

## **Corporate Presentation**

NASDAQ: CRIS



## Cautionary Note Regarding Forward Looking Statements

This presentation contains certain forward-looking statements about Curis, Inc. ("we," "us," or the "Company") within the meaning of the Private Securities Litigation Reform Act of 1995, as amended. Words such as "expect(s)," "believe(s)," "will," "may," "anticipate(s)," "focus(es)," "plans," "mission," "strategy," "potential," "estimate(s)", "intend," "project," "seek," "should," "would" and similar expressions are intended to identify forward-looking statements. Forward-looking statements are statements that are not historical facts, reflect management's expectations as of the date of this presentation, and involve important risks and uncertainties. Forward-looking statements herein include, but are not limited to, statements with respect to the timing and results of future clinical and pre-clinical milestones; the timing of future preclinical studies and clinical trials and results of these studies and trials; the clinical and therapeutic potential of our drug candidates; the proposed focus on emavusertib and management's ability to successfully achieve its goals. These forward-looking statements are based on our current expectations and may differ materially from actual results due to a variety of important factors including, without limitation, risks relating to: whether and when the U.S. Food and Drug Administration may remove the partial clinical hold on the combination therapy phase (Phase 1b) and the expansion phase (Phase 2a) of the Phase 1/2 TakeAim Leukemia trial, or may take further regulatory action with regard to this trial, whether any of our drug candidates will advance further in the clinical development process and whether and when, if at all, they will receive approval from the FDA or equivalent foreign regulatory agencies; whether historical preclinical results will be predictive of future clinical trial results; whether historical clinical trial results will be predictive of future trial results; whether any of our drug candidate discovery and development efforts will be successful; whether any of our drug candidates will be successfully marketed if approved; our ability to achieve the benefits contemplated by our collaboration agreements; management's ability to successfully achieve its goals; the sufficiency of our cash resources; our ability to raise additional capital to fund our operations on terms acceptable to us or the use of proceeds of any offering of securities or other financing; general economic conditions; competition; and the other risk factors contained in our periodic and interim reports filed with the Securities and Exchange Commission which are available on the SEC website at www.sec.gov. You are cautioned not to place undue reliance on these forward-looking statements that speak only as of the date hereof, and we do not undertake any obligation to revise and disseminate forward-looking statements to reflect events or circumstances after the date hereof, or to reflect the occurrence of or non-occurrence of any events, except as required by law.



