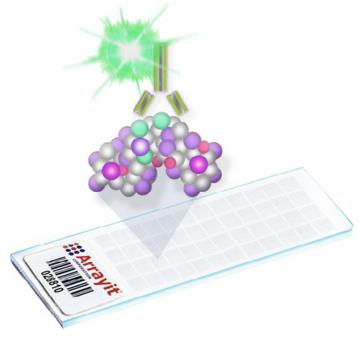


Protein Microarrays Journal Club

Arnaud Monnard



August 26th 2014 Aguzzi lab

Outline

Manufacturing of microarrays

Contact Printing

- Pin Printing
- Microstamping
- Flow Printing

Lithography

- Photolithography
- Electron Beam
- Dip Pen Nanolithography

Non Contact Printing

- Thermal
- Piezo
- Valve

Cell Free

- PISA
- NAPPA
- In Situ Puromycin Capture

Publications

Proof of Concepts

- MacBeath et al. Science 2000
- Chen et al. BioTechniques 2006

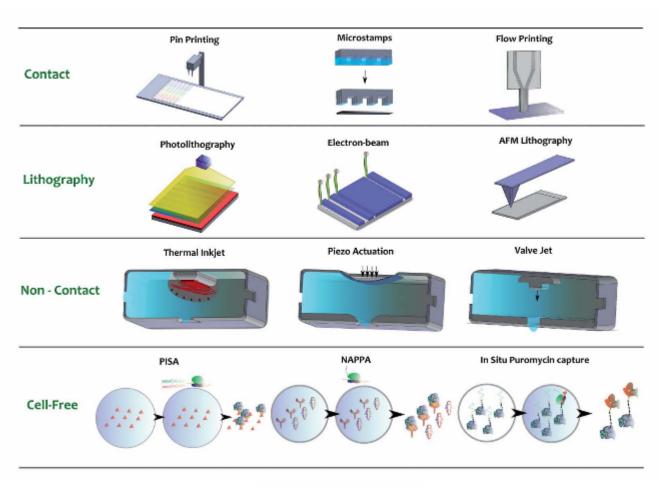
Applications

- Jones et al. Nature 2006
- Mok et al. Nature Protocols 2009
- Gagni et al. Biosensors and Bioelectronics 2013

Introduction

- First antibody array in 1983 followed by developement in the late 90's
- Use principles and technologies of DNA microarrays
 - Minimal sample consumption
 - High throughput
- Bypasses the limitation of DNA microarrays

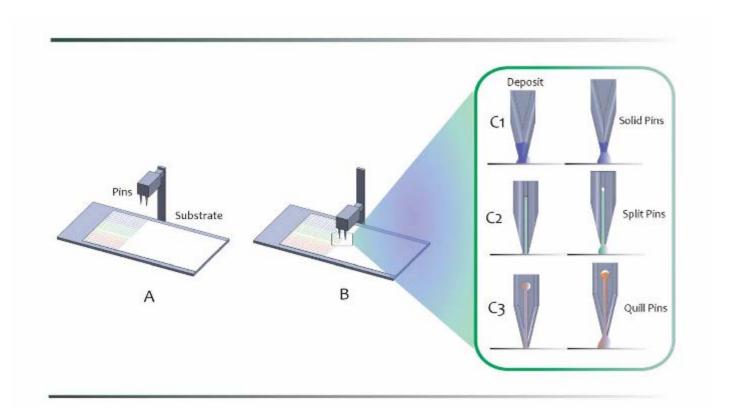
Protein Microarray Manufacturing Technologies



