DOI: 10.1002/ardp.202200473

FULL PAPER

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Replacing the oxidation-sensitive triaminoaryl chemotype of problematic K_V7 channel openers: Exploration of a nicotinamide scaffold

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Funding information

Deutsche Forschungsgemeinschaft, Grant/Award Numbers: BE 1287/6-2, LI 765/ 7-2

Abstract

 K_V7 channel openers have proven their therapeutic value in the treatment of pain as well as epilepsy and, moreover, they hold the potential to expand into additional indications with unmet medical needs. However, the clinically validated but meanwhile discontinued K_V7 channel openers flupirtine and retigabine bear an oxidation-sensitive triaminoraryl scaffold, which is suspected of causing adverse drug reactions via the formation of quinoid oxidation products. Here, we report the design and synthesis of nicotinamide analogs and related compounds that remediate the liability in the chemical structure of flupirtine and retigabine. Optimization of a nicotinamide lead structure yielded analogs with excellent $K_V7.2/3$ opening activity, as evidenced by EC_{50} values approaching the single-digit nanomolar range. On the other hand, weighted $K_V7.2/3$ opening activity data including inactive compounds allowed for the establishment of structure-activity relationships and a plausible binding mode hypothesis verified by docking and molecular dynamics simulations.

KEYWORDS flupirtine, KCNQ, K_v7, nicotinamide, retigabine

1 | INTRODUCTION

K_V7 (KCNQ) channels are homo- or heterotetrameric, voltage-gated potassium channels expressed in various tissues,^[1] whereby especially heterotetrameric neuronal K_V7 channels predominantly composed of K_V7.2 and K_V7.3 subunits are validated pharmacological targets.^[2] In general, K_V7 activation induces hyperpolarization of cell membranes, through which the ion channels contribute to controlling neuronal excitability. By increasing the action potential threshold^[3] and medium afterhyperpolarization while reducing spike frequency,^[4] K_V7 channels act as a "brake" for hyperexcitability.^[5] Moreover, their opening probability can be influenced by small-molecule ligands,^[6] making them attractive therapeutic targets, particularly for a range of neurological diseases.^[5]

For example, the administration of K_V7 channel openers was recently discussed for the therapy of various forms of brain damage, including chronic stress-induced brain injury (CSBI) as well as traumatic brain injury (TBI), for which currently no pharmacotherapeutic treatment options exist. In both cases, animal models suggest that reducing the underlying neuronal hyperexcitability by enhancing K_V7 -mediated potassium currents might offer a protective effect. Thus, K_V7 channel activation may be a novel therapeutic intervention

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strategy against post-TBI and CSBI brain damage.^[7] Furthermore, a recent review by Costi et al. examined the promising potential of K_V7 channel openers as novel antidepressants and concluded that existing preclinical and clinical studies provide initial evidence of a significant antidepressant effect.^[8] Consequently, their conclusions support the conduct of larger randomized controlled clinical trials to validate the potential of K_V7 channel openers as a treatment option for depressive disorders. These overall encouraging results are somewhat clouded by the fact that following the withdrawal of flupirtine (1, Figure 1) and retigabine (2, USAN: ezogabine) due to adverse drug reactions, no specific and safe K_V7 channel opener is currently available for therapeutic use in humans, which may hamper further development of this class of drugs as potential TBI/CSBI therapeutics or antidepressants.

Moreover, the failure of the two established drugs with their unique mechanism of action also leaves the desire for a replacement in the clinically validated indications of pain^[9] and epilepsy.^[10] For example, in the case of $K_{\rm V}7.2$ -related epilepsy due to loss-of-function variants, the use of K_V7 channel openers proved indispensable. As a result, there are currently calls for the reintroduction of retigabine for personalized treatment, despite its side effects that led to a withdrawal decision.^[11] A therapeutic gap with worrying consequences also remained in the field of pain therapy since metamizole, despite its known association with blood dyscrasias, is increasingly used in Germany as a substitute for flupirtine, which has led to a growing number of metamizole-induced neutropenia cases.^[12] Against this background, there is an urgent need for a safe replacement for flupirtine and retigabine to close the existing therapeutic gaps and enable the expansion of the therapeutic potential of K_V7 channel openers.

The reasons for the failure of flupirtine and retigabine have been analyzed in detail in our previous work.^[13,14] In essence, the adverse drug reactions responsible for withdrawal, particularly hepatotoxicity with flupirtine^[15] and discoloration of the skin and ocular tissues with retigabine,^[16] seem not related to the activation of K_v7 channels. Rather, they are most likely attributed to the oxidation-sensitive triaminoraryl scaffold both drugs have in common. As depicted in Figure 1, clear evidence points to the oxidative formation of reactive



FIGURE 1 Structures of flupirtine (**1**), retigabine (**2**), and their elusive *para* quinone diimine or azaquinone diimine oxidation products (**3**). *Ortho* quinone diimines or azaquinone diimines are also conceivable but not shown.

quinone diimine or azaquinone diimine metabolites such as 3 as the underlying cause of the adverse reactions in both cases.^[17,18] Consequently, a possible approach to obtain safer replacements of flupirtine and retigabine is to screen for entirely new chemotypes, as done in the case of the very recently published compound ZK-21 (8, Figure 2), which has a 4-aminotetrahydroquinoline scaffold.^[19] Moreover, the novel dual-mechanism K_V7 channel opener GRT-X (7) that activates both $K_{\rm V}$ 7 potassium channels and the mitochondrial translocator protein (TSPO) likewise has no noteworthy structural similarities with the triaminoaryl type K_V7 channel openers.^[20] Hence, new chemotypes such as 7 or 8 could represent a conceivable way to prevent the adverse effects that are presumably closely linked to the metabolically and chemically labile structure of flupirtine and retigabine. However, radical structural changes also increase the risk of unexpected new toxicities, as in the case of PF-04895162, a structural distinct K_V7 channel opener that was found to disrupt bile acid homeostasis and thus failed in phase I clinical trials.^[21]

A more conservative but straightforward approach to obtain safer K_V7 channel openers is to conduct minor structural changes in a ligand-based design. Such strategies have led to analogs with potent K_V7 opening activity like HN37 (4) or the retigabine analog **5**. Although both compounds demonstrated improved chemical stability compared to retigabine, their design does not entirely exclude the formation of quinone diimine oxidation products as it still includes a *para* diaminobenzene and a triaminobenzene structural motif, respectively.^[22]

Recently, our group also reported a ligand-based strategy, which in contrast to the approaches mentioned above, focused on completely avoiding structural motifs liable to quinoid metabolite formation. For this purpose, the substitution pattern of the central aromatic core of flupirtine and retigabine was redesigned.^[23] In



FIGURE 2 Selection of recently published K_V7 channel openers with scaffolds related to flupirtine and retigabine (4–6) and structurally distinct compounds with new chemotypes (7–9).

short, the oxidation-sensitive triaminoaryl structure was transformed into a nicotinamide scaffold without ortho/para-positioned nitrogen or oxygen atoms. However, further structural changes were necessary to maintain an acceptable Ky7.2/3 opening activity, such as the introduction of an additional methyl group together with a benzylic amide side chain. The resulting compound 6 demonstrated good $K_V 7.2/3$ opening activity with an EC₅₀ value of 0.310 µM, hence ranging between flupirtine and retigabine in terms of potency. Unfortunately, due to its lipophilic character $(\log D_{7.4} = 4.1)$ and low fraction of sp³ hybridized carbon atoms, the compound proved to be poorly soluble in water, which limited the toxicity testing and, thus, requires additional structural optimization. To overcome this shortcoming of analog 6 and further improve the $K_V 7.2/3$ opening activity, a hybridization approach was intended, using the water-soluble and highly potent analog 10 $(EC_{50} = 0.011 \,\mu\text{M})$ as a hybridization partner, which, however, is not devoid of the risk of azaquinone diimine formation (Figure 3).^[24] To anticipate a selected result of the biological testing, the hybridization product 18a showed promising K_V7.2/3 channel opening activity and thus served as a starting point for further structural modifications. For this purpose, the structure of **18a** was divided into five zones (Figure 3a-e), each of which was subjected to substitution with selected structural elements to investigate structure-activity relationships, shed light on a possible binding mode, and further improve the $K_V 7.2/3$ opening activity. Particular 6-morpholinonicotinamides such as 18a have been previously described in a patent by Grünenthal.^[25] Still, a comprehensive description of the underlying SARs and a basic toxicological characterization were lacking and could be provided by this study. In addition, we expanded the existing work to include new substituents not previously considered, some of which may not be covered by the referenced patent.

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2 | RESULTS AND DISCUSSION

2.1 | Chemistry

In the first step, the above-mentioned hybridization product 18a and analogous compounds derived from it with modifications predominantly located in zones A and E were synthesized. The conducted structural changes in zone A were primarily aimed at elucidating the role of the morpholine ring in molecular recognition by the K_V7.2/3 binding site. For this reason, the morpholine ring was replaced by carba analogs to help elucidate the possible role of the morpholine nitrogen atom. In particular, a tetrahydro-2H-pyran ring, a 3,6dihydro-2H-pyran substituent, and a pyridin-4-yl residue were considered as morpholine replacements by synthesizing analogs 17, 20, and 21 (Scheme 1). In general, while the morpholine ring is deemed a privileged structure with advantageous physicochemical, biological, and metabolic properties,^[26] in rare cases, it can be oxidized to reactive and potentially toxic iminium metabolites, as described for the multiple receptor tyrosine kinase inhibitor foretinib (not shown).^[27] Consequently, the carba analogs 17, 20, and 21 also represent alternatives that do not possess this potential metabolic lability. In addition, the strategy of substituting a heteroatom for a methine or methylene group was also applied by synthesizing the piperidine derivative 18e. Moreover, an attempt was made to further improve the water solubility of the analogs by reducing the compound lipophilicity. For this reason, a 2-oxa-6-azaspiro[3.3] heptane moiety was investigated in the case of analog 18f. This bioisosteric replacement for a terminal morpholine ring had been shown to lower the lipophilicity of a corresponding molecule effectively.^[28]

Regarding zone E, different strategies were followed. First, an additional effort was made to improve the aqueous solubility of the



FIGURE 3 Recently published K_V 7.2/3 openers 6 and 10, the hybridization approach conducted in this work, yielding analog 18a, and further structural modifications.

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SCHEME 1 Synthesis of nicotinamide analogs **14–21** with modifications in zones (a), (d), and (e). Reagents and conditions: (a) NaN₃, K₂CO₃, Cul, ethane-1,2-diamine, EtOH, Ar, reflux, 28 h, 54%; (b) 1. CDI, THF, 50°C, 1 h, 2. 4-fluorobenzylamine, THF, rt, 12 h, 35%; (c) morpholine, NMP, 165°C, µW irradiation, 1 h, 73%; (d) appropriate alcohol, NaH, THF, 0°C–70°C, 7–23 h, 96%–99%; (e) 1. (COCl)₂, DMF, DCM, 0°C–rt, 3 h, 2. amine, TEA, DCM, 0°C–rt, 16 h, 49%–61%; (f) pyridin-4-ylboronic acid, Pd(PPh₃)₄, Na₂CO₃, 1,4-dioxane, H₂O, Ar, 140°C, µW irradiation, 0.5 h, 59%; (g) amine, NMP, 165°C, 0.5–1 h, 28%–53%; (h) 1-cyclohexeneboronic acid pinacol ester, Pd(PPh₃)₄, Na₂CO₃, 1,4-dioxane, H₂O, Ar, 140°C, µW irradiation, 15 min, 86%; (i) H₂, Pd/C, MeOH, rt, 5 h, 74%.

analogs by reintroducing the primary amino group from flupirtine and retigabine in analog 14. Although the amino group, as a single ortho substituent of the amide function, led to inactive nicotinamide or benzamide analogs in our previous work, it has not yet been evaluated in combination with the advantageous ortho methyl group, which was also able to reconstitute the lost activity in the case of methoxy-substituted compounds such as 6.[23] Second, a different approach was aimed at improving the $K_V 7.2/3$ opening activity by varying the alkoxy residues. As described in our recent publication, the docking of **6** in combination with conformational analysis indicated that the methoxy group, together with the ortho methyl substituent, causes a favorable dihedral angle of the amide function.^[23] By synthesizing analogs **18b** and **18c**, it was investigated whether bulkier alkoxy substituents may further enhance the beneficial effect on K_V7.2/3 opening by increasing steric interactions and rotational energy barriers.

The initial series of nicotinamide derivatives was synthesized starting from 2,6-dichloro-4-methylnicotinic acid (**11**), whereby various substituents were introduced in position 2 of the pyridine ring in the first reaction step (Scheme 1). In particular, an amino group was attached via a copper-catalyzed Ullmann-type reaction by using sodium azide as a nitrogen source to obtain aminopyridine **12**. According to Zhao et al., the reaction is supposed to proceed via an intermediate azidopyridine (not shown), which is reduced directly in

situ to the corresponding primary amine **12**.^[29] In contrast, the introduction of alkoxy substituents, yielding **15a-c**, was performed via simple nucleophilic substitution reactions. The alcoholate, which served as the nucleophile, was generated in situ by reacting the corresponding alcohol with sodium hydride. Both reactions to introduce substituents at position 2 of the pyridine ring proceeded regioselectively at this specific position. The reason for the regioselectivity is probably a directing effect of the carboxyl function of **11**, which in both cases forms cyclic, five- or six-membered transition states, respectively, involving either a sodium alcoholate or a copper azide complex.^[29,30]

After introducing substituents at position 2 of the pyridine ring, the amide coupling allowed for the synthesis of all analogs. The activation of the nicotinic acid derivatives was performed either, as in the case of amino nicotinic acid derivative **12**, via the formation of an acyl imidazole after reaction with CDI or via generation of acyl chlorides in a DMF-catalyzed reaction with oxalyl chloride carried out in the case of **15a-c**. Both activated carboxylic acid species (not shown) were reacted directly, without prior isolation, with the corresponding amines to give the desired amides **13** and **16a-d**.

To introduce a substituent at position 6 of the pyridine ring, two different methods were principally applied. C-C bonds were formed via Suzuki coupling reactions by using the corresponding boronic acid or boronic acid pinacol ester to yield compounds **17** and **20**. The tetrahydropyran substituent in the case of analog 21 was obtained by catalytic hydrogenation of the dihydropyran precursor 20. In contrast, the introduction of amino substituents at position 6 in the case of compounds 18a-f was not carried out with a palladium catalyst but by nucleophilic substitution reactions, which were performed by microwave-assisted heating. In the case of all methoxy-substituted compounds (18a, d, e), a by-product was formed, causing low yields of the desired products. In the synthesis of compound **18a**, it was isolated and identified as the demethylated analog **19a**. Contrary to a methoxy residue, a 1-methyl-ethoxy and a 2,2,2-trifluoroethoxy substituent incorporated in compounds 18b/c were stable under identical reaction conditions. 2-Oxa-6-azaspiro [3.3]heptane oxalate required for synthesizing the spiro analog **18f** was prepared in a two-step synthesis according to a protocol shown).^[31] (not Briefly. described in the literature 4-methylbenzenesulfonamide was reacted with pentaerythritol tribromide to generate the spiro partial structure in a double-ring closure reaction. In the second step, the tosyl protective group was cleaved reductively with magnesium, and the resulting amine was precipitated as an oxalate salt. In the synthesis of compound **18f**, the free amine was released from its salt in situ by adding the stronger base DBU to the reaction mixture.

