

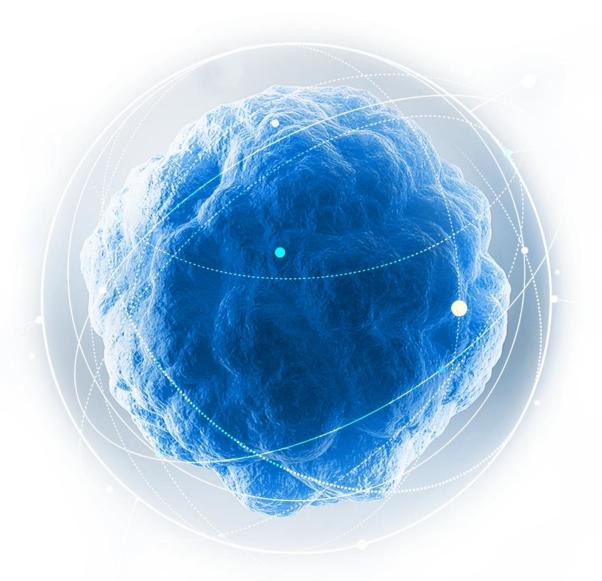


Mechanisms of BCL2i resistance in myeloid malignancies and the potential for emavusertib+venetoclax synergy

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Disclosures



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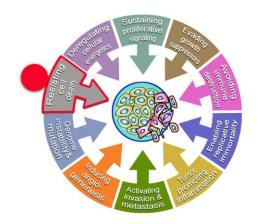
Stock options/Royalties:

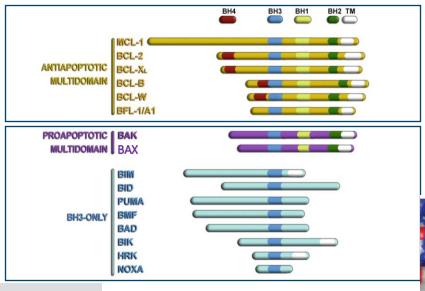
Reata Pharmaceutical



Resisting apoptosis is a hallmark of cancer and a primary cause of cancer resistance to therapy





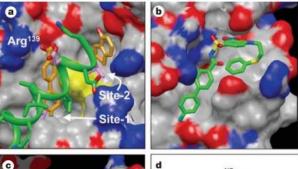


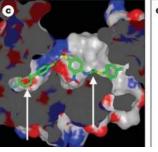
nature

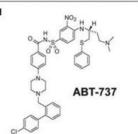
LETTERS

An inhibitor of Bcl-2 family proteins induces regression of solid tumours

Tilman Oltersdorf¹*, Steven W. Elmore²*, Alexander R. Shoemaker²*, Robert C. Armstrong¹, David J. Augeri², Barbara A. Belli¹, Milan Bruncko², Thomas L. Deckwerth¹, Jurgen Dinges², Philip J. Hajduk², Mary K. Joseph², Shinichi Kitada³, Stanley J. Korsmeyer^{4,5}, Aaron R. Kunzer², Anthony Letai⁵, Chi Li⁶, Michael J. Mitten², David G. Nettesheim², ShiChung Ng², Paul M. Nimmer², Jacqueline M. O'Connor², Anatol Oleksijew², Andrew M. Petros², John C. Reed³, Wang Shen², Stephen K. Tahir², Craig B. Thompson⁶, Kevin J. Tomaselli¹, Baole Wang², Michael D. Wendt², Haichao Zhang², Stephen W. Fesik² & Saul H. Rosenberg²



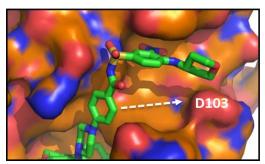


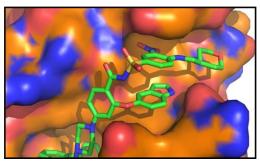


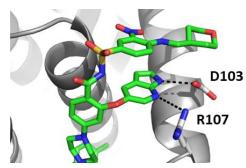


Structure-Based Design of BCL-2-Selective Inhibitor IRAK4 Symposium



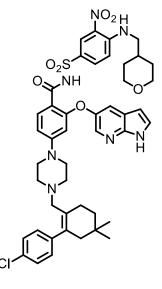






Insights from X-ray crystal structures drove the design of first-in-class BCL-2selective inhibitor, venetoclax

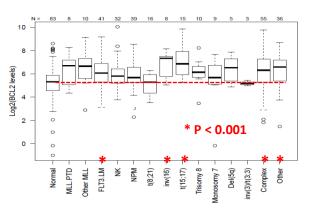
- Selective, high affinity for BCL-2
- Kills tumor cells but spares platelets
- Orally bioavailable



Venetoclax

BCL-2 as a therapeutic target in AML

BCL2 mRNA expression N = 370



- BCL-2 is highly expressed in AML blasts and stem /progenitor cells;
- Venetoclax effectively kills AML cells in vitro and in vivo
- **BCL2** targeting toxic to AML > CD34+ cells



Phase 3 trials validate BCL2 as a therapeutic target in AML

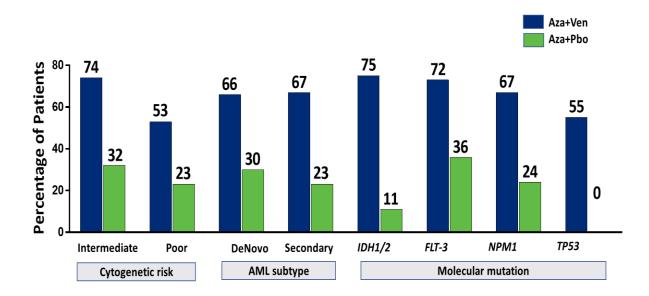


CR rate: 36.7% vs 17.9% (*P* < .001)

CR/CRi rate: 66.4% vs 28.3% (*P* < .001)

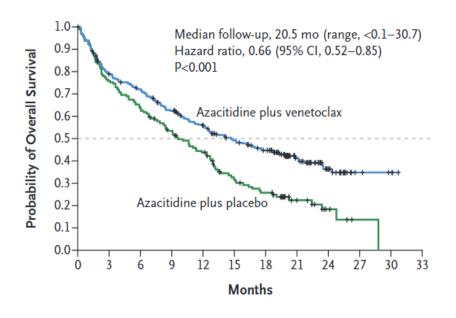
Median time to response: 1 vs 3 cycles (P < .001)

Improved responses occurred independent of high risk genomics



Significant OS improvement with venetoclax/azacitidine

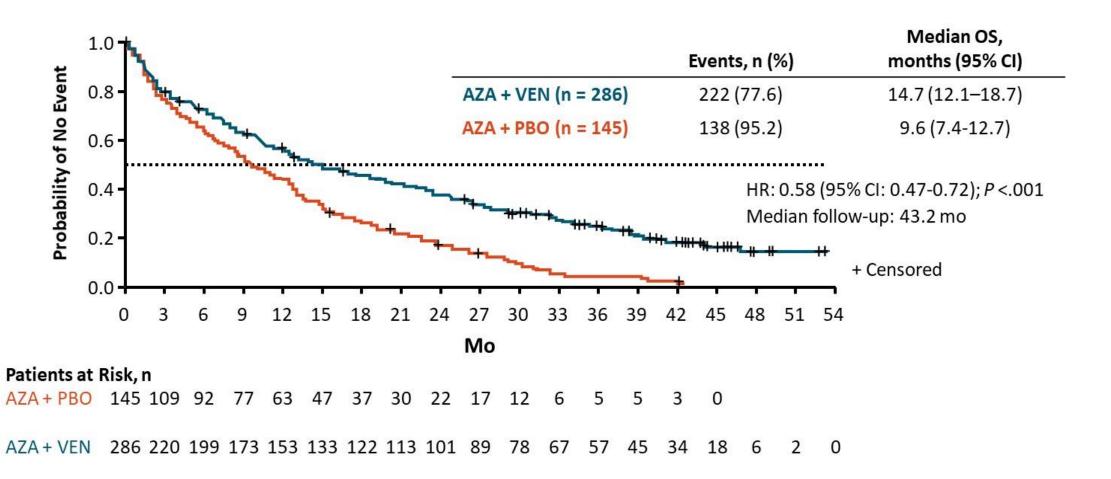
	Median OS, mo (95% CI)
VEN + AZA	14.7 (11.9-18.7)
AZA + placebo	9.6 (7.4-12.7)





Long-Term VIALE-A Follow Up: Cure remains rare and elusive

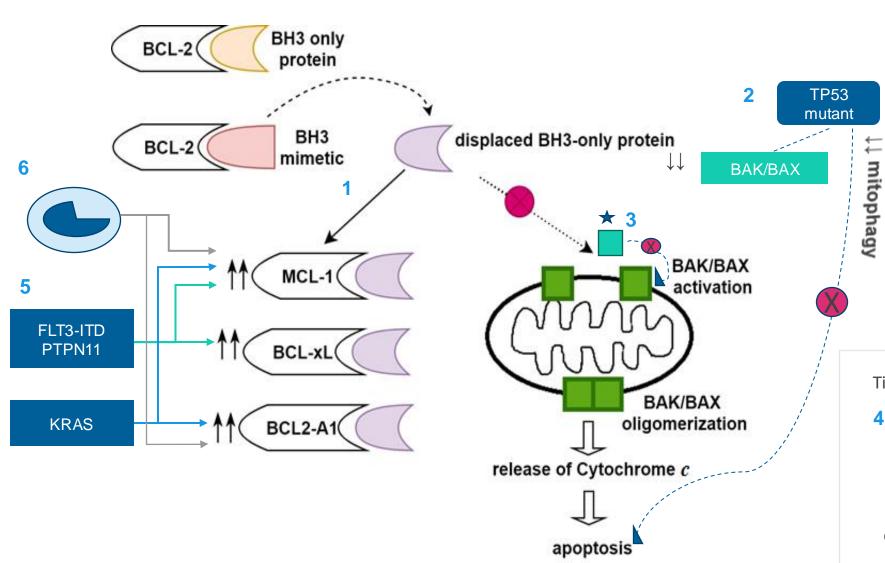






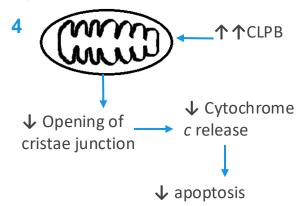
Multi-factorial Venetoclax Resistance in AML





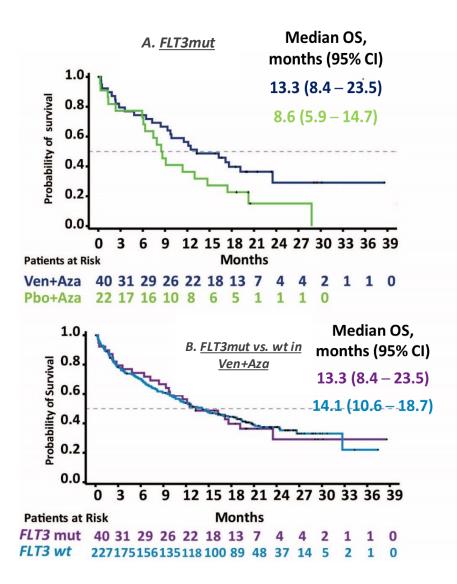
- 1. Other anti-apoptotic BCL-2 family proteins
- 2. TP53 mutation
- 3. BAX mutation
- **4.** Mitochondrial aberration
- Mutations in activating kinases
- **6.** Monocytic differentiation

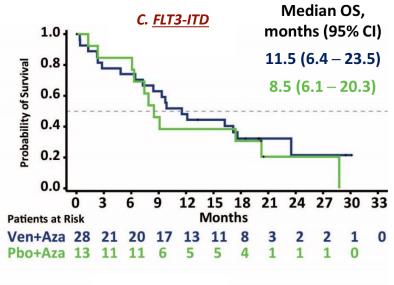
Tighter mitochondrial cristae lumen

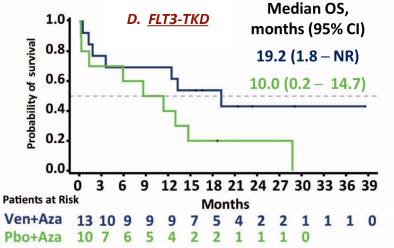


FLT3-mut AML: Subset Analysis Phase Ib and VIALE-A





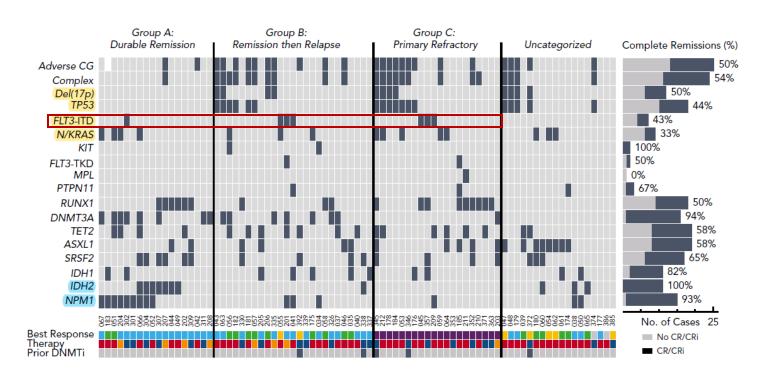


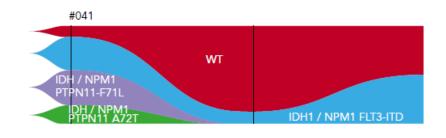


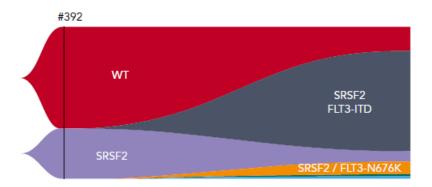


FLT3-ITD AML: Primary Refractory or Early Relapse



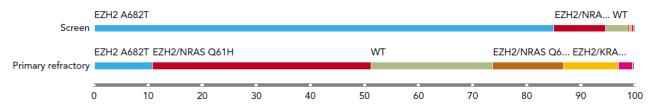






Patients treated at MDACC and The Alfred (n=81)

#064									
Sample	Date	Blast%	EZH2	EZH2/NRAS Q61H	WT	EZH2/NRAS Q61K	EZH2/KRAS Q61H	EZH2/KRAS G13D	EZH2/NRAS Q61R
Screen	03/31/2017	24	85.02%	9.57%	3.99%	0.59%	0.16%	0.55%	0.12%
Refractory	05/03/2017	53	10.85%	40.35%	22.52%	13.18%	10.12%	2.57%	0.41%

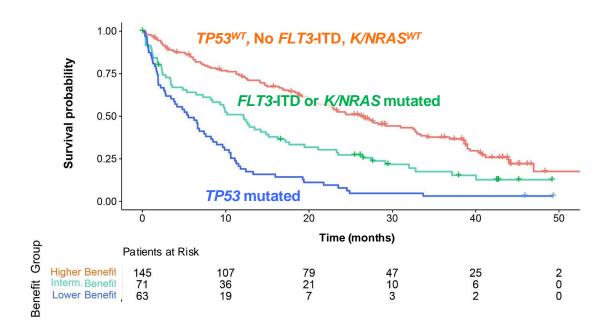




Ven+Aza: Genotype-specific Prognostic Model



- First a higher benefit group was identified, with a median OS > 24 months
- Subsequently a lower benefit group was determined, with a median OS < 6 months
- Patients fitting neither criteria were categorized as the intermediate benefit group, with a median OS of 12 months



Ven + Aza (N = 279)	n	Events	Median OS, months (95% CI)
Higher Benefit	145	96	26.51 (20.24, 32.69)
Intermediate Benefit	71	57	12.12 (7.26 – 15.15)
Lower Benefit	63	61	5.52 (2.79 – 7.59)

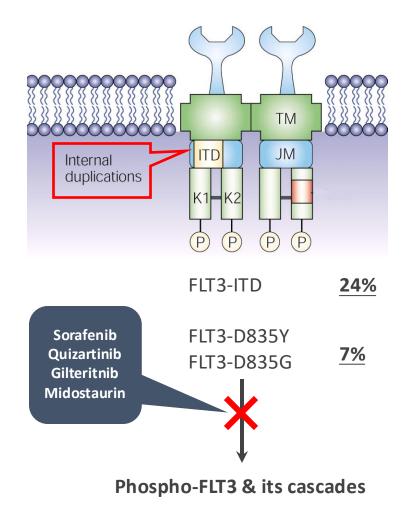
- Majority of patients in the Ven+Aza arm are in the higher benefit group: 52% (145/279)
- The remainder of the patients are distributed equally between the intermediate and lower benefit groups: 25.4% (71/279) and 22.6% (63/279), respectively

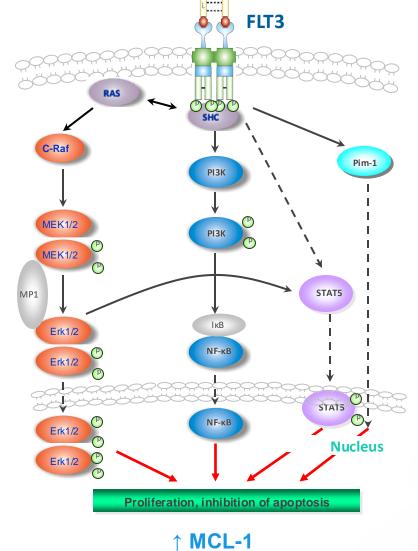


Dohner H et al, ASH 2022

Mutant FLT3: Oncogenic Signaling and MCL-1



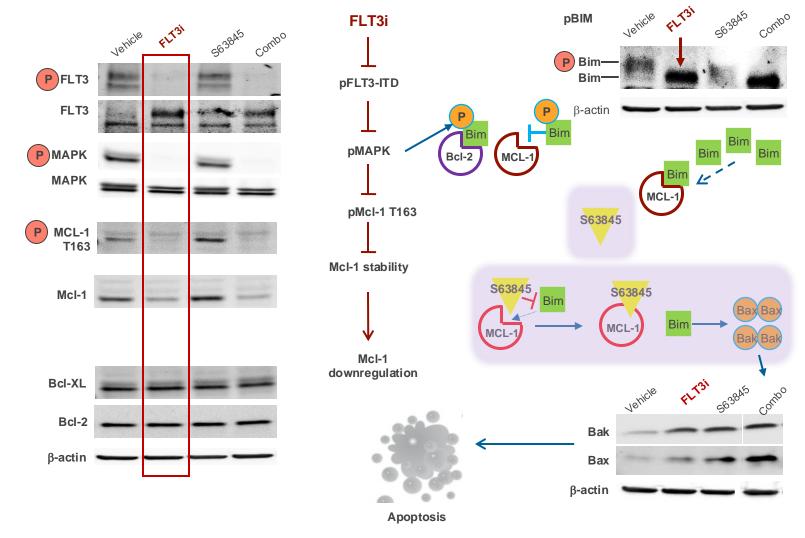






FLT3 inhibitor Targets both MCL-1 and BIM: Synergy with BCL-2 or MCL-1 inhibitors

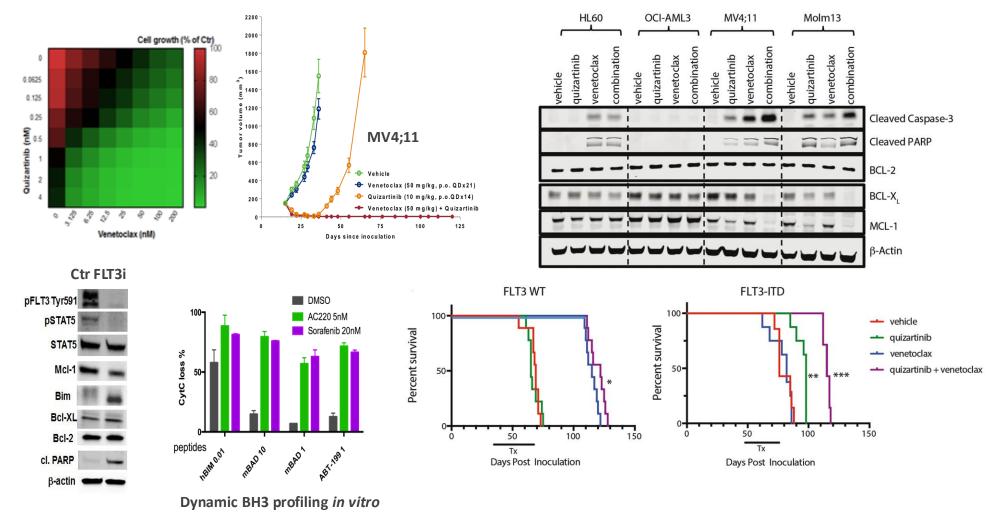






FLT3 Inhibitors and Venetoclax: Synergy and Priming

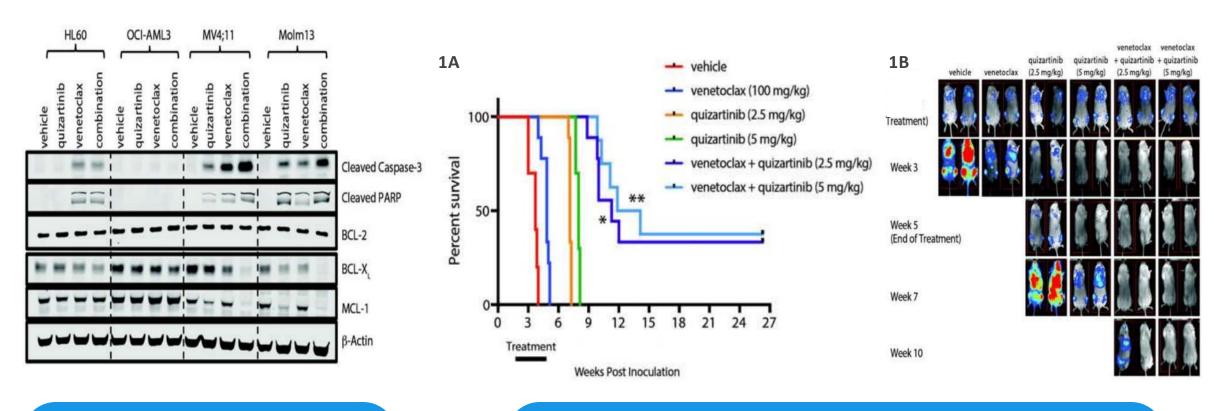






Venetoclax Combines Synergistically With FLT3i's





Cell lines were treated with combination

-↓ MCL-1, ↓ BCL-X_L

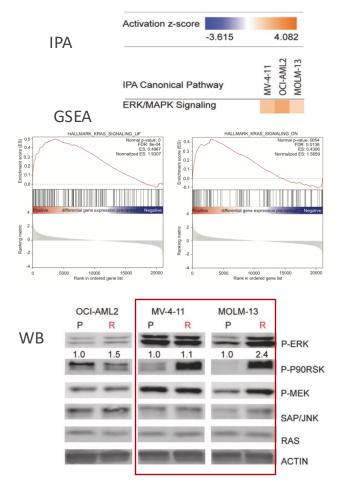
Venetoclax combined with quizartinib prolonged survival and reduced tumor burden in FLT3-ITD+ xenograft models



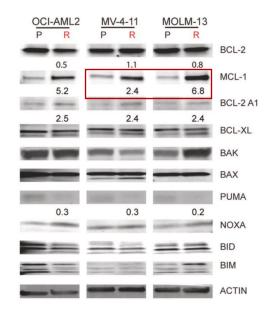
Adaptive Venetoclax Resistance in FLT3m AML: Activation of RAS/MAPK pathway and MCL-1 stabilization



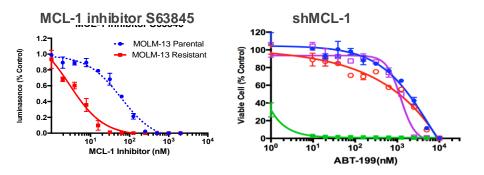
VEN-RE cells have enriched ERK/MAPK pathway



Increased MCL-1 expression



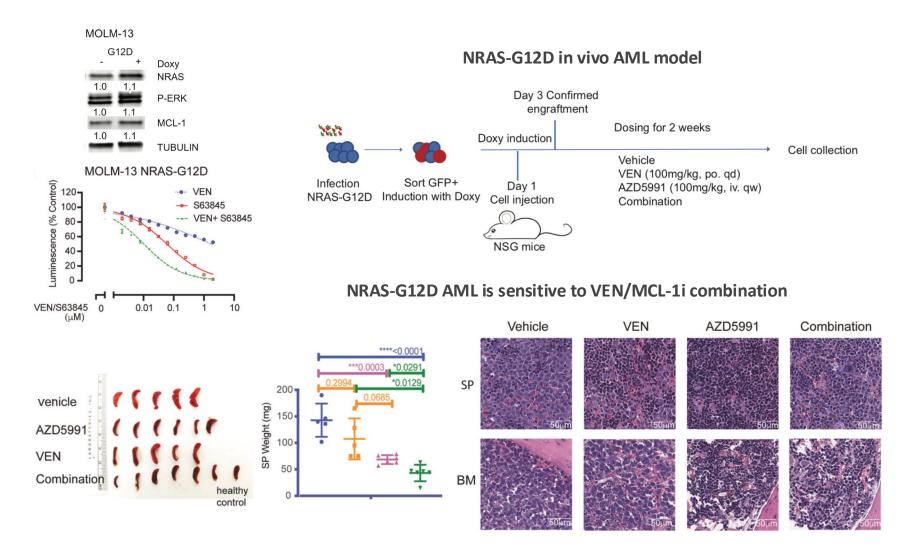
VEN-RE cells are sensitive to MCL-1 inhibition





VEN combined with MCL-1 inhibitor in VENresistant FLT3-ITD/NRASG12D AML in vivo

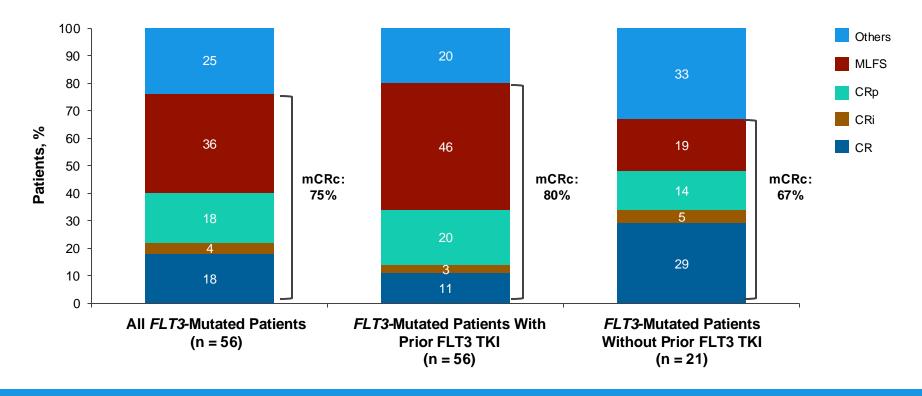






VEN + GILT: A Backbone to Build a Frontline Triplet ^{1,2}





Median salvage 2-3

Prior FLT3 TKI exposure: 60%

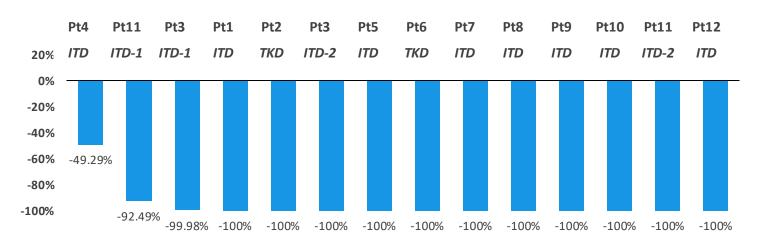
The mCRc rate in this study was 75%, whereas the CRc rate in the ADMIRAL phase 3 study for single-agent GILT was 54.3% (using the same response parameters)



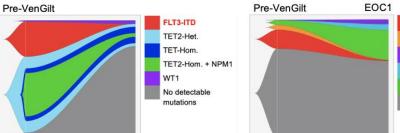
Decrease of *FLT3* clones with VenGilt treatment



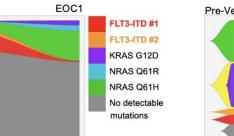
Maximal % Decrease from Baseline for FLT3-Containing Clones



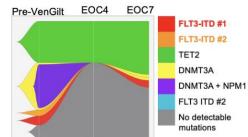
Decrease of FLT3-ITD to undetectable levels



