



CENTER FOR SCALABLE DATA ANALYTICS AND
ARTIFICIAL INTELLIGENCE

MDannotator – Visual Labeling of MD Simulations for Classification

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Gefördert durch:



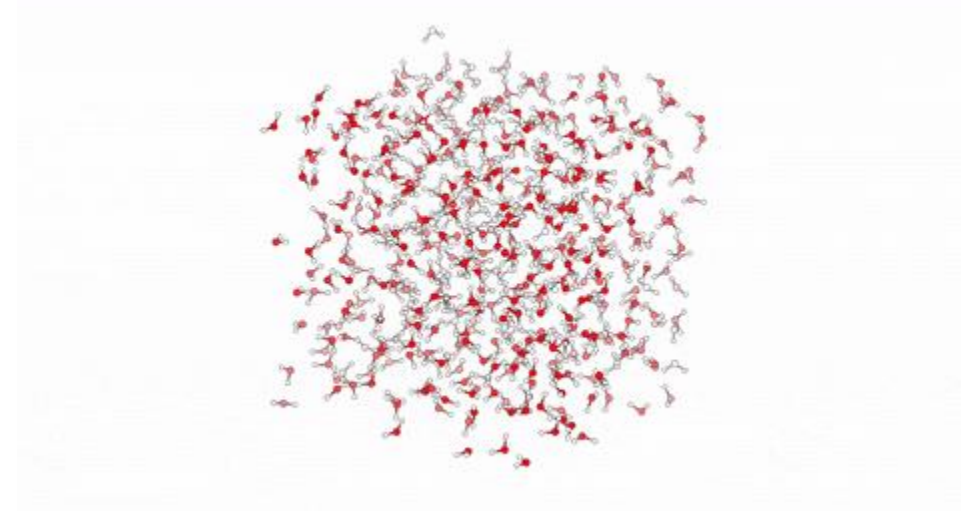
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Landtags beschlossenen Haushaltes.

MD Simulations?

- Structure prediction of molecules is cool, but their dynamic behavior can be really interesting:
 - Interaction with other molecules
 - Stable behavior in specific environments
 - Refolding events
- Can be very time-consuming
- Computationally still massive





MD Simulations?

Simulations can contain millions of frames.

- How to analyze and explore?
- **By Hand:**
 - Using poor PhD Students to force them to watch hours of trajectory data
 - Finding the needle in a hay stack

- **Metrics and Algorithms:**
 - Using poor PhD Students to pick the right metric
 - Might miss interesting spots



