

Building a Natural Product-inspired Macrocyclic Toolbox Platform for “Undruggable” Targets

Prabhat Arya

Founder and Chief Executive Officer, SignMod

www.signmod.org

Prabhat.arya@signmod.org



External

The information contained in this document is for the personnel (and the Organization) being contacted by SignMod only. Circulation or reproduction of this document is not permitted without seeking permission.

Confidential

The information contained in this document is proprietary and confidential. If you are not the intended recipient, please note that any use or circulation of this document is not permitted and may be cause for legal action.

Prabhat Arya

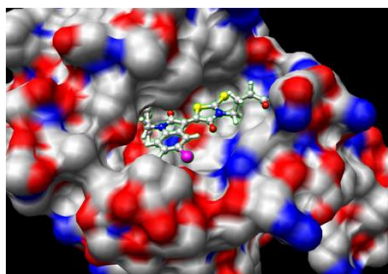
SignMod is building a highly unique and proprietary, natural product-inspired macrocyclic chemical toolbox platform for tackling challenging targets, currently defined as “undruggable”. These targets generally involve multiple protein-protein, DNA/RNA-protein-protein interactions and are difficult to handle by conventional, small molecule-based approaches. Our current focus is on cancer targets (and pathways), related to Wnt, c-Myc and K-Ras.

With the continual growth in the complexity of clinically-relevant cancer targets, there is a severe paucity in the small molecule arena having compounds capable of modulating protein-protein, DNA/RNA-protein interactions. Natural products have a proven track record when it comes to modulating protein-protein, DNA/RNA-protein interactions by small molecules. Due to several challenges that are associated with them, we are developing novel methods leading to macrocyclic compounds, classified as “natural product-inspired”. As with most macrocyclic natural products, these compounds have 3D shapes and are rich in sp³ centers. The presence of a large cyclic ring in their architectures also provide the possibility of interacting with the multiple sites. Due to the sufficient complexity of our compounds and an ease of the practical synthesis, they are highly attractive for further developing medicinal chemistry programs.

Build a proprietary, natural product-inspired macrocyclic toolbox for challenging “undruggable” targets that are difficult to handle by traditional small molecules.

Utilize this toolbox in finding novel functional small molecules for protein-protein, DNA/RNA-protein interactions, and in general, for targets related to Transcription and Translation machinery.

Classical Approaches: Simplistic View



Structural information on the target; finding the pocket

Lipinski's rules

Structural guided chem /med chem

Defining the target (could be enzymes or isolated protein(s))

Science (2003)

REVIEW

Assembly of Cell Regulatory Systems Through Protein Interaction Domains

Tony Pawson^{1,2*} and Piers Nash¹

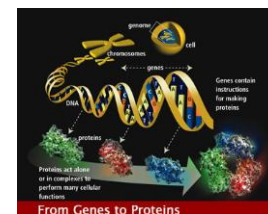
- Multiple protein-protein interactions
- Dynamic and temporal processes!
- Regulation (normal) and de-regulation (disease)

Post-genomic era

genes to pathways:



tough journey



Vol 437|22 September 2005

nature

NEWS & VIEWS FEATURE

PHARMACEUTICALS

A new grammar for drug discovery

Mark C. Fishman and Jeffery A. Porter

To realize the potential of the genome for identifying candidate drugs we must move beyond individual genes and proteins. The signalling pathways in cells provide the right level for such analyses.

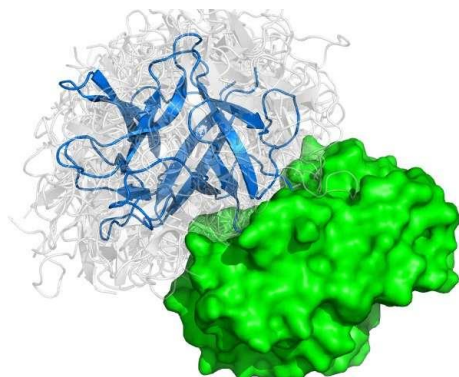
DRUG DISCOVERY

Sci Trans Med (2014)

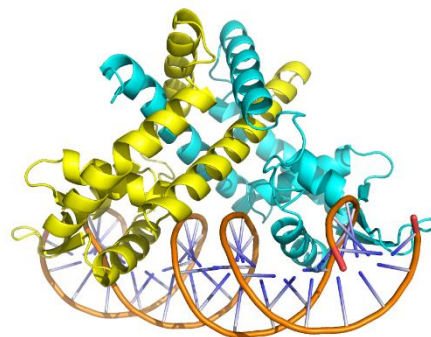
Turning the Titanic

Underestimated the complexity of human biology!

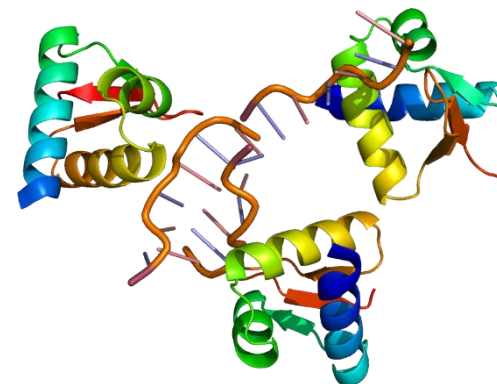
Protein-protein



DNA-protein



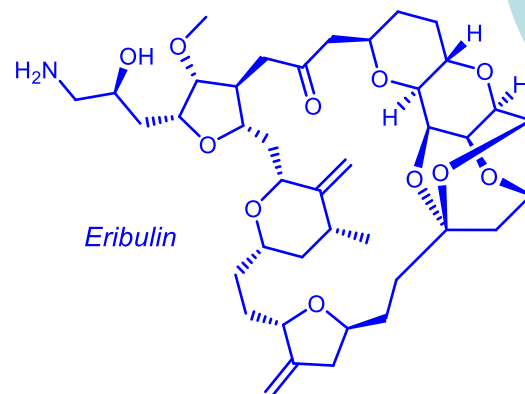
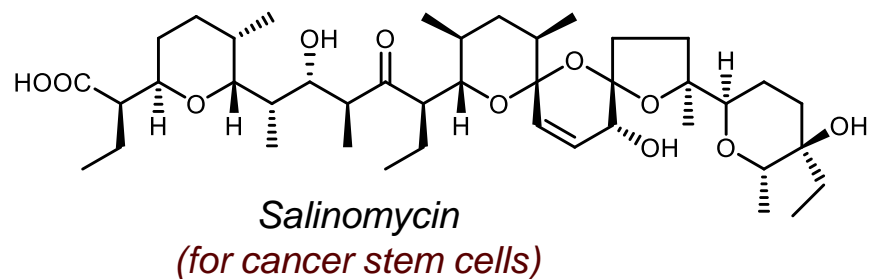
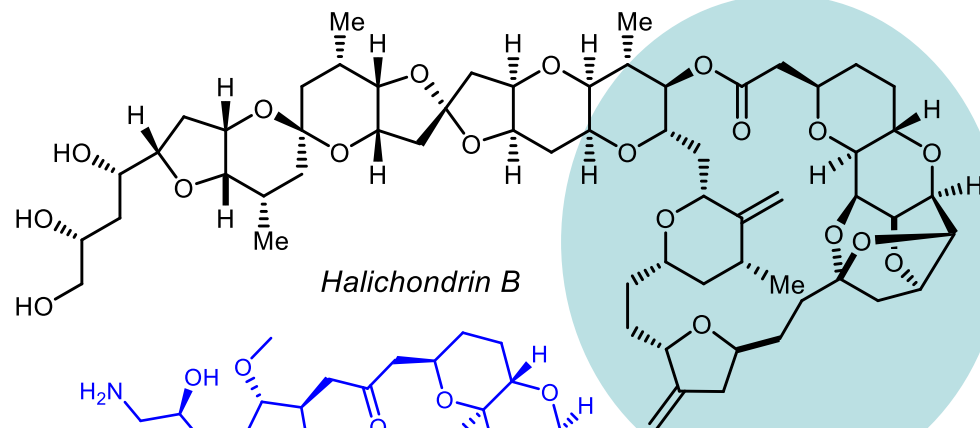
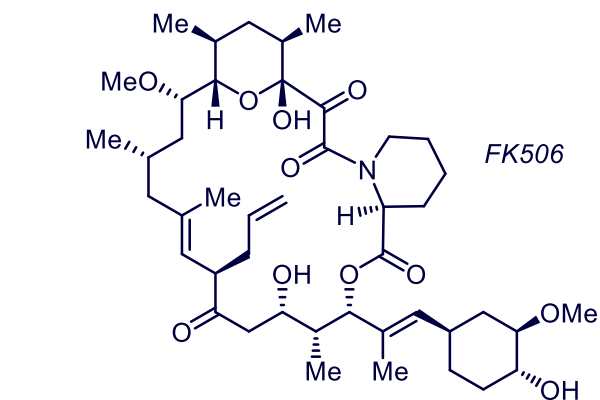
RNA-protein



