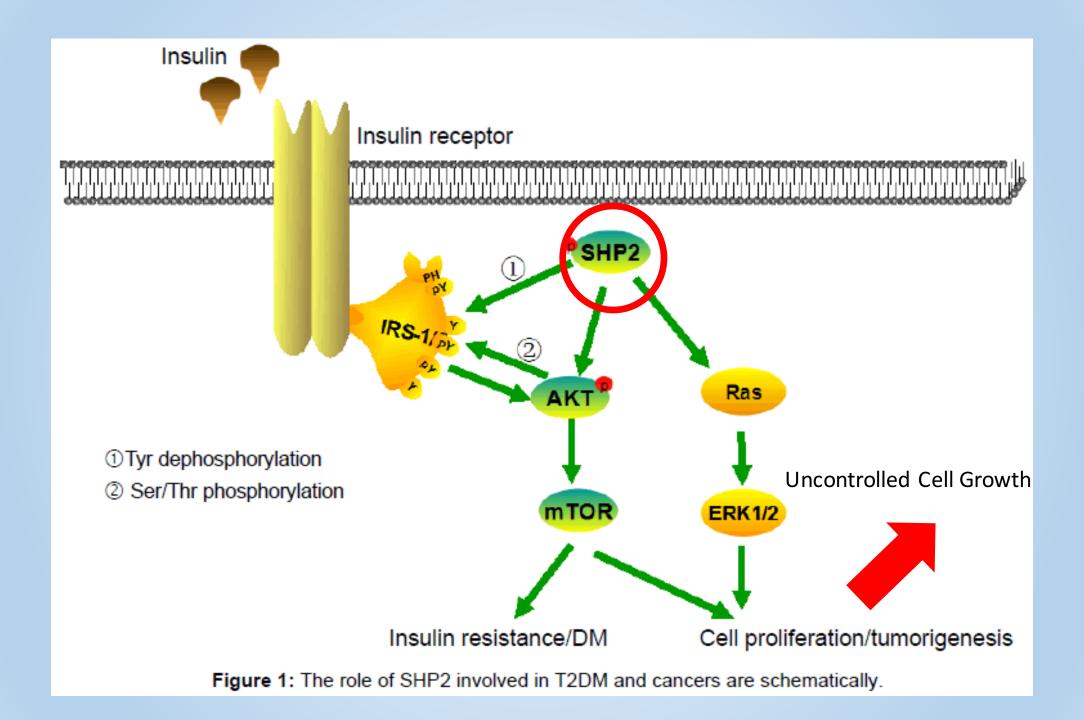
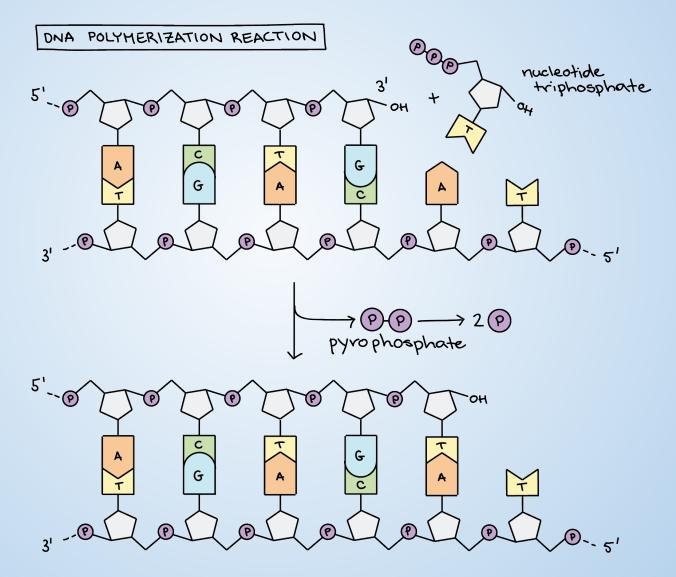


