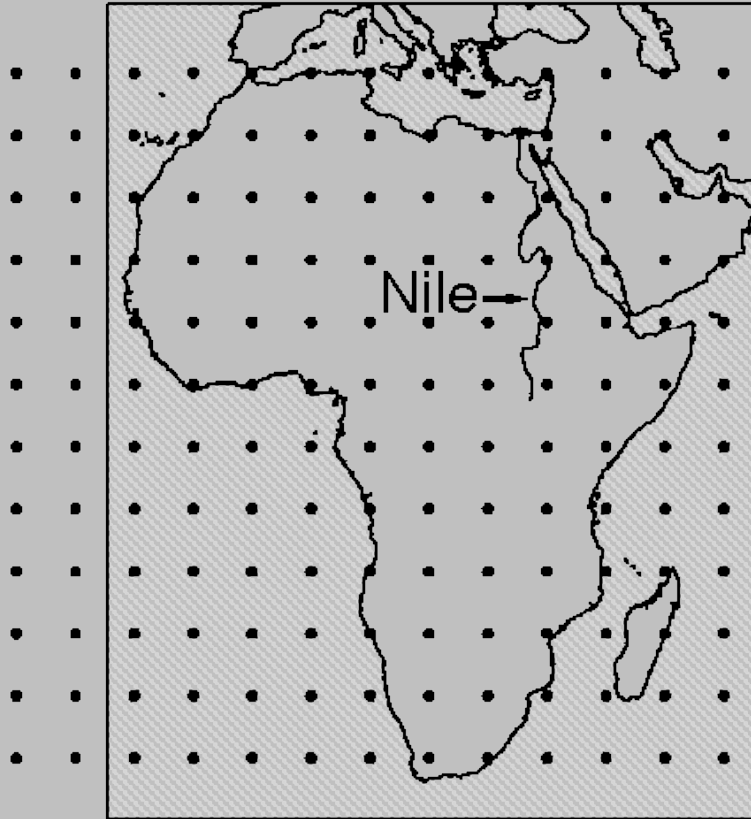
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Importance Sampling

Daan **Frenkel** & Berend **Smit**

Numerical Integration



Ensemble Average

Ensemble Average

$$\langle A \rangle_{NVT} = \frac{1}{Q_{NVT}} \frac{1}{\Lambda^{3N} N!} \int d\mathbf{r}^N A(\mathbf{r}^N) \exp[-\beta U(\mathbf{r}^N)]$$

Ensemble Average

$$\langle A \rangle_{NVT} = \frac{1}{Q_{NVT}} \frac{1}{\Lambda^{3N} N!} \int d\mathbf{r}^N A(\mathbf{r}^N) \exp[-\beta U(\mathbf{r}^N)]$$

$$P(\mathbf{r}^N) = \frac{\exp[-\beta U(\mathbf{r}^N)]}{Q_{NVT} \Lambda^{3N} N!}$$

Ensemble Average

$$\begin{aligned}\langle A \rangle_{NVT} &= \frac{1}{Q_{NVT}} \frac{1}{\Lambda^{3N} N!} \int d\mathbf{r}^N A(\mathbf{r}^N) \exp[-\beta U(\mathbf{r}^N)] \\ &= \frac{\int d\mathbf{r}^N A(\mathbf{r}^N) P(\mathbf{r}^N)}{\int d\mathbf{r}^N P(\mathbf{r}^N)}\end{aligned}$$

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Generate configurations using Monte Carlo moves

$$\{\mathbf{r}_1^N, \mathbf{r}_2^N, \mathbf{r}_3^N, \mathbf{r}_4^N \cdots, \mathbf{r}_M^N\}$$

Ensemble Average

$$\begin{aligned}\langle A \rangle_{NVT} &= \frac{1}{Q_{NVT}} \frac{1}{\Lambda^{3N} N!} \int d\mathbf{r}^N A(\mathbf{r}^N) \exp[-\beta U(\mathbf{r}^N)] \\ &= \frac{\int d\mathbf{r}^N A(\mathbf{r}^N) P(\mathbf{r}^N)}{\int d\mathbf{r}^N P(\mathbf{r}^N)} = \int d\mathbf{r}^N A(\mathbf{r}^N) P(\mathbf{r}^N) \\ &= \frac{\int d\mathbf{r}^N A(\mathbf{r}^N) C \exp[-\beta U(\mathbf{r}^N)]}{\int d\mathbf{r}^N C \exp[-\beta U(\mathbf{r}^N)]} = \frac{\int d\mathbf{r}^N A(\mathbf{r}^N) \exp[-\beta U(\mathbf{r}^N)]}{\int d\mathbf{r}^N \exp[-\beta U(\mathbf{r}^N)]}\end{aligned}$$

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$$\{\mathbf{r}_1^N, \mathbf{r}_2^N, \mathbf{r}_3^N, \mathbf{r}_4^N \cdots, \mathbf{r}_M^N\}$$

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$$P^{MC}(\mathbf{r}^N) = C^{MC} \exp[-\beta U(\mathbf{r}^N)]$$

Ensemble Average

$$\begin{aligned}\langle A \rangle_{NVT} &= \frac{1}{Q_{NVT}} \frac{1}{\Lambda^{3N} N!} \int d\mathbf{r}^N A(\mathbf{r}^N) \exp[-\beta U(\mathbf{r}^N)] \\ &= \frac{\int d\mathbf{r}^N A(\mathbf{r}^N) P(\mathbf{r}^N)}{\int d\mathbf{r}^N P(\mathbf{r}^N)} = \int d\mathbf{r}^N A(\mathbf{r}^N) P(\mathbf{r}^N) \\ &= \frac{\int d\mathbf{r}^N A(\mathbf{r}^N) C \exp[-\beta U(\mathbf{r}^N)]}{\int d\mathbf{r}^N C \exp[-\beta U(\mathbf{r}^N)]} = \frac{\int d\mathbf{r}^N A(\mathbf{r}^N) \exp[-\beta U(\mathbf{r}^N)]}{\int d\mathbf{r}^N \exp[-\beta U(\mathbf{r}^N)]}\end{aligned}$$

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$$\{\mathbf{r}_1^N, \mathbf{r}_2^N, \mathbf{r}_3^N, \mathbf{r}_4^N \dots, \mathbf{r}_M^N\} \quad \bar{A} = \frac{1}{M} \sum_{i=1}^M A(\mathbf{r}_i^N)$$

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Ensemble Average

$$\begin{aligned}\langle A \rangle_{NVT} &= \frac{1}{Q_{NVT}} \frac{1}{\Lambda^{3N} N!} \int d\mathbf{r}^N A(\mathbf{r}^N) \exp[-\beta U(\mathbf{r}^N)] \\ &= \frac{\int d\mathbf{r}^N A(\mathbf{r}^N) P(\mathbf{r}^N)}{\int d\mathbf{r}^N P(\mathbf{r}^N)} = \int d\mathbf{r}^N A(\mathbf{r}^N) P(\mathbf{r}^N) \\ &= \frac{\int d\mathbf{r}^N A(\mathbf{r}^N) C \exp[-\beta U(\mathbf{r}^N)]}{\int d\mathbf{r}^N C \exp[-\beta U(\mathbf{r}^N)]} = \frac{\int d\mathbf{r}^N A(\mathbf{r}^N) \exp[-\beta U(\mathbf{r}^N)]}{\int d\mathbf{r}^N \exp[-\beta U(\mathbf{r}^N)]}\end{aligned}$$

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Ensemble Average

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Ensemble Average

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Generate configurations using Monte Carlo moves

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$$\begin{aligned}
 &= \frac{\int d\mathbf{r}^N A(\mathbf{r}^N) C^{MC} \exp[-\beta U(\mathbf{r}^N)]}{\int d\mathbf{r}^N C^{MC} \exp[-\beta U(\mathbf{r}^N)]} \\
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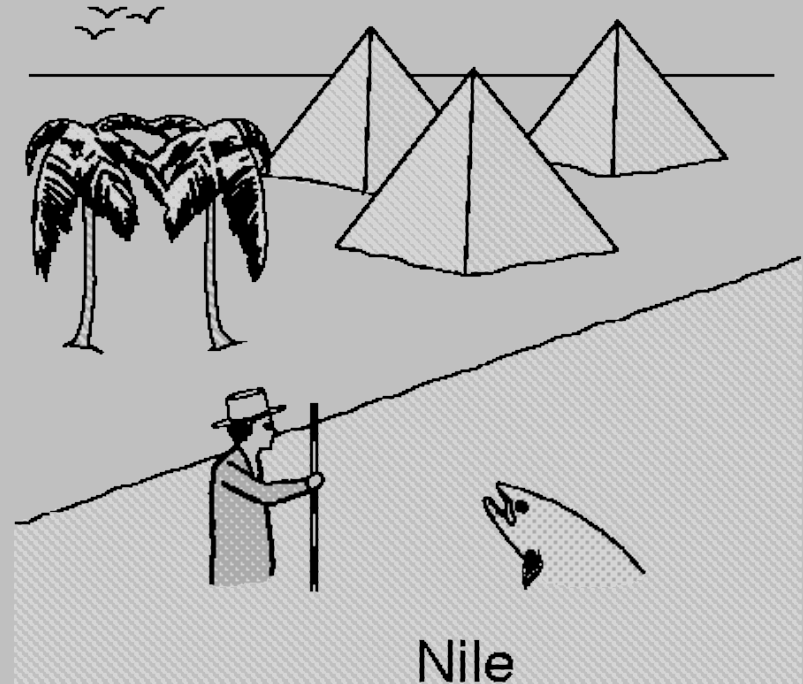
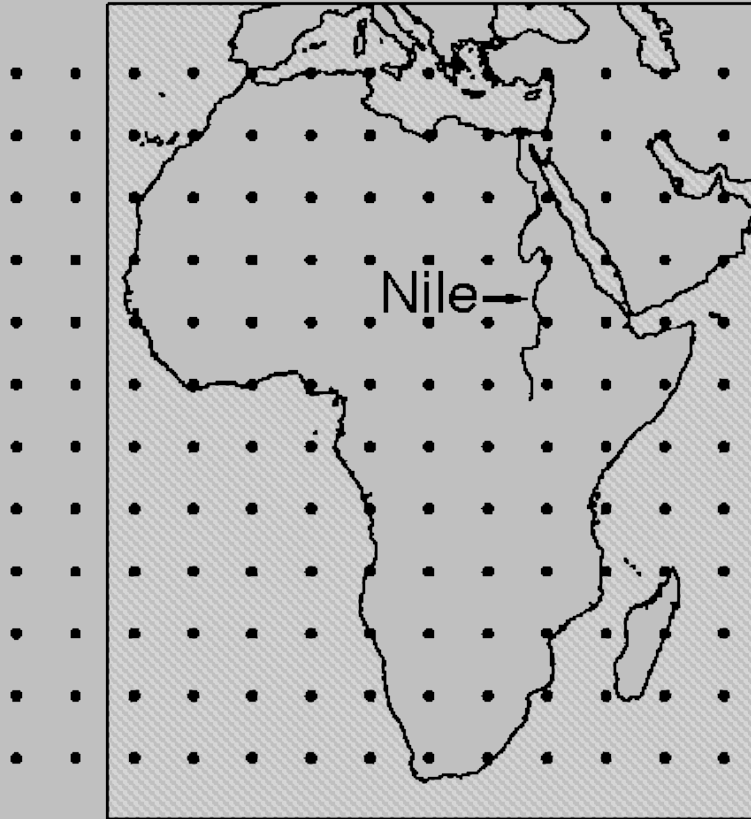
Generate configurations using Monte Carlo moves

$$\{\mathbf{r}_1^N, \mathbf{r}_2^N, \mathbf{r}_3^N, \mathbf{r}_4^N \dots, \mathbf{r}_M^N\} \quad \bar{A} = \frac{1}{M} \sum_{i=1}^M A(\mathbf{r}_i^N) = \frac{\int d\mathbf{r}^N A(\mathbf{r}^N) P^{MC}(\mathbf{r}^N)}{\int d\mathbf{r}^N P^{MC}(\mathbf{r}^N)}$$

with:

$$\begin{aligned}
 P^{MC}(\mathbf{r}^N) &= C^{MC} \exp[-\beta U(\mathbf{r}^N)] \\
 &= \frac{\int d\mathbf{r}^N A(\mathbf{r}^N) C^{MC} \exp[-\beta U(\mathbf{r}^N)]}{\int d\mathbf{r}^N C^{MC} \exp[-\beta U(\mathbf{r}^N)]} \\
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 \end{aligned}$$

Importance Sampling



Algorithm 1 (Basic Metropolis Algorithm)

<pre>PROGRAM mc</pre>	basic Metropolis algorithm
<pre>do icycl=1,ncycl</pre>	perform <code>ncycl</code> MC cycles
<pre> call mcmove</pre>	displace a particle
<pre> if (mod(icycl,nsamp).eq.0)</pre>	
<pre>+ call sample</pre>	sample averages
<pre> enddo</pre>	
<pre>end</pre>	

Comments to this algorithm:

1. Subroutine `mcmove` attempts to displace a randomly selected particle (see Algorithm 2).
2. Subroutine `sample` samples quantities every `nsamp`th cycle.

Algorithm 2 (Attempt to Displace a Particle)

<pre>SUBROUTINE mcmove o=int(ranf()*npart)+1 call ener(x(o), eno) xn=x(o)+(ranf()-0.5)*delx call ener(xn, enn) if (ranf().lt.exp(-beta + *(enn-eno)) x(o)=xn return end</pre>	attempts to displace a particle select a particle at random energy old configuration give particle random displacement energy new configuration acceptance rule (3.2.1) accepted: replace $x(o)$ by xn
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Comments to this algorithm:

1. Subroutine `ener` calculates the energy of a particle at the given position.
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3. The `ranf()` is a random number uniform in $[0, 1]$.

