

# AR-mTOR-26 – A Potent, Selective mTORC 1/2 Kinase Inhibitor for the Treatment of Malignancy Stefan D. Gross, Rui Xu, Mark Boys, Michael Burkard, Lisa De Meese, Walt E. DeWolf, John Fischer, Susan Gloor, Michael J. Humphries, Kelly Regal, Brad Fell, Kevin Condroski, Greg Miknis, Kevin Rasor, Mareli Rodriguez, Georg Topalov,

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### Introduction

The PI3K/Akt pathway is constitutively activated in a large proportion of human cancers. mTOR kinase plays a vital role in this pathway as the key component of two independent signaling complexes (mTORC1 and mTORC2). Inhibition of mTOR kinase could therefore serve as an effective means of abrogating PI3K-dependent signaling. In addition, the activity of TORC1 is often aberrantly activated in a PI3Kindependent manner enabling tumor cells to survive and proliferate despite the many negative influences of the tumor microenvironment. Therefore, targeting the activity of TORC1 and TORC2 would abrogate both PI3K signaling and the cancer cells' ability to survive in the harsh environment of the tumor. We report here the development of a potent and selective small molecule inhibitor of mTOR kinase designated AR-mTOR-26

On enzyme this compound exhibits low single digit nM activity against isolated mTOR kinase with >40-fold selectivity against the  $\alpha$ ,  $\beta$  and  $\delta$  isoforms of PI3K as well as a panel of over 250 ser/thr and tyrosine kinases.

 On cells AR-mTOR-26 potently inhibits TORC1/2-dependent readouts pAKT(S473). p4E-BP1(S37/46) and pS6(S235/6) yet is significantly inactive against the PI3Kdependent readout pAKT(T308) confirming its selectivity against Class I PI3K-kinases.

This cellular potency readily translates into broad anti-proliferative activity against a wide array of solid tumor and hematological cancer cell lines irrespective of mutational status (i.e. KRAS, PTEN, PIK3CA, etc.)

 AR-mTOR-26 possesses exceptional pharmacokinetic properties across multiple species, including mouse with exposures predicted to be biologically active.

The potential for in vivo activity was confirmed in two tumor xenograft models both in terms of tumor growth inhibition as well as inhibition of mTOR-relevant targets in the tumors

## **AR-mTOR-26 In Vitro Activity and Safety Profile**

Enzyme	AR-mTOR-26 IC <sub>50</sub> (nM)	Selectivity Data 261 Kinase Panel	AR-mTOR-26
mTOR	<0.5	>50% inhibition at 1 uM	ΜΚΚ7β (49)
p110α	21	58 Channel / Receptor Panel	
р110β	240	>50% inhibition at 10 uM (% Control in binding assay)	Adenosine A3 (80) [>100 uM]
p110γ	6	[Functional assay IC <sub>50</sub> ]	
p110δ	176	hERG channel (% inhibition)	14% at 1 uM 59% at 10 uM

•AR-mTOR-26 is potent inhibitor of mTOR enzyme that is >40-fold selective for the  $\alpha$ ,  $\beta$  and  $\delta$  isoforms of the Class I PI3Ks and substantially selective vs. 261 other protein kinases.

•AR-mTOR-26 exhibited minimal effects on a panel of 58 receptors and was only a modest inhibitor of the hERG cardiac channel.

## Preclinical Pharmacokinetics<sup>1</sup>

Species	Mouse		F	Rat	Dog		
Route	IV PO		IV	PO	IV	PO	
Dose (mg/kg)	2	10	2	10	2	10	
CL (ml/min/kg)	8.8		4.7		14		
Vss (l/kg)	3.2		4.0		8.0		
t <sub>1/2</sub> (h)	3.9		N/A		9.8		
AUC (mg/h/ml)	3.7	6.8	7.2	14.2	2.5	9.1	
Cmax (mg/ml)		0.9		0.5		0.4	
%F		37		39		75	

 Based upon its pharmacokinetic properties, both AR-mTOR-26 is predicted to produce biologically active exposures via PO dosing in mouse as well as other species.