Decrease but not elimination of FLT3-ITD



Decrease and subsequent rise of FLT3-ITD



- All 14 FLT3^{mut} clones decreased in size on therapy
- 11 clones from in 9 patients decreased to an undetectable level at maximum response
- 2 clones returned at a later timepoint
- Response was frequently rapid, with maximal decrease of FLT3 by cycle 1 day 28 of therapy in 5 of 8 evaluable patients
- 7 patients had matching timepoints where FLT3-ITD was evaluated by MyFLT3 specific MRD assay. Sensitivity of decrease ranged between 10⁻² to 10⁻⁶



Aza+Ven+Gilteritinib in Frontline FLT3-mutated AML: Healthier marrow, potentially more curative and better tolerated



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Induction

Azacitidine
75 mg/m² IV/SC on D1-7
Venetoclax R/U to goal 400mg D1-14
Gilteritinib 80 mg on D1-14
(--if blasts <5% on D14, hold both GV
--if blasts >5% on D14 continue GV and repeat BM in 1 week)

N = 30

CR 92% CRi 4% CR+CRi: 96%

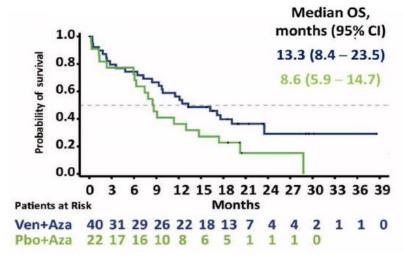
Recovery: ANC ≥0.5 37d Plt ≥50 25d Consolidation (up to 24 cycles)

Azacitidine 75 mg/m² IV/SC on D1-5

Venetoclax 400mg on D1-7

Gilteritinib 80 mg on D1-28

Historical perspective (Konopleva M et al CCR 2023)
AZA+VEN in FLT3m frontline AML (N=40)



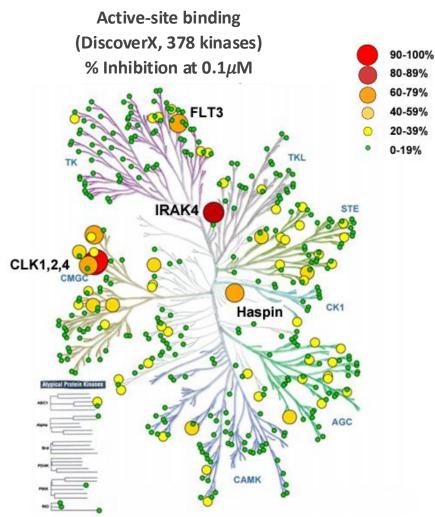


Short N, Daver N, et al. J Clin Oncol. Jan 2024

Emavusertib (CA-4948)

IRAK4 Symposium

A Small Molecule Kinase Inhibitor



CA-4948 Binding Affinity Activity

	DiscoverX
Kinase	K _d (nM)
IRAK4	23
IRAK1	12,000
IRAK2	>20,000
IRAK3	8,500

Other top hits:

raior top into				
CLK1	10			
CLK4	14			
CLK2	20			
FLT3	31			
DYRK1A	25			
Haspin (GSG2)	32			
TrkA	130			

- First-in-class IRAK4 and FLT3 inhibitor for cancer patients currently being investigated in Phase 1/2 dose-escalation trials
- Binds with high affinity to IRAK4, FLT3, CLK and DYRK1A
- Multi-kinase inhibitor with strong preclinical evidence to over come Venetoclax resistance.
- CA-4948 targets transcriptional and posttranslational regulation of MCL-1

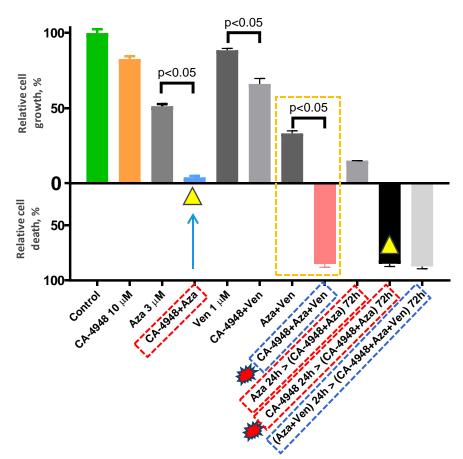


CA-4948

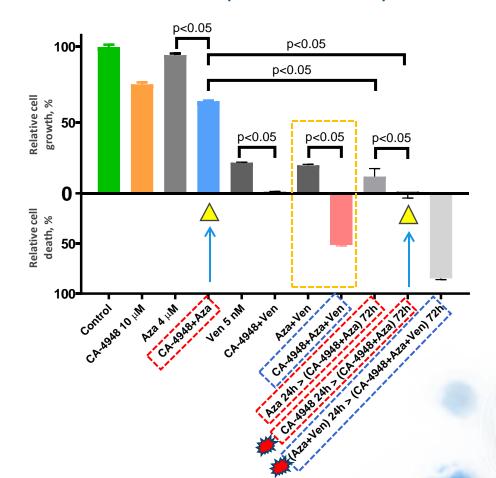


Combination of CA-4948 with azacitidine and venetoclax in FLT3-WT AML cell lines

THP-1 (venetoclax resistant)



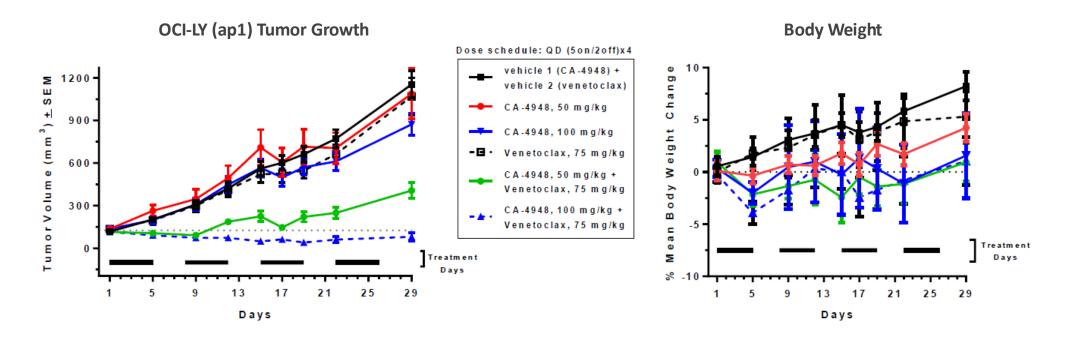
OCI-AML2 (azicitidine resistant)





Efficacy of CA-4948 + Venetoclax Combination Dosed 5on/2off in the DLBCL OCI-LY10 (ap1) Model





- OCI-Ly10 (ap1) cells: DLBCL-ABC, MYD88-L265P, CD79A
- CA-4948 + venetoclax combination on a 5on/2off dose schedule induced tumor stasis and regression with CA-4948 at low and high dose, respectively
- The CA-4948 + venetoclax combination was tolerated with the 5on/2off schedule, but minor skin/fur/rough coat signs were present in 50 and 100 mg/kg CA-4948 (single and combinations) treated animals on days 17 and 17/19, respectively.

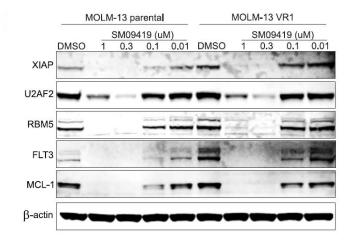


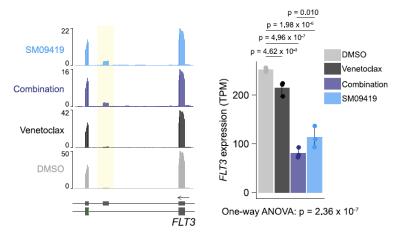
Modulation of RNA splicing enhances response to BCL2 inhibition in leukemia



Cancer Cell

CLK inhibition induces splicing alterations of key survival genes in AML





- CLK inhibition is associated with splicing changes by increased intron retention within the transcripts of RNA splicing factors - SRSF5, U2AF2, RBM17, and RBM5
- CLK inhibition promoted inclusion of an exon with an in-frame stop codon (a "poison exon," whose inclusion renders the transcript NMD sensitive) in the receptor tyrosine kinase FLT3
- CLK inhibition in AML cells led to downregulation of MCL-1, XIAP and FLT3 levels that confer resistance to BCL2 inhibitors
- Additional CLK inhibition targets are:
 - SMYD2 (a lysine methyltransferase recognized as a therapeutic target in AML)
 - DHODH (a metabolic enzyme and recent AML therapeutic target)
 - ATAD3A (a metabolic enzyme whose expression has been included in leukemia stem cell signatures)
 - 4. MYC target gene CDC16
 - RNA processing genes SRPK3, TRA2A, and DDX51

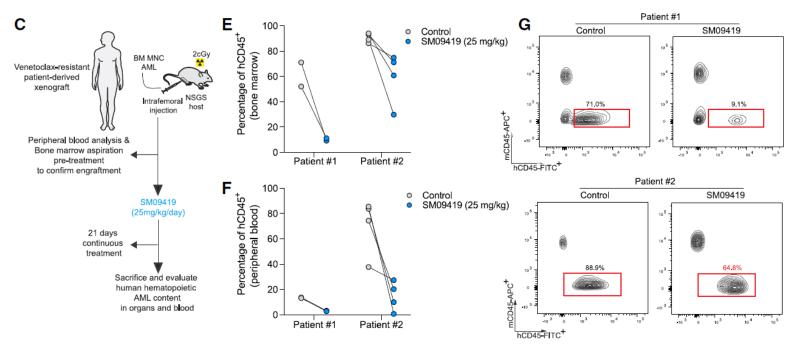


Modulation of RNA splicing enhances response to BCL2 inhibition in leukemia



Cancer Cell

CLK inhibition circumvents therapeutic resistance to venetoclax



Patient	Prior therapies	Reponse	Start of venetoclax	End of venetoclax	Sample date	Genetic alterations
1	VEN+Azacytidine	Primary refractory	01/18/2019	02/27/2019	06/05/2020	RUNX1, PTPN11, FLT3, SF3B1, STAG2
2	VEN+Low-dose Cytarabine	Primary refractory	01/03/2017	04/06/2017	04/10/2017	RUNX1, DNMT3A, IDH2



Summary



- Resistance to BCL2i is multifactorial, commonly associated with TP53 and "signaling" mutations and differentiation state:
 - Deficit of activator (BAX expression/mutations) or/and
 - Switch to other anti-apoptotics (MCL-1/BCL-xL)
- Inhibition of FLT3 signaling by FLT3 inhibitors or emavusertib reduces BCL-xL and MCL-1 expression via PI3K-AKT, RAS-MAPK and STAT5 pathways and sensitizes FLT3-mutated cells to venetoclax
- Inhibition of CLK (CDC like kinases) and DYRKs (dual-specificity tyrosine-regulated kinases) by emavusertib can overcome venetoclax resistance and shows strong potential to synergize with venetoclax
- Emavusertib in combination with azacitidine and venetoclax demonstrated significant antileukemic effects in all AML cell lines, including azacitidine- or venetoclax-resistant cell lines (Ugolkov EHA 2021)



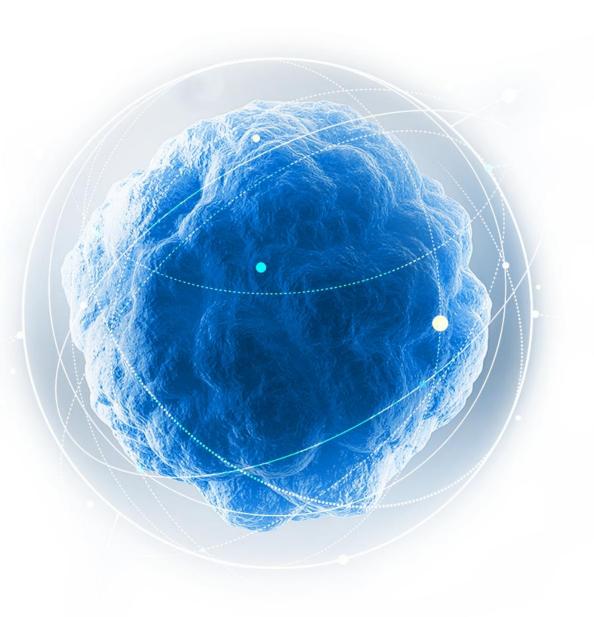


Incorporating IRAK inhibition in the treatment of myeloid disease

Guillermo Garcia-Manero MD

McCredie Professor Chief, Section of MDS Department of Leukemia MD Anderson Cancer Center Houston, TX





Agenda

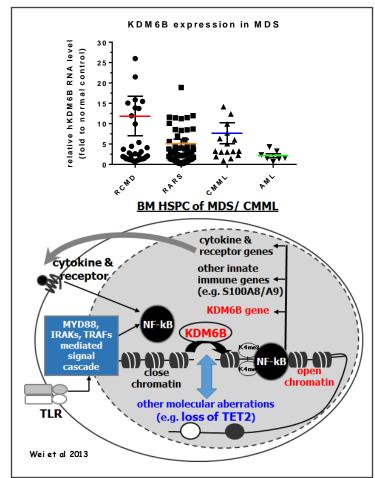


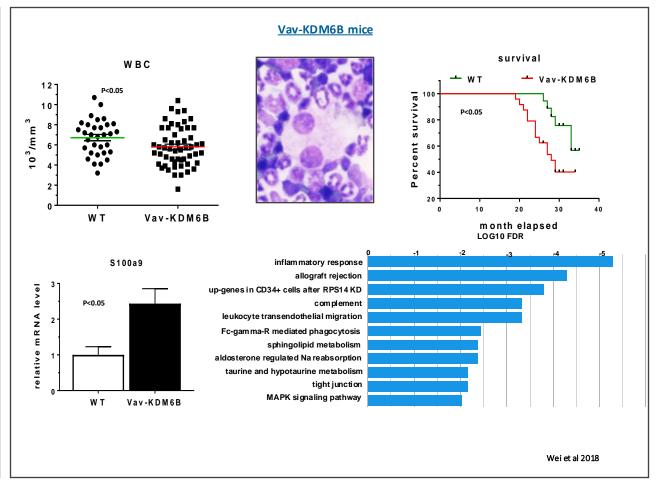
- Discuss basic concepts of MDS
- Review clinical activity of CA4849 in RR MDS and AML
- Discuss potential role of IRAK inhibition in MDS
- Discuss potential role of IRAK inhibition in AML



Innate Immune Signal Deregulation including Overexpression of KDM6B Drives Pathogenesis of MDS/ CMML





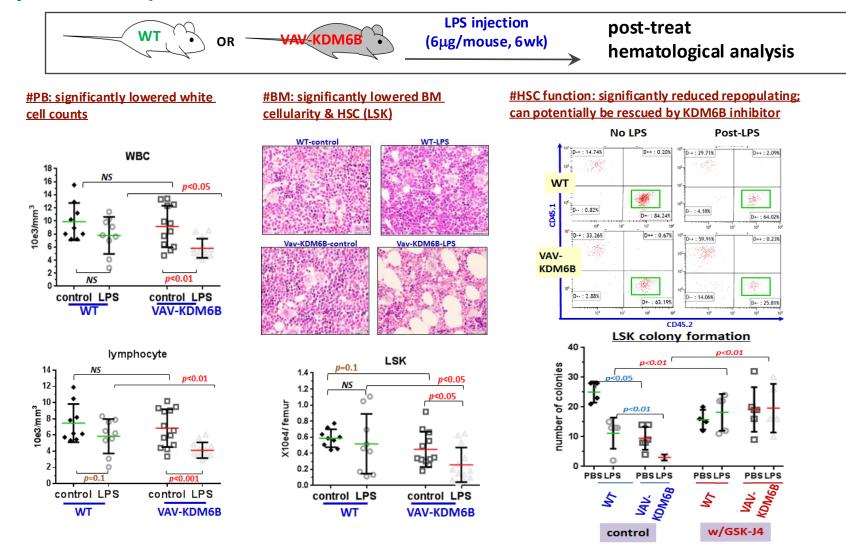




Vav-KDM6B(JMJD3)-Tg mice display MDS-like phenotype, particularly after exposure to chronic innate immune stimulation



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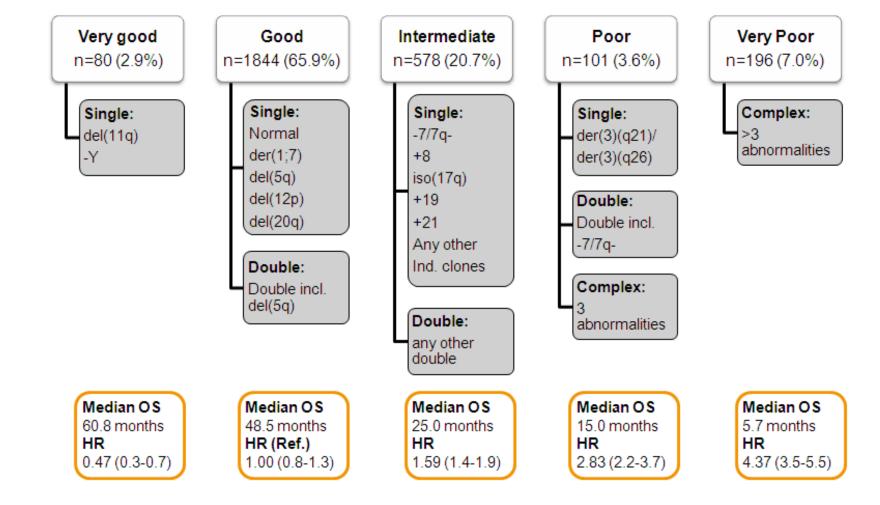


Wei et al Blood Adv 2019

Cytogenetic Scoring System in MDS



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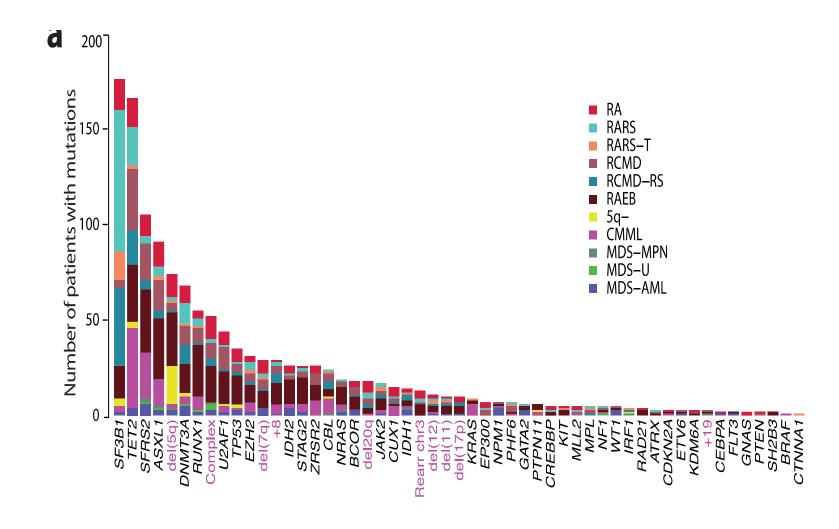




Schanz et al., JCO, 2011

Genomics of MDS



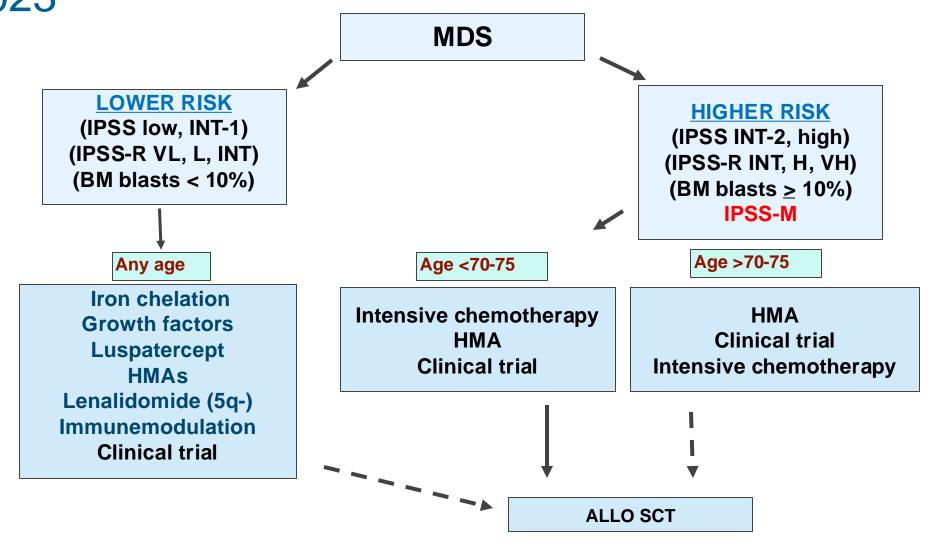




Proposed treatment algorithm for patients with MDS 2023

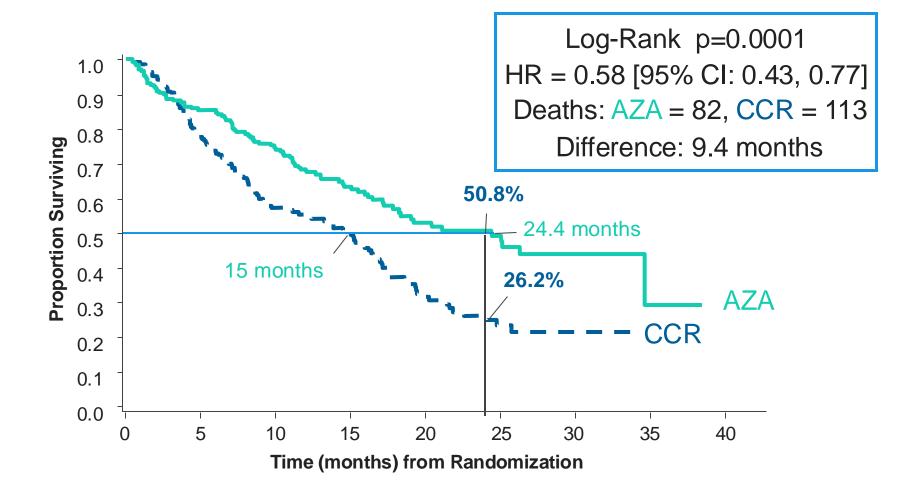


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Garcia-Manero et al AJH 2023





Doublets in Higher Risk MDS



- Panther (P-3001): azacytidine +/- pevonedistat
- Azacytidine + APR-246 for p53 mutated MDS
- HMA+ anti CD47
- HMA + venetoclax
- HMA + sabatolimab







A phase 1 study of azacitidine combined with venetoclax for myelodysplastic syndrome and chronic myelomonocytic leukemia

Alexandre Bazinet, MD, MSc, FRCPC Leukemia Fellow

Updated EHA 22 P757 Lancet Hematology 2022

Department of Leukemia
University of Texas MD Anderson Cancer Center



Phase I Azacitidine and Venetoclax for High-Risk MDS and CMML



Responses (N = 23 ITT analysis)

Response (Modified IWG)	All (n = 23) n (%) or median [range]	HMA-naïve (n = 17) n (%) or median [range]	HMA-failure (n = 6) n (%) or median [range]
ORR	20 (87)	14 (82)	6 (100)
CR	3 (13)	3 (18)	0 (0)
mCR	17 (74)	11 (65)	6 (100)
mCR + HI	5 (22)	5 (29)	0 (0)
mCR alone	12 (52)	6 (35)	6 (100)
Median DOR (months)		12.2	5.4
Median cycles given	3 [1–11]	3 [1–11]	5 [2–8]
Median cycles to response	1 [1–2]	1 [1–2]	1 [1–2]

Cytogenetic response rate in patients with baseline abnormality: 17% (2/12)

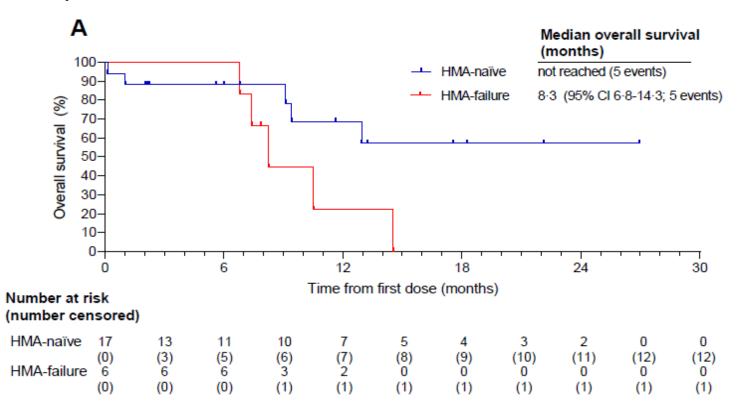


Phase I Azacitidine and Venetoclax for High-Risk MDS and CMML



Overall survival (N = 23)

Median follow-up: 13.2 months



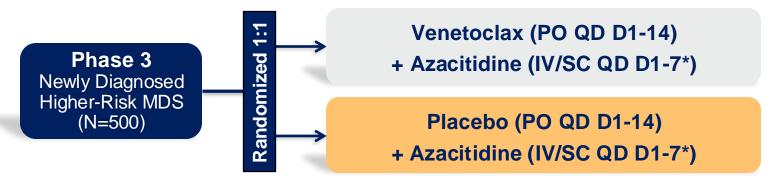


Phase 3 VERONA (NCT04401748)



Study Design and Endpoint

VERONA Study Design



*7 days within the first 9 calendar days/28 day cycle

Select Inclusion Criteria

- + ≥18 years old with newly diagnosed MDS according to 2016 WHO classification
- + <20% BM blasts
- + ECOG PS 0-2
- + IPSS-R score of >3 (Intermediate, High, Very High)
- No planned HSCT at the time of C1D1

Select Exclusion Criteria

- Prior therapy for MDS with HMA, chemotherapy, or allo-HSCT
- Prior diagnosis of therapy-related MDS, MDS evolved from MPN, MDS/MPN including CMML, aCML, JMML, and unclassifiable MDS/MPN

End Points

Primary: *CR, OS*

Secondary: *mOR, TI, ORR, fatigue score, physical functioning score, time to deterioration in physical functioning*

aCML=Atypical Chronic Myeloid Leukemia. allo-HSCT=Allogeneic Hematopoietic Stem Cell Transplant. AML=Acute Myeloid Leukemia. BM=Bone Marrow. C=Cycle. CMML=Chronic Myelomonocytic Leukemia. CR=Complete Remission. D=Day. ECOG PS=Eastern Cooperative Oncology Group Performance Status. HMA=Hypomethylating Agent. HSCT=Hematopoietic Stem Cell Transplantation. IPSS-R=Revised International Prognostic Scoring System. IV=Intravenous. JMML=Juvenile Myelomonocytic Leukemia. MDS=Myelodysplastic Syndrome. mOR=Modified Overall Response. MPN=Myeloproliferative Neoplasm. ORR=Overall Response Rate. OS=Overall Survival. PO=Oral. QD=Daily. SC=Subcutaneous. TI=Transfusion Independence. WHO=World Health Organization. 1. ClinicalTrials.gov. NCT04401748. https://clinicaltrials.gov/nCT04401748. Accessed July 2021



Targeted options in MDS



- IDH-2 (5-10%): enasidenib, venetoclax
- IDH-1 (5%): ivosidenib, venetoclax
- Flt-3 (15%): multiple agents
- TP53 (10%): anti-CD47?
- NPM1 (1%): ara-C based
- Splicing: IRAK4, H3BIO, Clk



Enasidenib in MDS: Response Rates

	Response	Arm A (Untreated)	Arm B (HMA- failure)
	Evaluable	AZA + ENA	ENA
	(N = 46)	(N = 25)	(N = 21)
ponse rate (ORR), n (%)	30 (68)	21 (84)	9 (43)
te remission (CR)	11 (24)	6 (24)	5 (24)
I remission (PR)	3 (7)	2 (8)	1 (5)
row CR (mCR)	12 (26)	11 (44)	1 (5)
ical improvement (HI) only	4 (9)	2 (8)	2 (10)
onse (NR), n (%)	16 (35)	4 (16)	12 (57)
e disease (SD)	14 (30)	4 (16)	10 (48)
sive disease (PD)	2 (4)	0 (0)	2 (10)

