### Targeting MAT2A in MTAP-deleted Cancers

Presented at the American Association for Cancer Research (AACR) Annual Meeting, April 14-18, 2018, Chicago, IL, USA

### Acknowledgements

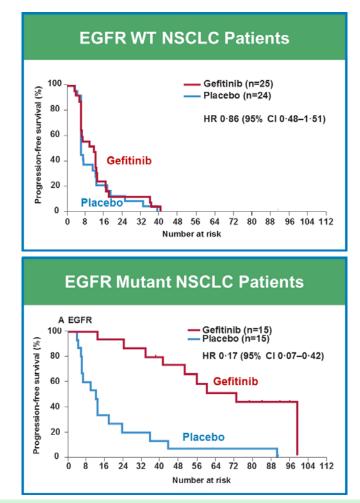
### **Agios 2017 Founders Day Retreat**



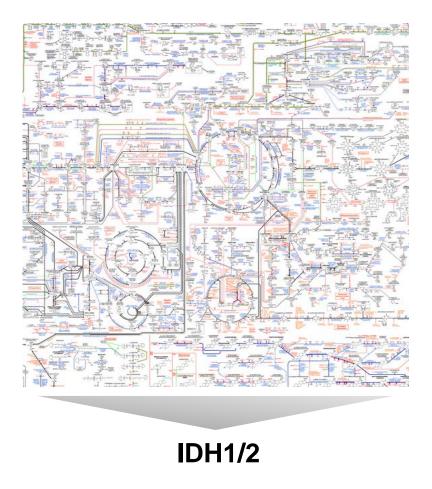
"Hire great people, think big, have fun, follow the science and do what's right for patients"

### The Challenge: Identifying Precision Medicine Approaches in Cancer Metabolism

Directly drugging 'driver mutations' has yielded transformative medicines

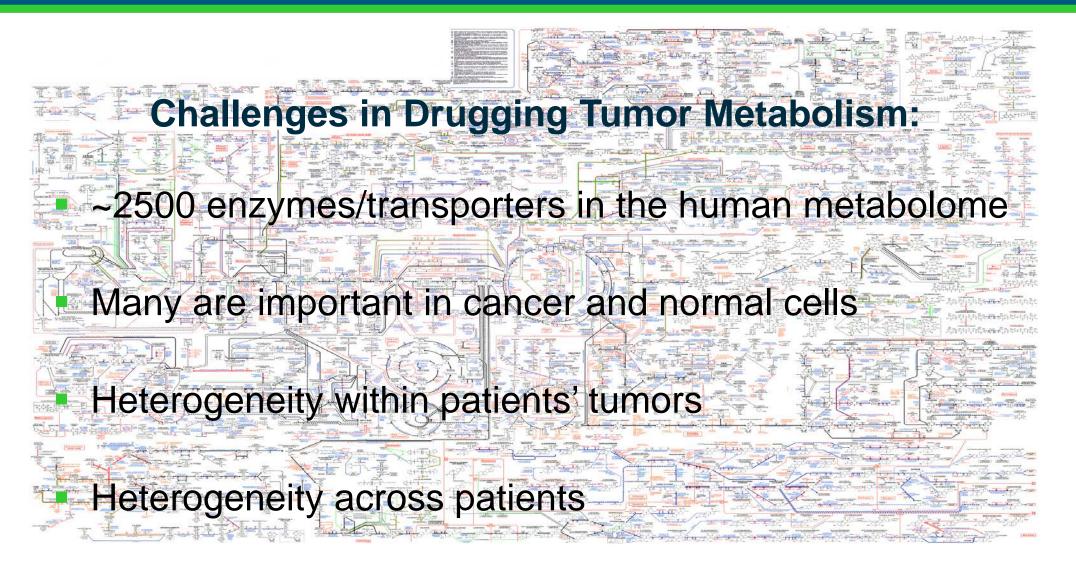


but...DNA sequencing has identified only 2 gain-of-function metabolic 'driver' mutations out of 2000+ metabolic genes

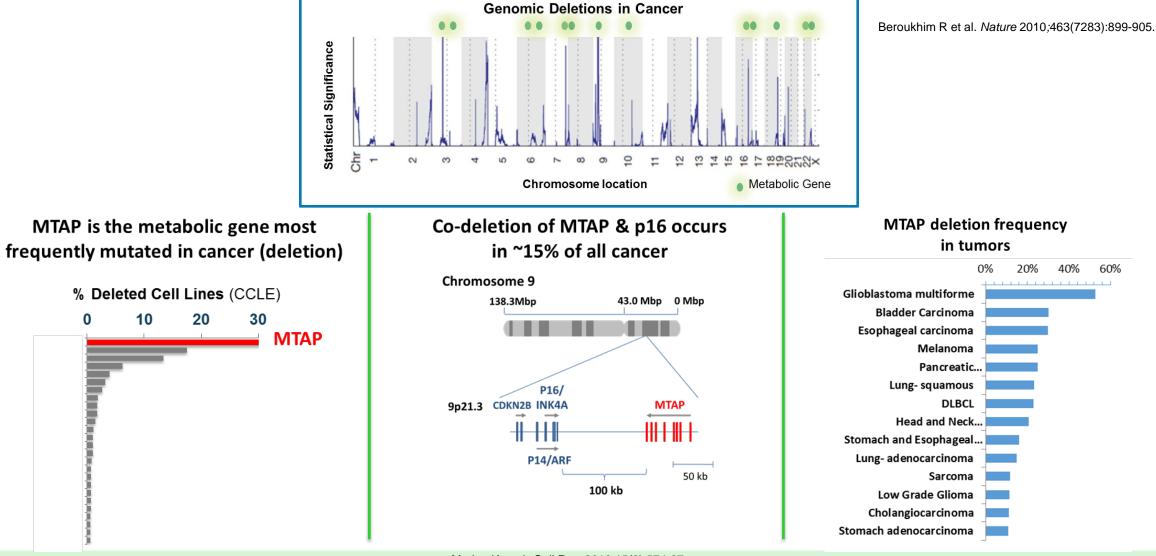


Zhang L et al. Lancet Oncol 2012;13(5):466-75.

