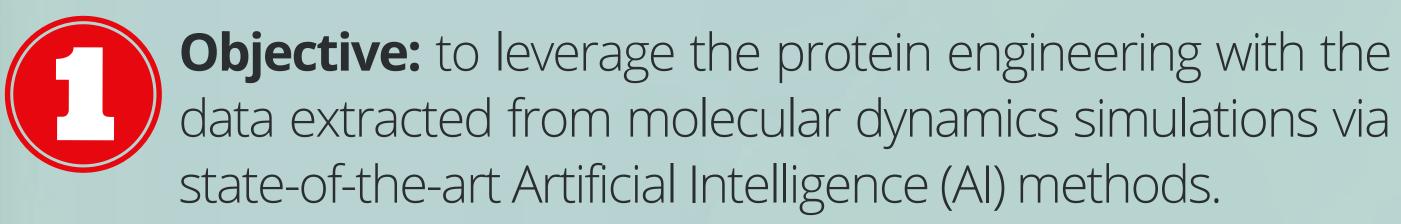
## Decoding Protein Dynamics using Artificial Intelligence

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and molecules.





Proteins form the basis of life, with all its great variety and efficiency. Wide range of functions in living organisms, from catalyzing metabolic reactions to building structures in cells and tissues.

Protein engineering develops advanced and valuable proteins for a vast range of applications in various fields.

Understating Dynamics — Understating Functionality

**Molecular Dynamics (MD)** a method for analyzing the physical motion of atoms

Results of MD are sequential snapshots states of the biomolecule at specific time intervals, represented 3D atomic coordinates.

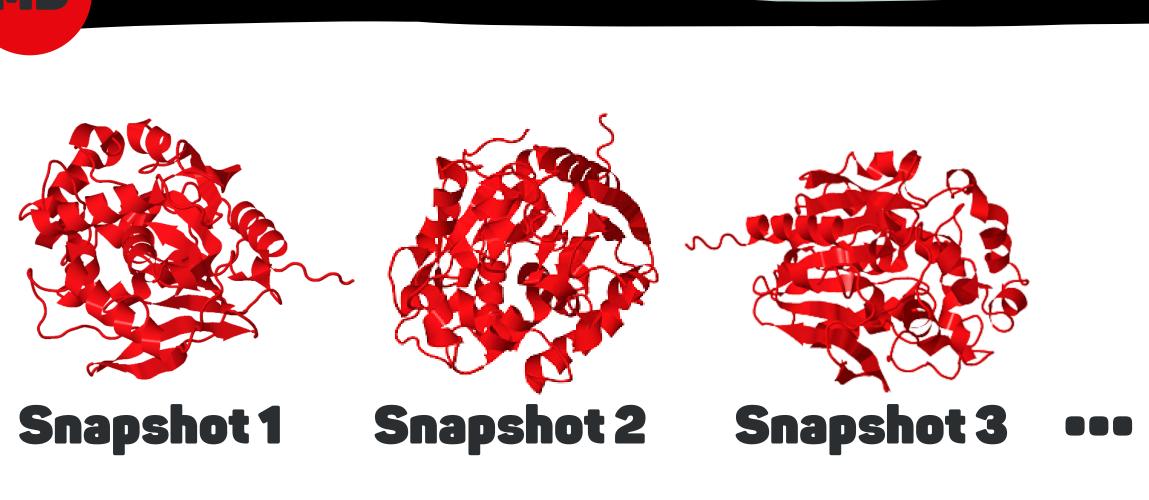
Proteins are moving!!!

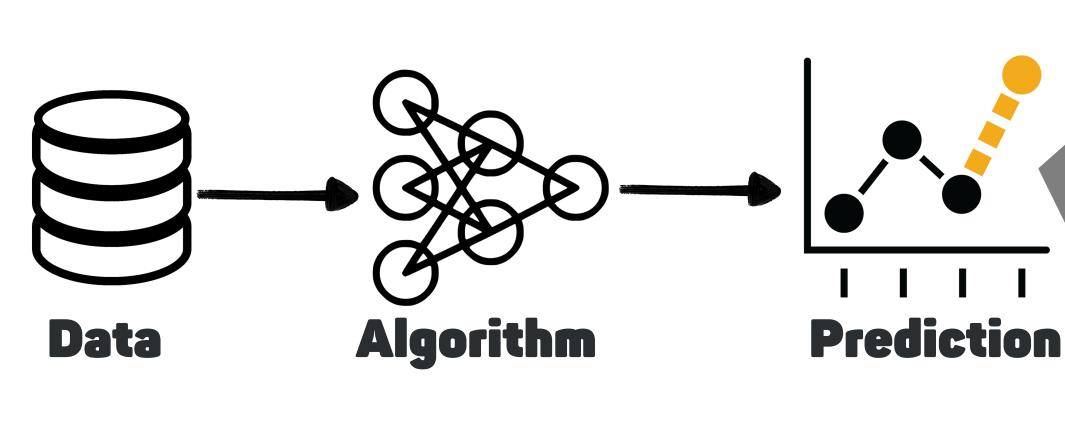
**Tremendous amount of data** Visual inspection insufficient Manually selected metrics **Human bias** 

what is the solution?









