

Protein-Protein Interactions (cont.) & Phosphorylation

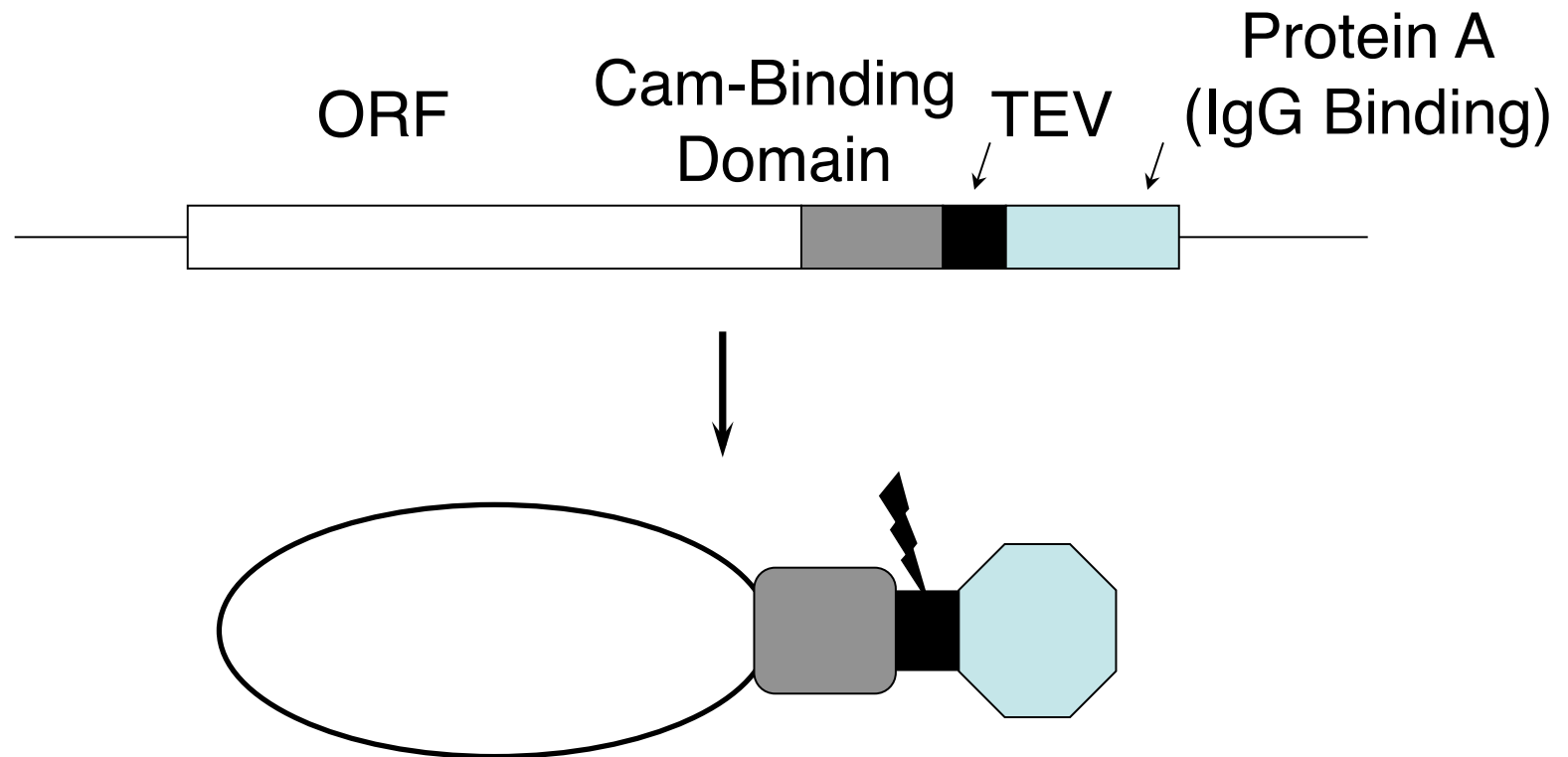
Michael Snyder

April 8, 2009

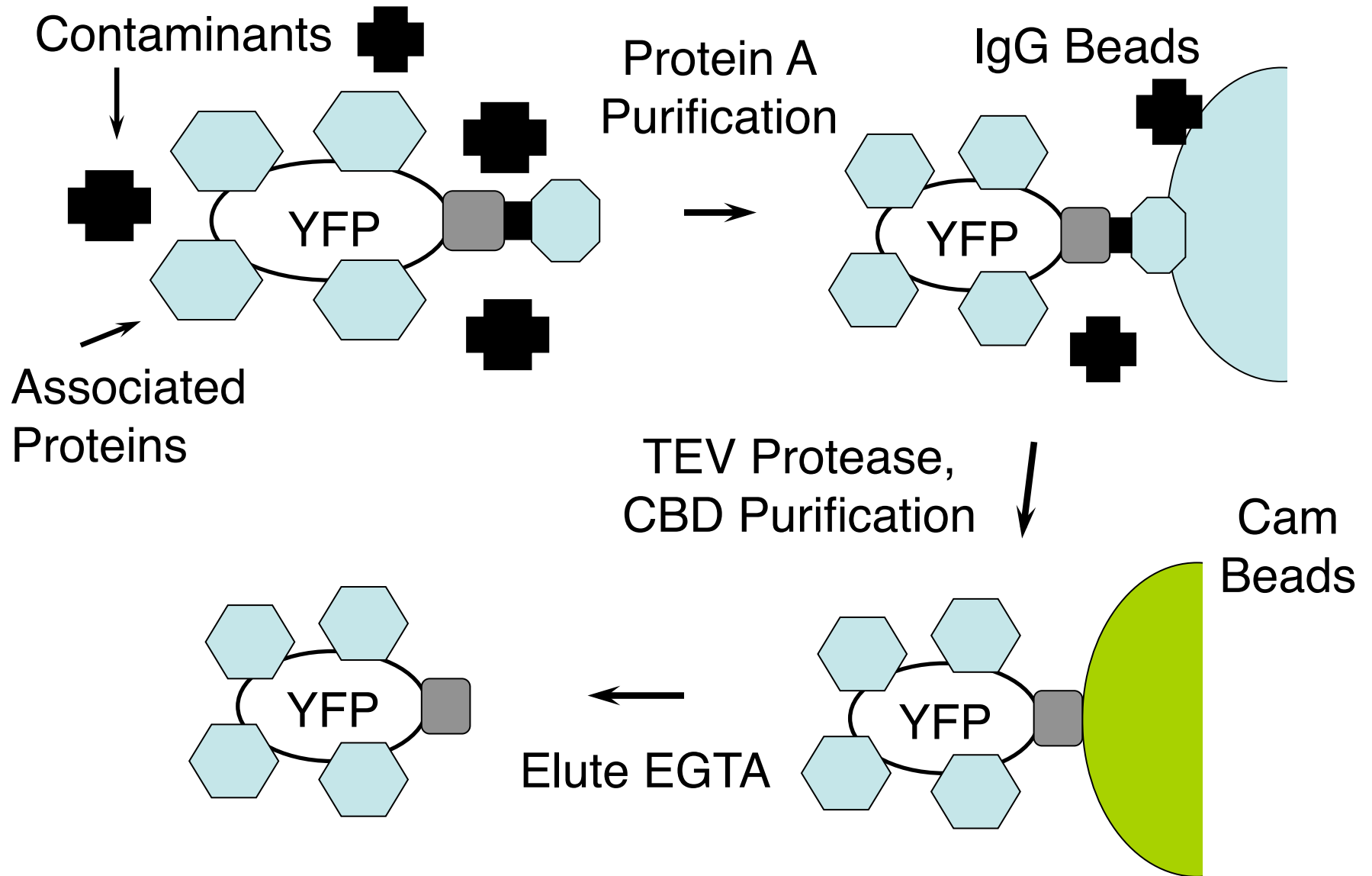
Three Methods:

- 1) Two Hybrid
- 2) Complex Analysis: Affinity tagging/Mass Spectrometry
- 3) Protein Chip

Tandem Affinity Purification (TAP) Tagging



TAP Approach

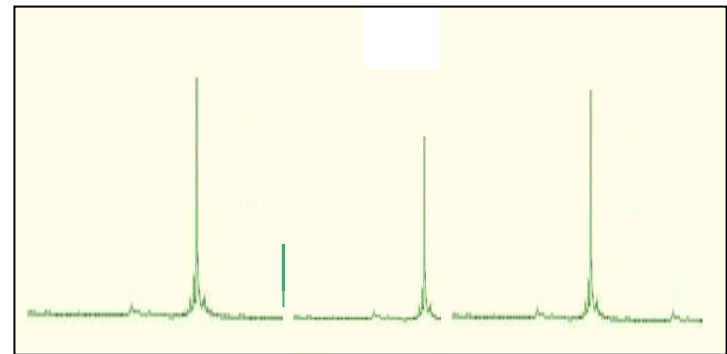


Identify Proteins by Mass Spec

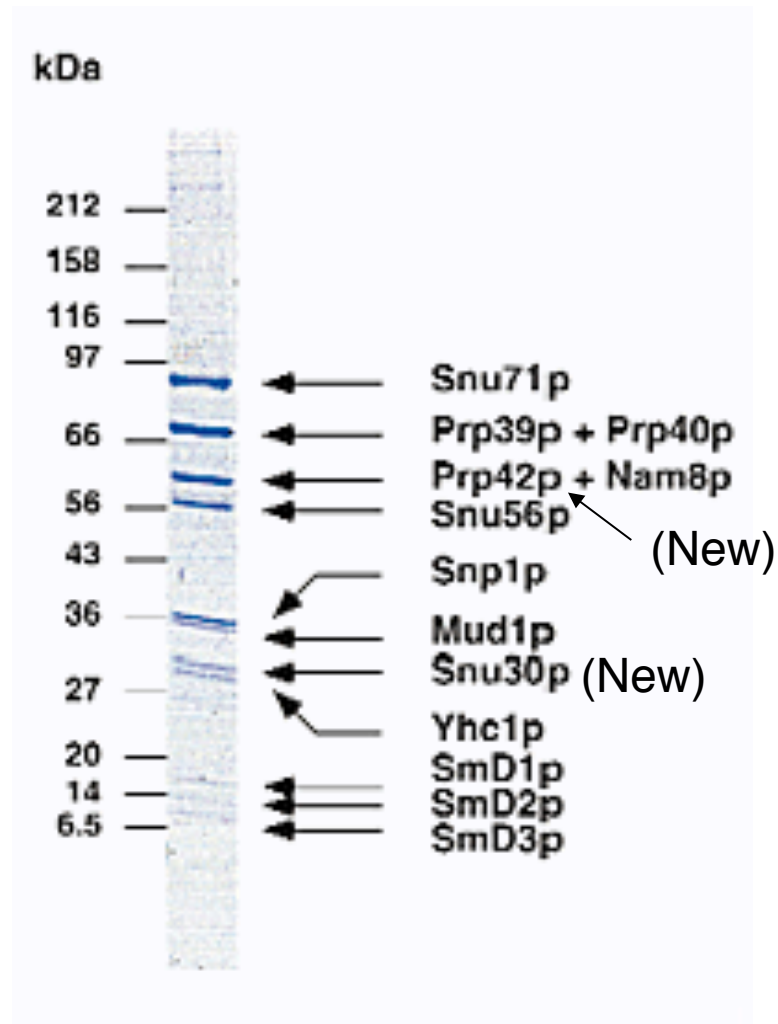
Load on SDS Gel



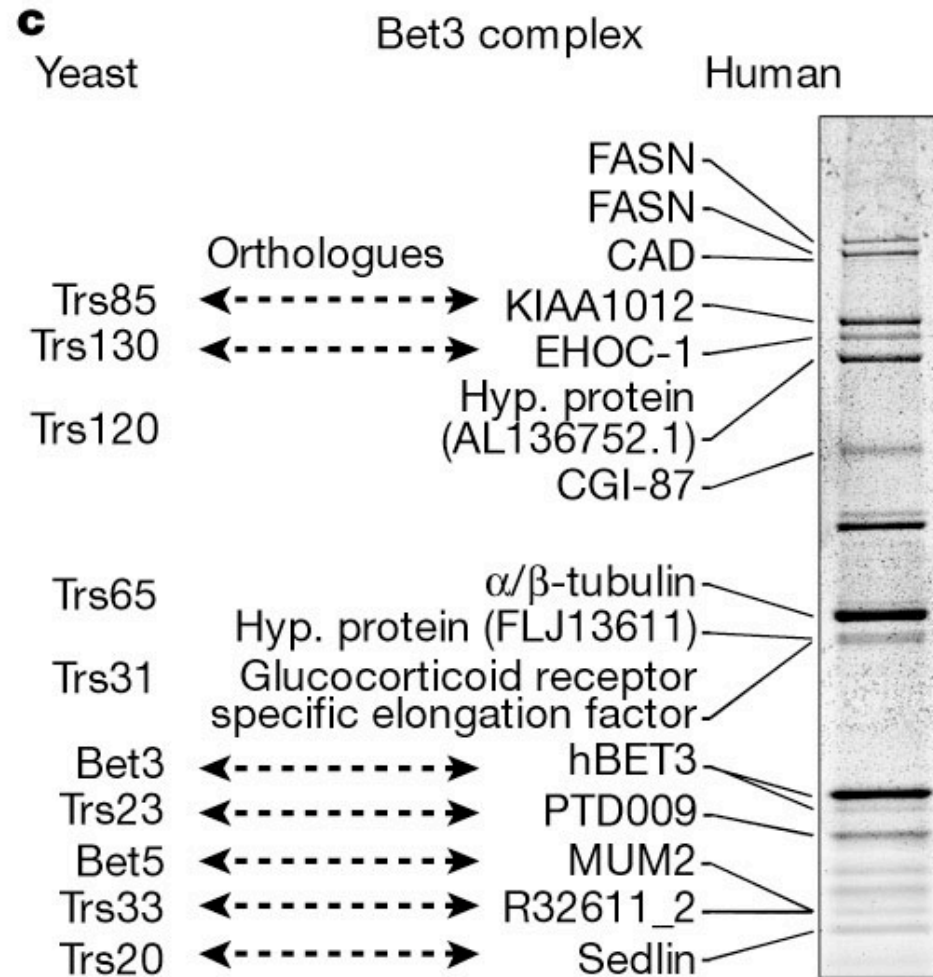
Excise Band;
Digest with Trypsin;
Run Mass Spec