Current Limitations

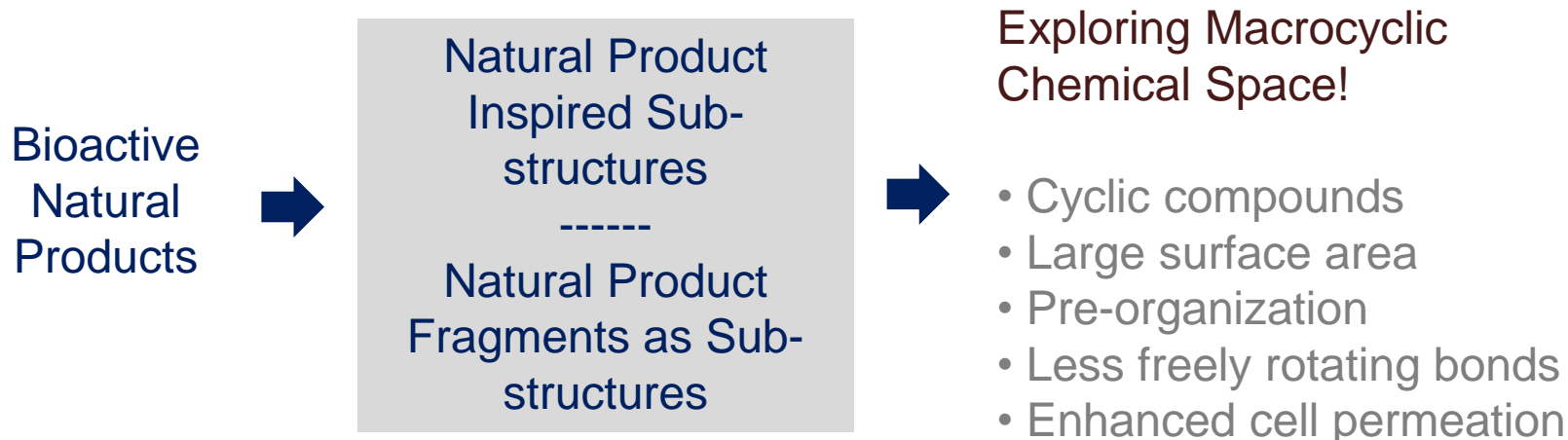
- Limited or no structural biology knowledge as the starting point
- Targets related to transcription and translation machinery
- Mapping large surface area by traditional small molecules is not effective by current approaches
- In general, small molecules are enriched with heterocyclic compounds; not attractive for targets related to multi protein-protein, DNA/RNA-protein interactions
- Lack of our ability to handle these challenging targets classifies them as “undruggable”
- Severely lack of “compounds” within pharma for undertaking “next generation targets”!

Why Looking at Natural Products?



The most complex synthetic compound as drug!

- Natural products (biologically validated chemical space) have an excellent track record as modulators of protein-protein interactions.
- Present complex 3D architectures and dense display of stereo-defined groups.
- Due to complex structures, present a challenging task in taking them forward.
- Can serve as a good source of inspiration for developing sufficiently complex, novel scaffolds in the drug discovery arena!



Key features in our design:

- 3D architectures; sufficient complexity
- stereochemical and skeletal diversity
- synthesis in a reasonable time-scale
- easy to follow-up medicinal chemistry studies
- ease of scalable synthesis is attractive in our designs

Our Chemical Toolbox Platform

*See these research papers from
Arya Research Team:*

Chem. & Biol. 163 (2005)
Curr. Opin. Chem. Biol. 247 (2005)
Chem. Rev. 1999 (2009)

Diversity-Based Organic Synthesis in the Era of Genomics and Proteomics**

Prabhat Arya,* Doug T. H. Chou, and Myung-Gi Baek

Angew. Chem. Int. Ed. **2001**

Natural-product-like chiral derivatives by solid-phase synthesis

Prabhat Arya* and Myung-Gi Baek

Curr. Opin. Chem. Biol. **2001**

Chemistry & Biology, Vol. 9, 145–156, February, 2002, ©2002 Elsevier Science Ltd. All rights reserved. PII S1074-5521(02)00105-9

**Toward High-Throughput Synthesis
of Complex Natural Product-Like Compounds
in the Genomics and Proteomics Age**

Review

RESCUING COMBICHEM

Diversity-oriented synthesis aims to pick up where traditional combinatorial chemistry left off

STU BORMAN, C&EN WASHINGTON

**Natural product-like chemical space: search for chemical
dissectors of macromolecular interactions**

Ayub Reayi and Prabhat Arya

Curr. Opin. Chem. Biol. **2005**

**The natural-product-like compounds produced in DOS have
a much better shot at interacting with desired molecular
targets and exhibiting interesting biological activity.**

Chem. & Eng. News **2004**
(statement from Prabhat Arya)

Chemistry & Biology, Vol. 12, 163–180, February, 2005, ©2005 Elsevier Ltd All rights reserved. DOI 10.1016/j.chembiol.2005.01.011

**Exploring New Chemical Space
by Stereocontrolled
Diversity-Oriented Synthesis**

Review

Chem. Rev. **2009**, *109*, 1999–2060

1999

**CHEMICAL
REVIEWS**

2014

Review

pubs.acs.org/CR

**Advances in Solution- and Solid-Phase Synthesis toward the Generation of
Natural Product-like Libraries**

Jyoti P. Nandy,[†] Michael Prakesch,^{†,‡} Shahriar Khadem,^{†,§} P. Thirupathi Reddy,[†] Utpal Sharma,[†] and Prabhat Arya^{*,†,‡}

**Small Molecule Modulators of Protein–Protein Interactions: Selected
Case Studies**

Madhu Aeluri,[†] Srinivas Chamakuri,[†] Bhanudas Dasari,[†] Shiva Krishna Reddy Guduru,[†] Ravikumar Jimmidi,[†] Srinivas Jogula,[†] and Prabhat Arya^{*}

Dr. Reddy's Institute of Life Sciences (DRILS), University of Hyderabad Campus Gachibowli, Hyderabad 500046, India

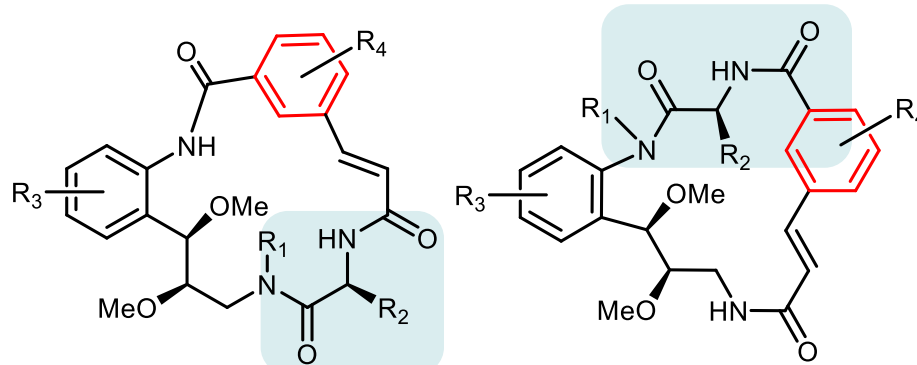
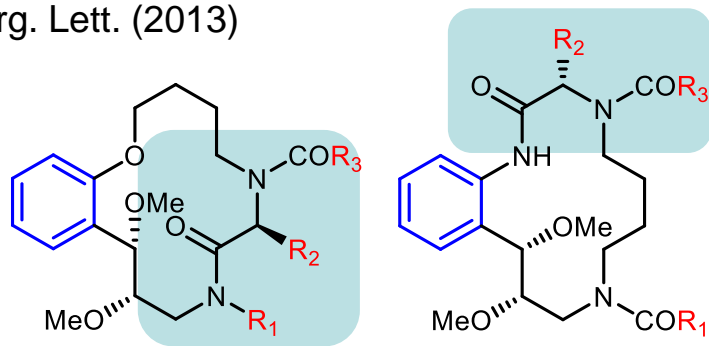
1 Natural Product-Inspired,
Functionalized 14- and 17-Membered Rings
Macrocyclic Toolbox