Interpretability

Machine learning (ML) can help us!

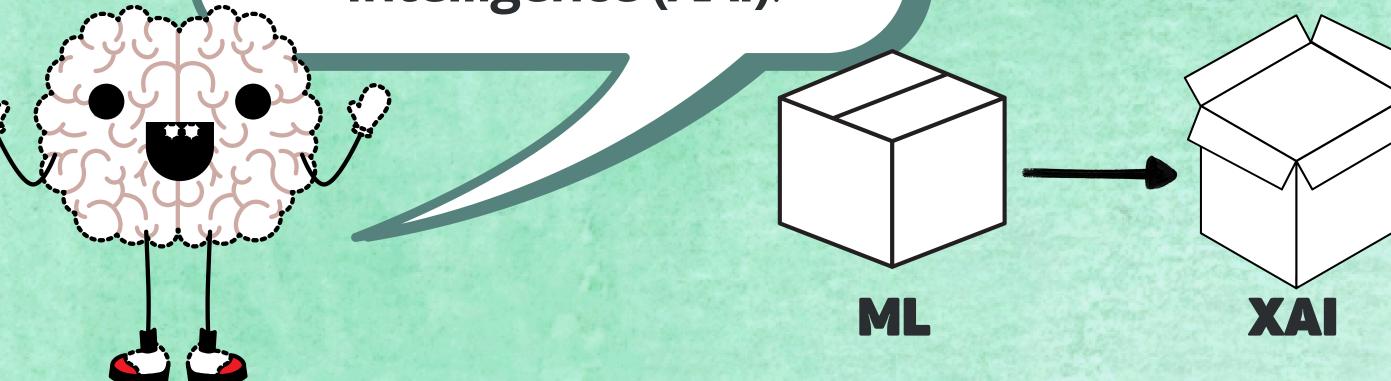
But we need another solution to find concealed features in the proteins!



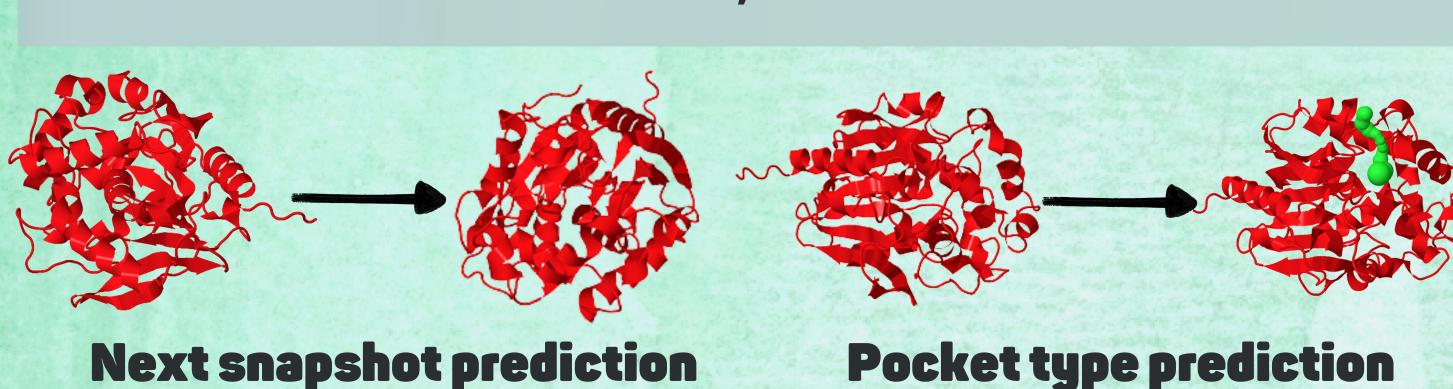
**Concealed features** 



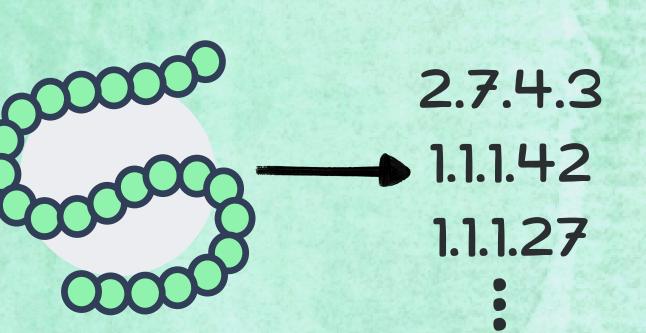
Yes! Explainable Artificial Intelligence (XAI)!