### Which Pathways and Targets to Drug?

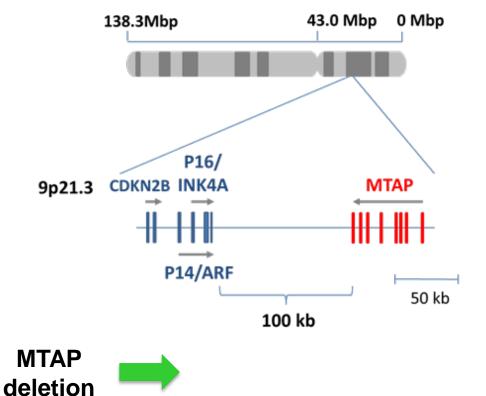


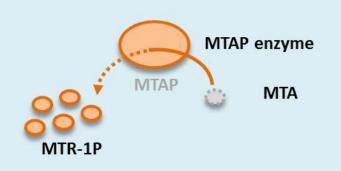
### **Homozygous Deletions of Metabolic Genes in Cancer**



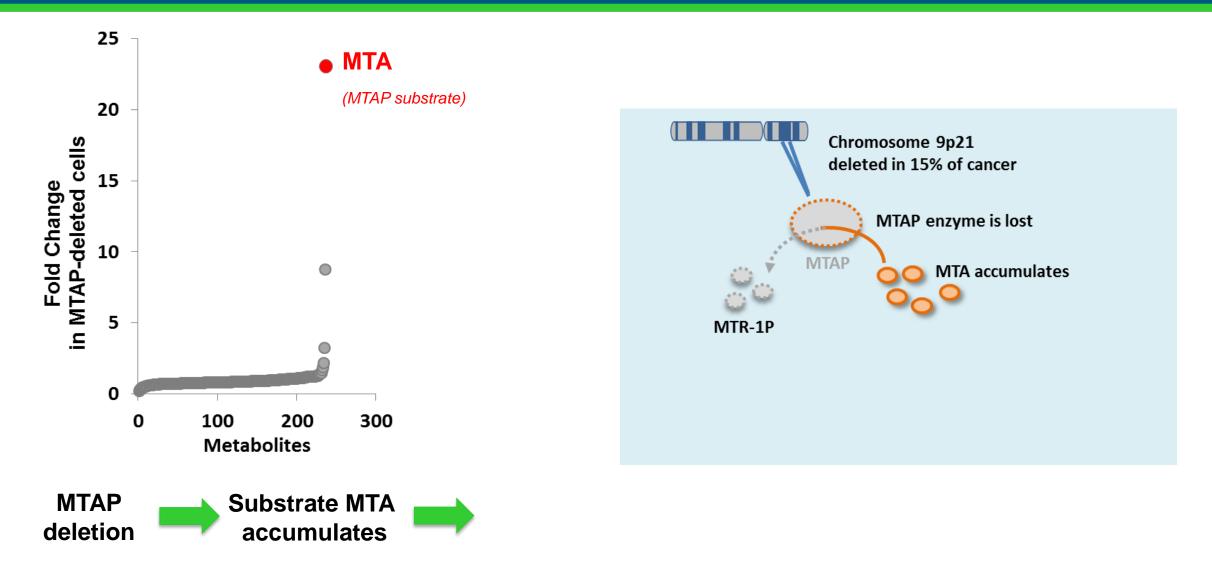
### Co-deletion of MTAP & p16 occurs in ~15% of all cancer