TAP Purification of The U1 Splicing Complex (Snu71p)



Many Complexes Are Conserved



Affinity Purification/Mass Spec Analysis of Complexes - Yeast

4,562 Purifications (Krogan et al. 2002)

2,357 Successful

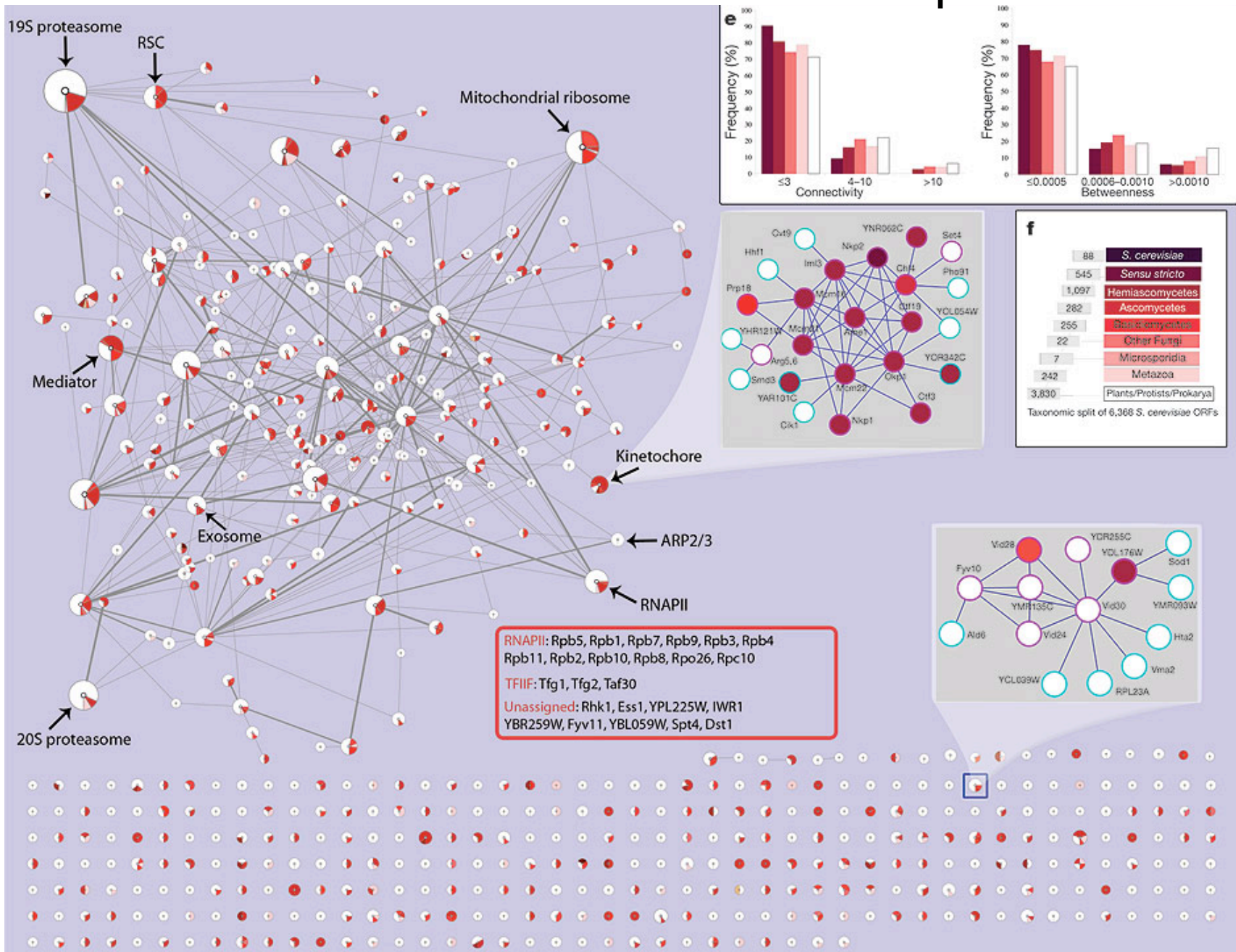
4,087 Interacting Proteins

7,123 Core Interactions (2,708 proteins)

14,317 Extended (3,672 proteins)

547 Complexes

Size and Conservation of the Complexes



TAP Tag Approach

Advantages

- In Vivo Assay
- Identifies Entire Complex

Disadvantages

- Interactions may be indirect
- Likely to miss some rare components
- Contaminants may copurify

Summary

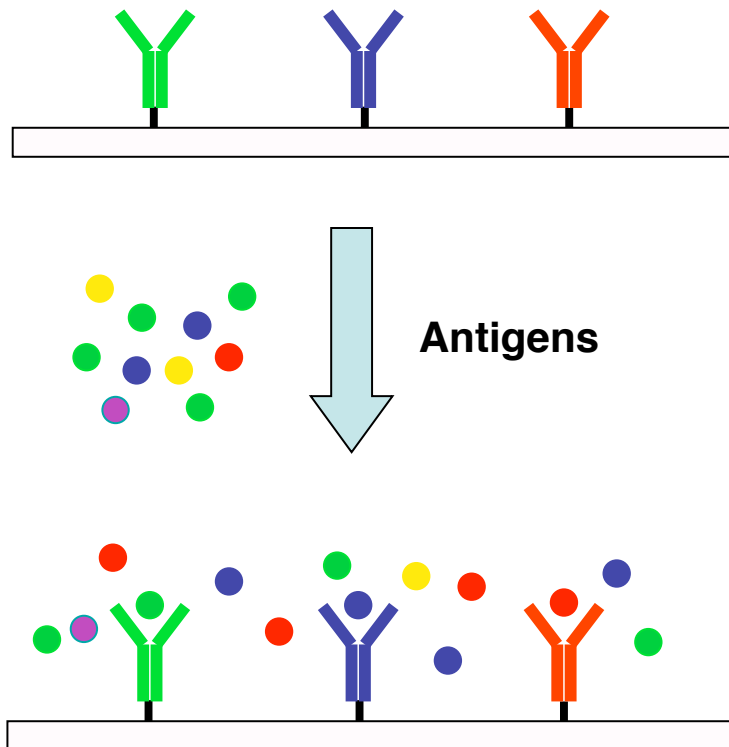
- Affinity Purification: ~10,000 High Confidence Interactions Among ~3-4000 Proteins
- Two Hybrid: >4,549 Interactions Among 3,278 Proteins
- >20,000 Interactions
- Combining Data = More Accuracy

What is a Protein Microarray?

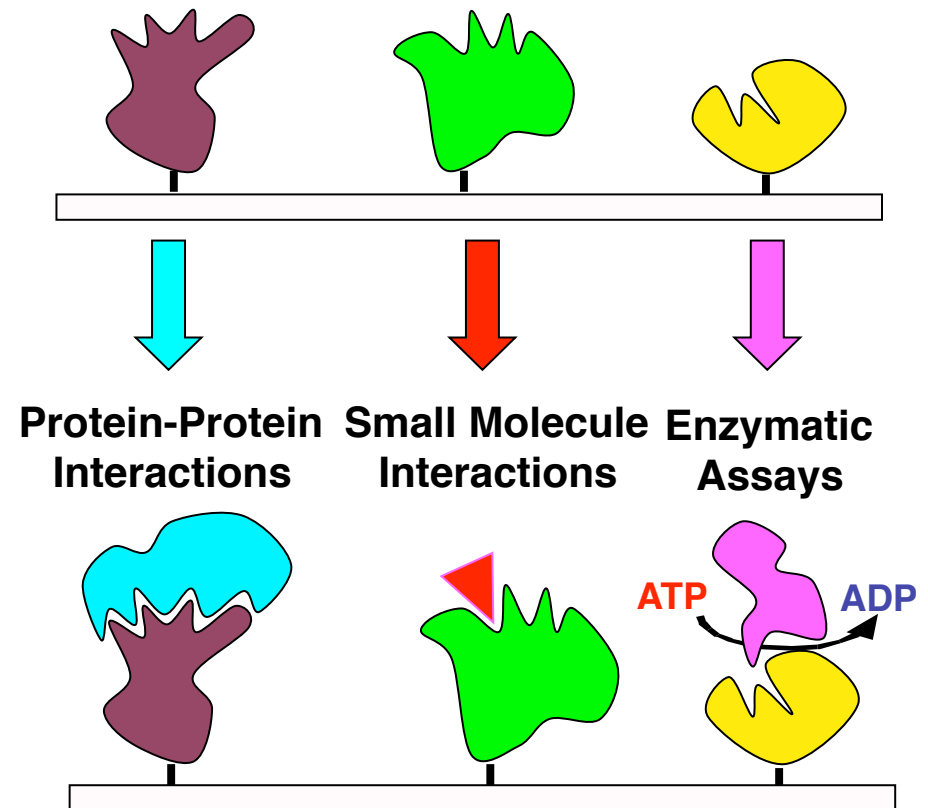
A high density array
containing 100s to many
thousands of proteins

Two Types of Protein Microarrays

Antibody Microarrays



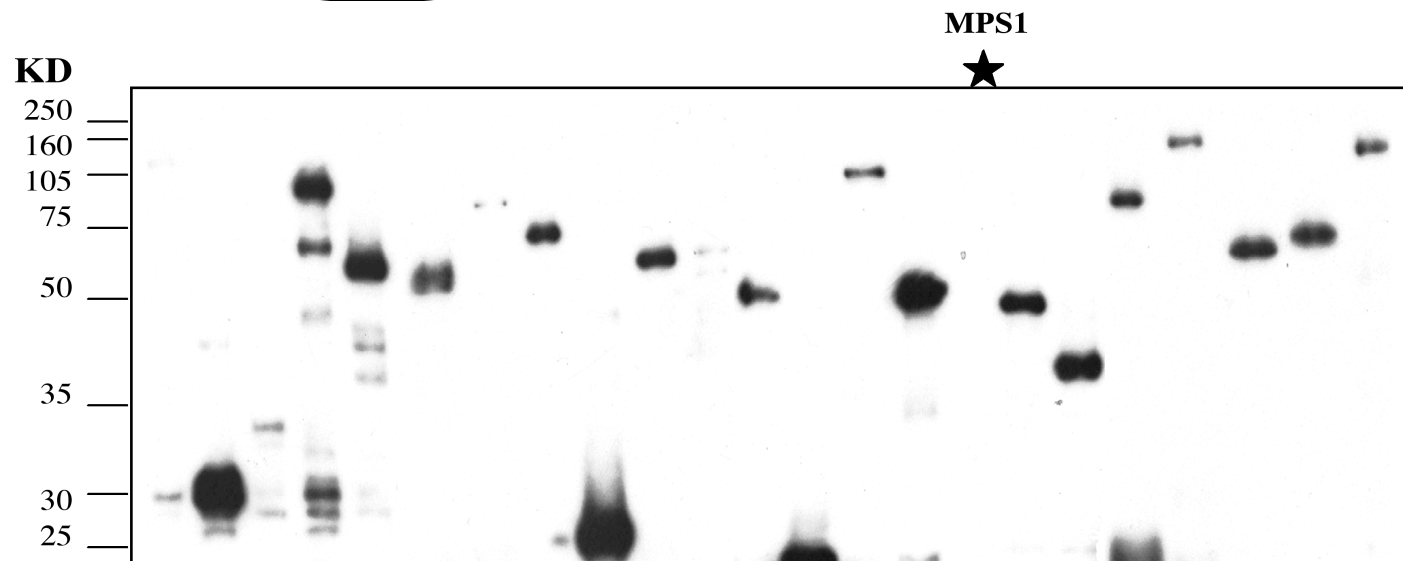
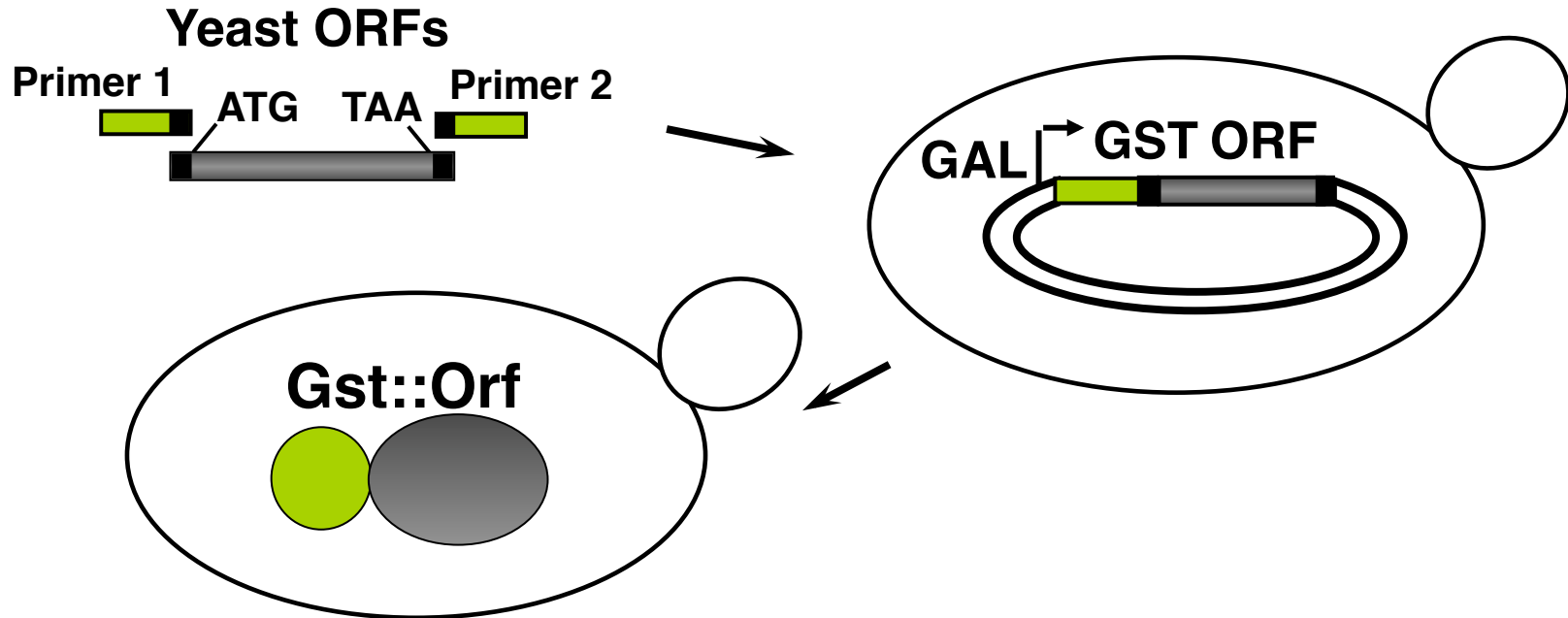
Functional Protein Microarrays



