

# Chemical approaches to drugging “undruggable” proteins



Diego A. Granados

May 5<sup>th</sup>, 2023

Knowles Group Literature Meeting

Princeton University

# Outline

- Introduction to undruggable proteins
  - What makes a protein “undruggable”?
  - Attempts to drug K-Ras mutations
- Activity-based approaches to finding “druggable” sites
- Success stories in covalent drugs
  - Ibrutinib and Bruton’s tyrosine kinase
  - Sotorasib and K-Ras G12C
- Conclusions
  - “Yet to be drugged” instead of “undruggable”

# Outline

- Introduction to undruggable proteins
  - What makes a protein “undruggable”?
  - Attempts to drug K-Ras mutations
- Activity-based approaches to finding “druggable” sites
- Success stories in covalent drugs
  - Ibrutinib and Bruton’s tyrosine kinase
  - Sotorasib and K-Ras G12C
- Conclusions
  - “Yet to be drugged” instead of “undruggable”

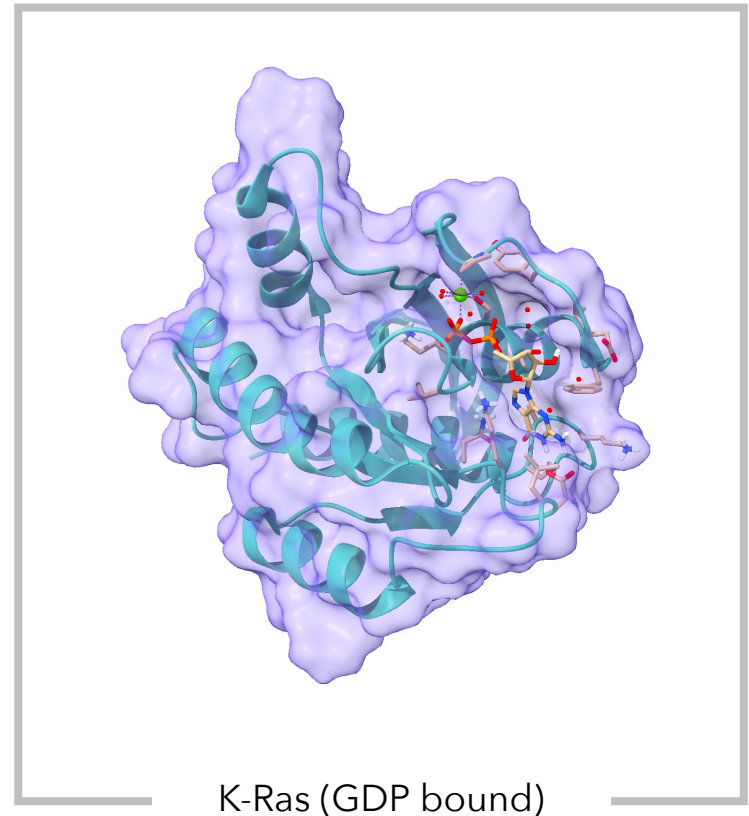
## What makes something “undruggable”?

A protein is considered **undruggable** when traditional pharmacological strategies have failed

# What makes something “undruggable”?

A protein is considered **undruggable** when traditional pharmacological strategies have failed

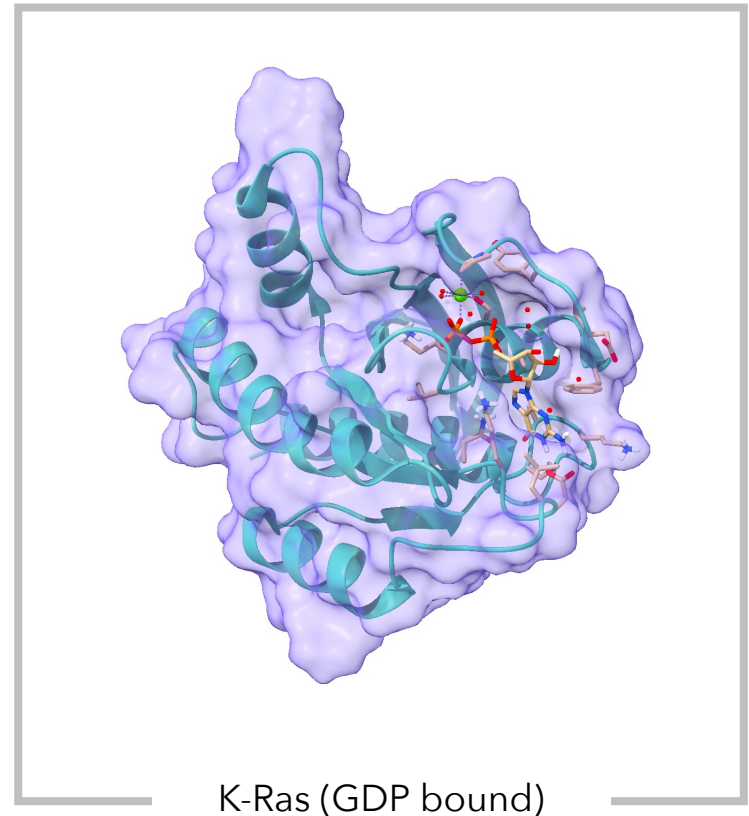
- **Kirsten rat sarcoma oncogene homologue (K-Ras)** mutations are implicated in ~25% of tumors
  - First discovered oncogene (1967)
  - GTP kinase (GTPase) involved in MAPK signal transduction (cell proliferation)



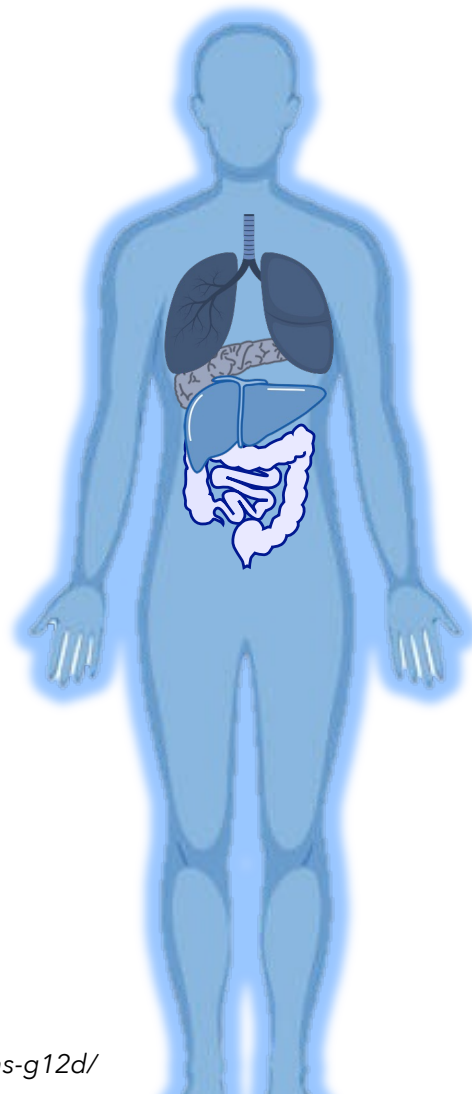
# What makes something “undruggable”?

A protein is considered **undruggable** when traditional pharmacological strategies have failed

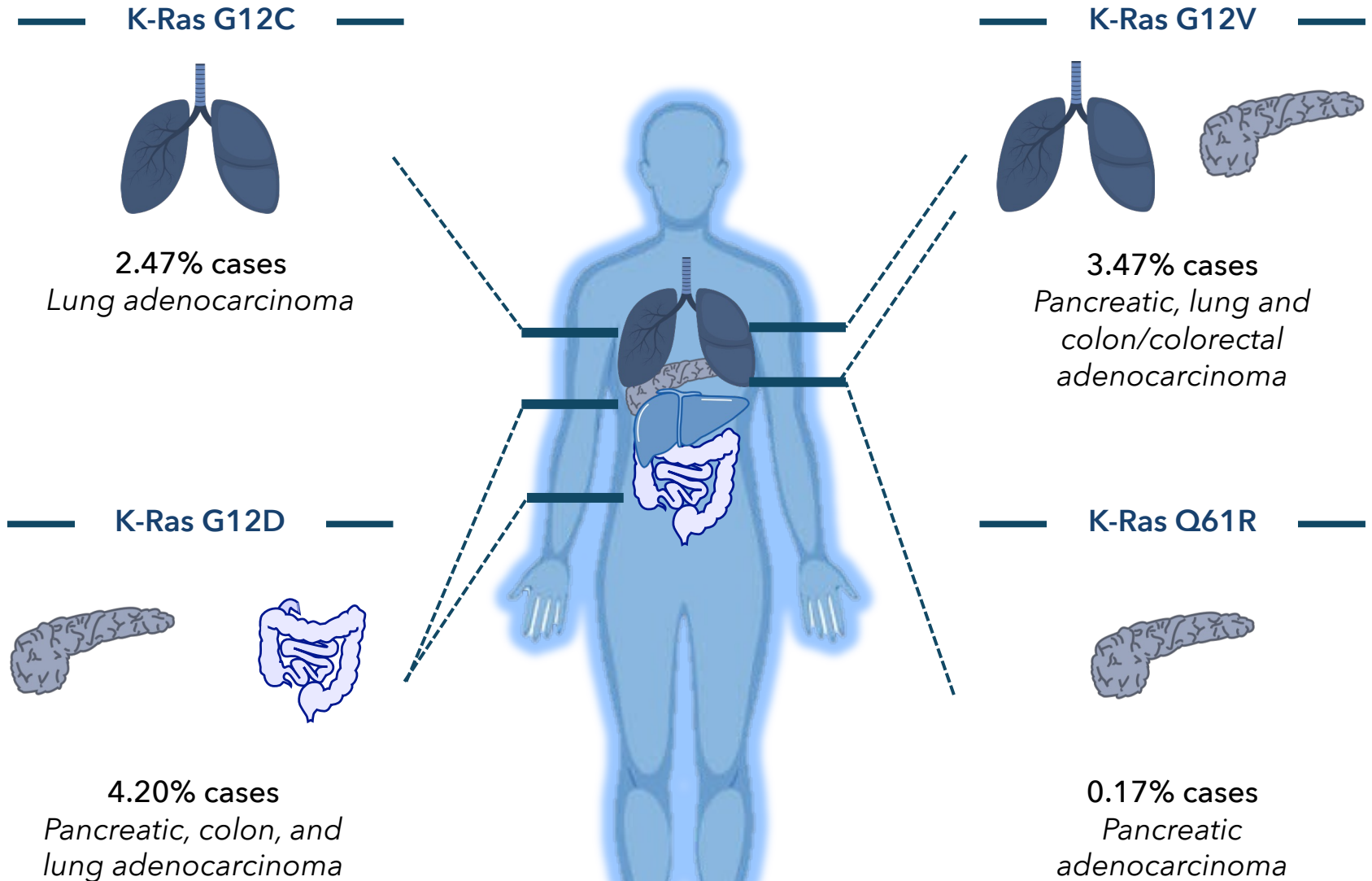
- **Kirsten rat sarcoma oncogene homologue (K-Ras)** mutations are implicated in ~25% of tumors
  - First discovered oncogene (1967)
  - GTP kinase (GTPase) involved in MAPK signal transduction (cell proliferation)
- Most common mutations are of the **Gly12** residue
  - G12D (33%)
  - G12V (20%)
  - G12C (10%)



K-Ras mutations are implicated in a variety of cancers



# K-Ras mutations are implicated in a variety of cancers

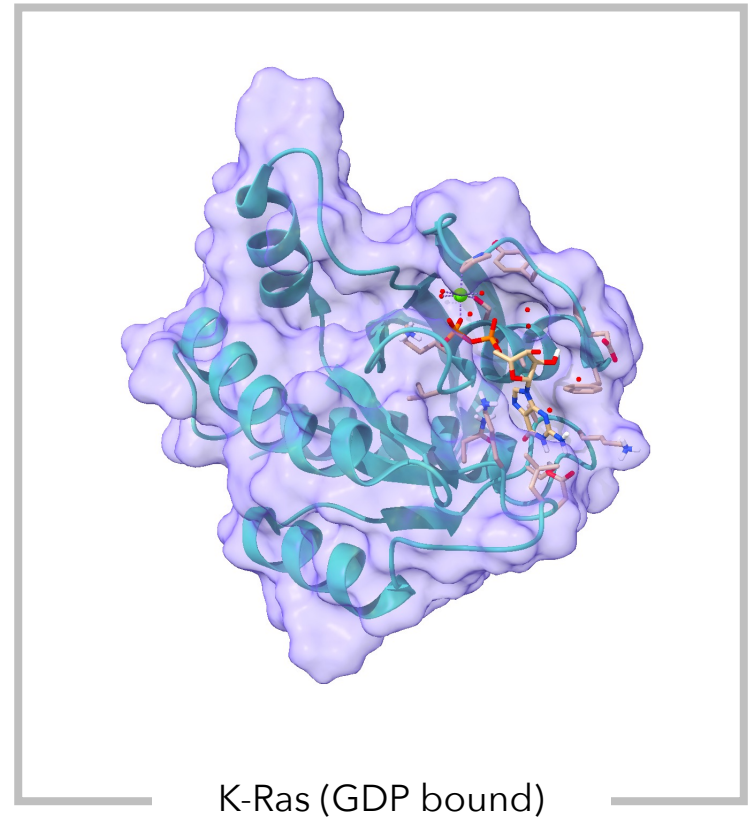




# What makes something “undruggable”?

A protein is considered **undruggable** when traditional pharmacological strategies have failed

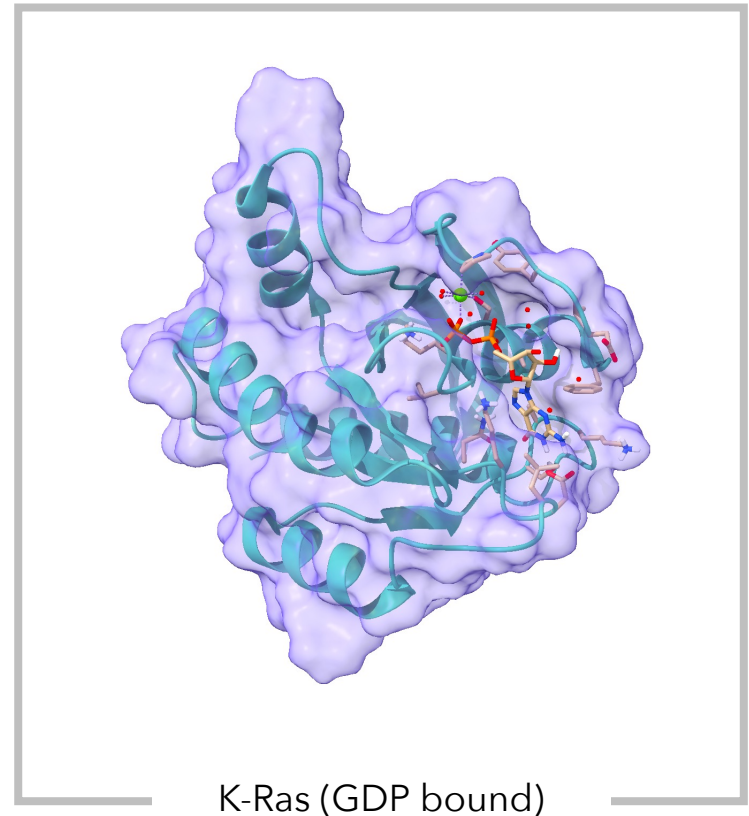
- **Kirsten rat sarcoma oncogene homologue (K-Ras)** mutations are implicated in ~25% of tumors
  - First discovered oncogene (1967)
  - GTP kinase (GTPase) involved in MAPK signal transduction (cell proliferation)
- Most common mutations are of the **Gly12** residue
  - G12D (33%)
  - G12V (20%)
  - G12C (10%)
- **Decades** of effort into drugging K-Ras



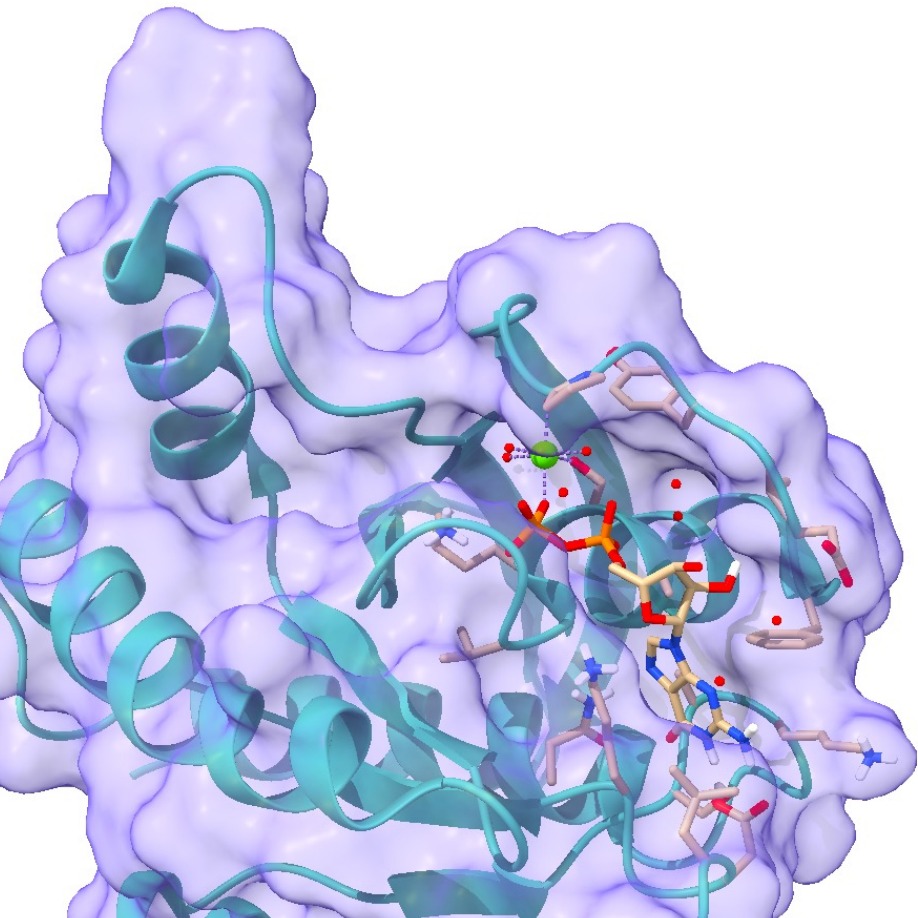
# What makes something “undruggable”?

A protein is considered **undruggable** when traditional pharmacological strategies have failed

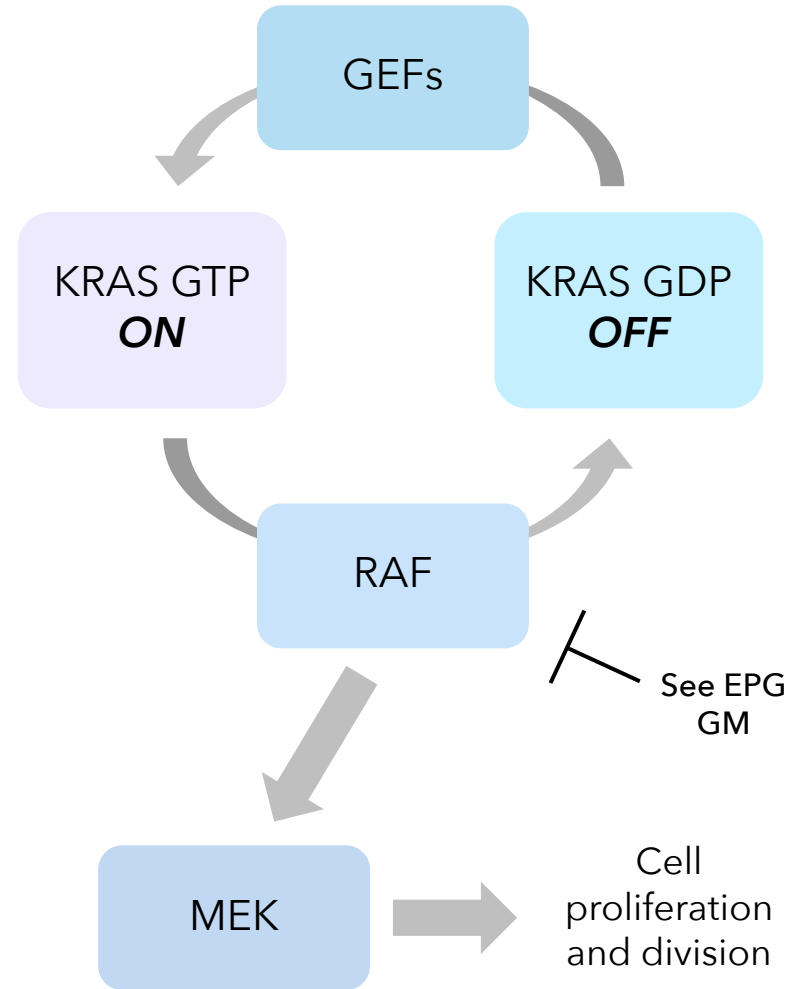
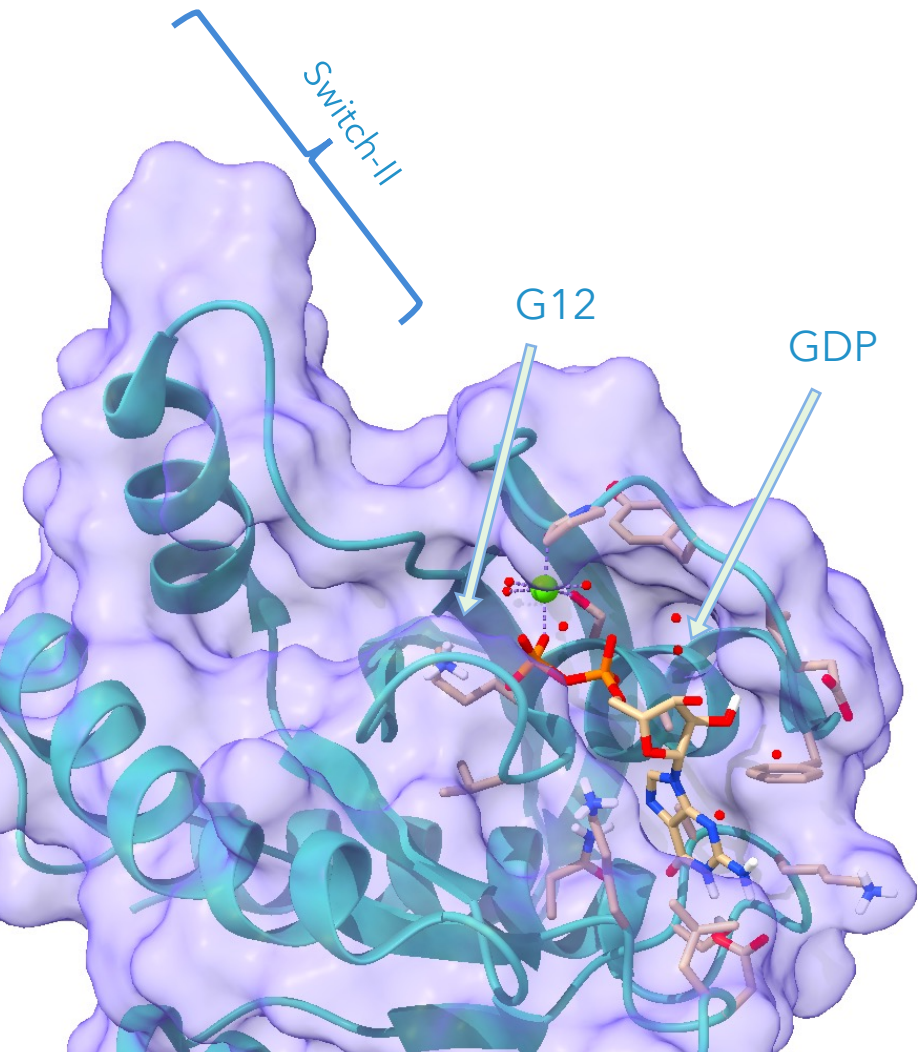
- **Kirsten rat sarcoma oncogene homologue (K-Ras)** mutations are implicated in ~25% of tumors
  - First discovered oncogene (1967)
  - GTP kinase (GTPase) involved in MAPK signal transduction (cell proliferation)
- Most common mutations are of the **Gly12** residue
  - G12D (33%)
  - G12V (20%)
  - G12C (10%)
- **Decades** of effort into drugging K-Ras
- Labeled as “undruggable”
  - **Why?**



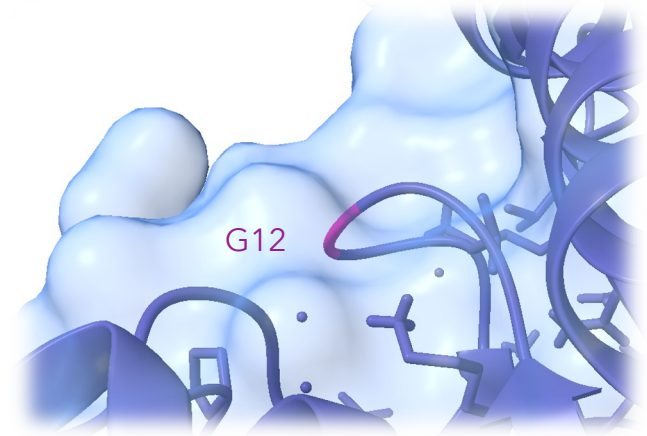
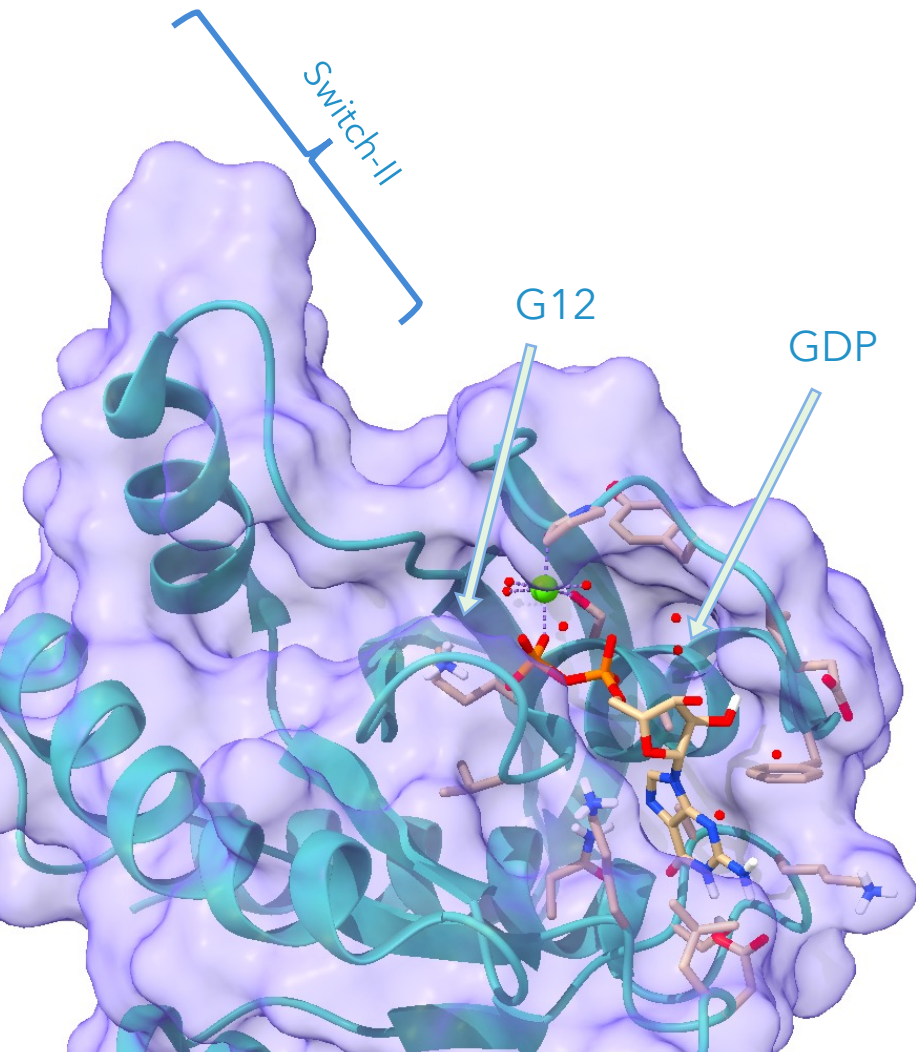
K-Ras mutations tend to affect phosphorylation state



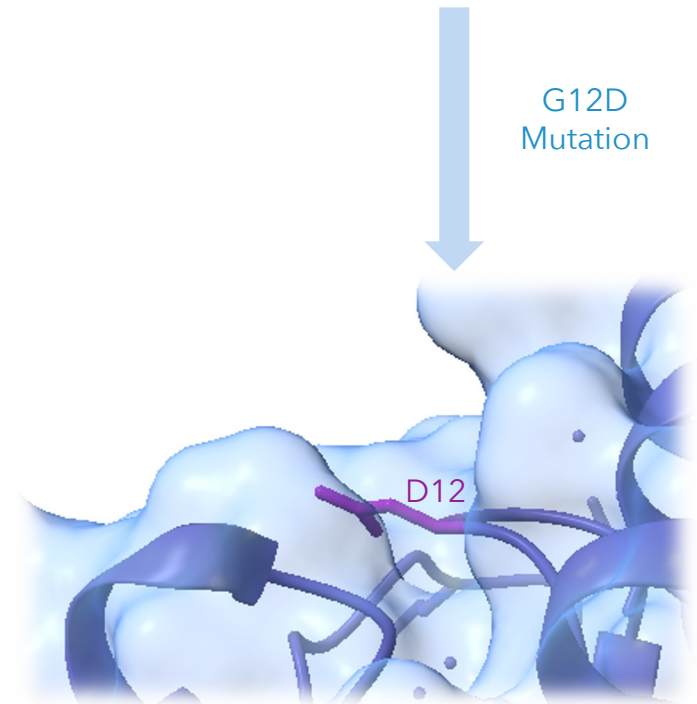
# K-Ras mutations tend to affect phosphorylation state



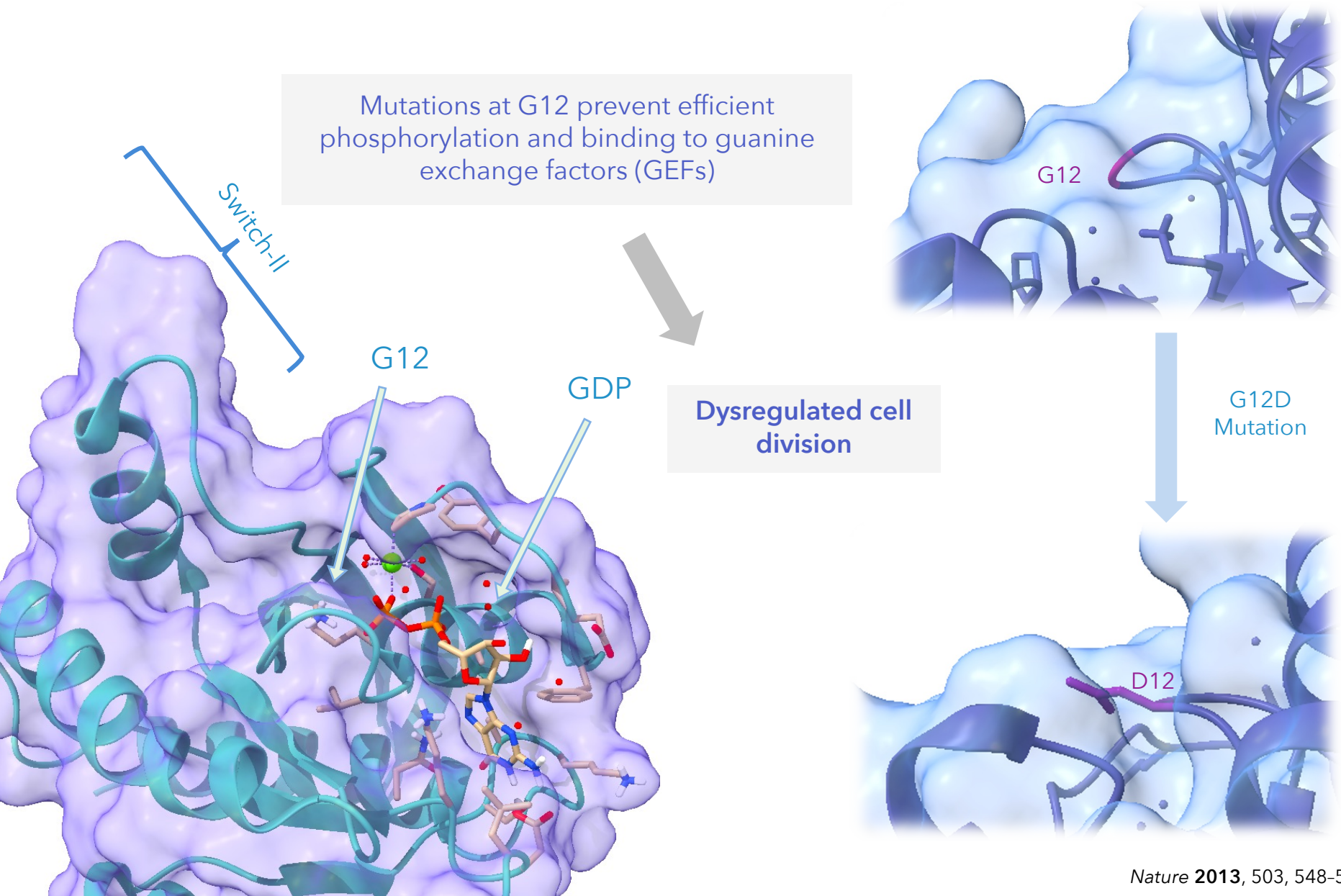
# K-Ras mutations tend to affect phosphorylation state



