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Lecture 7 - outline

- Importance sampling with $M(RT)^2$
- Reweighting
- Structural quantities: Density and distribution functions
- Master equation
- Fokker-Planck equation
- Smart Monte Carlo
- MD/MC code (for NVT Canonical Ensemble LJ simulations)

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Importance sampling with $M(RT)^2$

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- Sometimes, the greatest shortcoming of the Metropolis algorithm is that the usual random displacement sampled from $T(x|y)$ does not contain any information on $p(x)$, the probability density we need to sample. This leads to a higher rejection rate than necessary
- Suppose now that $p(x)$ can be factorized: $p(x)=d_1(x) \cdot d_2(x)$, and that we are already able to sample $d_1(x)$ directly, i.e., without the metropolis algorithm.
- We can try to use $d_1(x)$ in the construction of the random walk: the trial transition probability will consist in keeping $T(x|y)=d_1(x)$, what is the Metropolis acceptance of such a move?

$$A(x|y) = \min \left[1, \frac{T(y|x) \times p(x)}{T(x|y) \times p(y)} \right] = \min \left[1, \frac{d_1(y) \times d_1(x) d_2(x)}{d_1(x) \times d_1(y) d_2(y)} \right] = \min \left[1, \frac{d_2(x)}{d_2(y)} \right]$$

Thus in the "limit" $p(x)=d_1(x)$ and $d_2(x)=1$, we realize the "perfect" sampling: $A(x|y)=1$ always!

- More realistically, when $d_2(x) \neq 1$ we have reduced the computational effort in the calculation of $A(x|y)$ and being $T(x|y)$ more "similar" to $p(x)$, we have improved the efficiency in the algorithm, i.e., the acceptance $A(x|y)$ on average will grow

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Re-weighting

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- Suppose we have to compute the following series of integrals

$$I_a = \int_{\Omega} d\vec{x} g(\vec{x}) \frac{p_a(\vec{x})}{\int_{\Omega} d\vec{x} p_a(\vec{x})} \quad \text{with } p_a(\vec{x}) \geq 0 \quad \forall \vec{x} \in \Omega$$

where the probability density p_a depends on the parameter a .

- When we have computed the previous integral for a particular value of a ($a=\alpha$) via Monte Carlo by using points in \mathbb{R}^n sampled from the probability distribution $p_{a=\alpha}$, the re-weighting technique enables us to compute, in principle, the whole series of integrals I_a by using the same single set of points sampled from $p_{a=\alpha}$.
- In principle the technique is very powerful, however it has some limitations due to the variance of the calculations of the integrals I_a , for $a \neq \alpha$, which can grow in an uncontrolled way. This happens when p_a becomes too different from p_{α} for $a \neq \alpha$; in fact in this case we are implementing the "importance sampling" in a bad way thus inducing an increment of the variance in our calculation of I_a .
- One could have problems with the re-weighting technique also when p_a is "pathologic" (e.g. non continuous) as a function of the parameter a .

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- To implement the re-weighting technique, we need to know what are the new functions we have to evaluate, on the points sampled with p_α , in place of the function g . Such functions, obviously, must depend on the parameter α :

$$\begin{aligned}
 I_{a(\neq \alpha)} &= \int_{\Omega} d\vec{x} g(\vec{x}) \frac{p_a(\vec{x})}{\int_{\Omega} d\vec{x} p_a(\vec{x})} = \frac{\int_{\Omega} d\vec{x} g(\vec{x}) p_a(\vec{x})}{\int_{\Omega} d\vec{x} p_a(\vec{x})} = \\
 &= \frac{\int_{\Omega} d\vec{x} g(\vec{x}) \frac{p_a(\vec{x})}{p_\alpha(\vec{x})} p_\alpha(\vec{x})}{\int_{\Omega} d\vec{x} \frac{p_a(\vec{x})}{p_\alpha(\vec{x})} p_\alpha(\vec{x})} = \frac{\int_{\Omega} d\vec{x} [g(\vec{x}) w_a(\vec{x})] p_\alpha(\vec{x})}{\int_{\Omega} d\vec{x} w_a(\vec{x}) p_\alpha(\vec{x})} \Bigg|_{w_a(\vec{x}) = \frac{p_a(\vec{x})}{p_\alpha(\vec{x})}} = \\
 &= \frac{\int_{\Omega} d\vec{x} [g(\vec{x}) w_a(\vec{x})] \frac{p_\alpha(\vec{x})}{\int_{\Omega} d\vec{x} p_\alpha(\vec{x})}}{\int_{\Omega} d\vec{x} w_a(\vec{x}) \frac{p_\alpha(\vec{x})}{\int_{\Omega} d\vec{x} p_\alpha(\vec{x})}} \approx \frac{\frac{1}{N} \sum_{i=1}^N g(\vec{x}_i) w_a(\vec{x}_i)}{\frac{1}{N} \sum_{i=1}^N w_a(\vec{x}_i)} \Bigg|_{\vec{x}_i \text{ sampled with } p_\alpha(\vec{x})}
 \end{aligned}$$

Ratio of two Monte Carlo estimations

- Importance sampling $\Rightarrow w_a \approx 1$

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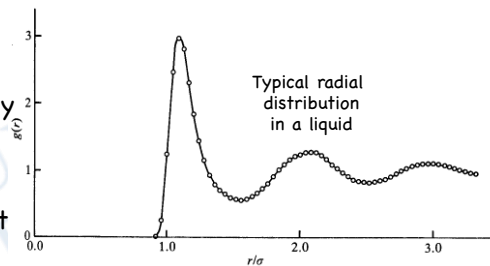
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Structural quantities

- The structure of simple monatomic fluids is characterized by a set of distribution functions for the atomic positions, the simplest of which is the **pair (or radial) distribution function** $g^{(2)}(\mathbf{r}_i, \mathbf{r}_j)$ or $g^{(2)}(r_{ij})$ or simply $g(r)$.
- This function gives the probability of finding a pair of atoms a distance r apart, relative to the probability expected for a completely random distribution at the same density.
- To define $g(r)$, we start from the **n-particles densities** which are obtained by integrating the configurational distribution function over the positions of all atoms except n , incorporating the appropriate normalization factors. In the canonical ensemble we have

$$\rho_N^{(n)}(\vec{r}_1, \dots, \vec{r}_n) = \frac{N!}{(N-n)!} \int d\vec{r}_{n+1} \dots d\vec{r}_N \frac{e^{-\beta V(\vec{r}_1, \dots, \vec{r}_N)}}{Z}$$

but of course this definition can be generalized to any configurational probability by substituting any desired $p(r_1, \dots, r_N)$ to $\exp(-\beta V)/Z$



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- The normalization implies that

$$\int d\vec{r}_1 \cdots d\vec{r}_n \rho_N^{(n)}(\vec{r}_1, \dots, \vec{r}_n) = \frac{N!}{(N-n)!} \int d\vec{r}_1 \cdots d\vec{r}_n \frac{e^{-\beta V(\vec{r}_1, \dots, \vec{r}_N)}}{Z} = \frac{N!}{(N-n)!}$$

so that $\int d\vec{r}_1 \rho_N^{(1)}(\vec{r}_1) = N$ and for an **homogeneous system** $\rho_N^{(1)}(\vec{r}_1) = \frac{N}{V}$