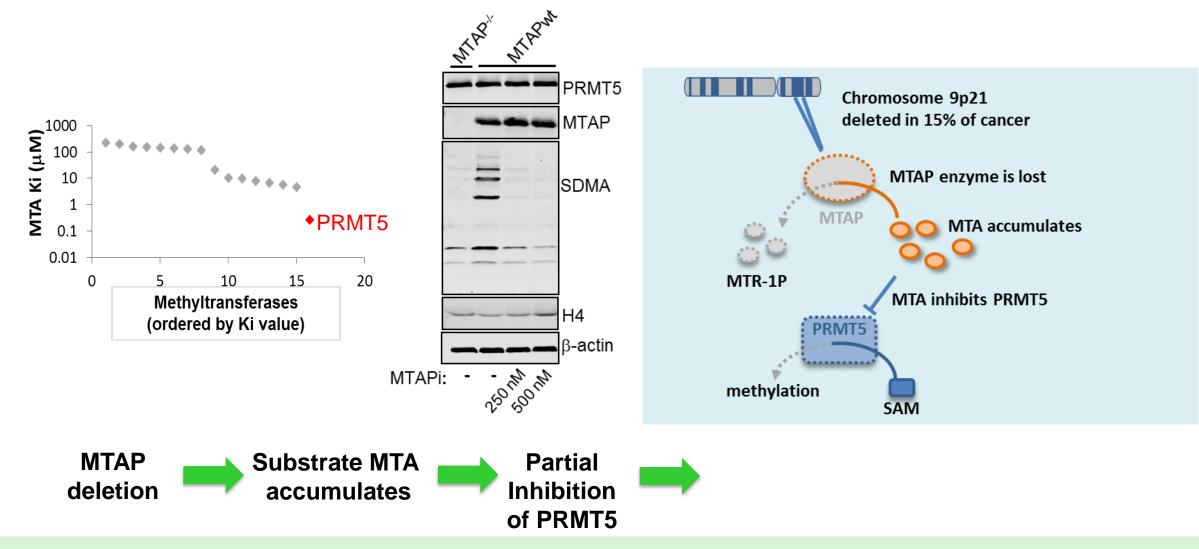
**Chromosome 9** 

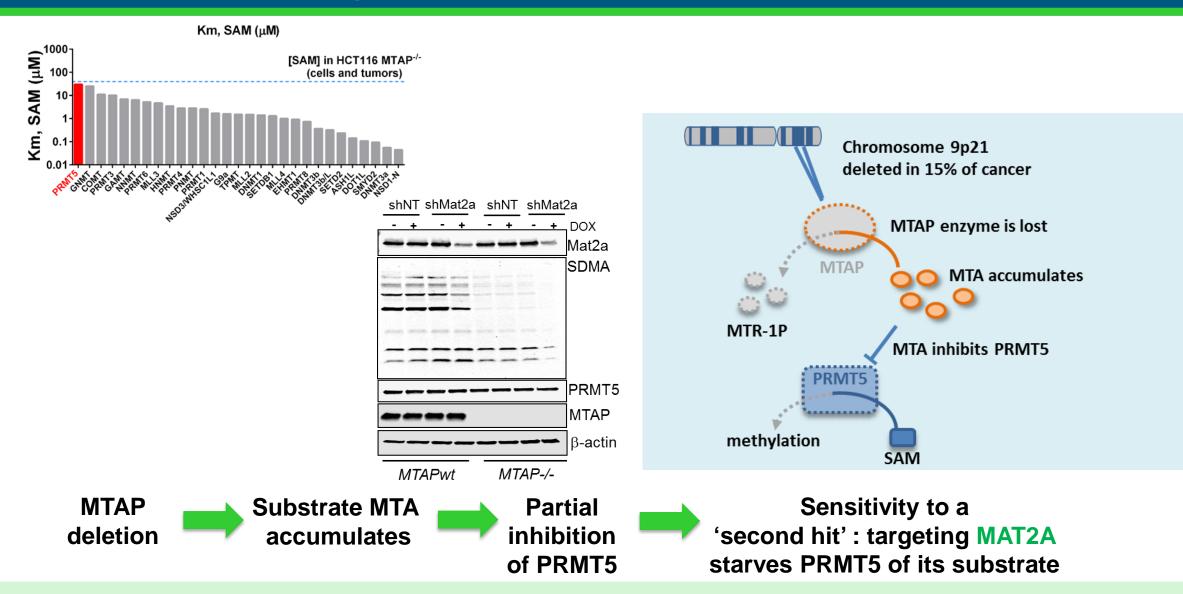




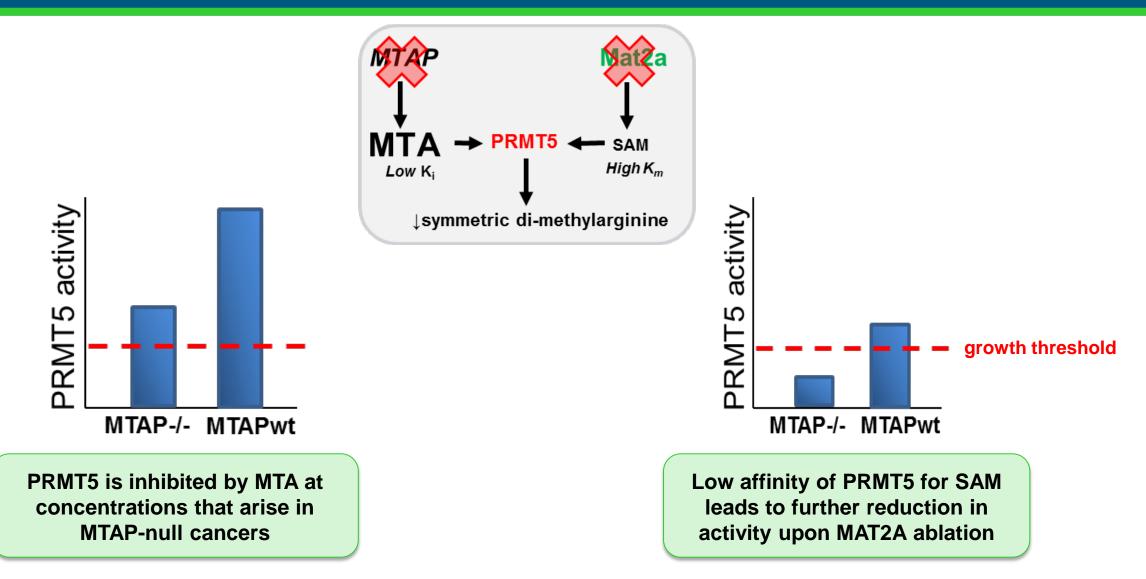
Marjon K et al. *Cell Rep.* 2016;15(3):574-87. © 2016 The Authors. Published by Elsevier Inc.