# **Corporate Overview**

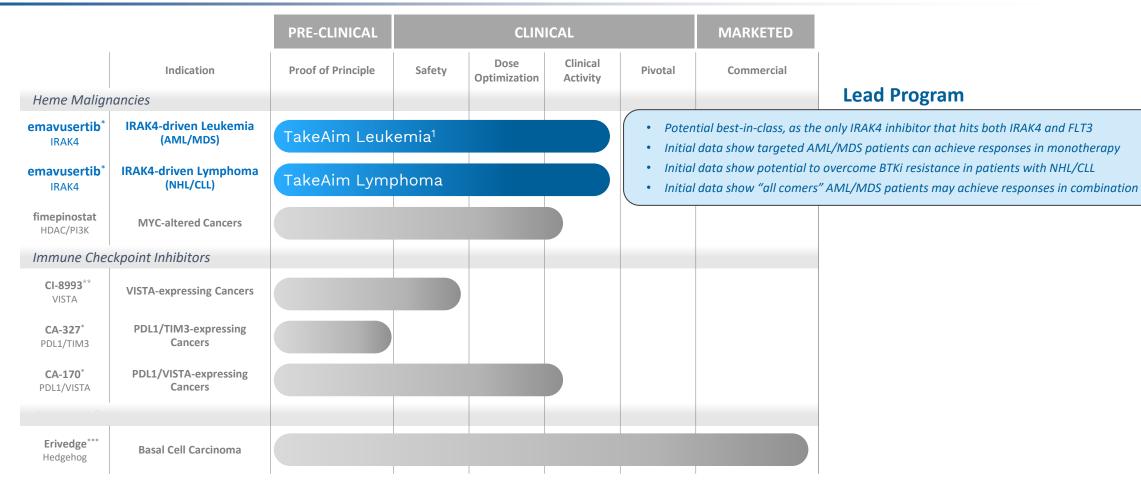
## Summary

Investment Thesis	Curis develops novel cancer therapeutics in areas of significant unmet patient need  Cash runway into 2025 – \$85.6M as of December 31, 2022	
Lead Program	<ul> <li>Emavusertib is positioned to become the cornerstone agent in heme malignancies</li> <li>IRAK4-L is the most prevalent driver of disease in AML/MDS<sup>1,2</sup></li> <li>IRAK4i has a synergistic effect when combined with ibrutinib in NHL</li> </ul>	
Market Opportunity	AML/MDS: 317K patients <sup>3</sup> (current standard of care is HMA)  NHL/CLL: 1.8M patients <sup>3</sup> (current standard of care is BTKi)	
2023 Milestones	Q3: Full release of clinical hold on TakeAim Leukemia Q3: Agreement with FDA on Recommended Phase 2 Dose Q4: Updated data from both the R/R Leukemia and R/R Lymphoma studies	



## **Pipeline**

### Curis develops first-in-class cancer therapeutics



<sup>&</sup>lt;sup>1</sup> In April 2022, the U.S. Food and Drug Administration ("FDA") placed the TakeAim Leukemia study on partial clinical hold. In August 2022, the FDA notified Curis that it may resume enrollment in the monotherapy dose finding phase of the study. The partial hold remains in place for the combination therapy and expansion phases of the study.



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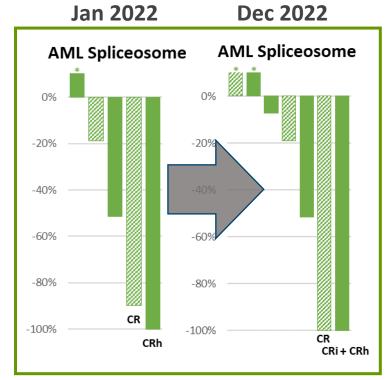
Emavusertib: Clinical Data Update at 2022 ASH

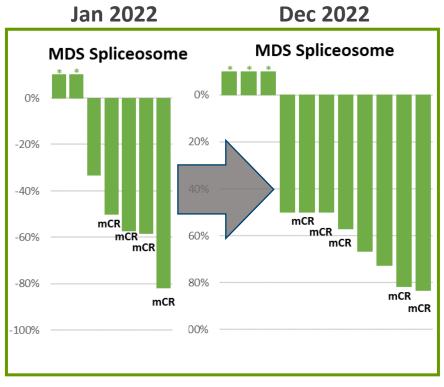
### **Emavusertib Current Data**

#### ASH 2022 Update Reflects Significant Progress in AML/MDS

- New data roughly doubles the dataset for targeted patient population (FLT3 or Spliceosome mutation)
- Data continue to show deep and durable responses
- Data continue to show competitive profile vs. currently available therapies











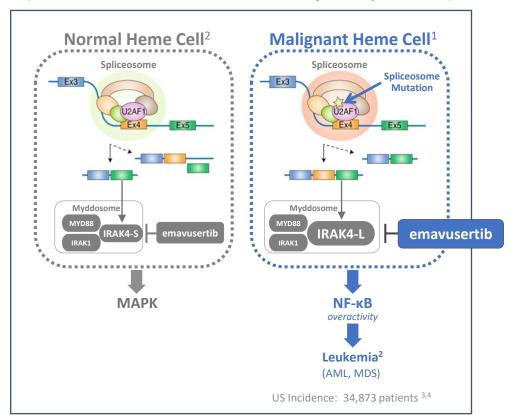
IRAK4 Biology and Mechanism of Action

### **Emavusertib Mechanism of Action**

#### IRAK4 is a novel and important target across multiple heme malignancies

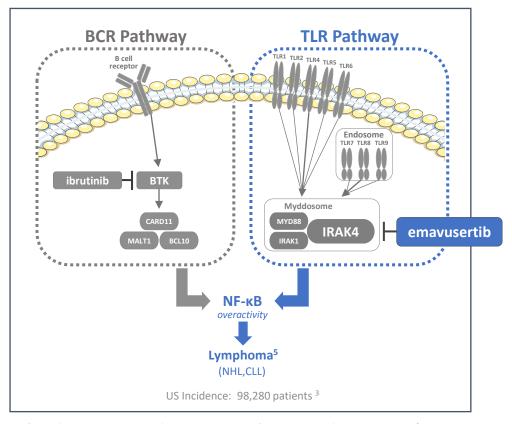
#### **IRAK4** in AML/MDS

Spliceosome mutation drives overexpression of IRAK4-L (which causes constitutive activation of the myddosome)