### What is Cancer?

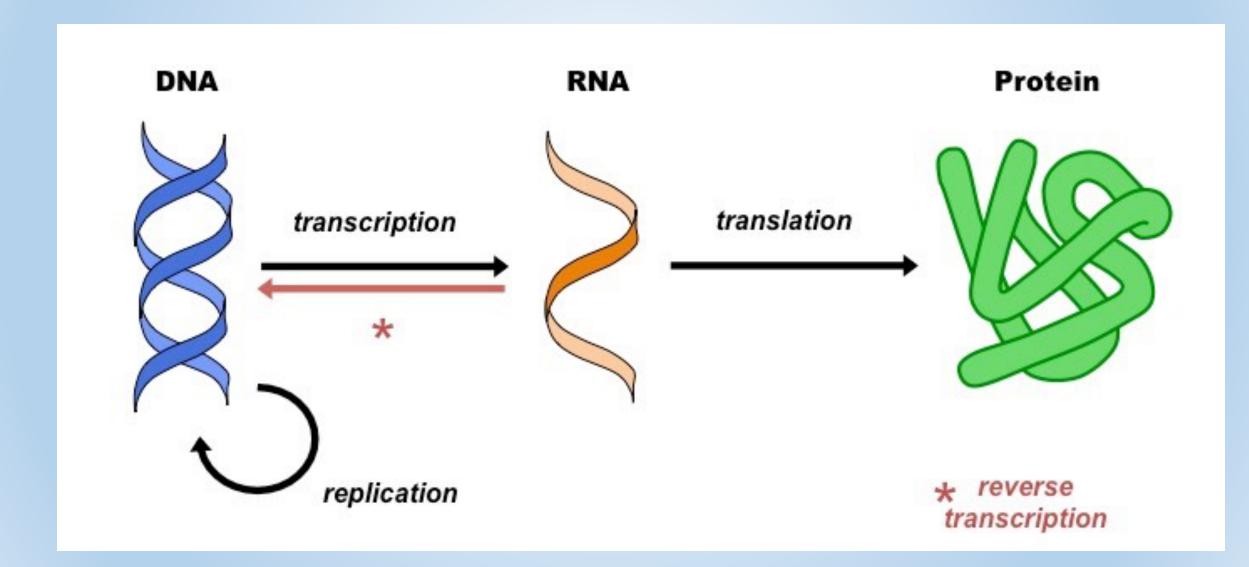
Uncontrollable Cell Growth



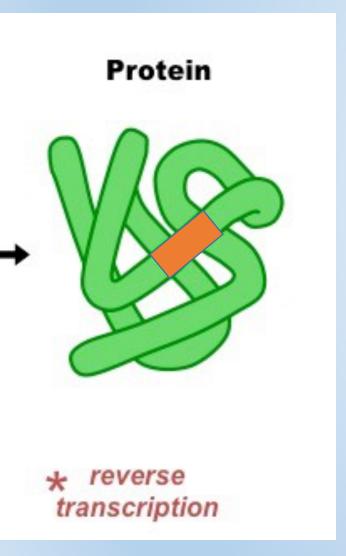
### What Causes Mutations?



