
Therapeutic Strategies for Human Papillomavirus-Associated Cancers

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Why consider therapeutic approaches for HPV-associated lesions and cancers?

500,000 new cases/288,000 cervical cancer deaths per year (worldwide)
(2nd most common cause of cancer death in women)

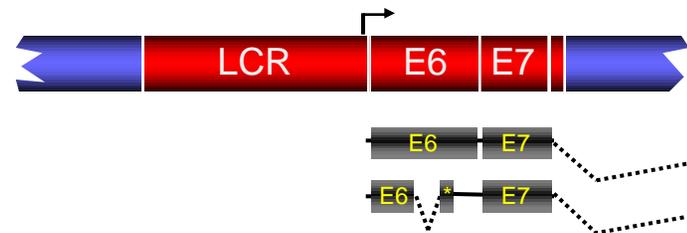
Prophylactic vaccines will only protect from new infections
by vaccine HPVs

Vaccine implementation has been challenging.
(US 2007: 25% of girls ages 13-17, 10% of all females ages 18-26
1.1% of Hispanic women)

Cervical cancer develops decades after the initial HPV infection

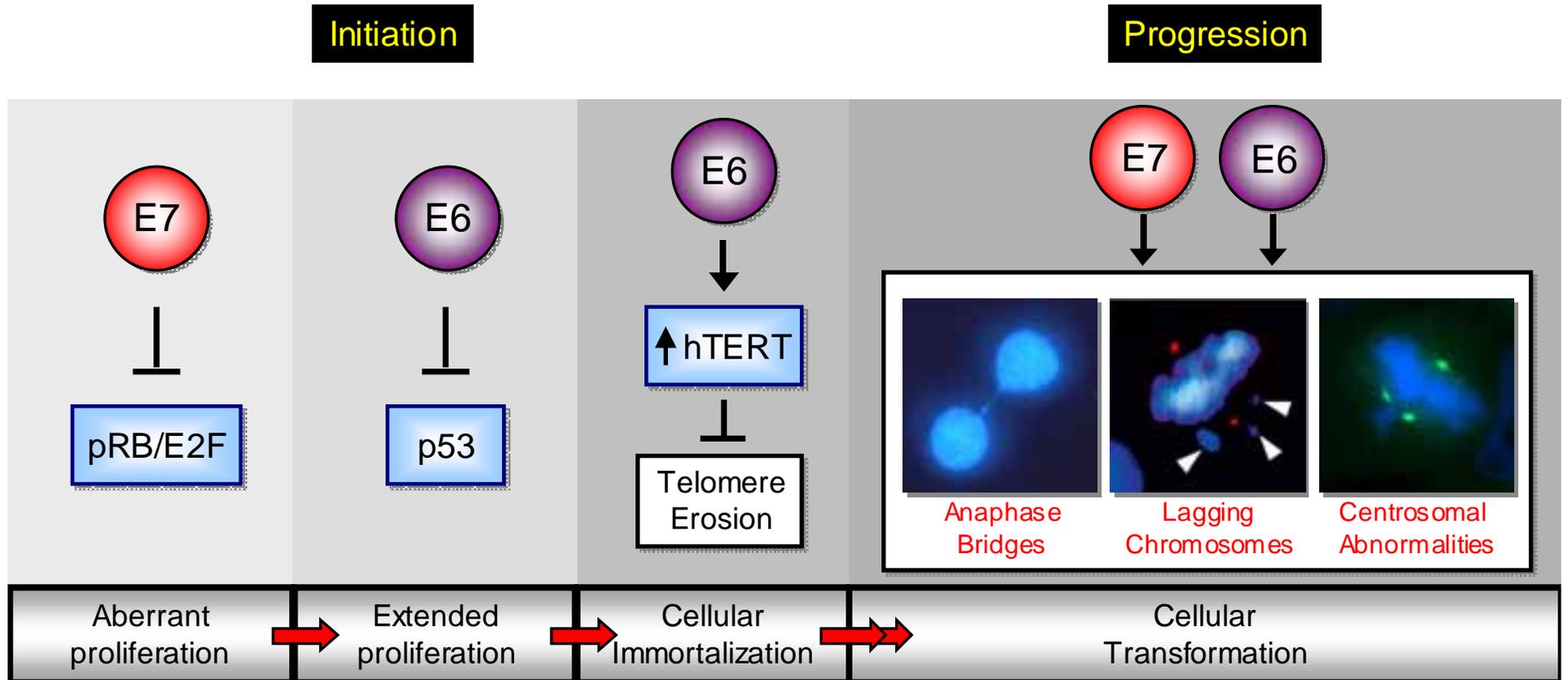
It will take decades before such vaccines will result in a measurable
decline of HPV-associated cancers