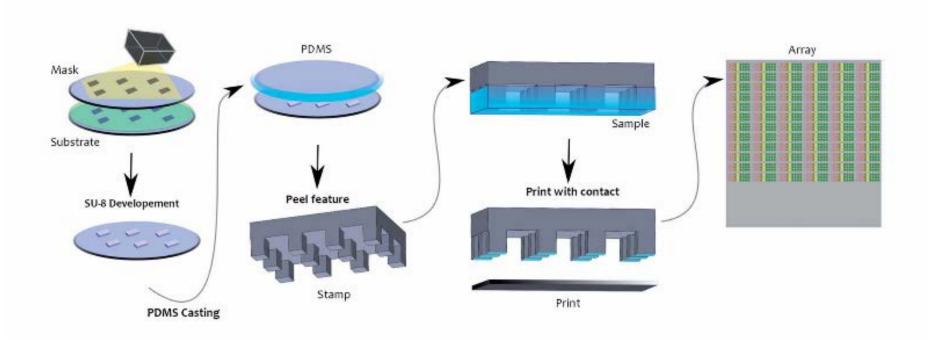
Contact Printing



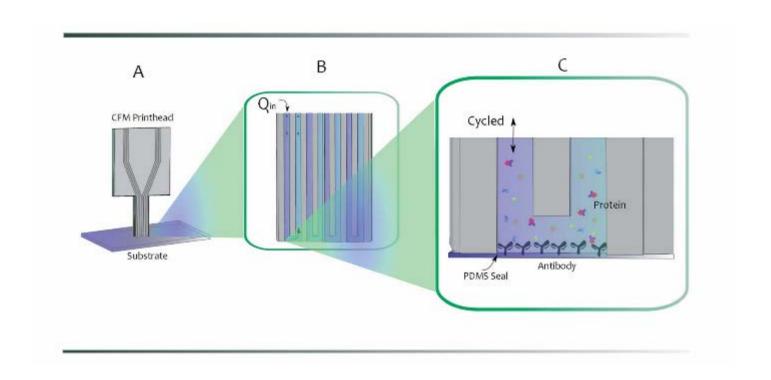
Pin Printing



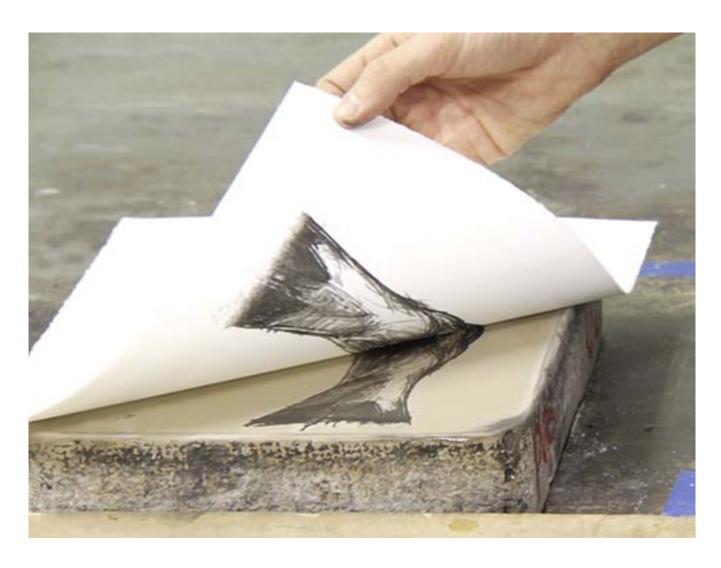
Microstamping



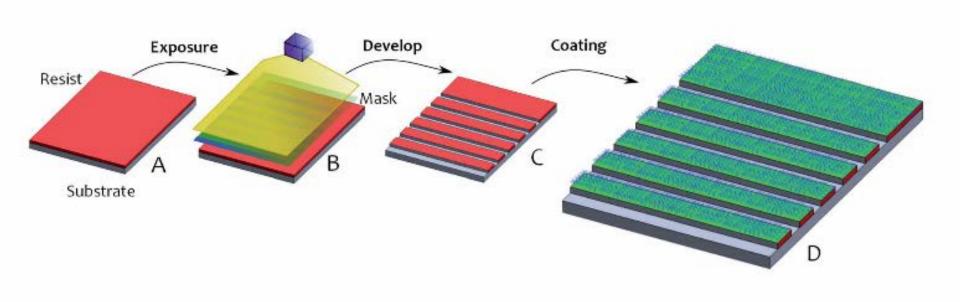
