pubs.acs.org/jmc

Development of a 1,2,4-Triazole-Based Lead Tankyrase Inhibitor: Part II

Ruben G. G. Leenders,[#] Shoshy Alam Brinch,[#] Sven T. Sowa, Enya Amundsen-Isaksen, Albert Galera-Prat, Sudarshan Murthy, Sjoerd Aertssen, Johannes N. Smits, Piotr Nieczypor, Eddy Damen, Anita Wegert, Marc Nazaré, Lari Lehtiö, Jo Waaler,[#] and Stefan Krauss^{*,#}



showed picomolar IC₅₀ inhibition in a cellular (HEK293) WNT/ β -catenin signaling reporter assay, no off-target liabilities, overall favorable absorption, distribution, metabolism, and excretion (ADME) properties, and an improved pharmacokinetic profile in mice. Moreover, treatment with compound **24** induced dose-dependent biomarker engagement and reduced cell growth in the colon cancer cell line COLO 320DM.

■ INTRODUCTION

Tankyrase 1 and tankyrase 2 (TNKS1/2) are members of the poly(ADP-ribose) polymerase (PARP) family of enzymes that regulate the turnover of specific target proteins through covalently linking the cellular redox metabolite NAD⁺ to target proteins in a process called poly-ADP-ribosylation (PARylation). The PAR chain produced in this post-translational modification is subsequently recognized by the E3 ubiquitin ligase ring finger protein 146 (RNF146) leading to polyubiquitination of the PARylated target proteins followed by proteasomal degradation.¹⁻⁴ Independent of their catalytic activity, TNKS1/2 also provides scaffolding functions that are important in the formation of protein complexes.^{5–8} TNKS1/2 PARylate a plethora of target proteins including peroxisome proliferator-activated receptor-gamma coactivator 1 α (PGC- 1α), telomeric repeat binding factor 1 (TRF1), phosphatase and tensin homologue (PTEN), AMP-activated protein kinase (AMPK), SRY-box transcription factor 9 (SOX9), and SH3 domain binding protein 2 (SH3BP2).^{3,9–15} In particular, TNKS1/2 regulate the turnover of AXIN1, AXIN2 (AXIN1/ 2) and of angiomotin (AMOT) proteins at the crossroad of the elementary wingless-type mammary tumor virus integration site $(WNT)/\beta$ -catenin and Hippo signaling pathways, respectively.^{3,14,16,17} Hence, controlling the catalytic activity of tankyrase by pharmacological intervention provides an attractive tool for reducing WNT/ $\beta\text{-}catenin$ and Hippo signaling. 18,19

Multiple potent TNKS1/2 inhibiting small molecules, including bicyclic lactams, compounds based on heteroalicyclic amide scaffolds, tricyclic fused ring systems and 1,2,4-triazole scaffolds, have been identified. These compounds block the catalytic domain of TNKS1/2, either by binding with high selectivity to the adenosine binding pocket (which differs from other members of the PARP family) or by binding to the more conserved nicotinamide pocket. The latter binding mode results in a lower selectivity across the PARP family. Other compounds target both pockets in the catalytic domain. ^{16,20–37} Inhibitors based on the 1,2,4-triazole scaffold such as JW74, ³⁸ G007-LK, ²¹ OD336, ²⁸ and OM-1700 (1)³⁹ target the adenosine binding pocket of the TNKS1/2 catalytic domain with high selectivity and are therefore able to display selectivity over other members of the PARP family.

Despite the significant progress in developing TNKS1/2 inhibitors, there is currently no viable TNKS1/2 specific

Received: July 15, 2021 Published: December 8, 2021



🚾 😳 💽

Article



inhibitor in clinical practice for any disease indication. Concerns that have hampered clinical trials of TNKS1/2 inhibitors include earlier reports indicating intestinal toxicity^{40,41} and bone loss in mouse models,¹³ although other studies show a beneficial effect of tankyrase inhibition on fracture healing.⁴² Nevertheless, mice that have been treated for an extended time with a moderate dose of a tankyrase inhibitor have not shown visible adverse effects.⁴³ Ongoing studies suggest that it is possible to overcome a potential biotarget toxicity and recently the tankyrase inhibitors E7449 (a dual inhibitor of PARP1/2 and TNKS1/2) and STP1002 have entered clinical trials with dose escalation studies in patients with advanced solid tumors.^{44,45} This clearly illustrates the potential of TNKS1/2 inhibitors and justifies the development of drugs directed toward TNKS1/2 inhibition with high potency and an optimized pharmacokinetic (PK) profile.

Here, we further optimize compounds based on the 1,2,4-triazole series as landmark compounds 21,28,38,39 and we present compounds reaching picomolar IC₅₀ values in a cellular WNT/ β -catenin signaling reporter assay, while also showing optimized absorption, distribution, metabolism, excretion (ADME), and pharmacokinetic (PK) properties.

RESULTS AND DISCUSSION

Compound Design Strategy. Starting from the previously described lead compound 1 (Figure 1)³⁹ and having

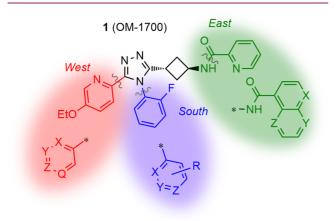


Figure 1. Lead compound 1 (OM-1700) and main modifications.

developed synthetic methodologies for preparing compound iterations in a modular fashion (see Supporting Experimentals for the general scheme of synthesis), we aimed at further improving the potency of tankyrase inhibition and optimizing ADME and pharmacokinetic properties of 1. Optimization was motivated by the highly potent compounds discovered during the development of 1 (e.g., compounds 16, 21, and 27 as numbered in our preceding paper,³⁹ see Chart S-1). In addition to this, the pharmacokinetic properties of 1, primarily its half-life and exposure, necessitated further improvements.

In the process of making next-generation derivatives based on these previous compounds,³⁹ we revisited South-, East-, and West-single-point mutations and hybrids thereof. From these structures, we selected six compounds based on their cellular activity and on the diversity of their molecular architecture. The properties of these six shortlisted compounds were evaluated in mouse peroral pharmacokinetic studies and compared with that of compound **1**. The best compound of this set was selected and further evaluated with respect to early ADME properties, off-target effects, binding affinity in the catalytic pocket of the TNKS2 protein, as well as inhibition of WNT/ β -catenin signaling and proliferation in the colon cancer cell line COLO 320DM. The results from these experiments culminated in the identification of a new 1,2,4-triazole based lead tankyrase inhibitor.

Structure–Activity Relationship (SAR) Investigation and Biological Evaluation. To further explore the SAR of the West- and South-regions of 1, single-point modifications departing from 1 were synthesized (Table 1). From these compounds, we concluded that a gain in potency in our *in vitro* TNKS2 assay and cell-based WNT/ β -catenin signaling assay cannot be attained with compounds possessing the East 2pyridyl group while changing South/West groups, and we therefore focused on compounds with annulated aromatic heterocycles such as naphthyridines and quinoxalines⁴⁶ (Table 2). Such variations enabled picomolar potencies as these moieties are able to form a significantly more efficient π – π stacking interaction with His1048 and an additional hydrophobic interaction with Phe1035 as evidenced by the obtained co-crystal structures.³⁹

First, we explored a series based on the East-side 1,5naphthyridine, either with or without a 3-fluoro substituent (Table 2). Since compound 15a was impaired by a high metabolic instability in mouse microsomes, we aimed at making the structure metabolically more stable by increasing the polarity with the introduction of a South-pyridylsubstitution and compounds 16-18 (Table 2) were prepared. In general, more polar compounds are expected to show greater metabolic stability. Compounds with such a Southpyridyl, however, proved to be less potent in the cellular WNT/ β -catenin signaling reporter assay compared to 15a and only compound 18b with an IC₅₀ value of 6 nM was selected for further characterization (Table 2). To improve potency of compounds **16–18** as earlier observed, a bicyclo[1.1.1]pentane core as an alternative to the cyclobutane influenced activity favorably (see, e.g., compound 19 as numbered in our preceding paper³⁹). Unfortunately, here, compounds built on this bicyclo[1.1.1]pentane core lacked picomolar cellular activity when combined with a South-pyridyl moiety (compounds 19-21, Table 2) and these South-pyridyl structures were abandoned. Furthermore, a South-methyl thiophene group was introduced as a phenyl bio-isostere47 (compounds 22 and 23, Table 2), resulting in potent inhibitors of which compound 22a was selected for further evaluation.

Finally, the introduction of an East quinoxaline moiety instead of the naphthyridine group (compound 24) again led to picomolar cellular activity ($IC_{50} = 0.63$ nM), and we prepared further variations based on a quinoxaline core to interrogate this part of the pharmacophore. Variations of lead 24 by four fluorine- and methyl-substituted quinoxalines showed similar inhibitory activity (compounds 25–28, Table 3) but by varying the West-moiety, highly efficacious inhibitors could be obtained with the quinoxaline East-group. Three of these compounds were selected for further profiling (compounds 30b, 31a, and 31b). This resulted in a short list of six inhibitors with relatively high structural diversity (Figure 2).

Next, the set of six shortlisted compounds (Figure 2) was subjected to mouse peroral pharmacokinetic (PK) analyses and compared with 1 (Table 4). Here, compounds 22a and 18b displayed a decreased exposure (area under the curve, AUC)

Table 1. West Variations of 1 (OM-1700)^a

ID	Structure		IC ₅₀ TNKS2 (nM)	IC ₅₀ HEK293 (nM)
1 ³⁹			14	19
2 ³⁹	N - Vate	F	29	127
3 ³⁹	Q. S. N. J. X.	F	9.2	12
4 ³⁹	N X	F	22	33
5 ³⁹	LN 24	F	16	37
6 ³⁹	F2HCO	F	8.8	36
7	A	F	8.6 (5.8-13)	49
8	Loly	F	12 (8.5-16)	110
9		F	220 (160-290)	789
10		F	93 (28-310)	355
11	Eto		5.2 (2.5-11)	65
12	Eto		13 (6.8-24)	83
13	Eto	T's	4.6 (3.4-6.4)	37
14	Eto	S CI	12 (4.9-28)	27

^aThe IC₅₀ values of the compounds of this work were determined with both the TNKS2 biochemical assay (quadruplicates used for each concentration tested, 95% confidence intervals are given in parentheses) and the cellular (HEK293) WNT/ β -catenin signaling reporter assay (triplicates used for each concentration tested, T/F-tests were performed for the IC₅₀ curve fitting; all p > 0.95).

while compound **31a** showed an AUC similar to **1**. In contrast, compound **30b** showed a high AUC, however, with a small volume of distribution (0.33 L/kg), and consequently, none of these compounds were selected for further evaluation.

Further, compounds 31b and 24 showed improved pharmacokinetic properties compared to 1 (Table 4). The AUCs for 31b and 24 were similar and about twice as high compared to the AUC for 1. The peak value (C_{max}) of exposure of 24 was approximately 3 times lower compared to the peak value for compound 31b. In addition, 24 showed a more favorable volume of distribution and solubility compared

to these data for compound **31b**. Compared to **1**, compound **24** possesses an AUC about twice as high and a peak value of about 60% of that of **1** resulting in a higher exposure with a lower peak level. In addition to this, efficacy in the cellular WNT/ β -catenin signaling reporter assay was approximately 30 times higher while the clearance of **24** was about half of that of **1**.

In summary, based on the overall improved mouse PK profile and the enhanced potency in the cellular WNT/ β -catenin signaling reporter assay, compound **24** was designated as the best derivative.

Table 2. East Naphthyridines and Quinoxalines⁴

ID	Structure			IC ₅₀ TNKS2 (nM)	IC ₅₀ HEK293 (nM)
15a (R=H) ³⁹ 15b (R=F) ³⁹	Eto	N ⁻ N L N F		4.32	0.63 0.17
16a (R=H) 16b (R=F)	E.			5.8 (3.5-9.5) 8.5 (5.1-14)	36 8.7
17a (R=H) 17b (R=F)	Ę,			4.5 (1.9-11) 5.8 (1.3-26)	16 9.1
18a (R=H) 18b* (R=F)	E.			5.4 (3.4-8.7) 2.9 (1.9-4.6)	13 6.4
19 (R=H)	E	₩		13 (6.8-25)	20
20 (R=H)	E.	₩		9.6 (3.9-24)	30
21 (R=H)	E.	⊢令⊣		9.4 (4.0-22)	43
22a* (R=H) 22b (R=F)	For			0.73 (0.41-1.3)	0.81 1.0
23	Fe		-X-N	2.4 (1.6-3.6)	1.0
24 * (OM-153)	F			2.0#	0.63#

^{*a*}The IC₅₀ values of the compounds of this work were determined with both the TNKS2 biochemical assay (quadruplicates used for each concentration tested, 95% confidence intervals are given in parentheses) and the cellular (HEK293) WNT/ β -catenin signaling reporter assay (triplicates used for each concentration tested, T/F-tests were performed for the IC₅₀ curve fitting; all p > 0.95). * indicates a shortlisted compound. [#] averages of multiple independent measurements; standard error of the means (SEMs) are shown in Table 5.

Subsequently, compound 24 was further characterized and compared to benchmark structure 1 with respect to early ADME properties and off-target effects. Tested ADME parameters such as kinetic solubility, Caco-2 permeability and efflux, microsomal stabilities, mouse plasma stability, and mouse plasma protein binding were found to be in line with 1, while solubility was somewhat lower although in an acceptable range (Table 5).

Next, we solved a co-crystal structure of TNKS2-24 demonstrating that 24 binds to the NAD⁺ cleft of the catalytic domain in a similar manner to the previously reported analogue 1^{39} (Figure 3). The observed electron density is clear for the compound except for an apparently mobile ethoxy group extending toward the nicotinamide site. The previously identified hydrogen bonds of the scaffold³⁹ were once more observed between 24 and the Tyr1060 and Asp1045 backbones, as well as with a water molecule. The quinoxaline moiety is forming $\pi - \pi$ stacking interactions with both His1048 and Phe1035 providing improved binding affinity compared with that of 1.

