

Discovery of Selective Transforming Growth Factor β Type II Receptor Inhibitors as Antifibrosis Agents

Shohei Miwa, Masahiro Yokota, Yoshifumi Ueyama, Katsuya Maeda, Yosuke Ogoshi, Noriyoshi Seki, Naoki Ogawa, Jun Nishihata, Akihiro Nomura, Tsuyoshi Adachi, Yuki Kitao, Keisuke Nozawa, Tomohiro Ishikawa, Yutaka Ukaji, and Makoto Shiozaki*

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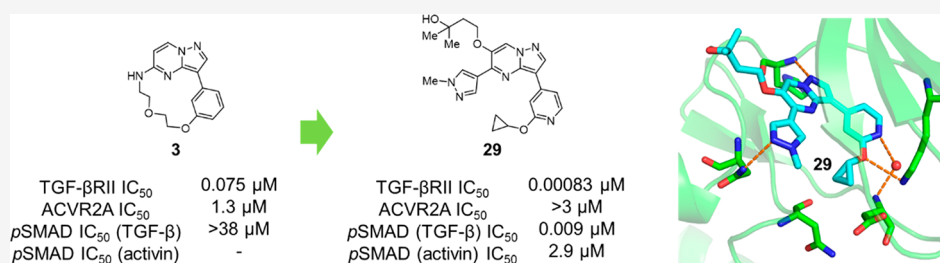
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ABSTRACT: Historically, modulation of transforming growth factor β (TGF- β) signaling has been deemed a rational strategy to treat many disorders, though few successful examples have been reported to date. This difficulty could be partially attributed to the challenges of achieving good specificity over many closely related enzymes that are implicated in distinct phenotypes in organ development and in tissue homeostasis. Recently, fresolimumab and disitertide, two peptidic TGF- β blockers, demonstrated significant therapeutic effects toward human skin fibrosis. Therefore, the selective blockage of TGF- β signaling assures a viable treatment option for fibrotic skin disorders such as systemic sclerosis (SSc). In this report, we disclose selective TGF- β type II receptor (TGF- β RII) inhibitors that exhibited high functional selectivity in cell-based assays. The representative compound **29** attenuated collagen type I alpha 1 chain (*COL1A1*) expression in a mouse fibrosis model, which suggests that selective inhibition of TGF- β RII-dependent signaling could be a new treatment for fibrotic disorders.

KEYWORDS: Selective kinase inhibitor, fibrosis, TGF- β RII, TGF- β signaling pathway, halogen dance rearrangement

Transforming growth factor β (TGF- β) is a pleiotropic cytokine family that comprises highly homologous isoforms TGF- β 1, TGF- β 2, and TGF- β 3. Since these cytokines play crucial roles in a variety of biological processes, aberrant regulation of this TGF- β signaling cascade often results in various pathologies, including cancer and fibrosis.^{1,2} Recently, two peptidic TGF- β inhibitors, fresolimumab and disitertide, were reported to exhibit good preventive effects against human skin fibrosis,^{3,4} suggesting that blocking a common TGF- β signaling pathway could be a viable option for treating fibrotic skin disorders such as systemic sclerosis (SSc).⁵ TGF- β signaling is initiated when the cytokine engages with TGF- β type II receptor (TGF- β RII), a transmembrane serine/threonine receptor kinase, which successively results in its complexation with another serine/threonine receptor kinase, TGF- β type I receptor (TGF- β RI, also known as ALKS). Upon formation of this complex, comprising a set of respective homodimers, two intracellular proteins, SMAD2 and SMAD3, are phosphorylated, leading to the formation of a heterotrimer with SMAD4.⁶ The resultant ternary complex then translocates into the nucleus, and transcription of several key fibrotic genes, such as those encoding collagens and fibronectin, are

subsequently triggered.^{7,8} Because of the complexity of TGF- β signaling, the precise mechanisms and functions of the respective receptors have been poorly understood. Previously, TGF- β RI inhibitors, including SM16 and GW788388, were proven to show antifibrotic effects,^{9–15} but cardiac side effects were also observed,^{16–19} pointing to its potential risk as a target for an antifibrosis agent. In contrast, there have been several selective TGF- β RII ligands. However, little has been clarified about the pharmacological role of TGF- β RII-dependent signals to date.