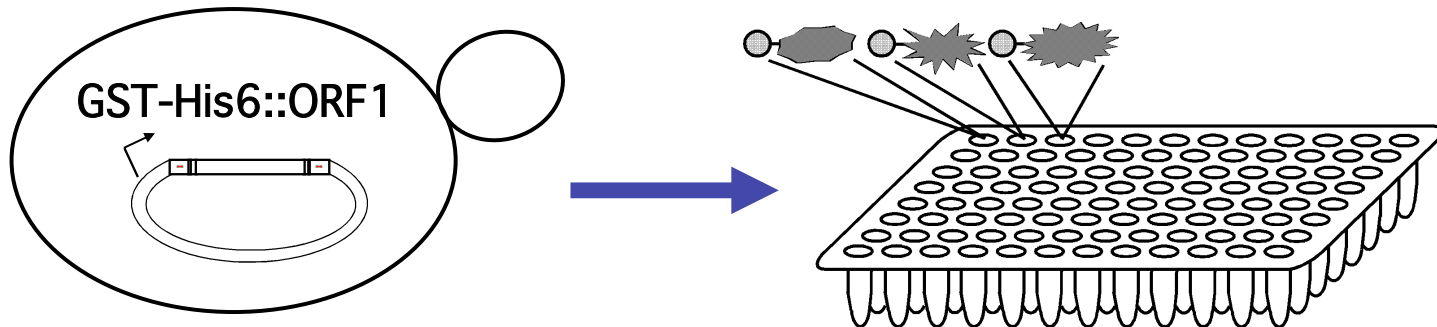
What Is Needed for Preparing and Screening Functional Protein Microarrays?

- A high quality expression library
- Methods for preparing large numbers of proteins
- Methods to array the proteins
- Assays for screening

Cloning & Expression Strategy

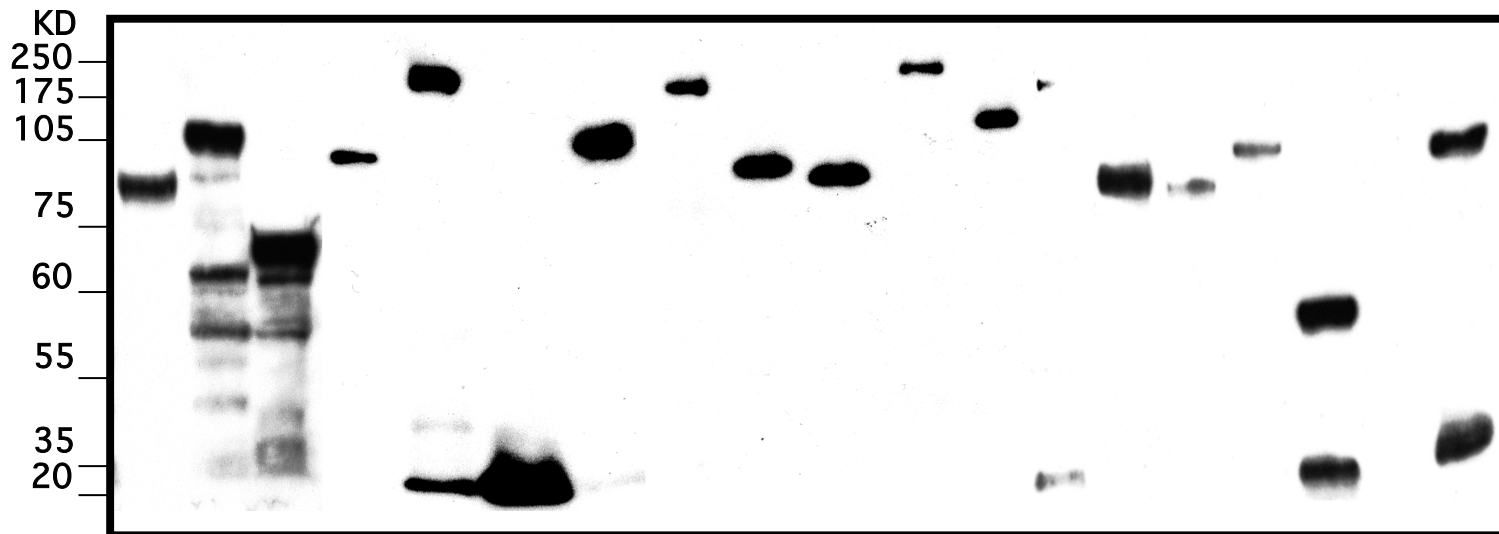


Producing the Yeast Proteome



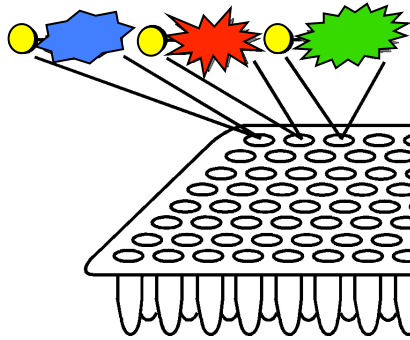
5800 expression clones 93.7%

~80% full-length proteins

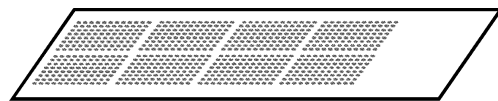
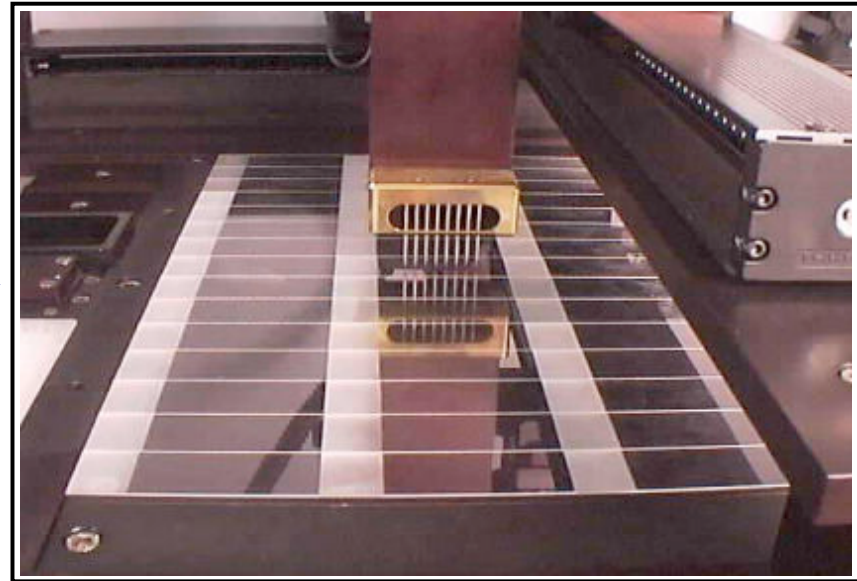