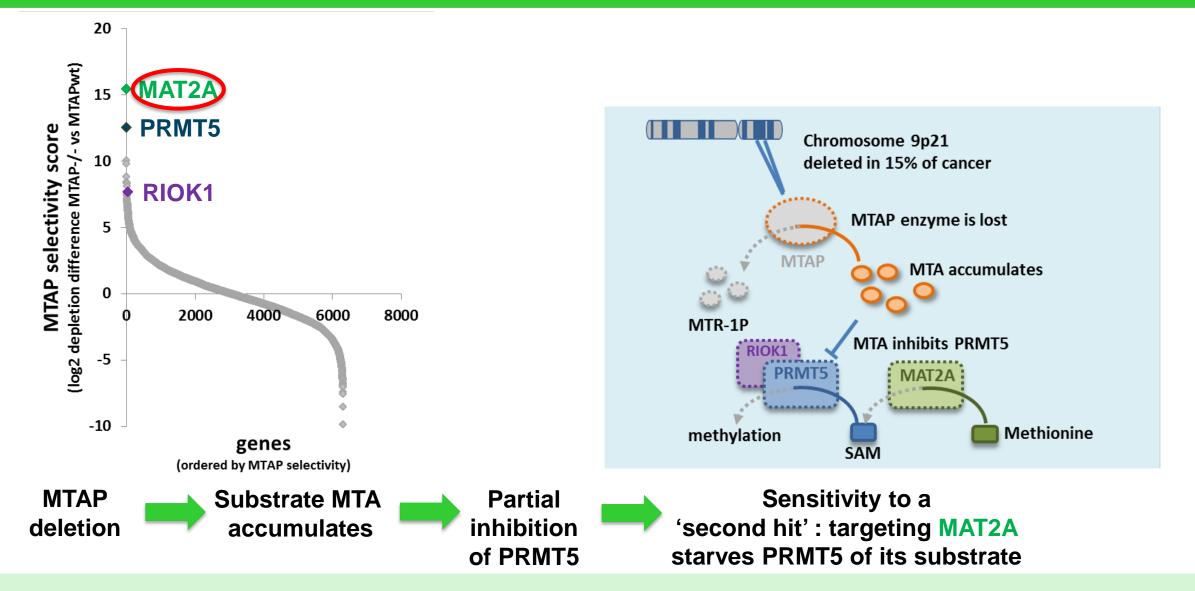






## Fortuitous Biochemical Features of PRMT5 Can Explain the Vulnerability of PRMT5 and MAT2A in MTAP-deleted Cancers

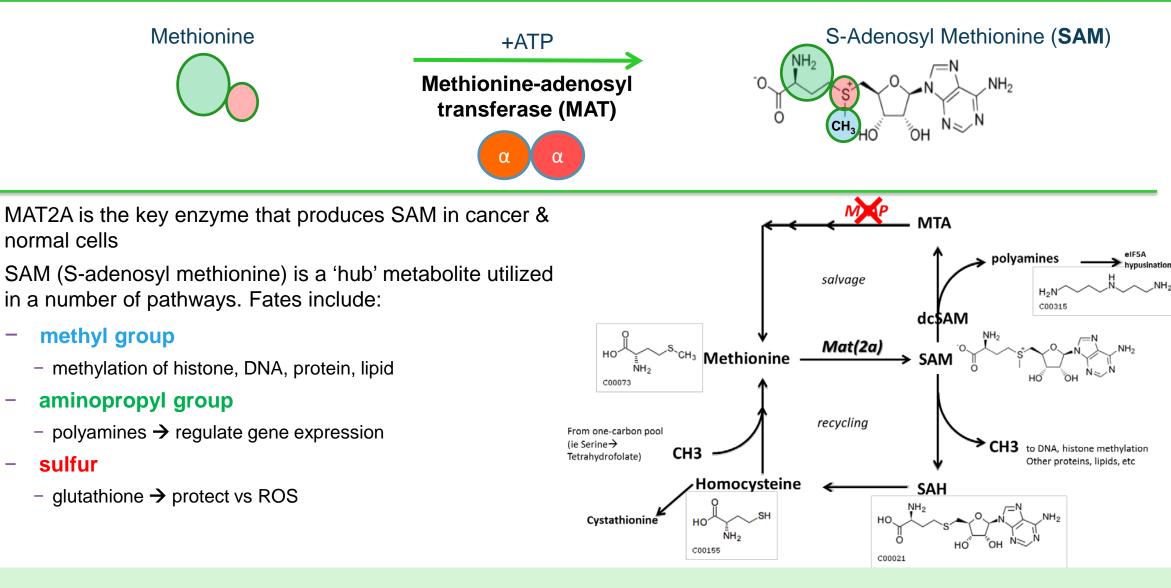




### MAT2A: Methionine Adenosyltransferase 2A

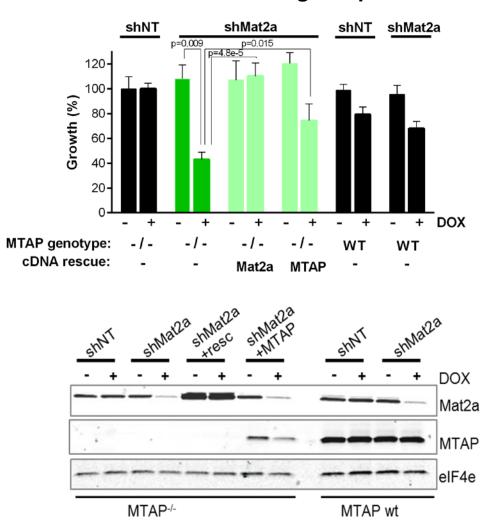
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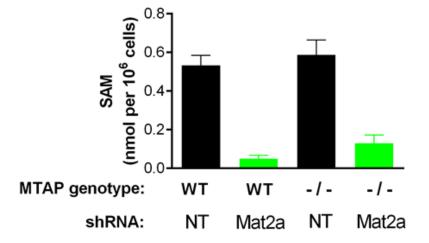
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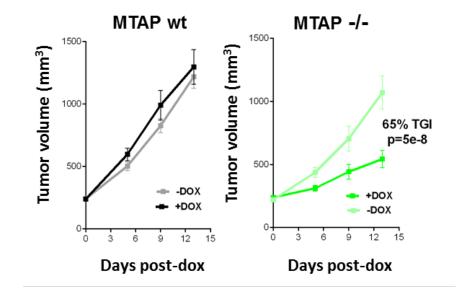


## Genetic Tools Validate MAT2A as a Selective Vulnerability in MTAP-deleted Cancers

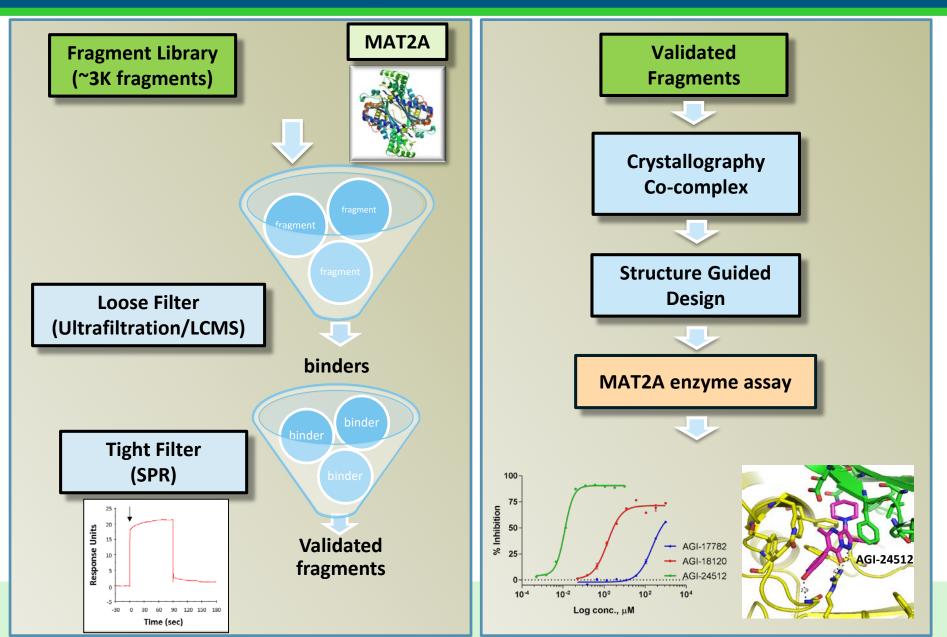
HCT116 MTAP isogenic pair



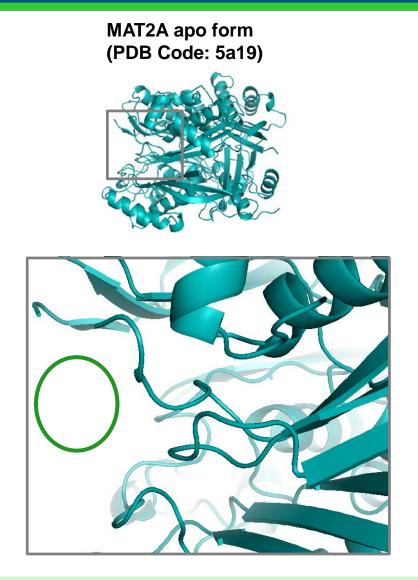




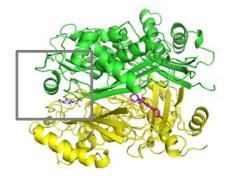
### Fragment-Based Screening for Discovery of MAT2A Inhibitors

