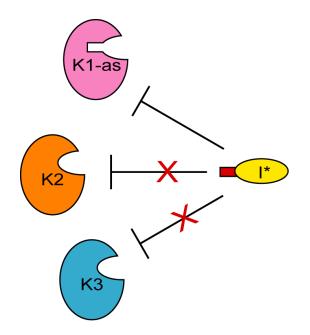
# Kinase Chemical Genetics



New Tools for Old Problems

MacMillan Group Meeting

July 1<sup>st</sup>, 2009

Alexander Warkentin

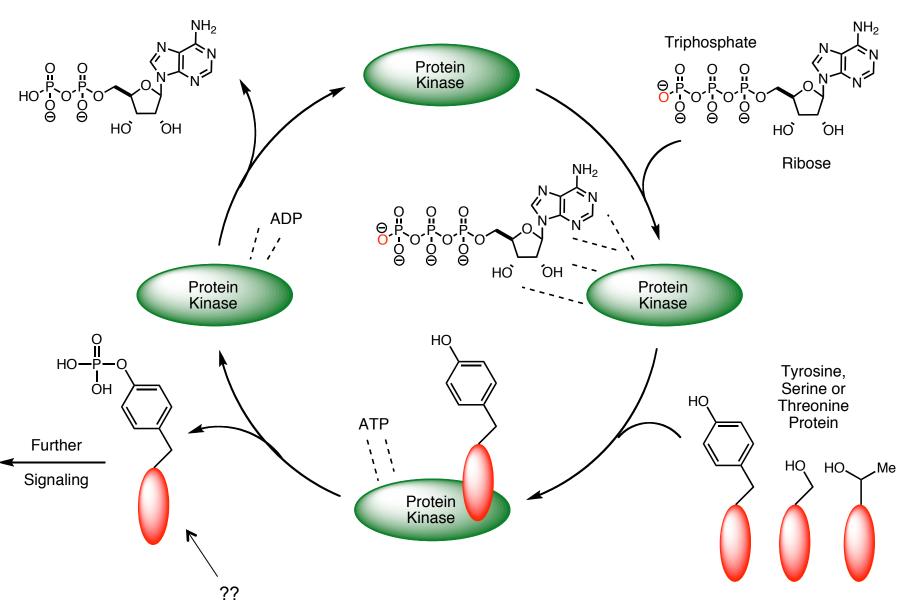
#### Outline

- Background biology notes
- Inspiration for kinase research
- Chemical genetics: inhibition, benefits for signaling subtleties
- Chemical genetics: A\*TP analogues, benefits for direct phosphoprotein substrate detection
- *in vivo* example

#### Kevan M. Shokat



- 1991 Ph.D. Chemistry, University of California, Berkeley
  - Advisor: P. G. Schultz; New Routes to Catalytic Antibodies
- 92' 94' Post-doctoral scholar, Stanford University
  - Advisor: C. C. Goodnow; Immune self tolerence in transgenic mice
- 94' 98' Assistant Professor of Chemistry and Molecular Biology, Princeton University
- 98' 99' Associate Professor of Chemistry and Molecular Biology, Princeton University
- 99'-01' Associate Professor of Chemistry, Berkeley; Pharmacology, UCSF
- 01' Professor of Chemistry, Berkeley; Pharmacology, UCSF
- 05' Investigator, Howard Hughes Medical Institute



Adenine

#### Kinases Covered

- CDK2 Cyclin-dependent kinase: critical for cell cycle
  - cdc28 yeast version of CDK2: used for easier analysis
- bcr, abl Together as a fusion protein kinase: leukemia
  - PYK2 Similar to bcr/abl in mode of inhibition
- v-Src First known kinase (Krebs, 1959): best studied
- EGFR Epithelial Growth Factor Receptor: breast, lung cancer
- Aurora B, Hck, Ire1, p110
- CAMKII Hippocampal long-term memory formation

Misregulation implicated in metastasis

# Background: Medium of Study

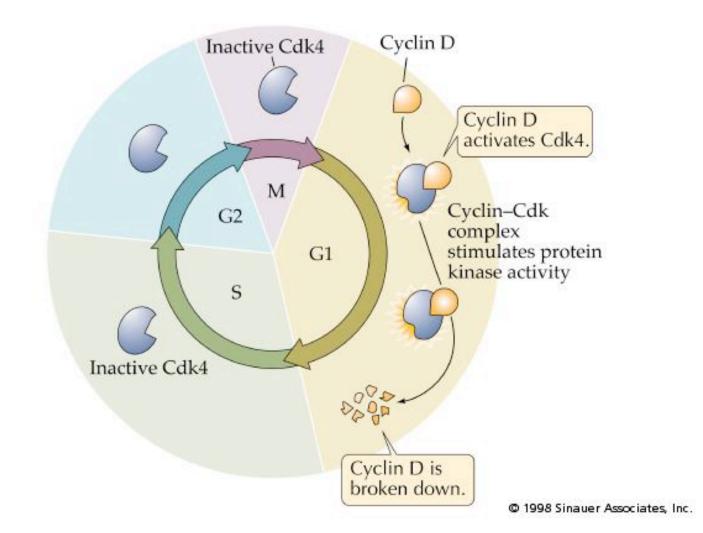
- *in vitro*: Isolated enzyme or other biomolecule.
  - First line of analysis; no interference or off target effects
  - Relatively fast; requires less than one milligram of small molecule
- *in lysate*: study occurs in cell-free environment in medium with other proteins
  - Not used very often
- In cell: mostly yeast cells
  - A non-trivial level of complexity already
- *in vivo*: Usually starts in the mouse
  - Mouse genome close to humans
  - Knockout experiments risk aborting embryogenesis; pharmacology risks off-target effects



