FEL applications: coulomb explosion imaging of proteins

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Structure and function





Serial Femtosecond Crystallography



Intense X-ray pulses destroys the structure



0.00 fs



Intense X-ray pulses destroys the structure



0 fs

35 fs

70 fs





Diffraction before destruction





Diffraction before destruction



Crystallography depends crystals





Single particle imaging using XFEL

Protein sample





Exploding sample







DIFFICTED BY PROFESSOR CALEMAN















HiPIP dimer and monomer



MS2 virus dimer symmetric and asymmetric



HiPIP dimer and monomer

TSNE - dimensionality reduction algorithm







TSNE - dimensionality reduction algorithm



MS2 virus dimer symmetric and asymmetric





TSNE - dimensionality reduction algorithm



MS2 virus dimer symmetric and asymmetric



The path forward is to combine diffraction, ion maps, alpha fold.

Can we even do without high resolutions (hard x-rays) diffraction?

Tack.