Printing the Yeast Proteome

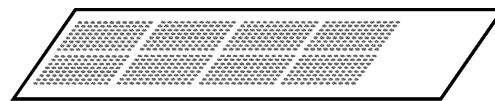
GST:P1 GST:P2 GST:P3



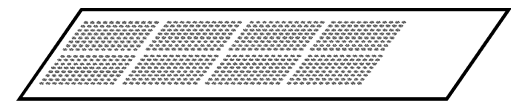
Source Plate



Protein-Protein



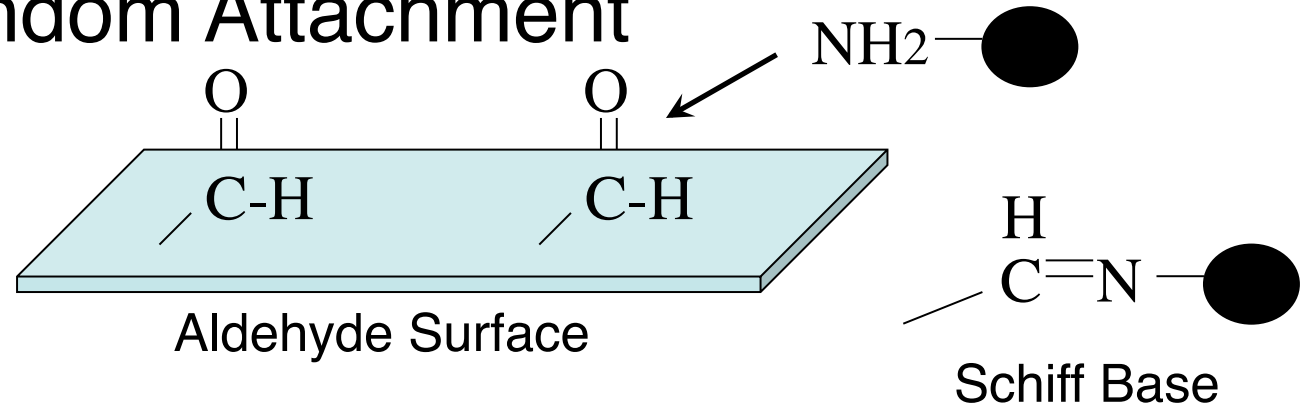
Protein-Lipid



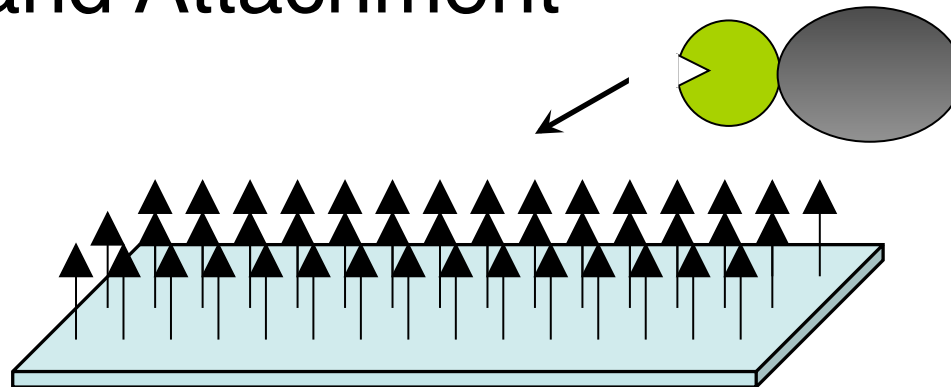
Protein-DNA

Glass Slides

1) Random Attachment



2) Ligand Attachment



The Yeast Proteome Chip

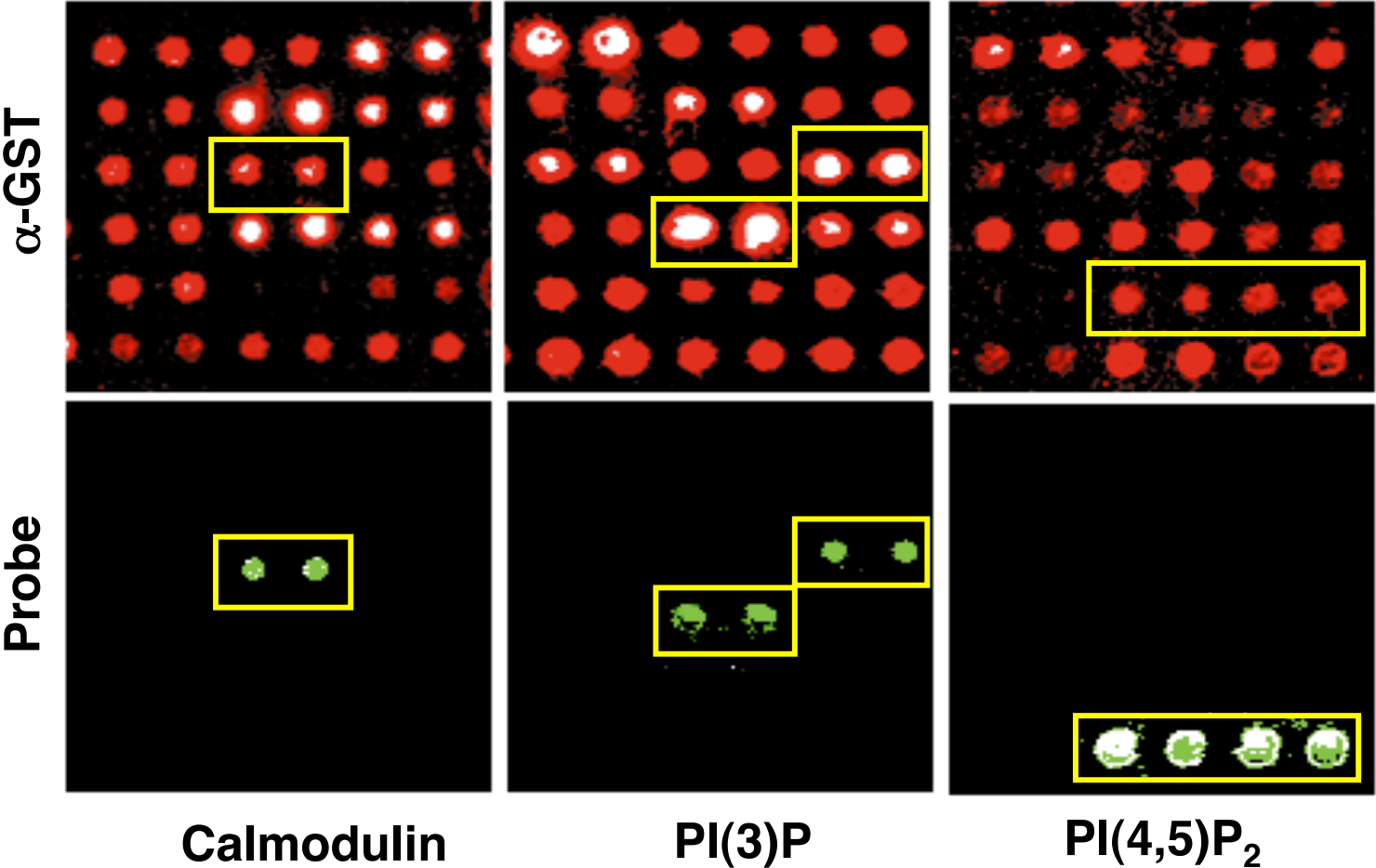


Probed With
Anti-GST Antibodies

Many Types of Applications

- Binding Assays
 - Protein-Protein Interactions
 - Protein-Lipid Interactions
 - Nucleic Acids (dsDNA, ssDNA, polyA-mRNA)
 - Small Molecule Screens
- Antibody Specificity
- Kinase Substrates and Posttranslational Modifications
- Viral Diagnostics

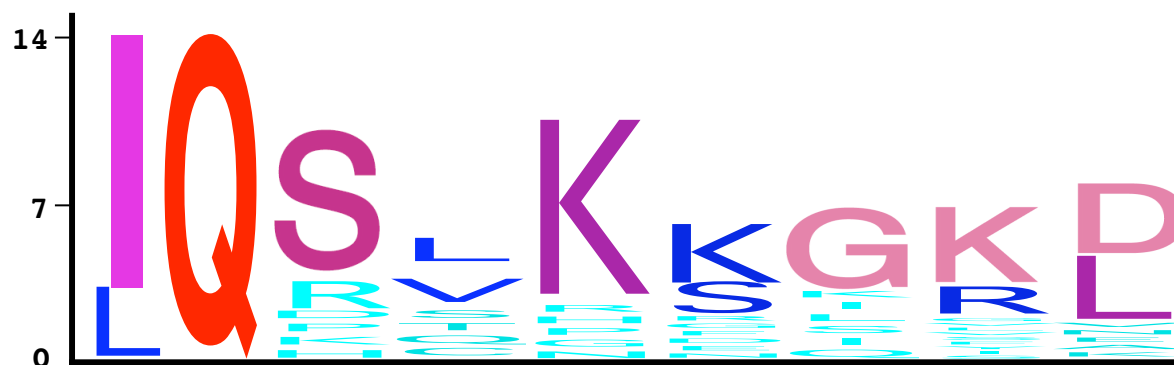
Biochemical Assays on Proteome Chips



Calmodulin-Binding Proteins

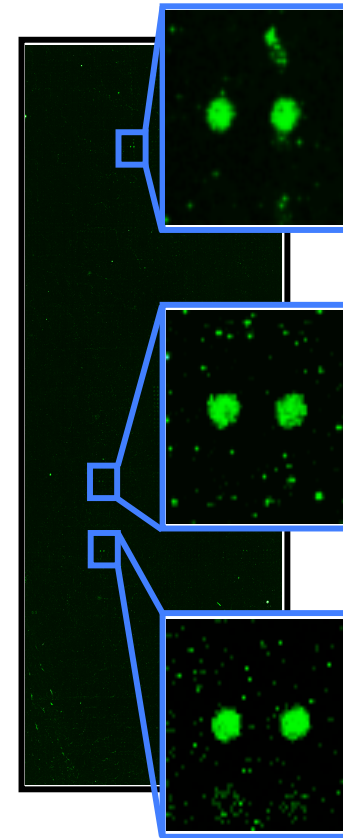
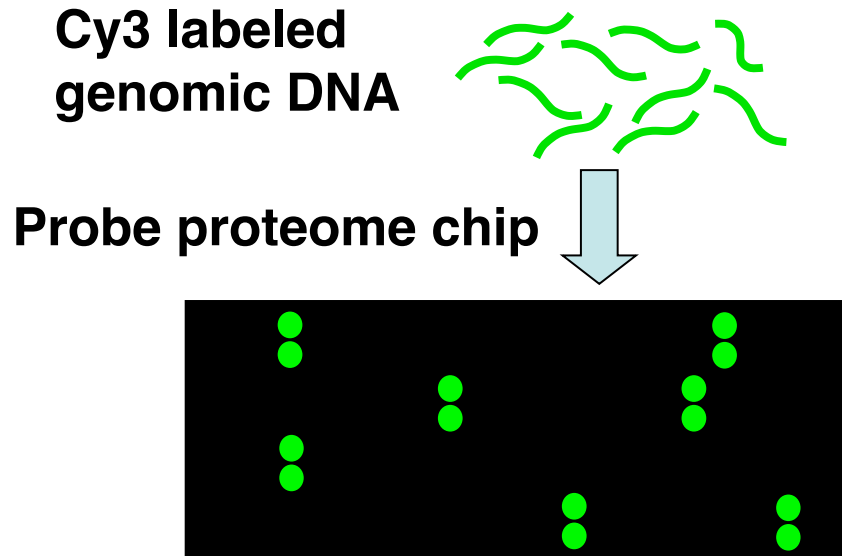
- 12 Known or Suspected Targets
- 33 New Binding Proteins
- Derived New Consensus Binding Site

Potential Calmodulin Binding Motif



YFL003C/MSH4	LKET	LOS	VK	SL	KD	AL	390	
YJR073C/OPI3	HSVD	LOS	SK	FQ	LA	IV	28	
YBR050C/REG2	DEHF	IQR	LP	ST	RL	NS	196	
YNL202W/SPS19	AKIE	LQR	LG	ST	RD	IA	246	
YOL016C/CMK2	DDL	LOS	QK	KG	GE	LT	395	
YBR011C/IPP1	LNPI	IQD	TK	KG	KL	RF	80	
YGR034W/RPL26B	RKAI	IQR	.	K	GG	KL	E.	129
YFR004W/RPN11	VDPE	IQS	VK	GK	VV	ID	154	
YIL021W/RPB3	GHPI	IQD	.	K	EG	NG	/L	143
YGL063W/PUS2	RVWG	IQP	VN	KK	FN	AR	103	
YDR292C/SRP101	LLRE	IQS	.	K	RS	KD	EE	388
YFR014C/CMK1	LNMK	IQK	LR	DL	YL	EQ	346	
YBR213W/MET8	DLFG	IQH	CH	N.	ID	/K	242	
YAL029C/MYO4	NGLI	IQS	SK	F	I	SK	/L	1167

Identification of New DNA Binding Activities



- ~200 bound DNA probe
Found New Activity
Arg5,6

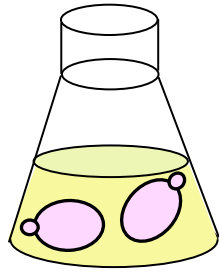
Yeast Phosphorylation

6,000 Proteins

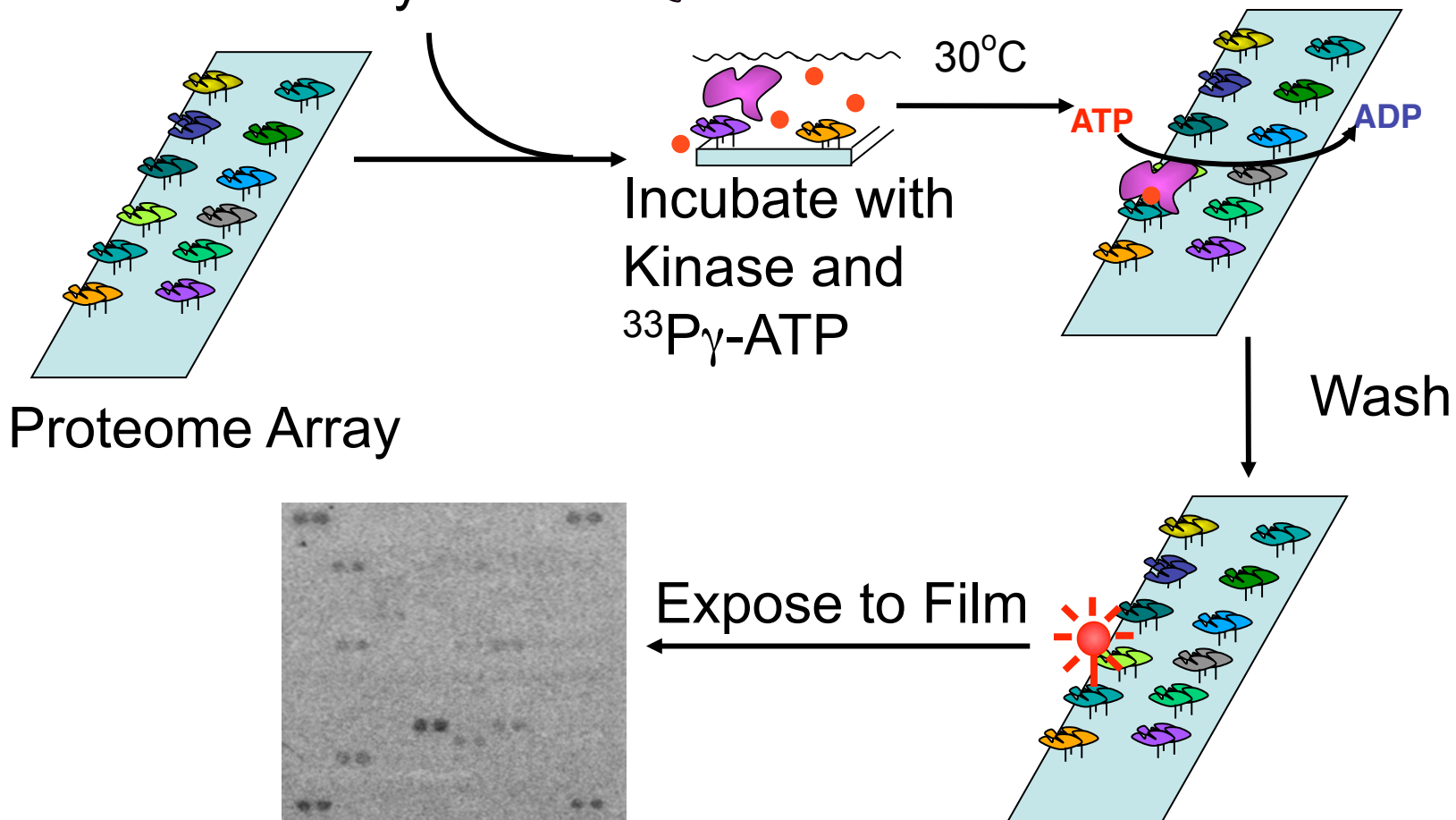
122 Protein Kinase Homologs

- 14 Uncharacterized
- 50% Have no known *in vivo* substrates
- <160 Known kinase-substrate phosphorylations