Since the pronounced structural change from a 4-fluorobenzylamino residue in **6** to a morpholino substituent in **18a** was possible without significantly affecting the activity, the question of the pharmacophore arises. Analog **29** (Scheme 2) was synthesized to clarify whether the morpholine ring is part of the core pharmacophore or, on the contrary, molecular recognition might even be possible without a morpholine ring. Analogously, derivatives **25** and **28a/b** were synthesized in which the methyl group or the alkoxy group were missing to clarify the role of the specific substituents, thus testing our hypothesis that ortho disubstitution of the amide group might be essential for K_v7.2/3 opening activity of nicotinamide analogs.

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In general, nicotinamide derivatives with missing substituents in positions 2, 4, or 6 of the pyridine ring were prepared by synthetic routes similar to those of the nicotinamide analogs 18a-f mentioned above. The synthesis of the nor-analog 25 was performed analogously to the synthesis of 18a with the difference that 2,6-dichloronicotinic acid (22) was used as starting material instead of 2,6-dichloro-4methylnicotinic acid (11). The amide coupling in the second reaction step to obtain 24 was done by using the coupling reagents DIC and HOBt (Scheme 2). In the case of the analogs 28a/b, in which the substituents in position 2 or positions 2 and 4 are absent, the introduction of the methoxy group in the first step was skipped. After the amide coupling, yielding 26a/b, the substitution reaction to introduce the morpholino substituent was carried out, which in this case required less drastic reaction conditions compared to the synthesis of analogs 18a-f. Conventional heating to 80°C instead of microwaveassisted heating to 165°C proved sufficient. Since two chloro substituents were accessible for the reaction, a mixture of two regioisomers was formed and subsequently separated by silica gel column chromatography to obtain the desired morpholino nicotinamides 27a/b and their corresponding regioisomers (not shown). Analytical discrimination of the regioisomers by NMR was not possible at this stage. However, after reductive cleavage of the chloro substituent in the following reaction step, confirmation of the identity of the desired analogs 28a/b based on ¹H-NMR spectroscopy was possible by analyzing the pyridine proton signals. In the case of 28a, two singlets were observed, unequivocally confirming the morpholino substituent in position 6 of the pyridine ring. The same applies to 28b, where a doublet with ${}^{3}J$ coupling (9.0 Hz), a doublet with ${}^{4}J$ coupling (2.5 Hz), and a doublet of doublets (J = 9.0, 2.5 Hz) verify the desired 6-morpholino nicotinamide structure while excluding morpholino substitution at position 2. Analogously, the chloro substituent of compound 17b was cleaved following the same catalytic hydrogenation procedure to obtain analog 29, which lacks a substituent in zone A.



SCHEME 2 Synthesis of nicotinamide analogs **25**, **28a/b**, and **29** lacking selected substituents of the central pyridine ring. Reagents and conditions: (a) MeOH, NaH, THF, 0°C–70°C, 7 h, 79%; (b) 4-fluorobenzylamine, DIC, HOBt, DMF, rt, 16 h, 65%; (c) morpholine, NMP, 165°C, μ W irradiation, 30 min, 36%; (d) 4-fluorobenzylamine, DIC, HOBt, DMF, rt, 16 h, 46%–86%; (e) morpholine, MeOH or 2-propanol, reflux, 1–3 d, 23%–26%; (f) H₂, Pd/C, TEA, MeOH, rt, 2–5 h, 64%–66%.

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Regarding zone B, a replacement of the central pyridine ring of the nicotinamide scaffold with a pyrimidine ring could be an interesting structural change since various examples exist where this bioisosteric exchange led to improved biological activity on different targets.^[32] In analogy to these literature cases, a pyrimidine ring with its altered electronic properties and reduced basicity may also influence the ligand binding to K_v7.2/3 channels. The envisioned pyrimidine analogs were accessible via the Biginelli multicomponent reaction, which in its original form, however, does not provide access to alkoxy-substituted compounds. For this reason, the resulting analogs **40a**-**c** contain an ethyl group instead of a methoxy substituent in zone E.

The initial Biginelli reaction, which involved an acid-catalyzed cyclocondensation of ethyl acetoacetate (32), propionaldehyde (31), and urea (30), yielded the 3,4-dihydropyrimidin-2(1H)-one 33, which was converted into the corresponding pyrimidin-2(1H)-one 34 in a subsequent oxidation step with nitric acid (Scheme 3). The following chlorination was carried out according to a modified procedure of Zhao et al. by heating 34 in excess phosphoryl chloride.^[33] In the course of the reaction, the 3,4dihydropyrimidin-2(1H)-one 34 probably forms an intermediate dichlorophosphate (not shown), as it was suggested for the chlorination of related guinazolin-4(3H)-ones.^[34] Subsequently, the dichlorophosphate was attacked by a chloride ion in an S_NAr reaction, yielding the chloropyrimidine **35**. The additional use of a base such as triethylamine. sometimes described in the literature for similar reactions,[35] has been evaluated but did not result in an improved yield. After the chlorination, two different synthetic routes were investigated. For the synthesis of analog 40a, the ethyl ester function of 35 was hydrolyzed under alkaline conditions, which gave the corresponding carboxylic acid 36. This was followed by a coupling reaction with HATU vielding amide **37**. In the final step, the morpholino substituent was introduced by applying the same microwave-assisted method used for the nicotinamides to obtain the final

analog **40a**. For the entire six-step synthesis of **40a**, a poor cumulative yield of only 2.8% was calculated. In retrospect, the ester hydrolysis in the first reaction step has proven particularly unfavorable because the yield was reduced by an undesired by-product. The side product was not isolated, but the chloropyrimidine structure of **35** was suspected to be affected by the alkaline hydrolysis leading to a pyrimidinone by-product. For this reason, the reaction sequence was rearranged, resulting in an alternative synthetic route for analogs **40b/c**. In particular, compared to the synthesis of **40a**, the morpholino substituent was introduced first to give compound **38**, while the hydrolysis of the ethyl ester to obtain carboxylic acid **39** took place in the following step since hydrolysis of the chloropyrimidine structure is thus not a concern. Overall, the modified reaction sequence and the use of DIC and HOBt instead of HATU as amide coupling reagents in the last step to obtain the final amides **40b/c** enabled an improved cumulative yield of 7.7%.

The carbamate partial structure of retigabine is supposedly involved in the formation of hydrogen bonds to the $K_{y}7.2$ binding site, which is also conserved in the case of nicotinamide derivative 6 as predicted by molecular docking.^[23,36] Apart from that, mainly hydrophobic interactions contribute to the binding of compound 6. Accordingly, the particular importance of the amide group can be anticipated, which was further explored by variation of the amide partial structure following three different approaches. The first structural change attempted in this direction was to replace the amide function with bioisosteric cyclic amide mimetics such as a 1,2,4-oxadiazole or a 1,2,4-triazole ring, resulting in analogs 47 and 54. A similar approach was recently used successfully in the optimization of a GPR88 agonist (not shown), where an amide bioisosteric replacement with a variety of azoles, followed by lead optimization, provided a potent and efficacious triazole-based GPR88 agonist.^[37] The second strategy was the incorporation of an amide-like structure into a fused ring system. Here, the isoxazolo[5,4-b]pyridin-3amine 44 was synthesized, representing a conformationally restricted



SCHEME 3 Synthesis of pyrimidine analogs **40a**-c. Reagents and conditions: (a) AcOH, EtOH, 90°C, 20 h, 32%; (b) HNO₃, H₂O, -10° C, 15 min, 87%; (c) POCl₃, 110°C, 1 h, 45%; (d) KOH, H₂O, THF, rt, 12 h, 72%; (e) 4-fluorobenzylamine, HATU, DIPEA, DMF, rt, 16 h, 42%; (f) morpholine, EtOH, 80°C, 2 h, 90%; (g) KOH, EtOH, H₂O, 80°C, 4 h, 84%; (h) morpholine, NMP, 165°C, μ W irradiation, 20 min, 73%; (i) amine, DIC, HOBt, DMF, rt, 16 h, 79%–81%.

analog. A comparable approach yielded, for example, highly potent benzo [d]isoxazol-3-amine-based sEH inhibitors with single-digit nanomolar IC₅₀ values from a benzamide lead structure.^[38] The third structural change carried out in this context was a shift of the amide group realized by transforming the *N*-benzylnicotinamide **18a** to the *N*-(pyridin-3-ylmethyl) benzamide analog **49**; thus altering the hydrogen-bonding ability as well as the flexibility and conformation in this part of the scaffold.

The synthesis of the isoxazolo[5,4-b]pyridin-3-amine analog 44 as a cyclic and conformationally restricted nicotinamide replacement could be realized in three steps (Scheme 4). The starting point was 2,6-dichloro-4methylnicotinonitrile (41), whose chloro substituent in position 6 was replaced by a morpholino residue through nucleophilic substitution. The reaction proceeded regioselectively with a preference for position 6. as reported by Antczak et al.^[39] Nevertheless, a small amount of the regioisomer substituted at position 2 was formed. The mixture of the two resulting regioisomers was separated by chromatography, and the desired regioisomer 42 was cyclized to afford the corresponding isoxazolo[5,4-b]pyridin-3-amine **43** by reaction with acetohydroxamic acid. This method involved a convenient one-pot procedure where the ortho chloro nicotinonitrile forms an N-[(3-cyanopyridin-2-yl)oxy]acetamide intermediate after nucleophilic substitution of the chloro substituent with a hydroxamate anion, followed by in situ base-catalyzed intramolecular cyclization and subsequent elimination of the acetyl residue.^[40] By reacting the amino group of compound 43 with 4-fluorobenzaldehyde, an imine (not shown) was formed in the last reaction step, which was reduced to the corresponding secondary amine without prior isolation by hydrosilvlation with triethylsilane and trifluoroacetic acid to obtain the final analog 44.

The synthesis of the oxadiazole derivative **47** began with the above-mentioned compound **42** (Scheme 4). In the first step, a methoxy group was introduced by substituting the remaining chloro substituent with sodium methoxide. The nitrile function of the resulting compound **45** was then converted to the corresponding amidoxime **46** by treatment with an aqueous solution of hydroxyl-amine. This reaction required an unusually long time (4 days) for

complete conversion, which is presumably attributed to the orthodisubstitution of the nitrile function and the resulting restricted steric accessibility. The electron-donating properties of the morpholino and methoxy substituents may also have reduced the reactivity of the nitrile function for nucleophilic attack by hydroxylamine. While most syntheses of oxadiazoles reported in the literature proceed via amidoxime precursors such as 46, the methods differ in the use of diverse activated carboxylic acid derivatives as acylating agents and in the reaction conditions applied for ring closure.^[41] In this particular case, the amidoxime 46 was acylated with phenylacetyl chloride. The corresponding acylation product (not shown) was not isolated, but a ring closure was induced directly by the addition of tetrabutylammonium fluoride (TBAF) as a catalyst. Mechanistically, the ring closure reaction of the O-acyl amidoxime intermediate to the final oxadiazole analog 47 is supposed to proceed via an intramolecular attack of the amidoxime nitrogen atom on the carbonyl carbon atom of the acyl group, followed by dehydration of the resulting dihydro-oxadiazolol. Both steps are facilitated by TBAF, whose fluoride ion acts as a strong base in polar aprotic solvents such as THF.^[42]

The *N*-(pyridin-3-ylmethyl)benzamide analog **49** with a shifted carbonyl group compared to the initial nicotinamide scaffold was synthesized from the nicotinonitrile **45** in two steps. First, the nitrile group of **45** was reduced by catalytic hydrogenation with Raney nickel as a catalyst to obtain the primary amine **48**. Then, in the second step, **48** was acylated by reaction with 4-fluorobenzoyl chloride, yielding the final analog **49**.

The above-mentioned nicotinic acid derivative **15a** was also used as a starting point for synthesizing the triazole derivative **54** (Scheme **5**). In the first step, the corresponding methyl ester **50** was obtained by alkylation of **15a** with iodomethane. This was followed by substituting the chlorine atom of **50** to afford the morpholino derivative **51**, whose methyl ester function was subjected to hydrazinolysis yielding the hydrazide **52**. The following triazole formation proved problematic since two attempts to achieve a ring closure reaction were initially unsuccessful. The first approach

SCHEME 4 Synthesis of analogs **44**, **47**, and **49** with modifications in zone C. Reagents and conditions: (a) morpholine, MeOH, 0°C-rt, 16 h, 70%; (b) acetohydroxamic acid, *t*-BuOK, Ar, DMF, RT–50°C, 5.5 h, 43%; (c) 4-fluorobenzaldehyde, $(C_2H_5)_3$ SiH, TFA, DCM, RT–60°C, 25 h, 80%; (d) NaOMe, MeOH, reflux, 24 h, 90%; (e) hydroxylamine (aq.), EtOH, reflux, 4 d, 99%; (f) 1. 2-phenylacetyl chloride, TEA, DCM, 0°C, 1 h, 2. TBAF, THF, rt, 2 h, 36%; (g) Ni, H₂, NH₃, MeOH, 50°C, 5 h, 71%; (h) 4-fluorobenzoyl chloride, TEA, DCM, rt, 16 h, 64%.



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SCHEME 5 Synthesis of the 1,2,4-triazole analog **54**. Reagents and conditions: (a) CH₃I, K₂CO₃, DMF, rt, 8 h, 98%; (b) morpholine, TEA, NMP, 90°C, 2 d, 39%; (c) N₂H₄ × H₂O, MeOH, reflux, 24 h, 67%; (d) NaOMe, MeOH, 0°C-rt, 24 h, 47%; I CuBr, Cs₂CO₃, DMSO, 120°C, 24 h; (f) K₂CO₃, *n*-BuOH, 150°C, μ W irradiation, 4 h, 47%; (g) 220°C, 0.1 mbar, 10 min; (h) HCl, MeOH, 0°C, 3 h, 58%; (i) NH₃, MeOH, rt, 24 h, 61%.

was based on the reaction of hydrazide **52** with the Pinner salt **56** to give the *N'*-(1-iminoalkyl)-hydrazide **53**. According to the literature, ring closure to obtain the triazole structure should occur by heating compounds like **53** to the melting point under reduced pressure without solvent.^[43] However, in this particular case, the procedure provided a complex mixture of products with little, if any, yield of the desired product according to TLC reaction control. A second unsuccessful attempt was based on a copper-catalyzed tandem addition-oxidative cyclization reaction described by Ueda and Nagasawa.^[44] Unfortunately, in this specific case, the reaction between nitrile **45** and amidinium salt **57**, previously prepared by ammonolysis of the Pinner salt **56**, did not lead to any product formation. Finally, the triazole synthesis succeeded by applying the method of Yeung et al., which includes a one-step, base-catalyzed condensation of the hydrazide **52** with 4-fluorobenzonitrile **(55)**.^[45]

2.2 | Pharmacology/biology/modeling

2.2.1 | Evaluation of K_V7.2/3 opening activity

The K_V7.2/3 opening activity was determined by applying a commercially available assay on HEK293 cells that overexpress the heterotetrameric K_V7.2/3 channel. The assay exploits the permeability of potassium channels for thallium ions and is based on a thallium-

sensitive fluorescent dye, which is trapped inside the cell after esterase cleavage. Thallium influx through K_V7.2/3 channels generates a fluorescence signal whose intensity correlates with the extent of channel opening. Based on the data generated this way, EC_{50} values were calculated, indicating the concentration at which the half-maximum fluorescence signal and, thus, the half-maximum K_V7.2/3 opening activity was achieved. In addition to EC_{50} values, the efficacy, that is, the maximum response, of the analogs relative to flupirtine was calculated. Both values can be found in Table 1 for all analogs. Based on these results, structure-activity relationships were derived, which are systematically discussed below.