Flow Printing



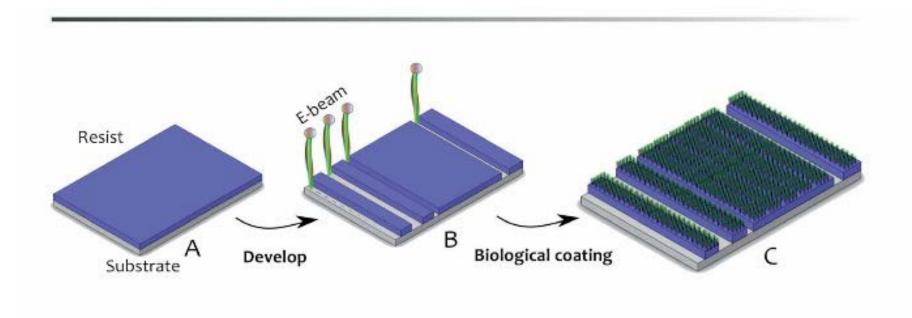
Lithography



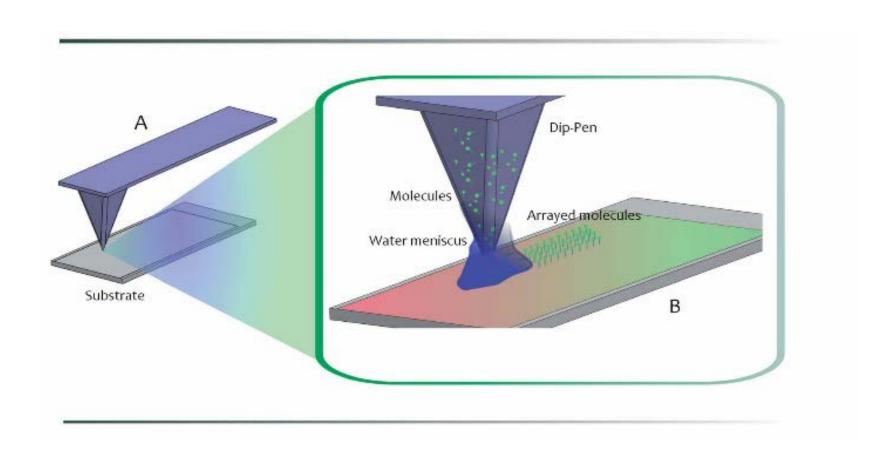
Photolithography



Electron Beam



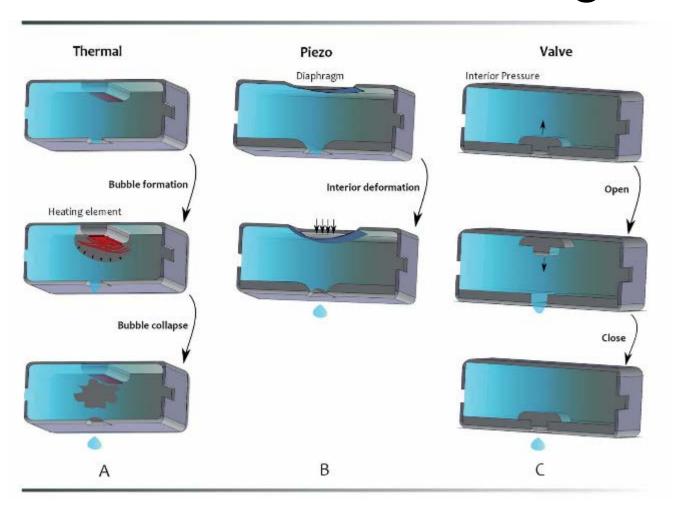
Dip Pen Nanolithography