Global Analysis of Kinase Substrates



Overexpress and Purify Kinase 



Proteome Array

Incubate with Kinase and $^{33}\text{P}_\gamma\text{-ATP}$

30°C

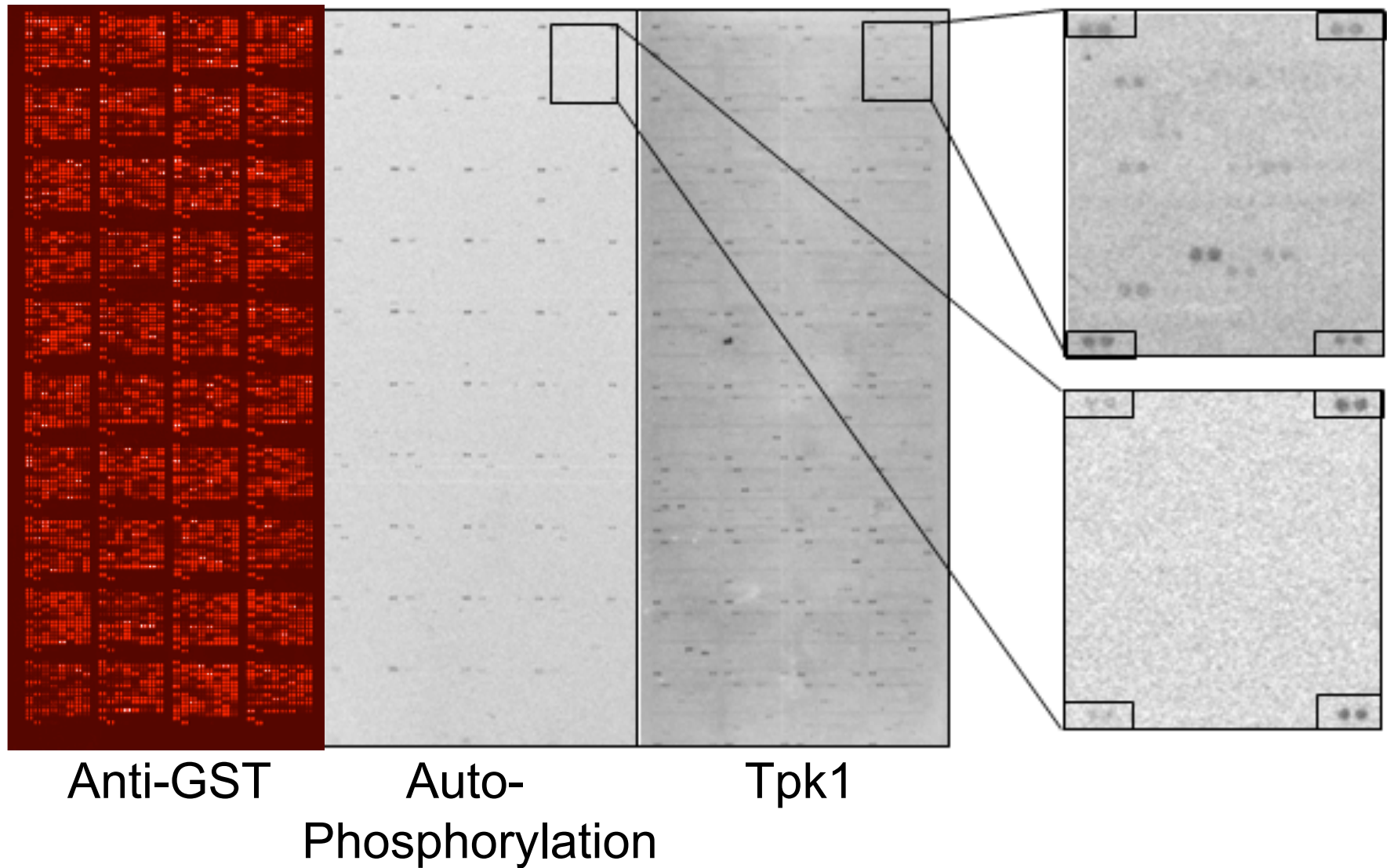
ATP

ADP

Wash

Expose to Film

Kinase Assays on Protein Chips



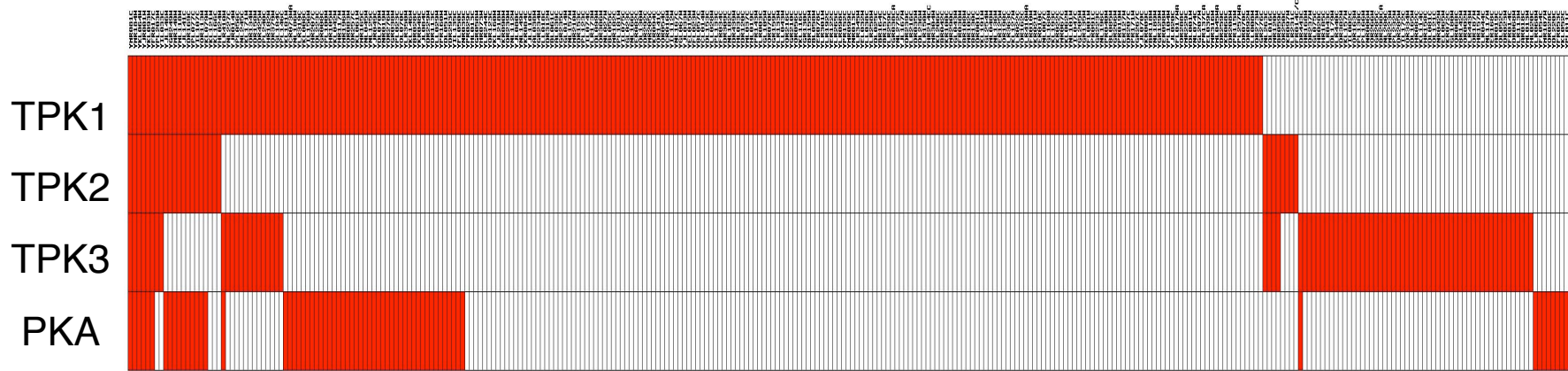
In Vitro Phosphorylome Summary

AKL1	IKS1	PKH2	STE20
ARK1	IME2	PRK1	SWE1
ATG1	IPL1	PRR1	TOS3
BCK1	IRE1	PRR2	TPK1
CDC15	KCC4	PTK2	TPK2
CDC5	KIN1	RAD53	TPK3
CKA1	KIN2	RCK1	VHS1
CLA4	KIN28	RCK2	YAK1
CMK1	KIN3	RIM11	YCK1
CMK2	KIN4	RIM15	YCK2
CTK1	KIN82	SAT4	YCK3
DBF2	KNS1	SKM1	YDR466W
DUN1	KSP1	SKS1	YGL059W
ELM1	KSS1	SKY1	YGR052W
FUN31	MCK1	SLT2	YKL171W
FUS3	MEK1	SNF1	YMR291W
GIN4	MKK1	SPS1	YOL128C
HAL5	MPS1	SRB10	YOR267C
HRR25	PAK1	SSK22	YPL141C
HSL1	PBS2	STE11	YPL150W
CDC28-CLN2	CDC28-CLB5	PHO85-ALONE	PHO85-PCL1
PHO85-PCL2	PHO85-PCL9	PHO85-PHO80	
RIM15 dead	DBF2 dead	HSL1 dead	RAD53 dead

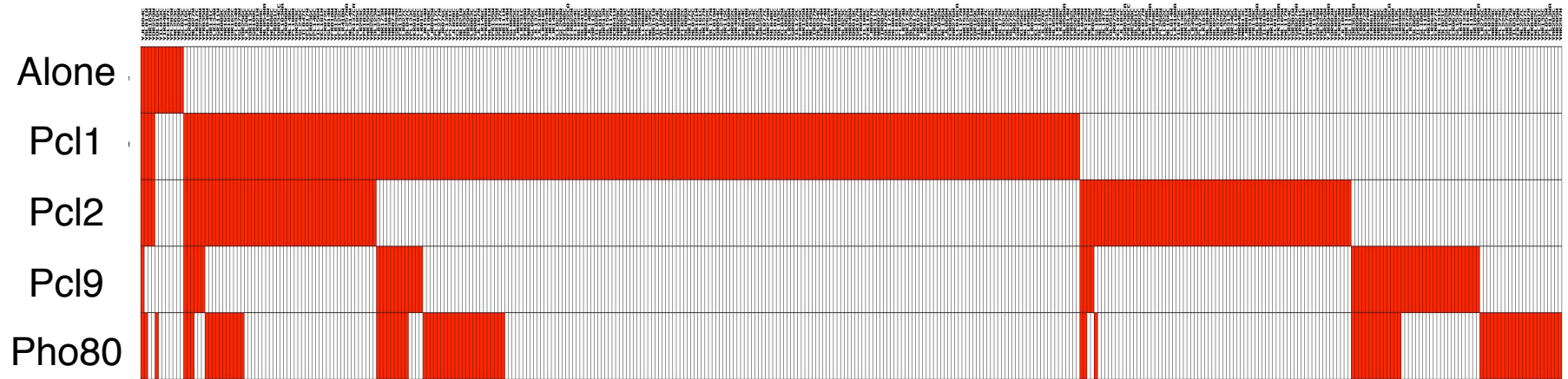
In Vitro Phosphorylome Summary

- 82 unique kinases and several CDKs with different cyclins (Cdc28; Pho85) for 87 specific hit lists
- 4200 total phosphorylation events on 1325 individual targets
- On average kinase phosphorylated 47 proteins on chip (Range 1- 250)
- 50% of substrates were phosphorylated by only one kinase
- Identified at least 21 known kinase-substrate phosphorylations

Substrate List Comparison



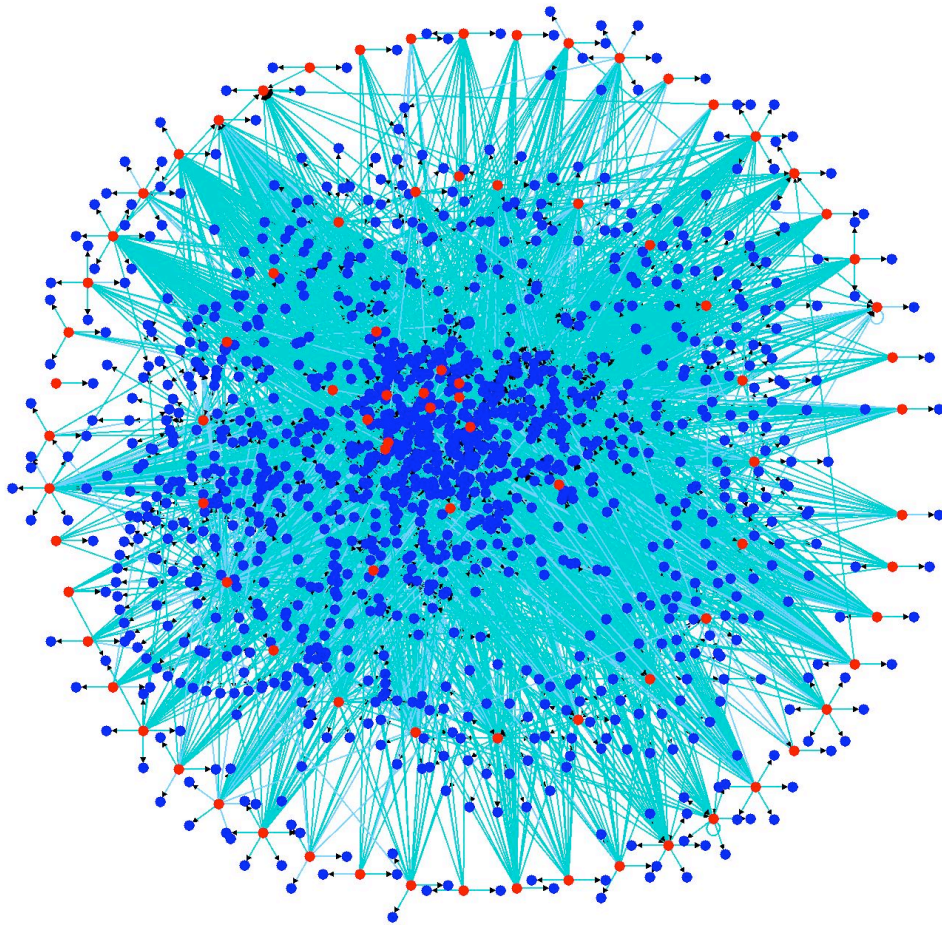
PHO85



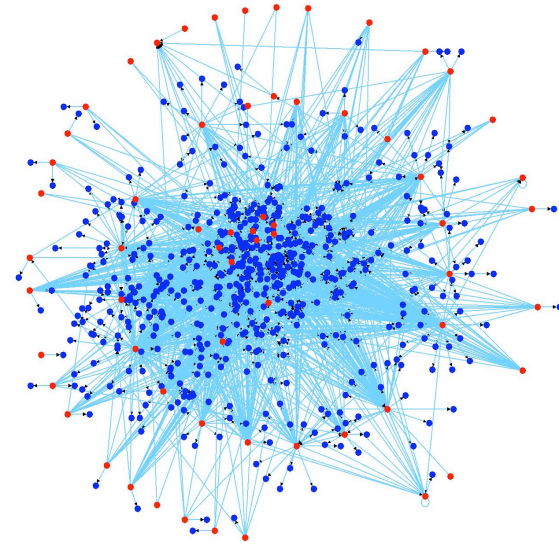
TPK1, PKA overlapped very well

Pcl9, Pho80 overlapped very well

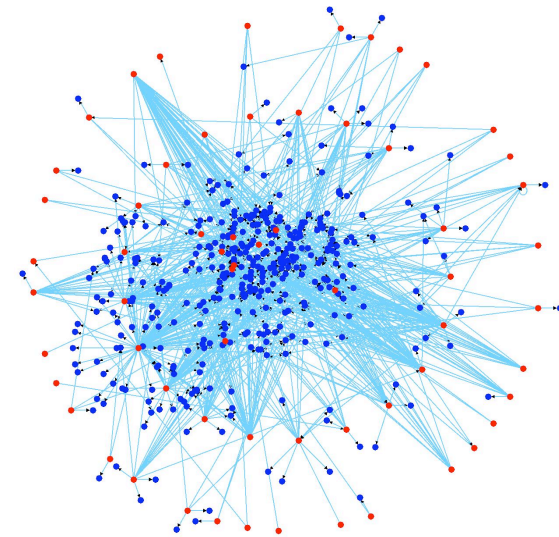
Phosphorylome Network



4200 Phosphorylations, 1325 Proteins



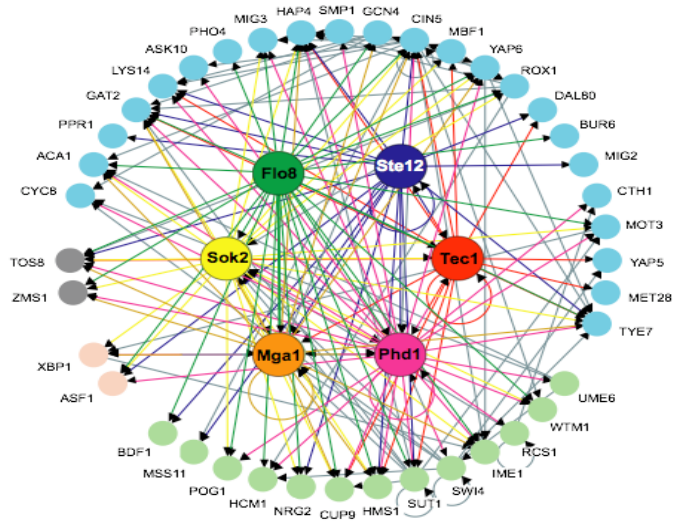
Same localization (33%; $p < E-99$)