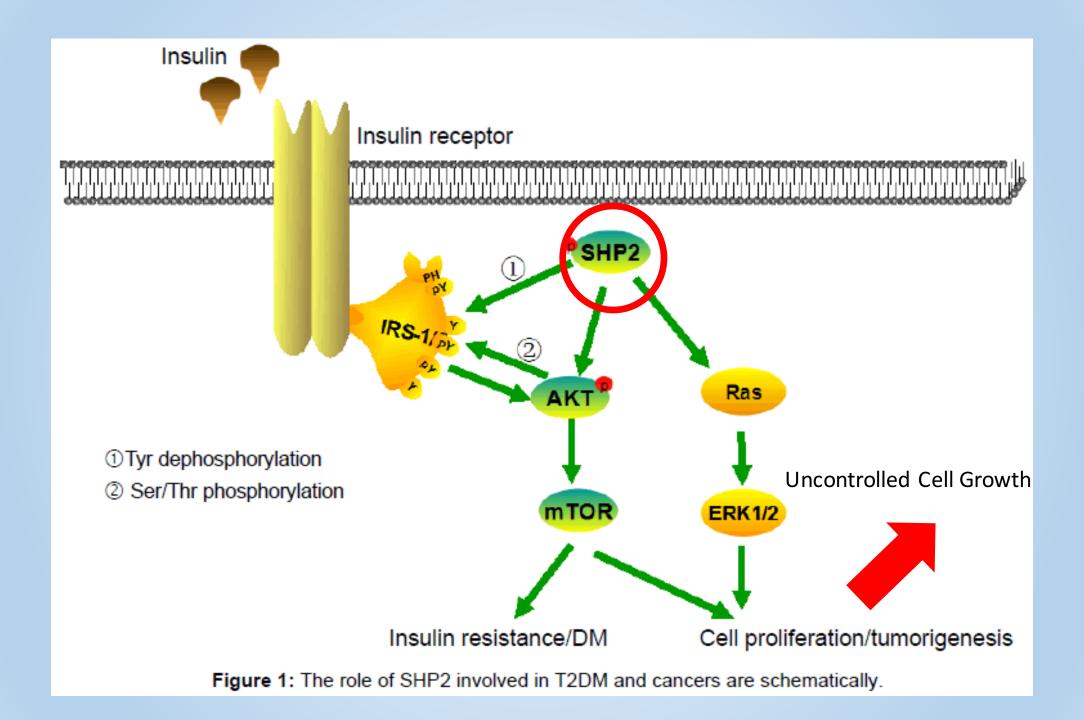
### Central Dogma



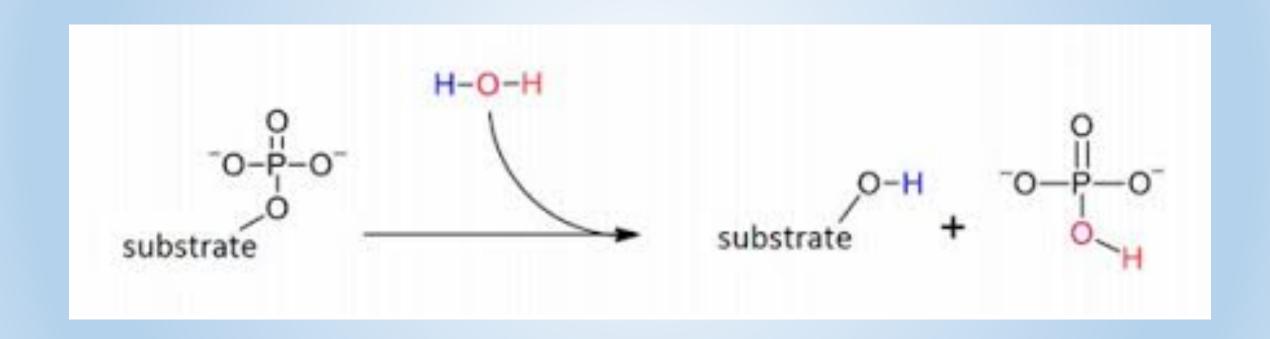
### Protein Mutation:



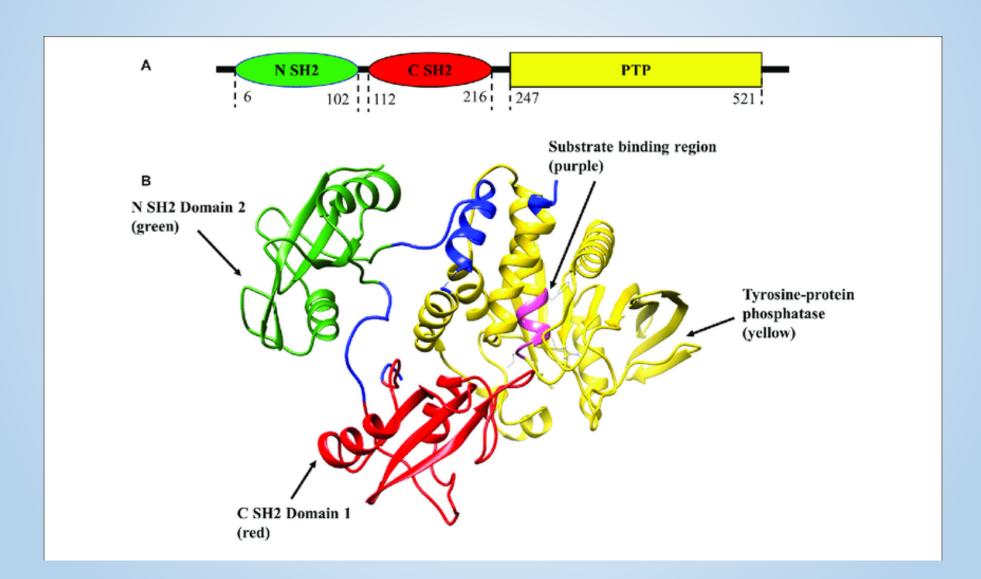
- May lead to new biochemical properties
- May loose ability for regulation