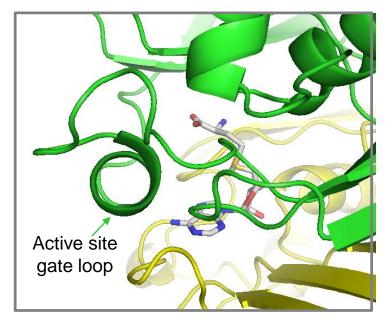


# MAT2A Inhibitor Binding in Allosteric Site Impacts Conformation of the Active Site Gate Loop

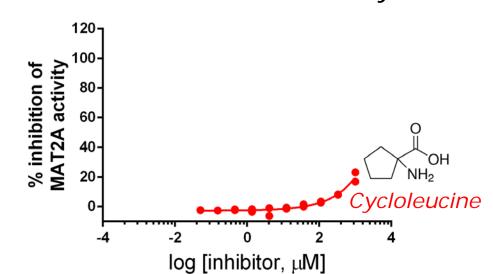


#### AGI-24512:MAT2A cocrystal

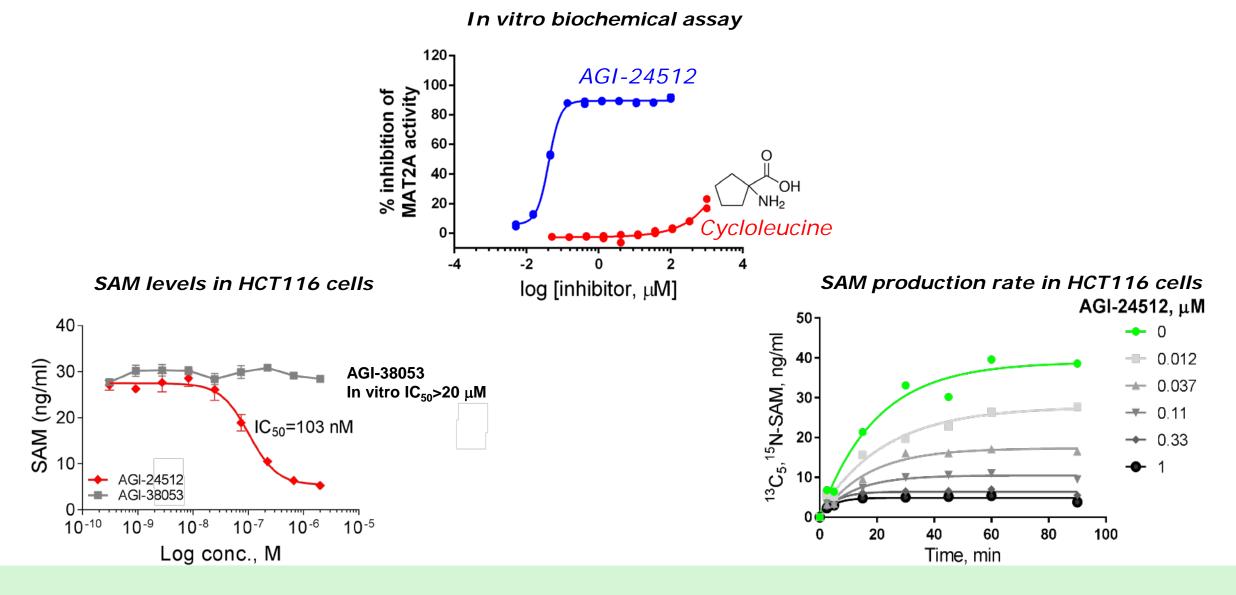




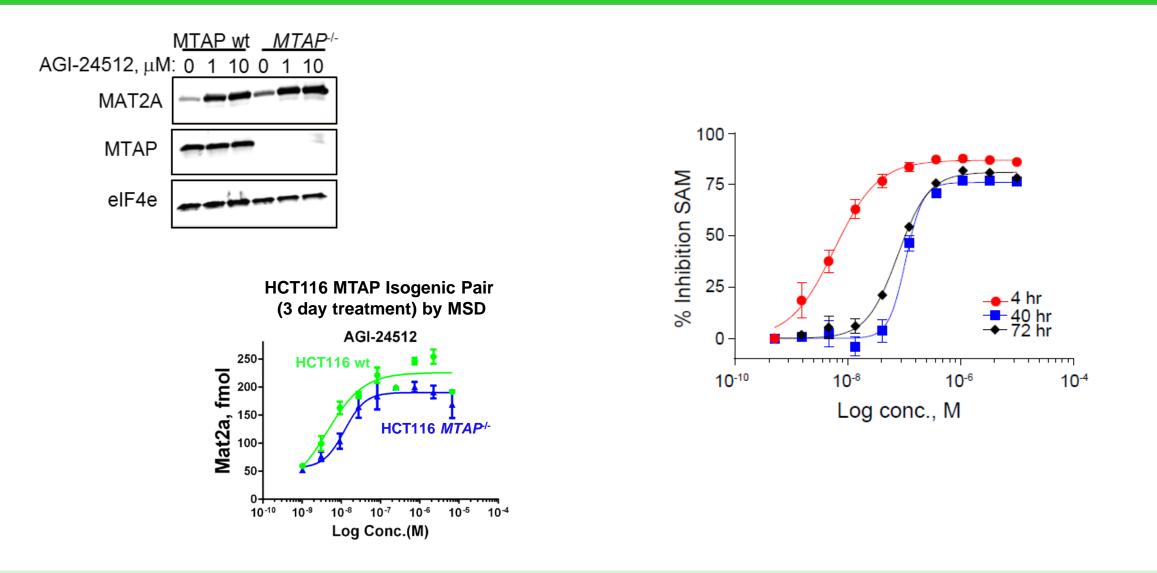
### In vitro and Cellular Activity of MAT2A Small Molecule Inhibitor



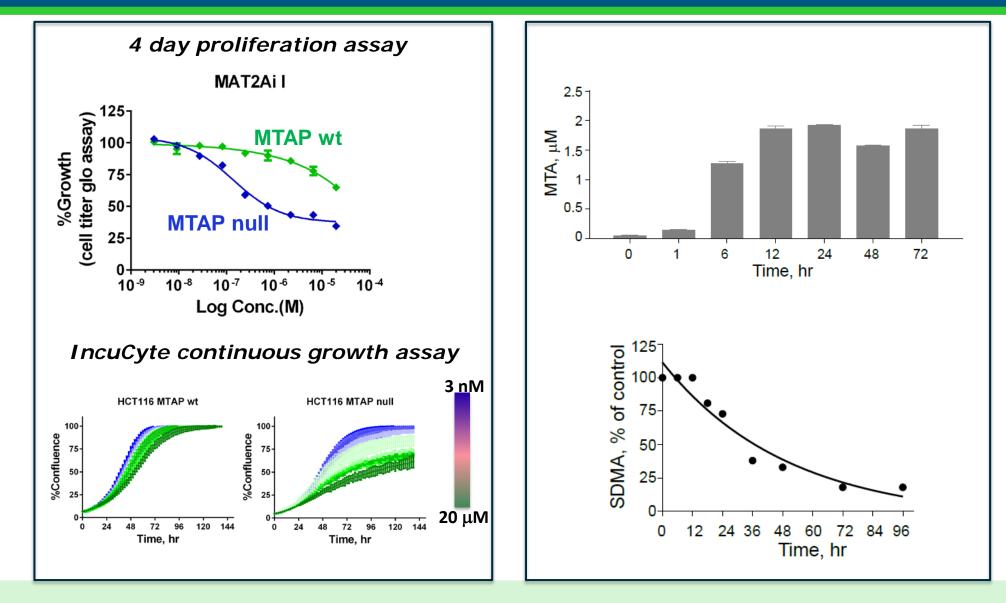
### In vitro and Cellular Activity of MAT2A Small Molecule Inhibitor



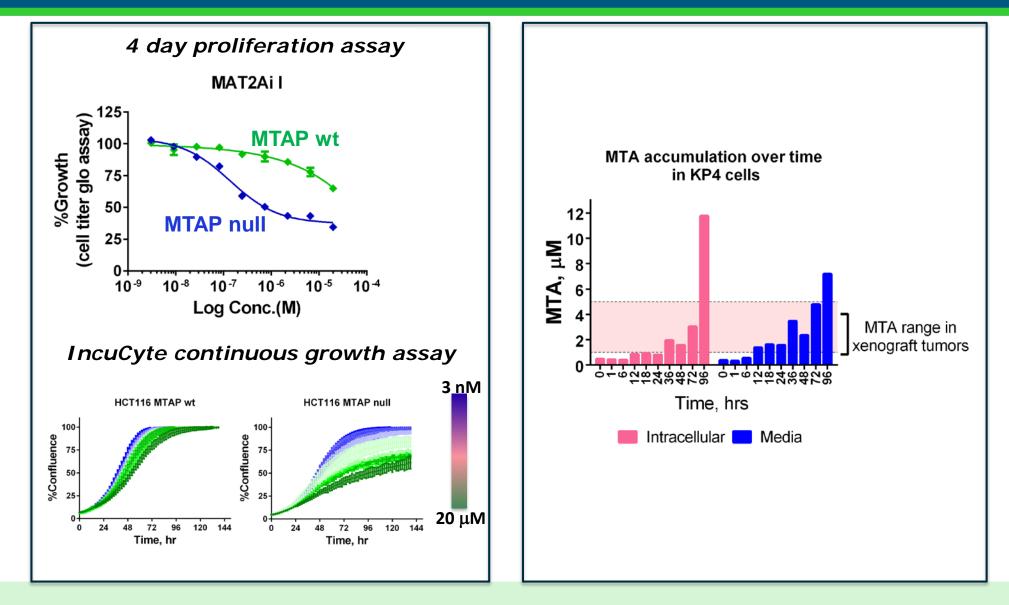
## MAT2A Inhibition Causes MAT2A Protein Upregulation and Potency Shift That is Limited and Surmountable