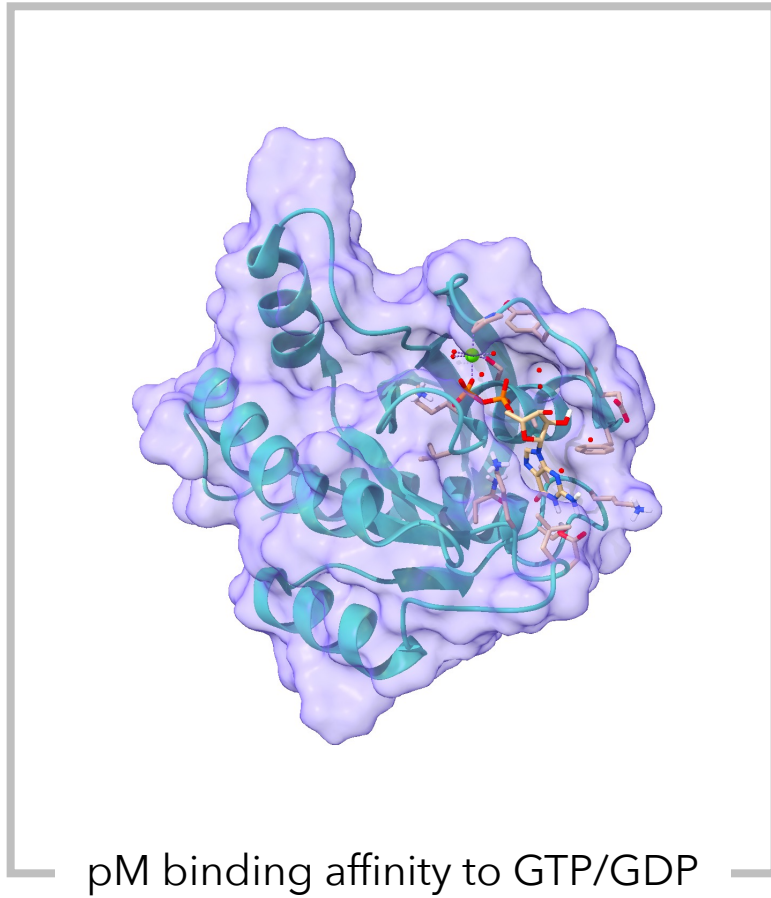
G12D  
Mutation



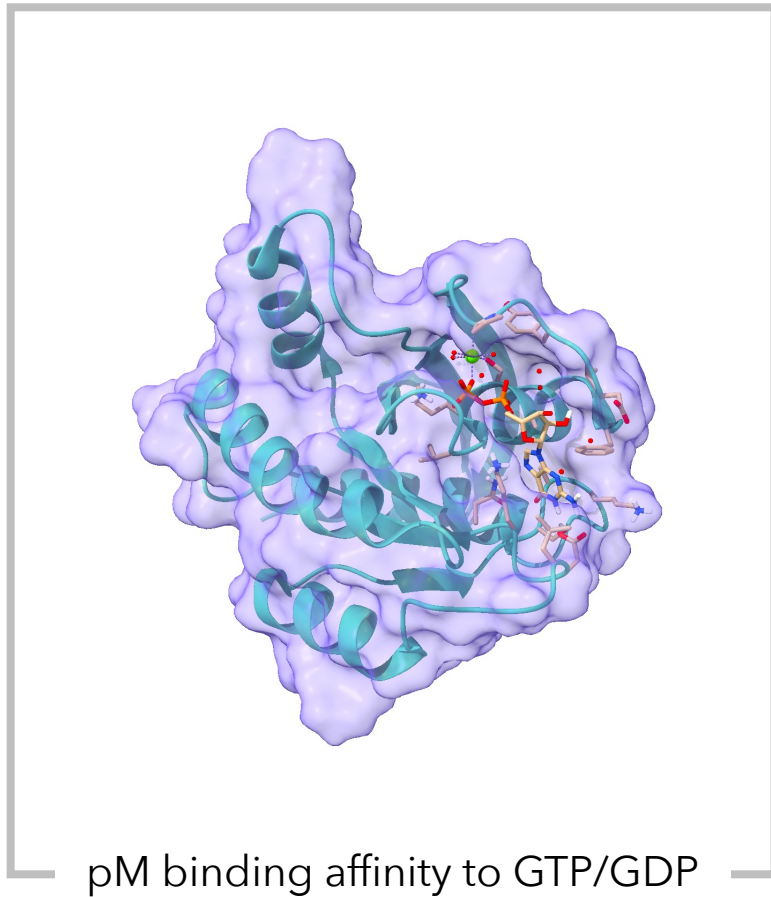
# K-Ras mutations tend to affect phosphorylation state



# Challenges in drugging K-Ras



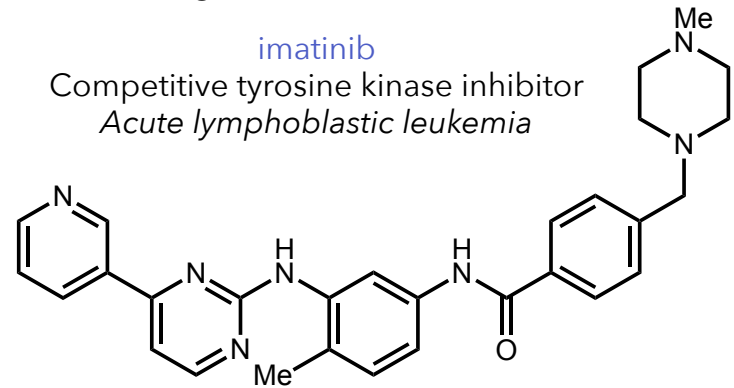
# Challenges in drugging K-Ras



- Previous strategies to drug ATPases involving ATP-mimics

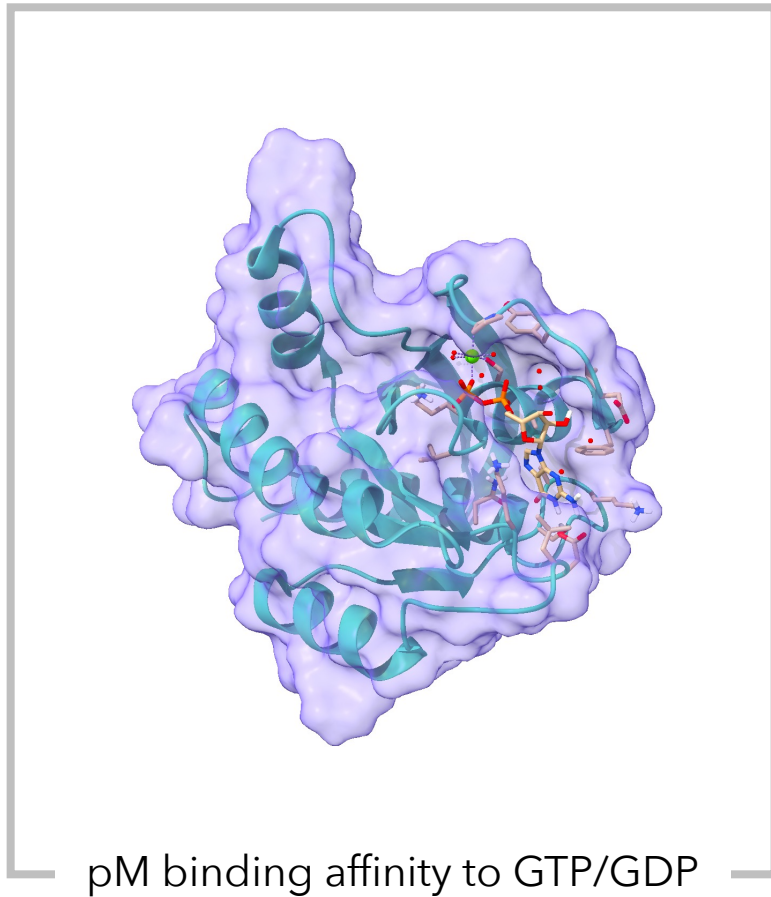
imatinib

Competitive tyrosine kinase inhibitor  
*Acute lymphoblastic leukemia*

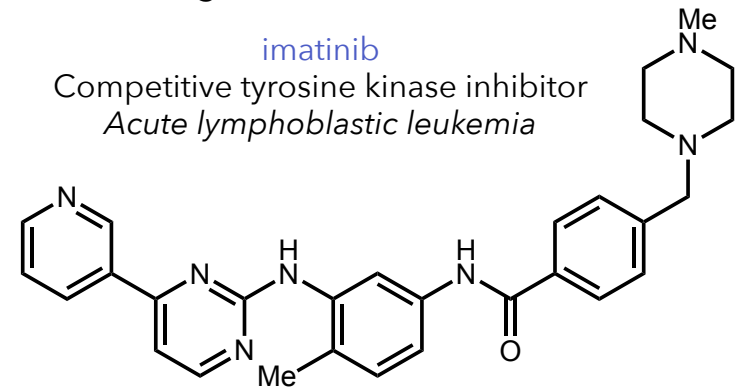




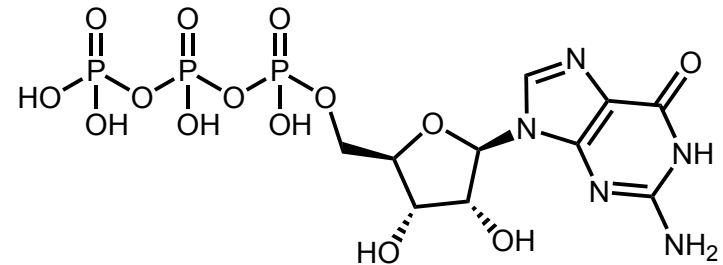
# Challenges in drugging K-Ras



- Previous strategies to drug ATPases involving ATP-mimics

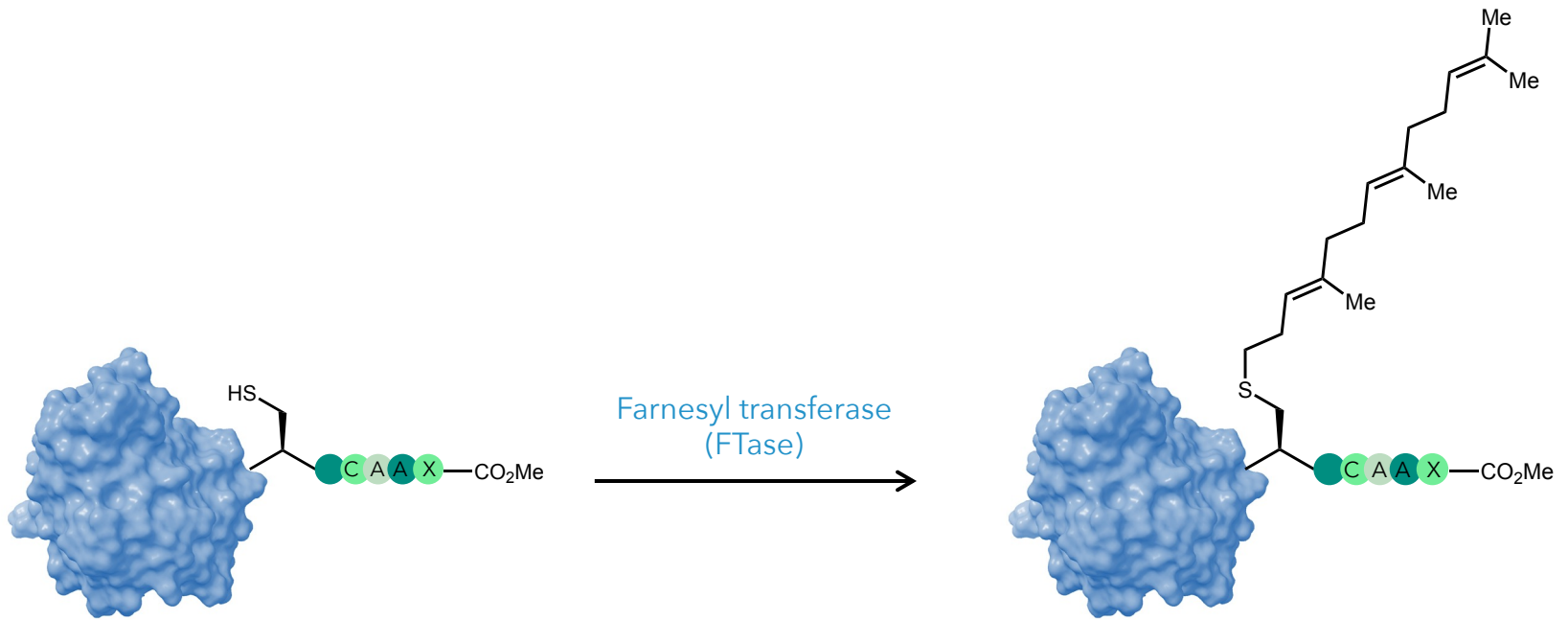


- Kinase competitive inhibitors are unsuccessful for K-Ras
- Unable to outcompete GTP binding



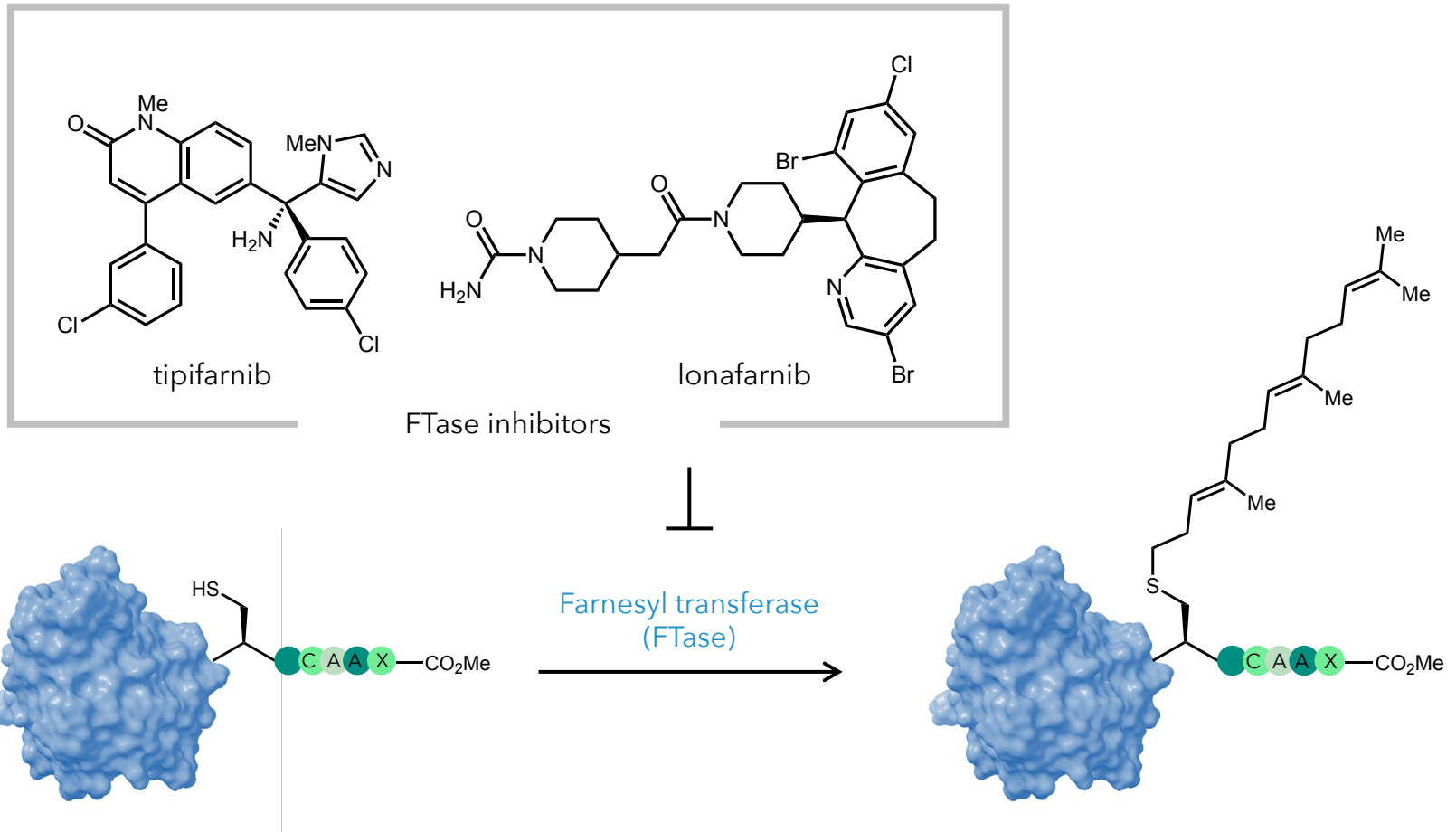
# Could inhibiting membrane localization drug K-Ras?

- Key finding: K-Ras is unable to function unless properly membrane localized



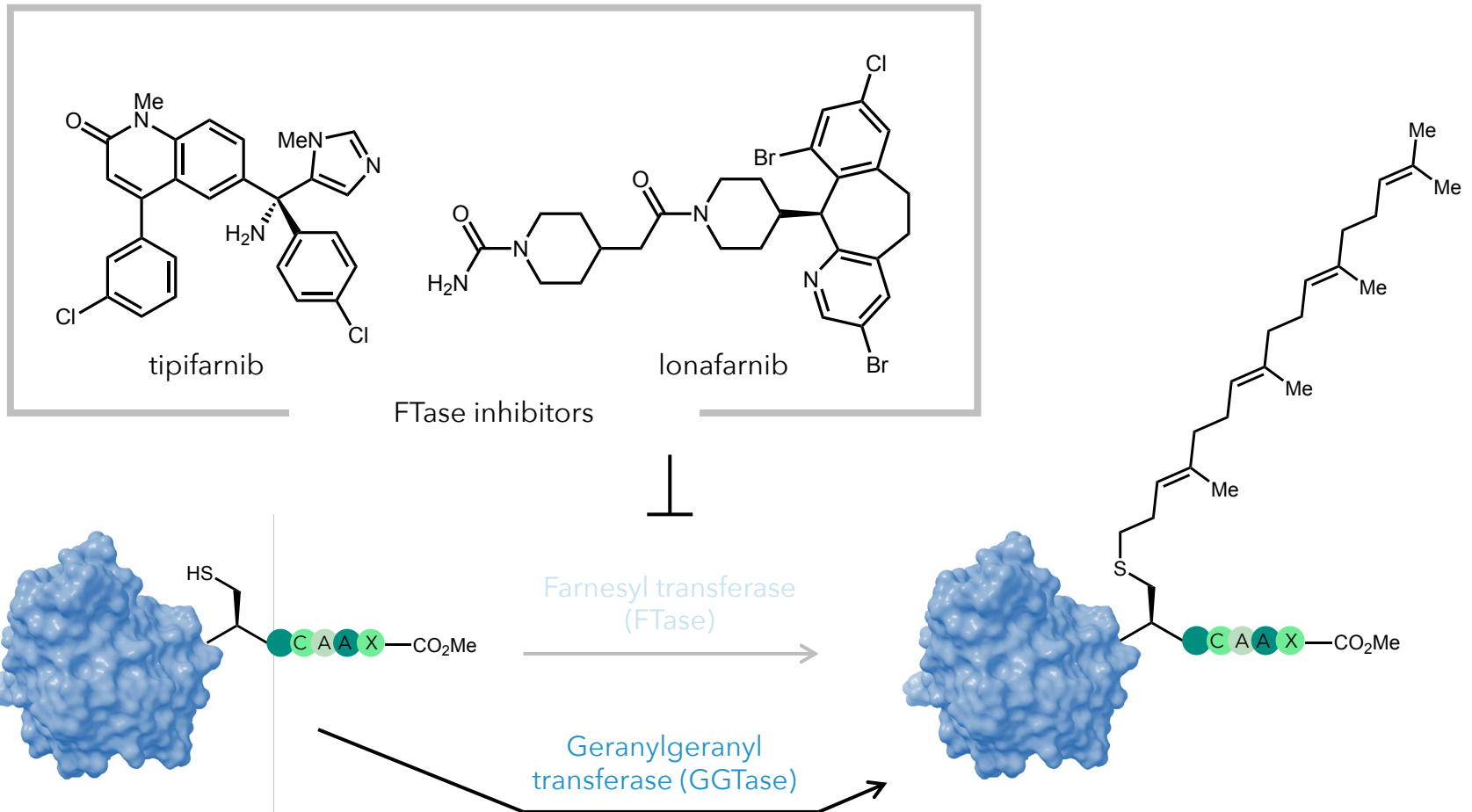
# Could inhibiting membrane localization drug K-Ras?

- Key finding: K-Ras is unable to function unless properly membrane localized



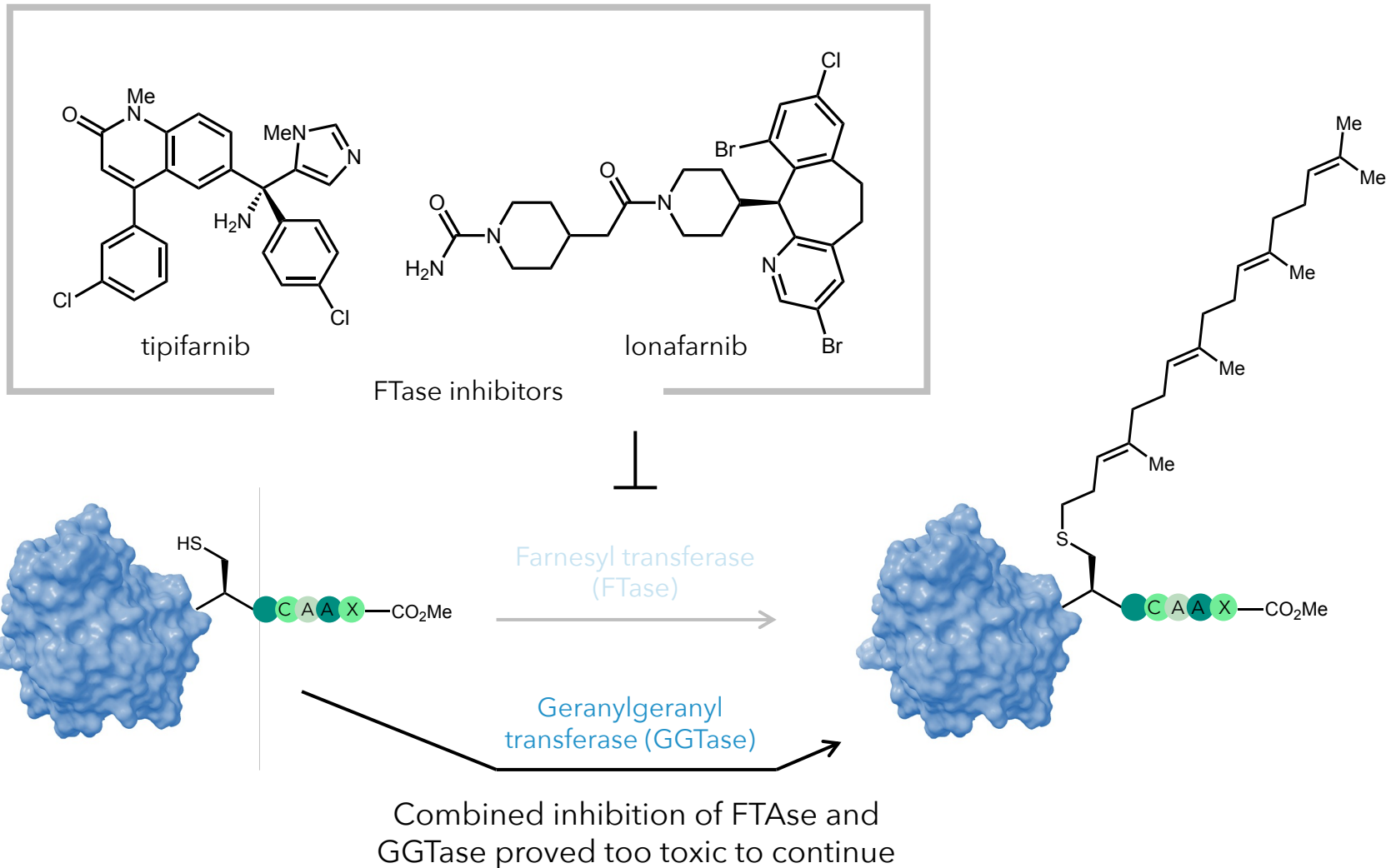
# Could inhibiting membrane localization drug K-Ras?

- Key finding: K-Ras is unable to function unless properly membrane localized

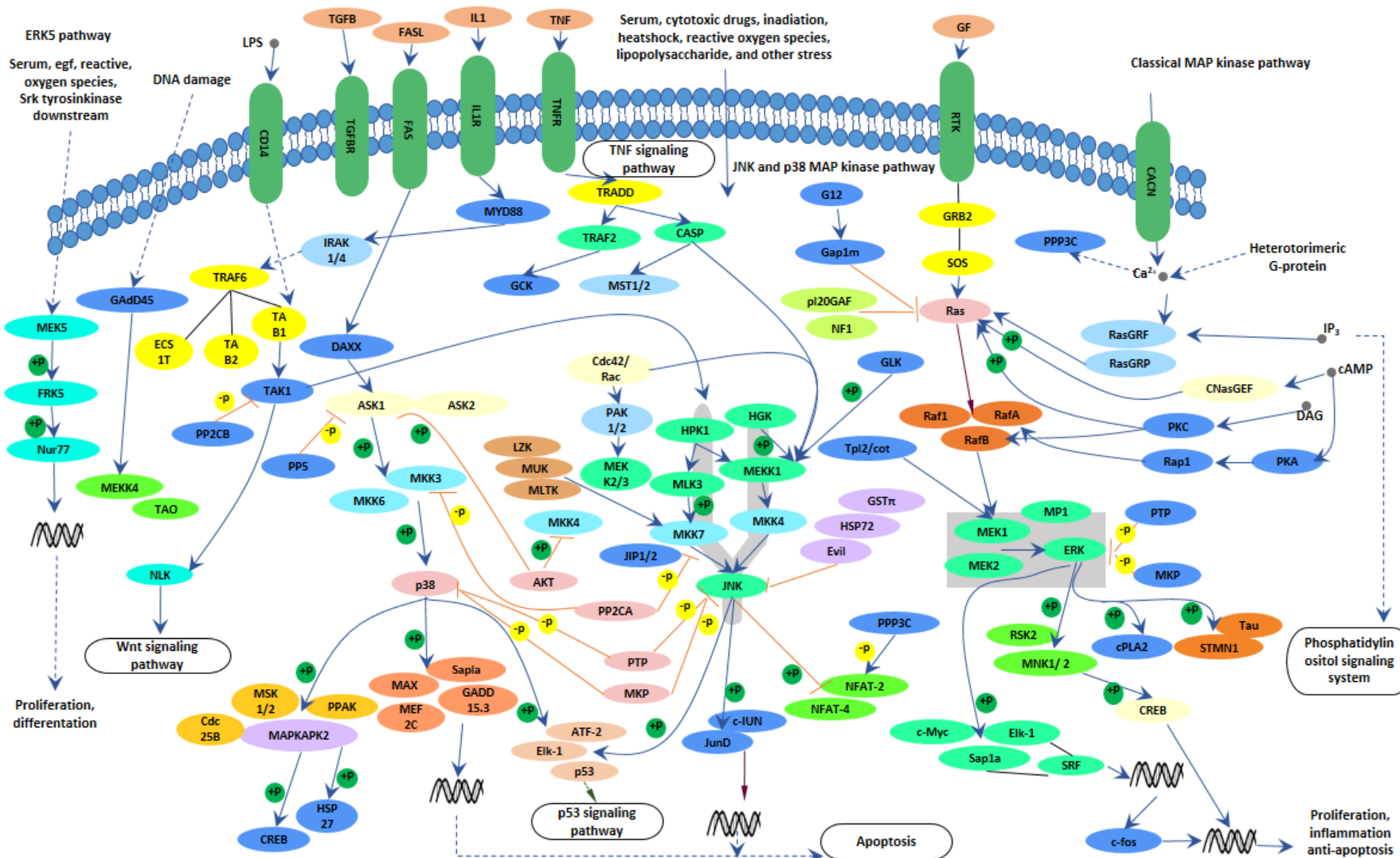


# Could inhibiting membrane localization drug K-Ras?

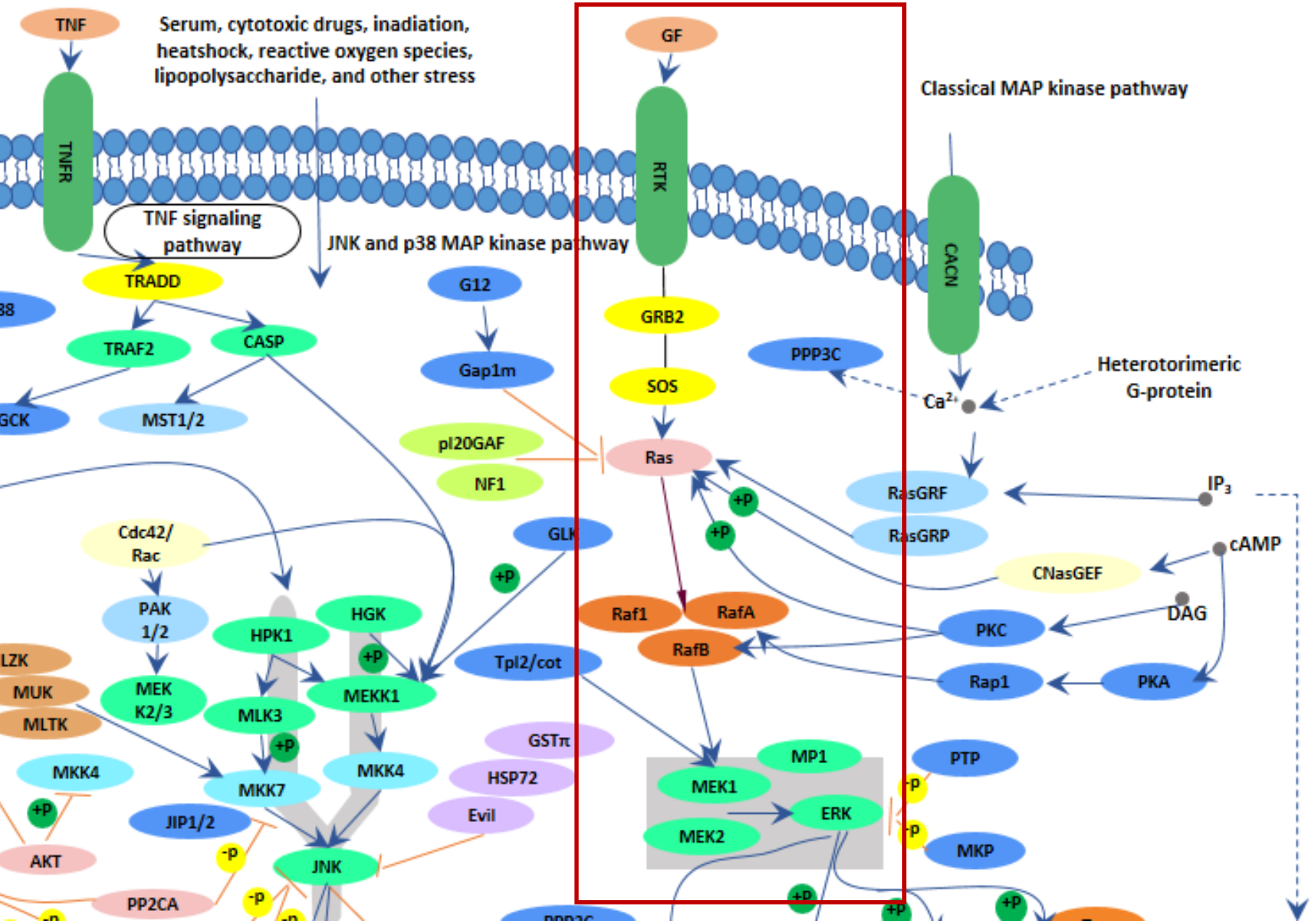
- Key finding: K-Ras is unable to function unless properly membrane localized



# The complexity of the MAPK signaling



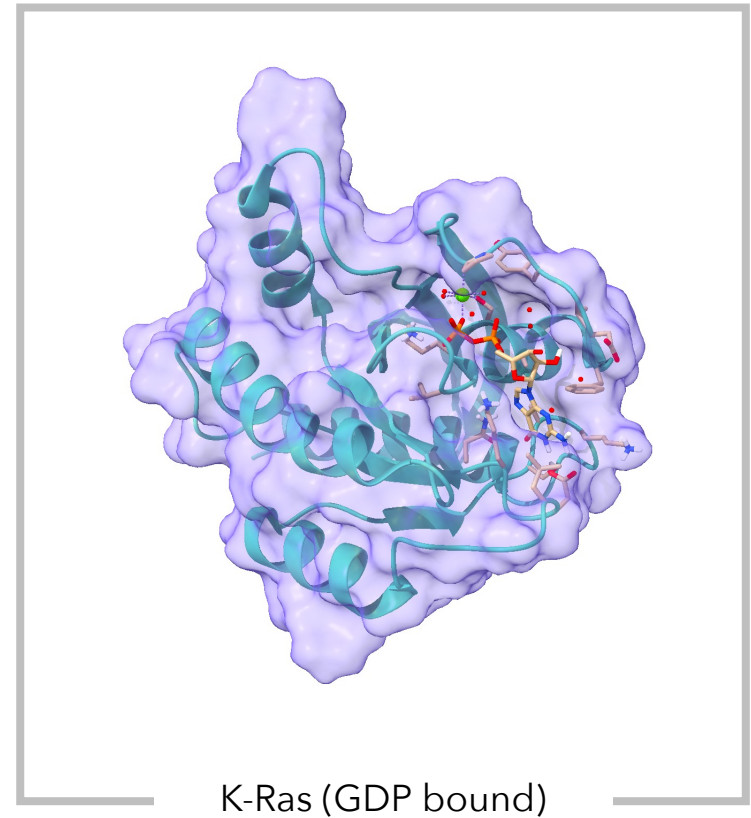
# The complexity of the MAPK signaling



# What makes something “undruggable”?

A protein is considered **undruggable** when traditional pharmacological strategies have failed

- Decades of effort into drugging K-Ras
- Quickly labeled as “undruggable”
  - Why?