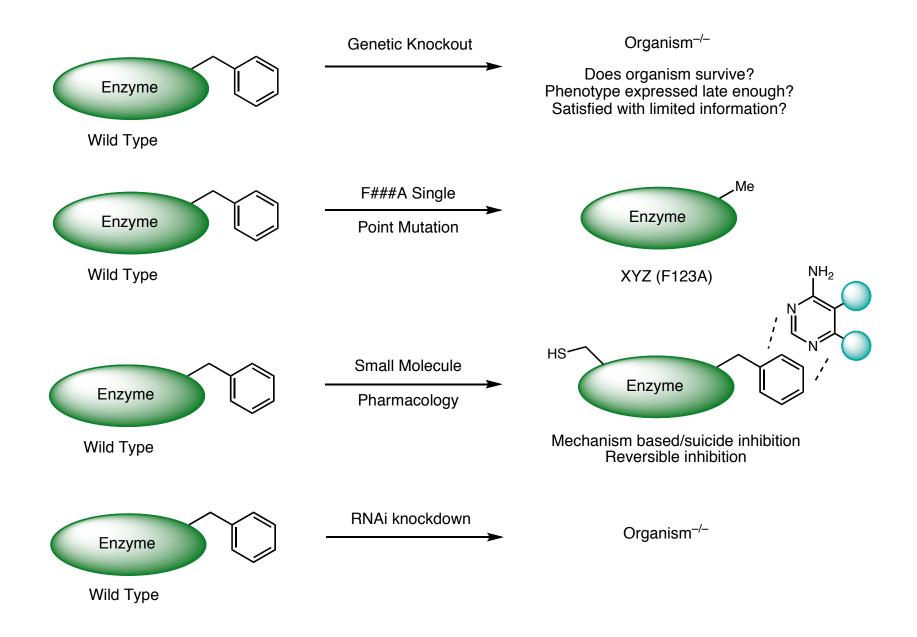




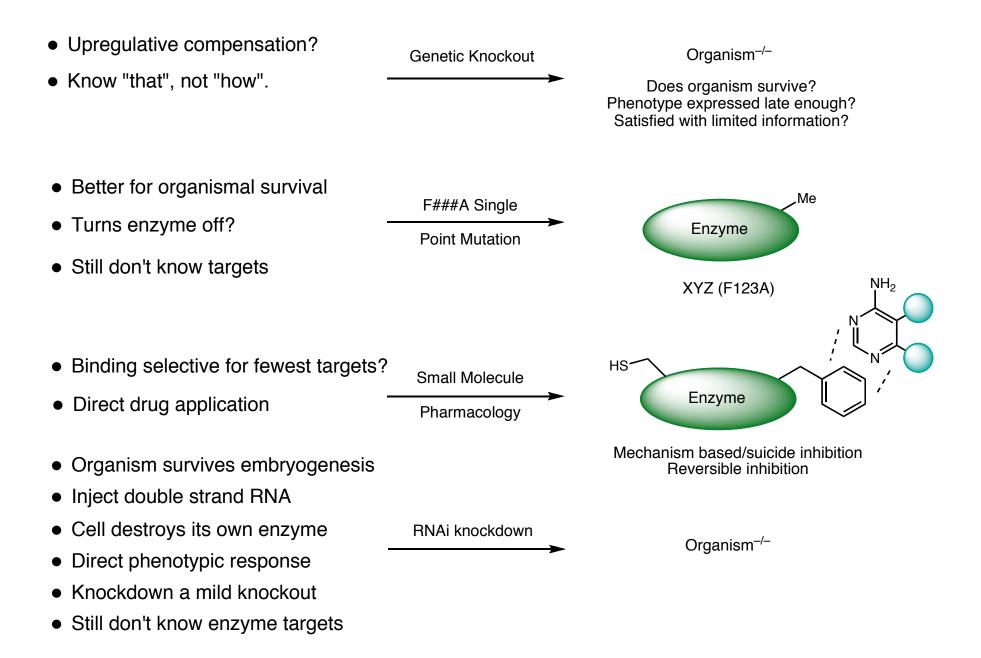
# Background: Cell Cycle



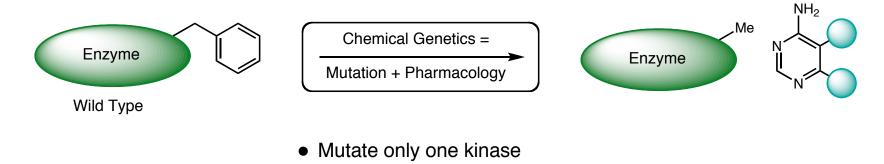
# Background: Experiments and Questions



# Background: Experiments and Questions

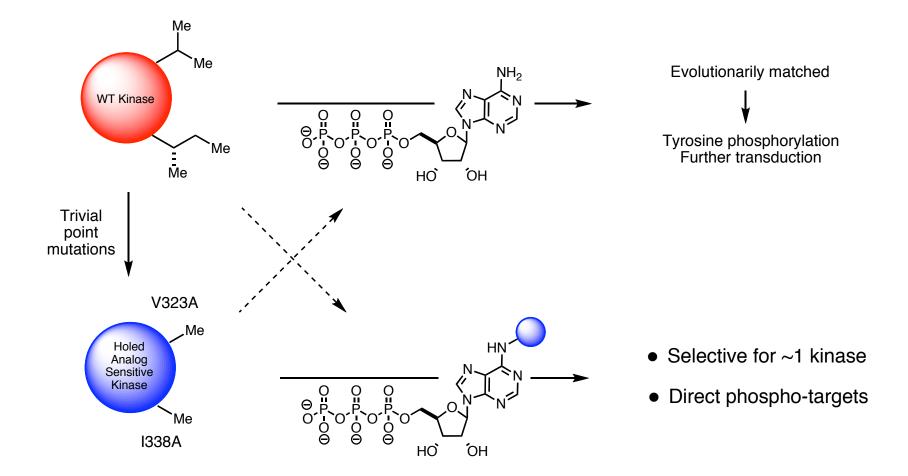


#### A New Experiment Combining Genetics and Pharmacology



- Match mutation with inhibitor
- Kinase specific information

#### A New Experiment Combining Genetics and Pharmacology



#### Gleevec (Imatinib) Heralded as the "Magic Bullet"

Gleevec treats Chronic Myelogenous Leukemia (CML)

At \$32,000 per year for a 400 mg per day dose, cited as a justifiably high cost for pharmaceutical innovation

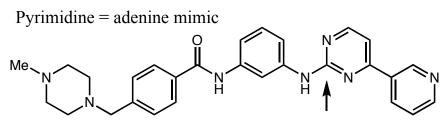
Novartis challenges Indian patent law: Madras High Court rejects claim

Sun Pharmaceuticals Industries Ltd. also challenges Novartis' US patent validity, which would set a decisive international precedent given the relative looseness of US patent law

Novartis wins: 1st world innovation saved; Sun wins: 3rd world affords drugs



April 21st, 2001



Gleevec (Imatinib)

# (Gleevec) Imatinib: Mode of Action

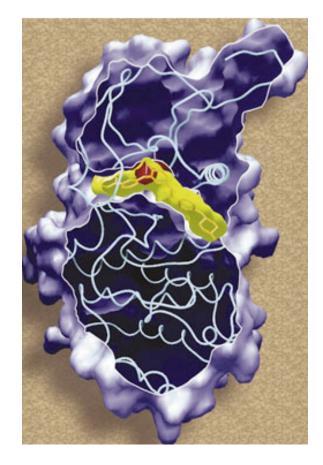
Gleevec treats Chronic Myelogenous Leukemia (CML) and has been approved for gastrointestinal and other malignancies

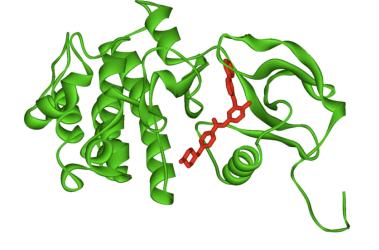
CML results from chromosomal mutation that effects translocation of the bcr and c-abl-encoding genes, the resulting fusion being termed the Philadelphia chromosome or bcr-abl oncogene

Expression of the bcr-abl fusion protein results in a myeloid cell line which is termed "growth factor independent for proliferation."

This thwarts apoptosis and leads to metastases

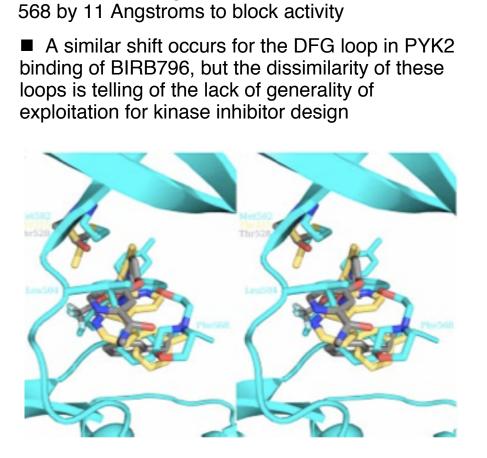
Treatable with bone marrow transplantation but only for 20% of patients due to age or compatibility





Druker B. J.; et al. Nature Medicine 1996, 2, 561.

# Explanation of Gleevec (Imatinib) Selectivity for Bcr-Able Fusion Kinase



Imatinib displays "bipartite" binding to both the

ATP pocket (conserved) and DFG-loop (not conserved)

Imatinib binding to Abl causes a unique shift of Phe-

Leu50 DFG-mot DFG-out DFG-in (inhibited) (active) Phe-568 Moves 11 Å

DFG = Aspartic Acid–Phenyl Alanine–Glycine

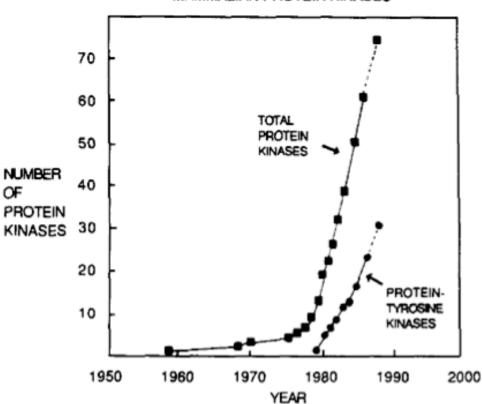
Han, S.; et al. J. Bio. Chem. 2009, 284, 13193.

High degree of homology has meant a rapid rate of discovery from molecular cloning of kinase genes

Appears that all eukaryotic protein Ser/Thr and Tyrkinases evolved from the same gene based on sequence

■ For example: cAMP dependent protein kinase shares300 amino acids with pp60 (v-Src) in catalytic domain alone

Downside is that pharmacology alone has trouble finding selective small molecule inhibitors



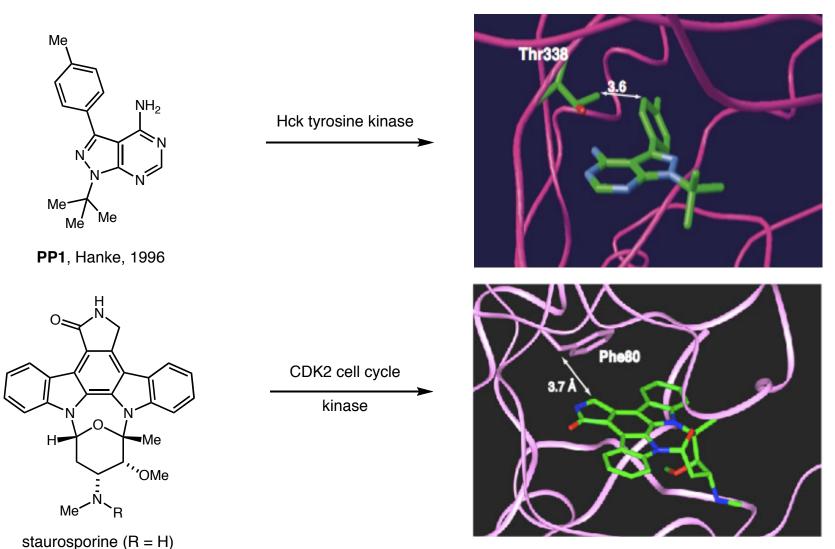
#### MAMMALIAN PROTEIN KINASES

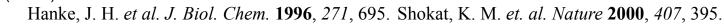
Hunter T. Cell 1987, 50, 823.

# PP1 is a Breakthrough in Src-Family Specificity

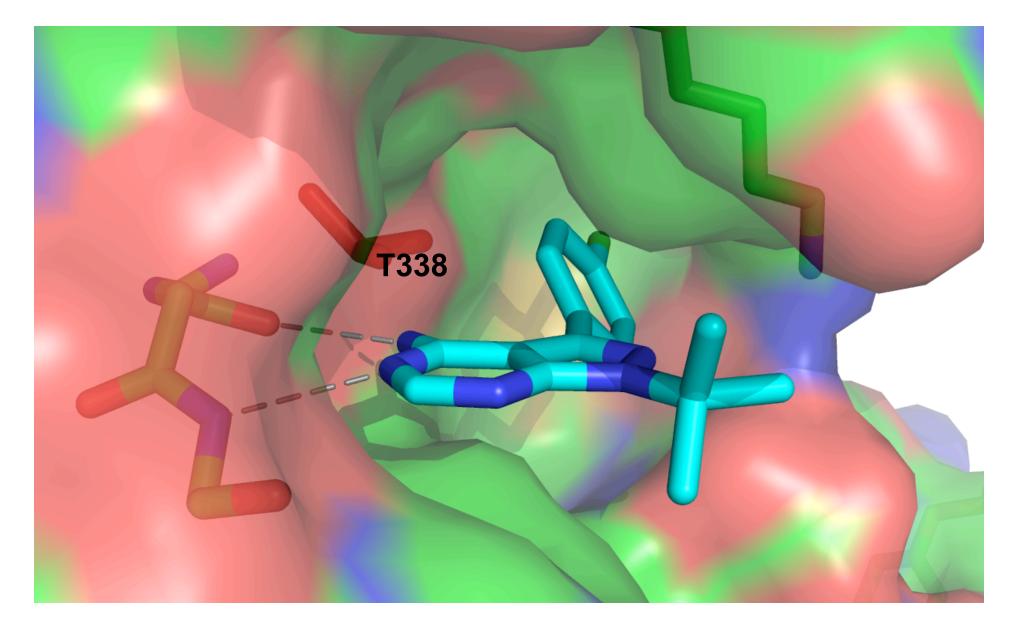
■ Hanke discovers PP1 series as selective kinase inhibitors in 1996 which is a more selective class for Src family kinases than previously reported staurospirine.