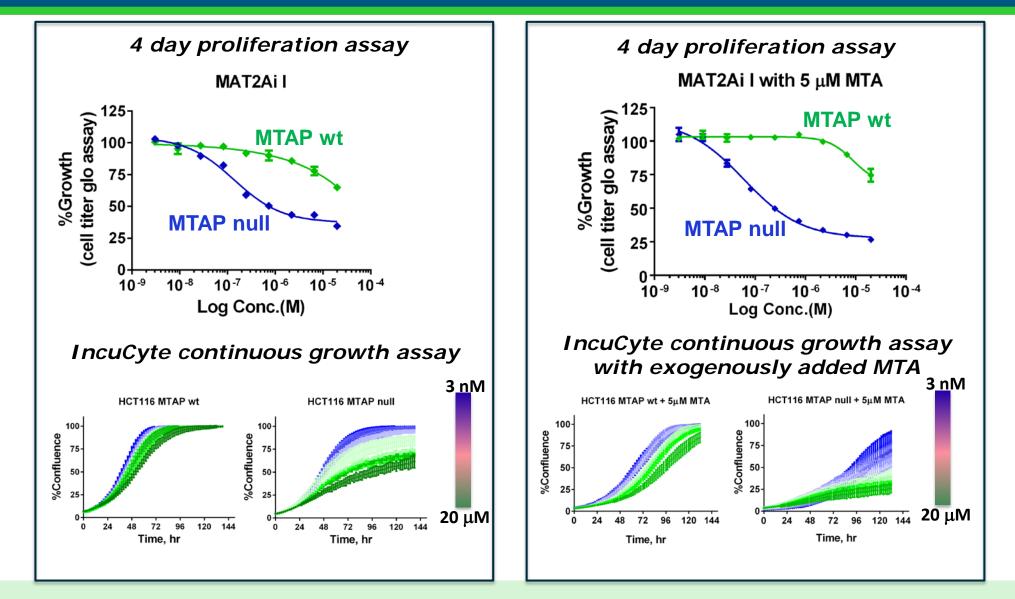
# MAT2A Inhibitors Selectively Block Growth of MTAP-deleted Cancer Cells *in vitro*



# MAT2A Inhibitors Selectively Block Growth of MTAP-deleted Cancer Cells *in vitro*



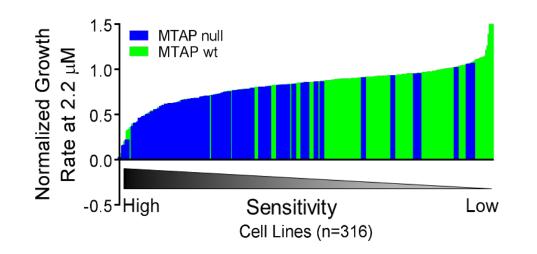
# MAT2A Inhibitors Selectively Block Growth of MTAP-deleted Cancer Cells *In Vitro*

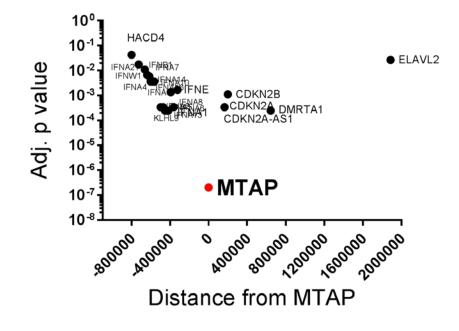


## MAT2A Inhibitors Selectively Block Growth of MTAP-deleted Cancer Cells *In Vitro*

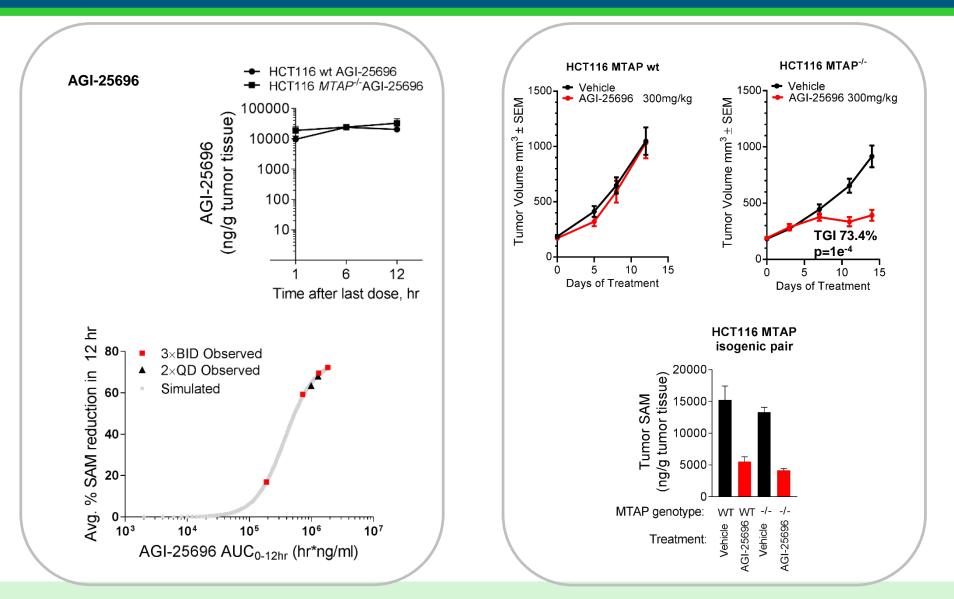
#### MTAP predicts sensitivity in Cell Panel with AGI-24512

(p=1.29e-15)

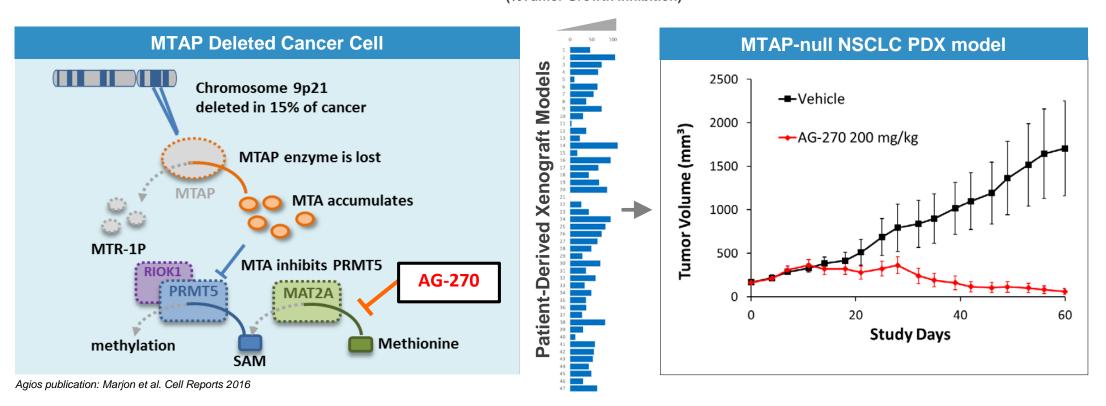




### Pharmacologic Targeting of MAT2A with *In Vivo* Tool Molecule Selectively Blocks Growth of MTAP-deleted Tumors *in vivo*