HPV genome integration is a hallmark of malignant progression



HPV-associated cancers only express two viral proteins, E6 and E7

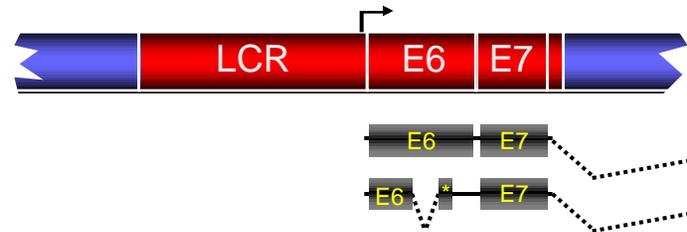
Mechanisms of HPV-associated carcinogenesis



HPV oncoproteins target cellular signaling pathways that are frequently mutated in human solid tumors:

pRB pathway >80%
p53 >60%
Telomere maintenance ~100%

Evidence that HPV E6/E7 expression is necessary and sufficient for cervical carcinogenesis



HPV16 E6/E7 expression in primary epithelial cells causes histopathological abnormalities reminiscent of CIN3

HPV16 E6/E7 expression causes cervical cancer in transgenic mice

Extinguishing HPV E6/E7 expression in cervical cancer cell lines causes cellular senescence/G1 arrest/apoptosis

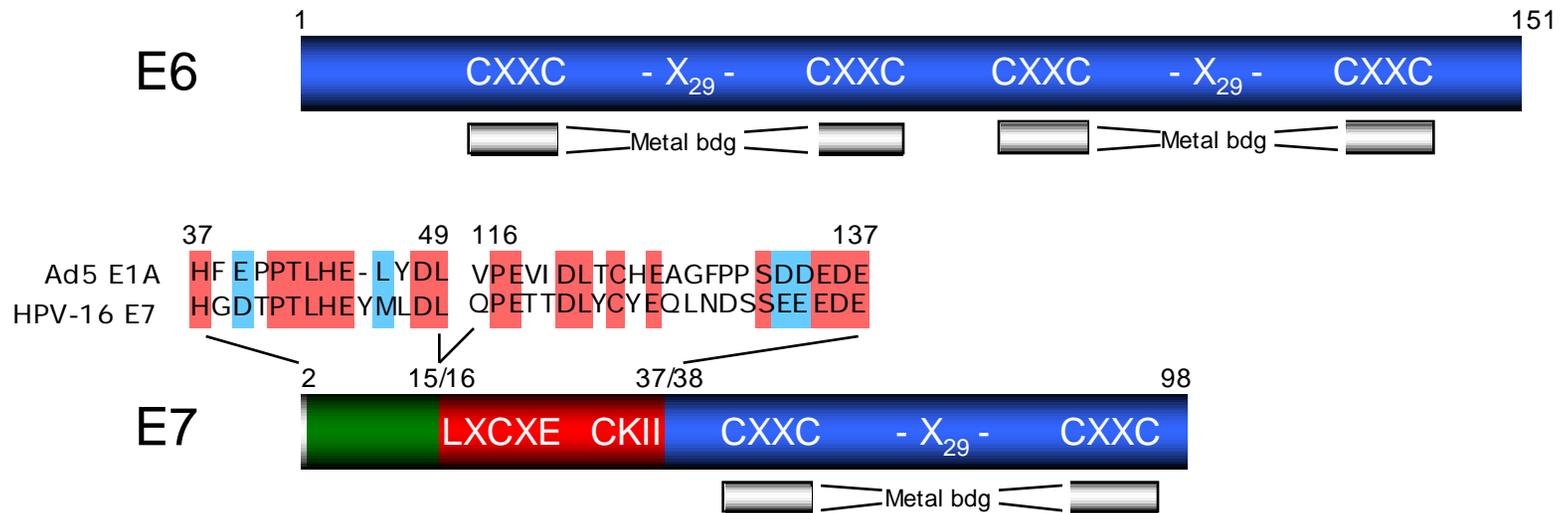
HPV-associated cancers are the only human solid tumors where the carcinogenic agent is known at a molecular level

Therapeutic opportunities

HPV-associated cancers are driven by expression of the E6 and E7 oncoproteins

HPV E6 and E7 oncoproteins should be excellent drug targets

HPV E6/E7 Oncoproteins



No cellular homologues

No intrinsic enzymatic activities

No specific DNA binding activities

Associate with and functionally modify host cellular protein complexes

Therapeutic opportunities

HPV E6 and E7 function through protein/protein interactions
(E6/p53; E7/pRB)

Protein/protein interactions are difficult to target by small molecule approaches

Identification of therapeutic targets

Are HPV oncoproteins associated with cellular enzymatic activities that are necessary for their activities?

Does expression of HPV oncoproteins induce perturbations of cellular signal transduction networks that may be harnessed for therapy?

Identification of therapeutic targets

Are HPV oncoproteins associated with cellular enzymatic activities that are necessary for their activities?

