

Hippo Pathway Relationship to RAS / EGFR Signaling; Therapeutic Potential

Hippo Pathway Targeted Drug Development Summit Jeffrey Ecsedy

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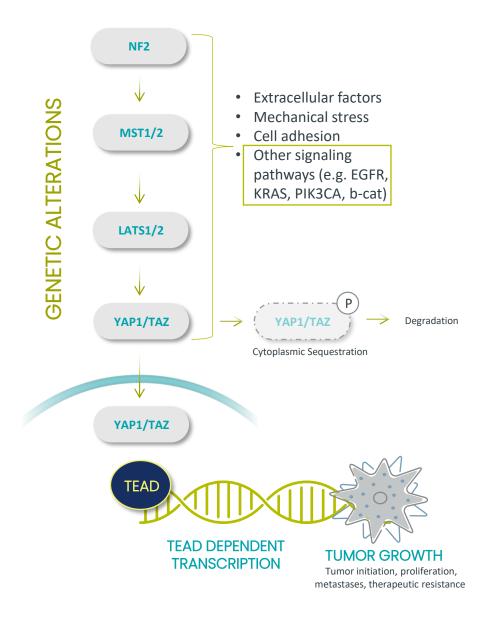
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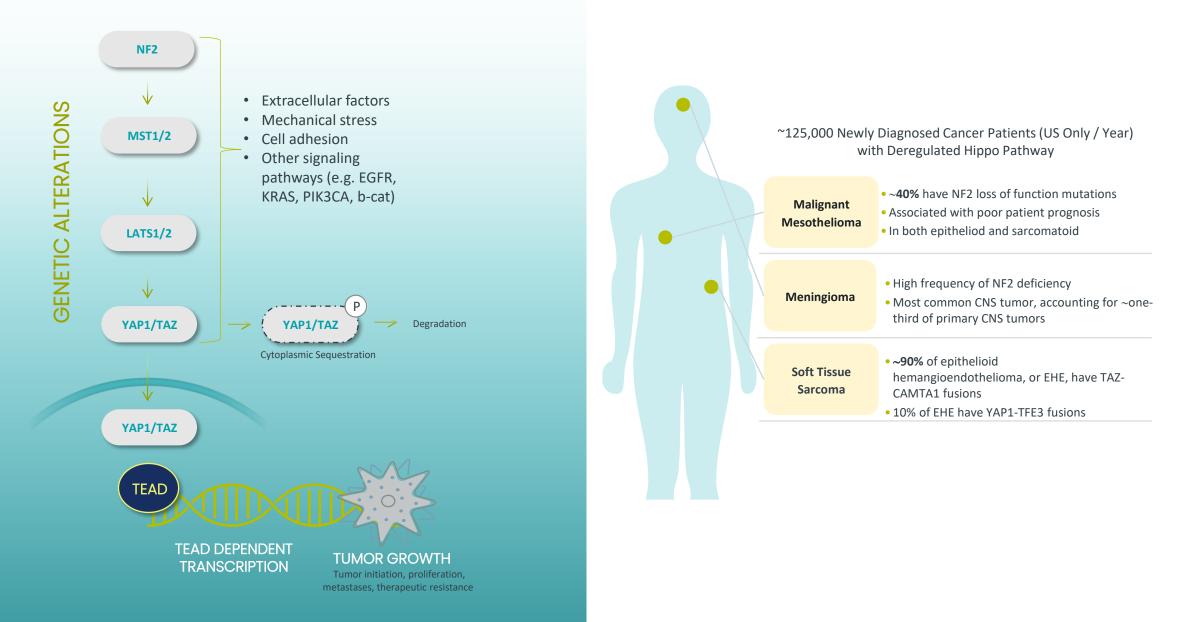
These forward-looking statements are based on the beliefs of our management as well as assumptions made by and information currently available to us. Although we believe the expectations reflected in such forward-looking statements are reasonable, we can give no assurance that such expectations will prove to be correct. If such assumptions do not fully materialize or prove incorrect, the events or circumstances referred to in the forward-looking statements may not occur. We undertake no obligation to update publicly any forward-looking statements for any reason after the date of this presentation to conform these statements to actual results or to changes in our expectations, except as required by law. Accordingly, readers are cautioned not to place undue reliance on these forward-looking statements. Additional risks and uncertainties that could affect our business are included under the caption "Risk Factors" in our most recent report filed with the Securities and Exchange Commission.