Both small molecules are well defined in their role in kinase inhibition; cocrystal structures obtained.





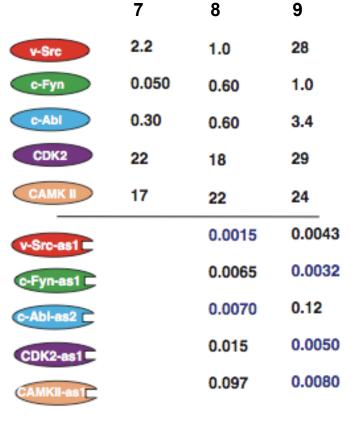
# Hck – PP2 Complex with Gatekeeper: T338



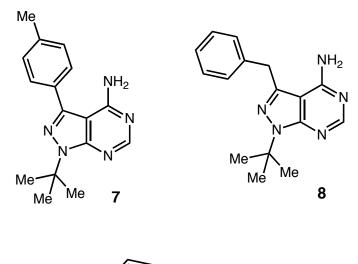
Schindler T., et al Mol.Cell 1999, 3, 639.

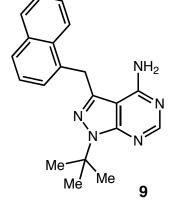
# New PP1 Analogues Display Unprecedented Binding Affinity and Specificity

- Analysis of cocrystal structure seggested extension of C7 substituent to increase selectivity
- Inhibitor 9 showed unprecedented activity toward kinase mutants relative to wild type (proof of princeiple)



Micromolar IC<sub>50</sub>

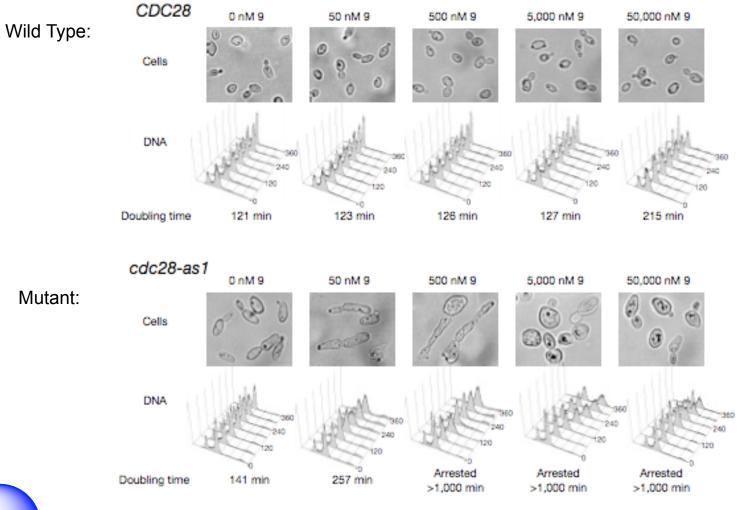




Shokat, K. M. et. al. Nature 2000, 407, 395.

### PP1 Analogues Show Cell Cycle Arrest of Cdc28-121 mutant

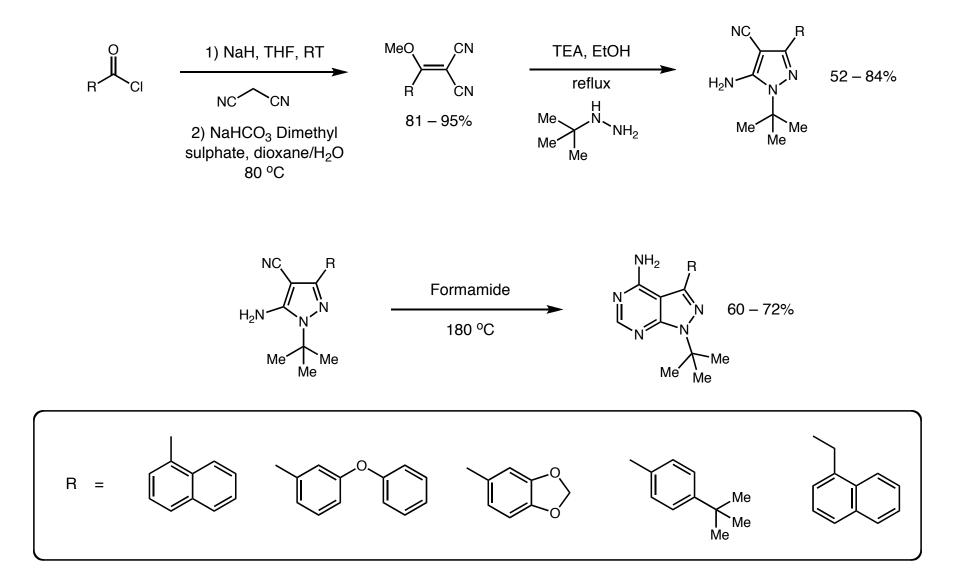
■ Cell cycle doubling in budding yeast affectively shut down while wild type cells are unaffected.



in cell

Shokat, K. M. et. al. Nature 2000, 407, 395.

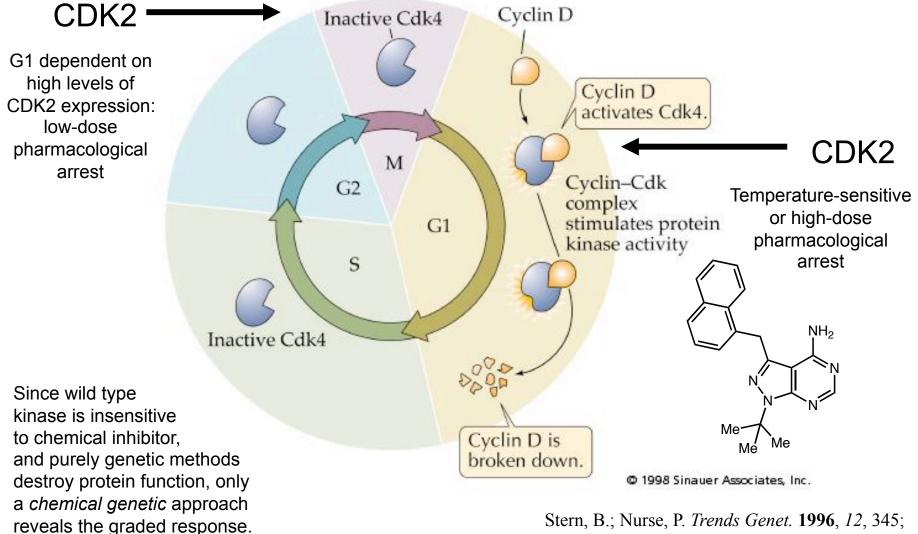
# PP1 Inhibitor Synthesis



Examples of Benefits of Chemical Genetics Techniques

# Gradational Response Discovered for Cell Cycle Inhibition with ASKA

Cell cycle progression assumed to be turned on by CDK2: by knockout of cdc28 or temp. shock
 Bishop and Shultz found a gradational response when PP1-derivative given to analog sensitive cdc28



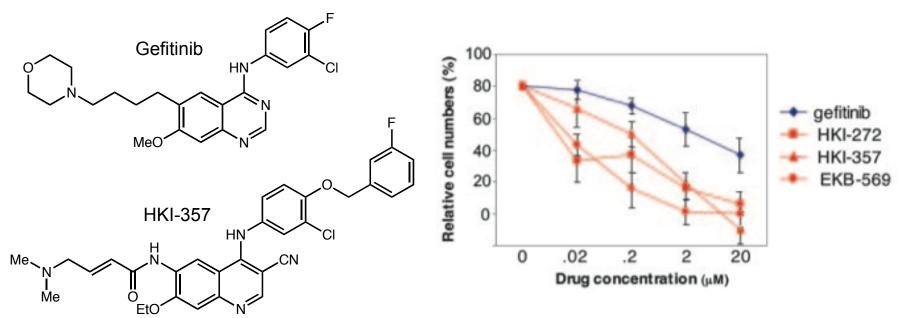
Bishop, D. P.; et al. Nature 2000, 407, 395.

#### Importance of Gradational Responce in Clinical use of Gefitinib

■ Gefitinib is an inhibitor of the tyrosine kinase domain of epidermal growth factor receptor (EGFR)

- EGFR misregulation implicated in several cancers inlcuding breast and non-small cell lung cancer
- Many patients have tumors resistent to gefitinib; tumors continue to grow
- Resistance is not due to EGFR mutations (as we would expect); tumor lines selected for resistance do not acquire mutations.