Herein we report the discovery of novel TGF- β RII inhibitors with superb selectivity over closely related isozymes. One of the best compounds achieved good functional selectivity in

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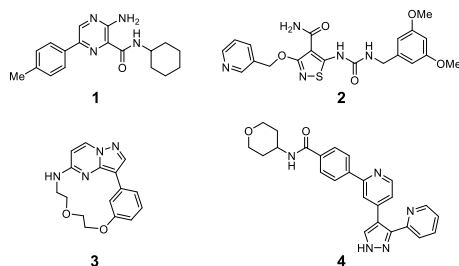
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cell-based assays and was subjected to *in vivo* experiments to understand TGF- β RII-dependent pharmacology.

Historically, there have been several TGF- β RII inhibitors that appeared in preceding reports,^{20–22} and representative compounds disclosed in those publications were initially characterized (Table 1). Among the distinct class of TGF-

Table 1. Preceding TGF- β RII Inhibitors



compd	enzyme IC ₅₀ (μ M) ^{a,b}			cell IC ₅₀ (μ M) ^{c,d}	
	TGF- β RII	ACVR2A	ALK5	TGF- β	activin
1	0.49	0.18	NT ^e	>38	>38
2	0.11	20	NT	10	3.3
3	0.075	1.3	4.4	>38	NT
4	NT	NT	0.018 ^f	0.28	0.52

^aKinase inhibitory activity against each receptor kinase. ^bIC₅₀ values are mean values determined from three replicates. ^cSuppressive effect on SMAD3 phosphorylation after TGF- β or activin stimulation in Expi293F cells. ^dIC₅₀ values are mean values determined from four replicates. ^eNT = not tested. ^fReported value in the literature.¹⁴

β RII inhibitors, compounds **2**²⁰ and **3**²¹ showed decent selectivity over activin receptor type 2A (ACVR2A), which is also a well-recognized receptor of the TGF- β superfamily with serine/threonine receptor kinase activity,²³ while compound **1**²² rarely revealed TGF- β RII selectivity. A significant reduction of cell potency was also notable for both **2** and **3**, but most disappointingly, **2** lost functional selectivity in view of SMAD3 phosphorylation in the cell-based assays (Table 1). Activin-initiated SMAD3 phosphorylation is known to occur through its binding to ACVR2A and subsequent phosphorylation of activin receptor-like kinase 4 (ALK4)/activin receptor-like kinase 7 (ALK7). Therefore, when the kinase inhibitory profiles of compound **2** with respect to TGF- β RII and ACVR2A were taken into consideration, the outcome was

unexpected. Contrastingly, the nonselective nature of compound **4** (GW788388) in the cell assays was quite understandable since this compound was known to inhibit ALK5, TGF- β RII, and ACVR2 (IC₅₀ values against TGF- β RII and ACVR2 were not specified in the literature). These data led us to examine their K_m values for ATP, and it turned out that ACVR2A has roughly 30-times lower affinity toward the natural substrate relative to TGF- β RII ($K_{m,ATP}$ = 9.54 μ M for ACVR2A vs 0.33 μ M for TGF- β RII).²⁴ Although we cannot fully exclude the role of other factors in SMAD3 phosphorylation, we proposed that ACVR2A inhibition had a greater impact on the suppressive effect on SMAD3 phosphorylation and that the several-hundred-fold TGF- β RII selectivity over this isozyme was not sufficient to realize the desired functional selectivity. To this end, we conducted SAR investigations to identify a TGF- β RII inhibitor of great specificity, which ultimately resulted in uncovering a new role of TGF- β RII selective signaling.

