

Paris, March 4-6, 2013

## IMMUNOMODULATORY ACTIVITY OF SGI-110, A SECOND GENERATION HYPOMETHYLATING AGENT

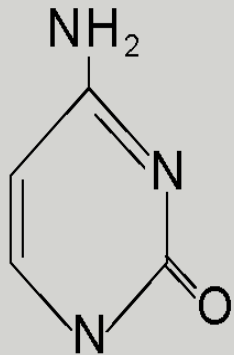
**Michele Maio**<sup>1</sup>, Alessia Covre<sup>1,2</sup>, Giulia Parisi<sup>1</sup>, Hugues JMG Nicolay<sup>1,2</sup>, Ester Fonsatti<sup>1</sup>, Sandra Coral<sup>1,2</sup>, Pietro Taverna<sup>3</sup>, Hagop Kantarjian<sup>4</sup>, **on behalf of SGI-110-1 Study Investigators**

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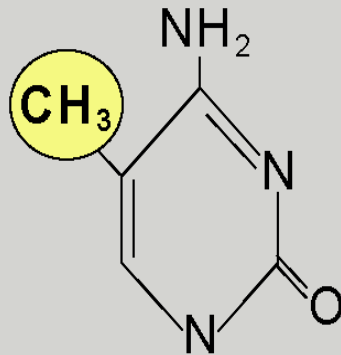
# DNA Methylation as a Therapeutic Target

- DNA methylation is abnormal in most cancers and affects the expression of key genes and pathways
- DNA methylation and epigenetic readers and writers are often mutated in cancer
  - In leukemias: DNMT3a, TET2, EZH2, ASXL1, MLL1-3, CBP etc.
- The cancer phenotype can be reversed by DNA methylation reprogramming
- DNMT inhibitors or Hypomethylating Agents (HMAs) demonstrated efficacy in the treatment of MDS and AML

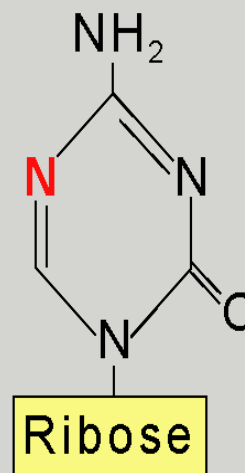
# Cytosine Analogues as HMAs



*Cytosine*

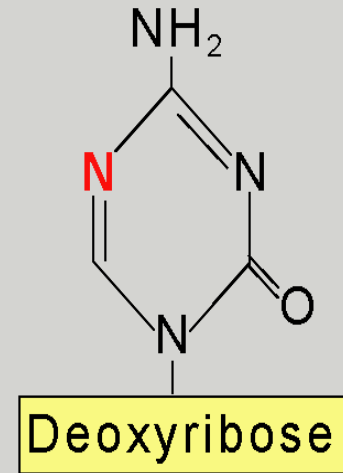


*5-methyl-  
cytosine*



*5-aza-  
cytidine*

**Azacitidine  
(2004)<sup>1</sup>**



*5-aza-2'-deoxy-  
cytidine*

**Decitabine  
(2006)<sup>1</sup> (2012)<sup>2</sup>**

<sup>1</sup>Year approved by FDA for MDS treatment

<sup>2</sup>Year approved by EMA for AML treatment

*Santini V, et al. Ann Intern Med. 2001;134(7):573-86*

# IMMUNOMODULATORY ACTIVITY OF DECITABINE

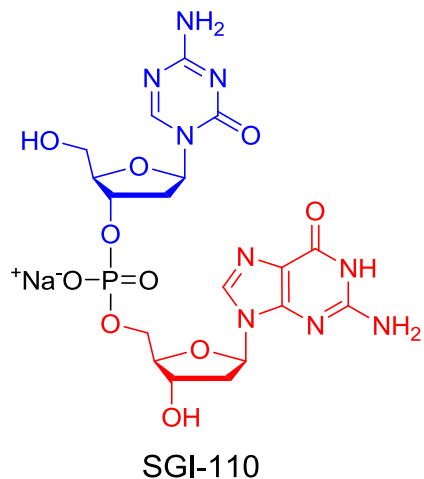
## *Pre-clinical*

- Induction/up-regulation of CTA expression in tumor cells of different histotype (Coral, Clin Cancer Res 2002)
- Up-regulation of the expression of HLA class I antigens and co-stimulatory molecules in tumor cells of different histotype (Fonsatti, Clin Cancer Res 2007)
- Increased recognition of cancer cells treated with decitabine by TAA-specific CTL (Sigalotti, Cancer Res 2004)
- Persistent induction/up-regulation of CTA expression in tumor xenografts (Coral, J Cell Physiol 2006)
- Generation of circulating anti-CTA antibodies in mice injected with decitabine-treated human melanoma cells (Coral, J Cell Physiol 2006)