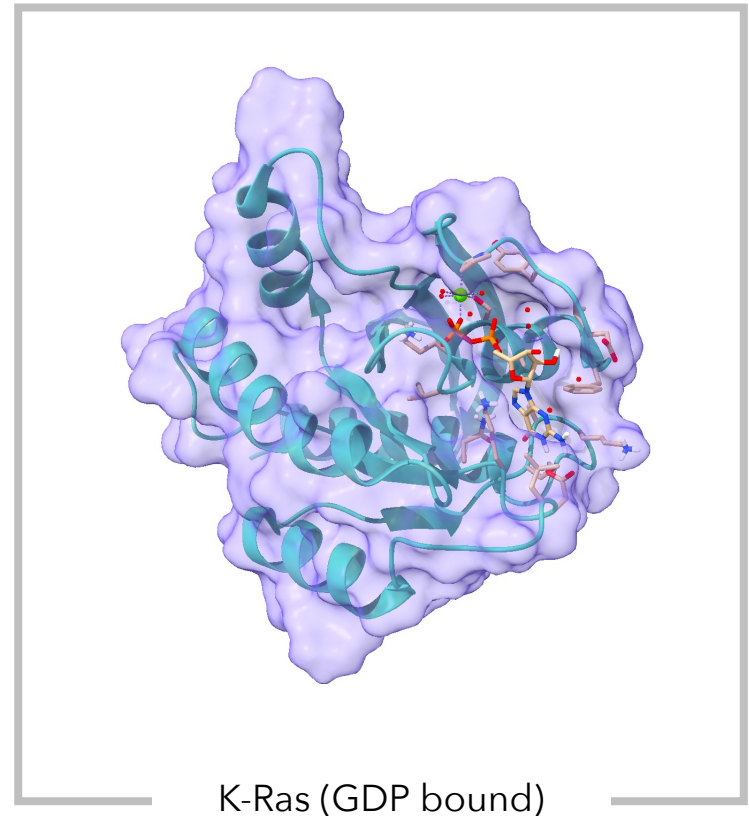




# What makes something “undruggable”?

A protein is considered **undruggable** when traditional pharmacological strategies have failed

- *Decades* of effort into drugging K-Ras
- Quickly labeled as “undruggable”
  - Why?
- High affinity for native GTP substrate
- Inhibition of membrane localization is ineffective
- K-Ras involved in highly complex signaling pathway - difficult to understand how knocking out one protein affects downstream effects!

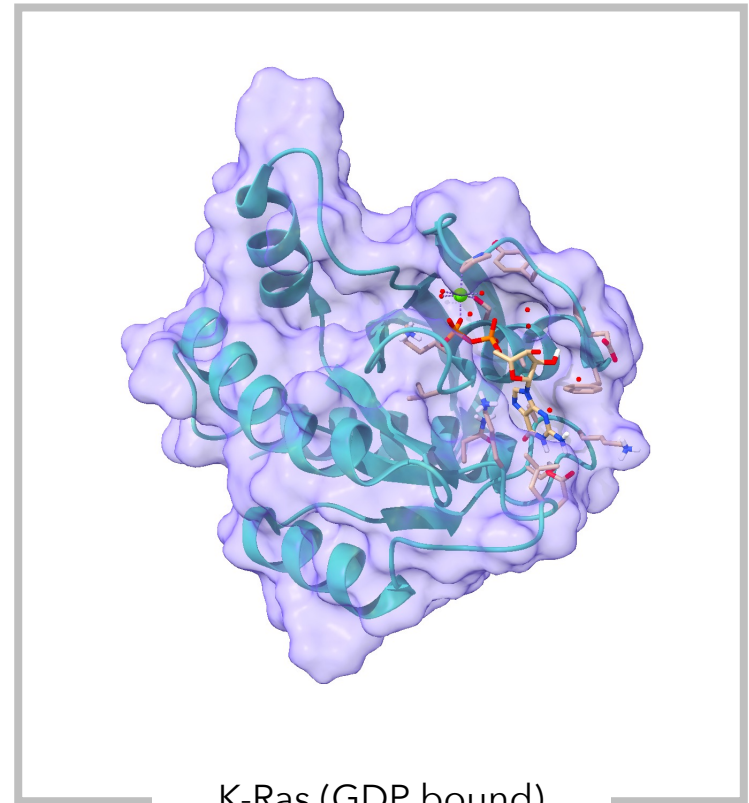


# What makes something “undruggable”?

A protein is considered **undruggable** when traditional pharmacological strategies have failed

- Decades of effort into drugging K-Ras
- Quickly labeled as “undruggable”
  - Why?
- High affinity for native GTP substrate
- Inhibition of membrane localization is ineffective
- K-Ras involved in highly complex signaling pathway - difficult to understand how knocking out one protein affects downstream effects!

New methods to find “druggable” sites on proteins are necessary



# Outline

- Introduction to undruggable proteins
  - What makes a protein “undruggable”?
  - Attempts to drug K-Ras mutations
- Activity-based approaches to finding “druggable” sites
- Success stories in covalent drugs
  - Ibrutinib and Bruton’s tyrosine kinase
  - Sotorasib and K-Ras G12C
- Conclusions
  - “Yet to be drugged” instead of “undruggable”

# The advent of activity-based protein profiling and covalent drugs



Prof. Benjamin Cravatt III  
Scripps Research Institute

- Known for **activity-based protein profiling (ABPP)**
- Allows for global analysis of **covalent** interactions with variety of proteins
- Proteomics (mass spectrometry) based approach focusing on *reactive* functionality rather than quantity

# The advent of activity-based protein profiling and covalent drugs



Prof. Benjamin Cravatt III  
Scripps Research Institute

- Known for **activity-based protein profiling (ABPP)**
- Allows for global analysis of **covalent** interactions with variety of proteins
- Proteomics (mass spectrometry) based approach focusing on *reactive* functionality rather than quantity

*To date, proteomics efforts have primarily been confined to recording variations in protein level rather than activity. The ability to profile classes of proteins on the basis of changes in their activity would greatly accelerate both the assignment of protein function and the identification of potential pharmaceutical targets.*

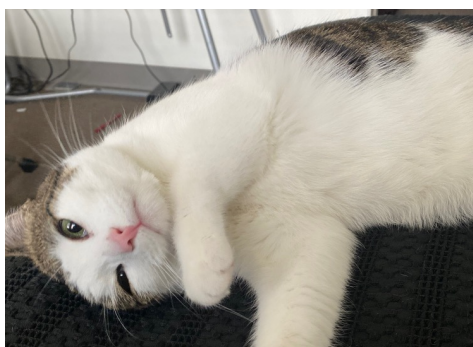
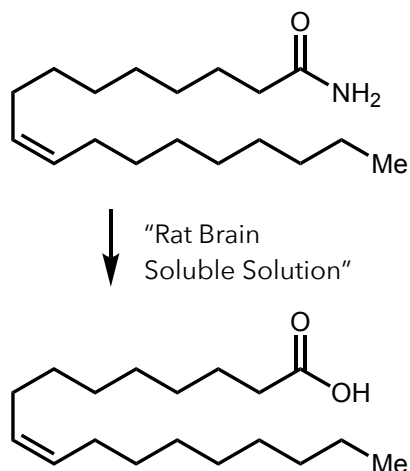
# The advent of activity-based protein profiling and covalent drugs



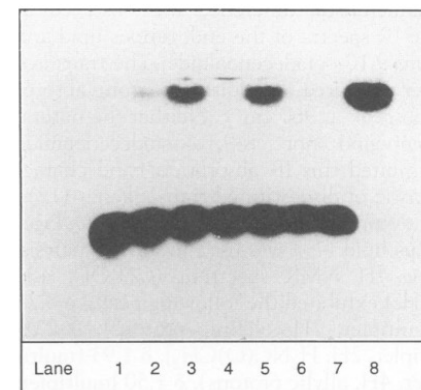
Prof. Benjamin Cravatt III  
Scripps Research Institute

- Known for **activity-based protein profiling (ABPP)**
- Allows for global analysis of **covalent** interactions with variety of proteins
- Proteomics (mass spectrometry) based approach focusing on *reactive* functionality rather than quantity

*To date, proteomics efforts have primarily been confined to recording variations in protein level rather than activity. The ability to profile classes of proteins on the basis of changes in their activity would greatly accelerate both the assignment of protein function and the identification of potential pharmaceutical targets.*



*Isolated from cerebrospinal fluids of sleep deprived cats  
(like Ophelia)*



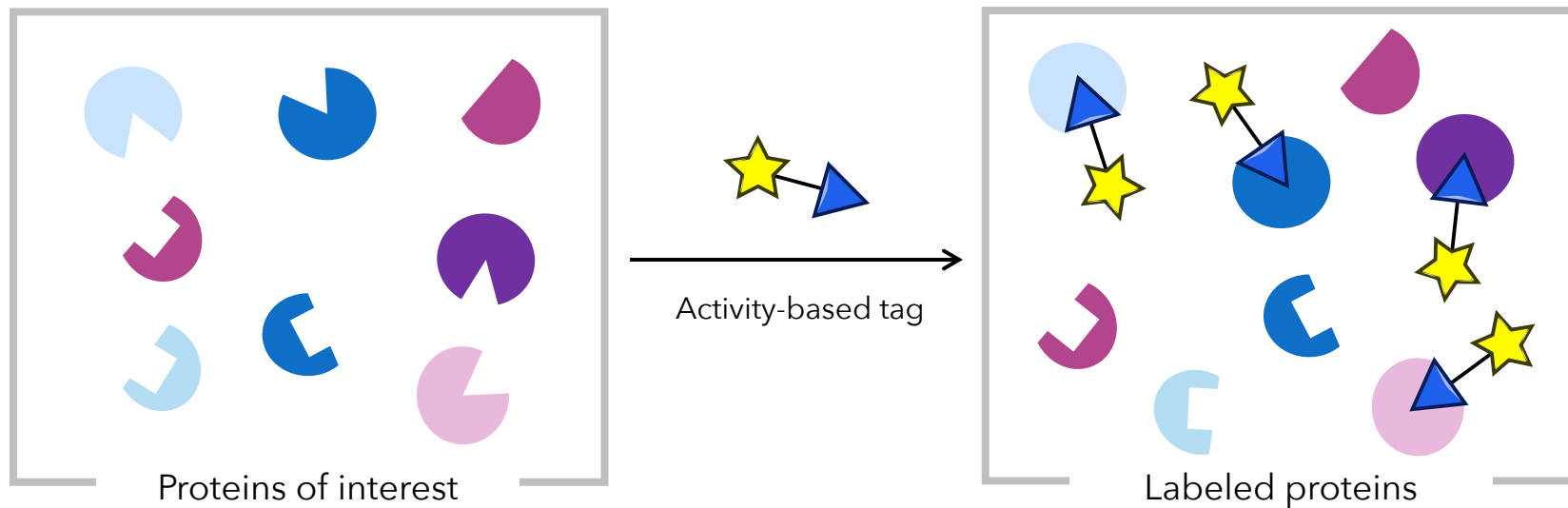
# Principles of activity-based proteomics

---



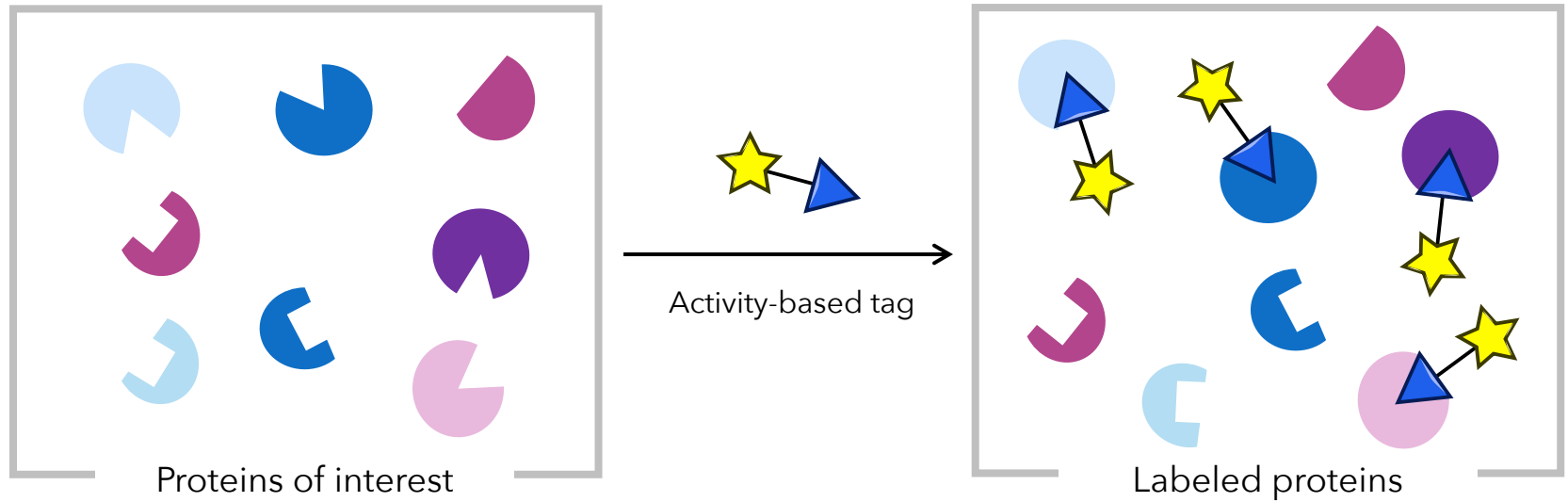
Proteins of interest

# Principles of activity-based proteomics





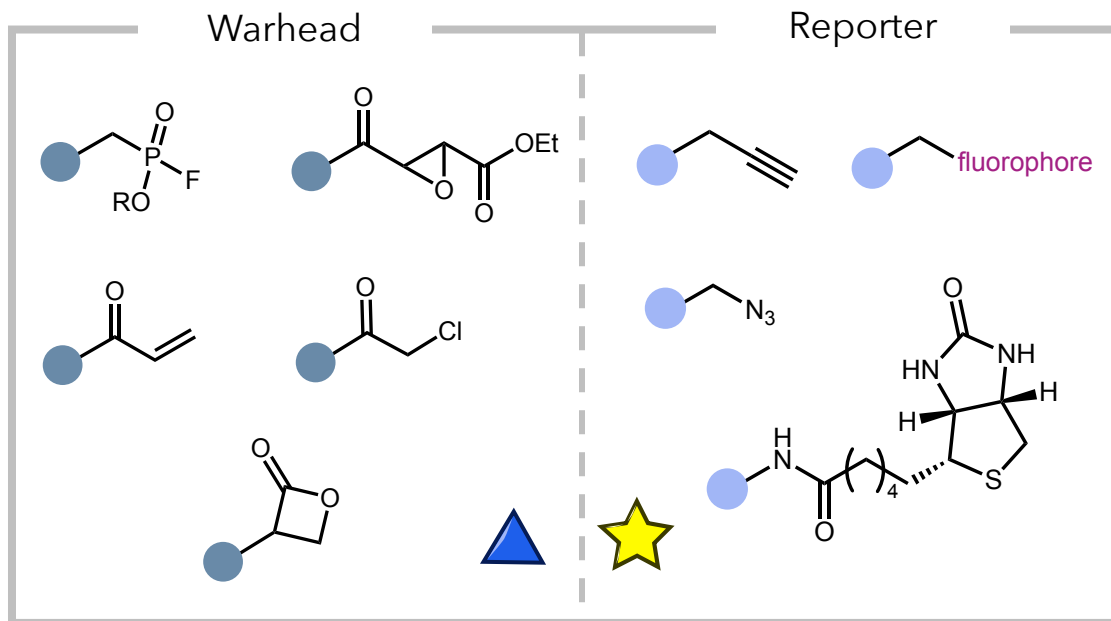
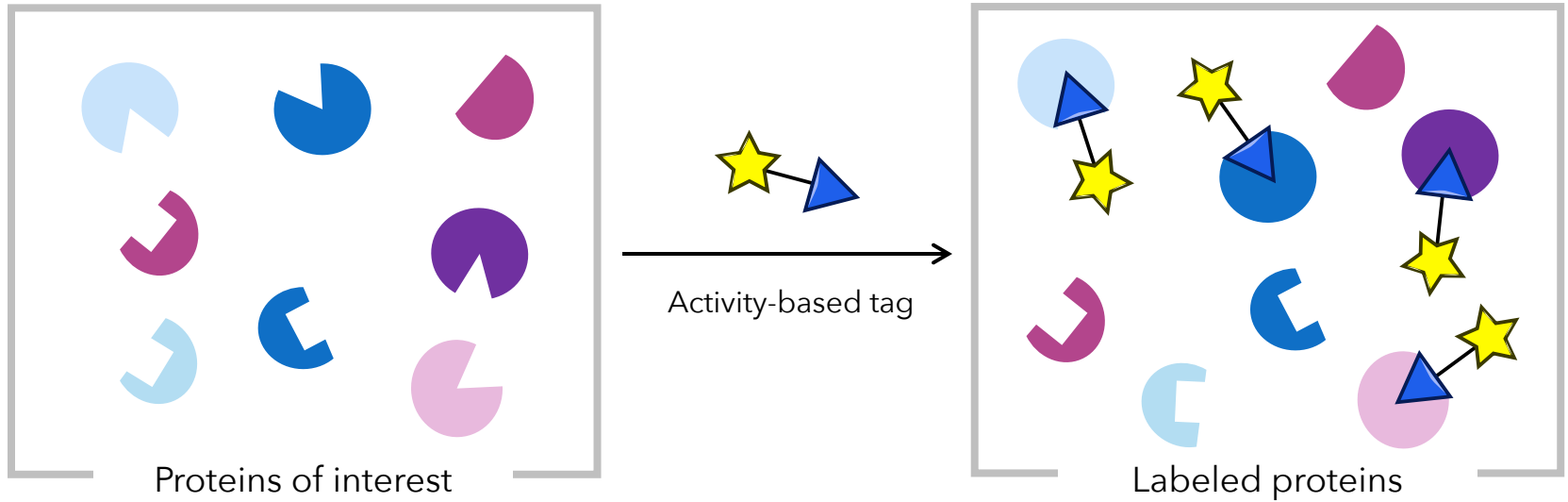
# Principles of activity-based proteomics



↓  
Visualization  
methods

- Fluorescence
- "Click"
- Streptavidin enrichment
- Mass spectrometry

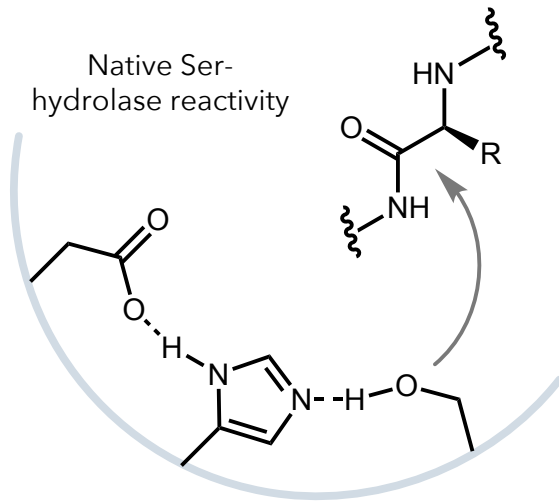
# Principles of activity-based proteomics



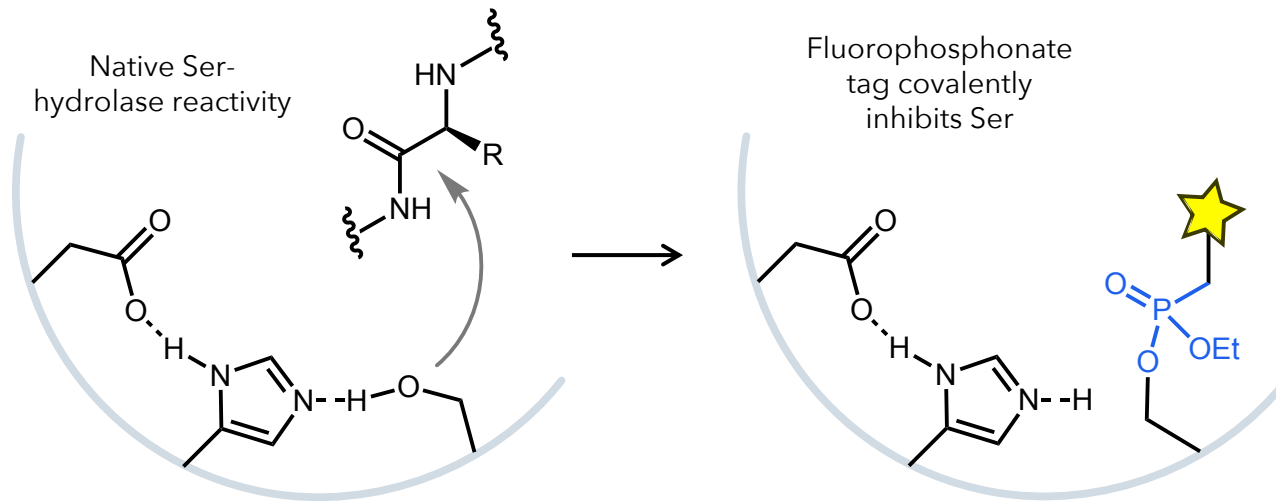
Visualization methods

- Fluorescence
- "Click"
- Streptavidin enrichment
- Mass spectrometry

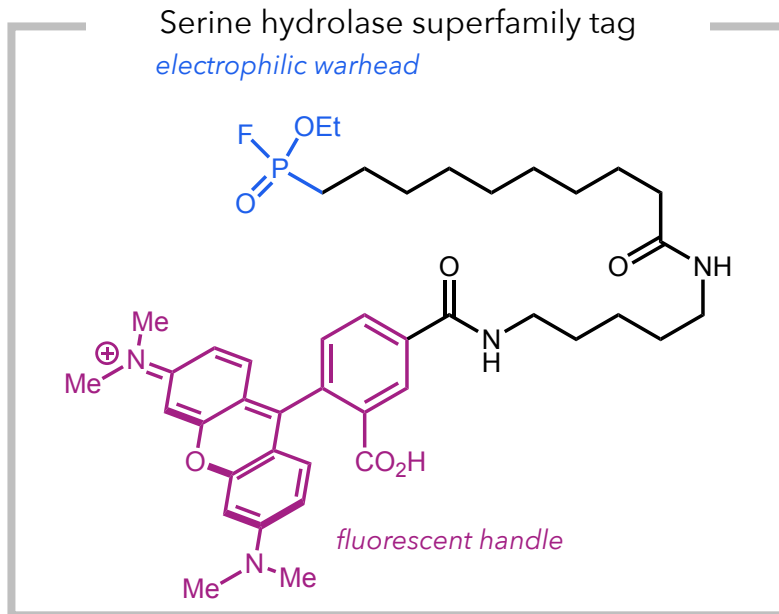
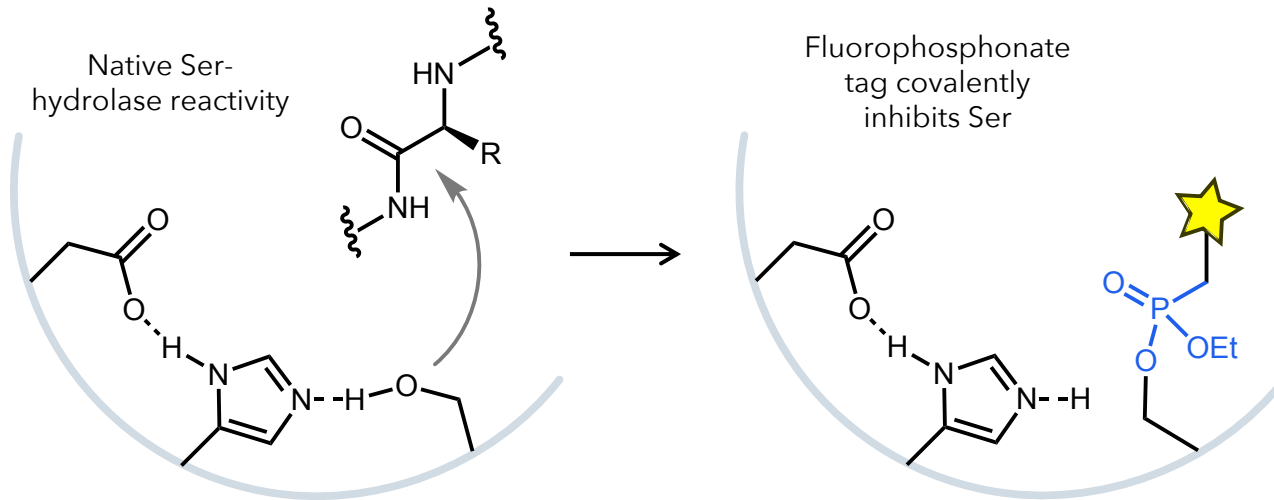
# ABPP characterization of serine hydrolases



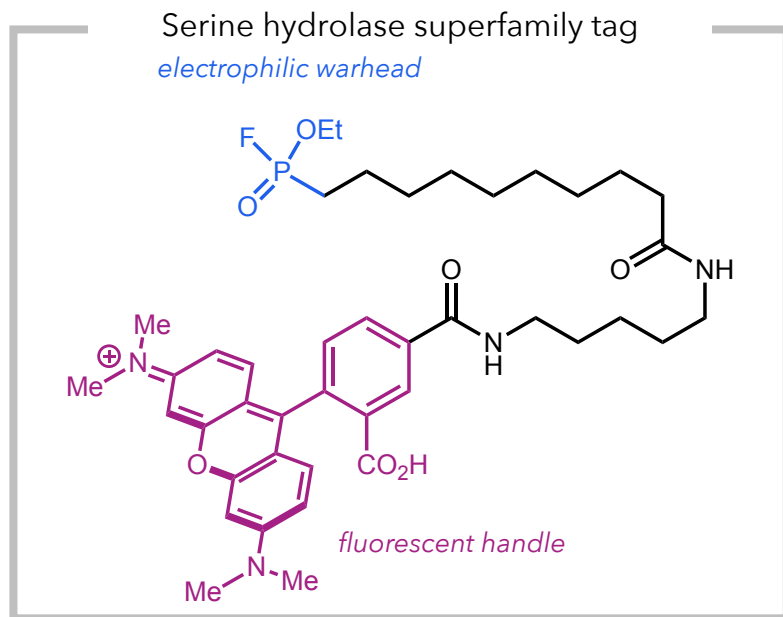
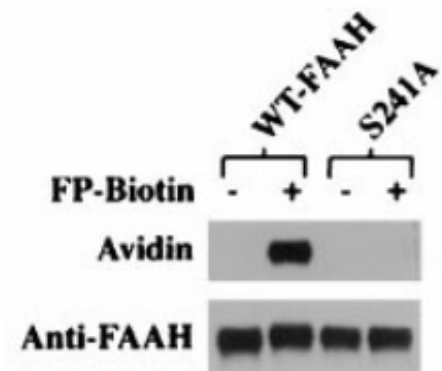
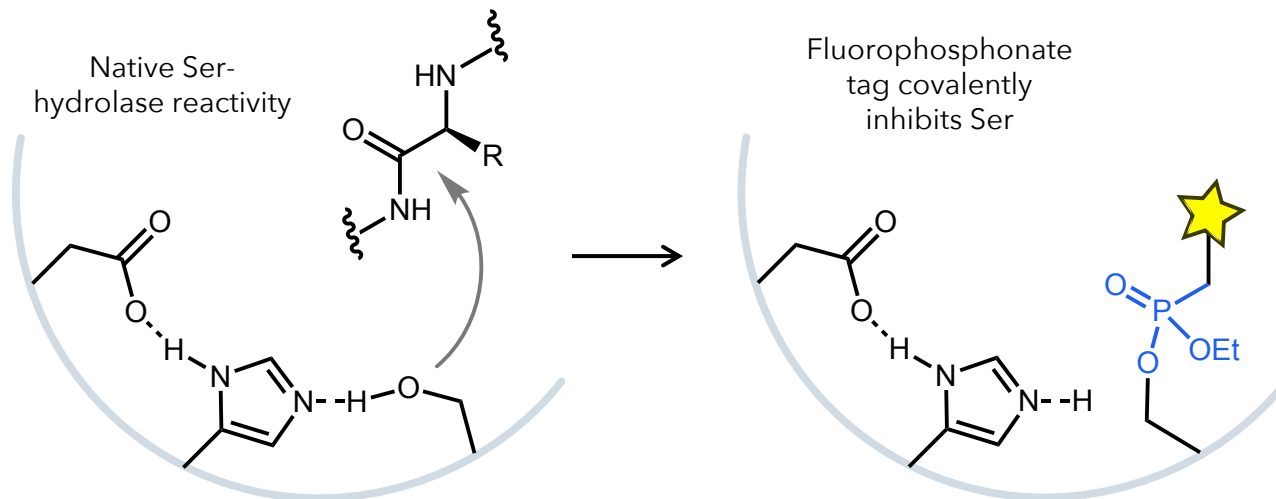
# ABPP characterization of serine hydrolases



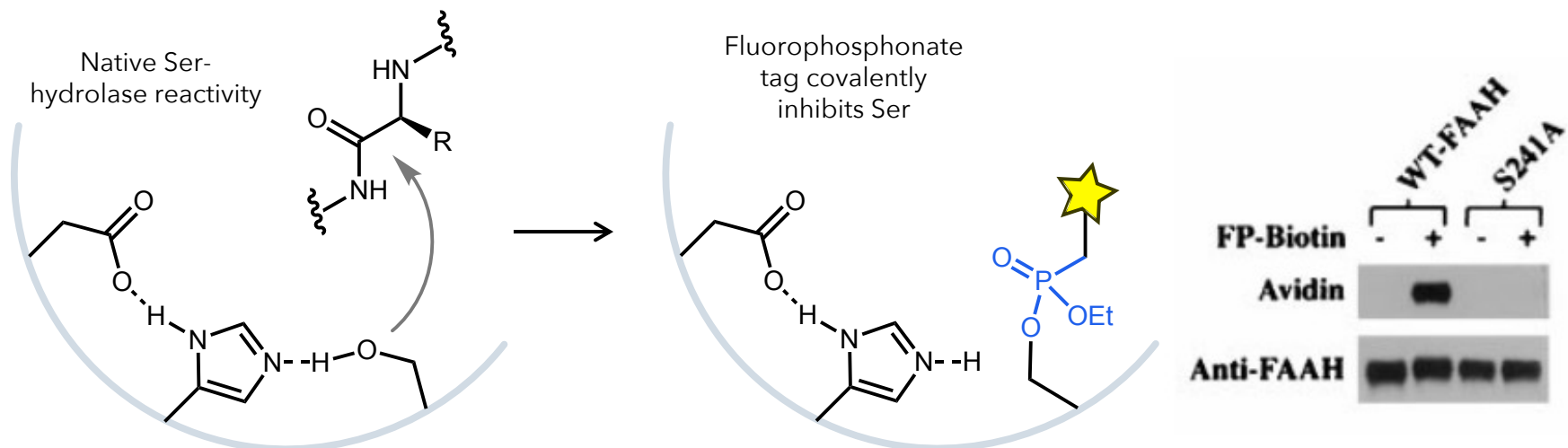
# ABPP characterization of serine hydrolases