2 Indoline / Tetrahydroquinoline (Alkaloids)
and Benzofuran (Flavonoids)-Inspired
Macrocyclic Toolbox

3 Glyco-based Macrocyclic Toolbox

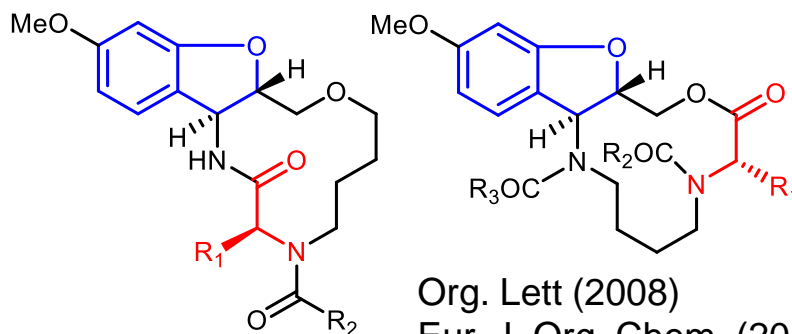
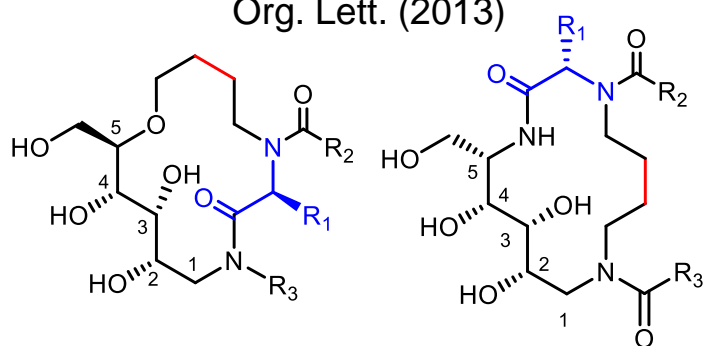
Examples of Our Early Work (Contd.)

Org. Lett. (2013)

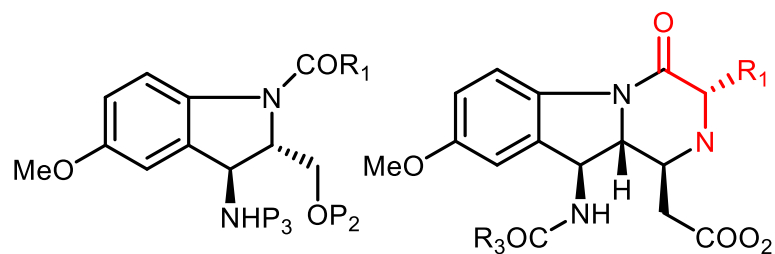


Eur. J. Org. Chem. (2013)

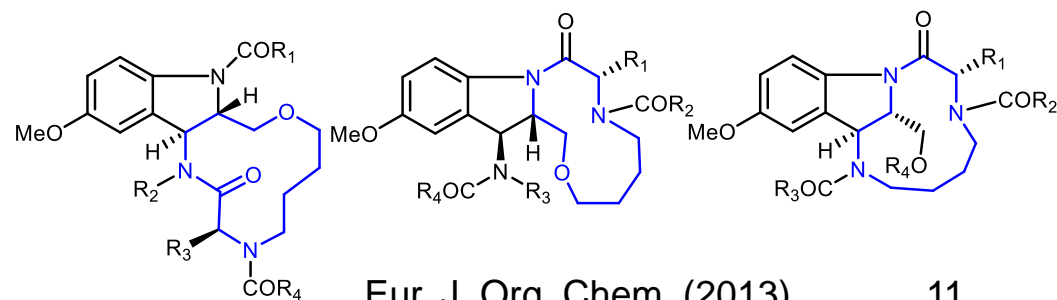
Org. Lett. (2013)



Org. Lett (2008)
Eur. J. Org. Chem. (2014)

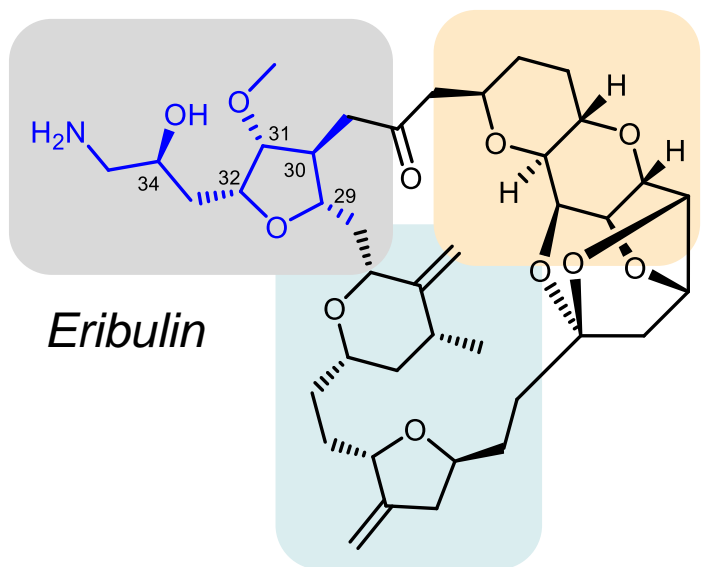


Angew Chem (2005)



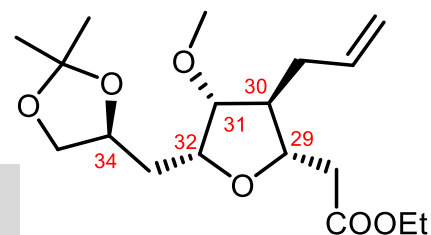
Eur. J. Org. Chem. (2013)

Example: Macrocyclic Toolbox based on Eribulin Sub-structures



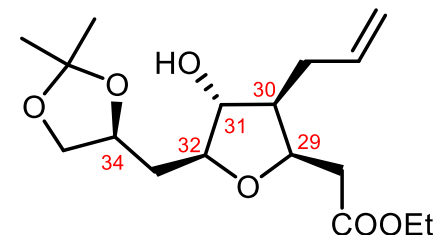
Practical and scalable synthesis

A

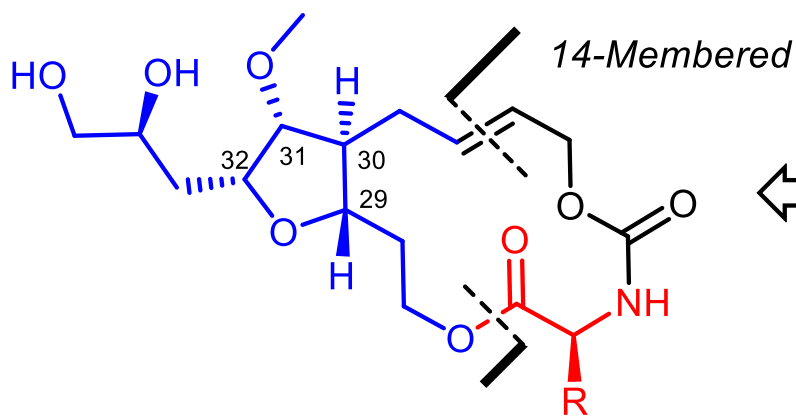


*Precise
fragment-based*

B

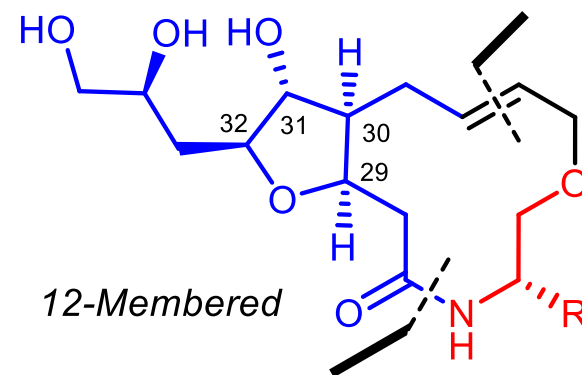


*Diastereomeric
fragment-based*



A

B



Org. Lett. 468-471 (2015)