Finally, to complete the analysis of compound 24, COLO 320DM cells, likely of neuroendocrinal origin,⁴⁸ were treated with various doses of 24 to evaluate the efficacy in reducing canonical WNT/ β -catenin signaling and the potential as an antiproliferative agent in this cancer cell line. Previously, we have shown that tankyrase inhibition can block WNT/ β catenin signaling and attenuate proliferation and viability in cancer cell lines in vitro and in vivo, including COLO 320DM that is commonly used for testing tankyrase inhibitors.^{28,39,49,50} Treatment of COLO 320DM cells with 24 decreased viability with a GI₅₀ value of 10.1 nM and a GI₂₅ value of 2.5 nM (for 1, these values were 650 and 94 nM, respectively³⁹), while APC^{wild-type} RKO cells as a control colon cancer line were only modestly affected by treatment with compound 24 (Figure 4a). As previously observed upon tankyrase inhibition,^{49,51} treatment with 24 dose-dependently either increased or decreased the TNKS1/2 protein levels (Figure 4b). Compound 24 also stabilized AXIN1 and AXIN2 proteins and reduced the level of transcriptionally active β -catenin (nonphosphorylated) in both cytoplasmic and nuclear fractions (Figure 4b). In addition, real-time qRT-PCR analyses revealed

Table 3. East Quinoxaline Variations^a

ID	Structure		IC ₅₀ TNKS2 (nM)	IC ₅₀ HEK293 (nM)
24 * (OM-153) (R ¹ , R ² , R ³ = H)	ر Na		1.6#	0.63#
	EtO	$R^2 R^3$		
25	R1=	=F, R ² =H, R ³ =H	1.5 (1.0-2.2)	1.5
26	R ¹	=H, R ² =Me, R ³ =H	1.5 (0.74-3.2)	2.2
27	R ¹ :	=H, R²=H, R³=Me	1.1 (0.46-2.5)	2.4
28	R1:	=H, R²=Me, R³=Me	2.6 (1.4-4.8)	4.3
29a	[N]X	R ¹ =H, R ² =H, R ³ =H	1.9 (1.2-2.9)	1.2
29b	F ₂ HCO	R ¹ =F, R ² =H, R ³ =H	2.4 (1.6-3.5)	0.51
30a	IN X	R ¹ =H, R ² =H, R ³ =H	1.6 (0.87-3.0)	0.29
30b*		R ¹ =F, R ² =H, R ³ =H	2.0 (1.0-3.7)	0.80
31a*	TN J	R ¹ =H, R ² =H, R ³ =H	1.7 (1.1-2.7)	1.5
31b*		R ¹ =F, R ² =H, R ³ =H	1.6 (0.97-2.7)	0.17

^{*a*}The IC₅₀ values of the compounds of this work were determined with both the TNKS2 biochemical assay (quadruplicates used for each concentration tested, 95% confidence intervals are given in parentheses) and the cellular (HEK293) WNT/ β -catenin signaling reporter assay (triplicates used for each concentration tested, T/F-tests were performed for the IC₅₀ curve fitting; all p > 0.95). * indicates shortlisted compound. # averages of multiple independent measurements; SEMs are shown in Table 5.

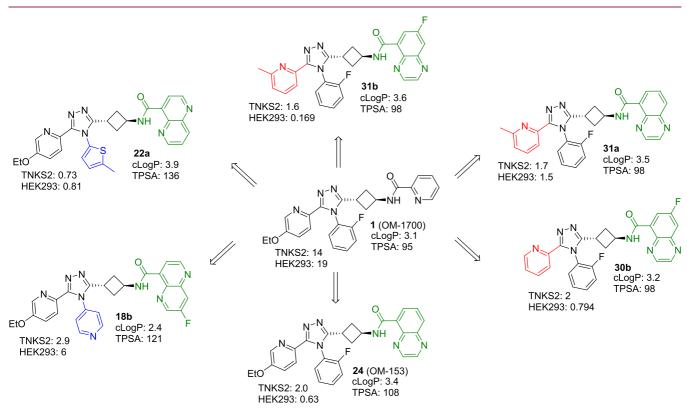


Figure 2. Short list of six compounds and 1 including their respective biochemical TNKS2 and cellular (HEK293) WNT/ β -catenin signaling reporter assays IC₅₀ values in nM. clog *P* and tPSA (in Å²) as calculated by DataWarrior v5.5.0. Moieties in color were different from 1.

reduced levels of transcripts of the WNT/ β -catenin signaling target genes *AXIN2, DKK1, NKD1,* and *APCDD1* in a dosedependent manner (Figure 4c). Collectively, these results showed that **24** both potently and specifically can inhibit

WNT/ β -catenin signaling activity and block proliferation in

COLO 320DM cells.

pubs.acs.org/jmc

Table 4. Mouse PO PK 5 mg/kg and Kinetic Solubility Data of 1 and the Six Shortlisted Compounds

compound	$\stackrel{t_{1/2}}{(\mathrm{h})}$	$t_{ m max}$ (h)	C _{max} (ng/mL)	AUC $0 \rightarrow t$ (ng/mL·h)	$\begin{array}{c} \text{AUC } 0 \rightarrow \infty \\ (\text{ng/mL} \cdot \text{h}) \end{array}$	$\begin{array}{c} \text{MRT } 0 \to \infty \\ (h) \end{array}$	V _d (L/kg)	CL (L/h/kg)	solubility (µM)
1 (OM-1700) ³⁹	0.67	0.25	3203	2384	2388	0.69	2.03	2.09	>80
18b	1.00	0.25	285	319	322	1.56	22.4	15.6	>80
22a	1.17	0.25	779	543	547	1.39	15.5	9.14	50
24 (OM-153)	1.50	0.5	1967	4945	5038	2.39	2.15	0.99	31
30b	0.69	0.5	6512	15 083	15 105	1.93	0.33	0.33	>80
31a	0.76	0.25	2770	3401	3404	1.24	1.61	1.47	>80
31b	0.59	0.25	5796	5313	5316	1.47	0.80	0.94	13

Table 5. Profiling of Compound 24

parameter	1 (OM-1700)	24 (OM-153)
efficacy		
TNKS1 $(IC_{50}, nM (pIC_{50} \pm SEM))^{a}$	$127 (6.90 \pm 0.05)$	$13 (7.90 \pm 0.054)$
TNKS2 $(IC_{50}, nM (pIC_{50} \pm SEM))^{a}$	$14 (7.85 \pm 0.04)$	$2.0 (8.71 \pm 0.069)$
HEK293 reporter assay (IC ₅₀ , nM (pIC ₅₀ ± SEM))	$19 (7.75 \pm 0.067)$	$0.63 \ (9.22 \pm 0.037)$
COLO 320DM/RKO cells (GI ₅₀ , nM)	650/>10 000	10/>10 000
ADME		
kinetic solubility PBS pH = 7 (μ M)	>80	31
Caco-2 A–B: P_{app} (10 ⁻⁶ cm/s)	39.5	40.5
Caco-2 efflux ratio	0.61	0.64
microsomal stability human/mouse/dog $ ext{CL}_{ ext{int}}$ ($\mu ext{L/min/mg}$ protein)	<5/27/nd	18/22/3.8
mouse plasma stability $t_{1/2}$ (min)	>120	>120
mouse PPB (%)	93.92	98.58
off-target		
PARPs ^b PARP1/2/3/4/10/12/14/15 (IC ₅₀ , µM)	>10	>10
hERG inhibition (IC ₅₀ , μ M)	>25	>25
Ames test	nongenotoxic	nongenotoxic
CYP3A4 inhibition $(IC_{50}, \mu M)$	>25	>25
CYP induction (human PXR)	nd	nonactivator ^{bc}
Cerep Safety panel 44 targets@10 μ M (inhibition)	clean, (A2A, 53%)	clean, (all <50%)
mouse pharmacokinetics		
PO PK mouse $t_{1/2}$ (h)	0.67	1.5
PO PK mouse C_{max} (ng/mL)	3202	1967
PO PK mouse CL (L/h/kg)	2.09	0.99
PO PK mouse V _d (L/kg)	2.03	2.15
PO PK mouse AUC $0 \rightarrow t (ng/mL)$	2384	4945
calculated properties ^d		
MW (g/mol)	458.5	509.6
clog P	3.1	3.4
tPSA	95	108

^aSee Figure 1, Supporting Information. ^bSee Table 1, Supporting Information. ^cHighest concentration, 100 μ M. ^dCalculated by DataWarrior v5.5.0.

CONCLUSIONS

In this further development of our structure-guided lead optimization program of 1,2,4-triazole-based tankyrase inhibitors, we have extensively improved previous lead compound 1 to lead candidate compound 24.⁵² We showed that compound 24 possesses improved binding affinity in the catalytic pocket of the TNKS2 protein with concurrently modest selectivity over TNKS1, picomolar IC₅₀ activity in the cellular WNT/ β -catenin signaling reporter assay, a clean off-target safety profile, good ADME properties, an optimized mouse PK profile, and potent inhibition of WNT/ β -catenin signaling and proliferation in COLO 320DM. These results justify testing compound 24 in a pharmacodynamics setting as well as in toxicity models (publication pending).

EXPERIMENTAL SECTION

General Methods. NMR spectra were recorded on a 400 MHz spectrometer with tetramethylsilane as internal standards. Coupling constants are given in hertz. Peaks are reported as singlet (s), doublet (d), triplet (t), quartet (q), quintet (p), sextet (h), septet (hept), multiplet (m), or a combination thereof; br stands for broad.

Liquid chromatography/mass spectroscopy (LC/MS) chromatograms mass spectra were recorded using electrospray ionization (ESI) in positive or negative ionization mode on Agilent 1260 Bin: pump, G1312B, degasser; autosampler; ColCom; DAD G1315C; MSD G6130B ESI; eluent A, acetonitrile; eluent B, 10 mM ammonium bicarbonate in water (base mode) or 0.1% formic acid in water (acid mode). High-resolution mass spectra (HRMS) were recorded with an LC-MS Q Exactive Focus high-resolution mass spectrometer (Thermo Scientific). Calibration was done with the Pierce calibration solutions containing 1-butylamine, caffeine, MRFA, and Ultramark 1621 (positive mode) and the Pierce calibration solution containing sodium dodecyl sulfate, sodium taurocholate, and Ultramark 1621

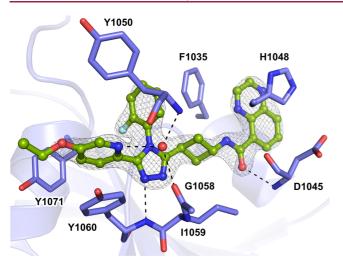


Figure 3. Co-crystal structure of TNKS2 with **24** (PDB 706X). The protein is shown in blue, and **24** in green. The dashed lines in black represent hydrogen bonds, and the red spheres represent water molecules. The σ_A weighted $2F_o - F_c$ electron density maps around the ligands are contoured at 1.8 σ .

(negative mode). Analysis: 1 μ L of a 10 μ g/mL sample in MeCN/ DMSO 99:1 is injected and data are acquired under full MS mode (resolution 70 000 FWHM at 200 Da) over the mass range m/z of pubs.acs.org/jmc

150–2000. Standard ESI conditions compatible with the flow rate are applied: spray voltage, 3.5 kV; auxiliary gas heater temperature, 463 °C; capillary temperature, 280 °C; sheath gas, 58; auxiliary gas, 16; sweep gas, 3; S-lens radio frequency (RF) level, 50. Mass scan range is 150–2000 m/z. Mass resolution is set at 70 000 (<3 ppm mass accuracy). Data are evaluated using Xcalibur Qual Browser version 4.2.47 (Thermo Fisher).

All test compounds were >95% pure by LC/MS and ¹H NMR analyses. All spectra as well as preparation of the intermediates and general procedures are given in the Supporting Experimentals.

N-(*trans*-3-(5-(5-Cyclopropoxypyridin-2-yl)-4-(2-fluorophenyl)-4H-1,2,4-triazol-3-yl)cyclobutyl)picolinamide (**7**). The title compound was prepared according to general procedure F as a white solid (24.3 mg, 75%). LC/MS (ESI) *m*/*z* for C₂₆H₂₃N₆O₂F 470 (calculated) 471 ([M + H]⁺, found). ¹H NMR (400 MHz, CDCl₃) δ 8.56-8.49 (m, 1H), 8.26-8.13 (m, 3H), 7.99 (s, 1H), 7.83 (td, *J* = 7.7, 1.7 Hz, 1H), 7.50-7.37 (m, 3H), 7.24-7.14 (m, 3H), 4.82-4.71 (m, 1H), 3.74 (tt, *J* = 6.1, 3.1 Hz, 1H), 3.46 (tt, *J* = 10.3, 5.3 Hz, 1H), 3.11-2.96 (m, 2H), 2.52-2.35 (m, 2H), 0.84-0.71 (m, 4H). HRMS *m*/*z* [M + H]⁺: 471.19393 (calculated), 471.1929 (found), Δ = -2.28 ppm.