Regarding zone A, the replacement of the 4-fluorobenzylamino residue of 6 with a morpholine ring in 18a improved the biological activity as intended by the hybridization approach. In particular, the EC_{50} value of 18a decreased compared to 6 from 0.310 to 0.117 μ M, and at the same time, the efficacy increased from 105% to 144%. However, the difference in potency does not appear to be significant, and the reported outstanding K_V7.2/3 opening activity of the second hybridization partner 10 was not quite reached. In contrast, the complete deletion of a substituent in zone A resulted in a total loss of activity in the case of analog 29, which clearly confirmed the morpholine ring as part of the core pharmacophore.

Both heteroatoms of the morpholino substituent, on the other hand, do not appear to be of essential importance for $K_V7.2/3$ opening. This was evident in the case of the morpholine oxygen atom, where a replacement with a methylene group even led to improved $K_V7.2/3$ opening. The corresponding piperidine derivative 18e exhibited a 2.9-fold reduced EC₅₀ value compared to the direct morpholino counterpart 18a. This observation correlates with molecular docking results (Figure 4b,c), which predicted the morpholine ring to occupy a hydrophobic cavity with no involvement of the oxygen atom in any hydrogen bond interactions.

In contrast, when considering only the activity data, it was initially presumed that the morpholine nitrogen atom might contribute substantially to binding since the corresponding ydroxy analog 21, bearing a tetrahydropyran ring, showed a considerably 20-fold decrease in potency compared to the direct morpholino congener 18a. However, the dihydropyran derivative 20, which also has a carbon atom in place of the morpholine nitrogen atom, was approximately as potent as 18a. This observation, combined with molecular docking results displayed in Figure 4, in turn, rather implies that the tertiary amino group of the morpholine ring does not participate in direct interactions with the Ky7.2/3 binding site but instead might favor an advantageous molecular geometry. Specifically, the morpholine nitrogen nonbonding pair of electrons interacts with the electron-deficient pyridine π -system and thus probably favors a coplanar conformation, which is also assumed to be probable for the dihydropyran moiety of 20 since the double bond is in conjugation with the adjacent pyridine ring. These conformational considerations were confirmed by quantum mechanical calculations using density functional theory (DFT), which revealed energetic minima at a dihedral angle of approximately 0/180° and high rotational energy barriers for a morpholino as well as a dihydropyran

TABLE 1 K_V7.2/3 channel opening activity, in vitro toxicity, and log $D_{7.4}$ values of the synthesized compounds **14–54** in comparison to flupirtine and retigabine^[a]

Entry	LogD _{7.4}	HEK-293 EC ₅₀ ^[b] (μM)	Efficacy (%)	ΤΑΜΗ LD ₅₀ ^[c] (μΜ)	LD ₂₅ ^[d] (µM)	HEP-G2 LD ₅₀ ^[c] (μM)	LD ₂₅ ^[d] (µM)	LD ₂₅ /EC ₅₀ ^[f]
Flu ^[g]	2.0	1.837 ± 0.844	100	487 ± 51	103 ± 47	547 ± 111	74 ± 40	40
Ret ^[g]	2.1	0.249 ± 0.052	134 ± 16	>400	>400	>400	269 ± 166	1080
6 ^[g]	4.1	0.310 ± 0.119	105 ± 12	>63	>63	>16	>16	52
10 ^[g]	3.6	0.011 ± 0.004	111 ± 7	>500	212 ± 140	>500	231 ± 141	19273
14	2.2	6.858 ± 1.319	113 ± 28	>500	349 ± 36	>500	123 ± 73	18
17	2.7	0.500 ± 0.105	112 ± 11	>63	>63	>63	26 ± 20	52
18a	3.0	0.117 ± 0.029	144 ± 11	>63	>63	>63	>63	539
18b	3.8	0.017 ± 0.009	132 ± 16	>31	>31	>31	15±1	882
18c	3.3	0.012 ± 0.004	117 ± 19	>15	>15	>15	>15	1250
18d	2.6	3.799 ± 1.730	170 ± 4	>250	>250	>250	>250	66
18e	4.7	0.040 ± 0.007	127 ± 1	>31	>31	>31	32±11	775
18 f	3.8	_ [e]	_ [e]	>63	>63	>63	>63	-
19a	2.4	_ [e]	_ [e]	>63	30 ± 17	>63	28±16	-
20	3.1	0.143 ± 0.003	111 ± 11	>125	>125	>63	>63	440
21	3.1	2.402 ± 0.759	129 ± 16	>125	>125	>125	>125	52
25	3.8	_ [e]	_ [e]	>125	77 ± 35	>125	75 ± 24	-
28a	2.4	8.632 ± 1.876	76±19	>250	169 ± 29	>250	126 ± 36	15
28b	2.3	_ [e]	_ [e]	>125	>125	>125	>125	-
29	2.6	_ [e]	_ [e]	>125	13±3	>125	>125	-
40a	3.0	0.126 ± 0.035	114 ± 10	>31	>31	>31	20±12	159
40b	3.5	0.035 ± 0.028	104 ± 12	>31	>31	>31	>31	886
40c	3.4	2.134 ± 0.591	52±11	>63	>63	>63	>63	30
44	3.2	_ [e]	_ [e]	>125	>125	>125	>125	-
47	4.7	1.179 ± 0.193	45 ± 4	>30	>30	>30	>30	25
49	3.6	_ [e]	_ [e]	>31	>31	>63	29 ± 5	-
54	3.3	2.245 ± 0.338	149 ± 25	>30	>30	>30	>30	13

Note: [a] LogD_{7.4} values were estimated by employing an HPLC-based method. HEK293 cells overexpressing the K_V7.2/3 channel were used to obtain the EC₅₀ values by applying a fluorimetric assay. LD values were determined by an MTT assay in TAMH and HEP-G2 cell lines after 24 h of exposure. EC₅₀ and LD values are means of \geq 3 independent determinations ± standard deviations. [b] Necessary concentration to reach half-maximal K_V7.2/3 channel opening activity. [c] Concentration required to reduce cell viability to 50% compared to untreated controls. [d] Concentration required to reduce cell viability to 75% compared to untreated controls. [e] No K_V7.2/3 channel opening activity up to a concentration of 20 µM. [f] The lower of both LD₂₅ values was used to calculate the LD₂₅/EC₅₀ ratio. If no LD₂₅ value could be determined, the maximum tested concentration was used for calculation. [g] Previously published values.^[14,23,24]

substituent (Figure 5a,d). In contrast, a twisted conformation is probably energetically preferred by the 2-(tetrahydro-2*H*-pyran-4-yl) pyridine moiety of **21** owing to the sp³ hybridized ydroxy carbon atom, as verified by the dihedral scanning plot, which shows a global minimum at 60° and low rotational energy barriers (Figure 5c). Viewed in correlation with the corresponding docking poses, which also indicate a coplanar orientation of the pyridine ring with the adjacent heterocyclic substituent, the different molecular geometries suggest that the inferior $K_V7.2/3$ opening activity of **21** compared to **18a** and **20** is mainly due to a conformational preorganization of **18a** and **20**, which may reduce the entropic costs of ligand binding.

Regarding the other zone A substituents, an aromatic residue, such as a pyridine ring in analog **17**, was also found to be tolerated since the incorporated biaryl structure is expected to favor a coplanar conformation, too (Figure 5e). However, despite the beneficial coplanarity, compound **17** was still 4.3-fold less potent than the

DPhG Arch Pharm Archiv der Pharmazie (a) F305 F305 (b) (C) F305 F240 F740 F740 L338 L338 1339 1339 L338 1339 W236 W236 W236 F343 F343 F343 L225 L225 L221 L225 L221 L221

Predicted binding poses of retigabine (a), 14 (b), and 18a (c). The hydrogen bonds to W236 and S342 are maintained for all FIGURE 4 compounds, although the central pyridine ring is slightly displaced compared to retigabine. Therefore, the primary amino group of 14 does not participate in hydrogen bond formation with S342 as predicted for retigabine. Instead, it is shifted into a more hydrophobic cavity, which is also occupied by the methyl substituent of 18a. The 4-fluorobenzyl group binds to a larger hydrophobic pocket formed by L221, L225, and F343 as previously reported for other derivatives.^[23] The π - π interactions to W236 (represented as blue dashed lines) can be observed for all three ligands during molecular dynamics simulations, depending on the distinct orientation of the aromatic rings. The color of the secondary structure elements represents the respective chains of $K_{y}7.2$ (gold) and $K_{y}7.3$ (silver) in the $K_{y}7.2/3$ -heterotetramer.



FIGURE 5 Dihedral angle scanning for various pyridine scaffolds using B3LYP-D3/6-31G(d,p). The calculations were performed with Jaguar version 11.5 (Schrödinger, LLC, 2022).^[46]

direct morpholino analog 18a, but this may be attributed to other reasons, such as the flat and rigid geometry of the 4-pyridyl moiety, which deviates considerably from the chair conformation expected for a morpholine ring.^[47] Consequently, this specific conformation of the morpholino substituent, which a pyridine substituent cannot adopt, might be required for good Ky7.2/3 channel opening since it is also observed in molecular dynamics simulations of corresponding analogs (Figure 4b,c).

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Despite the presence of a tertiary amino partial structure as with a morpholino substituent, the spiro analog 18f showed no Ky7.2/3

opening activity up to 20 µM. Presumably, the 2-oxa-6-azaspiro[3.3] heptane substituent is slightly too long for an adequate fit into the $K_{\rm V}7.2/3$ binding pocket since the distance between the nitrogen and the oxygen atom is 4.3 Å compared to 2.8 Å for morpholine.^[28] Moreover, the substituent has a rather linear geometry and differs significantly from the chair conformation of morpholine, which is presumed to be favorable for Ky7.2/3 opening. Regarding physicochemical properties, contrary to cases reported in the literature, the introduction of the 2-oxa-6-azaspiro[3.3]heptane substituent did not result in a decreased lipophilicity, as originally intended by this

structural change. Indeed, as with the corresponding morpholino substituted compound **18b**, the $logD_{7,4}$ value of the spiro analog **18f** was determined to be 3.8.

Concerning zone B, very potent pyrimidine analogs with nanomolar $K_V7.2/3$ opening activity were obtained (**40a/b**), indicating that a central pyrimidine ring is well tolerated. However, since the corresponding analogs also bear an alkyl residue in zone E instead of an alkoxy substituent due to the chosen synthesis route via the Biginelli reaction, no final, conclusive statement can be made with certainty about the influence of the pyrimidine ring. Nevertheless, the almost identical EC₅₀ values of the closely related analogs **18a** and **40a** suggest that the pyrimidine ring did not significantly affect the $K_V7.2/3$ opening activity compared to a pyridine core, which is also supported by very similar docking poses of the corresponding analogs with no specific interaction of the additional pyrimidine nitrogen atom.

Contrary to the possible role of the pyrimidine ring, however, a valid judgment can be made on the importance of the methyl substituent in zone B, which must be considered in conjunction with the second ortho substituent of the amide group in zone E. In particular, compound 25, which differs from the submicromolar active analog **18a** only in the absence of the methyl group, as well as compound 28b, which lacks both ortho substituents of the amide group, were inactive up to a concentration of 20 µM. In contrast, compound 28a, which has a methyl substituent attached to the pyridine core but lacks the second ortho substituent in zone E, was still active, albeit weakly potent, with a 70-fold reduced EC₅₀ value compared to 18a. Based on these observations, two conclusions can be drawn concerning the possible role of the methyl group in zone B. First, viewed in isolation, the methyl group seems to be slightly more important for Ky7.2/3 opening than the second ortho substituent of the amide group since the methylated analog 28a was still weakly active while 25 and 28b, both lacking the methyl group, were completely inactive. Second, however, only the presence of both amide ortho substituents, as in the case of 18a, allows sufficient $K_{V}7.2/3$ opening activity with submicromolar potency. Essentially, these results complete the evidence supporting our earlier hypothesis that the ortho disubstitution of the amide function is necessary to favor a molecular geometry that matches the bound-state conformation of the nicotinamide analogs in which the amide group is rotated out of the aromatic plane.^[23] In contrast, a methoxy residue as the only amide ortho substituent, as in 25, probably stabilizes an unfavorable coplanar orientation of the amide group through intramolecular hydrogen bonding. This negative effect is impossible with a methyl group as the sole ortho substituent, which is why the methylated analog 28a, in contrast to compound 25 bearing a methoxy residue, was at least weakly active.

By modifying zone C, the possible role of the N-substituted amide group was investigated, which is presumed to be a crucial structural element of the initial nicotinamide scaffold. A first structural change based on the incorporation of an amideanalogous partial structure into an isoxazolo[5,4-*b*]ydroxyl-3-amine scaffold proved clearly detrimental to $K_V7.2/3$ opening since the

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corresponding analog **44** was inactive up to a concentration of $20 \,\mu$ M. This observation essentially confirms the hypothesis formulated above that the amide group in the bound state conformation is probably rotated out of the pyridine plane, which of course, is not possible in the case of the conformationally restricted isoxazolo[5,4-*b*]pyridine analog **44**.

A different attempt to replace the amide function of 18a with heterocyclic amide bioisosteres was slightly more successful, yielding the weakly to moderately active compounds 47 and 54, which provided valuable SAR insights despite their overall inferior Ky7.2/3 opening activity. As suggested by docking and molecular dynamics simulations, the amide group of nicotinamide analogs could be involved in direct interactions with the $K_V 7.2/3$ binding site as both a hydrogen bond donor and acceptor. In particular, the role of the amide function as a hydrogen bond donor was partially elucidated by the cyclic amide bioisosteres since the triazole ring of analog 54 provides a hydrogen atom similar to an amide group. In contrast, the oxadiazole structure of 47 is not able to act as a hydrogen bond donor. Consistent with this assumption, docking predicted the triazole ring of 54 to mimic the original amide group by forming both hydrogen bond interactions, and moreover, it may even be involved in a third hydrogen bond to S342 (Figure 6b). Contrary, the docking pose of the oxadiazole analog 47 shows a different orientation due to the lack of hydrogen bonding with I339 and S342 (Figure 6a). Albeit the hydrogen bond donor ability seems not essential for Ky7.2/3 opening, it may still play an important role, as the triazole analog 54 had a superior efficacy of 149%, whereas the oxadiazole analog 47 was on the verge of inactivity with an efficacy of only 45%. Presumably, the putative hydrogen bond interactions of the triazole derivative 54 result in a slight shift of the S6' alpha helix. which is part of the pore-forming domain, thereby enhancing the $K_{V}7.2/3$ opening of 54 compared to the oxadiazole analog 47.