Questions

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Questions

Desired distribution: NVT ensemble

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Markov Processes

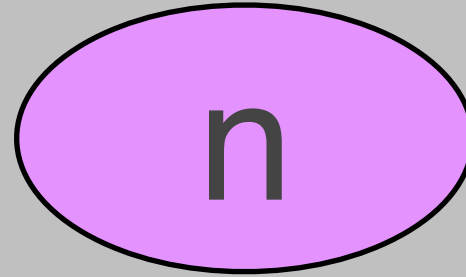
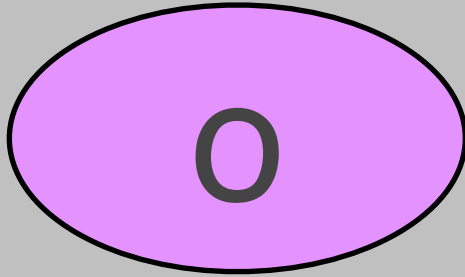
Markov Process

- Next step only depends on the current state
- Ergodic: all possible states can be reached by a set of single steps
- Detailed balance
- * Process will approach a limiting distribution

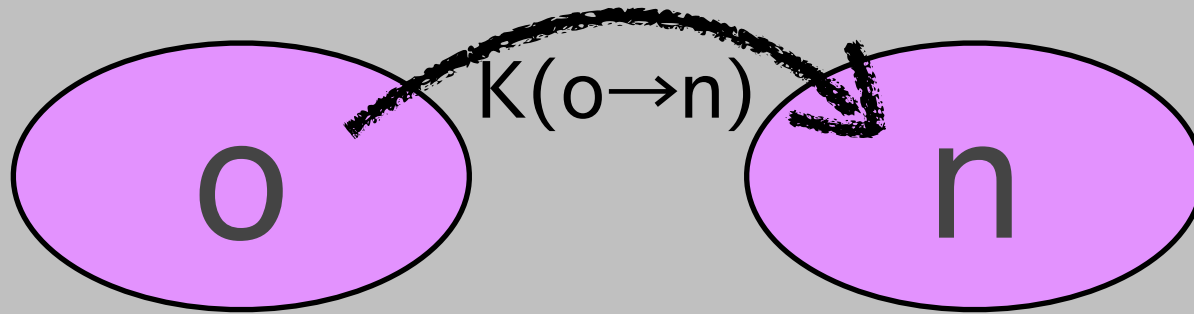
Ensemble - probability

- $P(o)$: probability to find the state o
- Ensemble: take a very large number (M) of identical systems: $N(o) = M \times P(o)$; the total number of systems in the state o

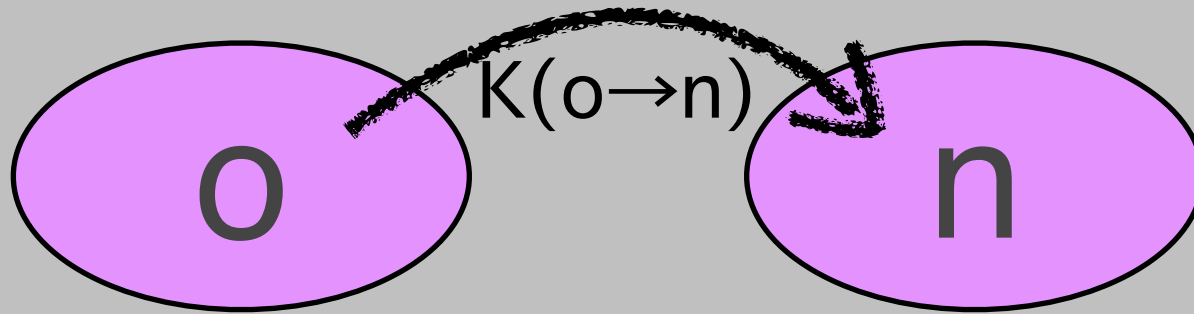
Markov Processes - Detailed Balance



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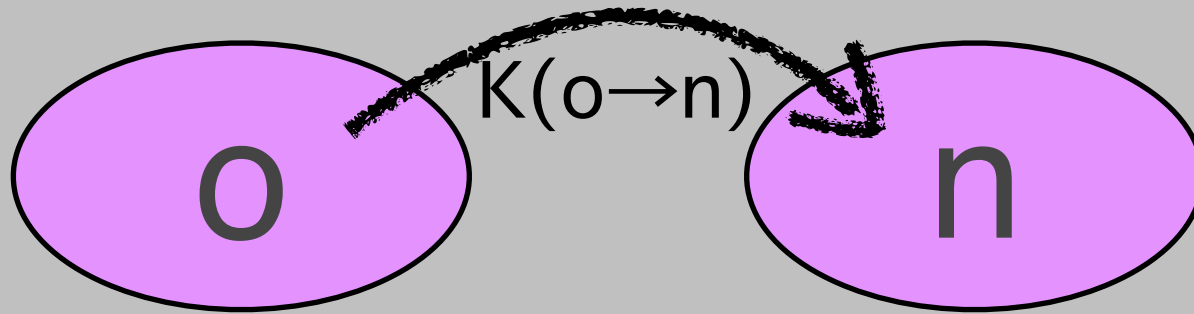


Markov Processes - Detailed Balance



$K(o \rightarrow n)$: total number of systems in our ensemble that move $o \rightarrow n$

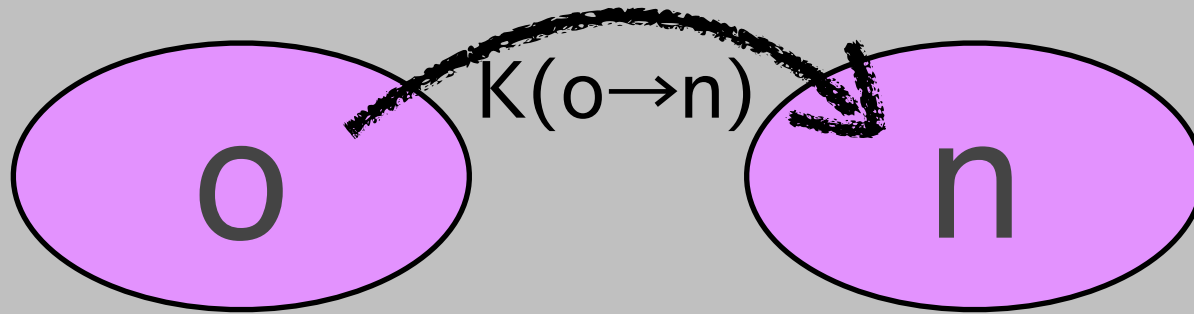
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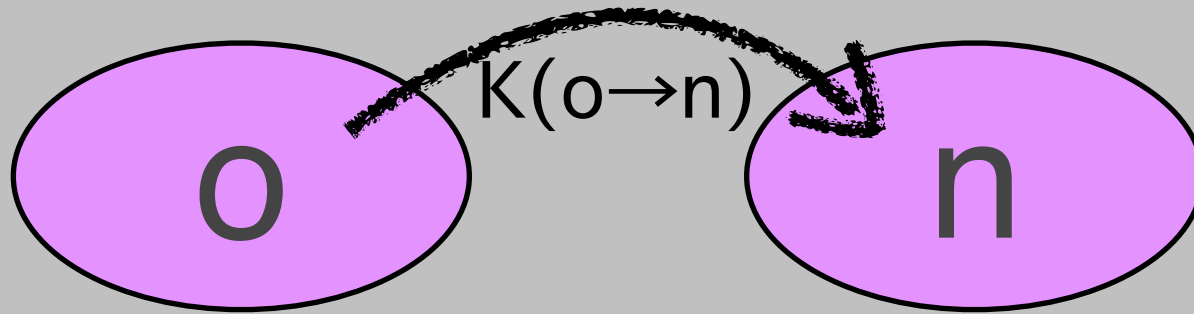
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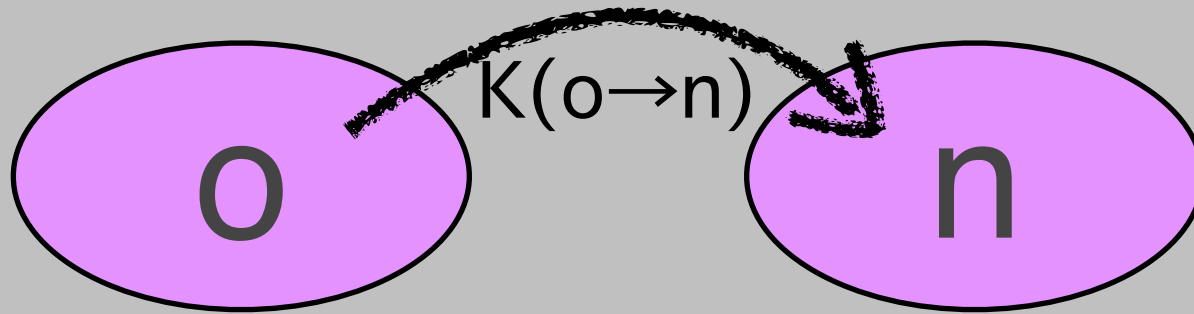


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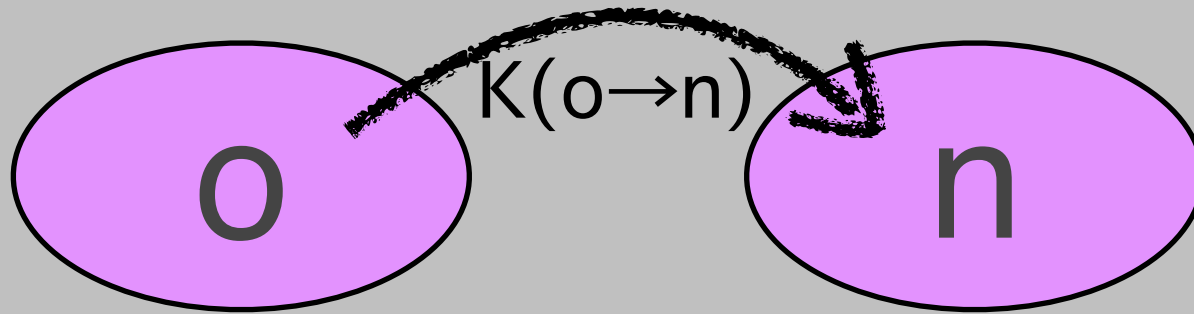


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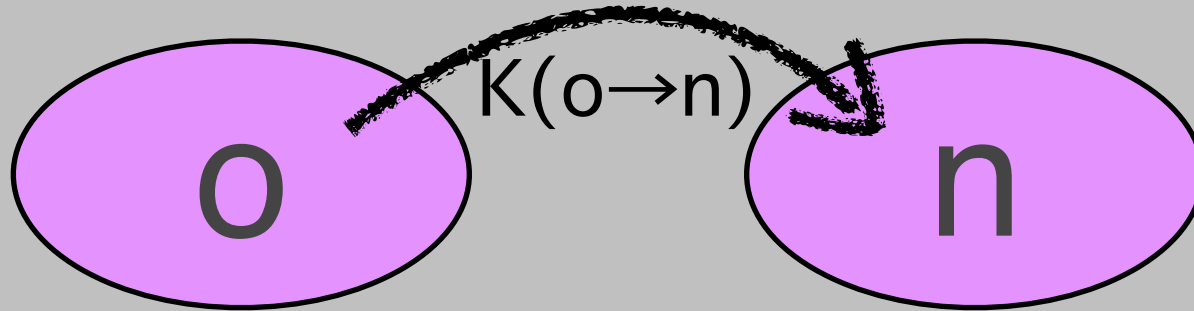


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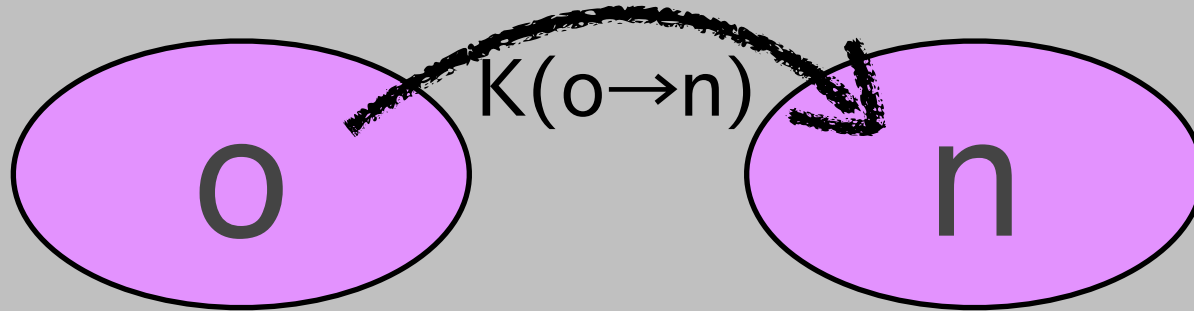
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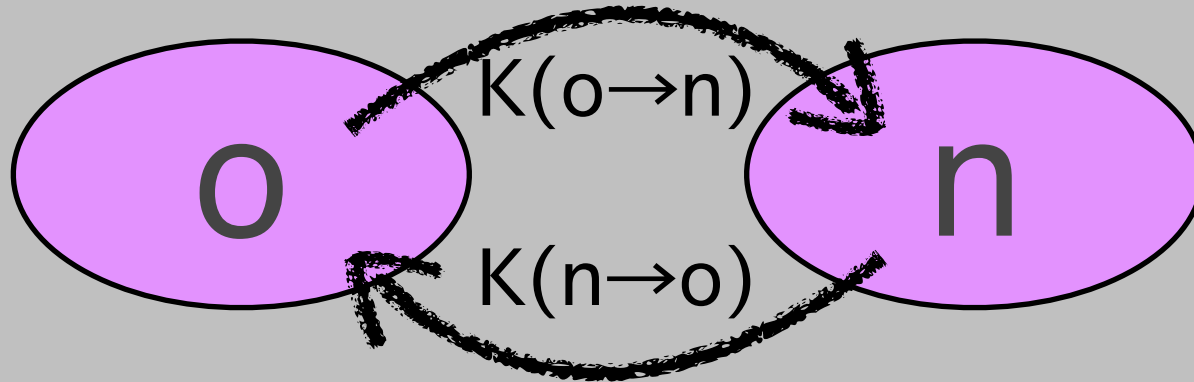


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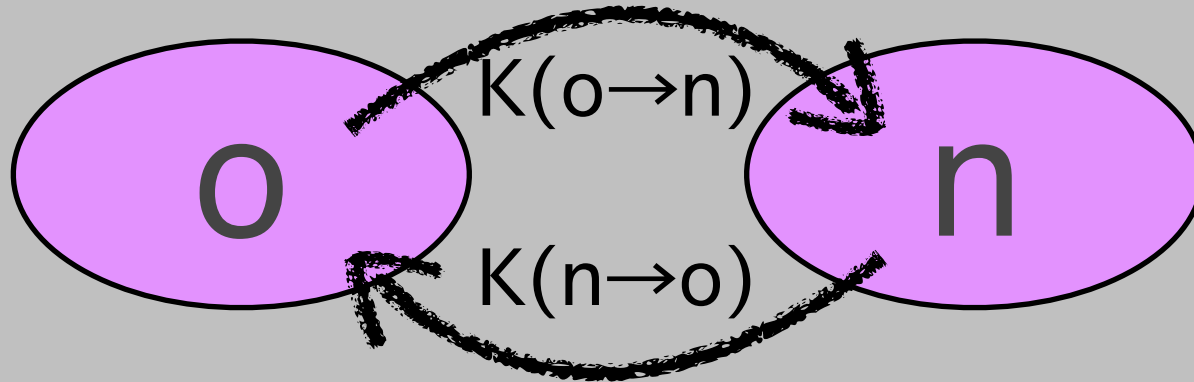
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Markov Processes - Detailed Balance