### What's Happening at the Chemical Level?



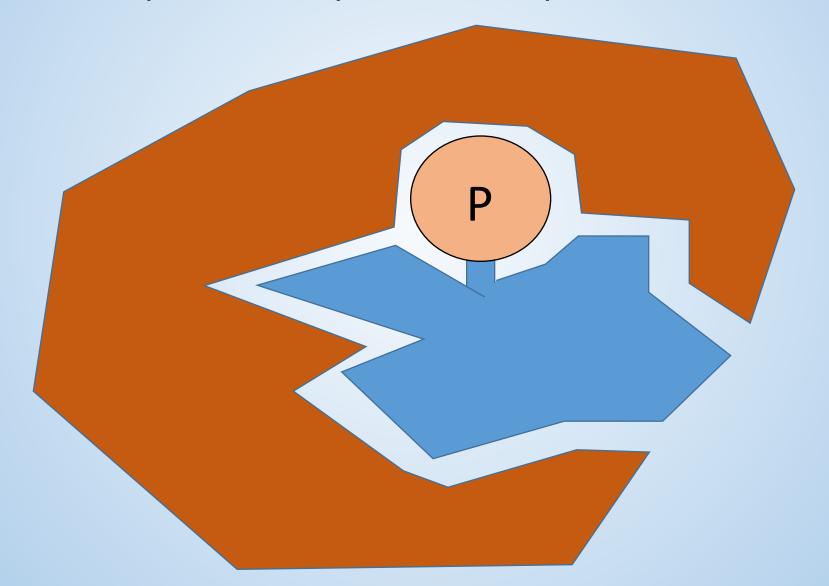
### Regulation in Phosphatases



How do we stop constitutive activation?

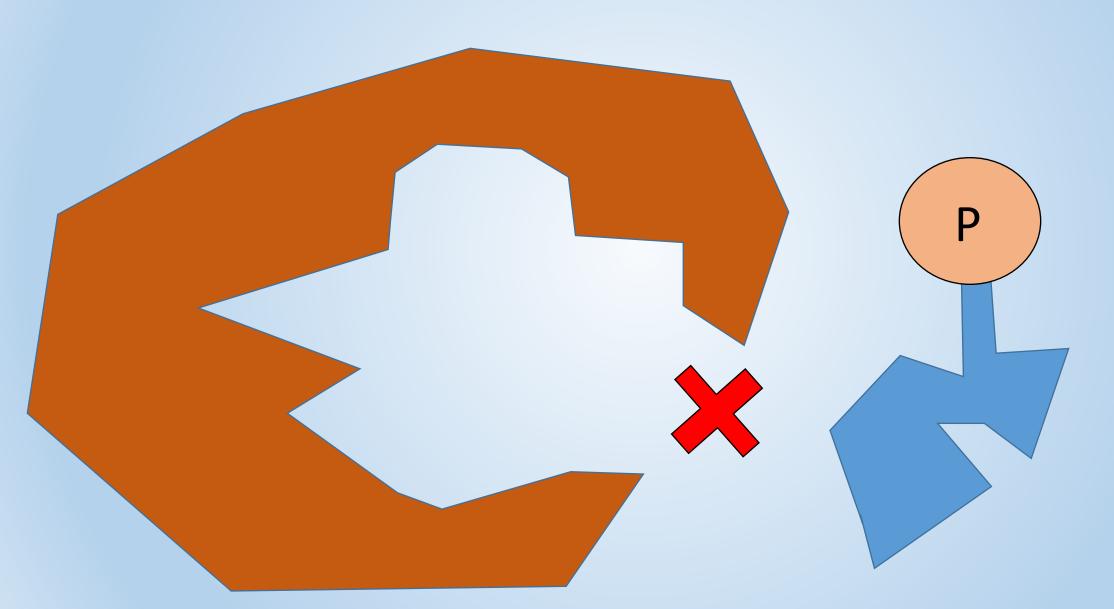
Drugs!

## Substrate Specificity of Phosphatases



# Substrate Specificity of Phosphatases

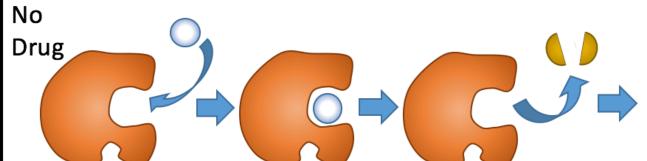
### Substrate Specificity of Phosphatases



# Making a Drug!



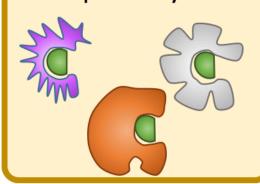
### 1A: Traditional Active Site Targeting



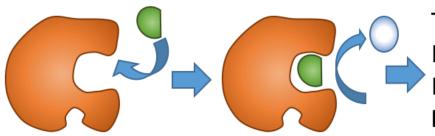
Excessive tyrosine phosphorylation: Uncontrolled cell growth

The Problem With Active Site Targeting:

PTP active sites cannot be targeted with specificity!