Non-Contact Printing

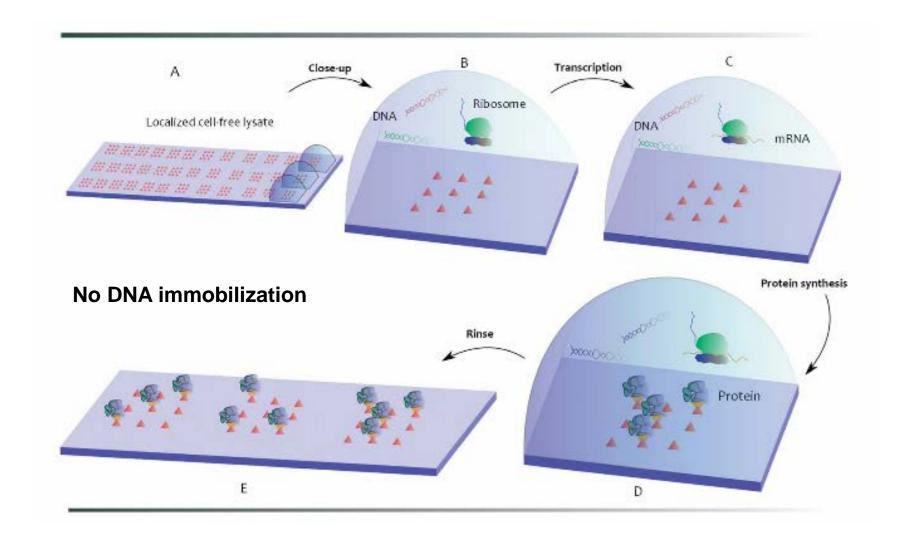


Non contact Printing

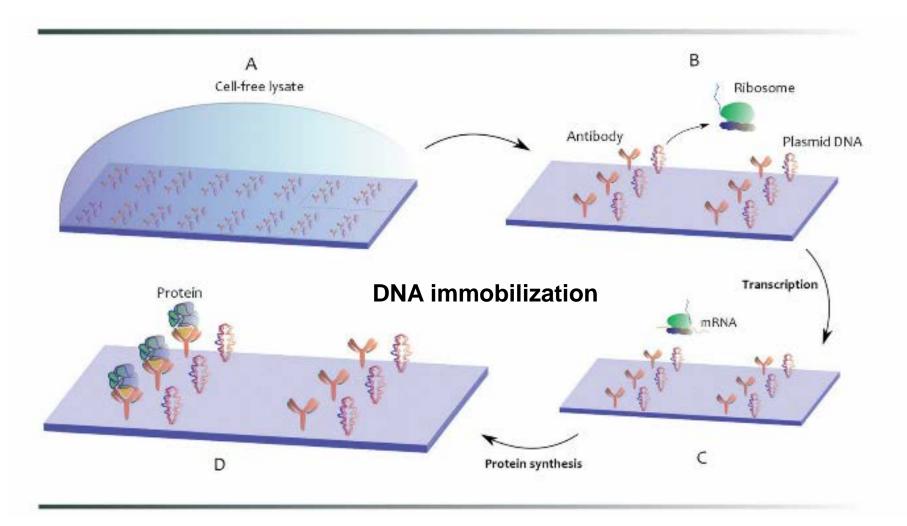


Cell Free Systems

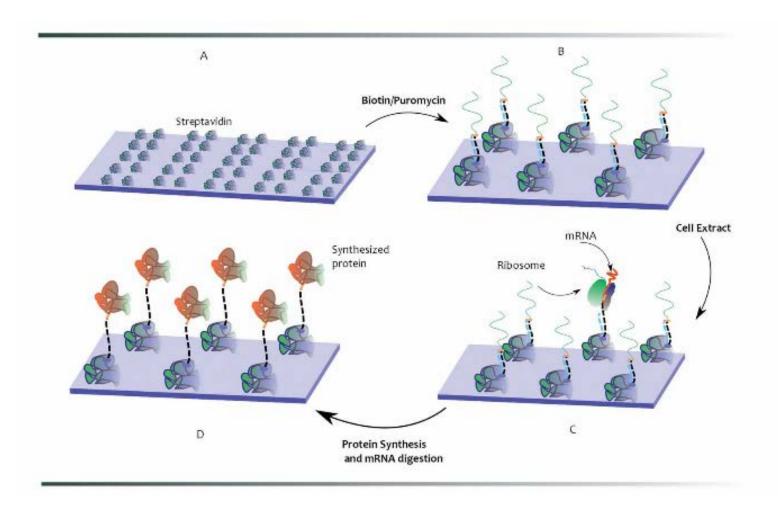
Protein In Situ Array (PISA)



