

- A brief overview of the reflectometry technique
- Discussion of reflectometry analysis methodology

Introduction to Reflectometry: With a Focus on Analysis

FASEM for Life Science — 2024/03/12



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♂ (he/him)

Just a few housekeeping rules.

- between the accent and my speed of speech, I may be hard to understand — let me know.
- let's make this a dialogue; ask questions if you have them and I will ask you questions.
- unfortunately, I need to leave, to return to Bristol, straight after lunch, if you want to chat please drop me an email.

- A brief overview of the reflectometry technique
- Discussion of reflectometry analysis methodology
- Outline of some life science examples of reflectometry
- The opportunity to perform some reflectometry analysis using Python

What is the difference between reflectometry and reflectivity?

- reflectometry: the technique used to measure reflectivity
- reflectivity: the quantity measured by reflectometry

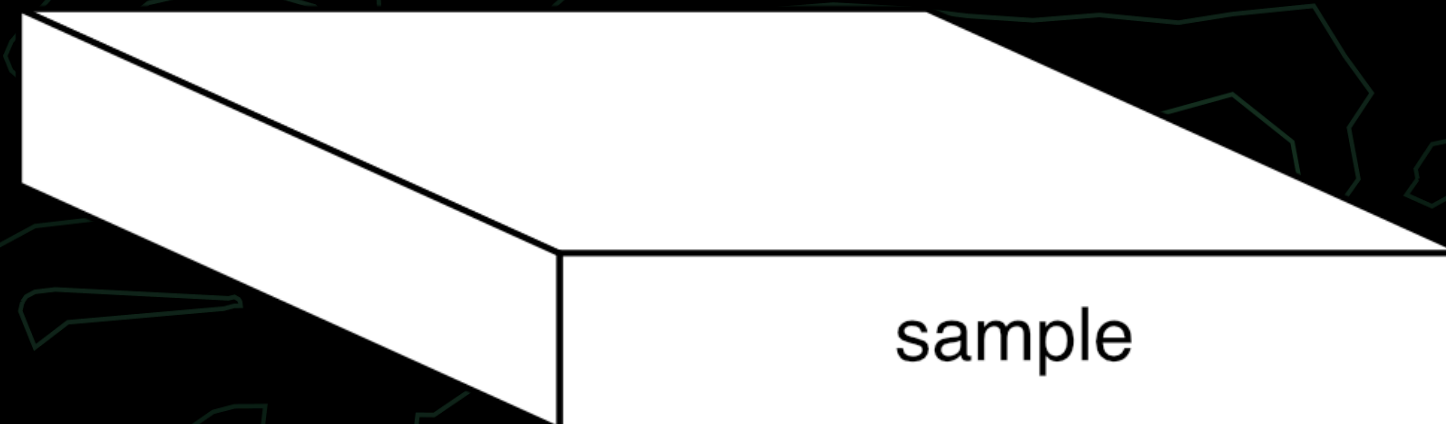
Reflectometry is a surface sensitive analysis technique.

"God made solids, but surfaces are the work of the devil",

Wolfgang Pauli

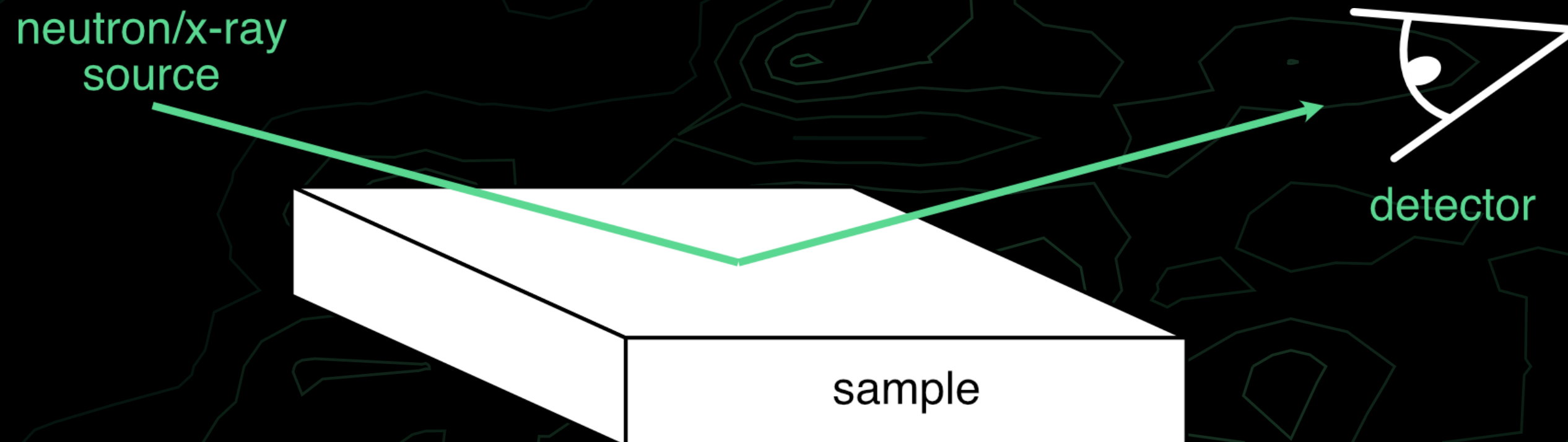
Reflectometry is a surface sensitive analysis technique.

- neutron or x-rays
- reflection geometry: angle of incidence = angle of reflection
 - ratio of incidence to reflected intensity



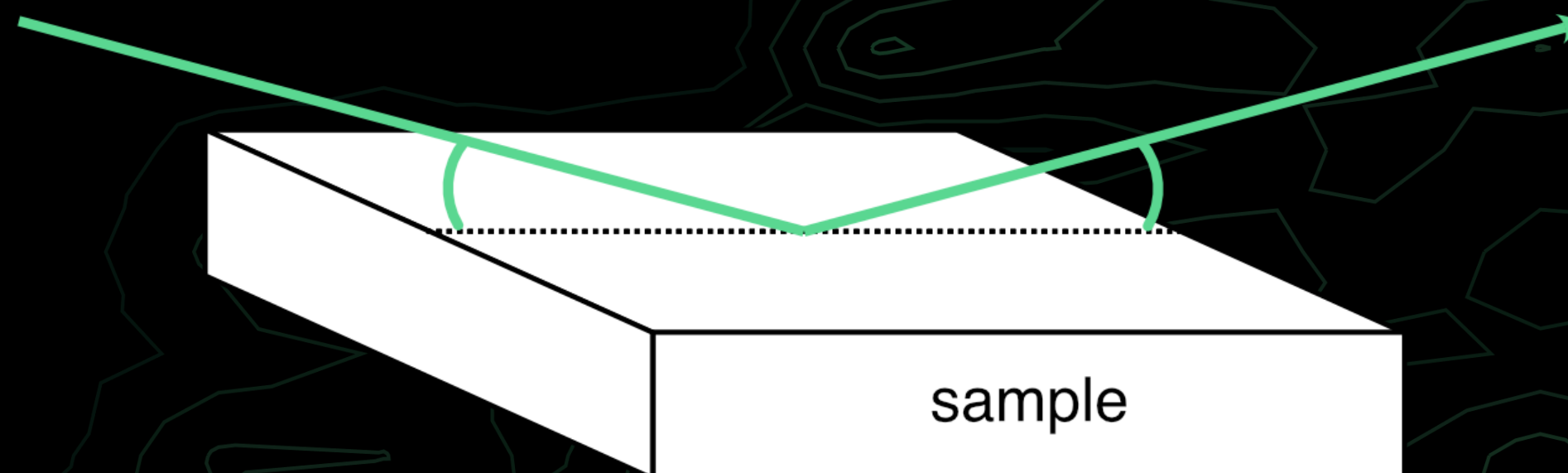
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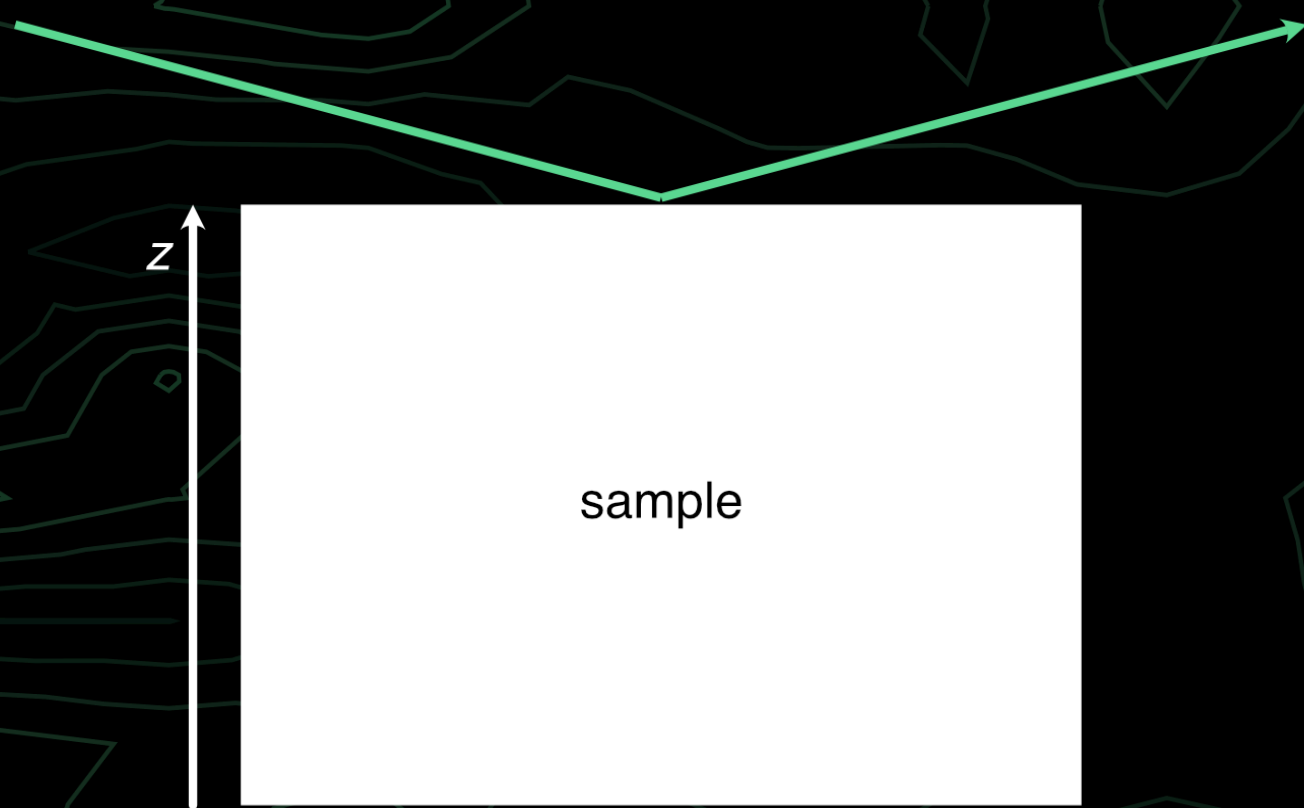
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$$q = \frac{4\pi}{\lambda} \sin \theta$$

