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On the cover: Structure of lipid-bound apoE, zooming in on novel glycosylation site, Ser²⁹⁰. The apoE structure was kindly provided by K. Weisgraber. The protein backbone is rendered in ribbon representation, with the coloring following the chain from blue for the N-terminus to red for the C-terminus. Animation was generated using Discovery Studio 2.0 by Accelrys Software Inc with rendering by Pov-Ray. For further details see Lee et al, pages 1968–1981.

- 1898 **Comprehensive Characterization of Methicillin-resistant *Staphylococcus aureus* subsp. *aureus* COL Secretome by Two-dimensional Liquid Chromatography and Mass Spectrometry**
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- 1937 **Quantitative Proteomics Analysis of the Nuclear Fraction of Human CD4⁺ Cells in the Early Phases of IL-4-induced Th2 Differentiation**
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- 1991 **Identification of Cellular Targets in Human Intrahepatic Cholangiocarcinoma Using Laser Microdissection and Accurate Mass and Time Tag Proteomics**
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