Preclinical studies characterize tumor type sensitivity to FASN inhibition and the mechanism and efficacy of novel drug combinations with TVB-2640

Timothy S. Heuer, Richard Ventura, Julie Lai, Joanna Waszczuk, Kasia Mordec, Claudia Rubio, Glenn Hammonds, Marie O' Farrell, Douglas Buckley and George Kemble 3-V Biosciences, Menlo Park, CA

#4743

Introduction

- 3-V Biosciences' lead, oral FASN inhibitor is in Phase I clinical development for the treatment of solid tumors
- Fatty acid synthase (FASN) catalyzes the synthesis of palmitate from acetyl-CoA, malonyl-CoA, and NADPH
- Tumor cells have an increased dependence on FASN-synthesized palmitate compared to non-tumor cells
- FASN Inhibition decreases cellular levels of palmitate and saturated fatty acids and induces tumor cell apoptosis
- In vitro and in vivo studies have identified multiple mechanisms of action for FASN inhibition efficacy that include: (1) Membrane and lipid raft architecture remodeling; (2) Tumor cell gene expression reprogramming; (3) Signal transduction pathway inhibition
- Studies to understand the mechanisms of action and biological consequences of FASN inhibition are guiding the discovery of tumors highly dependent on FASN and biomarkers for assessment of pharmacodynamic activity and patient selection

Results

Classification of Tumor Lipogenic and Epithelial Gene Expression Aligns with FASN-Inhibitor Sensitivity In Vitro

- Published literature reports gene expression signatures of lipogenic, glycolytic, epithelial, and mesenchymal phenotypes
 - Lipogenic and epithelial phenotypes co-segregate
 - Glycolytic and mesenchymal phenotypes co-segregate
- Application to 3-V In vitro data shows association with FASN inhibitor sensitivity
- Application to TCGA tumor data (8588 samples) to identify candidate tumor types and biomarkers of FASN inhibitor sensitivity
 - Stratification and analysis of molecular genetic features of tumors

References

1. Collisson et al. Nat Med. 2011 Apr;17(4):500-3. Epub 2011 Apr 3. PMID: 21460848

2. Daemen et al. Proc Natl Acad Sci U S A. 2015 Aug 11;112(32):E4410-7. Epub 2015 Jul 27. PMID: 26216984



Figure 1. Tumor cell line classification by gene expression analysis. Methods described in the Collisson et al and Daemen et al reports were followed using in vitro viability data (cell titer glo) with TVB-3166. Gene expression data were from the CCLE (Broad Institute, Cambridge, MA).



Figure 2. Classification of 8558 TCGA tumor sample data using the Daemen et al 2015 methods. Data were downloaded, processed, and analyzed at 3-V Biosciences. In figure 3 below, tumor cell line xenografts representing (1) liver tumors (LIHC) and (2) lung adenocarcinoma tumors (LUAD) were assessed for tumor growth inhibition efficacy using TVB-3166.

Tumor Cell Line Xenograft Sensitivity Aligns with TCGA Tumor Classification





Figure 3. In vivo mouse tumor xenograft efficacy analysis of the HEPG2 liver tumor and A549 NSCLC tumor mode Liver tumors classify as the most highly lipogenic by gene expression analysis (GEA); lung adenocarcinomas are moderately lipogenic by GEA. HEPG2 and A549 tumor cell lines classify as undetermined; thus, not as highly lipogenic noma tumor cell lines. In-life analysis of tumor growth efficacy was conducted by as other liver or lung adenocarcir Crown Biosciences (Santa Clara, CA)

KRAS-Mutant NSCLC Tumors are More Lipogenic

- Normalized gene expression scores
- W-Score: Weighted sum of 4 gene sets plus VIM and FASN. Weights determined by multi-variable linear least squares regression for best match to FASNi sensitivity.
- **VIM:** Vimentin, high mesenchymal expression, low epithelial expression.
- Alpha: FASN sensitivity-derived gene set. Difference between mean z-score of 17 positive correlation genes and mean z-score of 20 negative correlation genes.
- MES: PDAC-derived mesenchymal/epithelial gene set. Difference between mean z-score of 20 mesenchymal genes and mean z-score of 22 epithelial genes.
- **FASN**: TVB-2640 target. Synthesizes palmitate that provides a substrate for complex lipids that function in cell signaling and metabolism.

Cell Lines



Figure 4. Cell Titer Glo in vitro assay was used to asses cell viability in response to treatment with the FASN inhibitor TVB-3166. Cell were treated for 7 days with TVB-3166.