NAPPA



In Situ Puromycin Capture



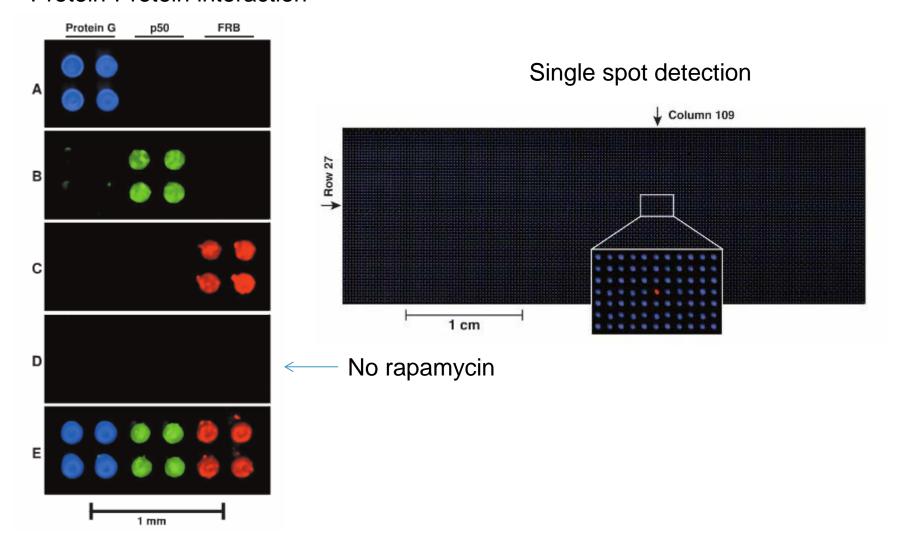


Printing Proteins as Microarrays for High-Throughput Function Determination

Gavin MacBeath and Stuart L. Schreiber Science 289, 1760 (2000); DOI: 10.1126/science.289.5485.1760

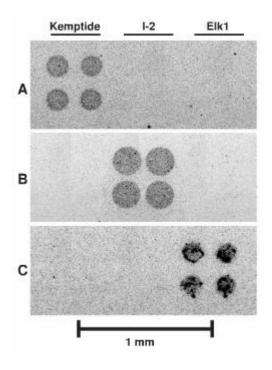
Proof of Concept

Protein-Protein interaction

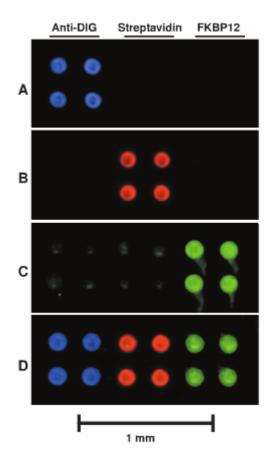


Proof of Concept

Detection of kinase substrates



Detection of small molecules targets



Techniques Essay

Protein Microarrays

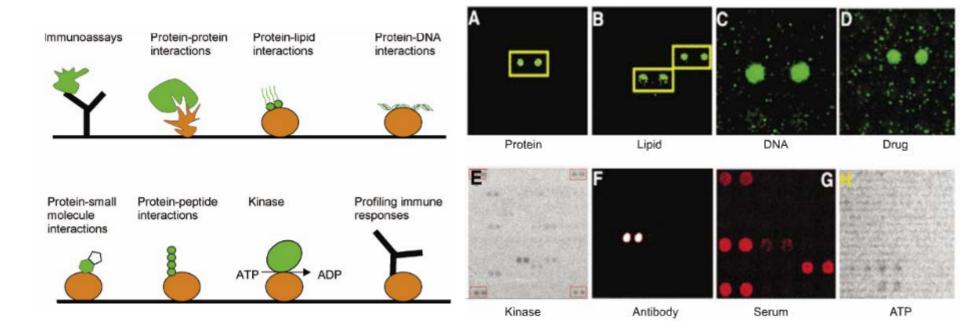
Chien-Sheng Chen and Heng Zhu

Department of Pharmacology and Molecular Sciences/High-Throughput Biology Center, Johns Hopkins University School of Medicine, Baltimore, MD, USA

Vol. 40, No. 4 (2006)