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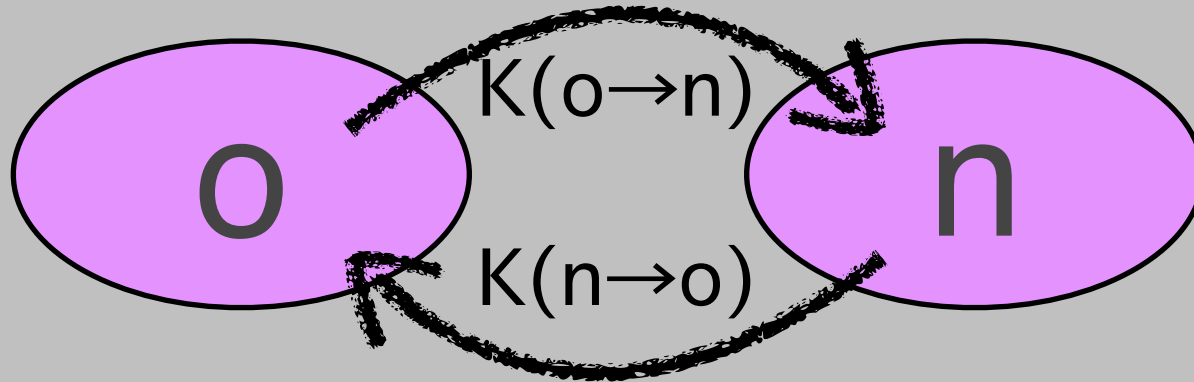
Markov Processes - Detailed Balance



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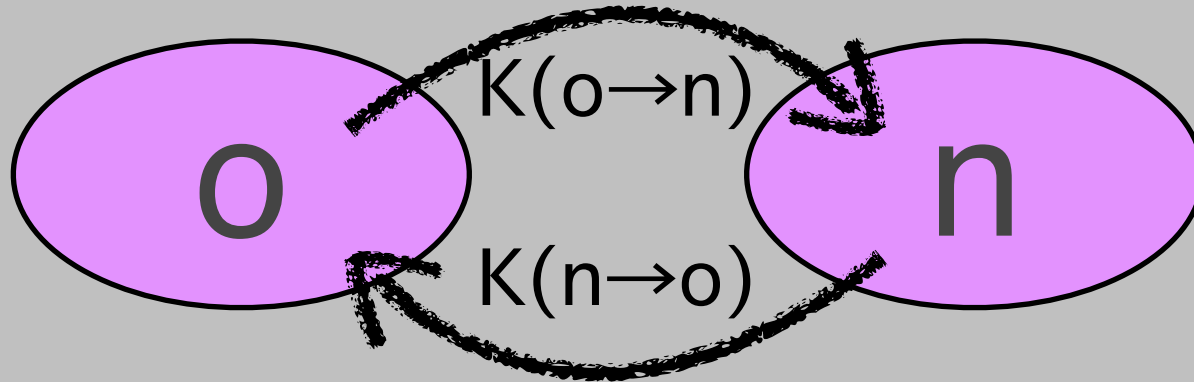


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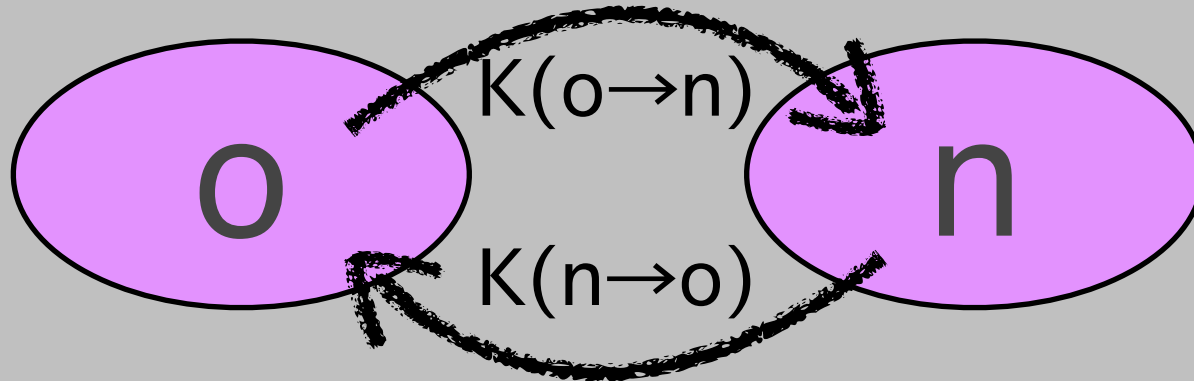
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Which gives as condition for the acceptance rule:

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Metropolis et al.

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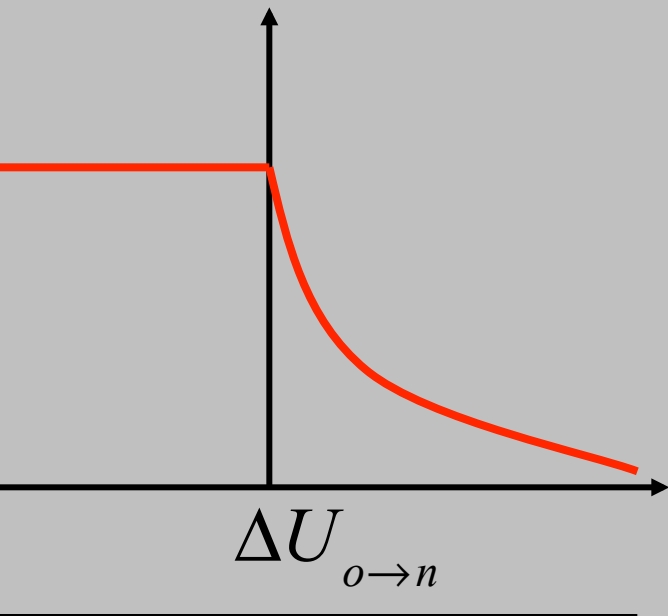
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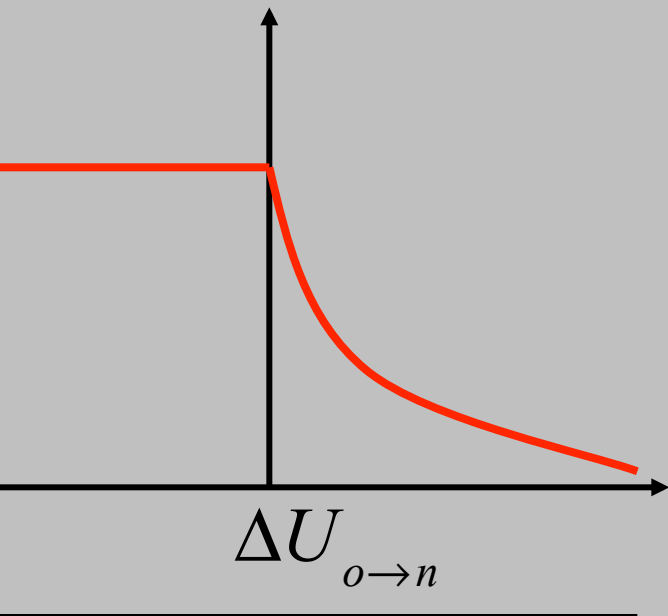
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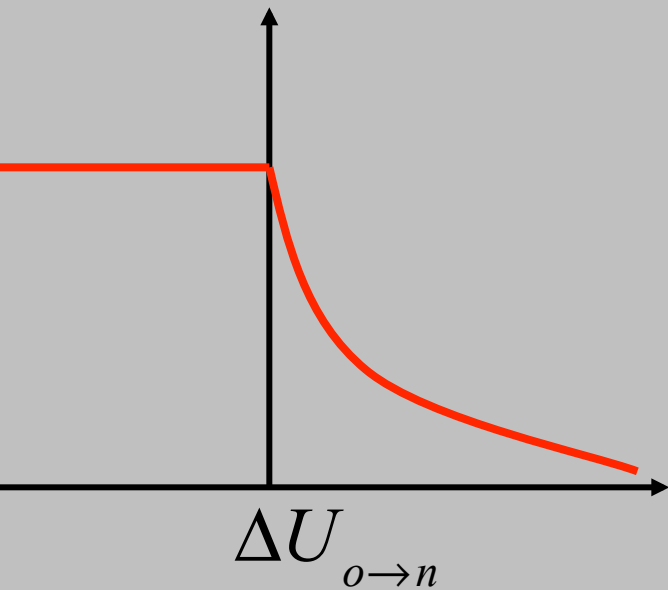


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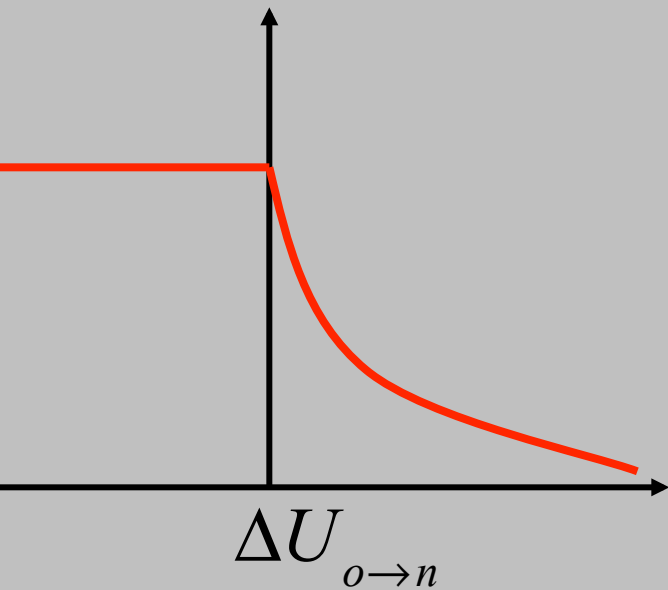
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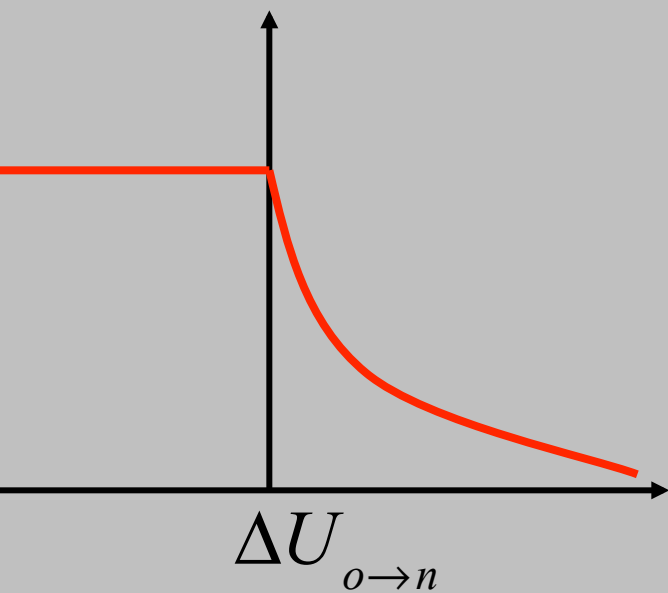
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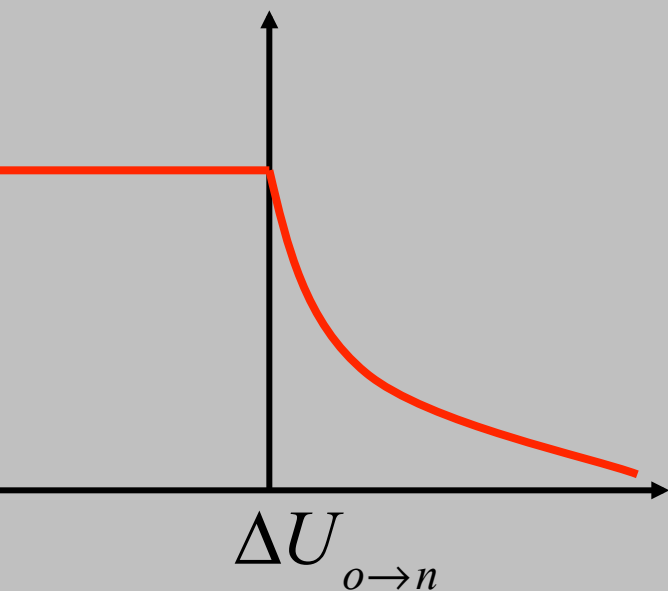
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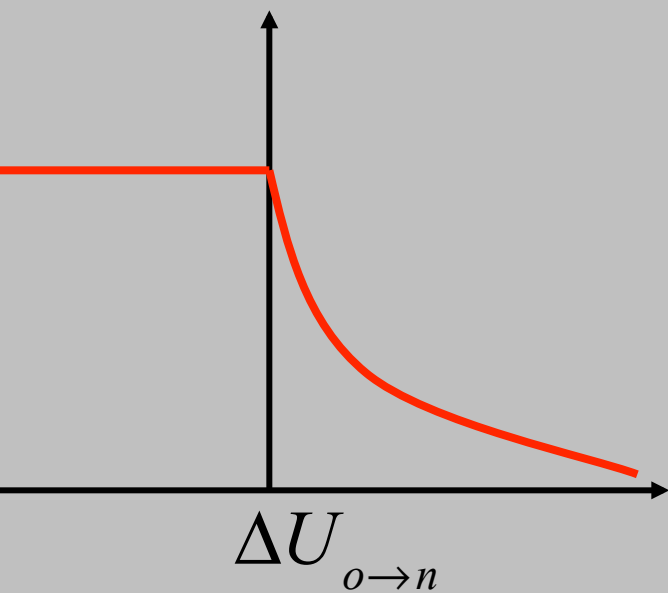
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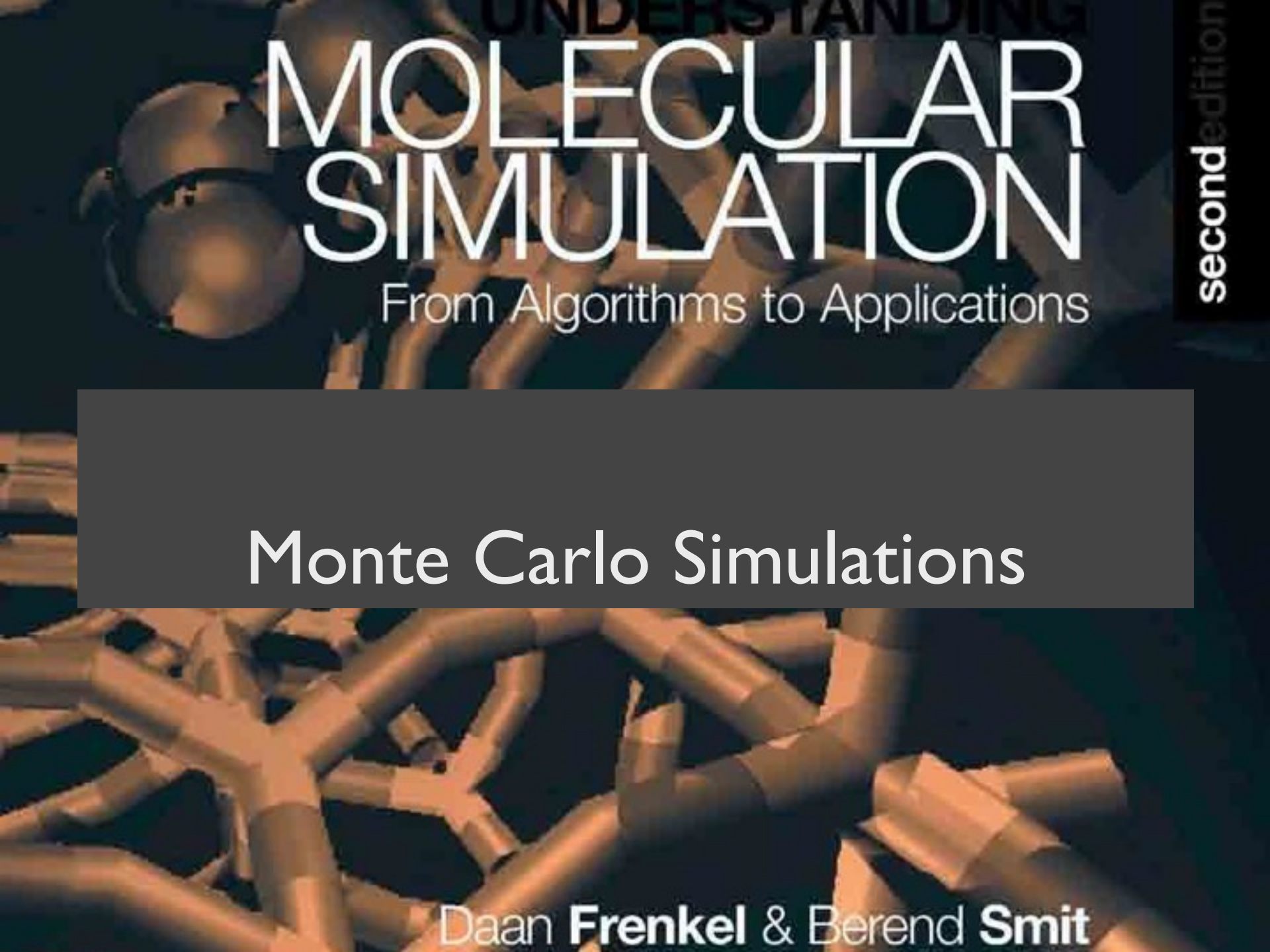
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The background of the slide is a 3D molecular simulation. It features a complex network of orange and yellow rods representing atoms or molecules, with some spheres visible. The overall color scheme is dark with warm, glowing highlights from the simulation.