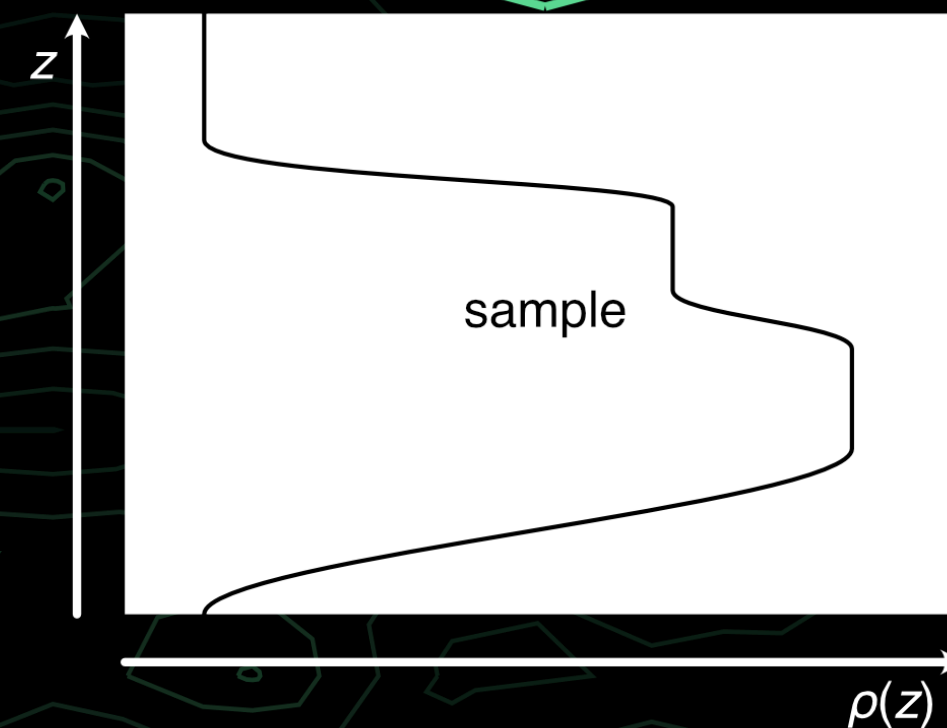
Reflectometry measurements contain information about the material structure perpendicular to the interface.

- information about the scattering length density: $\rho = \frac{\sum_{i=1}^N b_i}{V_m}$
 - ρ is a material property
- we relate ρ to our understanding of the system



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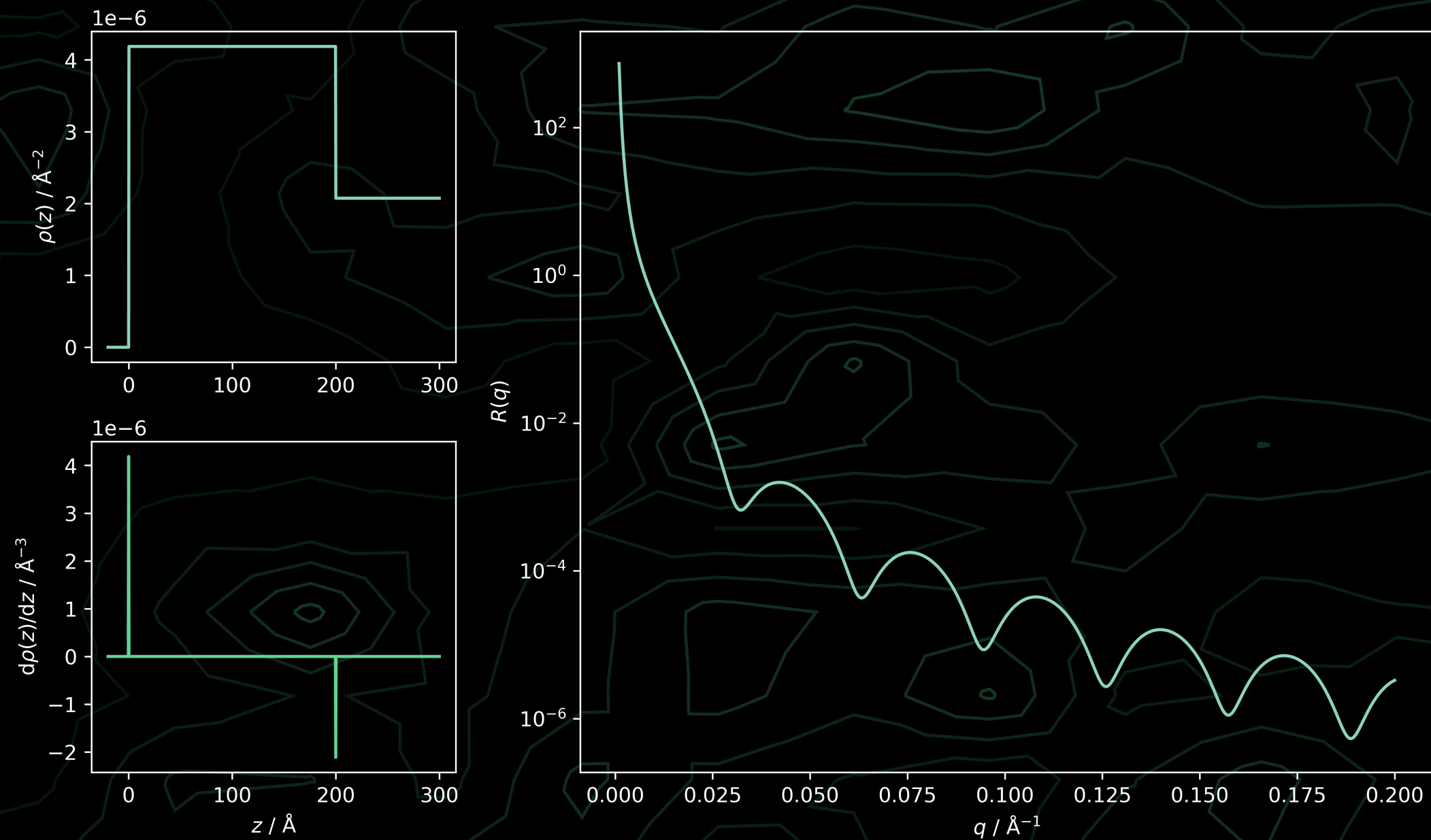


Under the Born approximation, the reflectometry profile can be estimated from a Fourier transform of the scattering length density profile.

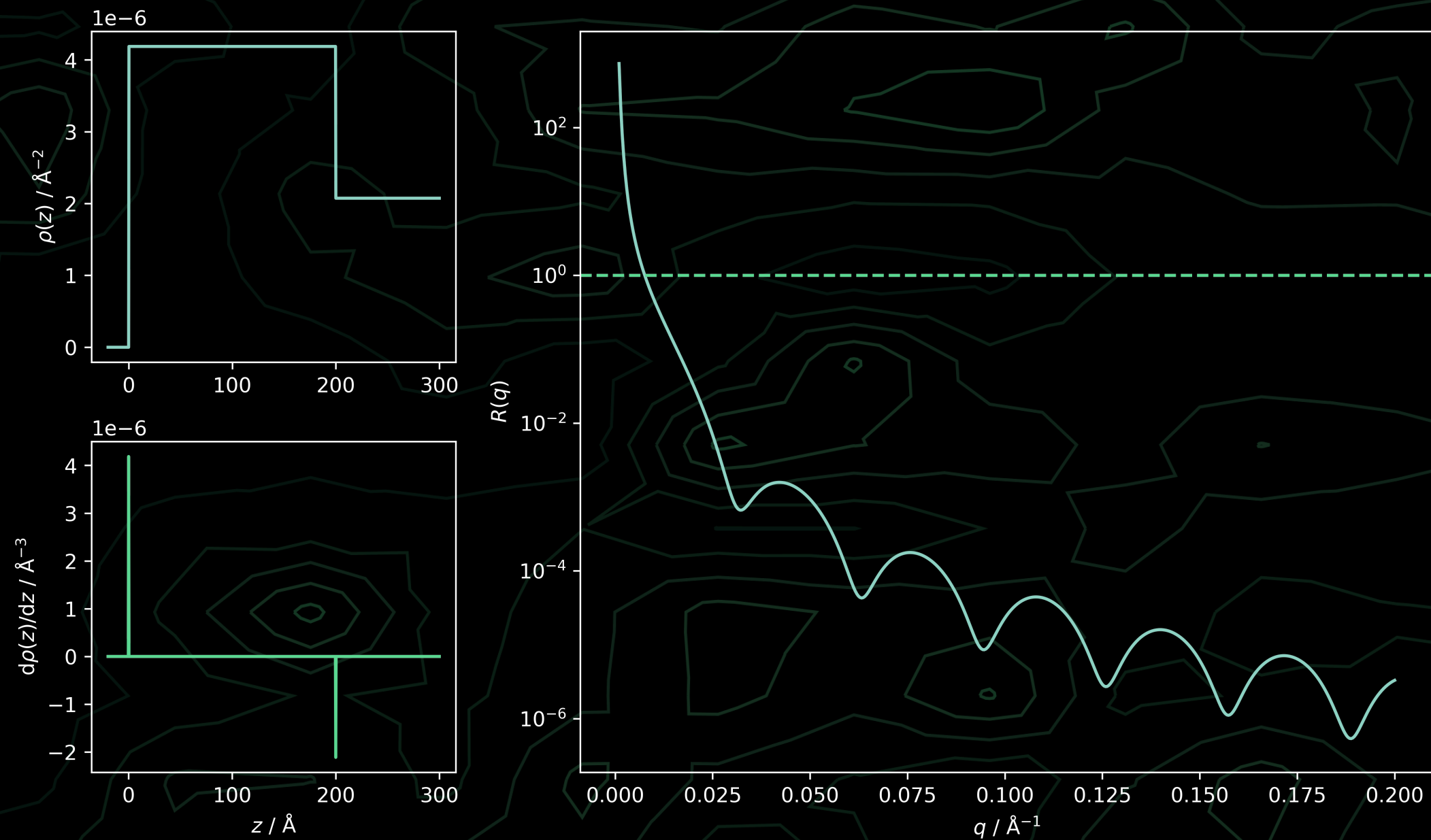
- specifically, the Fourier transform of the derivative (or change) in the scattering length density:

$$R(q) \approx \frac{16\pi^2}{q^4} \left| \int_{-\infty}^{\infty} \frac{d\rho}{dz} \exp(izq) dz \right|^2$$

