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# *N*-(3-fluoro-4-(2-arylthieno[3,2-*b*]pyridin-7-yloxy)phenyl)-2-oxo-3-phenylimidazolidine-1-carboxamides: A novel series of dual c-Met/VEGFR2 receptor tyrosine kinase inhibitors

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## ABSTRACT

A series of *N*-(3-fluoro-4-(2-arylthieno[3,2-*b*]pyridin-7-yloxy)phenyl)-2-oxo-3-phenylimidazolidine-1carboxamides targeting c-Met and VEGFR2 tyrosine kinases was designed and synthesized. The compounds were potent against these two enzymes with  $IC_{50}$  values in the low nanomolar range in vitro, possessed favorable pharmacokinetic profiles and showed high efficacy in vivo in several human tumor xenograft models in mice.

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Receptor tyrosine kinases (RTKs) play crucial roles in numerous signal transduction pathways and cellular processes. Many are implicated in cancer,<sup>1</sup> including the Hepatocyte Growth Factor Receptor c-Met,<sup>2</sup> which is over-expressed and/or mutated in various human tumor types, and the Vascular Endothelial Growth Factor Receptors (VEGFR), which play key roles in tumor angiogenesis.<sup>3</sup> Dysregulation of these RTKs affects cell proliferation, survival and motility, leading to tumor growth, angiogenesis, and metastasis. The combined inhibition of several RTKs is a promising approach for cancer therapy, as it targets multiple pathways involved in angiogenesis and tumor survival.<sup>4</sup>

Figure 1 highlights some of the known inhibitors of c-Met<sup>5</sup> and/ or VEGFR<sup>6</sup> enzymes from the patent and scientific literature.

Sutent (1),<sup>7</sup> a marketed anticancer drug for patients with advanced renal cancer and gastrointestinal stromal tumors, is a potent inhibitor of VEGFR2 but does not inhibit c-Met. Conversely, PHA-665752 ( $\mathbf{2}$ )<sup>8</sup> is a potent inhibitor of c-Met but a weak inhibitor of VEGFR2. In contrast to both molecules, representative quinoline compounds of Kirin ( $\mathbf{3}$ )<sup>9</sup> and Exelixis ( $\mathbf{4}$ )<sup>10</sup> are effective against both enzymes.

Recently, we reported on a series of 2-substituted thieno[3,2*b*]pyridine-based c-Met/VEGFR2 inhibitors exemplified by compounds **5–10** (Table 1) bearing acyclic structural fragments, such as arylacetyl thioureas, arylacetyl ureas or various arylmalonamides.<sup>11</sup> Of particular interest was compound **8**, which showed the same potency against both c-Met and VEGFR2 as its desmethyl analogue **7**. We hypothesized that connecting the *N*-methyl with the central methylene of the malonamide head group of compound **8** via a CH<sub>2</sub>-linker, as illustrated in the *N*-phenylpyrrolidone-type compounds **11** and **12**, could preserve the same kinase inhibitory properties. It turned out that compound **11** was indeed similar to compound **8** (Table 1).<sup>12,13</sup> Moving the carbonyl-group (compound **13**) or creating a quarternary center (compound **14**) resulted in a significant loss of activity against VEGFR2 for both compounds compared to compound **11**. The activities of both

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Figure 1. c-Met and/or VEGFR inhibitors with their IC<sub>50</sub> values (generated in house).

#### Table 1

Enzymatic c-Met and VEGFR2 and cellular TPR-Met phosphorylation assay results for compounds 5-15 of general structure



Compound	R <sup>1</sup>	R <sup>2</sup>	c-Met IC <sub>50</sub> (µM)	VEGFR2 IC50 (µM)	Phospho-TPR-Met $IC_{50}$ ( $\mu$ M)
5	Me	is solution	0.037	0.026	0.05
6	Me	ζξ Ο Ο Ο	0.025	0.037	n.a.
7	Me	i c l l l l l l l l l l l l l l l l l l	0.028	0.015	0.31
8	Me	<sup>v</sup> <sup>2</sup> <sup>5</sup> ↓ N ↓ ↓ ↓ ↓ ↓ ↓ ↓ ↓ ↓ ↓ ↓ ↓ ↓ ↓ ↓ ↓ ↓	0.043	0.010	0.16
9	Me		0.028	0.009	0.06
10	Me	X <sup>2</sup> O O O	1	0.057	1.44
11	Me	×	0.035	0.039	0.37
12	<i>i-</i> Pr	× (n-()	0.039	0.016	0.21
13	i-Pr		0.3	0.8	n.a.
14	Me		0.067	0.23	1.00
15	Ме	25 N N N	0.023	0.008	0.12