#### **IRAK4 in NHL/CLL**

TLR Pathway is dependent upon IRAK4 for function (2<sup>nd</sup> pathway driving NF-кВ overactivity)



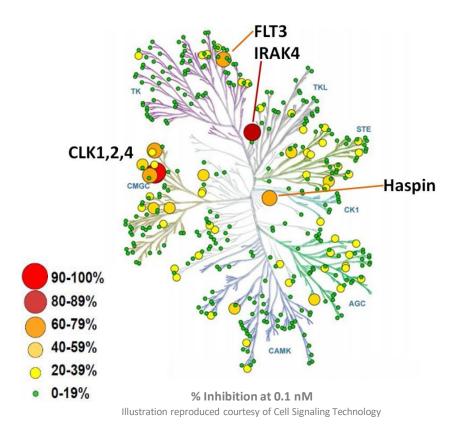
<sup>1)</sup> Guillamot et al. Nat Cell Biol 2019; 2) Smith et al. Nat Cell Biol 2019; 3) American Cancer Society, Cancer Facts & Figures 2020; 4) Leukemia & Lymphoma Society, Facts and Statistics Overview; 5) IMBRUVICA Package Insert. Rev 08/2018



## Emavusertib Unique Molecular Fingerprint

Targeted design specifically engineered to hit key oncogenic targets





# **Emavusertib Binding Affinity**

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

DiscoverX Kinase Panel (378 kinases screened)



NCI Selection for IRAK4

Binds with high affinity to IRAK4

Dual targeting of IRAK4 and FLT3 confers potential efficacy advantage vs. other IRAK4 inhibitors

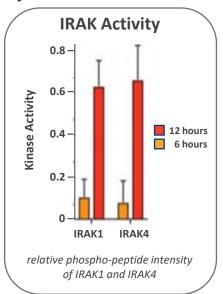


## Additional Competitive Advantage in FLT3 AML

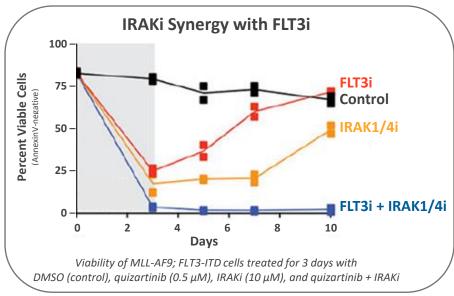
Dual targeting of IRAK4 and FLT3 specifically designed to provide efficacy advantage in AML

"Concomitant targeting of IRAK1 or IRAK4, alongside FLT3, is the most effective means to overcome the adaptive resistance incurred when targeting FLT3"

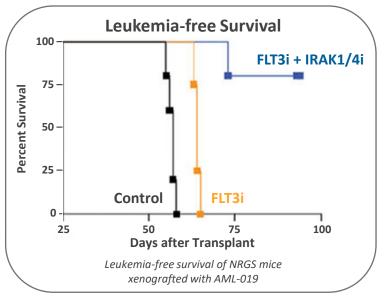
IRAK activity increases after treatment with FLT3i



IRAK/FLT3 combination is synergistically cytotoxic

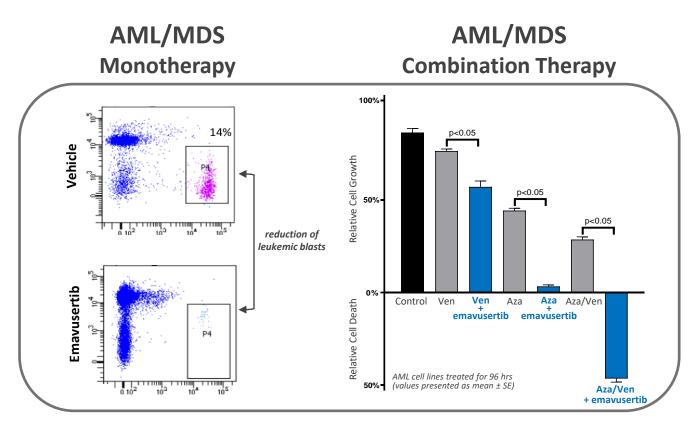


Mice die if treated FLT3i alone, but survive if treated with IRAK/FLT3 combination