Given that **3** displayed the highest ligand efficiency (LE)²⁵ value among the historical TGF- β RII inhibitors **1–3** (0.37, 0.30, and 0.44 for compounds **1**, **2**, and **3**, respectively), we chose this candidate as a lead compound and undertook synthetic explorations (Table 2). First, we prepared compound **5**, a surrogate molecule of **3**, and then examined the SAR of its two aromatic portions. Simple installation of a nitrogen atom at the 4-position of the benzene ring (X = N) boosted the TGF- β RII inhibitory activity over 200-fold, and more gratifyingly, this compound **6** achieved good selectivity over two closely related kinases, ACVR2A and ALK5. In accordance with such enzyme selectivity, we observed moderate functional selectivity in the cell-based assay, and **6** suppressed TGF- β -initiated SMAD3 phosphorylation roughly 3-fold stronger than the phosphorylation triggered by activin initiation. Since this platform suggested further gains in TGF- β RII selectivity, we continued successive efforts, leaving intact the critical methoxypyridine moiety. Examination of the effects of substitution at the 6-position of the pyrazolo[1,5-*a*]pyrimidine ring revealed a rather flat SAR with virtually no deviation of TGF- β RII inhibitory activity, as shown in Table 2 (7–11). With regard to TGF- β RII selectivity, this series of compounds possessed a general tendency to inhibit ACVR2A at levels stronger than ALK5. Therefore, TGF- β RII selectivity would be assured as far as inhibitory activity against ACVR2A is concerned. On the basis of both its cell-based potency and

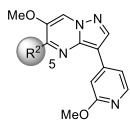
Table 2. Initial SAR Exploration








compd	R ¹	X	enzyme IC ₅₀ (μ M) ^a			cell IC ₅₀ (μ M) ^b	
			TGF- β RII	ACVR2A	ALK5	TGF- β	activin
5	H	CH	1.0	>20	NT ^c	>38	NT
6	H	N	0.0039	1.1	4.1	4.8	14
7	Me	N	0.0029	1.3	2.0	5.4	22
8	vinyl	N	<0.003 (82%)	0.19	1.4	0.38	4.1
9	Et	N	0.0015	0.70	1.9	1.3	11
10	<i>c</i> -Pr	N	0.0037	0.63	8.9	6.9	12
11	MeO	N	0.0010	0.31	0.71	0.33	4.8

^aIC₅₀ values are mean values determined from three replicates. ^bValues of IC₅₀ are mean values determined from four replicates. ^cNT = not tested.

chemical stability, **11** was chosen for further investigation, and the 5-position on the pyrazolo[1,5-*a*]pyrimidine ring was investigated (Table 3). Simple deletion of the methylamino

Table 3. SAR of the 5-Position on the Pyrazolo[1,5-*a*]pyrimidine Ring



Compd	R ²	Enzyme IC ₅₀ (μM) ^d		Cell IC ₅₀ (μM) ^b		MS (%) ^f	
		TGF-βRII	ACVR2A	TGF-β	activin	human	rat
11	NHMe	0.0010	0.31	0.33	4.8	63	74
12	H	0.015	1.2	11	7.6	2	10
13	NMe ₂	0.0079	3.0	0.37	>3.8	16	3
14	Me	0.0050	1.2	4.4	29	44	13
15	Et	0.036	0.98	0.22	>3.8	NT ^d	NT ^d
16	<i>i</i> -Pr	0.0010	3.6	0.12	12	40	14
17		0.0033	2.7	0.23	12	66	92
18		0.0012	2.9	0.14	>38	32	7
19^e		0.0011	1.9	0.026	9.1	29	34
20		<0.0030 (90%)	4.1	0.020	2.2	21	30
21		0.0036	5.3	1.8	15	13	30
22		0.0029	>3.0	1.6	>38	31	62
23		<0.0010 (72%)	0.53	0.054	>38	43	56

^aIC₅₀ values are mean values determined from three replicates. ^bIC₅₀ values are mean values determined from four replicates. ^cMetabolic stability in liver microsomes after incubation for 60 min. ^dNot tested because of insolubility problems. ^eRacemate.