- For an **ideal gas** we have

$$V(\vec{r}_1, \dots, \vec{r}_N) = 0 \quad \text{and} \quad Z_N = V^N \Rightarrow \rho_N^{(n)}(\vec{r}_1, \dots, \vec{r}_n) = \frac{N! \rho^n}{N^n (N-n)!}$$

it follows that the probability to find a particle in a certain position having fixed the position of another particle in a system with N particles is

$$\rho_N^{(2)}(\text{ideal}) = \rho^2 \frac{N(N-1)}{N^2} = \rho \frac{N-1}{V}$$

which is proportional to (N-1)/V as it should

- The **n-particles distribution function is defined as**

$$g_N^{(n)}(\vec{r}_1, \dots, \vec{r}_n) = \frac{\rho_N^{(n)}(\vec{r}_1, \dots, \vec{r}_n)}{\prod_{i=1}^n \rho_N^{(1)}(\vec{r}_i)}$$

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Radial distribution function

- The **pair distribution function** is useful, not only because it provides insight into the liquid structure, but also because the ensemble average of **any pair function may be computed directly by knowing it**
- Consider for example a pair additive potential energy:

$$V(\vec{r}_1, \dots, \vec{r}_N) = \frac{1}{2} \sum_{i=1}^N \sum_{\substack{j=1 \\ (i \neq j)}}^N v(|\vec{r}_i - \vec{r}_j|)$$

- The expectation value of the potential energy is

$$\begin{aligned} \langle V \rangle &= \int d\vec{r}_1 \cdots d\vec{r}_N \frac{e^{-\beta V(\vec{r}_1, \dots, \vec{r}_N)}}{Z_N} V(\vec{r}_1, \dots, \vec{r}_N) = \dots \text{for homogeneous fluids} \dots = \\ &= \frac{1}{2} N(N-1) \int d\vec{r}_1 d\vec{r}_2 v(|\vec{r}_1 - \vec{r}_2|) \int d\vec{r}_3 \cdots d\vec{r}_N \frac{e^{-\beta V(\vec{r}_1, \dots, \vec{r}_N)}}{Z_N} = \\ &= \frac{1}{2} \int d\vec{r}_1 d\vec{r}_2 v(|\vec{r}_1 - \vec{r}_2|) \rho_N^{(2)}(\vec{r}_1, \vec{r}_2) = \frac{\rho^2}{2} \int d\vec{r}_1 d\vec{r}_2 v(|\vec{r}_1 - \vec{r}_2|) g^{(2)}(\vec{r}_1, \vec{r}_2) = \\ &= \frac{N^2}{2V^2} \int d\vec{r}_1 d\vec{r}_2 v(|\vec{r}_1 - \vec{r}_2|) g(|\vec{r}_1 - \vec{r}_2|) = \frac{N^2}{2V} \int d\vec{r} v(r) g(r) = \\ &= 2\pi\rho N \int_0^{+\infty} dr r^2 v(r) g(r) \end{aligned}$$

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Tail corrections

- Thus the knowledge of the radial distribution function $g(r)$ in principle give access to the expectation value of any pair function of the coordinates. See also the relevant [connection with a scattering experiment](#) in the supplementary material: $g(r)$ can be indirectly measured via such kind of experiments
- Computer simulations frequently use a potential with a **spherical cutoff** at a distance r_c . It becomes useful to **correct** the results of simulations **to compensate for the missing long-range part of the potential**
- Contributions to the energy, pressure etc. for $r > r_c$ are frequently estimated by assuming that $g(r) \approx 1$ in this region:

$$\langle V \rangle_{tail} = 2\pi\rho N \int_{r_c}^{+\infty} dr r^2 v(r) g(r) \approx 2\pi\rho N \int_{r_c}^{+\infty} dr r^2 v(r)$$

- For example, the Lennard-Jones potential gives

$$\begin{aligned} \frac{\langle V_{LJ} \rangle_{tail}}{N} &\approx 2\pi\rho \int_{r_c}^{+\infty} dr r^2 4\epsilon \left[\left(\frac{\sigma}{r} \right)^{12} - \left(\frac{\sigma}{r} \right)^6 \right] = 8\pi\rho\epsilon \left[-\frac{\sigma^{12}}{9r^9} + \frac{\sigma^6}{3r^3} \right]_{r_c}^{+\infty} = \\ &= 8\pi \left(\frac{\rho}{\sigma^3} \right) \epsilon \left[-\frac{1}{9} \left(\frac{\sigma}{r} \right)^9 + \frac{1}{3} \left(\frac{\sigma}{r} \right)^3 \right]_{r_c}^{+\infty} = 8\pi\rho_{LJ}\epsilon \left[\frac{1}{9} \left(\frac{\sigma}{r_c} \right)^9 - \frac{1}{3} \left(\frac{\sigma}{r_c} \right)^3 \right] = \frac{8\pi\rho_{LJ}\epsilon}{9r_{c(LJ)}^9} - \frac{8\pi\rho_{LJ}\epsilon}{3r_{c(LJ)}^3} \end{aligned}$$

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Tail corrections for pressure

- We thus can compute the tail correction to the pressure:

$$\begin{aligned} \frac{\langle w_{LJ} \rangle_{tail}}{3N\epsilon} &\approx \frac{2}{3}\pi\rho \int_{r_c}^{+\infty} dr r^2 48 \left[\left(\frac{\sigma}{r} \right)^{12} - \frac{1}{2} \left(\frac{\sigma}{r} \right)^6 \right] = 32\pi\rho \left[-\frac{\sigma^{12}}{9r^9} + \frac{\sigma^6}{6r^3} \right]_{r_c}^{+\infty} = \\ &= 32\pi \left(\frac{\rho}{\sigma^3} \right) \left[-\frac{1}{9} \left(\frac{\sigma}{r} \right)^9 + \frac{1}{6} \left(\frac{\sigma}{r} \right)^3 \right]_{r_c}^{+\infty} = 32\pi\rho_{LJ} \left[\frac{1}{9} \left(\frac{\sigma}{r_c} \right)^9 - \frac{1}{6} \left(\frac{\sigma}{r_c} \right)^3 \right] = \\ &= 32\pi\rho_{LJ} \left[\frac{1}{9r_{c(LJ)}^9} - \frac{1}{6r_{c(LJ)}^3} \right] \end{aligned}$$

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g(r) : the algorithm

- We have to deduce an algorithm for the calculation of the radial distribution function g(r). We observe that

$$\begin{aligned}\langle \delta(\vec{r} - \vec{r}_1) \rangle &= \int \delta(\vec{r} - \vec{r}_1) \frac{e^{-\beta V(\vec{r}_1, \dots, \vec{r}_N)}}{Z_N} d\vec{r}_1 \dots d\vec{r}_N = \int \frac{e^{-\beta V(\vec{r}, \vec{r}_2, \dots, \vec{r}_N)}}{Z_N} d\vec{r}_2 \dots d\vec{r}_N \\ \Rightarrow \rho_N^{(1)}(\vec{r}) &= N \int \frac{e^{-\beta V(\vec{r}, \vec{r}_2, \dots, \vec{r}_N)}}{Z_N} d\vec{r}_2 \dots d\vec{r}_N = \left\langle \sum_{i=1}^N \delta(\vec{r} - \vec{r}_i) \right\rangle\end{aligned}$$

this is an algorithm for the **one-particle density**.