MOLECULAR SIMULATION

From Algorithms to Applications

second edition

Monte Carlo Simulations

Daan **Frenkel** & Berend **Smit**

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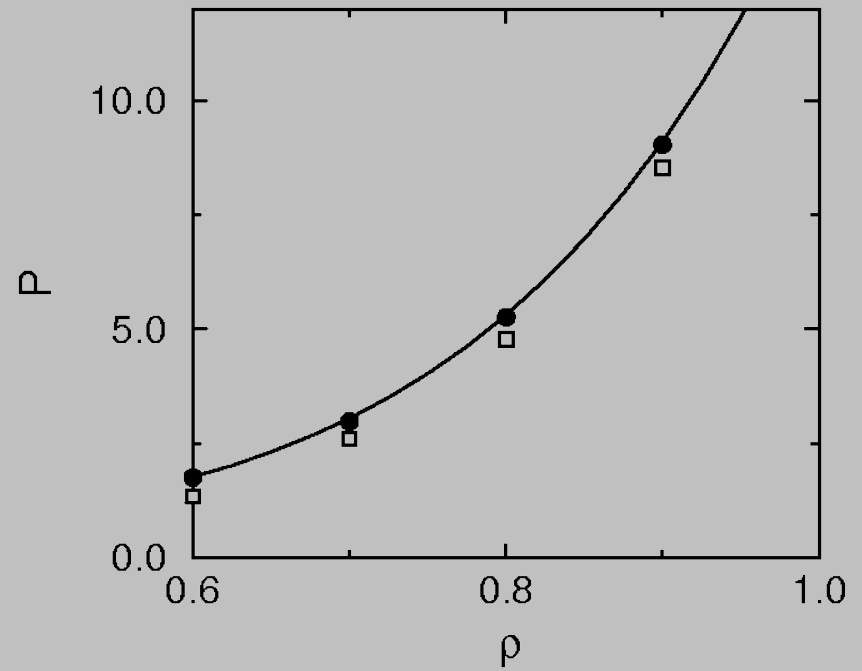
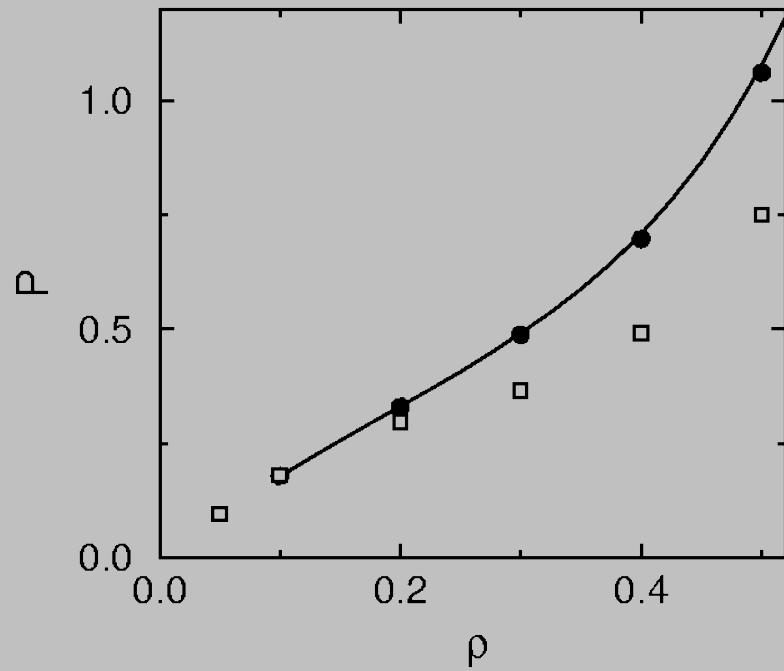


If we do not keep the old configuration:



Independent of the
temperature

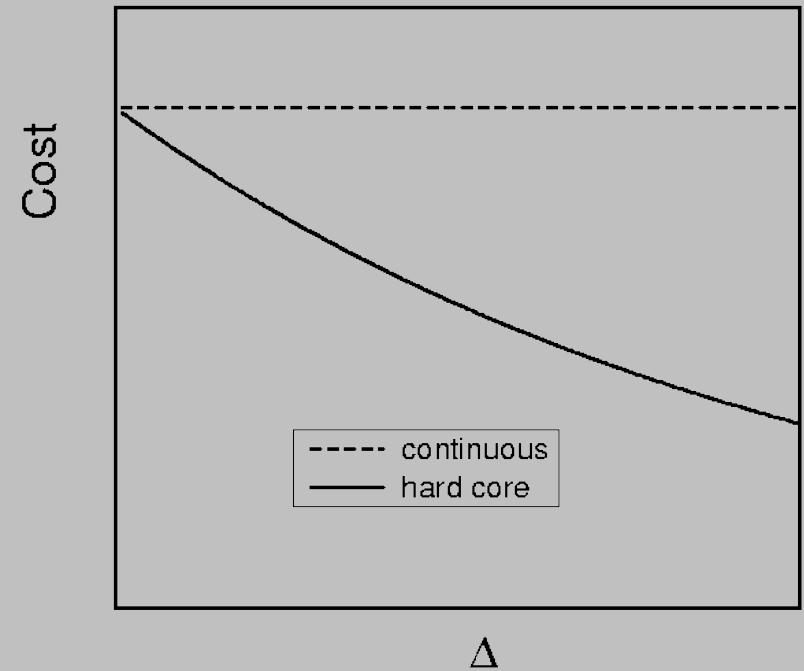
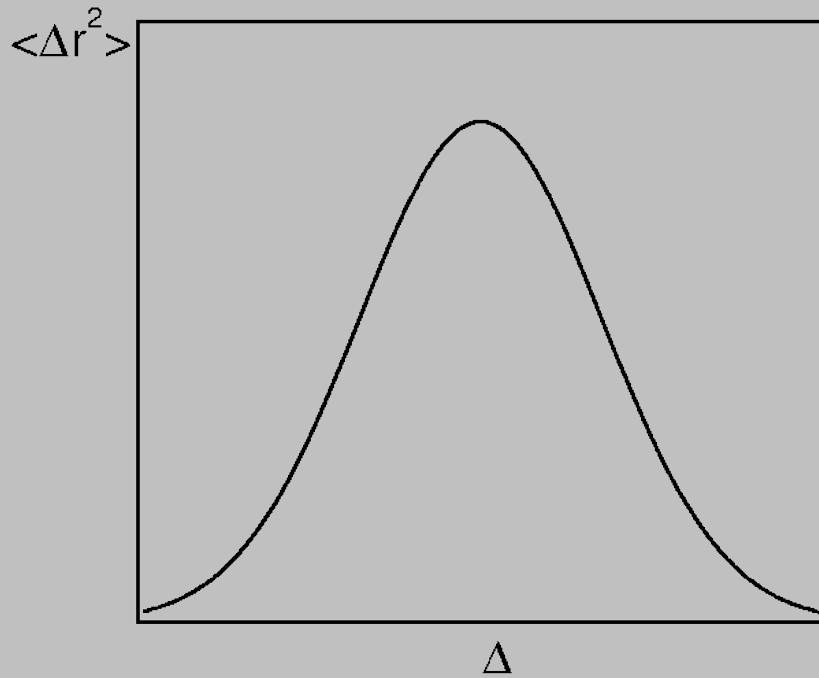
Lennard Jones fluid

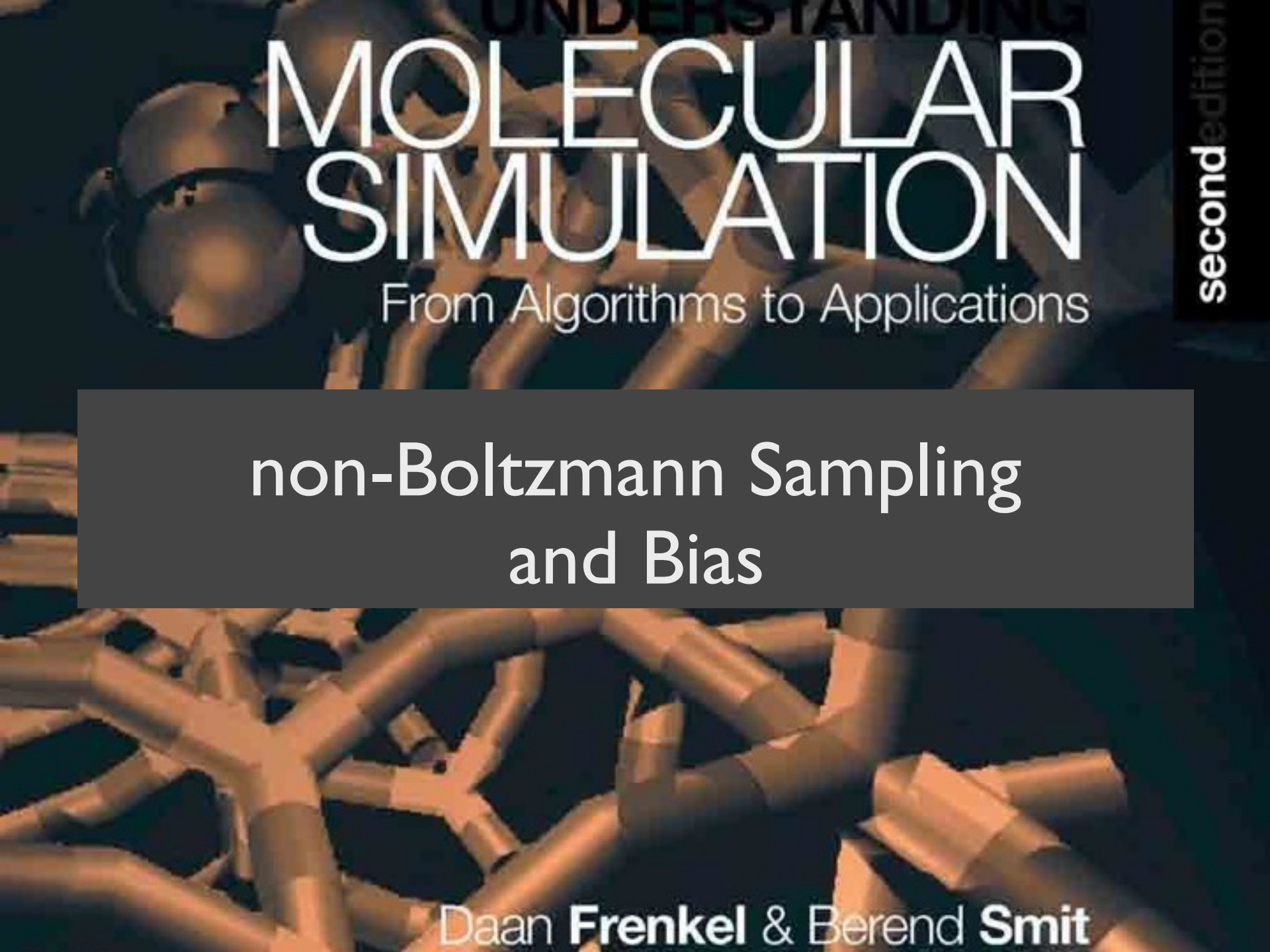


Questions

- How can we prove that this scheme generates the desired distribution of configurations?
- Why make a random selection of the particle to be displaced?
- Why do we need to take the old configuration again?
- **How large should we take: Δx ?**

Not too big Not too small



The background of the slide is a 3D molecular simulation. It features a complex network of orange and yellow rods representing atoms and bonds, set against a dark blue background. In the upper left, there is a cluster of spheres, some colored red and others blue, representing different types of atoms. The overall image has a scientific and technical feel.