group (**12**) led to a >10-fold reduction of TGF-βRII inhibitory activity, and introduction of a dimethylamino (**13**), methyl (**14**), or ethyl substituent (**15**) could not fully revive the TGF-βRII potency. Meanwhile, a branched alkyl substituent exemplified by an isopropyl group (**16**) conferred potency comparable to that of **11**, although subsequent incorporation of a hydroxyl group into **16** to produce the corresponding tertiary alcohol **17** led to a reduction of the potency. Driven by the result with **16**, compounds possessing a cyclic motif at the 5-position were next synthesized. The enzyme potencies of a

compound having a cyclopentane ring (**18**) and one with a tetrahydrofuran ring (**19**) were similar to each other, having IC₅₀ values comparable to that of **16**. Simultaneously, their functional potencies in the cell assay diverged, with **19** being 5 times more potent than **18** in the TGF-β-initiated SMAD3 phosphorylation assay. Further enlargement at the terminus of the tetrahydrofuran ring (**20**) retained the enzyme potency, even though its enantiomer **21** showed substantially lower potency. This outcome suggested that this region was still encapsulated within the pocket with strict discrimination of the chirality.

Analogues of **20** possessing an aromatic ring at the R² position were also synthesized (**22** and **23**), and **23** reached a subnanomolar value of IC₅₀ in the enzyme assay. Rather disappointingly, the strong TGF-βRII inhibitory activity of **23** did not directly translate into cell-based potency. However, the >700-fold increase of functional selectivity against activin-initiated SMAD3 phosphorylation was quite noteworthy, so the final optimization was carried out with this compound. Among the compounds listed in Table 3, **20** gave a clear cocrystal with the kinase domain of TGF-βRII, and the structural information we obtained facilitated final optimization of this series of compounds. The data pointed out that the nitrogen atom at the 1-position of the pyrazolo[1,5-*a*]pyrimidine ring and the oxygen atom of the tetrahydrofuran made firm hydrogen bonds with His328 and Asn332, respectively, while the methoxypyridine moiety and Lys277 and Asp397 formed a hydrogen-bonding network with the aid of a water molecule. More importantly, there seemed to be sufficient space in the front pocket region along with an additional small cavity near the pyridine ring, so we undertook further investigation of these two sites (Figure 1a).

Driven by such information, we first examined small alkoxy groups as the R³ substituent (Table 4). Unfortunately, no additional gain in the enzyme potency was observed with compounds **24**–**27**, although the cell-based potency was improved, and **26** scored the best cell potency among those analogues of **23**. On the other hand, the selectivity over activin-initiated SMAD3 phosphorylation was diminished, which led us to focus on another aspect of the structure. Since the 6-position of the core ring was present in the front pocket region projecting toward the outside of the pocket, several glycol motifs, one of which anchored to the core part, were examined. To mitigate the potential risk of oxidation and/or hazardous conjugate formation, a dimethyl substituent was simultaneously incorporated within the glycol motifs.

Among analogues of **26** possessing dimethylated glycol pendants (**28**–**30**), compound **29** showed the best functional selectivity for SMAD3 phosphorylation, as indicated by its superb TGF-βRII selectivity over ACVR2A. In order to understand the high selectivity of **29**, a docking model was generated by superimposition of this compound with the cocrystallized ligand **20** (Figure 1a). Two closely related receptor kinases, ACVR2A and ALK5 (roughly 100-fold selectivity over this receptor was confirmed as the result of the kinase panel study shown in the Supporting Information), were chosen as representative off targets, and their crystal structures were successively superimposed with the corresponding residues of TGF-βRII. As demonstrated in Figure 1b, it is hypothesized that both ACVR2A and ALK5 cannot get optimal interactions in two subsites where Asn332 and Cys396 are present in TGF-βRII, since they commonly possess smaller amino acids (Ser and Ala, respectively) in those locations.

