## 2020 EORTC-NCI-AACR Drug Design Abstract ENA20-0018 Poster Number 143

# Discovery and Characterization of Potent and Selective AXL Receptor Tyrosine Kinase Inhibitors for Cancer Therapy

## **OVERVIEW**

- ✤ AXL receptor tyrosine kinase (AXL) is a transmembrane protein which is overexpressed in a variety of cancers and has been implicated in the development of resistance to chemotherapy and immunotherapy.
- \* Arcus Biosciences is developing novel and selective AXL receptor tyrosine kinase inhibitors as potential anti-cancer drugs.
- This poster describes the discovery and characterization of two novel inhibitors, developed through study of earlier literature compounds. These compounds are characterized by various in vitro assays and show good potency and selectivity toward AXL in relation to similar kinases. Initial pharmacokinetic profiles are promising for future in vivo efficacy studies.

## **AXL BIOLOGY & DOWNSTREAM EFFECTS**

- ✤ Binding of Gas6 to AXL on the tumor cell surface induces AXL autophosphorylation.<sup>1</sup>
- ✤ AXL activation induces PI3K-AKT pathway signaling, resulting in upregulation of genes associated with EMT<sup>2</sup> & DNA damage repair<sup>3</sup> leading to the rapeutic resistance to multiple agents.



Figure 1. Gas6 induced AXL Signaling. Binding of Gas6 to AXL leads to AXL autophosphorylation and signaling through the PI3K-AKT pathway, inducing genes that drive EMT & DNA damage repair

#### Structural Biology and Rationale of AXL Receptor Tyrosine Kinase Inhibition

Upon GAS6-induced AXL dimerization, tyrosine residues throughout the intracellular domain are phosphorylated by action of the intracellular kinase domain, resulting in various biological effects.<sup>4</sup> Dimerization can be prevented extracellularly through use of an antibody or AXL 'decoy receptor' (Figure 2). Alternatively, phosphorylation is mitigated through small molecule inhibition of the intracellular kinase domain (Figures 2C and 3) We embarked upon the design of novel AXL inhibitors, utilizing a pharmacophore mapping approach based on existing commercial and academic molecules. Aside from pan-kinase selectivity, we specifically sought to avoid MERTK and TYRO3 inhibition to reduce any unwanted off-target effects.<sup>5</sup> Due to the high homology of these catalytic domains, design of AXL-selective inhibitors is challenging.<sup>6</sup>



Figure 2. Selected strategies for inhibiting AXL autophosphorylation. a) Blocking antibody. b) AXL 'decoy receptor' to sequester GAS6. c) Small molecule inhibition of kinase domain.

#### Assay

hAXL HTRF IC<sub>50</sub> (biochemical, nM)

mAXL HTRF IC<sub>50</sub> (biochemical, nM)

hMERTK / hTYRO3 HTRF selectivity (biochemical, enzyme  $IC_{50}$  over AXL  $IC_{50}$ )

hAXL NanoBRET<sup>TM</sup>  $K_{D}$  (cellular, nM)

hAXL PathHunter<sup>®</sup> IC<sub>50</sub> (cellular, nM)

hAXL PathHunter<sup>®</sup> IC<sub>50</sub> (cellular, nM) 100% serum

**Table 1.** Characterization and comparison of optimized Arcus and benchmark AXL inhibitors. Kinase activity of AXL, MERTK and TYRO3 were tested using HTRF KinEASE – TK kit (CisBio) in the presence of 700 µM ATP. Inhibitor engagement to intracellularly expressed AXL kinase domain was detected using AXL NanoBRET<sup>™</sup> TE intracellular kinase assay (Promega) with transiently transfected HEK293 cells. Compound activity in inhibiting SH2 domain translocation to phosphorylated AXL cytoplasmic domain upon AXL activation was tested using PathHunter<sup>®</sup> U2OS AXL functional assay (Eurofins DiscoverX). Cells were preincubated with inhibitors for 1 hr followed by 3 hrs Gas6 (1 µg/mL) induction. Assay was carried out in either a serum-free or 100% human serum medium.



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## **INITIAL DESIGN, OPTIMIZATION, AND CHARACTERIZATION OF ARCUS AXL INHIBITORS**



Figure 3. X-ray structure of AXL/smal molecule inhibitor complex (PDB: 5U6B).<sup>6</sup>

## **Pan-Kinase Selectivity of Arcus AXL Inhibitors**

In addition to being selective against MERTK and TYRO3, Compound B was tested against a broad non-mutant kinase panel to determine any other potential off-target effects. Using a qPCR-based competition binding experiment, Compound B showed <20% of control activity (indicating stronger binding) against 10 other kinase targets, excluding MERTK and TYRO3. Follow-up  $K_d$  values were determined through qPCR-based competition binding dose response.



Figure 5. Kinome map (403 total) showing significant kinase/Compound B interactions as red circles.

