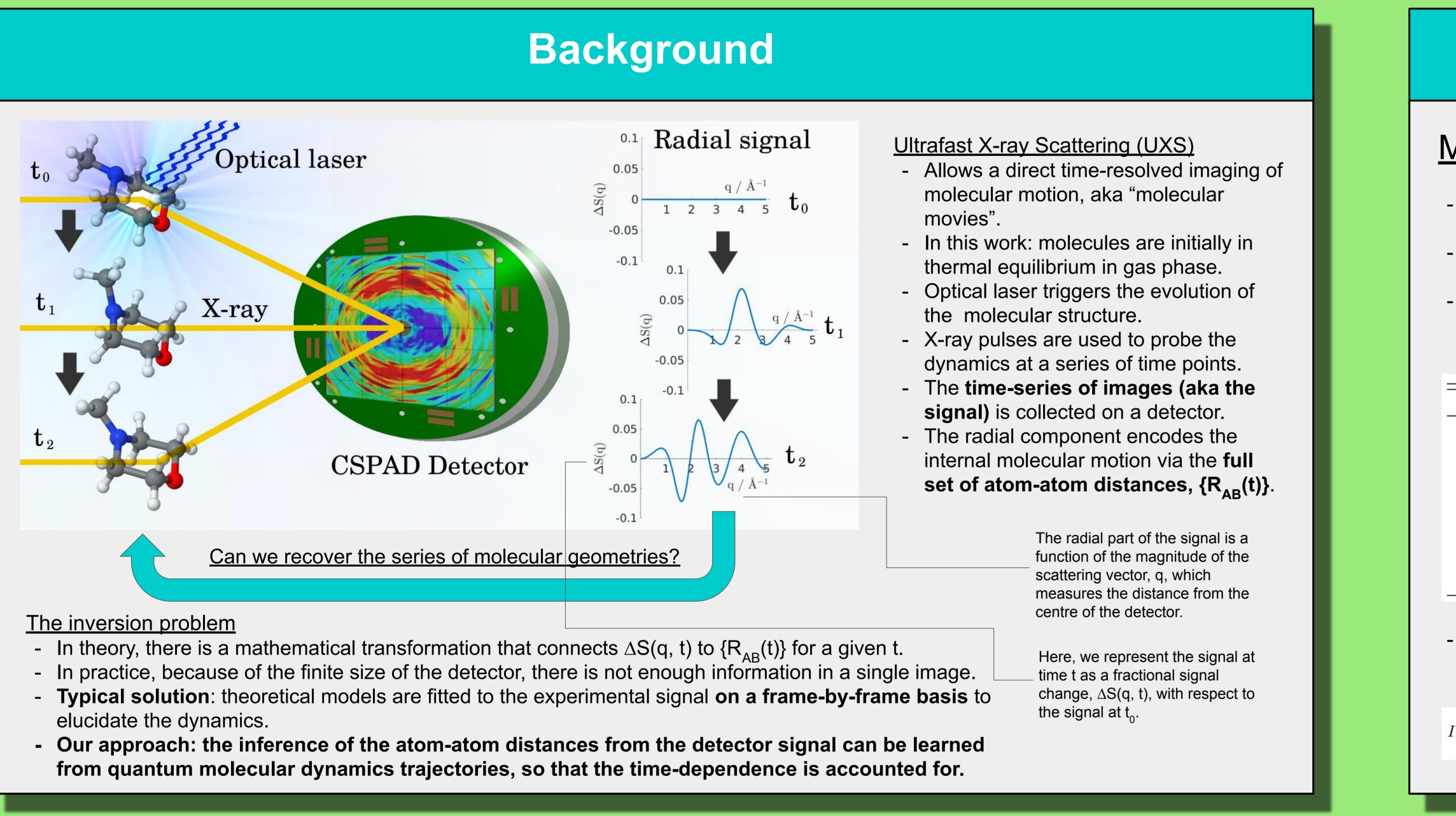
Inversion of Ultrafast X-ray Scattering with Dynamics Constraints