In addition to these findings regarding the possible role of a hydrogen bond donor ability, analog **54** also provided a novel substance class of $K_V7.2/3$ openers since, to our knowledge, **54** is the first 1,2,4-triazole derivative described to address this target. Indeed, there is still room for improvement in terms of potency, but considering the superior efficacy of **54**, it was even slightly more active than flupirtine, indicated by an overall left-shifted concentration-activity curve.

The third structural modification affecting the amide region again resulted in an inactive compound. The *N*-(ydroxyl-3-ylmethyl) benzamide analog **49** with a displaced amide carbonyl group compared to the original nicotinamide scaffold demonstrated no K_v7.2/3 opening activity up to 20 μ M, thus underlining the crucial importance of the amide group for K_v7.2/3 opening and confirming the postulated binding mode. By shifting the carbonyl group of the original amide partial structure while maintaining the position of the amide NH moiety, two important mechanisms contributing to ligand binding are disrupted at once. In detail, steric interactions of the amide group with ortho substituents on the pyridine core are vastly reduced, and one of the two presumed hydrogen bonds of the amide function is also prevented.



FIGURE 6 Predicted binding poses of **47** (a) and **54** (b). The triazole ring forms hydrogen bonds to the S342 sidechain and the I339 backbone carbonyl oxygen atom, probably affecting the S6' alpha helix, which is part of the pore-forming domains. The color of the secondary structure elements represents the respective chains of $K_V7.2$ (gold) and $K_V7.3$ (silver) in the $K_V7.2/3$ -heterotetramer.

According to molecular docking, the amide side chain in zone D is likely to occupy a hydrophobic cavity formed essentially by L221, L225, and F343, which has so far been mainly addressed with benzyl moieties. However, from a physicochemical point of view, this type of amide side chain, inherited from lead 6, remains a certain hindrance to the desired increased aqueous solubility. Therefore, while incorporating polar structural elements in this region of the scaffold did not seem expedient, attempts were made to replace the benzyl mojety with aliphatic side chains to increase the overall sp^3 fraction and thus possibly improve the water solubility. The corresponding compounds 18d and 40c were both found to be moderately active with EC₅₀ values slightly inferior to flupirtine but showed a remarkable difference in efficacy. Whereas analog 40c bearing a bulky 3,3-dimethylbutyl sidechain achieved only 52% efficacy, an nbutyl sidechain in 18d resulted in an impressive 170% efficacy, which was the best of all compounds tested in this study (Figure 7). Consequently, despite the clear negative impact on the EC₅₀ value, an n-butyl sidechain may still represent an attractive structural element for future designs since analog 18d not only showed the best efficacy but was also one of the most soluble substances in the toxicity tests carried out where it could be tested up to 250 µM. The detrimental effect of the *n*-butyl amide side chain on the EC_{50} value might be partially compensated when combined with successful substituents in zones A and E of the scaffold. However, considering only the Ky7.2/3 opening activity while leaving aside physicochemical properties, a benzylic amide side chain was still clearly superior. The beneficial effect could even be enhanced by replacing the initially used 4-fluorobenzyl moiety with a 3-(trifluoromethyl) benzyl residue, as revealed by a 40a and 40b comparison, where this structural change led to a 3.6-fold increase in potency. The observed boost in K_V7.2/3 opening activity might be attributed to a π - π interaction between the benzyl side chain and F343 being strengthened upon trifluoromethylation. This is consistent with computational studies by Mottishaw and Sun, which demonstrated that trifluoromethylation of an aromatic core results in improved π - π



FIGURE 7 The concentration-activity curves for K_V7.2/3 opening of analogs **18c** and **18d** in comparison to flupirtine demonstrate the excellent potency of **18c** (EC₅₀ = 0.012 μ M) as well as the superior efficacy of **18d** (E_{max} = 170%).

interactions compared to direct monofluorination, as indicated by increased intermolecular interaction energies and reduced π - π distances.^[48]

Regarding the investigated substituents in zone E, the impression emerged that sterically more demanding substituents with increased lipophilicity favor K_V7.2/3 opening. A replacement of the methoxy group of **18a** with a 2,2,2-trifluoroethoxy or a 1-methyl-ethoxy substituent increased the potency of the corresponding analogs **18b** and **18c** significantly by a factor of 6.9 and 9.8, respectively, leading to an excellent EC_{50} value of 0.012 μ M in the case of **18c** (Figure 7). This is noteworthy since the predicted binding poses of the nicotinamide analogs indicate that the alkoxy substituents do not interact directly with the binding pocket but rather point in the opposite direction toward the vicinity of the K_V7.2/3 channel (Figure 4c). Consistent with the assumption about the orientation of the alkoxy substituents, an exchange of the methoxy group for an isosteric ethyl residue in analog **40a** was also tolerated well, as there are no specific interactions of the alkoxy oxygen atom with the K_V7.2/3 binding site predicted.

However, without direct contact with the binding site, the question arises of how the bulky and lipophilic alkoxy substituents were able to improve K_V7.2/3 opening. Remarkably, the position of the $K_V 7.2/3$ binding site may be relevant in this case since it is situated in the transmembrane region of the channel on the proteinphospholipid interface (Figure 8). Hence, the bulky and lipophilic alkoxy substituents in zone E, which are presumed to face in the direction of the cell membrane, may improve possible interactions with adjacent lipid tails. Such ligand-lipid interactions have so far been underestimated as influencing factors of drug activity. However, they are becoming increasingly important in drug design as a growing number of intramembrane binding sites are revealed by X-ray crystallography and cryogenic electron microscopy.^[49] Consequently, for various small molecules and targets, it is assumed that ligand-lipid interactions are significantly involved in ligand binding. A prominent and impressive example of ligand-lipid interactions is ivacaftor (not shown), a potentiator of the cystic fibrosis transmembrane conductance regulator (CFTR) that binds to a site within

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the transmembrane region of the CFTR chloride channel. The molecule bears two *tert*-butyl groups that interact with lipids of the adjacent membrane and shield the polar part of the ligand, which is responsible for molecular recognition via hydrogen bonds, against the phospholipid interface.^[49] Conceivably, the lipophilic alkoxy substituents of **18b** and **18c** function in a similar way as the *tert*-butyl groups of ivacaftor. To confirm the postulated ligand–lipid interactions, a molecular dynamics simulation was performed with **18c** and a K_v7.2/3 channel, which was inserted in a phosphatidylcholine bilayer simulating the cell membrane. The result presented in Figure 8 clearly shows that the trifluoroethoxy group, together with the 4-fluorophenyl ring, is able to shield the polar part of the molecule, that is, the amide group and the pyridine nitrogen atom, from the adjacent lipid layer, thus improving hydrophobic interactions with lipid tails.

In contrast to the alkoxy substituents, the considerably more hydrophilic primary amino function of 14 resulted in a reduced logD_{7.4} value and, as intended, in an improved solubility in toxicity testing, but at the same time caused an unexpected drastic loss of $K_{\rm V}7.2/3$ opening activity. Consequently, the question arises why the primary amino function is well tolerated in flupirtine and retigabine, whereas it appears to be clearly disadvantageous in nicotinamide analogs. In this case, there are probably several reasons to consider. First, molecular dynamics simulations indicate that the amino function of 14 is slightly shifted in the binding pocket compared to the amino group of retigabine. As a result, the hydrogen bond to \$342, which is assumed to be advantageous, is not predicted as with retigabine (Figure 4b). Second, in the case of 14, the methyl group is located on the lipid-facing side of the molecule. Therefore, the shielding against the cell membrane is likely reduced compared to bulky lipophilic alkoxy residues of analogs 18b/c. Finally, with a logD_{7.4} value of 2.2, compound 14 is also the most hydrophilic of the



FIGURE 8 Predicted binding mode of **18c** from combined ligand docking and all-atom molecular dynamics simulations. The trifluoroethoxy group points toward the membrane and shields it from the hydrophilic parts of the ligand.

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nicotinamide analogs, which may inhibit diffusion through the lipid layer and thus impede the compound from reaching the intramembrane binding site.

The same may also apply to analog 19a, which has a ydroxyl group in place of the primary amino substituent of 14. Remarkably, in contrast to the still weakly active compound 14, analog 19a turned out to be completely inactive up to a concentration of 20 µM. The even worse performance of a ydroxyl group compared to the amino substituent of 14 might be attributed to a tautomeric relationship of 19a, whose 2-hydroxypyridine form is likely in equilibrium with the 2-pyridone form. This is particularly relevant as docking suggests that the central pyridine ring of nicotinamide analogs is involved in π - π interactions with the W236 indole side chain, similar to the phenyl ring of retigabine (Figure 4). This specific interaction is presumed to be highly relevant for ligand binding since it could be shown that a mutation of W236 completely abolishes the effect of retigabine.^[50] Consequently, this crucial interaction might be weakened in the case of the nonaromatic 2-pyridone tautomer of **19a**, hence leading to the observed inactivity.

2.2.2 | Evaluation of in vitro toxicity

Flupirtine-induced severe hepatotoxicity is a very rare phenomenon with a reporting rate of 1.68 cases per 100,000 patient years.^[51] For this reason, it was not reported in any in vitro or in vivo toxicity studies during preclinical and clinical development, and thus flupirtine has long been considered a well-tolerated analgesic.^[9,52] In accordance, in vitro LD₅₀ values of flupirtine were determined in the range of 500 µM, hence significantly exceeding therapeutically relevant concentrations, which are in the low single-digit micromolar range.^[53] In general, idiosyncratic toxicity, as suspected for flupirtine, is very difficult to reproduce in an in vitro toxicity model due to the multifactorial causes,^[54] which in the case of flupirtine probably include both the formation of potentially toxic metabolites and involvement of the adaptive immune system.^[18,55] Despite all efforts, currently, no in vitro or in vivo model exists that reliably predicts idiosyncratic drug-induced liver injury (DILI),^[56] which is reflected in the fact that idiosyncratic hepatotoxicity is still one of the two most common reasons for drug withdrawals, restrictions and project terminations.^[57] For these reasons, since flupirtine usually behaves uncritically in standard toxicity tests and no adequate assay for idiosyncratic hepatotoxicity exists, the toxicological evaluation carried out in this work should not be understood as a proof of concept to verify the design hypothesis but rather as a general examination of the new scaffolds for potential intrinsic toxicity. This standard toxicological characterization was carried out on two hepatic cell lines with an MTT assay, which essentially measures metabolic activity as an indicator of cell viability. The method is based on the mitochondrial reduction of 3-(4,5-dimethylthiazol-2-yl)-2,5diphenyltetrazolium bromide (MTT) to the corresponding formazan, which can be quantified colorimetrically.^[58] The human HEP-G2 and mouse TAMH cell lines used for this assay are well established for in vitro hepatotoxicity testing.^[59]

A limitation in toxicity testing that inherently demands that much higher concentrations are applied than in activity testing continued to be the relatively poor water solubility of most analogs, which was already problematic for compound 6.^[23] Remarkably, the introduction of a morpholino substituent in **18a** did not result in significantly improved solubility. Like analog 6, the morpholino derivative 18a could only be tested up to a concentration of $63 \,\mu$ M, which was surprising since the $log D_{7.4}$ value was reduced from 4.1 to 3.0, and at the same time, the fraction of sp³ hybridized carbon atoms was increased from 0.18 to 0.37. However, the three-dimensional structure of 18a and the other morpholino-substituted analogs is probably still rather flat due to the coplanar conformation of the morpholino substituent and the central aromatic ring, which might favor aggregation and thus prevent noticeably improved water solubility. This hypothesis is supported by the increased aqueous solubility of analog 21, which could be tested up to a concentration of 125 µM and was thus about twice as soluble as 18a. The reason for this difference is probably the tetrahydropyran ring of 21, which does not prefer a coplanar conformation like the morpholine ring of 18a, as discussed above. Since no LD₅₀ value could be determined for any analog due to the limited water solubility, LD₂₅ values were calculated, which indicate the concentration that was necessary to reduce the cell viability to 75% to enable a comparison of the analogs regarding their hepatotoxic potential. Nevertheless, for 12 of the 22 analogs, no LD₂₅ value could be determined for either cell line because the required concentrations could not be reached. However, apart from the poor water solubility, the lack of LD values principally indicates that most of the analogs do not possess a pronounced level of in vitro hepatotoxicity at concentrations required for Ky7.2/3 opening.

While it was possible to derive structure-activity relationships for $K_{\rm V}7.2/3$ opening, an analysis of the toxicity data displayed in Table 1 does not allow a clear correlation with certain structural elements to set up any valid structure-toxicity relationships. However, the nicotinamide scaffold per se does not appear to be of toxicological concern. This becomes apparent when the analogs with low water solubility are initially left aside, and instead, the focus is placed on the more soluble substances such as 14, 18d, 21, or 28a/ b. These analogs displayed rather uncritical levels of in vitro hepatotoxicity with higher LD₂₅ values compared to flupirtine. Nevertheless, focusing on the remaining analogs, there are some compounds that appear to perform worse than flupirtine and retigabine. This applies, for example, to the nanomolar active analogs 18b and 18e, for which LD₂₅ values of 15 and 32 µM, respectively, were determined for the HEP-G2 cell line. However, the structural similarity of the compounds to the significantly less toxic analogs 14 and 18d suggests that the specific chemical structure may not be responsible for the reduced LD₂₅ values. Instead, the reason could be the overall increased lipophilicity of 18b and 18e, which was among the highest of the current analogs as indicated by logD7.4 values of 3.8 and 4.7, respectively. This would be consistent with an analysis of 1036 FDA-approved drugs suggesting that compound lipophilicity is statistically significantly associated with DILI risk.^[60]

However, it must be emphasized that it is not possible to define a general LD limit at which a substance is to be classified as hepatotoxic. Tham et al. reported LD₅₀ values of 10 proven hepatotoxic drugs determined by the MTT assay on HEP-G2 cells, which extend over a wide concentration range from $25 \,\mu\text{M}$ to 20 mM.^[61] Therefore, LD values on their own are not very meaningful and must always be considered in relation to the biological activity, which among other factors, ultimately determines the necessary therapeutic concentration. For this reason, LD₂₅/EC₅₀ ratios were calculated for all active analogs, thus enabling a preliminary estimation of the therapeutic range. Considering these LD₂₅/EC₅₀ ratios, the highly potent analogs 18b and 18e appear to have significantly improved therapeutic safety windows compared to flupirtine despite lower absolute LD₂₅ values. The same applies to compound **18c** with an LD₂₅/ EC₅₀ ratio of 1,250 that even surpasses retigabine, for which no hepatotoxic effect is known. Overall, only the weakly to moderately active analogs 14, 28a, 40c, 47, and 54 have a poorer preliminary therapeutic range than flupirtine, whereby the exact LD₂₅/EC₅₀ ratios of 40c, 47, and 54 are actually unknown since the maximum soluble concentrations were used for the calculation due to the lack of LD₂₅ values; therefore the calculated LD₂₅/EC₅₀ ratios represent worst-case scenarios. In summary, the potential to cause intrinsic hepatotoxicity appears to be low for most analogs based on preliminary in vitro toxicity data. However, as mentioned above, the results of the MTT assay may not be predictive of idiosyncratic hepatotoxicity.