For the **two-particles density** we see that

$$\begin{aligned}\langle \delta(\vec{r} - \vec{r}_1) \delta(\vec{r}' - \vec{r}_2) \rangle &= \int \frac{e^{-\beta V(\vec{r}, \vec{r}', \vec{r}_3, \dots, \vec{r}_N)}}{Z_N} d\vec{r}_3 \dots d\vec{r}_N \\ \Rightarrow \rho_N^{(2)}(\vec{r}, \vec{r}') &= N(N-1) \int \frac{e^{-\beta V(\vec{r}, \vec{r}', \vec{r}_3, \dots, \vec{r}_N)}}{Z_N} d\vec{r}_3 \dots d\vec{r}_N = \left\langle \sum_{i \neq j=1}^N \delta(\vec{r} - \vec{r}_i) \delta(\vec{r}' - \vec{r}_j) \right\rangle\end{aligned}$$

- In a **homogeneous liquid** where $g^{(2)}(\mathbf{r}_1, \mathbf{r}_2)$ is simply a function of the modulus of the difference between the two vectors \mathbf{r}_1 and \mathbf{r}_2 :
 $g^{(2)}(\mathbf{r}_1, \mathbf{r}_2) = g^{(2)}(|\mathbf{r}_1 - \mathbf{r}_2|) = g(r)$

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it is not the case to compute directly the one-particle and the two-particles densities to compute g(r) as their ratio, instead we observe

$$\begin{aligned}\left\langle \frac{1}{N} \sum_{i \neq j=1}^N \delta(\vec{r} + \vec{r}_j - \vec{r}_i) \right\rangle &= \left\langle \frac{1}{N} \int \sum_{i \neq j=1}^N \delta(\vec{r} + \vec{r}' - \vec{r}_i) \delta(\vec{r}' - \vec{r}_j) d\vec{r}' \right\rangle = \\ &= \frac{1}{N} \int \rho_N^{(2)}(\vec{r} + \vec{r}', \vec{r}') d\vec{r}' = \frac{1}{N} \int \rho_N^{(2)}(|\vec{r} + \vec{r}' - \vec{r}'|) d\vec{r}' = \frac{1}{\rho} \rho_N^{(2)}(|\vec{r}|) = \rho g(r) \\ \Rightarrow g(\vec{r}) &= \frac{1}{\rho N} \left\langle \sum_{i \neq j=1}^N \delta[\vec{r} - (\vec{r}_i - \vec{r}_j)] \right\rangle \\ \Rightarrow g(r) &= \left\langle g(\vec{r}) \right\rangle_{\substack{r \in [r, r+dr] \\ \theta \in [0, \pi] \\ \varphi \in [0, 2\pi]}} = \frac{1}{\rho N \Delta V(r)} \left\langle \sum_{i \neq j=1}^N \delta[|\vec{r}| - |\vec{r}_i - \vec{r}_j|] \right\rangle\end{aligned}$$

$$\text{with } \Delta V(r) = \frac{4\pi}{3} [(r+dr)^3 - r^3]$$

- In practice, during the simulation, one should "fill" an histogram by increasing the bin by 2 at r whenever he finds two particles at distance between r and r+dr. At the end of the simulation he should normalize the histogram with the quantity $\rho N \Delta V(r)$.

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Derivation of the Master equation

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- Let's go back to a **Markov process** which, given two positive normalized functions p_1 and $p_{1|1}$, is fully characterized by
 - The **Chapman-Kolmogorov** equation

$$p_{1|1}(x_3, t_3 | x_1, t_1) = \int p_{1|1}(x_3, t_3 | x_2, t_2) p_{1|1}(x_2, t_2 | x_1, t_1) dx_2 \quad \text{for } t_3 \geq t_2 \geq t_1$$
 - The equation:

$$p_1(x_2, t_2) = \int p_{1|1}(x_2, t_2 | x_1, t_1) p_1(x_1, t_1) dx_1$$
- According to these equations, the **conditional probability** is the crucial quantity determining in a Markov process the **evolution with time of the probability distribution** $p_1(x, t)$
- Let us consider the last equation for times t and $t+\tau$, where τ is an infinitesimally short time interval. Proceeding in this way, we obtain the short-time evolution equation

$$p_1(x, t + \tau) = \int dy p_{1|1}(x, t + \tau | y, t) p_1(y, t)$$

- Next we can construct time derivatives of p_1

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$$\frac{\partial p_1(x, t)}{\partial t} = \lim_{\tau \rightarrow 0} \frac{p_1(x, t + \tau) - p_1(x, t)}{\tau} =$$

$$= \lim_{\tau \rightarrow 0} \frac{1}{\tau} \left[\int p_{1|1}(x, t + \tau | y, t) p_1(y, t) dy - p_1(x, t) \right] = \dots$$

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- Since we will take the limit $\tau \rightarrow 0$, we can expand the transition probability $p_{1|1}$ in a power series in τ and keep only the lowest-order term. In order to conserve probability at all times, its most general form is:

$$p_{1|1}(x, t + \tau | y, t) \approx_{\tau \rightarrow 0} \delta(y - x) \left[1 - \tau \int \frac{\partial p_{1|1}(z, t' | y, t)}{\partial t'} \Big|_{t'=t} dz \right] + \tau \frac{\partial p_{1|1}(x, t' | y, t)}{\partial t'} \Big|_{t'=t}$$

probability that no transition occurs during the time interval $t \rightarrow t+\tau$

probability of a transition from y to x during the time interval $t \rightarrow t+\tau$

$$p_{1|1}(x, t + \tau | y, t) \approx_{\tau \rightarrow 0} \delta(y - x) \left[1 - \tau \int dz R_t(z | x) \right] + \tau R_t(x | y)$$

transition rates

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- If we now substitute back p_{11} we obtain

$$\begin{aligned}\frac{\partial p_1(x, t)}{\partial t} &= \lim_{\tau \rightarrow 0} \frac{1}{\tau} \left\{ \int \left[\delta(y - x) \left[1 - \tau \int dz R_t(z|x) \right] + \tau R_t(x|y) \right] p_1(y, t) dy - p_1(x, t) \right\} = \\ &= \int dy p_1(y, t) \left[R_t(x|y) - \delta(y - x) \int dz R_t(z|x) \right] = \int dy R_t(x|y) p_1(y, t) - \int dz R_t(z|x) p_1(x, t)\end{aligned}$$

- Which lead us to the **Master equation**:

$$\frac{\partial p(x, t)}{\partial t} = \int dy \left[R_t(x|y) p(y, t) - R_t(y|x) p(x, t) \right]$$

- The quantities $R_t(x|y)$ are transition probabilities per time unit or **transition rates**. The Master equation can be read as a balance equation for probability flows: **the change per unit of time of the probability of state x is the sum of two terms with opposite effects**. Firstly, there is a probability flux from all other states y into the state x. Secondly, there is a probability flux out of state x into all other states y. The change per time unit of the probability p is caused by the difference of those probability fluxes.
- Discrete Master equation:

$$\frac{\partial p_n(t)}{\partial t} = \sum_m \left[R_t^{nm} p_m(t) - R_t^{mn} p_n(t) \right]$$

- The **master equation** is thus an equivalent form of the Chapman-Kolmogorov equation for Markov processes, but it is easier to handle and **more directly related to physical concepts**.
- In the **master equation** the quantities representing the knowledge about the system are the **transition rates**. One considers them as given by **the specific system**, and then has an equation for the probabilities which **determine the statistical evolution** of that system under conditions of uncertainty or restricted information.
- In **social sciences**, the transition rates can often be inferred from phenomenological considerations based on plausibility arguments.
- In **natural sciences**, fundamental considerations may be the starting point in the construction of the transition rates or the transition rates may be derived from first principles. Given the transition rates, the master equation is an instrument to model equilibrium and out of equilibrium statistical systems
- Let's make a step further and derive another relevant equation connected with a Markov process ...

Kramers-Moyal expansion

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Let us now set the stage for a further approximation of the master equation

$$\frac{\partial p(x, t)}{\partial t} = \int dy \left[R_+(x|y) p(y, t) - R_-(y|x) p(x, t) \right]$$

and make the following assumptions:

- First express the transition probability R_+ as a function of the size of the jumps and of the starting point:
we write $R_+(x|y) = R_+(x|x-r) =: R_+(x-r; r)$, with the 'jump distance' $r := x-y$. Similarly, $R_-(y|x) = R_-(x-r|x) =: R_-(x; -r)$.