Proteomic analysis of HPV16 E7 associated host cellular protein complexes



Construction of stable cell lines
expressing double tagged E7



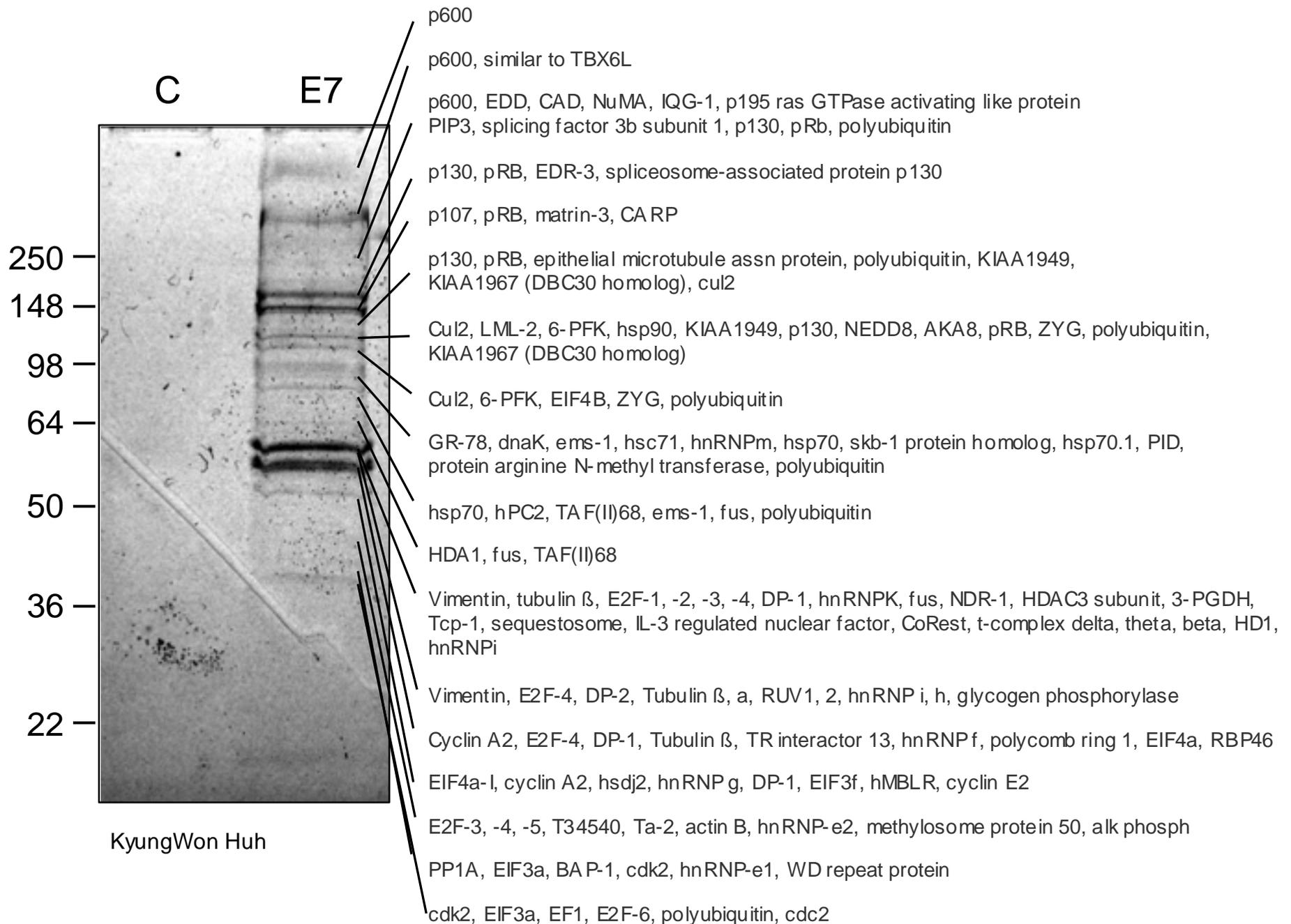
Sequential immunoaffinity purification with
anti-FLAG and anti-HA resin