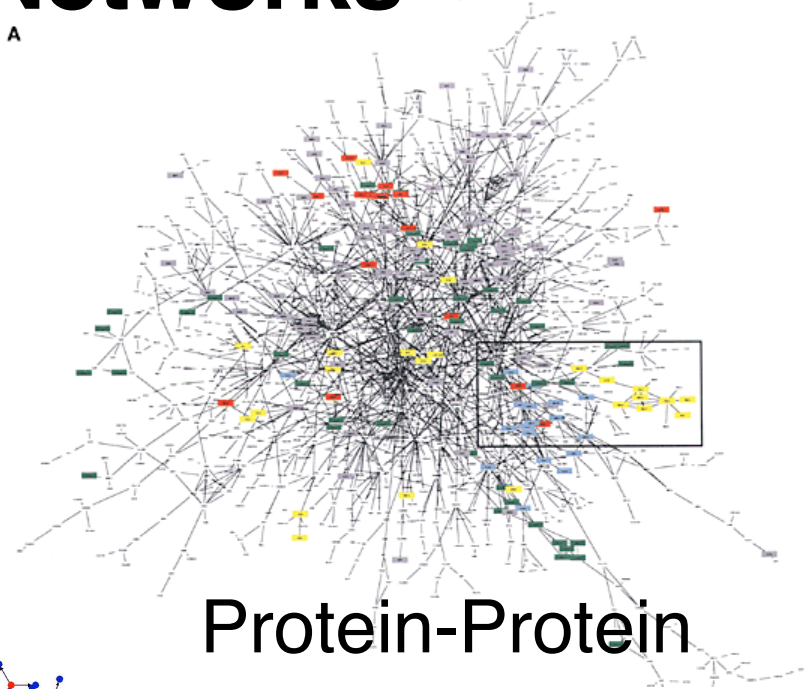
Same function (18.3%; $p < E-99$)

Regulatory Networks

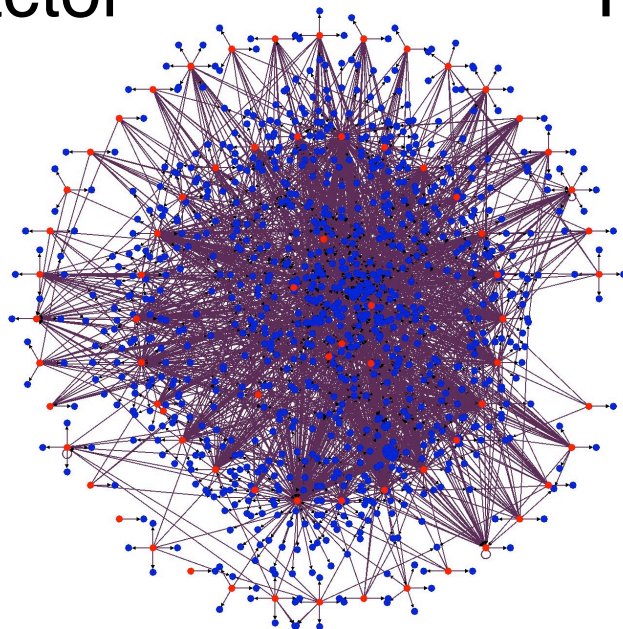
A



Transcription Factor
Binding

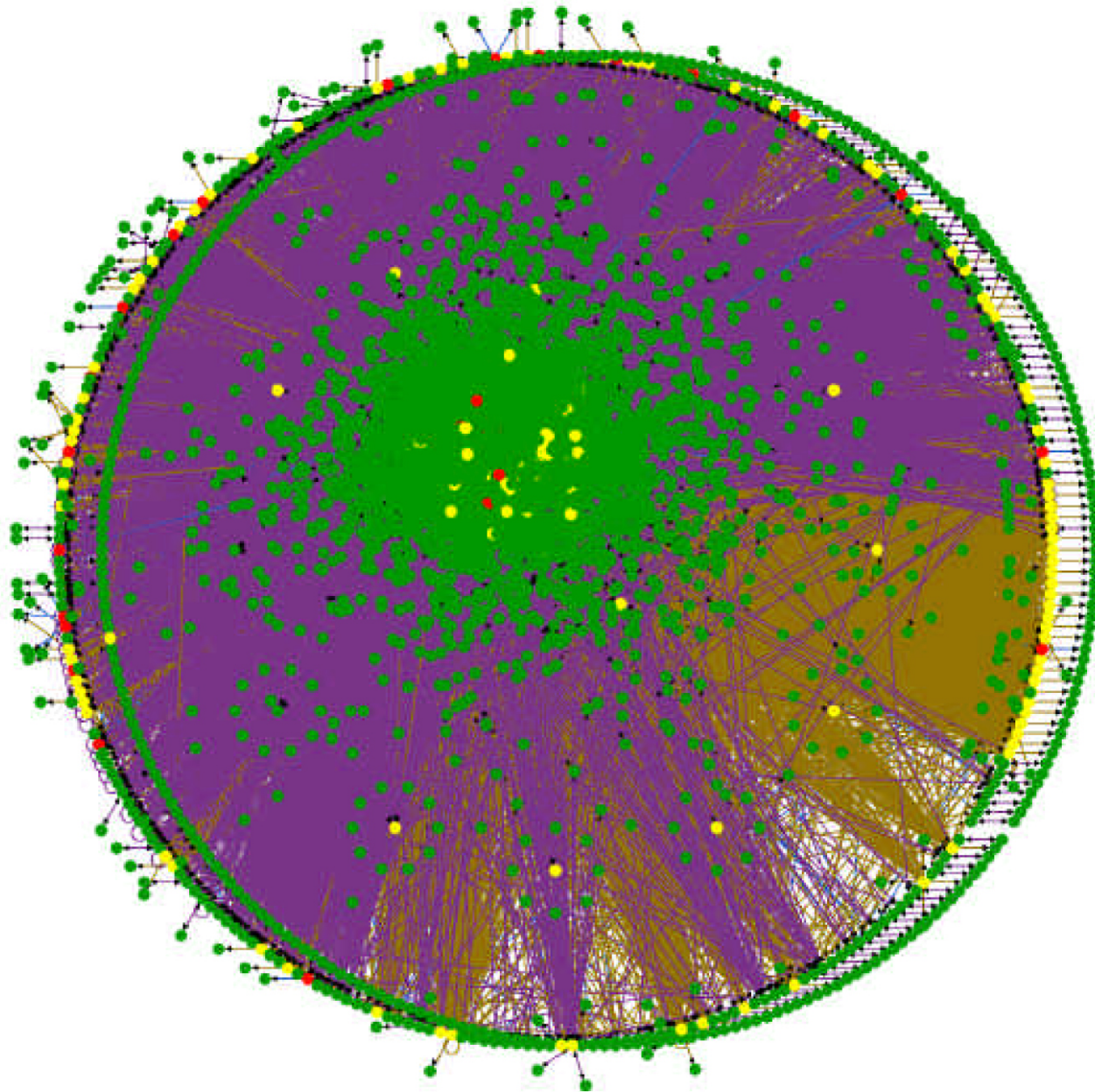


Protein-Protein
Interaction

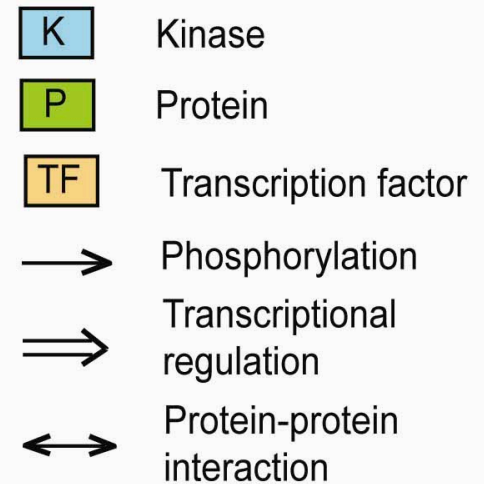
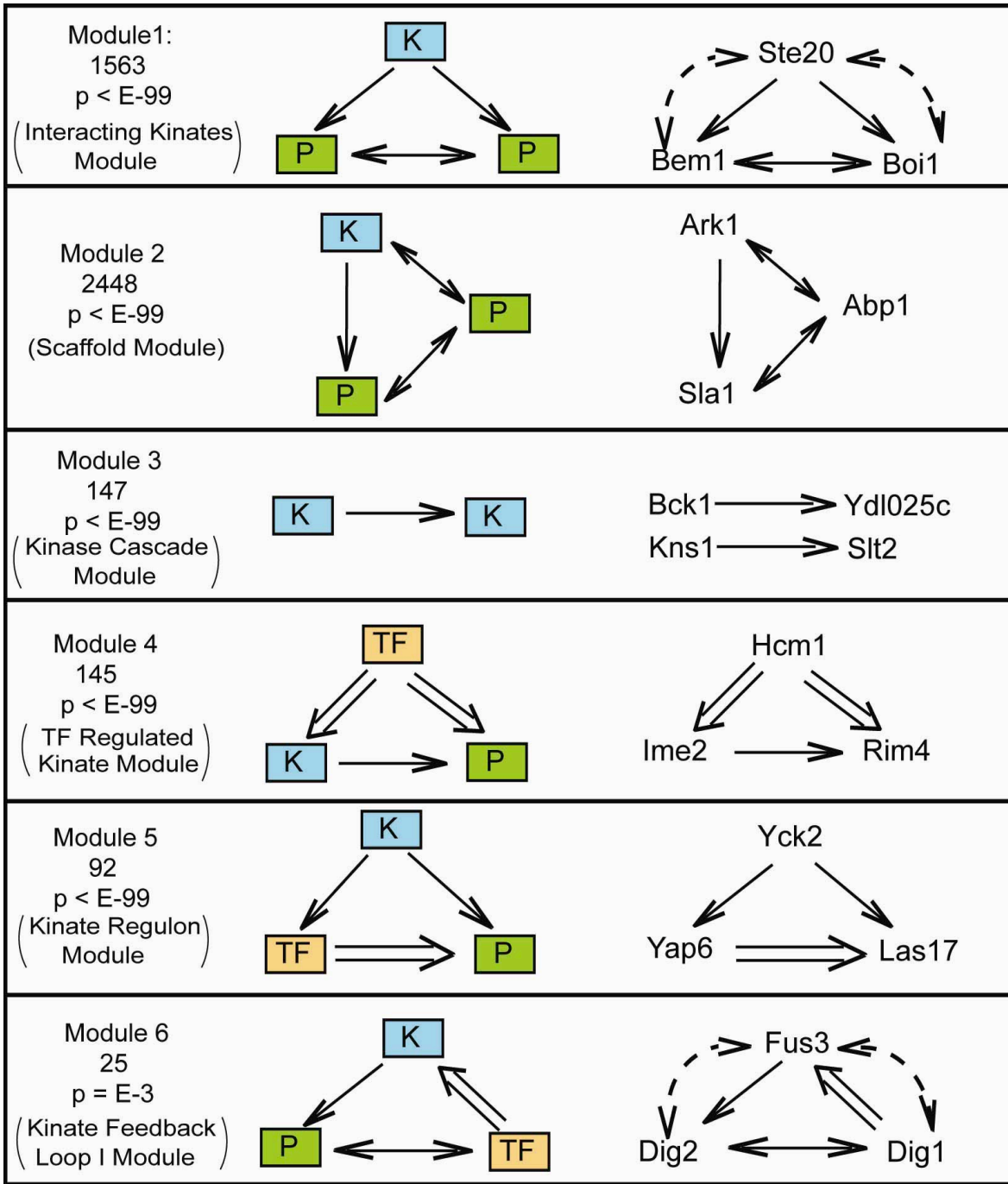


Phosphorylation

Meta Network



Regulatory Modules



Advantages of Protein Chips

- Can screen many proteins simultaneously
- Small amounts of proteins and reagents
- High throughput
- Diverse applications-biochemical assays, posttranslational modifications, small molecule screening