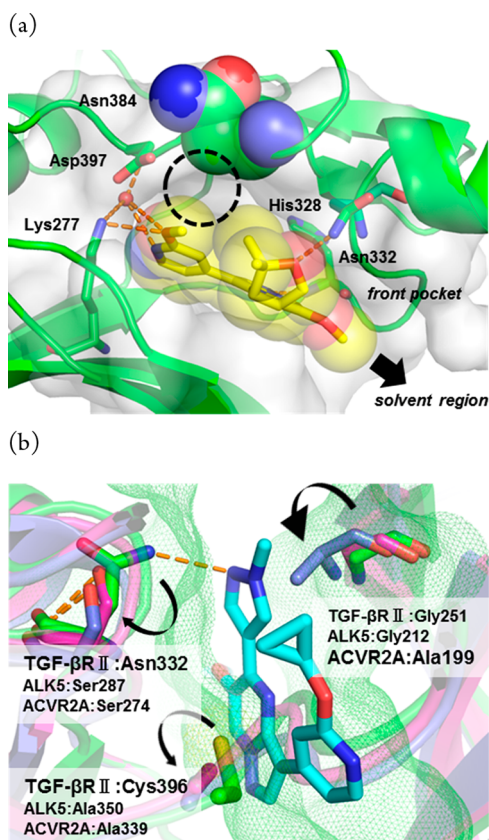


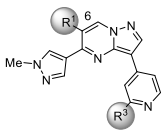
Figure 1. (a) Cocystal structure of **20** in TGF- β R11 (PDB code 7DV6). Hydrogen bonds are depicted as dashed lines (orange), and the water molecule is shown as spheres (red). The CPK representation of **20** is shown to clarify unfulfilled spaces in the binding pocket. The dashed circle indicates a small cavity near the pyridine ring. (b) Docking model of **29** in TGF- β R11 (green; PDB code 7DV6), ALK5 (magenta; PDB code 1VJY), and ACVR2A (slate blue; PDB code 3Q4T). Protein surfaces of TGF- β R11 are depicted as meshed lines. Key amino acids for realizing selectivity are displayed as sticks (green for TGF- β R11, magenta for ALK5, and slate blue for ACVR2A, respectively).

Moreover, Ala199 of ACVR2A could cause a steric clash with the methyl substituent on the pyrazole ring, rationalizing the

significant loss of ACVR2A potency. To this end, a lead compound that was subjected to *in vivo* experiments was selected among the TGF- β R11 inhibitors listed in Table 4. Obviously, high selectivity and good PK profiles are crucial features of the probe molecule to clarify TGF- β R11-dependent signaling *in vivo*. We thus prioritized functional selectivity (TGF- β over activin) in the cell assays with metabolic stability as the criteria for the selection. Among the most potent compounds in the cell (**24**, **26**, **27**, **28**, and **29**), compound **29** stood out for having the highest metabolic stability and best functional selectivity by over 300-fold. As expected from its physicochemical properties, the solubility and caco2 cell permeability of **29** were generally good, which surely reflected favorable PK profiles, fulfilling the good quality of a probe molecule (Figure 2a). The toxicity-related off-target potencies, including hERG (IC_{50} = 22.8 μ M) and CYP (IC_{50} > 10 μ M, CYP3A4, CYP2C8, CYP2D6, CYP1A2, CYP2C19; IC_{50} = 7.8 μ M, CYP2C9), were also sparing (Figure 2b). Therefore, this compound was determined to be a suitable molecule to clarify the role of TGF- β R11-dependent signaling. To assess disease-amelioration effects *in vivo*, the TGF- β -induced mouse fibrosis model was selected, and three doses of **29** were administered orally to the tested subjects (Figure 2c). The skin concentration of collagen type I alpha 1 chain (COL1A1) was monitored as a plasma marker of fibrosis, and significant reduction of the mRNA level was observed in a dose-dependent manner with no sign of cardiac side effects. Moreover, the 10 mg/kg dose groups reached full efficacy, being comparable in effectiveness with the known ALK5 inhibitor (SB525334), indicating that selective TGF- β R11 inhibition could afford a new option to treat skin diseases, including SSC.

Compound **29** was synthesized following the procedures depicted in Scheme 1. *O*-Alkylation of **31** was conducted, and acetal formation to install SEM protection was successfully carried out. Unfortunately, subsequent attempts to directly transform **35** were unsuccessful, and only the undesired regioisomer **34** was obtained. Eventually, base-assisted iodine migration solved the problem, and **35** was prepared effectively by treatment of **34** with Knochel–Hauser base.^{26,27} This compound was further subjected to Suzuki coupling with 1-methylpyrazole-3-borate to afford **36**, which was coupled with