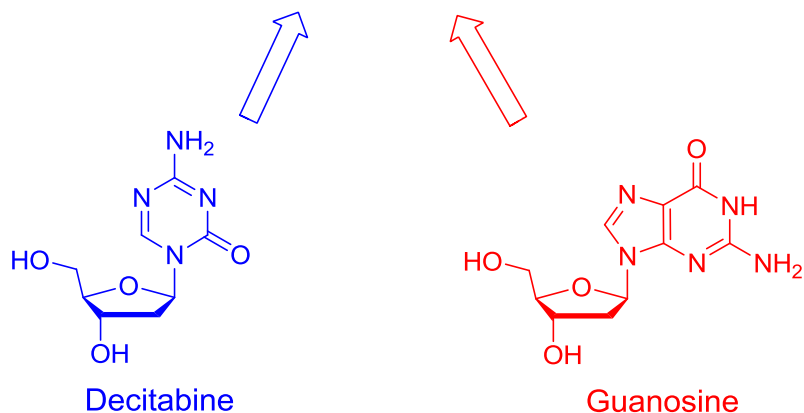
## *Clinical*

- Induction of CTA expression in AML and MDS patients (Sigalotti et al, Blood 2003)
- Post-treatment generation of circulating anti-CTA antibodies in patients with thoracic malignancies (Schrump, Clin Cancer Res 2006)
- Complete remission following decitabine/dendritic cell vaccine in a case of relapsed neuroblastoma (Krishnadas, Pediatrics, 2012)

# New DNMT Inhibitor: SGI-110

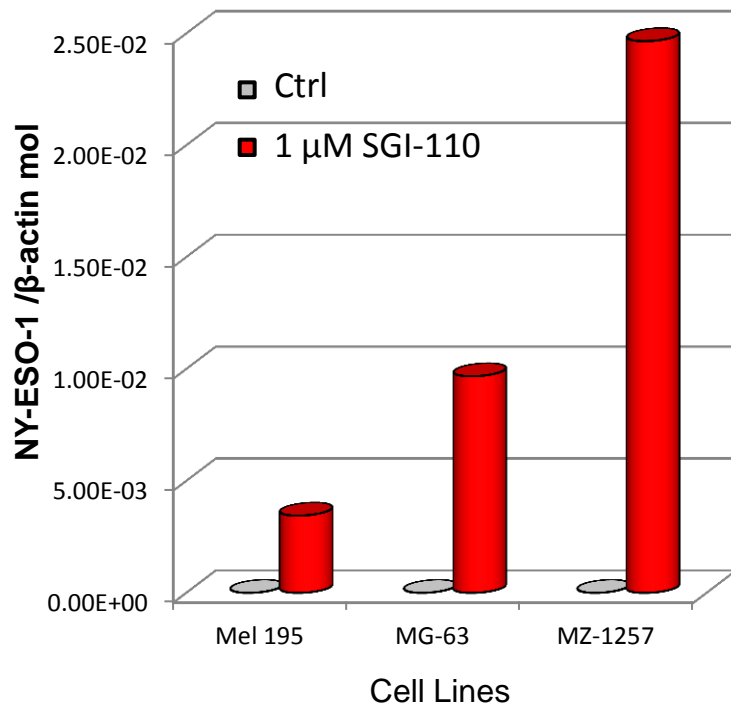


- Decitabine is rapidly eliminated by Cytidine Deaminase, limiting drug exposure time to cancer cells *in vivo*
- SGI-110 is a Dinucleotide of Decitabine and Deoxyguanosine that increases the *in vivo* exposure of decitabine by protecting it from deamination