MOLECULAR SIMULATION

From Algorithms to Applications

second edition

non-Boltzmann Sampling and Bias

Daan **Frenkel** & Berend **Smit**

Non-Boltzmann sampling

$$\langle A \rangle_{NVT_1} = \frac{1}{Q_{NVT_1}} \frac{1}{\Lambda^{3N} N!} \int d\mathbf{r}^N A(\mathbf{r}^N) \exp[-\beta_1 U(\mathbf{r}^N)]$$

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We perform a simulation at $T=T_2$ and we determine A at $T=T_1$

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T_1 is
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We only
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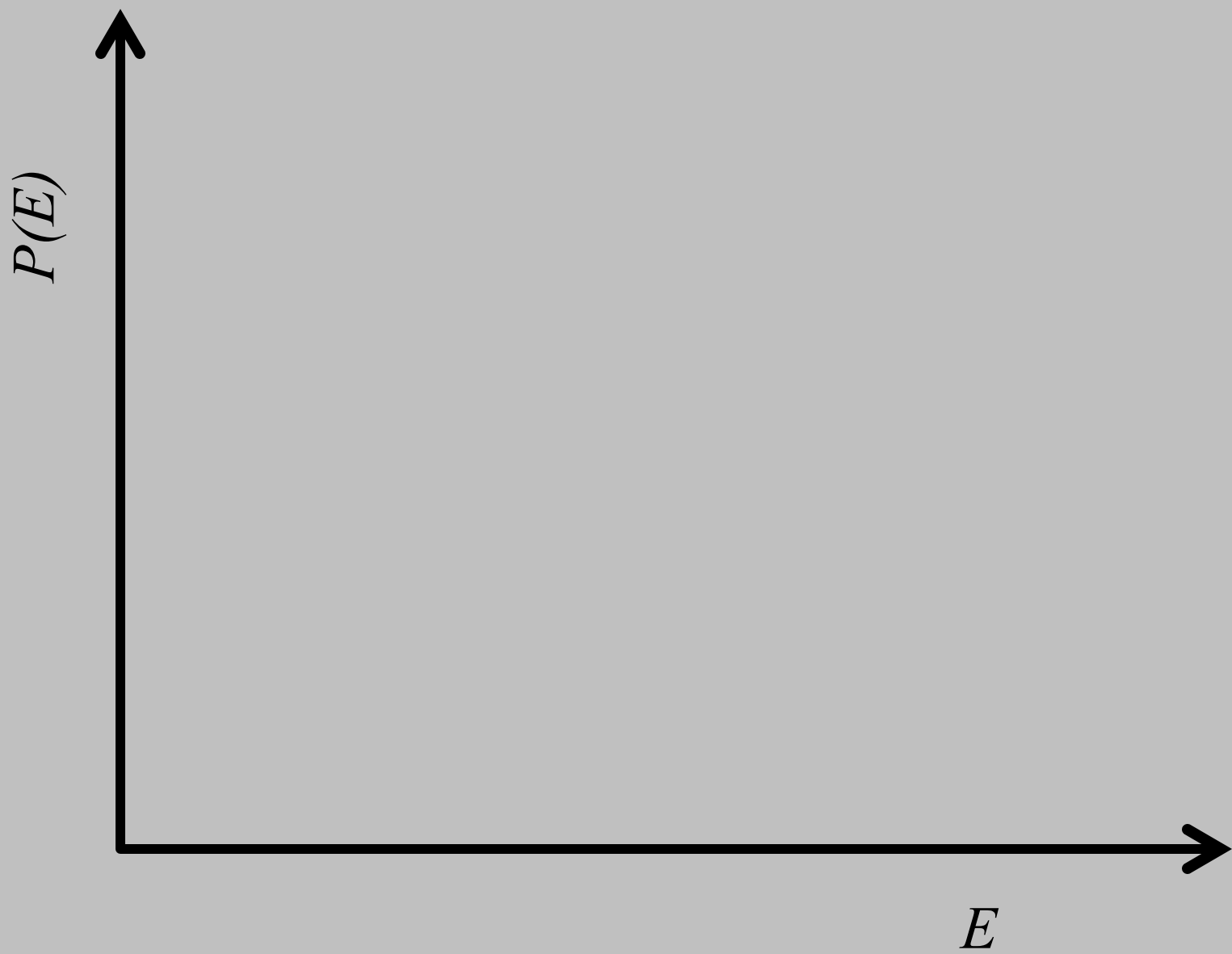
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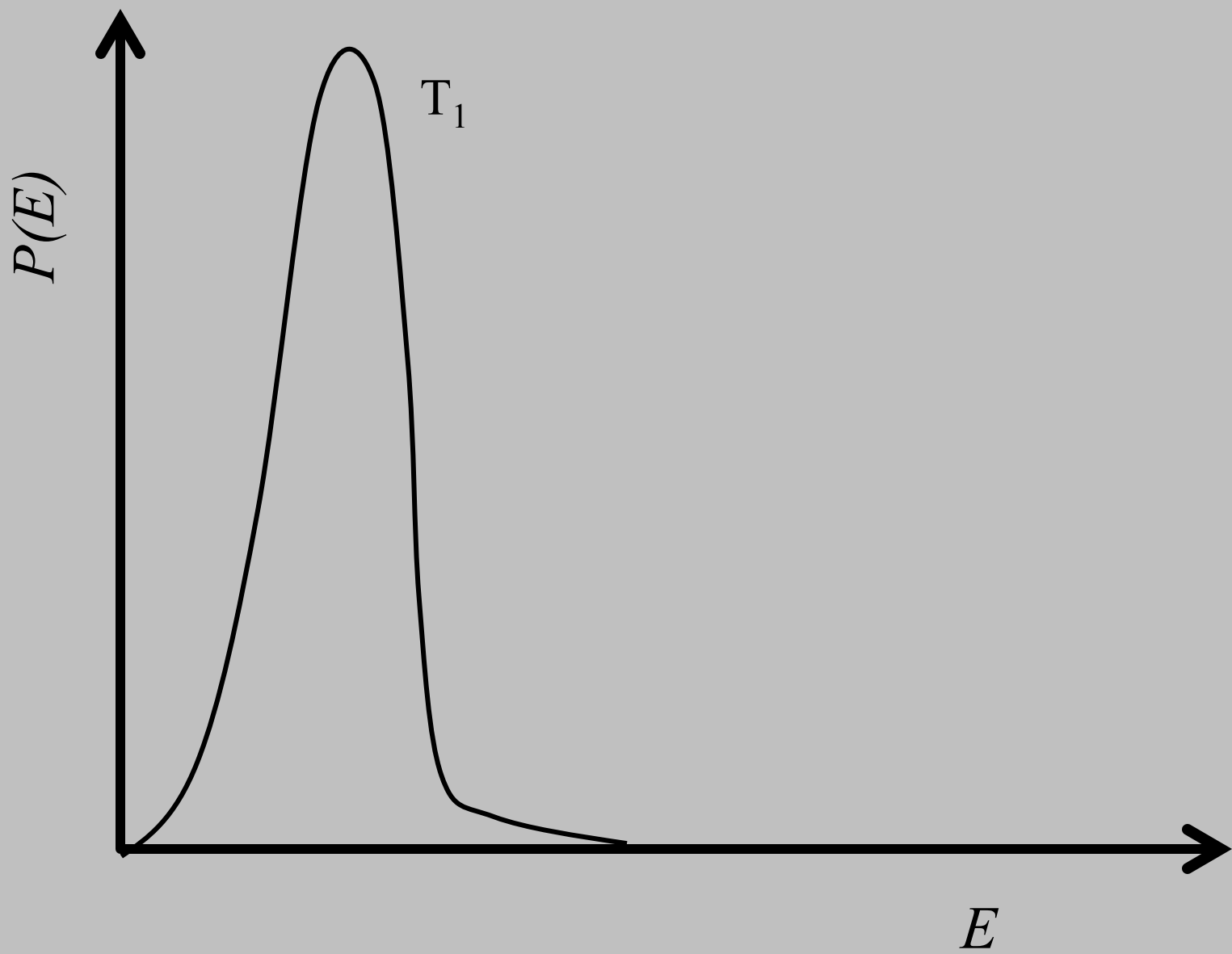
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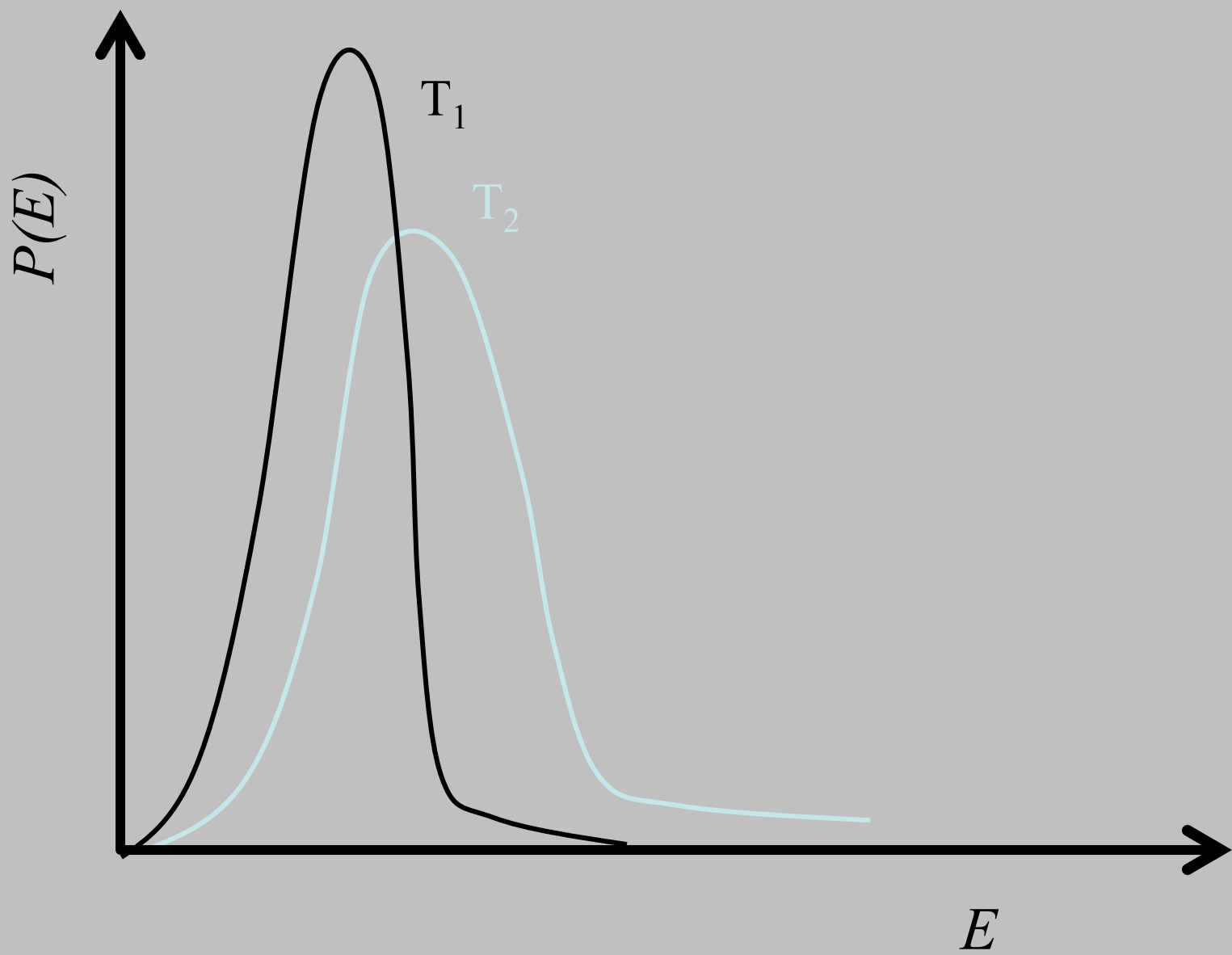
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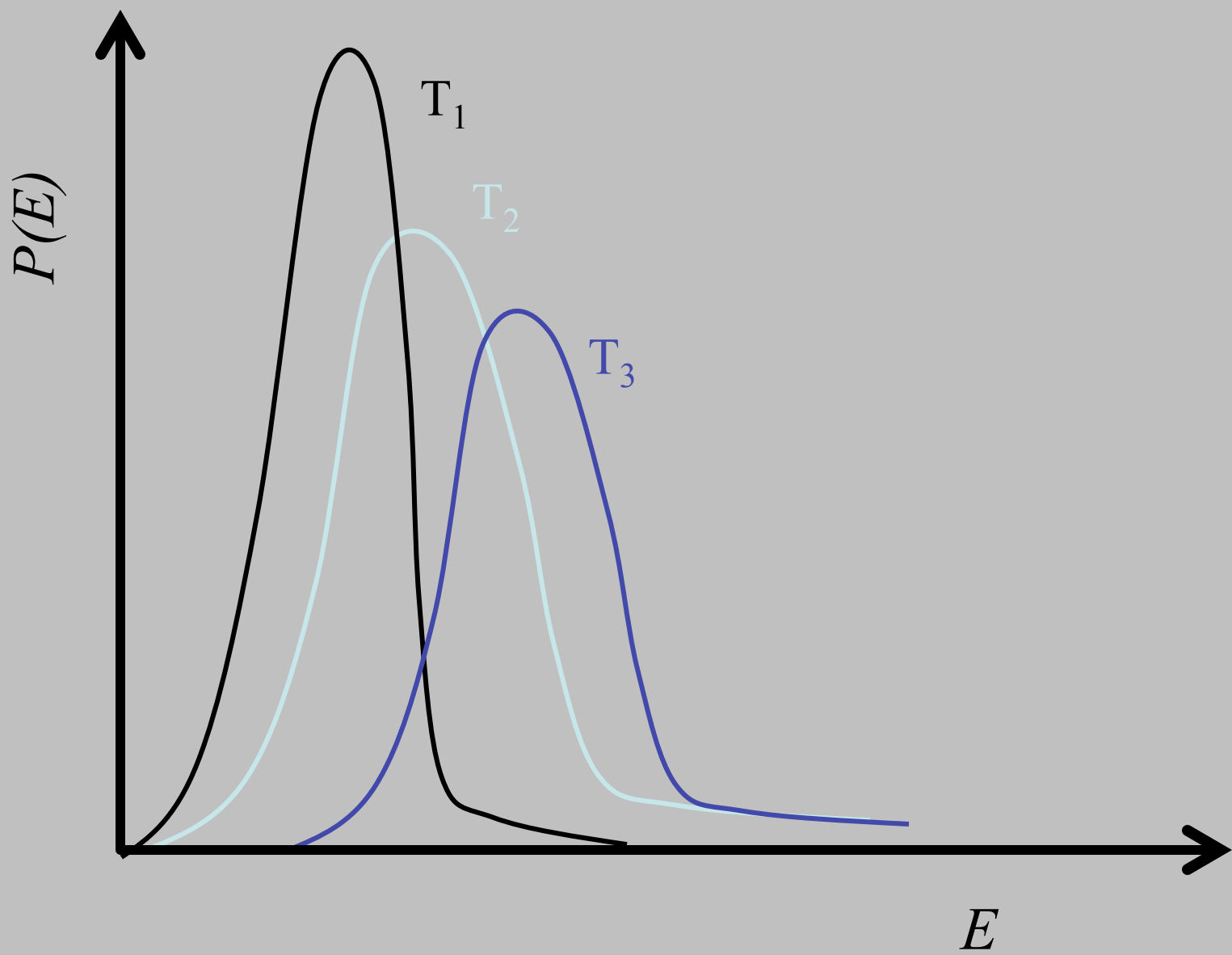
Why are we not using this?