Disadvantage

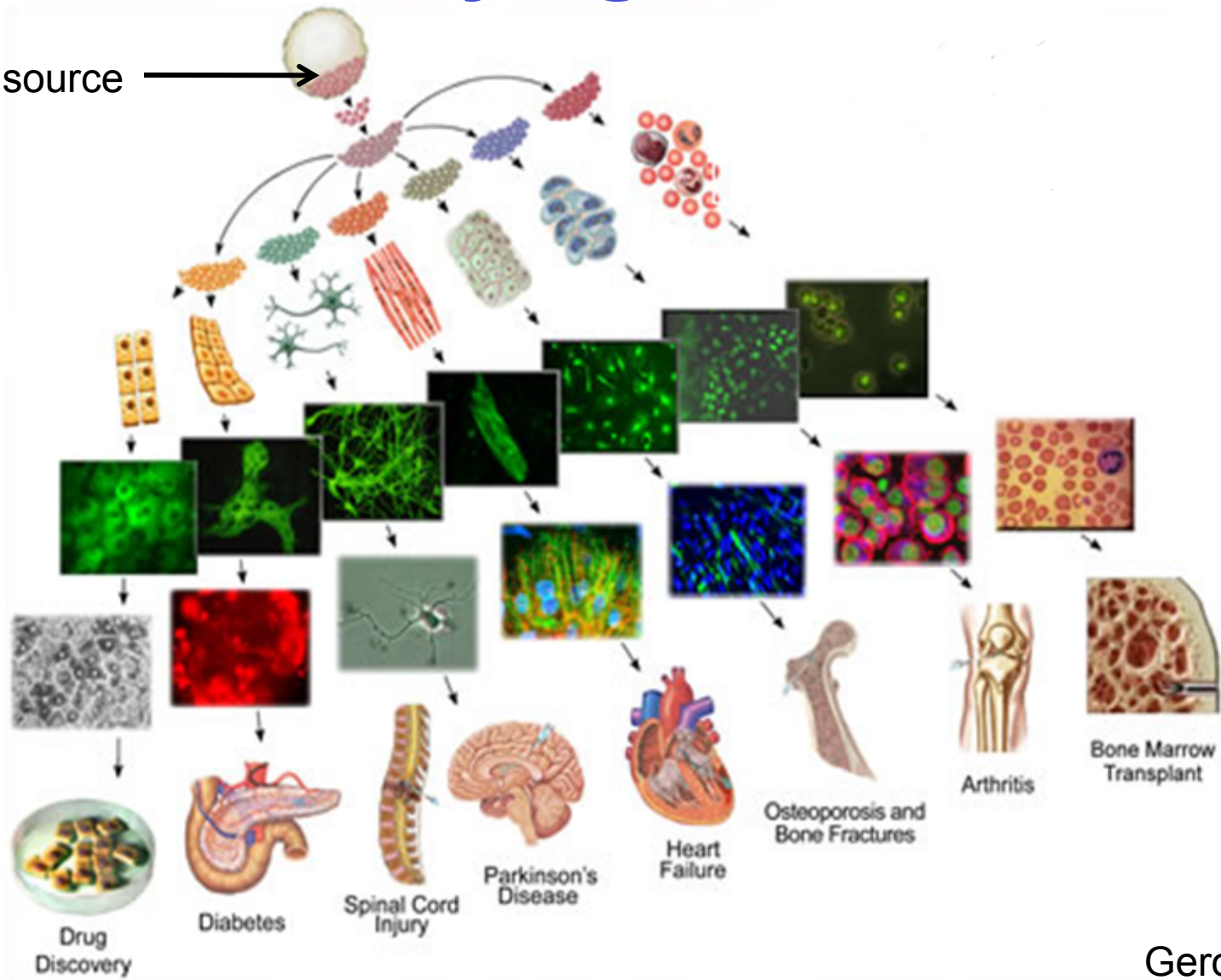
- In Vitro Assay - Must validate In vivo

Many Transcription Factors Remain Uncharacterized

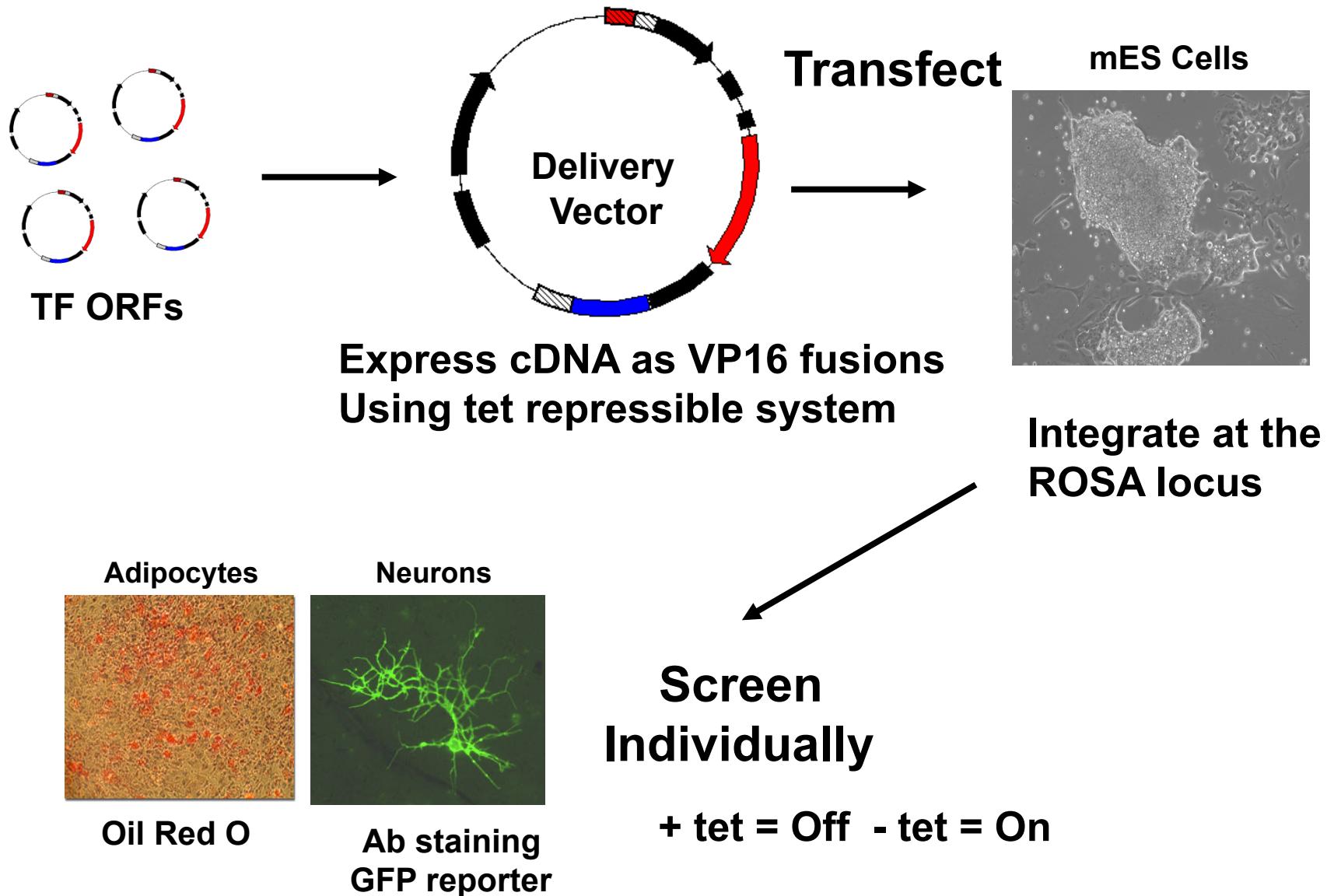
- ❖ 1400-1700 human transcription factors
- ❖ Methods of studying TF function in differentiation
 - Map binding sites
 - Tissue/cell localization
 - Knock down
 - Ectopic expression
- ❖ Ectopically expressed TFs with the ability to activate differentiation programs
 - MyoD (muscle), PPARgamma (fat), Sox 5, 6, 9 (cartilage)

Embryonic Stem Cells are a 'Blank Slate' for Studying Gene Function

ES cell source



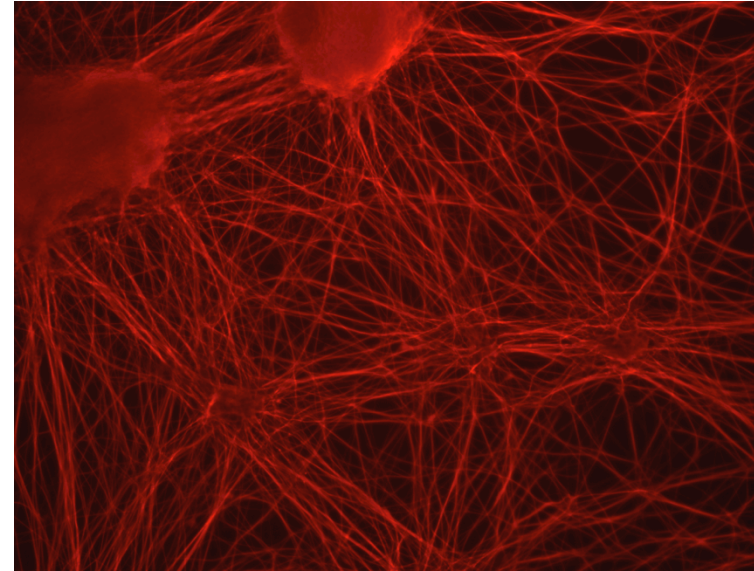
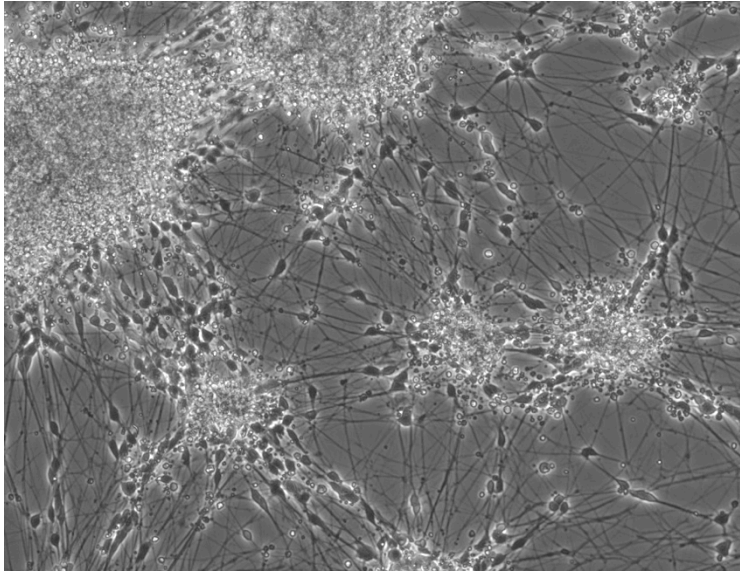
Screen 700 Human TFs in mES Cells



24 TFs Induce Differentiation

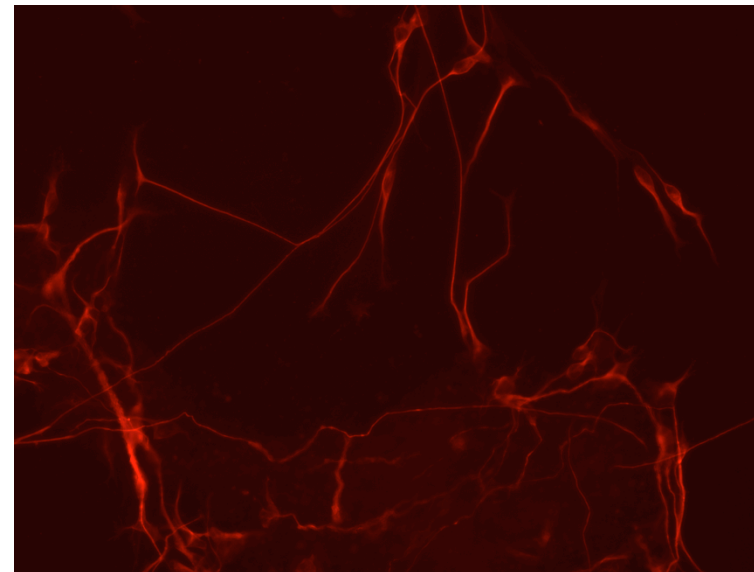
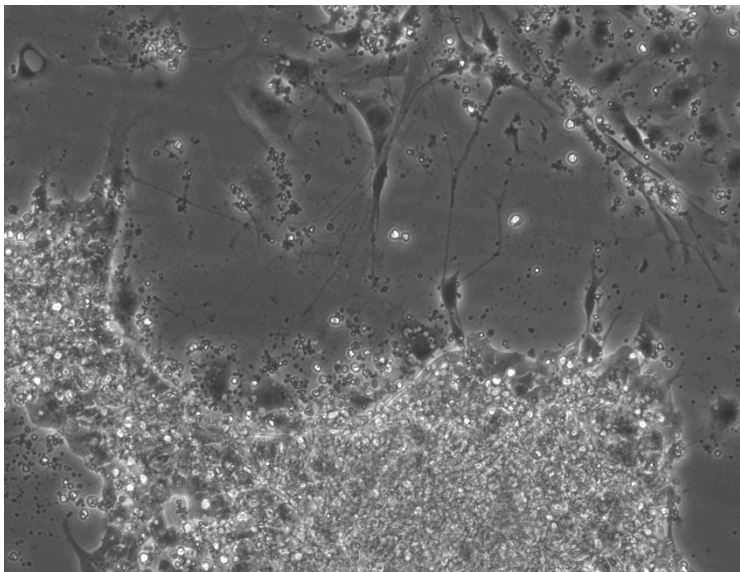
Phase

anti-beta III tubulin



Neuro
G1

-Tc
Neuro
Media
96hrs



ZNF205

24 TFs Induce Neural Differentiation

Hits with VP16

NeuroD1	KLF15
NeuroD6	Braf35
NeuroG1	Meox2
Nkx2.5	Oct-2 *
Tead2	KLF12
Tcf4	ZNF43
MTGR1	ZNF205
Six1	ZNF37A
DMRTC2	ZFP64
Hoxd3	ZNF435
Hoxa3	ZNF408
SATB1	WDR34

