Molecular & Cellular PROTEOMICS
Vol. 7, No. 1 January 2008

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Research

On the cover, new strategies for characterization of interacting partners of protein complexes using affinity purification and SILAC-based quantitative MS analysis. Cells stably expressing Rpn11-HTBH were grown in light (L) medium, while cells stably expressing the HTBH tag alone were grown in heavy (H) medium. Two sample preparation strategies were employed: (*a*) PAM (purification after mixing) and (*b*) MAP (mixing after purification). Four types of proteins are classified here based on characteristic patterns of their SILAC ratios. For details, see the article by Wang and Huang, pages 46–57.

[S] Online version of this article contains supplemental material.

HUPO Views

Information