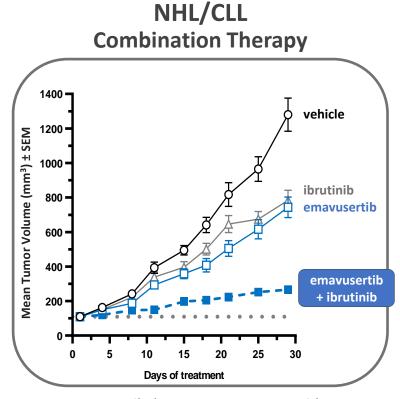
### **Emavusertib Preclinical Data**

Clear anti-cancer activity suggests broad potential across heme malignancies



emavusertib demonstrates monotherapy activity in patient-derived xenografts<sup>1</sup>

emavusertib demonstrates synergy with both azacitidine and venetoclax in THP-1 model<sup>2</sup>

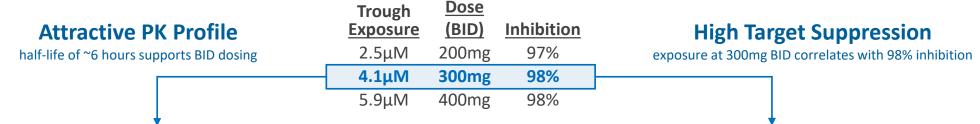


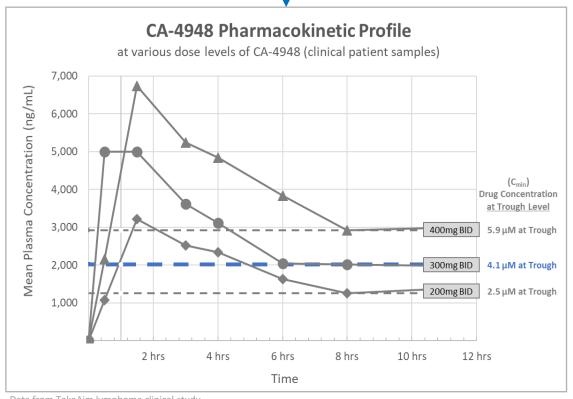
emavusertib demonstrates synergy with ibrutinib in OCI-Lv10 model<sup>3</sup>