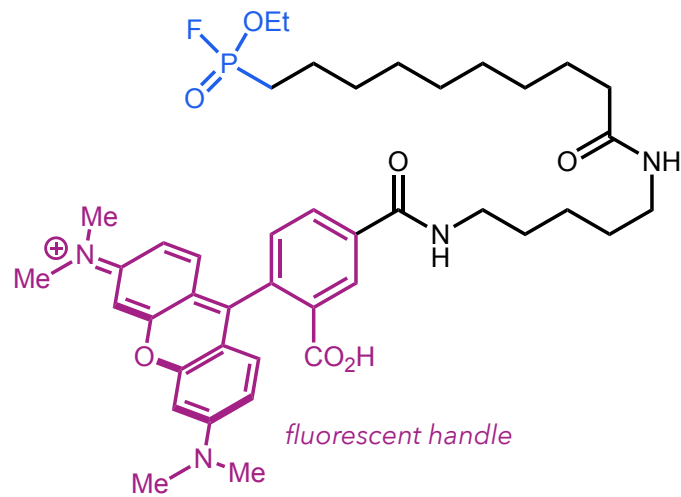
# ABPP characterization of serine hydrolases



# ABPP characterization of serine hydrolases

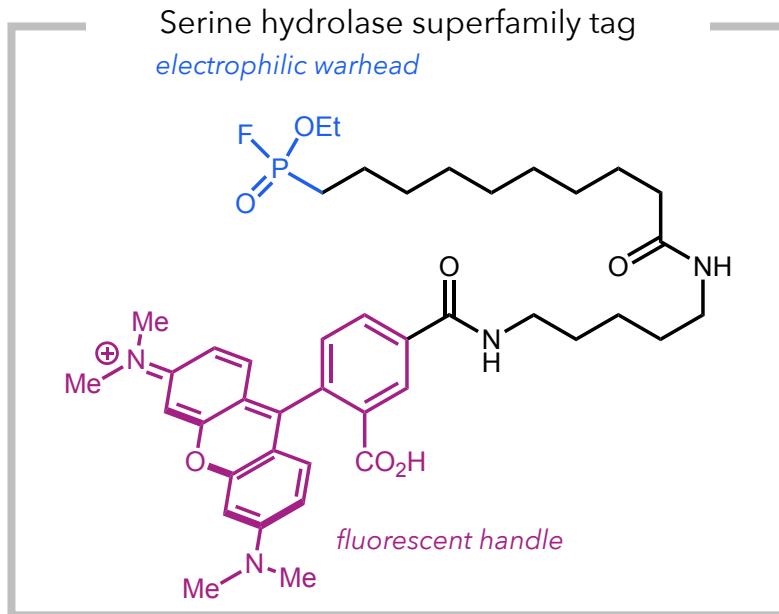
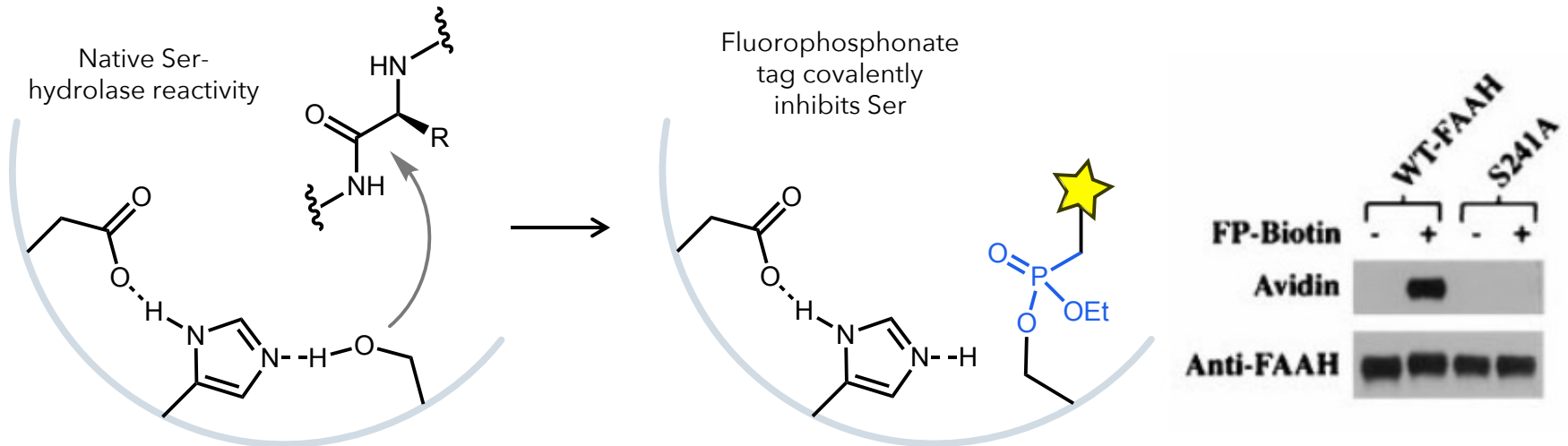


Serine hydrolase superfamily tag  
*electrophilic warhead*



- Serine hydrolases are involved in blood coagulation, inflammation, angiogenesis, amongst other processes

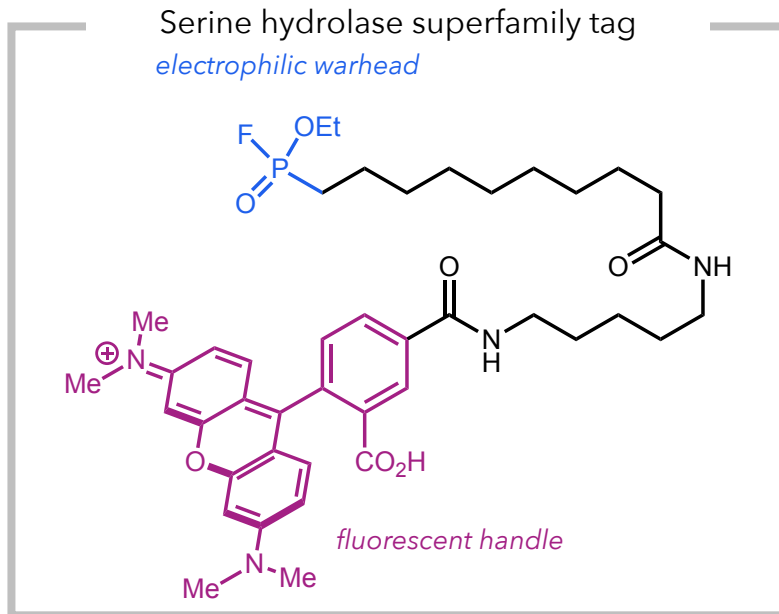
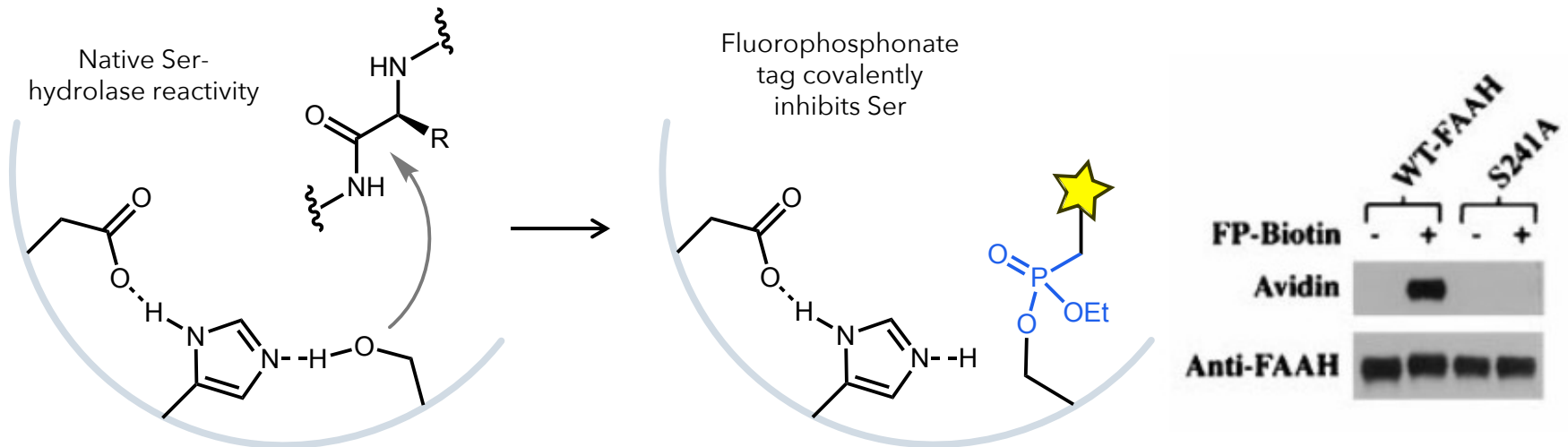
# ABPP characterization of serine hydrolases



- Serine hydrolases are involved in blood coagulation, inflammation, angiogenesis, amongst other processes
- Frequently implicated in cancer, emphysema
- APBB allows for profiling of multiple potential hydrolases in **few experiments**



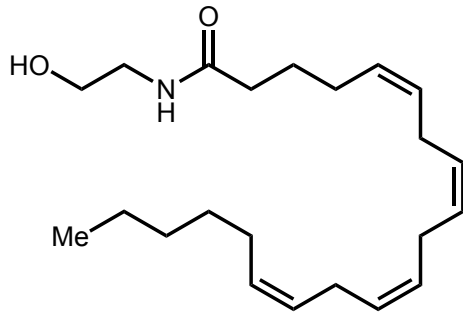
# ABPP characterization of serine hydrolases



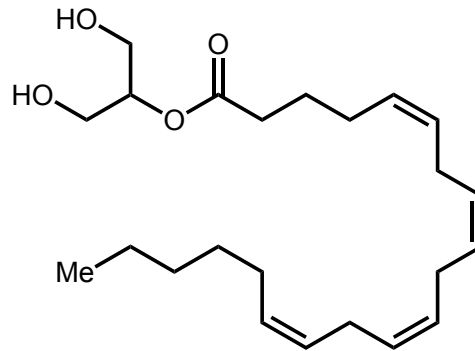
- Serine hydrolases are involved in blood coagulation, inflammation, angiogenesis, amongst other processes
- Frequently implicated in cancer, emphysema
- APBB allows for profiling of multiple potential hydrolases in **few experiments**
- This strategy is general and allows for characterization of specific protein classes dependent on activity

## Fatty Acid Amide Hydrolase (FAAH)

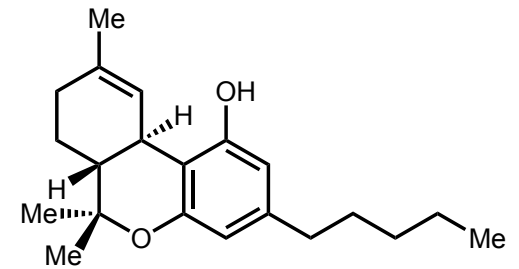
- Fatty acid amide hydrolase are serine hydrolases involved in lipid hydrolysis
- Endocannabinoid signaling pathway invoked in pain response, fear, and anxiety



anandamide



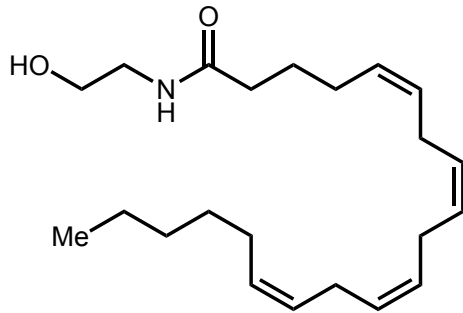
arachidonoyl glycerol



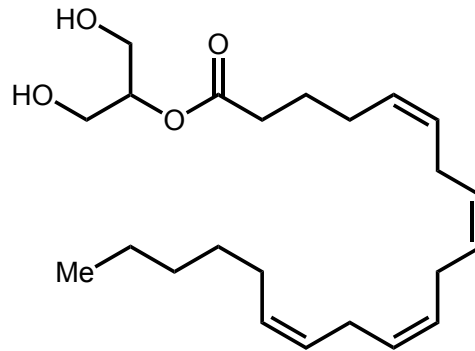
tetrahydrocannabinol

# Fatty Acid Amide Hydrolase (FAAH)

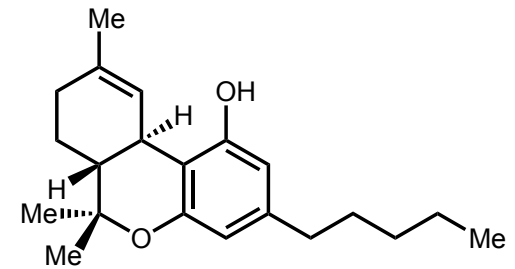
- Fatty acid amide hydrolase are serine hydrolases involved in lipid hydrolysis
- Endocannabinoid signaling pathway invoked in pain response, fear, and anxiety



anandamide



arachidonoyl glycerol



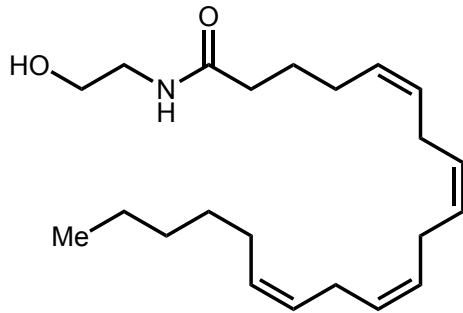
tetrahydrocannabinol



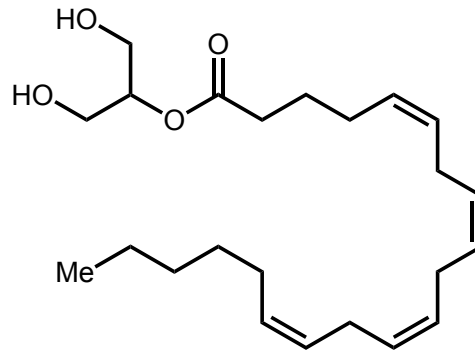
- Jo Cameron, 71, has never felt pain, fear, or anxiety
- Genetic studies reveal knockout of FAAH gene, with high levels of anandamide in brain
- Burns and cuts healed quicker than average

# Fatty Acid Amide Hydrolase (FAAH)

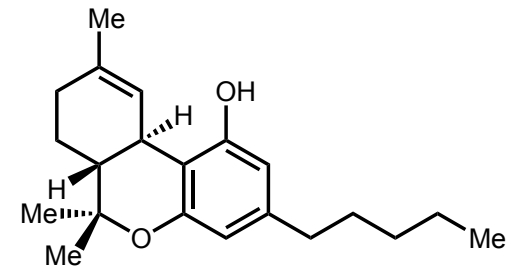
- Fatty acid amide hydrolase are serine hydrolases involved in lipid hydrolysis
- Endocannabinoid signaling pathway invoked in pain response, fear, and anxiety



anandamide



arachidonoyl glycerol



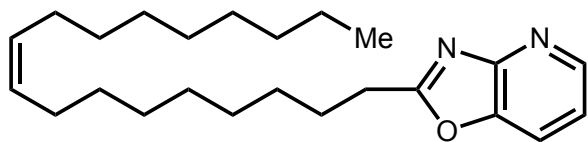
tetrahydrocannabinol



- Jo Cameron, 71, has never felt pain, fear, or anxiety
- Genetic studies reveal knockout of FAAH gene, with high levels of anandamide in brain
- Burns and cuts healed quicker than average

Is FAAH inhibition a strategy to treat pain?

## Fatty Acid Amide Hydrolase (FAAH) Inhibitors

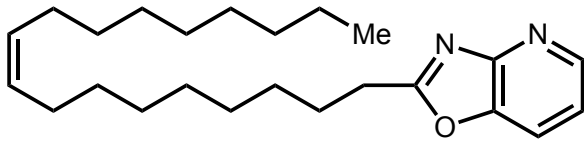


reversible FAAH inhibitor



Binding affinity not strong enough,  
non-specific targeting

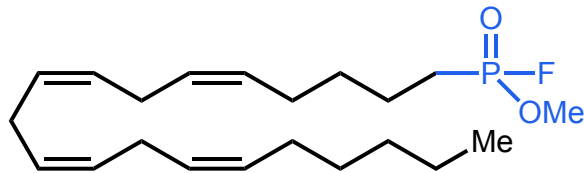
## Fatty Acid Amide Hydrolase (FAAH) Inhibitors



reversible FAAH inhibitor



Binding affinity not strong enough,  
non-specific targeting

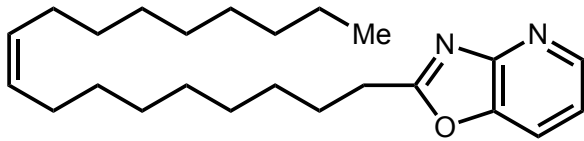


covalent FAAH inhibitor



Good binding affinity, too much off-  
target Ser-hydrolase inhibition

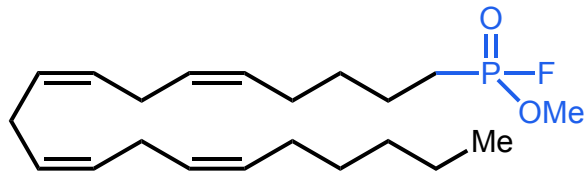
# Fatty Acid Amide Hydrolase (FAAH) Inhibitors



reversible FAAH inhibitor



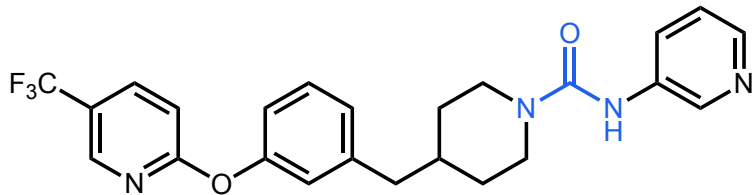
Binding affinity not strong enough,  
non-specific targeting



covalent FAAH inhibitor



Good binding affinity, too much off-  
target Ser-hydrolase inhibition

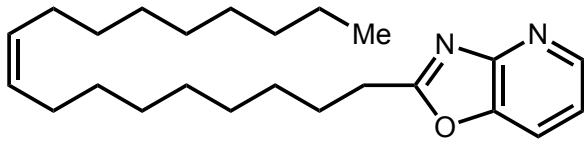


Pfizer PF-3845



ABPP shows remarkable selectivity  
and potent inhibition of FAAH

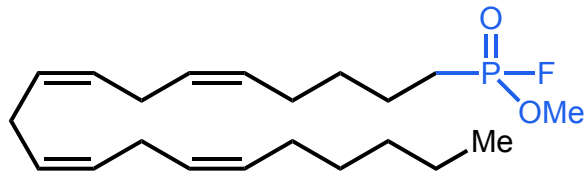
# Fatty Acid Amide Hydrolase (FAAH) Inhibitors



reversible FAAH inhibitor



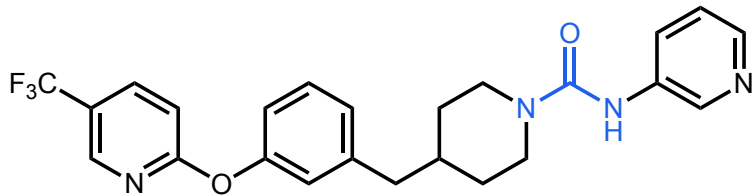
Binding affinity not strong enough,  
non-specific targeting



covalent FAAH inhibitor



Good binding affinity, too much off-  
target Ser-hydrolase inhibition



Pfizer PF-3845

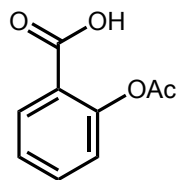


ABPP shows remarkable selectivity  
and potent inhibition of FAAH

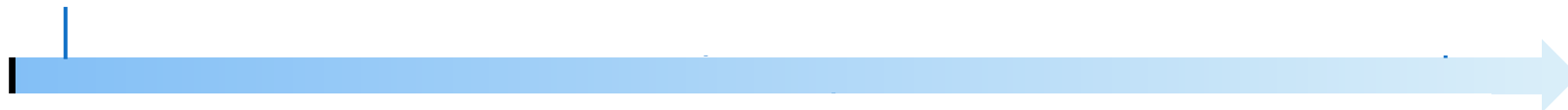
Could covalent inhibitors serve as viable  
candidates for drug discovery?



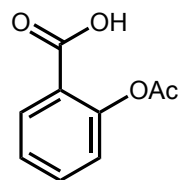
# Are covalent inhibitors a viable strategy for drug discovery?



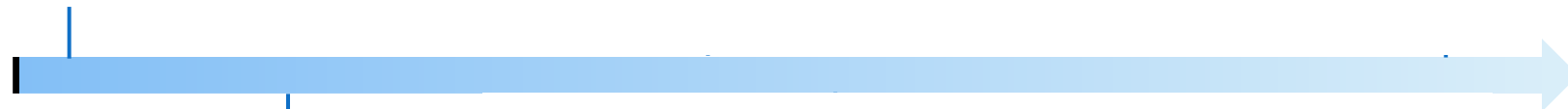
1890s - aspirin



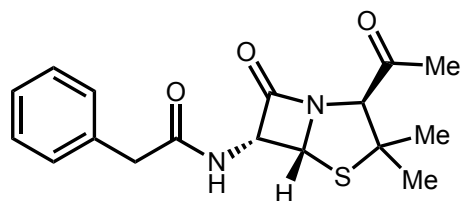
# Are covalent inhibitors a viable strategy for drug discovery?



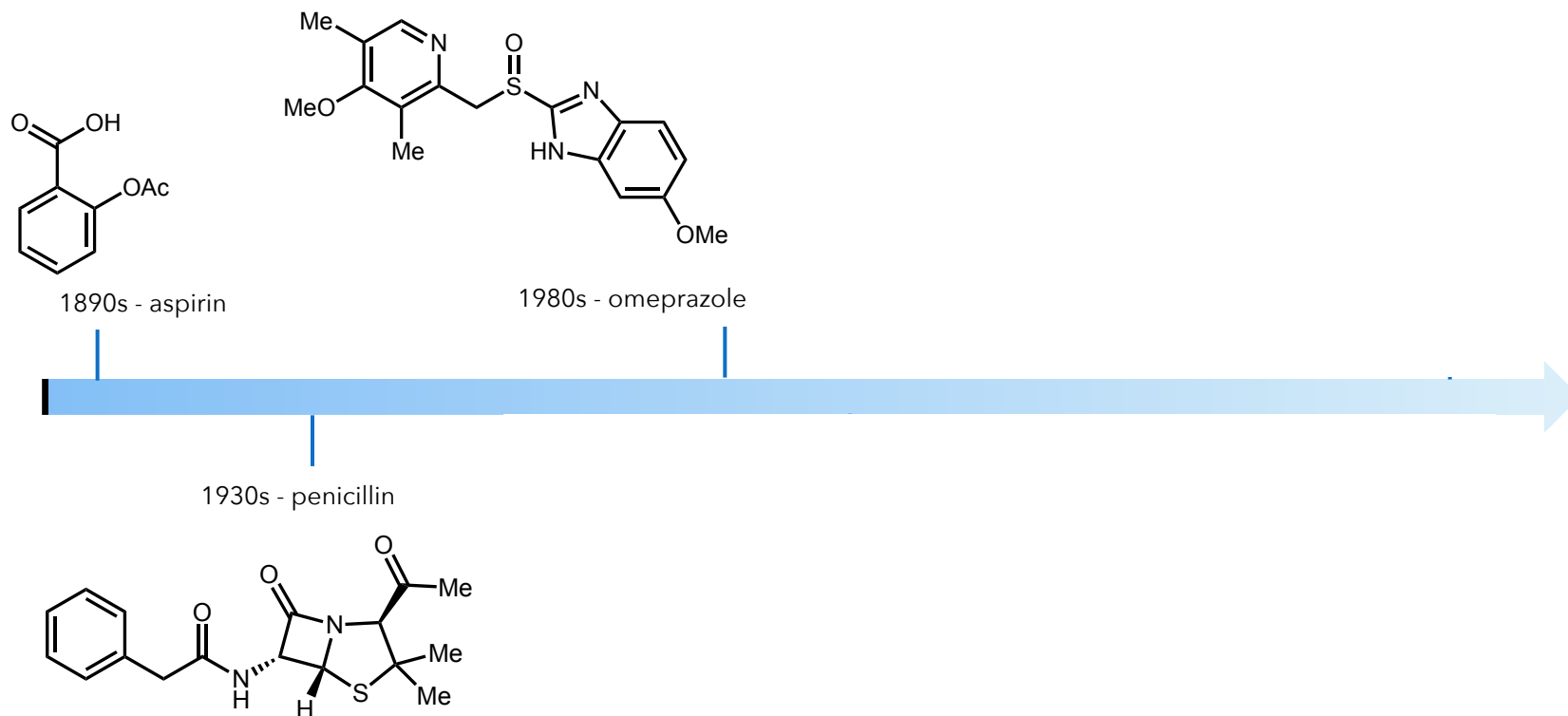
1890s - aspirin



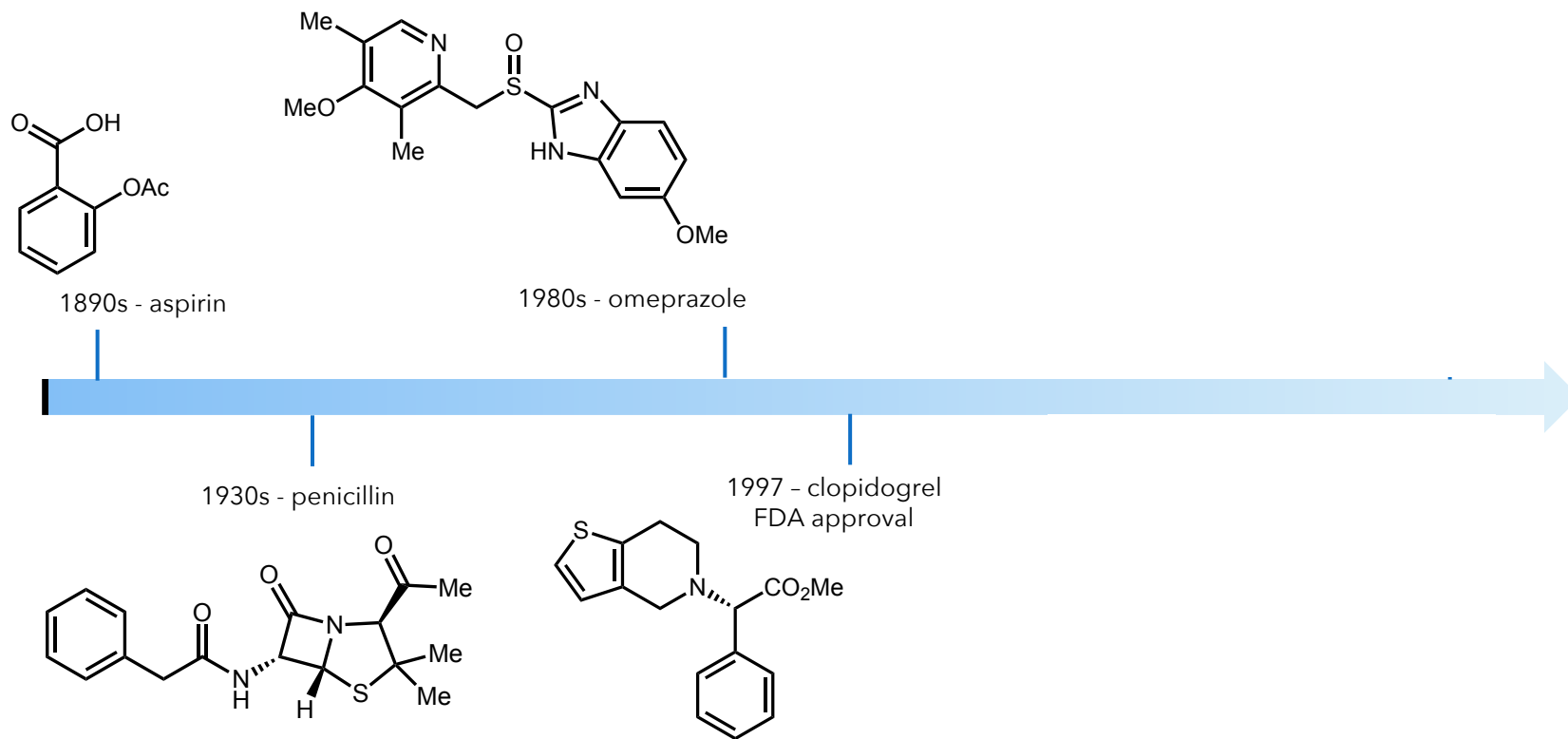
1930s - penicillin



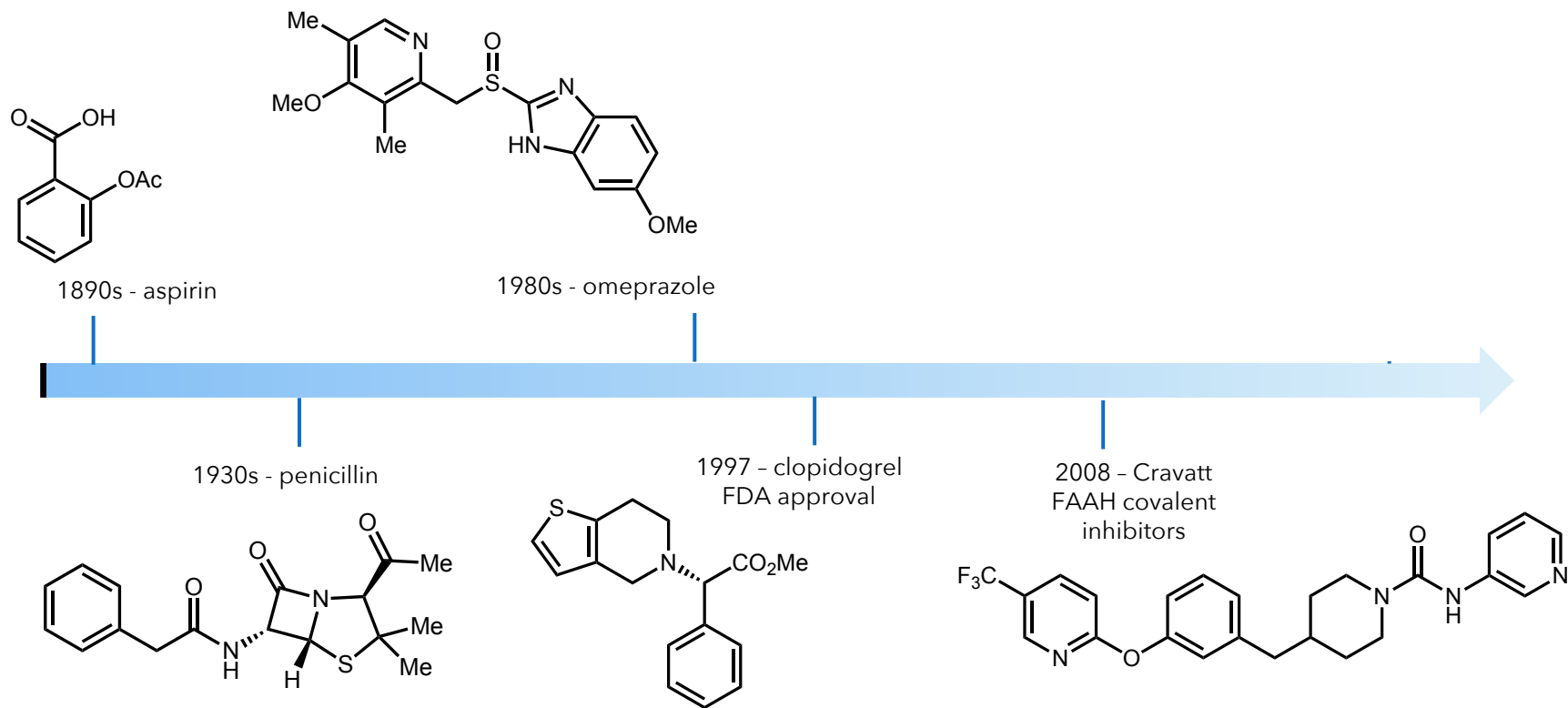
# Are covalent inhibitors a viable strategy for drug discovery?