Scheme 1. Synthesis of 11 and 15. Reagents and conditions: (a) i–*n*-BuLi, THF, –78 °C; ii–ZnCl<sub>2</sub>, THF, –78 °C to rt; iii–Pd(PPh<sub>3</sub>)<sub>4</sub>, 4-iodo-1-methyl-1*H*-imidazole, THF, reflux. (b) 2-Fluoro-4-nitrophenol, K<sub>2</sub>CO<sub>3</sub>, Ph<sub>2</sub>O, 180 °C. (c) NiCl<sub>2</sub>, NaBH<sub>4</sub>, MeOH. (d) HATU reagent, DIPEA, DMF. (e) DIPEA, DCM.

compounds against c-Met changed as well—only slightly for compound **14** but more significantly for compound **13**. We then replaced the stereogenic center in the pyrrolidone fragment of compound **11** with a nitrogen atom to form the corresponding cyclic urea analogue **15**. This compound turned out to be equipotent to compound **8** at both the enzymatic<sup>14</sup> and cellular<sup>15</sup> level. Thus, rigidification of the arylmalonamide fragment into the arylimidazolidine motif is well tolerated, giving rise to a novel class of c-Met/ VEGFR2 kinase inhibitors, the *N*-(3-fluoro-4-(2-arylthieno[3,2-*b*]pyridin-7-yloxy)phenyl)-2-oxo-3-phenylimidazolidine-1-carboxamides (e.g. compound **15**).<sup>12,13</sup>

Synthesis of compounds **11** and **15** is shown in Scheme 1. 7-Chlorothieno[3,2-*b*]pyridine and 4-iodo-1-methyl-1*H*-imidazole underwent a *Negishi* coupling reaction to form a methylimidazolyl substituted thienopyridine **A**.<sup>16</sup> This material was then reacted with 2-fluoro-4-nitrophenol under basic conditions at high temperature to form the nitro compound **B**. The nitro group of **B** was reduced using a mixture of sodium borohydride and nickel chloride in methanol to form the aniline **C**. The intermediate **C** was treated either with 2-oxo-1-phenylpyrrolidine-3-carboxylic acid in the presence of HATU reagent and diisopropylethylamine to form compound **11**, or with 2-oxo-3-phenylimidazolidine-1-carbonyl chloride<sup>17</sup> and diisopropylethylamine to give the cyclic urea **15**.

To study the SAR of compound **15**, we first explored the substituents at position 2 of the thieno[3,2-*b*]pyridine template (Table 2). Elongation of the chain (compound **16**) and/or incorporation of a basic center (compounds **17–19**) were well tolerated and provided potent molecular entities. Changing the orientation of the imidazolyl ring system—proceeding from compound **15** and **16** to compounds **20** and **21**—did not affect potency against c-Met but were slightly less active against VEGFR2. Replacement of the imidazole ring by a *para*-substituted phenyl ring (compounds **22–25**) was well tolerated. However, having three basic nitrogens, as in compound **25** resulted in a dramatic decrease in cell-based activity. The *m*-acetophenyl group in compound **26** proved to be detrimental and resulted in reduced enzyme inhibitory activity. A last modification which was explored was the replacement of the heteroaryl or aryl appendages by either an amide (compound **27**)

or a tetrahydropyridine (compound **28**). Interestingly, compound **28** was significantly less active against VEGFR2.