N-(*trans*-3-(4-(2-*Fluorophenyl*)-5-(5-*isopropoxypyridin*-2-*yl*)-4H-1,2,4-*triazol*-3-*yl*)*cyclobutyl*)*picolinamide* (8). The title compound was prepared according to general procedure F as a white solid (18.3 mg, 77%). LC/MS (ESI) *m*/*z* for $C_{26}H_{25}N_6O_2F$ 472 (calculated) 473 ([M + H]⁺, found). ¹H NMR (400 MHz, CDCl₃) δ 8.57–8.48 (m, 1H), 8.21 (d, J = 6.9 Hz, 1H), 8.18–8.08 (m, 2H), 7.91–7.79 (m, 2H), 7.48–7.38 (m, 2H), 7.25–7.14 (m, 4H), 4.76 (h, J = 7.2 Hz,

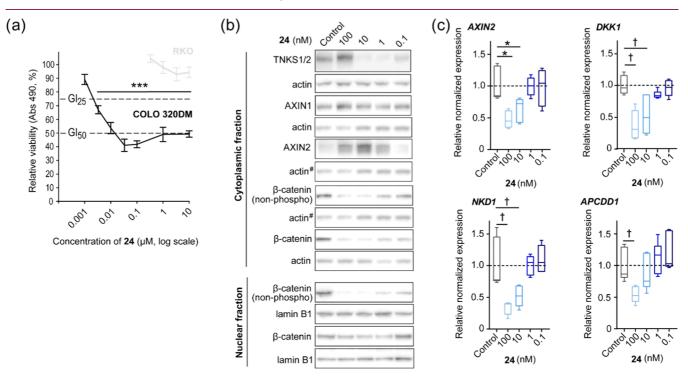


Figure 4. Compound 24 decreased cell growth and inhibited WNT/β-catenin signaling activity in COLO 320DM cells. (a) 3-(4,5-Dimethylthiazol-2-yl)-5-(3-carboxymethoxyphenyl)-2-(4-sulfophenyl)-2H-tetrazolium (MTS) colorimetric cell growth assay for various doses of 24 in APC^{mutated} COLO 320DM (black) and APC^{wild-type} RKO (gray) cells. After 5 days, the antiproliferative effect of compound treatment was measured at 490 nm. Mean value ± standard deviation (SD) for one representative experiment of more than three repeated assays, each with six replicates, are shown. Dotted lines depict 50% (GI₅₀-value) and 25% (GI₂₅-value) growth inhibition levels and control = 100% (0.1% dimethyl sulfoxide (DMSO)). (b) Representative immunoblots of cytoplasmic TNKS1/2, AXIN1, AXIN2, and cytoplasmic and nuclear transcriptionally active β-catenin (non-phospho) and β-catenin. Actin and lamin B1 show equal protein loading, while [#] indicates that the same actin immunoblot is used as loading control for both AXIN2 and β-catenin. For (b) and (c), control = 0.001% DMSO. (c) Real-time RT-qPCR analyses of WNT/β-catenin signaling target genes (*AXIN2, DKK1, NKD1,* and *APCDD1*). Boxplots show median, first and third quartiles, and maximum and minimum whiskers for combined data from three independent experiments with three replicates each. Dotted lines depict the control mean value = 1. For (a) and (c), analysis of variance (ANOVA) tests (Holm–Sidak method, versus control) are indicated by *** (*p* < 0.001) and * (*p* < 0.05), while ANOVA on ranks tests (Dunn's method, versus control) are indicated by [†] (*p* < 0.05).

1H), 4.54 (h, *J* = 6.0 Hz, 1H), 3.47 (tt, *J* = 10.1, 5.8 Hz, 1H), 3.11–2.97 (m, 2H), 2.52–2.36 (m, 2H), 1.32 (d, *J* = 6.0 Hz, 6H). HRMS $m/z \,[M + H]^+: 473.20958$ (calculated), 473.2089 (found), $\Delta = -1.42$ ppm.

N-(*trans*-3-(4-(2-*Fluorophenyl*)-5-*phenyl*-4*H*-1,2,4-*triazol*-3-*yl*)*cyclobutyl*)*picolinamide* (9). The title compound was prepared according to general procedure F as a white solid (21 mg, 84%). LC/ MS (ESI) *m*/*z* for C₂₄H₂₀N₅OF 413 (calculated) 414 ([M + H]⁺, found). ¹H NMR (400 MHz, CDCl₃) δ 8.56–8.50 (m, 1H), 8.23 (d, *J* = 6.8 Hz, 1H), 8.17 (dt, *J* = 7.9, 1.1 Hz, 1H), 7.84 (td, *J* = 7.7, 1.7 Hz, 1H), 7.53–7.46 (m, 1H), 7.46–7.39 (m, 3H), 7.39–7.32 (m, 1H), 7.32–7.27 (m, 2H), 7.26–7.21 (m, 2H), 7.21–7.13 (m, 1H), 4.83– 4.70 (m, 1H), 3.49 (tt, *J* = 10.2, 5.6 Hz, 1H), 3.11–2.95 (m, 2H), 2.52–2.40 (m, 2H). HRMS *m*/*z* [M + H]⁺: 414.17246 (calculated), 414.1714 (found), Δ = -2.46 ppm.

N-(*trans*-3-(4-(-2-*Fluorophenyl*)-5-(*pyrazin*-2-*yl*)-4H-1,2,4-*triazol*-3-*yl*)*cyclobutyl*)*picolinamide* (**10**). The title compound was prepared according to general procedure F as a white solid (7.9 mg, 21%). LC/MS (ESI) *m/z* for C₂₂H₁₈FN₇O 415 (calculated) 416 ([M + H]⁺, found). ¹H NMR (400 MHz, CDCl₃) δ 9.52 (d, *J* = 1.8 Hz, 1H), 8.53 (d, *J* = 4.4 Hz, 1H), 8.49 (d, *J* = 2.5 Hz, 1H), 8.22–8.14 (m, 2H), 7.84 (td, *J* = 7.7, 1.7 Hz, 1H), 7.54–7.39 (m, 2H), 7.22 (td, *J* = 8.1, 7.5, 1.9 Hz, 3H), 4.83–4.74 (m, 1H), 3.52–3.46 (m, 1H), 3.07–3.02 (m, 1H), 2.56–2.41 (m, 2H), 1.55 (s, 2H). HRMS *m/z* [M + H]⁺: 416.16296 (calculated), 416.1620 (found), Δ = -2.40 ppm.

N-(*trans*-3-(5-(5-*Ethoxypyridin*-2-*yl*)-4-*phenyl*-4*H*-1,2,4-*triazol*-3-*yl*)*cyclobutyl*)*picolinamide* (11). The title compound was prepared according to general procedure F as a white solid (30 mg, 52%). LC/ MS (ESI) *m/z* for C₂₅H₂₄N₆O₂ 440 (calculated) 441 ([M + H]⁺, found). ¹H NMR (400 MHz, CDCl₃) δ 8.53 (dq, *J* = 4.8, 1.1 Hz, 1H), 8.21 (d, *J* = 7.0 Hz, 1H), 8.16 (dt, *J* = 7.8, 1.2 Hz, 1H), 8.01 (d, *J* = 8.8 Hz, 1H), 7.94 (d, *J* = 2.9 Hz, 1H), 7.84 (td, *J* = 7.7, 1.7 Hz, 1H), 7.46-7.38 (m, 4H), 7.21 (dd, *J* = 8.7, 2.9 Hz, 1H), 7.17-7.12 (m, 2H), 4.76 (h, *J* = 6.5 Hz, 1H), 4.04 (q, *J* = 7.0 Hz, 2H), 3.50 (dddd, *J* = 10.4, 8.9, 6.2, 5.0 Hz, 1H), 3.09-2.98 (m, 2H), 2.47-2.37 (m, 2H), 1.40 (t, *J* = 7.0 Hz, 3H). HRMS *m/z* [M + H]⁺: 441.20335 (calculated), 441.2025 (found), Δ = -1.86 ppm.

N-(*trans*-3-(4-(3-Chlorophenyl)-5-(5-ethoxypyridin-2-yl)-4H-1,2,4-triazol-3-yl)cyclobutyl)picolinamide (**12**). The title compound was prepared according to general procedure F as a white solid (14.3 mg, 74%). LC/MS (ESI) *m*/*z* for C₂₅H₂₃N₆O₂Cl 474/476 (calculated) 475/477 ([M + H]⁺, found). ¹H NMR (400 MHz, CDCl₃) δ 8.53 (dt, *J* = 4.7, 1.3 Hz, 1H), 8.22 (d, *J* = 7.0 Hz, 1H), 8.17 (dt, *J* = 7.7, 1.0 Hz, 1H), 8.10 (d, *J* = 8.7 Hz, 1H), 7.93 (s, 1H), 7.84 (td, *J* = 7.7, 1.7 Hz, 1H), 7.46–7.40 (m, 2H), 7.36 (t, *J* = 8.0 Hz, 1H), 7.23 (dd, *J* = 8.7, 2.7 Hz, 1H), 7.18 (t, *J* = 2.0 Hz, 1H), 7.07 (dt, *J* = 8.0, 1.4 Hz, 1H), 4.78 (h, *J* = 7.0 Hz, 1H), 4.05 (q, *J* = 7.0 Hz, 2H), 3.53–3.42 (m, 1H), 3.09–2.97 (m, 2H), 2.51–2.39 (m, 2H), 1.41 (t, *J* = 7.0 Hz, 3H). HRMS *m*/*z* [M + H]⁺: 475.16438 (calculated), 475.1636 (found), Δ = -1.73 ppm.

N-(*trans*-3-(5-(5-*Ethoxypyrid*in-2-*yl*)-4-(5-*methylthiophen*-2-*yl*)-4*H*-1,2,4-*triazol*-3-*yl*)*cyclobutyl*)*picolinamide* (13). The title compound was prepared according to general procedure F as a white solid (19 mg, 85%). LC/MS (ESI) *m*/*z* for C₂₄H₂₄N₆O₂S 460 (calculated) 461 ([M + H]⁺, found). ¹H NMR (400 MHz, CDCl₃) δ 8.57–8.52 (m, 1H), 8.25 (d, *J* = 7.1 Hz, 1H), 8.18 (dt, *J* = 7.8, 1.1 Hz, 1H), 8.12 (d, *J* = 2.9 Hz, 1H), 7.95 (d, *J* = 8.7 Hz, 1H), 7.85 (td, *J* = 7.7, 1.7 Hz, 1H), 7.43 (ddd, *J* = 7.7, 4.8, 1.3 Hz, 1H), 7.22 (dd, *J* = 8.7, 3.0 Hz, 1H), 6.70 (d, *J* = 3.6 Hz, 1H), 6.62 (dd, *J* = 3.6, 1.3 Hz, 1H), 4.78 (q, *J* = 7.1 Hz, 1H), 4.08 (q, *J* = 6.9 Hz, 2H), 3.65–3.56 (m, 1H), 3.11– 3.01 (m, 2H), 2.56–2.49 (m, 2H), 2.47 (d, *J* = 1.1 Hz, 3H), 1.43 (t, *J* = 7.0 Hz, 3H). HRMS *m*/*z* [M + H]⁺: 461.17542 (calculated), 461.1746 (found), Δ = -1.85 ppm.

N-(*trans-3*-(4-(5-Chlorothiophen-2-yl)-5-(5-ethoxypyridin-2-yl)-4H-1,2,4-triazol-3-yl)cyclobutyl)picolinamide (14). The title compound was prepared according to general procedure F as a white solid (14.9 mg, 103%). LC/MS (ESI) m/z for C₂₃H₂₁N₆O₂SCl 480/482 (calculated) 481/483 ([M + H]⁺, found). ¹H NMR (400 MHz, CDCl₃) δ 8.55 (dt, *J* = 4.7, 1.4 Hz, 1H), 8.26 (d, *J* = 6.9 Hz, 1H), 8.18 (dt, J = 7.8, 1.1 Hz, 1H), 8.09 (d, J = 2.9 Hz, 1H), 8.06 (d, J = 8.8 Hz, 1H), 7.85 (td, J = 7.7, 1.7 Hz, 1H), 7.43 (ddd, J = 7.8, 4.7, 1.3 Hz, 1H), 7.25–7.20 (m, 1H), 6.82 (d, J = 4.0 Hz, 1H), 6.72 (d, J = 4.0 Hz, 1H), 4.79 (q, J = 7.2 Hz, 1H), 4.09 (q, J = 6.9 Hz, 2H), 3.63–3.56 (m, 1H), 3.11–3.01 (m, 2H), 2.59–2.48 (m, 2H), 1.43 (t, J = 7.0 Hz, 3H). HRMS m/z [M + H]⁺: 481.12080 (calculated), 481.1201 (found), $\Delta = -1.14$ ppm.

N-(*trans*-3-(5-(5-*Ethoxypyridin*-2-*yl*)-4-(*pyridin*-2-*yl*)-4H-1,2,4-*triazol*-3-*yl*)*cyclobutyl*)-1,5-*naphthyridine*-4-*carboxamide* (**16***a*). The title compound was prepared according to general procedure F as an off-white solid (13.7 mg, 55%). LC/MS (ESI) *m*/*z* for C₂₇H₂₄N₈O₂ 492 (calculated) 493 ([M + H]⁺, found). ¹H NMR (400 MHz, CDCl₃) δ 11.32 (d, *J* = 6.0 Hz, 1H), 9.14 (d, *J* = 4.4 Hz, 1H), 8.99 (dd, *J* = 4.3, 1.8 Hz, 1H), 8.59–8.52 (m, 3H), 8.17 (d, *J* = 8.8 Hz, 1H), 7.87 (d, *J* = 2.9 Hz, 1H), 7.79 (td, *J* = 7.7, 1.9 Hz, 1H), 7.75 (dd, *J* = 8.5, 4.2 Hz, 1H), 7.37 (ddd, *J* = 7.5, 4.9, 1.0 Hz, 1H), 7.26–7.22 (m, 1H), 7.21–7.16 (m, 1H), 4.86–4.73 (m, 1H), 4.05 (q, *J* = 6.9 Hz, 2H), 3.80–3.69 (m, 1H), 3.11–3.01 (m, 2H), 2.57–2.46 (m, 2H), 1.41 (t, *J* = 7.0 Hz, 3H). HRMS *m*/*z* [M + H]⁺: 493.20950 (calculated), 493.2085 (found), Δ = –1.99 ppm.