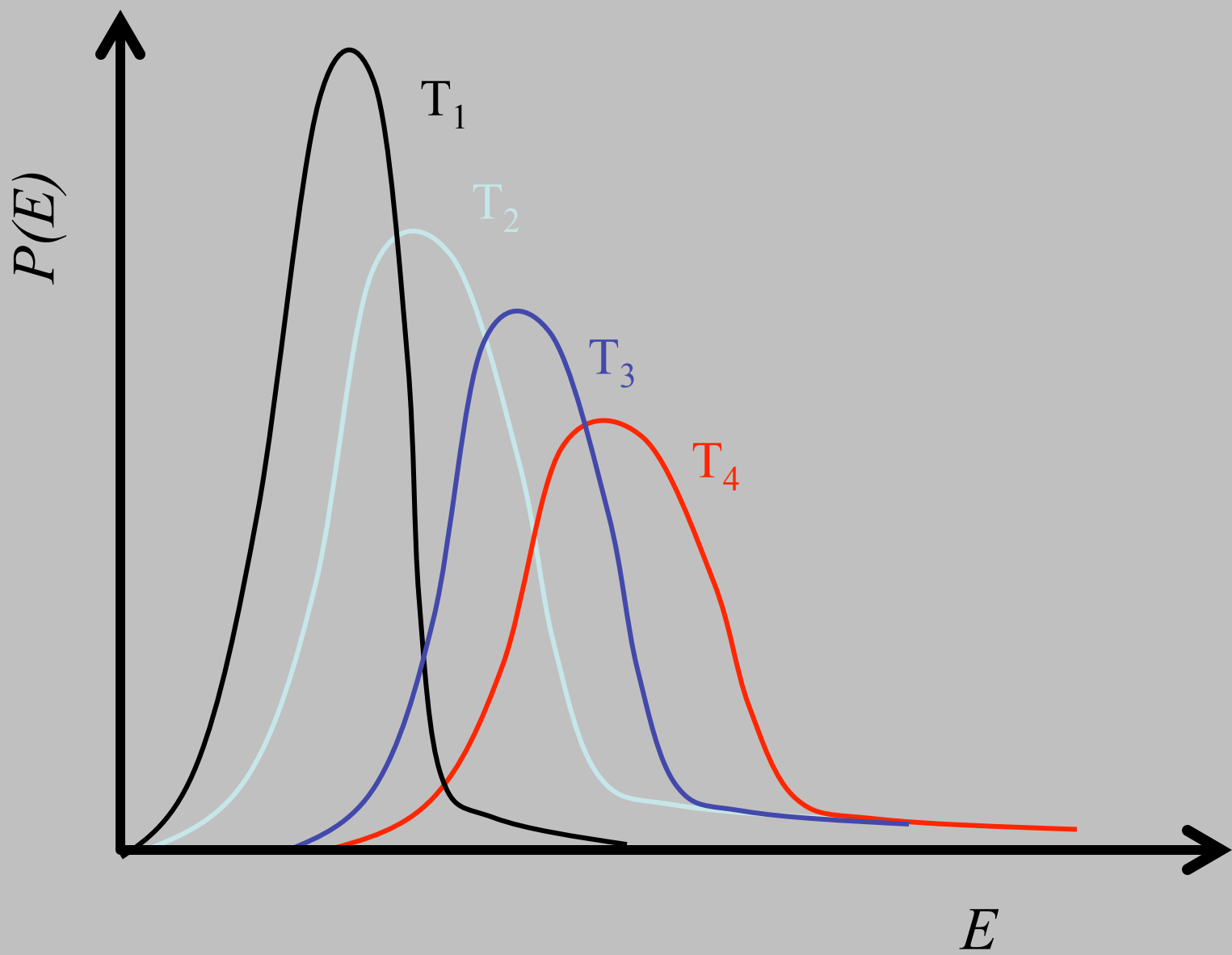
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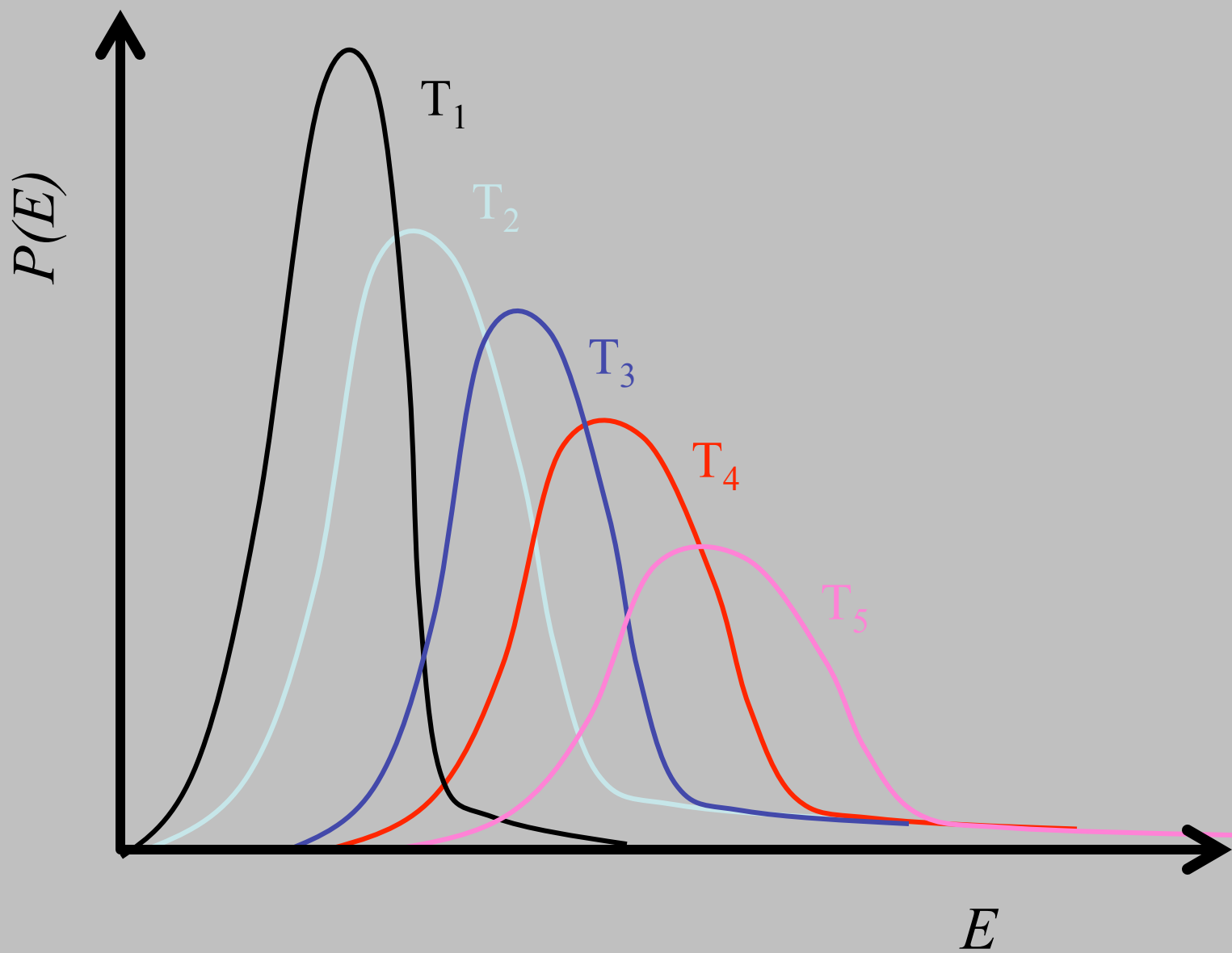