We next investigated the presence of a small substituent on the phenyl ring of the phenylimidazolidine fragment of the ethyl substituted analogues of compound **16**. (Table 3). The ortho- and para-substituted fluoride atom allowed us to slightly improve VEGFR2 inhibitory activity (compounds **29** and **30**, compared to compound **16**). On the other hand, the presence of an *ortho*-fluoro group (compound **30**) had a slightly negative impact on cell-based activity. Finally, although compound **31** with an *ortho*-methoxy group was equipotent to compound **16**, it showed reduced activity in our c-Met-driven cellular assays (data not shown).

We also examined the replacement of the oxygen atoms by sulfur atoms in the dicarbonyl framework of compound **15** (Table 4). Compound **32** (an analogue of compound **5**) was as potent as compound **15**, while compound **33** was inactive against c-Met and VEGFR2.

Rat pharmacokinetic analyses were performed on a small set of compounds (Table 5). Compounds with *N*-methylimidazole fragments (e.g. compounds **11** and **15**) showed a short to medium half-life and poor bioavailability. Results were encouraging, particularly for compound **29** as it displayed a long half-life and decent bioavailability, an appreciable improvement when compared to compound **5**. Compounds that had basic substituents generally had a shorter half-life and lower bioavailability (data not shown).

Compound **29** was further evaluated in other cell-based assays. First, it was tested for its ability to inhibit two c-Met-specific functional endpoints (Table 6). Compound **29** dramatically impaired HGF-induced epithelial cell migration and scattering, showing that it is able to inhibit c-Met-dependent cell motility events. Second, compound **29** was tested in a variety of VEGF-dependent cellular assays (Table 6). It potently inhibited the VEGF-induced phosphorylation of ERK, a downstream effector of the VEGF pathway, and also impeded the VEGF-dependent proliferation of human umbilical vein endothelial cells (HUVEC). In an *in vitro* angiogenesis assay, which measures the formation of tubules generated by a co-culture of endothelial and fibroblast cells (Angiokit<sup>TM</sup>, TCS Cellworks), compound **29** potently inhibited tubule growth and was also able to almost completely inhibit tubule formation at the 100 nM dose

### Table 2

Enzymatic c-Met and VEGFR2 and cellular TPR-Met phosphorylation assay results for compounds **16–28** of general structure



Compound	R	c-Met IC <sub>50</sub> (µM)	VEGFR2 IC <sub>50</sub> (µM)	Phospho-TPR-Met $IC_{50}$ ( $\mu$ M)
16	Et N	0.025	0.012	0.04
17	Me <sub>2</sub> N N - 5-	0.034	0.014	0.06
18		0.035	0.02	0.32
19	$\sum_{N}$	0.021	0.008	0.30
20	∫ N→ξ- N Me	0.026	0.031	0.12
21	$ \prod_{\substack{N \\ Et}}^{N} \xi^{-}$	0.024	0.026	0.14
22		0.022	0.011	0.42
23	ν Me'	0.029	0.012	0.25
24	MeHN	0.016	0.008	0.08
25	-N_Nξ-	0.028	0.003	0.83
26		0.2	0.88	0.27
27	Me <sub>2</sub> N <sup>1,1</sup> →-ξ-	0.042	0.023	0.23
28	Me - N	0.033	0.17	0.30

#### Table 3

Enzymatic c-Met and VEGFR2 and cellular TPR-Met phosphorylation assay results for compounds **29–31** of general structure



Compound	R	c-Met IC <sub>50</sub> (µM)	VEGFR2 IC <sub>50</sub> (µM)	Phospho- TPR-Met IC <sub>50</sub> (µM)
16	Н	0.025	0.012	0.04
29	p-F	0.027	0.007	0.05
30	o-F	0.021	0.003	0.10
31	0-	0.026	0.014	0.06
	OMe			

#### Table 4

Enzymatic c-Met and VEGFR2 and cellular TPR-Met phosphorylation assay results for compounds  ${\bf 32}$  and  ${\bf 33}$  of general structure



-			(μM)	(μM)	(μM)	
15	0	0	0.023	0.008	0.12	
32	S	0	0.024	0.012	0.08	
33	0	S	2	1.8	>10	

#### Table 5

Rat PK analysis: iv half-life and bioavailability (F%) for select compounds with general structure



Compound	R <sup>1</sup>	Х	R <sup>2</sup>	<i>t</i> <sub>1/2</sub> iv (h) <sup>a</sup>	F (%) <sup>b</sup>
5	Me			1.2	12
11	Me	CH	Н	1.1	14
15	Me	Ν	Н	2.3	13
29	Et	Ν	F	6.3	37

<sup>a</sup> iv doses 2.17-2.86 mg/kg.

