Targeting MET in preclinical models to support the clinical development of AZD6094/HMPL-504 (Volitinib) in NSCLC

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Introduction:

- Met is a receptor tyrosine kinase that is deregulated (amplification mutation and over-expression) across multiple cancer types
- Aberrant HGF/Met pathway activation leads to uncontrolled tumor cell growth, invasion and survival

AZ is partnering with Hutchison MediPharma in the development of AZD6094/HMPL-504 (Volitinib) a potent (IC50 4 nM) and selective (>650 fold selectivity over 265 kinases) small molecule inhibitor of Met

AZD6094/HMPL-504 is a monotherapy for Met-amplified cancers (e.g. subsets of gastrointestinal and lung cancers), with opportunities to combine with EGFRi or chemotherapy in Met-driven cancers

AZD6094/HMPL-504 is active in both HGF dependent and independent disease settings. AZD6094/HMPL-504 activity in these settings is distinct from Met and HGF antibodies (onartuzumab and rilotumumab) that are only active in HGF dependent disease. In addition, AZD6094/HMPL-504 selectivity may provide a more favorable therapeutic index than multi-kinase inhibitors (crizotinib and cabozantinib)

Recent evaluation of AZD6094/HMPL-504 across a panel of cancer cell lines demonstrated selectivity for Met-driven disease, with Met amplified cell lines being most sensitive.

Using patient-derived xenograft (PDX) and cell lines models, we demonstrate that AZD6094/HMPL-504 has significant anti-tumor activity as a monotherapy or in combination with taxotere or gefitinib.



AZD6094/HMPL-504 is a potent and selective Met inhibitor active in HGF dep and indep models Figure 1. Sanger cell line pharmacology



Figure 1: A) 972 cell lines collapsed by disease type and shaped by MET copy number lines with <10uM (~In2.4) are highlighted for AZD6094/HMPL-504; B) Crizotinib, a multikinase inhibitor (AnLK, Met, Ros) appears far less potent than AZD6094/HMPL-504. Cel lines marked are those that overlap with AZD6094/HMPL-504 <10uM cell lines. Red line, In0=1uM

Figure 2. Gastric and lung cancer cell lines are sensitive to AZD6094/HMPI -504

A. Gastric and Lung B. Amp vs. WT MET

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Figure 3. Lung cancer cell lines with Met amplification validated for AZD6094/HMPI -504



Figure 4. A) Anti-tumor activity in EBC-1 xenografts is shown at 4 doses, 2.5, 5, 10 and 30 mpk qd. Maximal anti-tumor activity of 93% is achieved with 30 mpk AZD6094/HMPL-504. B) Body weight measurements throughout study demonstrate that AZD6094/HMPL-504 is well tolerated

AZD6094/HMPL-504 is efficacious in panel of **NSCLC** xenografts

Figure 5. Panel of cell line and patient derived xenografts characterized for Met status

	HLX-033	Calu-3	LG0645	HLX-036	HLX-036LN	HLX-0567	H441	H1993
MET				a Cali				
MET-SR/ CEP7-S0	37			2.68	290		1.24	
	HUX-033	Calu3	LG-0645	HEXE-036	HIXE-036LN	LG-0567	NCI-441	NCI-H1993
IHC	·ve	20% 3+, 80% 2+	20% 3+, 70% 2+	60% 3+, 30% 2+	90% 3+, 10% 2+	50% 3+, 30% 2+	90% 3+	80% 3+
Aug. CN	2.2	2.1	2.6	3.3	3.7	4.0	4.0	>50
Gene/CEP	1.1	0.84	1.02	0.95	1.33	1.2	<2	>10
KRAS	MT	WT	WT	WT	WT	MT	MT	WT
EGFR	wr	WT	WT	wr	wr	wr	WT (OE)	WT(OE)
HGF expression	Med	Recomb. HGF	Med/H	Low	Low	V.Low	Coimplant	Ligand Indep
Volitinib%TGI		22%	41%		154%	70%	77%	65%

Figure 6. H1993 efficacy correlates with p-Met inhibition



Figure 6. A) Anti-tumor activity in H1993 xenografts is shown at 4 doses; 1, 3, 10 and 30 mpk qd. Maximal anti-tumor activity of 65% is achieved with 30 mpk AZD6094/HMPL-504 . B) Pharmacodynamic marker, p-Met was measured by ELISA and demonstrates maximal inhibition at 4 hrs with continued inhibition at 10 hrs with 30 mpk, the most efficacious dose.

Figure 7. AZD6094/HMPL-504 is efficacious in PDX model with <4 copies Met. HLXF-036LN



Figure 7. HLXF-036LN (Met 3.72 copies, Met/CEP ratio 1.33) A) Tumor regressions with AZD6094/HMPL-504 therapy alone in model with <4 copy number and 90% 3+ IHC for Met. B) Tumor HGF (mRNA, RNAscope) and p-Met (IHC) expression are both detectable C) Analysis of tumor samples shows maximal inhibition of p-Met after 6-12 hrs AZD6094/HMPL 504 with simultaneous down-regulation of pERK1/2 signaling and caspase 3 induction

Figure 8. AZD6094/HMPL-504 and Taxotere provide durable antitumor response in PDX model with < 4 copies Met, LG0645



Figure 8. LG0645 (Met 2.64 copies, Met/CEP 1.02) A) Tumor regressions and durable esponses are demonstrated with AZD6094/HMPI -504/Taxotere combination B) Tumor HGE (mRNA, RNAScope) expression is high C) Tumor samples shows inhibition of p-Met at 4 hrs and back by 24 hrs

AZD6094/HMPL-504 is active in a model of **EGFR TKI resistance**

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Figure 9. Xenograft of EGFRm, TKI-R, Met-Amp responds to AZD6094/HMPL-504 and gefitinib combination



Figure 9. HCC827-C4R, EGFR TKI resistant (Met 4 GCN). A) Efficacy study demonstrates tumor regressions with AZD6094/HMPL-504 and Gefitinib in combination. B) Analysis of tumor samples shows inhibition of p-Met at 2-4 hrs with down regulation of pMET, p-EGFR and greater inhibition of pERK in the combination group. Additional data to support AZD6094/HMPL-504 in EGFRm Met Amp TKI resistance were previously reported (F. Zhou AACR 2013 #971)

Summary:

We have provided a platform of evidence for AZD6094/HMPL-504(Volitinib) treatment in NSCLC by selecting preclinical models representative of selected patient Met FISH or IHC scores to inform design of clinical trials and support development of rational combinations

Response to AZD6094/HMPL-504 monotherapy in WT EGFR NSCLC is seen in a subset of cell lines and patient derived xenografts with Met amp or over expression.

· Preclinical data support the combination of AZD6094/HMPL-504 and taxotere for reducing tumor burden in EGFR WT NSCLC.

AZD6094/HMPL-504, in combination with gefitinib, is efficacious in EGFRm relapsed disease, supporting Met's role in resistance in this setting. Preclinical support, internal and external, for combining with EGFR TKI is strong.

AZD6094/HMPL-504 is more potent and selective than other Met inhibitors (Crizotinib) and may have activity in subsets of NSCLC that may not respond to HGF and Met antibodies (onartuzumab and rilotumumab) due to the ability to inhibit constitutively active, non HGF dependent disease.

References:

Sanger Institute: Website ref is http://www.cancerRxgene.org http://nar.oxfordjournals.org/content/41/D1/D955

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- 2) Systematic identification of genomic markers of drug sensitivity in cancer cells. Nature. Volume:483, Pages:570-575 (29 March 2012)