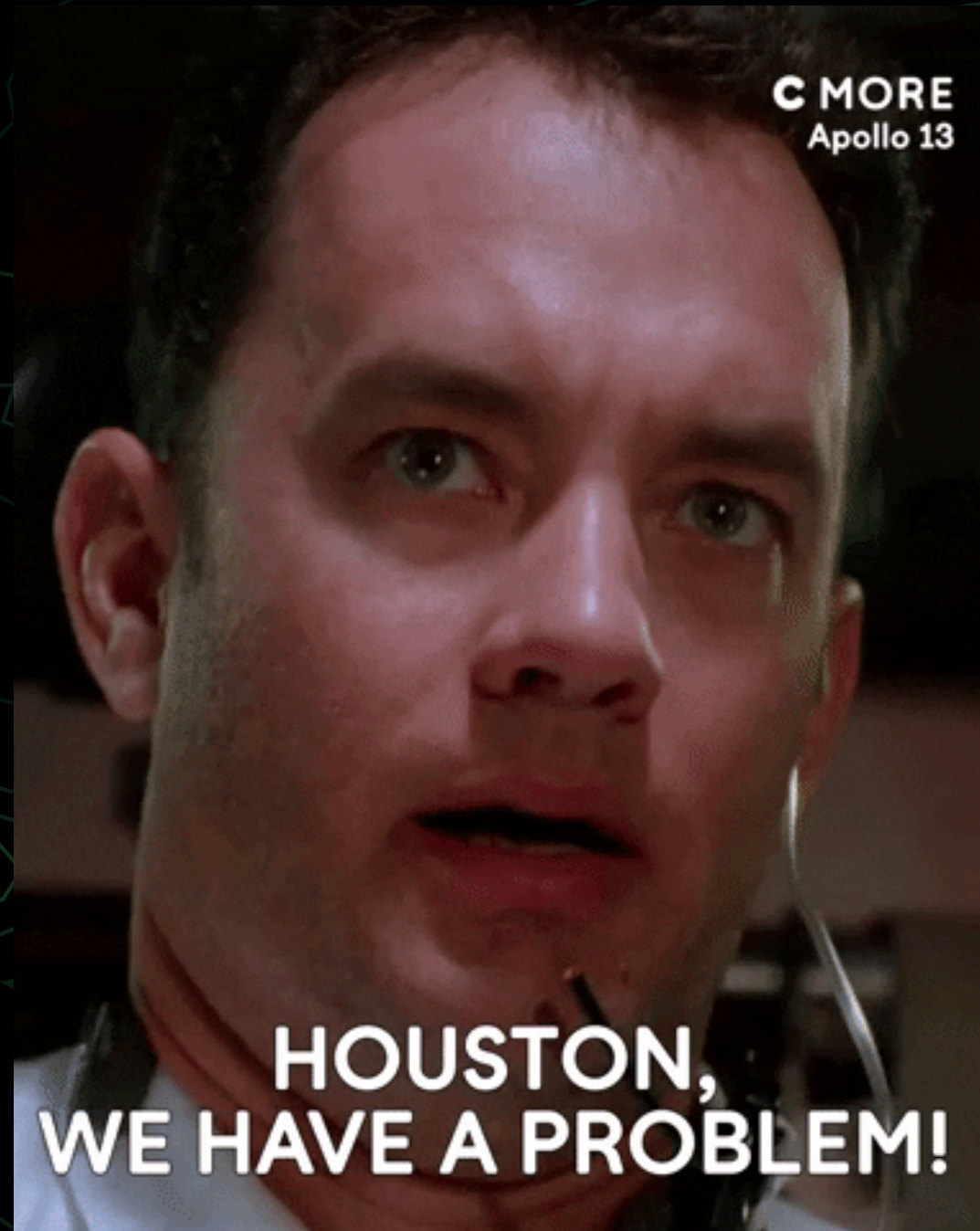
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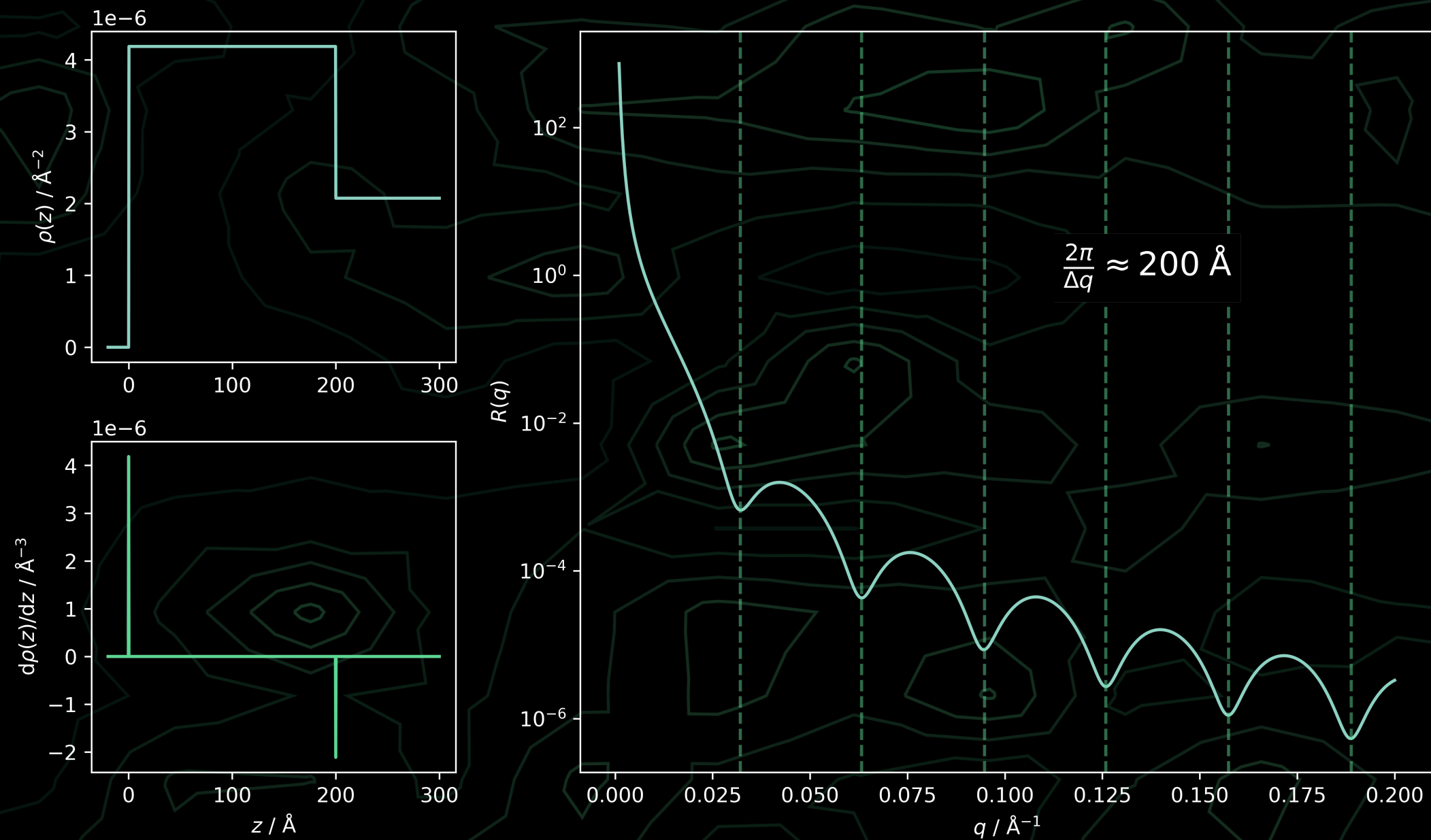
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The Born approximation appears not to work for reflectometry.

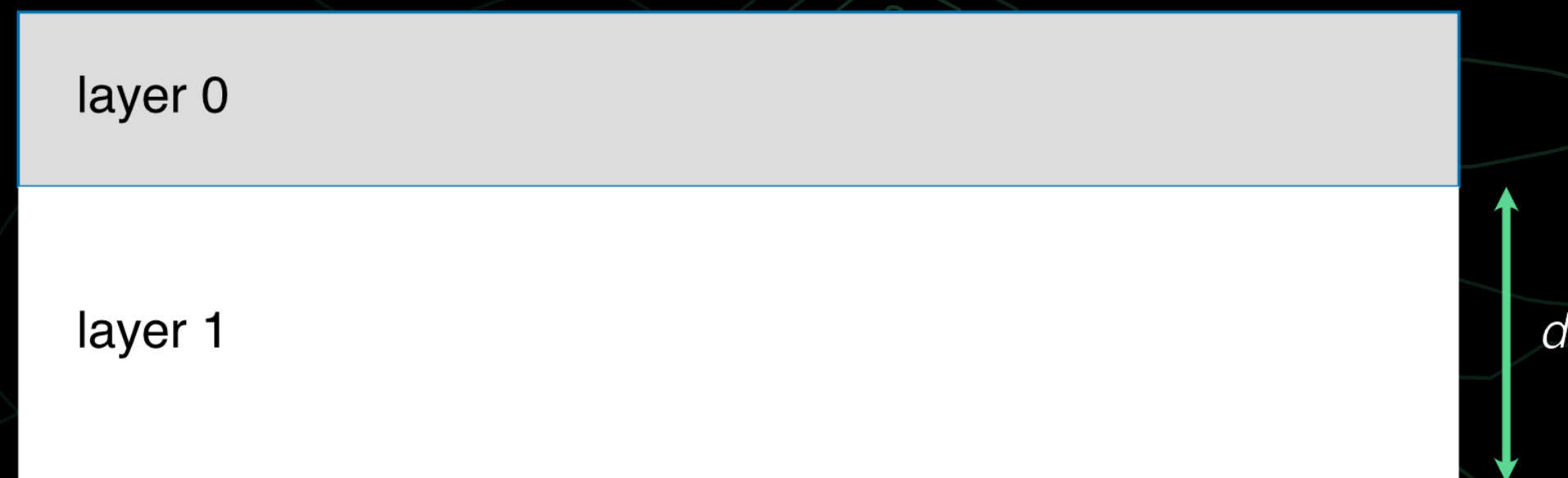


Before we leave the Born approximation behind, it is important to acknowledge all hope is not lost.