- There are only small jumps, i.e., $R_+(x-r; r)$ as a function of r is a sharply peaked function around $r=0$:

$$\exists \delta > 0 : R_+(x-r; r) \approx 0, \quad |r| > \delta$$

- $R_+(x-r; r)$ is a slowly varying function of its first argument

$$\exists \delta' > 0 : R_+(x-r; r) \approx R_+(x; r), \quad |r| < \delta'$$

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- Moreover R_+ and p are sufficiently smooth functions of both arguments.
- We rewrite the Master equation as

$$\frac{\partial p(x, t)}{\partial t} = \int dr R_+(x-r; r) p(x-r, t) - p(x, t) \int dr R_-(x; -r)$$

- We can now perform a Taylor expansion in $x-r$ around $r=0$ in the first integral on the right-hand side:

$$\frac{\partial p(x, t)}{\partial t} = \int dr \left\{ R_+(x|r) p(x, t) - r \frac{\partial}{\partial x} [R_+(x|r) p(x, t)] + \frac{1}{2} r^2 \frac{\partial^2}{\partial x^2} [R_+(x|r) p(x, t)] + \dots \right\} +$$

Yelding

$$\frac{\partial p(x, t)}{\partial t} = \sum_{n=1}^{\infty} \frac{(-1)^n}{n!} \frac{\partial^n}{\partial x^n} \left[a_n(x) p(x, t) \right] - p(x, t) \int dr R_-(x; -r)$$

$$a_n(x) = \int_{-\infty}^{+\infty} dr r^n R_+(x; r)$$

- Which is the Kramers-Moyal expansion of the master equation. As such it is only a rewriting of the master equation, changing it from an integro-differential equation into a partial differential equation of infinite order.

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Fokker-Planck Equation

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- In deriving the Kramers-Moyal expansion, we have not yet made use of our assumption that the transition rates $R_i(x, r)$ are supposed to be slowly varying functions of their first argument. For these **slowly varying functions** (which will give rise to slowly varying probability densities $p(x, t)$) we can make the **assumption that we can truncate the Kramers-Moyal expansion at a certain order of derivatives**.
- A truncation after the second order leads to the so-called **Fokker-Planck equation**:

$$\frac{\partial p(x, t)}{\partial t} = -\frac{\partial}{\partial x} \left[p(x, t) \int dr r R_t(x; r) \right] + \frac{1}{2} \frac{\partial^2}{\partial x^2} \left[p(x, t) \int dr r^2 R_t(x; r) \right]$$

$$\frac{\partial p(x, t)}{\partial t} = -\frac{\partial}{\partial x} \left[a_1(x) p(x, t) \right] + \frac{1}{2} \frac{\partial^2}{\partial x^2} \left[a_2(x) p(x, t) \right]$$

a_1 is called the **drift coefficient** and a_2 the **diffusion coefficient**.

- Note that $[a_n(x)\Delta t]$ are the n^{th} moment of the "distribution of step size in time Δt " (which depends on the starting point x)

$$a_n(x)\Delta t = \int_{-\infty}^{+\infty} dr r^n R_t(x; r)\Delta t$$

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Brownian motion: again Diffusion

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- In fact, let's go back to the **Brownian motion** (without drift) introduced in Lecture 3. Its coordinate x may be treated on a coarse time scale as a Markov process:

$$x(t + \Delta t) = x(t) + \sigma Z \sqrt{\Delta t} \quad \text{being } Z \sim \mathcal{N}(0; 1)$$

- Accordingly we have the picture of a particle that makes random jumps back and forth over the x -axis. The jumps may have any length, but the probability for large jumps falls off rapidly. Moreover the probability is symmetrical and independent of the starting point. Hence

$$a_1 = \int_{-\infty}^{+\infty} dr r R_t(x; r) = 0 \quad a_2 \Delta t = \int_{-\infty}^{+\infty} dr r^2 R_t(x; r) \Delta t = \sigma^2 \Delta t$$

- The Fokker-Planck equation for $p(x, t)$ is therefore the **diffusion equation**:

$$\frac{\partial p(x, t)}{\partial t} = \frac{a_2}{2} \frac{\partial^2}{\partial x^2} p(x, t) \quad \Rightarrow \quad a_2 = \sigma^2 = 2D$$

Consequently $a_2/2$ correspond to the diffusion constant D

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Drift force & drift velocity

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- We see that respect to the simpler Diffusion equation, the Fokker-Planck equation contains the possible presence of an **external drift force**.
- We now want to make a **more general contact between the Fokker-Planck equation** for the probability density of a Markov process and the **equation governing the time development of the sample path $x(t)$** of this process considering the addition of a drift velocity term to the Wiener process.
- A feasible form for this **new term** can be derived assuming that (think about the Brownian motion) in presence of an external drift force F and a **finite viscosity, η , of the dense fluid** the the Brownian particle will rapidly acquire a **drift velocity**:

$$m\ddot{x} = F - k\eta v \quad \ddot{x} = 0 \Rightarrow v_{\infty} = \frac{1}{k\eta} F = -\frac{1}{k\eta} \frac{dU}{dx} = \mu$$

- By adding a drift velocity term to the **Brownian motion** we obtain

$$x(t + \Delta t) = x(t) + \mu \Delta t + Z\sigma\sqrt{\Delta t} \quad \text{being } Z \sim \mathcal{N}(0;1)$$

as already discussed in lecture 3, and $a_1 \Delta t = \mu \Delta t$.

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- Let's go back now to the Metropolis algorithm and consider cases where the efficiency of the sampling obtained by proposing random moves via simple $T(x|y)$ is too much inefficient because, for example, the target joint distribution is different from zero only on a intricate domain. Have we other possibilities to propose "intelligent" trial moves in these cases? Yes, via what is called **"smart Monte Carlo" or "Langevin Monte Carlo"**.
- This different approach includes **importance sampling** based on the **Fokker-Planck equation**: Consider a **diffusion process** characterized by a time-dependent probability density $p(x,t)$.
- It is known that the probability density of a diffusion processes with drift obeys the following **multivariate Fokker-Planck equation**:

$$\frac{\partial p(\vec{x}, t)}{\partial t} = \sum_i \frac{\partial}{\partial x_i} \left[D \frac{\partial}{\partial x_i} p(\vec{x}, t) - \gamma F_i(\vec{x}) p(\vec{x}, t) \right]$$

where D is the **diffusion constant** and F_i is the **i -th component of a drift force** caused by an external potential.

- As usual, we wish to converge to the stationary probability density $p(x)$

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- An **unchanging probability**, for which $\partial p / \partial t = 0$, may be obtained by setting the left-hand side of the previous equation to zero, namely

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$$\sum_i D \left\{ \frac{\partial^2 p(\vec{x})}{\partial x_i^2} - \frac{\gamma}{D} \frac{\partial}{\partial x_i} [F_i(\vec{x}) p(\vec{x})] \right\} = 0$$

- This equation can most readily be satisfied if **each term of the previous sum vanishes**, yielding

$$\frac{\partial^2 p(\vec{x})}{\partial x_i^2} = \frac{\gamma}{D} \left[p(\vec{x}) \frac{\partial}{\partial x_i} F_i(\vec{x}) + F_i(\vec{x}) \frac{\partial}{\partial x_i} p(\vec{x}) \right]$$

- The **drift velocity F** therefore must be of the form $F_i(\vec{x}) = \frac{D}{\gamma} g(p) \frac{\partial p(\vec{x})}{\partial x_i}$ in order to obtain a second derivative of p on the right-hand side. If we substitute this form of F into the previous equation we find

$$\frac{\partial^2 p}{\partial x_i^2} = p \frac{\partial g}{\partial p} \left(\frac{\partial p}{\partial x_i} \right)^2 + p g \frac{\partial^2 p}{\partial x_i^2} + g \left(\frac{\partial p}{\partial x_i} \right)^2$$

- **Cancellation of the second derivative terms requires** that $g=1/p$. This choice **also leads to cancellation of the first derivative terms**.