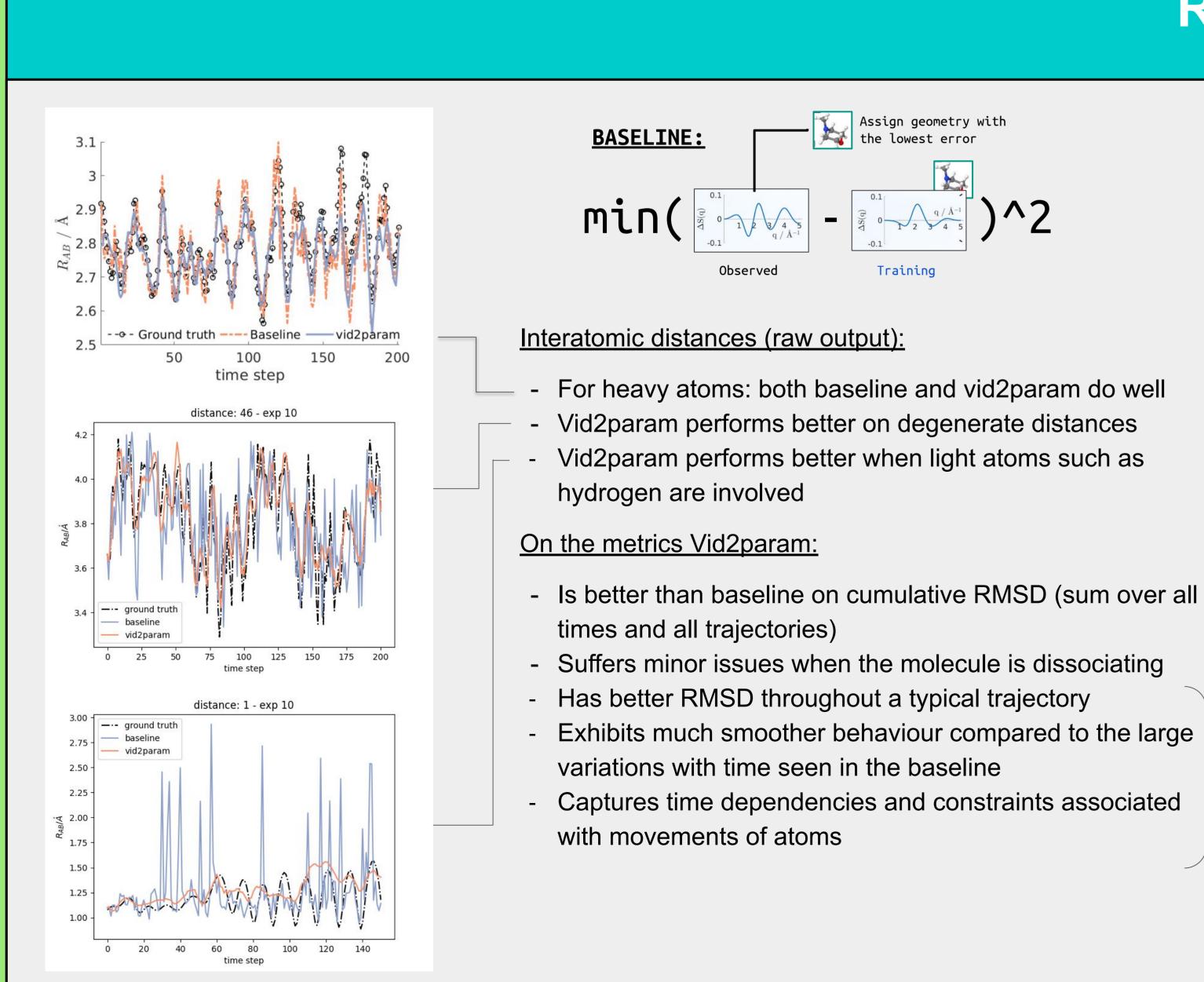
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Results

🐨 the lowest error Method $\left(\log_{10}\right)$ vid2param (*) indicates that dissociation has baseline been removed from the data set. Training C_2H_4 * C_2H_4 CS_2^* NMM RMSD vid2param 0.03 ----- SSR vid2param 0.02 BMSD nic <u>5 0.01</u> 50