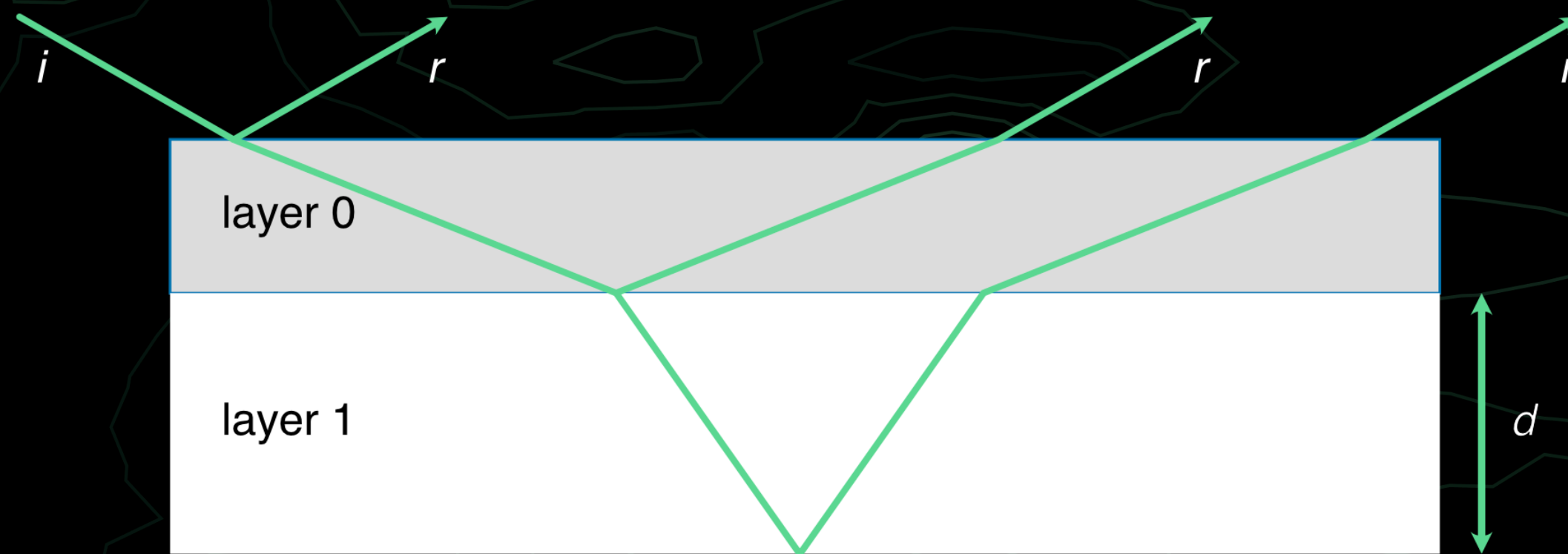
A different approach to understand reflectometry treats the system as a series of layers with optical properties.

- each layer has a given thickness, scattering length density, with a roughness parameter between each layer



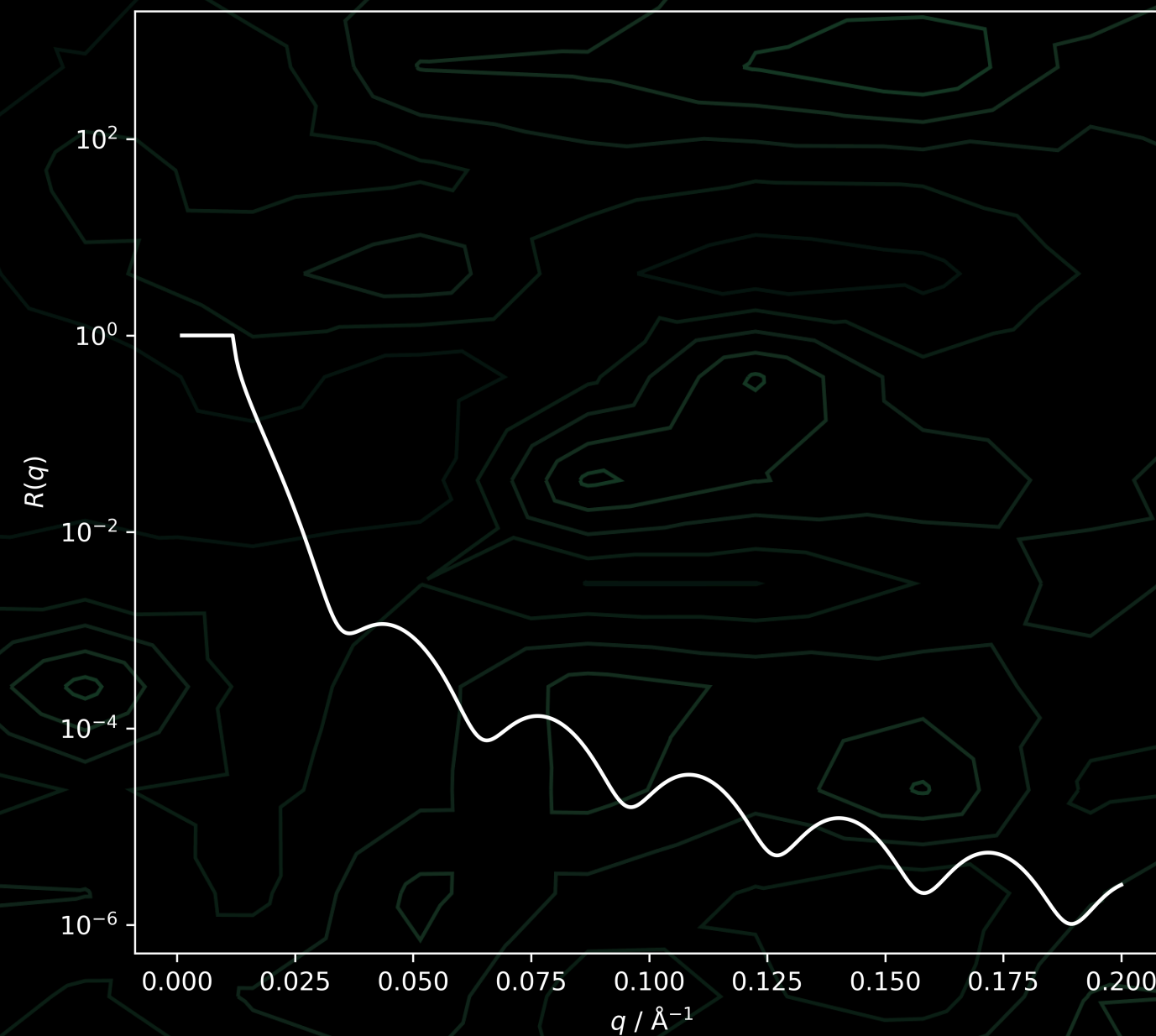
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- optical matrix formalism computes the propagation of the neutron/x-ray waves through each material