- These tumors simply afford resistance by increasing their threshold for EGFR inhibition as a result.
- Since the mode of action is technically still the same, suicide inhibitors are then effective at inhibition.



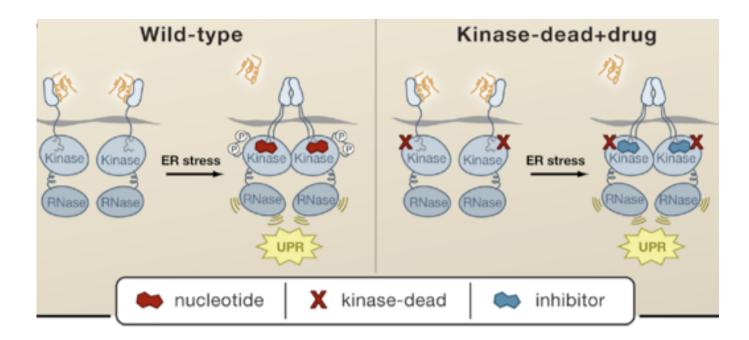
Kwak, E. L.; et al. Proc. Natl. Acad. Sci. USA 2005, 102, 7665.

# PP1 Uncovers a Mode of Action for the Unfolded Protein Response

■ Knockout of Ire1 or expression of a kinase-dead allele blocks the unfolded protein response (UPR)

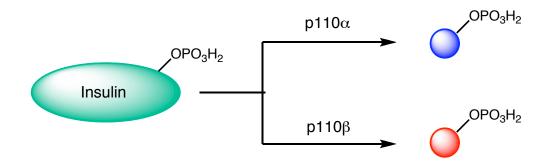
- An ATP competitive inhibitor of the kinase-dead allele rescues the UPR
- This means that  $PP_1$  acts as an  $Ire_1$  agonist rather than an  $Ire_1$  inhibitor, even though it binds to the  $Ire_1$  active site and directly blocks kinase activity (!).

Explanation: An ATP competitive ligand for the Ire1 kinase domain allosterically activates the Ire1 RNAse domain during the UPR



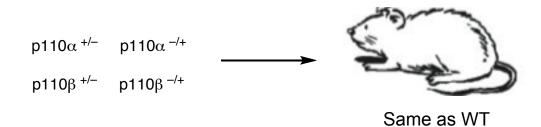
Patil, C., Walter, P. *Curr. Opin. Cell Biol.* **2001**, *13*, 349; Papa, F. R., Zhang, C., Shokat, K. M.; Walter, P. *Science* **2003**, *302*, 1533.

Both PI3 kinases, p110 $\alpha$  and p110 $\beta$ , carry signals from the growth factor, insulin

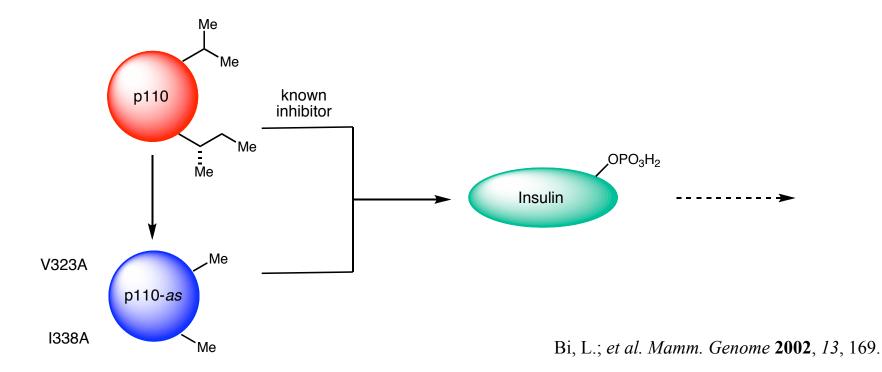


Bi, L.; et al. Mamm. Genome 2002, 13, 169.

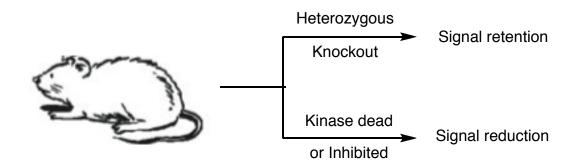
- Both PI3 kinases, p110 $\alpha$  and p110 $\beta$ , carry signals from the growth factor, insulin
- Because knockout of either kinase isoform kills mice early in development we know that they cannot compensate for each other but don't know their roles
- Heterozygous deletion of either produces no phenotype; deletion of p85, the p110 binding partner, paradoxically increases insulin signaling



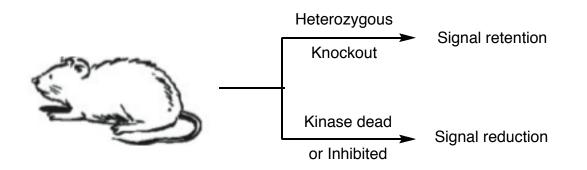
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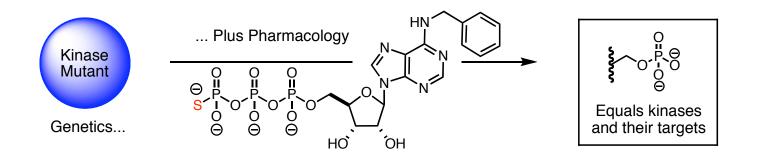
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- Resulting Model: since p85 can function as a negative regulator of p110, insulin transduction controlled by relative stoichiometry of p110 to p85 rather than absolute amounts.



Chemical Genetics Techniques for Substrate Identification

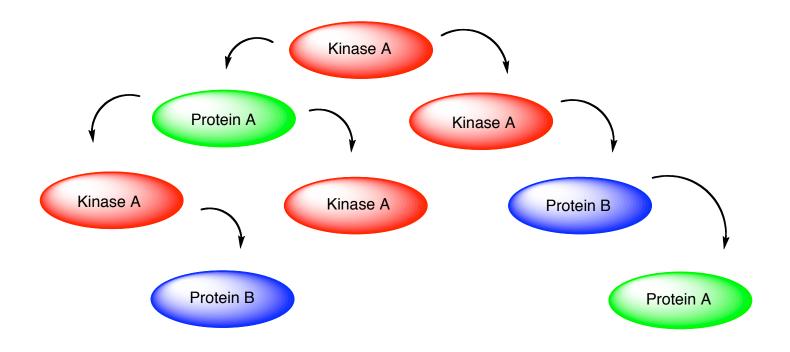


### Previous Phospho-Protein Substrate Identification Methodology

■ MALDI-TOF mass spectrometry of cell lysates requires high concentration of desired phosphoprotein substrate relative to non-desired phosphate-containing lysates

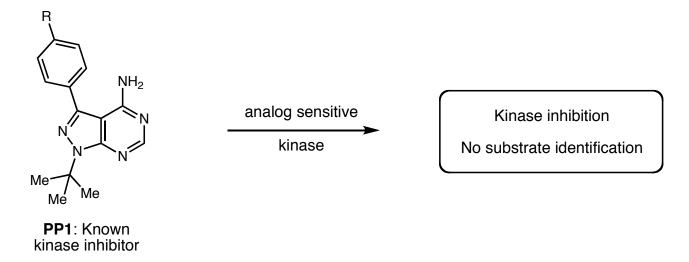
■ Protein lysate chromatography involves analyzing individual column fractions for their activity against a particular kinase; tedious and has problem that kinases are promiscuous in vitro

■ Yeast two hybrid screen not applicable when a third party protein is involved

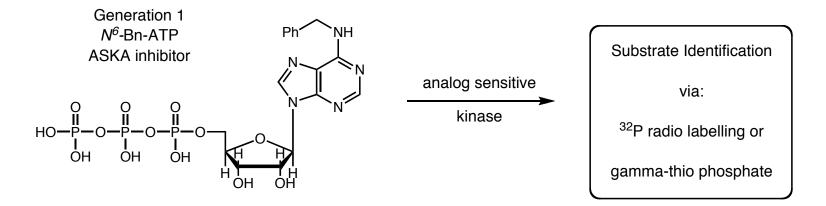


PP1 Inhibitor Usage Contrasted with Bumped N-Bn ATP Analog Usage

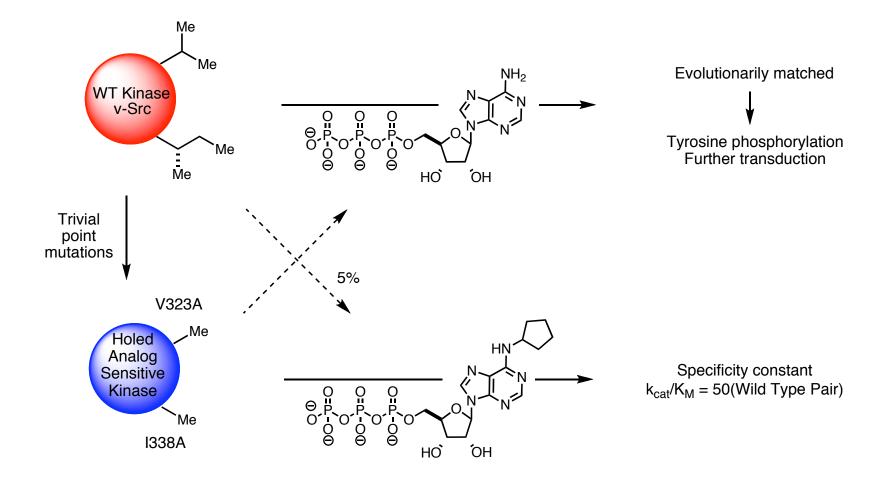
■ PP1 inhibitor derivatives serve to study kinase signaling pathways by inhibition