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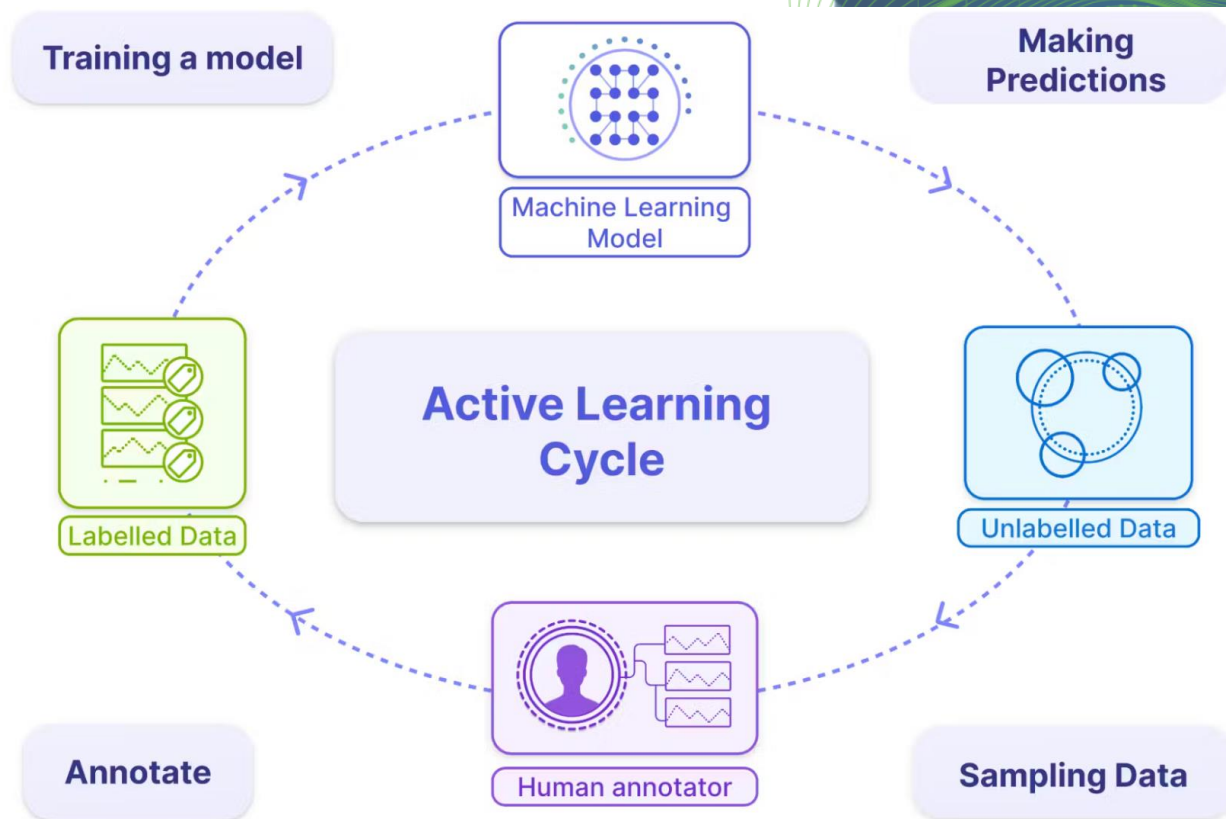
Might miss an interesting spot in the data due to boredom!

→ Can we use AI for making PhD students happy?

Data Sources?

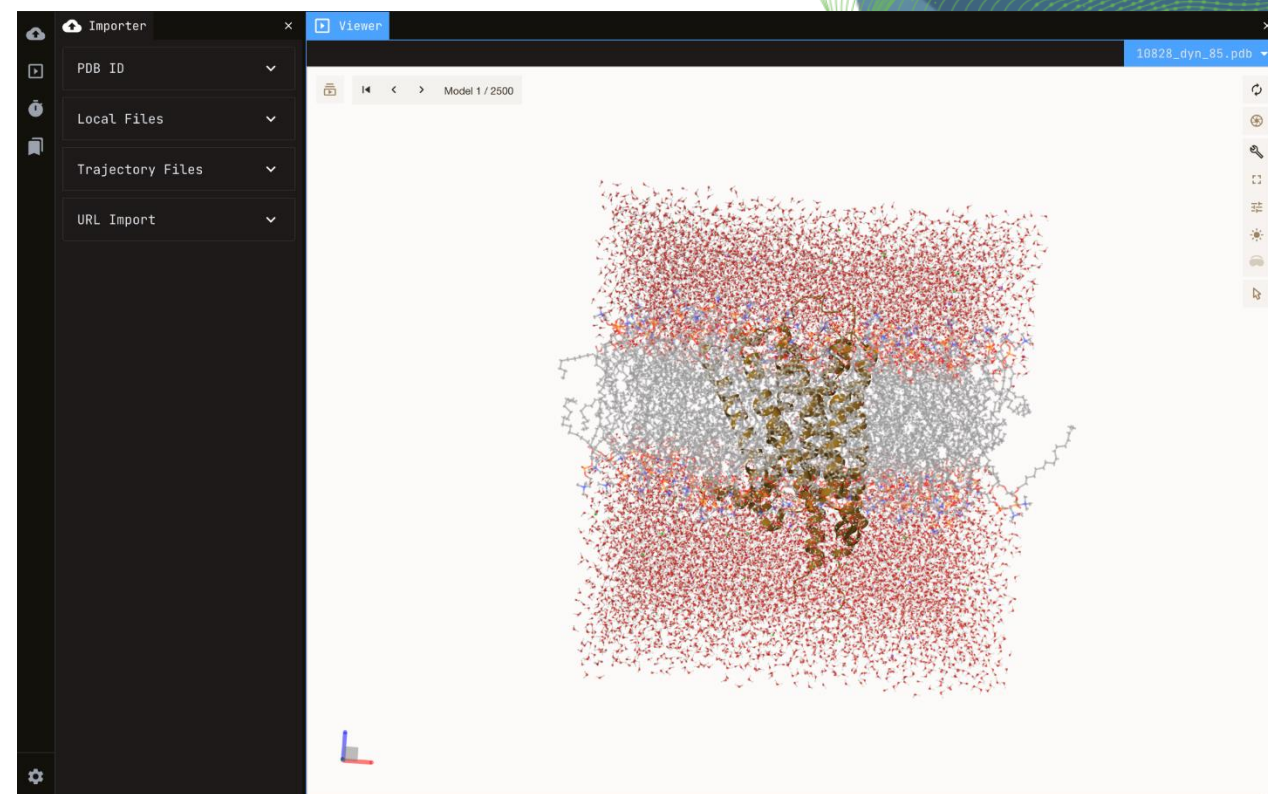
- We have large databases for MD simulations
- But: No labeled data for machine learning
- Use PhD students to create labeled data for a ML model:
 - Iteration loop with zero shot model and human annotator
 - Use metrics to detect uncertain areas in the latent space