3 | CONCLUSION

The $K_{V}7.2/3$ opening activity of nicotinamide lead compound **6** could be successfully improved by introducing a morpholino substituent and a 2,2,2-trifluoroethoxy group. The resulting compound **18c** was 150 times more potent than flupirtine and 20 times more potent than retigabine while also demonstrating a superior toxicity/activity ratio. The water solubility of **18c** and related compounds was still low, but considering the nanomolar activity, it might be sufficient for further development, which in perspective also includes in vivo assays. Moreover, the $K_V 7.2/3$ activity data of the structurally distinct analogs provided profound structure-activity relationships, which could be correlated with docking and molecular dynamics simulations to hypothesize a plausible binding mechanism for the nicotinamide derivatives. As a result, the ortho disubstitution of the amide function was confirmed as an essential structural feature. In addition, the role of a cyclic substituent in position 6 of the pyridine ring was clarified, which must be aligned coplanar to the central pyridine ring but still has to allow for a flexible conformation to fit into the binding pocket adequately. Finally, a bulky alkoxy group is presumed to better shield the analogs in their intramembrane binding site from the adjacent lipid layer by improving ligand-lipid interactions. Taken together, the SAR findings obtained in this work provide a strong basis for future drug design.

4.1 | Chemistry

4.1.1 | General remarks

The starting materials, reagents, and solvents were commercially available and purchased from Sigma Aldrich, VWR, TCI, or ABCR. All chemicals were used as received unless specified otherwise. Anhydrous solvents were obtained from Acros Organics, except THF, which was dried by refluxing over sodium. Microwavesupported syntheses were conducted using an Anton Paar Monowave 300 reactor in closed vessel mode with an integrated IR sensor for temperature control. NMR spectra were recorded on a Bruker Avance III device at 400 MHz (¹H) and 100 MHz (¹³C), respectively, using CDCl₃, DMSO- d_6 , or MeOH- d_4 as solvents. The chemical shifts were referenced to the internal standard tetramethylsilane (TMS) and reported in parts per million (ppm). The coupling constants (J) are in Hz, and the following abbreviations were used to designate the multiplicities: br = broad, s = singlet, d = doublet, t = triplet, q = quartet, m = multiplet, and combinations thereof. MIR spectroscopy was performed with an ALPHA FT-IR instrument from Bruker Optics equipped with a diamond ATR accessory unit. A Bruker Elute UHPLC with Bruker compact QTOF-MS, a Bruker maXis LC-QTOF-MS, or a Shimadzu LCMS-IT-TOF system, each operated with ESI ionization, were used to measure the HRAM-MS data. HPLC analysis with UV detection at 220 nm using the 100% method determined the purity of all final compounds to be >95%. The melting points were measured with an automated Büchi Melting Point M-565 device. Analytical thin-layer chromatography was carried out on silica gel 60 F254 aluminum plates obtained from Merck, and visualization was accomplished with UV light. Column chromatography on silica gel was performed using silica gel 60 from Carl Roth with a particle size of 20-45 µm. Flash chromatographic separations were conducted using the Sepacore system from Büchi with 25 or 50 g Biotage SNAP KP-SIL columns, or alternatively, an Interchim puriFlash XS 520Plus system in combination with 80 g puriFlash 30SI-HP or 25 g puriFlash 15SI-HP columns. The InChI codes of the investigated analogs are provided as Supporting Information. The Supporting Information also contains ¹H-NMR and ¹³C-NMR spectra of all synthesized compounds as well as HPLC traces of the final compounds. All new compounds were fully characterized, including purity by HPLC and HRAM-MS data within 4 ppm accuracy.

4.1.2 | Synthesis and characterization

General procedure A: introduction of alkoxy substituents

A 60% suspension of NaH in mineral oil (2.5 equiv., 1.21 mmol/ml) was suspended in dry THF under an argon atmosphere, and the resulting suspension was cooled to 0°C. A solution of the chosen alcohol (1.2 equiv., 0.58 mmol/ml) in dry THF and a solution of the required chloropyridine (5.2–24.3 mmol, 0.49 mmol/ml) in dry THF

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were added successively. Afterward, the reaction mixture was heated to 70°C and stirred at 70°C until the TLC control indicated complete conversion (7–23 h). The reaction was quenched by the addition of water (100 ml). The resulting aq. mixture was adjusted to pH 12 by the addition of a 2 M aq. NaOH solution and extracted with ethyl acetate (100 ml). The organic phase was discarded, and the aq. phase was adjusted to pH 2–3 with conc. HCl. Subsequently, it was extracted with ethyl acetate again (2 × 50–235 ml). Finally, the combined organic phases were washed with brine, dried over Na₂SO₄, filtrated, and concentrated under reduced pressure.

General procedure B: Amide coupling with oxalyl chloride

Under an argon atmosphere, the required nicotinic acid derivative (2.5-12.1 mmol, 0.2 mmol/ml) was dissolved in dry DCM, and a catalytic amount of dry DMF was added. The mixture was cooled to 0°C, and a solution of oxalyl chloride (3.0 equiv., 2.0 mmol/ml) in dry DCM was added dropwise. After complete addition, the cooling was discontinued, and the reaction mixture was stirred at room temperature for 3 h. Subsequently, all volatiles were removed under reduced pressure, and the residue was redissolved in dry DCM (10-48 ml). The resulting solution was added dropwise to a solution of amine reactant (1.2 equiv., 0.3 mmol/ml) and triethylamine (2.0 equiv., 0.5 mmol/ml) in dry DCM at 0°C under stirring. Afterward, the cooling was discontinued, and the reaction mixture was stirred at room temperature. After 16 h, additional DCM (50-240 ml) was added. The resulting solution was extracted successively with an equal volume of a saturated aq. NaHCO₃ solution and a 2 M aq. HCl solution. Finally, the organic phase was washed with brine, dried over Na₂SO₄, filtrated, and concentrated under reduced pressure to obtain the crude product.

General procedure C: Microwave-assisted nucleophilic substitution The required chloropyridine (0.9–3.0 mmol, 0.33 mmol/ml) was dissolved in NMP. Subsequently, the corresponding amine (5.0 equiv.) was added, and the mixture was heated in a microwave reactor at 165°C in a closed vessel under stirring. After complete conversion (30–60 min), the reaction mixture was cooled to room temperature, dissolved in ethyl acetate (100 ml), and extracted with water (2 × 100 ml). Finally, the organic phase was washed with brine, dried over Na₂SO₄, filtrated, and concentrated under reduced pressure to obtain the crude product.

General procedure D: Amide coupling with DIC and HOBt

The required carboxylic acid (1.0-5.8 mmol, 0.15 mmol/ml) was dissolved in DMF. HOBt (2.0 equiv.), DIC (2.0 equiv.) and the corresponding amine (1.5 equiv.) were added successively, and the reaction mixture was stirred at room temperature. After 16 h, the solution was partitioned between ethyl acetate (50-290 ml) and an equal volume of water. The organic phase was extracted with the same volume of water again (2×), washed with brine, dried over Na₂SO₄, filtrated, and concentrated under reduced pressure to obtain the crude product.

General procedure E: Dechlorination by catalytic hydrogenation The required chloropyridine (0.3-3.0 mmol, 0.02 mmol/ml) was dissolved in methanol. Pd/C (10% Pd, 50% water wet, 0.1 equiv.) and triethylamine (1.5 equiv.) were added. Subsequently, the suspension was carefully set under a hydrogen atmosphere (balloon pressure) and stirred at room temperature. After complete conversion (2-5 h), the reaction mixture was filtered through a pad of celite, and the filtrate was concentrated under reduced pressure to obtain the crude product.

2-Amino-6-chloro-4-methylnicotinic acid (12)

2,6-Dichloro-4-methylnicotinic acid (950 mg, 4.61 mmol), K₂CO₃ (1275 mg, 9.22 mmol, 2.0 equiv.), Cul (176 mg, 0.92 mmol, 0.2 equiv.), NaN₃ (1199 mg, 18.44 mmol, 4.0 equiv.), and ethane-1,2diamine (62 µl, 0.92 mmol, 0.2 equiv.) were successively dissolved/suspended in ethanol (90 ml). The reaction mixture was set under an atmosphere of argon and stirred under reflux. After 23 h, additional amounts of K₂CO₃ (382 mg, 2.77 mmol, 0.6 equiv.), Cul (53 mg, 0.28 mmol, 0,06 equiv.), NaN₃ (360 mg, 5.5 mmol, 1.2 equiv.) and ethane-1,2-diamine (19 µl, 0.28 mmol, 0.06 equiv.) were added, and the reaction was continued for 5 h under the same conditions. Afterward, the reaction mixture was filtered through a pad of silica gel. Subsequently, the silica gel pad was rinsed with ethanol, and the combined eluates were concentrated under reduced pressure. The residue was dissolved in a 1 M ag. KOH solution and the product was precipitated by the addition of conc. HCl. Finally, the precipitate was filtered off to obtain **12** as a brown solid (467 mg, 2.50 mmol, 54%). R_f = 0.63 (toluene/ethanol/AcOH 5:4:1); mp = 178°C; ¹H-NMR (400 MHz, DMSO-d₄): δ (ppm) = 13.36 (s. 1H), 7.20 (s. 2H), 6.54 (s. 1H), 2.38 (s, 3H); ¹³C-NMR (100 MHz, DMSO-d₆): δ(ppm) = 168.8, 159.7, 154.0, 151.1, 113.86, 106.23, 22.21; IR (ATR): v = 3406, 3282 (m, v_{N-H}), 3300–2500 (b, v_{O-H}), 1681 (s, $v_{C=O}$), 1619 (s, δ_{N-H}).

2-Amino-6-chloro-N-(4-fluorobenzyl)-4-methylnicotinamide (13)

Compound 12 (424 mg, 2.27 mmol) was dissolved in THF (22 ml). CDI (737 mg, 4.56 mmol, 2.0 equiv.) was added in one portion, and the mixture was stirred at 50°C. After 1 h, the solution was cooled to room temperature, 4-fluorobenzylamine (1040 µl, 9.09 mmol) was added, and the mixture was continued to stir at room temperature. After 12 h, ethyl acetate (100 ml) was added. The resulting solution was extracted with a saturated aq. NaHCO₃ solution (100 ml), washed with brine, dried over Na₂SO₄, filtrated, and concentrated under reduced pressure. Finally, the crude residue was purified by silica gel column chromatography (ethyl acetate/n-hexane 2:3) to obtain 13 as a slightly yellow solid (236 mg, 0.80 mmol, 35%). $R_{\rm f}$ = 0.70 (ethyl acetate/*n*-hexane 7:3); mp = 169°C; ¹H-NMR (400 MHz, DMSO-d₆): δ(ppm) = 8.87 (t, J = 6.0 Hz, 1H), 7.43-7.33 (m, 2H), 7.22-7.11 (m, 2H), 6.49 (s, 1H), 6.14 (s, 2H), 4.41 (d, J = 5.9 Hz, 2H), 2.10 (s, 3H); ¹³C-NMR (100 MHz, DMSO-d₆): δ (ppm) = 166.3, 161.2 (d, J = 242.4 Hz), 156.6, 148.0, 147.9, 135.4 (d, J = 3.1 Hz), 129.6 (d, J = 8.2 Hz), 115.2, 115.0 (d, J = 21.3 Hz), 112.4, 41.9, 18.8; IR (ATR): ṽ = 3441, 3278 (m, v_{N-H}), 1613 (s, v_{C=O}).

2-Amino-N-(4-fluorobenzyl)-4-methyl-6morpholinonicotinamide (14)

The synthesis was conducted from **13** (350 mg, 1.19 mmol) and morpholine in accordance with general procedure C. The crude residue was purified by silica gel column chromatography (ethyl acetate/n-hexane 7:3) and successive recrystallization (methanol/water) to obtain **14** as a slightly yellow solid (300 mg, 0.87 mmol, 73%). $R_f = 0.34$ (ethyl acetate/n-hexane 7:3); mp = 177°C; ¹H-NMR (400 MHz, DMSO-d_6): δ (ppm) = 8.41 (t, *J* = 6.0 Hz, 1H), 7.41–7.31 (m, 2H), 7.20–7.10 (m, 2H), 5.86 (s, 1H), 5.63 (s, 2H), 4.39 (d, *J* = 6.0 Hz, 2H), 3.67–3.60 (m, 4H), 3.41–3.34 (m, 4H), 2.14 (s, 3H); ¹³C-NMR (100 MHz, DMSO-d_6): δ (ppm) = 168.1, 161.1 (d, *J* = 242.0 Hz), 157.9, 156.2, 146.5, 135.9 (d, *J* = 2.9 Hz), 129.4 (d, *J* = 8.1 Hz), 115.0 (d, *J* = 21.3 Hz), 105.6, 96.7, 66.0, 44.9, 41.8, 20.5; IR (ATR): \tilde{v} = 3458, 3346, 3254 (m, v_{N-H}), 3049, 2974 (w, v_{C-H}), 1583 (s, v_{C=O}), 1531 (m, δ_{N-H}); ESI-HRAM-MS (*m*/z): calcd. for [C₁₈H₂₁N₄O₂F+H]⁺ 345.1721, found 345.1708; cpd purity (220 nm): 99.6%.

6-Chloro-2-methoxy-4-methylnicotinic acid (15a)

The synthesis was conducted from 2,6-dichloro-4-methylnicotinic acid (5.00 g, 24.3 mmol) and methanol in accordance with general procedure A to obtain **15a** as a beige-colored solid (4.72 g, 23.4 mmol, 96%). $R_{\rm f}$ = 0.78 (*n*-butanol/AcOH/water 8:1:1); mp: 166°C; ¹H-NMR (400 MHz, DMSO-d₆): δ (ppm) = 13.39 (s, 1H), 7.09 (s, 1H), 3.87 (s, 3H), 2.17 (s, 3H); ¹³C-NMR (100 MHz, DMSO-d₆): δ (ppm) = 166.6, 159.4, 149.8, 147.0, 118.0, 117.4, 54.3, 18.4; IR (ATR): \tilde{v} = 3002, 2954 (w, v_{C-H}), 3300–2500 (b, v_{O-H}), 1686 (s, v_{C=O}).