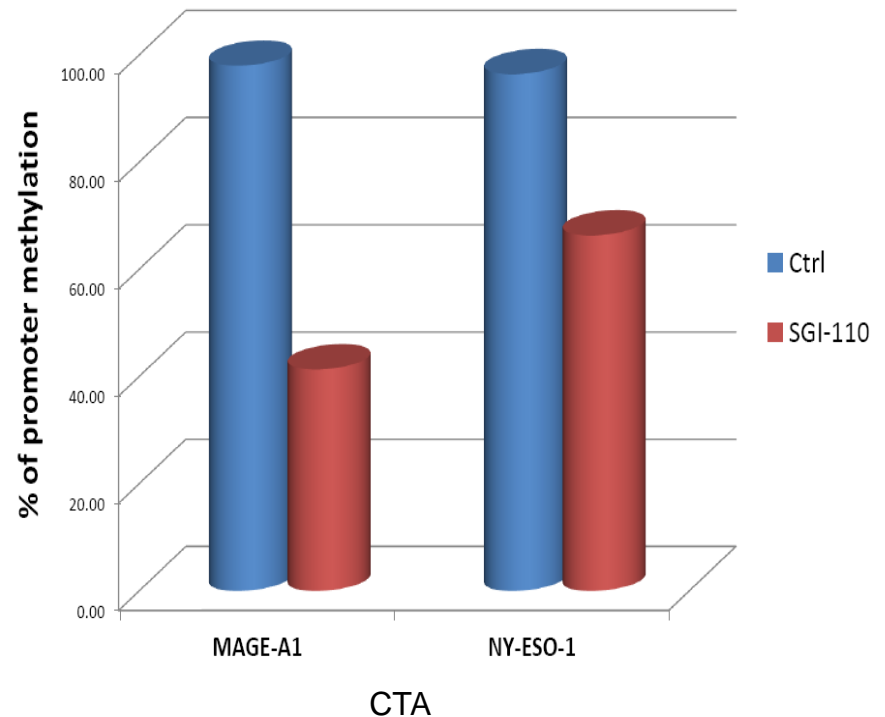


# SGI-110 Modulates CTA Expression and Methylation in Cancer Cells

## NY-ESO-1 Induction

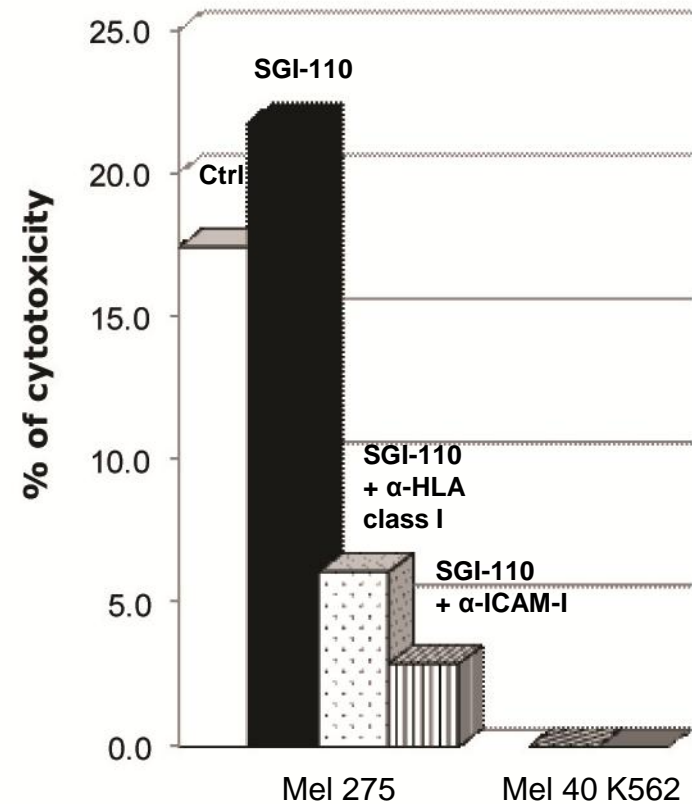
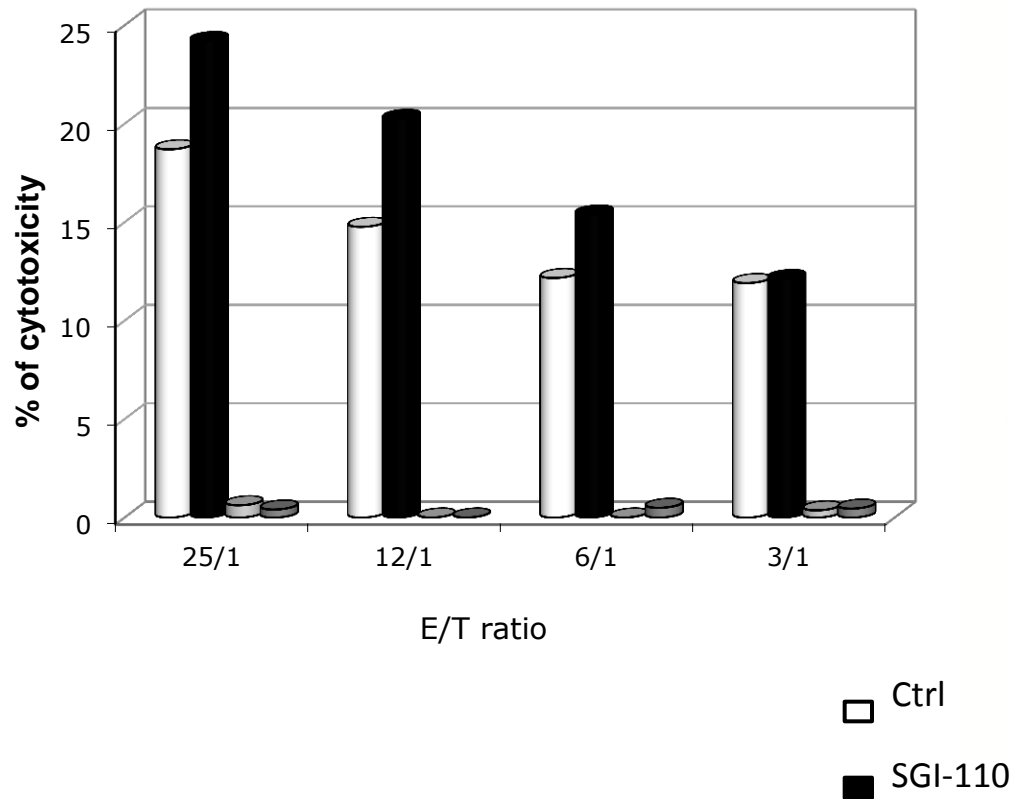


## CTA Promoter Methylation in Mel195 Cells



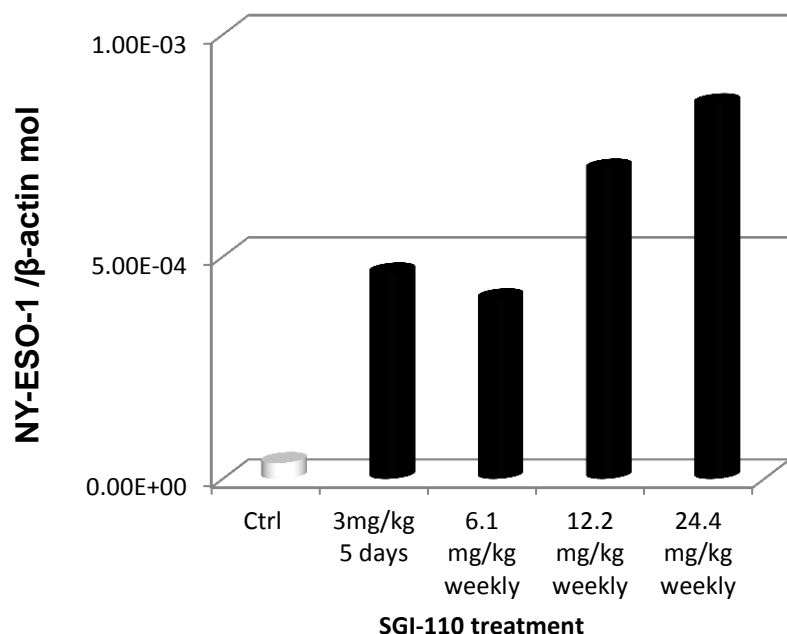
**SGI-110 induces the demethylation of CTA promoters and induces their expression**