Methods

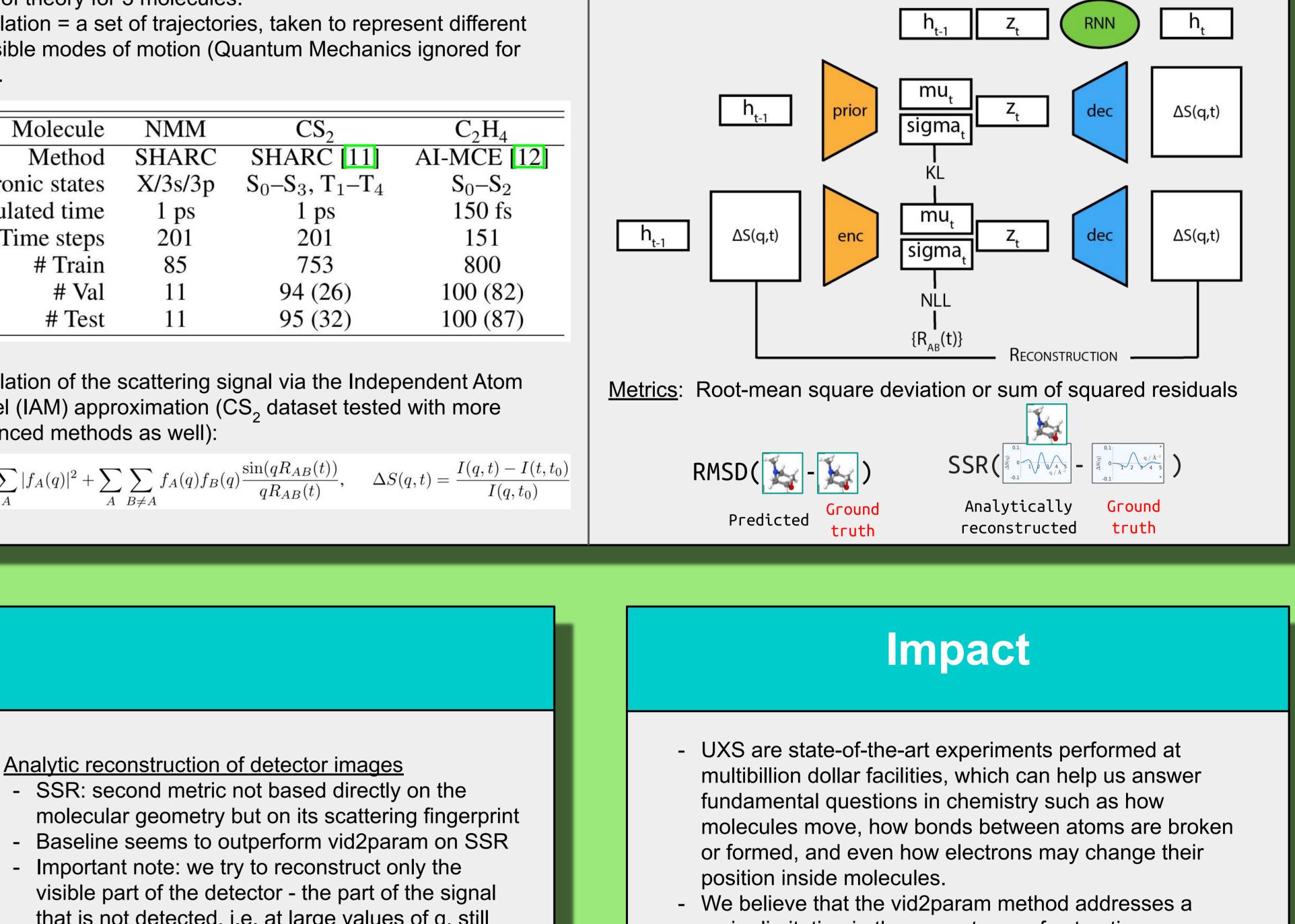
Moleclar dynamics and X-ray scattering

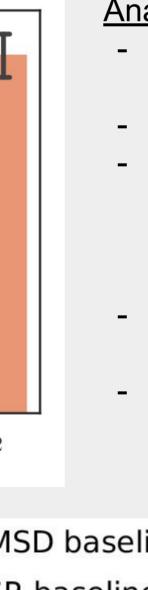
- We utilise simulated data to train, validate and test our model (separately for each molecule).
- Quantum molecular dynamics simulations based in CASSCF level of theory for 3 molecules.
- Simulation = a set of trajectories, taken to represent different plausible modes of motion (Quantum Mechanics ignored for now).

Molecule	NMM	CS ₂	C_2H_4
Method	SHARC	SHARC [11]	AI-MCE 12
Electronic states	X/3s/3p	$S_0 - S_3, T_1 - T_4$	S_0-S_2
Simulated time	1 ps	1 ps	150 fs
Time steps	201	201	151
# Train	85	753	800
# Val	11	94 (26)	100 (82)
# Test	11	95 (32)	100 (87)

Simulation of the scattering signal via the Independent Atom Model (IAM) approximation (CS₂ dataset tested with more advanced methods as well):

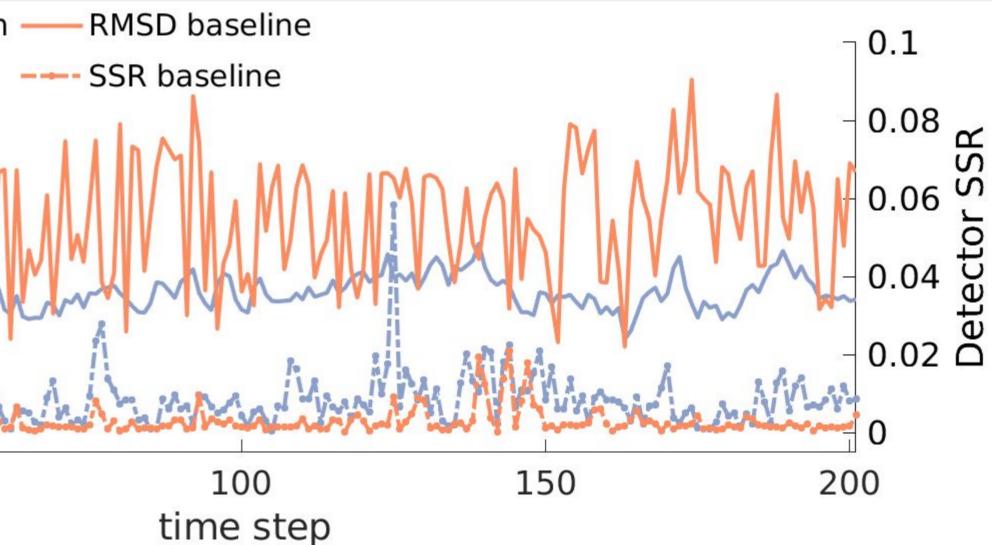
$$q,t) = \sum_{A} |f_A(q)|^2 + \sum_{A} \sum_{B \neq A} f_A(q) f_B(q) \frac{\sin(qR_{AB}(t))}{qR_{AB}(t)}, \qquad \Delta S(q,t) = \frac{I(q,t) - I(t,t)}{I(q,t_0)}$$





visible part of the detector - the part of the signal that is not detected, i.e. at large values of q, still contains vital information. - Better SSR in the visible part does not guarantee better SSR overall

 vid2param learns from previous time points, hence, producing a better estimate for the molecular structure (RMSD) regardless of the SSR



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Inversion with dynamics constraints

- Encode detector signal into a latent space z_t - Include information about previous dynamics h_{t-1} - Enforce part of that latent space to be atom-to-atom distances

major limitation in the current way of extracting information from UXS data.

Immediate applications in other experimental

techniques for studying ultrafast processes where direct (analytical) inversion is not possible.

Acknowledgements

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Datasets, expanded results and source code can be seen at https://sites.google.com/view/mlscattering/