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On the cover: Multidisciplinary approaches are rapidly emerging that can be used to interface the data gathered by interaction proteomics and structural biology. As an illustrative example, depicted on the cover, the composition, topology and structure of the yeast vacuolar ATPase, which is a large multi-subunit protein complex that can acidify a variety of intracellular compartments, was elucidated by a combination of quantitative proteomics, electron microscopy, NMR, mass spectrometry and computational molecular modeling. Other examples of such multidisciplinary approaches are featured throughout this special issue.

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