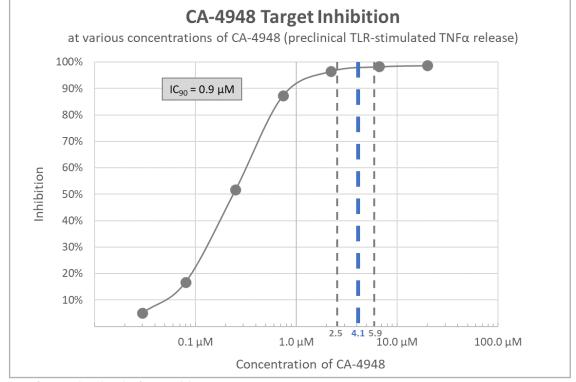


### Emavusertib PK/PD

#### Attractive PK profile supports BID dosing and high target suppression







Data from TakeAim lymphoma clinical study

Data from preclinical study of target inhibition

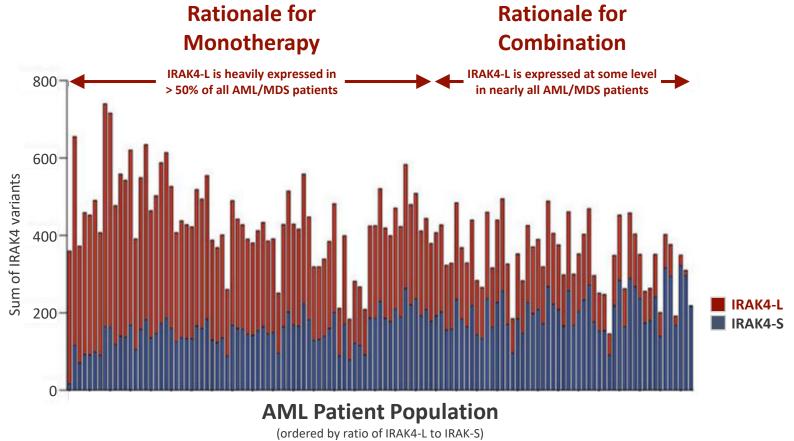


### Emavusertib in AML/MDS

#### Genomic data provide clear rationale for monotherapy & combination

# IRAK4-L is the most prevalent disease driver in AML/MDS<sup>1,2</sup>

<u>Disease Driver</u>	% of Patient <u>Population</u>
IRAK4-L	> 50% <sup>1</sup>
FLT3	> 25% <sup>2</sup>
TET2	10-20% <sup>3</sup>
IDH2	9-13%4
IDH1	6-10%4
CEBPA	~10%3



- ο IRAK4 activation stimulates NF-κB and an array of anti-apoptotic factors
- o emavusertib enhances the anti-cancer efficacy of azacitidine and venetoclax in preclinical models



## **Emavusertib Core Development Strategy**

#### Strategic Priorities

#### Leukemia

#### **Monotherapy**

Possibility for rapid development in a genetically-defined population (patients with FLT3, U2AF1, or SF3B1 mutation)

- R/R AML with FLT3
- o R/R AML with Spliceosome
- o R/R MDS with Spliceosome

possibility for full approval with single-arm study

#### **Combination**

Possibility to expand treatable population to include all patients with AML/MDS, due to frequency of IRAK4-L expression

- R/R AML aza ± emavusertib (or triplet)
- R/R MDS aza ± emavusertib (or triplet)

### Lymphoma

#### Combination

Possibility to gain first lymphoma approval with a defined study in a small patient population with significant unmet need

R/R PCNSL BTKi ± emavusertib

Possibility to expand treatable population to include all patients with NHL/CLL, due to synergy of blocking BCR and TLR pathways

- R/R MZL BTKi ± emavusertib
- R/R MCL BTKi ± emavusertib
- R/R CLL BTKi ± emavusertib
- o etc



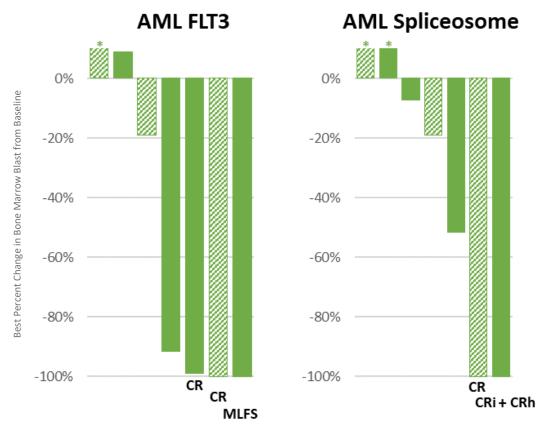




Emavusertib in Leukemia (AML/MDS)

### **Emavusertib Initial Clinical Data**

Potential for clear regulatory path in two target populations: AML FLT3 and AML Spliceosome



<sup>\*</sup> indicates the graphic cutoff as 10%

3 patients have both a FLT3 and spliceosome mutation and are included in both populations

	CR/CRh Rate		
FLT3 AML		2/7	(29%)
Spliceosome AML		2/9	(22%)

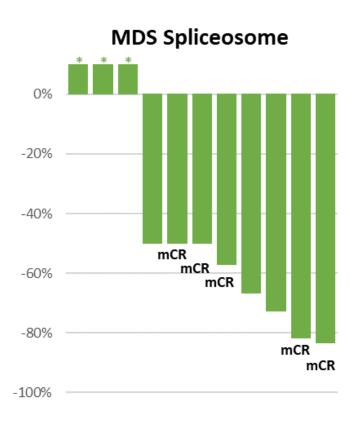
- All patients were R/R, most patients received prior HMA
  - Prognosis for this population is very poor; mOS is 2-4 months<sup>1</sup>
- Full approval has been granted for several AML drugs with a single-arm study using CR/CRh rate as primary endpoint
  - 21% CR/CRh rate for gilteritinib
  - 23% CR/CRh rate for enasidenib

Note: 2 of the 9 AML patients with spliceosome mutation had no post-treatment bone marrow assessment, but reported progressive disease and were considered as response evaluable