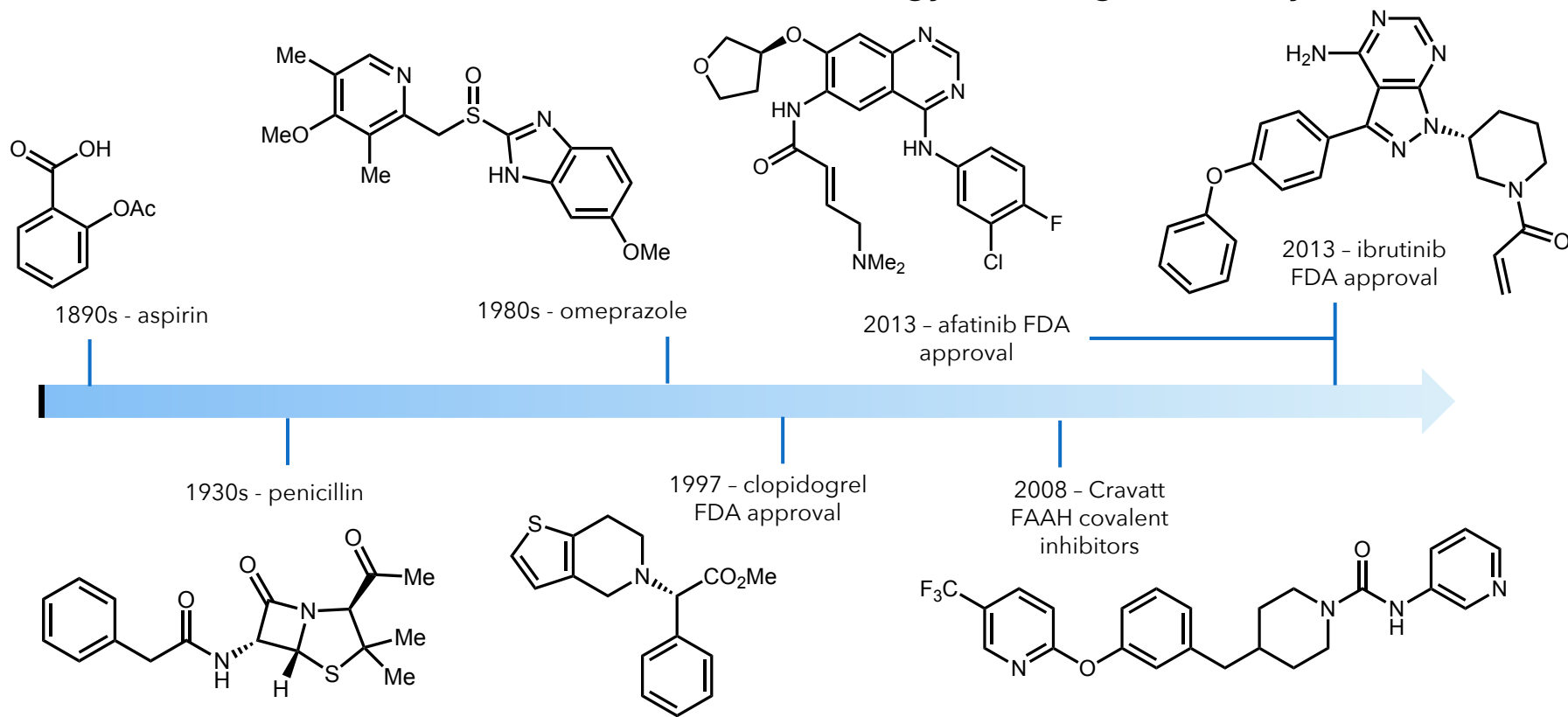
# Are covalent inhibitors a viable strategy for drug discovery?



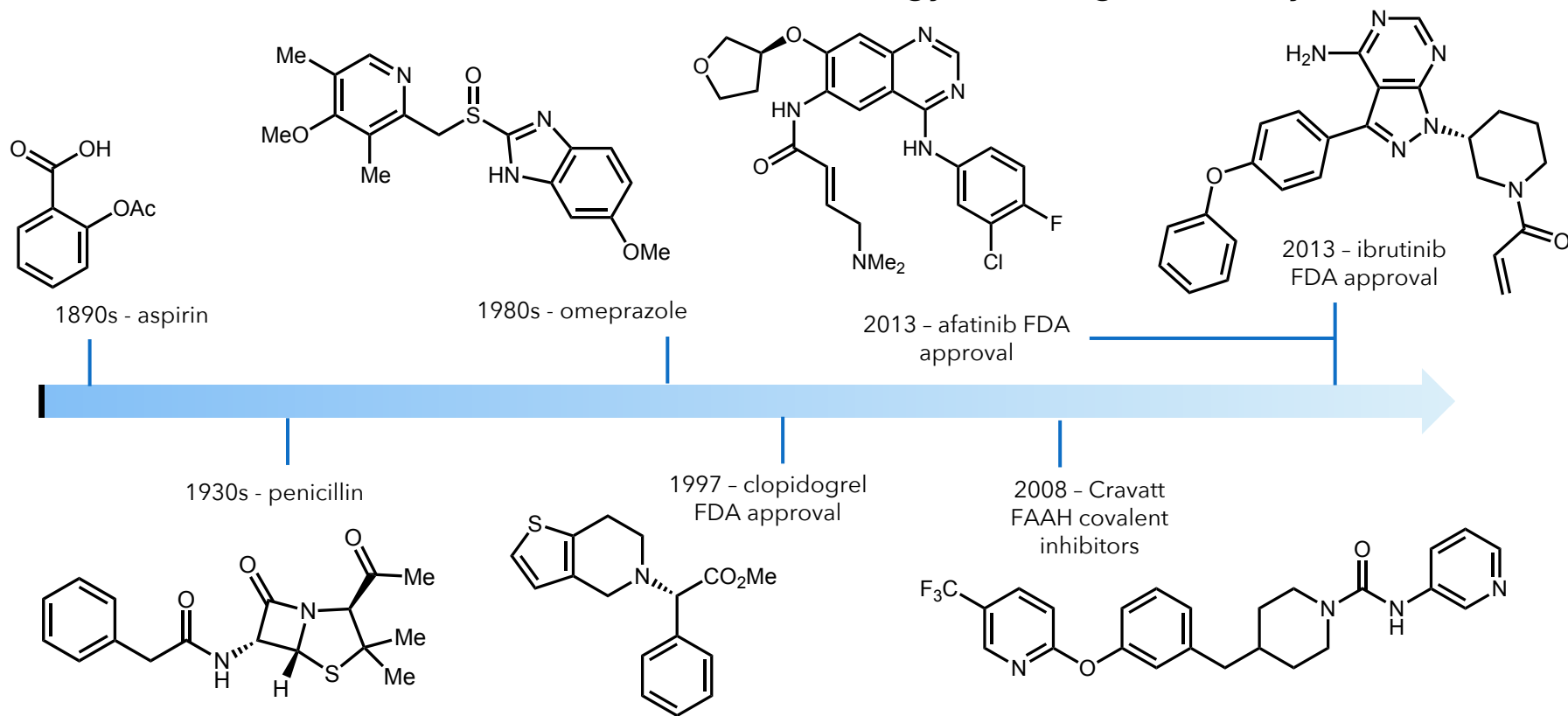
# Are covalent inhibitors a viable strategy for drug discovery?



# Are covalent inhibitors a viable strategy for drug discovery?

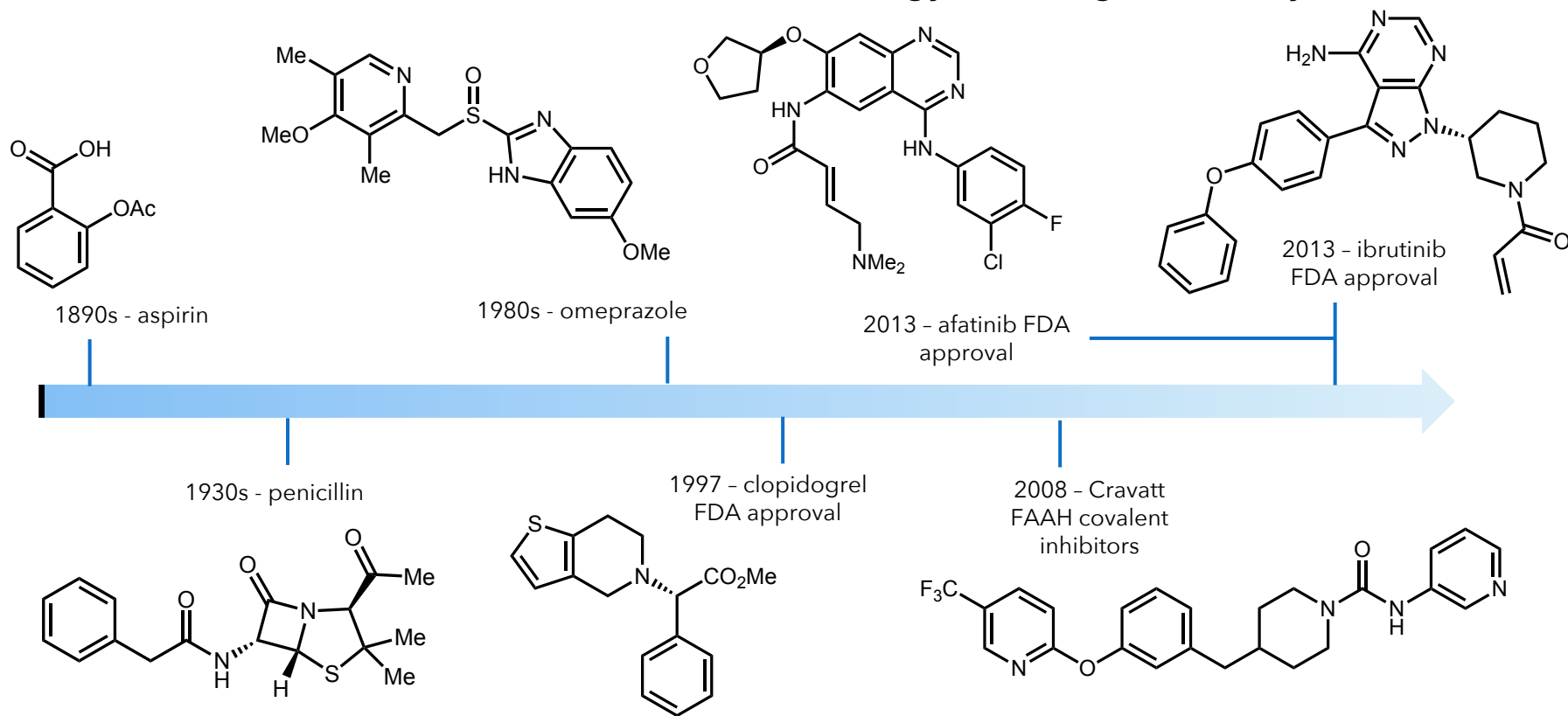


# Are covalent inhibitors a viable strategy for drug discovery?



- Meta-analysis of idiosyncratic drug toxicity reveals most "structural alerts" in drug design are over exaggerated

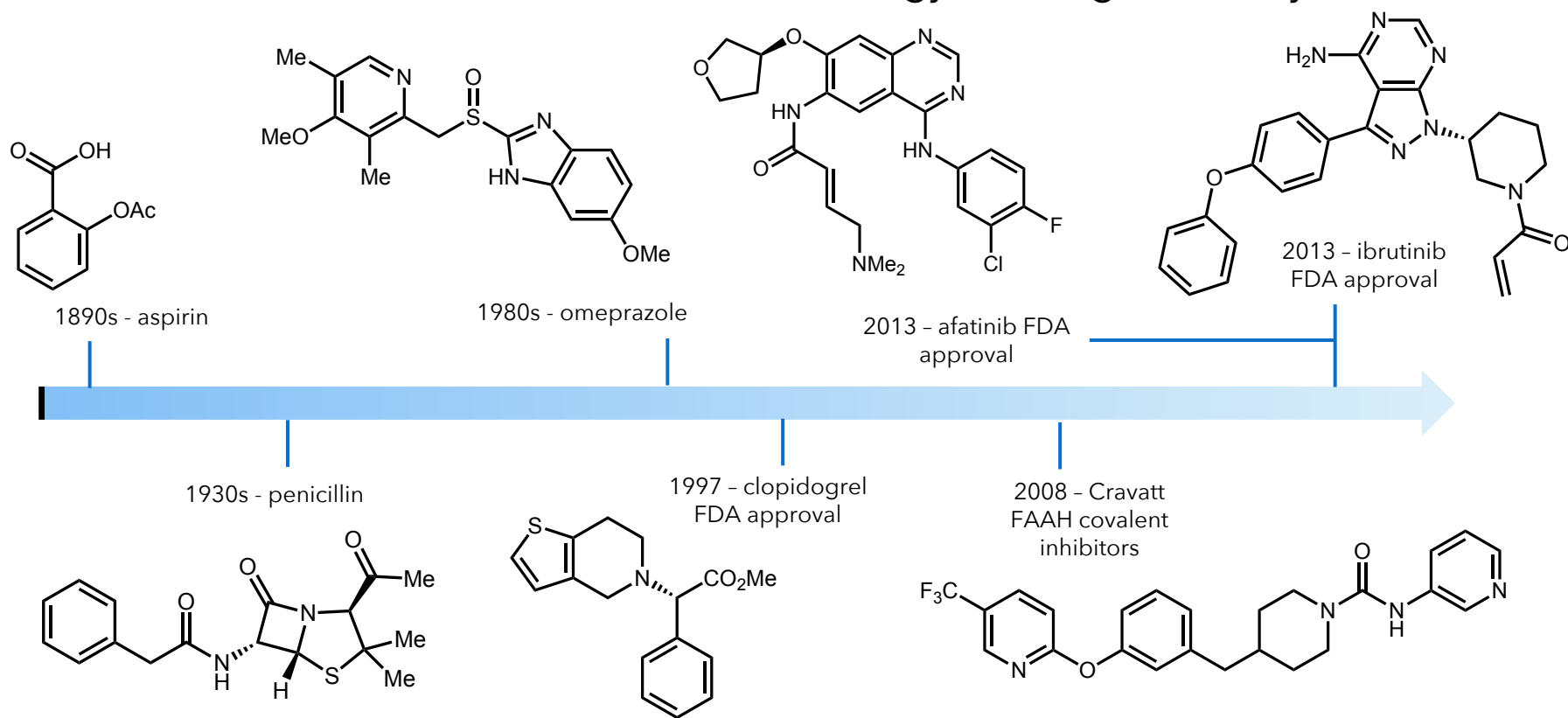
# Are covalent inhibitors a viable strategy for drug discovery?



- Meta-analysis of idiosyncratic drug toxicity reveals most “structural alerts” in drug design are over exaggerated
- The single biggest link to toxicity were high dosages (several hundred mgs) rather than specific mechanisms or structures



# Are covalent inhibitors a viable strategy for drug discovery?

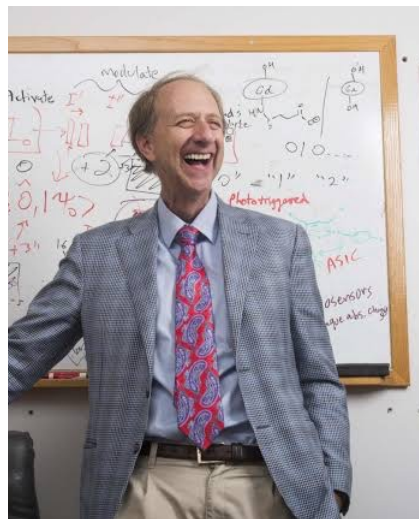


*Prior to such methods [ABPP], the question of selectivity of covalent drug candidates was pure speculation, which led to excessive concern. Once covalent drug candidates were shown to modify proteins with very high selectivity by ABPP, it gave medicinal chemists more confidence that this modality could be safe.*

# Outline

- Introduction to undruggable proteins
  - What makes a protein “undruggable”?
  - Attempts to drug K-Ras mutations
- Activity-based approaches to finding “druggable” sites
- Success stories in covalent drugs
  - Ibrutinib and Bruton’s tyrosine kinase
  - Sotorasib and K-Ras G12C
- Conclusions
  - “Yet to be drugged” instead of “undruggable”

Everything is bigger in Texas

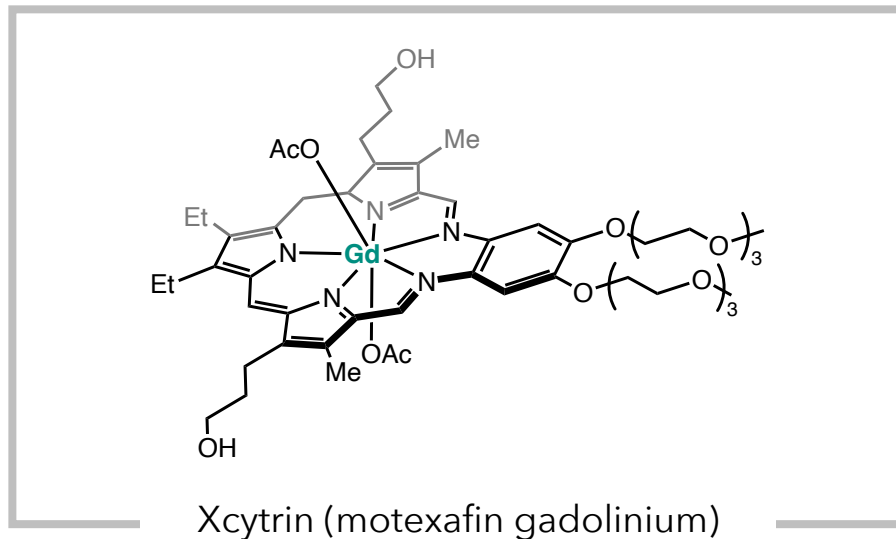
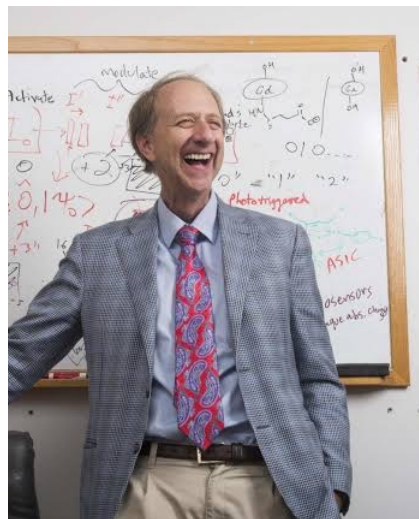


Prof. Jonathan Sessler  
**University of Texas at Austin**



Known for: synthesis of  
"texaphyrins",  
supramolecular chemistry

Everything is bigger in Texas

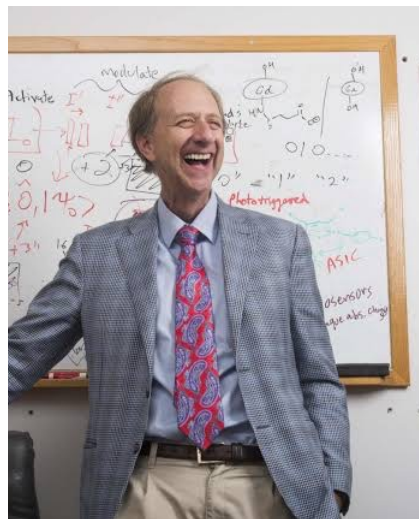


Prof. Jonathan Sessler  
**University of Texas at Austin**



Known for: synthesis of  
"texaphyrins",  
supramolecular chemistry

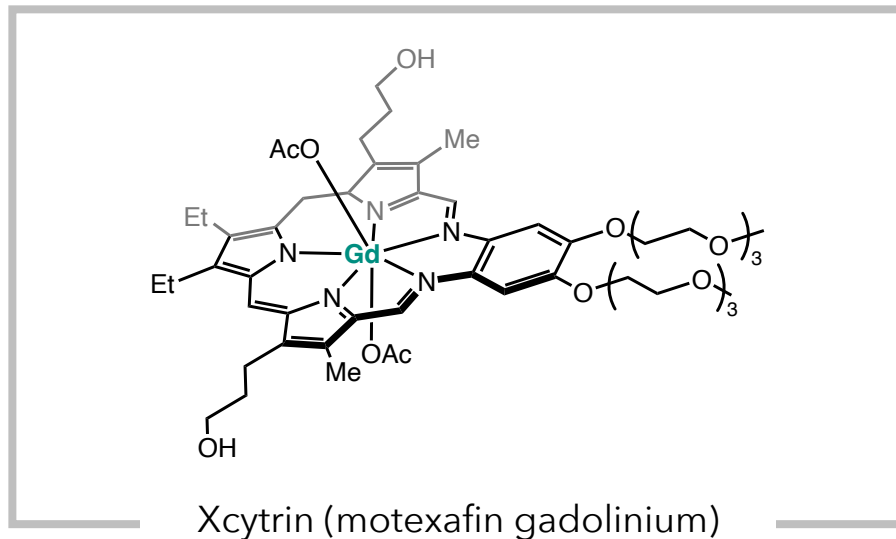
# Everything is bigger in Texas



Prof. Jonathan Sessler  
University of Texas at Austin



Known for: synthesis of  
"texaphyrins",  
supramolecular chemistry



Localization in tumors  
for more efficient  
radiation treatment

Attempt to develop treatment  
for brain metastases for lung  
cancer



## Pharmacyclics acquisition of Celera Genomics

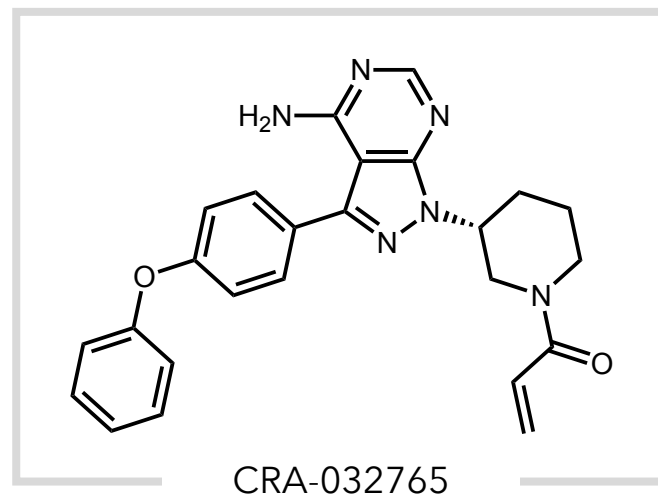
- In 2006, Pharmacyclics had acquired several small molecule probes from now defunct Celera Genomics, mostly focusing on **histone deacetylase inhibitors (HDACs)**

## Pharmacyclics acquisition of Celera Genomics

- In 2006, Pharmacyclics had acquired several small molecule probes from now defunct Celera Genomics, mostly focusing on **histone deacetylase inhibitors (HDACs)**
- Thrown in together with the deal, essentially for free, was an **early ABPP type probe for Bruton's tyrosine kinase (CRA-032765)**

## Pharmacyclics acquisition of Celera Genomics

- In 2006, Pharmacyclics had acquired several small molecule probes from now defunct Celera Genomics, mostly focusing on **histone deacetylase inhibitors (HDACs)**
- Thrown in together with the deal, essentially for free, was an **early ABPP type probe for Bruton's tyrosine kinase (CRA-032765)**

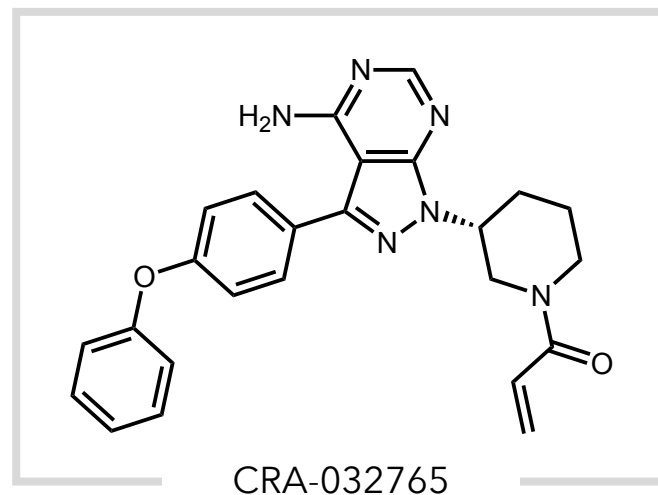




## Pharmacyclics acquisition of Celera Genomics

- In 2006, Pharmacyclics had acquired several small molecule probes from now defunct Celera Genomics, mostly focusing on **histone deacetylase inhibitors (HDACs)**
- Thrown in together with the deal, essentially for free, was an **early ABPP type probe for Bruton's tyrosine kinase (CRA-032765)**

*In the \$6.6 million transaction, the Celera team ascribed **next to no value to the BTK inhibitors...** Celera didn't even bother to secure future milestone payments... **The company essentially include CRA-032765 in the deal for nothing.***

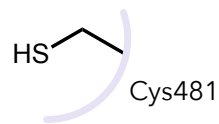
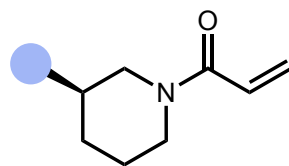
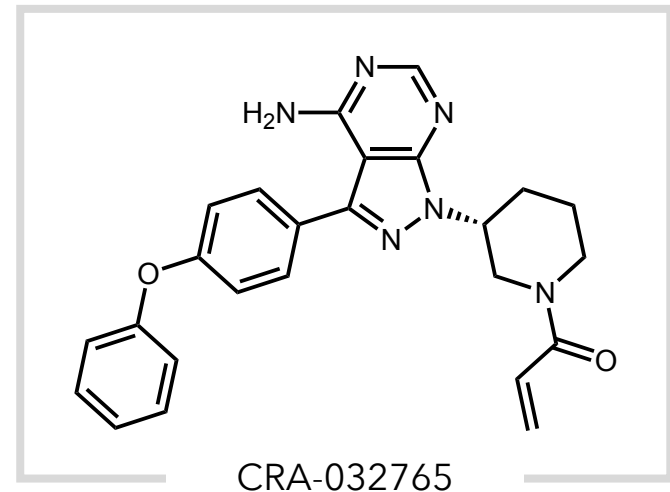


# Pharmacyclics acquisition of Celera Genomics

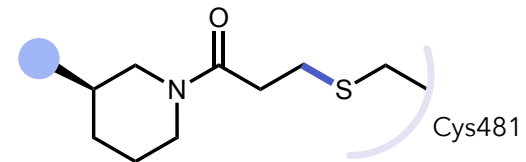
- In 2006, Pharmacyclics had acquired several small molecule probes from now defunct Celera Genomics, mostly focusing on **histone deacetylase inhibitors (HDACs)**
- Thrown in together with the deal, essentially for free, was an **early ABPP type probe for Bruton's tyrosine kinase (CRA-032765)**

*In the \$6.6 million transaction, the Celera team ascribed **next to no value to the BTK inhibitors...** Celera didn't even bother to secure future milestone payments... **The company essentially include CRA-032765 in the deal for nothing.***

- Early studies showed CRA-032765 effectively inhibited BTK, but concerns about covalent binding mechanism led to it being cast aside

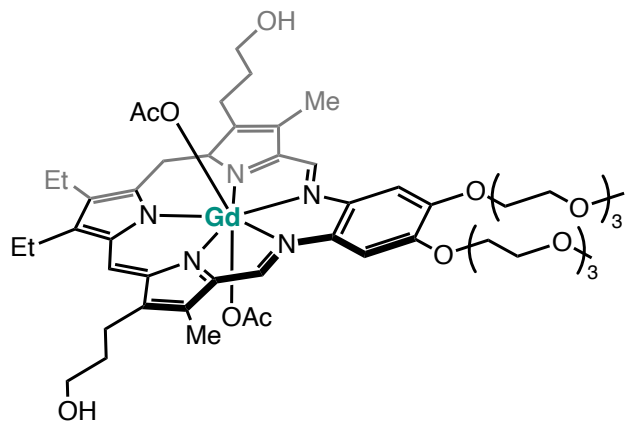


Bruton's tyrosine kinase (BTK)

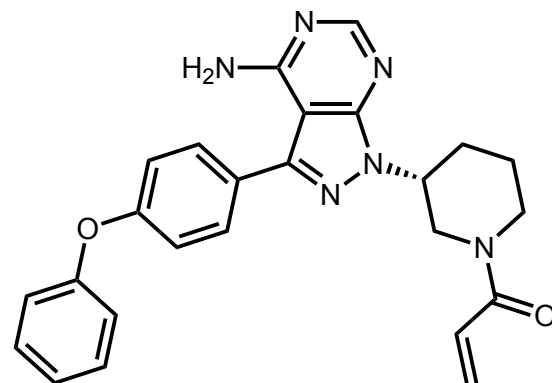


covalent inhibition of BTK  
prevent B-cell proliferation

## Pharmacyclics and Xcytrin

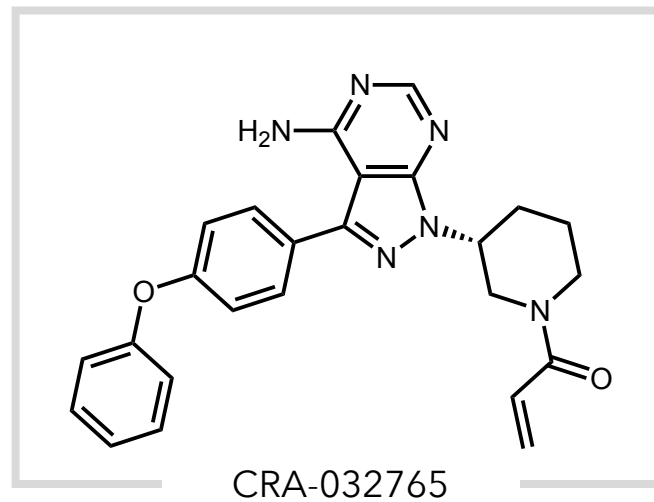
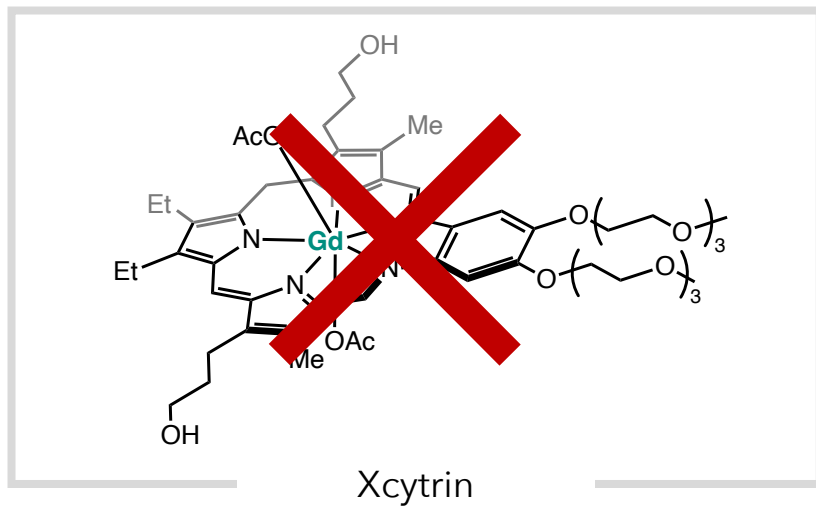


Xcytrin



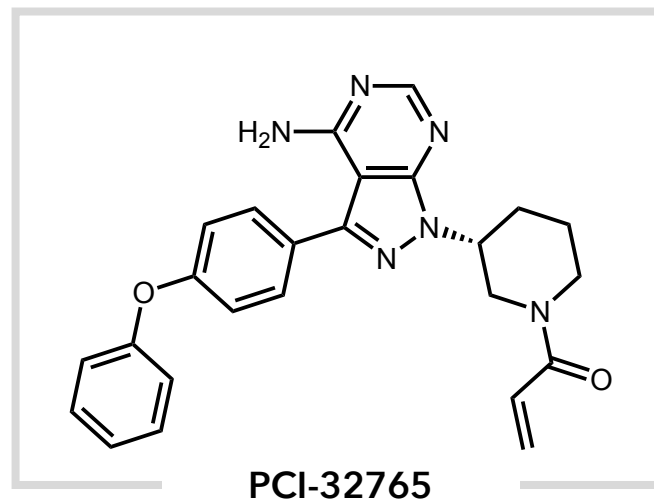
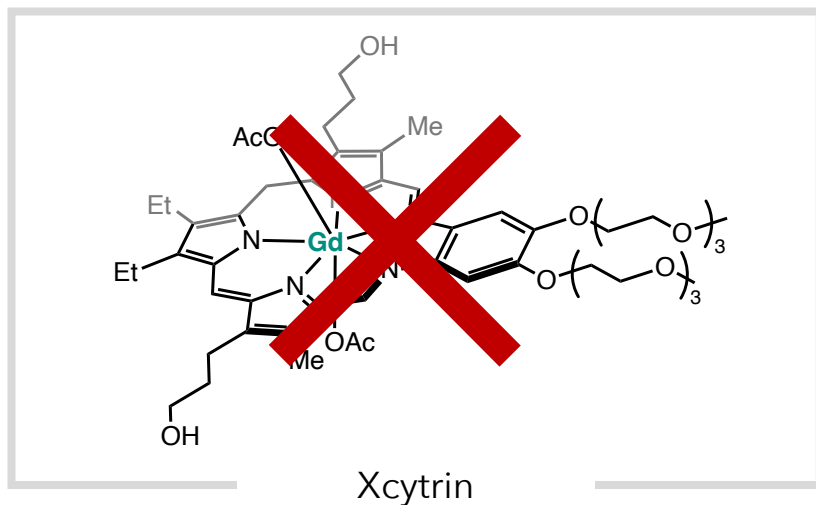
CRA-032765