# Recognition of SGI-110-treated Mel 275 melanoma cells by gp100-specific CTL

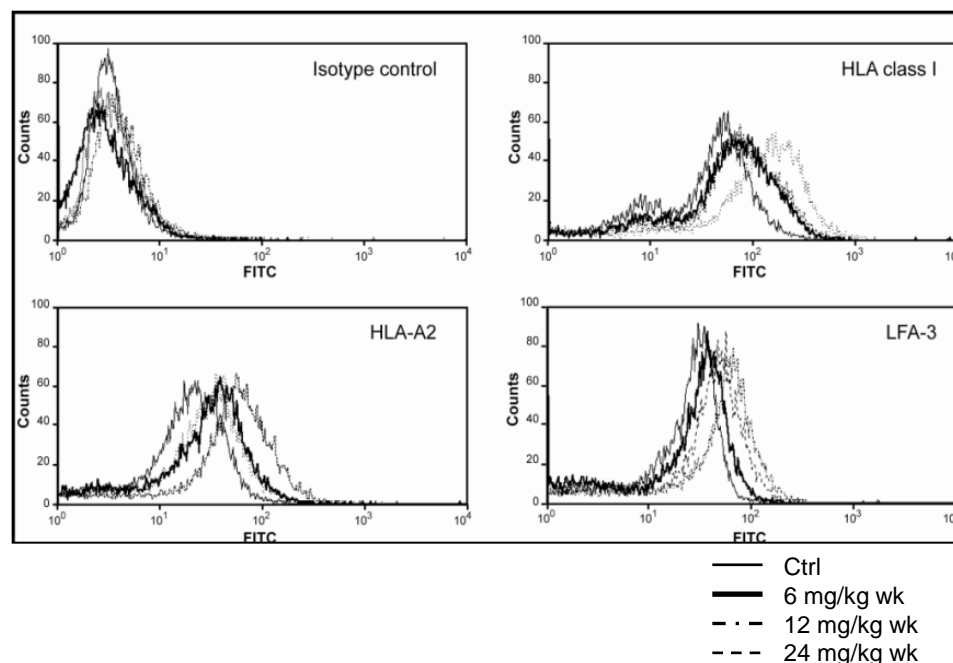


# SGI-110 Modulates CTA Expression and Immune Phenotype of Melanoma Xenografts

**NY-ESO-1 Induction in Mel313 xenografts**



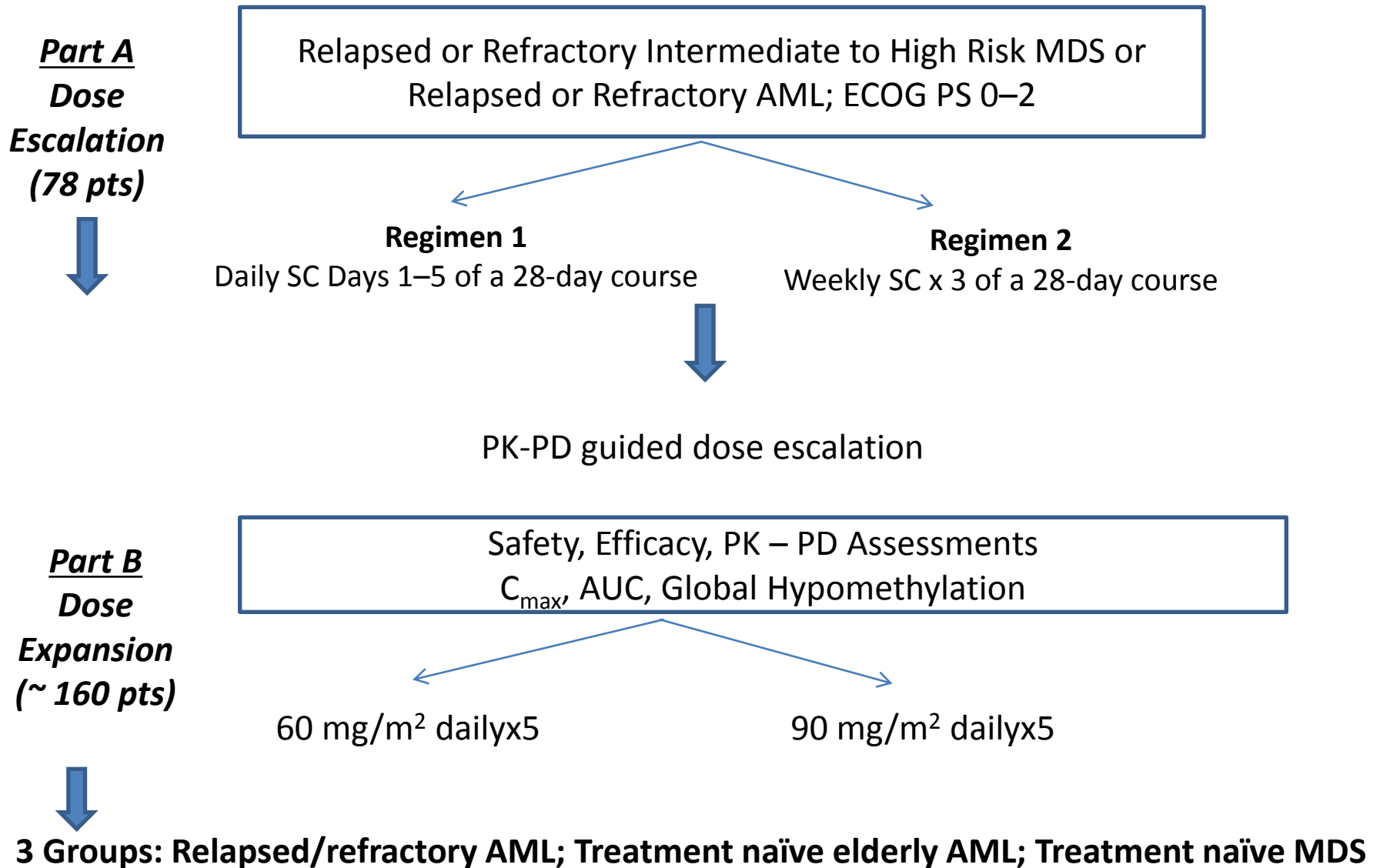
**HLA Class I and co-stimulatory molecules induction in Mel195 xenografts**