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- Therefore the stationary probability density $p(x)$ results from choosing the drift vector to be:

$$\vec{F}(\vec{x}) = \frac{D}{\gamma} g(p) \vec{\nabla} p(\vec{x}) = \frac{D}{\gamma} \frac{1}{p(\vec{x})} \vec{\nabla} p(\vec{x}) = \frac{D}{\gamma} \vec{\nabla} \ln p(\vec{x})$$

Clearly **this drift causes the move to be "biased" by $p(x)$** . This biased diffusion process incorporates importance sampling.

- We now have a diffusion equation which gives the desired probability distribution, but **how do we implement it using Monte Carlo?**
- We have seen that Fokker-Planck trajectories are generated by means of Brownian motion with drift:

$$\vec{x}(t + \Delta t) = \vec{x}(t) + D \Delta t \vec{\nabla} \ln p(\vec{x}) + \vec{\chi}$$

here $\vec{\chi}$ is a randomly fluctuating displacement which is distributed according to a **multi-dimensional Gaussian with a mean of zero and a variance of $2D\Delta t$** , i.e., proportional to what is called a multivariate Gaussian white noise.

- By using a discretized finite time step we have introduced a **bias** into the stochastic dynamics for any $\Delta t > 0$ due to the approximations introduced to derive the Fokker-Planck equation.

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Smart Monte Carlo method

- Do we have a method to correct this bias and obtain the correct sampling of $p(x)$? Yes, again the Metropolis!
- In fact, we have obtained a dynamical equation for generating guided trajectories containing a drift force: $F(x) = (D/\gamma) \nabla \ln p(x)$. This dynamical equation directs "walkers" toward regions of large $p(x)$, but suffers by being exact only at $\Delta t = 0$. We will now show that by combining this equation with a Metropolis algorithm the time step bias can be eliminated.
- From the generalized Metropolis method recall that the complete transition probability $K(x|y)$ is composed by a trial transition probability $T(x|y)$ times an acceptance probability $A(x|y)$ of the proposed move $y \rightarrow x$. We can use this approach, and choose the $T(x|y)$ which corresponds to the previously discretized Langevin equation.
- In order to be able to do this, we have to find the functional form of $T(x|y)$ which corresponds to the previous Brownian motion difference equation with the added condition that $T(x|y, \Delta t=0) = \delta(x-y)$

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- We realize that if χ is a multivariate Gaussian random variable with zero mean and variance $2D\Delta t$, also

$$\vec{x} - \vec{y} - (D\Delta t) \vec{\nabla} \ln p(\vec{y}) = \vec{\chi}$$

is such a Gaussian random variable; therefore we can choose

$$T(\vec{x}|\vec{y}) \propto e^{-\frac{\{\vec{x} - \vec{y} - (D\Delta t) \vec{\nabla} \ln p(\vec{y})\}^2}{4D\Delta t}}$$

Thus χ is a Gaussian random variable with mean $y + (D\Delta t) \nabla \ln p(y)$ and variance $2D\Delta t$; we have already learned how to sample such Gaussian random variables via the Box-Muller method.

- Note that in general, being $T(x|y)$ different from $T(y|x)$, for the Metropolis acceptance probability we need the calculation of :

$$\frac{T(\vec{y}|\vec{x})}{T(\vec{x}|\vec{y})} = \frac{e^{-\frac{\{\vec{y} - \vec{x} - D\Delta t \vec{\nabla} \ln p(\vec{x})\}^2}{4D\Delta t}}}{e^{-\frac{\{\vec{x} - \vec{y} - D\Delta t \vec{\nabla} \ln p(\vec{y})\}^2}{4D\Delta t}}}$$

- $(D\Delta t)$ turns out to be an adjustable parameter to force $A(x|y) \approx 50\%$

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Lecture 7: Suggested books

- M.P. Allen and D.J. Tildesley, *Computer Simulation of Liquids* – Clarendon Press
- D. Frenkel and B. Schmidt, *Understanding Molecular Simulation, From Algorithms to Applications* – Academic Press
- L. Hammond, W. A. Lester, Jr., P. J. Reynolds, *Monte Carlo Methods in Ab Initio Quantum Chemistry* – World Scientific (1994)
- W. Paul, J. Baschnagel, *Stochastic Processes. From Physics to Finance*, Springer (2000)
- N.G. Van Kampen, *Stochastic Processes in Physics and Chemistry*, Elsevier (2007)

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Numerical Simulation Laboratory

MD/MC code
(the same introduced in Lecture 4)

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Molecular Dynamics/Monte Carlo code: `main`

```
#include <iostream>
#include <fstream>
#include <ostream>
#include <cmath>
#include <iomanip>
#include "MD_MC.h"

using namespace std;

int main(){
    Input(); //Inizialization
    int nconf = 1;
    for(int iblk=1; iblk <= nblk; iblk++){ //Data blocking
        Reset(iblk); //Reset block averages
        for(int istep=1; istep <= nstep; istep++){
            Move(); //Move part with Verlet or Metropolis algorithm
            Measure(); //Mesure properties
            Accumulate(); //Update block averages
            if(istep%10 == 0){
                ConfXYZ(nconf); //Write conf in XYZ format; ommented to avoid "filesystem full"!
                nconf += 1;
            }
        }
        Averages(iblk); //Print results for current block
    }
    ConfFinal(); //Write final configuration
    return 0;
}
```

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MD/MC code: `Input ()`

```
#include <iostream>
#include <fstream>
#include <ostream>
#include <cmath>
#include <iomanip>
#include "MD_MC.h"

using namespace std;

int main(){
    Input(); //Inizialization
    int nconf = 1;
    for(int iblk=1; iblk <= nblk; iblk++){ //Data blocking
        Reset(iblk); //Reset block average
        for(int istep=1; istep <= nstep; istep++){
            Move(); //Move part with Verlet
            Measure(); //Mesure properties
            Accumulate(); //Update block averages
            if(istep%10 == 0){
                ConfXYZ(nconf); //Write conf in XYZ format; ommented
                nconf += 1;
            }
        }
        Averages(iblk); //Print results for current block
    }
    ConfFinal(); //Write final configuration
    return 0;
}
```

`Input.dat`

```
1      <- for MC run
0
1.1
108
0.8
2.5
0.2    <- delta Metropolis
20
2000

ReadInput >> iNVET; 0=MD(NVE)
ReadInput >> restart;
ReadInput >> temp;
ReadInput >> npart;
ReadInput >> rho;
ReadInput >> rcut;
ReadInput >> delta;
ReadInput >> nblk;
ReadInput >> nstep;
```

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MD/MC code: Move ()

```
#include <iostream>
#include <fstream>
#include <ostream>
#include <cmath>
#include <iomanip>
#include "MD_MC.h"

using namespace std;

int main(){
    Input(); //Inizialization
    int nconf = 1;
    for(int iblk=1; iblk <= nblk; iblk++){ //Data blocking
        Reset(iblk); //Reset block averages
        for(int istep=1; istep <= nstep; istep++){
            Move(); //Move part with Verlet or Metropolis algorithm
            Measure(); //Mesure properties
            Accumulate(); //Update block averages
            if(istep%10 == 0){
                ConfXYZ(nconf); //Write conf in XYZ format; ommented to avoid "filesystem full"!
                nconf += 1;
            }
        }
        Averages(iblk); //Print results for current block
    }
    ConfFinal(); //Write final configuration
    return 0;
}
```