## Pharmacyclics and Xcytrin



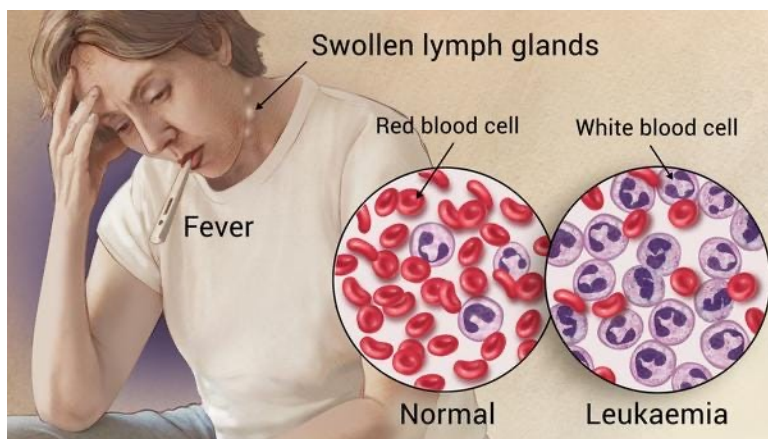
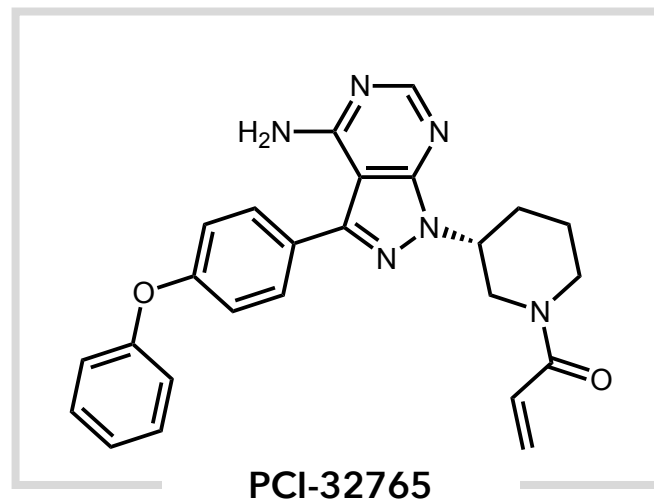
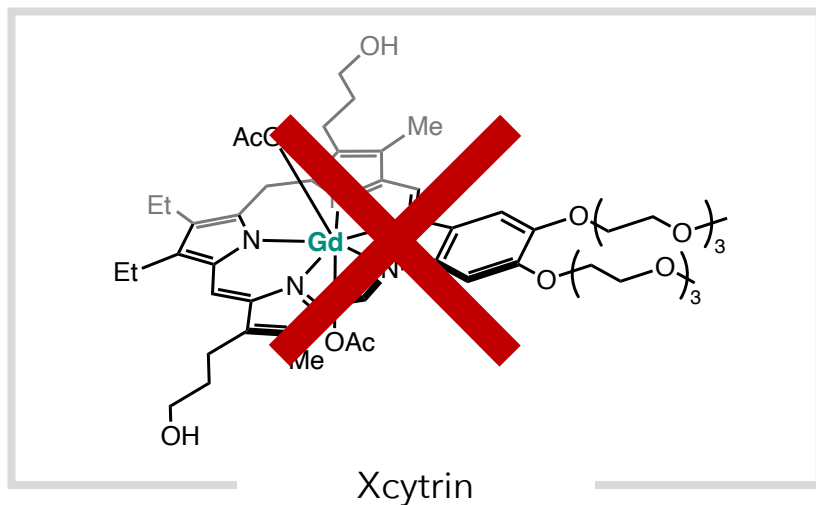
- After years of attempts, Pharmacyclics received a non-approvable letter from the FDA in 2007

## Pharmacyclics and Xcytrin



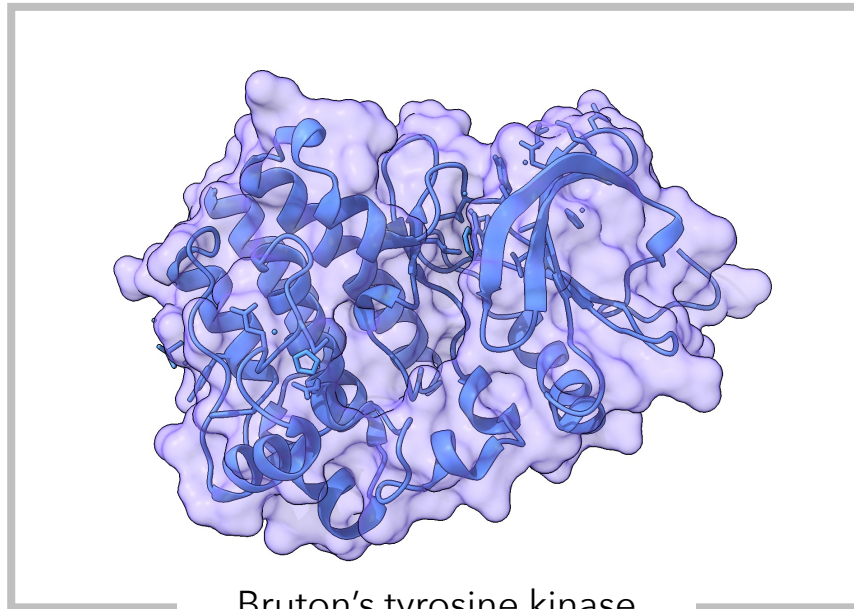
- After years of attempts, Pharmacyclics received a non-approvable letter from the FDA in 2007
- Scrambling during a financial crisis, Pharmacyclics invested some effort into studying CRA-032765, now given codename **PCI-32765**

## Pharmacyclics and Xcytrin

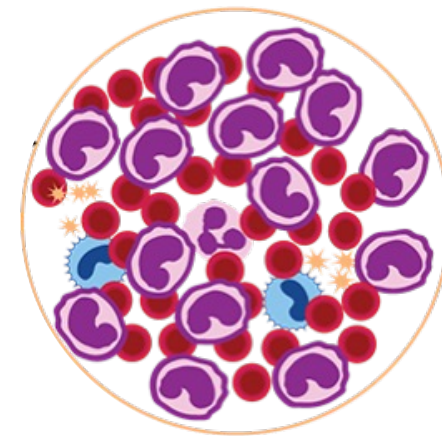
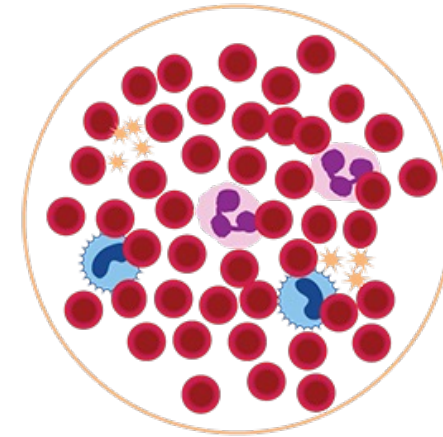


- After years of attempts, Pharmacyclics received a non-approvable letter from the FDA in 2007
- Scrambling during a financial crisis, Pharmacyclics invested some effort into studying CRA-032765, now given codename **PCI-32765**
- Early clinical trials showed reduction of tumors in patients with **chronic lymphocytic leukemia (CLL)**, the most common form of adult leukemia

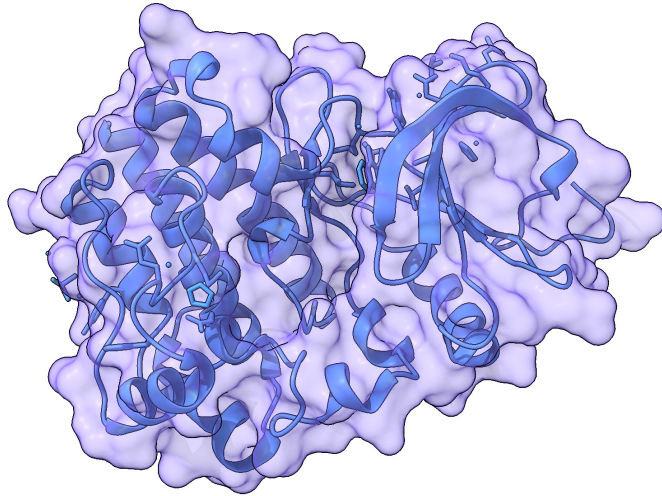
## Bruton's tyrosine kinase (BTK) and B-cell cancers



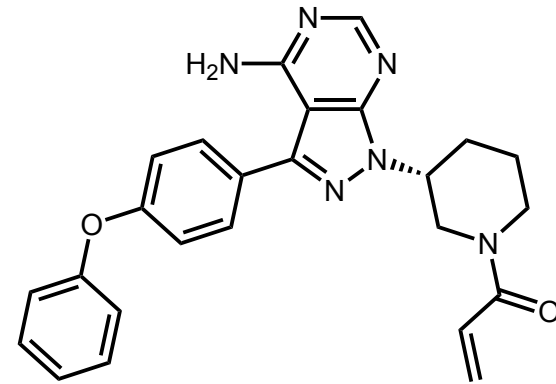
- Mutations cause various B-cell malignancies
- Involved in non-Hodgkin lymphomas, chronic lymphocytic leukemia, multiple myeloma,
- Traditional treatments include harsh chemotherapies



## Bruton's tyrosine kinase (BTK) and B-cell cancers

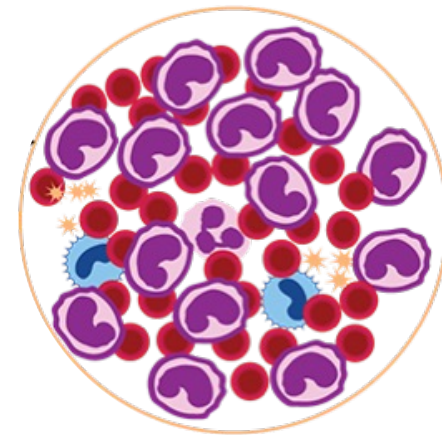


Bruton's tyrosine kinase



PCI-32765

- Mutations cause various B-cell malignancies
- Involved in non-Hodgkin lymphomas, chronic lymphocytic leukemia, multiple myeloma,
- Traditional treatments include harsh chemotherapies
- **PCI-32765 shows remarkable selectivity for BTK inhibition**



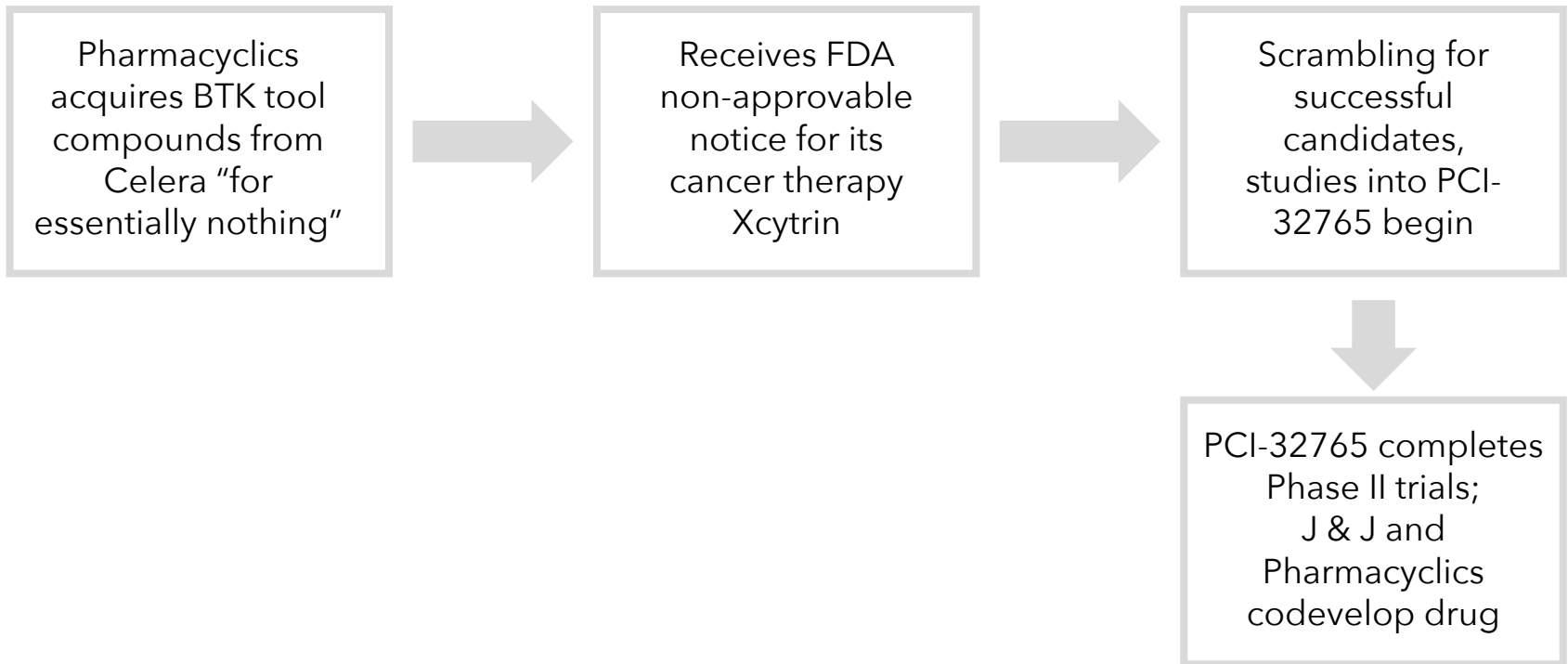
Leukemia



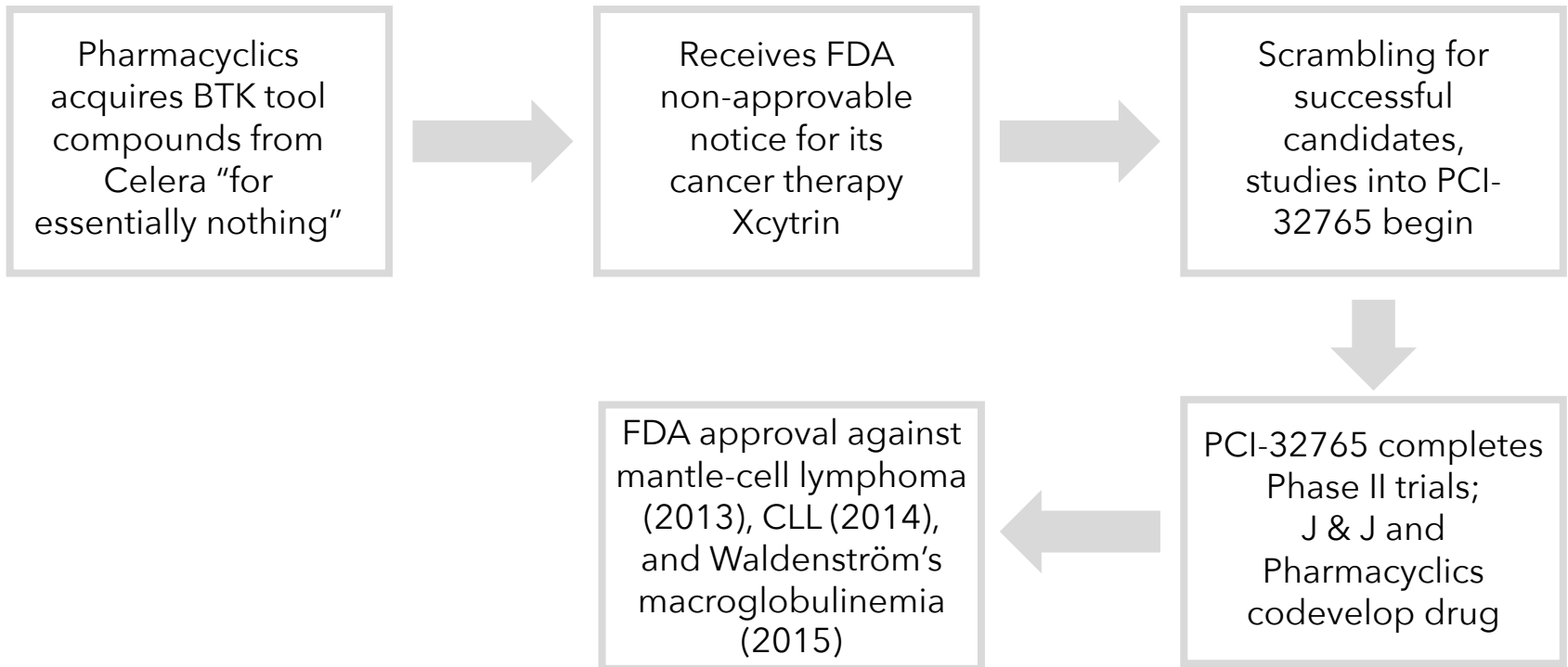
## The remarkable story of ibrutinib



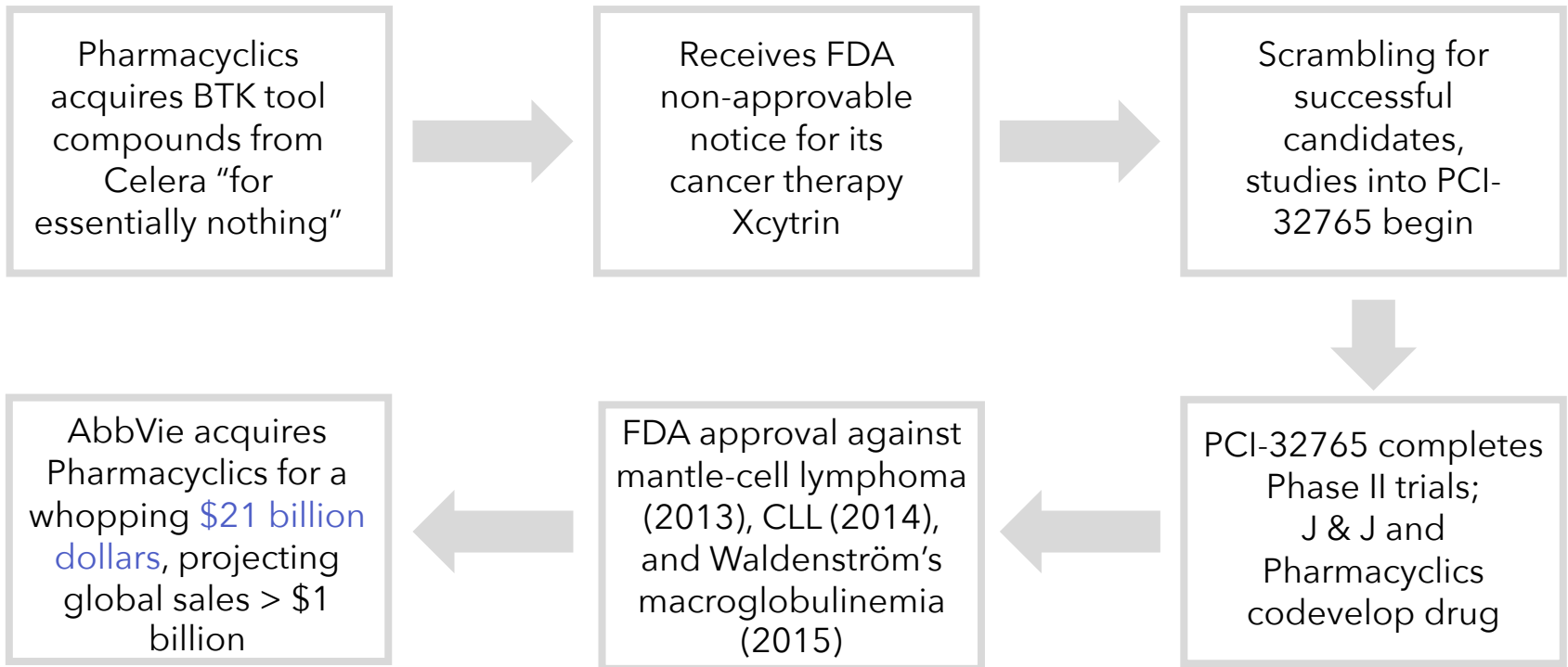
## The remarkable story of ibrutinib



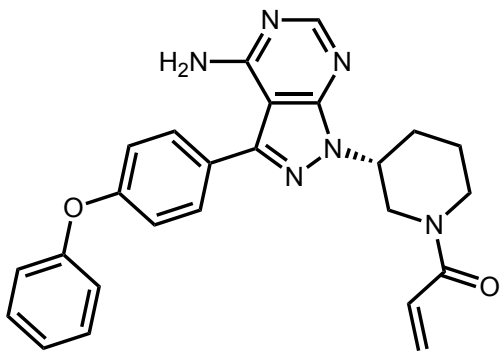
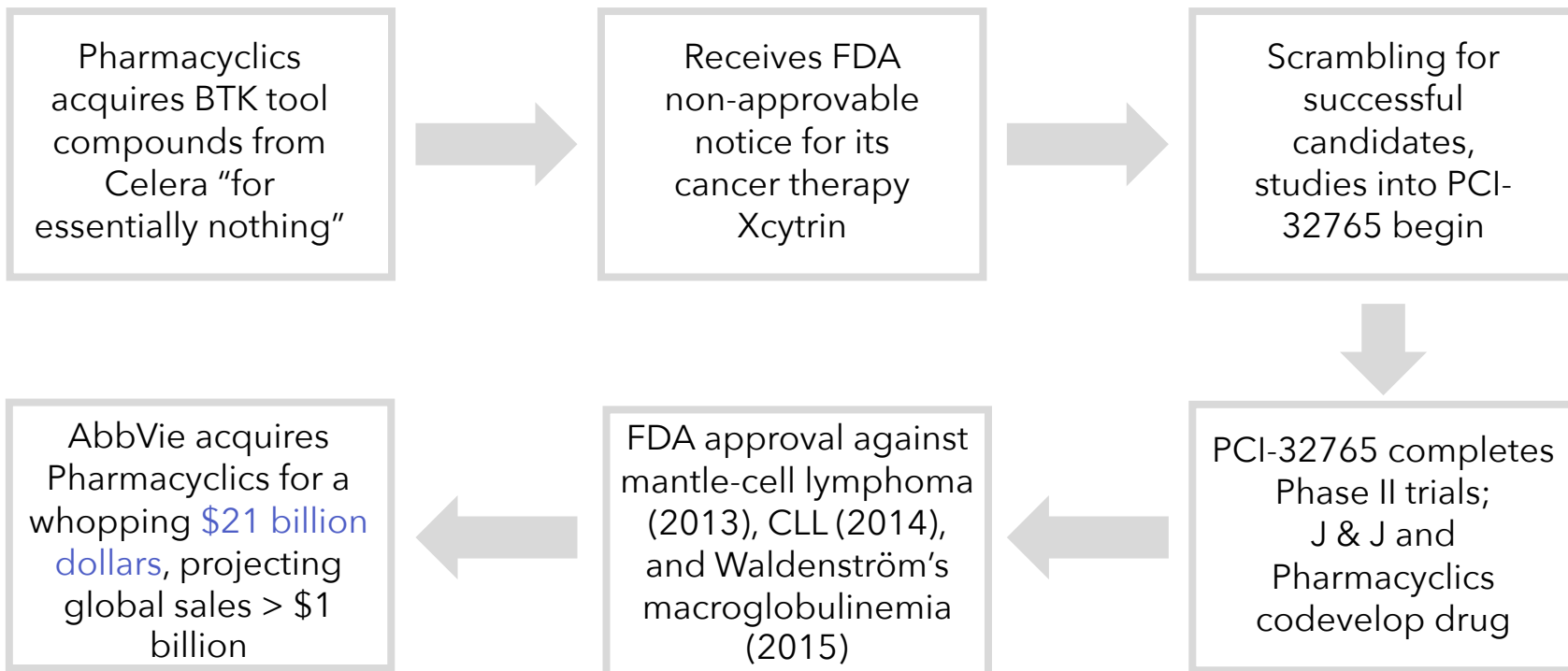
## The remarkable story of ibrutinib



## The remarkable story of ibrutinib



## The remarkable story of ibrutinib



### Imbruvica® (ibrutinib)

- Irreversibly binds Bruton's tyrosine kinase, inhibiting the B-cell receptor pathway
- Proves covalent inhibition is a viable strategy for drug design
- In 2022, global Imbruvica net revenues were \$1.115 billion

Happy Cinco de Mayo!



...and why most people in Mexico don't celebrate

- Cinco de Mayo does not celebrate Mexico's independence!
- Mexican independence is September 16<sup>th</sup> (Huge holiday in Mexico!)
- Instead corresponds to Battle of Puebla against France in 1862
  - *"...the victory was short-lived—the French later captured Mexico City and took over the country..."*

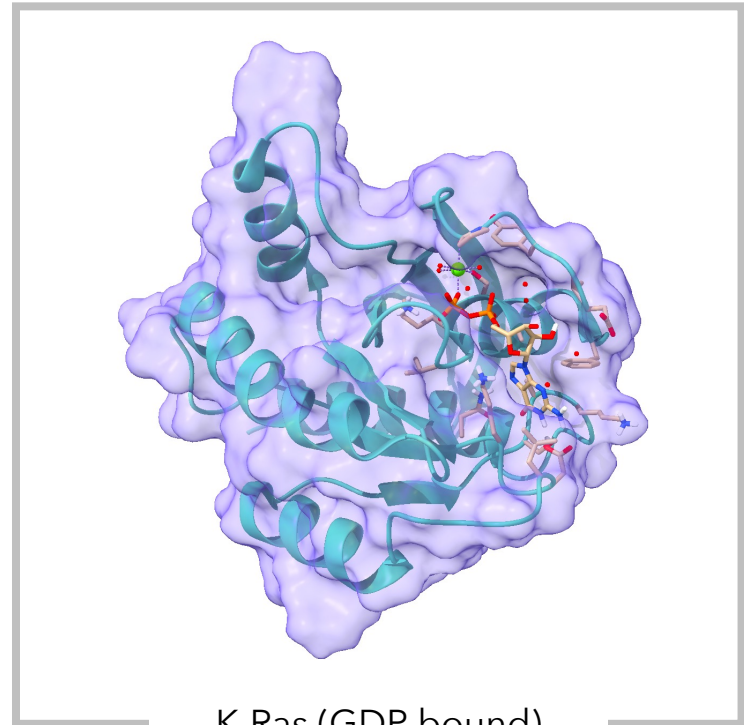


- Honestly, Cinco de Mayo is not really a thing in Mexico...
- Has become a much bigger thing in the US as a celebration of Mexican-American heritage
- Enjoy a margarita or some tacos today!

# Revisiting K-Ras and the undruggable proteome



Prof. Kevan Shokat  
UCSF



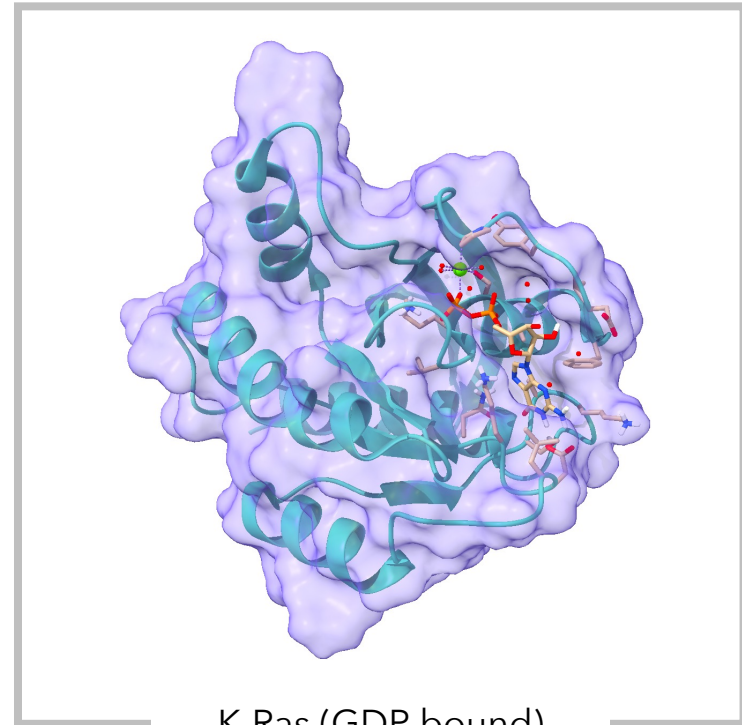
K-Ras (GDP bound)



# Revisiting K-Ras and the undruggable proteome



Prof. Kevan Shokat  
UCSF

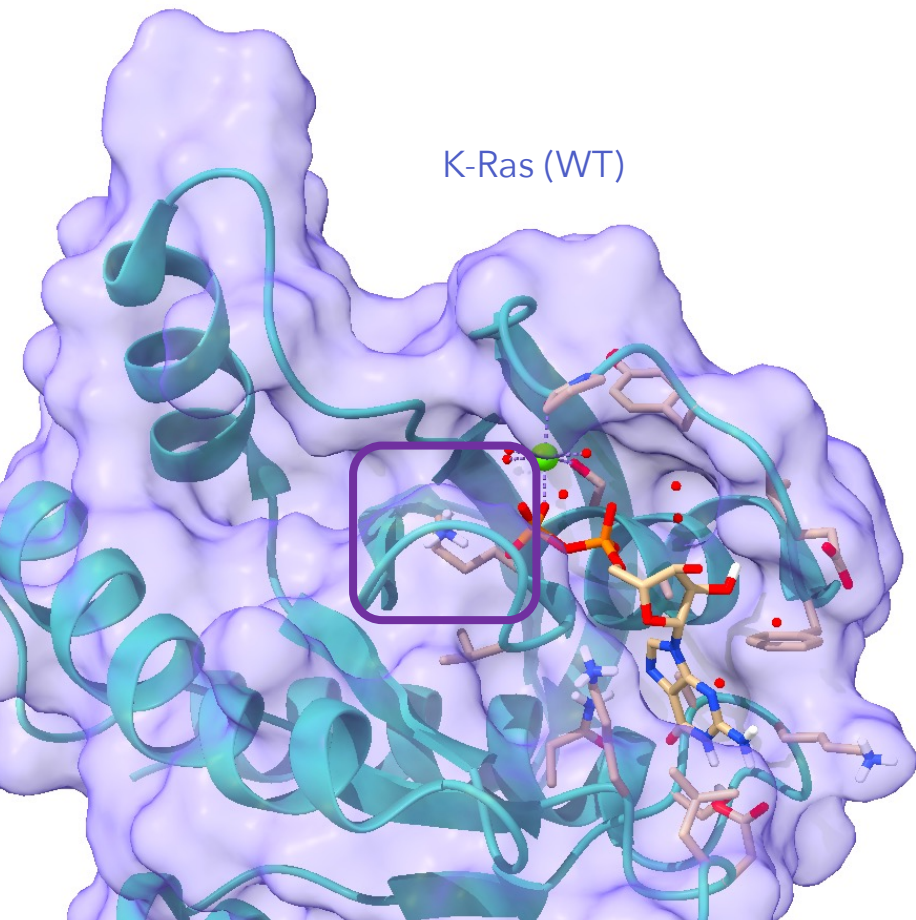


K-Ras (GDP bound)

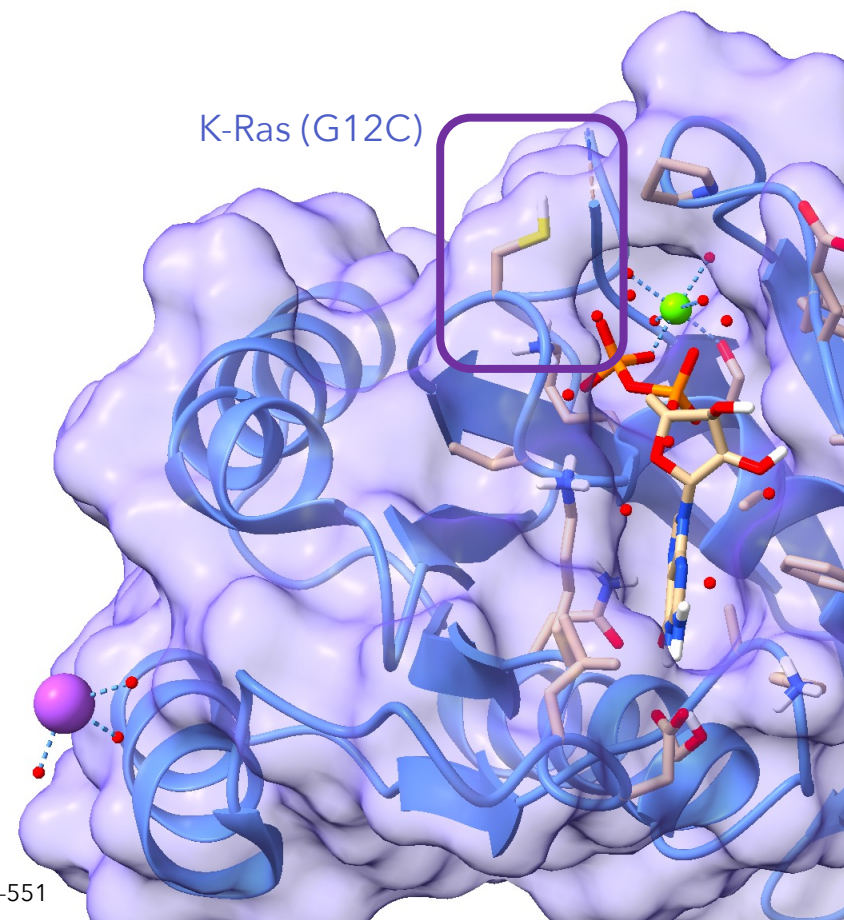
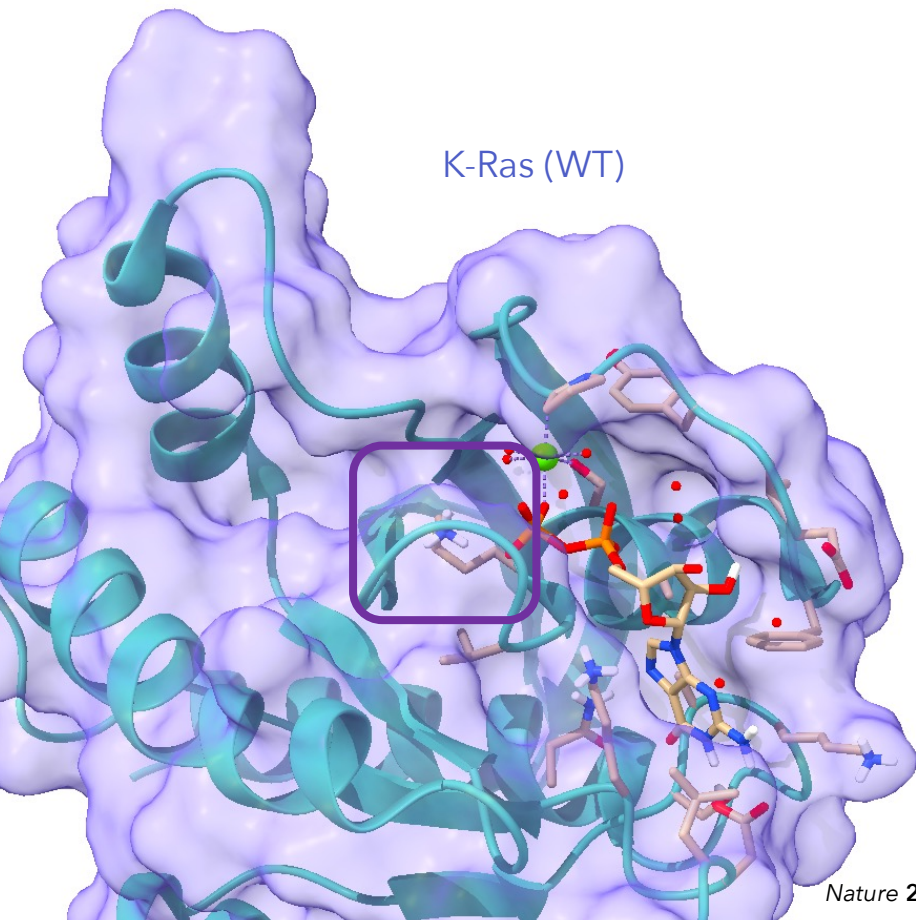
## Reasons K-Ras was deemed “undruggable”

- High affinity for native GTP substrate
- Inhibition of membrane localization is ineffective
- K-Ras involved in highly complex signaling pathway - difficult to understand how knocking out one protein affects downstream effects!