We need an easy to use web based tool to convince people to do this → MDannotator



MDannotator

- Using MDsrv [1] as starting point:
 - Implementing a collaboration platform to share MD simulations
 - Comparison of simulation runs
 - Labeling mechanisms for presentation
 - Integrate metrics for exploration to support users
 - Develop a common format to describe events in MD simulations

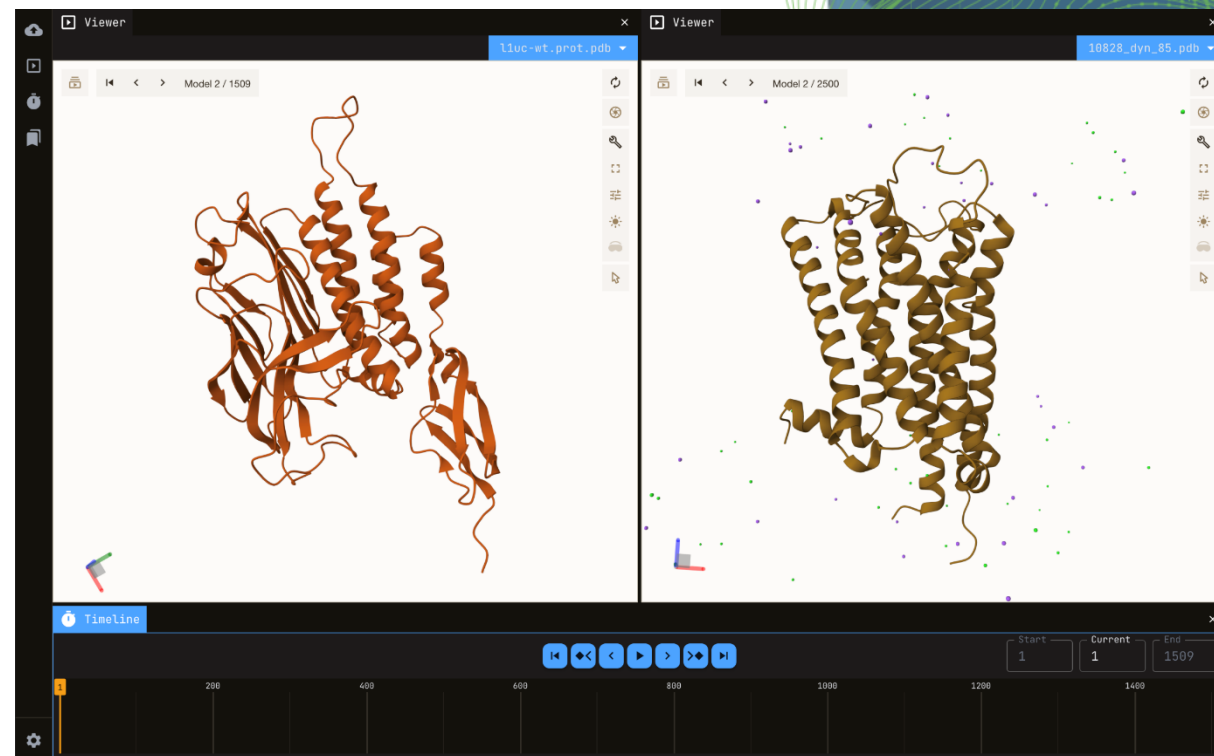


[1] Kampfrath, Michelle, et al. "MDsrv: visual sharing and analysis of molecular dynamics simulations." *Nucleic Acids Research* 50.W1 (2022): W483-W489.

MDannotator – Comparison of Simulations

- **Coupled view with shared interactions:**

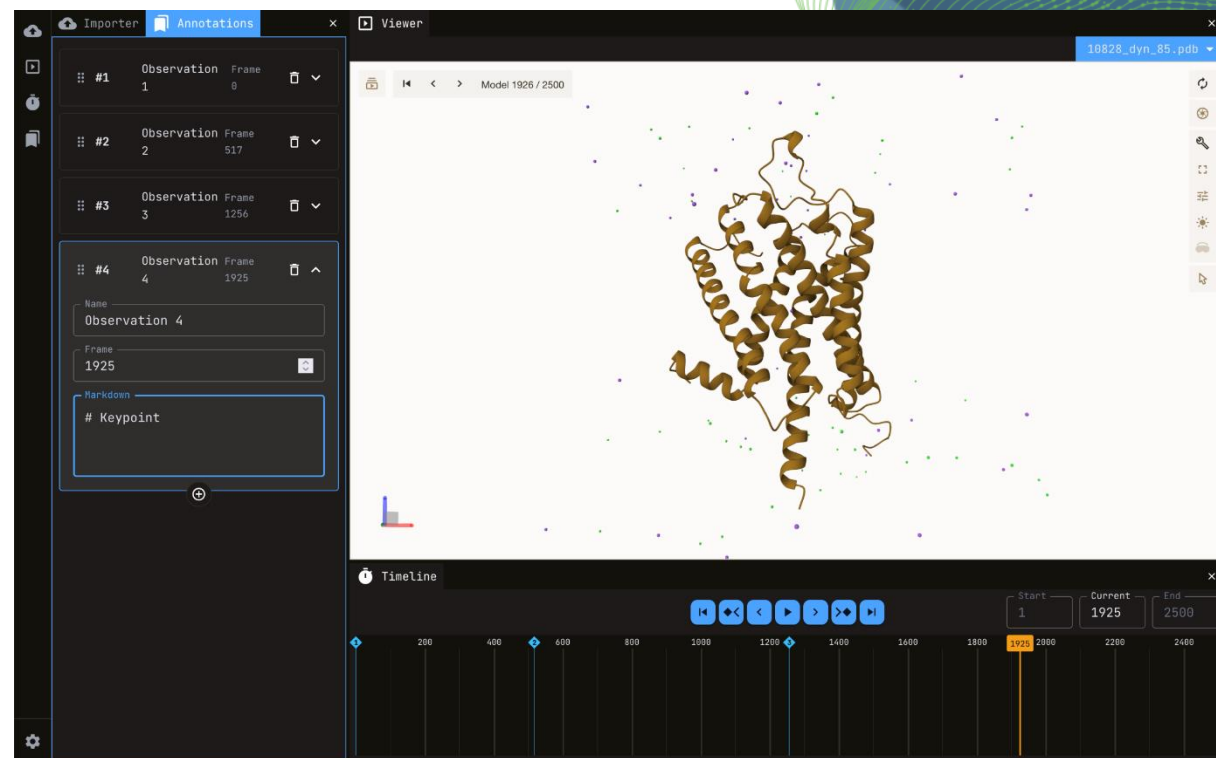
- Shared timeline as coordinate system between different simulations
 - Using it as an overview for labeled events
- Interpolation between different space coordinate systems