Hits without VP16

Nkx2.5
NeuroD1
Six1
NeuroD6
Hoxd3
Hoxa3
ZNF205
NeuroG1
KLF15



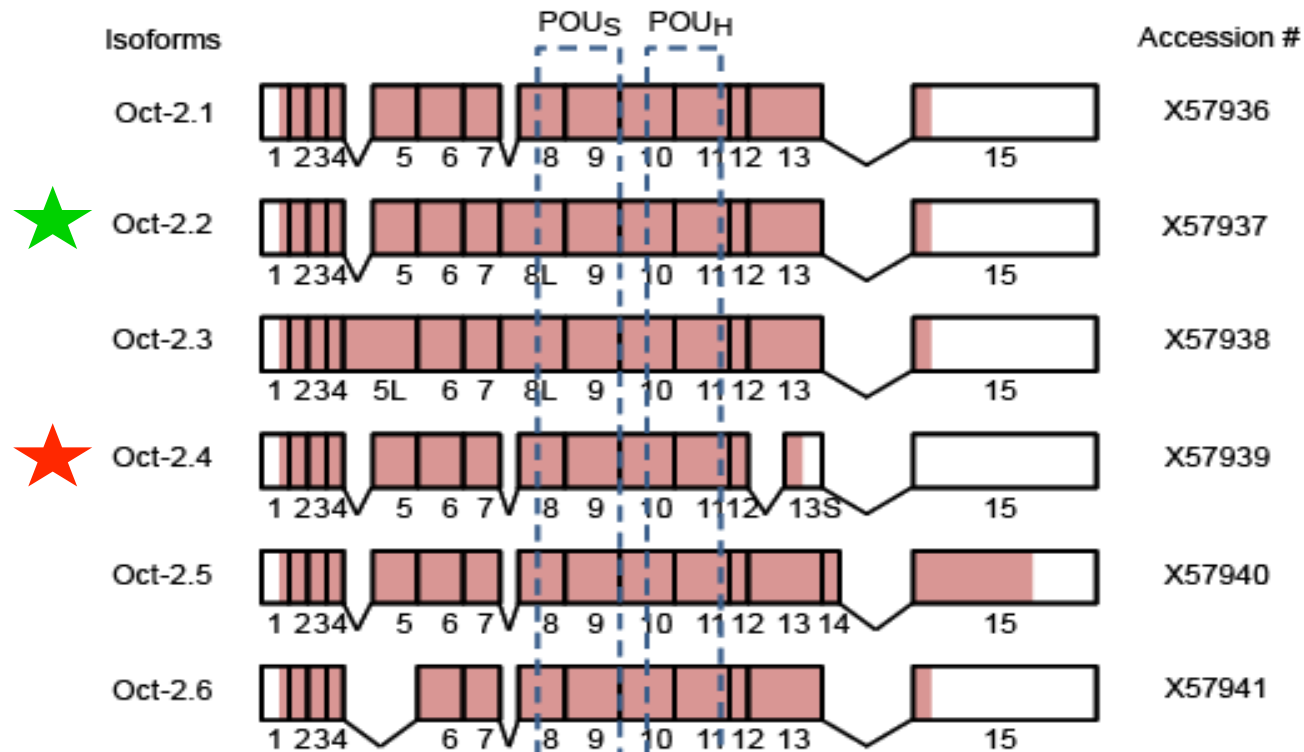
+ Control
Implicated
Repress
Novel

Oct-2 Many Isoforms

Activators Oct-2.1, 2.2, 2.5



Repressor Oct-2.4

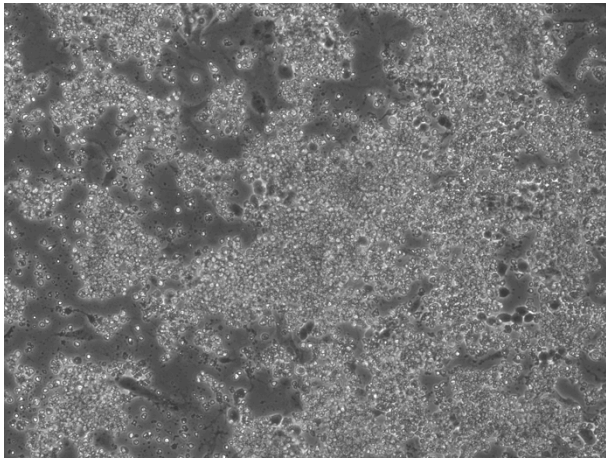


Adapted from Dong et al., 2007

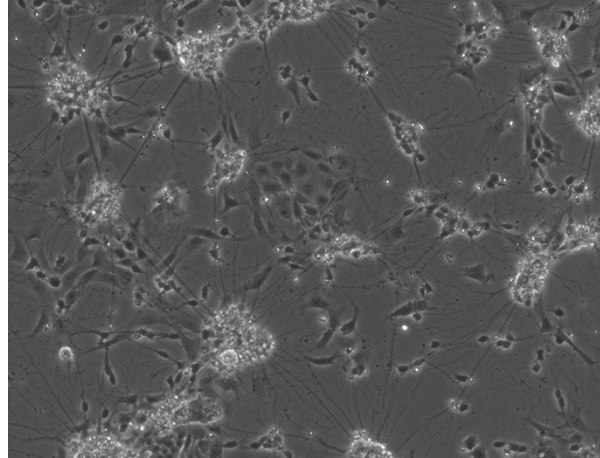
Oct-2.4 Inhibits Neuronal Differentiation

EBs dissociated and plated onto poly-ornithine/laminin coated plates

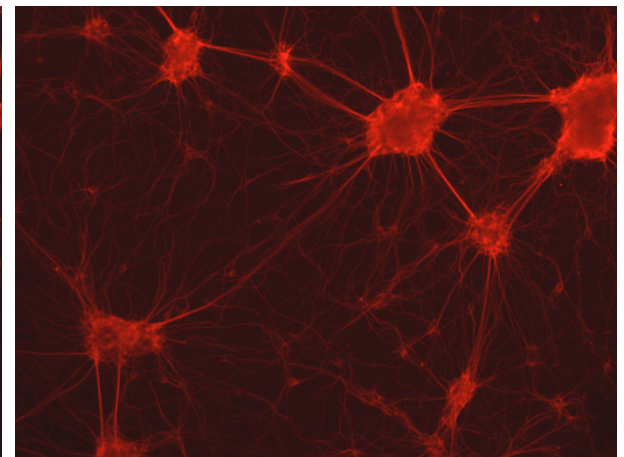
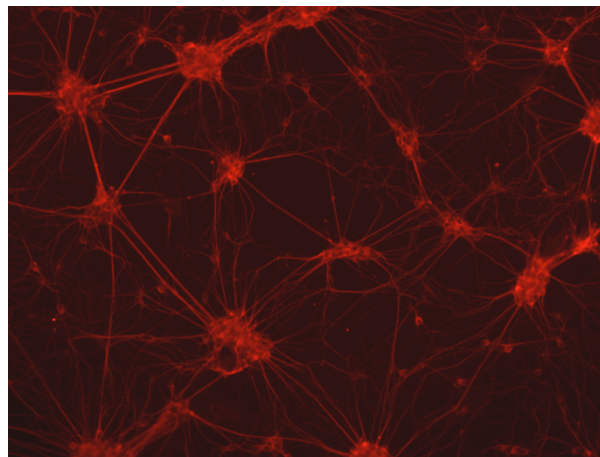
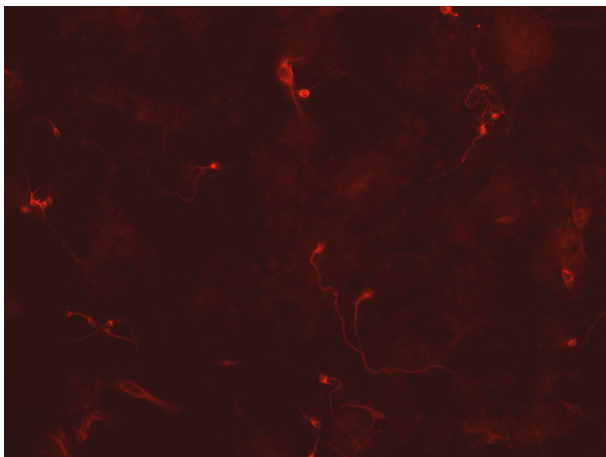
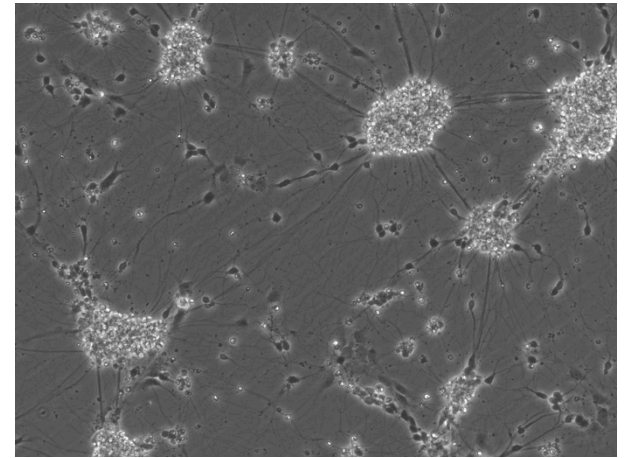
Oct-2.4



EBRTcH3 (WT)



Oct-2.4::VP16



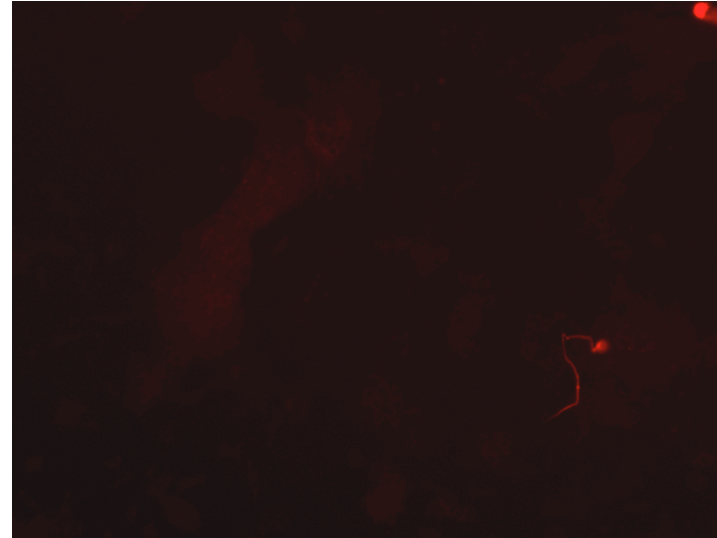
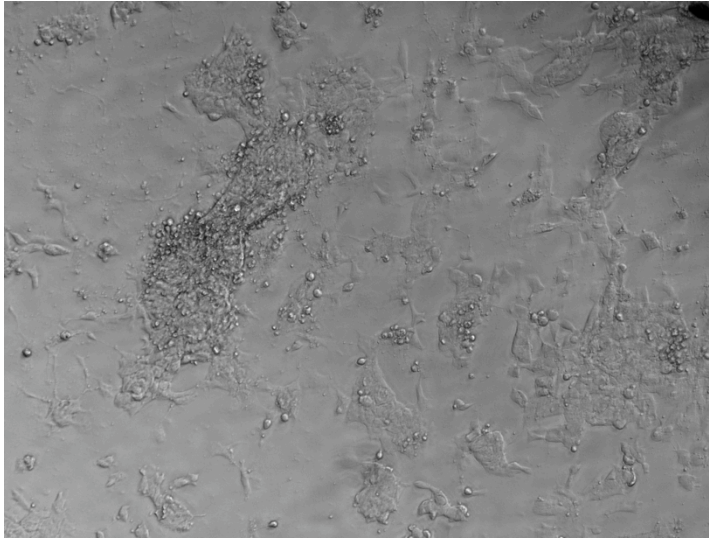
← anti-βIII tubulin →

Oct-2.2 Isoform Induces Differentiation

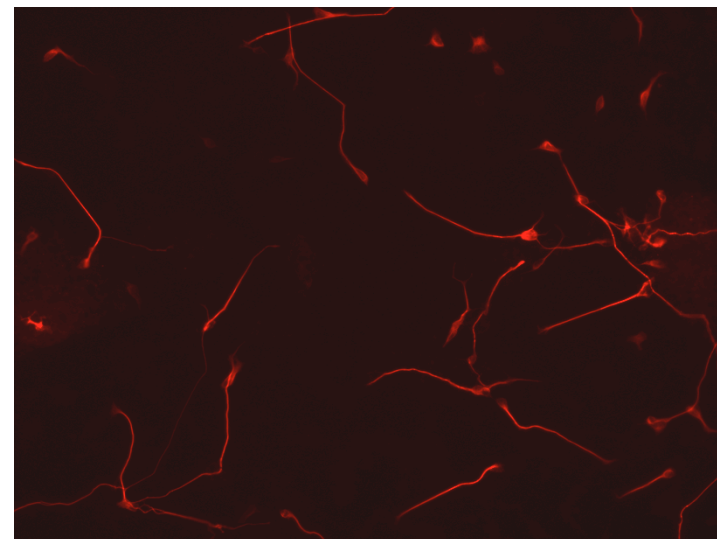
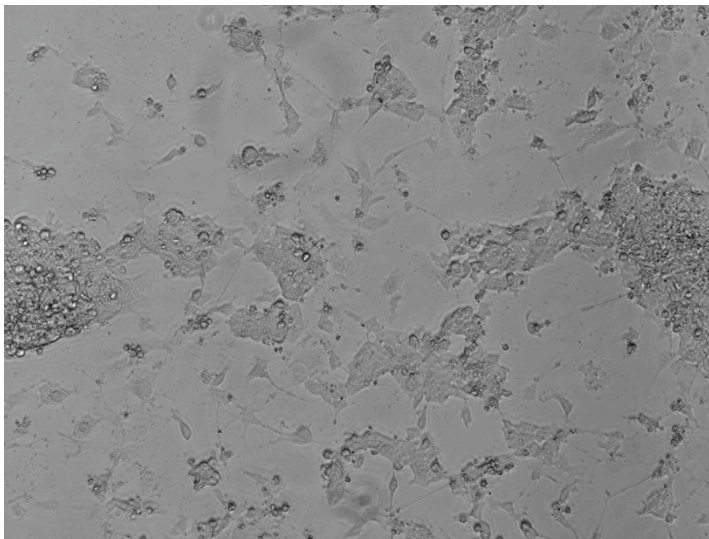
Phase

anti- β III tubulin

Uninduced
(+Tc)



Induced
(-Tc)



Oct-2.2

Oct-2 Differentiation Model