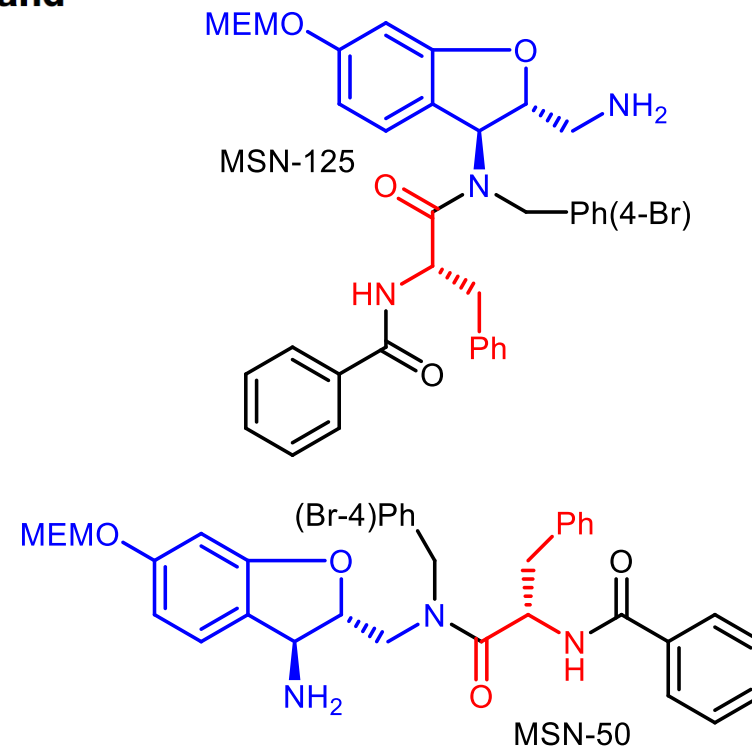
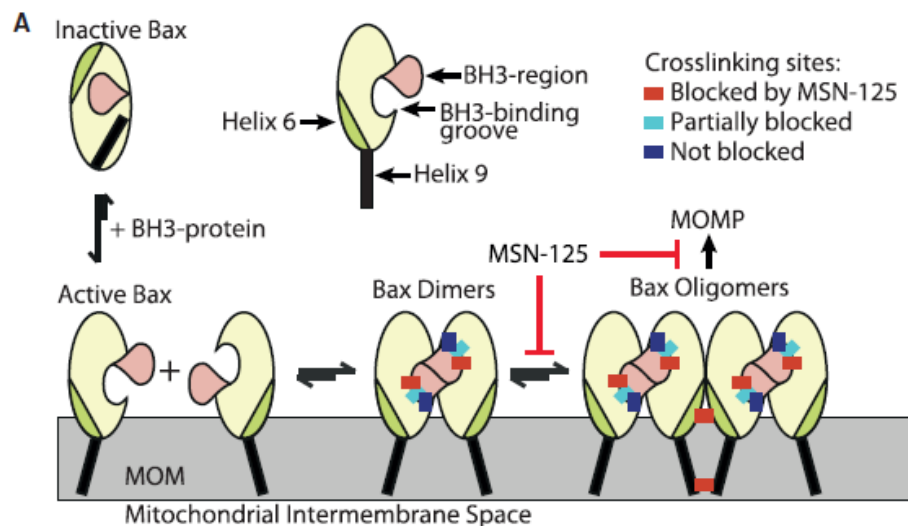
Applications of Our Chemical Toolbox in Finding
Active Small Molecules (as Early Stage Drug
Candidates) for Challenging “Undruggable” Targets:

Three Case Studies

Cell Chemical Biology

A Small-Molecule Inhibitor of Bax and Bak Oligomerization Prevents Genotoxic Cell Death and Promotes Neuroprotection

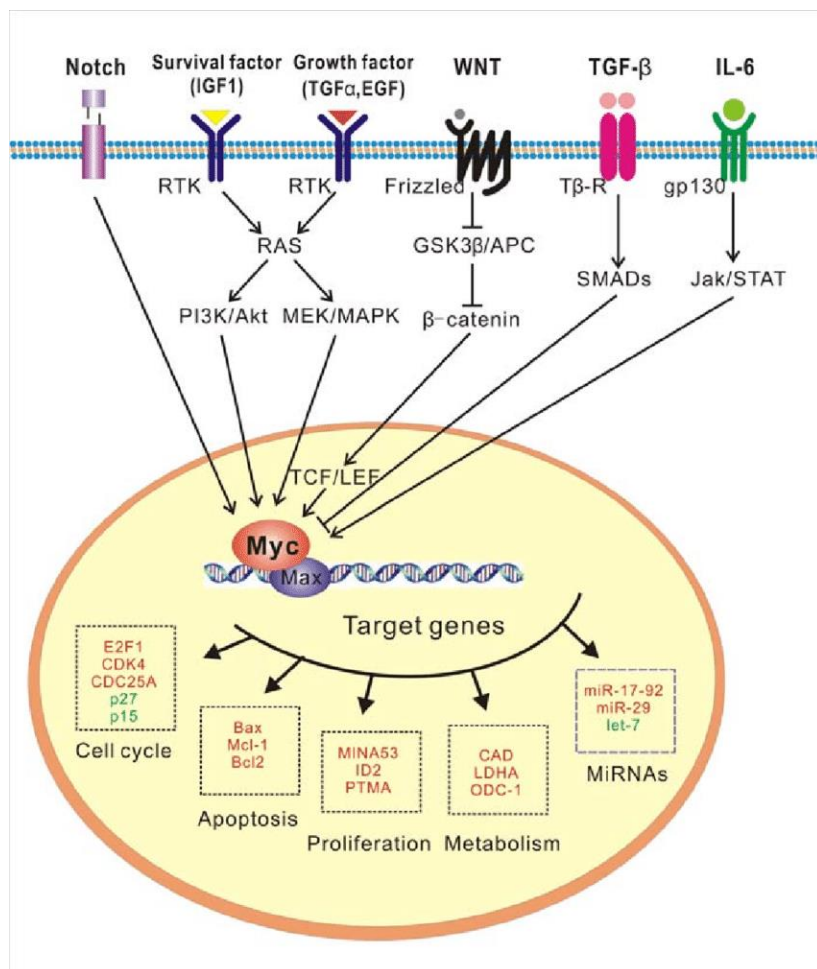
2017



In collaboration with with David Andrews Lab, Univ of Toronto

During the academic tenure (from Arya Group)

Going in for c-Myc: Rationale

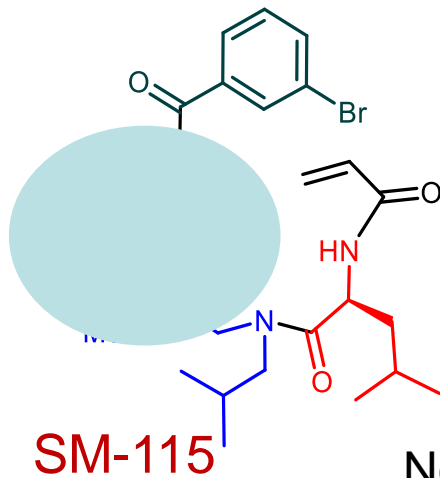
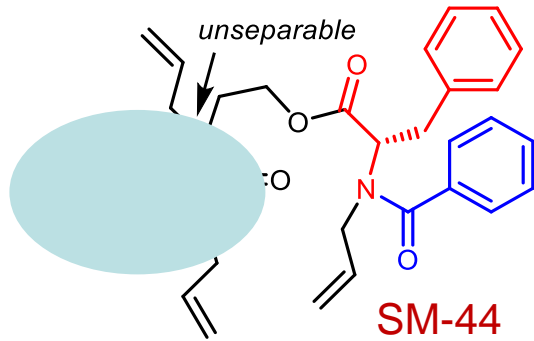


- “Undruggable”
- Frequently mutated and overexpressed in many human tumors
- A transcription factor
- Responsible for upregulation of several genes related to cell cycle, apoptosis, and cellular transformation which are directly associated with cancer
- Despite working on this target for more than 3 decades, the current chemistry approaches have not led to producing the effective drug candidates!