Hippo Signal Transduction Pathway in Cancer



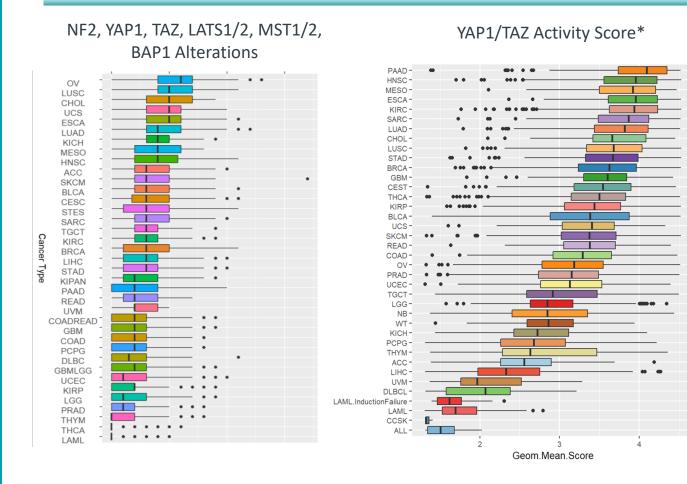
- Multiple activating signals drive YAP/TAZ nuclear localization ---> TEAD binding ---> gene expression of proliferation / pro-survival pathways
- TEAD transcription dysregulated in many cancers
 Numerous tumor suppressor / oncogenes lead to TEAD activation
 Increased nuclear YAP1/TAZ, TEAD activity associated with poor outcome
- Key mechanism of therapeutic resistance

Genetic Alterations in Hippo Pathway Drive Oncogenesis in Patients Across Multiple Indications



Translational Data to Drive Indication Selection

Bioinformatics Analyses

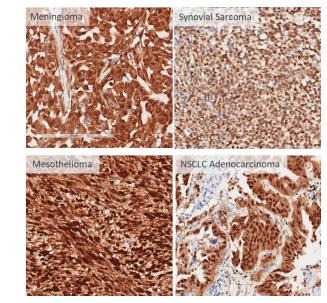


Overlapping Tumors

MESO, HNSCC, CHOL, NSCLC, ESCA, KIRC/CH

*Signature derived from Pham et al 2021

YAP/TAZ Nuclear Localization

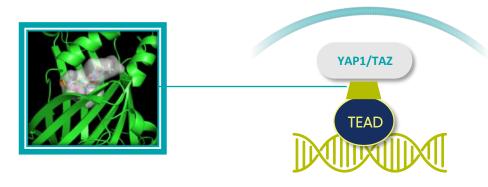


High YAP1 nuclear protein expression indicative of pathway activation in select indications

	%YAP1 +2 +3	%TAZ +2 +3
Meningioma	76	8
Sarcoma	56	11
Mesothelioma	46	19
HNSCC	43	1
Cholangiocarcinoma	31	4
NSCLC	25	10
Pancreas	20	4
Thymoma	10	5
Liver/Hepatocellular	3	1

IK-930 is an Oral, Selective, Potent TEAD Inhibitor

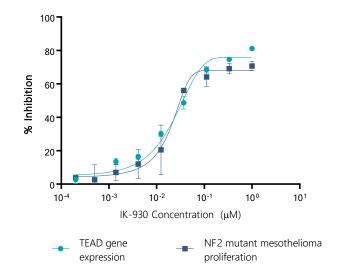
Binding the Central Lipid Pocket of TEAD