N-(*trans*-3-(5-(5-*Ethoxypyridin*-2-*yl*)-4-(*pyridin*-2-*yl*)-4*H*-1,2,4-*triazol*-3-*yl*)*cyclobutyl*)-7-*fluoro*-1,5-*naphthyridine*-4-*carboxamide* (**16b**). The title compound was prepared according to general procedure F as a white solid (17.9 mg, 69%). LC/MS (ESI) *m/z* for $C_{27}H_{23}N_8O_2F$ 510 (calculated) 511 ([M + H]⁺, found). ¹H NMR (400 MHz, CDCl₃) δ 10.81 (d, *J* = 6.0 Hz, 1H), 9.15 (d, *J* = 4.5 Hz, 1H), 8.92 (d, *J* = 2.9 Hz, 1H), 8.56 (dd, *J* = 5.1, 1.9 Hz, 1H), 8.52 (d, *J* = 4.4 Hz, 1H), 8.22–8.18 (m, 1H), 8.17 (d, *J* = 9.0 Hz, 1H), 7.87 (d, *J* = 2.8 Hz, 1H), 7.79 (td, *J* = 7.7, 1.9 Hz, 1H), 7.37 (dd, *J* = 7.5, 4.8 Hz, 1H), 7.26–7.21 (m, 1H), 7.18 (d, *J* = 8.0 Hz, 1H), 4.80 (h, *J* = 6.9 Hz, 1H), 4.05 (q, *J* = 7.0 Hz, 2H), 3.72 (tt, *J* = 10.2, 5.3 Hz, 1H), 3.06 (ddd, *J* = 13.2, 8.0, 5.3 Hz, 2H), 2.50 (ddd, *J* = 12.6, 9.5, 6.2 Hz, 2H), 1.41 (t, *J* = 7.0 Hz, 3H). HRMS *m/z* [M + H]⁺: \$11.20008 (calculated), \$11.1991 (found), Δ = -1.86 ppm.

N-(*trans*-3-(5-(5-*Ethoxypyridin*-2-*yl*)-4-(*pyridin*-3-*yl*)-4*H*-1,2,4-*triazol*-3-*yl*)*cyclobutyl*)-1,5-*naphthyridine*-4-*carboxamide* (**17a**). The title compound was prepared according to general procedure F as a white solid (12.6 mg, 50%). LC/MS (ESI) *m/z* for C₂₇H₂₄N₈O₂ 492 (calculated) 493 ([M + H]⁺, found). ¹H NMR (400 MHz, CDCl₃) δ 11.31 (d, *J* = 6.0 Hz, 1H), 9.14 (d, *J* = 4.5 Hz, 1H), 8.98 (dd, *J* = 4.3, 1.8 Hz, 1H), 8.68 (dd, *J* = 4.8, 1.5 Hz, 1H), 8.59–8.52 (m, 2H), 8.48 (d, *J* = 2.5 Hz, 1H), 8.17 (d, *J* = 8.8 Hz, 1H), 7.87 (d, *J* = 2.9 Hz, 1H), 7.74 (dd, *J* = 8.6, 4.2 Hz, 1H), 7.58 (ddd, *J* = 8.1, 2.5, 1.5 Hz, 1H), 7.41 (dd, *J* = 8.0, 4.7 Hz, 1H), 7.24 (dd, *J* = 8.7, 2.9 Hz, 1H), 4.86 (dtd, *J* = 8.7, 6.9, 5.2 Hz, 1H), 4.05 (q, *J* = 7.0 Hz, 2H), 3.58–3.47 (m, 1H), 3.14–3.03 (m, 2H), 2.62–2.51 (m, 2H), 1.41 (t, *J* = 7.0 Hz, 3H). HRMS *m/z* [M + H]⁺: 493.20950 (calculated), 493.2084 (found), Δ = –2.32 ppm.

N-(*trans*-3-(5-(5-*Ethoxypyridin*-2-*yl*)-4-(*pyridin*-3-*yl*)-4H-1,2,4-*triazol*-3-*yl*)*cyclobutyl*)-7-*fluoro*-1,5-*naphthyridine*-4-*carboxamide* (**17b**). The title compound was prepared according to general procedure F as a white solid (22 mg, 85%). LC/MS (ESI) *m/z* for $C_{27}H_{23}N_8O_2F$ 510 (calculated) 511 ([M + H]⁺, found). ¹H NMR (400 MHz, CDCl₃) δ 10.83 (d, *J* = 6.0 Hz, 1H), 9.15 (d, *J* = 4.4 Hz, 1H), 8.91 (d, *J* = 2.8 Hz, 1H), 8.69 (dd, *J* = 4.8, 1.5 Hz, 1H), 8.52 (d, *J* = 4.4 Hz, 1H), 8.48 (d, *J* = 2.5 Hz, 1H), 8.23–8.18 (m, 1H), 8.17 (d, *J* = 8.5 Hz, 1H), 7.87 (d, *J* = 2.8 Hz, 1H), 7.57 (dt, *J* = 8.1, 2.0 Hz, 1H), 7.41 (dd, *J* = 8.2, 4.8 Hz, 1H), 7.26–7.21 (m, 1H), 4.92–4.81 (m, 1H), 4.05 (q, *J* = 7.0 Hz, 2H), 3.56–3.45 (m, 1H), 3.14–3.03 (m, 2H), 2.60–2.50 (m, 2H), 1.41 (t, *J* = 7.0 Hz, 3H). HRMS *m/z* [M + H]⁺: \$11.20008 (calculated), \$11.1991 (found), Δ = -1.95 ppm.

N-(*trans*-3-(5-(5-*Ethoxypyridin*-2-*yl*)-4-(*pyridin*-4-*yl*)-4*H*-1,2,4-*triazol*-3-*yl*)*cyclobutyl*)-1,5-*naphthyridine*-4-*carboxamide* (**18***a*). The title compound was prepared according to general procedure F as a white solid (12.7 mg, 51%). LC/MS (ESI) *m*/*z* for C₂₇H₂₄N₈O₂ 492 (calculated) 493 ([M + H]⁺, found). ¹H NMR (400 MHz, CDCl₃) δ 11.31 (d, *J* = 6.0 Hz, 1H), 9.15 (d, *J* = 4.4 Hz, 1H), 8.97 (dd, *J* = 4.3, 1.8 Hz, 1H), 8.75–8.69 (m, 2H), 8.59–8.52 (m, 2H), 8.17 (d, *J* = 8.8 Hz, 1H), 7.88 (d, *J* = 2.8 Hz, 1H), 7.74 (dd, *J* = 8.5, 4.2 Hz, 1H), 7.26–7.22 (m, 1H), 7.17–7.11 (m, 2H), 4.92–4.81 (m, 1H), 4.06 (q, *J* = 7.0 Hz, 2H), 3.58–3.47 (m, 1H), 3.14–3.04 (m, 2H), 2.63–2.52 (m, 2H), 1.42 (t, *J* = 7.0 Hz, 3H). HRMS m/z [M + H]⁺: 493.20950 (calculated), 493.2084 (found), Δ = -2.16 ppm.

N-(*trans*-3-(5-(5-*E*thoxypyridin-2-yl)-4-(pyridin-4-yl)-4H-1,2,4-triazol-3-yl)cyclobutyl)-7-fluoro-1,5-naphthyridine-4-carboxamide (**18b**). The title compound was prepared according to general procedure F as a white solid (19.2 mg, 73%). LC/MS (ESI) *m/z* for $C_{27}H_{23}N_8O_2F$ 510 (calculated) 511 ([M + H]⁺, found). ¹H NMR (400 MHz, CDCl₃) δ 10.83 (d, *J* = 6.0 Hz, 1H), 9.16 (d, *J* = 4.5 Hz, 1H), 8.91 (d, *J* = 2.9 Hz, 1H), 8.76–8.69 (m, 2H), 8.52 (d, *J* = 4.5 Hz, 1H), 8.20 (dd, *J* = 8.7, 2.9 Hz, 1H), 8.17 (d, *J* = 8.6 Hz, 1H), 7.91–7.85 (m, 1H), 7.26–7.22 (m, 1H), 7.17–7.10 (m, 2H), 4.87 (h, *J* = 7.1 Hz, 1H), 4.06 (q, *J* = 6.9 Hz, 2H), 3.56–3.45 (m, 1H), 3.14– 3.03 (m, 2H), 2.62–2.51 (m, 2H), 1.42 (t, *J* = 7.0 Hz, 3H). HRMS *m/z* [M + H]⁺: 511.20008 (calculated), 511.1992 (found), Δ = –1.69 ppm.

N-(3-(5-(5-*Ethoxypyridin*-2-*yl*)-4-(*pyridin*-2-*yl*)-4*H*-1,2,4-*triazol*-3-*yl*)*bicyclo*[1.1.1]*pentan*-1-*yl*)-1,5-*naphthyridine*-4-*carboxamide* (**19**). The title compound was prepared according to general procedure F as a white solid (12.1 mg, 47%). LC/MS (ESI) *m/z* for C₂₈H₂₄N₈O₂ 504 (calculated) 505 ([M + H]⁺, found). ¹H NMR (400 MHz, CDCl₃) δ 11.46 (s, 1H), 9.14 (d, *J* = 4.5 Hz, 1H), 9.00 (dd, *J* = 4.3, 1.7 Hz, 1H), 8.63 (dd, *J* = 5.1, 1.8 Hz, 1H), 8.55 (dd, *J* = 8.6, 1.7 Hz, 1H), 8.50 (d, *J* = 4.4 Hz, 1H), 8.15 (d, *J* = 8.8 Hz, 1H), 7.87 (td, *J* = 7.7, 1.9 Hz, 1H), 7.84 (d, *J* = 3.0 Hz, 1H), 7.75 (dd, *J* = 8.6, 4.3 Hz, 1H), 7.46 (ddd, *J* = 7.6, 4.8, 1.0 Hz, 1H), 7.34 (d, *J* = 7.9 Hz, 1H), 7.21 (dd, *J* = 8.7, 2.9 Hz, 1H), 4.03 (q, *J* = 7.0 Hz, 2H), 2.48 (s, 6H), 1.40 (t, *J* = 7.0 Hz, 3H). HRMS *m/z* [M + H]⁺: 505.20950 (calculated), 505.2086 (found), Δ = -1.87 ppm.

N-(3-(5-(5-*Ethoxypyridin*-2-*yl*)-4-(*pyridin*-3-*yl*)-4*H*-1,2,4-*triazol*-3-*yl*)*bicyclo*[1.1.1]*pentan*-1-*yl*)-1,5-*naphthyridine*-4-*carboxamide* (**20**). The title compound was prepared according to general procedure F as a white solid (20.2 mg, 78%). LC/MS (ESI) *m/z* for C₂₈H₂₄N₈O₂ 504 (calculated) 505 ($[M + H]^+$, found). ¹H NMR (400 MHz, CDCl₃) δ 11.45 (s, 1H), 9.14 (d, *J* = 4.5 Hz, 1H), 8.99 (dd, *J* = 4.1, 1.8 Hz, 1H), 8.74 (dd, *J* = 4.8, 1.5 Hz, 1H), 8.58–8.52 (m, 2H), 8.49 (d, *J* = 4.4 Hz, 1H), 8.14 (d, *J* = 8.8 Hz, 1H), 7.87 (d, *J* = 2.9 Hz, 1H), 7.74 (dd, *J* = 8.6, 4.3 Hz, 1H), 7.69 (ddd, *J* = 8.1, 2.5, 1.5 Hz, 1H), 7.46 (dd, *J* = 8.0, 4.7 Hz, 1H), 7.22 (dd, *J* = 8.8, 2.9 Hz, 1H), 4.04 (q, *J* = 6.9 Hz, 2H), 2.49 (s, 6H), 1.40 (t, *J* = 7.0 Hz, 3H). HRMS *m*/*z* [M + H]⁺: 505.20950 (calculated), 505.2085 (found), Δ = -2.01 ppm.

N-(3-(5-(5-*E*thoxypyridin-2-yl)-4-(pyridin-4-yl)-4H-1,2,4-triazol-3-yl)bicyclo[1.1.1]pentan-1-yl)-1,5-naphthyridine-4-carboxamide (**21**). The title compound was prepared according to general procedure F as a white solid (10.2 mg, 62%). LC/MS (ESI) *m*/*z* for C₂₈H₂₄N₈O₂ 504 (calculated) 505 ([M + H]⁺, found). ¹H NMR (400 MHz, CDCl₃) δ 11.45 (s, 1H), 9.14 (d, *J* = 4.5 Hz, 1H), 8.99 (dd, *J* = 4.2, 1.8 Hz, 1H), 8.82–8.75 (m, 2H), 8.55 (dd, *J* = 8.6, 1.8 Hz, 1H), 8.50 (d, *J* = 4.4 Hz, 1H), 8.14 (d, *J* = 8.8 Hz, 1H), 7.86 (d, *J* = 2.8 Hz, 1H), 7.74 (dd, *J* = 8.5, 4.2 Hz, 1H), 7.27 (d, *J* = 1.7 Hz, 2H), 7.23 (dd, *J* = 8.7, 2.9 Hz, 1H), 4.04 (q, *J* = 6.9 Hz, 2H), 2.51 (s, 6H), 1.41 (t, *J* = 7.0 Hz, 3H). HRMS *m*/*z* [M + H]⁺: 505.20950 (calculated), 505.2085 (found), Δ = -2.02 ppm.