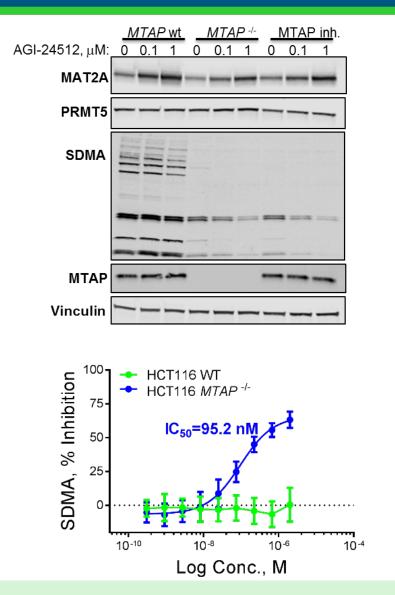
### AG-270 Active in Wide Variety of MTAP-deleted Cancer Models

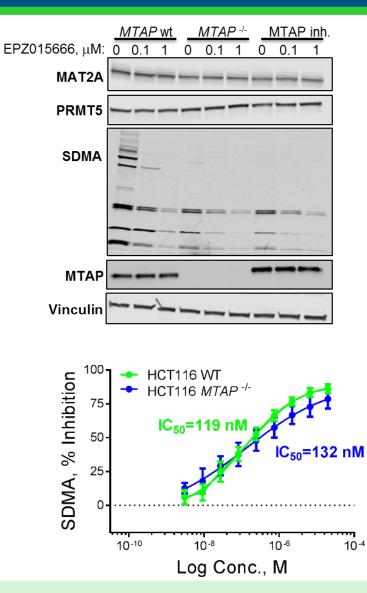


Efficacy (%Tumor Growth Inhibition)

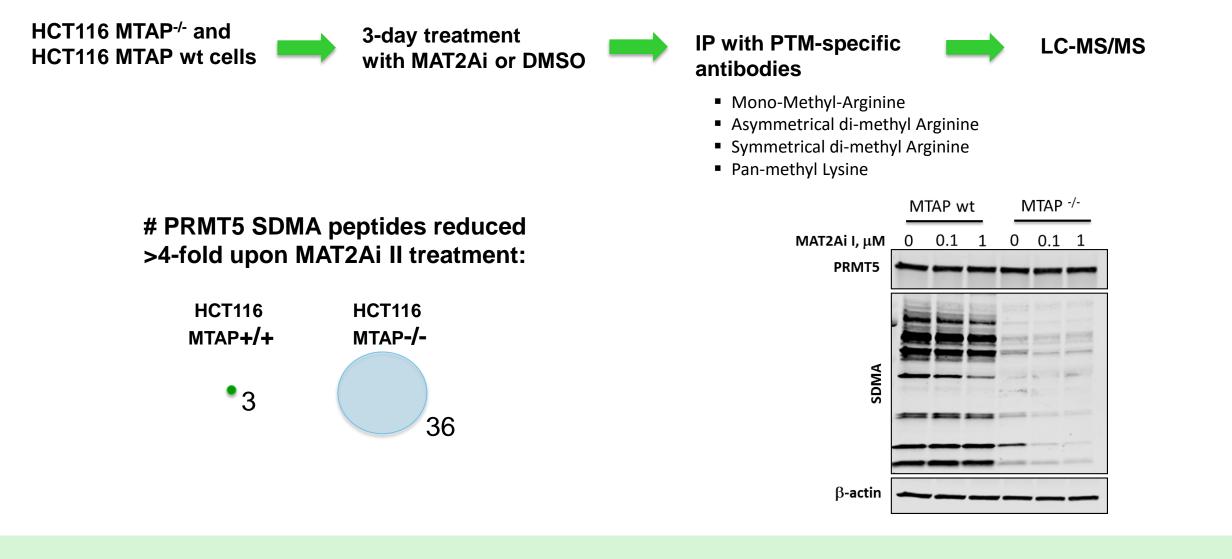
### First-in-human Phase 1 dose-escalation clinical trial started Q1 2018

# MAT2A Inhibition Selectively Impacts PRMT5 Activity in MTAP-deleted Background



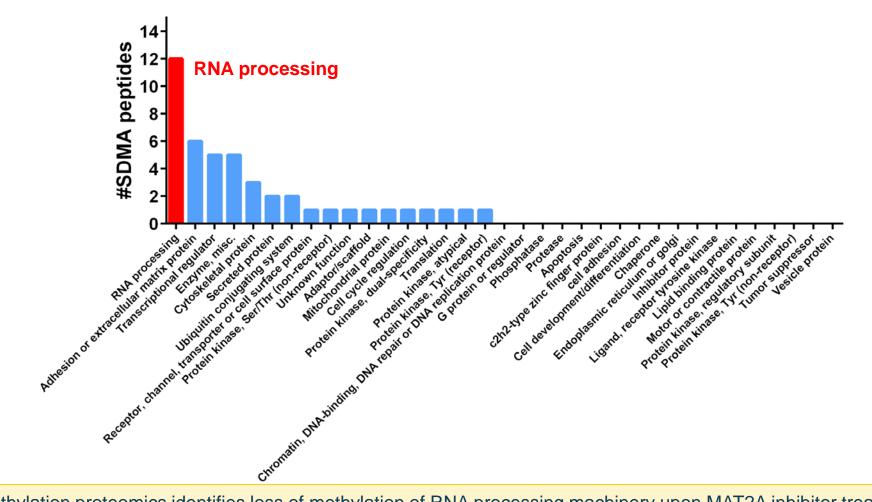


### Methylation Proteomics Corroborates Role for PRMT5 as a Key Downstream Mediator of MAT2Ai in MTAP-deleted Cells



### Methylation Proteomics Indicates MAT2A Inhibition Reduces Methylation of RNA Processing Machinery in MTAP-deleted Cells

**#SDMA** peptides that decrease upon MAT2A inhibition in HCT116 MTAP<sup>-/-</sup>

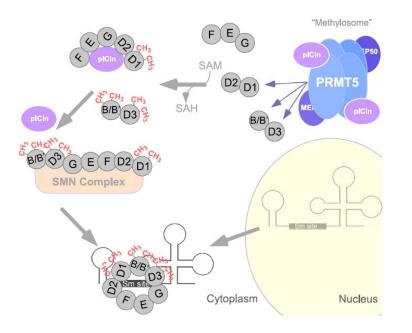


Methylation proteomics identifies loss of methylation of RNA processing machinery upon MAT2A inhibitor treatment