BioTechniques 423

Applications

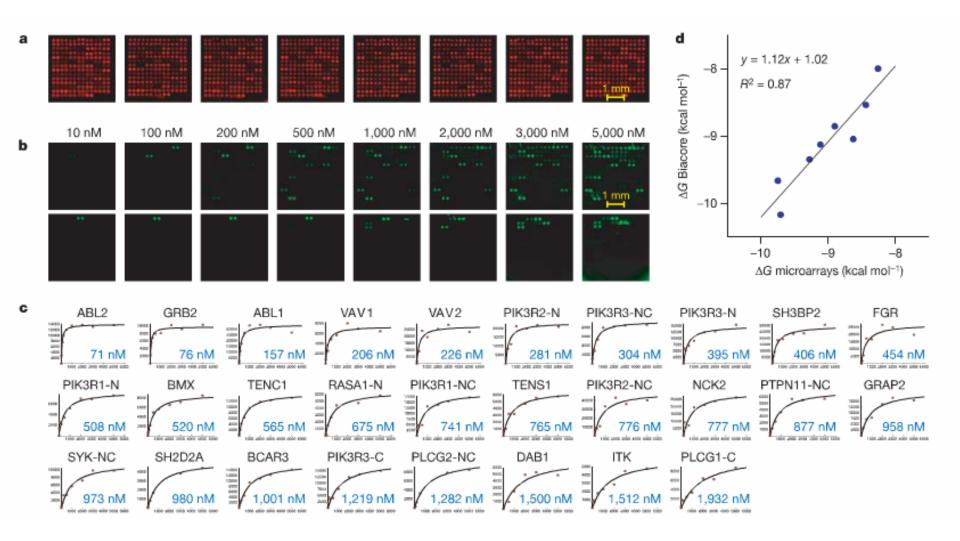


ARTICLES

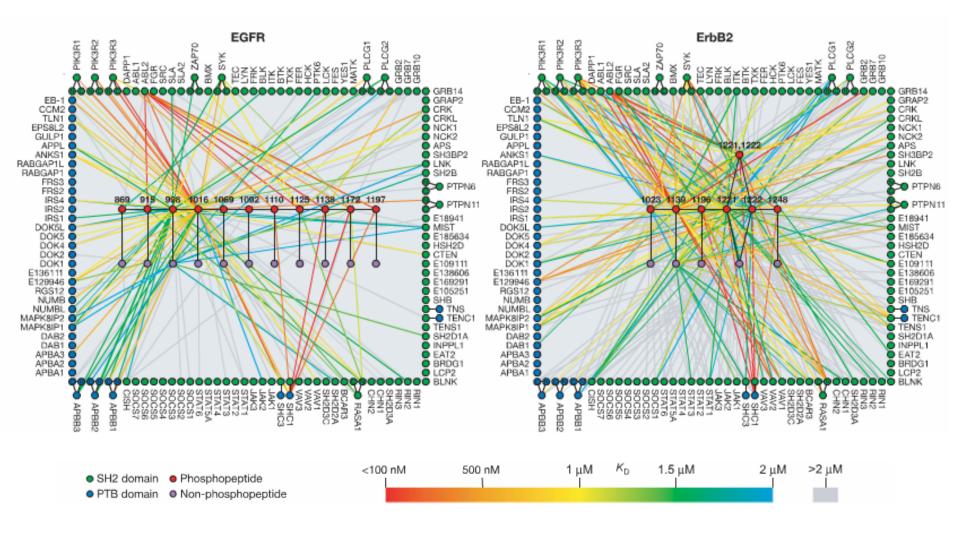
A quantitative protein interaction network for the ErbB receptors using protein microarrays