# Chemical strategies to drug K-Ras

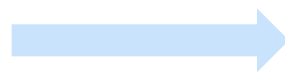


# Chemical strategies to drug K-Ras

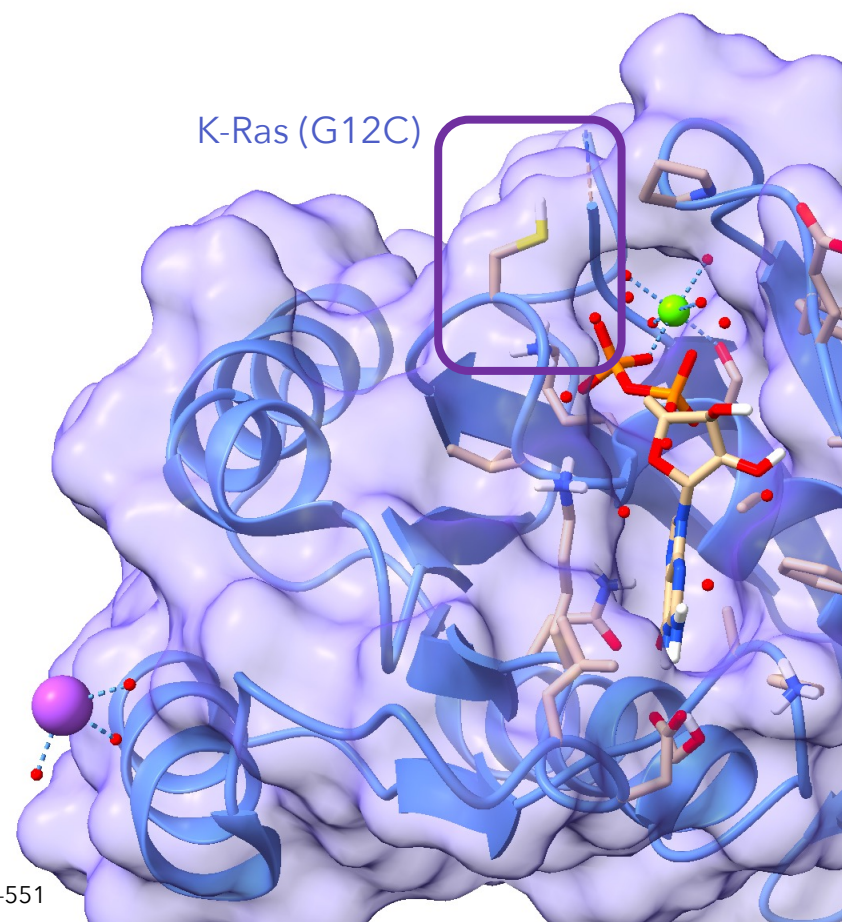
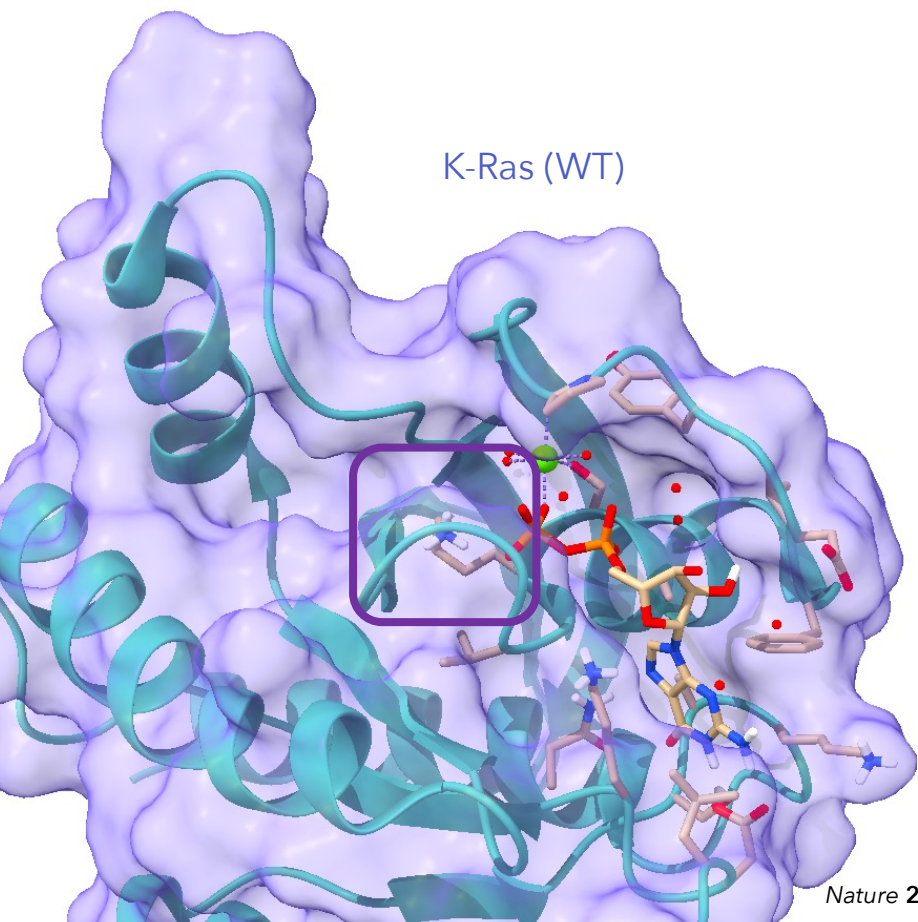


# Chemical strategies to drug K-Ras

**Strategy:** use cysteine-reactive functional groups in an ABPP-type approach to determine what ligands bind to K-Ras

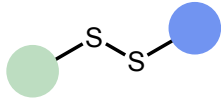


Determine allosteric sites while simultaneously only targeting mutated Ras owing to Cys12 reactivity



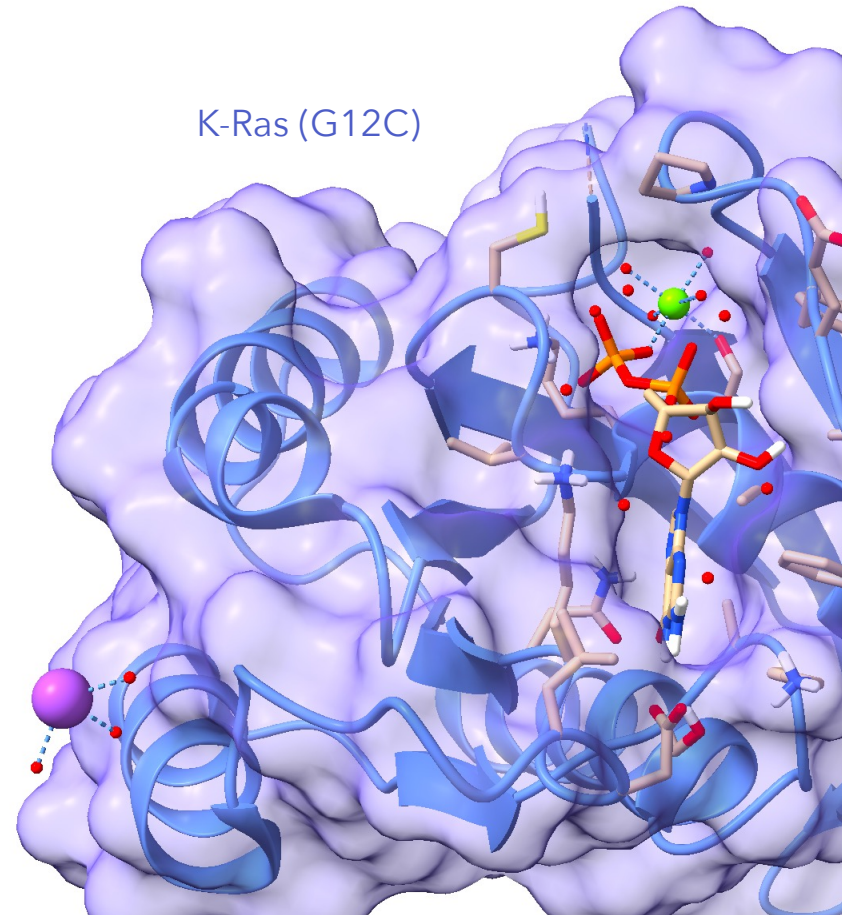
# Chemical strategies to drug K-Ras

Disulfide "tethering"



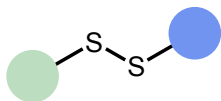
Determine allosteric sites while simultaneously only targeting mutated Ras owing to Cys12 reactivity

K-Ras (G12C)



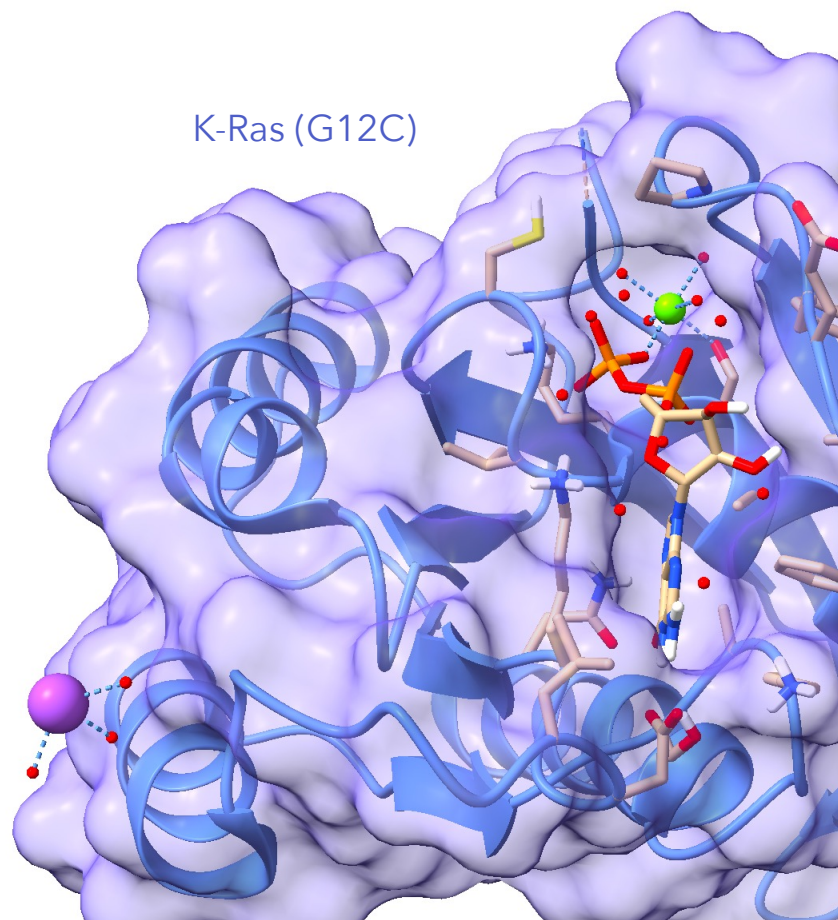
# Chemical strategies to drug K-Ras

Disulfide "tethering"



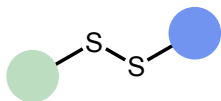
480 compounds  
screened

Determine allosteric sites  
while simultaneously  
only targeting mutated  
Ras owing to Cys12  
reactivity

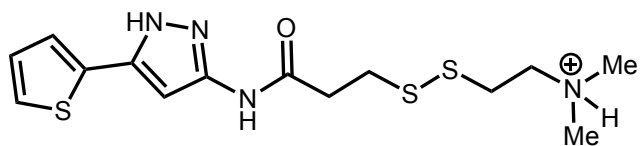


# Chemical strategies to drug K-Ras

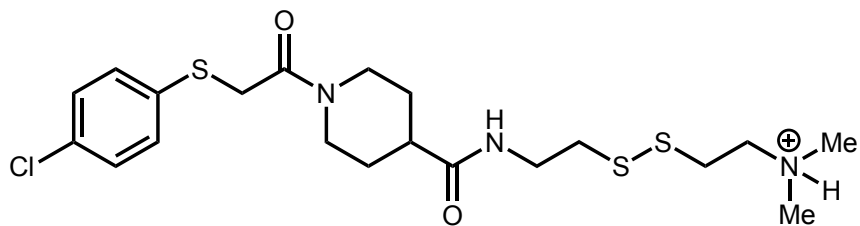
Disulfide "tethering"



480 compounds  
screened



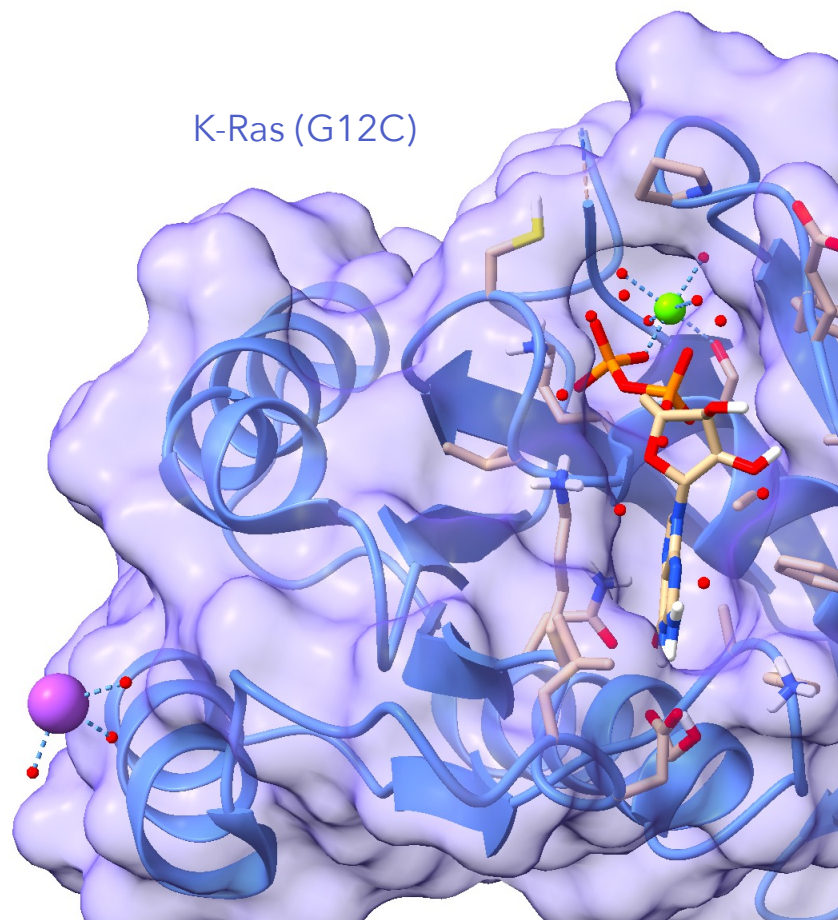
2E07



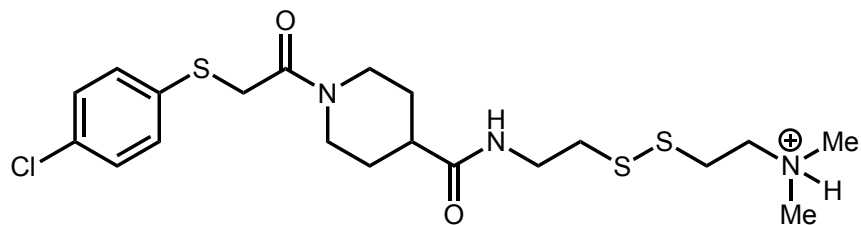
6H05

Determine allosteric sites  
while simultaneously  
only targeting mutated  
Ras owing to Cys12  
reactivity

K-Ras (G12C)



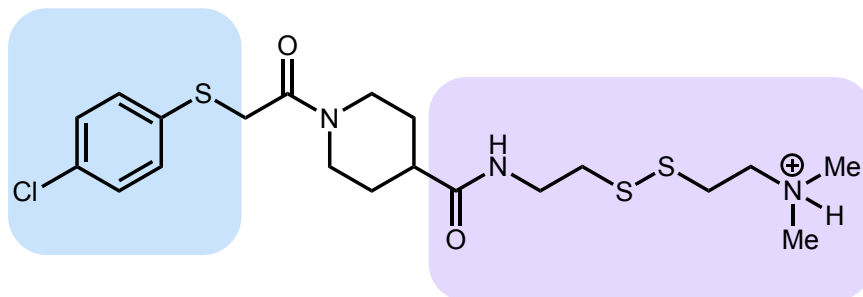
# A structure-activity campaign in understanding K-Ras inhibition



6H05



# A structure-activity campaign in understanding K-Ras inhibition



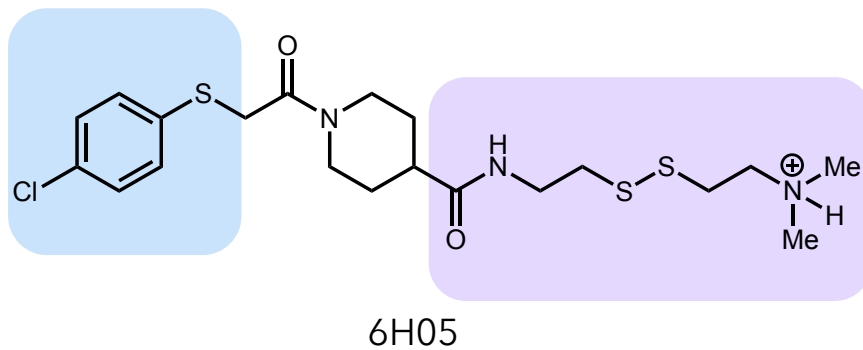
6H05

fragment A

fragment B

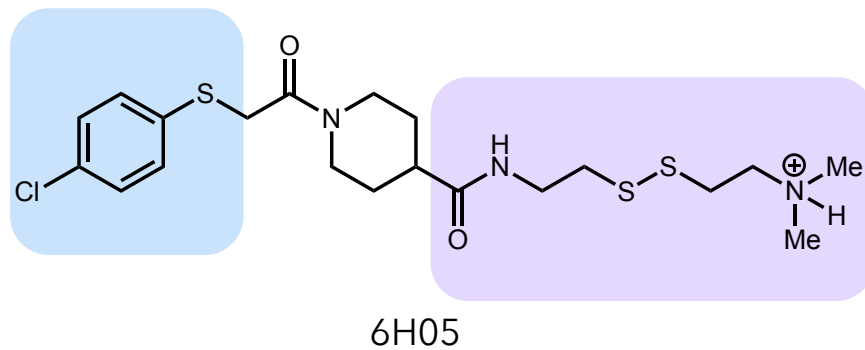
relative potency

# A structure-activity campaign in understanding K-Ras inhibition



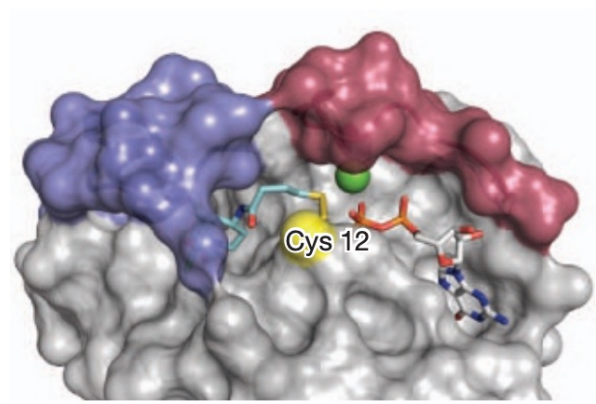
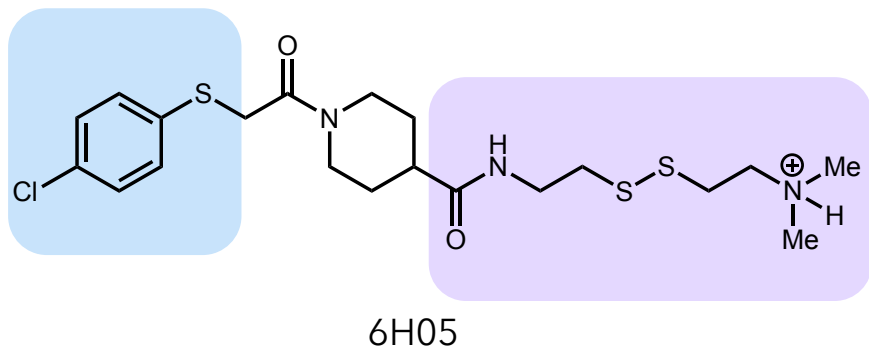
fragment A	fragment B	relative potency
		<0.1
		0.34

# A structure-activity campaign in understanding K-Ras inhibition

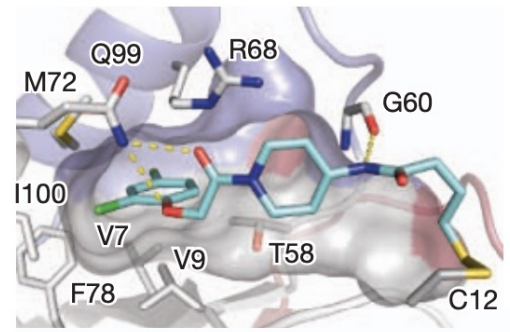
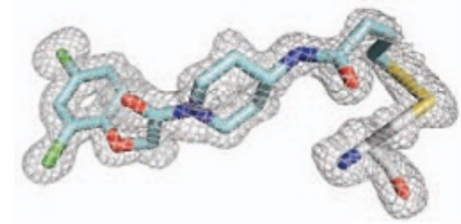


fragment A	fragment B	relative potency
		<0.1
		0.34
		1.2
		4.2

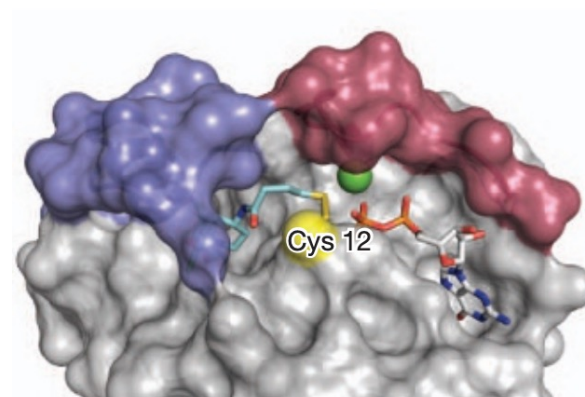
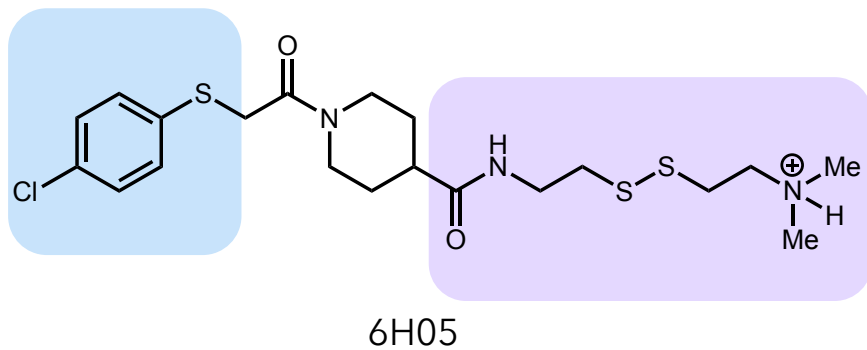
# A structure-activity campaign in understanding K-Ras inhibition



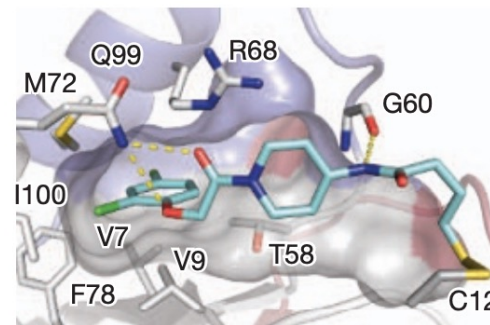
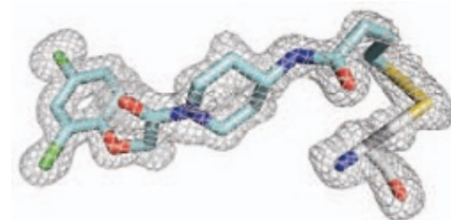
fragment A	fragment B	relative potency
		<0.1
		0.34
		1.2
		4.2



# A structure-activity campaign in understanding K-Ras inhibition

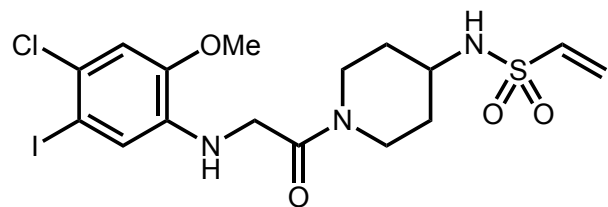


fragment A	fragment B	relative potency
		<0.1
		0.34
		1.2
		4.2

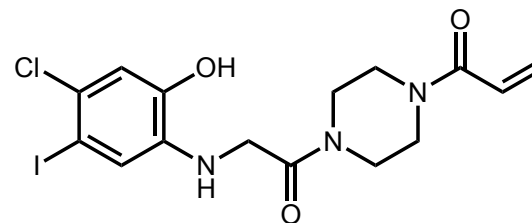


Activity-based approach reveals allosteric "Switch-II" pocket

## Switching to irreversible activity-based probes

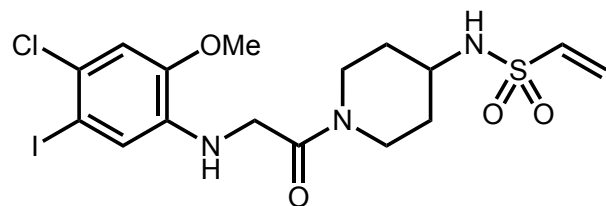


compound 9

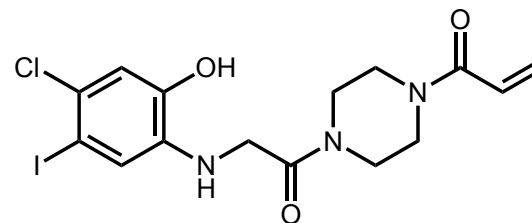


compound 12

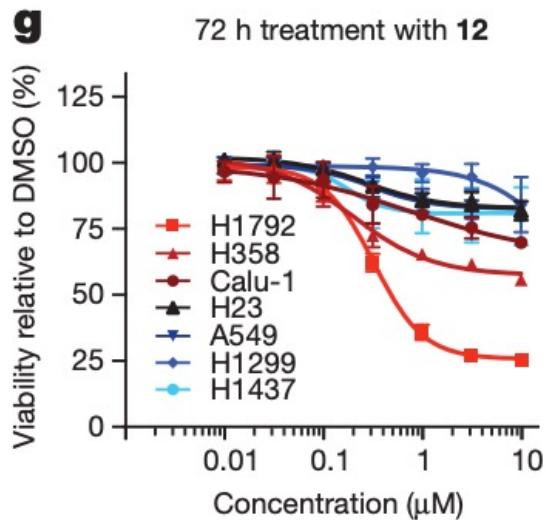
## Switching to irreversible activity-based probes



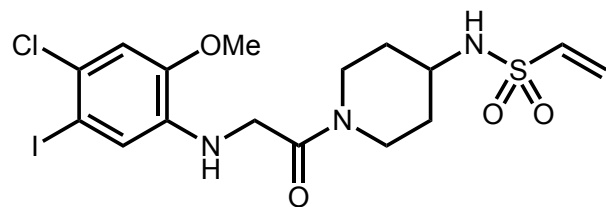
compound 9



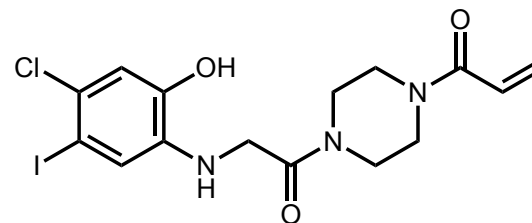
compound 12



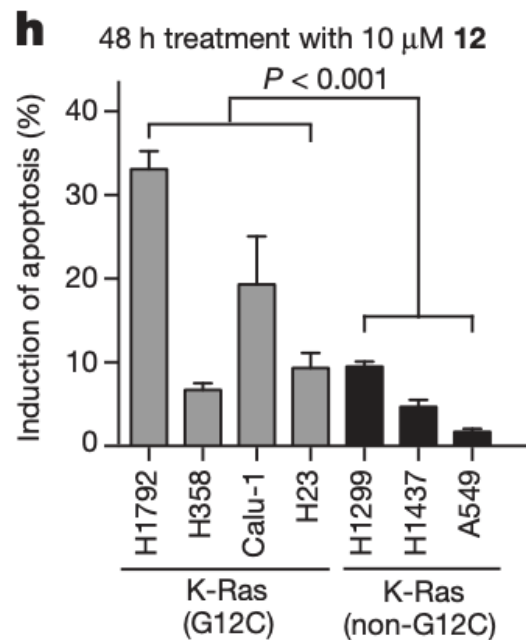
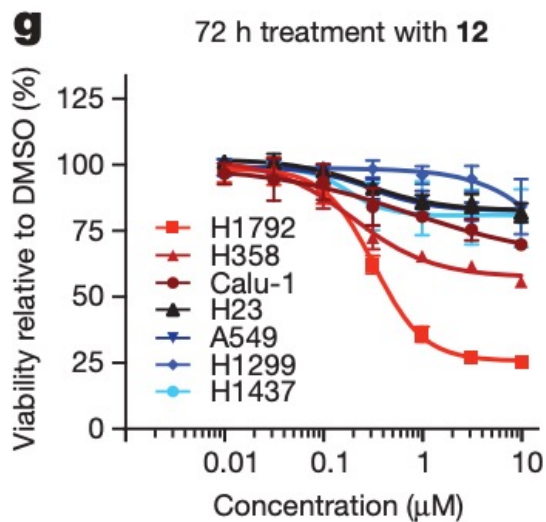
# Switching to irreversible activity-based probes



compound 9

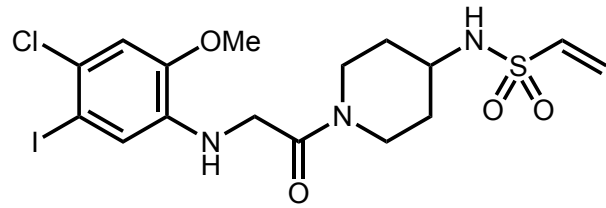


compound 12

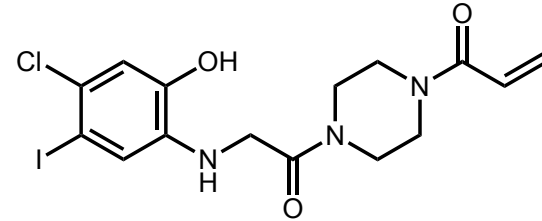




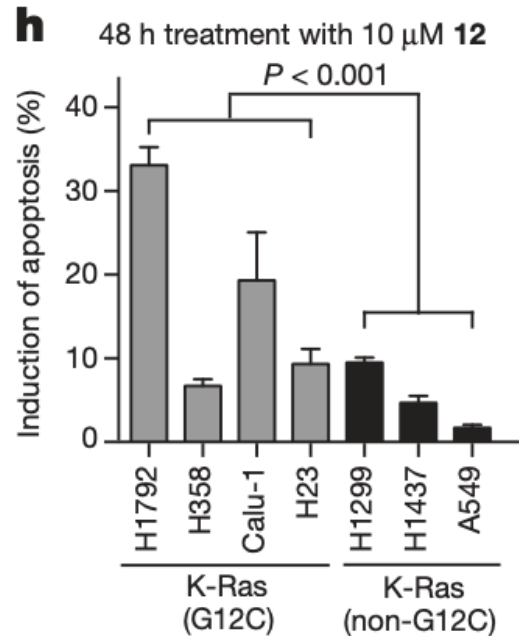
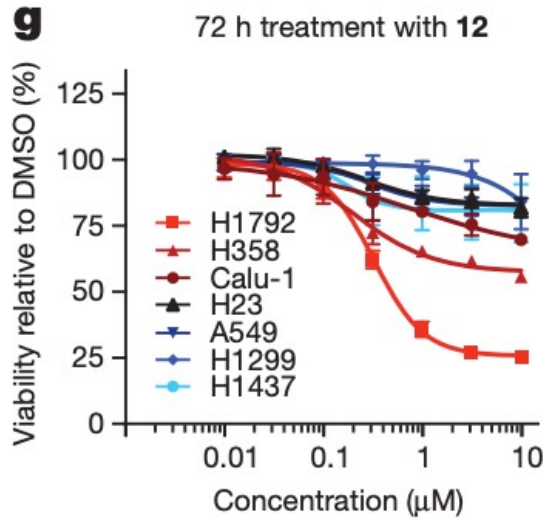
# Switching to irreversible activity-based probes



compound 9



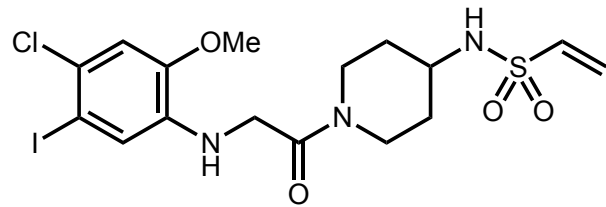
compound 12



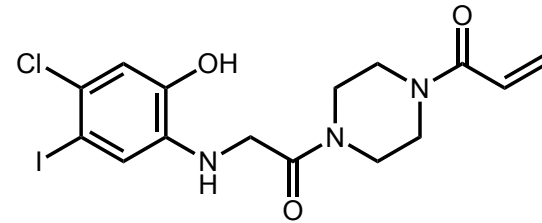
Compound 12 served as an efficient covalent, irreversible inhibitor of K-Ras G12C