## **AXL Engagement with Gas6 Results in Decreased Soluble AXL** and Is Rescued by AXL Inhibition



Figure 4. A) Inhibition of AXL, MERTK, and TYRO3 by Compound A in the HTRF biochemical assay. B) Inhibition of AXL by Compound A, Compound B, and Bemcentinib in the AXL PathHunter® cellular assay with 100% serum.





Figure 6. Prototype AXL inhibitors Compound A and Compound B rescue soluble AXL levels that are reduced by Gas6. A549 human lung carcinoma cells (top) and 4T1 murine breast cancer cells (bottom) were treated with 1 – 1,000 nM (A549) or 0.1 – 300 nM (4T1) of Compound A or Compound B for 1 h. 200 ng/mL hGas6 (A549 cells) or 500 ng/mL mGas6 (4T1) was added 1 h after addition of AXL inhibitors and cells were cultured for 72 h. Supernatants were collected and soluble AXL levels were determined by ELISA.



	Kinase	<i>K</i> d	Kinase	K <sub>d</sub>
CK1 AGC	BMPR1B	8.4 nM (221x)	MAP4K5	1.7 nM (44x)
	DRAK1	4.5 nM (118x)	SGK	2.3 nM (61x)
	HPK1	4.5 nM (118x)	SLK	84 nM (2210x)
	MAP4K2	13 nM (342x)	STK16	13 nM (342x)
	MAP4K3	4.7 nM (123x)	TNIK	14 nM (368x)

**Table 2.** K<sub>d</sub> values for selected kinases with Compound B. Selectivity values in parentheses defined as kinase-inhibitor  $K_{d}$  over by AXL-inhibitor  $K_{d}$  (0.038)

## **PHARMACOKINETICS and SAFETY**

### Pharmacokinetic and Safety Characterization

Both Compound A and Compound B exhibit favorable in vitro pharmacokinetic profiles with low intrinsic clearance in both rat and human hepatocytes (Table 3). When dosed in rat models, Compound A and Compound B show moderate to low clearance (Table 4). Additionally, Compound B exhibits a good safety profile, with acceptable CYP isoform and hERG inhibition (Table 5) as well as no time-dependent CYP inhibition (data not shown).

	CL <sub>int</sub> (µL/min/10 <sup>6</sup> cells)				
Compound	Mouse	Rat	Dog	Human	
Compound A		3.8		2.6	
Compound B	13.8	<0.7	<0.7	3.1	
Table 3. Summary of hepatocyte stability in various species.					

#### *In Vivo* Rat Pharmacokinetics

Compound	<b>CL</b> (L/h/kg)	V <sub>ss</sub> (L/kg)	<b>T<sub>1/2</sub></b> (h)
Compound A	1.6	3.3	1.9
Compound B	0.66	2.2	4.8
Table 4. Summarv	of experimental rat	PK parameters. Rate	s were dosed 0.25

mg/kg IV in DMAC:Ethanol:PG (ca. 1:1:1).

### **CYP Isoform and hERG Inhibition of Compound B**

IC <sub>50</sub> 2C8	IC <sub>50</sub> 2C9	IC <sub>50</sub> 2C19	IC <sub>50</sub> 2D6	IC <sub>50</sub> 3A4	hERG inh.
(μΜ)	(μΜ)	(μΜ)	(μΜ)	(μΜ)	at 10 μΜ
>40	>40	38	31	11	31%

**Table 5.** In vitro evaluation of Compound B for potential to inhibit major human
 drug metabolizing enzymes of the cytochrome P450 family and hERG.

- 1 and Figure 4).
- selectivity ratio is for MAP4K5, at 44-fold (Figure 5 and Table 2).
- concentrations (Figure 6).

## REFERENCES

- 1) Linger, et al. Adv. Cancer Res. 2008, 100:35.
- 2) Wang, et al. Theranostics. 2016, 6(8):1205.
- 3) Balaji, et al. Mol. Cancer Res. 2016, 15(1):45. 4) Zhu, et al. Mol. Cancer 2019, 18:153.
- Graham, et al. Nat. Rev. Cancer 2014, 14(12):769.

### Hepatocyte Stability

## **SUMMARY**

Two unique prototype AXL inhibitors have been discovered and characterized in functional biochemical and cell-based assays. When taken together, these compounds show improved potency and selectivity compared to previous small-molecule AXL inhibitors (Table

Compound B shows a good pan-kinase selectivity profile, suggesting minimal off-target effects within this inhibitor class. The lowest

✤ A decrease in soluble AXL levels is observed upon addition of GAS6 in both lung and breast cancer cells. Addition of either Compounds A or B resulted in an increase of soluble AXL levels to pre-GAS6

✤ Both Arcus inhibitors show promising pharmacokinetic profiles with low intrinsic and in vivo clearances. Compound B shows a good safety profile, with minimal CYP and hERG liabilities (Tables 3-5).

> 6) Gajiwala, et al. J. Biol. Chem. 2017, 292(38):15705. 7) Data generated in-house. Compounds purchased from Synnovator (Bemcentinib), Advanced ChemBlocks (Dubermatinib), and Cayman Chemical (Sitravatinib).