Response (Modified IWG)	All (n = 23) n (%) or median [range]	HMA-naïve (n = 17) n (%) or median [range]
ORR	20 (87)	14 (82)
CR	3 (13)	3 (18)
mCR	17 (74)	11 (65)
mCR + HI	5 (22)	5 (29)
mCR alone	12 (52)	6 (35)
Median DOR (months)		12.2
Median cycles given	3 [1–11]	3 [1–11]
Median cycles to response	1 [1–2]	1 [1–2]



DiNardo et al Blood Advances 2022



TAKEAIM LEUKEMIA- A PHASE 1/2A STUDY OF THE IRAK4 INHIBITOR EMAVUSERTIB (CA-4948) AS MONOTHERAPY OR IN COMBINATION WITH AZACITIDINE OR VENETOCLAX IN RELAPSED/REFRACTORY AML OR MDS

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Date: June 11, 2022

Session Title: Novel insights into AML treatment

Abstract: S129

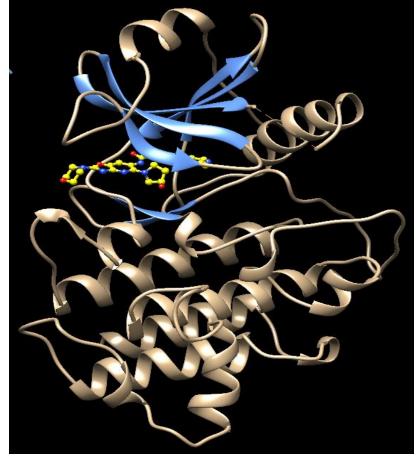


Emavusertib, An Oral IRAK4 Inhibitor

Emavusertib:

- Selective, small molecule inhibitor of IRAK4
- ATP-competitive, type 1 inhibitor, reversible
- Excellent drug-like properties:
 - Orally bioavailable (>100% dog/mouse)
 - Moderate plasma binding (77% human)
 - Stable in plasma, liver microsomes, hepatocytes
 - No inhibition of 7 major CYP450s
 - No significant metabolism in vitro
 - Humans: rapid absorption/clearance, T1/2 6 hr, no accumulation with QD dosing

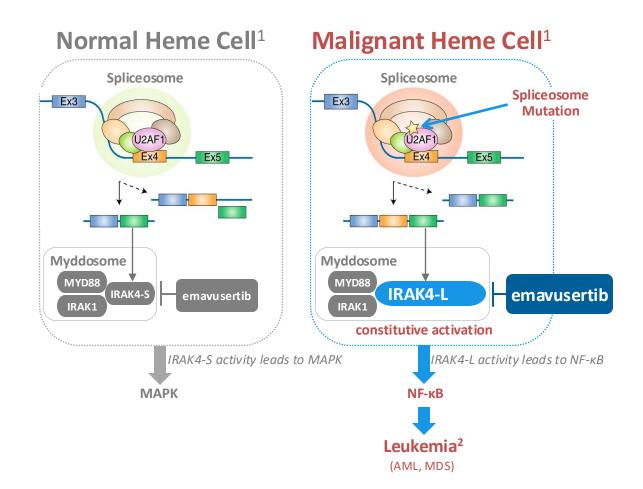
IRAK4/Emavusertib Co-crystal Structure



2.4Å resolution







- Emavusertib (CA-4948), a novel oral IRAK4 inhibitor has potential anti-leukemia activity
- Specific genetic mutations (SF3B1, U2AF1) in the spliceosome drive overexpression of IRAK4 long isoform (IRAK4-L)
- IRAK4-L then causes constitutive activation of the myddosome, leading to overactivity of NF-kB
- Therefore, this drug can target patients with splicing mutations
- Emavusertib also targets FLT3 and has shown potential synergetic activity with other drugs

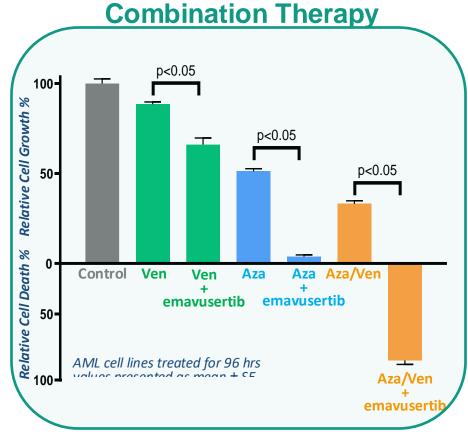




Emavusertib: Preclinical Activity in AML and MDS

Monotherapy Specimen_001-7-3068 14% Vehicle Émavusertib reduces *leukemic blasts* Specimen_001-1-3067 in monotherapy **Emavusertib**

Emavusertib demonstrates monotherapy activity in patient-derived xenografts¹

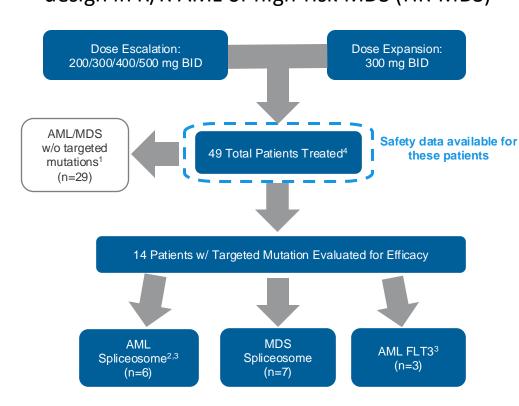


Emavusertib demonstrates synergy with both azacitidine and venetoclax in THP-1 model



Emavusertib: Study Design

TakeAim Leukemia (NCT #04278768): Open-label, single arm, Phase 1/2 dose escalation and expansion 3+3 study design in R/R AML or high-risk MDS (HR-MDS)



Study Objectives

- 1º: Determine maximum tolerated dose
 Determine recommended Phase 2 dose
- 2º: Pharmacokinetic (PK) profile Preliminary anti-cancer activity

Study Population

- Relapsed/Refractory AML or high-risk MDS
- ECOG performance Status of ≤ 2
- Age ≥ 18 years

All the data was extracted on Dec 16, 2021. Patients began enrollment into the combination therapy portion of the study in November 2021.

- 1. These are non-targeted patients, due to lack of spliceosome or *FLT3* mutation, this population will be addressed in the combination therapy study
- 2.One patient was not response evaluable because of discontinuation due to patient decision
- 3.Two AML patients have both a spliceosome and *FLT3* mutation and are included in both populations (there are 13 total evaluable patients with spliceosome or *FLT3* mutation)
- 4.Six patients did not start treatment by September 30th, 2021, which did not allow 2 on-study disease assessments



Dosing

- · Oral, BID Dosing
- 28-day cycles



Emavusertib: Baseline Characteristics

		Allmotionto		AML/MDS Subsets	
		All patients (n=49)	AML Spliceosome ¹ (n=6)	MDS Spliceosome (n=7)	AML FLT3 ¹ (n=3)
Female n (%): N	Male n (%)	16 (33) : 33 (67)	0 (0) : 6 (100)	5 (71) : 2 (29)	0 (0): 3 (100)
Age (yrs): media	n (range)	74 (32, 87)	76 (60, 84)	74 (61, 80)	80 (78, 87)
ECOG: n 0/1/2		11/30/8	0/4/2	2/5/0	0/1/2
Median platelets (10³/mm³) (range)		30 (4, 275)	28 (21, 80)	16 (7, 146)	21 (9, 23)
Median ANC (1	Median ANC (10 ³ /mm ³) (range)		0.23 (0, 3.3)	1.85 (0.15, 11.0)	0.05 (0, 0.11)
Median bone marrow blasts (%) (range)		-	33 (20, 95)	8 (3, 12)	60 (39, 95)
Median lines of prior therapy (range)		2 (1, 5)	2.5 (1, 4)	2 (1, 4)	2 (1, 4)
Prior thorapy	HMA ²	-	6 (100)	7 (100)	3 (100)
Prior therapy,	Chemotherapy ³	-	3 (50)	0 (0)	1 (33)
n (%)	Venetoclax	-	4 (67)	1 (14)	3 (100)

- Two AML patients have both a spliceosome and FLT3 mutation and are included in both populations (there are 13 total evaluable patients with spliceosome or FLT3 mutation)
- HMA includes azacitidine, decitabine, and guadecitabine
- Chemotherapy includes cytarabine



Emavusertib: Toxicities Profile

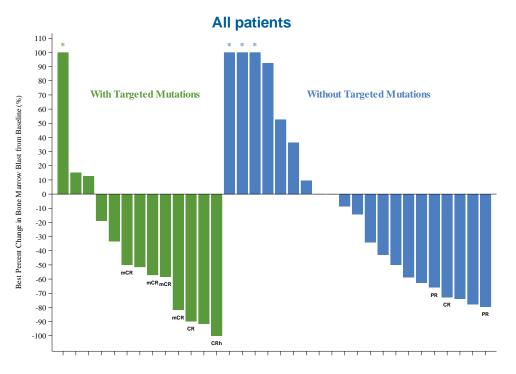
• During the initial dose escalation phase, no DLT was observed in 200-400 mg BID dose levels. Additional patients were enrolled at 300 mg and 400 mg BID to further explore the safety profile.

Grade 3+ Treatment-Related Adverse Event	200 mg BID (N = 3)	300 mg BID $(N = 26)^1$	400 mg BID (N = 17)	500 mg BID (N = 3)
	n (%)	n (%)	n (%)	n (%)
Number of patients having grade 3+ TRAEs	1 (33.3)	6 (23.1)	6 (35.3)	2 (66.7)
Alanine aminotransferase increased	1 (33.3)			
Blood creatine phosphokinase increased		1 (3.8)		
Dizziness	1 (33.3)			
Dyspnoea			1 (5.9)	
Enterobacter infection			1 (5.9)	
Fatigue			1 (5.9)	
Gastrointestinal haemorrhage		1 (3.8)		
Hypophosphataemia		1 (3.8)		
Hypotension		1 (3.8)		
Lipase increased		2 (7.7)		
Platelet count decreased		1 (3.8)		
Presyncope			1 (5.9)	
Rhabdomyolysis		1 (3.8)	2 (11.8)	1 (33.3)
Syncope				1 (33.3)

- 1. Data for the two patients that have escalated from 300 mg BID to 400 mg BID were included in the 400 mg BID dose group.
- 2. One death occurred after the data extraction date, currently under review.



Emavusertib: Single-agent Activity in AML and HR-MDS



Only evaluable patients with baseline and post-treatment bone marrow blast counts are included in the waterfall plot; among the patients w/o targeted mutations (SF3B1 / U2AF1 / FLT3 mutation), 1 reached CR and 2 PR

Subset of patients with targeted mutations (SF3B1 / U2AF1 / FLT3 mutation)

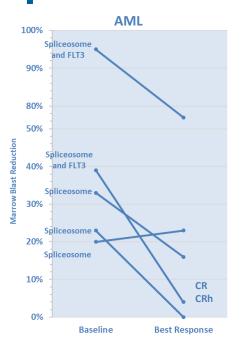
Best Response	Effica	ісу
Population #1: AML Spliceosome Patients ^{1, 2}		
CR/CRh Rate	2/5	(40%)
CR	1/5	(20%)
CRh	1/5	(20%)
Population #2: MDS Spliceosome Patients		
Objective Response Rate (ORR)	4/7	(57%)
CR	0/7	(0%)
mCR	4/7	(57%)
Population #3: AML FLT3 Patients ¹		
CR/CRh Rate	1/3	(33%)
CR	1/3	(33%)
CRh	0/3	(0%)

- 1. Two AML patients have both a spliceosome and *FLT3* mutation and are included in both populations (there are 13 total evaluable patients with spliceosome or *FLT3* mutation)
- 2. One patient was not response evaluable because of discontinuation due to patient decision



^{*} Indicates the best percentage change from baseline >100%

Emavusertib: Single-agent Activity in R/R AML with Spliceosome Mutation



Dose (BID)	Risk (ELN)	Baseline Molecular Mutations	# of Prior Therapies	Duration on emavusertib (mos)	Blasts Baseline	Blasts Best Response ¹	% Change
300 mg	Intermediate	SF3B1, RUNX1, WT1,	1	7	23	0	-100% (CRh)
300 mg	Intermediate	U2AF1, FLT3, BCOR, WT1	1	6+	39	4	-90% (CR)
300 mg	Intermediate	U2AF1, NRAS	4	2.5	33	16	-52%
300 mg	Adverse	FLT3, SF3B1, NRAS, PTPN11, RAD21, RUNX1, TET2, GATA, STAT3	4	2.6	95	77	-19%
400 mg	Adverse	SF3B1, DNMT3A, P53	1	2	20	23	15%

Data extraction date: Dec 16, 2021; "+" in Duration of Treatment indicates the patient remains on treatment as of the date of data extraction.

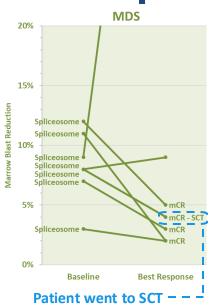
1. Two AML patients have both a spliceosome and FLT3 mutation and are included in both populations (there are 13 total evaluable patients with spliceosome or FLT3 mutation).

Emavusertib achieved 40% CR/CRh rate, despite transformed AML being historically highly resistant to treatment

AML Spliceosome Mutation







Dose (BID)	IPSS-R	Baseline Molecular Mutations	# of Prior Therapies	Duration on emavusertib (mos)	Blasts Baseline	Blasts Best Response	% Change
200 mg	Very High Risk	U2AF1 ,ASXL1, NF1, PHF6, GFI1, KDM6A, TET2	1	5.7	11	2	-82% (mCR)
300 mg	Very High Risk	U2AF1, DNMT3A, BCOR, STAG2, BCORL1, ETV6, SETBP1	1	3.3+	12	5	-58% (mCR)
400 mg	Very High Risk	SF3B1, RUNX1, NFE2	2	4.3	7	3	-57% (mCR)
300 mg	High Risk	SF3B1, DNMT3A, ASXL1, TET2, EZH2	2	0.9	8	4	-50% (mCR)
300 mg	High Risk	U2AF1, ASXL1	4	5.3+	3	2	-33%
300 mg	Very High Risk	SF3B1, ASXL1, NF1, SH2B3, RUNX1, PHF6, CBL, GFI1, EZH2	3	1.6	8	9	13%
400 mg	Very High Risk	U2AF1, ASXL1, BCOR, DNMTA, GATA2, SETBP1	1	1.2	9	62	>100%

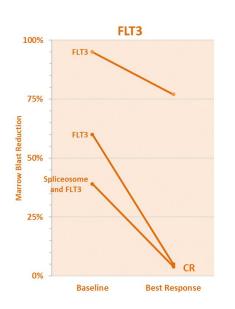
Data extraction date: Dec 16, 2021; "+" in Duration of Treatment indicates the patient remains on treatment as of the date of data extraction.

Emavusertib achieved 57% ORR, including one patient who was able to proceed to transplant

MDS Spliceosome Mutation



Emavusertib: Single-agent Activity in R/R AML with FLT3 Mutation



Dose (BID)	Risk (ELN)	Baseline Molecular Mutations	# of Prior Therapies	Duration on emavusertib (mos)	Blasts Baseline	Blasts Best Response ¹	% Change
400 mg	Adverse	FLT3 (eradicated at C3D1), ASXL1, BCOR, CEBPA (eradicated at C3D1), CSF3R, EZH2, NRAS, RUNX1 (X3), STAG2, TET2(X2,1) (eradicated at C3D1)	2	5.1	60	5	-92%
300 mg	Intermediate	FLT3 (eradicated at C4D1), BCOR (eradicated at C4D1), U2AF1 (decreased to 1.3 VAF at C4D1), WT1 (eradicated at C4D1)	1	6.2+	39	4	-90% (CR)
300 mg	Adverse	FLT3, SF3B1, NRAS, PTPN11, RAD21, RUNX1, TET2, GATA, STAT3	4	2.6	95	77	-19%

Data extraction date: Dec 16, 2021; "+" in Duration of Treatment indicates the patient remains on treatment as of the date of data extraction.

Emavusertib achieved 33% CR rate, and FLT3 mutation eradicated in 2 out of 3 patients

AML FLT3 Mutation



^{1.} Two AML patients have both a spliceosome and *FLT3* mutation and are included in both populations (there are 13 total evaluable patients with spliceosome or *FLT3* mutation).





Summary



- Emavusertib has a manageable safety profile
- Demonstrates oral, single-agent, anti-cancer activity in heavily pretreated AML and HR-MDS patients with targeted mutations (*U2AF1*, *SF3B1*, or *FLT3*)
- Potential candidate for use in combination therapy for all AML/HR-MDS patients, including patients without a targeted mutation

Next Steps:

- Correlative analysis ongoing
- Trials in lymphoma and solid tumors are being explored

We would like to thank the patients, their families and caregivers for their invaluable contribution and participation in this study.



Phase 2a Dose Expansion: Cohort 4



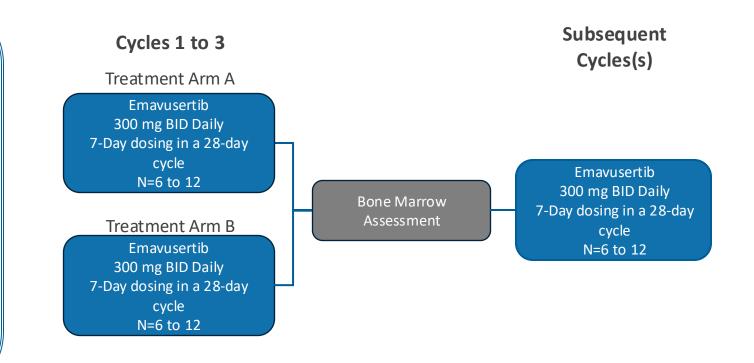
• Cohort 4 will include 2 treatment arms of orally administered CA-4948 as monotherapy to assess the safety, tolerability, and effect of 2 different dosing regimens of CA-4948 on the BM blast count and hematological parameters in patients with R/R hrMDS

Phase 2a Cohort 4: Adjusted Dosing Strategy for R/R hrMDS

Dose and Treatment Arms: Patients will receive 300 mg BID of CA-4948 with a shortened treatment duration of either 7 days (Arm A) or 14 days (Arm B) in a 28-day cycle, allowing for a drug holiday to promote normal hematological recovery.

Potential Benefits: This approach aims to maintain disease control while reducing transfusion needs and minimizing cytopenia-related AEs, potentially enhancing quality of life and prolonging survival.

Crossover Opportunity: After 3 cycles or 2 bone marrow assessments in Arm A, patients may switch to 14 days of treatment if early disease progression or lack of hematologic improvement is observed.





Phase 2a Dose Expansion: Cohort 4



Objectives

Primary

assess safety and tolerability of the 2 different dosing regimens of CA-4948 in patients with R/R hrMDS

Secondary

assess the anti-cancer activity of the 2 different dosing regimens of CA-4948 in patients with R/R hrMDS

Exploratory

To further assess the anti-cancer activity of the 2 different dosing regimens of CA-4948 in patients with R/R hrMDS

Endpoints

Primary

Safety measured by AEs, ECGs, chemistry and hematology laboratory values, vital signs, and physical examination

Secondary

Proportion of patients that achieve mCR or CR
• Reduction in frequency of transfusion versus baseline, including - Proportion of patients that achieve transfusion independence

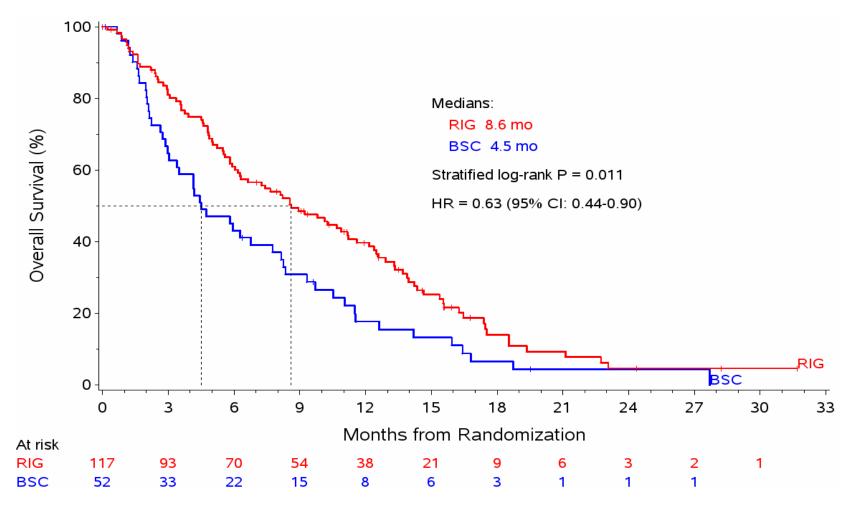
Exploratory

- •Proportion of patients that achieve HI: HI-E or HI-P or HI-N
- Proportion of patients who achieve CR, or PR, or mCR with HI
 - DOR
 - Time to responses
 - OS



ONTIME Trial: Median Overall Survival for Pts with Primary HMA Failure - Blinded, Centralized Assessment







Garcia-Manero. Lancet Oncology 2016

Potential role of IRAK inhibition in MDS



- Examples HR MDS front line
 - Doublet HMA+ CA4948
 - Triplet HMA+venetoclax+ CA4948
- Monotherapy in HMA failure HR MDS
- Consider potential role in lower risk MDS



Potential role of IRAK inhibition in AML



- Role as a Flt-3 inhibitor
- Similar path as in AML?

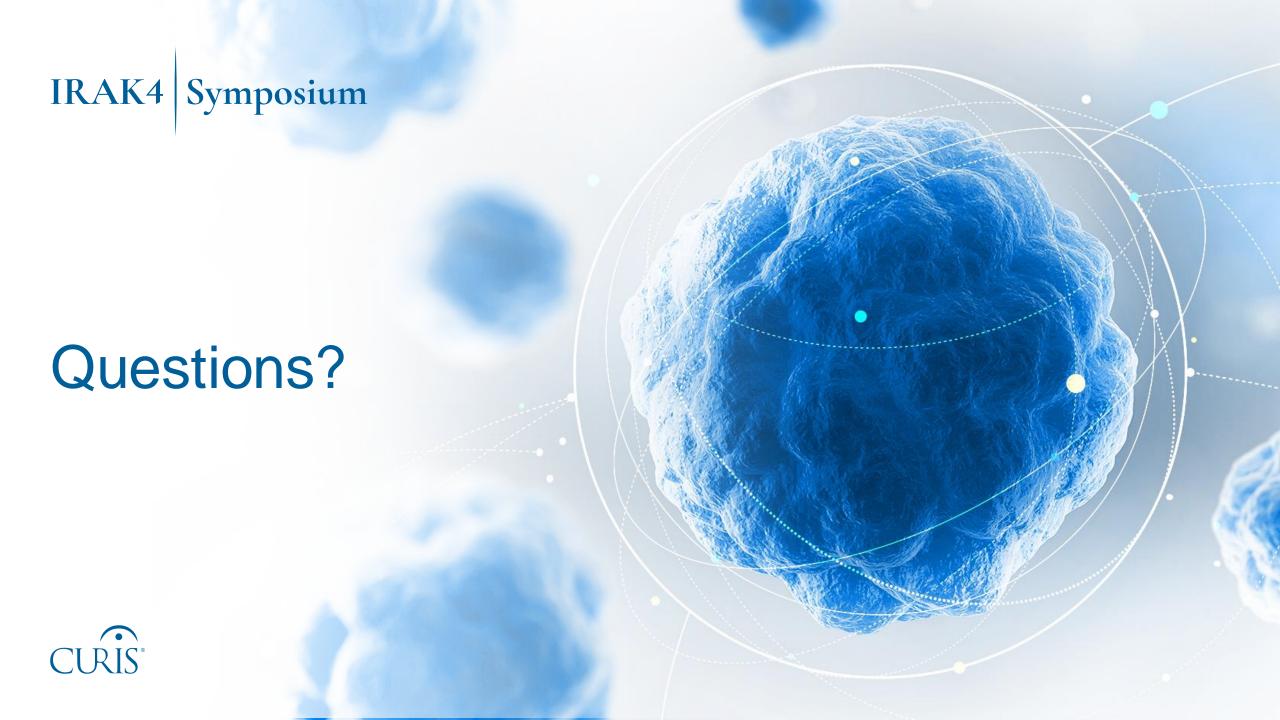


Thank you



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Department of Leukemia
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Emavusertib for Acute Myeloid Leukemia

Eric S. Winer, MDSymposium Co-Chair

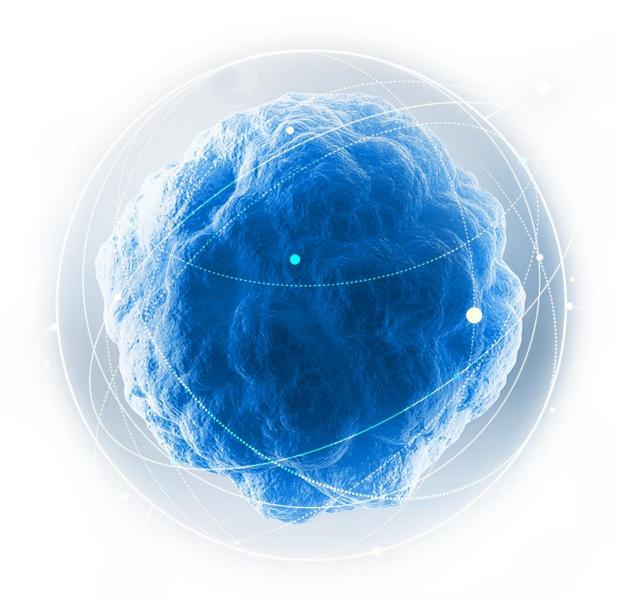
Clinical Director, Adult Leukemia

Dana Farber Cancer Institute

Assistant Professor of Medicine

Harvard Medical School

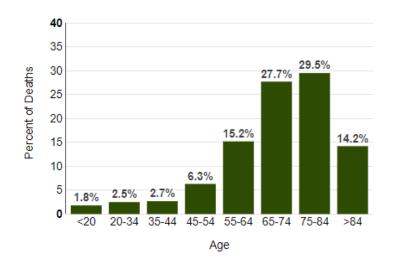


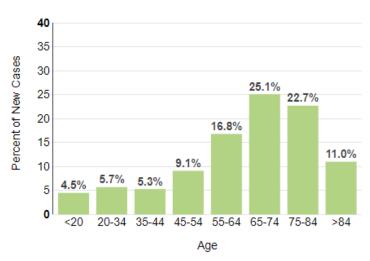


The State of Acute Myeloid Leukemia

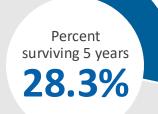
IRAK4 Symposium

- Accounts for only 1.2% of all cancer deaths
 - Estimated new cases 2022: 20,050
 - Estimated Deaths 2022: 11,540
 - 5-year relative survival (2018): 28.3%
- Median age is 68 years of age
- Risk Factors:
 - Age
 - Prior Chemotherapy
 - Ionizing Radiation
 - Chemicals/Solvents













https://seer.cancer.gov/statfacts/html/amyl.html

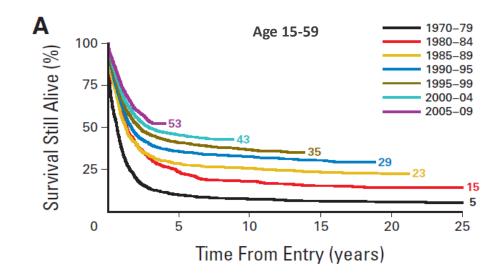
The State of AML – novel approved therapies

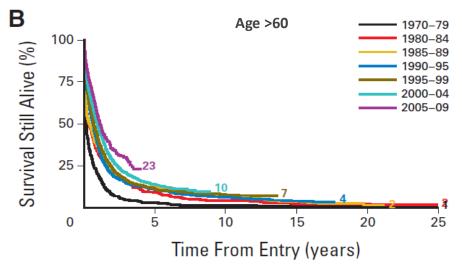


64

Incremental improvements

- Midostaurin (induction)
 - Midostaurin 74.7 months, Placebo 25.6 months
- Vyxeos (CPX-351) (induction)
 - Vyxeos 9.6 months, 7+3 5.95 months
- Venetoclax (with Aza)
 - Aza/Ven 14.7 months, Aza alone 9.6 months
- Glasdegib (with Low dose cytarabine)
 - Glasdegib/LoDAC 8.8 months, LODAC 4.9 months
- Gilteritinib
 - Gilteritinib 9.3 months, Salvage 5.6 months
- Ivosidenib (with aza)
 - 24 months (Aza/Ivo) v 7.9 months Aza
- Quizartinib (induction)
 - Quizartinib 31.9 months v placebo 15.1 months
- Aza/Ven/Gilt (non-randomized)
 - 18 mo RFS and OS 71 and 72%