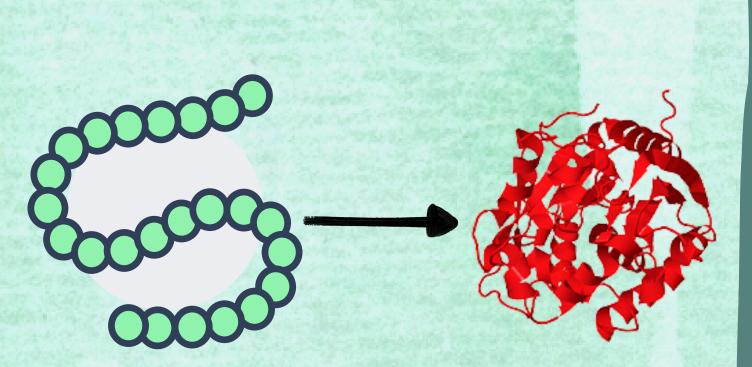
ML architecture transforms raw data into a training set to learn patterns and make decisions (predictions). Different ML architecture: CNN, RNN...



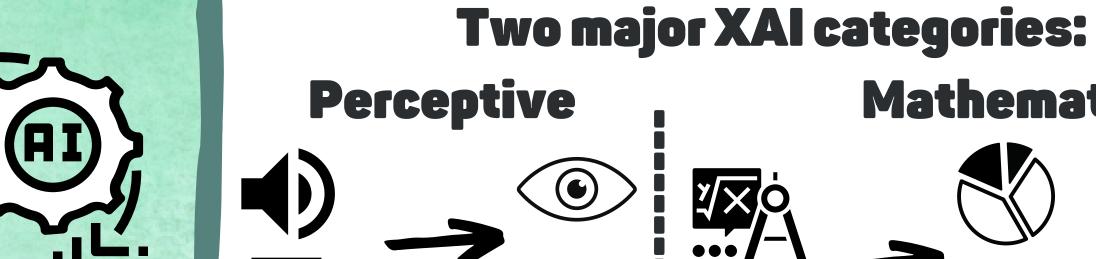
Next snapshot prediction

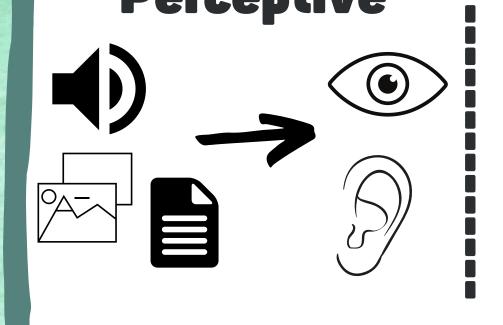


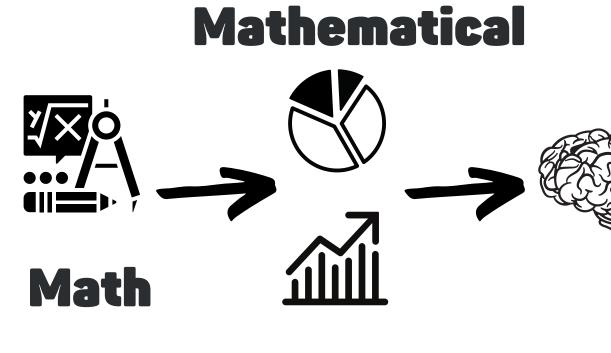
EC number prediction



3D structure prediction







Caver rapid, accurate, and fully automated calculation of protein tunnels.

Current version does not benefit from available MD simulations. **Next steps:** conduct MD simulations and their ML-based analyses

- tunnel motions
- various types of tunnels prediction
- best hotspots for protein engineering identification





LoopGrafter transplants loops between two GRAFTER structurally similar proteins.

Current version uses a simple B-factor model based on the parameters calculated in X-ray crystal structures to analyze the dynamics of the loops.

**Next steps:** conduct MD simulations and their ML-based analyses

- identification of interlinked loops
- loop ranking according to the desired property