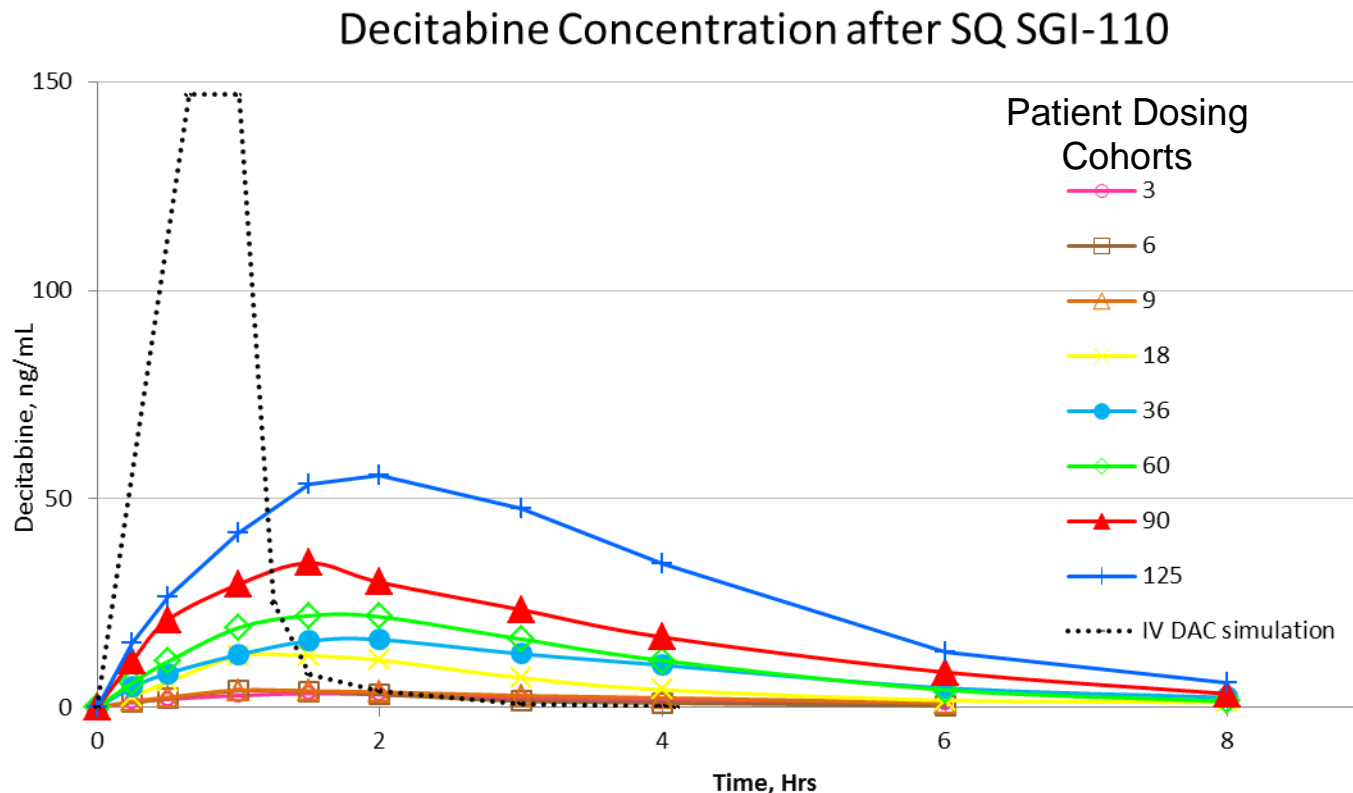
**Tolerated doses and schedules of SQ SGI-110 induces CTAs, HLA class I antigens, HLA-A2 alleles, and the co-stimulatory molecules LFA-3 and ICAM-1 in melanoma xenografts**