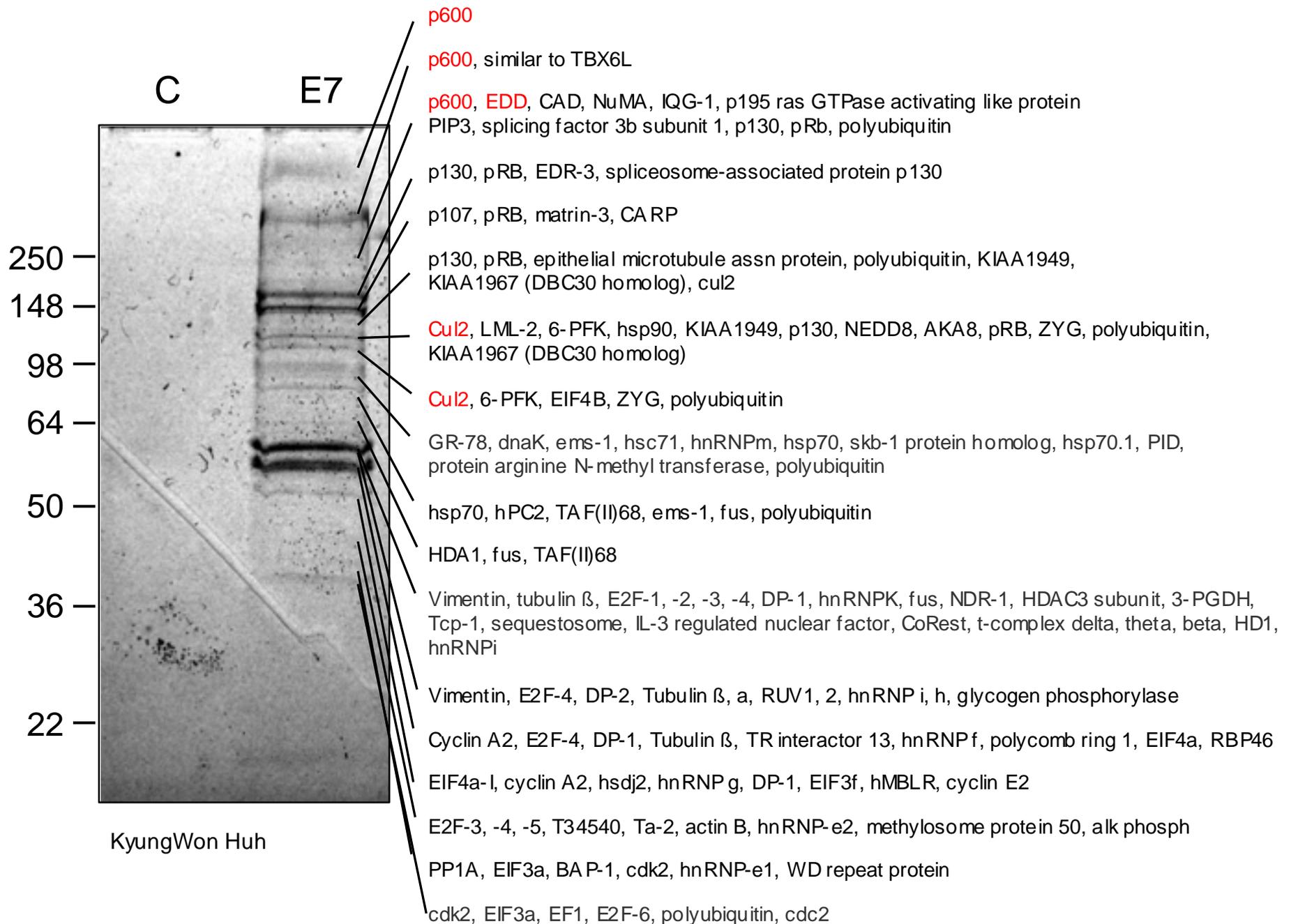


SDS-PAGE and stain

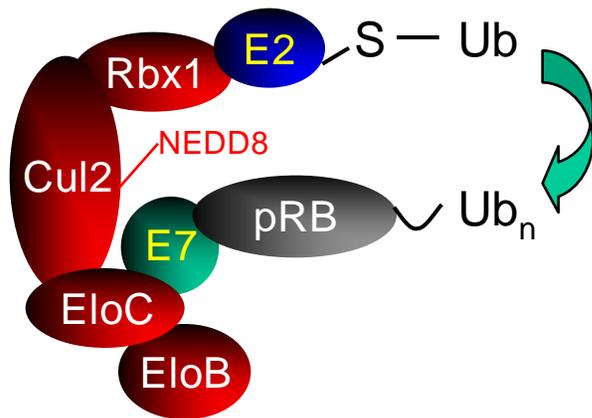


Identification of protein components by
mass spectrometry

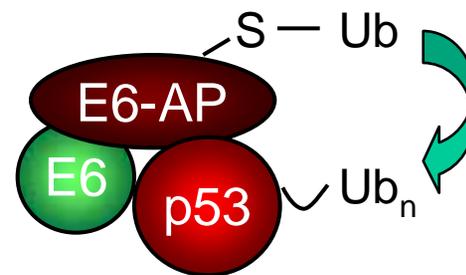




HPV E6 and E7 oncoproteins associate with and reprogram cellular enzymes



HPV16 E6 retargets the cellular cullin 2 ubiquitin ligase complex to the retinoblastoma tumor suppressor protein, pRB



HPV16 E6 retargets the cellular E6AP ubiquitin ligase to the p53 tumor suppressor protein

HPV E6 and E7 oncoproteins reprogram cellular ubiquitin ligases to target associated cellular tumor suppressors for degradation

Are HPV-positive cancer lines sensitive to the proteasome inhibitor, Bortezomib?

EC50s:

CaSki (HPV16): ~10-15 nM

SiHa (HPV16): ~30-35 nM

HeLa (HPV18): ~25 nM

Will test on HPV16 oncogene expressing primary human epithelial cells

Karin Hellner

collaboration with Jochen Lorch/Marshall Posner

Identification of therapeutic targets

Does expression of HPV oncoproteins induce perturbations of cellular signal transduction networks that may be harnessed for therapy?

