

## N-(3-Ethynyl-2,4-difluorophenyl)sulfonamide Derivatives as Selective Raf Inhibitors

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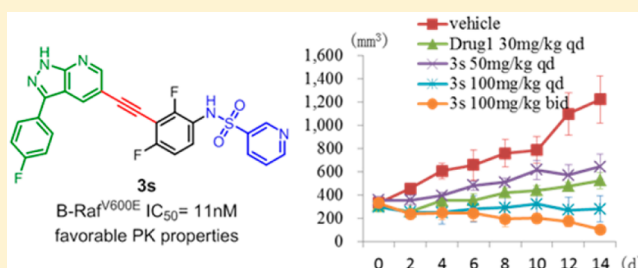
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**S** Supporting Information

**ABSTRACT:** A series of N-(3-ethynyl-2,4-difluorophenyl)-sulfonamides were identified as new selective Raf inhibitors. The compounds potently inhibit B-Raf<sup>V600E</sup> with low nanomolar IC<sub>50</sub> values and exhibit excellent target specificity in a selectivity profiling investigation against 468 kinases. They strongly suppress proliferation of a panel of human cancer cell lines and patient-derived melanoma cells with B-Raf<sup>V600E</sup> mutation while being significantly less potent to the cells with B-Raf<sup>WT</sup>. The compounds also display favorable pharmacokinetic properties with a preferred example (3s) demonstrating significant *in vivo* antitumor efficacy in a xenograft mouse model of B-Raf<sup>V600E</sup> mutated Colo205 human colorectal cancer cells, supporting it as a promising lead compound for further anticancer drug discovery.

**KEYWORDS:** B-Raf, colon cancer, melanoma, kinase inhibitor, targeted therapy



The Raf serine/threonine kinases are key components of mitogen-activated protein kinase (MAPK) cascade.<sup>1</sup> Among three isoforms of the family (i.e., A-Raf, B-Raf, and C-Raf), mutations of B-Raf are the most frequently detected in human cancers,<sup>2</sup> including 50–80% of melanoma,<sup>3</sup> ~100% of hairy cell leukemia,<sup>4</sup> 45–50% of papillary thyroid carcinoma,<sup>5</sup> 11% of colorectal cancer,<sup>3</sup> and others. Notably, over 90% of B-Raf mutations are the substitution of valine to glutamate at residue 600 (V600E). This substitution mimics phosphorylation of the activation loop, elevating the *in vitro* kinase activity by up to 500–700-fold compared with its wide-type counterpart.<sup>6</sup> Therefore, selectively inhibiting B-Raf becomes an attractive strategy for the clinical treatment of B-Raf<sup>V600E</sup>-driven human cancers.<sup>7</sup>

Several classes of B-Raf inhibitors have been discovered,<sup>8,9</sup> among which vemurafenib (1, Figure 1) and dabrafenib (2, Supporting Information) have been approved by US FDA for the treatment of metastatic and unresectable melanoma harboring B-Raf mutations.<sup>10,11</sup> Many other clinical investigations of B-Raf inhibitors alone or in combination with other kinase inhibitors, immunotherapies, or conventional chemotherapies, are ongoing, for the treatment of metastatic

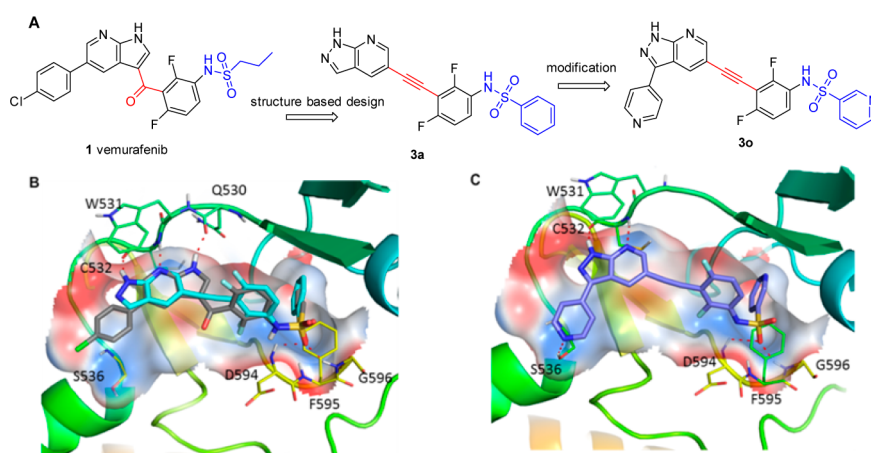
colorectal cancer, papillary thyroid carcinoma, and metastatic nonsmall cell lung cancer.<sup>12</sup>