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MD/MC code: Move ()

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```
void Move(){
    int o;
    double p, energy_old, energy_new;
    double xnew, ynew, znew;
    if(iNVET){ // Monte Carlo (NVT) move
        for(int i=0; i<npart; ++i){
            //Select randomly a particle
            o = (int)(rnd.Rannyu()*npart);
            //Old
            energy_old = Boltzmann(x[o],y[o],z[o],o);
            //New
            x[o] = Pbc( x[o] + delta*(rnd.Rannyu() - 0.5) );
            y[o] = Pbc( y[o] + delta*(rnd.Rannyu() - 0.5) );
            z[o] = Pbc( z[o] + delta*(rnd.Rannyu() - 0.5) );
            energy_new = Boltzmann(x[o],y[o],z[o],o);
            //Metropolis test
            p = exp(beta*(energy_old-energy_new));
            if(p >= rnd.Rannyu()){ //Update
                xold[o] = x[o];
                yold[o] = y[o];
                zold[o] = z[o];
                accepted = accepted + 1.0;
            } else {
                x[o] = xold[o];
                y[o] = yold[o];
                z[o] = zold[o];
                attempted = attempted + 1.0;
            }
        }
    } else { // Molecular Dynamics (NVE) move
        //...
    }
    return;
}
```

```
double Boltzmann(double xx, double yy,
double zz, int ip)
{
    double ene=0.0;
    double dx, dy, dz, dr;
    for (int i=0; i<npart; ++i)
    {
        if(i != ip)
        {
            // distance ip-i in pbc
            dx = Pbc(xx - x[i]);
            dy = Pbc(yy - y[i]);
            dz = Pbc(zz - z[i]);

            dr = dx*dx + dy*dy + dz*dz;
            dr = sqrt(dr);

            if(dr < rcut)
            {
                ene += 1.0/pow(dr,12)-1.0/pow(dr,6);
            }
        }
    }
    return 4.0*ene;
}
```

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MD/MC code: Measure ()

```
#include <iostream>
#include <fstream>
#include <ostream>
#include <cmath>
#include <iomanip>
#include "MD_MC.h"

using namespace std;

int main(){
    Input(); //Inizialization
    int nconf = 1;
    for(int iblk=1; iblk <= nblk; iblk++){ //Data blocking
        Reset(iblk); //Reset block averages
        for(int istep=1; istep <= nstep; istep++){
            Move(); //Move part with Verlet or Metropolis algorithm
            Measure(); //Mesure properties
            Accumulate(); //Update block averages
            if(istep%10 == 0){
                ConfXYZ(nconf); //Write conf in XYZ format; ommented to avoid "filesystem full"!
                nconf += 1;
            }
        }
        Averages(iblk); //Print results for current block
    }
    ConfFinal(); //Write final configuration
    return 0;
}
```

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MD/MC code: Measure ()

```
void Measure(){ //Properties measurement
    double v = 0.0, kin=0.0;
    double vij;
    double dx, dy, dz, dr;

    //cycle over pairs of particles
    for (int i=0; i<npart-1; ++i){
        for (int j=i+1; j<npart; ++j){
            // distance i-j in pbc = Minimum Image Convention
            dx = Pbc(x[i] - x[j]);
            dy = Pbc(y[i] - y[j]);
            dz = Pbc(z[i] - z[j]);
            //update of the histogram of g(r)
            dr = dx*dx + dy*dy + dz*dz;
            dr = sqrt(dr);
            if(dr < rcut){ //Potential cut-off
                vij = 1.0/pow(dr,12) - 1.0/pow(dr,6);
                v += vij;
            }
        }
    }

    //Measure Kinetic energy
    for (int i=0; i<npart; ++i) kin += 0.5 * (vx[i]*vx[i] + vy[i]*vy[i] + vz[i]*vz[i]);

    walker[iv] = 4.0 * v; // Potential energy
    walker[ik] = kin; // Kinetic energy
    walker[it] = (2.0 / 3.0) * kin/(double)npart; // Temperature
    walker[ie] = 4.0 * v + kin; // Total energy;

    return;
}
```

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MD/MC code: Averages (iblk)

```
#include <iostream>
#include <fstream>
#include <ostream>
#include <cmath>
#include <iomanip>
#include "MD_MC.h"

using namespace std;

int main(){
    Input(); //Inizialization
    int nconf = 1;
    for(int iblk=1; iblk <= nblk; iblk++){ //Data blocking
        Reset(iblk); //Reset block averages
        for(int istep=1; istep <= nstep; istep++){
            Move(); //Move part with Verlet or Metropolis algorithm
            Measure(); //Mesure properties
            Accumulate(); //Update block averages
            if(istep%10 == 0){
                ConfXYZ(nconf); //Write conf in XYZ format; ommented to avoid "filesystem full"!
                nconf += 1;
            }
        }
        Averages(iblk); //Print results for current block
    }
    ConfFinal(); //Write final configuration

    return 0;
}
```

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MD/MC code: Averages (iblk) 1.

```
void Averages(int iblk) { //Print results for current block
    ofstream Epot, Ekin, Etot, Temp;
    const int wd=12;

    cout << "Block number " << iblk << endl;
    cout << "Acceptance rate " << accepted/attempted << endl << endl;

    Epot.open("output_epot.dat",ios::app);
    Ekin.open("output_ekin.dat",ios::app);
    Temp.open("output_temp.dat",ios::app);
    Etot.open("output_etot.dat",ios::app);

    stima_pot = blk_av[iv]/blk_norm/(double)npart; //Potential energy
    glob_av[iv] += stima_pot;
    glob_av2[iv] += stima_pot*stima_pot;
    err_pot=Error(glob_av[iv],glob_av2[iv],iblk);

    stima_kin = blk_av[ik]/blk_norm/(double)npart; //Kinetic energy
    glob_av[ik] += stima_kin;
    glob_av2[ik] += stima_kin*stima_kin;
    err_kin=Error(glob_av[ik],glob_av2[ik],iblk);
}
```

Add tail corrections



... continues in the next slide ...

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MD/MC code: Averages (iblk) 2.

```
stima_etot = blk_av[ie]/blk_norm/(double)npart; //Total energy
glob_av[ie] += stima_etot;
glob_av2[ie] += stima_etot*stima_etot;
err_etot=Error(glob_av[ie],glob_av2[ie],iblk);

stima_temp = blk_av[it]/blk_norm; //Temperature
glob_av[it] += stima_temp;
glob_av2[it] += stima_temp*stima_temp;
err_temp=Error(glob_av[it],glob_av2[it],iblk);