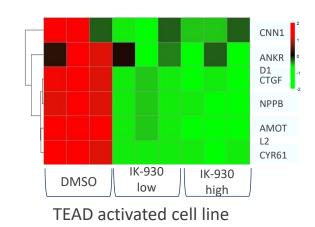


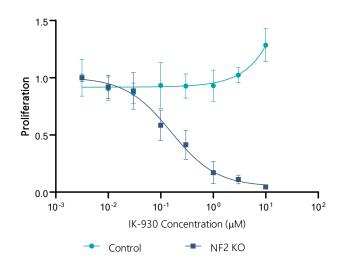
Potent TEAD Inhibition

Robust Inhibition TEAD Target Gene Expression









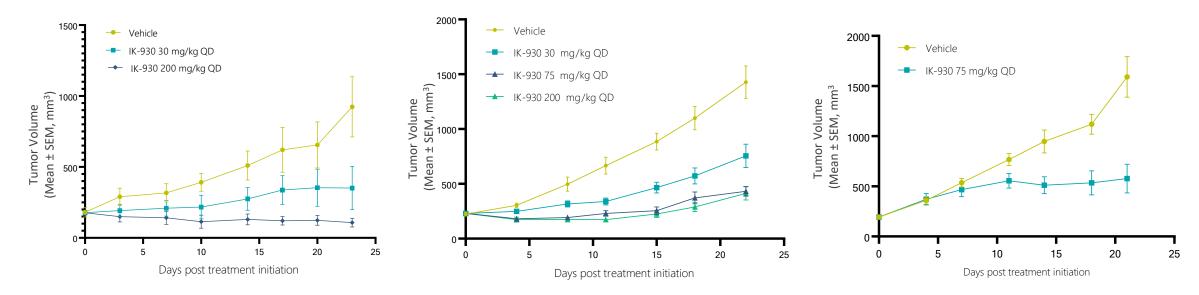
IK-930 Demonstrated Anti-Tumor Activity in Tumor Models with Hippo Pathway Mutations



NF2 Deficient Mesothelioma Model

LATS1/LATS2 Mutated Mesothelioma Model

YAP1 Amplified HNSCC Model



IK-930 is a Well Tolerated TEAD Inhibitor with Favorable ADME/PK Profile

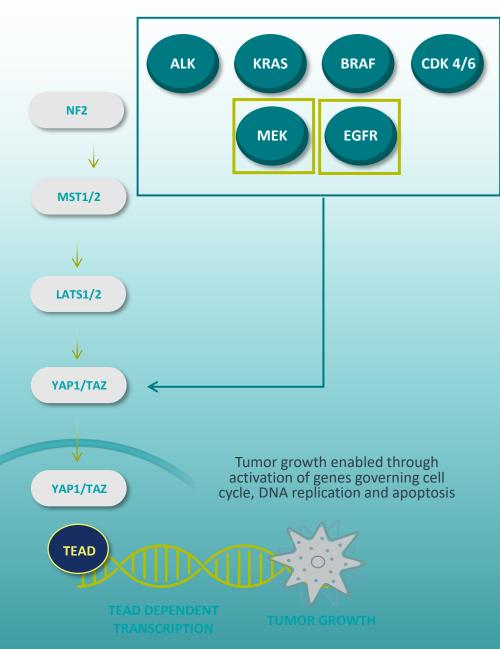
- ✓ Highly selective across a receptor, enzyme, ion channel safety panel (> 50 fold over H226 IC₅₀)
- ✓ Minimal inhibition of hERG in automated patch clamp assay (IC₅₀ > 200 fold over H226 IC₅₀)
- Minimal Cyp inhibition low potential to drug-drug-interactions
- ✓ Not a substrate of P-gp or BCRP transporters
- ✓ Moderate and similar plasma protein binding across species
- ✓ Very good oral bioavailability in mouse, rat, dog, and monkey
- ✓ Brain penetrant

Cyp, hERG and Safety Panel Profiling Suggest Low Risk for Drug-drug Interaction and Off Target Toxicity Concerns