“Under the Streetlight” Approach

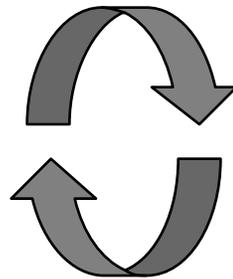


“You study what you can see”

“Unbiased” approach



“You see what you can study”



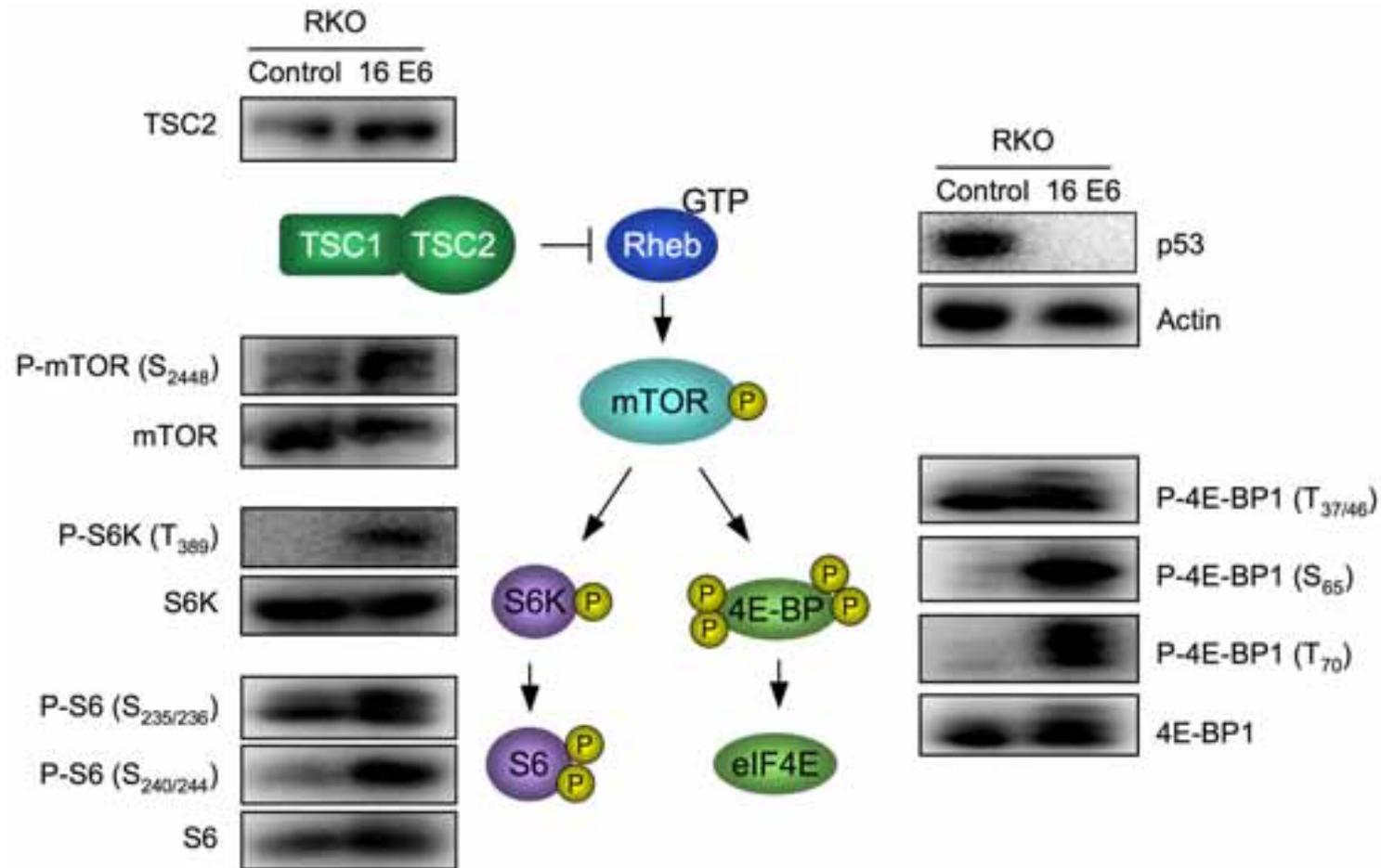
Identification of therapeutic targets

“Under the Streetlight” Approach



“You study what you can see”

HPV E6 expression activates mTOR signaling



Jennifer Spangle based on
Lu et al, JBC 279: 35664-70, 2004

Are HPV-positive cancer lines susceptible to mTOR inhibitors?

EC50s for RAD001:

CaSki (HPV16): ~15 μ M

SiHa (HPV16): ~25 μ M

HeLa (HPV18): ~27 μ M

Will test on HPV16 oncogene expressing primary human epithelial cells

Will test combination with 2-deoxyglucose

Karin Hellner

collaboration with Jochen Lorch/Marshall Posner

Identification of therapeutic targets

“Unbiased” approach



Genetic Screens

Systems Biology

“You see what you can study”

Identification of therapeutic targets

“Unbiased” approach

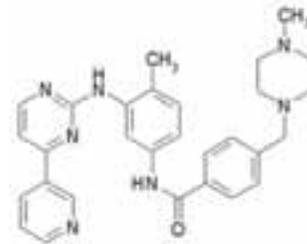


Genetic Screens

“You see what you can study”

Essential kinases for cervical cancers

Precedence: Gleevec® inhibits the bcr/abl kinase that is critical for the growth of certain leukemias (CML) that have the Philadelphia translocation



Can we identify a kinase target for a “cervical cancer-specific Gleevec”?