■ ATP analogs allow for substrate identification through gamma phosphate manipulation

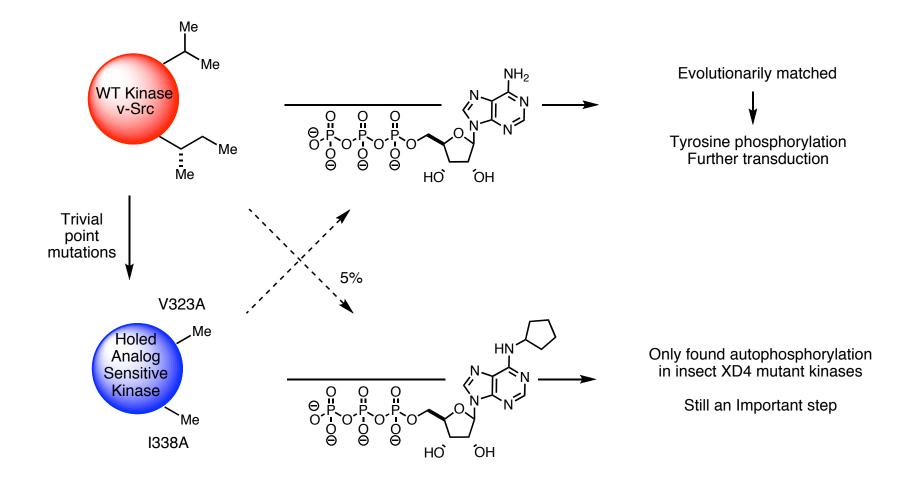


#### First Example of Potent and Selective ATP Analog



Shokat, K, M,; et al. Proc. Natl. Acad. Sci. USA 1997, 94, 3565.

#### First Example of Potent and Selective ATP Analog



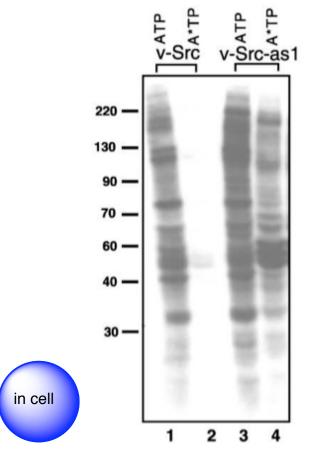
Shokat, K, M,; et al. Proc. Natl. Acad. Sci. USA 1997, 94, 3565.

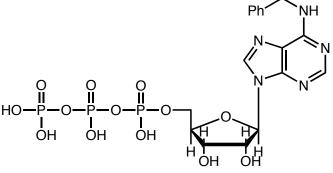
# ASKA Strategy Leads to Identification of Novel v-Src Targets

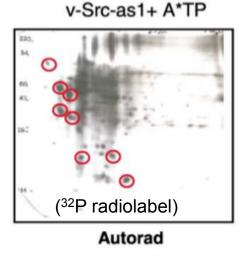
■ v-Src phosphorylates tyrosine on 50 proteins but any could come from phosphorylation events from other kinases that v-Src interacts with.

■ Radiolabeling of an ATP analog reveals precise phosphotyrosine substrates since the interaction orthogonally adds a <sup>32</sup>PO<sub>4</sub><sup>2−</sup> radiolabel only to v-Src-as1 (analog sensitive mutant) targets.

■ Cofilin not a known phosphotyrosine product of v-Src; results presumed faulty. ASKA strategy confirms Cofilin and calumenin as novel targets of oncogene v-Src.



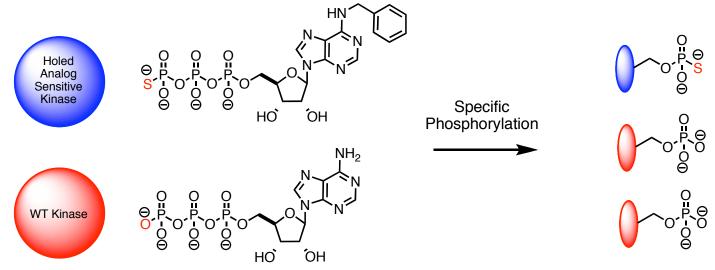




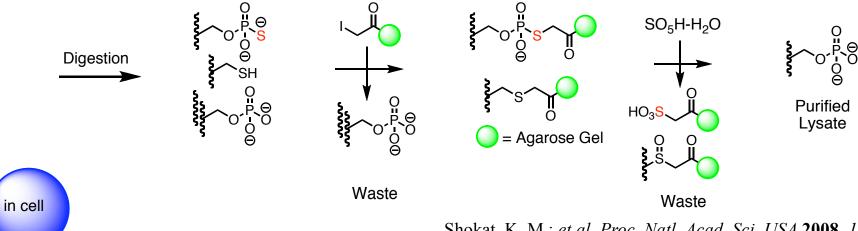
Shah, K.; Shokat, K. M. Chem. Biol. 2002, 9, 35.

# Thio-Phosphate Tagging of as-Kinase Direct Phosphorylation Products

■ Gamma thio *N*-Bn ATP selectively binds to Cdk1-cyclin B (F80G) then yielding chalco-differentiated tyrosine phosphorylation targets



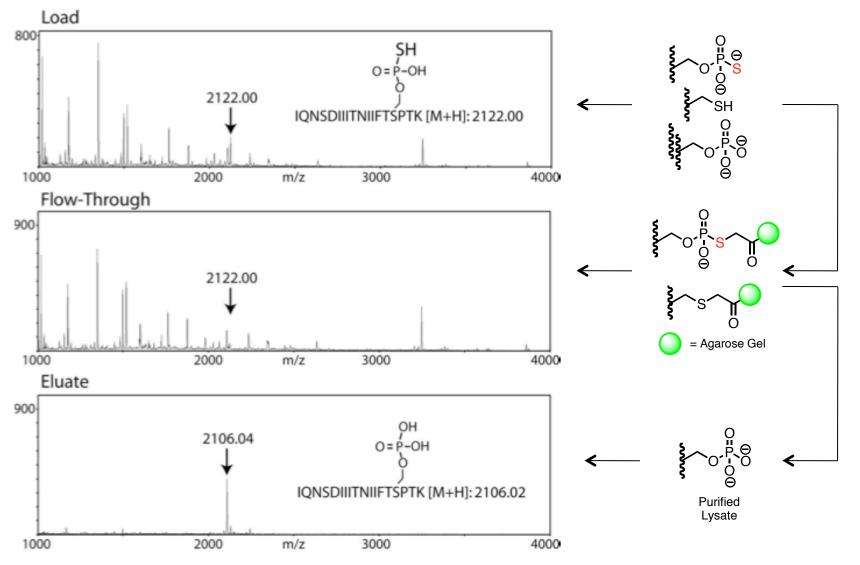
After lysis, oxo-phosphate derivatives are removed by selective alkylation; cystein products are washed away via Oxone oxidation, concommitantly re-oxidizing thio-phosphate



Shokat, K, M,; et al. Proc. Natl. Acad. Sci. USA 2008, 105, 1442.

Thio-Phosphate Tagging of as-Kinase Direct Phosphorylation Products

MALDI-TOF analysis of each stage of substrate purification



Shokat, K, M,; et al. Proc. Natl. Acad. Sci. USA 2008, 105, 1442.

#### Chromosome **Proteolysis: Endocytosis:** GAP and VPS9 Segregation: **TPP1 201** domains 1 987 **TPX2 738** mRNA Unclassified: Subcellular Transcription, Desmoyokin 2833 Localization: **Processing and EF hand domain** Nup358 2251 family member D2 74 Regulation: **Signal Sequence NUCKS 181** Amida 180 **Receptor Alpha 268 APRIN 1370** PYM Protein 72 **Protein Folding: SAMHD 1 592** HMG I/Y 52 \* Calnexin 583 Zinc Finger CCCH-type hnRNP-K 216 ERp57 456 containing 11A 321 HuR 202 Ubiguitinylation: RNA Pol II 1934 Suppresor of Ty 5 NICE-4 454 homolog 666 **NIPA 395 SRRM2 866** Ubp14 234 SRRM2 1413 Chromatin Structure: U2AF65 79 Histone H1a 17 rRNA Metabolism: Histone H1e 17 Antisense ERCC1 287 NSBP1 31 NOLC1 607 Zinc Finger Nucleolin 83 Protein 261 826 Nucleophosmin 70 Protein Nucleophosmin 237 \* **Phosphorylation:** Treacle 156 Signal **AAK1 389** Treacle 506 Transduction: PKL2 535 Treacle 906 **GYF2 30 PP1 Regulatory** IIRS2 391 Cellular Structure: Subunit 12A 409/ **Biosynthesis and OGFR 378** Cortactin 405 PP1-Alpha 320 \* Metabolism: PI4K Beta 266 Diaphanous-Related ADE2PUR6 26 Formin 1 747 Nuclear Structure: **DNA Replication** dUTPase 99 KI-67 761 Dynamin 1and Repair: **GMP-synthase 318** Related 616 Lamin A 22\* **DNA Ligase 176** HMG-CoA synthase 495 MAP7 209 Lamin A 392 \* DNA Topo. II Pyrroline-5-carboxylate Palladin 641 Lamin B 23\* Alpha 1213 \* reductase 2 304 Palladin-Related 155 Lamina-Associated