Table 4. Final Optimization of **23**



compd	R ¹	R ³	enzyme IC_{50} (μ M) ^a		cell IC_{50} (μ M) ^b		MS (%) ^c	
			TGF- β R11	ACVR2A	TGF- β	activin	human	rat
23	MeO	MeO	0.00017	0.53	0.054	>38	43	56
24	MeO	<i>i</i> -PrO	0.00058	2.3	0.011	0.43	32	50
25	MeO	<i>t</i> -BuO	0.0070	0.80	0.024	0.92	35	61
26	MeO	<i>c</i> -PrO	0.0016	0.86	0.005	0.39	47	32
27	MeO	<i>c</i> -BuO	0.00061	2.4	0.014	>1.3	NT ^d	NT ^d
28	HOC(Me) ₂ CH ₂ O	<i>c</i> -PrO	0.00031	2.3	0.014	0.87	53	81
29	HOC(Me) ₂ (CH ₂) ₂ O	<i>c</i> -PrO	0.00083	>3.0	0.009	2.9	74	84
30	HOC(Me) ₂ (CH ₂) ₃ O	<i>c</i> -PrO	0.0020	2.9	0.035	0.63	54	64

^a IC_{50} values are mean values determined from three replicates. ^b IC_{50} values are mean values determined from four replicates. ^cMetabolic stability in liver microsomes after incubation for 60 min. ^dNot tested because of insolubility problems.

(a)

Sol.	cLogP	Caco2 Papp	PB human	rat PK			mouse PK		
				iv (0.3mg/kg)	po (1mg/kg)	po (1mg/kg)	AUC	T _{1/2}	F
(μM)	(cm^2/sec)	(%)	MRT (hr)	CL _{tot} (L/hr/kg)	Vd _{ss} (L/kg)	AUC ($\mu\text{M}\cdot\text{hr}$)	T _{1/2} (hr)	F (%)	
>475	2.36	14	99	5.5	0.49	2.7	2.3	4.6	48

(b)

CYP 450 inhibition IC ₅₀ (μM)					
CYP3A4	CYP2C8	CYP2D6	CYP1A2	CYP2C19	CYP2C9
>10	>10	>10	>10	>10	7.8

(c)

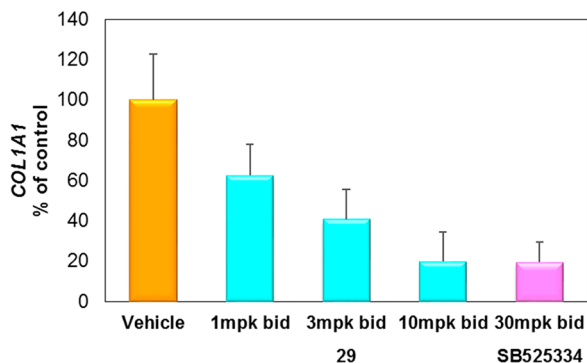
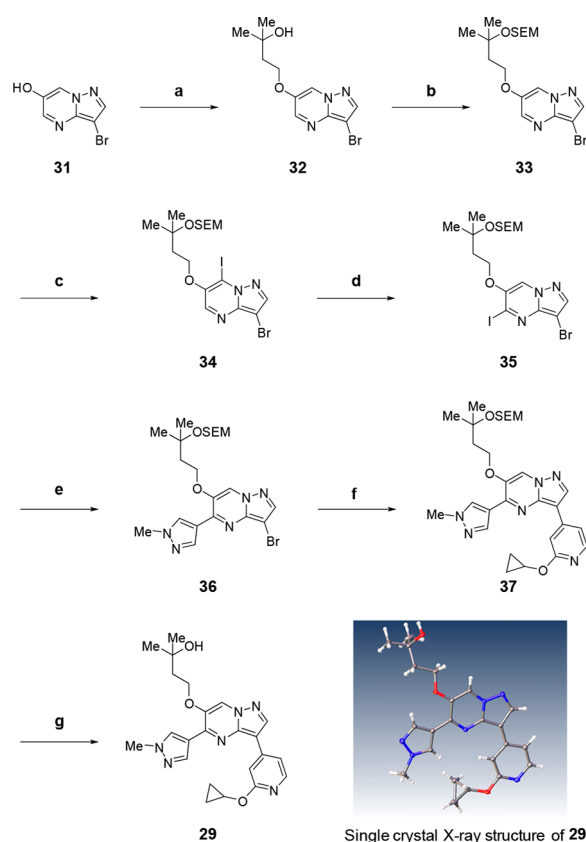


