

Research

- 1046 **Unconventional Secretion is a Major Contributor of Cancer Cell Line Secretomes**
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On the cover: Microarrays of SH2 and PTB domains were used to generate unbiased, quantitative interaction maps for most of the human Receptor Tyrosine Kinases, as well as upstream adaptor proteins. In addition to uncovering many previously unrecognized interactions, a systematic analysis of the data revealed a significant link between the connectivity of receptors and adaptor proteins and their intrinsic oncogenicity. For details, see article by Grigoriy Koytiger, *et al.*, pages 1204–1213.

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