However, intrinsic or acquired resistance against current B-Raf inhibitor therapies becomes a major challenge.<sup>13</sup> For instance, a majority of colorectal cancer patients display inherent resistance against vemurafenib, although they were detected to harbor B-Raf<sup>V600E</sup> mutation.<sup>14</sup> The exact mechanism for the resistance remains elusive, but development of new inhibitors with differentiated chemical scaffolds may be a valuable strategy to overcome or delay the occurrence of this challenging medical dilemma. Herein, we describe the design and optimization of N-(3-ethynyl-2,4-difluorophenyl)-sulfonamide derivatives as new selective Raf inhibitors.

An X-ray crystallographic analysis reveals that vemurafenib binds to B-Raf with a DFG-in and  $\alpha$ C-helix-out inactive conformation.<sup>11</sup> The 1H-pyrrolo[2,3-b]pyridine core forms two pairs of hydrogen bonds with the backbone amide of Cys532 and the backbone carbonyl of Gln530, respectively (Figure 1B). The sulfonamide may form two extra hydrogen bonds with

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**Figure 1.** Design of new Raf inhibitors (A). Predicted binding model of **3a** (blue) superimposed to vemurafenib (gray) in a crystal structure of B-Raf (PDB: 3OG7) (B). Predicted binding model of **3o** with B-Raf (C).

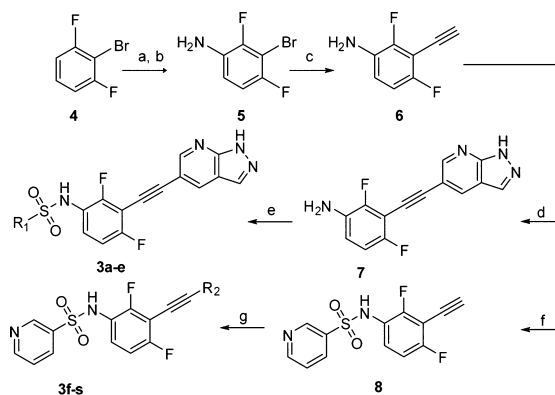
Asp594 and Gly596, respectively, and the propyl group is critical for selectivity by occupying a small lipophilic pocket enlarged by an outward shift of  $\alpha$ C-helix of the protein. On the basis of these observations, we rationally designed a series of *N*-(3-ethynyl-2, 4-difluorophenyl)sulfonamides (**3**) as new B-Raf inhibitors, in which a druglike scaffold 1*H*-pyrazolo[3,4-*b*]pyridine was utilized as the crucial hinge-binding moiety; a conformation and distance favorable ethynyl linker was introduced between the hinge-binding core and a difluorophenyl group; and the sulfonamide moiety was preserved to form the important hydrogen bonding networks with the protein. Instead of using an alkyl substituent in the sulfonamide (i.e., a propyl group in vemurafenib), an aryl group was introduced to capture a potential CH- $\pi$  interaction with Phe595 (Figure 1B).

The designed compounds were synthesized by using Sonogashira coupling as the key steps (Scheme 1). Briefly, the intermediate **6** was readily prepared from 2-bromo-1,3-difluorobenzene (**4**) through consequent procedures including

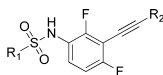
a standard nitration, a stannous chloride mediated reduction, a modified Sonogashira coupling, and a detrimethylsilylation.<sup>15</sup> The hinge-binding 1*H*-pyrazolo-[3,4-*b*]pyridine was then introduced via another Sonogashira reaction. A direct arylsulfonation of **7** with different arylsulfonfyl chlorides yielded the final sulfonamides **3a–3e** with good to moderate yields. Compounds **3f–3s** were prepared through similar protocols (Supporting Information).