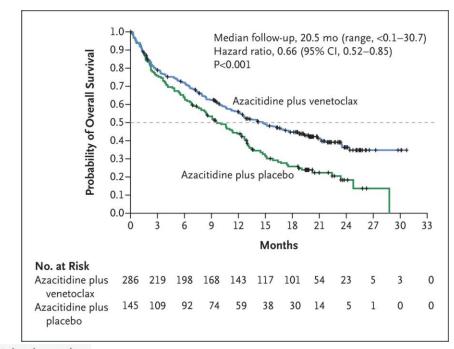


Burnett A et al. *J Clin Oncol 29:487-494*

Venetoclax + Azacitidine AML - VIALE-A

IRAK4 Symposium

- Randomized trial in patients ineligible for induction
- Median age 76 years old
- Endpoint of Overall Survival
- OS:
 - 14.7 mo Aza/Ven v. 9.6 mo Aza
- CR:
 - 36.7% Aza/Ven v. 17.9% Aza
- CRc:
 - 66.4% Aza/Ven v. 28.3% Aza
- Primary and adaptive resistance seen in activating signaling pathways (i.e. *FLT3*, *RAS*, *TP53*)



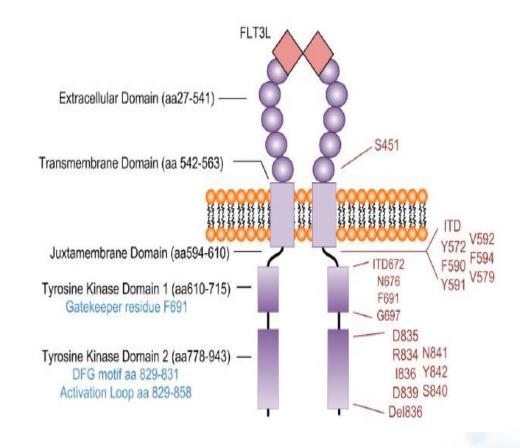
Molecular marker			
FLT3	19/29 (65.5)	19/22 (86.4)	⊢ ■
IDH1	15/23 (65.2)	11/11 (100.0)	⊢
IDH2	15/40 (37.5)	14/18 (77.8)	⊢
IDH1 or IDH2	29/61 (47.5)	24/28 (85.7)	⊢ ■
TP53	34/38 (89.5)	13/14 (92.9)	- ■
NPM1	16/27 (59.3)	14/17 (82.4)	⊢



FLT and AML

IRAK4 Symposium

- FLT3 most common mutated gene in AML (30-35%)
- Often associated with normal karyotype
- Different mutations
 - Internal Tandem Duplicates
 - Tyrosine Kinase Domain
- Proliferative phenotype
- Allelic ratio important for pathogenicity/prognosis

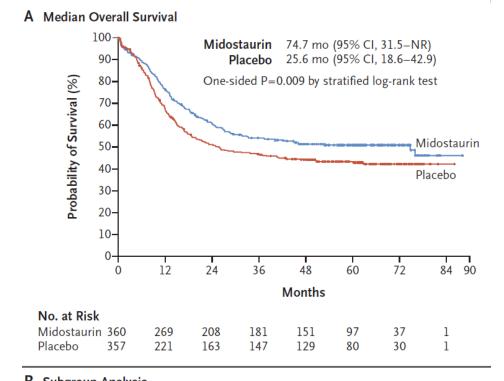


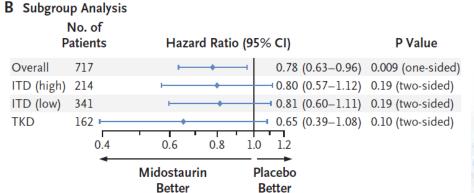


7+3+ Midostaurin for FLT3 positive AML

IRAK4 Symposium

- Patients screened for FLT3 Mutation (ITD or TKD)
 - 3277 screened for 717 participants
- Randomized to either 7+3 or 7+3 + midostaurin (days 8-21)
- Day 21 marrow for aplasia
- Consolidation with HiDAC +/- midostaurin
- Maintenance phase



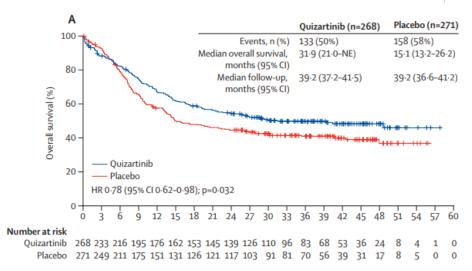


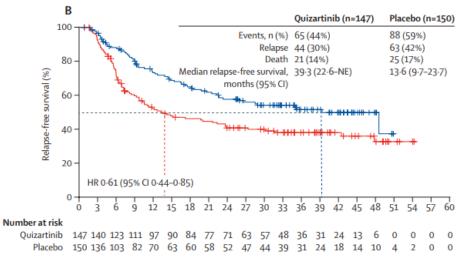


7+3+Quizartinib for AML (QuANTUM-First)



- Enrolled 539 patients with FLT3-ITD positive AML randomized to 7+3+Quizartinib v. 7+3+placebo
- Median age 56 years
- Median overall survival:
 - Quizartinib: 31.9 months (72% CCR)
 - Placebo: 15.1 months (65% CCR)
- MRD (post hoc) <10⁻⁴
 - Quizartinib: 42%
 - Placebo: 38%





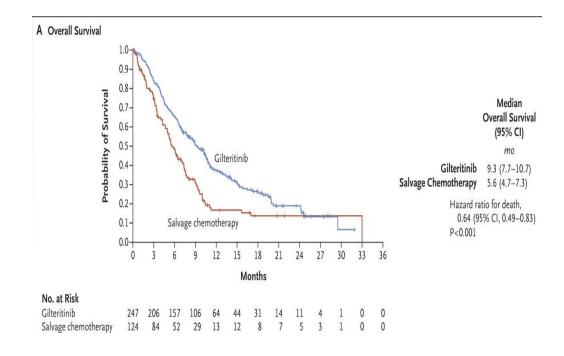


68

Gilteritinib



- Phase 3 study randomizing 2:1 Gilteritinib
 v. salvage chemotherapy (MEC, FLAG-IDA,
 Low dose cytarabine, azacitidine)
- Gilteritinib improved median overall survival of 9.3 months v. 5.6 months
- EFS 2.8 months v. 0.7 months
- CR+CRi 34% v. 15.3%
- Hazard ratio for death 0.64 (p<0.001)



Previous therapy for AML — no. (%)

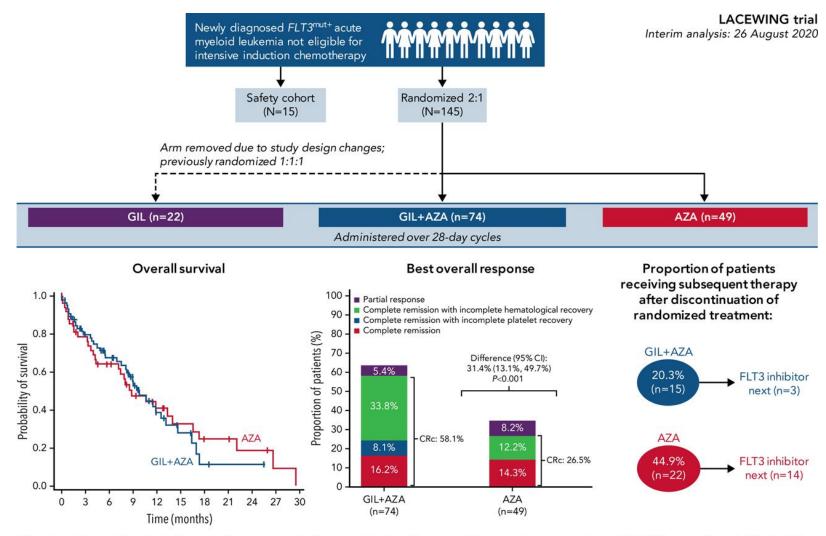
Anthracycline	311 (83.8)	205 (83.0)	106 (85.5)
FLT3 inhibitor	49 (13.2)	34 (13.8)	15 (12.1)
HSCT	74 (19.9)	48 (19.4)	26 (21.0)



69

Azacitidine + Gilteritinib v. Azacitidine





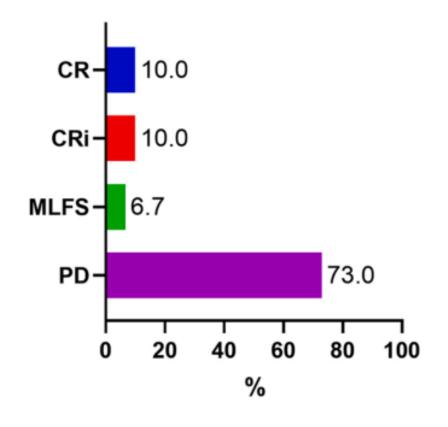
Abbreviations: AZA, azacitidine 75 mg/m² intravenously or subcutaneously daily on days 1–7; CI, confidence interval; CRc, composite complete remission; FLT3, FMS-like tyrosine kinase 3; GIL, gilteritinib 120 mg orally daily on days 1–28; HR, hazard ratio.



Landscape of response in relapsed FLT-3 AML



- Retrospective study evaluating response to FLT-3 patients previously treated with venetoclax
- ORR to FLT3 inhibitors was 26.7%
- Median overall survival was 6.7 months among all patients

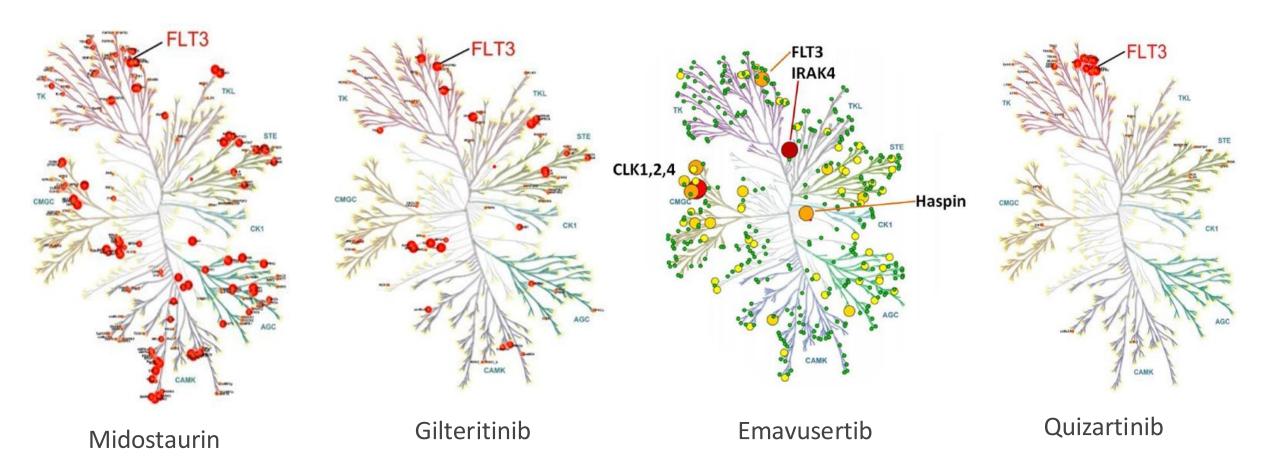




Bewerdorf, et al. Leuk Res. 2022.

FLT3 Kinome comparisons







CA-4948: Emavusertib – an IRAK4/FLT3 kinase inhibitor



Emavusertib with interaction points on IRAK4

Potent inhibition of IRAK4 with significant inhibition of pIRAK1 in PK-PD studies

Gate-keeper Met265 Hinge Tvr264

Demonstrates activity in other kinases such as CLK1, CLK2, CLK4, FLT3, DYRK1A, DYRK1B, TrkA, TrkB, Haspin, and NEK11

Emavusertib Binding Affinity

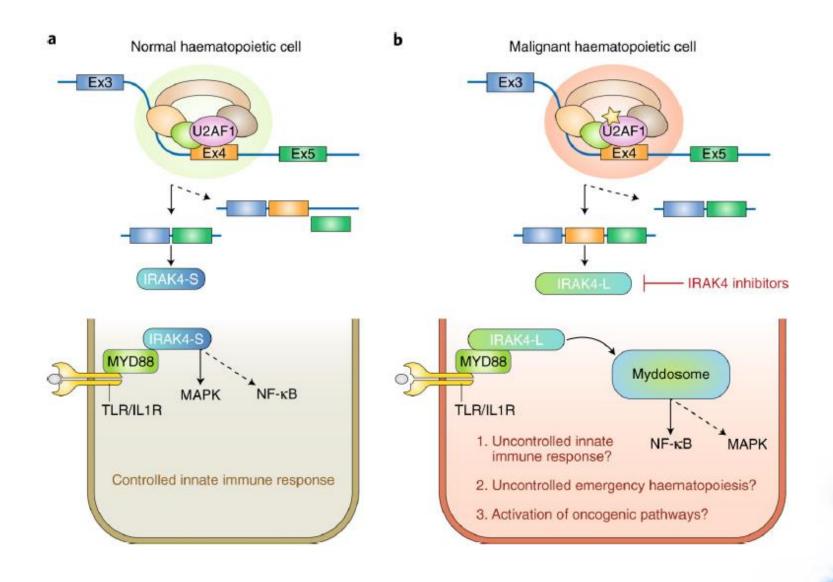
	Target	K_d nM	
	IRAK1	12,000	
	IRAK2	>20,000	
	IRAK3	8,500	
	IRAK4	23	
Ī	DYRK1A	25	
	FLT3 wt	31	
	FLT3 (D835H)	5	
	FLT3 (D835V)	44	
	FLT3 (D835Y)	3	
	FLT3 (ITD)	8	
	FLT3 (K663Q)	47	
	FLT3 (N841I)	16	
Ī	Haspin (GSG2)	32	
	CLK1	10	
	CLK2	20	
	CLK3	>20,000	
	CLK4	14	
	TrkA	130	

DiscoverX Kinase Panel (378 kinases screened)



IRAK4: The long and the short



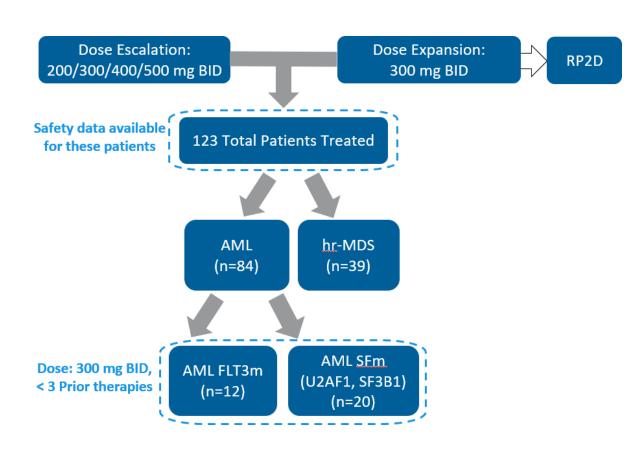


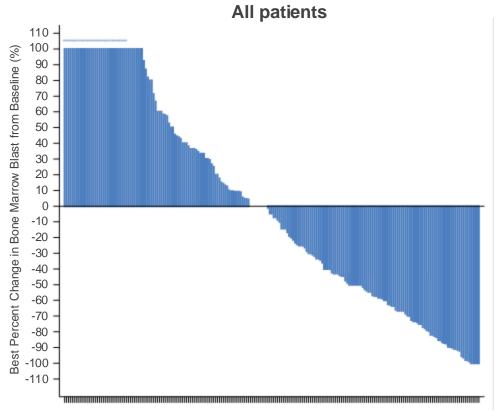


TakeAim Leukemia study:

IRAK4 Symposium

Schema





Includes all patients that had baseline and post-treatment bone marrow blast assessments. *Indicates best percentage change from baseline >100%.



TakeAim Leukemia: Adverse Events



Treatment-related adverse events (TRAEs) of Grade ≥3 in all TakeAim Leukemia trial patients

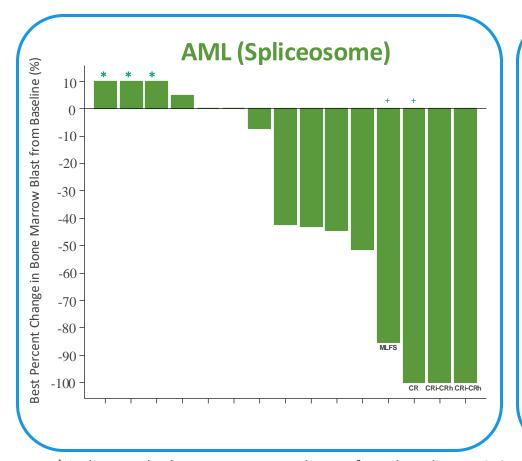
Grade 3+ Treatment-Related Adverse Event reported in >1 patients, n (%)	200 mg BID (N = 27)	300 mg BID (N = 78)	400 mg BID (N = 15)	500 mg BID (N = 3)	Total (N=123)
# of patients having grade 3+ TRAEs	4 (14.8)	21 (26.9)	7 (46.7)	2 (66.7)	34 (27.6)
# of patients having non-hematological grade 3+ TRAEs	3 (11.1)	17 (21.8)	6 (40)	2 (66.7)	28 (22.8)
Blood creatine phosphokinase increased	0	6 (7.7)	0	0	6 (4.9)
Platelet count decreased	1 (3.7)	3 (3.8)	2 (13.3)	0	6 (4.9)
Rhabdomyolysis ^{a,b}	0	2 (2.6)	1 (6.7)	1 (33.3)	4 (3.3)
Anemia	0	3 (3.8)	0	0	3 (2.4)
Aspartate aminotransferase increased	1 (3.7)	2 (2.6)	0	0	3 (2.4)
Alanine aminotransferase increased	2 (7.4)	0	0	0	2 (1.6)
Dizziness	1 (3.7)	1 (1.3)	0	0	2 (1.6)
Febrile neutropenia	0	2 (2.6)	0	0	2 (1.6)
Lipase increased	0	2 (2.6)	0	0	2 (1.6)
Neutropenia	0	1 (1.3)	1 (6.7)	0	2 (1.6)
Neutrophil count decreased	0	1 (1.3)	1 (6.7)	0	2 (1.6)
Syncope	0	1 (1.3)	0	1 (33.3)	2 (1.6)

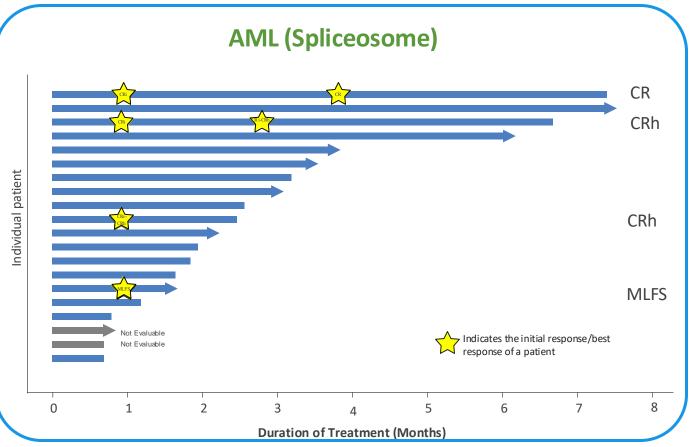
^aAfter discussion with regulatory authorities of investigator-reported AEs, objective laboratory criteria for the determination of rhabdomyolysis were adopted from existing approved drug labels (creatine phosphokinase >10 x upper limit of normal and SCr ≥ 1.5 x upper limit of normal). Previously, reported events of rhabdomyolysis were determined by subjective criteria. Using the objective criteria, rhabdomyolysis was reported in 1/123 patients. ^bOne patient receiving 300 mg BID emavusertib died with an investigator-reported cause of multi-organ failure and rhabdomyolysis, with the latter assessed by the investigator as likely related to study drug. The Leukemia and Lymphoma Society's independent safety board adjudicated that the fatal outcome in this patient was unrelated to treatment but instead due to multi-organ failure from disease progression.



TakeAim Leukemia: Spliceosome mutation (SF3B1 and U2AF1 mutations)





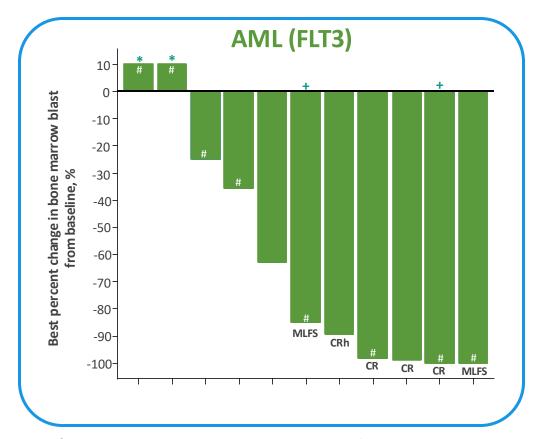


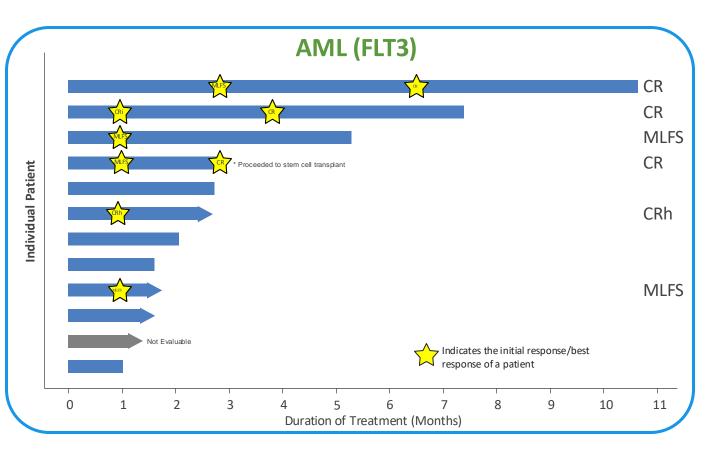
- * indicates the best percentage change from baseline >10%
- + indicates concomitant FLT3 mutation



TakeAim Leukemia: FLT3m Cohort







- * indicates the best percentage change from baseline >10%
- # demonstrates previous venetoclax exposure
- + indicates concomitant spliceosome mutation

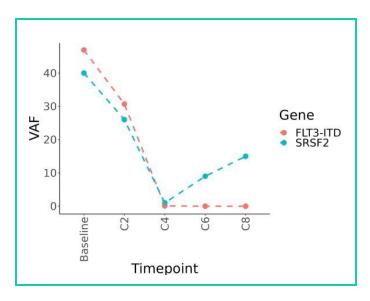


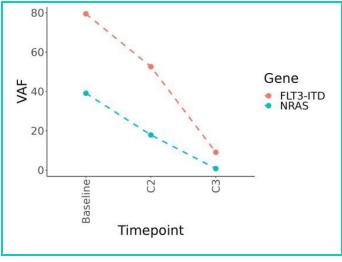
TakeAim: FLT3m responders



Clinical activity in responders with R/R AML - FLT3m

#	Age	Sex	ELN risk per 2017	FLT3 mutation	# prior therapy	Prior BCL2i	Prior HMA	Prior FLT3i	Best response	Co-mutations at baseline
1	80	M	Intermediate	ITD	1	Υ	Υ	N	CR	U2AF1, BCOR, WT1
2	44	M	Adverse	ITD	2	Υ	Ν	Υ	CR	NRAS, WT1
3	74	M	Adverse	Not available	2	Υ	Υ	N	MLFS	SF3B1, GATA2, PHF6, RUNX1, CBLC
4	78	F	Adverse	ITD	2	Υ	Υ	Υ	MLFS	Not available
5	79	F	Intermediate	ITD	2	N	Υ	N	CR	DMNT3A, SRSF2
6	74	M	Intermediate	ITD	1	N	Υ	Υ	CRh	Not available



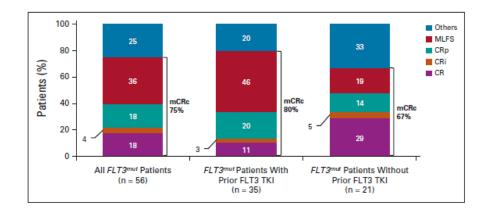


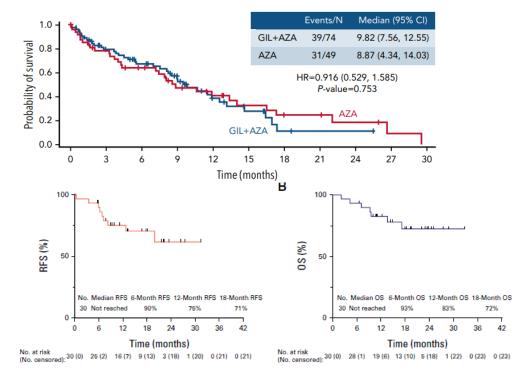


Doublets and Triplets – More is better?



- Venetoclax + Gilteritinib
 - CR 18%, CRi 4%, CRp 18%, MLFS 36%
 - mDOR 4.9 months
- Azacitidine + Gilteritinib v. Azacitidine
 - No difference in OS (9.82 v 8.87 mo)
- Azacitidine + Venetoclax + Gilteritinib
 - Frontline CR 90%, CRi 6%
 - 97% of patients >60
 - Median number of cycles: 3
 - 43% patients went to alloBMT







Emavusertib with other agents



- Cell line data showed increased cell death in:
 - 2 of 4 cell lines with Venetoclax
 - 3 of 4 cell lines with Azacitidine
 - 1 of 4 cell lines with decitabine
 - 4 of 4 cell lines with Azacitidine and Venetoclax
- Did not seem to have significant activity with daunorubicin and/or cytarabine.

AML cell line	Decitabine + CA-4948	Azacitidine + CA-4948	Venetoclax + CA-4948	Venetoclax + Decitabine + CA-4948	Venetoclax + Azacitidine + CA-4948
THP-1	NS	++	+	NS	+++
F-36P	NS	++	NS	NS	++
OCI-AML2	++	+	++	NS	+++
GDM-1	NS	++	NS	NS	++

AML cell line	Ara-C + CA-4948	Venetoclax + Ara-C + CA-4948	Daunorubicin + CA-4948	Daunorubicin + Ara-C + CA-4948
THP-1	+	++	NS	++
F-36P	++	NS	NS	NS
OCI-AML2	+	NS	NS	NS
GDM-1	NS	NS	NS	NS

- + <50% growth inhibition, P<0.05
- ++ 50 <100% growth inhibition p<0.05
- +++ 100% growth inhibition and induction of cell death. P<0.05

NS: Not significant



IRAK4 Symposium

Eligibility: Untreated AML

Adequate organ function

AND

Age ≥60

OR

sAML

OR

Co-morbidities that preclude intensive chemotherapy.

Level -1: Azacitidine 75mg/m2 IV or SC days 1-7 Venetoclax 400mg PO daily days 1-28* Emavusertib 50 mg PO daily days 1-28*

Level 1: Azacitidine 75mg/m2 IV or SC days 1-7
Venetoclax 400mg PO daily days 1-28*
Emavusertib 100mg PO daily days 1-28*

Level 2: Azacitidine 75mg/m2 IV or SC days 1-7
Venetoclax 400mg PO daily days 1-28*
Emavusertib 200mg PO daily days 1-28*

Level 3: Azacitidine 75mg/m2 IV or SC days 1-7 Venetoclax 400mg PO daily days 1-28* Emavusertib 300mg PO daily days 1-28*



^{*}Bone marrow biopsy to be evaluated day 14-18.