CYP Inhibition, IC ₅₀				
Cyp1A2	>10 uM			
Cyp2B6	>10 uM			
Cyp2C9	>10 uM			
Cyp2C19	7.6 uM			
Cyp2D6	>10 uM			
СурЗА4-М	9.0 uM			
0 0047	>10 uM			
СурЗА4-Т	>10 uivi			
Plasma protei free frac	n binding,			
Plasma protei	n binding,			
Plasma protei free frac	n binding, tion			
Plasma protei free frac Mouse	n binding, tion 2.8%			
Plasma protei free frac Mouse Rat	n binding, tion 2.8% 1.7%			
Plasma protei free frac Mouse Rat Dog	n binding, tion 2.8% 1.7% 2.1%			

Nonclinical PK Summary				
a	T1/2	1.6 h		
Mouse	Vd	2.7 L/kg		
2	Oral bioavailability	55%		
	T1/2	1.7 h		
Rat	Vd	2.8 L/kg		
	Oral bioavailability	56%		
	T1/2	1.8 h		
Dog	Vd	3.1 L/kg		
	Oral bioavailability	52%		
λ	T1/2	2.2 h		
Monkey	Vd	2.8 L/Kg		
Σ	Oral bioavailability	49%		

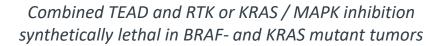
Role of Hippo Pathway in Therapeutic Resistance; Multiple Opportunities for Combination with IK-930

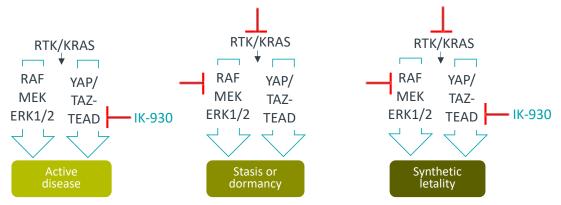


Screens identifying Hippo-mediated resistance					
Cancer	Condition	Hit	Format	Reference	
Melanoma	BRAFi	NF2	CRISPR	Shalem, O. et al. (2014) Science, 343, 84	
Melanoma	BRAFi	EMICERI*	CRISPR	Joung, J. et al. (2017) Nature, 548, 343	
BRAF mut lung	BRAFi	YAP	shRNA	Lin, et al., (2015) Nat Genet, Mar; 47(3): 250	
Kras mut CRC	Kras KD	YAP	cDNA	Shao et al., (2014) Cell, 3;158(1):171	
PDAC	Kras KO	YAP amp	GEMM	Kapoor, A. et al. (2014) Cell, 158, 185	
NSCLC	EGFRi	TEAD Gene signature	RNASeq	Kurppa, K et al. (2020) Cell, 37 (104-22)	
NSCLC	EGFRi	NF2	CRISPR	Zeng, H. et al (2019 Elife, 8:e50223	

Screens identifying Hippo-mediated resistance

* EMICERI : Increase MOB3B (component of MST1/2 and LATS1/2 inhibitor complex) expression

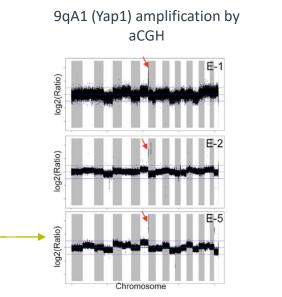




Adapted from Lin, et al., (2015) Nat Genet, Mar; 47(3): 250

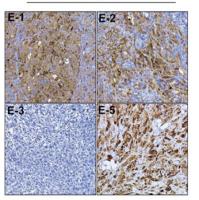
Role of YAP1/TEAD in KRAS Mutated Pancreatic Cancer

- Yap required for progression to invasive PDAC in mutant Kras or Kras:Trp53 mice (Zhang Sci Signal, 2014)
- YAP1/TEAD mediates tumor recurrence in PDAC GEMM upon KRAS^{G12D} withdrawal (Kapoor et al Cancer Cell 2014)
 - KRAS^{G12D} extinction induced rapid tumor regression
 - Subset of relapsed tumors lacking KRAS^{G12D} mediated by YAP1/TEAD
- YAP1 activation mediates pancreatic cancer progression upon wild-type KRAS allele loss (Yan et al Oncogene 2021)