<sup>b</sup> po doses 4.28-4.98 mg/kg.

where cells were purposely supplemented with VEGF (Table 6). These results show that compound **29** can potently inhibit VEGF-dependent functional activity in cells.

#### Table 7

Effect of compound 29 on human cancer cell proliferation

Compound	MKN-45 IC <sub>50</sub>	MNNG-HOS	HCT 116 IC <sub>50</sub>	MDA MB-231
	(μM)	IC <sub>50</sub> (μM)	(µM)	IC <sub>50</sub> (μM)
29	0.01	0.033	13	20

Third, compound **29** was evaluated for its ability to inhibit the proliferation of a variety of human cancer cell lines, using a 72 h MTT assay. It displayed nanomolar IC<sub>50</sub>s in tumor lines whose growth is known to be sensitive to c-Met inhibition, namely the MKN-45 gastric carcinoma and the TPR-Met-expressing MNNG-HOS cell lines. In contrast, significantly higher concentrations of compound **29** were required to inhibit the growth of c-Met-independent HCT116 colon carcinoma and MDA-MB-231 breast adenocarcinoma cells (Table 7). These results show that compound **29** specifically targets c-Met-driven cancer cell proliferation in culture.

Lastly, compound **29** was also profiled against a small set of kinases [using Millipore's Kinaseprofiler<sup>TM</sup> assay services] (Table 8). It strongly inhibited VEGFR1, VEGFR3, Ron and Tie-2, moderately inhibited Flt-3 and was not active against Aurora A.

Finally, compound **29** was tested in a mouse xenograft model bearing tumors derived from the c-Met-driven MKN-45 cells (Fig. 2). Interestingly, oral administration of a 40 mg/kg dose once daily for 12 days caused significant regression of MKN-45 tumor growth.

In conclusion, a series of 2-oxo-3-aryl-*N*-(4-(thieno[3,2-*b*]pyridin-7-yloxy)phenyl)imidazolidine-1-carboxamides was designed and synthesized. The compounds show low nanomolar inhibitory activity of both c-Met and VEGFR2 enzymes.

## Table 8

% enzyme inhibition at 100 nM of 29

VEGFR-1	VEGFR-3	Ron	Tie-2	FLT-3	Aurora A
99	99	100	99	80	9



**Figure 2.** Oral anti-tumor activity of compound **29** in the MKN-45 tumor xenograft model. Compound **29** was dosed po, once daily, at 40 mg/kg (6 mice per group) starting on Day 0.

Table 6			
Effect of compound 29 on c-Met and	l VEGF-mediated	cellular	endpoints

Compound	A549 wound healing	DU145 scattering	VEGF-dept phospho-	VEGF- dept HUVEC	Angiokit, Tubule	Angiokit, Tubule length
	inh. IC <sub>50</sub> (μM)	inh. IC <sub>50</sub> (µM)	ERK IC <sub>50</sub> (µM)	proliferation IC <sub>50</sub> (µM)	length IC <sub>50</sub> (μM)	with VEGF (% inh.)
29	0.24	0.08	0.001	0.006	0.006	98

Compound **29**, the lead molecule, shows an excellent in vitro profile and favorable pharmacokinetic characteristics and exhibits significant in vivo efficacy in the MKN-45 human xenograft mouse model. Efforts to further optimize pharmacokinetic characteristics and physicochemical properties of *N*-(3-fluoro-4-(2-arylthie-no[3,2-*b*]pyridin-7-yloxy)phenyl)-2-oxo-3-phenylimidazolidine-1-carboxamides as well as attempts to expand this series of compounds towards chemical entities with a different kinase inhibitory profile is in progress.