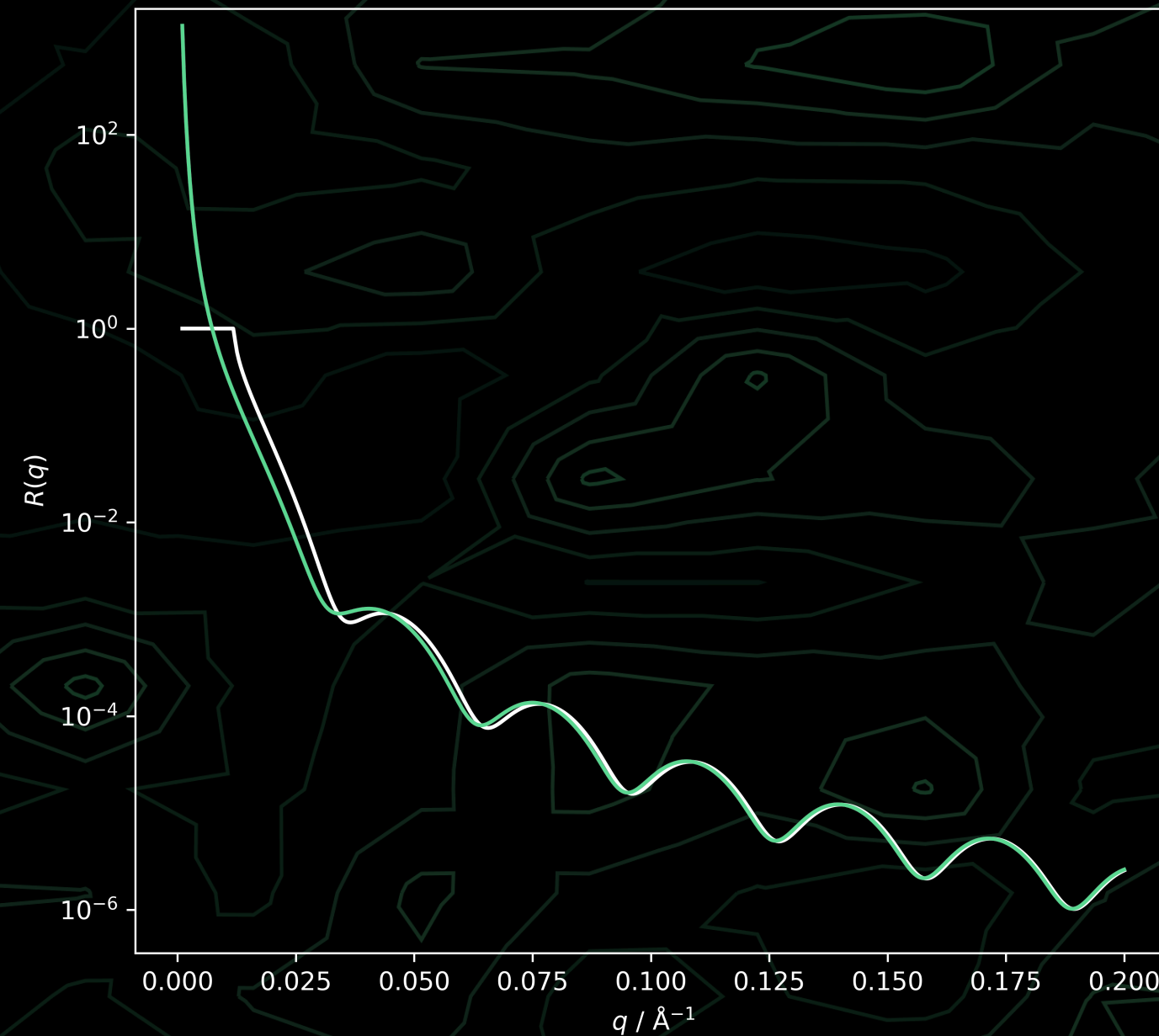
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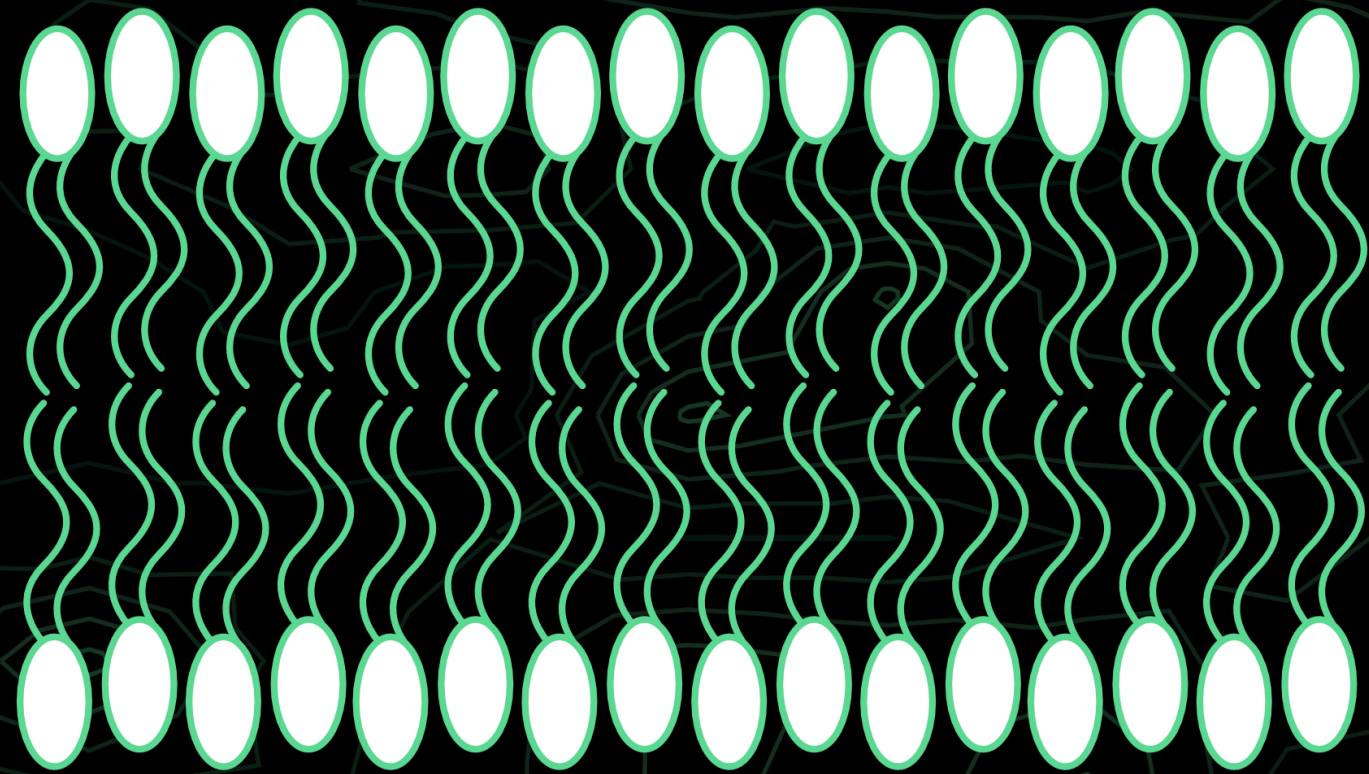
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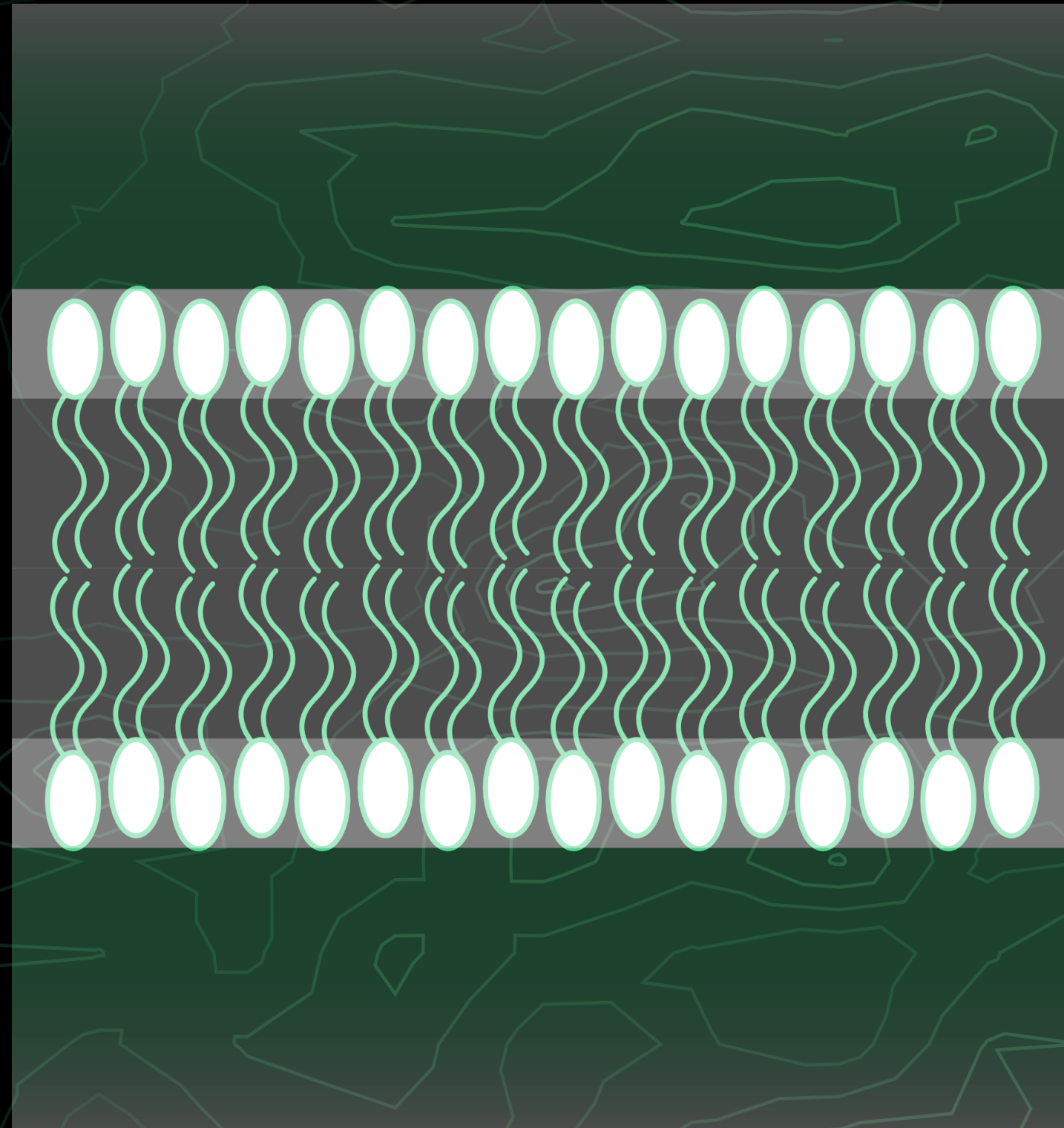
The difference between the Born approximation and the optical matrix formalism may be from the reflection geometry.

- perhaps multiple scattering due to the long path length from shallow angles
 - though this is debated in the community

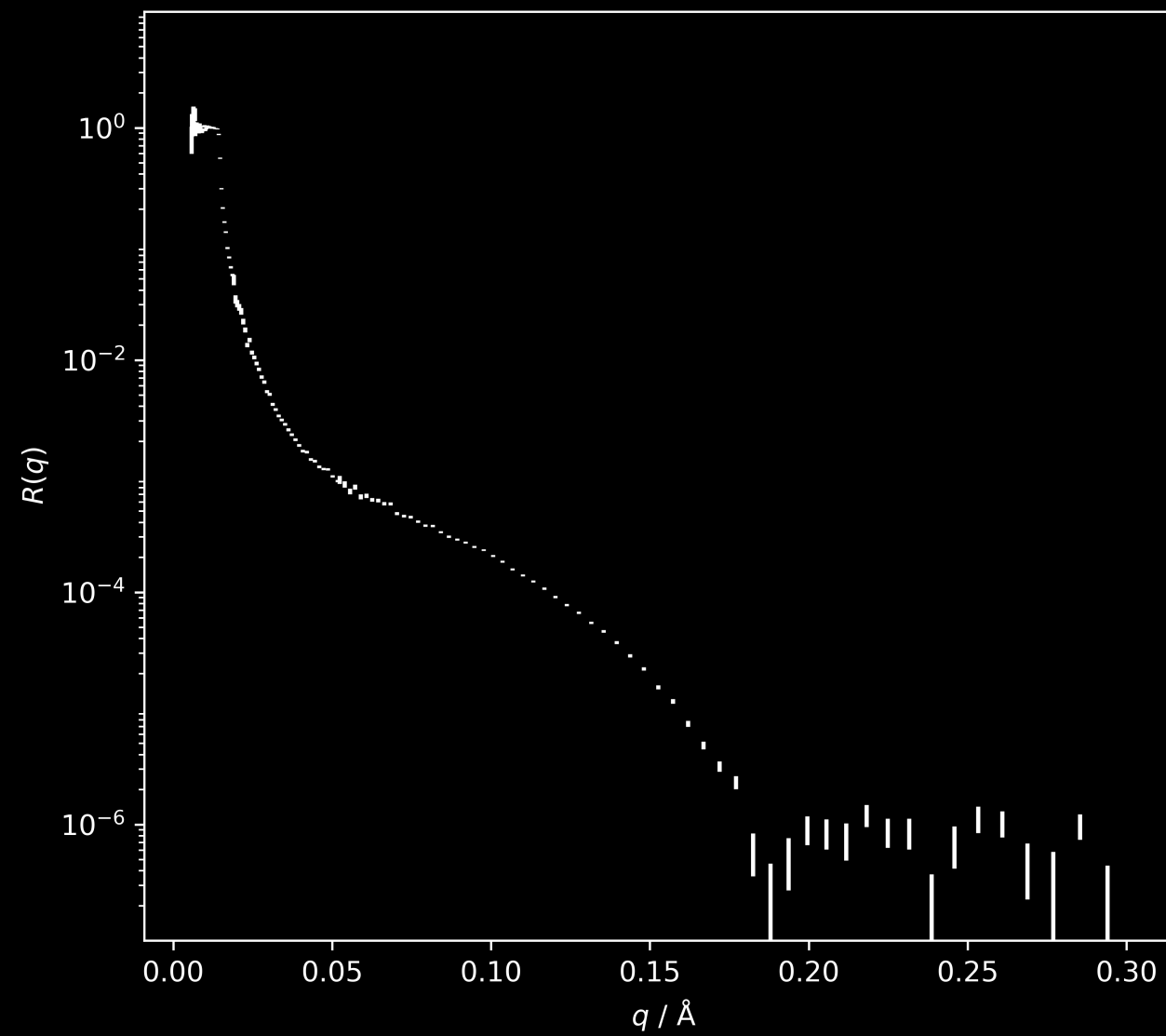
This means we must be able to describe our system as a series of distinct layers.