· · · · ·	Compound	MUA	pso	р4Е-ВРТ	ракт-5473	PAKt-1308	
A	R-mTOR-26	mTOR kinase	11	89	15	427	
	AR472303	mTOR kinase	37	215	44	124	
	AR454415	mTOR/PI3K	7	83	20	69	
	AR453517	mTOR/PI3K	142	485	65	172	
	AR466979	ΡΙ3Κα/δ	560	2666	186	251	
	Conc. (	PI-103	00 333 111 37	AR-mTOR-2	6 4 0.5 0.2 0.0	Control 5 0.02 0	
	p30 (3	37/46)					
	pAKT (	(T308)					
	pAKT (	(\$473)					
	tı	ubulin					

Cellular pathway effects of AR-mTOR-26

In-cell western analysis of pathway inhibition [IC<sub>50</sub> (nM)]

 In cells, AR-mTOR-26 is a potent inhibitor of TORC1 and TORC2dependent readouts (pS6, p4E-BP1 and pAKT-S473)

•The compound also effects a substantial but incomplete loss of pAKT(T308) phosphorylation in marked contrast to PI3K inhibitor AR00466979 which is equipotent on both readouts

Methods. LNCaP cells were used for in-cell western analysis employing phospho-specific antibodies to the indicated TORC1, TORC2 or PI3K-dependent substrates. PC3 cells were used for the immunoblotting analysis employing the same antibodies

## mTOR vs. PI3K selectivity in cells



 AR-mTOR-26 potently inhibits the TORC2 readout pAKT-S473 but is substantially inactive against the PI3K-dependent readout pAKT-T308.

 Over the dose range analyzed, data indicate that AR-mTOR-26 actually activates the PI3K pathway likely due to abrogation of a downstream negative feedback loop.

•This profile is in marked contrast to PI3K inhibitor AR00466979 which potently inhibits both readouts.

Methods. To measure the cellular activities of PI3K and TORC2 on AKT independently, DNA constructs expressing mutant forms of AKT substituted with glutamate at one of the regulatory phosphorylation sites T308 or S473 were generated. Expression constructs were then transfected into HEK-293T cells, treated with compound and processed for in-cell western analysis probing for either pAKT(S473) or pAKT(T308). Upper panel depicts typical dose/response curves for AR-mTOR-26 and AR00466979. Lower panel is a table of representative IC<sub>50</sub> values obtained for the indicated compounds in the pAKT T308E and pAKT S473E cell

Solid Tu	imor l	_ines		G	M)						
Cell Line:	BxPC -3	DU145	Malme -3M	OVCAR -8	PC3	LNCaP	DLD -1	BT-474	22RV1	H460	HCT116
Mutation:	WT	LKB1	BRAF VODE	ERBB2 GTTEV	PTEN -	PTEN -	KRAS G13D	PI3KCA KIIIN	PI3KCA Ester	PI3KCA ES45K KRAS GETH	PI3KCA HID47R KRAS G12D
AR-mTOR-26	34	46	34	76	21	17	38	100	37	34	40
AR454415	54	81	60	51	36	21	17	85	34	44	102
AR453517	1942	233	109	305	115	295	134	596	402	366	324
AR466979	316	710	354	1685	360	415	458	312	761	604	767

Cell Line:	K562	GDM -1	THP1	HEL	MOLT4	MO7e	MV4-11	MOLM13	MOLT3	HL -60
Mutation:	BCR - Abl	NRAS	BRAF VSDOE	ERBB2 GTTEV	PTEN -	PTEN -	KRAS G13D	PI3KCA KIIIN	PI3KCA ESACR	PI3KCA Est KRAS GET
AR-mTOR- 26	41	76	69	41	17	31	13	14	20	90
AR454415	51	231	154	76	57	59	52	24	55	248
AR453517	1068	620	492	1319	151	728	295	143	158	164
AR466979	6789	1030	778	5647	207	3820	349	344	271	169

Across both solid tumor and leukemic cell lines, the mTOR-selective

status of the PI3K or RAS/RAF/Mek/Erk pathways

Methods. Indicated cell lines were plated out at a density of 5-10K cells/well of a 96-well plate, treated with compound for 3-5 days and then assessed for viability/proliferation using Celltiterblue (Promega).



Methods. Cell cycle analysis: NCI-H460 (NSCLC) and HEL 92.1.7 (Leukemia) cells were treated for 24h with either 170 nM (H460) or 200 nM (HEL) AR-mTOR-26 for 24h and assessed for cell cycle distribution by propidium iodide staining and subsequent FACS analysis [*left panel*]. Apoptosis assays: HEL and H460 cells vere treated with compound (same concentrations as for cell cycle) for 24-72h and stained using Annexin V or monitored for caspase 3/7 activation (CaspaseGlo 3/7, Promega) [right panel]

Inhibition of TORC1 and TORC2 signaling in vivo

<sup>1</sup>All in vivo studies were performed in accordance with IACUC guidelines and in harmony with the Guide for Laboratory Animal Care and Use