A preliminary computational study suggested that **3a** might bind to the DFG-in B-Raf<sup>V600E</sup> with a similar mode to that of vemurafenib (Figure 1B). A biological evaluation showed that **3a** indeed inhibited B-Raf<sup>V600E</sup> kinase with an IC<sub>50</sub> value of 0.149  $\mu$ M in an FRET-based Z'Lyte assay.<sup>16</sup> It also selectively suppresses the proliferation of Colo205 human colorectal cancer cells harboring B-Raf<sup>V600E</sup> mutation with an IC<sub>50</sub> value of 2.103  $\mu$ M but is obviously less potent against the growth of HCT116 colorectal carcinoma cells with B-Raf<sup>WT</sup> (IC<sub>50</sub> > 10  $\mu$ M) (Table 1). These results support that **3a** may serve as a lead compound for new B-Raf<sup>V600E</sup> inhibitor discovery. An extensive structure–activity relationship investigation was then conducted, and the results revealed that an F substitution at different positions of R<sub>1</sub>-phenyl ring in **3a** could achieve diverse impact on the B-Raf<sup>V600E</sup> inhibitory activity. For instance, *ortho*-F substituted analogue **3b** displays a similar B-Raf<sup>V600E</sup> inhibitory activity to that of **3a**, while the *meta*-fluoro compound **3c** is 2–3-fold more potent, with an IC<sub>50</sub> value of 0.068  $\mu$ M. However, when an F atom was introduced in the para-position, the resulting compound **3d** was 6-fold less potent. Further study showed that the *meta*-fluoro phenyl group in **3c** could be replaced by a 3-pyridyl moiety (**3f**) to display almost identical potency against B-Raf<sup>V600E</sup>, with an IC<sub>50</sub> value of 0.051  $\mu$ M. However, the 2-pyridyl analogue (**3e**) is about 10-times less potent. It is hypothesized that **3c** and **3f** may induce a conformational rearrangement of the protein to generate additional hydrogen bonds between the F or N atom and the backbone of the DFG motif. A future cocrystallographic investigation will be highly valuable to demonstrate the precise interactions. Our docking study has suggested that the two pairs of hydrogen bonds formed by 1*H*-pyrazolo[3,4-*b*]pyridine in **3f** with Cys532 contribute greatly to the compound's B-Raf<sup>V600E</sup> inhibition. Indeed, compound **3h** lacking a hydrogen bond acceptor showed an IC<sub>50</sub> value against B-Raf<sup>V600E</sup> of 4.31  $\mu$ M, which was 85-times less potent than **3f**. However, the removal of a hydrogen bond donor (**3i**)

#### Scheme 1. Synthesis of Compounds **3a–3s**<sup>a</sup>



<sup>a</sup>Reagents and conditions: (a) con. H<sub>2</sub>SO<sub>4</sub>, KNO<sub>3</sub>, rt, 2.0 h, 91%; (b) SnCl<sub>2</sub>, con. HCl, EtOH, reflux, 93%; (c) (i) Pd(dba)<sub>2</sub>, CuI, ethynyltrimethylsilane, *t*-(Bu)<sub>3</sub>P, K<sub>2</sub>CO<sub>3</sub>, dry THF, 120 °C, 48 h; (ii) *n*-(Bu)<sub>4</sub>NF, THF, rt, 1 h, 46%; (d) 6-bromo-3*H*-pyrazolo[3,4-*b*]pyridine, Pd(dba)<sub>2</sub>, CuI, *t*-(Bu)<sub>3</sub>P, K<sub>2</sub>CO<sub>3</sub>, dry THF, 120 °C, 24 h, 63%; (e) R<sub>1</sub>Cl, pyridine, dry DCM, rt, overnight, 39–90%; (f) pyridine-3-sulfonyl chloride, pyridine, dry DCM, rt, overnight, 93%; (g) R<sub>2</sub>Br, Pd(dba)<sub>2</sub>, CuI, *t*-(Bu)<sub>3</sub>P, K<sub>2</sub>CO<sub>3</sub>, dry THF, 120 °C, 24 h, 47–70%.