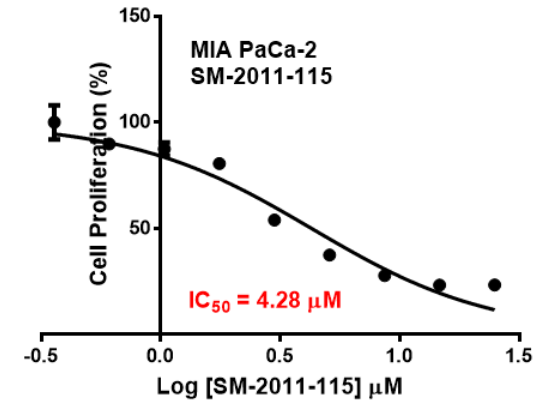
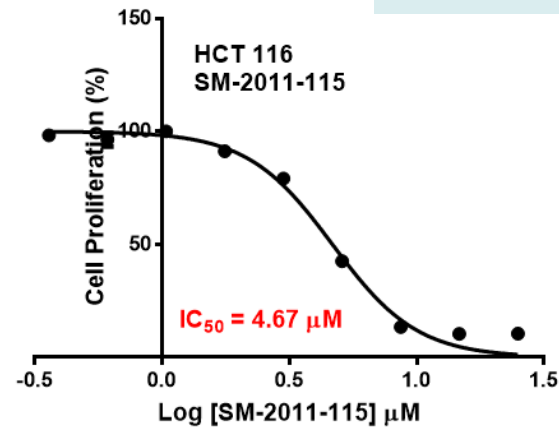
Note: Given the track record of working with “undruggable” targets (for ex. BAX), our chemical toolbox platform provide an excellent opportunity to tackle this validated target which is challenging the community for long!

A Glimpse of Our Data

WST1 Screening Assay

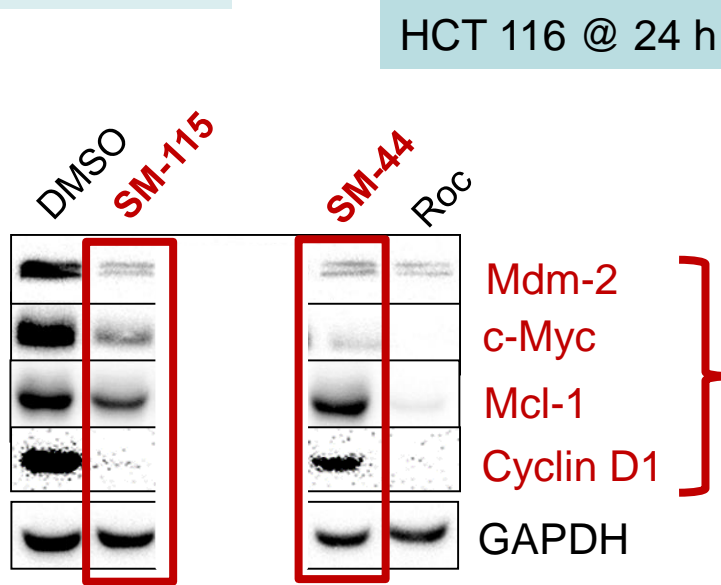


with SM-115



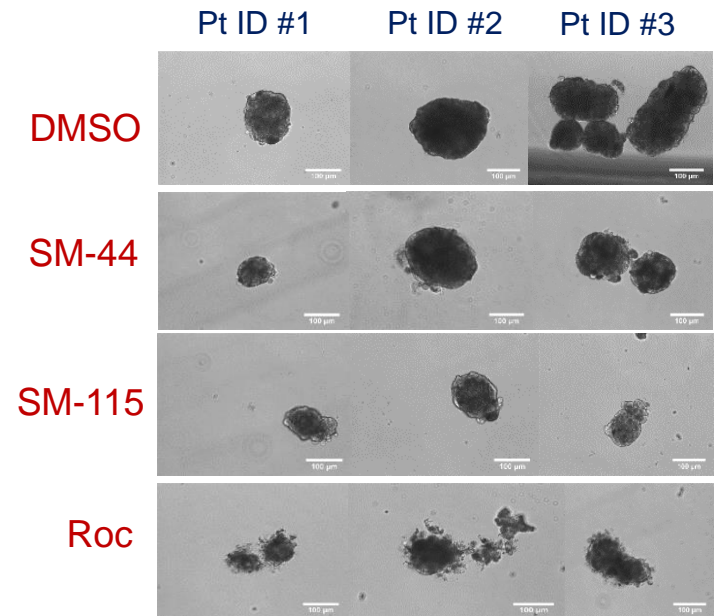
Note: First-in-class small molecule inhibitors of c-Myc translation

Western Blots



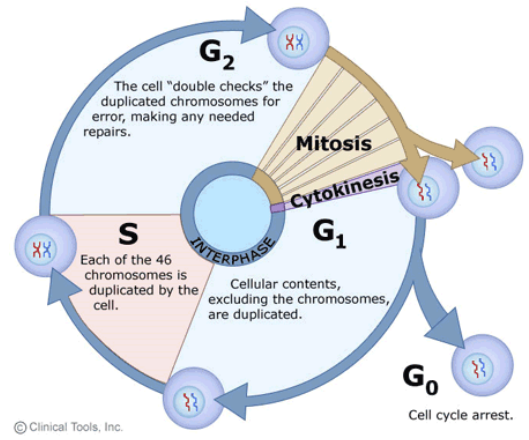
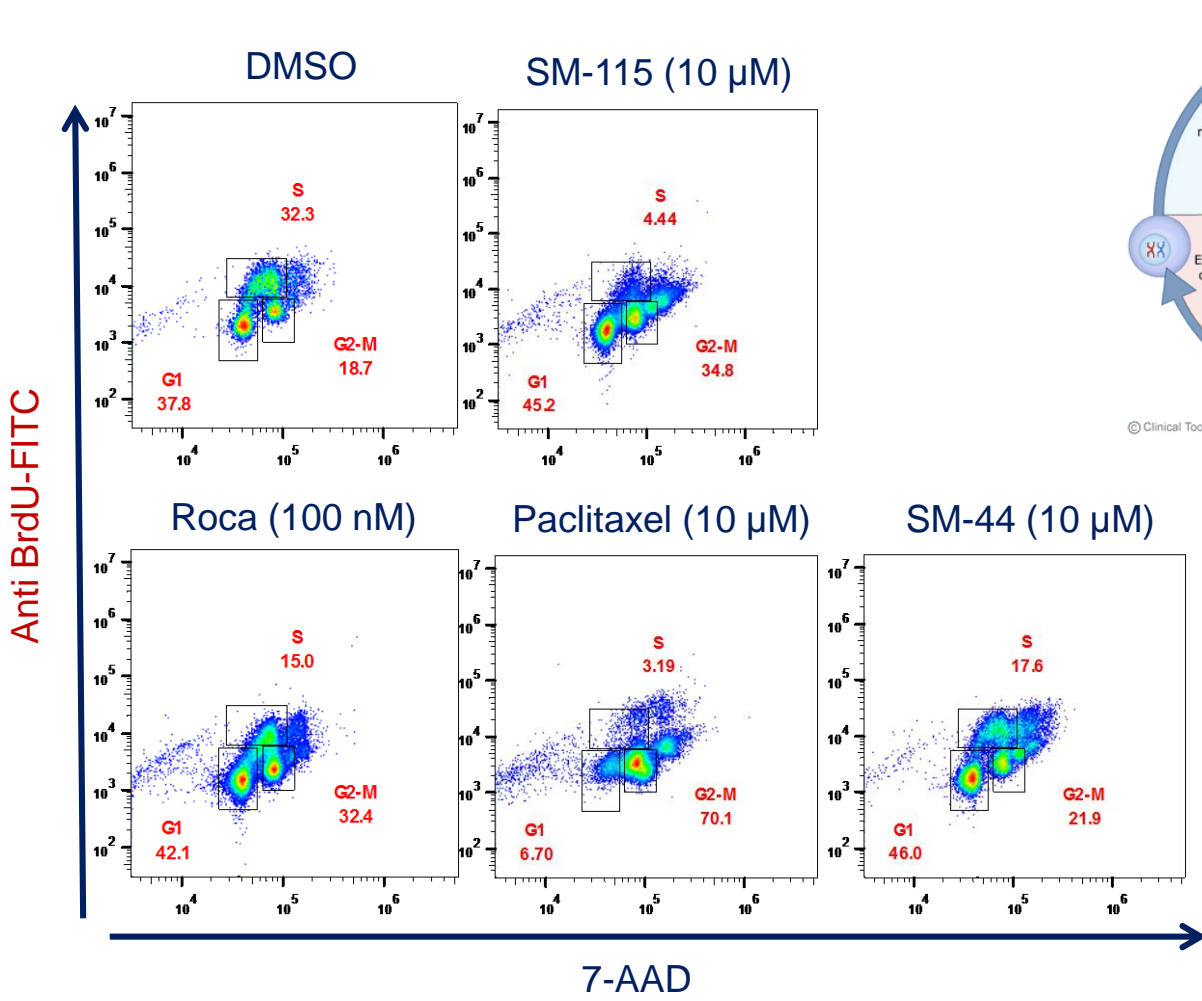
Serious decrease in expression level of c-Myc and target associated molecules like MDM-2, Mcl-1 and Cyclin D1!