# SGI-110-01 Phase 1/2 Clinical Trial Design



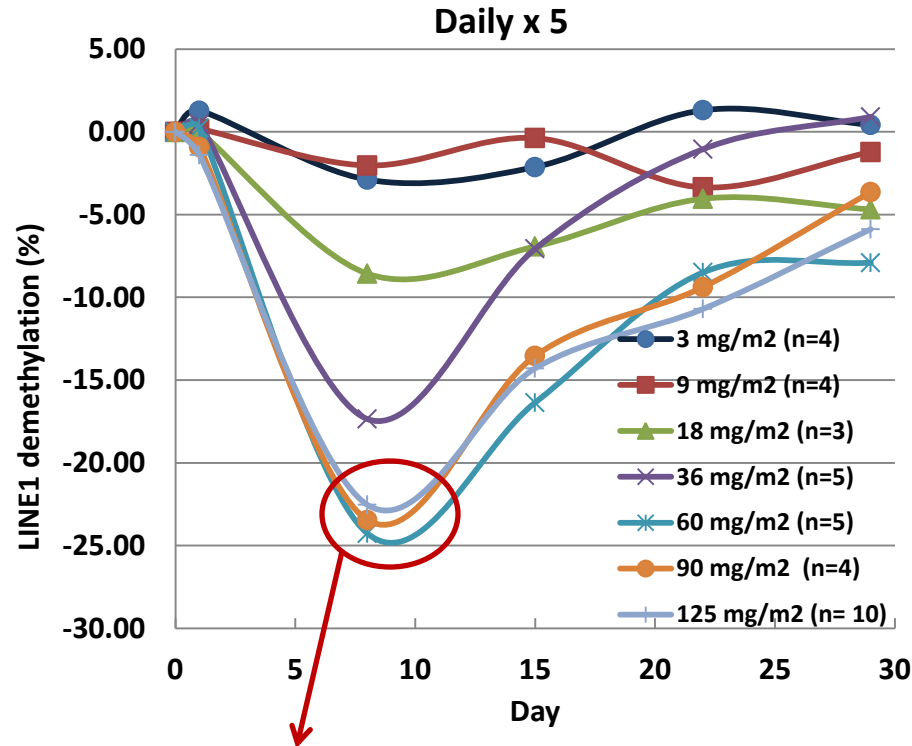
# PK of decitabine delivered by SGI-110 SQ injection



## Compared to Dacogen IV (DAC IV):

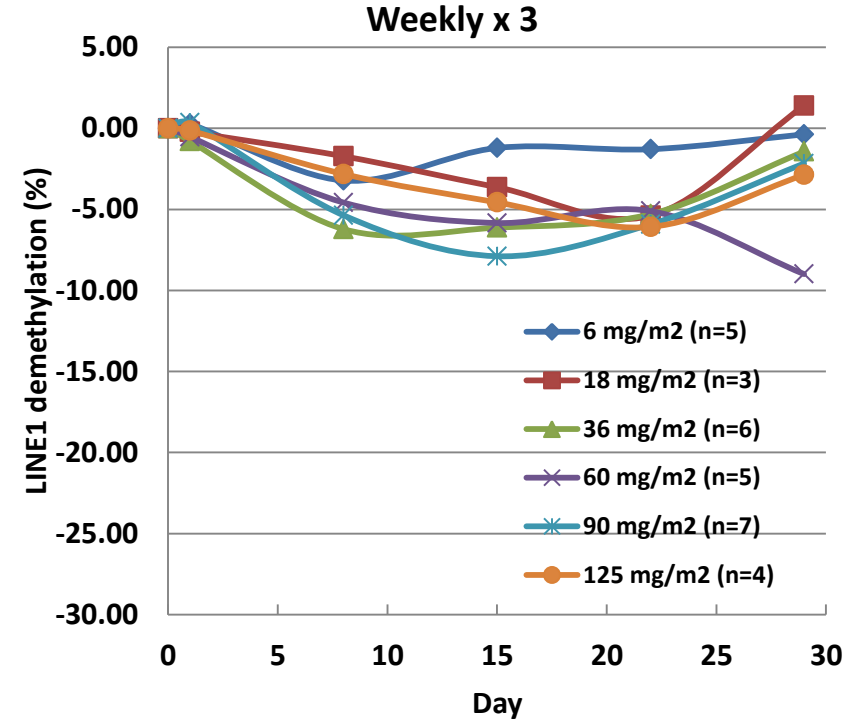
- Doubled exposure window to decitabine (8+ hrs vs. 3-4 hrs)
- Up to 4-fold longer half life of decitabine (1.5-2.5 hrs vs. 35 minutes)
- Cmax less than half of decitabine

# LINE1 Demethylation by Cohort



**BED: 60 mg/m<sup>2</sup> dailyx5**

The BED defined as the smallest dose that achieves a maximum global hypomethylation in at least three successive dose levels



# AML Responses correlated with demethylation extent

LINE1 Demethylation	Number Treated <sup>1</sup>	Responders (CR/CRi/CRp)	Percent
< 10%	31	0	0%
≥ 10%	19	5	26%
Total	50	5	10%