**Table 1. Enzymatic and Cellular Activities of Compounds 3a–3s**


Compd	R1	R2	B-Raf <sup>V600E</sup> (IC <sub>50</sub> , μM) <sup>a</sup>	Anti-proliferation <sup>b</sup> (IC <sub>50</sub> , μM)	
				Colo205 B-Raf <sup>V600E</sup>	HCT116 B-Raf <sup>WT</sup>
<b>1</b>			0.033	0.309	>10
<b>3a</b>			0.149	2.103	>10
<b>3b</b>			0.177	0.534	>10
<b>3c</b>			0.068	0.758	>10
<b>3d</b>			0.917	6.780	>10
<b>3e</b>			0.593	2.670	>10
<b>3f</b>			0.051	0.445	>10
<b>3g</b>			0.048	1.167	>10
<b>3h</b>			4.31	>10	>10
<b>3i</b>			0.533	1.840	~10
<b>3j</b>			0.017	0.089	1.280
<b>3k</b>			0.122	1.976	~10
<b>3l</b>			0.025	0.148	4.626
<b>3m</b>			0.010	0.061	0.584
<b>3n</b>			0.016	0.200	>10
<b>3o</b>			0.003	0.078	>10
<b>3p</b>			0.131	3.604	>10
<b>3q</b>			0.107	>10	>10
<b>3r</b>			0.021	0.318	>10
<b>3s</b>			0.011	0.166	>10

<sup>a</sup>Kinase activity assays were performed by a FRET-based Z'-Lyte assay. <sup>b</sup>The antiproliferative activities were evaluated using a MTT assay. Data are means of three independent experiments, and the variations are <20%.

also caused a significant loss of potency. Not surprisingly, 1H-pyrrolo[3,4-b]pyridine hinge-binding moiety could be re-

placed by a similar bivalent 1H-pyrrolo[2,3-b]pyridine (**3g**) without obviously affecting the B-Raf<sup>V600E</sup> inhibitory potency. Further investigation revealed that 3-position of the 1H-pyrrolo[3,4-b]pyridine core could be substituted by a methoxyl (**3j**), methyl (**3l**), cyclopropyl (**3m**), or phenyl (**3n**) group to achieve 2–5-fold potency improvement. However, the 3-ethoxyl compound **3k** is 2-times less potent than **3f**. Encouragingly, when the position was substituted by a 4-pyridyl group, the resulting compound **3o** displayed an IC<sub>50</sub> value of 3.0 nM against B-Raf<sup>V600E</sup>, which is 10-times more potent than vemurafenib in a parallel comparison. Compound **3o** also displayed a 4-time stronger antiproliferative activity, with an IC<sub>50</sub> value of 78 nM against B-Raf<sup>V600E</sup> mutated Colo205 cancer cells and favorable selectivity with >10 μM IC<sub>50</sub> value against HCT116 cancer cells with B-Raf<sup>WT</sup>. It was predicted an extra hydrogen bond between N atom of the 4-pyridinyl group and the side chain of Ser536 may contribute to this potency improvement (Figure 1C). The *para*-fluorophenyl analogue **3s** also displayed a slightly increased potency, whereas the *ortho*-Cl or *meta*-Cl substituted compounds **3p** and **3q** are 8- and 7-fold less potent than **3n**, respectively. Thus, **3o** and **3s** were selected as the representatives for further investigations.