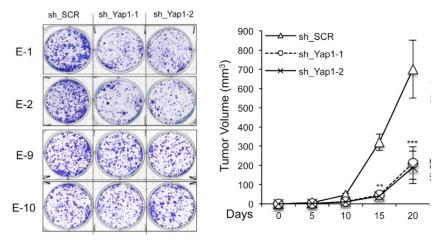






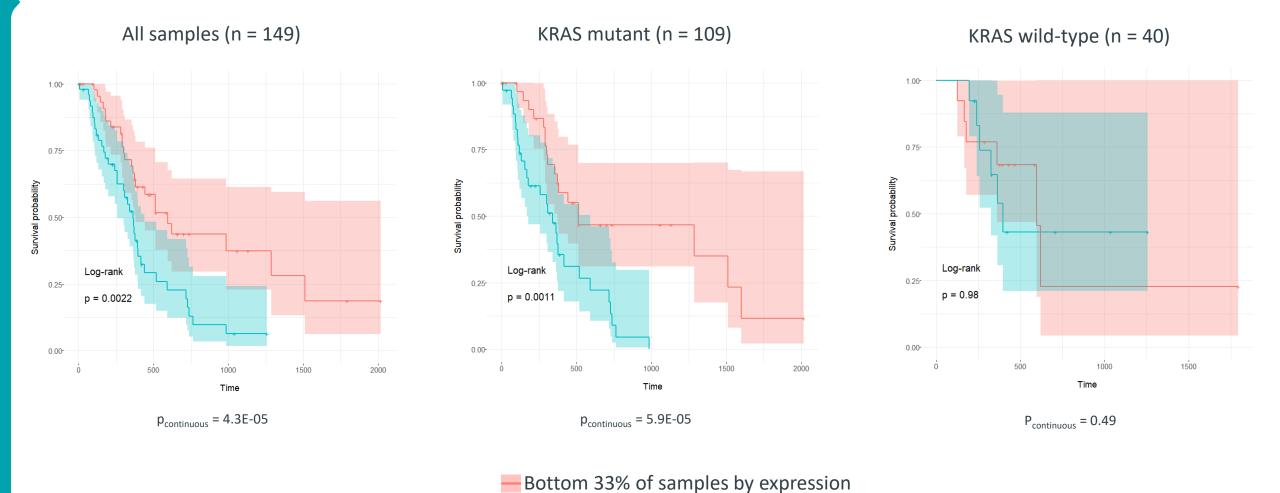


YAP1 dependent tumor growth in KRAS independent tumors (E-1, E-2)



Kapoor et al Cancer Cell 2014 (DOI: 10.1016/j.cell.2014.06.003)

Hippo/YAP1 Transcriptional Signature Associated with Poor Outcome in KRAS Mutated Pancreatic Cancer



— Top 33% of samples by expression

Signature also associated with decreased OS in KRAS mutated NSCLC

Hippo signature derived from Pham et al Can Dis 2021

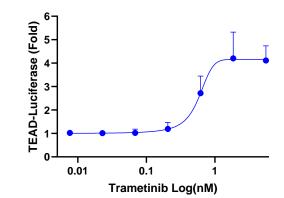
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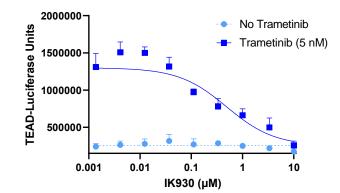
MEK Inhibitor Induces YAP1 Nuclear Localization and TEAD Dependent Transcription

HCT116 cells (KRAS G13D)









Tametinibulic Lanctinit (5 nM) (2 nM) (6 nM) (A nM)