Effect on Patient-derived Tumorspheres at Day 12



Note: First-in-class small molecule inhibitors of c-Myc translation

Flow Cytometry (BrdU Probe for Cell Cycle)



Cyclin D1
CDK4

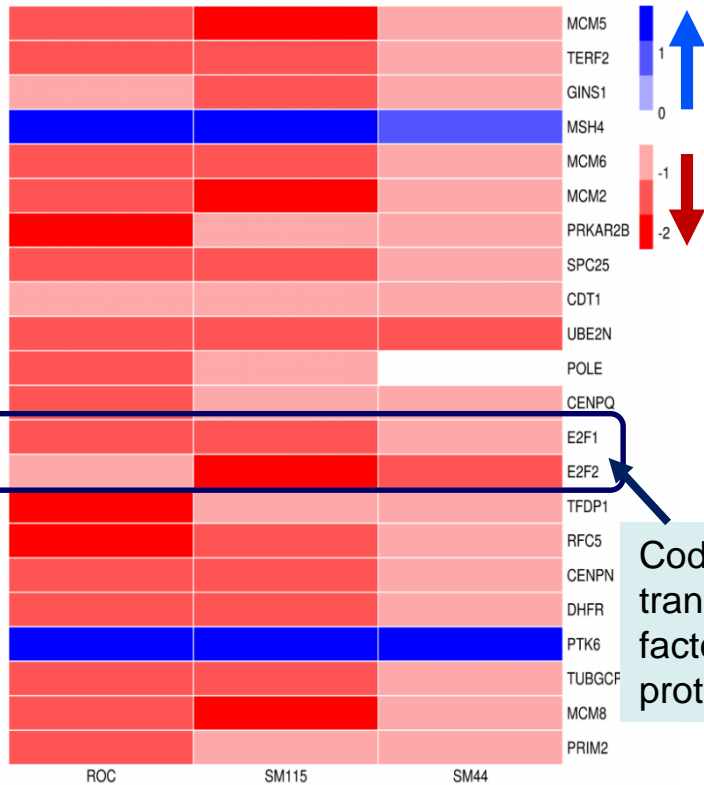
	G1	S	G2/M
DMSO	36.6	32.3	18.6
SM-115	43	4.44	34.5
SM-44	43.2	17.6	21.7
Rocaglamide	36	15	31.1
Paclitaxel	6.19	3.19	69.4

Roca = Rocaglamide

Note: G1 arrest – possible downregulation of Cyclin D1 and CDK4

Our Transcriptomic Study

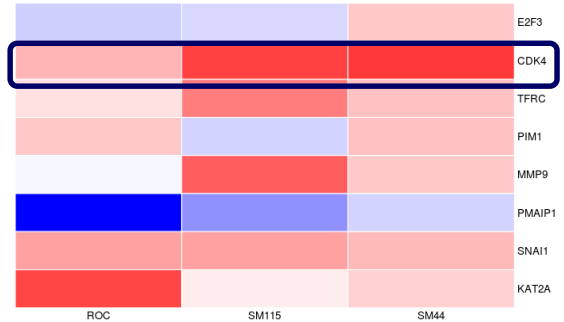
Cell Cycle



Codes for transcription factor proteins

Note: A similar pattern in cell cycle gene set (see ROC vs SM115 and SM44)

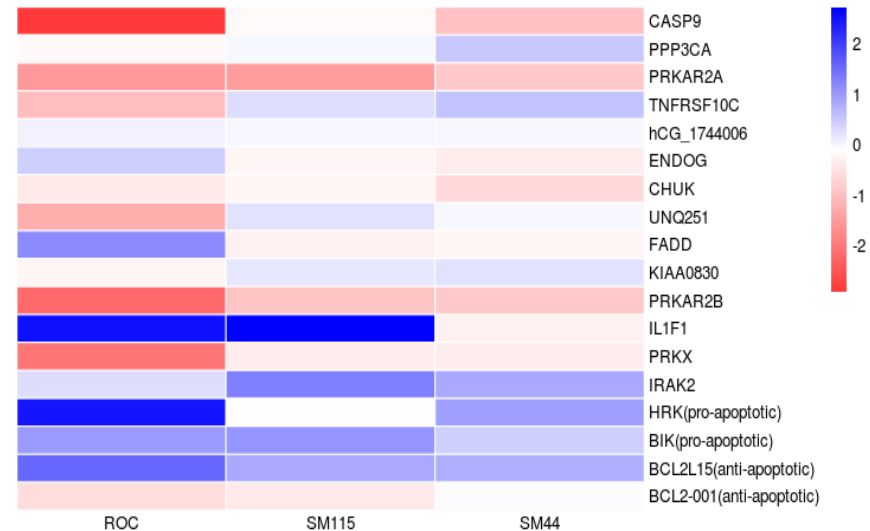
MYC



CDK4 gene crucial in cell cycle from G1 to S phase

Note: Downregulation of CDK4 with SM115 and SM44

APOPTOSIS



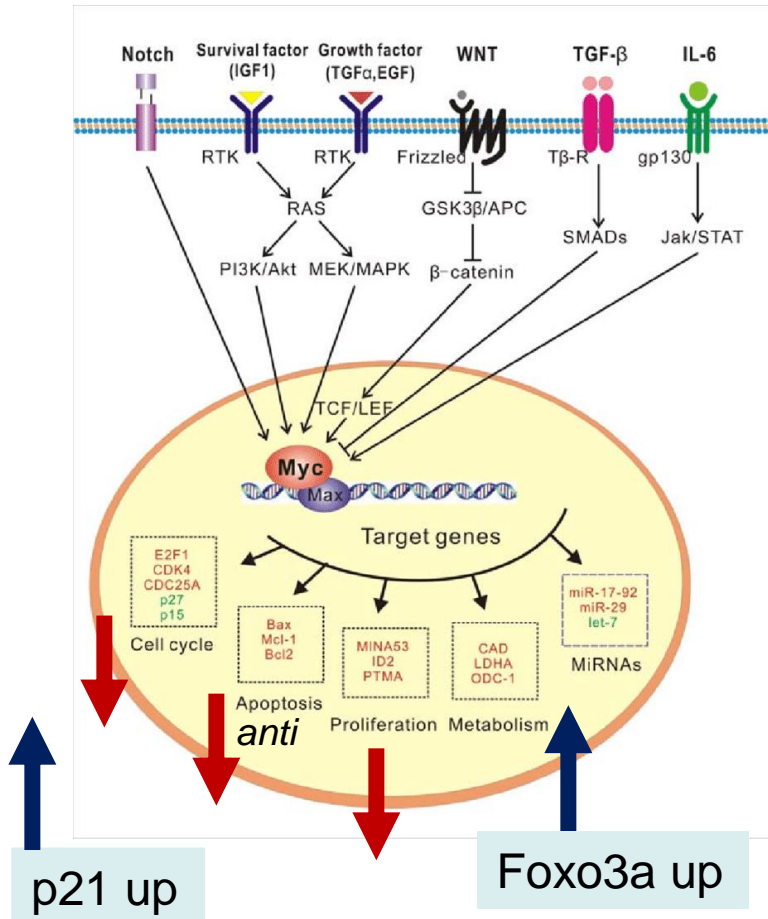
Down regulation of anti-apoptotic gene

Upregulation of pro-apoptotic genes

Note: similar upregulation for BIK (pro-apoptotic gene), BCL2-001 (anti-apoptotic gene)