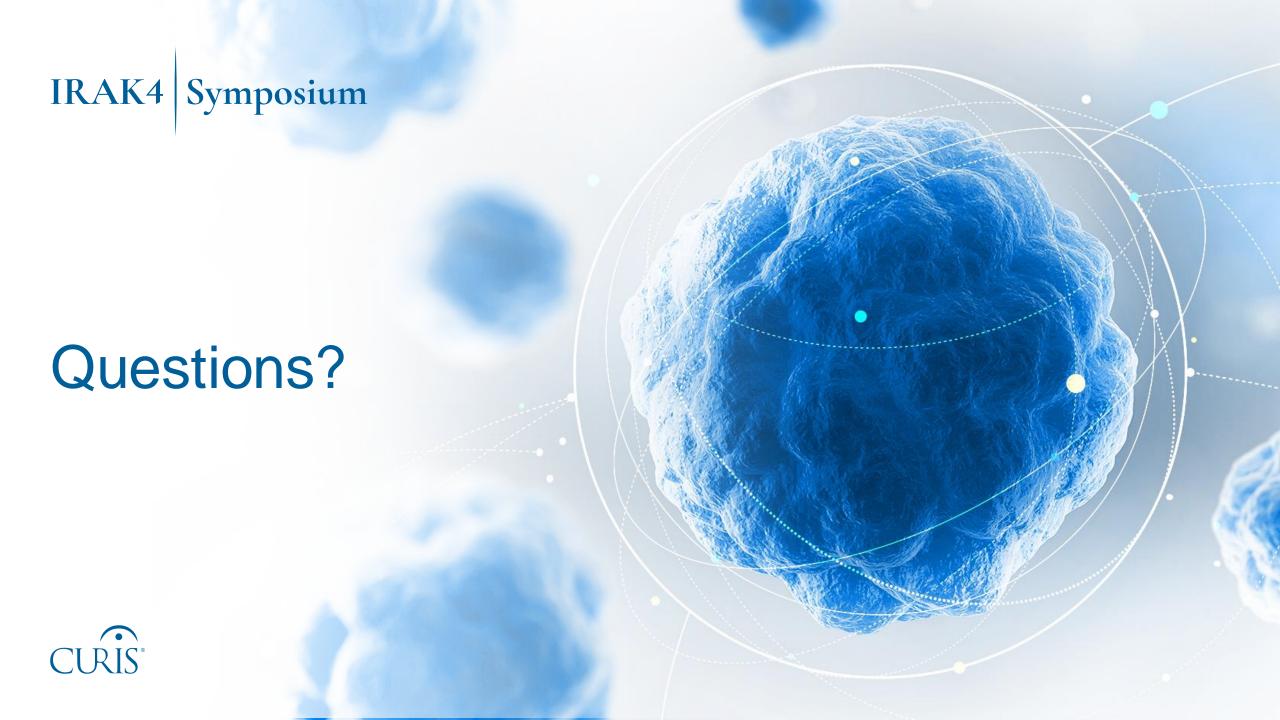
If marrow is ablated, then Venetoclax and Emavusertib held until count recovery.

Conclusions: Emavusertib in AML



- Emavusertib demonstrates an acceptable and manageable safety profile in patients with AML
- The mutational response of emavusertib in AML patients as a single agent demonstrate the on-target efficacy, particularly in FLT3 mutated and spliceosome mutated AML
- Emavusertib monotherapy has shown responses in FLT3 patients who have been refractory to venetoclax and hypomethylating agents and prior FLT3 inhibitors
- A trial will open shortly using emavusertib in combination with azacitidine and venetoclax in untreated AML in patients either >60 years of age, secondary AML, or unfit for intensive induction chemotherapy





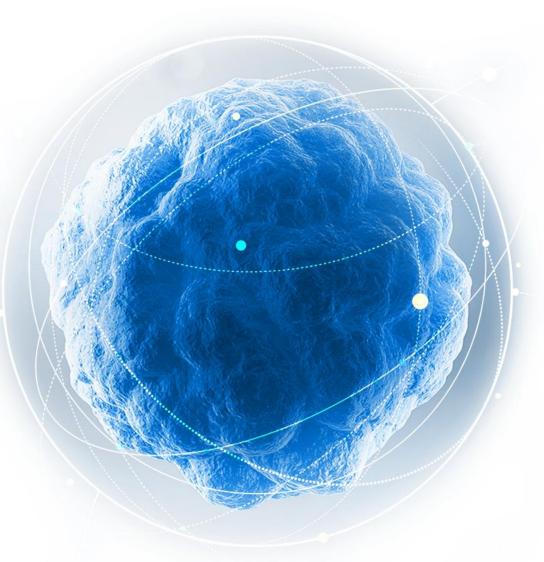


Emavusertib in combination with azacitidine and venetoclax in patients with AML with MRD+ CR

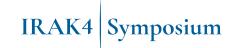
Klaus H Metzeler, MD

Professor of Translational Hematology, University of Leipzig





Background: AML with MRD+



- AML is a heterogenous disease that exhibits a dynamic mutational landscape as the disease progresses 1,2
- Achieving MRD– is an important treatment goal in AML³
 - In the VIALE-A study, patients with AML with CRc and MRD of <10⁻³ or MRD- had longer DOR, event-free survival, and OS than patients who achieved CRc but with MRD+³
- Patients with AML who are older or not suitable for intensive chemotherapy have limited treatment options and resistance to venetoclax-based combinations is common⁴⁻⁶
 - In the 1L setting, the current standard of care is azacitidine + venetoclax^{5,6}
 - Primary and adaptive resistance to venetoclax-based combinations has been associated with acquisition or expansion of clones with *FLT3-ITD* mutations⁴
 - MCL-1 has also been shown to drive resistance to venetoclax⁷
 - While FLT3-driven relapses are common,⁴ there is no established role for a FLT3 inhibitor

DiNardo CD, et al. N Engl J Med. 2020;383:617-629. 6. Short NJ, et al. J Clin Oncol; 2024; 42:1499-1508. 7. Thijssen R, et al. Haematologica. 2015;100(8):e302-6.

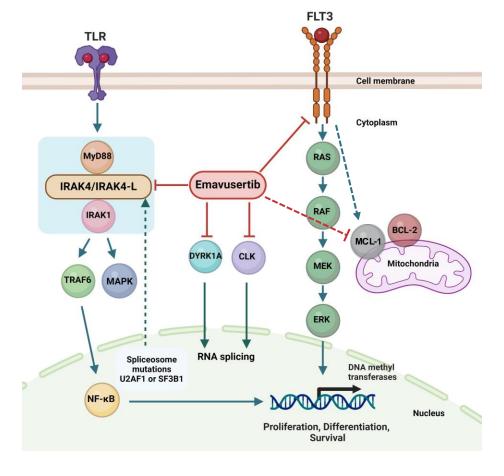


Emavusertib is a novel inhibitor of IRAK4, FLT3, and CLKs



- IRAK4 is a serine/threonine kinase that mediates signaling downstream of TLRs, resulting in NF-κB and MAPK activation, and is associated with cell survival, proinflammatory cytokine production, and activation of the innate immune system¹⁻⁴
- Emavusertib is a novel potent oral inhibitor of IRAK4 with additional inhibitory activity against FLT3 and CLK1/2/4⁵
- IRAK1/4 inhibition has been shown to result in reduced MCL-1 stability and abundance⁶

Emavusertib mechanism of action^{5–7}



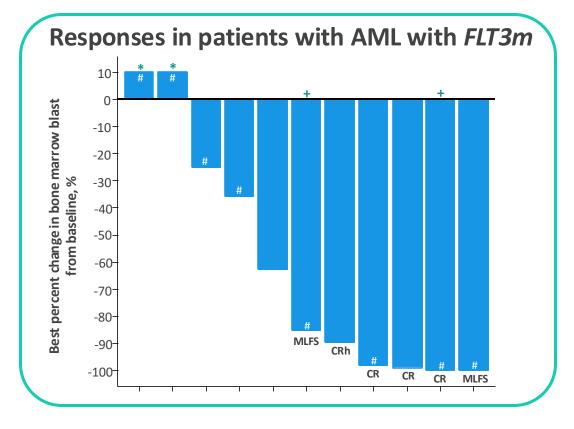


The efficacy of emavusertib monotherapy has been demonstrated in patients with AML



In the Phase 1 TakeAim Leukemia trial, emavusertib monotherapy has demonstrated both CRs and a significant reduction in blasts in patients with R/R AML with *FLT3*m who received prior therapy with an HMA and/or a FLT3 inhibitor

- Patients with AML with FLT3m received 300 mg BID emavusertib; 6 of 11 patients achieved a response of CR, CRh, or MLFS
- Of 6 responders:
 - 4 had prior BCL2 inhibitor therapy
 - 5 had prior HMA therapy
 - 3 had prior FLT3 inhibitor therapy
 - 2/6 had both prior HMA and FLT3 inhibitor therapy





Emavusertib monotherapy had an acceptable and tolerable safety profile



In the Phase 1 TakeAim Leukemia trial, emavusertib monotherapy had an acceptable and tolerable safety profile across the full patient population

Treatment-related adverse events (TRAEs) of Grade ≥3 in all TakeAim Leukemia trial patients

Grade 3+ Treatment-Related Adverse Event reported in > 1 patients, n (%)	200 mg BID (N = 27)	300 mg BID (N = 78)	400 mg BID (N = 15)	500 mg BID (N = 3)	Total (N=123)
# of patients having grade 3+ TRAEs	4 (14.8)	21 (26.9)	7 (46.7)	2 (66.7)	34 (27.6)
# of patients having non-hematological grade 3+ TRAEs	3 (11.1)	17 (21.8)	6 (40)	2 (66.7)	28 (22.8)
Blood creatine phosphokinase increased	0	6 (7.7)	0	0	6 (4.9)
Platelet count decreased	1 (3.7)	3 (3.8)	2 (13.3)	0	6 (4.9)
Rhabdomyolysis ^{a,b}	0	2 (2.6)	1 (6.7)	1 (33.3)	4 (3.3)
Anemia	0	3 (3.8)	0	0	3 (2.4)
Aspartate aminotransferase increased	1 (3.7)	2 (2.6)	0	0	3 (2.4)
Alanine aminotransferase increased	2 (7.4)	0	0	0	2 (1.6)
Dizziness	1 (3.7)	1 (1.3)	0	0	2 (1.6)
Febrile neutropenia	0	2 (2.6)	0	0	2 (1.6)
Lipase increased	0	2 (2.6)	0	0	2 (1.6)
Neutropenia	0	1 (1.3)	1 (6.7)	0	2 (1.6)
Neutrophil count decreased	0	1 (1.3)	1 (6.7)	0	2 (1.6)
Syncope	0	1 (1.3)	0	1 (33.3)	2 (1.6)

^aAfter discussion with regulatory authorities of investigator-reported AEs, objective laboratory criteria for the determination of rhabdomyolysis were adopted from existing approved drug labels (creatine phosphokinase >10 x upper limit of normal and SCr ≥ 1.5 x upper limit of normal). Previously, reported events of rhabdomyolysis were determined by subjective criteria. **Using the objective criteria**, **rhabdomyolysis was reported in 1/123 patients.** Done patient receiving 300 mg BID emavusertib died with an investigator-reported cause of multi-organ failure and rhabdomyolysis, with the latter assessed by the investigator as likely related to study drug. The Leukemia and Lymphoma Society's independent safety board adjudicated that the fatal outcome in this patient was unrelated to treatment but instead due to multi-organ failure from disease progression.

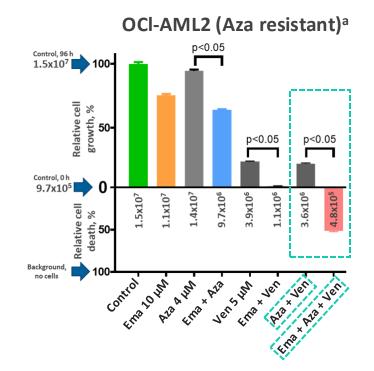


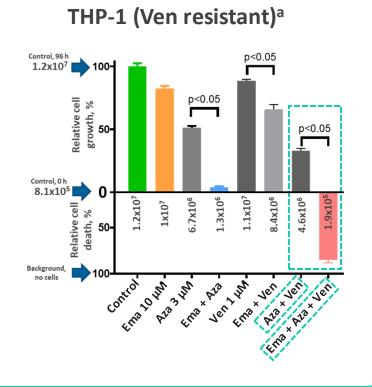
Addition of emavusertib can restore anti-leukemic activity in resistant AML cell lines



Emavusertib in combination with azacitidine and venetoclax has demonstrated significant anti-leukemic effects in AML cell lines, including azacitidine- or venetoclax-resistant cell lines

- AML cell lines were treated for 96 hours
- Relative cell viability was measured at 0 and 96 hours





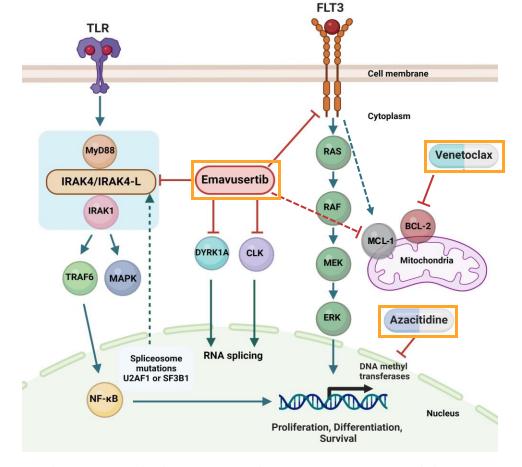


Preclinical and clinical data of emavusertib support further investigation in patients with AML



- FLT3 mutations and MCL-1 have significant roles in conferring resistance to azacitidine + venetoclax^{1,2}
- The activity of emavusertib against FLT3m, preclinical evidence of synergy with azacitidine and venetoclax, and evidence of IRAK4 inhibition resulting in reduced MCL-1 support clinical evaluation of a triplet regimen³⁻⁶
- Adding emavusertib to the azacitidine/venetoclax doublet in patients in MRD+ CR may enable patients to achieve MRD negativity without significant added toxicity
- The phase 1b CA-4948-104 study was designed to assess the efficacy of emavusertib in combination with venetoclax and azacitidine in AML patients in MRD+ CR (EUCTR#2023-505828-58)⁷

Emavusertib + venetoclax + azacitidine mechanism of action^{6–8}





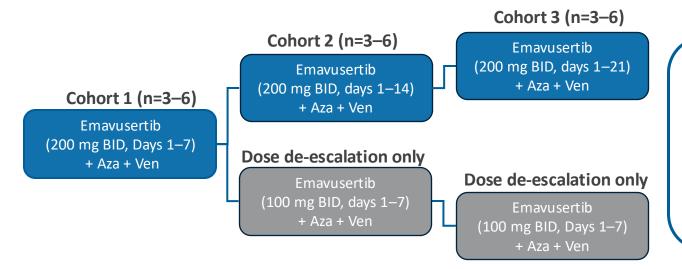
CA-4948-104: Study design



• CA-4948-104 is a Phase 1b, single-arm, open label study assessing emavusertib + venetoclax + azacitidine in patients aged ≥60 years with AML who achieved CR or CRh with MRD+ on venetoclax + azacitidine as 1L therapy (EUCTR#2023-505828-58)^{1,2}

Key Eligibility Criteria

- Aged ≥60 years
- Confirmed diagnosis of AML
- Ongoing 1L ven + aza, ≤6 cycles
- Documented CR or CRh within
 28 days prior to Cycle 1, Day 1
- Documented bone marrow MRD positivity



Treatment until

- Withdrawal
- Disease progression
- Intolerable toxicity
- Not achieving MRD negative status within 6 cycles

Emavusertib is administered orally 200 mg BID on days 1–7, 1–14, or 1–21 of a 28-day cycle Aza and ven are administered at the same dose schedule as when patient achieved CR or CRh



CA-4948-104: Objectives and endpoints



Objectives¹

Endpoints^{1,2}

Primary

• Evaluate safety and tolerability of different dose schedules of emavusertib as an add-on agent to the combo of aza + ven (triplet regimen)

Safety, including AEs

Secondary

- Evaluate conversion of MRD+ to MRD- status with triplet regimen
- Characterize PK profiles of emavusertib, azacitidine, and venetoclax
- Assess effects of triplet regimen on dynamics of MRD status and relationship to outcomes
- Evaluate continuous anti-cancer activity of the triplet regimen

- Rate of MRD conversion by flow cytometry
- PK parameters
- Time to MRD conversion from first dose of triplet regimen
- Duration of MRD negativity
- OS

Exploratory

- Evaluate the molecular profile of peripheral blood at baseline and following treatment with the triplet regimen
- Changes in expression profiles
- Identification of biomarkers



CA-4948-104: Patient eligibility



01 Identifying Eligible Patients

- Newly diagnosed patient with AML who is ≥60 years old
- Planned first-line aza + ven or currently receiving aza + ven with ≤6 cycles
- Follow response assessments through cycle 6 for 1st CR/CRh with MRD+

02 Patient tracking

HAS first response of CR/CRh with MRD+ via bone marrow

03 Toxicity

NOT experiencing ≥G2 aza/ven toxicity

04 Stem cell transplant

NOT an immediate candidate for allogeneic stem cell transplant

05 ECOG PS, organ function & CPK

WITH ECOG PS ≤2 | WITH acceptable organ function and CPK Level (≤2.5 x ULN)

06 Exclusion Criteria

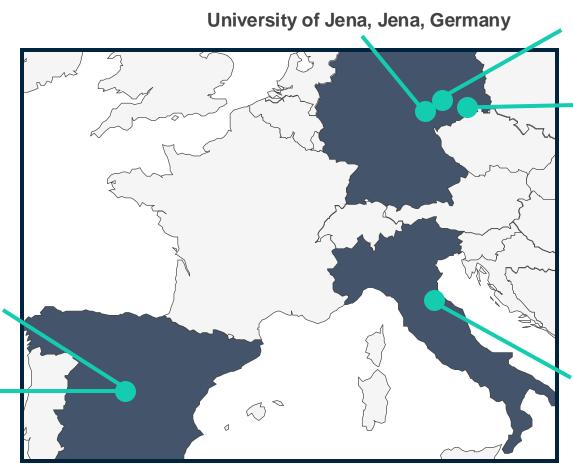
DOES NOT meet exclusion criteria





Study sites currently enrolling





University Hospital Leipzig, Leipzig, Germany

University of Dresden, Dresden, Germany

Istituto Romagnolo per lo Studio dei Tumori, Meldola, Italy



Hospital Gregorio Marañon,

MD Anderson, Madrid, Spain

Madrid, Spain

Summary



- Emavusertib monotherapy demonstrated promising efficacy and has been well tolerated in patients with R/R AML
- Emavusertib in combination with azacitidine and venetoclax demonstrated synergistic anti-leukemic effects in AML cell lines
- Adding emavusertib to the azacitidine/venetoclax doublet in MRD+ CR may enable patients to achieve MRD negativity without significant toxicity
- This triplet combination has a potential to become a new option in 1L therapy for older/unfit patients with AML, regardless of mutation status
- Exploratory biomarkers will be analyzed to determine predictive biomarkers of response
- The study is currently enrolling patients





Current and Future Opportunities For IRAK4 Inhibition in Leukemia and MDS Roundtable

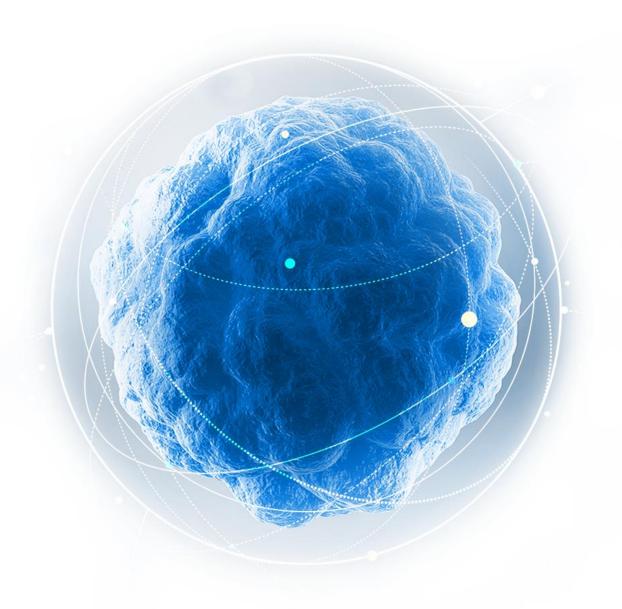
Eric S. Winer (moderator)

Guillermo Garcia-Manero

Klaus Metzeler

Marina Konopleva





Discussion Questions



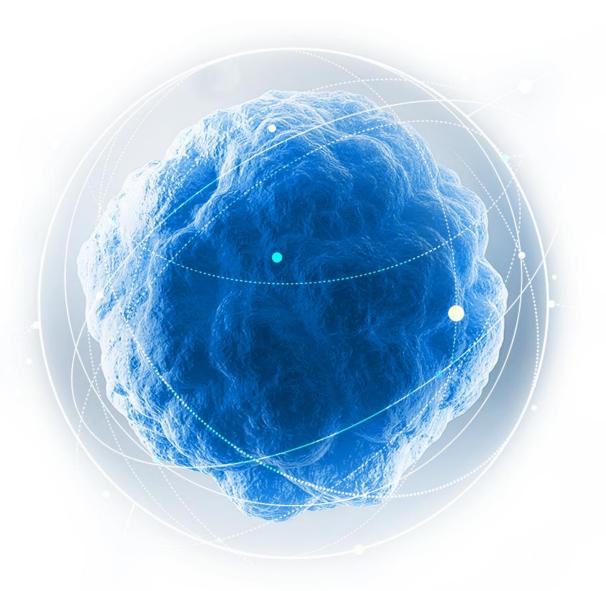
- What is your general approach to the older patient with AML/MDS? Do you envision some of the combinations that we are discussing supplanting the "standard" treatment of 7+3 intensive chemotherapy.
- What are key unmet needs in myeloid malignancies, besides TP53 mutated MDS/AML?
- What is the optimal combination strategy for IRAK4 inhibition in myeloid malignancies? Do you see a benefit in combining emavusertib with other FLT3 inhibitors? With other agents such as HMA, venetoclax or both?
 - Given current treatment approaches, in which patients and contexts might addition of an IRAK4 inhibitor be an option?
- Which genetic subtypes of myeloid malignancies are likely to be most sensitive to IRAK4 inhibition? Are there biomarkers for response other than mutations?
- What are the expected benefits for reduced dose schedules of emavusertib, such as the 7-day and 14-day dose schedules being tested in MDS? In which patient populations would assessing a modified dose be suitable? Do you see a role for this in AML?
- Is there potential for emavusertib to be a component of treatment either broadly or in specific populations as 1L therapy for MDS or AML? What would need to be demonstrated to support this? What about as monotherapy in 2L, such as following HMA failure?





IRAK4 Targeting in PCNSL

Han W. Tun, MD
Professor of Medicine
Department of Hematology/Oncology
Department of Cancer Biology
Mayo Clinic Florida





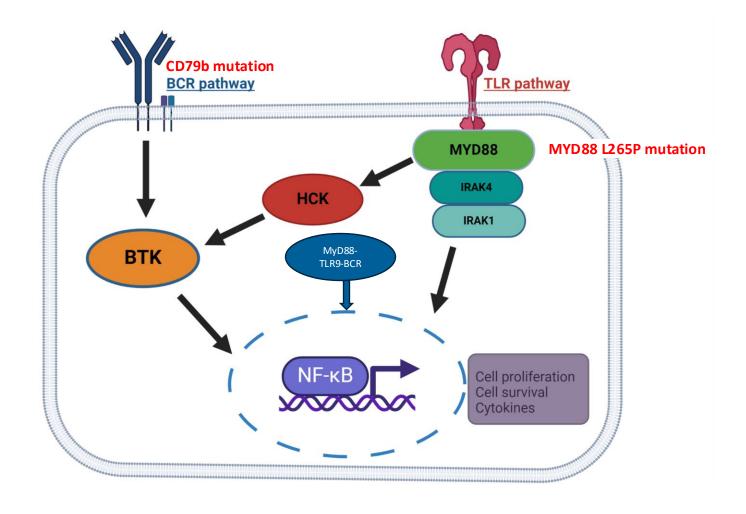
Interleukin-1 Receptor-associated Kinase 4 (IRAK4)

A serine/threonine kinase in the Toll-like receptor and interleukin-1 receptor pathways (Innate immune response)

Recruited to MYD88 upon activation with downstream activation of NFKB and MAPK



Constitutive activation of BCR and TLR pathways in MYD88 mutant B cell lymphomas





MYD88 L265P mutant B-cell non-Hodgkin lymphomas

Non-germinal center diffuse large B cell lymphoma (33%)

Primary cutaneous DLBCL, leg type (70%)

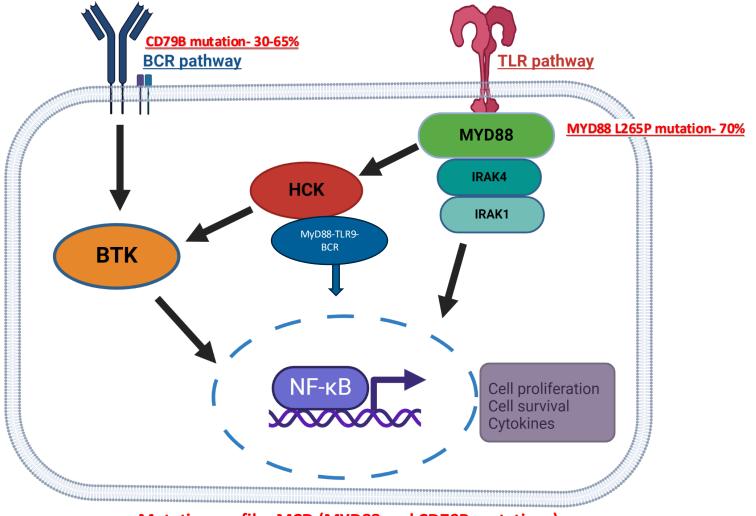
Primary CNS lymphoma (70%)

Primary testicular lymphoma (70-80%)

Waldenstrom macroglobulinemia/lymphoplasmacytic lymphoma (95-97%)



PCNSL



Mutation profile- MCD (MYD88 and CD79B mutations)
Pathway signature- Oncogenic constitutive activation of BCR and TLR pathways
Activated NFKB signaling



Genomic Landscape of PCNSL



- Genomic instability (CDKN2A deletion)
- Oncogenic Toll-like receptor signaling (MYD88 L265P mutation)
- Oncogenic B cell receptor signaling (CD79 mutation)
- Dysregulation of BCL6 (Mutations and translocation involving BCL6)
- Immune evasion (PD-L1/PD-L2 gains and translocation)



Primary CNS lymphoma

Diffuse large B cell lymphoma confined to the central nervous system (brain, leptomeningeal, eyes, spinal cord, cranial nerves)

Incidence ~ 1800 new cases per year in US

Potentially curable with chemoimmunotherapy followed by autologous stem cell transplant



25792 Patients with Non-HIV PCNSL (SEER + CBTRUS- 1973-2013)



Age	Incidence
>50	79%
>60	62%
>70	26%

Age	Median OS
<50 (21%)	83 M
50-69 (53%)	25 M
>70 (26%)	6 M

Median OS for the whole group doubled from 12.5 M in 1970s to 26 M in 2010s

Median OS for age >70 has not changed last 40 years – 6 months



Ibrutinib monotherapy in R/R PCNSL/SCNSL

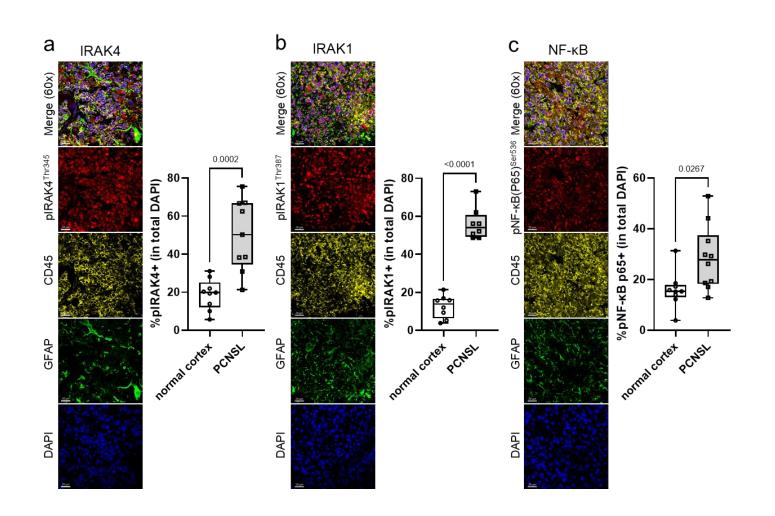
Clinical Trial	Phase	Treatment	Median F/U	Total accrual	ORR/CR+CRu	mPFS	mOS
Soussain et al.	II	Ibrutinib 560 mg qd	25.7 M	52 with R/R PCNSL or PVRL	52%/19%	3.3 M	14.4 M
Grommes et al.	al. I Ibrutinib 560- 840 mg qd	Ibrutinib 560-	15.7 M	13 with R/R PCNSL	77%/38%	4.6 M	15 M
Grommes et al.		T2./ IAI	7 with R/R PCNSL	71%/57%	7.4 M	NR	