### **Emavusertib Initial Clinical Data**

Clear unmet need in R/R hrMDS – there are no approved therapies





- All patients on study were R/R, most patients received prior HMA
  - Prognosis for this population is very poor; mOS is 4-6 months<sup>1</sup>
  - There are no approved therapies for patients who are R/R to HMA
- U2AF1/SF3B1 are the most prominent mutations in MDS
  - 30% of MDS patients have one of these two spliceosome mutations<sup>2</sup>



<sup>\*</sup> indicates the graphic cutoff as 10% data include all response evaluable patients with baseline and post-treatment bone marrow assessments at data cutoff

### Emavusertib in AML/MDS

#### Clinical data support strategy for monotherapy & combination

#### In patients with a targeted mutation:

 Enhanced monotherapy efficacy in a genetically-defined population suggests potential for clear path to first NDA submission

#### In patients without a targeted mutation:

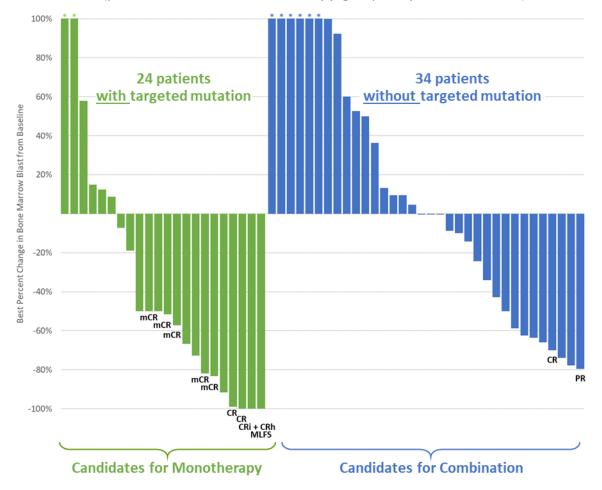
 Initial data show broad anti-cancer activity in non-targeted patients, suggesting emavusertib could enhance the efficacy of other therapies when used in combination

## Spliceosome mutations can be used to identify patients with higher levels of IRAK4-L expression

- U2AF1 and SF3B1 mutations cause overexpression of IRAK4-L<sup>1</sup>
- Genetic screening enables the utilization of existing gene panels to identify candidates for monotherapy

#### **Initial Clinical Data in AML/MDS**

(patients treated with monotherapy, grouped by mutation status)



 <sup>\*</sup> Indicates the graphic cutoff as 100%





<sup>2</sup> additional AML patients with spliceosome mutation had no post-treatment bone marrow assessment, but reported progressive disease and were considered as response evaluable

## Emavusertib in Leukemia (AML/MDS)

The ideal candidate to address the two largest genetically-defined populations in AML/MDS

Emavusertib addresses the two largest genetically-defined populations in AML/MDS:

- 1) IRAK4-L (>50% of population)<sup>1</sup>
- 2) FLT3 (>25% of population)<sup>2</sup>
- Clear single-agent activity
- Genetically-defined patient population
- Dual targeting of IRAK4 and FLT3 may prevent adaptive resistance observed with current FLT3 inhibitors<sup>3</sup>



#### **Next Steps**

#### TakeAim Leukemia Study

- Monotherapy: targeted patients (FLT3 Spliceosome)
- Combination: all other patients







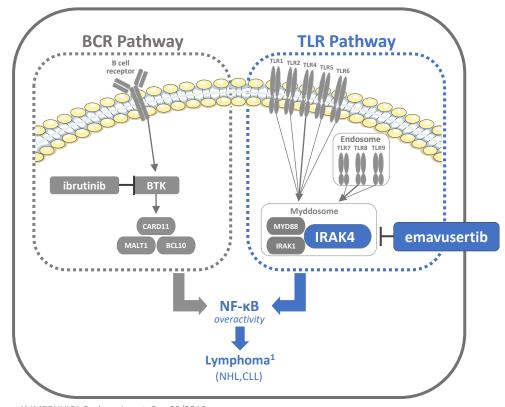
Emavusertib in Lymphoma (NHL/CLL)

### **Emavusertib in B Cell Cancers**

Combination therapy provides complimentary inhibition of two pathways that drive NF-κB