6-Chloro-2-isopropoxy-4-methylnicotinic acid (15b)

The synthesis was carried out from 2,6-dichloro-4-methylnicotinic acid (2.50 g, 12.1 mmol) and 2-propanol following general procedure A to obtain **15b** as a beige-colored solid (2.75 g, 12.0 mmol, 99%), which was used for the following reaction without any further characterization and purification.

6-Chloro-4-methyl-2-(2,2,2-trifluoroethoxy)nicotinic acid (15c)

The synthesis was conducted from 2,6-dichloro-4-methylnicotinic acid (1.25 g, 6.1 mmol) and 2,2,2-trifluoroethanol in accordance with general procedure A to obtain **15c** as a beige-colored solid (1.62 g, 6.0 mmol, 99%), which was used for the following reaction without any further characterization and purification.

6-Chloro-N-(4-fluorobenzyl)-2-methoxy-4-methylnicotinamide (16a) The synthesis was carried out from **15a** (500 mg, 2.48 mmol) and 4fluorobenzylamine according to general procedure B. The crude residue was purified by silica gel column chromatography (ethyl acetate/*n*-hexane 3:2), which yielded **16a** as a colorless solid (402 mg, 1.30 mmol, 53%). R_f = 0.29 (ethyl acetate/*n*-hexane 1:1); mp: 127°C; ¹H-NMR (400 MHz, DMSO-d₆): δ (ppm) = 8.89 (t, *J* = 6.0 Hz, 1H), 7.43–7.33 (m, 2H), 7.24–7.08 (m, 2H), 7.05 (d, *J* = 0.6 Hz, 1H), 4.42 (d, *J* = 6.0 Hz, 2H), 3.86 (s, 3H), 2.18 (d, *J* = 0.6 Hz, 3H); ¹³C-NMR (100 MHz, DMSO-d₆): δ (ppm) = 164.5, 161.2 (d, *J* = 242.1 Hz), 159.6, 149.9, 146.4, 135.3 (d, *J* = 3.0 Hz), 129.1 (d, *J* = 8.1 Hz), 119.9, 117.9, 115.0 (d, J = 21.3 Hz), 54.1, 41.5, 18.0; IR (ATR): $\tilde{v} = 3310$ (m, v_{N-H}), 3067, 2954 (w, v_{C-H}), 1635 (s, $v_{C=O}$), 1604 (m, δ_{N-H}).

6-Chloro-N-(4-fluorobenzyl)-2-(propan-2-yloxy)-4methylnicotinamide (**16b**)

The synthesis was carried out from **15b** (2.78 g, 12.1 mmol) and 4-fluorobenzylamine according to general procedure B. The crude residue was purified by flash chromatography (mobile phase: ethyl acetate/*n*-hexane with 0%–30% ethyl acetate), which yielded **16b** as a colorless solid (2.10 g, 6.2 mmol, 51%). $R_f = 0.83$ (ethyl acetate/*n*-hexane 2:1); mp: 122°C; ¹H-NMR (400 MHz, DMSO-d₆): δ (ppm) = 8.82 (t, *J* = 6.1 Hz, 1H), 7.45–7.35 (m, 2H), 7.21–7.11 (m, 2H), 7.00 (s, 1H), 5.16 (sept, *J* = 6.2 Hz, 1H), 4.42 (d, *J* = 6.1 Hz, 2H), 2.19 (s, 3H), 1.26 (d, *J* = 6.1 Hz, 6H); ¹³C-NMR (100 MHz, DMSO-d₆): δ (ppm) = 164.6, 161.2 (d, *J* = 242.0 Hz), 158.9, 149.9, 146.3, 135.4 (d, *J* = 3.0 Hz), 129.0 (d, *J* = 8.1 Hz), 120.4, 117.4, 114.8 (d, *J* = 21.0 Hz), 69.4, 41.3, 21.7, 18.0; (ATR): \tilde{v} = 3267 (m, v_{N-H}), 3078, 2982 (w, v_{C-H}), 1632 (s, v_{C=O}).

6-Chloro-N-(4-fluorobenzyl)-4-methyl-2-(2,2,2-trifluoroethoxy) nicotinamide (**16c**)

The synthesis was conducted from **15c** (1.25 g, 4.64 mmol) and 4-fluorobenzylamine following general procedure B. The crude residue was purified by silica gel column chromatography (ethyl acetate/*n*-hexane 2:3), which yielded **16c** as a beige-colored solid (855 mg, 2.27 mmol, 49%). $R_f = 0.67$ (ethyl acetate/*n*-hexane 1:1); mp: 127°C; ¹H-NMR (400 MHz, DMSO-d₆): δ (ppm) = 8.98 (t, J = 6.0 Hz, 1H), 7.43–7.32 (m, 2H), 7.21 (d, J = 0.6 Hz, 1H), 7.18–7.07 (m, 2H), 4.96 (q, J = 9.0 Hz, 2H), 4.44 (d, J = 6.0 Hz, 2H), 2.24 (d, J = 0.6 Hz, 3H); ¹³C-NMR (100 MHz, DMSO-d₆): δ (ppm) = 163.7, 161.2 (d, J = 242.1 Hz), 157.1, 151.1, 146.1, 135.1 (d, J = 3.0 Hz), 129.0 (d, J = 8.1 Hz), 123.7 (q, J = 277.7 Hz), 120.0, 119.6, 114.9 (d, J = 21.3 Hz), 62.3 (q, J = 35 Hz), 41.5, 18.1; (ATR): $\tilde{v} = 3404$ (m, v_{N+H}), 3013, 2924 (w, v_{C+H}), 1613 (s, $v_{C=O}$).

N-Butyl-6-chloro-2-methoxy-4-methylnicotinamide (16d)

The synthesis was carried out from **15a** (2.19 g, 10.87 mmol) and *n*butylamine according to general procedure B. The crude residue was purified by silica gel column chromatography (ethyl acetate/*n*-hexane 3:7), which yielded **16d** as a colorless solid (1.71 g, 6.7 mmol, 61%). $R_{\rm f} = 0.81$ (ethyl acetate/*n*-hexane 2:1); mp: 57°C; ¹H-NMR (400 MHz, DMSO-d₆): δ (ppm) = 8.30 (t, *J* = 5.7 Hz, 1H), 7.04 (s, 1H), 3.82 (s, 3H), 3.19 (td, *J* = 6.9, 5.7 Hz, 2H), 2.19 (s, 3H), 1.51–1.40 (m, 2H), 1.43–1.27 (m, 2H), 0.89 (t, *J* = 7.3 Hz, 3H); ¹³C-NMR (100 MHz, DMSO-d₆): δ (ppm) = 164.1, 159.6, 149.7, 146.1, 120.4, 117.8, 54.0, 38.3, 31.0, 19.5, 18.0, 13.6; IR (ATR): $\tilde{v} = 3220$ (m, $v_{\rm N-H}$), 3068, 2961 (w, $v_{\rm C-H}$), 1645 (s, $v_{\rm C=O}$), 1624 (m, $\delta_{\rm N-H}$).

N-(4-Fluorobenzyl)-6-methoxy-4-methyl-(2,4'-bipyridine)-5carboxamide (**17**)

In a microwave vessel, compound **16a** (210 mg, 0.68 mmol) was dissolved in 1,4-dioxane (2 ml). Eight hundred and eighty microliters of a 2 M aq. solution of Na_2CO_3 (1.76 mmol, 2.6 equiv.), tetrakis (triphenylphosphine)palladium(0) (79 mg, 0.07 mmol, 0.1 equiv.) and

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pyridin-4-ylboronic acid (100 mg, 0.82 mmol, 1.2 equiv.) were added. Argon was passed through the reaction mixture for 30 min. Afterward, the mixture was heated in a microwave reactor at 140°C under stirring. After 30 min, the reaction mixture was cooled to room temperature, water (100 ml) was added, and the aq. suspension was extracted with ethyl acetate (100 ml). The organic phase was washed with brine, dried over Na₂SO₄, filtrated, and concentrated under reduced pressure. The crude residue was purified by flash chromatography (ethyl acetate/n-hexane, 70%-90% ethyl acetate) and subsequent recrystallization (methanol/water), which yielded 17 as a beige-colored solid (140 mg, 0.40 mmol, 59%). $R_{\rm f}$ = 0.29 (ethyl acetate/*n*-hexane 3:1); mp: 139°C; ¹H-NMR (400 MHz, DMSO-d₆): δ (ppm) = 8.93 (t, J = 6.1 Hz, 1H), 8.74–8.68 (m, 2H), 8.10-8.04 (m, 2H), 7.69 (d, J = 0.7 Hz, 1H), 7.47-7.38 (m, 2H), 7.26-7.15 (m, 2H), 4.46 (d, J = 6.0 Hz, 2H), 4.01 (s, 3H), 2.29 (s, 3H); 13 C-NMR (100 MHz, DMSO-d₆): δ (ppm) = 165.2, 161.2 (d, J = 242.1 Hz), 159.8, 150.3, 150.0, 147.7, 144.8, 135.5 (d, J = 3.1 Hz), 129.1 (d, J = 8.2 Hz), 121.5, 120.6, 116.0, 115.0 (d, J = 21.3 Hz), 53.4, 41.5, 18.4; (ATR): ṽ = 3330 (m, v_{N-H}), 3023, 2951 (w, v_{C-H}), 1633 (s, $v_{C=O}$), 1594 (m, δ_{N-H}); ESI-HRAM-MS (m/z): calcd. for [C₂₀H₁₉N₃O₂F+H]⁺ 352.1456, found 352.1463; cpd purity (220 nm): 99.5%.

N-(4-Fluorobenzyl)-2-methoxy-4-methyl-6-

morpholinonicotinamide (18a)

The synthesis was conducted from **16a** (300 mg, 0.97 mmol) and morpholine in accordance with general procedure C. The crude residue was purified by flash chromatography (mobile phase: ethyl acetate/*n*-hexane with 50%–100% ethyl acetate) and successive recrystallization (methanol/water), which yielded **18a** as a colorless solid (126 mg, 0.35 mmol, 36%). R_f = 0.60 (ethyl acetate/*n*-hexane 3:2); mp: 192°C; ¹H-NMR (400 MHz, DMSO-d₆): δ (ppm) = 8.52 (t, *J* = 6.2 Hz, 1H), 7.43–7.33 (m, 2H), 7.21–7.10 (m, 2H), 6.22 (s, 1H), 4.38 (d, *J* = 6.1 Hz, 2H), 3.80 (s, 3H), 3.73–3.65 (m, 4H), 3.47–3.40 (m, 4H), 2.13 (s, 3H); ¹³C-NMR (100 MHz, DMSO-d₆): δ (ppm) = 166.1, 161.1 (d, *J* = 241.7 Hz), 158.8, 157.0, 148.6, 135.9 (d, *J* = 2.9 Hz), 128.9 (d, *J* = 8.1 Hz), 114.9 (d, *J* = 21.0 Hz), 109.6, 99.5, 65.9, 52.7, 45.0, 41.5, 19.2; IR (ATR): \tilde{v} = 3330 (m, v_{N+H}), 3049, 2997 (w, v_{C-H}), 1627 (s, v_{C=O}); ESI-HRAM-MS (*m/z*): calcd. for [C₁₉H₂₃N₃O₃F+H]⁺ 360.1718, found 360.1717; cpd purity (220 nm): 100.0%.

N-(4-Fluorobenzyl)-2-(propyl-2-oxy)-4-methyl-6morpholinonicotinamide (**18b**)

The synthesis was conducted from **16b** (596 mg, 1.77 mmol) and morpholine in accordance with general procedure C. The crude residue was purified by silica gel column chromatography (ethyl acetate/*n*-hexane 1:1) and successive recrystallization (ethanol/water), which yielded **18b** as a colorless solid (356 mg, 0.92 mmol, 52%). $R_f = 0.54$ (ethyl acetate/*n*-hexane 2:1); mp: 134°C; ¹H-NMR (400 MHz, DMSO-d₆): δ (ppm) = 8.41 (t, J = 6.2 Hz, 1H), 7.45–7.36 (m, 2H), 7.19–7.09 (m, 2H), 6.20 (s, 1H), 5.16 (sept, J = 6.2 Hz, 1H), 4.39 (d, J = 6.1 Hz, 2H), 3.73–3.63 (m, 4H), 3.47–3.36 (m, 4H), 2.14 (s, 3H), 1.24 (d, J = 6.2 Hz, 6H); ¹³C-NMR (100 MHz, DMSO-d₆): δ (ppm) = 166.2, 161.0 (d, J = 241.7 Hz), 158.1,

157.0, 148.6, 135.9 (d, J = 3.0 Hz), 128.9 (d, J = 8.0 Hz), 114.7 (d, J = 21.2 Hz), 110.0, 99.4, 67.6, 65.8, 45.0, 41.4, 22.0, 19.2; (ATR): $\tilde{v} = 3277$ (m, v_{N-H}), 2967 (w, v_{C-H}), 1619 (s, $v_{C=O}$), 1591 (m, δ_{N-H}); ESI-HRAM-MS (*m*/*z*): calcd. for $[C_{21}H_{27}N_3O_3F+H]^+$ 388.2031, found 388.2030; cpd purity (220 nm): 99.8%.

N-(4-Fluorobenzyl)-4-methyl-6-morpholino-2-(2,2,2-trifluoroethoxy) nicotinamide (**18c**)

The synthesis was conducted from **16c** (500 mg, 1.33 mmol) and morpholine in accordance with general procedure C. The crude residue was purified by silica gel column chromatography (ethyl acetate/*n*-hexane 1:1) and successive recrystallization (methanol/water), which yielded **18c** as a colorless solid (212 mg, 0.50 mmol, 37%). $R_f = 0.56$ (ethyl acetate/*n*-hexane 3:2); mp: 164°C; ¹H-NMR (400 MHz, DMSO-d₆): δ (ppm) = 8.61 (t, J = 6.1 Hz), 7.42–7.32 (m, 2H), 7.17–7.05 (m, 2H), 6.35 (s, 1H), 4.89 (q, J = 9.2 Hz, 2H), 4.39 (d, J = 6.0 Hz, 2H), 3.72–3.65 (m, 4H), 3.49–3.42 (m, 4H), 2.16 (s, 3H); ¹³C-NMR (100 MHz, DMSO-d₆): δ (ppm) = 165.38, 161.1 (d, J = 241.8 Hz), 156.7, 156.3, 149.4, 135.6 (d, J = 2.9 Hz), 128.8 (d, J = 8.0 Hz), 124.1 (q, J = 278.1 Hz), 114.7 (d, J = 21.2 Hz), 109.4, 101.0, 65.8, 61.2 (q, J = 34.5 Hz), 44.9, 41.5, 19.1; (ATR): $\tilde{v} = 3291$ (m, v_{N-H}), 2978 (w, v_{C-H}), 1617 (s, $v_{C=O}$), 1602 (s, δ_{N-H}); ESI-HRAM-MS (*m*/z): calcd. for $[C_{20}H_{22}N_3O_3F_4+H]^+$ 428.1592, found 428.1591; cpd purity (220 nm): 98.2%.