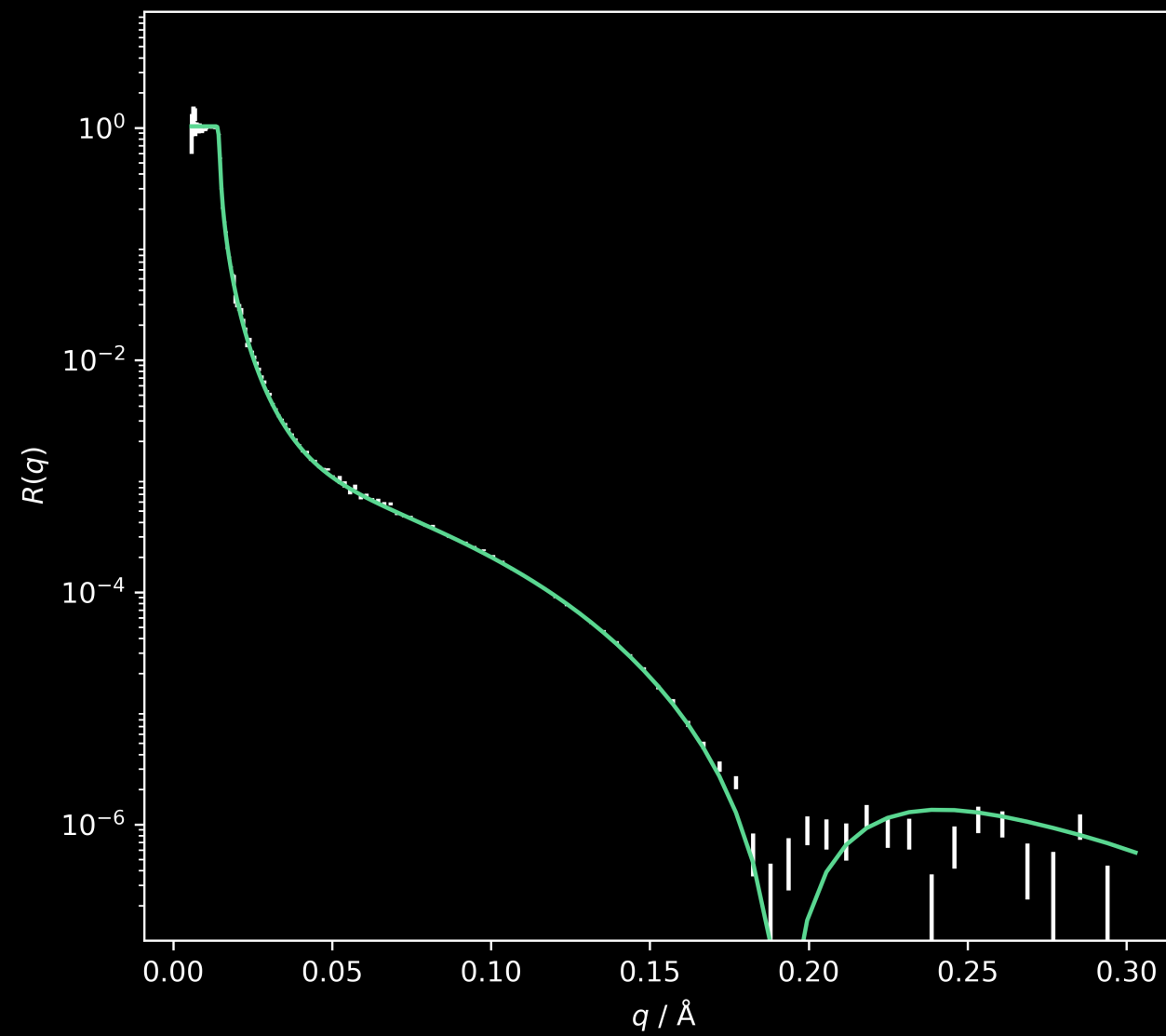
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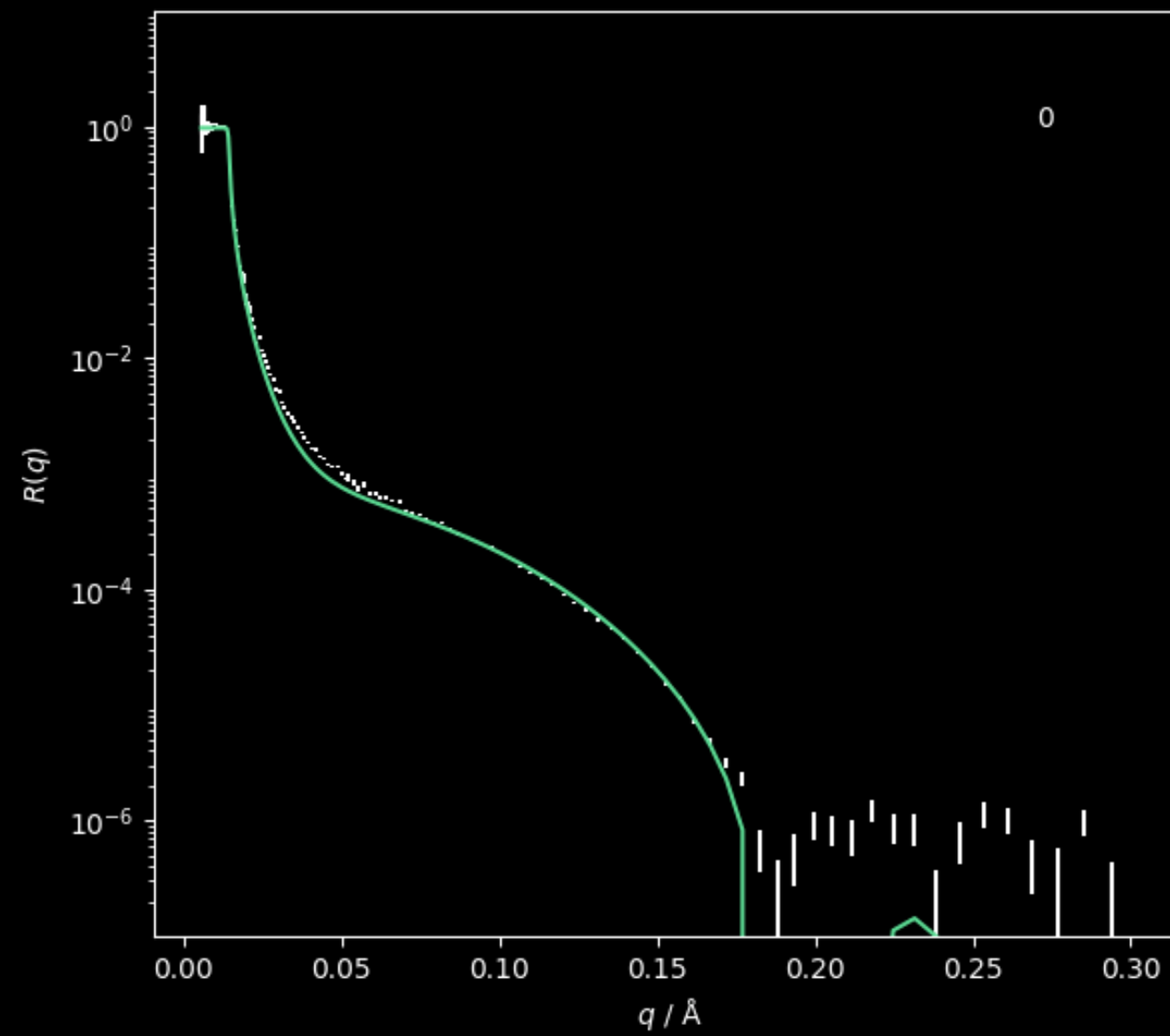
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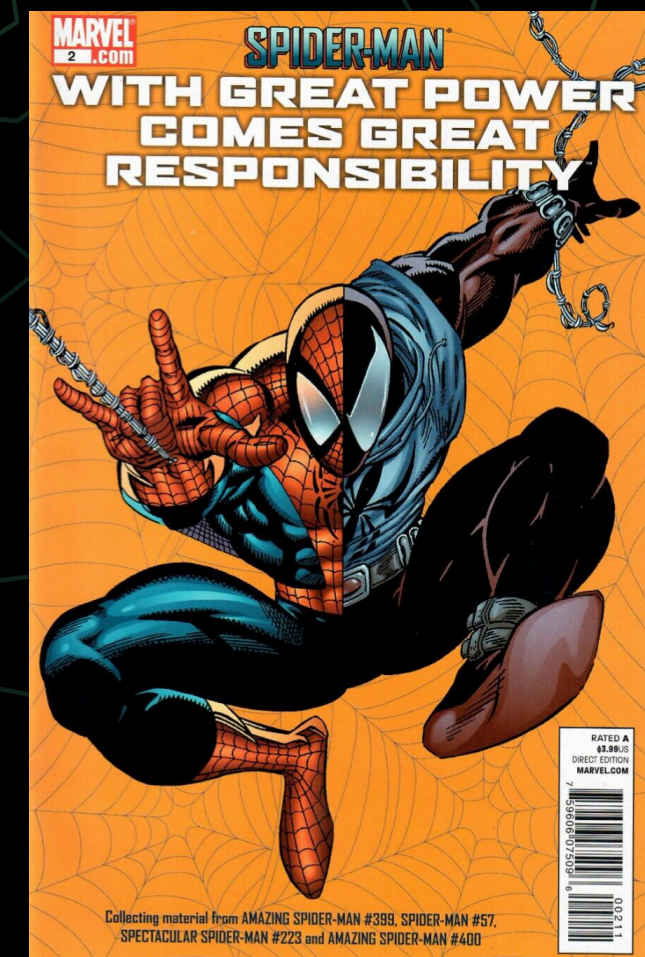


Once we have a model, it needs to be optimised with respect to the data.