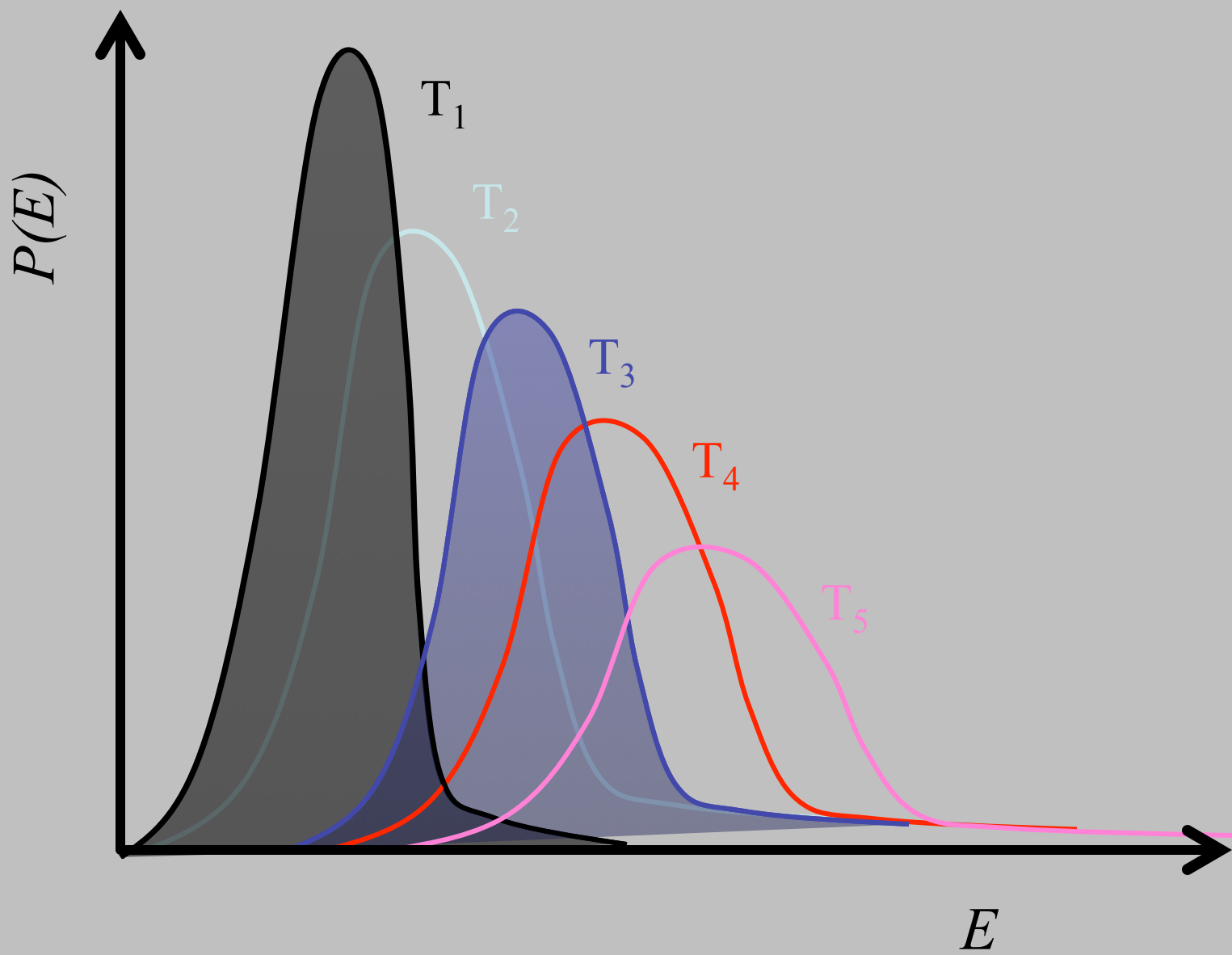


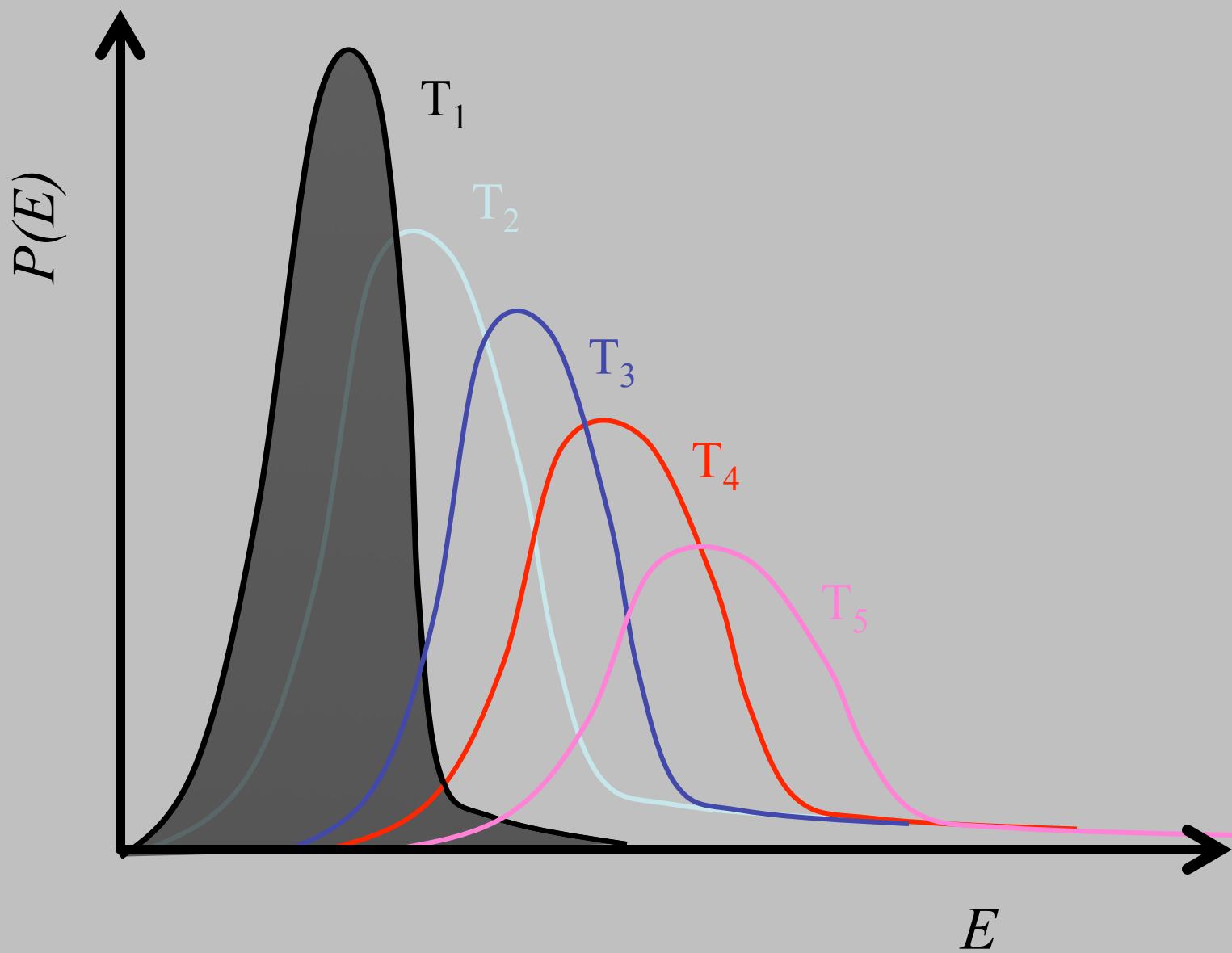


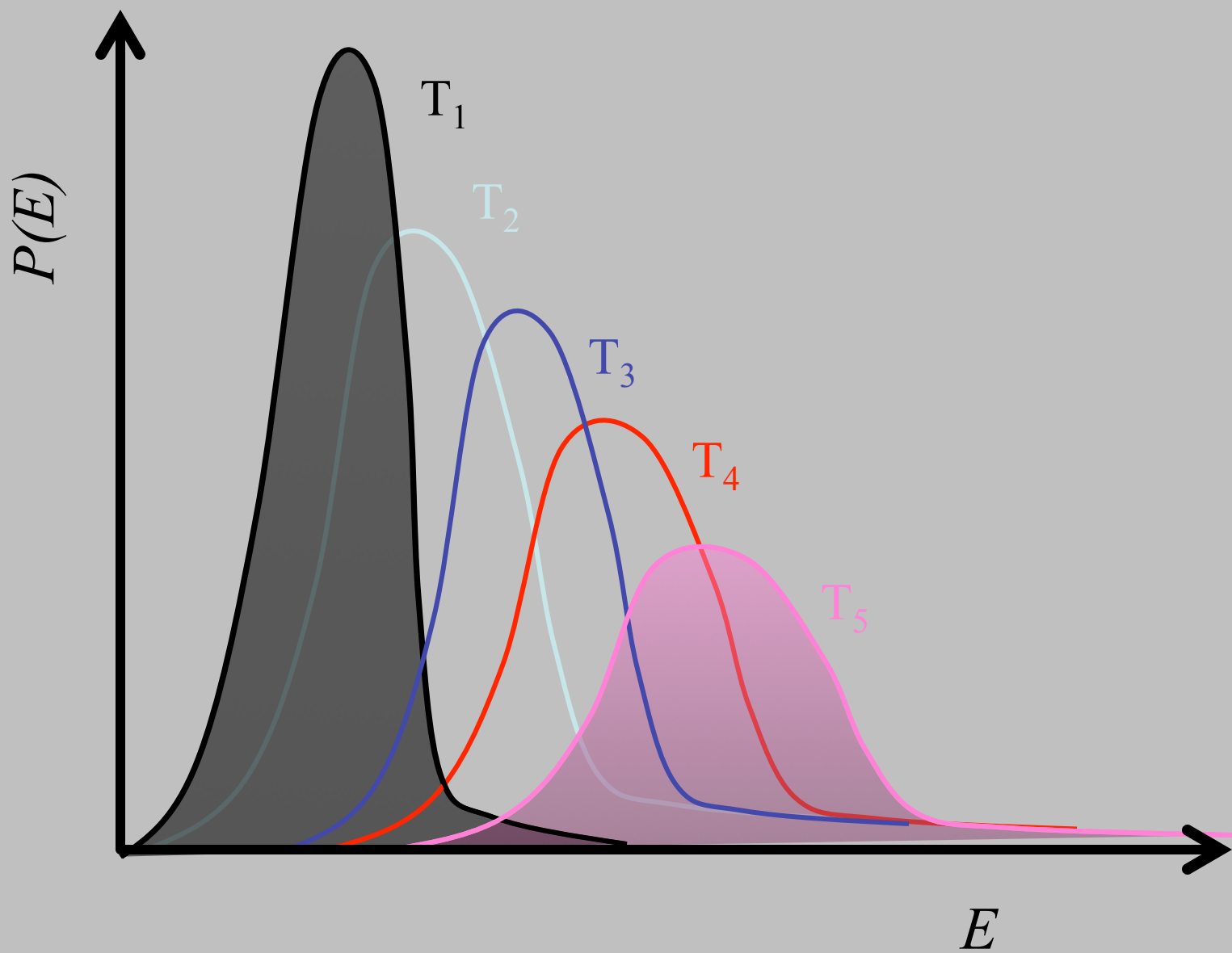


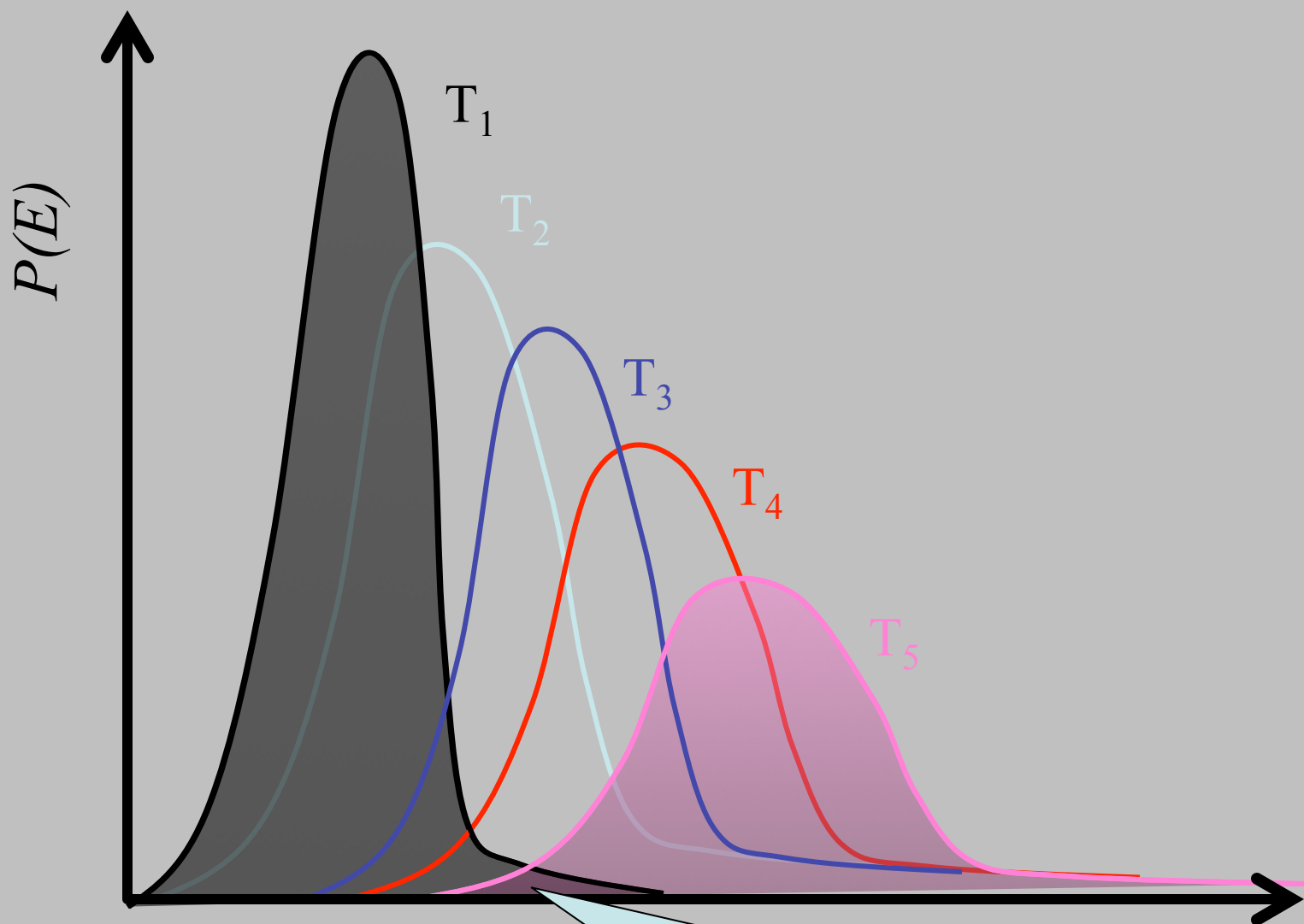






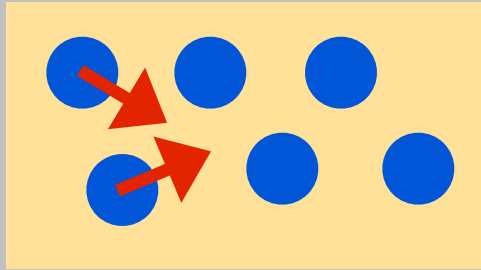




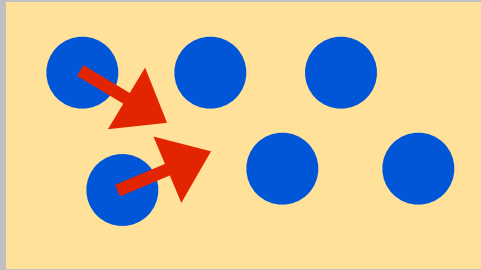


Overlap becomes very small

Parallel Monte Carlo

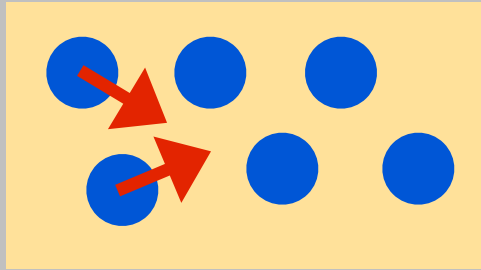


Parallel Monte Carlo



How to do a Monte Carlo simulation in parallel?

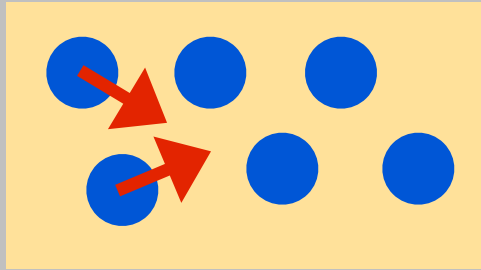
Parallel Monte Carlo



How to do a Monte Carlo simulation in parallel?

- (trivial but works best) Use an ensemble of systems with different seeds for the random number generator

Parallel Monte Carlo



How to do a Monte Carlo simulation in parallel?

- (trivial but works best) Use an ensemble of systems with different seeds for the random number generator
- Is it possible to do Monte Carlo in parallel?
 - Monte Carlo is sequential!
 - We first have to know the fait of the current move before we can continue!

Parallel Monte Carlo - algorithm

Parallel Monte Carlo - algorithm

Naive (and wrong)

Parallel Monte Carlo - algorithm

Naive (and wrong)

- 1 .Generate k trial configurations in parallel

Parallel Monte Carlo - algorithm

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1. Generate k trial configurations in parallel
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Parallel Monte Carlo - algorithm

Naive (and wrong)

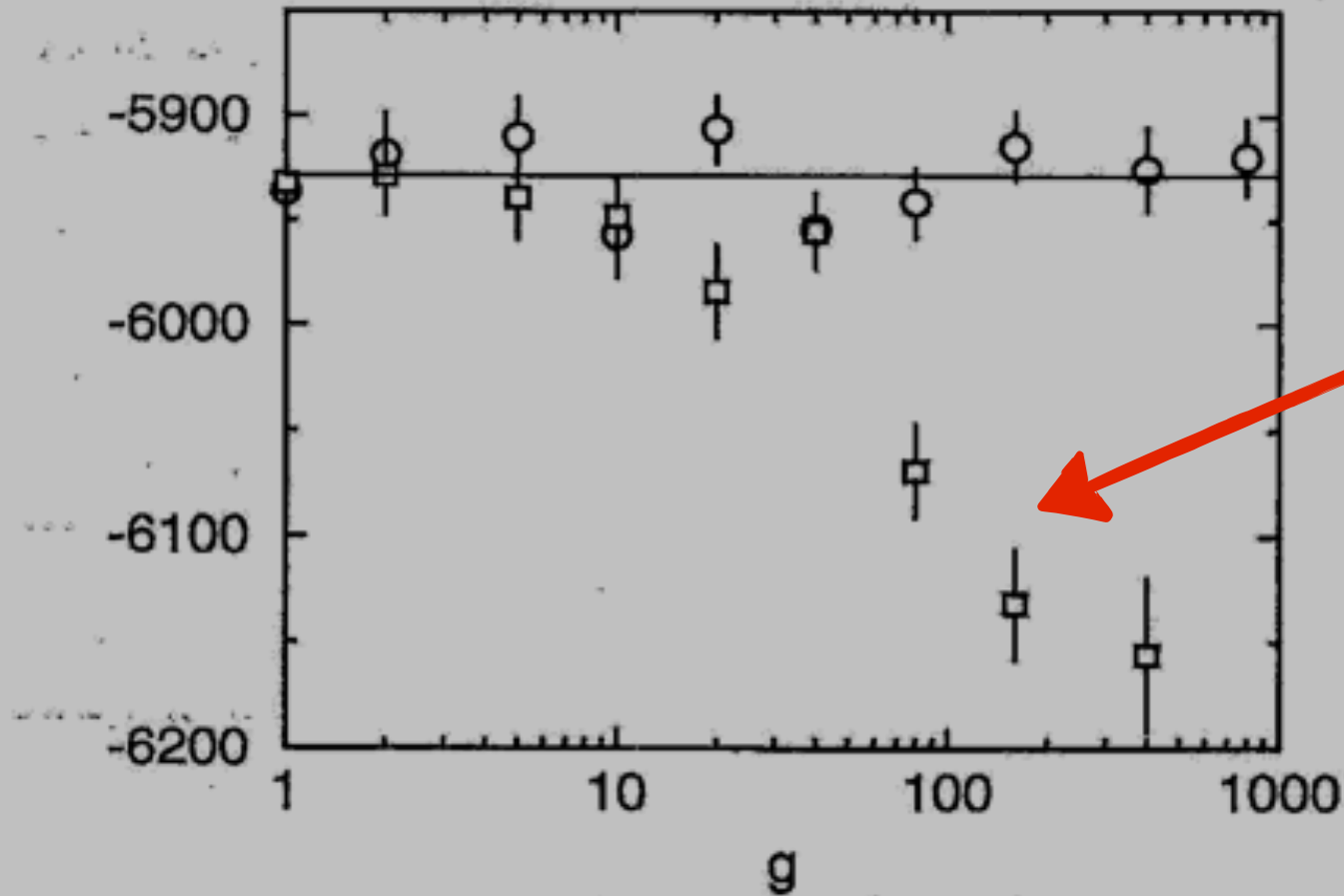
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3. Accept and reject using normal Monte Carlo rule:

$$\text{acc}(o \rightarrow n) = \exp[-\beta(U_n - U_o)]$$

Conventional acceptance rules



The conventional acceptance rules give a bias

What went wrong?