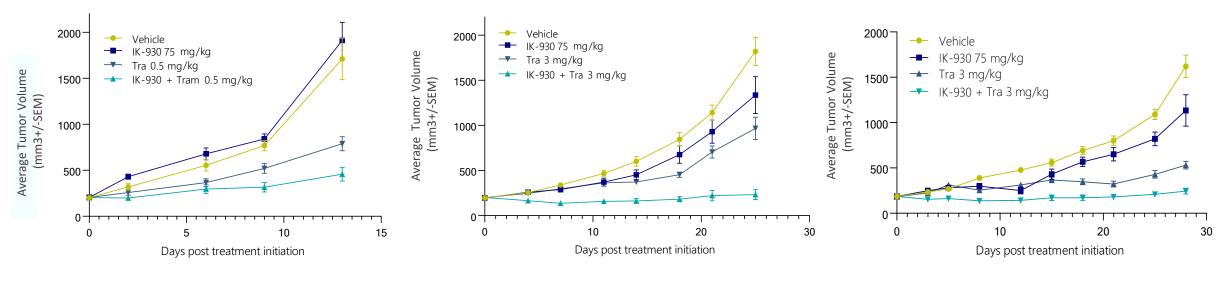
HCT116 cells (KRAS G13D)

IK-930 Enhances Apoptosis in Trametinib-Treated KRAS Mutant Cells

LOVO: Human KRAS G13D CRC HCT116: KRAS G13D CRC 1.5×10-3 Normalized Apoptosis Green Object Count Per Well / Phase Area Per Well / Im³/Well) 6×104-Tra 30nM+930 1 µM Normalized Apoptosis Green Object Count Per Well / Phase Area Confluence (1/Well / %) ra 30nM+930 0.3 µM ra 30nM+930 0.1 µM Tra 30nM+930 0.3 µM 1×10-3-Tra 30nM+930 0.1 µM 4×104-Tra 30 nM --5×10-4-Tra 30 nM • 2×104 930 1 μM DMSO 930 1 µM DMSO 0 20 Λ 40 60 20 60 40 80 Hours Hours A549: KRAS G12S NSCLC Calu-1: KRAS G12C NSCLC 0.025-5×103--**■**- Tra 100nM+930 1 µM Normalized Apoptosis Green Object Count Per Image / Green Object Average (1/Image / µm²) ase Area Object Average (1/Image / µm²) - 0000 - Tra 100nM + 930 1 uM Normalized Apoptosis Green Object Count Per Well / bhase Area Confluence (1/Well / & 4×10³ Tra 100nM + 930 0.1 uM Tra 100nM+930 0.1 µM 3×103 Tra 100 nM Tra 100nM -0-2×10 Green Obj Phase Area 🕂 930 1 µM 0.005 1×10 DMSO -930 1 uM DMSO 0 -0.000 20 0 40 60 60 80 100 120 140 0 20 40 Hours Hours

Increased Anti-Tumor Effect of IK-930 in Combination with MEK Inhibitor in KRAS Mutant Tumors In Vivo





A549: KRAS G12S NSCLC

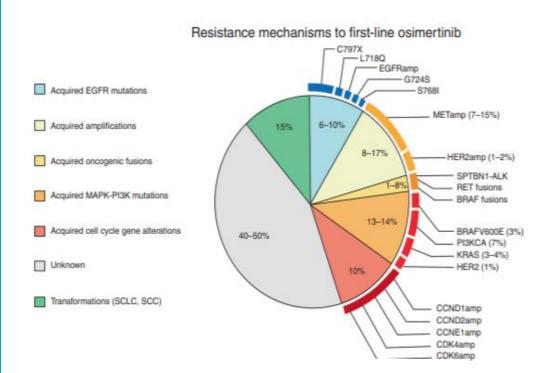
HCT116: KRAS G13D CRC

LOVO: Human KRAS G13D CRC

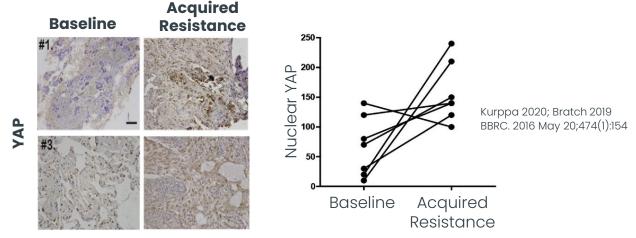
Model	HCT116	A549	Lovo
In vivo TGI Combination	83% (1mg/kg MEKi)	78% (0.5mg/kg MEKi)	75% (1mg/kg MEKi)

