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On the cover: Proteomic and electron microscopy survey of large assemblies in macrophage cytoplasm. Sucrose gradient fractions of macrophage cell lysate were analyzed to identify constituent proteins by mass spectrometry and visualize them by electron microscopy. The image shows the superposition of (1) background: transmission (metal shadowed replica) electron micrograph of macrophage cell interior; (2) center, grey boxes: example two-dimensional averages of putative macromolecular complexes as analysed by single particle electron microscopy; and (3) colored balls connected by lines: example functional interaction network of components of a fraction (generated using String protein interactions database). For details, see article by Bohumil Maco *et al.*, pages M111.008763, 1–9.

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