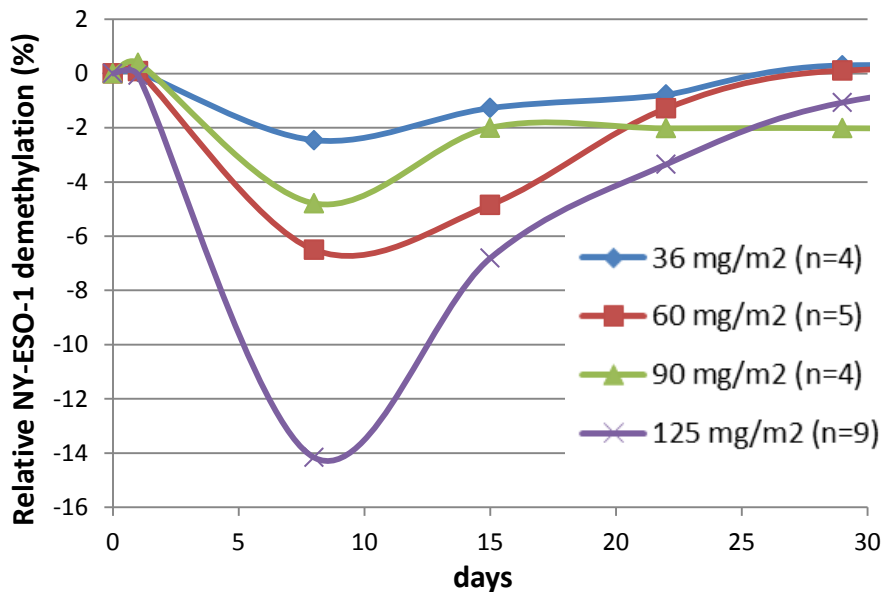
<sup>1</sup> All 50 r/r AML patients with LINE1 data

5 responses in MDS patients with prior HMA treatment  
5 responses in AML patients regardless of prior HMA treatment

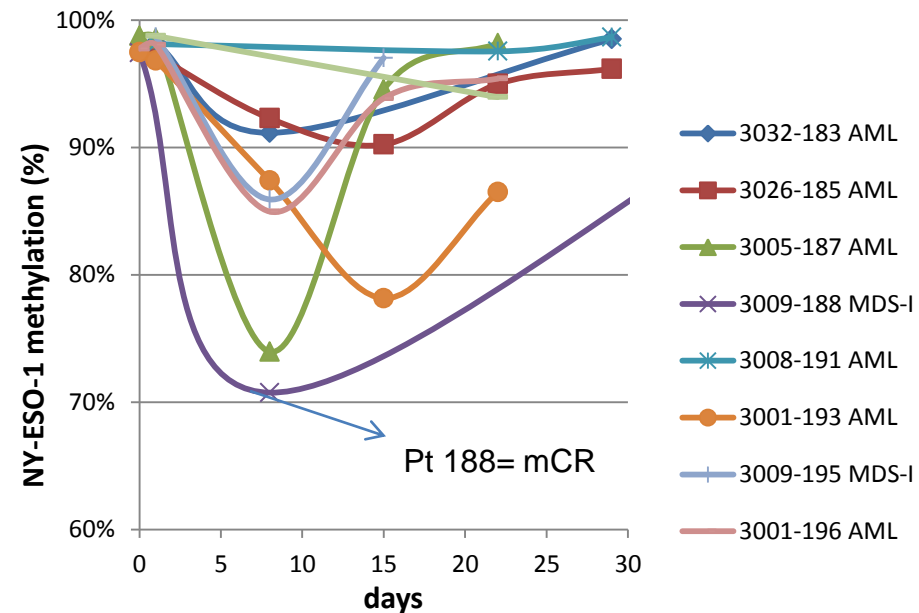
**EPIGENETIC MODULATION OF CTA  
IN BLOOD SAMPLES  
FROM PATIENTS ENROLLED IN STUDY SGI-110-01**