The antiproliferative activity of **3o** and **3s** were examined against a panel of cancer cell lines and primary cancer cells derived from New Zealand metastatic melanoma patients harboring differing status of B-Raf.<sup>17,18</sup> It was shown that **3s** displayed similar antiproliferative effects against B-Raf<sup>V600E</sup> mutated cancer cells to that of vemurafenib, whereas compound **3o** is moderately more potent (Table 2). In contrast, none of

**Table 2. Antiproliferative Activities of the Compounds against a Panel of Cancer Cells with Different Status of B-Raf**

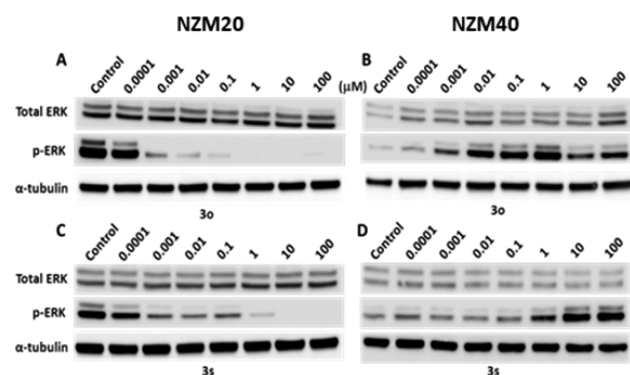
cell lines	antiproliferation (IC <sub>50</sub> , μM)		
	3o	3s	1
Lovo <sup>a</sup>	>10	>10	>10
SK-MEL-2 <sup>a</sup>	>10	>10	>10
NZM09 <sup>a,c</sup>	>10	>10	8.33
NZM40 <sup>a,c</sup>	4.44 ± 1.09	4.35 ± 1.31	3.01 ± 0.31
HT29 <sup>b</sup>	0.263 ± 0.02	0.384 ± 0.07	0.601 ± 0.08
SK-MEL-1 <sup>b</sup>	0.202 ± 0.07	0.448 ± 0.07	1.499 ± 0.04
SK-MEL-28 <sup>b</sup>	0.075 ± 0.03	0.257 ± 0.11	0.381 ± 0.15
A375 <sup>b</sup>	0.046 ± 0.02	0.101 ± 0.05	0.079 ± 0.02
NZM20 <sup>b,c</sup>	0.002 ± 0.001	0.023 ± 0.004	0.024 ± 0.003
NZM07 <sup>b,c</sup>	0.003 ± 0.001	0.024 ± 0.006	0.036 ± 0.008

<sup>a</sup>The cells harbor B-Raf<sup>WT</sup>. <sup>b</sup>The cells express B-Raf<sup>V600E</sup>. <sup>c</sup>Primary cell lines derived from New Zealand metastatic melanoma patients. The IC<sub>50</sub> values were determined using MTT<sup>a,b</sup> or sulforhodamine B<sup>c</sup> assay. Data are mean values ± standard deviation (SD) of three independent experiments.

the compounds showed obvious growth inhibition against cell lines expressing B-Raf<sup>WT</sup>. It is also noteworthy that the patient-sourced melanoma cells with B-Raf<sup>V600E</sup> mutation are more sensitive to the inhibitors. Both **3o** and **3s** inhibit the growth of NZM20 and NZM07 cells with low nanomolar IC<sub>50</sub> values.

The B-Raf inhibition of **3o** and **3s** was further validated by investigating their suppressive abilities on the activation of MAPK signal pathway in NZM20 primary melanoma cancer cells expressing B-Raf<sup>V600E</sup> and the corresponding NZM40 cells with B-Raf<sup>WT</sup> (Figure 2). Both compounds displayed dose-dependent inhibition against the phosphorylation of ERK in



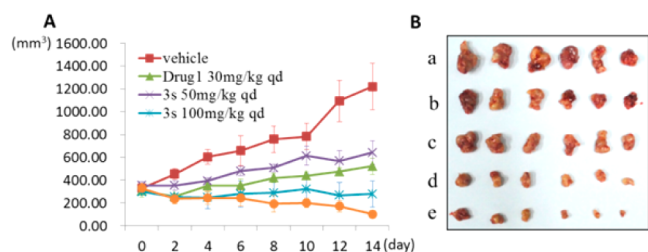


**Figure 2.** Compounds **3o** and **3s** dose-dependently inhibit the activation of ERK in NZM20 cells with B-Raf<sup>V600E</sup> (A,C) but elevate the pERK levels in NZM40 harboring B-Raf<sup>WT</sup> (B,D).