Optimisation is achieved through an optimisation algorithm that aims to maximise the agreement between the model and the data.

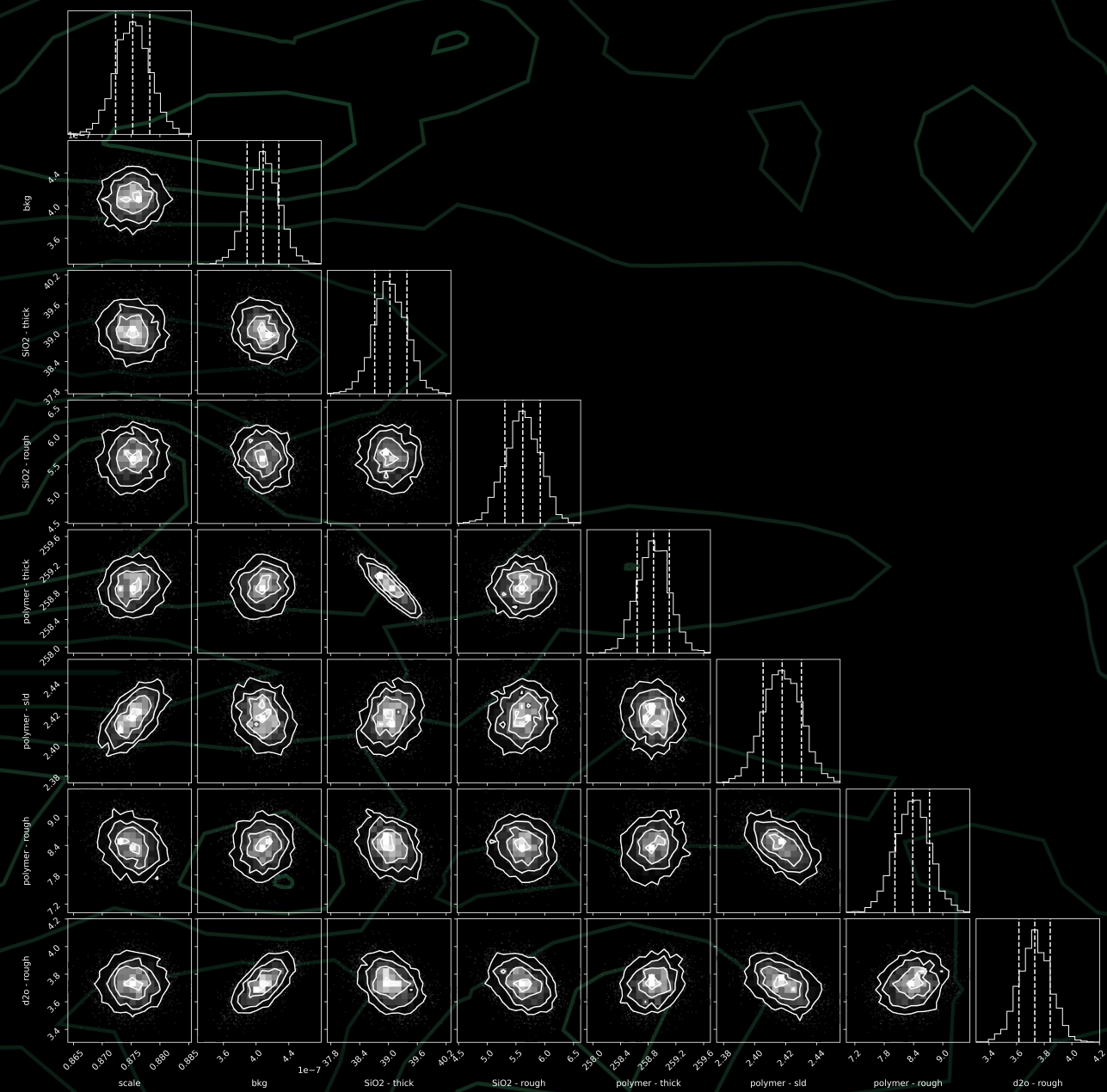
- this optimisation comes with great responsibility
- if you let a parameter vary, the algorithm will try and optimise its value
- popular algorithms: gradient descent methods, differential evolution, DREAM



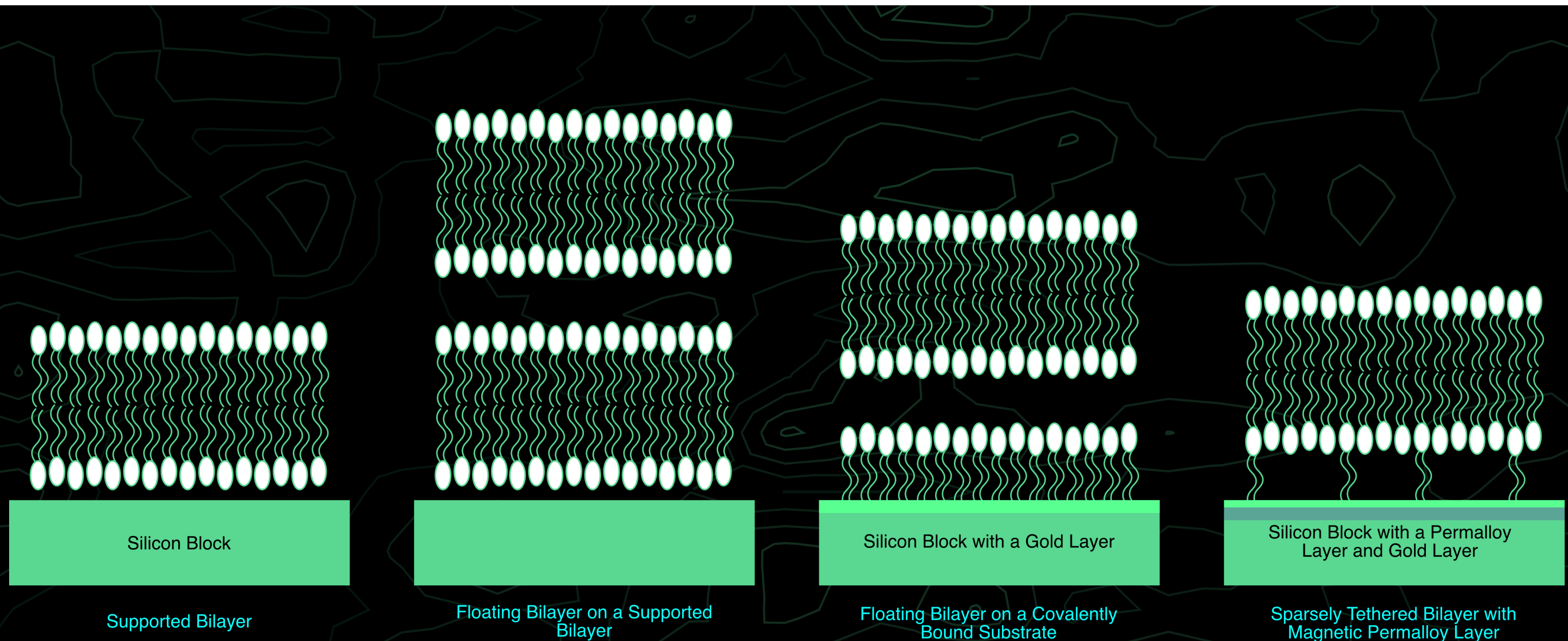
A. R. McCluskey, et al. *Mach. Learn.: Sci. Technol.*, 1(3), 035002, 2020. DOI: 10.1088/2632-2153/ab94c4.

Sampling the probability distribution is becoming a common approach in reflectometry analysis.

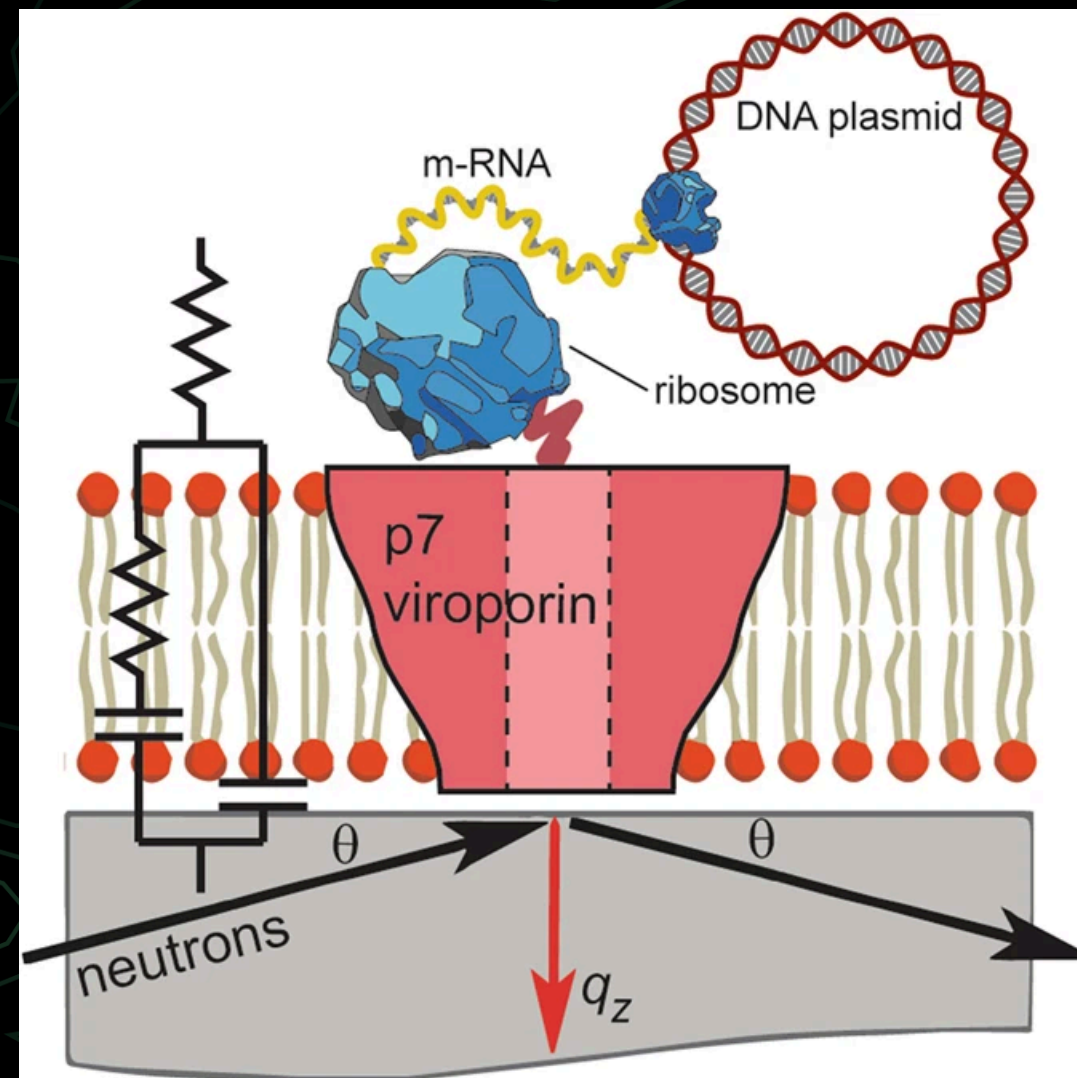
- reflectometry needs this, it is ill-posed
- gives a more complete picture of the parameter uncertainties and correlations
- this comes alongside the popularity of Bayesian modelling