Detailed balance!

$$K(o \rightarrow n) = K(n \rightarrow o)$$

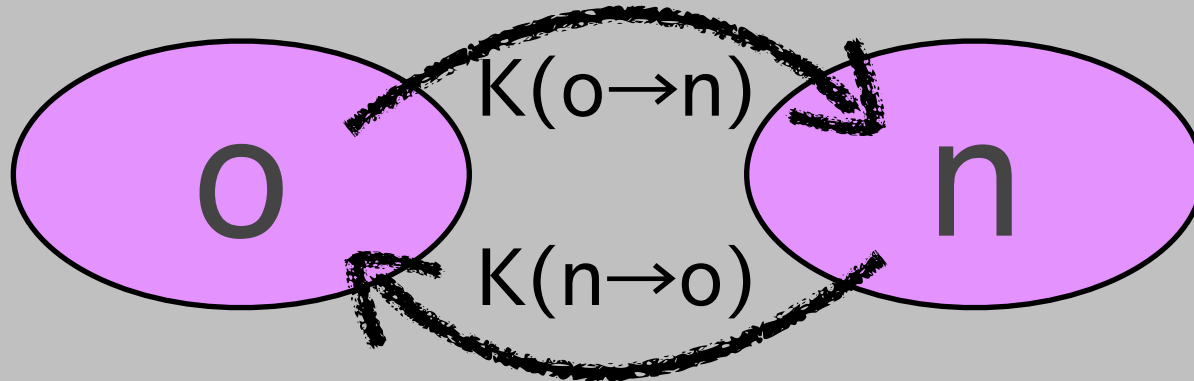
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Markov Processes - Detailed Balance



Condition of detailed balance:

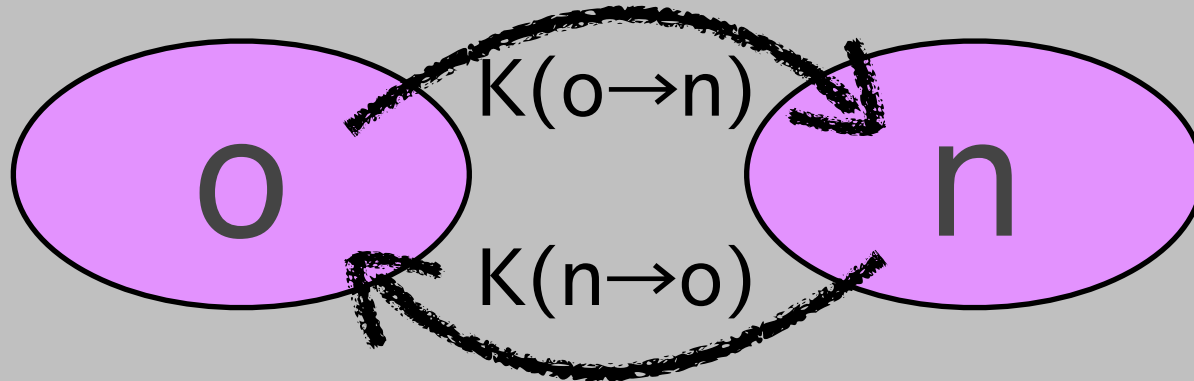
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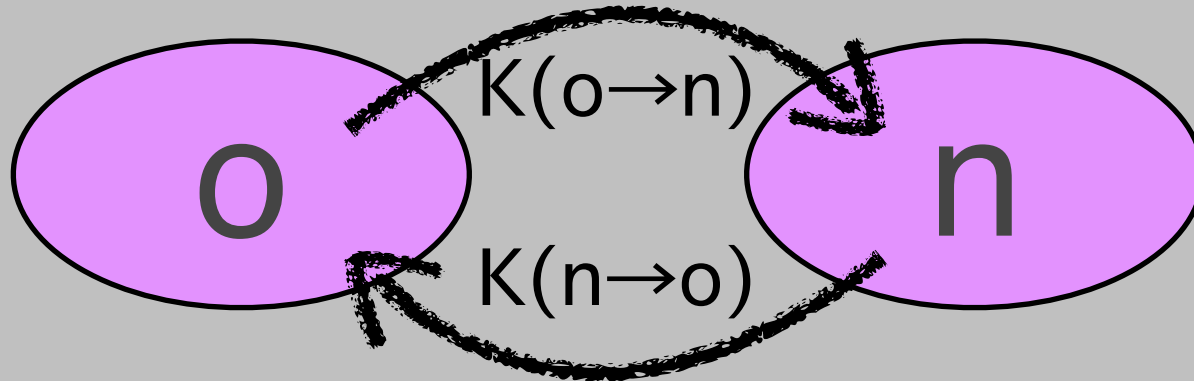
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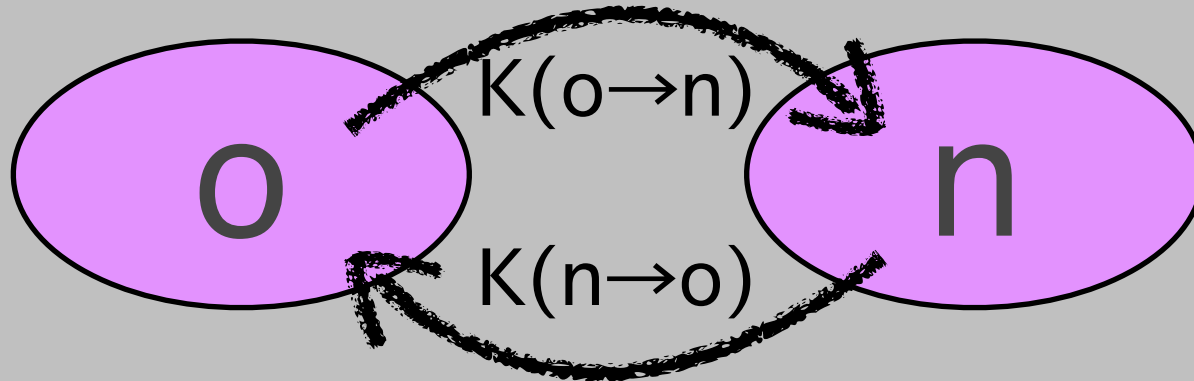
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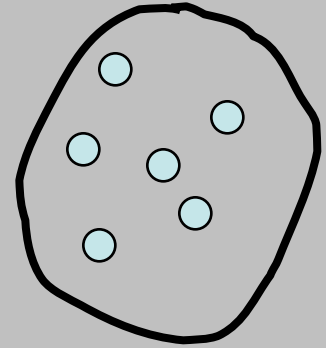
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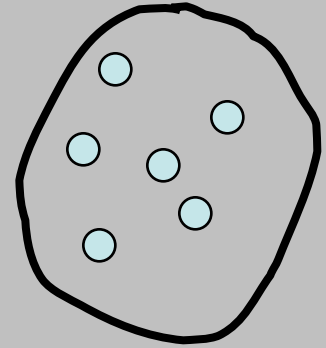
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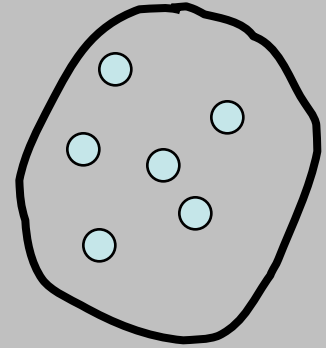
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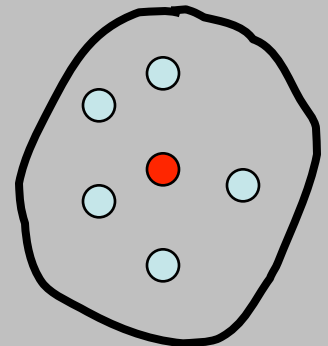
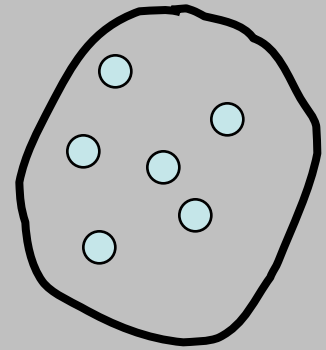
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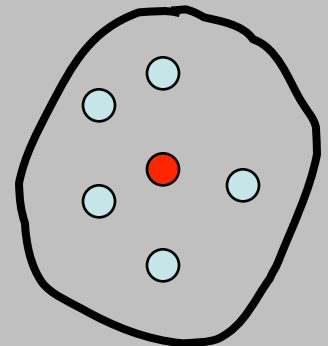
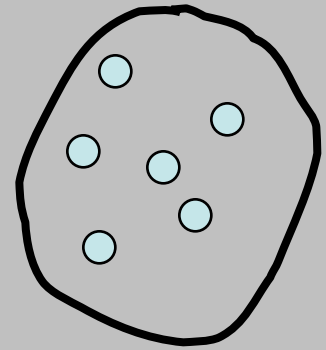


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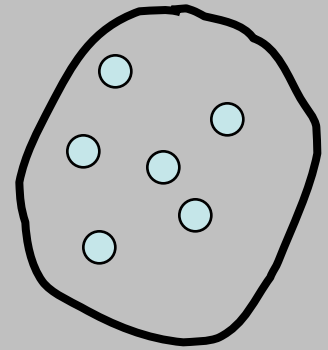
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$$\alpha(n \rightarrow o) = \frac{\exp[-\beta(U_o)]}{\sum_{j=1}^g \exp[-\beta(U_j)]}$$



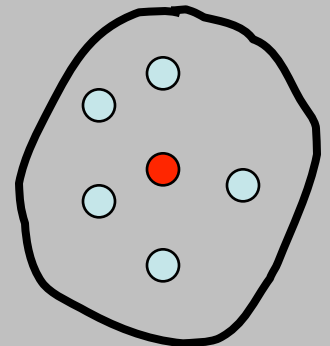
$$K(o \rightarrow n) = N(o) \times \alpha(o \rightarrow n) \times \text{acc}(o \rightarrow n)$$

$$\alpha(o \rightarrow n) = \frac{\exp[-\beta(U_n)]}{\sum_{j=1}^g \exp[-\beta(U_j)]}$$



$$\alpha(o \rightarrow n) = \frac{\exp[-\beta(U_n)]}{W(\textcolor{red}{n})}$$

$$\alpha(n \rightarrow o) = \frac{\exp[-\beta(U_o)]}{\sum_{j=1}^g \exp[-\beta(U_j)]}$$



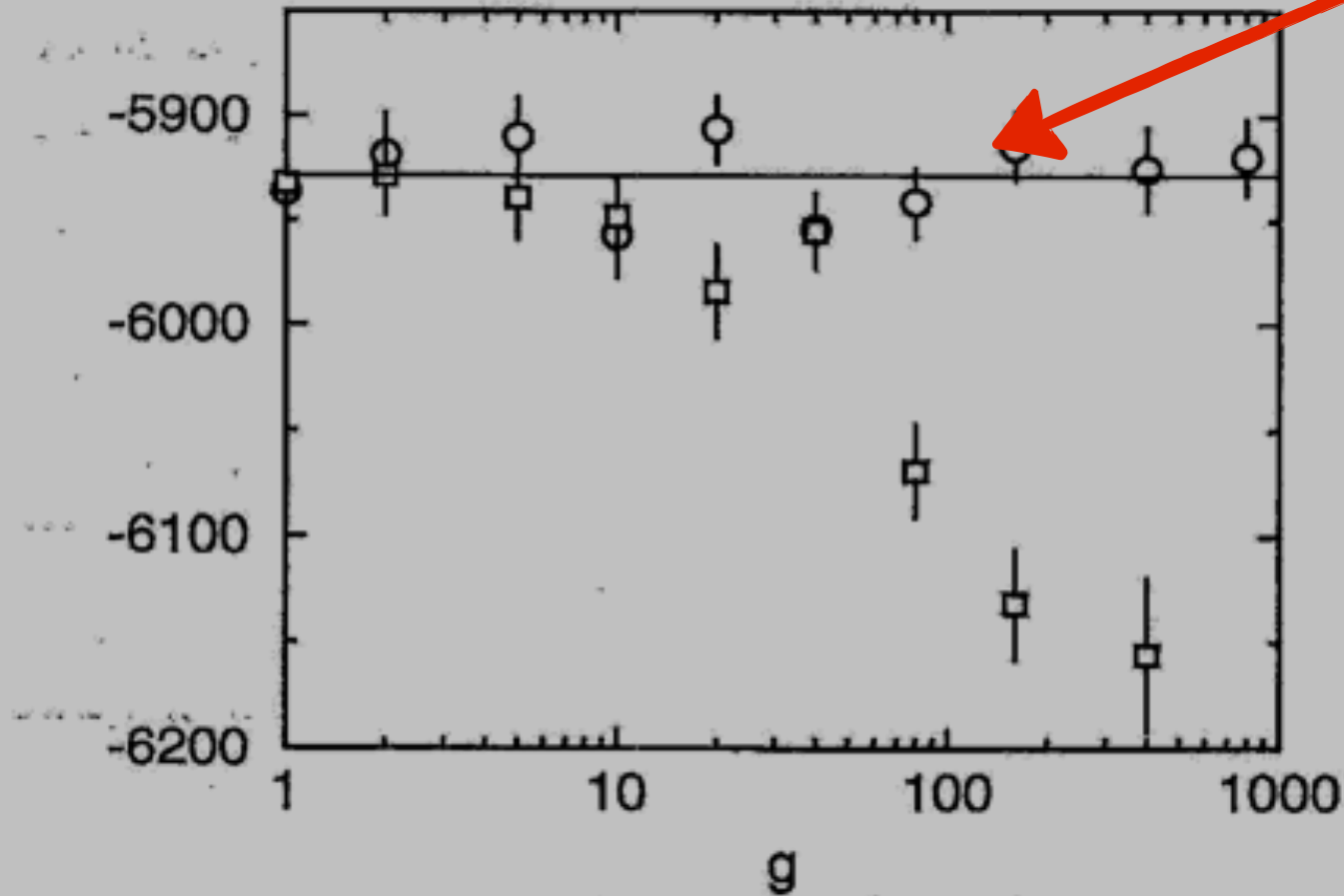
$$\alpha(n \rightarrow o) = \frac{\exp[-\beta(U_o)]}{W(\textcolor{red}{o})}$$

$$\frac{\text{acc}(o \rightarrow n)}{\text{acc}(n \rightarrow o)} = \frac{N(n) \times \alpha(n \rightarrow o)}{N(o) \times \alpha(o \rightarrow n)} = \frac{N(n)}{N(o)}$$

$$\alpha(o \rightarrow n) = \frac{\exp[-\beta(U_n)]}{W(\textcolor{red}{n})} \quad \alpha(n \rightarrow o) = \frac{\exp[-\beta(U_o)]}{W(\textcolor{red}{o})}$$

$$\frac{\text{acc}(o \rightarrow n)}{\text{acc}(n \rightarrow o)} = \frac{N(n) \times \frac{\exp[-\beta(U_o)]}{W(\textcolor{red}{o})}}{N(o) \times \frac{\exp[-\beta(U_n)]}{W(\textcolor{red}{n})}} = \frac{W(n)}{W(o)}$$

Conventional acceptance rules



Modified acceptance rules remove the bias exactly