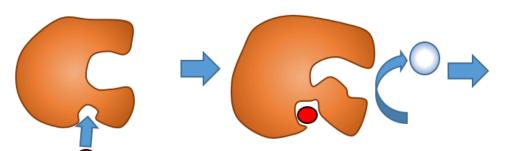
With Traditional Drug



Tyrosine blocked from PTP binding,
No phosphorylation,
No uncontrolled growth

### 1B: Allosteric Site Targeting

With Allosteric Inhibitor



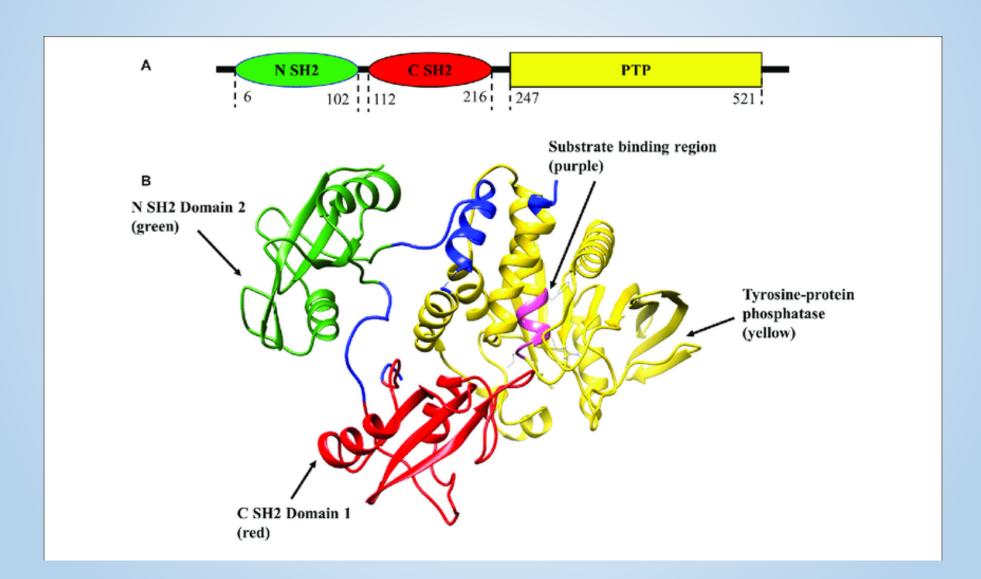
PTP active site unable to bind tyrosine,
No phosphorylation,
No uncontrolled growth
Other PTPs unaffected

# Targeting a Cryptic Allosteric Site for Selective Inhibition of the Oncogenic Protein Tyrosine Phosphatase Shp2

Cynthia M. Chio, Christopher S. Lim, and Anthony C. Bishop\*

Protein tyrosine phosphatases (PTPs) have been the subject of considerable pharmaceutical-design efforts because of the ubiquitous connections between misregulation of PTP activity and human disease. PTPinhibitor discovery has been hampered, however, by the difficulty in identifying cell-permeable compounds that can selectively target PTP active sites, and no PTP inhibitors have progressed to the clinic. The identification of allosteric sites on target PTPs therefore represents a potentially attractive solution to the druggability problem of PTPs. Here we report that the oncogenic PTP Shp2 contains an allostericinhibition site that renders the enzyme sensitive to potent and selective inhibition by cell-permeable biarsenical compounds. Because Shp2 contains no canonical tetracysteine biarsenical-binding motif, the enzyme's inhibitor-binding site is not readily predictable from its primary or three-dimensional structure. Intriguingly, however, Shp2's PTP domain does contain a cysteine residue (C333) at a position that is removed from the active site and is occupied by proline in other classical PTPs. We show that Shp2's unusual cysteine residue constitutes part of a Shp2-specific allosteric-inhibition site, and that Shp2's consistivity to himmonicals is demandant on the massage of the notiveally accomming C222. The determinative

### Regulation in Phosphatases



# For Research Confidentiality Reasons I Can't Share the Experimental data with you.

Below is link to paper I talked about in class:

https://www.ncbi.nlm.nih.gov/pmc/articles/PMC4303306/

Allosteric Site-Directed Drugs are the Future

Thank You