18th June 2025 - 2:00 p.m. CFEL-bldg. 99, seminar room IV

Carl Caleman

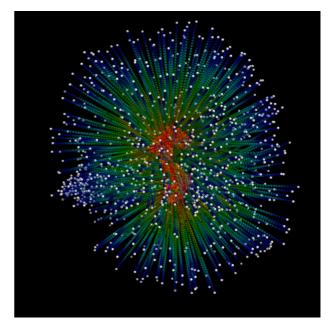
Department of Physics and Astronomy, Uppsala University, Uppsala, Sweden

Protein structure classification based on X-ray-laser-induced Coulomb explosion

We present simulations of Coulomb explosion dynamics due to fast ionization induced by high-intensity x-rays in six proteins that share similar atomic content and shape. We followed and projected the trajectory of the fragments onto a virtual detector, providing a unique explosion footprint. After collecting 500 explosion footprints for each protein, we utilized principal component analysis and *t*distributed stochastic neighbor embedding to classify these. Results show that the classification algorithms were able to separate proteins on the basis of explosion footprints

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from structurally similar proteins into distinct groups. The explosion footprints, therefore, provide a unique identifier for each protein. We envision that method could be used concurrently with single-particle coherent imaging experiments to provide additional information on shape, mass, or conformation.