N-(*trans*-3-(5-(5-*Ethoxypyridin*-2-*yl*)-4-(5-*methylthiophen*-2-*yl*)-4*H*-1,2,4-*triazol*-3-*yl*)*cyclobutyl*)-1,5-*naphthyridine*-4-*carboxamide* (**22a**). The title compound was prepared according to general procedure F as a white solid (8.4 mg, 33%). LC/MS (ESI) *m/z* for $C_{27}H_{25}N_7O_2S$ 511 (calculated) 512 ([M + H]⁺, found). ¹H NMR (400 MHz, CDCl₃) δ 11.31 (d, *J* = 6.0 Hz, 1H), 9.15 (d, *J* = 4.4 Hz, 1H), 9.00 (dd, *J* = 4.3, 1.8 Hz, 1H), 8.60–8.53 (m, 2H), 8.12 (d, *J* = 2.8 Hz, 1H), 7.97 (d, *J* = 8.7 Hz, 1H), 7.75 (dd, *J* = 8.5, 4.2 Hz, 1H), 7.23–7.20 (m, 1H), 6.72 (d, *J* = 3.7 Hz, 1H), 6.61 (dd, *J* = 3.7, 1.3 Hz, 1H), 4.86 (q, *J* = 7.0 Hz, 1H), 4.08 (q, *J* = 7.0 Hz, 2H), 3.72– 3.62 (m, 1H), 3.16–3.05 (m, 2H), 2.69–2.58 (m, 2H), 2.47 (d, *J* = 1.1 Hz, 3H), 1.43 (t, *J* = 7.0 Hz, 3H). HRMS *m/z* [M + H]⁺: 512.18632 (calculated), 512.1854 (found), Δ = –1.73 ppm.

N-(trans-3-(5-(5-Ethoxypyridin-2-yl)-4-(5-methylthiophen-2-yl)-4H-1,2,4-triazol-3-yl)cyclobutyl)-7-fluoro-1,5-naphthyridine-4-carboxamide (**22b**). The title compound was prepared according to general procedure F as a white solid (13 mg, 43%). LC/MS (ESI) m/ *z* for C₂₇H₂₄N₇O₂SF 529 (calculated) 530 ([M + H]⁺, found). ¹H NMR (400 MHz, CDCl₃) δ 10.84 (d, *J* = 6.0 Hz, 1H), 9.16 (d, *J* = 4.4 Hz, 1H), 8.93 (d, *J* = 2.9 Hz, 1H), 8.54 (d, *J* = 4.4 Hz, 1H), 8.20 (dd, *J* = 8.7, 2.9 Hz, 1H), 8.12 (d, *J* = 2.9 Hz, 1H), 7.97 (d, *J* = 8.8 Hz, 1H), 7.23–7.19 (m, 1H), 6.72 (d, *J* = 3.7 Hz, 1H), 6.65–6.59 (m, 1H), 4.94–4.82 (m, 1H), 4.08 (q, *J* = 7.1 Hz, 2H), 3.72–3.60 (m, 1H), 3.15–3.06 (m, 2H), 2.66–2.57 (m, 2H), 2.47 (d, *J* = 1.1 Hz, 3H), 1.43 (t, *J* = 7.0 Hz, 3H). HRMS *m*/*z* [M + H]⁺: 530.17690 (calculated), 530.1761 (found), $\Delta = -1.59$ ppm.

N-(*trans*-3-(5-(5-*E*thoxypyridin-2-yl)-4-(5-methylthiophen-2-yl)-4H-1,2,4-triazol-3-yl)cyclobutyl)quinoline-8-carboxamide (**23**). The title compound was prepared according to general procedure F as a white solid (12 mg, 41%). LC/MS (ESI) *m*/*z* for C₂₈H₂₆N₆O₂S 510 (calculated) 511 ([M + H]⁺, found). ¹H NMR (400 MHz, CDCl₃) δ 11.57 (d, *J* = 5.9 Hz, 1H), 8.94 (dd, *J* = 4.2, 1.9 Hz, 1H), 8.85 (dd, *J* = 7.4, 1.6 Hz, 1H), 8.29 (dd, *J* = 8.3, 1.8 Hz, 1H), 8.12 (d, *J* = 2.8 Hz, 1H), 7.96 (dd, *J* = 8.3, 1.6 Hz, 2H), 7.68 (t, *J* = 7.7 Hz, 1H), 7.53– 7.47 (m, 1H), 7.22–7.19 (m, 1H), 6.72 (d, *J* = 3.7 Hz, 1H), 6.63– 6.58 (m, 1H), 4.83 (q, *J* = 7.0 Hz, 1H), 4.08 (q, *J* = 7.0 Hz, 2H), 3.77–3.60 (m, 1H), 3.12–3.06 (m, 2H), 2.67–2.60 (m, 2H), 2.46 (d, *J* = 1.2 Hz, 3H), 1.43 (t, *J* = 6.9 Hz, 3H). HRMS *m*/*z* [M + H]⁺: 511.19107 (calculated), 511.1902 (found), Δ = -1.63 ppm.

N-(*trans*-3-(5-(5-*Ethoxypyridin*-2-*yl*)-4-(2-*fluorophenyl*)-4H-1,2,4*triazol*-3-*yl*)*cyclobutyl*)*quinoxaline*-5-*carboxamide* (**24**, OM-153). The title compound was prepared according to general procedure F as a white solid (20.4 mg, 79%). LC/MS (ESI) *m/z* for C₂₈H₂₄N₇O₂F 509 (calculated) 510 ([M + H]⁺, found). ¹H NMR (400 MHz, CDCl₃) δ 10.68 (d, *J* = 5.9 Hz, 1H), 8.97 (d, *J* = 1.9 Hz, 1H), 8.89– 8.83 (m, 2H), 8.26 (dd, *J* = 8.3, 1.6 Hz, 1H), 8.16 (d, *J* = 8.7 Hz, 1H), 7.94–7.86 (m, 2H), 7.47–7.39 (m, 1H), 7.25–7.13 (m, 4H), 4.83 (dqd, *J* = 8.5, 6.8, 5.3 Hz, 1H), 4.04 (q, *J* = 7.0 Hz, 2H), 3.53 (dp, *J* = 10.1, 5.5 Hz, 1H), 3.08 (dtd, *J* = 12.0, 5.8, 3.9 Hz, 2H), 2.53 (ddtt, *J* = 19.4, 9.6, 6.2, 3.0 Hz, 2H), 1.40 (t, *J* = 7.0 Hz, 3H). HRMS *m/z* [M + H]⁺: \$10.20483 (calculated), \$10.2040 (found), Δ = -1.66 ppm.

N-(*trans*-3-(5-(5-*Ethoxypyridin*-2-*y*))-4-(2-*fluorophenyl*)-4H-1,2,4*triazo*]-3-*y*])*cyclobuty*])-7-*fluoroquinoxaline*-5-*carboxamide* (25). The title compound was prepared according to general procedure F as a white solid (18.2 mg, 68%). LC/MS (ESI) *m*/*z* for C₂₈H₂₃N₇O₂F 527 (calculated) 528 ([M + H]⁺, found). ¹H NMR (400 MHz, CDCl₃) δ 10.62 (d, *J* = 5.8 Hz, 1H), 8.96 (d, *J* = 1.8 Hz, 1H), 8.82 (d, *J* = 1.9 Hz, 1H), 8.64 (dd, *J* = 9.6, 3.1 Hz, 1H), 8.24–8.11 (m, 1H), 7.87 (dd, *J* = 7.9, 3.1 Hz, 2H), 7.48–7.39 (m, 1H), 7.25–7.14 (m, 4H), 4.83 (h, *J* = 6.8 Hz, 1H), 4.04 (q, *J* = 6.9 Hz, 2H), 3.57–3.46 (m, 1H), 3.13–3.01 (m, 2H), 2.60–2.45 (m, 2H), 1.40 (t, *J* = 6.9 Hz, 3H). HRMS *m*/*z* [M + H]⁺: 528.19541 (calculated), 528.1946 (found), $\Delta = -1.59$ ppm.

N-(*trans*-3-(5-(5-*Ethoxppyridin*-2-*yl*)-4-(2-fluorophenyl)-4H-1,2,4*triazol*-3-*yl*)*cyclobutyl*)-3-*methylquinoxaline*-5-*carboxamide* (**26**). The title compound was prepared according to general procedure F as a slightly pink solid (93.1 mg, 65%). LC/MS (ESI) *m/z* for C₂₉H₂₆N₇O₂F 523 (calculated) 524 ($[M + H]^+$, found). ¹H NMR (400 MHz, DMSO) δ 10.42 (d, *J* = 7.1 Hz, 1H), 8.98 (s, 1H), 8.41 (dd, *J* = 7.3, 1.5 Hz, 1H), 8.21 (dd, *J* = 8.3, 1.5 Hz, 1H), 8.06 (d, *J* = 8.7 Hz, 1H), 7.94 (d, *J* = 3.0 Hz, 1H), 7.87 (dd, *J* = 8.3, 7.3 Hz, 1H), 7.61–7.53 (m, 2H), 7.50 (dd, *J* = 8.8, 3.0 Hz, 1H), 7.47–7.39 (m, 1H), 7.36–7.29 (m, 1H), 4.75 (q, *J* = 7.3 Hz, 1H), 4.10 (q, *J* = 7.0 Hz, 2H), 3.40 (dt, *J* = 9.4, 4.6 Hz, 1H), 2.92–2.82 (m, 1H), 2.75 (s, 3H), 2.72–2.66 (m, 1H), 2.45–2.29 (m, 2H), 1.31 (t, *J* = 6.9 Hz, 3H). HRMS *m/z* [M + H]⁺: S24.22048 (calculated), S24.2197 (found), Δ = -1.56 ppm.

N-(*trans-3*-(5-(5-*Ethoxypyridin-2-yl*)-4-(2-*fluorophenyl*)-4H-1,2,4*triazol-3-yl*)*cyclobutyl*)-2-*methylquinoxaline-5-carboxamide* (27). The title compound was prepared according to general procedure F as an off-white solid (20.1 mg, 71%). LC/MS (ESI) *m/z* for $C_{29}H_{26}N_7O_2F$ 523 (calculated) 524 ($[M + H]^+$, found). ¹H NMR (400 MHz, DMSO) δ 10.42 (d, *J* = 7.1 Hz, 1H), 8.98 (s, 1H), 8.41 (dd, *J* = 7.3, 1.6 Hz, 1H), 8.21 (dd, *J* = 8.4, 1.5 Hz, 1H), 8.06 (d, *J* = 8.8 Hz, 1H), 7.93 (d, *J* = 2.8 Hz, 1H), 7.91–7.84 (m, 1H), 7.63–7.52 (m, 2H), 7.50 (dd, *J* = 8.8, 3.0 Hz, 1H), 7.47–7.39 (m, 1H), 7.36–7.29 (m, 1H), 4.75 (q, *J* = 7.2 Hz, 1H), 4.10 (q, *J* = 7.0 Hz, 2H), 3.44–3.36 (m, 1H), 2.87 (q, *J* = 3.8 Hz, 1H), 2.75 (s, 3H), 2.68 (s, 1H), 2.45–2.28 (m, 2H), 1.31 (t, *J* = 6.9 Hz, 3H). HRMS *m*/*z* [M + H]⁺: 524.22048 (calculated), 524.2198 (found), Δ = –1.36 ppm.

N-(*trans*-3-(5-(5-*Ethoxypyridin*-2-*y*))-4-(2-*fluorophenyl*)-4H-1,2,4*triazol*-3-*y*)(*cyclobutyl*)-2,3-*dimethylquinoxaline*-5-*carboxamide* (**28**). The title compound was prepared according to general procedure F as an off-white solid (29.7 mg, 43%). LC/MS (ESI) *m*/*z* for C₃₀H₂₈N₇O₂F 537 (calculated) 538 ([M + H]⁺, found). ¹H NMR (400 MHz, DMSO) δ 10.59 (d, *J* = 7.2 Hz, 1H), 8.36 (dd, *J* = 7.4, 1.6 Hz, 1H), 8.18–8.01 (m, 2H), 7.94 (d, *J* = 3.0 Hz, 1H), 7.82 (t, *J* = 7.8 Hz, 1H), 7.56 (td, *J* = 8.2, 3.1 Hz, 2H), 7.50 (dd, *J* = 8.8, 2.9 Hz, 1H), 7.43 (t, *J* = 9.2 Hz, 1H), 7.33 (t, *J* = 7.7 Hz, 1H), 4.75 (q, *J* = 7.2 Hz, 1H), 4.10 (q, *J* = 7.0 Hz, 2H), 2.86 (d, *J* = 9.7 Hz, 1H), 2.72 (d, *J* = 3.6 Hz, 7H), 2.45–2.28 (m, 2H), 1.31 (t, *J* = 7.0 Hz, 3H). HRMS *m*/*z* [M + H]⁺: 538.23613 (calculated), 538.2349 (found), Δ = -2.27 ppm.

N-(*trans*-3-(5-(5-(*Difluoromethoxy*)*pyridin*-2-*yl*)-4-(2-*fluorophen*-*yl*)-4*H*-1,2,4-*triazol*-3-*yl*)*cyclobutyl*)*quinoxaline*-5-*carboxamide* (**29a**). The title compound was prepared according to general procedure F as a white solid (23 mg, 86%). LC/MS (ESI) *m/z* for $C_{27}H_{20}N_7O_2F_3$ 531 (calculated) 532 ([M + H]⁺, found). ¹H NMR (400 MHz, CDCl₃) δ 10.67 (d, *J* = 5.9 Hz, 1H), 8.97 (d, *J* = 1.8 Hz, 1H), 8.89–8.83 (m, 2H), 8.33 (d, *J* = 8.8 Hz, 1H), 8.26 (dd, *J* = 8.4, 1.6 Hz, 1H), 8.07 (d, *J* = 2.8 Hz, 1H), 7.90 (dd, *J* = 8.4, 7.4 Hz, 1H), 7.55 (dd, *J* = 8.8, 2.8 Hz, 1H), 7.51–7.42 (m, 1H), 7.26–7.16 (m, 3H), 6.52 (t, *J* = 72.5 Hz, 1H), 4.90–4.79 (m, 1H), 3.57–3.47 (m, 1H), 3.14–3.02 (m, 2H), 2.63–2.47 (m, 2H). HRMS *m/z* [M + H]⁺: 532.17033 (calculated), 532.1692 (found), Δ = –2.12 ppm.

