## **ULTRA FAST IMAGING**

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The current bottleneck in the atomic resolution imaging of biological systems is a fundamental need for crystals. This limits the scope of detailed structural studies to macromolecules and macromolecular assemblies which can be crystallised. X-ray freeelectron lasers have the potential of changing this picture. Theory predicts that with very intense and very short X-ray pulses, a single diffraction image could be recorded from a macromolecule, virus, or a protein cluster before damage-induced movements destroy the sample. This type of imaging would not require a crystal. Three-dimensional reconstruction of the structure may be possible with reproducible samples, using a new sample molecule for each shot. When a reproducible sample scatters a sufficiently large number of photons for its orientation to be determined, averaging techniques can be applied to extend the resolution in a redundant data set. The talk will describe concepts, models and progress with the project.