#### NFκB Biology: Two Pathways Drive B Cell Cancers

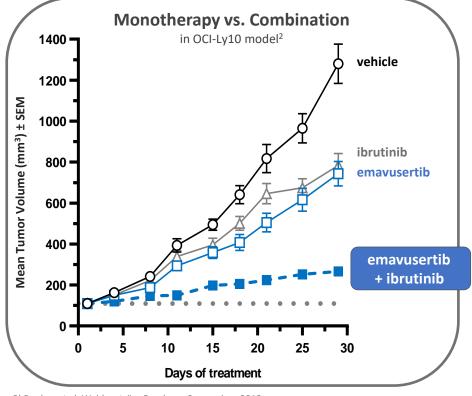
BCR and TLR Pathways independently drive NF-κB overactivity (and NF-κB drives B Cell Cancers)



#### 1) IMBRUVICA Package Insert. Rev 08/2018

#### **Clinical Strategy: Use Combination Therapy**

In preclinical testing, blocking both IRAK4 and BTK drove tumor reduction better than blocking either one alone



2) Booher et al. Waldenström Roadmap Symposium 2019

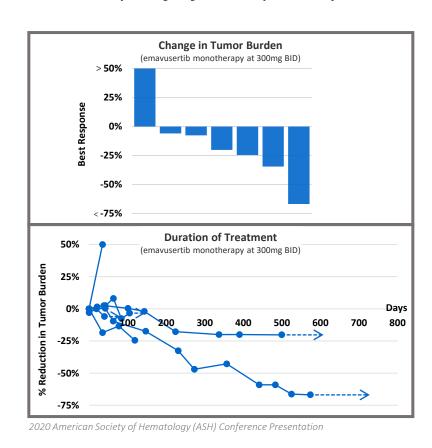


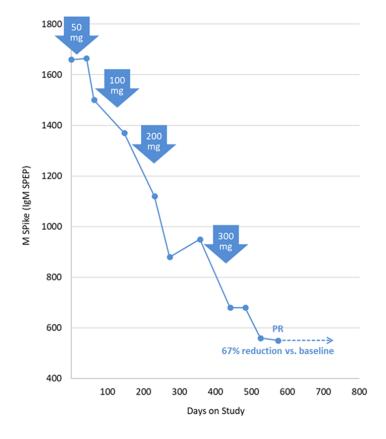
## Emavusertib Initial Clinical Data in Lymphoma – Monotherapy

*Proof-of-Concept demonstrated with monotherapy* 

#### **Monotherapy**

Ph1 proof-of-concept study demonstrated durable tumor reduction in monotherapy

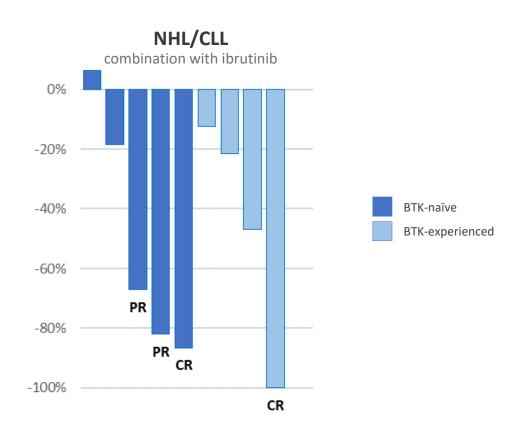






## Emavusertib Initial Clinical Data in Lymphoma – Combination

Majority of patients achieved decreases in tumor burden, including complete responses



response evaluable patients with baseline and post-treatment disease assessment at data cutoff

#### **Change in Tumor Burden Over Time** 100% 80% % Change in Tumor Burden from Baseline 60% 40% 20% 200 250 300 100 -20% -60% -80% -100% **Study Day**



## Emavusertib in Lymphoma

The ideal candidate to combine with BTKi to maximize downregulation of NF-κB

Patients are currently treated with BTKi because it downregulates NF-κB



- Two pathways drive NF-κB:
  - 1) BCR Pathway: addressed by blocking BTK
  - 2) TLR Pathway: addressed by blocking IRAK4
- Initial clinical data suggest blocking both pathways may overcome resistance to ibrutinib