# Symmetric Arginine Methylation of Spliceosome Components by PRMT5 is Important for Spliceosome Maturation

#### Published substrates include:

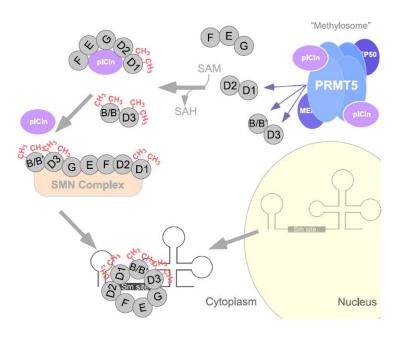
- SmD1,SmD3,SmB/B' (Brahms, RNA 2001 and Friesen Mol Cell 2001)
  - Methylation is required for interaction w/ SMN
- PRMT5 KO mouse NPCs have splicing defects (Bezzi, Genes Dev 2013)



### **PRMT5 Modulation Impacts Various Splicing Modalities**

#### Published substrates include:

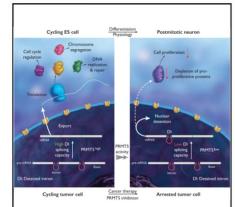
- SmD1,SmD3,SmB/B' (Brahms, RNA 2001 and Friesen Mol Cell 2001)
  - Methylation is required for interaction w/ SMN
- PRMT5 KO mouse NPCs have splicing defects (Bezzi, Genes Dev 2013)



#### Cancer Cell

Coordinated Splicing of Regulatory Detained Introns within Oncogenic Transcripts Creates an Exploitable Vulnerability in Malignant Glioma

#### Graphical Abstract



Christian J. Braun, Monica Stanciu, Paul L. Boutz, ..., Phillip A. Sharp, Michael T. Hemann, Jacqueline A. Lees

Article

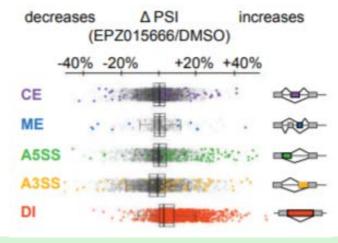
#### Correspondence

hemann@mit.edu (M.T.H.), jalees@mit.edu (J.A.L.)

#### In Brief

Authors

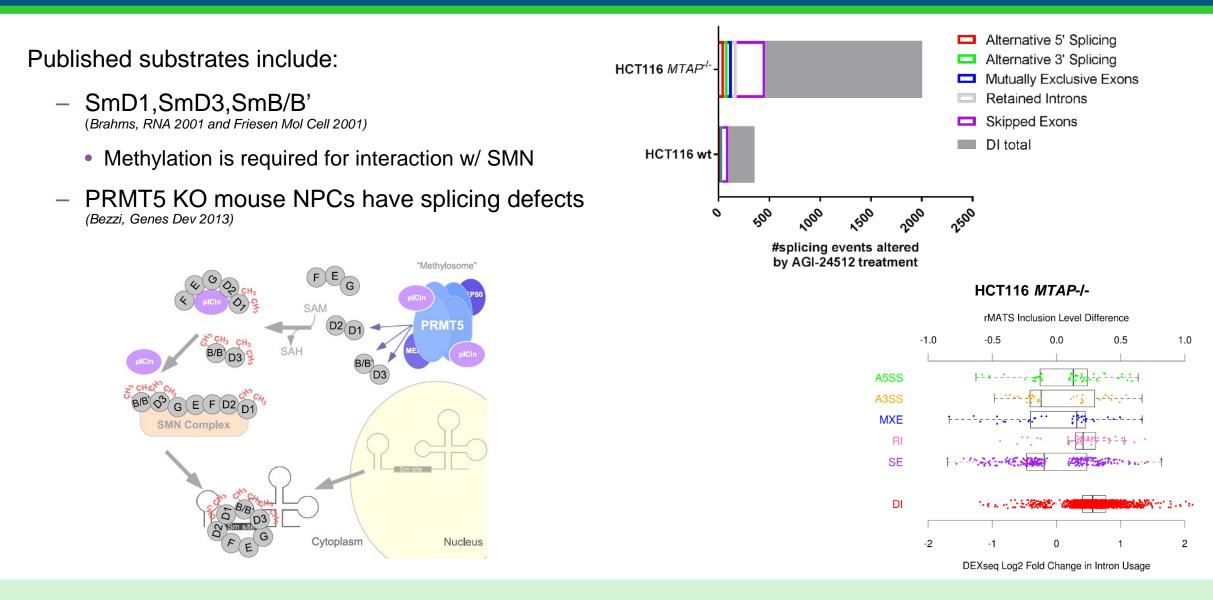
Braun et al. show that glioblastoma is selectively sensitive to the inhibition of PRMT5 and identify a predictive biomarker for this sensitivity. PRMT5 inhibition primarily disrupts the removal of detained introns, which results in the reduction of functional transcripts of mainly proliferation-associated genes.



Braun CJ et al. Cancer Cell 2017;32(4):411-26.

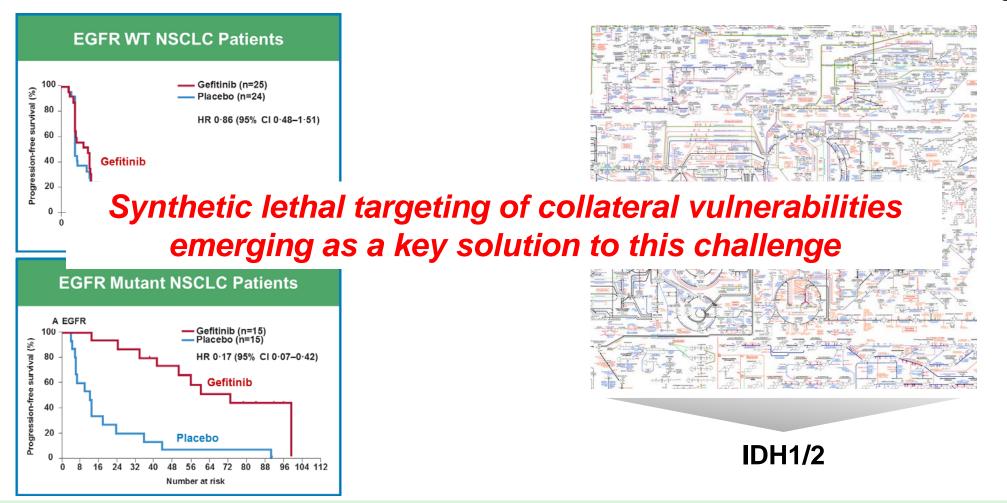
#### 29

### MAT2A Inhibition Selectively Modulates Splicing in MTAP-deleted Background



### The Challenge: Identifying Precision Medicine Approaches in Cancer Metabolism

Directly drugging 'driver mutations' has yielded transformative medicines



but...DNA sequencing has identified only 2 gain-of-function

metabolic 'driver' mutations out of 2000+ metabolic genes

Zhang L et al. Lancet Oncol 2012;13(5):466-75.

- Agios discovered potent, cell and *in vivo* active small molecule inhibitors of MAT2A
  - MAT2A inhibitors selectively block growth of MTAP-deleted cancer cells and tumors
  - Phase I clinical trial was recently initiated for AG-270

 Surprising specificity of cellular effects following reduction of universal donor of methyl groups SAM is mediated at least in part via impact on PRMT5 activity and downstream splicing biology