IK-930 Opportunity to Address Emerging Resistance Associated With Early use of Osimertinib in EGFRm NSCLC

Resistance mechanisms to osimertinib in EGFR-mutated non-small cell... A Leonetti et al.

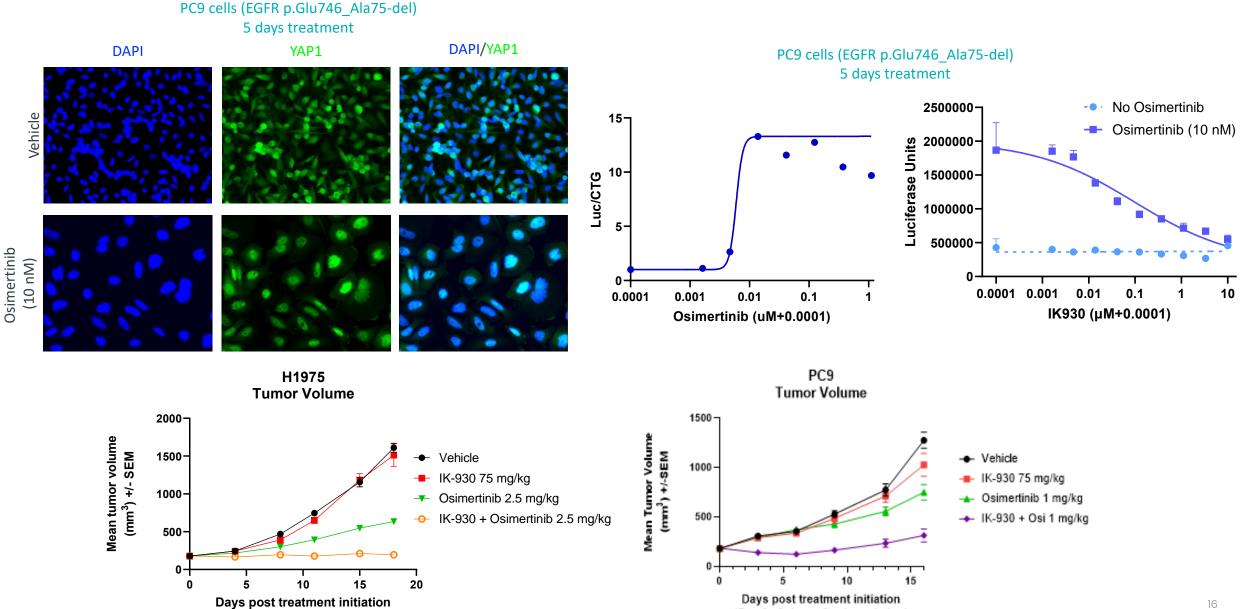


- Success of osimertinib (3rd Gen EGFRi) as 1L and adjuvant tx of EGFRm NSCLC
- Resistance is related to acquired EGFRm, MET amplifications, oncogenic fusions, histological transformation, BUT...
- At least 40% of these cases do not have an identified mechanism of resistance and represents a significant unmet medical need
- Literature suggests that Hippo/Yap1 activation associated with acquired resistance



- Opportunity for IK-930 based combinations to address acquired Osimertinib resistance (post 1L /adjuvant)
- Opportunity to identify subset of patients in whom addition of IK930 combo can delay/prevent the emergence of resistance (1L)

EGFR Inhibitor Induces YAP1 Nuclear Localization and TEAD Dependent Transcription