N-(*trans*-3-(5-(5-(2,2-*Difluoroethoxy*)*pyridin*-2-*yl*)-4-((5)-2-*fluorophenyl*)-4*H*-1,2,4-*triazol*-3-*yl*)*cyclobutyl*)-7-*fluoroquinoxaline*-5-*carboxamide* (**29b**). The title compound was prepared according to general procedure F as a white solid (19.8 mg, 71%). LC/MS (ESI) *m/z* for C₂₇H₁₉N₇O₂F₄ 549 (calculated) 550 ([M + H]⁺, found). ¹H NMR (400 MHz, CDCl₃) δ 10.63 (d, *J* = 5.9 Hz, 1H), 8.96 (d, *J* = 2.0 Hz, 1H), 8.82 (d, *J* = 1.9 Hz, 1H), 8.64 (dd, *J* = 9.5, 3.1 Hz, 1H), 8.33 (d, *J* = 8.7 Hz, 1H), 8.07 (d, *J* = 2.6 Hz, 1H), 7.87 (dd, *J* = 7.8, 3.1 Hz, 1H), 7.55 (dd, *J* = 8.8, 2.8 Hz, 1H), 7.51–7.42 (m, 1H), 7.25–7.16 (m, 3H), 6.52 (t, *J* = 72.5 Hz, 1H), 4.91–4.79 (m, 1H), 3.56–3.45 (m, 1H), 3.14–3.01 (m, 2H), 2.62–2.47 (m, 2H). HRMS *m/z* [M + H]⁺: \$50.16091 (calculated), \$50.1597 (found), Δ = -2.25 ppm.

N-(*trans*-3-(4-(2-*Fluorophenyl*)-5-(*pyridin*-2-*yl*)-4H-1,2,4-*triazol*-3-*yl*)*cyclobutyl*)*quinoxaline*-5-*carboxamide* (**30***a*). The title compound was prepared according to general procedure F as a white solid (19.3 mg, 83%). LC/MS (ESI) *m*/*z* for C₂₆H₂₀N₇OG 465 (calculated) 466 ([M + H]⁺, found). ¹H NMR (400 MHz, CDCl₃) δ 10.68 (d, *J* = 5.8 Hz, 1H), 8.97 (d, *J* = 1.8 Hz, 1H), 8.89–8.83 (m, 2H), 8.26 (dd, *J* = 8.3, 1.5 Hz, 2H), 8.24–8.18 (m, 1H), 7.94–7.87 (m, 1H), 7.76 (td, *J* = 7.8, 1.7 Hz, 1H), 7.49–7.40 (m, 1H), 7.25–7.14 (m, 4H), 4.90–4.78 (m, 1H), 3.59–3.49 (m, 1H), 3.15–3.03 (m, 2H), 2.62–2.47 (m, 2H). HRMS *m*/*z* [M + H]⁺: 466.17861 (calculated), 466.1775 (found), Δ = –2.34 ppm.

N-(*trans*-3-(5-(5-(6-*Methylpyridin*-2-*y*))-4-(5)-2-fluorophenyl)-4H-1,2,4-triazol-3-*y*)/*cyclobutyl*)-7-fluoroquinoxaline-5-carboxamide (**30b**). The title compound was prepared according to general procedure F as a white solid (16 mg, 65%). LC/MS (ESI) *m/z* for $C_{26}H_{19}N_7OF_2$ 483 (calculated) 484 ([M + H]⁺, found). ¹H NMR (400 MHz, CDCl₃) δ 10.63 (d, *J* = 5.8 Hz, 1H), 8.96 (d, *J* = 1.8 Hz, 1H), 8.83 (d, *J* = 1.8 Hz, 1H), 8.64 (dd, *J* = 9.6, 3.1 Hz, 1H), 8.26 (dt, *J* = 8.1, 1.1 Hz, 1H), 8.22 (dd, *J* = 4.8, 1.6 Hz, 1H), 7.87 (dd, *J* = 7.8, 3.1 Hz, 1H), 7.76 (td, *J* = 7.8, 1.8 Hz, 1H), 7.49–7.40 (m, 1H), 7.25– 7.14 (m, 4H), 4.84 (ht, *J* = 7.1, 1.5 Hz, 1H), 3.53 (tt, *J* = 9.3, 5.3 Hz, 1H), 3.14–3.02 (sym. m, 2H), 2.55 (dddd, *J* = 16.5, 13.1, 9.8, 6.5 Hz, 2H). HRMS *m/z* [M + H]⁺: 484.16919 (calculated), 484.1684 (found), Δ = -1.65 ppm.

N-(*trans*-3-(4-(2-*Fluorophenyl*)-5-(6-*methylpyridin*-2-*yl*)-4H-1,2,4-*triazol*-3-*yl*)*cyclobutyl*)*quinoxaline*-5-*carboxamide* (**31a**). The title compound was prepared according to general procedure F as a white solid (20.7 mg, 86%). LC/MS (ESI) *m/z* for C₂₇H₂₂N₇OF 479 (calculated) 480 ([M + H]⁺, found). ¹H NMR (400 MHz, CDCl₃) δ 10.68 (d, *J* = 5.7 Hz, 1H), 8.96 (d, *J* = 1.9 Hz, 1H), 8.90– 8.82 (m, 2H), 8.26 (dd, *J* = 8.4, 1.6 Hz, 1H), 8.07 (d, *J* = 7.8 Hz, 1H), 7.90 (dd, J = 8.4, 7.4 Hz, 1H), 7.63 (t, J = 7.8 Hz, 1H), 7.49–7.41 (m, 1H), 7.26–7.14 (m, 3H), 7.03 (d, J = 7.7 Hz, 1H), 4.89–4.78 (m, 1H), 3.61–3.50 (m, 1H), 3.15–3.05 (m, 2H), 2.62–2.48 (m, 2H), 2.07 (s, 3H). HRMS m/z [M + H]⁺: 480.19426 (calculated), 480.1934 (found), $\Delta = -1.75$ ppm.

N-(*trans*-3-(4-(2-*Fluorophenyl*)-5-(6-*methylpyridin*-2-*yl*)-4H-1,2,4-*triazol*-3-*yl*)*cyclobutyl*)-7-*fluoroquinoxaline*-5-*carboxamide* (**31b**). The title compound was prepared according to general procedure F as a white solid (18.2 mg, 72%). LC/MS (ESI) *m/z* for $C_{27}H_{21}N_7OF_2$ 497 (calculated) 498 ([M + H]⁺, found). ¹H NMR (400 MHz, CDCl₃) δ 10.63 (d, *J* = 5.9 Hz, 1H), 8.96 (d, *J* = 1.9 Hz, 1H), 8.83 (d, *J* = 1.8 Hz, 1H), 8.64 (dd, *J* = 9.5, 3.1 Hz, 1H), 8.06 (d, *J* = 7.8 Hz, 1H), 7.87 (dd, *J* = 7.8, 3.1 Hz, 1H), 7.63 (t, *J* = 7.8 Hz, 1H), 7.49–7.41 (m, 1H), 7.24–7.15 (m, 3H), 7.03 (d, *J* = 7.7 Hz, 1H), 4.91–4.78 (m, 1H), 3.60–3.49 (m, 1H), 3.15–3.03 (m, 2H), 2.62–2.48 (m, 2H), 2.07 (s, 3H). HRMS *m/z* [M + H]⁺: 498.18484 (calculated), 498.1838 (found), Δ = -2.11 ppm.

Biochemical Assay. Recombinantly expressed human tankyrase active constructs for TNKS1 (residues 1030-1317) and TNKS2 (residues 873-1162) and other PARP enzymes used for biochemical assays were produced as previously described.³⁹ The enzymatic assay measures unreacted NAD⁺, which is chemically reacted into a fluorescent compound.53 The fluorescence intensity was measured with excitation/emission wavelengths of 372 and 444 nm, respectively, using Tecan Infinity M1000 Pro. New compounds were prepared in half-log dilution series, and the reactions were carried out in quadruplicates with protein and compound controls to exclude the effect of compound autofluorescence. All reactions were performed at ambient temperature. TNKS1 (20 nM) or TNKS2 (5 nM) was incubated for 20 h in assay buffer (50 mM Bis-Tris propane (BTP), pH 7.0, 0.5 mM Tris(2-carboxyethyl)phosphine (TCEP), 0.01% Triton X-100) with compound and 10 µM or 500 nM NAD⁺, respectively.

The assay conditions for the other PARP enzymes were used as previously described.³⁹ Reference compound OD336 behaved in this assay as previously reported.²⁸

For single IC_{50} curves, the 95% confidence interval (asymptotic) was calculated in GraphPad Prism 8.

WNT/ β -Catenin Signaling Reporter Assay. The luciferase based WNT/ β -catenin signaling pathway reporter assay in human HEK293 cells was performed in triplicates as previously described²¹ also taking OD336 as a reference compound, which behaved in this assay as previously described.²⁸

ADME. Kinetic solubility, microsomal stability, and plasma stability were determined following our internal protocol (Symeres). The Caco-2 and PPB determinations were performed by Cyprotex.

Off-Target. A safety panel (Cerep) of 44 selected targets (n = 2) including hERG Inhibition using 10 μ M 24 was performed by Eurofins. Ames and CYP3A4 inhibition assays were performed by Cyprotex, and the PXR assay (CYP induction) was performed at Admescope. The PARP assays are described below.

Mouse Pharmacokinetic Analysis. The pharmacokinetic analyses in mice were performed according to the standard protocols of Medicilon as previously described²¹ following approval by local animal experiment authorities (Shanghai, China) and in compliance with FELASA guidelines and recommendations. Three IRC mice were used per treatment group: 5 mg/kg 24 in 5% DMSO, 50% PEG400 (both Sigma-Aldrich), and 45% saline as vehicle. Samples were collected after 5, 15, and 30 min and 1, 2, 4, 8, and 24 h.

Crystallography. The co-crystallization of human TNKS2 catalytic domain (residues 952–1161) in complex with **24** was done in the presence of chymotrypsin (1:100) and based on crystallization efforts previously described.³⁹ Protein (5.3 mg/mL) was mixed with 0.4 mM compound from a 10 mM stock solution in DMSO. The droplets for crystallization were set up using the sitting-drop vapor diffusion method by mixing 200 nL of protein with 100 nL of precipitant solution (0.1 M Bicine, pH 8.5–9.0, 7.5–25% PEG6000). All steps were performed at room temperature. Rod-shaped crystals appeared within 24 h and were cryoprotected using the precipitant solution containing 25% PEG6000 and 20% glycerol.

Data were collected at Diamond Light Source on beamline I04. Diffraction data were processed using the XDS package.⁵⁴ The substructure was solved using molecular replacement with PHASER⁵⁵ using the structure of TNKS2 (PDB code: 5NOB) as starting model. Model building and refinement was done using Coot⁵⁶ and Refmac5,⁵⁷ respectively. Preparation of the crystal structure images was done with The PyMOL Molecular Graphics System (PyMOL, version 1.8.4.0).

Proliferation Assay. Antiproliferative assays using colon cancer cell lines COLO 320DM and RKO were performed as previously described. 39

Western Blot Analysis. Western blot analysis of nuclear and cytoplasmic lysates from compound-treated COLO 320DM cells was performed as previously described.^{39,50}

RNA Isolation and Real-Time qRT-PCR. RNA isolation and real-time qRT-PCR were performed as previously described.^{39,50}

ASSOCIATED CONTENT

Supporting Information

The Supporting Information is available free of charge at https://pubs.acs.org/doi/10.1021/acs.jmedchem.1c01264.

Molecular formula strings and data (CSV)

General and specific synthetic procedures and spectra for all compounds and inhibition data and crystallography; atomic coordinates and structure factors have been deposited to the Protein Data Bank under accession number 7O6X, and raw diffraction images are available at IDA (https://doi.org/10.23729/ 4d448eb0-c6ef-4b7d-a557-3758f5f68d47). The authors will release the atomic coordinates upon article publication (PDF)

AUTHOR INFORMATION

Corresponding Author

Stefan Krauss – Hybrid Technology Hub - Centre of Excellence, Institute of Basic Medical Sciences, University of Oslo, 0317 Oslo, Norway; Department of Immunology and Transfusion Medicine, Oslo University Hospital, 0424 Oslo, Norway; Phone: +47 97610063; Email: s.j.k.krauss@ medisin.uio.no

Authors

- Ruben G. G. Leenders Symeres, 6546 BB Nijmegen, The Netherlands; orcid.org/0000-0001-6140-5405
- Shoshy Alam Brinch Hybrid Technology Hub Centre of Excellence, Institute of Basic Medical Sciences, University of Oslo, 0317 Oslo, Norway; Department of Immunology and Transfusion Medicine, Oslo University Hospital, 0424 Oslo, Norway

Sven T. Sowa – Faculty of Biochemistry and Molecular Medicine, Biocenter Oulu, University of Oulu, 90014 Oulu, Finland

Enya Amundsen-Isaksen – Hybrid Technology Hub - Centre of Excellence, Institute of Basic Medical Sciences, University of Oslo, 0317 Oslo, Norway; Department of Immunology and Transfusion Medicine, Oslo University Hospital, 0424 Oslo, Norway

Albert Galera-Prat – Faculty of Biochemistry and Molecular Medicine, Biocenter Oulu, University of Oulu, 90014 Oulu, Finland

Sudarshan Murthy – Faculty of Biochemistry and Molecular Medicine, Biocenter Oulu, University of Oulu, 90014 Oulu, Finland

- Sjoerd Aertssen Symeres, 6546 BB Nijmegen, The Netherlands
- Johannes N. Smits Symeres, 6546 BB Nijmegen, The Netherlands
- **Piotr Nieczypor** Symeres, 6546 BB Nijmegen, The Netherlands
- Eddy Damen Symeres, 6546 BB Nijmegen, The Netherlands
- Anita Wegert Symeres, 6546 BB Nijmegen, The Netherlands
- Marc Nazaré Medicinal Chemistry, Leibniz-Forschungsinstitut für Molekulare Pharmakologie (FMP), 13125 Berlin, Germany; ⊚ orcid.org/0000-0002-1602-2330
- Lari Lehtiö Faculty of Biochemistry and Molecular Medicine, Biocenter Oulu, University of Oulu, 90014 Oulu, Finland; orcid.org/0000-0001-7250-832X
- Jo Waaler Hybrid Technology Hub Centre of Excellence, Institute of Basic Medical Sciences, University of Oslo, 0317 Oslo, Norway; Department of Immunology and Transfusion Medicine, Oslo University Hospital, 0424 Oslo, Norway; orcid.org/0000-0002-8501-6225

Complete contact information is available at: https://pubs.acs.org/10.1021/acs.jmedchem.1c01264

Author Contributions

[#]R.G.G.L. and S.A.B. contributed equally to this article, while J.W. and S.K. contributed equally as co-senior authors of this article. The manuscript was written through contributions of all authors, and all authors have given approval to the final version of the manuscript.