Pre-clinical evaluation of Emavusertib (IRAK-4 inhibitor)



Myddosome expression in human PCNSL



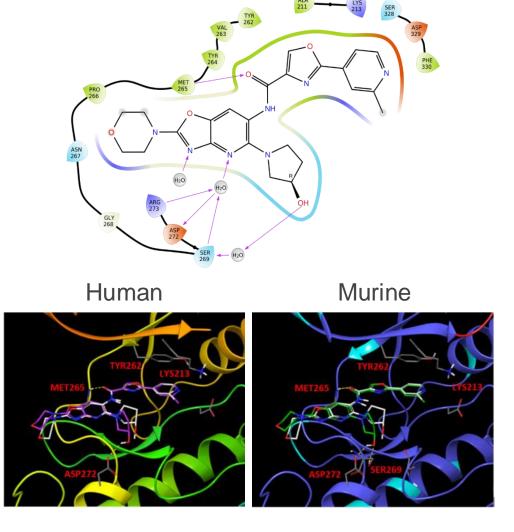


- High levels of p-IRAK4,p-IRAK1, and p-NF-кВ
- High CD45 infiltration



Emavusertib (CA-4948)

- First-in-class inhibitor
- High binding affinity to human IRAK4 (23 nM), high predicted binding affinity to murine IRAK4
- Well tolerated; safety profile allows long-term treatment and combination with other therapies

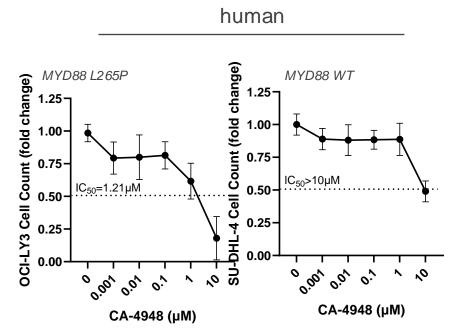


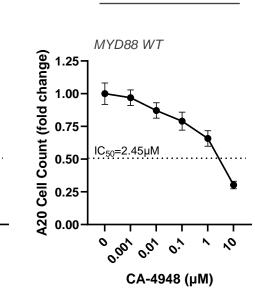


Emavusertib anti-lymphoma activity



- Dose-dependent decrease in lymphoma proliferation
- MYD88 L265P sensitivity
- Anti-tumor activity in immune-competent MYD88 WT lymphoma





murine

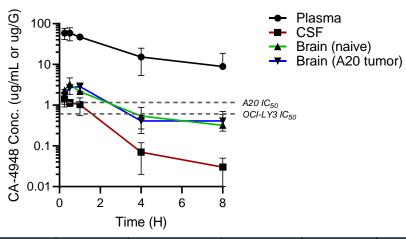


CNS penetration (preclinical)



- Emavusertib can cross the BBB
- Relevant therapeutic dose levels detected in naïve parenchyma and CSF
- No notable changes in permeability in tumor-bearing mice

LC-MS/MS detection of CA-4948 in murine CNS

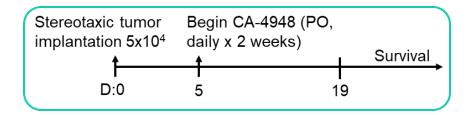


Parameter	Units	Plasma	CSF (Naïve)	Brain (Naïve)	Brain (Tumor)	
C _{max}	μg/mL or μg/g	60.3 ± 19.26	1.42±0.52	3.25±1.41	3.22±0.18	
T _{max}	h	0.38 ± 0.14	0.25	0.5	0.83±0.29	
T _{1/2}	h	2.73	1.33	1.39	1.19	
AUC _{0-8 h}	h*µg/mL or h*µg/g	189.51	2.91	8.09	8.68	
AUC _{0-∞}	h*µg/mL or h*µg/g	224.46	2.96	8.72	9.39	
Brain to plasma ratio	%		1.53	4.26	4.95	

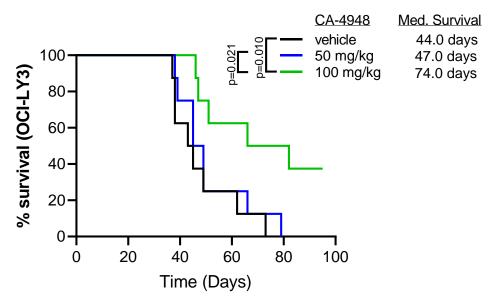


Emavusertib Preclinical PCNSL anti-tumor activity

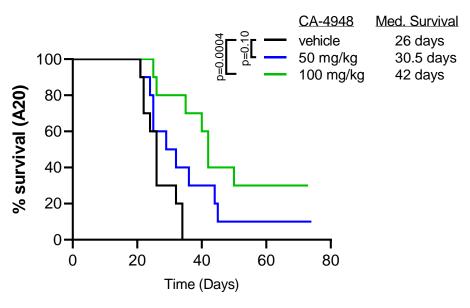




Human MYD88 L265P



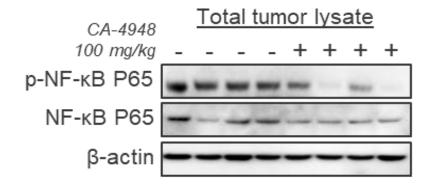
Murine MYD88 WT

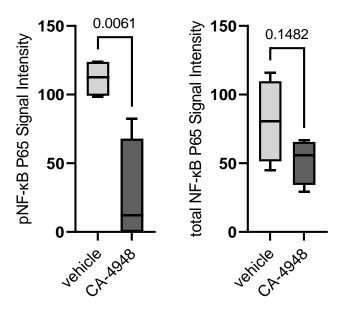




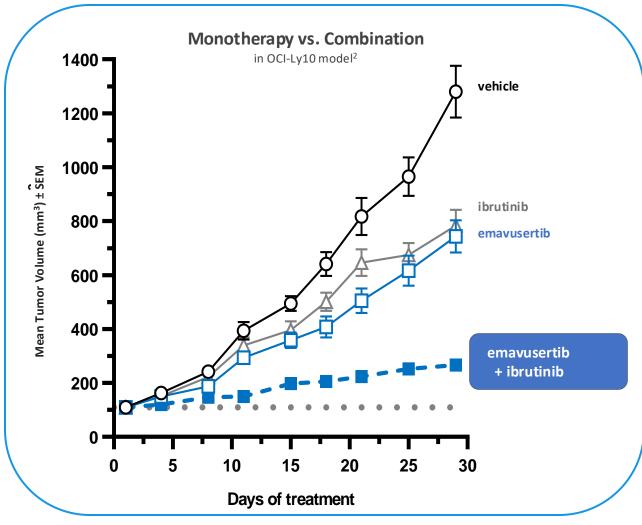
NF-kB biomarker downregulation











Booher et al. Waldenström Roadmap Symposium 2019







TakeAim Lymphoma

Open-label expansion trial evaluating the safety, PK/PD, and clinical activity of emavusertib (CA-4948) + ibrutinib in R/R primary CNS lymphoma

NCT03328078



TakeAim Lymphoma



Study Progress - Enrolling into PCNSL Expansion

Complete

Part A1
Monotherapy Dose Escalation
(MTD)



Complete

Part A2
Combination Dose Escalation

emavusertib (100 - 300 mg BID)

ibrutinib (420 - 560 mg QD)

First patient enrolled: February 2021

Enrolling

Part B
PCNSL Expansion Cohort

emavusertib (100 - 200 mg BID)

ibrutinib (560 mg QD)

First patient enrolled: October 2023



TakeAim Lymphoma PSCNL Expansion Study Design



Study Objectives

Study Population

Primary CNS Lymphoma

ECOG ≤ 2

Age ≥ 18 years

Dosing

Oral, BID

28-day cycles emavusertib (100 - 200 mg BID)

ibrutinib (560 mg QD)

Primary

MTD

RP2D

Safety

Secondary

PK profile

Preliminary anti-cancer activity

Study Phases



Part A1 Monotherapy
Dose Escalation



Part A2 Combination
Dose Escalation

Part B PCNSL Expansion Cohort [ONGOING]



TakeAim Lymphoma



Well-tolerated and manageable adverse event profile at multiple dose levels

Treatment-related adverse events (TRAEs) Grade ≥ 3 in all TakeAim Lymphoma trial patients

Grade 3+ Treatment-Related Adverse Event Occurred in >1 Patient	100 mg BID+IBR	200 mg BID+IBR	300 mg BID+IBR	Total
N (%)	(N=2)	(N=10)	(N=7)	(N=19)
# patients having grade 3+ TRAEs	1 (50)	7 (70)	6 (86)	14 (74)
Platelet count decreased		2 (20)	1 (14)	3 (16)
Alanine aminotransferase increased		1 (10)	1 (14)	2 (11)
Aspartate aminotransferase increased		1 (10)	1 (14)	2 (11)
Fatigue		1 (10)	1 (14)	2 (11)
Hyponatraemia		2 (20)		2 (11)
Lipase increased	1 (50)	1 (10)		2 (11)

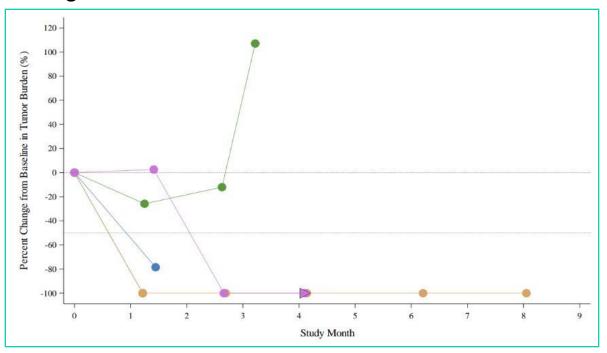
Emavusertib in combination with ibrutinib continues to exhibit a manageable and acceptable safety profile, in heavily pretreated patients, including BTKi-naïve and BTKi-experienced patients.



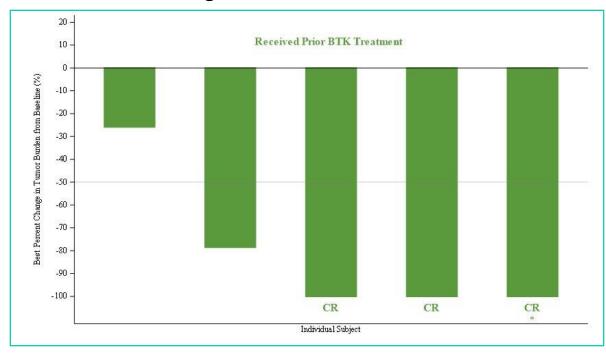
TakeAim Lymphoma: Clinical Outcomes PCNSL



Change in tumor burden over time



Best Percent change in tumor burden



At the October 12, 2023, data cutoff

- 6 patients with PCNSL had received treatment
- Of 5 evaluable patients, all previously treated with BTKi, 3 achieved CR (3/5, 60%)
- One patient had a durable response for approximately 7 months



PCNSL Patient Case Study



PCNSL Expansion – CA-4948 100 mg BID + Ibrutinib 560 mg QD

Patient Background:

• 53 year old, Male

Diagnosis: PCNSL diagnosed on 30June2020

Past Medical History: Unremarkable

Condition:

Baseline: Depression, elevated LFTs, loss of appetite, cerebral edema, mixed

IBS, essential hypertension, hiatal hernia, GERD and obstructive sleep

apnea

Con Meds: ondansetron, sertraline, tadalafil and OTCs for gastroesophageal reflux

and nausea

Prior Tx: (1) MTX, rituximab, Ara-C, thiotepa, high dose BCNU (PR), WBRT &

ASCT

(2) ibrutinib (CR then relapse)

Relapsed: 29Nov2022 with 1 target lesion 13 x 12 mm

Treatment Regimen

Date	Intervention
Feb 24, 2023	Started emavusertib 100 mg BID and ibrutinib 560 mg QD
Jun 30 – Jul 9, 2023	Emavusertib was interrupted due to Gr 3 elevated lipase; resumed at same dose level
Feb 6 – Feb 9, 2024	Emavusertib was interrupted due to Gr 1 vomiting; resumed at same dose level

TEAEs:

Gr 3 amylase increased, and Gr 3 lipase increased: dose interruptions

Outcome:

- Amylase increase recovered to Gr 1 and lipase resolved
- Brain lesion was SD at the end of cycle 2, and disappeared by the end of cycle 4 and onward
- Most recent MRI was performed in Feb '24 and patient is still CR

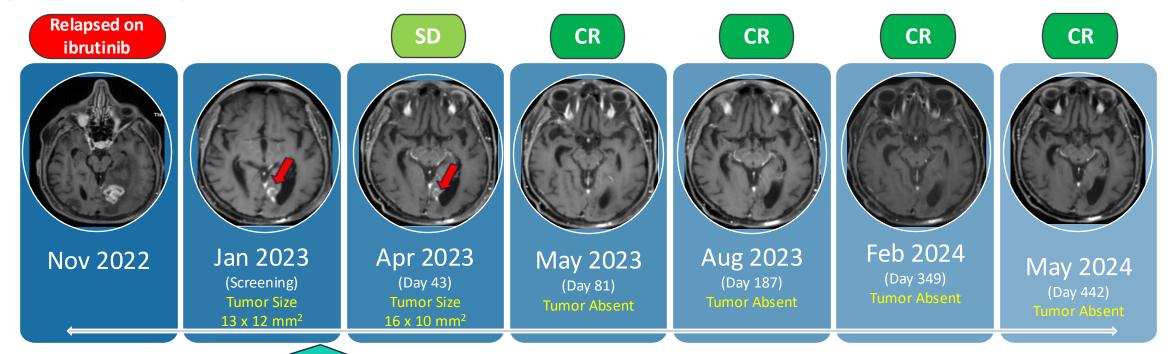
- Time on the study treatment for > 400 days, well tolerated without major long-term safety concerns
- Duration of CR exceeded ~8 months



PCNSL patient who achieved CR on ibrutinib + emavusertib



Previous treatments: MATRIX, HD BCNU/Thiotepa-ASCT, whole brain radiation, ibrutinib



emavusertib + ibrutinib C1D1 2/24/2023

Axial Magnetic Resonance images (MRI) showing pre-treatment and post-treatment PCNSL brain images from one R/R PCNSL patient. After two cycles of emavusertib + ibrutinib, the patient showed stable disease (SD). Complete responses (CR) with absent lesions have been seen after cycle 4.

Consistent with our previous findings, these data support the hypothesis that emavusertib can re-sensitize patients to BTKi therapy, marking a significant advancement in R/R PCNSL treatment.



Genomic profile (GTC Hematology Plus)



Detected Genomic Alterations										
CD79B	ETV6 (2 mutations)	MYD88	KMT2D	PBRM1						
TBL1XR1	IDH1	Chromosomal structural analysis shows: 3p-, 8q-, +9 with bi-allelic deletion of CDKN2A/B), +12, +13, +16, 18q+, and +21.	Expression profiling suggests ABC cell of origin, more aggressive subtype.							

Results Summary

- -Mutations in CD79B, ETV6 (2 mutations), MYD88, KMT2D, PBRM1, TBL1XR1, and IDH1 genes
 - -Chromosomal structural analysis shows: 3p-, 8q-, +9 with bi-allelic deletion of CDKN2A/B), +12, +13,
 - +16, 18q+, and +21.
 - -Increased B-cell markers
 - -No significant increase in BCL1, BCL2, or MYC mRNA.
 - -Expression profiling suggests ABC cell of origin, more aggressive subtype.
 - -These findings are consistent with diffuse large B-cell lymphoma, ABC cell of origin, more aggressive subtype.



Future directions for BTKi + IRAK4i in PCNSL

Concurrent therapy in relapsed/refractory PCNSL

Concurrent therapy in newly diagnosed PCNSL

Maintenance therapy in high-risk patients post-induction, ASCT, CAR-T

Combination with other therapeutic agents- IMiDs and chemotherapy



Ibrutinib monotherapy in newly diagnosed PCNSL

Patients	Continuous CR
#1	76 M
#2	61 M
#3	18 M
#4	7 M



Genomic Profile of Patient #1 with CCR for 76 months (GTC Hematology Plus)



Detected Genomic Alterations									
PIM1 (10 mutations)	RET	SOCS1	CD79B	TET2					
GNAS	ETV6	MYD88	Chromosomal structural analysis shows: 1q+, 2q-, +3, -5, -6, 8q-, 11q+, 12q +, 17p-(TP53 bi- allelic deletion), 19p-, 19q+, and 22q						

Heterogeneity

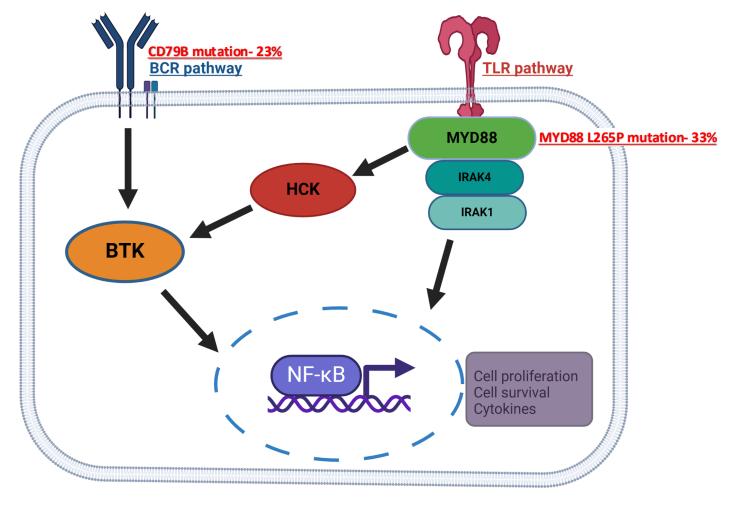
There is a dominant abnormal clone with PIM1 (p.Lys186Asn) mutation. The PIM1 (9 mutations), RET, SOCS1, CD79B, TET2, GNAS, ETV6, and MYD88 mutations are detected in subclones.

Expression	
Increased B-cell markers	Marked increase in Ki67 mRNA
No significant increase in BCL1, BCL2 or MYC mRNA	No increase in ALK mRNA



ABC-DLBCL





MCD and N1 subtypes of DLBCL highly sensitive to ibrutinib
CARD11 and TNFAIP3 mutations in BCR pathway associated with resistance to ibrutinib



Ibrutinib monotherapy in R/R ABC-DLBCL



	N=38
ORR	37%
CR	16%
DOR	4.83 M
mPFS	1.64 M
mOS	6.41 M

4/6 complete responders in remission longer than 1 Y



Young patients (age ≤60 years) with non-GCB DLBCL (Phoenix trial)



Young patients with MCD or N1 subtypes of DLBCL-

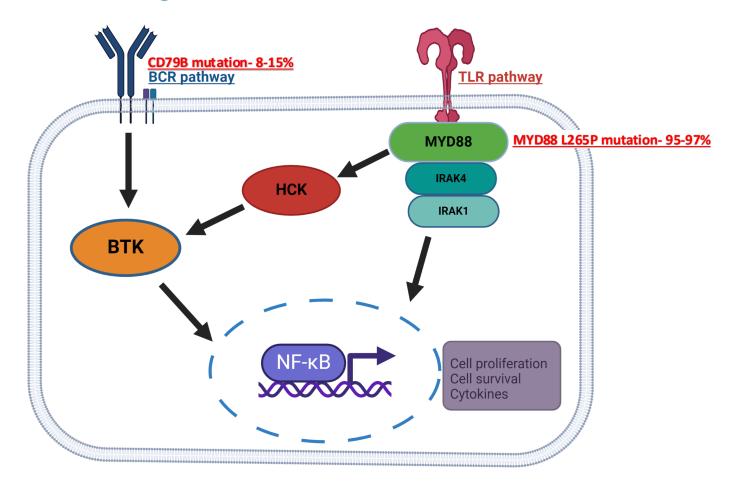
Genetic subtype	3Y Event-free survival				
MCD	IRCHOP- 100% RCHOP- 42.9% (p- 0.0105)				
N1	IRCHOP- 100% RCHOP- 50% (p-0.0161)				



Cancer Cell 2021;39:1643-1653

Waldenstrom Macroglobulinemia





CXCR4 mutation (30-40%) promotes resistance to ibrutinib.

Pathway signature- Oncogenic constitutive activation of BCR and TLR pathways

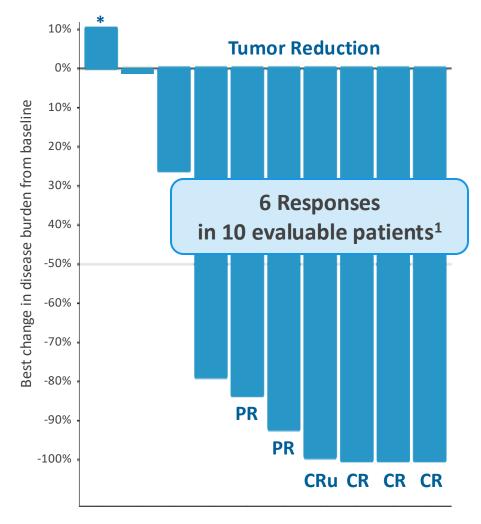
Activated NFKB signaling

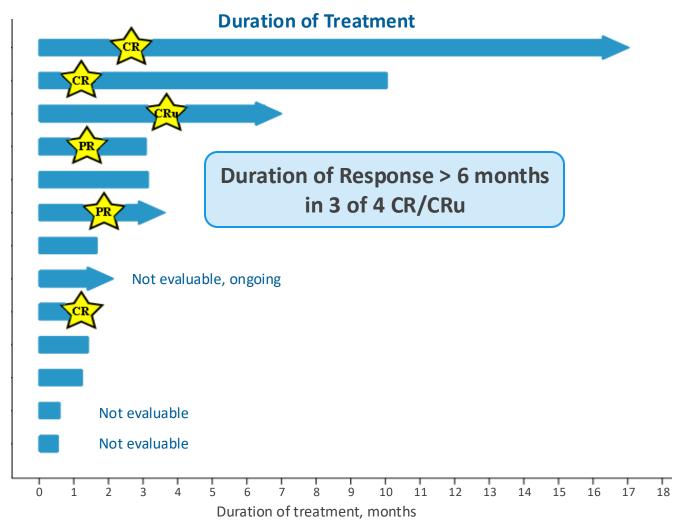


TakeAim Lymphoma

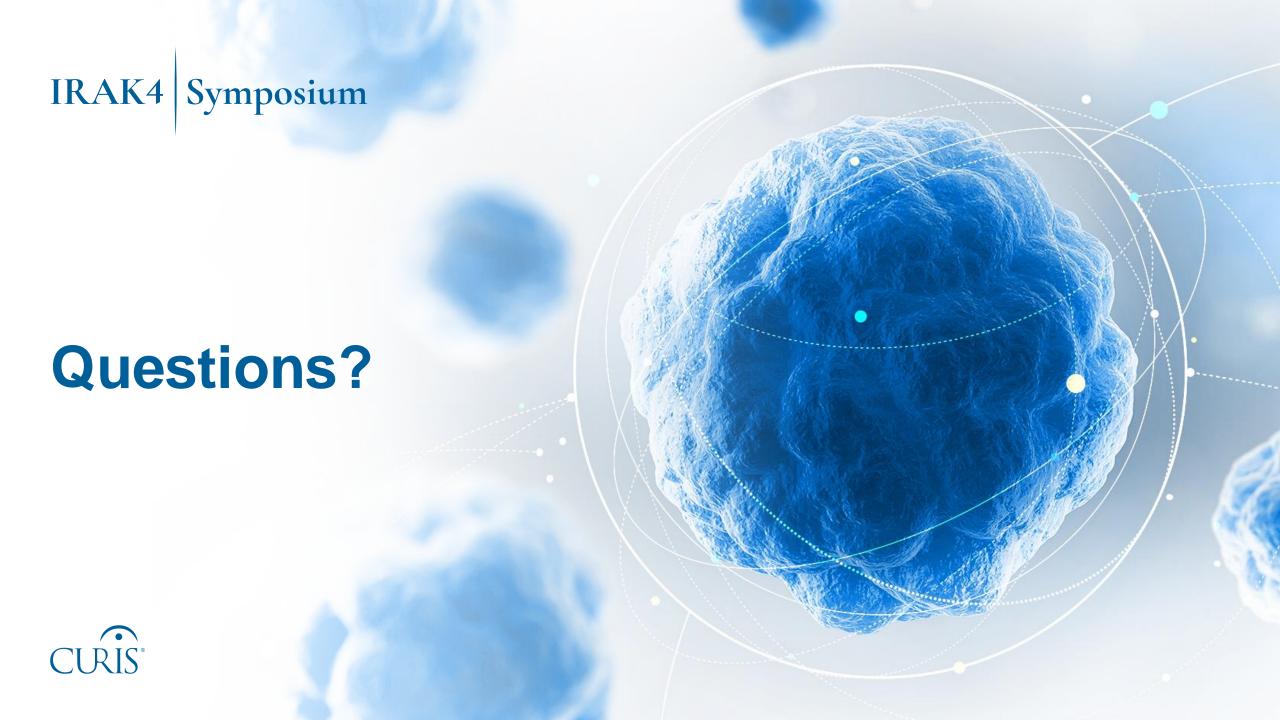


Results in 10 evaluable patients¹ with R/R PCNSL (BTKi-experienced) treated with emavusertib + ibrutinib





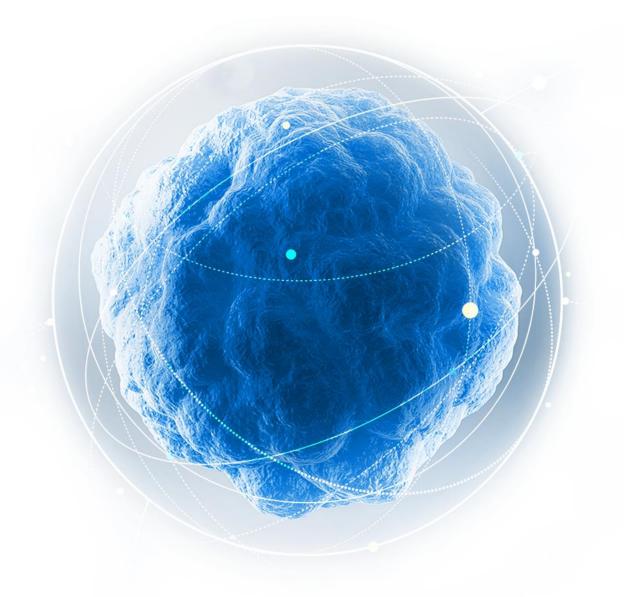






We will now take a short break

Please return in 10 minutes







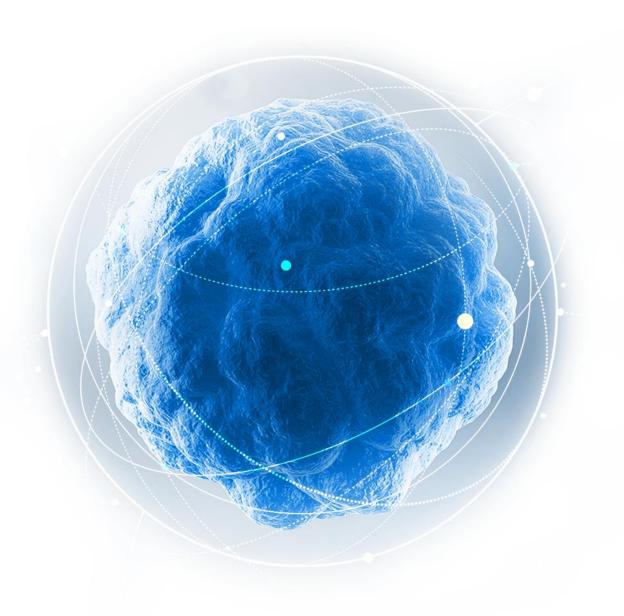
Current and Future Opportunities For IRAK4 Inhibition in Lymphoma Roundtable

Lakshmi Nayak (moderator)

Andrés Ferreri

Han Tun





Discussion Questions

- What are unmet needs for patients with NHLs, especially PCNSL?
- By what mechanism does the BTK inhibitor and IRAK4 inhibitor combination potentiate NF-κB blockade?
- How does blocking the TLR-IRAK4 pathway affect BTKi resistance?
- What are current treatment approaches in lymphomas driven by activated MYD88? What are their limitations?
- Do you see potential for emavusertib to overcome BTK resistance in non-MYD88 lymphomas such as mantle cell lymphoma?
- In what specific populations (line of therapy, prior treatment) is emavusertib + ibrutinib being tested in for PCNSL, and why? What are next steps if the expected signal is seen?
- How would a positive result in the PCSNL expansion trial impact 1L therapies, therapy options for elderly patients, or therapy options for the first-relapse population?
 - Would you expect emavusertib to be combined directly with ibrutinib or only after ibrutinib failure?
- Beyond PCNSL, in which other NHLs do you see potential for emavusertib therapy? Are there particular patient
 populations you would prioritize for this investigation?