Ed Harlow



Dorre
Grueneberg



Karl Munger



Amy
Baldwin



Karin
Hellner



Miranda
Grace



Haoxuan
Tong

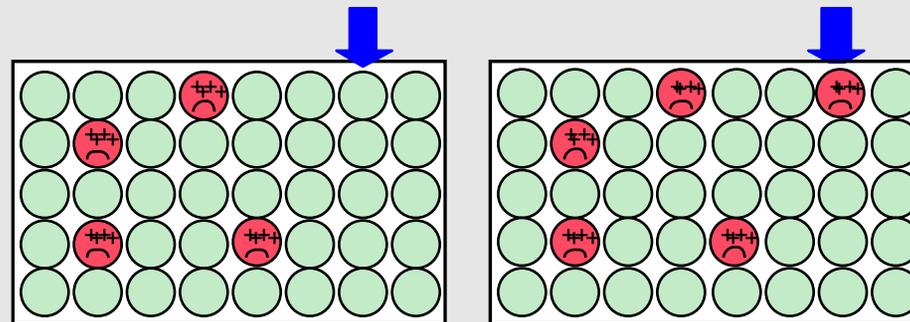
Essential kinases for cervical cancers: Experimental Strategy

Normal
Keratinocytes

Cervical
Carcinoma

“100 Hits” Kinase shRNA library
*Targets a subset of 80 kinases that
potently kill a variety of cancer cell lines*

Identify essential kinases for each cell population



*Normal
Keratinocytes*

*Cervical
Carcinoma*

ANPb

CDK7

EPHB1

HER3

HIPK2

IRR

JNK3

KHS1

MELK

MYO3B

PAK3

PAK6

PCTAIRE1

PDHK2

PITSLRE

PLK1

ROS

RSK2

SGK2

TSSK

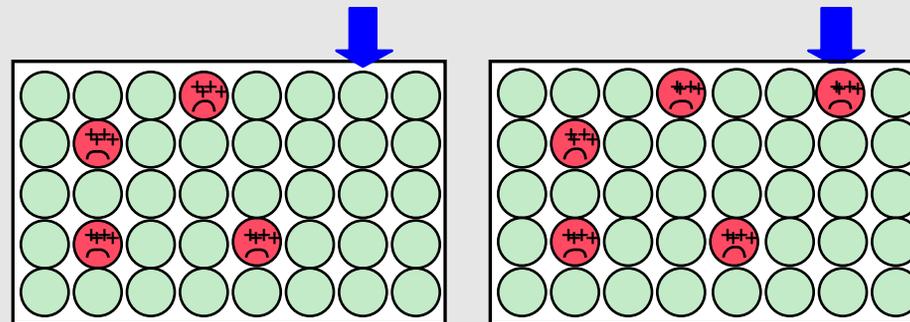
Essential kinases for cervical cancers: Experimental Strategy

Normal Keratinocytes	HPV E6 Expression	HPV16 Immortalized	Cervical Carcinoma
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“100 Hits” Kinase shRNA library

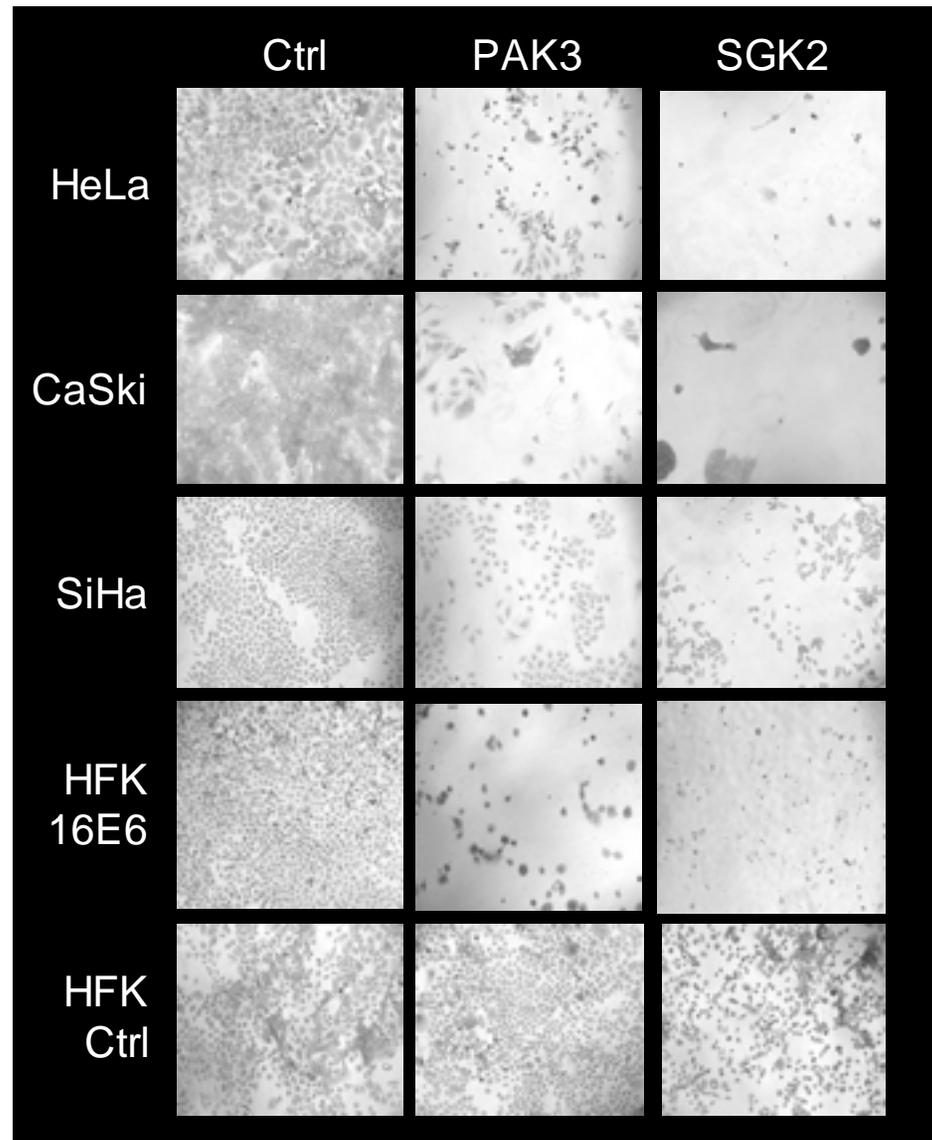
*Targets a subset of 80 kinases that
potently kill a variety of cancer cell lines*