Figure 2. ADME profiles and *in vivo* results for **29**. (a) *In vitro* ADME parameters and PK parameters. (b) CYP inhibition profile. (c) The skin (ear) concentration of collagen type I alpha 1 chain (COL1A1) was monitored as a plasma marker of fibrosis. Oral administration of compound **29** resulted in significant reduction of the mRNA level in a dose-dependent manner. The 10 mg/kg dose groups reached full efficacy, being comparable in effectiveness with the known ALK5 inhibitor SB525334.¹⁰

O-alkylated pyridine-4-borate, followed by final acidic workup to give the target molecule **29**.

In summary, synthetic explorations were carried out to generate novel molecules that selectively inhibited TGF- β RII-dependent signaling. Starting from a less potent TGF- β RII inhibitor with no cell potency (**3**), significant improvement was achieved, and the best compound, **29**, demonstrated a nanomolar IC₅₀ against TGF- β -initiated SMAD3 phosphorylation. More importantly, its 300-fold functional selectivity over the effect on activin-initiated SMAD3 phosphorylation (IC₅₀ = 9 nM for TGF- β vs 2900 nM for activin) was notable. To assess the kinase selectivity, typical kinases were chosen within eight branches of the kinome, and the inhibitory activity of **29** was determined against them. Generally, high selectivity (over roughly 100-fold) was observed, except for DYRK2 (75% inhibition at 0.1 μM), which is not found in SMAD signaling and would have little effect on clarifying TGF- β RII pharmacology.²⁸

The new finding regarding the “halogen dance rearrangement” should also be emphasized since it successfully allowed us to access 3,6-disubstituted-5-iodopyrazolo[1,5-*a*]pyrimidines. The concept of halogen dance rearrangement was originally proposed in 1951,²⁹ but there have been unexpectedly few examples of its use in past drug discovery programs. As a matter of fact, there have been no reports applying such methodology to obtain drug-like molecules with pyrazolo[1,5-*a*]pyrimidine, and the application of this type of reaction could accelerate the discovery of future clinical

Scheme 1. Synthesis of **29**^a

^aReagents and conditions: (a) 4-bromo-2-methylbutan-2-ol, Cs₂CO₃, DMF, r.t.; (b) SEMCl, DIPEA, CH₂Cl₂, r.t., 91% for two steps; (c) (i) TMPMgCl-LiCl, THF, -78 °C, (ii) I₂, THF, -78 to 0 °C, 66%; (d) (i) TMPMgCl-LiCl, THF, -78 °C, (ii) AcOH, THF, -78 to 0 °C, 94%; (e) 1-methyl-4-(4,4,5,5-tetramethyl-1,3,2-dioxaborolan-2-yl)-1H-pyrazole, Pd(PPh₃)₄, K₃PO₄, DME, H₂O, 100 °C, 58%; (f) 2-cyclopropoxy-4-(4,4,5,5-tetramethyl-1,3,2-dioxaborolan-2-yl)pyridine, RuPhos Pd G2, K₃PO₄, DME, H₂O, 100 °C, 26%; (g) TFA, CH₂Cl₂, r.t., 40%.

candidates since heteroaromatics, including pyrazolo[1,5-*a*]pyrimidine, are found within many bioactive molecules. The TGF- β signaling pathway has had a long history as the site of promising targets typified in cancer therapy. Meanwhile, a lack of selective ligands has hampered a clear understanding of the respective roles of TGF- β receptors with complex phenotypes in cells and *in vivo*. For instance, SB525334, known as a pan-inhibitor of ALK4 and ALK5,¹⁰ demonstrated similar antifibrotic effects regardless of different fibrotic initiators, while TGF- β and bleomycin, though selective TGF- β RII inhibitors, were found to show reduced potency in the bleomycin-induced model (data not shown). As far as we know, this is the first example to demonstrate small-molecule-based inhibition of TGF- β RII signaling leading to an *in vivo* antifibrotic effect. It is known that receptors of the TGF- β superfamily work modularly and mediate a variety of signals by forming distinct oligomers. Because of this complex nature, downstream signaling of each receptor is so diverse and hardly interpretable. Suffice it to say that **29** is an ideal molecule, conferring subnanomolar enzyme inhibition, excellent cell potency, and functional selectivity. Although further understanding of selective TGF- β RII inhibition, especially at the gene regulation level, is beyond the scope of this study, the

discovery of selective ligands for each TGF- β receptor symbolized by this compound will definitely accelerate future clarification of the TGF- β superfamily signaling pathway.