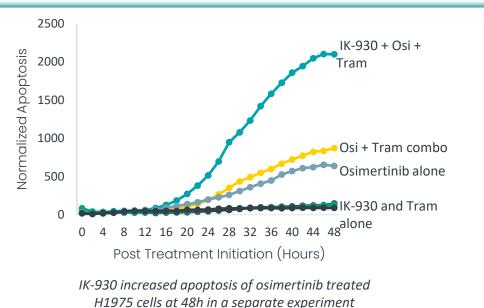
(Day 0= 8 days after inoculation)

Anti-Tumor Activity of IK-930 Combined with EGFR and MEK Inhibitors

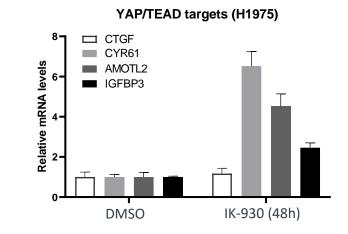
YAP1 nuclear localization in EGFR mutant H1975DMSOOsi + TramOsi + TramImage: Colspan="2">Image: Colspan="2" Image: Colspa

IF: α-YAP1

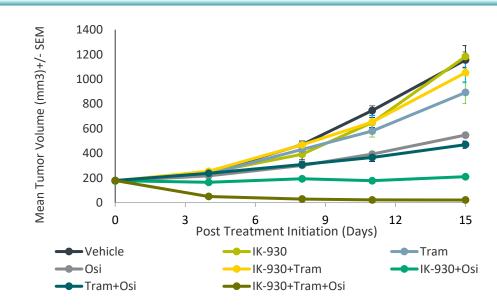
IK-930 enhances apoptosis in vitro in EGFR mutant H1975



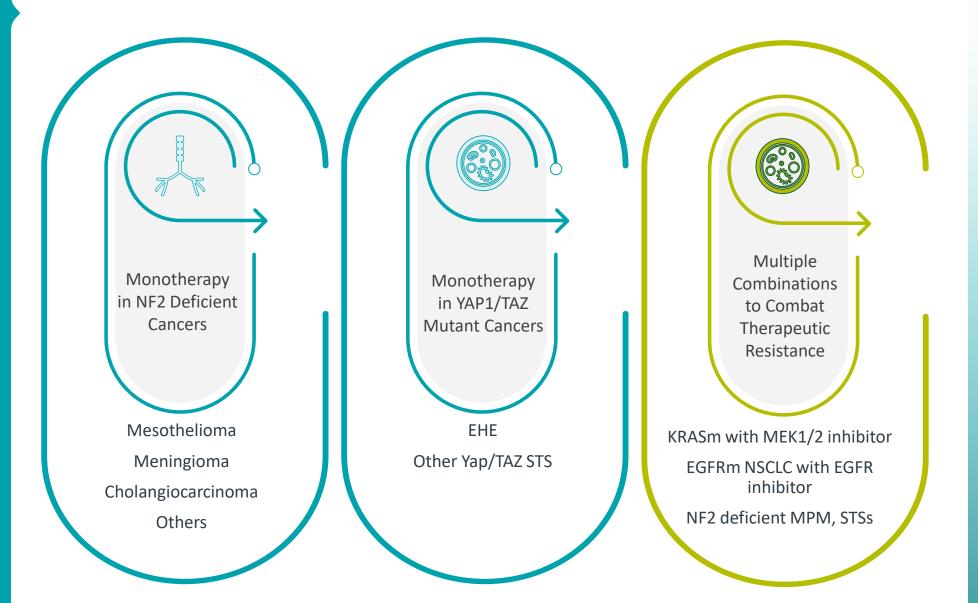
TEAD-dependent genes induced by EGFRi + MEKi in H1975



IK-930 synergy with EGFRi and MEKi in vivo



Developing First-in-Class TEAD Inhibitor for Genetically Altered Cancers and Therapeutic Resistance



Monotherapy strategy focused on NF2- deficient orphan indications including NF2 deficient MPM, EHE and other solid tumors with prevalent YAP/TAZ fusion genes

Combination strategy to explore multiple with targeted agent combos to reverse mechanism of resistance in broader indications

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