# NY-ESO-1 Promoter Demethylation after SGI-110 in AML and MDS patients

Average NY-ESO-1 Demethylation  
Daily regimen

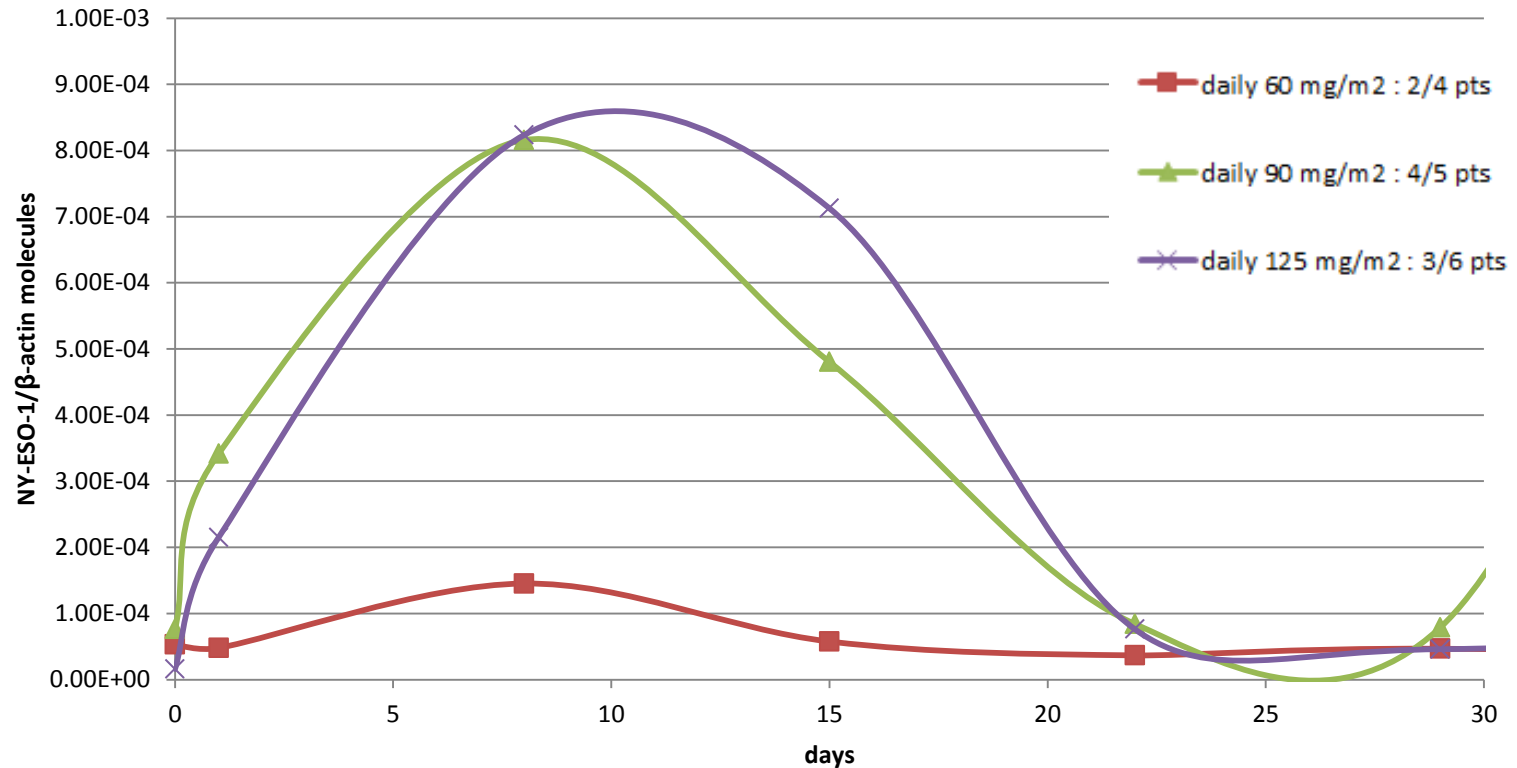


NY-ESO-1 Demethylation  
125 mg/m<sup>2</sup> daily x 5



- SGI-110 induces a dose-dependent demethylation of NY-ESO-1 promoter
- Similar extent of demethylation observed also for MAGE-A1 promoter

# NY-ESO-1 Induction (cut-off $\geq 1E-05$ ) after SGI-110 in AML and MDS patients



- NY-ESO-1 transcript was induced in 9 of 15 evaluable patients treated at SGI-110 BED
- 4 and 5 of the 15 patients induced also MAGE-A1 and -A3 respectively

# Summary

- Excellent LINE1 hypomethylation induction with dailyx5; BED is 60 mg/m<sup>2</sup> dailyx5
- Well tolerated; most common AE's were Injection site pain (mostly Grade 1) and myelosuppression (neutropenia/neutropenic fever; anemia; thrombocytopenia)
- Major responses were observed in relapsed/refractory AML when adequate hypomethylation achieved (regardless of regimen)
- SGI-110 reduced the constitutive methylation levels in promoters of NY-ESO-1 and MAGE-A1 in a dose-dependent manner
- The induction and/or up-regulation of NY-ESO-1, MAGE-A1, MAGE-A3 expression was observed in 9/15, 4/15 and 5/15 patients treated with SGI-110 biologically effective doses
- These immunomodulatory properties and its favorable PK/PD profile make SGI-110 an active agent to implement new and more effective combined chemo-immunotherapeutic approaches



# Acknowledgements: Clinical Study SGI-110-01



Hagop Kantarjian, MD  
Guillermo Garcia-Manero, MD  
Farhad Ravandi, MD



Casey O'Connell, MD  
Anthony El Khoueiry, MD



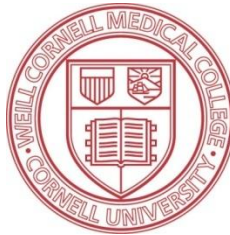
David Rizzieri, MD  
Arati Rao, MD  
Carlos Decastro, MD



Katherine Walsh, MD  
William Blum, MD



Raoul Tibes, MD, PhD  
Ruben Mesa, MD



Gail Roboz, MD  
Eric Feldman, MD  
Ellen Ritchie, MD



Steve Baylin, MD  
Peter Jones, PhD  
Jean Pierre Issa, MD



Mohammad Azab, MD  
Gavin Choy, PharmD  
Sue Naim  
Aram Oganesian, PhD  
Sanjeev Redkar, PhD



Elizabeth Griffiths, MD



Karen Yee, MD  
Aaron Schimmer, MD



Jean Pierre Issa, MD  
Woonbok Chung PhD

# Acknowledgements

## Immunomodulatory Activity of SGI-110

MEDICAL ONCOLOGY AND IMMUNOTHERAPY  
DEPT. OF MEDICAL ONCOLOGY  
UNIVERSITY HOSPITAL OF SIENA



- Alessia Covre
- Giulia Parisi
- Hugues Nicolay
- Sandra Coral
- Ester Fonsatti

**Epigen Therapeutics™**  
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- Sandra Coral
- Alessia Covre
- Hugues Nicolay