Richard B. Jones¹*, Andrew Gordus^{1,2}*, Jordan A. Krall¹ & Gavin MacBeath¹

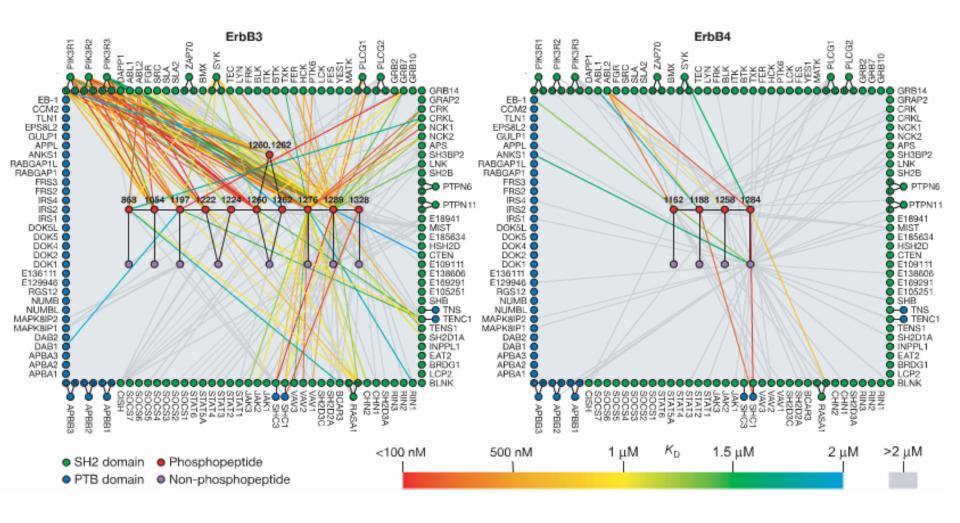
Binding Affinity



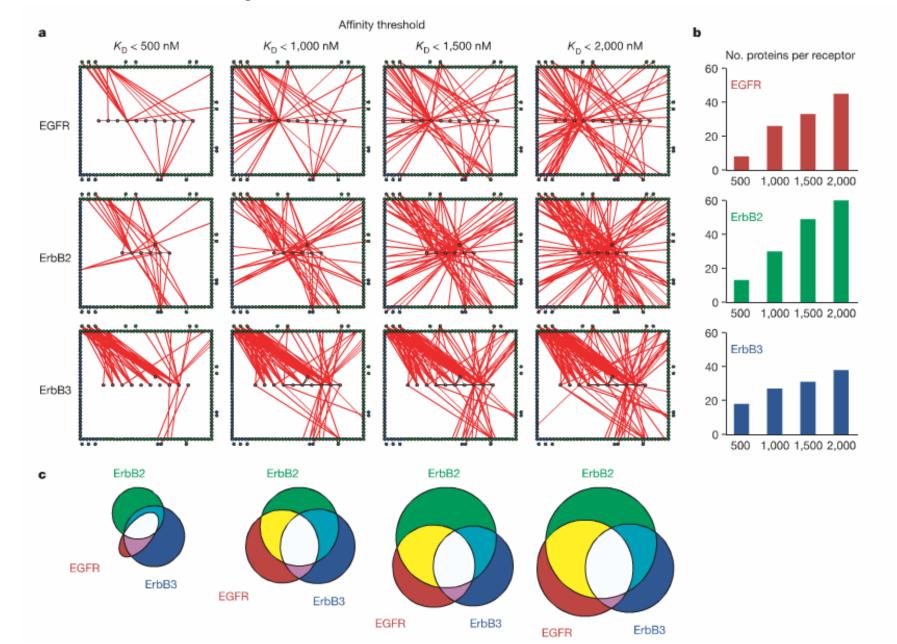
Quantitative Interaction



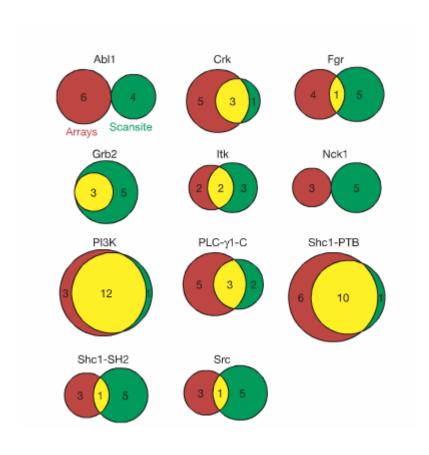
Quantitative Interaction



System-level View



Available Domains



PROTOCOL

Global identification of protein kinase substrates by protein microarray analysis

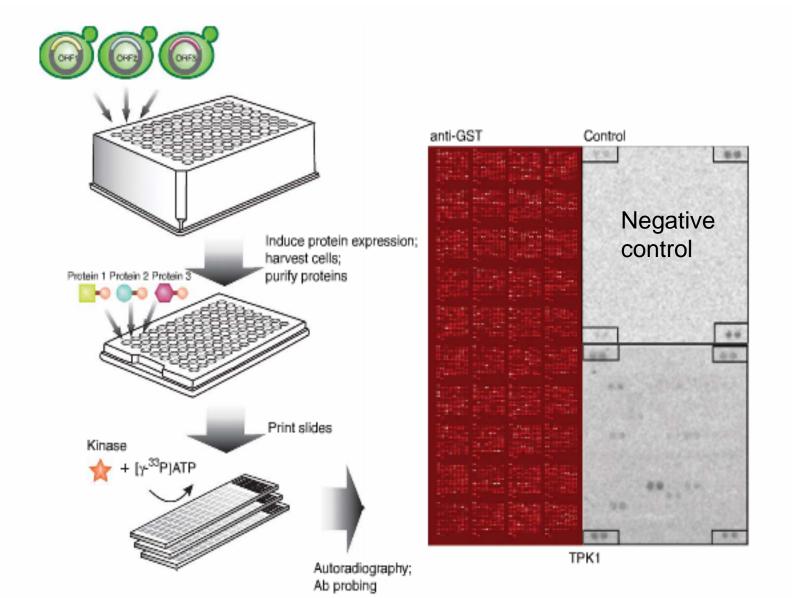
Janine Mok1-3, Hogune Im1-3 & Michael Snyder1,2