Identify essential kinases for each cell population



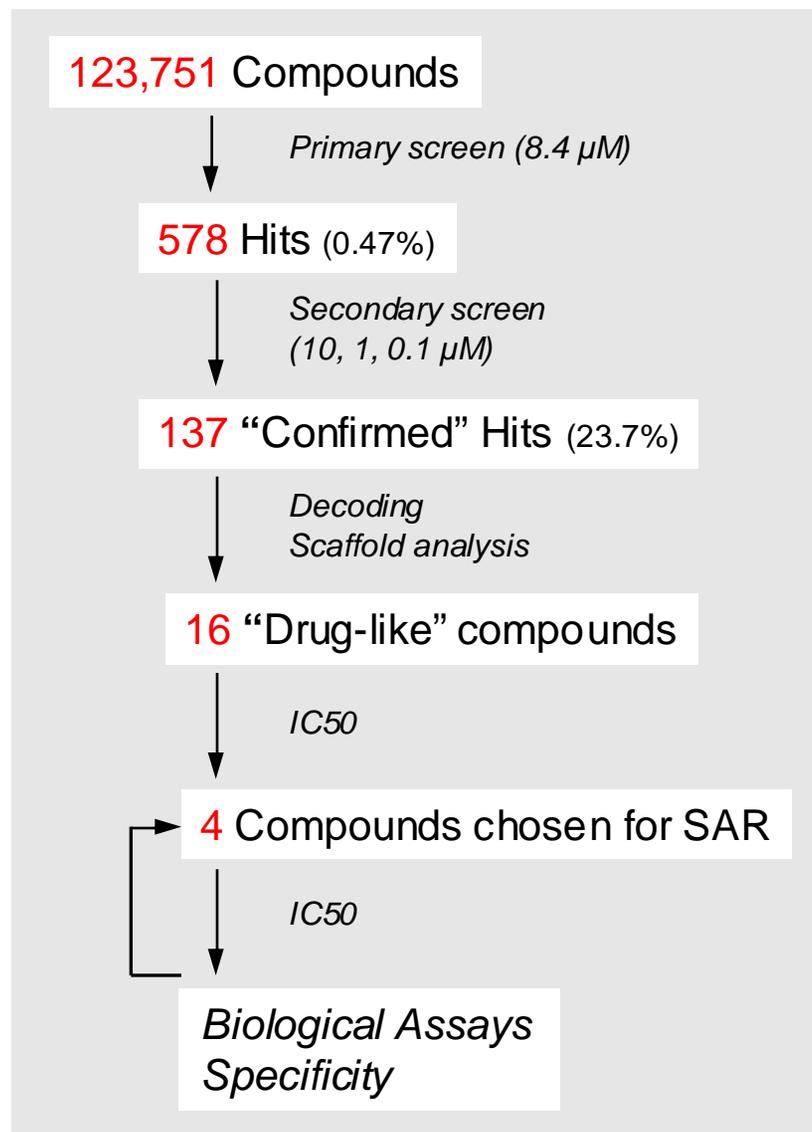
<i>Normal Keratinocytes</i>	<i>HPV E6 Expression</i>	<i>HPV16 Immortalized</i>	<i>Cervical Carcinoma</i>
		ANPb	ANPb
		CDK7	CDK7
			EPHB1
		HER3	HER3
		HIPK2	HIPK2
		IRR	IRR
		JNK3	JNK3
			KHS1
			MELK
		MYO3B	MYO3B
	PAK3	PAK3	PAK3
		PAK6	PAK6
		PCTAIRE1	PCTAIRE1
		PDHK2	PDHK2
		PITSLRE	PITSLRE
		PLK1	PLK1
			ROS
		RSK2	RSK2
	SGK2	SGK2	SGK2
		TSSK2	TSSK