#### **Next Steps**

#### **TakeAim Lymphoma Study**

- Targeted Patients: pCNSL
- All Comers: patients resistant to BTKi



# **Corporate Overview**

## Summary

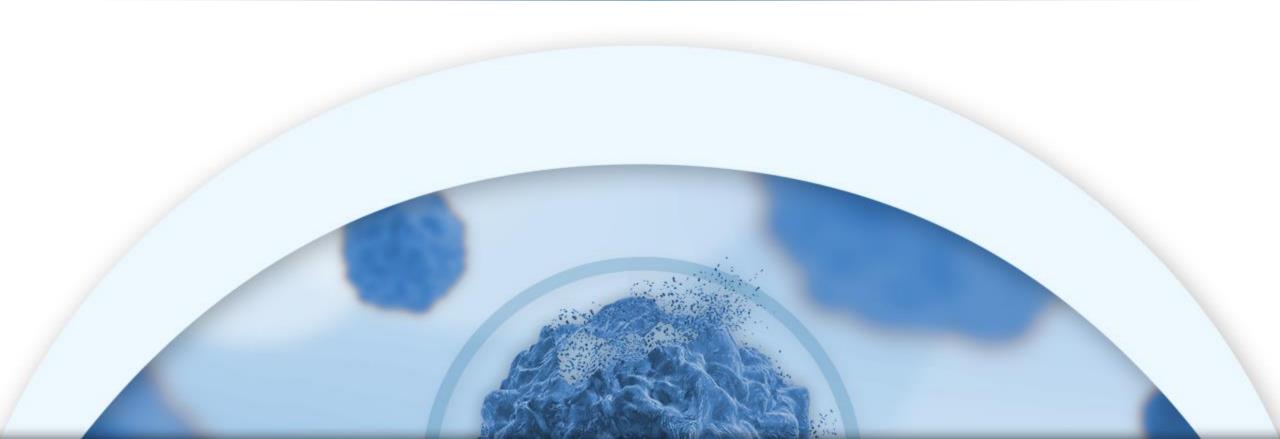
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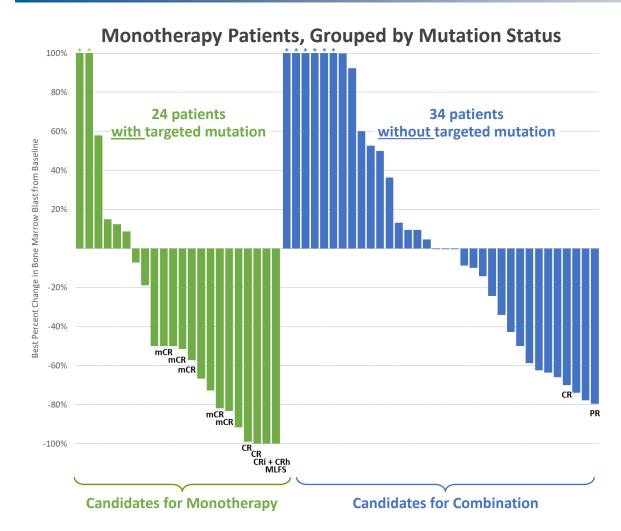
## **End of Corporate Presentation**

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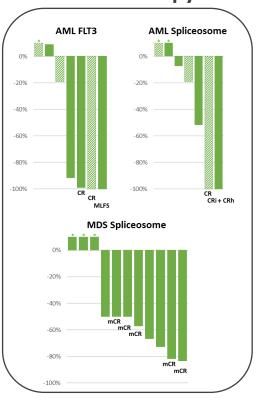
### **Emavusertib Overview**

### Initial clinical data support development strategy for novel IRAK4 inhibitor

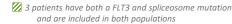


<sup>\*</sup> Indicates the graphic cutoff as 100%

## with targeted mutations Monotherapy

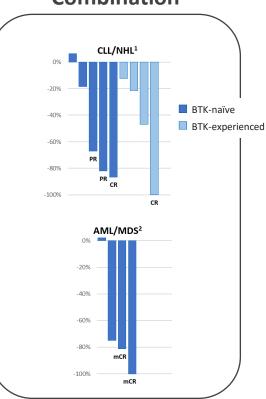






<sup>2</sup> additional AML patients with spliceosome mutation had no post-treatment bone marrow assessment, but reported progressive disease and were considered as response evaluable

## all comers Combination



<sup>&</sup>lt;sup>1</sup> in combination with ibrutinib



<sup>2</sup> additional AML patients with spliceosome mutation had no post-treatment bone marrow assessment, but reported progressive disease and were considered as response evaluable

<sup>&</sup>lt;sup>2</sup> in combination with venetoclax