//Potential energy per particle
Epot << setw(wd) << iblk << setw(wd) << stima_pot << setw(wd) <<
glob_av[iv]/(double)iblk << setw(wd) << err_pot << endl;
//Kinetic energy
Ekin << setw(wd) << iblk << setw(wd) << stima_kin << setw(wd) <<
glob_av[ik]/(double)iblk << setw(wd) << err_kin << endl;
//Total energy
Etot << setw(wd) << iblk << setw(wd) << stima_etot << setw(wd) <<
glob_av[ie]/(double)iblk << setw(wd) << err_etot << endl;
//Temperature
Temp << setw(wd) << iblk << setw(wd) << stima_temp << setw(wd) <<
glob_av[it]/(double)iblk << setw(wd) << err_temp << endl;
//g(r)
```

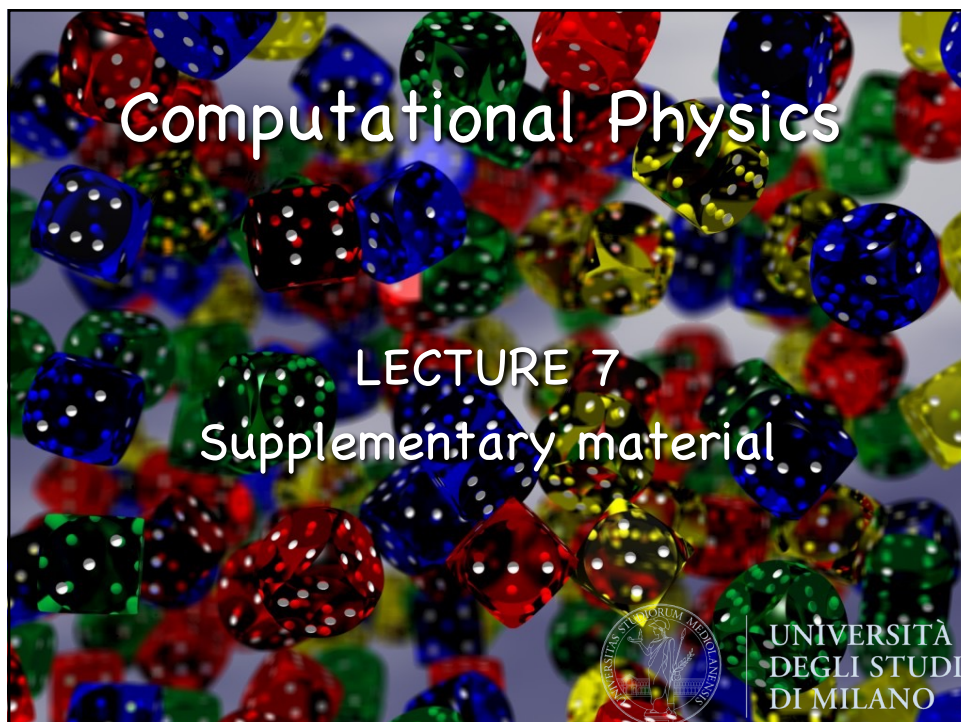
← insert your code here

```
cout << "-----" << endl << endl;