PAK3 and SGK2 become essential upon HPV E6 expression



PAK3 Inhibitor Screen

Collaboration with Greg Cuny/Jun Xian (PCDD)



Karin Hellner

Identification of therapeutic targets

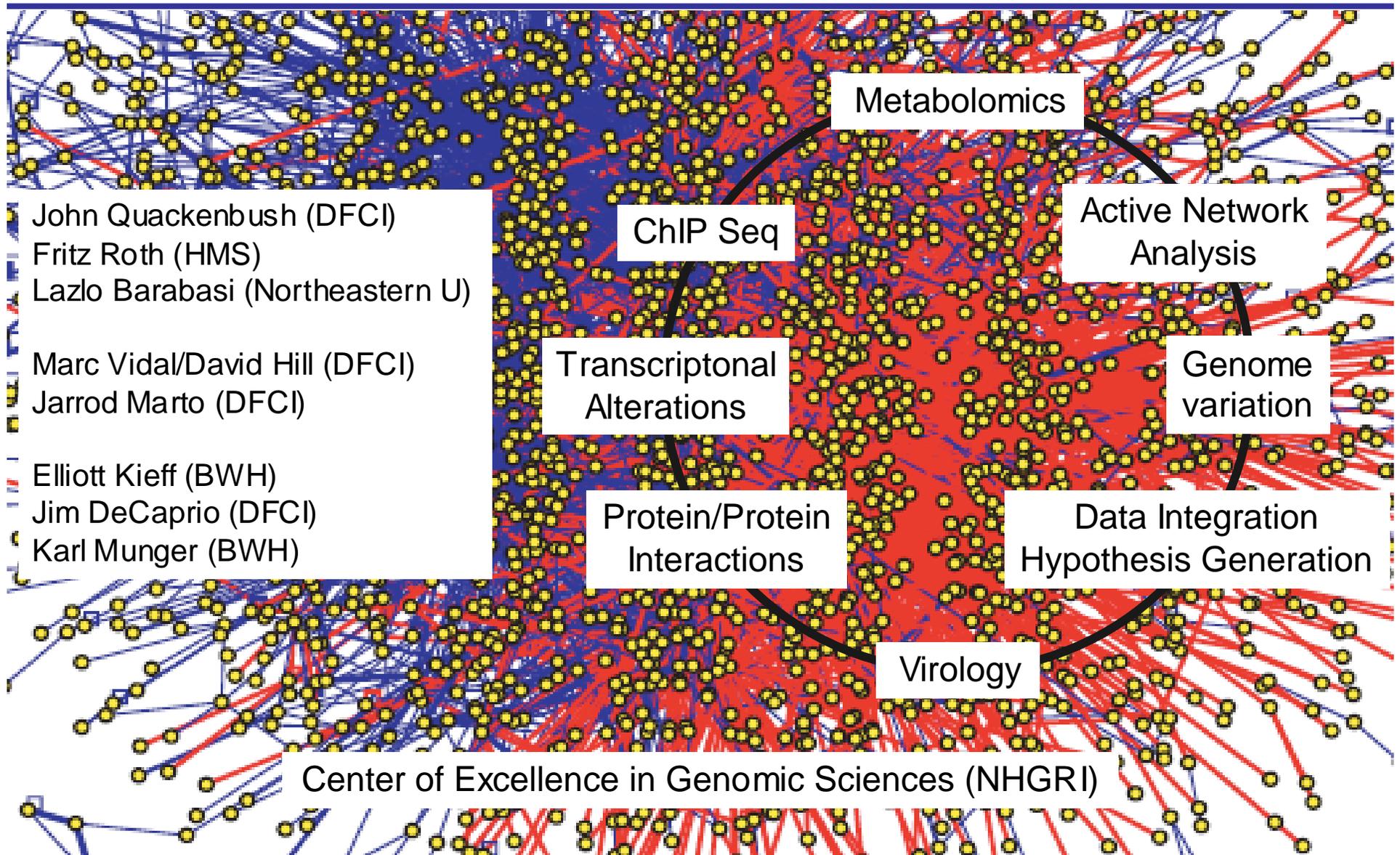
“Unbiased” approach



Systems Biology

“You see what you can study”

An integrative approach to identify cellular network perturbations induced by viral oncoprotein expression





Munger Lab

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Miranda Grace
Karin Hellner
Sergio Ita
Molly McLaughlin-Drubin
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Lessons and Take Home Messages

- Virus associated cancers offer unique opportunities for prevention, diagnosis and therapy
- Some mechanistic insights obtained with HPV-associated cancers should be generally applicable to non-virus associated human cancers
- Unbiased genetic screens and integrative, system biology based approaches will define novel therapeutic targets