IRAK4 as a therapeutic target in GI cancers

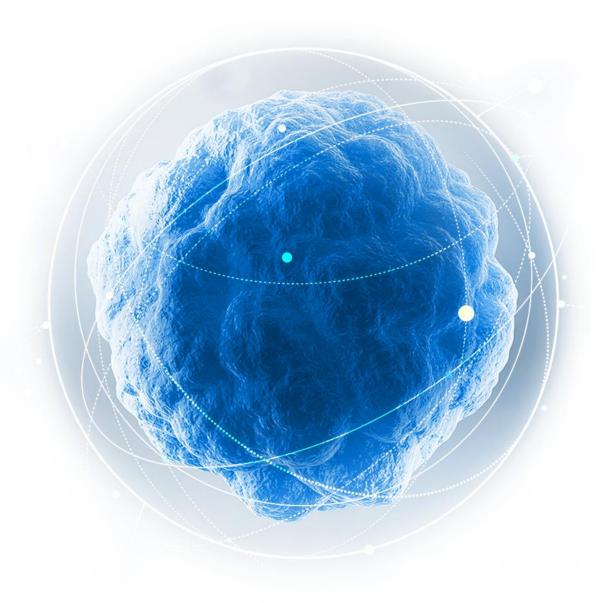
Kian Lim, MD PhD

Professor

Director of GI Oncology

Division of Oncology, Department of Internal Medicine

Washington University School of Medicine in St. Louis





Disclaimer

- Clinical trials: Merck, BMS, Verastem, Takeda, Biomed Valley, Ipsen, Celgene, AstraZeneca, CURIS, PanCan.
- Advisory: Jacobio, Genentech



Outline

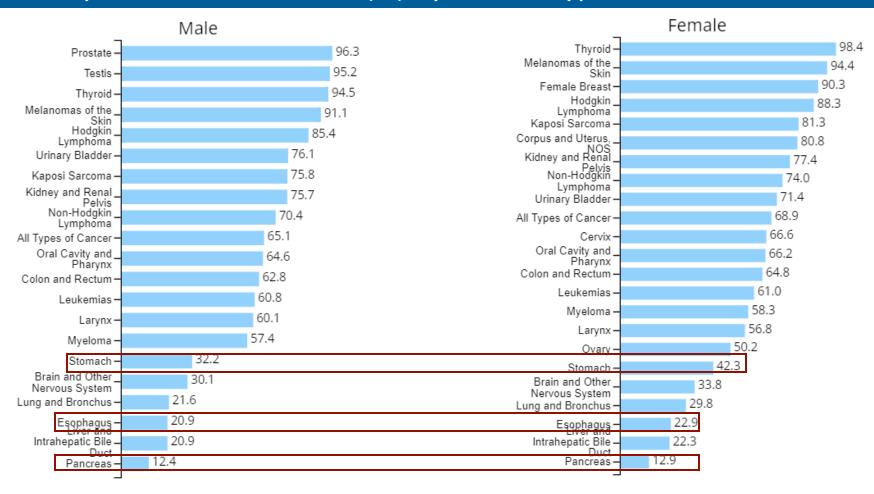


- Introduction
- Role of IRAK4 in GI cancers
 - Pancreatic ductal adenocarcinoma (PDAC)
 - Colorectal cancer (CRC)
 - Gastric cancer





5-year Relative Survival (%) by Cancer Type, United States





PDAC: We still use chemotherapy for all patients



FOLFIRINOX versus Gemcitabine for Metastatic Pancreatic Cancer

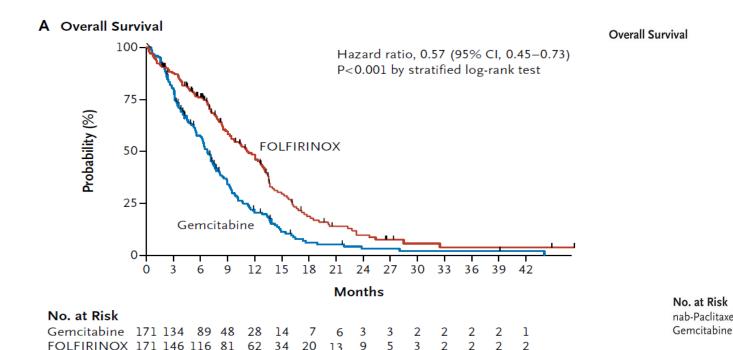
N ENGL J MED 364;19 NEJM.ORG MAY 12, 2011

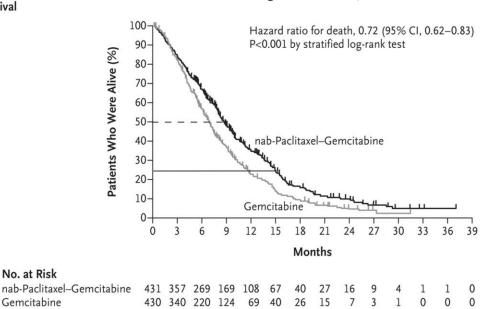
The NEW ENGLAND JOURNAL of MEDICINE

ORIGINAL ARTICLE

Increased Survival in Pancreatic Cancer with nab-Paclitaxel plus Gemcitabine

N Engl J Med 2013; 369:1691-1703



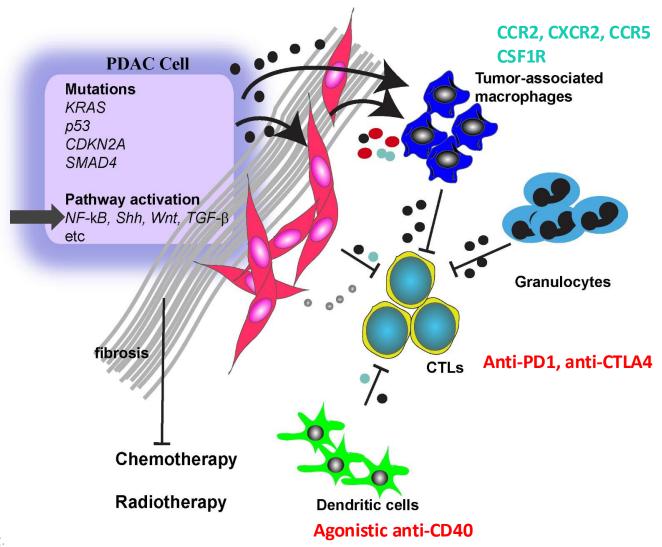


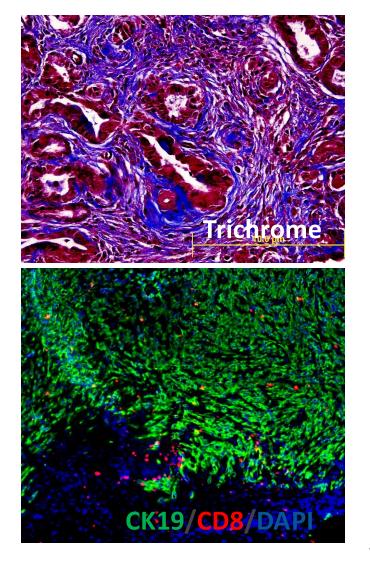
Combination chemotherapies are not curative and have significant side effects



PDAC cells are driven by powerful oncogenes and shielded by a fibrotic stroma







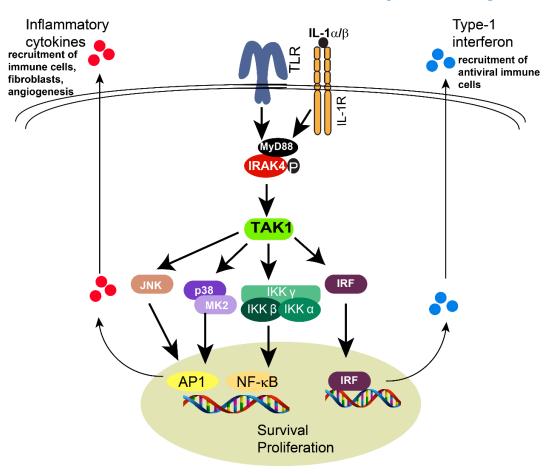


The NF-κB pathway is a therapeutic target in PDAC



- RELA, the p65 subunit of NF-κB, was constitutively nuclear (activated) in **67**% of PDAC and associated with poor prognosis
- Associated with treatment resistance, invasion, metastasis and immune evasion
- Targeted deletion of $IKK\beta$ abrogates PDAC development in KRAS/Ink4A genetic mouse model
- IKK inhibitors are toxic in clinical trials.
- <u>Interleukin-1 Receptor-Associated Kinase 4</u> (IRAK4) is potentially the new target

The innate inflammation pathway



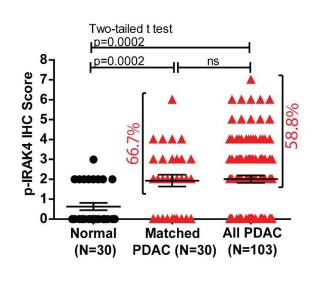
Lim, et al. Cold Spring Harb Perspect Biol. 2013;5(1):a011247.

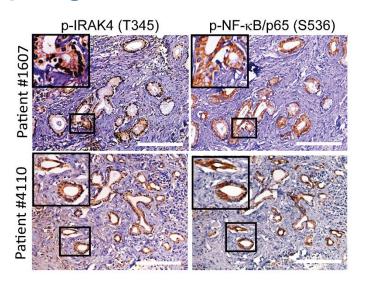


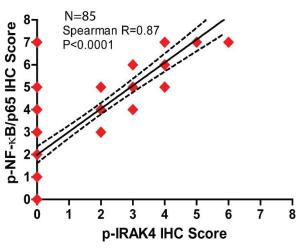
Activated IRAK4 positively correlates with phospho-RELA/NF-κB1 and poor prognosis in PDAC

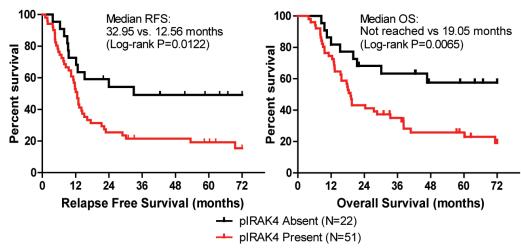


143











Zhang, et al. Clin Cancer Res. 2017;23(7):1748-1759.

IRAK4 kinase activity is required to activate RELA and NF-κB pathway in PDAC



144

	PANC-1						Capan-1					
scramble:	+						+					
shIRAK1:		1	2			1		1	2			1
shIRAK4:				1	2	1				1	2	1
Total IRAK1	_	-	-	-	-				-	•	-	
Total IRAK4	-	-	-	Sic.	100	6.7		-	-	-	***************************************	
p-IRAK4 (T345/S346)	-	44	-	_	-		Marie .	ria.	Eles.		incom	44
p-IKKα/β (S176/180)	=	1	-	· fa	-	1		1	-		-	-
p-NF-κB/p65 (S536)	-	tent .	***	-	-	1007	-	.–	-			Corre
β-actin		•				1	-	_	_	_	-	-



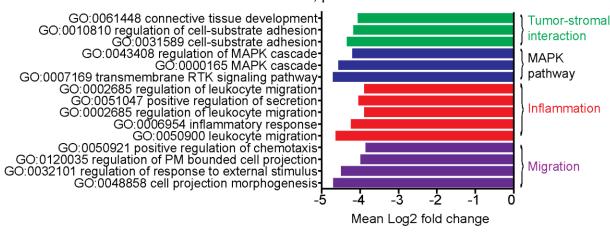
Zhang, et al. *Clin Cancer Res.* 2017;23(7):1748-1759.

IRAK4 controls several suppressive chemokines and checkpoint ligands



KP2 cells (from a KPC mouse)

IRAK4 KO vs. WT: Selected top 15 downregulated GO signatures All FDR<0.001, p<0.0001



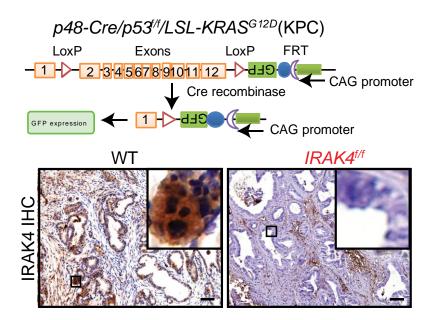
Downregulated in <i>IRAK4</i> KO cells, all FDR<0.05 Independently confirmed by <i>IRAK4</i> rescue and IRAK4 inhibitor					
Gene	log ₂ FC	adj.P.Val	Reported function in PDAC		
CCL2	-0.5411205	1.12E-03	Recruits monocytes, granulocytes		
CCL20	-1.1869978	6.63E-03	Recruits Treg, DC		
CXCL1	-2.0180684	2.24E-08	Recruits monocytes, DC		
CXCL16	-0.7801257	6.98E-05	Directs activated stellate cells		
CXCL3	-4.3381343	5.91E-05	Promotes metastasis		
CXCL5	-3.1309933	4.93E-07	Promotes angiogenesis		
IL-33	-4.1958493	6.11E-10	Promotes M2 TAM polarization		
CD274	-0.6596101	2.28E-02	T cell checkpoint		
Nectin2	-1.5205052	9.58E-08	T and NK cell checkpoint		
TIGIT	-2.3387297	5.74E-04	T and NK cell checkpoint		

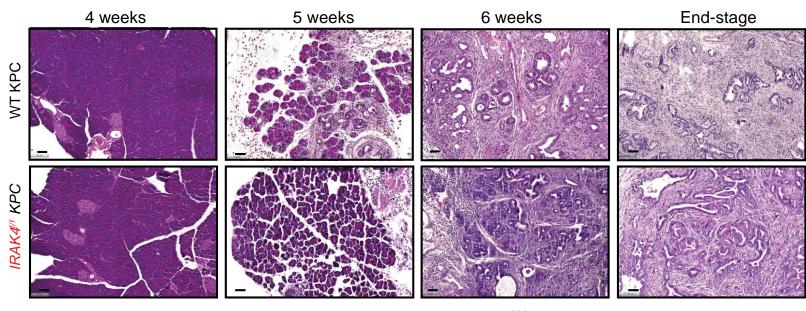
Q: Does tumor *IRAK4* affect T cell response?

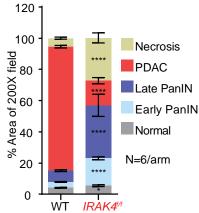


Conditional IRAK4flox/flox KPC





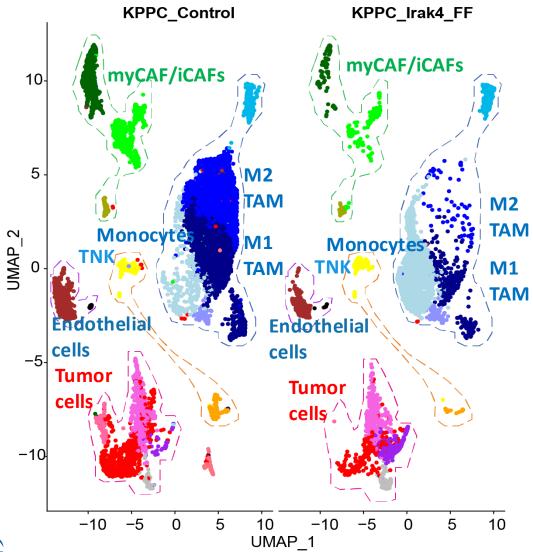


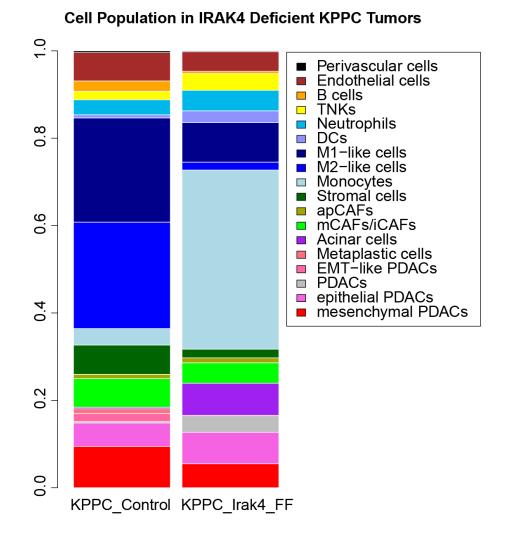




Conditional *IRAK4*^{flox/flox} KPC mice have lower intratumoral TAMs and CAFs



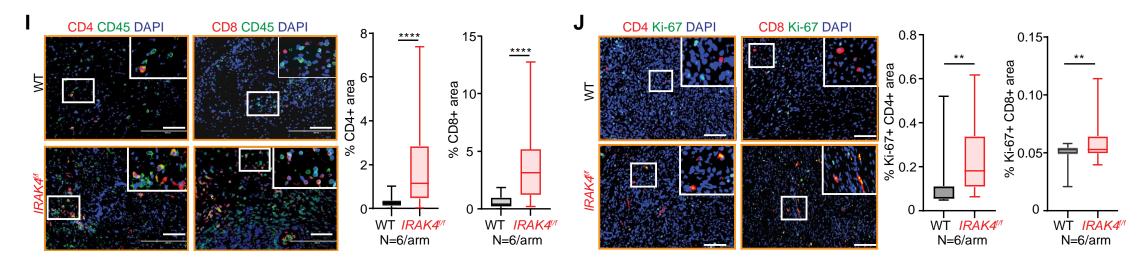




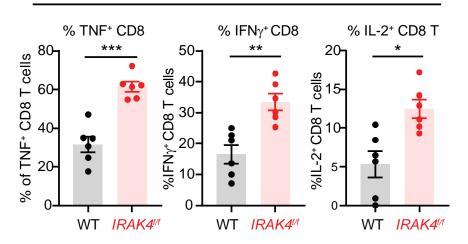


Conditional *IRAK4*^{flox/flox} KPC mice have higher intratumoral and systemic activated T cells





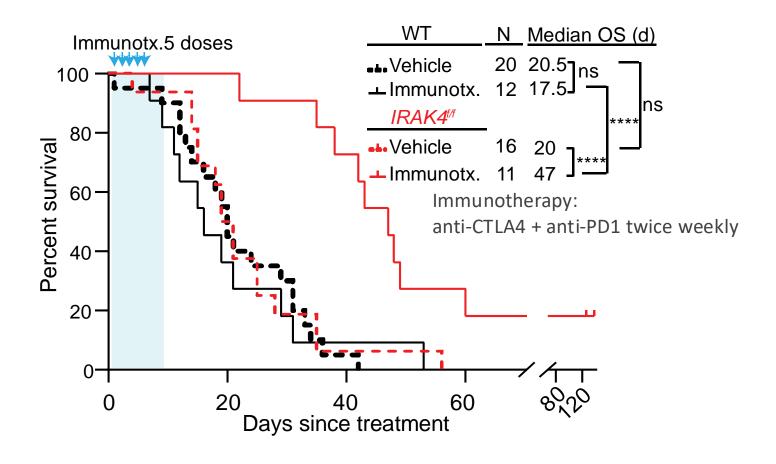
Peripheral blood CD8 T cells





Conditional *IRAK4*^{flox/flox} KPC mice respond to checkpoint immunotherapy



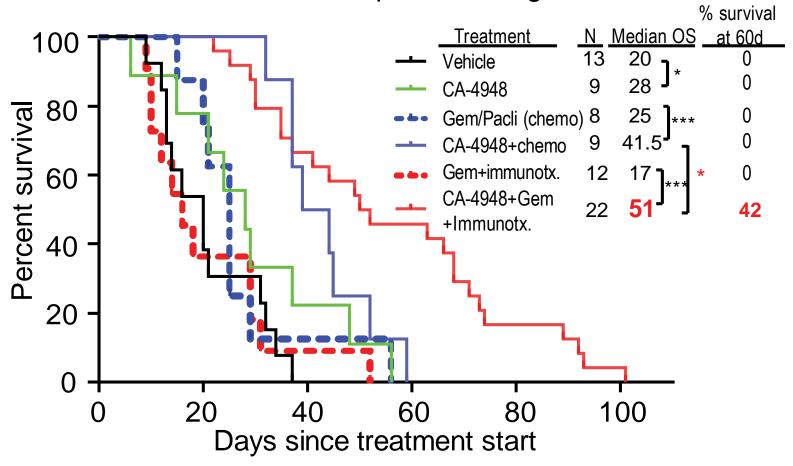




CA-4948 plus chemo-immunotherapy is an effective combination



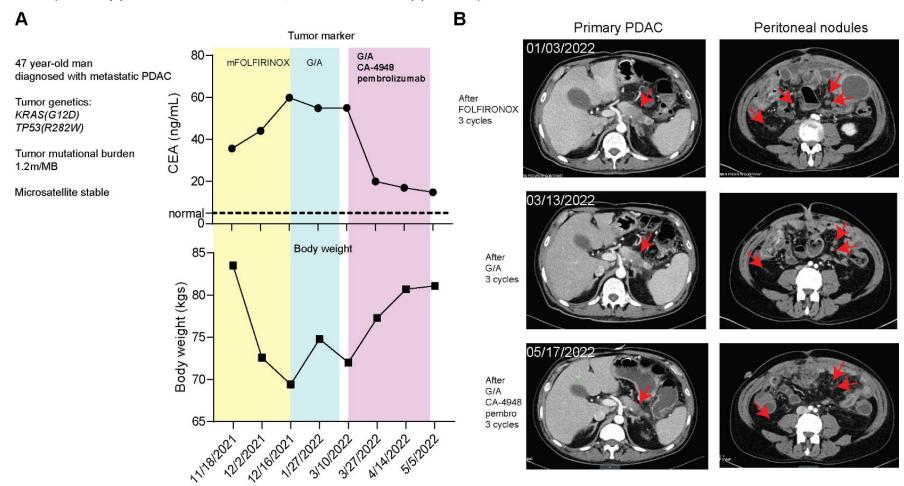
Autochtonous KPC pharmacologic treatment





Single patient compassionate use

Single patient compassionate use of Gemcitabine/Abraxane plus CA-4948 and pembrolizumab (FDA approved IND#161391, Wash U IRB approved)





NCI ETCTN 10522: A Phase 1 Clinical Trial of CA-4948 in Combination with Gemcitabine and Nab-Paclitaxel in Metastatic or Unresectable Pancreatic Ductal Carcinoma (original proposal was to include anti-PD1)



PI (Lead Organization): Patrick Grierson, MD PhD (Wash U) Translational PI: Kian-Huat Lim, MD PhD. (Wash U)

Sites: Columbia, UF, Yale, Vanderbilt, Wash U, UCI....

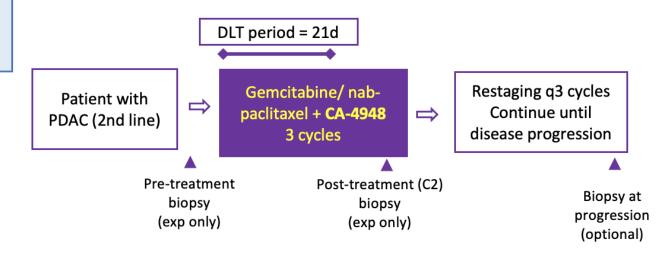


Part A. Dose Escalation (N=up to 18)

CA4948 + gemcitabine/nabpaclitaxel (D1, 8 q21d)

Dose level	CA-4948 (PO BID)
-1	100mg
0 (starting)	150mg
1	200mg
2	250mg
3	MTD (-)1 level
4	MTD

Part B. Dose Confirmation/Expansion (N= up to 18)





^{*}For Dose Levels 3 and 4, gem/nab-paclitaxel will be given D1, 8, 15 q28d

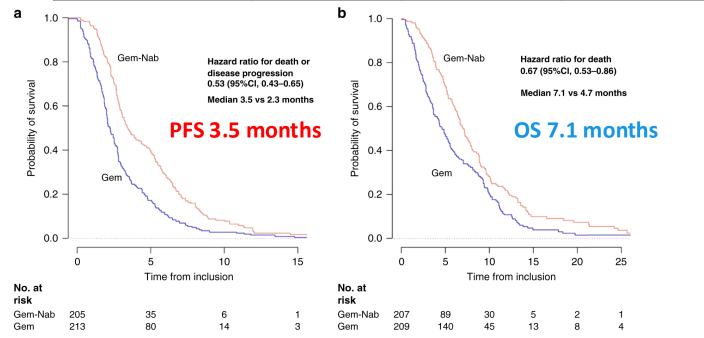
Gemcitabine + Nab-paclitaxel or Gemcitabine alone after FOLFIRINOX failure in patients with metastatic pancreatic adenocarcinoma: a real-world AGEO study

IRAK4 Symposium

Sonia Zaibet¹, Vincent Hautefeuille², Edouard Auclin³,⁴, Astrid Lièvre⁵, David Tougeron ⊙6, Mathieu Sarabi², Marine Gilabert8, Julie Wasselin², Julien Edeline ⊙9, Pascal Artru¹0, Dominique Bechade¹¹, Clémence Morin², Agnes Ducoulombier¹², Julien Taieb ⊙¹ and Simon Pernot ⊙¹¹™

18 French centers

Best response	Gem-Nab <i>(N</i> = 219)	Gem (<i>N</i> = 208)	
Partial response—N (%)	22 (11.28%)	13 (8.28%)	
Stable disease—N (%)	87 (44.62%)	38 (24.2%)	D 4 0 001
Progressive disease—N (%)	86 (44.1%)	106 (67.52%)	<i>P</i> < 0.001
NA	24	51	







Gemcitabine plus nab-paclitaxel for advanced pancreatic cancer after first-line FOLFIRINOX: single institution retrospective review of efficacy and toxicity

Yue Zhang^{1*}, Howard Hochster², Stacey Stein² and Jill Lacy²

Table 1 Patient characteristics at initiation of gemcitabine and nab-paclitaxel

Age, years	61 (50-74)
Sex, no (%)	
Male	11 (39.3 %)
Female	17 (60.7 %)
Disease, no (%)	
Metastatic	23 (82.1 %)
Locally advanced	5 (17.9 %)
ECOG performance status, no (%)	
≤1	27 (96.4 %)
≥2	1 (3.6 %)
Median no of FOLFIRINOX cycles (range)	12 (5-46)
Median interval from last FOLFRINOX to initiation of $G + Nab-P$, weeks (range)	5.4 (1.7-40.3)
N = 28	

Table 3 Efficacy of second line gemcitabine and nab-paclitaxel

tune:	PFS 3 months	
Median time to treatment failure, weeks	12.0 (2.0–36.0)	
Median overall survival, weeks	23.0 (2.1-85.4)	
Response by RECIST, n (%)	OS 5.7 months	
Partial response	5 (17.9)	
Stable disease	8 (28.6)	
Progressive disease	12 (42.8)	
Inevaluable	3 (10.7)	
Serologic response, n (%) ^a		
CA 19-9	13 (46.4)	
CEA	11 (39.3)	

^a >30 % decrease from pre-treatment baseline



Exp Hematol Oncol (2015) 4:29 154

Summary and next steps for PDAC

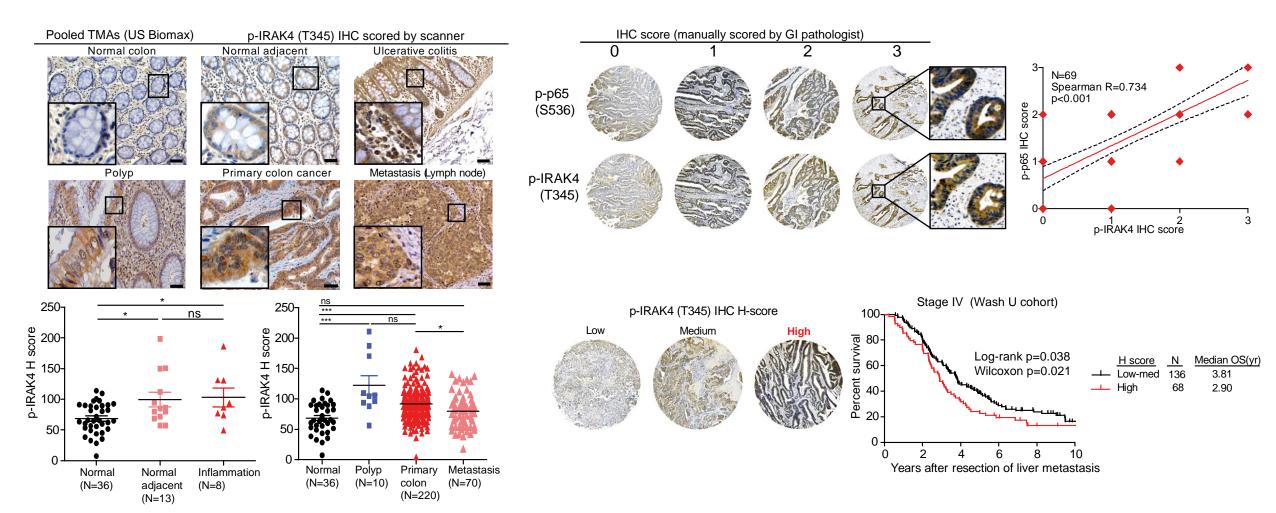
- 1. Adding CA-4948 to G/A is tolerable without added toxicities beyond chemo
 - Dose escalation is ongoing to achieve DLT → dose expansion

2. Pre- and post-treatment tumor samples will be obtained from dose expansion to study immunological changes \rightarrow justification of adding anti-PD1



Activated IRAK4 positively correlates with phospho-RELA and poor prognosis in colon cancer



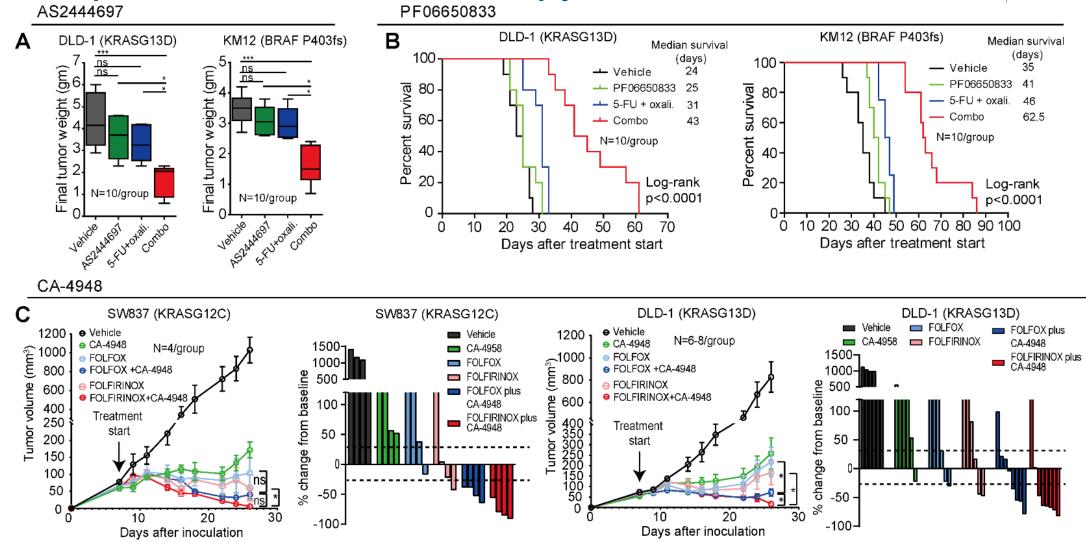




Li, et al. JCI Insight. 2019;4(19):e130867.