■ ASSOCIATED CONTENT

SI Supporting Information

The Supporting Information is available free of charge at <https://pubs.acs.org/doi/10.1021/acsmchemlett.0c00679>.

Synthetic schemes, procedures, experimental data, biological protocols, and crystallography details (PDF)

■ AUTHOR INFORMATION

Corresponding Author

Makoto Shiozaki – Central Pharmaceutical Research Institute, Japan Tobacco Inc., Takatsuki, Osaka 569-1125, Japan; orcid.org/0000-0001-5519-6547; Phone: +81-72-681-9700; Email: makoto.shiozaki@jt.com

Authors

Shohei Miwa – Central Pharmaceutical Research Institute, Japan Tobacco Inc., Takatsuki, Osaka 569-1125, Japan; Division of Material Sciences, Graduate School of Natural Science and Technology, Kanazawa University, Kakuma, Kanazawa 920-1192, Japan

Masahiro Yokota – Central Pharmaceutical Research Institute, Japan Tobacco Inc., Takatsuki, Osaka 569-1125, Japan

Yoshifumi Ueyama – Central Pharmaceutical Research Institute, Japan Tobacco Inc., Takatsuki, Osaka 569-1125, Japan

Katsuya Maeda – Central Pharmaceutical Research Institute, Japan Tobacco Inc., Takatsuki, Osaka 569-1125, Japan

Yosuke Ogoshi – Central Pharmaceutical Research Institute, Japan Tobacco Inc., Takatsuki, Osaka 569-1125, Japan; orcid.org/0000-0002-1440-6701

Noriyoshi Seki – Central Pharmaceutical Research Institute, Japan Tobacco Inc., Takatsuki, Osaka 569-1125, Japan

Naoki Ogawa – Central Pharmaceutical Research Institute, Japan Tobacco Inc., Takatsuki, Osaka 569-1125, Japan

Jun Nishihata – Central Pharmaceutical Research Institute, Japan Tobacco Inc., Takatsuki, Osaka 569-1125, Japan

Akihiro Nomura – Central Pharmaceutical Research Institute, Japan Tobacco Inc., Takatsuki, Osaka 569-1125, Japan

Tsuyoshi Adachi – Central Pharmaceutical Research Institute, Japan Tobacco Inc., Takatsuki, Osaka 569-1125, Japan

Yuki Kitao – Central Pharmaceutical Research Institute, Japan Tobacco Inc., Takatsuki, Osaka 569-1125, Japan

Keisuke Nozawa – Central Pharmaceutical Research Institute, Japan Tobacco Inc., Takatsuki, Osaka 569-1125, Japan

Tomohiro Ishikawa – Central Pharmaceutical Research Institute, Japan Tobacco Inc., Takatsuki, Osaka 569-1125, Japan

Yutaka Ukaji – Division of Material Sciences, Graduate School of Natural Science and Technology, Kanazawa University, Kakuma, Kanazawa 920-1192, Japan; orcid.org/0000-0003-3185-3113

Complete contact information is available at: <https://pubs.acs.org/doi/10.1021/acsmchemlett.0c00679>

Notes

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■ ABBREVIATIONS

CYP, cytochrome P450; DIPEA, *N,N*-diisopropylethylamine; hERG, human ether-a-go-go-related gene; PK, pharmacokinetics; SAR, structure–activity relationship; SEM, 2-(trimethylsilyl)ethoxymethyl; SMAD, mothers against decapentaplegic homologue; RuPhos Pd G2, chloro(2-dicyclohexylphosphino-2',6'-diisopropoxy-1,1'-biphenyl)[2-(2'-amino-1,1'-biphenyl)]palladium(II); TFA, trifluoroacetic acid; TMP, tetramethylpiperidinyl

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