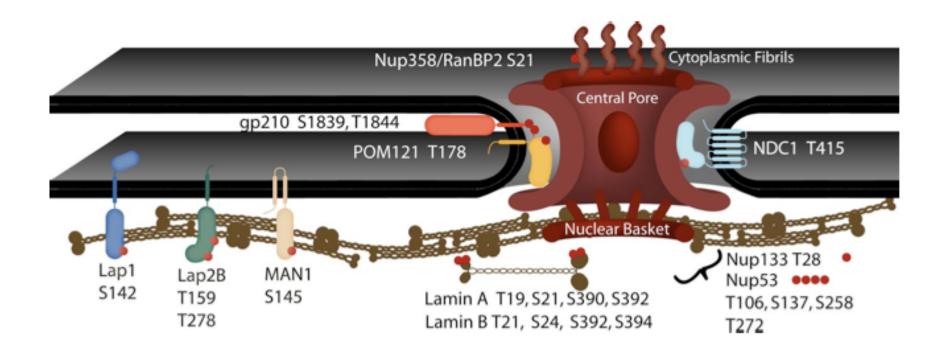
#### Thio-Phosphate Tagging of as-Kinase Direct Phosphorylation Products



# Thio-Phosphate Tagging of as-Kinase Direct Phosphorylation Products

■ Protocol next used to study phosphorylation sites on the nuclear pore complex (NPC) and nuclear lamina, both comprising the nuclear envelope (known target of Cdk activity)

Rediscovered known phosphorylation sites and discovered new ones

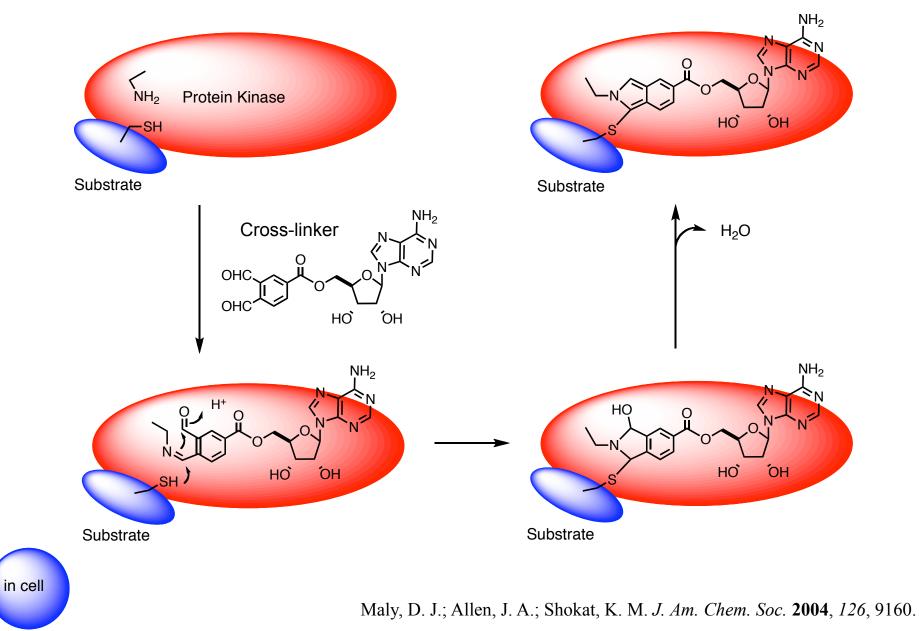




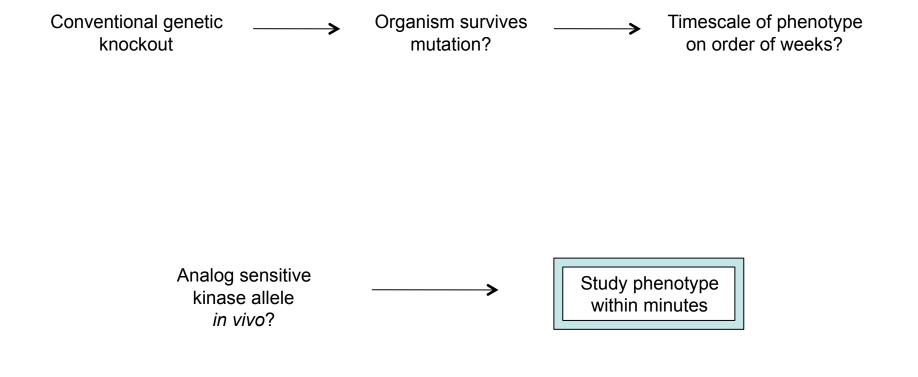
Shokat, K, M,; et al. Proc. Natl. Acad. Sci. USA 2008, 105, 1442.

## Variable Reversal: Hold Substrate Constant

■ Mutate substrate of interest (cysteine) and cross-link to discover kinase target

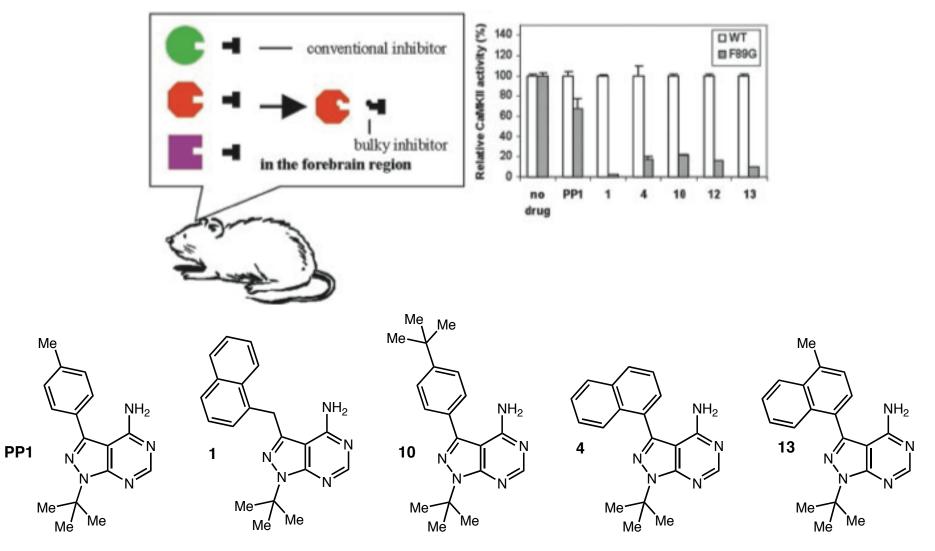


True Power of Chemical Genetics: Temporal Control



# True Power of Chemical Genetics: Temporal Control

- Can we direct the knockout at the protein level rather than at the DNA level?
- PP1 derivative found to be potent for of  $\alpha$ -Ca<sup>2+</sup>/calmodulin-dependent protein kinase II ( $\alpha$ CAMKII)

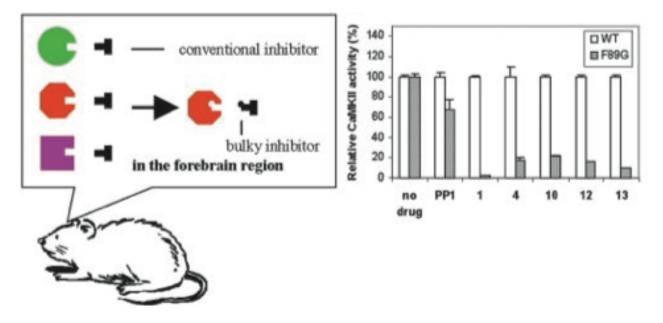


Shokat, K. M.; Tsien, J. Z.; et al. Proc. Natl. Acad. Sci. USA 2003, 100, 4287.

in vitro

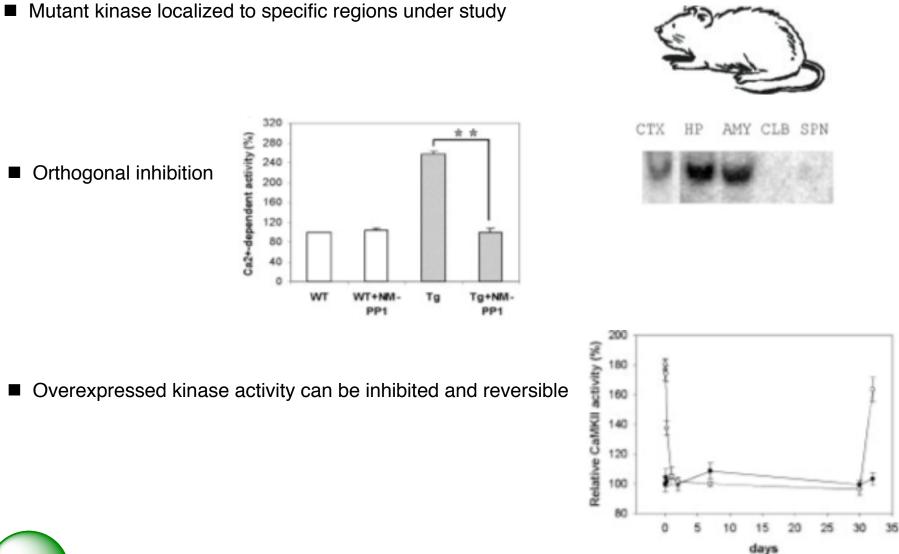
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- PP1 derivative found to be potent for of  $\alpha$ -Ca<sup>2+</sup>/calmodulin-dependent protein kinase II ( $\alpha$ CAMKII)



- αCAMKII studied in relation to contextual (hippocampal) and fear-based (amygdala-based) conditioning
- Produced mice overexpressing αCAMKII-F89G in forebrain, hippocampus and amygdala (mRNA levels)
- Tritium incorporated NM-PP1 (1) enters forbrain in 3 5 min, peaks at 20 min, bases out at 45 min.