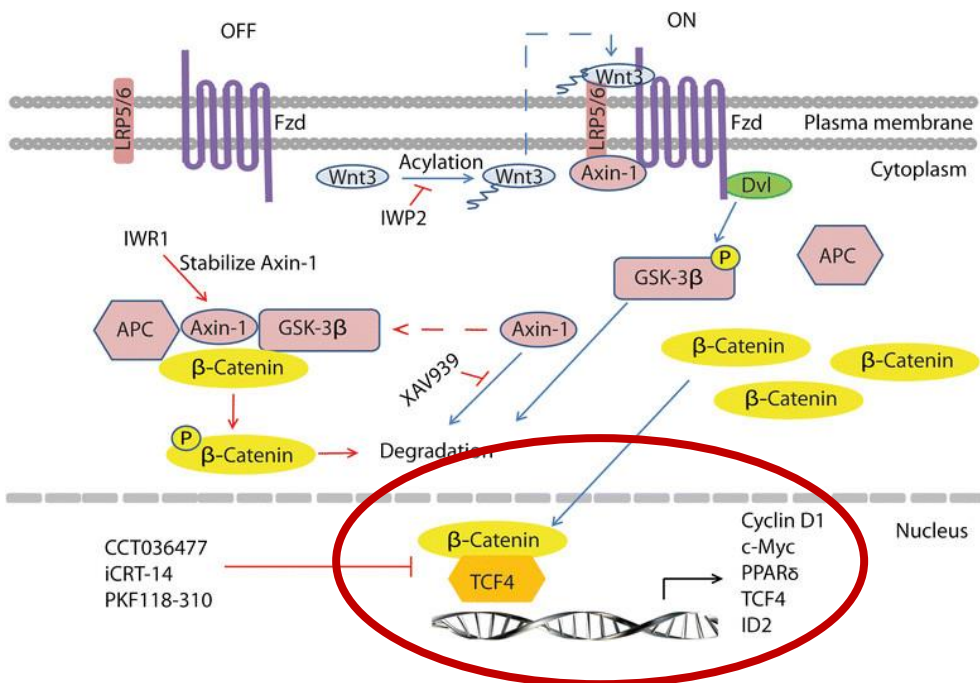
Comparison of data with Rocaglamide (ROC) and our two actives (SM115 and SM44)

Highlights of Our Program



1. Discovered two novel families of small molecules as **c-Myc** translation inhibitors.
2. Our actives are effective in inhibiting the expression levels of c-Myc, MDM-2, Mcl-1 and Cyclin D1!
3. Our actives are promoters of apoptosis and also exhibit the cell cycle arrest at the G1 phase.
4. Our actives can be considered **as the functional mimics of Rocaglamide**, a natural product, well-known as the c-Myc translation inhibitor.
5. Our cellular studies are supported by the NGS data (for ex, related to seeking info on mode of action)
6. To our knowledge, there are no small molecules known to date; our work opens up a new direction in the field of c-Myc translation-based cancer drug discovery!

Going in for Wnt: Rationale



- Undruggable
- Play a key role in stemness, EMT, cancer stem cells and metastasis.
- Aberrant regulation is a common theme seen across many tumor types.
- A highly attractive target due to its role in development and maintenance of cancer stem cells.
- Several small molecules are in clinic but most are not very effective.
- ICG001 is a small molecule going forward in clinic as the Wnt transcription inhibitor.

Small molecule inhibitors of Wnt transcription

Note: no drugs are developed to date

Nat Rev Drug Disc (2014)

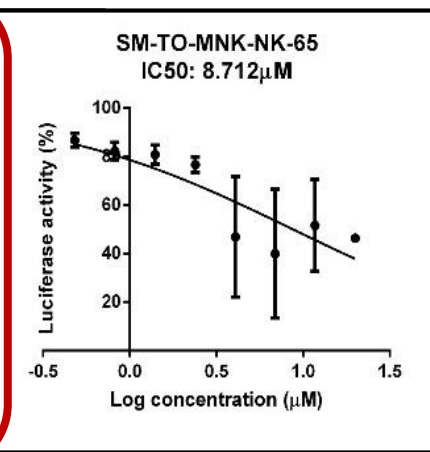
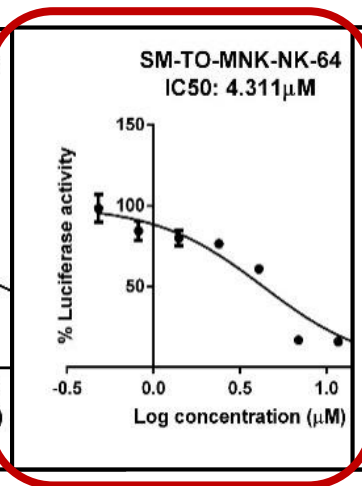
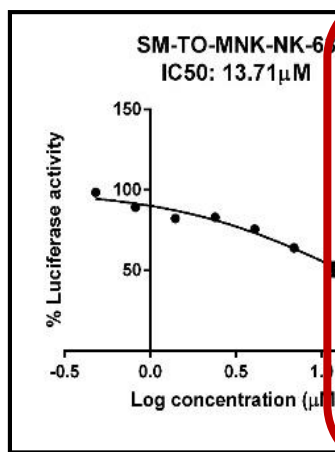
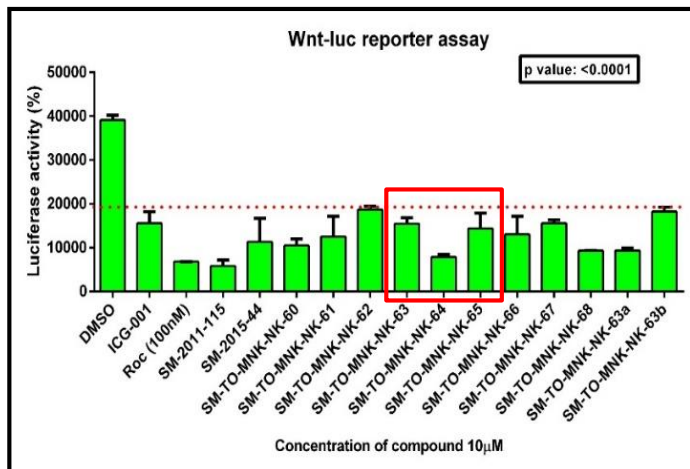
Note: Given the track record of working with “undruggable” targets (for ex. BAX), our chemical toolbox platform provide an excellent opportunity to tackle this target which is challenging the community for long!

Wnt Program: A Glimpse of Our Data

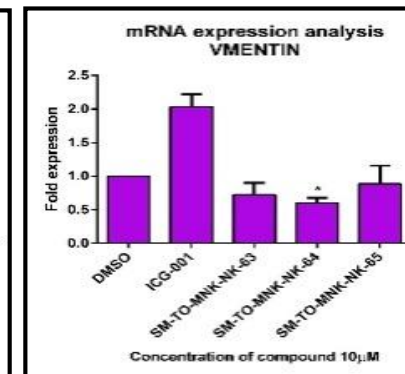
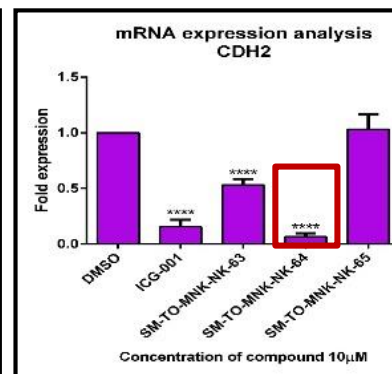
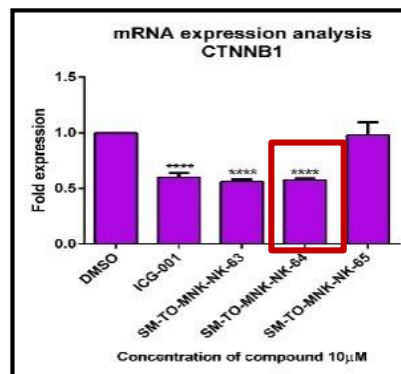
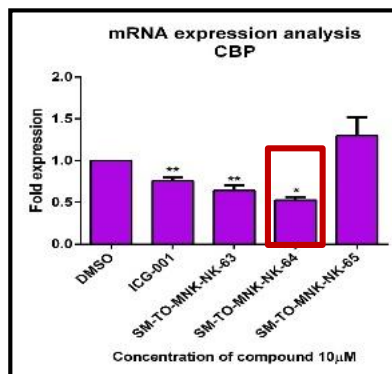