Funding

S.K. was supported by the Research Council of Norway (grant nos. 262613 and 267639) and by South-Eastern Norway Regional Health Authority (grant nos. 16/00528-9, 15/00779-2, and 2015012). S.A.B. and J.W. were supported by the South-Eastern Norway Regional Health Authority (grant nos. 2019090 and 2021035). L.L., S.T.S., A.G.-P., and S.M. were supported by the Jane and Aatos Erkko Foundation, Sigrid Jusélius Foundation and Academy of Finland (grant nos. 287063, 294085, and 319299).

Notes

The authors declare the following competing financial interest(s): J.W., M.N., L.L., A.W., R.G.G.L., and S.K. hold patents related to tankyrase inhibitor therapy, and these authors declare no additional interests. The remaining authors declare no competing interests.

ACKNOWLEDGMENTS

Heli Alanen is acknowledged for her contribution in performing some of the biochemical assays. Ilonka Meerts and Eef van den Elzen, both of Symeres, are acknowledged for their contributions with the ADME assays and the HRMS. Peter Zenhorst (Symeres) is acknowledged for the development of a CMC route for compound **24**. Protein crystallography experiments were performed at the Diamond Light Source (Didcot, U.K.) on beamline I04. The authors are grateful to local contacts for providing assistance in using the beamlines. The use of the facilities and expertise of the Biocenter Oulu Protein Crystallography core facility, a member of Biocenter Finland and Instruct-FI, is gratefully acknowledged.

ABBREVIATIONS USED

ADME, absorption, distribution, metabolism, and excretion; ADP, adenosine 5'-diphosphate; AKT, serine/threonine kinase; AMOT, angiomotin; AMP, adenosine monophosphate; AMPK, AMP-activated protein kinase; APC, adenomatous polyposis coli; APCDD1, APC downregulated 1; AUC, area under the curve; AXIN, axis inhibition protein; CL, clearance; CMC, chemistry, manufacturing, and control; DKK1, dickkopf WNT signaling pathway inhibitor 1; DMSO, dimethyl sulfoxide; HEK293, human embryonic kidney 293; hERG, human ether-à-go-go-related gene; LC/MS, liquid chromatography/mass spectroscopy; MTS, 3-(4,5-dimethylthiazol-2-yl)-5-(3-carboxymethoxyphenyl)-2-(4-sulfophenyl)-2H-tetrazolium; NKD1, NKD inhibitor of WNT signaling pathway 1; NAD, nicotinamide adenine dinucleotide; NMR, nuclear magnetic resonance; PARP, poly-ADP-ribose polymerase; PARsylate, poly(ADP-ribose)sylate; PGC-1 α , peroxisome proliferator-activated receptor- γ coactivator 1 α ; PK, pharmacokinetics; PO, per oral; PPB, plasma protein binding; PTEN, phosphatase and tensin homolog; RT-qPCR, reverse transcription quantitative polymerase chain reaction; RNF146, ring finger protein 146; SD, standard deviation; SEM, standard error of the mean; SH3BP2, sarcoma homology 3 (SH3) domain binding protein 2; SOX9, sex-determining region Y (SRY)-box transcription factor 9; $t_{1/2}$, half-life; TNKS, telomeric repeat factor (TRF1)-interacting ankyrin-related ADP-ribose polymerases, tankyrase; TNKS1/2, tankyrase 1 and tankyrase 2; tPSA, total polar surface area; TRF, telomeric repeat factor; V_{d} , volume of distribution; WNT, wingless-type mammary tumor virus integration site

REFERENCES

(1) Zhang, Y.; Liu, S.; Mickanin, C.; Feng, Y.; Charlat, O.; Michaud, G. A.; Schirle, M.; Shi, X.; Hild, M.; Bauer, A.; Myer, V. E.; Finan, P. M.; Porter, J. A.; Huang, S.-M. A.; Cong, F. RNF146 Is a Poly(ADP-Ribose)-Directed E3 Ligase That Regulates Axin Degradation and Wnt Signalling. *Nat. Cell Biol.* **2011**, *13*, 623–629.

(2) Callow, M. G.; Tran, H.; Phu, L.; Lau, T.; Lee, J.; Sandoval, W. N.; Liu, P. S.; Bheddah, S.; Tao, J.; Lill, J. R.; Hongo, J.-A.; Davis, D.; Kirkpatrick, D. S.; Polakis, P.; Costa, M. Ubiquitin Ligase RNF146 Regulates Tankyrase and Axin to Promote Wnt Signaling. *PLoS One* **2011**, *6*, e22595–e22608.

(3) Haikarainen, T.; Kraus, S.; Lehtiö, L. Tankyrases: Structure, Function and Therapeutic Implications in Cancer. *Curr. Pharm. Des.* **2014**, *20*, 6472–6488.

(4) Nie, L.; Wang, C.; Li, N.; Feng, X.; Lee, N.; Su, D.; Tang, M.; Yao, F.; Chen, J. Proteome-Wide Analysis Reveals Substrates of E3 Ligase RNF146 Targeted for Degradation. *Mol. Cell. Proteomics* **2020**, *19*, 2015–2030.

(5) Seimiya, H.; Smith, S. The Telomeric Poly(ADP-Ribose) Polymerase, Tankyrase 1, Contains Multiple Binding Sites for Telomeric Repeat Binding Factor 1 (TRF1) and a Novel Acceptor, 182-KDa Tankyrase-Binding Protein (TAB182)*. J. Biol. Chem. 2002, 277, 14116–14126.

(6) Mariotti, L.; Templeton, C. M.; Ranes, M.; Paracuellos, P.; Cronin, N.; Beuron, F.; Morris, E.; Guettler, S. Tankyrase Requires SAM Domain-Dependent Polymerization to Support Wnt- β -Catenin Signaling. *Mol. Cell* **2016**, *63*, 498–513.

(7) Pollock, K.; Liu, M.; Zaleska, M.; Meniconi, M.; Pfuhl, M.; Collins, I.; Guettler, S. Fragment-Based Screening Identifies Molecules Targeting the Substrate-Binding Ankyrin Repeat Domains of Tankyrase. *Sci. Rep.* **2019**, *9*, No. 19130.

(8) Perdreau-Dahl, H.; Progida, C.; Barfeld, S. J.; Guldsten, H.; Thiede, B.; Arntzen, M.; Bakke, O.; Mills, I. G.; Krauss, S.; Morth, J. P. Sjögren Syndrome/Scleroderma Autoantigen 1 Is a Direct Tankyrase Binding Partner in Cancer Cells. Commun. Biol. 2020, 3, No. 123.

(9) Wang, H.; Kuusela, S.; Rinnankoski-Tuikka, R.; Dumont, V.; Bouslama, R.; Ramadan, U. A.; Waaler, J.; Linden, A.-M.; Chi, N.-W.; Krauss, S.; Pirinen, E.; Lehtonen, S. Tankyrase Inhibition Ameliorates Lipid Disorder via Suppression of PGC-1 α PARylation in Db/Db Mice. *Int. J. Obes.* **2020**, *44*, 1691–1702.

(10) Azarm, K.; Bhardwaj, A.; Kim, E.; Smith, S. Persistent Telomere Cohesion Protects Aged Cells from Premature Senescence. *Nat. Commun.* **2020**, *11*, No. 3321.

(11) Li, N.; Zhang, Y.; Han, X.; Liang, K.; Wang, J.; Feng, L.; Wang, W.; Songyang, Z.; Lin, C.; Yang, L.; Yu, Y.; Chen, J. Poly-ADP Ribosylation of PTEN by Tankyrases Promotes PTEN Degradation and Tumor Growth. *Genes Dev.* **2015**, *29*, 157–170.

(12) Kim, S.; Han, S.; Kim, Y.; Kim, H.-S.; Gu, Y.-R.; Kang, D.; Cho, Y.; Kim, H.; Lee, J.; Seo, Y.; Chang, M. J.; Chang, C. B.; Kang, S.-B.; Kim, J.-H. Tankyrase Inhibition Preserves Osteoarthritic Cartilage by Coordinating Cartilage Matrix Anabolism via Effects on SOX9 PARylation. *Nat. Commun.* **2019**, *10*, No. 4898.

(13) Mukai, T.; Fujita, S.; Morita, Y. Tankyrase (PARP5) Inhibition Induces Bone Loss through Accumulation of Its Substrate SH3BP2. *Cells* **2019**, *8*, 195–208.

(14) Peters, X. Q.; Malinga, T. H.; Agoni, C.; Olotu, F. A.; Soliman, M. E. S. Zoning in on Tankyrases: A Brief Review on the Past, Present and Prospective Studies. *Anti-Cancer Agents Med. Chem.* **2020**, *19*, 1920–1934.

(15) Zimmerlin, L.; Zambidis, E. T. Pleiotropic Roles of Tankyrase/ PARP Proteins in the Establishment and Maintenance of Human Naïve Pluripotency. *Exp. Cell Res.* **2020**, *390*, 111935–111945.

(16) Huang, S.-M. A.; Mishina, Y. M.; Liu, S.; Cheung, A.; Stegmeier, F.; Michaud, G. A.; Charlat, O.; Wiellette, E.; Zhang, Y.; Wiessner, S.; Hild, M.; Shi, X.; Wilson, C. J.; Mickanin, C.; Myer, V.; Fazal, A.; Tomlinson, R.; Serluca, F.; Shao, W.; Cheng, H.; Shultz, M.; Rau, C.; Schirle, M.; Schlegl, J.; Ghidelli, S.; Fawell, S.; Lu, C.; Curtis, D.; Kirschner, M. W.; Lengauer, C.; Finan, P. M.; Tallarico, J. A.; Bouwmeester, T.; Porter, J. A.; Bauer, A.; Cong, F. Tankyrase Inhibition Stabilizes Axin and Antagonizes Wnt Signalling. *Nature* **2009**, *461*, 614–620.

(17) Wang, Y.; Jiang, W.; Liu, X.; Zhang, Y. Tankyrase 2 (TNKS2) Polymorphism Associated with Risk in Developing Non-Small Cell Lung Cancer in a Chinese Population. *Pathol., Res. Pract.* **2015**, *211*, 766–771.

(18) Zamudio-Martinez, E.; Herrera-Campos, A. B.; Muñoz, A.; Rodríguez-Vargas, J. M.; Oliver, F. J. Tankyrases as Modulators of Pro-Tumoral Functions: Molecular Insights and Therapeutic Opportunities. J. Exp. Clin. Cancer Res. **2021**, 40, No. 144.

(19) Liu, Z.; Wang, P.; Wold, E. A.; Song, Q.; Zhao, C.; Wang, C.; Zhou, J. Small-Molecule Inhibitors Targeting the Canonical WNT Signaling Pathway for the Treatment of Cancer. *J. Med. Chem.* **2021**, *64*, 4257–4288.

(20) Chen, B.; Dodge, M. E.; Tang, W.; Lu, J.; Ma, Z.; Fan, C.-W.; Wei, S.; Hao, W.; Kilgore, J.; Williams, N. S.; Roth, M. G.; Amatruda, J. F.; Chen, C.; Lum, L. Small Molecule–Mediated Disruption of Wnt-Dependent Signaling in Tissue Regeneration and Cancer. *Nat. Chem. Biol.* **2009**, *S*, 100–107.

(21) Voronkov, A.; Holsworth, D. D.; Waaler, J.; Wilson, S. R.; Ekblad, B.; Perdreau-Dahl, H.; Dinh, H.; Drewes, G.; Hopf, C.; Morth, J. P.; Krauss, S. Structural Basis and SAR for G007-LK, a Lead Stage 1,2,4-Triazole Based Specific Tankyrase 1/2 Inhibitor. *J. Med. Chem.* **2013**, *56*, 3012–3023.

(22) Bregman, H.; Chakka, N.; Guzman-Perez, A.; Gunaydin, H.;
Gu, Y.; Huang, X.; Berry, V.; Liu, J.; Teffera, Y.; Huang, L.; Egge, B.;
Mullady, E. L.; Schneider, S.; Andrews, P. S.; Mishra, A.; Newcomb,
J.; Serafino, R.; Strathdee, C. A.; Turci, S. M.; Wilson, C.; DiMauro, E.
F. Discovery of Novel, Induced-Pocket Binding Oxazolidinones as
Potent, Selective, and Orally Bioavailable Tankyrase Inhibitors. *J. Med. Chem.* 2013, *56*, 4320–4342.

(23) Hua, Z.; Bregman, H.; Buchanan, J. L.; Chakka, N.; Guzman-Perez, A.; Gunaydin, H.; Huang, X.; Gu, Y.; Berry, V.; Liu, J.; Teffera, Y.; Huang, L.; Egge, B.; Emkey, R.; Mullady, E. L.; Schneider, S.; Andrews, P. S.; Acquaviva, L.; Dovey, J.; Mishra, A.; Newcomb, J.; Saffran, D.; Serafino, R.; Strathdee, C. A.; Turci, S. M.; Stanton, M.; Wilson, C.; DiMauro, E. F. Development of Novel Dual Binders as Potent, Selective, and Orally Bioavailable Tankyrase Inhibitors. *J. Med. Chem.* **2013**, *56*, 10003–10015.