Pre-Behavioral Studies Showing Orthogonality

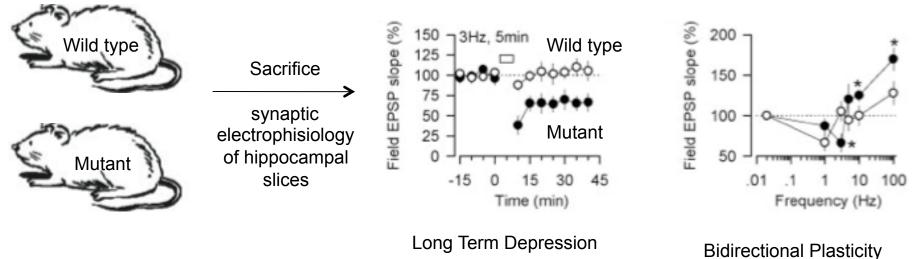


in vivo

Shokat, K. M.; Tsien, J. Z.; et al. Proc. Natl. Acad. Sci. USA 2003, 100, 4287.

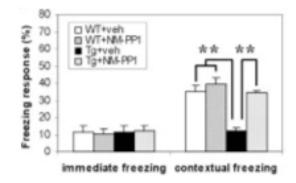
# Pre-Behavioral Studies Showing Electrophisiological Significance

Significant long term depression and bidirectional shifting for kinase mutant mice was encouraging



Response Curve

NM-PP1 (1) inhibitor rescues fear-based freezing response





Shokat, K. M.; Tsien, J. Z.; et al. Proc. Natl. Acad. Sci. USA 2003, 100, 4287.

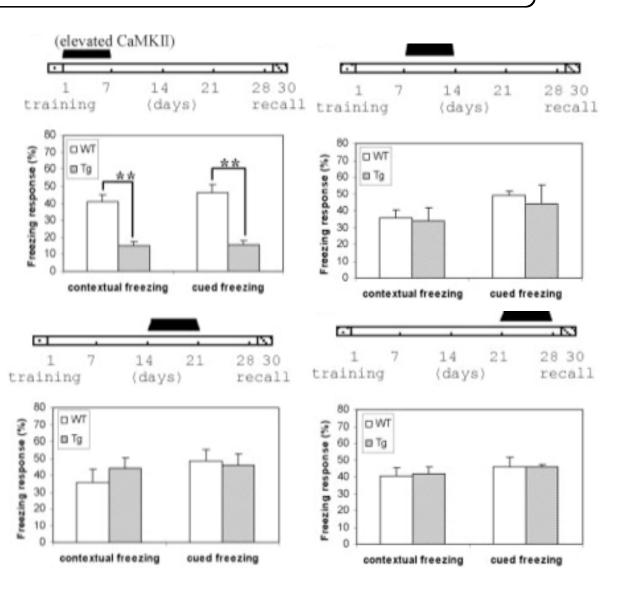
# Behavioral Studies: Temporal Response Crucial for Research and Discovery

■ Fear-based learning for this mechanism is limited to the first week of post-trauma

CaMKII involved in learning by Calcium channel activity

Each of 4 mice allowed elevated CaMKII levels in a different week (rest supressed by NM-PP1 inhibitor

Because elevated CaMKII levels only dampen fear response in first week, contenxtual and cued response limited to that week



Shokat, K. M.; Tsien, J. Z.; et al. Proc. Natl. Acad. Sci. USA 2003, 100, 4287.



#### References

Druker B. J.; *et al. Nature Medicine* 1996, *2*, 561. Han, S.; *et al. J. Bio. Chem.* 2009, *284*, 13193.
Hunter T. *Cell* 1987, *50*, 823.
Schultz, P. G.; *et al. Science* 1998, *281*, 533.
Hanke, J. H. *et al. J. Biol. Chem.* 1996, *271*, 695. Shokat, K. M. *et. al. Nature* 2000, *407*, 395.
Schindler T., *et al Mol.Cell* 1999, *3*, 639.
Shokat, K. M. *et. al. Nature* 2000, *407*, 395.

Stern, B.; Nurse, P. *Trends Genet.* 1996, *12*, 345;
Bishop, D. P.; *et al. Nature* 2000, *407*, 395.
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Patil, C., Walter, P. *Curr. Opin. Cell Biol.* 2001, *13*, 349;
Papa, F. R., Zhang, C., Shokat, K. M.; Walter, P. *Science* 2003, *302*, 1533.
Crackower, M. A. *et al. Cell* 2002, *110*, 737; Patrucco, E. *et al. Cell* 2004, *118*, 375.
Bi, L.; *et al. Mamm. Genome* 2002, *13*, 169.

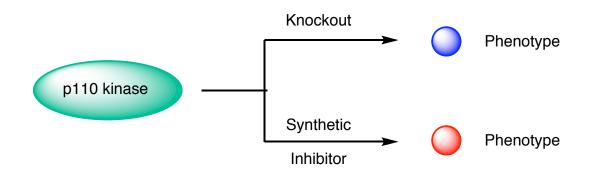
Knight, Z. A.; Shokat, K. M. Cell 2007, 123, 425.

Shokat, K, M,; *et al. Proc. Natl. Acad. Sci. USA* **1997**, *94*, 3565. Shah, K.; Shokat, K. M. *Chem. Biol.* **2002**, *9*, 35. Shokat, K, M,; *et al. Proc. Natl. Acad. Sci. USA* **2008**, *105*, 1442. Maly, D. J.; Allen, J. A.; Shokat, K. M. *J. Am. Chem. Soc.* **2004**, *126*, 9160.

Shokat, K. M.; Tsien, J. Z.; et al. Proc. Natl. Acad. Sci. USA 2003, 100, 4287.

#### Kinase Inhibition that Leaves Protein Function Intact

■ Small molecule inhibitors and knockouts of the same protein produce different phenotypes



Crackower, M. A. et al. Cell 2002, 110, 737; Patrucco, E. et al. Cell 2004, 118, 375.

#### Kinase Inhibition that Leaves Protein Function Intact

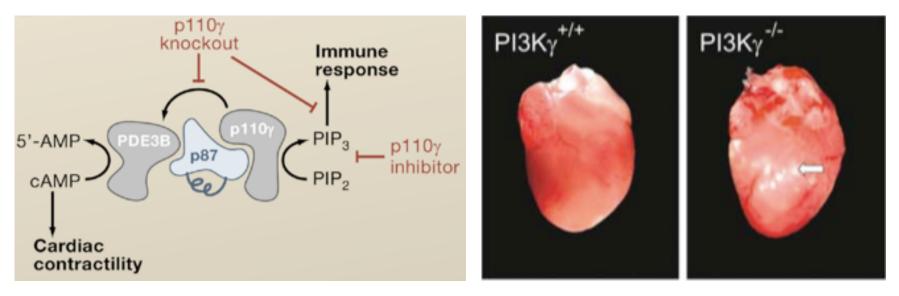
■ Small molecule inhibitors and knockouts of the same protein produce different phenotypes

p110γ mediates leukocyte reaction to the inflammatory response; mice lacking it have more dampened immune response, leading to the search for anti-inflammatory drugs



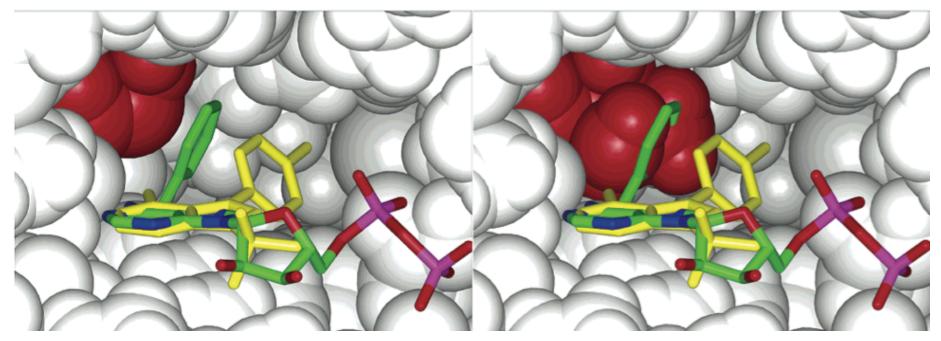
## Kinase Inhibition that Leaves Protein Function Intact

- Small molecule inhibitors and knockouts of the same protein produce different phenotypes
- p110γ mediates leukocyte reaction to the inflammatory response; mice lacking it have more dampened immune response, leading to the search for anti-inflammatory drugs
- While the p110γ knockout mice had elevated myocardial damage from aortic constriction, the mice with overexpressed kinase-dead (mutant) p110γ protects them from this damage
- Why does p110γ deletion produce a different phenotype from kinase-dead overexpression?
- Knockin mice with kinase-dead levels at wild-type concentration retain immune deficit but with normal heart tissue --> cardiac effect not due to loss of kinase activity
- p110g allosterically binds to an enzyme that catalyzes cAMP destruction ([cAMP] proportional to pathological cardiac response); knockouts prevent this, inhibitors miss the point

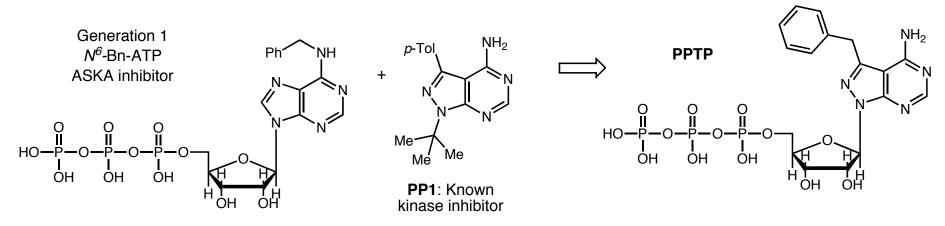


Crackower, M. A. et al. Cell 2002, 110, 737; Patrucco, E. et al. Cell 2004, 118, 375.