Epot.close();
Ekin.close();
Etot.close();
Temp.close();
}
```

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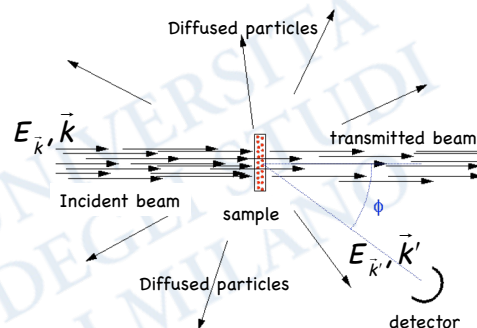
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Scattering experiments

- In a typical scattering (diffusion) experiment a beam of "particles" (photons, neutrons, electrons,...) with linear momentum $\hbar\vec{k}$ and energy E_k is radiated over a sample.



- The scattering experiments analyses how these "particles" are diffused, i.e. one counts how many "particles" of the incident beam come out with linear momentum $\hbar\vec{k}'$ and energy $E_{k'}$

$$\text{Exchanged momentum: } \hbar\vec{q} = \hbar(\vec{k} - \vec{k}')$$

$$\text{Exchanged energy: } \hbar\omega = E_k - E_{k'}$$

- Hypothesis: the "particles" of the incident beam interact weakly with the sample

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1st Born approximation

- This means that most of the "probe-particles" do not exchange energy or momentum with the sample, thus they are found in the transmitted beam
- In such case we can treat the interaction term among "particles" and the sample at first order of perturbative theory, which is called **1st Born approximation**
- The 1st Born approximation, in practice, assumes that the **field inside the scattering volume is constant and equal to the incident field**
- This is the case of **thermal neutrons** ($E_k \approx k_B T$), and of **photons** (visible light and X-rays) scattered by a **transparent sample**, i.e. matter which is not reflecting like a metal for visible light and that does not present absorption for the considered photon frequency
- To be concrete we will discuss an incident beam of non-relativistic massive particles, i.e. the ("thermal") neutron scattering case

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Neutron scattering

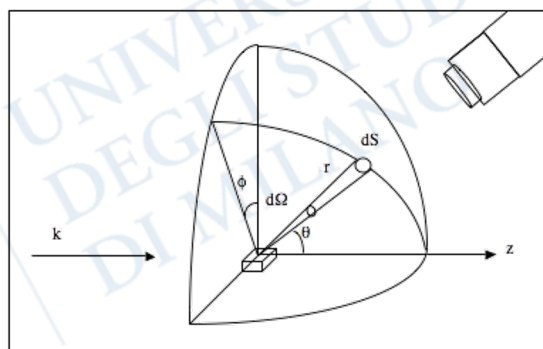
- Since **neutrons** are neutral, their **interaction with the nuclei is short-ranged**; in contrast to electrons, they penetrate deep into the solid.
- Furthermore, due to their magnetic moment and the associated dipole interaction with magnetic moments of the solid, neutrons can also be **used to investigate magnetic properties** like magnetic order in the system or dynamical properties: spin waves
- Therefore, the usefulness of neutron scattering originates from the comparatively simple and weak interaction of the neutrons with condensed matter and the well adapted "mechanical properties" ($\lambda_{DB} \approx$ inter-atomic distance for "thermal" energies) of the neutron beams to the microscopic and **energetic structure of matter**
- The weakness of the interaction renders the system in the ground state for elastic scattering or in a well defined excited state for inelastic scattering

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Scattering cross sections

- Consider a **beam of thermal** (i.e. cooled down [moderated] to thermal energy 2-100 meV [$30 < T < 1100$ K]) **neutrons**, all with the same energy E (**monochromatic beam**) incident on a target
- Suppose we set up a **neutron counter** and measure the number of neutrons scattered in a **given direction** as function of their energy E' .
- The distance of the counter from the target is assumed to be large compared with the dimensions of the counter and the target, so that the small angle $d\Omega$ subtended by the counter at the target is well defined. Let the directions of the scattered neutron be θ and ϕ



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- The **partial differential cross section** is defined as:

$$\frac{d^2\sigma}{d\Omega dE'} = \frac{\text{(number of neutrons scattered per second into a small solid angle } d\Omega \text{ in the direction } \theta \text{ and } \phi \text{ with final energy between } E' \text{ and } E'+dE')}{(\Phi d\Omega dE')}$$

where Φ is the flux of the incident neutrons, i.e. the number through unit area per second

- Suppose we do not analyse the energy of the scattered neutrons, but simply count all the neutrons scattered into a solid angle $d\Omega$ in the direction θ and ϕ . The cross section corresponding to these measurements, known as the **differential cross section**, is defined by

$$\frac{d\sigma}{d\Omega} = \frac{\text{(number of neutrons scattered per second into } d\Omega \text{ in the direction } \theta, \phi)}{(\Phi d\Omega)}$$

- The **total scattering cross section** is defined by the equation

$$\sigma_{tot} = \frac{\text{(total number of neutrons scattered per second)}}{(\Phi)}$$

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- From their definitions the three cross-sections are related by the following equations

$$\frac{d\sigma}{d\Omega} = \int_0^\infty dE' \left(\frac{d^2\sigma}{d\Omega dE'} \right) ; \quad \sigma_{tot} = \int d\Omega \left(\frac{d\sigma}{d\Omega} \right)$$

- The **dimensions of the total cross-section is [area]**, and more precisely is an area rescaled with the ratio of two pure numbers: the number of scattered neutrons divided by the number of incoming neutrons, i.e. the percentage of the scattered neutrons; it is a sort of **measure of the "surface" able to deviate the beam**. The definition of cross-sections apply to any kind of scattering
- The nuclear forces which cause the scattering have a range of about 10^{-14} to 10^{-15} m; the wavelength of thermal neutrons is of the order of 10^{-10} m, and is thus much larger than this range. In these circumstances the scattering, analysed in terms of partial waves, comes entirely from **s waves** ($l=0$)
- In fact, if waves of any kind are scattered by an object small compared to the wavelength of the waves, then the scattered wave is **spherically symmetric**

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The dynamic structure factor

- In the 1st Born approximation where the Fermi's golden rule applies one can easily obtain that:

$$\frac{d^2\sigma}{d\Omega dE'} = \frac{1}{N} \frac{k'}{k} \frac{a^2}{2\pi\hbar} \int_{-\infty}^{\infty} dt e^{i\omega t} \left\langle \sum_j e^{-i\vec{q}\cdot\vec{R}_j(t)} \sum_j e^{i\vec{q}\cdot\vec{R}_j} \right\rangle = \frac{k'}{k} \frac{a^2}{\hbar} S(\vec{q}, \omega)$$

where we have introduced the **dynamic structure factor** defined as

$$S(\vec{q}, \omega) = \frac{1}{2\pi N} \int_{-\infty}^{\infty} dt e^{i\omega t} \left\langle \sum_j e^{-i\vec{q}\cdot\vec{R}_j(t)} \sum_j e^{i\vec{q}\cdot\vec{R}_j} \right\rangle = \frac{1}{2\pi N} \int_{-\infty}^{\infty} dt e^{i\omega t} \langle \hat{\rho}_{\vec{q}}(t) \hat{\rho}_{\vec{q}}^{\dagger}(0) \rangle$$

- It is very easy to see that $\rho_{\vec{q}}$, which is generally called the **density fluctuation operator**, is the Fourier transform of the number density operator:

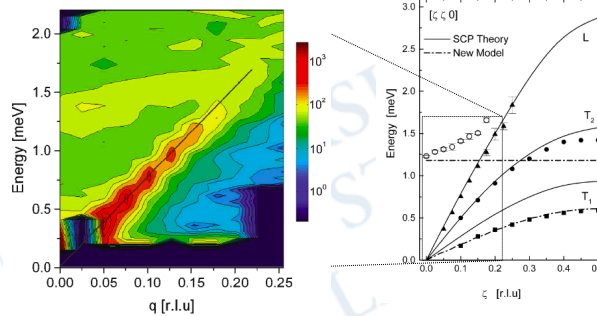
$$\hat{\rho}(\vec{r}, t) = \sum_j \delta(\vec{r} - \vec{r}_j(t)) \Rightarrow \hat{\rho}_{\vec{q}}(t) = \int d\vec{r} e^{-i\vec{q}\cdot\vec{r}} \hat{\rho}(\vec{r}, t) = \sum_j e^{-i\vec{q}\cdot\vec{r}_j(t)}$$

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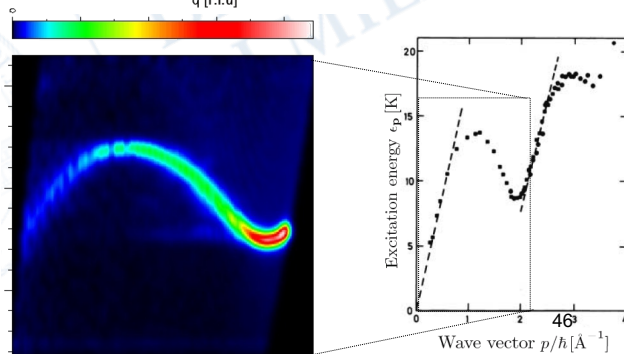
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- We give her two examples:

Inelastic scattering:
 $S(\vec{q}, \omega)$ in solids (phonons)



Inelastic scattering:
 $S(\vec{q}, \omega)$ in He-II



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The static structure factor

- The **differential scattering cross-section** is connected with a scattering experiment in which one registers all events related to a momentum transfer equal to $\hbar\mathbf{q}$ but **independently on any particular energy transfer**
- It is obtained by integrating with respect to the energy the partial differential scattering cross-section:

$$\frac{d\sigma}{d\Omega} = \int dE \frac{d^2\sigma}{d\Omega dE} = \hbar \int_{-\infty}^{\infty} d\omega \frac{k'}{k} \frac{a^2}{\hbar} S(\vec{q}, \omega) = \frac{k'}{k} a^2 \int_{-\infty}^{\infty} d\omega S(\vec{q}, \omega) = \frac{k'}{k} a^2 S(\vec{q})$$

it then proportional to the function **$S(\mathbf{q})$** which is called **static structure factor**

- This function is also easily obtained in a scattering experiment and it is very important because give information on the static structure of the system:

$$S(\vec{q}) = \int_{-\infty}^{\infty} d\omega S(\vec{q}, \omega) = \int_{-\infty}^{\infty} d\omega \frac{1}{2\pi N} \int_{-\infty}^{\infty} dt e^{i\omega t} \left\langle \hat{\rho}_{\vec{q}}(t) \hat{\rho}_{\vec{q}}^+(0) \right\rangle = \dots$$

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- Thus

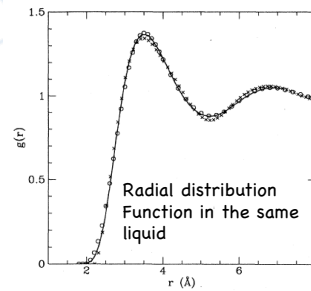
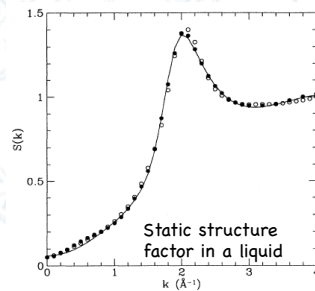
$$S(\vec{q}) = \frac{1}{N} \int_{-\infty}^{\infty} dt \delta(t) \left\langle \hat{\rho}_{\vec{q}}(t) \hat{\rho}_{\vec{q}}^+(0) \right\rangle = \frac{1}{N} \left\langle \hat{\rho}_{\vec{q}} \hat{\rho}_{\vec{q}}^+ \right\rangle$$

- In an isotropic and homogeneous system there is a **direct relation** between **$S(\mathbf{q})$** and the **radial (2-particles) distribution function**:

$$S(\vec{q}) = \frac{1}{N} \left\langle \sum_j e^{-i\vec{q} \cdot \vec{r}_j} \sum_l e^{i\vec{q} \cdot \vec{r}_l} \right\rangle = 1 + \frac{1}{N} \left\langle \sum_{j \neq l} e^{-i\vec{q} \cdot (\vec{r}_j - \vec{r}_l)} \right\rangle = \dots$$

$$= 1 + \frac{1}{N} \left\langle \sum_{j \neq l} \int d\vec{r} d\vec{r}' e^{-i\vec{q} \cdot (\vec{r} - \vec{r}')} \delta(\vec{r} - \vec{r}_j) \delta(\vec{r}' - \vec{r}_l) \right\rangle = 1 + \frac{1}{N} \int d\vec{r} d\vec{r}' e^{-i\vec{q} \cdot (\vec{r} - \vec{r}')} \rho^{(2)}(\vec{r}, \vec{r}') =$$

Uniform system $\longrightarrow = 1 + \rho \int d(\vec{r} - \vec{r}') e^{-i\vec{q} \cdot (\vec{r} - \vec{r}')} g^{(2)}(\vec{r} - \vec{r}') = 1 + \rho \int d\vec{r} e^{-i\vec{q} \cdot \vec{r}} g(\vec{r})$



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