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On the cover: This map of the HSP90 interactome was generated using immobilized broad spectrum kinase inhibitors and HSP90 co-immunoprecipitation in conjunction with HSP90 inhibition and quantitative mass spectrometry. HSP90 isoforms are shown in the center, surrounded by co-chaperones, kinase in the next layer and other proteins of diverse functions in the outer layer. For details, see article by Bernhard Kuster *et al.*, pages M111.016675, pages 1–14.

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