#### Structural Analysis Leads to a Rationally Designed ASKA Inhibitor



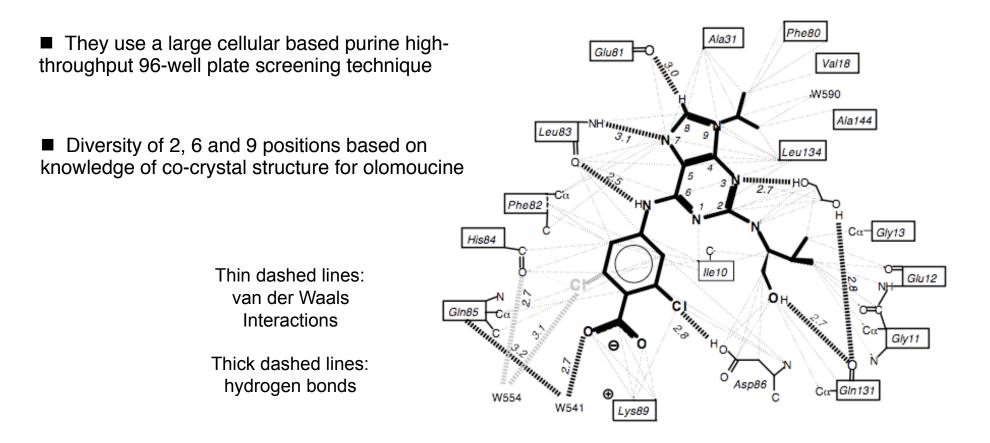
Shokat noticed naturally occuring hydrophobic pocket to be exploited, requiring a new inhibitor
 New inhibitor desing challenging: "requirements for substrate recognition/transition state stabilization different versus inhibitor binding at the same active site."



Kraybill, B. C.; Ilkin, L. L.; Blethrow, J. D.; Morgan, D. O.; Shokat, K. M. J. Am. Chem. Soc. 2002, 124, 12118.

#### An Apex of Traditional Pharmacology

Schultz and coworkers discover Purvalanol B, a potent and moderately selective inhibitor of cyclin dependent kinase (CDK2 in humans cdc28 in yiest)

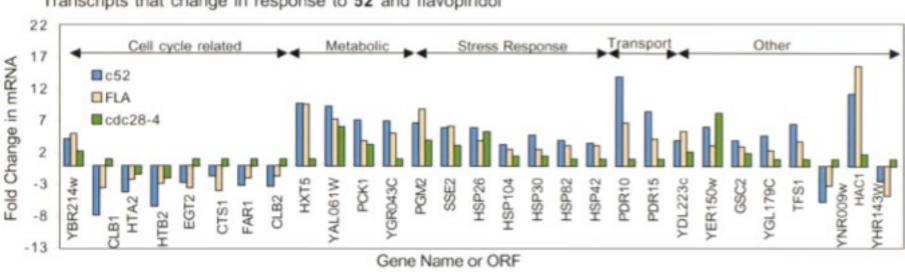


Schultz, P. G.; et al. Science 1998, 281, 533.

# Traditional Pharmacology Has Limits

■ For the yeast case (cdc28) Schultz could measure mRNA levels of nearly all yeast genes as determined by high-density oligonucleotide expression arrays

mRNA level changes are representative of the degree of up or down-regulation of that gene 



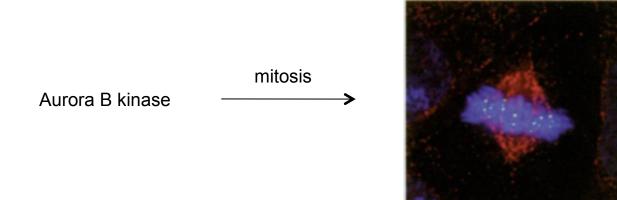
Transcripts that change in response to 52 and flavopiridol

While discovering a potent and moderately selective new inhibitor of CDK2, and a wealth of information regarding yeast genetic up/down regulation, there was one drawback:

> "Our current experimental design does not allow us to definitively Identify the primary target or targets of inhibition by [purvalanol]."

# The Scaffolding Function of Aurora B Kinase

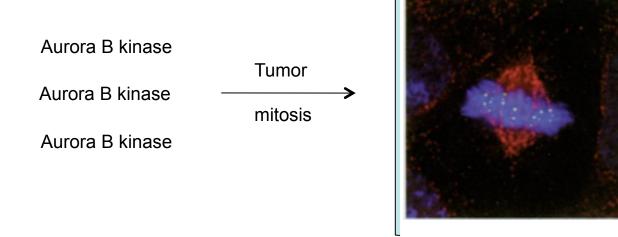
- Aurora kinases regulate spindle assembly and chromosomal alignment during mitosis
- Many tumors overexpress Aurora kinases which lead to interest in inhibitor development
- RNAi of Aurora B leads to major chromosomal alignment problems due to disruption of Aurora B interacting with Survivin at the centromere
- When RNAi treated Aurora B is then inhibited with ZM447439, Survivin is then localized correctly
- Therefore, the kinase is involved in a "scaffolding" effect not directly tied to tyrosine phosphorylation



Ditchfield, C.; et al. Cell Biol. 2003, 161, 267.

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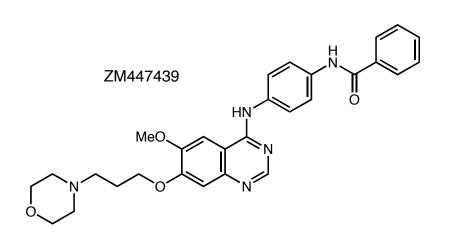


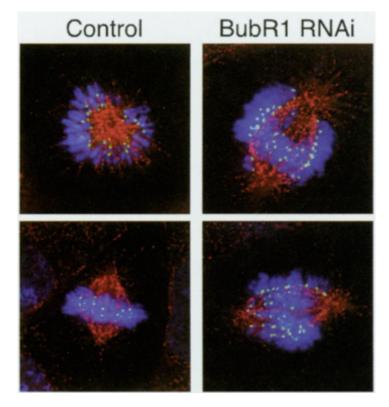
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Ditchfield, C.; et al. Cell Biol. 2003, 161, 267.

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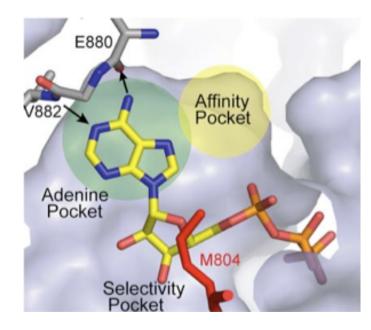


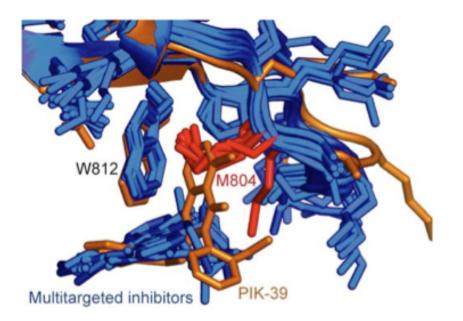


Ditchfield, C.; et al. Cell Biol. 2003, 161, 267.

#### Importance of Relative Enzyme Stoichiometry Masked by Knockout

- Both PI3 kinases, p110 $\alpha$  and p110 $\beta$ , carry signals from the growth factor, insulin
- Because knockout of either kinase isoform kills mice early in development we know that they cannot compensate for each other but don't know their roles
- Heterozygous deletion of either produces no phenotype; deletion of p85, the p110 binding partner, paradoxically increases insulin signaling
- Both knockin kinase-dead mice as well as mice treated with a p110 $\alpha$ -selective inhibitor show reduced insulin signaling
- Paradox: why do mice lacking p110 or p85 show normal or increased insulin signaling while kinasedead mutants (or their equivalent p110 $\alpha$ -inhibited mice) show a reduction?
- Resulting Model: since p85 can function as a negative regulator of p110, insulin transduction controlled by relative stoichiometry of p110 to p85 rather than absolute amounts.



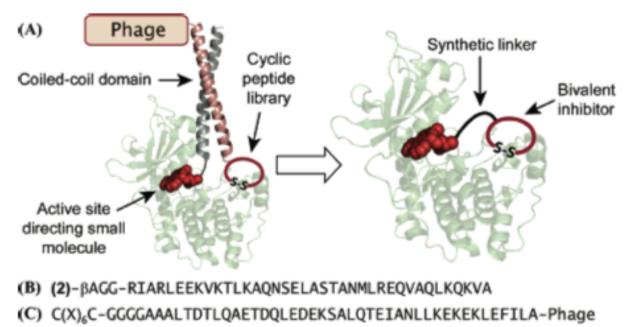


Shokat, K. M.; et al. Cell 2006, 125, 733.

# Tethering Small Molecules to a Phage-Display Library

■ If the phosphorylated protein target of a specific kinase is known, an abbreviated peptide analog of that protein can be synthetically appended to an ATP-competitive inhibitor.

- Ghosh used a phage display library approach to discover the right match.
- Phage display is a method for the study of protein-protein, protein peptide and DNA interactions that uses bacteriophages to connect proteins with the genetic information that encodes them



(D) cyclo(CTFRVFGC)G [3]

#### Bivalency and Synergystic Affinity and Selectivity

■ By seeking a bivalent inhibitor, a synergistic binding with warhead and cyclic peptide was employed to acheive selective inhibition of cAMP-dependent protein Kinase A (PKA).