Wnt Luciferase Assay

Use of ICG001 as the control



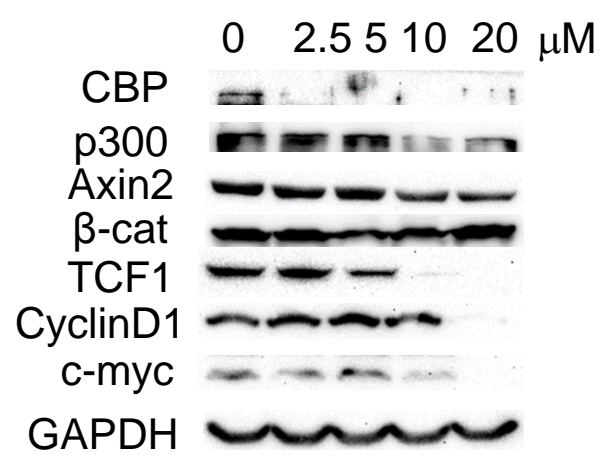
RT-PCR



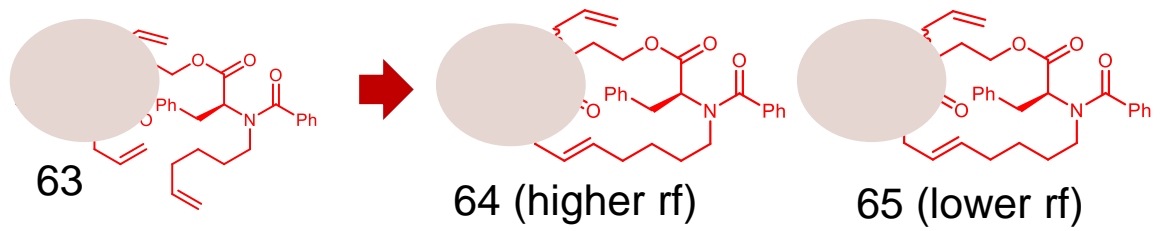
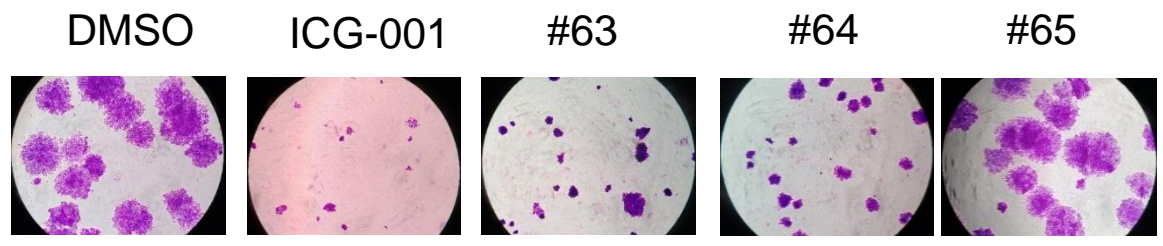
Wnt Program: A Glimpse of Our Data (contd.)



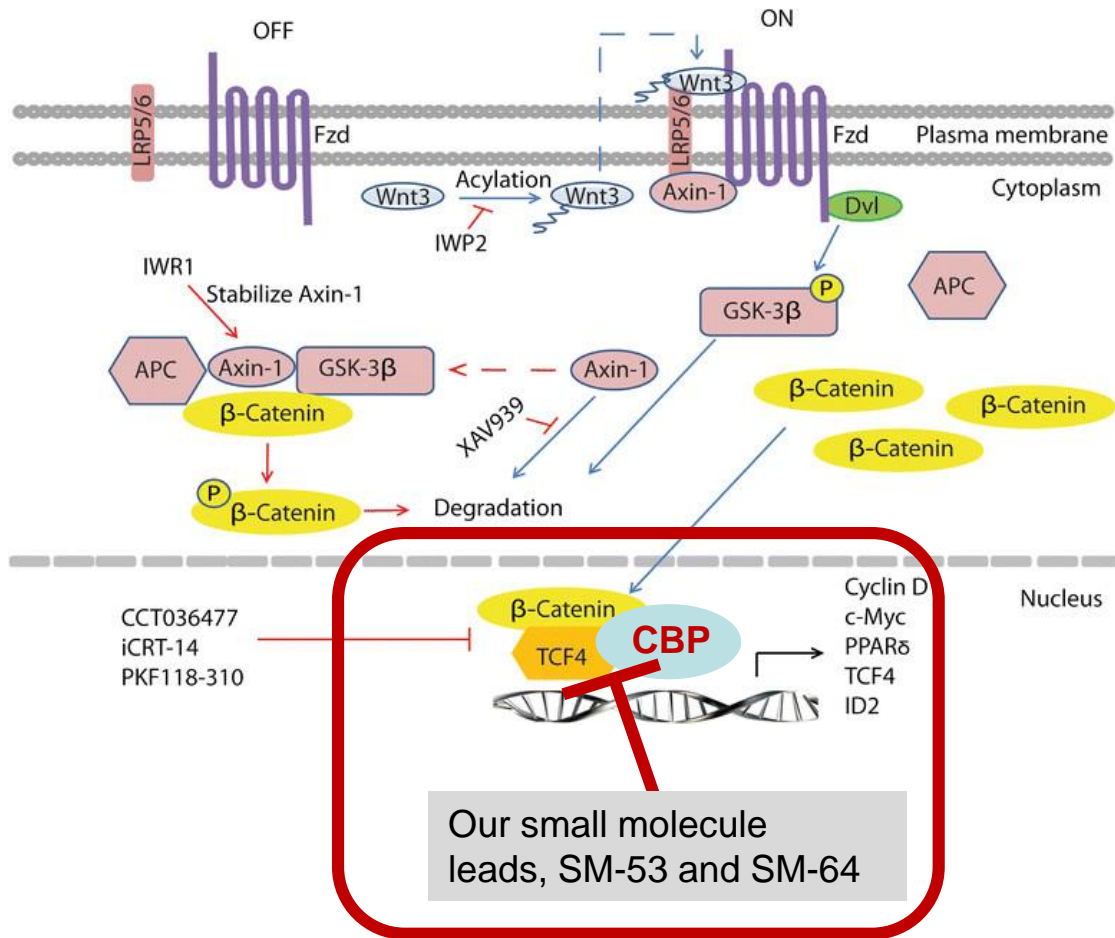
Western Blots



Colony Formation Assay



64 and 65 are diastereomers



The Next Steps:

- Hit to lead studies (almost finished)
- Thorough evaluation on patient-derived tumorspheres and organoids
- Precise Target ID information
- Detailed studies with tumor metastasis, migration, stem cell maintenance, EMT related to cancer stem cells.
- Finish all pre-clinical studies; reaching the IND stage

Wnt Transcription Inhibitors

1. Natural Product-Inspired Macrocyclic Compounds as Wnt Transcription Inhibitors. US Patent Application No. 62862147, Filed June 17, 2019.
2. Natural Product-Inspired Compounds as the Inhibitors of Wnt Transcription Machinery. US Patent Application No. 62863287, Filed June 19, 2019.
3. Substituted furan-based macrocyclic compounds as the Inhibitors Wnt Transcription, **Patent in progress**, expected filing date: Sept-Oct 2021
4. Isatin-based 16-membered, macrocyclic compounds as the Inhibitors Wnt Transcription, **Patent in progress**, expected filing date: Sept-Oct 2021

c-Myc Translation Inhibitors

1. Natural Product-inspired Acyclic and Macrocyclic Compounds as c-Myc Translation Inhibitors. US Patent Application No. 62938342, Nov 21, 2019.
2. Inhibition of c-Myc Translation by Natural Product-inspired Macrocyclic Compounds. US Patent Application No. 62947551, Filed Dec 13, 2019.

- ✓ **Chemical Toolbox:** Cutting edge leadership; working in this domain >20 years.
- ✓ 52 PDFs and 12 PhD students contributed to developing novel synthesis methods.
- ✓ Ownership in several novel scaffolds: 50 published; 250 unpublished scaffolds.
- ✓ Our compounds are highly unique (not me too type) and useful to mapping a large surface area for challenging targets.
- ✓ Identified novel small molecules as Wnt transcription and Myc translation inhibitors (filed 4 US prov. patents).
- ✓ Working with BI and one academic group for building a focused macrocyclic toolbox for their needs.



- ❖ Seeking funds to building a macrocyclic toolbox with 20-40K novel small molecules (highly attractive from IP direction) over 3 years period.
- ❖ In addition to utilizing these small molecules in our own drug discovery programs, they can also be shared with others through various business models.
- ❖ Could also build a focused toolbox (through consultation) matching the needs of specific programs of others.

Note: the detailed information on our published and unpublished scaffold data base can be made available as required