MDannotator – Labeling

- **Labels:**

- Single or multiple frames
- Metric values like RMSD
- Binding events of local components
-

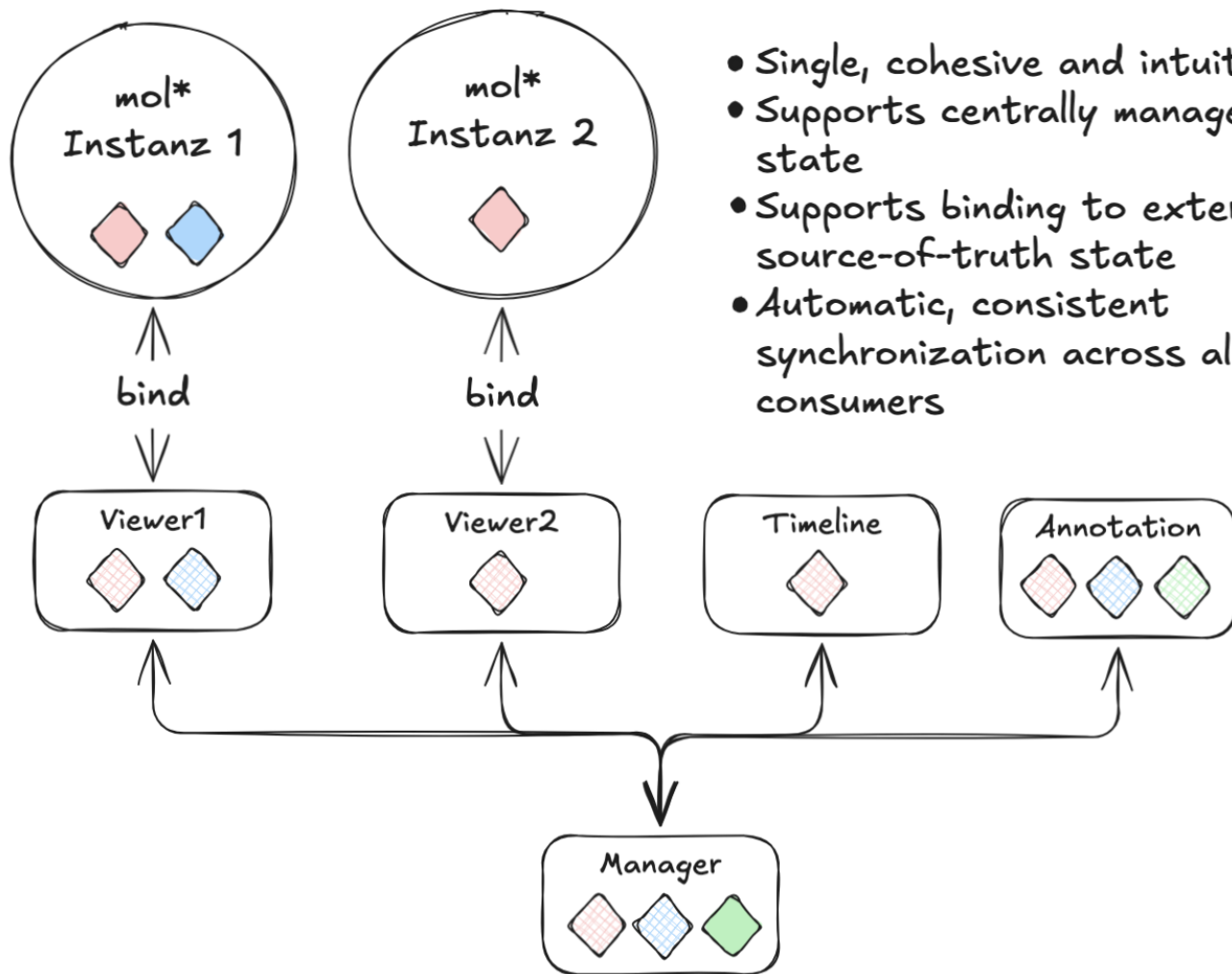


MDannotator - Architecture

Unified State Management with Optional Binding

- **We need a state management:**

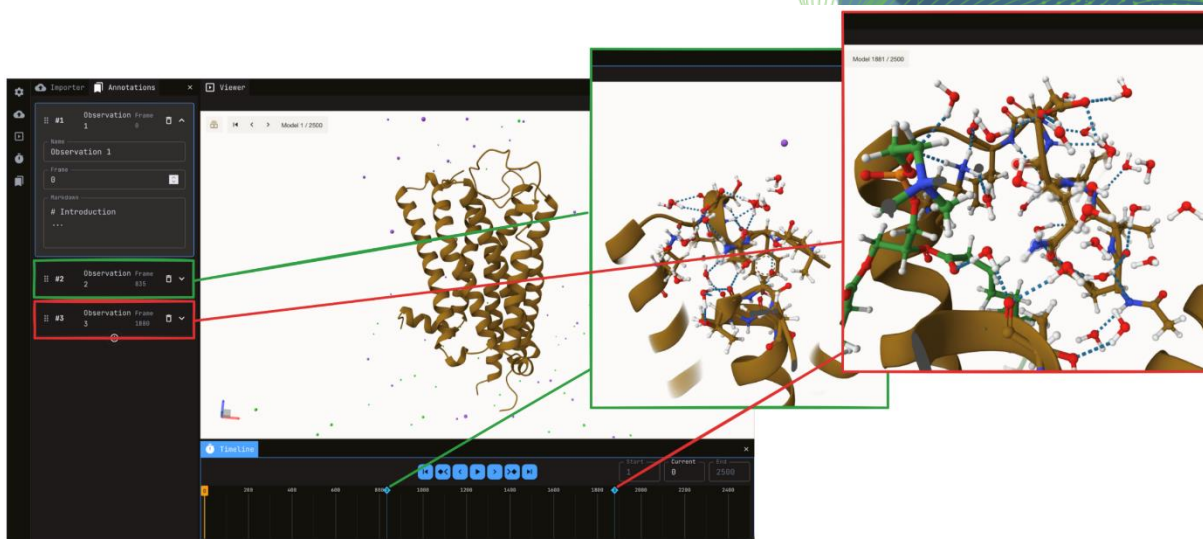
- Multiple views with different data
- Interaction can modify multiple states
- States can exist for varying lengths of time

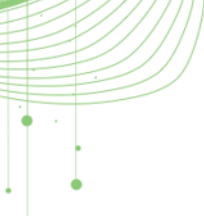


- Single, cohesive and intuitive API
- Supports centrally managed atom state
- Supports binding to external source-of-truth state
- Automatic, consistent synchronization across all consumers

MDannotator - Outlook

- Use user defined labels for the Active Learning approach to fine-tune a zero-shot approach
- Visualize predicted labels for inspection
- Generate uncertain labels for user correction
- Establish an machine readable format for MD simulation events for presentation and classification





Thanks for listening!

Questions?

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