

## Prof. Brandon Ruotolo

University of Michigan

Structural Biology in the Gas Phase: New Approaches for Conformationally-selective Inhibitor Screening and Multiprotein Topology

Within each living organism proteins are at work, carrying out activities which impact every aspect of cellular function from synthesis to cell death. A key factor in varied protein functionality is their ability to self-assemble, creating vast molecular 'machines' capable of performing intricate, highly specialized tasks. To rapidly discover the structural properties of such assemblies, especially when bound to therapeutic small molecules, is a grand challenge for structural biology which is nearly insurmountable with current tools. In this presentation, I discuss recent developments in ion mobilitymass spectrometry (IM-MS) methods aimed at bridging this gap in basic technology. The IM-MS approaches developed by my group exploit a series of highly-accurate, gas-phase measurements of protein size and mass to rapidly determine multiprotein architectures and stabilities when bound to small molecules, without the need for covalent labels or tagging, and consuming 10-100 times less protein Recent IM-MS than almost any other label-free technology. applications, including: Multi-protein structure modelling, highthroughput discovery of conformationally-selective inhibitors and the rapid structural characterization of biotherapeutics will be discussed.

Thursday November 19

12:15 p.m. 1315 Chemistry

Coffee & cookies at 12 p.m. outside 1315