N-Butyl-2-methoxy-4-methyl-6-morpholinonicotinamide (18d)

The synthesis was conducted from **16d** (770 mg, 3.0 mmol) and morpholine in accordance with general procedure C. The crude residue was purified by flash chromatography (mobile phase: ethyl acetate/*n*-hexane with 0%–100% ethyl acetate) and successive recrystallization (ethanol/water), which yielded **18d** as a colorless solid (250 mg, 0.85 mmol, 28%). $R_f = 0.43$ (ethyl acetate/*n*-hexane 2:1); mp: 135°C; ¹H-NMR (400 MHz, DMSO-d₆): δ (ppm) = 7.93 (t, *J* = 5.7 Hz, 1H), 6.20 (s, 1H), 3.76 (s, 3H), 3.71–3.65 (m, 4H), 3.46–3.39 (m, 4H), 3.15 (td, *J* = 6.9, 5.7 Hz, 2H), 2.13 (s, 3H), 1.50–1.38 (m, 2H), 1.42–1.26 (m, 2H), 0.89 (t, *J* = 7.3 Hz, 3H); ¹³C-NMR (100 MHz, DMSO-d₆): δ (ppm) = 165.7, 158.7, 156.9, 148.3, 110.3, 99.4, 65.9, 52.6, 45.1, 38.4, 31.1, 19.5, 19.1, 13.7; (ATR): \tilde{v} = 3303 (m, v_{N-H}), 3067, 2967 (w, v_{C-H}), 1625 (s, v_{C=O}), 1594 (m, δ_{N-H}); ESI-HRAM-MS (*m*/*z*): calcd. for [C₁₆H₂₆N₃O₃+H]⁺ 308.1969, found 308.1968; cpd purity (220 nm): 99.7%.

N-(4-Fluorobenzyl)-2-methoxy-4-methyl-6-(piperidin-1-yl) nicotinamide (**18e**)

The synthesis was conducted from **16a** (400 mg, 1.30 mmol) and piperidine in accordance with general procedure C. The crude residue was purified by flash chromatography (mobile phase: ethyl acetate/*n*-hexane with 30%–70% ethyl acetate) and successive recrystallization (methanol/water), which yielded **18e** as a colorless solid (161 mg, 0.45 mmol, 35%). R_f = 0.76 (ethyl acetate/*n*-hexane 1:1); mp: 147°C; ¹H-NMR (400 MHz, DMSO-d₆): δ (ppm) = 8.49 (t, *J* = 6.2 Hz, 1H), 7.48–7.30 (m, 2H), 7.26–7.05 (m, 2H), 6.19 (s, 1H), 4.38 (d, *J* = 6.1 Hz, 2H), 3.79 (s, 3H), 3.65–3.40 (m, 4H), 2.13 (s, 3H), 1.67–1.57 (m, 2H), 1.57–1.48 (m, 4H); ¹³C-NMR (100 MHz, DMSO-d₆): δ (ppm) = 166.2, 161.0 (d, *J* = 241.8 Hz), 158.9, 156.7, 148.6, 136.0 (d, *J* = 2.9 Hz), 128.9 (d,

$$\begin{split} J &= 8.2 \text{ Hz}), \ 114.8 \ (d, J = 21.1 \text{ Hz}), \ 108.1, \ 99.4, \ 52.6, \ 45.5, \ 41.5, \ 24.9, \ 24.3, \\ 19.3; \ (ATR): \ \bar{\nu} &= 3335 \ (m, \ v_{N-H}), \ 2948 \ (w, \ v_{C-H}), \ 1626 \ (s, \ v_{C=O}), \ 1593 \ (s, \ \delta_{N-H}), \\ H; \ ESI-HRAM-MS \ (m/z): \ calcd. \ for \ [C_{20}H_{25}N_3O_2F+H]^+ \ 358.1925, \ found \\ 358.1954; \ cpd \ purity \ (220 \ nm): \ 99.7\%. \end{split}$$

N-(4-Fluorobenzyl)-2-isopropoxy-4-methyl-6-(2-oxa-6-azaspiro[3.3] heptan-6-yl)nicotinamide (**18f**)

The synthesis was conducted from 16b (500 mg, 1.49 mmol) and 2-oxa-6-azaspiro[3.3]heptane oxalate (856 mg, 2.97 mmol, 2.0 equiv.) following general procedure C. Deviating from general procedure C, DBU (665 µl, 4.45 mmol, 3.0 equiv.) was added to the reaction mixture. The crude residue was purified by flash chromatography (mobile phase: ethyl acetate/n-hexane with 60%-100% ethyl acetate), which yielded 18f as a slightly yellow solid (314 mg, 0.79 mmol, 53%). R_f = 0.32 (ethyl acetate/nhexane 1:1); mp: 146°C; ¹H-NMR (400 MHz, DMSO-d₆): δ(ppm) = 8.39 (t, J = 6.2 Hz, 1H), 7.45–7.36 (m, 2H), 7.19–7.09 (m, 2H), 5.77 (s, 1H), 5.17 (sept, J = 6.2 Hz, 1H), 4.71 (s, 4H), 4.38 (d, J = 6.1 Hz, 2H), 4.06 (s, 4H), 2.11 (s, 3H), 1.25 (d, J = 6.2 Hz, 6H); ¹³C-NMR (100 MHz, DMSO-d₆): δ (ppm) = 166.3, 161.0 (d, J = 241.6 Hz), 158.7, 158.5, 148.0, 135.9 (d, J = 2.9 Hz), 128.9 (d, J = 8.0 Hz), 114.7 (d, J = 21.1 Hz), 109.7, 98.3, 79.9, 67.5, 59.6, 41.4, 38.3, 22.0, 18.9; (ATR): v = 3329 (m, v_{N-H}), 2932 (m, v_{C-H}), 1628 (s, v_{C=O}), 1591 (s, δ_{N-H}); ESI-HRAM-MS (m/z): calcd. for [C₂₂H₂₇N₃O₃F+H]⁺ 400.2031, found 400.2033; cpd purity (220 nm): 98.4%.

N-(4-Fluorobenzyl)-4-methyl-6-morpholino-2-oxo-1,2dihydropyridine-3-carboxamide (**19a**)

Compound **19a** was a side product in the synthesis of **18a**. It was separated from the main product by flash chromatography and further purified by recrystallization (methanol/water). The title compound was obtained as a colorless solid (132 mg, 0.38 mmol, 39%). $R_f = 0.42$ (ethyl acetate); mp: 225°C (decomp.); ¹H-NMR (400 MHz, DMSO-d_6): δ (ppm) = 11.12 (s, 1H), 9.73 (s, 1H), 7.39–7.29 (m, 2H), 7.18–7.08 (m, 2H), 5.76 (s, 1H), 4.40 (d, J = 5.9 Hz, 2H), 3.70–3.63 (m, 4H), 3.42–3.35 (m, 4H), 2.41 (s, 3H); ¹³C-NMR (100 MHz, DMSO-d_6): δ (ppm) = 166.3, 162.4, 161.0 (d, J = 241.8 Hz), 155.8, 152.8, 136.3 (d, J = 3.0 Hz), 129.0 (d, J = 8.0 Hz), 114.9 (d, J = 21.2 Hz), 106.3, 95.2, 65.5, 46.0, 41.2, 22.7; IR (ATR): $\tilde{v} = 3303$ (m, v_{N-H}), 3060, 2924 (w, v_{C-H}), 1656 (s, $v_{C=O}$); ESI-HRAM-MS (*m*/z): calcd. for [$C_{18}H_{20}N_3O_3F+H$]⁺ 346.1561, found 346.1564; cpd purity (220 nm): 100.0%.

6-(3,6-Dihydro-2H-pyran-4-yl)-N-(4-fluorobenzyl)-2-methoxy-4methylnicotinamide (**20**)

In a microwave vessel, **16a** (600 mg, 1.94 mmol) was dissolved in 1,4dioxane (4 ml). 1.76 ml of a 2 M aq. solution of Na_2CO_3 (3.5 mmol, 1.8 equiv.), tetrakis(triphenylphosphine)palladium(0) (112 mg, 0.10 mmol, 0.05 equiv.) and 2-(cyclohex-1-en-1-yl)-4,4,5,5-tetramethyl-1,3,2dioxaborolane (490 mg, 2.33 mmol, 1.2 equiv.) were added. Argon was passed through the reaction mixture for 30 min. Afterward, the mixture was heated in a microwave reactor at 140°C. After 15 min, the reaction mixture was cooled to room temperature, water (100 ml) was added, and the aq. suspension was extracted with ethyl acetate (100 ml). The organic phase was washed with brine, dried over Na₂SO₄, filtrated, and concentrated under reduced pressure. The crude residue was purified by flash chromatography (ethyl acetate/*n*-hexane, 10%–30% ethyl acetate) and successive recrystallization (methanol/water), which yielded **20** as a colorless solid (595 mg, 1.67 mmol, 86%). R_f = 0.50 (ethyl acetate/*n*-hexane 1:1); mp: 167°C; ¹H-NMR (400 MHz, CDCl₃): δ (ppm) = 7.48–7.30 (m, 2H), 7.14–6.96 (m, 2H), 6.89–6.78 (m, 2H), 6.58 (s, 1H), 4.64 (d, *J* = 5.8 Hz, 2H), 4.49–4.32 (m, 2H), 3.98 (s, 3H), 3.94 (t, *J* = 5.5 Hz, 2H), 2.65–2.54 (m, 2H), 2.45 (s, 3H); ¹³C-NMR (100 MHz, CDCl₃): δ (ppm) = 166.5, 162.4 (d, *J* = 245.4 Hz), 159.8, 154.3, 150.1, 134.3 (d, *J* = 3.3 Hz), 133.4, 129.5 (d, *J* = 8.1 Hz), 127.1, 117.2, 115.7 (d, *J* = 21.4 Hz), 114.7, 66.0, 64.5, 53.7, 43.2, 25.8, 20.4; (ATR): \bar{v} = 3325 (m, v_{N-H}), 1631 (s, v_{C=O}), 1590 (m, δ_{N-H}); ESI-HRAM-MS (*m*/*z*): calcd. for [C₂₀H₂₂N₂O₃F+H]⁺ 357.1609, found 357.1607; cpd purity (220 nm): 99.5%.

N-(4-Fluorobenzyl)-2-methoxy-4-methyl-6-(oxan-4-yl) nicotinamide (21)

Compound 20 (250 mg, 0.70 mmol) was dissolved in methanol (20 ml). Pd/C (10% Pd, 50% water wet, 150 mg) was added, the suspension was carefully set under a hydrogen atmosphere (balloon pressure), and the reaction mixture was stirred at room temperature. After 5 h, the resulting mixture was filtered through a pad of celite, and the filtrate was concentrated under reduced pressure. The crude residue was purified by silica gel column chromatography (ethyl acetate/n-hexane 2:3) and successive recrystallization (methanol/water) to obtain 21 as a colorless solid (186 mg, 0.52 mmol, 74%). R_f = 0.46 (ethyl acetate/nhexane 1:1); mp: 124°C; ¹H-NMR (400 MHz, DMSO-d₆): δ(ppm) = 8.81 (t, J = 6.1 Hz, 1H), 7.50-7.31 (m, 2H), 7.28-7.06 (m, 2H), 6.77 (s, 1H), 4.41 (d, J = 6.1 Hz, 2H), 4.02-3.90 (m, 2H), 3.86 (s, 3H), 3.50-3.36 (m, 2H), 2.91-2.73 (m, 1H), 2.16 (s, 3H), 1.83-1.70 (m, 4H); ¹³C-NMR (100 MHz, DMSO-d₆): δ(ppm) = 165.6, 161.3, 160.8 (d, J = 241.0 Hz), 159.2, 146.7, 135.6 (d, J = 3.0 Hz), 129.0 (d, J = 8.1 Hz), 118.8, 115.5, 114.9 (d, J = 21.2 Hz), 67.0, 53.0, 41.8, 41.4, 31.7, 18.2; (ATR): \tilde{v} = 3351 (m, v_{N-H}), 3051, 2949 (w, v_{C-H}), 1634 (s, $v_{C=O}$), 1596 (m, δ_{N-H}); ESI-HRAM-MS (m/z): calcd. for [C₂₀H₂₄N₂O₃F+H]⁺ 359.1765, found 359.1759; cpd purity (220 nm): 100.0%.

6-Chloro-2-methoxynicotinic acid (23)

The synthesis was conducted from 2,6-dichloronicotinic acid (1.00 g, 5.2 mmol) and methanol in accordance with general procedure A. The crude residue was purified by recrystallization (toluene/*n*-hexane) to obtain **23** as a slightly pink-colored solid (768 mg, 4.09 mmol, 79%). $R_{\rm f} = 0.68$ (ethyl acetate/toluene/acetic acid 5:5:1); mp: 176°C; ¹H-NMR (400 MHz, DMSO-d₆): δ (ppm) = 13.14 (s, 1H), 8.16 (d, *J* = 7.8 Hz, 1H), 7.18 (d, *J* = 7.9 Hz, 1H), 3.92 (s, 3H); ¹³C-NMR (100 MHz, DMSO-d₆): δ (ppm) = 165.1, 161.4, 150.2, 143.9, 116.5, 113.8, 54.4; (ATR): $\bar{v} = 2958$ (w, v_{C-H}), 3200–2500 (b, v_{O-H}), 1687 (s, v_{C=O}).

6-Chloro-N-(4-fluorobenzyl)-2-methoxynicotinamide (24)

The synthesis was conducted from **23** (300 mg, 1.60 mmol) and 4-fluorobenzylamine in accordance with general procedure D. The crude residue was purified by flash chromatography (mobile phase: ethyl acetate/n-hexane with 10–70% ethyl acetate) and successive

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recrystallization (methanol/water), which yielded **24** as a beige-colored solid (308 mg, 1.05 mmol, 65%). $R_{\rm f}$ = 0.52 (ethyl acetate/*n*-hexane 1:3); mp: 101°C; ¹H-NMR (400 MHz, DMSO-d₆): δ (ppm) = 8.80 (t, *J* = 6.0 Hz, 1H), 8.15 (d, *J* = 7.9 Hz, 1H), 7.42–7.32 (m, 2H), 7.22 (d, *J* = 7.9 Hz, 1H), 7.21–7.11 (m, 2H), 4.48 (d, *J* = 6.1 Hz, 2H), 3.98 (s, 3H); ¹³C-NMR (100 MHz, DMSO-d₆): δ (ppm) = 162.9, 161.1 (d, *J* = 241.6 Hz), 159.9, 148.8, 142.9, 135.5 (d, *J* = 3.0 Hz), 129.1 (d, *J* = 8.1 Hz), 117.0, 116.6, 115.0 (d, *J* = 21.4 Hz), 54.6, 42.0; (ATR): \bar{v} = 3416 (m, v_{N-H}), 3091, 2953 (w, v_{C-H}), 1652 (s, v_{C=O}), 1582 (m, δ_{N-H}).