KRAS G12 Mutation Increases NSCLC Lipogenic Gene Expression

Gene or Gene Set

RAS Mutation Associated with FASN Sensitivity in NSCLC

G12 Mutant

Wild Type

FASN Inhibition Blocks RAS-Associated Signaling Pathways Raft Membrane Microdomaiı



Ventura et al. EBioMedicine. 2015 Jul 2;2(8):806-22 PMID: 26425687; 3-V Biosciences.

TVB-3664 Inhibits Increased Palmitoylation of RAS-Associated Signaling Proteins in KRAS-Mutant NSCLC Lines

KRas genotype NSCLC Cell Line	G1 A5	2S 49	G12 A42	D 27	G12C H23	Q6 CAL	1K U-6	W H5	Т 20	W H13	T 895		WТ H52
		2-E	Bromo	o-Paln	nitate (20 μN	1)						TVB
2-BP or TVB-3664	-	+	-	+	-	-	+	-	+	-	+	-	
Palmitoyl-EGFR		I	-	percent	-	-	-			pieros .	Annual		
Palmitoyl-Akt	1	-	-	-	-	-	-		-			=	
Palmitoyl-α-tubulin	I	I	I	1	-	-	-	-	-			I	-
Palmitoyl-β-tubulin			-	-	-								0.000
Palmitoyl-NRas		-	-	_	-	- No. of Concession, Name		-electricity	and the second	(antopican)	terret open	1	-
Palmitoyl-Cav2	-	1	1	-	-	-	1			-	_		

Figure 5. In vitro acyl-biotinyl exchange and Western blot analysis of protein palmitoylation and inhibition by treatment with 2-bromopalmitate or the FASN inhibitor TVB-3664. TVB-3664 treatment for 72 hours. 2-BP treatment for 16 hours.

Oncogenic KRAS Reprograms Tumor Cell Metabolism

Increased Glutamine metabolism

Reductive glutamine metabolism can promote lipid synthesis via pyruvate and NADPH production Increased Glucose metabolism

- Glycolytic pyruvate anabolism can promote lipid synthesis
- Increased non-oxidative PPP biosynthesis and NADPH production



Figure 6. Activated KRAS reprograms tumor cell metabolism to increase utilization of glutamine and glucose. One of the consequences is high levels of pyruvate production. Son et al. 2013 Nature; Ying et al. 2012 Cell. A possible fate of pyruvate is metabolism to citrate and acetyl CoA, the substrate for palmitate synthesis by FASN.

Glutamine Sensitizes KRAS-Mutant Tumor Cells to FASN Inhibition



Figure 7. CALU-6 NSCLC cells respond to FASN inhibition in the presence of 1% glutamine but not glutamine-free media. Similar results observed in PANC-1 tumor cells. CASP1 and ELOVL6 gene expression by qPCR are representative of more global gene expression effects.

3-V BIOSCIENCES

G12S
A549





Combined FASN Inhibition with PD1 Immunotherapy

Figure 8. FASN inhibition sensitizes the LLC/LL NSCLC mouse tumor model to anti-PD immunotherapy. In vitro, FASN inhibition induces PD-L1 expression in NSCLC tumor cell lines

Combined FASN and Angiogenesis Inhibition Enhances Efficacy





Figure 9. TVB-3664 combined with bevacizumab shows increased tumor growth inhibition compared to single agent treatment. In vitro assays measuring endothelial cell tube formation (Incucyte analysis, Essen biosciences) shows that media conditioned with COLO-205 cells treated with TVB-3644 inhibits tube length in combination with axitinib better that axitinib alone or COLO-205 cells treated with vehicle (DMSO).

TVB-2640 Biomarker Analysis in CLIN-002 Tumor RNA



Tumor mRNA expression of genes in the Wnt/b-catenin, mitotic cell cycle, VEGFR and Ras signaling pathways segment responding and non-responding patients treated with TVB-2640

Further investigation in larger set of patients

Avasdi (Menlo Park, CA) Core **Topological Data Analysis**

Conclusions

- Lipogenic gene expression signatures classify FASNi sensitivity
- Lipogenic gene expression is increased in Kras-mutant NSCL tumors; Krasmutant NSCLC cell lines have increased FASNi sensitivity
- Palmitoylation of Ras-associated signaling proteins is increased in Krasmutant NSCLC cell lines and is inhibited by FASN inhibition with TVB-3664
- FASN inhibition combines with PD1 immunotherapy or VEGF inhibition (bevacizumab) to increase significantly in vivo tumor xenograft efficacy
- TVB-2640 CLIN-002 tumor mRNA shows b-catenin, mitotic cell cycle, VEGFR and Ras pathway dysregulation in responsive, not unresponsive, patients