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- 12. The details for the synthesis and the characterization of all the new compounds are described in US 2007/0004675 A1 and WO 2007/107005 A1.
- 13. The present work was first presented (poster session) at the International Conference on Medicinal Chemistry. Drug Discovery and Selection. 43<sup>èmes</sup> Rencontres internationales de Chimie Thérapeutique. July 4–6, 2007, Faculté de Pharmacie, Lille, France.
- 14. In Vitro Kinase Assays (c-Met and VEGFR-2/KDR): Preparation of GST fusion proteins: recombinant baculovirus containing the catalytic domain of c-Met and of the VEGFR-2/KDR receptor fused to glutathione S-transferase (GST) fusion genes were used to infect High five (c-Met) or Sf9 (VEGFR-2/KDR) cells at a multiplicity of infection of 1 or 0.1, respectively. Cell lysates were prepared after  $\sim$ 72 h of infection in 1% Triton X-100, 2 µg of leupeptin/mL, and 2 µg of aprotinin/mL after ~72 h of infection in phosphate-buffered saline, and the fusion proteins were purified over glutathione agarose (Sigma) according to the manufacturer's instructions. Biochemical kinase assays for IC<sub>50</sub> determination and kinetic studies: Inhibition of c-Met and VEGFR-2/KDR was measured in a DELFIA<sup>TM</sup> assay (Perkin Elmer). The substrate poly(Glu<sub>4</sub>,Tyr) was immobilized onto black high-binding polystyrene 96-well plates (Nunc Maxisorp). The c-Met kinase reaction was conducted in 25 mM Hepes pH 7.5 containing 20 mM NaCl, 10 mM MgCl<sub>2</sub>, 5 mM β-Mercaptoethanol, 0.1 mg/mL bovine serum albumin (BSA) and 20 µM vanadate, while the VEGFR-2/KDR reaction was conducted in 60 mM Hepes pH 7.5 containing 3 mM MgCl<sub>2</sub>, 3 mM MnCl<sub>2</sub>, 1.2 mM β-Mercaptoethanol, 0.1 mg/mL BSA and 3 uM vanadate. ATP concentrations in the assay were 10  $\mu$ M for c-Met (5× the  $K_m$ ) and 0.6  $\mu$ M for VEGFR-2/KDR ( $2 \times$  the  $K_m$ ). Enzyme concentration was 25 nM (c-Met) or 5 nM (VEGFR-2/KDR).The recombinant enzymes were pre-incubated with inhibitor and Mg-ATP on ice in polypropylene 96-well plates for 4 min, and then transferred to the substrate coated plates. The subsequent kinase reaction took place at 30 °C for 30 min. (c-Met) or 10 min. (VEGFR-2/KDR). After incubation, the kinase reactions were quenched with EDTA and the plates were washed. Phosphorylated product was detected by incubation with Europium-labeled anti-phosphotyrosine MoAb. After washing the plates, bound MoAb was detected by time-resolved fluorescence in a Gemini SpectraMax reader (Molecular Devices). Inhibitors were tested at seven different concentrations each in triplicate. IC<sub>50</sub>s were calculated in a four parameters equation curve plotting inhibition (%).
- 15. A cellular clone of 293T cells stably expressing TPR-Met (Park, M.; Dean, M.; Cooper, C. S.; Schmidt, M.; O'Brien, S. J.; Blair, D. G.; Vande Woude, G. F. *Cell* 1986, 45, 895), the activated mutated form of the receptor Met, under a CMV promoter was derived. Cells were treated with compounds dilutions for 150 min and lysate samples from treatment wells were transferred to high binding white polysterene 96-well plates (Corning). TPR-Met autophosphorylated levels were detected by ELISA using the primary antibodies anti-phospho-Tyrosine (Millipore, 4G10) and a reporter antibody, anti-mouse horseradish peroxidase (Sigma). Plates were washed on a plate washer (SkanWasher, Molecular Devises) and subsequently incubated with chemiluminescent substrate solution (ECL, Roche). Luminescence signal was captured on a Polar Star Optima apparatus (BMG LabTech). Average values of triplicate treatment points were used to prepare IC<sub>50</sub> curves using a 4-parameter fit model. These curves were calculated using GraFit 5.0 software.
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