NZM20 cancer cells by Western blot analysis. Alternately, an elevation of p-ERK level was observed for the compounds in NZM40 cells harboring B-Raf<sup>WT</sup> and NRAS<sup>N61H</sup> activating mutation, suggesting that inhibition of B-Raf<sup>WT</sup> in this cell line may result in activation of a compensatory positive feedback loop, leading to increased ERK p44/p42 phosphorylation, as previously observed for vemurafenib, a phenomena that may be related to the appearance of benign skin lesions.<sup>19</sup> The MAPK signal inhibition of the compounds was also validated in a pair of human colorectal cancer cells including B-Raf<sup>V600E</sup> mutated Colo205 cells and HCT-116 cells with B-Raf<sup>WT</sup> (Supporting Information).

Compound **3s** possesses favorable pharmacokinetic profiles with high oral exposures (AUC<sub>(0-∞)</sub> of over 30,000 μg/L·h and F% of 54.0% at a 25 mg/kg oral dose) and acceptable half-life (Supporting Information). It also displayed excellent target specificity in a kinase profiling investigation against 468 kinases with a selectivity score (S<sub>10</sub>) of 0.012 at 1.0 μM, which is about 48-fold of its K<sub>d</sub> value against B-Raf<sup>V600E</sup> (Supporting Information).<sup>20</sup>

The *in vivo* antitumor efficacy of **3s** was further evaluated using a xenograft mouse model of Colo205 cancer cells. The animals were repeatedly administrated with **3s** once or twice daily via oral gavage for 14 consecutive days. Vemurafenib was used as a positive control (Figure 3). Compound **3s** exhibited dose-dependent antitumor efficacy and was well tolerated in all of the tested groups with no mortality or significant loss of body weight (<5% relative to the vehicle-matched controls) observed during treatment (Supporting Information). The



**Figure 3.** (A) Compound **3s** dose-dependently inhibits the growth of Colo205 xenograft tumors following 14 consecutive days of administration. Days postinitiation treatment (d; y-axis) is plotted against the corresponding tumor volume (mm<sup>3</sup>; x-axis). (B) Tumor volumes after the last administration of drugs: a, vehicle; b, vemurafenib (drug 1) 30 mg/kg qd; c, **3s** 50 mg/kg qd; d, **3s** 100 mg/kg qd; e, **3s** 100 mg/kg bid.

tumor growth inhibition (TGI) values are 47.3% and 77.1% at dosages of 50 and 100 mg/kg/day, respectively. More significantly, when the animals were treated with 100 mg/kg of **3s** twice a day, 3 out of 6 mice achieved significant tumor regression, suggesting that sustained exposure of B-Raf inhibitor may be an efficient approach to treat B-Raf<sup>V600E</sup> mutated human colorectal cancer.

In summary, a series of *N*-(3-ethynyl-2,4-difluorophenyl)-sulfonamides were designed and synthesized as new selective B-Raf inhibitors. The compounds potently inhibit B-Raf<sup>V600E</sup> kinase with low nanomolar IC<sub>50</sub> values and selectively suppress the proliferation of a panel of human cancer cell lines with B-Raf<sup>V600E</sup> mutation. One of the most promising compounds **3s** demonstrates favorable pharmacokinetic properties and induces significant tumor regressions in a xenograft mouse model of B-Raf<sup>V600E</sup> mutated Colo205 human colorectal cancer cells without significant sign of toxicity. This compound may serve as a new lead compound for further drug discovery targeting B-Raf<sup>V600E</sup> mutation driven human cancers.

## ■ ASSOCIATED CONTENT

### Supporting Information

Experimental procedures for the syntheses, <sup>1</sup>H NMR and <sup>13</sup>C NMR for final compounds, kinase selectivity, and details of *in vitro* and *in vivo* assays. This material is available free of charge via the Internet at <http://pubs.acs.org>.

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### Notes

The authors declare no competing financial interest.

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