Insights gained from activity-based profiling

## Switching to irreversible activity-based probes



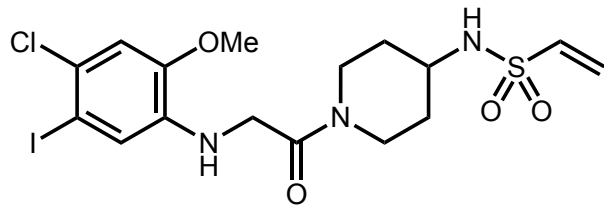
compound 9



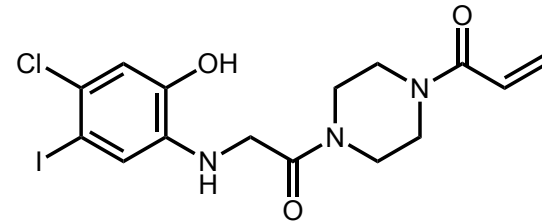
compound 12

**AMGEN**

## Switching to irreversible activity-based probes



compound 9



compound 12

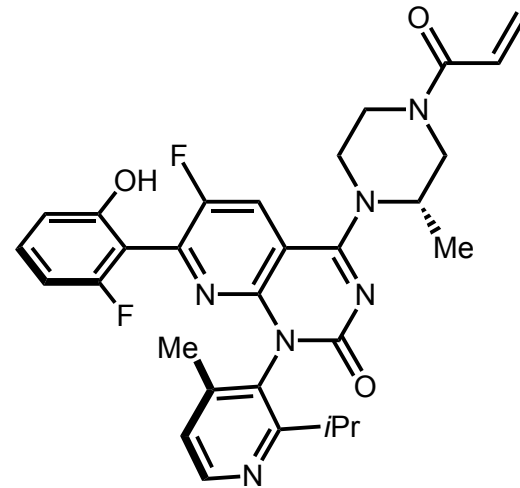


**AMGEN**

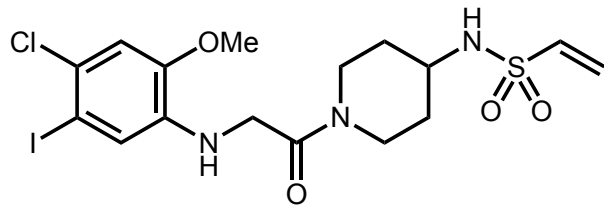
**Sotorasib**

FDA approval in 2021

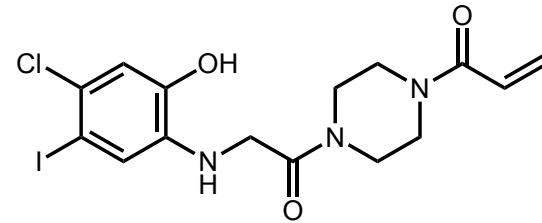
Treatment of K-Ras G12C  
mutations



## Switching to irreversible activity-based probes



compound 9



compound 12



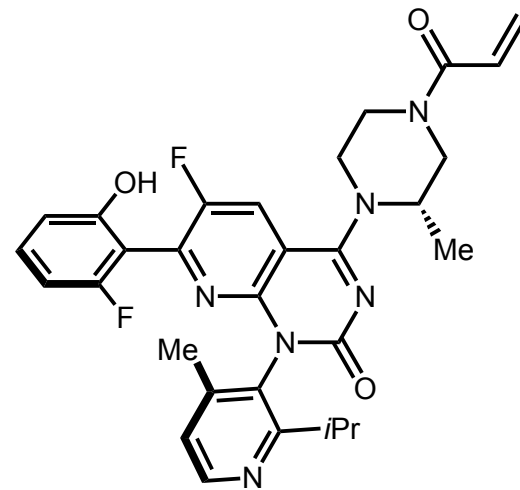
**AMGEN**

**Sotorasib**

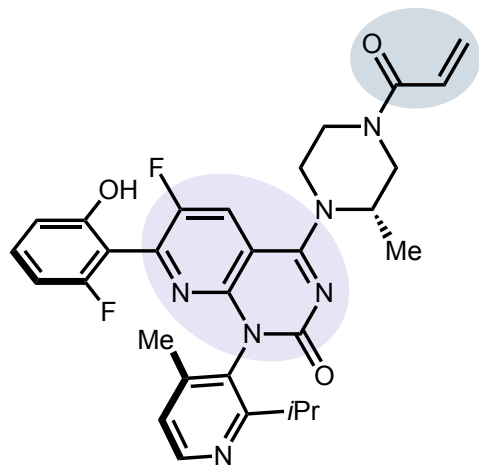
FDA approval in 2021

Treatment of K-Ras G12C  
mutations

Successful drugging on an  
"undruggable" oncogene!

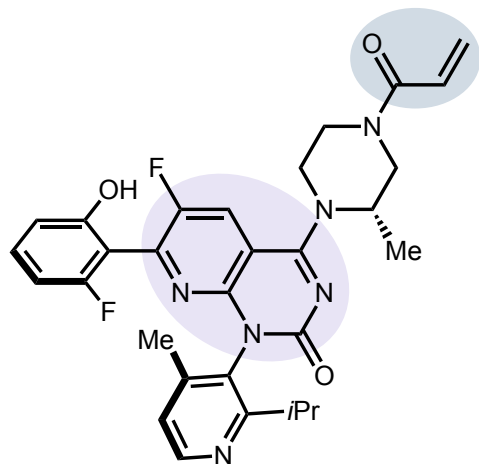


## Drugging the “undruggable” K-Ras G12 mutations



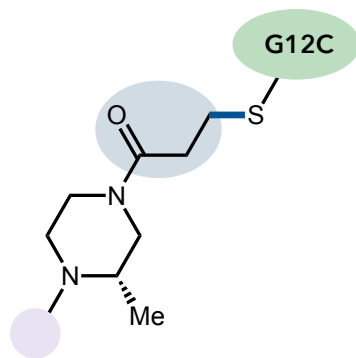
Sotorasib

# Drugging the “undruggable” K-Ras G12 mutations

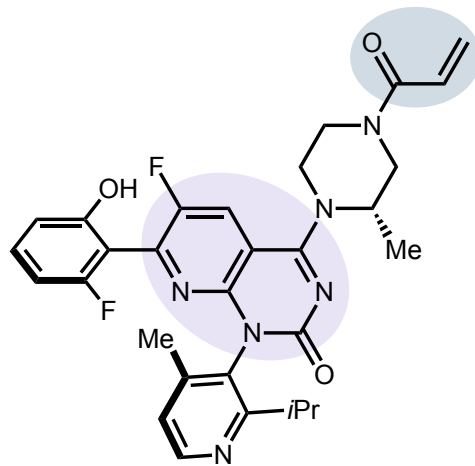


**Sotorasib**

FDA approval in 2021

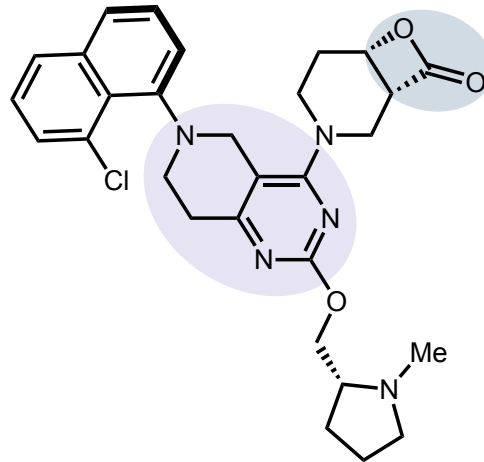
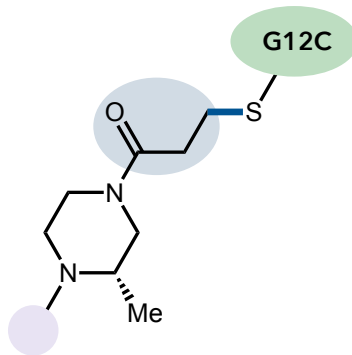


# Drugging the “undruggable” K-Ras G12 mutations

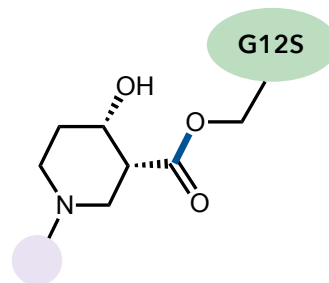


**Sotorasib**

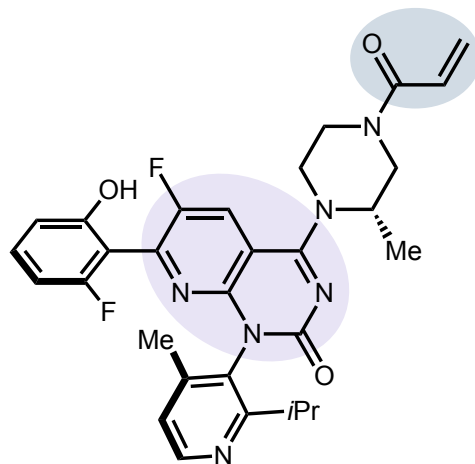
FDA approval in 2021



Shokat (2022)

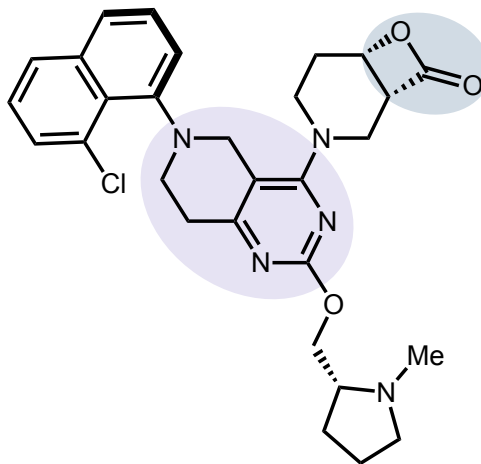
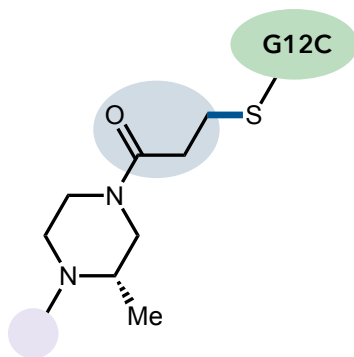


# Drugging the “undruggable” K-Ras G12 mutations

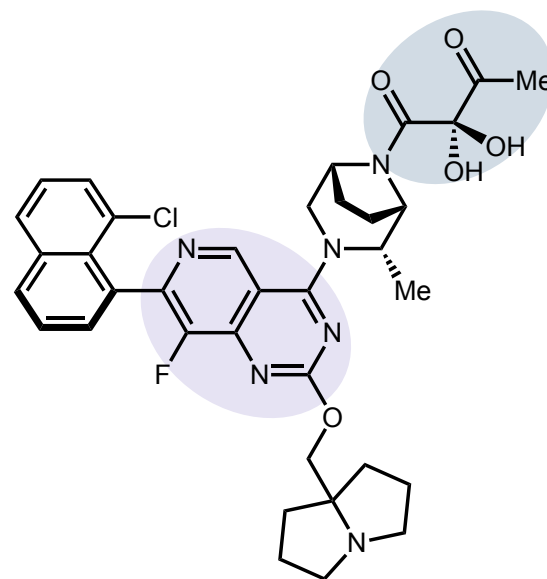
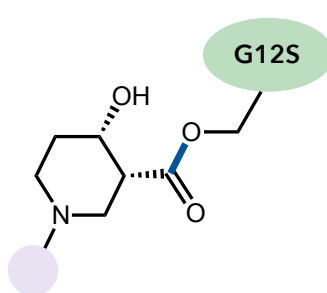


Sotorasib

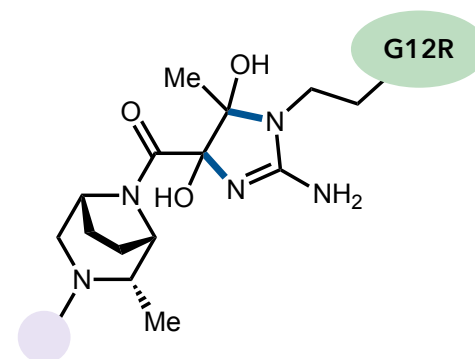
FDA approval in 2021



Shokat (2022)

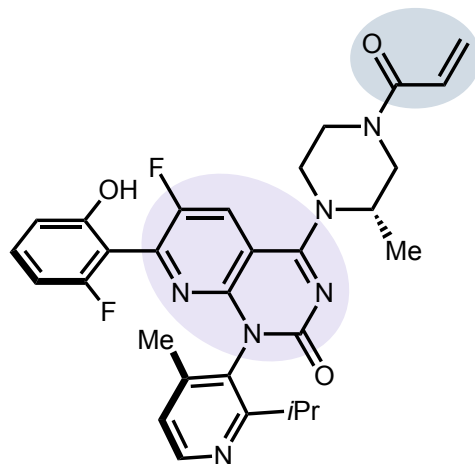


Shokat (2022)



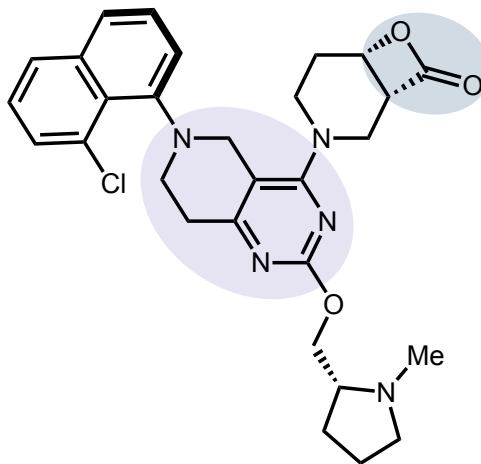


# Drugging the “undruggable” K-Ras G12 mutations

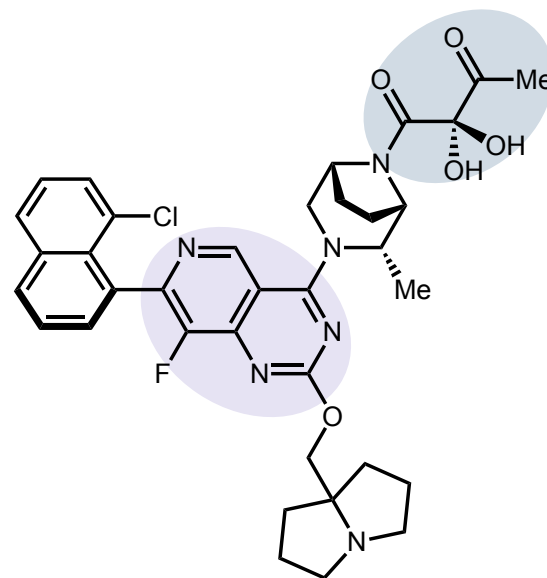


**Sotorasib**

FDA approval in 2021

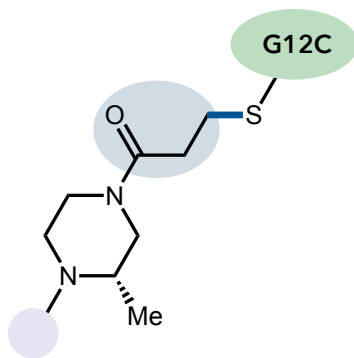


Shokat (2022)

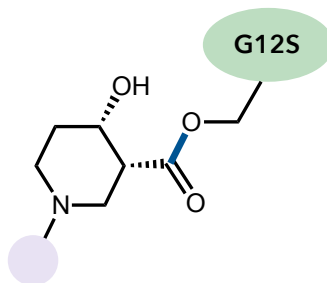


Shokat (2022)

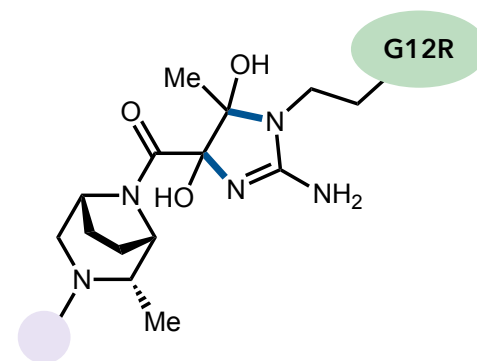
Successful K-Ras inhibitors dependent on covalent inhibition



G12C

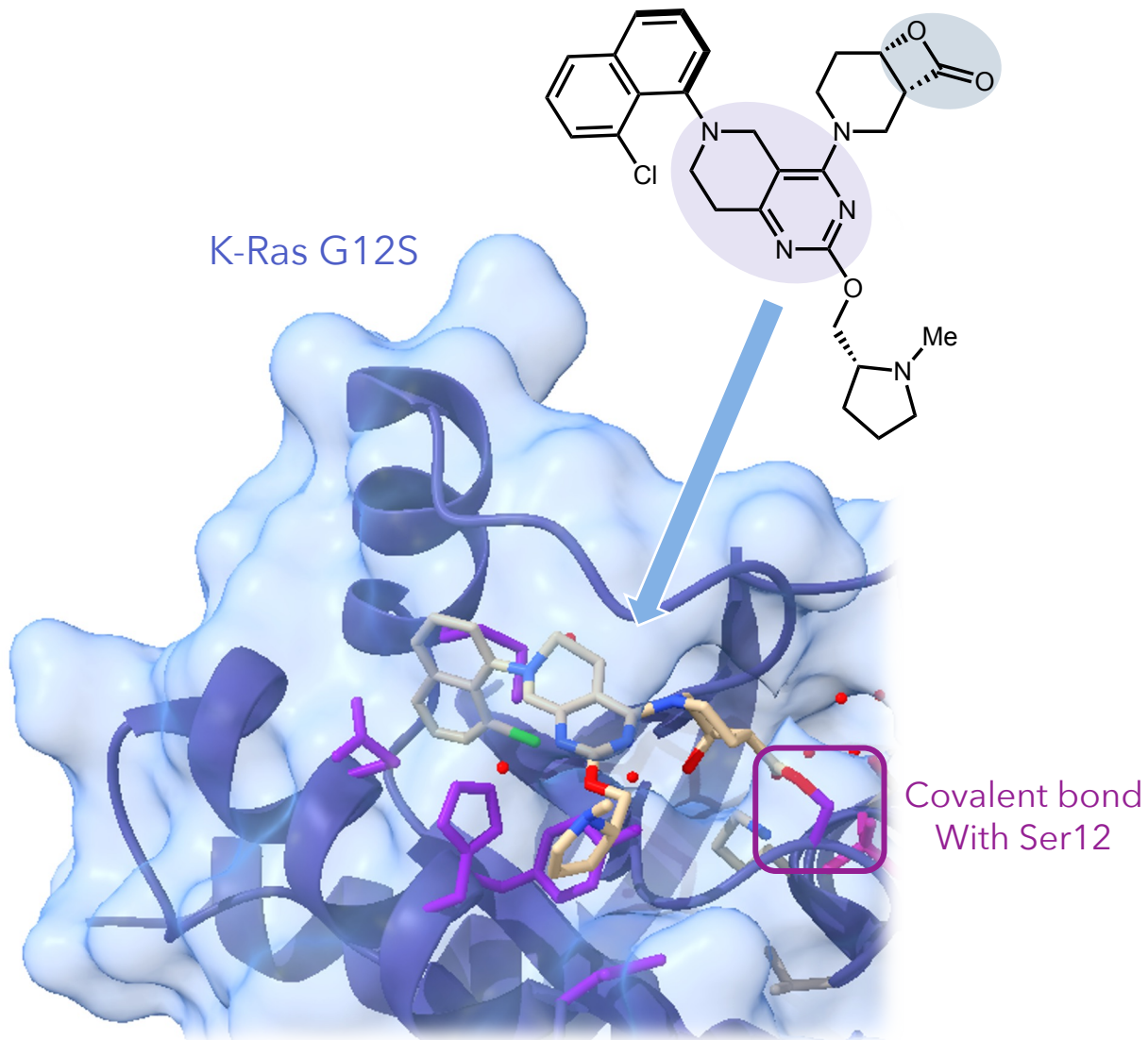


G12S

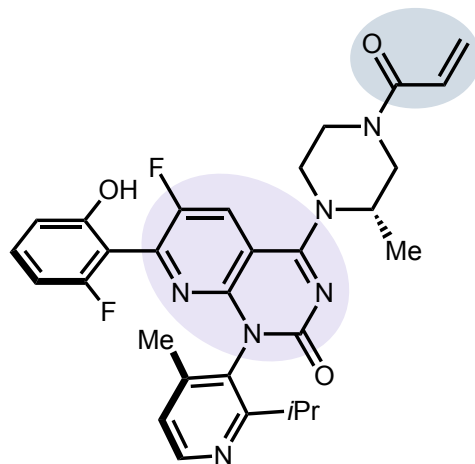


G12R

The switch-II pocket is flexible and inhibitors dock and covalently bind

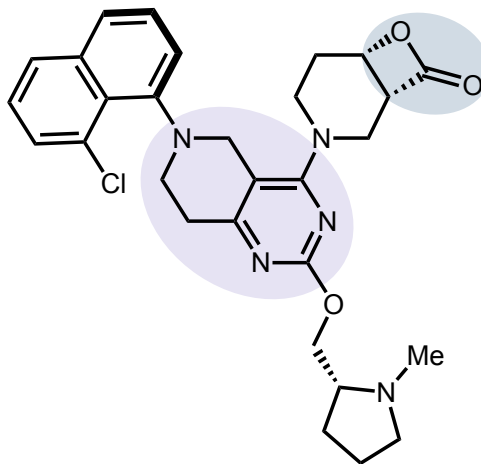


# Expanding beyond G12C with different covalent strategies

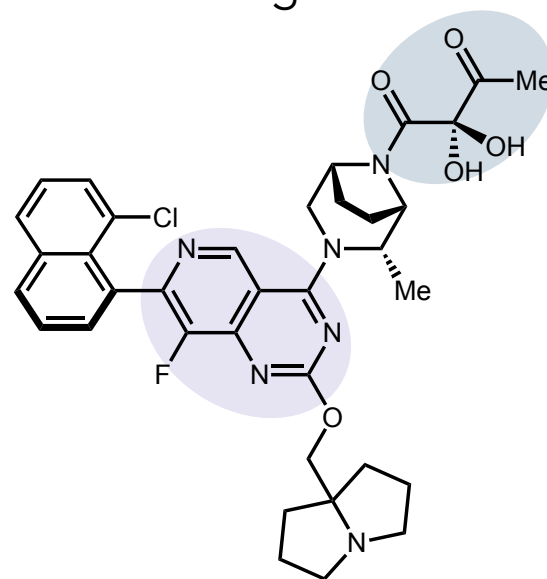


**Sotorasib**

FDA approval in 2021

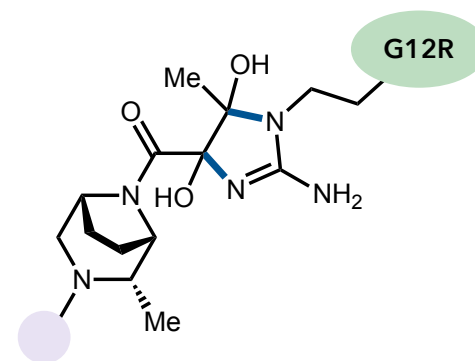
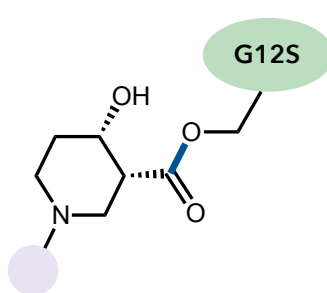
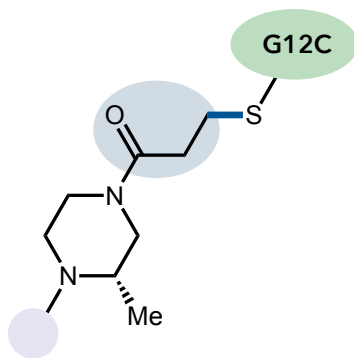


Shokat (2022)



Shokat (2022)

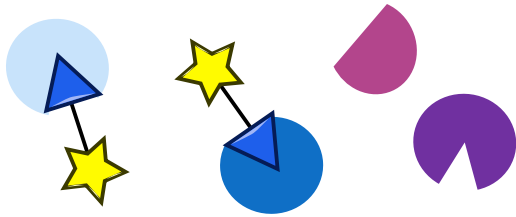
Successful K-Ras inhibitors dependent on covalent inhibition



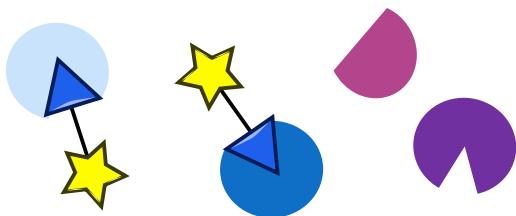
# Outline

- Introduction to undruggable proteins
  - What makes a protein “undruggable”?
  - Attempts to drug K-Ras mutations
- Activity-based approaches to finding “druggable” sites
- Success stories in covalent drugs
  - Ibrutinib and Bruton’s tyrosine kinase
  - Sotorasib and K-Ras G12C
- Conclusions
  - “Yet to be drugged” instead of “undruggable”

**Activity-based protein profiling** approaches and the advancement of proteomics have allowed for identification of variety of new covalent candidates and druggable sites



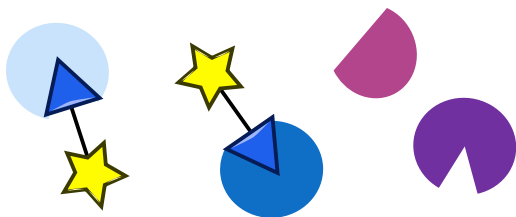
**Activity-based protein profiling** approaches and the advancement of proteomics have allowed for identification of variety of new covalent candidates and druggable sites



Analysis of **drug toxicity** concerns reveals that toxicity is mostly correlated to dosage, not mechanism of binding or structural alerts



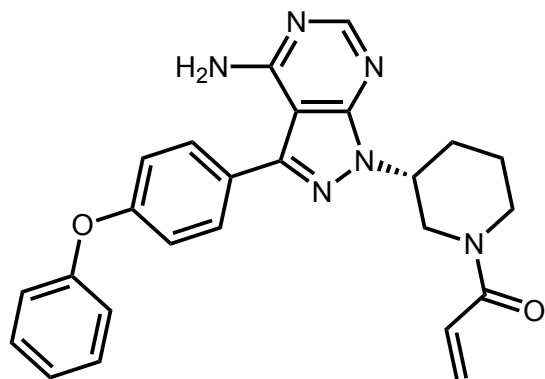
**Activity-based protein profiling** approaches and the advancement of proteomics have allowed for identification of variety of new covalent candidates and druggable sites



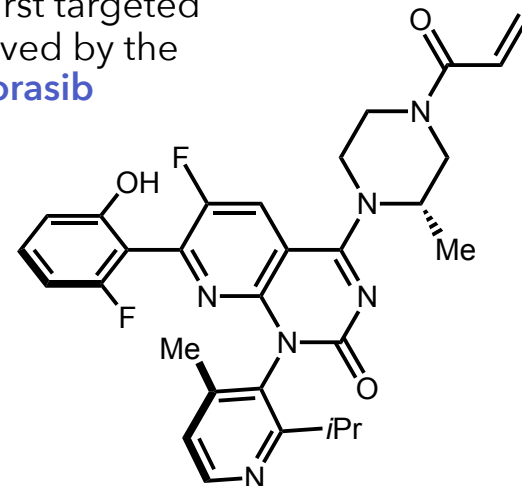
Analysis of **drug toxicity** concerns reveals that toxicity is mostly correlated to dosage, not mechanism of binding or structural alerts



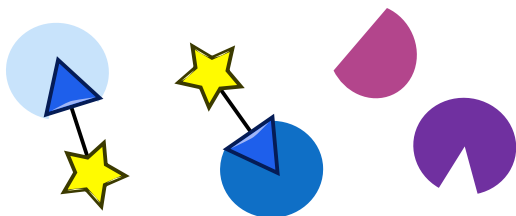
**Ibrutinib** represented a “behind-the-scenes” development of a **covalent inhibitor** that made its way into the market



Activity-based approaches allowed for **identification of a druggable site on K-Ras**, leading to the first targeted therapy approved by the FDA, **sotorasib**



**Activity-based protein profiling** approaches and the advancement of proteomics have allowed for identification of variety of new covalent candidates and druggable sites

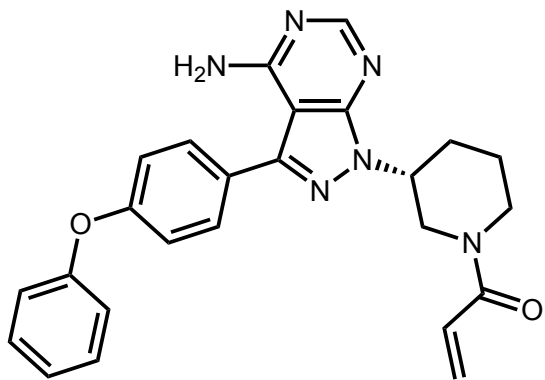


Analysis of **drug toxicity** concerns reveals that toxicity is mostly correlated to dosage, not mechanism of binding or structural alerts

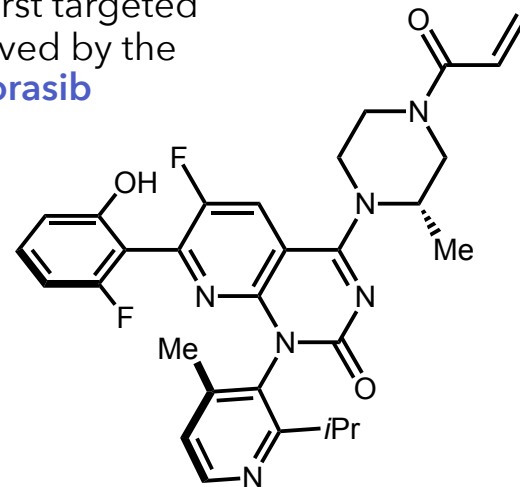


## Chemical strategies to drug "undruggable" proteins

**Ibrutinib** represented a "behind-the-scenes" development of a **covalent inhibitor** that made its way into the market



Activity-based approaches allowed for **identification of a druggable site on K-Ras**, leading to the first targeted therapy approved by the FDA, **sotorasib**





Thank you!