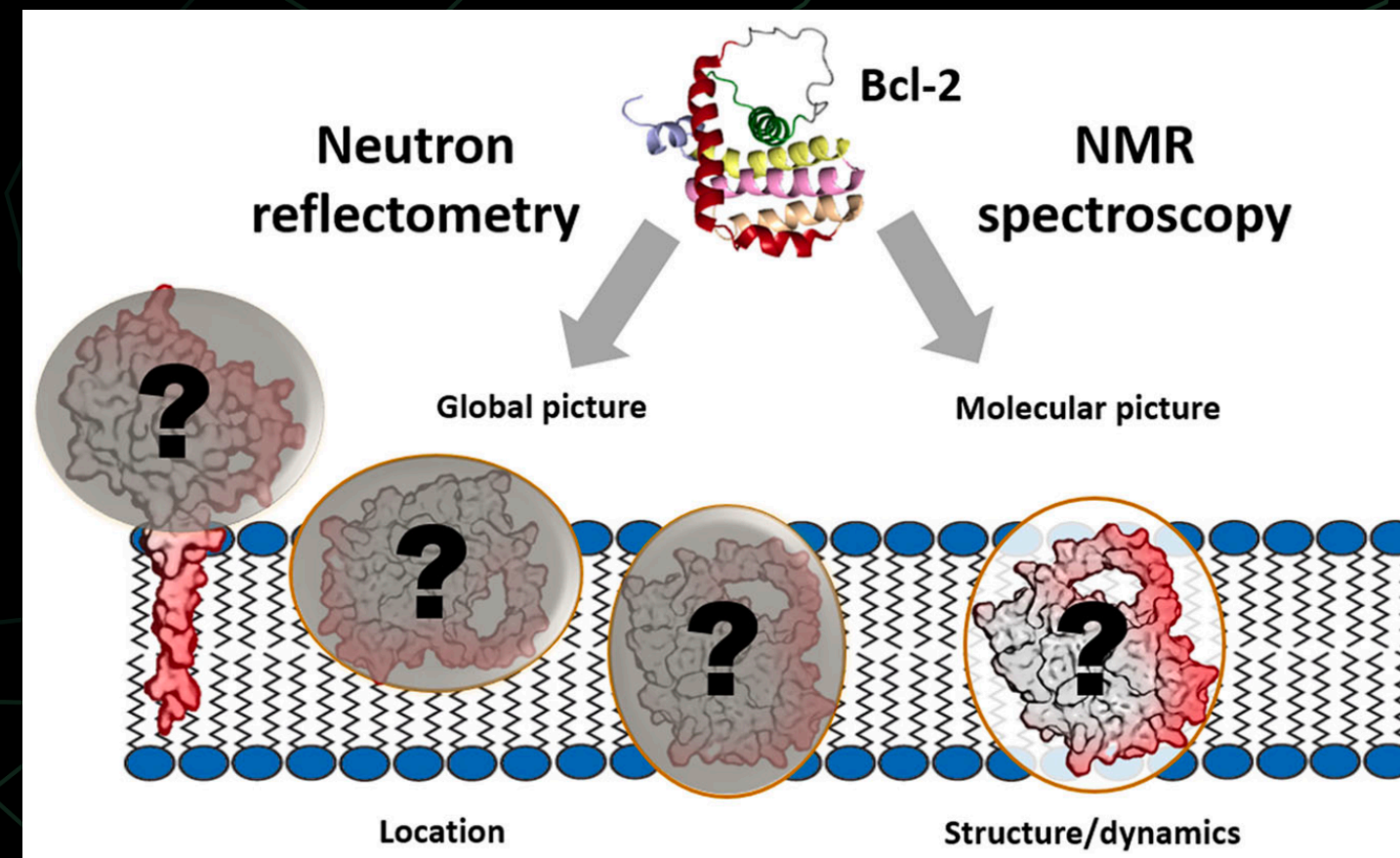
In life sciences, reflectometry methods are particularly valuable for the study of biological, or near biological, membranes.



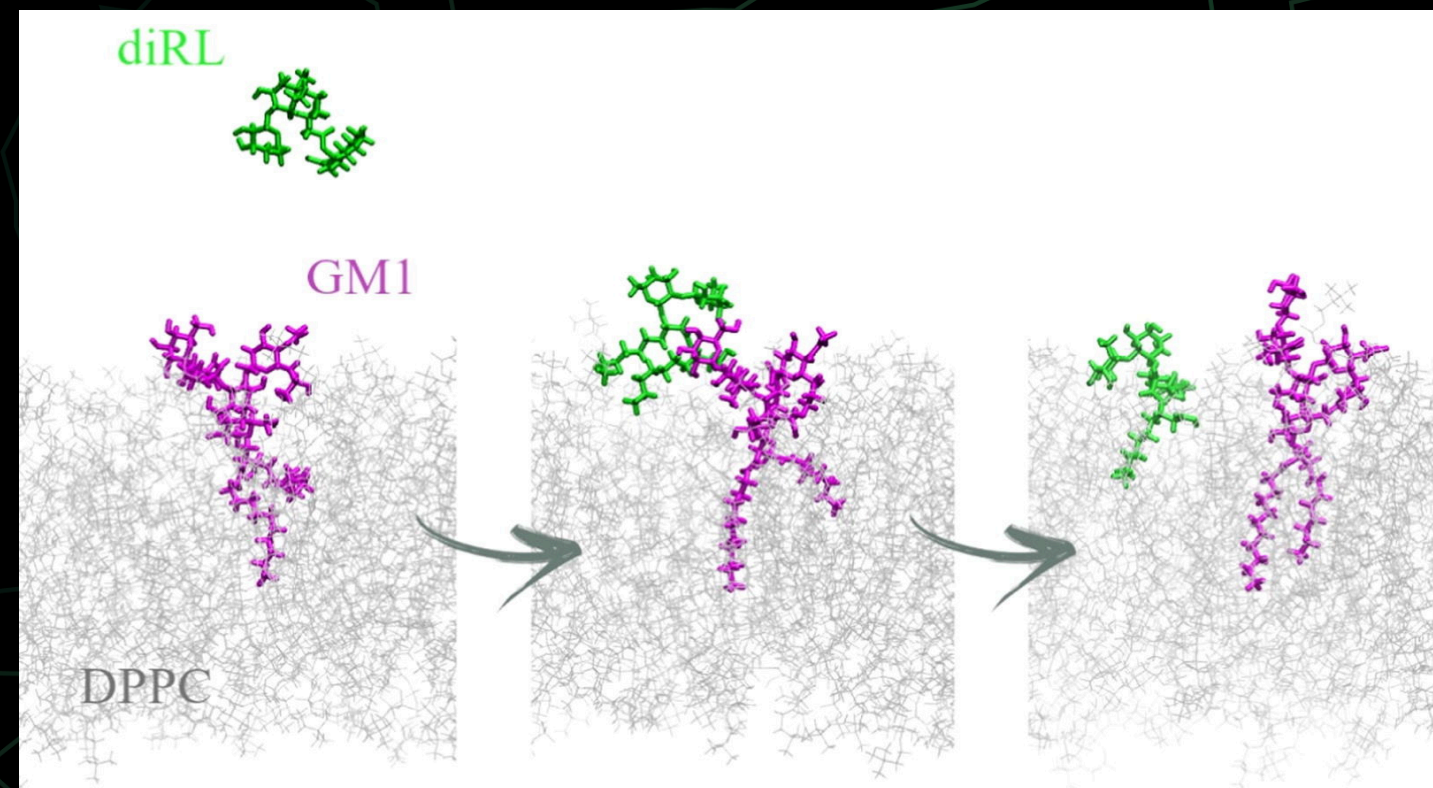
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Rondelli, V., et al.. *J. Colloid Interface Sci.* 616, 739, 2022. DOI: 10.1016/j.jcis.2022.02.120

Today we will be using Python and a Jupyter Notebook to analyse some reflectometry data.

- there are many Python packages to analyse reflectometry data: `refl1d`, `refnx`, **`anaklasis`**, `BornAgain`, `EasyReflectometry`, `genx`
- today we will use `refnx`, but the ideas you will learn are transferable

Today we will be using Python and a Jupyter Notebook to analyse some reflectometry data.

visit: <https://github.com/arm61/fasem>

click the "launch | binder" button

1. open and read/work through the `simple_fitting.ipynb` notebook, independently, use the "Run" button at the top to run code cells
2. together we will review the `simple_fitting.ipynb` notebook
3. work in pairs on the `bilayer_analysis.ipynb` notebook, completing the exercises where requested

ask questions!

thanks for listening!
questions?

- ORSO Reflectometry Tutorials: <http://reflectometry.org/learn/>
- Sivia, D. S. (2011). *Elementary Scattering Theory: For X-ray and Neutron Users*. Oxford University Press.
- Clifton, L.A., *et al.*. Lipid-Protein Interactions: Methods and Practices, **2003**, 201. 2019. DOI: 10.1007/978-1-4939-9512-7_11
- Gerelli, Y., *EPJ Web of Conference*, **236**, 04002, 2020. DOI: 10.1051/epjconf/202023604002

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