¹Department of Molecular, Cellular and Developmental Biology, Yale University, New Haven, Connecticut, USA. ²Present addresses: Stanford Genome Technology Center, Stanford University, Palo Alto, California, USA (J.M.); Department of Genetics, Stanford University, Palo Alto, California, USA (H.I. and M.S.). ³These authors contributed equally to this work. Correspondence should be addressed to M.S. (MPsnyder@stanford.edu).

Published online 19 November 2009; doi:10.1038/nprot.2009.194

| VOL.4 NO.12 | 2009 | NATURE PROTOCOLS

Yeast Kinase Identification





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journal homepage: www.elsevier.com/locate/bios



Development of a high-sensitivity immunoassay for amyloid-beta 1–42 using a silicon microarray platform



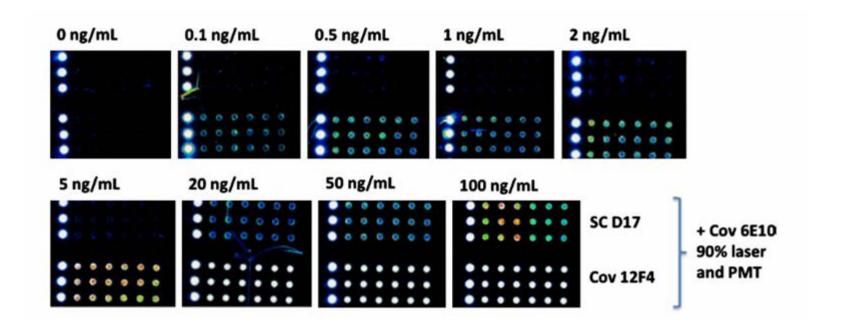
Paola Gagni, Laura Sola, Marina Cretich*, Marcella Chiari

Consiglio Nazionale delle Ricerche, Istituto di Chimica del Riconoscimento Molecolare (ICRM), Via Mario Bianco, 9 20131 Milano, Italy

Capture Ab selection

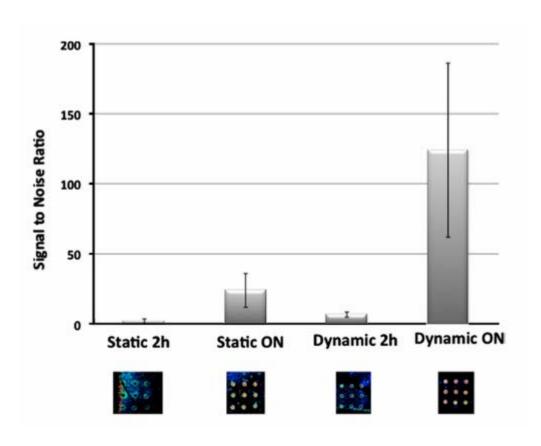
]	Incubation 1 hour/Cov-6E10 (80% laser and PMT)		
		A-Beta 1-42 100 ng/mL	A-Beta 1-39 100 ng/mL	Blank
Capture Ab	SC-D17	5540±1274	42±26	13±11
	NT-11H3	2290±412	51±37	23±10
	NT-8G7	506±210	47±8	47±13
	Cov-4G8	63219±751	13818±260	1208±117
	Cov-12F4	21010±1698	260±65	275±166

AB 42 Detection



Artificial CSF

Signal to Noise ratio



Human CSF

Conclusions

- 15 years after the first landmark papers and proof of concepts the technique is limited to a few applications
- Use of arrays is discouraged by
 - The laborious preparation processes
 - Cost and equipment
 - Complexity of data generated
- Protein folding and missfolding remain a challenge and an obstacle for protein chip
- Protein chips are already available commercially, but major publications validating their use are still missing

Questions?