IRAK4i potentiates chemotherapy







Study Rationale/Hypothesis



- Activated IRAK4 is a driver of NF-kB activity, leading to chemo-resistance and poor prognosis
- Our preclinical data strongly support combining CA-4948 with chemotherapy and ICB in gastrointestinal cancer
- Standard first-line therapy includes FOLFOX-based chemotherapy in several GI cancer, including colorectal cancer and gastroesophageal cancer
- Improving efficacy of first-line therapy by adding a novel agent with no overlapping toxicities may improve long-term survival in this population
- ➤ We hypothesize that combining CA-4948 with FOLFOX and bevacizumab or nivolumab will be well-tolerated while improving anti-tumor efficacy



NCI ETCTN 10655: Phase 1 Clinical Trial of CA-4948 in Combination with FOLFOX + Bevacizumab as Frontline Treatment in Patients with Metastatic Colorectal Cancer



PI: Susanna Ulahannan, M.D., Stephenson Cancer Center at Oklahoma University

CrD Mentor: Thomas George, M.D., University of Florida in Gainesville

Translational PI: Kian-Huat Lim, M.D., Ph.D., Washington University in St. Louis

Status: Trial protocol writing in final stage

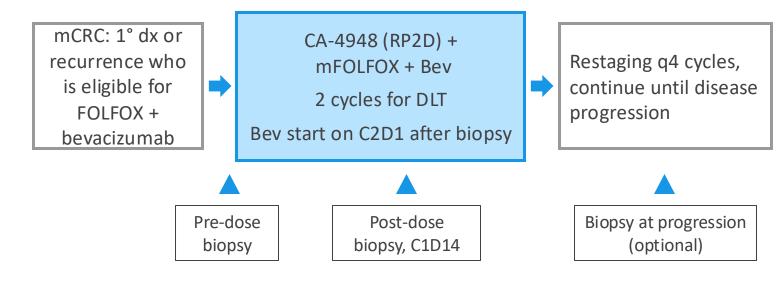
Target population: Newly diagnosed metastatic CRC patients

Part A. Dose Finding Phase (N=up to 18)

CA-4948 (PO, BID, D1-14) + mFOLFOX6+Bev, D1, q2w, DLT period: 28 days

Dose level	CA-4948 Dose (PO, BID)
-1	100 mg
1 (starting)	150 mg
2	200 mg
3	250 mg

Part B. Dose Expansion (N=up to 12)





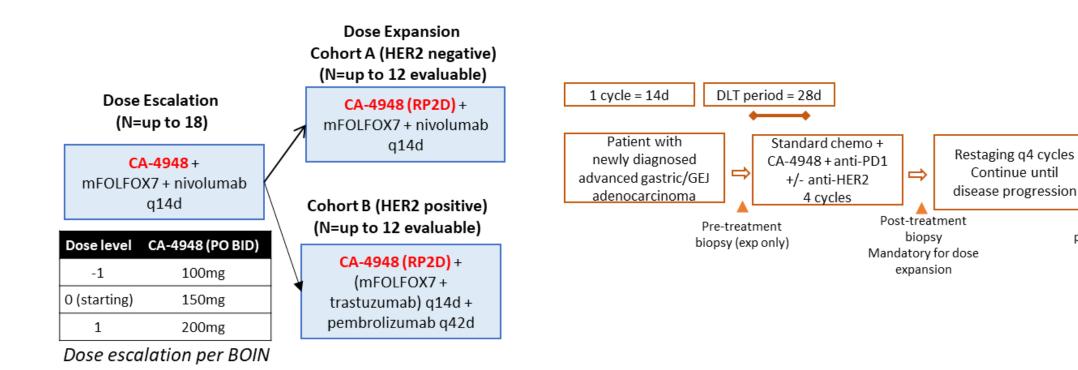
IRAK4 Symposium

Biopsy at

progression

(optional)

Wash U IIT: Phase I trial of CA-4948 in combination with FOLFOX/PD-1 inhibitor +/- trastuzumab for untreated unresectable gastric and esophageal cancer





First-line nivolumab plus chemotherapy versus chemotherapy alone for advanced gastric, gastro-oesophageal junction, and oesophageal adenocarcinoma (CheckMate 649): a randomised, open-label, phase 3 trial

Yelena Y Janjigian*, Kohei Shitara*, Markus Moehler, Marcelo Garrido, Pamela Salman, Lin Shen, Lucjan Wyrwicz, Kensei Yamaguchi, Tomasz Skoczylas, Arinilda Campos Bragagnoli, Tianshu Liu, Michael Schenker, Patricio Yanez, Mustapha Tehfe, Ruben Kowalyszyn, Michalis V Karamouzis, Ricardo Bruges, Thomas Zander, Roberto Pazo-Cid, Erika Hitre, Kynan Feeney, James M Cleary, Valerie Poulart, Dana Cullen, Ming Lei, Hong Xiao, Kaoru Kondo, Mingshun Li, Jaffer A Ajani

CheckMate 649

1581 patients, non-HER2 positive Chemo (N=792) FOLFOX q2weeks or CAPOX q3weeks

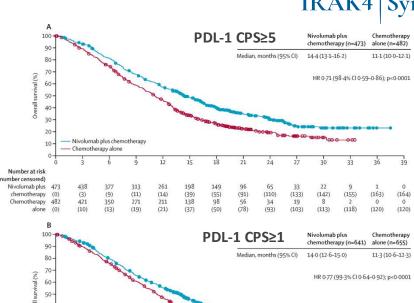
Chemo + anti-PD1 (N=789)
Nivolumab 240mg q2weeks or 360mg q3weeks

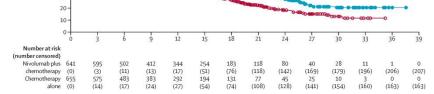
ORR 58%
Median PFS 7.7 months
Median OS 13.8 months

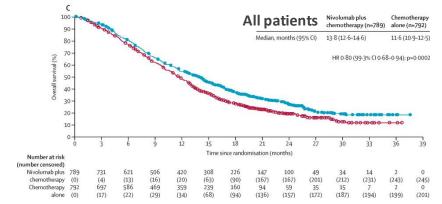




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Lancet. 2021;398(10294):27.

IRAK4 is a novel therapeutic target in GI cancers



- Tumor intrinsic IRAK4 activation drives MAPK and NF-κB activity → treatment resistance
- Tumor IRAK4 activation drives secretion of suppressive chemokines and checkpoint ligands (PD-L1, Nectin2) that collectively exhaust T cells
- Targeting IRAK4 represents a promising strategy to potentiate chemo- and immunotherapies

Ongoing work

- Role of IRAK4 in other cancer types: Lung, colon
- Cell type specific role: myeloid (*CSF1R-Cre:IRAK4*^{f/f})
- Downstream signaling targets driven by IRAK4



Thanks for your attention!

Acknowledgements

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- Ashenafi Bulle, Sapana Bansod, Timothy Chen
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- Huaping Li, Yutong Geng

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- Caron Ridgen, MD
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Biostatistics Core

Esther Lu







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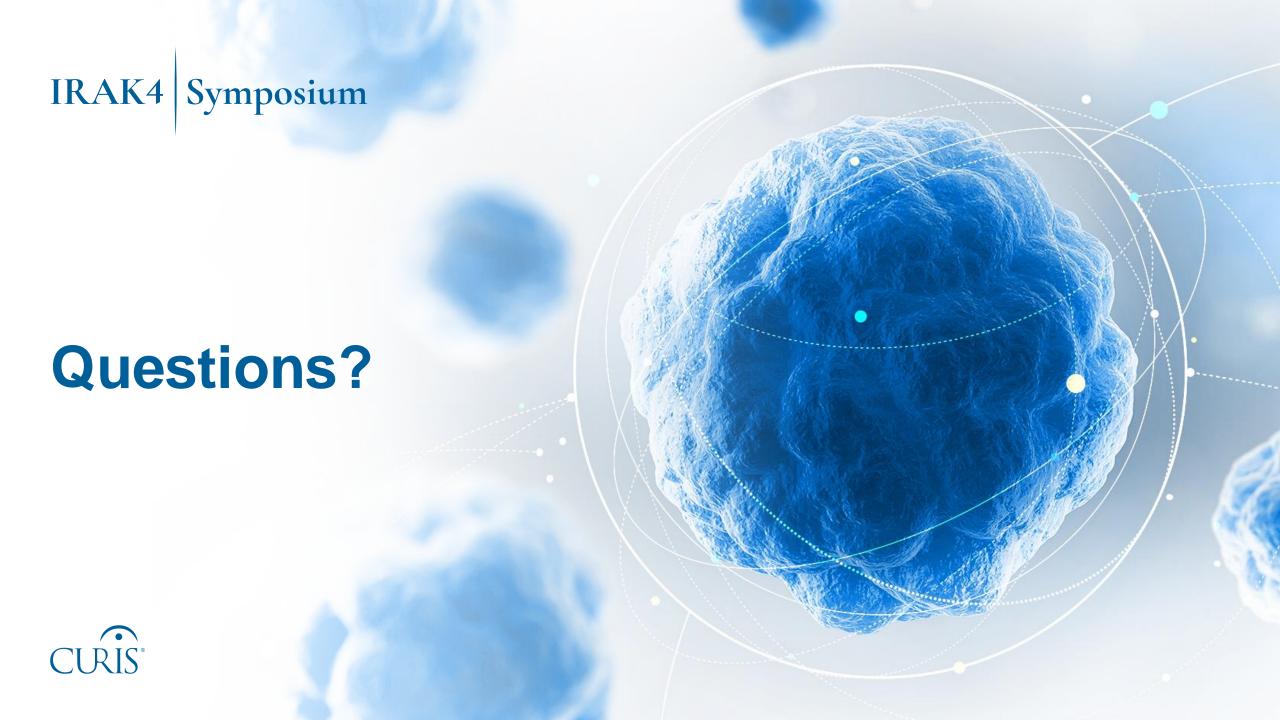














Emavusertib plus Pembrolizumab for Melanoma Brain Metastases

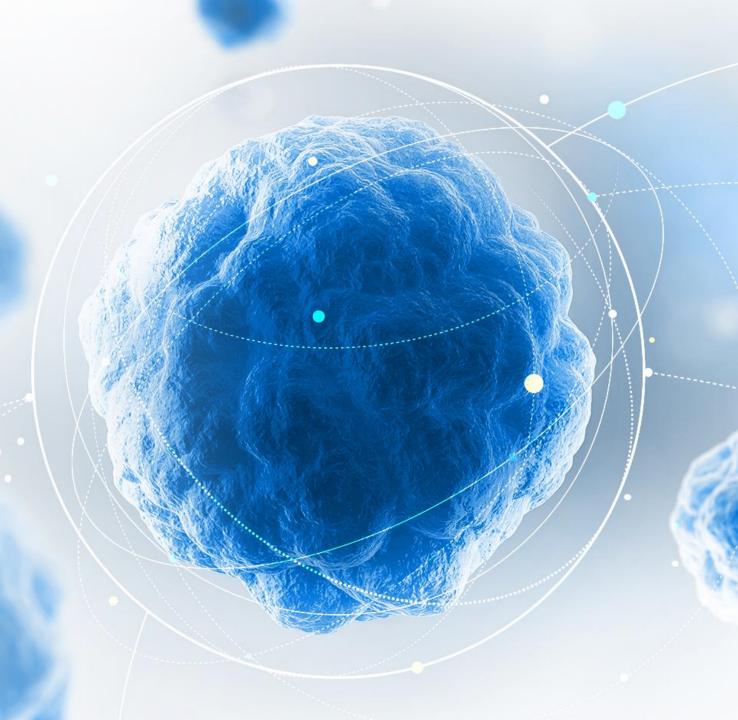
Dr. Bently Doonan, MD, MS

Assistant Professor

University of Florida Health Cancer Center

UF Wells Brain Tumor Immunotherapy Center

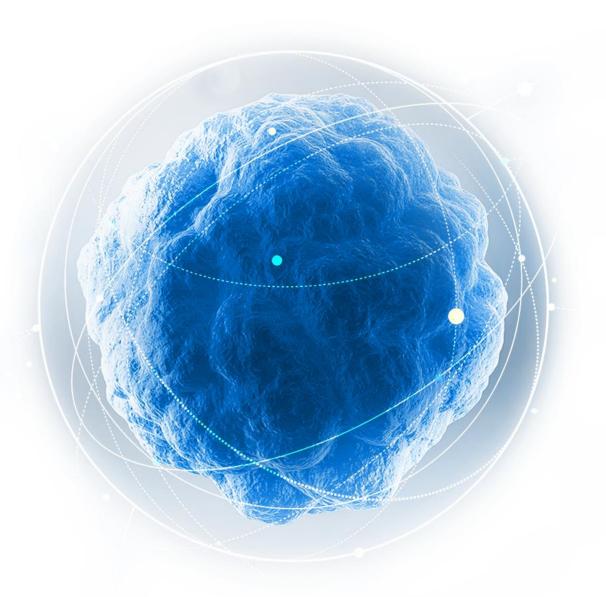






Disclosures

- Preclinical research support from Curis, Inc
- Clinical trial research support from Merck, Inc
- Clinical trial drug support from Curis, Inc
- Advisory board: Immunocore, Inc; Pfizer, Inc; Replimune

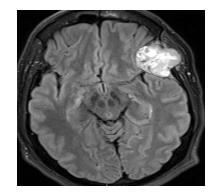




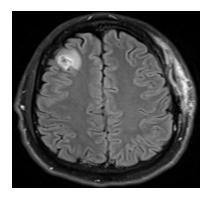
Melanoma Brain Metastases (MBM)

IRAK4 Symposium

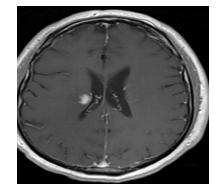
- Melanoma diagnoses are expected to continue to increase over the next 20 years, particularly in men^[1, 2]
- Metastatic melanoma caries a 60-75% risk of development of brain metastases during the course of the disease^[3, 4]
- Standard of care options include surgical resection, stereotactic radiosurgery, systemic immunotherapy, or targeted therapy in BRAF mutated patients^[5-8]
- Even with maximally tolerated therapy, roughly 50% of patients will experience intracranial relapse within the first 3 years^[9]
- Resistance to therapy develops at an increased rate intracranially and is often the only site of progression^[10]



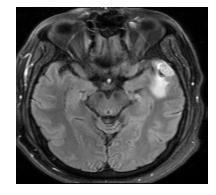
35 yo previously stage IIIC with intracranial recurrence as de novo metastasis



MBM recurrence following lpi/Nivo induction + Surgery + SRS



50 yo 10 months after Ipi/Nivo induction, while in CR oral TKI for C-KIT mutation

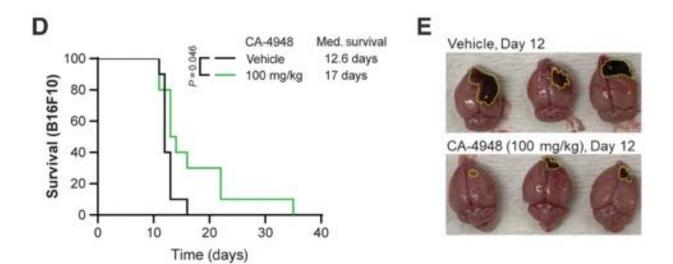


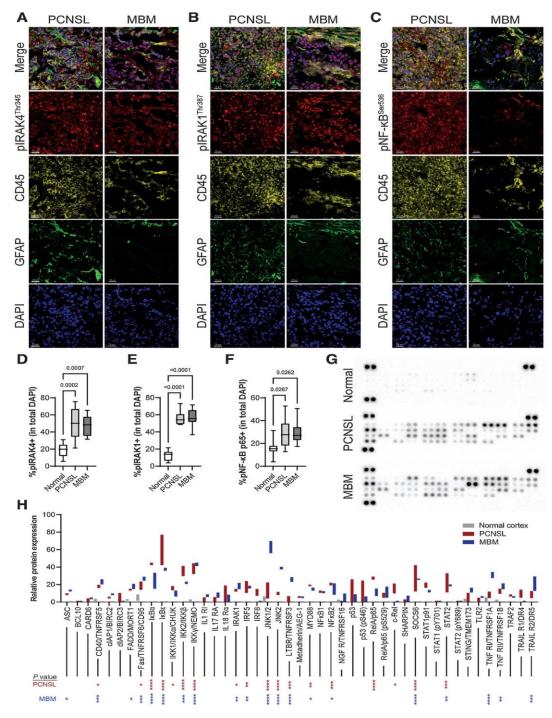
56 yo 6 months after Ipi/Nivo induction, while on second line Opdualag



IRAK 4 as Target in MBM

- We have previously shown the upregulation in IRAK4, Myd88 and downstream transcription factors in MBM^[11]
- We have also shown the intracranial permissibility and penetrance of Emavusertib through the BBB^[11]
- Emavusertib has modest single agent activity in aggressive MBM models^[11]

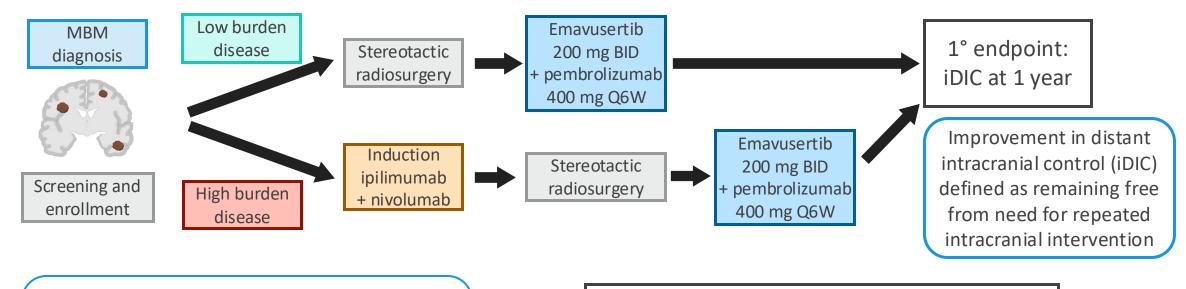






Phase 1/2 Study of Emavusertib in combination with pembrolizumab IRAK4 Symposium following SRS in patients with melanoma brain metastases

PI: Dr. Bently Doonan. Sponsors: Merck, Inc and Curis, Inc: NCT05669352



- Simon's 2 stage design
- Phase I safety run in (fixed dose emavusertib)
- No max limit of MBM, inclusion characteristic is ability to receive definitive SRS by institutional standard

Brain MRI q3 months for response monitoring RANO-BM



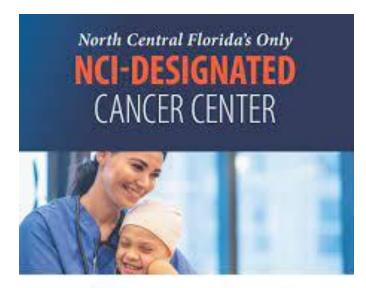
Questions and Thanks

IRAK4 Symposium

- Team and Collaborators
 - Dr. Duane Mitchell
 - Dr. Christina Von Roemeling
 - Dr. Changlin Yang
 - Jeet Patel
 - Savannah Carpenter
 - Skye Speakman
 - Dr. Thom George
 - Dr. Ji-Hyun Lee
 - Dr. April Salama (Duke University)
 - Entire UFBTIP

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- Merck, Inc MISP Portal
- James and Esther King Biomedical Research Program
- Gatorade
- Adam Rosen Foundation

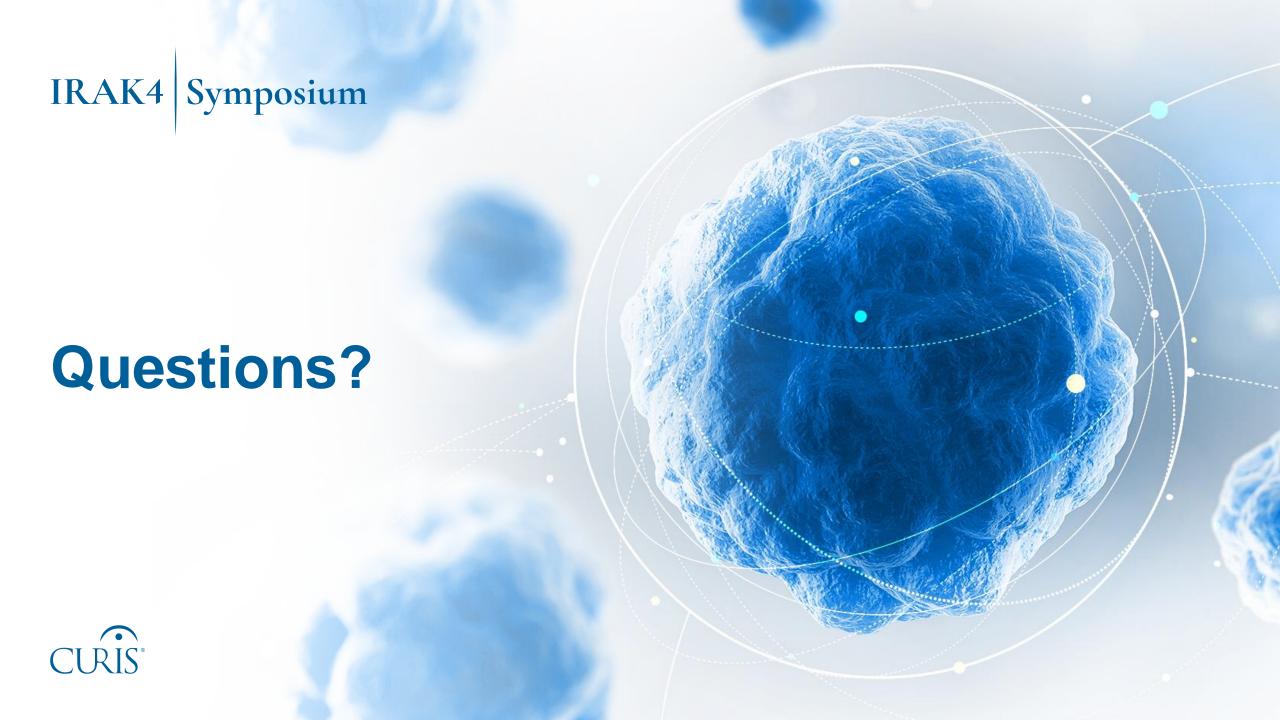












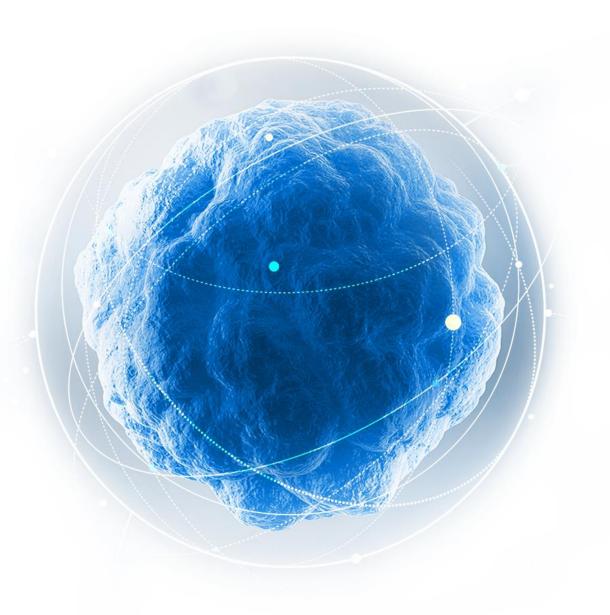


Panel Discussion: Advances and Next Steps

Eric S. Winer (moderator)

All speakers invited to join





Discussion Questions

- What do you think is the main role of IRAK4 inhibition as an anti-cancer mechanism? A signaling inhibitor, an anti-inflammatory agent, a T-cell exhaustion fixer, or something else?
- Do you see most benefit for emavusertib as a monotherapy or as part of a combination regimen? What combinations?
- What are the most important safety signals to consider in developing emavusertib alone and in combination?
 - Which patients might be considered for a modified dosing approach?
- Are there additional malignancies outside those discussed in this symposium where TLR-IRAK4 pathway activation may play a role in either initial pathogenesis or treatment resistance?
- How do you see the treatment landscape for the malignancies discussed today changing? How would emavusertib fit into this landscape?
- Do you expect specific mutationally defined subtypes to be more likely to respond to emavusertib?
- What advances in treatment do you hope to see in the next 5 years?