(24) McGonigle, S.; Chen, Z.; Wu, J.; Chang, P.; Kolber-Simonds, D.; Ackermann, K.; Twine, N. C.; Shie, J.; Miu, J. T.; Huang, K.-C.; Moniz, G. A.; Nomoto, K. E7449: A Dual Inhibitor of PARP1/2 and Tankyrase1/2 Inhibits Growth of DNA Repair Deficient Tumors and Antagonizes Wnt Signaling. *Oncotarget* **2015**, *6*, 41307–41323.

(25) Paine, H. A.; Nathubhai, A.; Woon, E. C. Y.; Sunderland, P. T.; Wood, P. J.; Mahon, M. F.; Lloyd, M. D.; Thompson, A. S.; Haikarainen, T.; Narwal, M.; Lehtiö, L.; Threadgill, M. D. Exploration of the Nicotinamide-Binding Site of the Tankyrases, Identifying 3-Arylisoquinolin-1-Ones as Potent and Selective Inhibitors in Vitro. *Bioorg. Med. Chem.* **2015**, *23*, 5891–5908.

(26) Nkizinkiko, Y.; Suneel Kumar, B. V. S.; Jeankumar, V. U.; Haikarainen, T.; Koivunen, J.; Madhuri, C.; Yogeeswari, P.; Venkannagari, H.; Obaji, E.; Pihlajaniemi, T.; Sriram, D.; Lehtiö, L. Discovery of Potent and Selective Nonplanar Tankyrase Inhibiting Nicotinamide Mimics. *Bioorg. Med. Chem.* **2015**, *23*, 4139–4149.

(27) Haikarainen, T.; Waaler, J.; Ignatev, A.; Nkizinkiko, Y.; Venkannagari, H.; Obaji, E.; Krauss, S.; Lehtiö, L. Development and Structural Analysis of Adenosine Site Binding Tankyrase Inhibitors. *Bioorg. Med. Chem. Lett.* **2016**, *26*, 328–333.

(28) Anumala, U. R.; Waaler, J.; Nkizinkiko, Y.; Ignatev, A.; Lazarow, K.; Lindemann, P.; Olsen, P. A.; Murthy, S.; Obaji, E.; Majouga, A. G.; Leonov, S.; von Kries, J. P.; Lehtiö, L.; Krauss, S.; Nazaré, M. Discovery of a Novel Series of Tankyrase Inhibitors by a Hybridization Approach. J. Med. Chem. **2017**, 60, 10013–10025.

(29) Ferri, M.; Liscio, P.; Carotti, A.; Asciutti, S.; Sardella, R.; Macchiarulo, A.; Camaioni, E. Targeting Wnt-Driven Cancers: Discovery of Novel Tankyrase Inhibitors. *Eur. J. Med. Chem.* **2017**, *142*, 506–522.

(30) Di Micco, S.; Pulvirenti, L.; Bruno, I.; Terracciano, S.; Russo, A.; Vaccaro, M. C.; Ruggiero, D.; Muccilli, V.; Cardullo, N.; Tringali, C.; Riccio, R.; Bifulco, G. Identification by Inverse Virtual Screening of Magnolol-Based Scaffold as New Tankyrase-2 Inhibitors. *Bioorg. Med. Chem.* **2018**, *26*, 3953–3957.

(31) Mizutani, A.; Yashiroda, Y.; Muramatsu, Y.; Yoshida, H.; Chikada, T.; Tsumura, T.; Okue, M.; Shirai, F.; Fukami, T.; Yoshida, M.; Seimiya, H. RK-287107, a Potent and Specific Tankyrase Inhibitor, Blocks Colorectal Cancer Cell Growth in a Preclinical Model. *Cancer Sci.* **2018**, *109*, 4003–4014.

(32) Menon, M.; Elliott, R.; Bowers, L.; Balan, N.; Rafiq, R.; Costa-Cabral, S.; Munkonge, F.; Trinidade, I.; Porter, R.; Campbell, A. D.; Johnson, E. R.; Esdar, C.; Buchstaller, H.-P.; Leuthner, B.; Rohdich, F.; Schneider, R.; Sansom, O.; Wienke, D.; Ashworth, A.; Lord, C. J. A Novel Tankyrase Inhibitor, MSC2504877, Enhances the Effects of Clinical CDK4/6 Inhibitors. *Sci. Rep.* **2019**, *9*, No. 201216.

(33) Shirai, F.; Tsumura, T.; Yashiroda, Y.; Yuki, H.; Niwa, H.; Sato, S.; Chikada, T.; Koda, Y.; Washizuka, K.; Yoshimoto, N.; Abe, M.; Onuki, T.; Mazaki, Y.; Hirama, C.; Fukami, T.; Watanabe, H.; Honma, T.; Umehara, T.; Shirouzu, M.; Okue, M.; Kano, Y.; Watanabe, T.; Kitamura, K.; Shitara, E.; Muramatsu, Y.; Yoshida, H.; Mizutani, A.; Seimiya, H.; Yoshida, M.; Koyama, H. Discovery of Novel Spiroindoline Derivatives as Selective Tankyrase Inhibitors. *J. Med. Chem.* **2019**, *62*, 3407–3427.

(34) Buchstaller, H.-P.; Anlauf, U.; Dorsch, D.; Kuhn, D.; Lehmann, M.; Leuthner, B.; Musil, D.; Radtki, D.; Ritzert, C.; Rohdich, F.; Schneider, R.; Esdar, C. Discovery and Optimization of 2-Arylquinazolin-4-Ones into a Potent and Selective Tankyrase Inhibitor Modulating Wnt Pathway Activity. *J. Med. Chem.* **2019**, 62, 7897–7909.

(35) Sabnis, R. W. Novel 4-Heteroarylcarbonyl-N-(Phenyl or Heteroaryl) Piperidine-1-Carboxamides as Tankyrase Inhibitors. *ACS Med. Chem. Lett.* **2020**, *11*, 1676–1677.

(36) Kinosada, H.; Okada-Iwasaki, R.; Kunieda, K.; Suzuki-Imaizumi, M.; Takahashi, Y.; Miyagi, H.; Suzuki, M.; Motosawa, K.; Watanabe, M.; Mie, M.; Ishii, T.; Ishida, H.; Saito, J.-I.; Nakai, R. The Dual Pocket Binding Novel Tankyrase Inhibitor K-476 Enhances the Efficacy of Immune Checkpoint Inhibitor by Attracting CD8+ T Cells to Tumors. *Am. J. Cancer Res.* **2021**, *11*, 264–276.

pubs.acs.org/jmc

(37) Mehta, C. C.; Bhatt, H. G. Tankyrase Inhibitors as Antitumor Agents: A Patent Update (2013 – 2020). *Expert Opin. Ther. Pat.* **2021**, 31, 645–661.

(38) Waaler, J.; Machon, O.; Kries, J. P.; von Wilson, S. R.; Lundenes, E.; Wedlich, D.; Gradl, D.; Paulsen, J. E.; Machonova, O.; Dembinski, J. L.; Dinh, H.; Krauss, S. Novel Synthetic Antagonists of Canonical Wnt Signaling Inhibit Colorectal Cancer Cell Growth. *Cancer Res.* **2011**, *71*, 197–205.

(39) Waaler, J.; Leenders, R. G. G.; Sowa, S. T.; Alam Brinch, S.; Lycke, M.; Nieczypor, P.; Aertssen, S.; Murthy, S.; Galera-Prat, A.; Damen, E.; Wegert, A.; Nazaré, M.; Lehtiö, L.; Krauss, S. Preclinical Lead Optimization of a 1,2,4-Triazole Based Tankyrase Inhibitor. *J. Med. Chem.* **2020**, *63*, 6834–6846.

(40) Zhong, Y.; Katavolos, P.; Nguyen, T.; Lau, T.; Boggs, J.; Sambrone, A.; Kan, D.; Merchant, M.; Harstad, E.; Diaz, D.; Costa, M.; Schutten, M. Tankyrase Inhibition Causes Reversible Intestinal Toxicity in Mice with a Therapeutic Index <1. *Toxicol. Pathol.* **2016**, 44, 267–278.

(41) Qin, D.; Lin, X.; Liu, Z.; Chen, Y.; Zhang, Z.; Wu, C.; Liu, L.; Pan, Y.; Laquerre, S.; Emery, J.; Fergusson, J.; Roland, K.; Keenan, R.; Oliff, A.; Kumar, S.; Cheung, M.; Su, D.-S. Discovery of Orally Bioavailable Ligand Efficient Quinazolindiones as Potent and Selective Tankyrases Inhibitors. *ACS Med. Chem. Lett.* **2021**, *12*, 1005–1010.

(42) Kwak, Y. H.; Barrientos, T.; Furman, B.; Zhang, H.; Puviindran, V.; Cutcliffe, H.; Herfarth, J.; Nwankwo, E.; Alman, B. A. Pharmacologic Targeting of β -Catenin Improves Fracture Healing in Old Mice. *Sci. Rep.* **2019**, *9*, No. 9005.

(43) Zhong, L.; Ding, Y.; Bandyopadhyay, G.; Waaler, J.; Börgeson, E.; Smith, S.; Zhang, M.; Phillips, S. A.; Mahooti, S.; Mahata, S. K.; Shao, J.; Krauss, S.; Chi, N.-W. The PARsylation Activity of Tankyrase in Adipose Tissue Modulates Systemic Glucose Metabolism in Mice. *Diabetologia* **2016**, *59*, 582–591.

(44) Plummer, R.; Dua, D.; Cresti, N.; Drew, Y.; Stephens, P.; Foegh, M.; Knudsen, S.; Sachdev, P.; Mistry, B. M.; Dixit, V.; McGonigle, S.; Hall, N.; Matijevic, M.; McGrath, S.; Sarker, D. Firstin-Human Study of the PARP/Tankyrase Inhibitor E7449 in Patients with Advanced Solid Tumours and Evaluation of a Novel Drug-Response Predictor. *Br. J. Cancer* **2020**, *123*, 525–533.

(45) Clinical-Trials-Registry/NCT01618136 and NCT04505839.

(46) Pereira, J. A.; Pessoa, A. M.; Cordeiro, M. N. D. S.; Fernandes, R.; Prudêncio, C.; Noronha, J. P.; Vieira, M. Quinoxaline, Its Derivatives and Applications: A State of the Art Review. *Eur. J. Med. Chem.* **2015**, *97*, 664–672.

(47) Baumeister, S.; Schepmann, D.; Wünsch, B. Synthesis and Receptor Binding of Thiophene Bioisosteres of Potent GluN2B Ligands with a Benzo[7]Annulene-Scaffold. *Med. Chem. Commun.* **2019**, *10*, 315–325.

(48) Quinn, L. A.; Moore, G. E.; Morgan, R. T.; Woods, L. K. Cell Lines from Human Colon Carcinoma with Unusual Cell Products, Double Minutes, and Homogeneously Staining Regions. *Cancer Res.* **1979**, *39*, 4914–4924.

(49) Lau, T.; Chan, E.; Callow, M.; Waaler, J.; Boggs, J.; Blake, R. A.; Magnuson, S.; Sambrone, A.; Schutten, M.; Firestein, R.; Machon, O.; Korinek, V.; Choo, E.; Diaz, D.; Merchant, M.; Polakis, P.; Holsworth, D. D.; Krauss, S.; Costa, M. A Novel Tankyrase Small-Molecule Inhibitor Suppresses APC Mutation–Driven Colorectal Tumor Growth. *Cancer Res.* **2013**, *73*, 3132–3144.

(50) Solberg, N. T.; Waaler, J.; Lund, K.; Mygland, L.; Olsen, P. A.; Krauss, S. TANKYRASE Inhibition Enhances the Antiproliferative Effect of PI3K and EGFR Inhibition, Mutually Affecting β -CATENIN and AKT Signaling in Colorectal Cancer. *Mol Cancer Res.* **2018**, *16*, 543–553.

Journal of Medicinal Chemistry

pubs.acs.org/jmc

(51) Waaler, J.; Machon, O.; Tumova, L.; Dinh, H.; Korinek, V.; Wilson, S. R.; Paulsen, J. E.; Pedersen, N. M.; Eide, T. J.; Machonova, O.; Gradl, D.; Voronkov, A.; Kries, J. P.; von Krauss, S. A Novel Tankyrase Inhibitor Decreases Canonical Wnt Signaling in Colon Carcinoma Cells and Reduces Tumor Growth in Conditional APC Mutant Mice. *Cancer Res.* **2012**, *72*, 2822–2832.

(52) WO2019243822A1.

(53) Narwal, M.; Fallarero, A.; Vuorela, P.; Lehtiö, L. Homogeneous Screening Assay for Human Tankyrase. *J. Biomol. Screen* **2012**, *17*, 593–604.

(54) Kabsch, W. XDS. Acta Crystallogr., Sect. D: Biol. Crystallogr. 2010, 66, 125–132.

(55) McCoy, A. J.; Grosse-Kunstleve, R. W.; Adams, P. D.; Winn, M. D.; Storoni, L. C.; Read, R. J. Phaser Crystallographic Software. *J. Appl. Crystallogr.* **2007**, *40*, 658–674.

(56) Émsley, P.; Cowtan, K. Coot: Model-Building Tools for Molecular Graphics. Acta Crystallogr., Sect. D: Biol. Crystallogr. 2004, 60, 2126–2132.

(57) Murshudov, G. N.; Skubák, P.; Lebedev, A. A.; Pannu, N. S.; Steiner, R. A.; Nicholls, R. A.; Winn, M. D.; Long, F.; Vagin, A. A. REFMAC5 for the Refinement of Macromolecular Crystal Structures. *Acta Crystallogr., Sect. D: Biol. Crystallogr.* **2011**, *67*, 355–367.