N-(4-Fluorobenzyl)-2-methoxy-6-morpholinonicotinamide (25)

The synthesis was conducted from **24** (250 mg, 0.85 mmol) and morpholine in accordance with general procedure C. The crude residue was purified by flash chromatography (mobile phase: ethyl acetate/*n*-hexane with 70%–100% ethyl acetate) and successive recrystallization (methanol/water), which yielded **25** as a colorless solid (105 mg, 0.30 mmol, 36%). R_f = 0.46 (ethyl acetate/*n*-hexane 1:1); mp: 129°C; ¹H-NMR (400 MHz, DMSO-d₆): δ (ppm) = 8.41 (t, *J* = 6.1 Hz, 1H), 8.05 (d, *J* = 8.5 Hz, 1H), 7.42–7.26 (m, 2H), 7.20–7.08 (m, 2H), 6.46 (d, *J* = 8.7 Hz, 1H), 4.46 (d, *J* = 6.1 Hz, 2H), 3.94 (s, 3H), 3.76–3.63 (m, 4H), 3.62–3.49 (m, 4H); ¹³C-NMR (100 MHz, DMSO-d₆): δ (ppm) = 163.7, 161.0 (d, *J* = 242.0 Hz), 159.7, 158.5, 142.3, 136.3 (d, *J* = 3.0 Hz), 129.0 (d, *J* = 8.1 Hz), 114.9 (d, *J* = 21.2 Hz), 103.4, 98.8, 65.8, 53.2, 44.6, 41.8; (ATR): \tilde{v} = 3403 (m, v_{N-H}), 2987 (w, v_{C-H}), 1642 (s, v_{C=O}), 1596 (s, δ_{N-H}); ESI-HRAM-MS (*m*/*z*): calcd. for [C₁₈H₂₁N₃O₃F+H]⁺ 346.1561, found 346.1558; cpd purity (220 nm): 98.7%.

2,6-Dichloro-N-(4-fluorobenzyl)-4-methylnicotinamide (26a)

The synthesis was conducted from 2,6-dichloro-4-methylnicotinic acid (1.19 g, 5.8 mmol) and 4-fluorobenzylamine in accordance with general procedure D. The crude residue was purified by flash chromatography (mobile phase: ethyl acetate/*n*-hexane with 10%–50% ethyl acetate) and successive recrystallization (acetone/water), which yielded **26a** as a pale yellow solid (835 mg, 2.66 mmol, 46%). $R_{\rm f}$ = 0.77 (ethyl acetate/*n*-hexane 1:1); mp: 161°C; ¹H-NMR (400 MHz, DMSO-d₆): δ (ppm) = 9.17 (t, *J* = 5.9 Hz, 1H), 7.57 (d, *J* = 0.8 Hz, 1H), 7.46–7.36 (m, 2H), 7.25–7.14 (m, 2H), 4.46 (d, *J* = 5.9 Hz, 2H), 2.26 (d, *J* = 0.7 Hz, 3H); ¹³C-NMR (100 MHz, DMSO-d₆): δ (ppm) = 163.9, 161.3 (d, *J* = 242.5 Hz), 151.2, 148.2, 145.4, 134.8 (d, *J* = 3.0 Hz), 132.6, 129.6 (d, *J* = 8.2 Hz), 124.6, 115.1 (d, *J* = 21.3 Hz), 41.8; (ATR): \tilde{v} = 3249 (m, v_{N-H}), 3078 (w, v_{C-H}), 1638 (s, v_{C=O}).

2,6-Dichloro-N-(4-fluorobenzyl)nicotinamide (26b)

The synthesis was conducted from 2,6-dichloronicotinic acid (500 mg, 2.60 mmol) and 4-fluorobenzylamine in accordance with general procedure D. The crude residue was purified by silica gel column chromatography (ethyl acetate/*n*-hexane 1:1) and successive recrystallization (methanol/water), which yielded **26b** as a colorless solid (673 mg, 2.25 mmol, 86%). R_f = 0.63 (ethyl acetate/*n*-hexane 1:1); mp: 168°C; ¹H-NMR (400 MHz, DMSO-d₆): δ (ppm) = 9.16 (t, *J* = 5.9 Hz, 1H), 8.03 (d, *J* = 8.0 Hz, 1H), 7.67 (d, *J* = 8.0 Hz, 1H), 7.50-7.32 (m, 2H), 7.26-7.11 (m, 2H), 4.46 (d, *J* = 5.9 Hz, 2H); ¹³C-NMR (100 MHz, DMSO-d₆): δ (ppm) = 164.2, 161.3 (d, *J* = 242.5 Hz),

149.2,145.8, 141.2, 134.8 (d, J = 3.0 Hz), 132.1, 129.3 (d, J = 8.1 Hz), 123.6, 115.1 (d, J = 21.3 Hz), 41.9; (ATR): \tilde{v} = 3253 (m, v_{N-H}), 3052 (w, v_{C-H}), 1638 (s, $v_{C=O}$), 1574 (m, δ_{N-H}).

2-Chloro-N-(4-fluorobenzyl)-4-methyl-6morpholinonicotinamide (**27a**)

Compound **26a** (600 mg, 1.92 mmol) was dissolved in 2-propanol (50 ml). Morpholine (835 μ l, 9.58 mmol, 5.0 equiv.) was added, and the reaction mixture was stirred under reflux. After 3 d, it was cooled to room temperature, and all volatiles were removed under reduced pressure. The crude residue was purified by flash chromatography (mobile phase: ethyl acetate/*n*-hexane with 50%–100% ethyl acetate), which yielded **27a** as a pale yellow solid (160 mg, 0.44 mmol, 23%). R_f = 0.26 (ethyl acetate/*n*-hexane 1:1); mp: 186°C; ¹H-NMR (400 MHz, DMSO-d_6): δ (ppm) = 8.88 (t, *J* = 6.0 Hz, 1H), 7.45–7.33 (m, 2H), 7.22–7.12 (m, 2H), 6.70 (s, 1H), 4.40 (d, *J* = 6.0 Hz, 2H), 3.73–3.60 (m, 4H), 3.51–3.38 (m, 4H), 2.14 (s, 3H); ¹³C-NMR (100 MHz, DMSO-d_6): δ (ppm) = 165.5, 161.2 (d, *J* = 242.1 Hz), 157.9, 148.5, 144.4, 135.4 (d, *J* = 3.0 Hz), 129.4 (d, *J* = 8.1 Hz), 122.5, 115.0 (d, *J* = 21.3 Hz), 106.1, 65.8, 44.9, 41.7, 19.1.

2-Chloro-N-(4-fluorobenzyl)-6-morpholinonicotinamide (27b)

Compound **26b** (600 mg, 2.01 mmol) was dissolved in methanol (30 ml). Morpholine (437 µl, 5.02 mmol, 2.5 equiv.) was added, and the reaction mixture was stirred under reflux. After 24 h, it was cooled to room temperature, and all volatiles were removed under reduced pressure. The crude residue was purified by flash chromatography (mobile phase: ethyl acetate/*n*-hexane with 50%–100% ethyl acetate), which yielded **27b** as a pale yellow solid (180 mg, 0.51 mmol, 26%). $R_f = 0.33$ (ethyl acetate/*n*-hexane 1:1); mp: 153°C; ¹H-NMR (400 MHz, DMSO-d₆): δ (ppm) = 8.77 (t, *J* = 6.1 Hz, 1H), 7.67 (d, *J* = 8.6 Hz, 1H), 7.40–7.31 (m, 2H), 7.20–7.10 (m, 2H), 6.81 (d, *J* = 8.6 Hz, 1H), 4.39 (d, *J* = 6.0 Hz, 2H), 3.70–3.63 (m, 4H), 3.53–3.44 (m, 4H); ¹³C-NMR (100 MHz, DMSO-d₆): δ (ppm) = 165.4, 161.2 (d, *J* = 242.1 Hz), 158.4, 145.1, 139.9, 135.4 (d, *J* = 3.0 Hz), 129.2 (d, *J* = 8.3 Hz), 120.2, 115.0 (d, *J* = 21.3 Hz), 104.8, 65.7, 44.7, 41.9; (ATR): \tilde{v} = 3261 (m, v_{N-H}), 2970 (w, v_{C-H}), 1623 (s, v_{C=O}), 1556 (m, δ_{N-H}).

N-(4-Fluorobenzyl)-4-methyl-6-morpholinonicotinamide (28a)

The synthesis was conducted from **27a** (130 mg, 0.36 mmol) following general procedure E. The crude residue was purified by silica gel column chromatography (ethyl acetate) and successive recrystallization (methanol/water) to obtain **28a** as a colorless solid (75 mg, 0.23 mmol, 64%). $R_f = 0.34$ (ethyl acetate/*n*-hexane 3:1); mp: 192°C; ¹H-NMR (400 MHz, DMSO-d₆): δ (ppm) = 8.71 (t, *J* = 6.0 Hz, 1H), 8.23 (s, 1H), 7.41–7.31 (m, 2H), 7.21–7.11 (m, 2H), 6.70 (s, 1H), 4.40 (d, *J* = 6.0 Hz, 2H), 3.72–3.65 (m, 4H), 3.53–3.46 (m, 4H), 2.34 (s, 3H); ¹³C-NMR (100 MHz, DMSO-d₆): δ (ppm) = 167.1, 161.1 (d, *J* = 242.1 Hz), 159.4, 147.3, 146.9, 135.9 (d, *J* = 3.2 Hz), 129.0 (d, *J* = 8.1 Hz), 121.9, 115.0 (d, *J* = 21.3 Hz), 107.8, 65.9, 44.8, 41.7, 20.0; (ATR): \tilde{v} = 3285 (m, v_{N-H}), 2930 (w, v_{C-H}), 1621 (s, v_{C=O}), 1604 (m, δ_{N-H}); ESI-HRAM-MS (*m*/*z*): calcd. for $[C_{18}H_{21}N_3O_2F+H]^+$ 330.1612, found 330.1616; cpd purity (220 nm): 99.6%.

N-(4-Fluorobenzyl)-6-morpholinonicotinamide (28b)

The synthesis was conducted from **27b** (110 mg, 0.32 mmol) following general procedure E. The crude residue was purified by silica gel column chromatography (ethyl acetate) and successive recrystallization (methanol/water) to obtain **28b** as a colorless solid (64 mg, 0.20 mmol, 65%). $R_{\rm f} = 0.43$ (ethyl acetate/*n*-hexane 3:1); mp: 212°C; ¹H-NMR (400 MHz, DMSO-d₆): δ (ppm) = 8.83 (t, *J* = 6.0 Hz, 1H), 8.66 (d, *J* = 2.5 Hz, 1H), 8.01 (dd, *J* = 9.0, 2.5 Hz, 1H), 7.39–7.29 (m, 2H), 7.19–7.09 (m, 2H), 6.86 (d, *J* = 9.0 Hz, 1H), 4.43 (d, *J* = 5.9 Hz, 2H), 3.72–3.65 (m, 4H), 3.59–3.52 (m, 4H); ¹³C-NMR (100 MHz, DMSO-d₆): δ (ppm) = 164.9, 161.1 (d, *J* = 241.0 Hz), 160.0, 147.9, 136.5, 136.0 (d, *J* = 3.0 Hz), 129.2 (d, *J* = 8.1 Hz), 118.8, 114.9 (d, *J* = 21.2 Hz), 105.6, 65.9, 44.7, 41.7; (ATR): $\bar{v} = 3298$ (m, $v_{\rm N-H}$), 2966 (w, $v_{\rm C-H}$), 1627 (s, $v_{\rm C=O}$), 1597 (s, $\delta_{\rm N-H}$); ESI-HRAM-MS (*m*/z): calcd. for [C₁₇H₁₉N₃O₂F+H]⁺ 316.1456, found 316.1463; cpd purity (220 nm): 97.9%.

N-(4-Fluorobenzyl)-2-isopropoxy-4-methylnicotinamide (29)

The synthesis was conducted from **16b** (1.00 g, 3.0 mmol) following general procedure E. The crude residue was purified by silica gel column chromatography (ethyl acetate/*n*-hexane 2:3) and successive recrystallization (methanol/water) to obtain **29** as a colorless solid (591 mg, 1.96 mmol, 66%). $R_f = 0.57$ (ethyl acetate/*n*-hexane 1:1); mp: 107°C; ¹H-NMR (400 MHz, DMSO-d₆): δ (ppm) = 8.78 (t, J = 6.2 Hz, 1H), 8.02 (d, J = 5.2 Hz, 1H), 7.48–7.38 (m, 2H), 7.21–7.10 (m, 2H), 6.84 (d, J = 5.3 Hz, 1H), 5.24 (sept, J = 6.2 Hz, 1H), 4.43 (d, J = 6.1 Hz, 2H), 2.18 (s, 3H), 1.26 (d, J = 6.2 Hz, 6H); ¹³C-NMR (100 MHz, DMSO-d₆): δ (ppm) = 165.5, 161.1 (d, J = 241.8 Hz), 159.2, 146.2, 146.0, 135.6 (d, J = 3.0 Hz), 128.9 (d, J = 8.1 Hz), 121.5, 118.3, 114.8 (d, J = 21.2 Hz), 67.9, 41.3, 21.9, 18.1; (ATR): $\bar{v} = 3283$ (m, v_{N-H}), 3066, 2992 (w, v_{C-H}), 1629 (s, $v_{C=O}$), 1588 (m, δ_{N-H}); ESI-HRAM-MS (*m*/*z*): calcd. for $[C_{17}H_{20}N_2O_3F+H]^+$ 303.1503, found 303.1498; cpd purity (220 nm): 99.4%.

Ethyl 4-ethyl-6-methyl-2-oxo-1,2,3,4-tetrahydropyrimidine-5carboxylate (**33**)

Urea (16.5 g, 275 mmol, 1.1 equiv.) was suspended in ethanol (60 ml). Afterward, ethyl 3-oxobutanoate (31.6 ml, 250 mmol), propanal (19.7 ml, 275 mmol, 1.1 equiv.), and a catalytic amount of acetic acid were added. The mixture was stirred in a closed vessel at 90°C. After 20 h, the reaction mixture was poured into water. The resulting precipitate was filtered off and recrystallized from ethanol. Finally the recrystallized product was washed with a small amount of a mixture of ethanol and water (1:1) to obtain **33** as a colorless solid (17.0 g, 80 mmol 32%). $R_f = 0.41$ (ethyl acetate); mp: 183°C (lit mp: 179°C–181°C)^[62]; ¹H-NMR (400 MHz, DMSO-d₆): δ (ppm) = 8.91 (s, 1H), 7.28 (s, 1H), 4.15–3.99 (m, 3H), 2.17 (s, 3H), 1.42 (qd, *J* = 7.4, 5.3 Hz, 2H), 1.19 (t, *J* = 7.1 Hz, 3H), 0.79 (t, *J* = 7.4 Hz, 3H); ¹³C-NMR (100 MHz, DMSO-d₆): δ (ppm) = 165.5, 152.8, 148.4, 98.8, 59.0, 51.3, 29.6, 17.7, 14.2, 8.5; IR (ATR): \bar{v} = 3241, 3117 (m, v_{N-H}), 2960 (w, v_{C-H}), 1720, 1701 (s, v_{C=O}).

Ethyl 6-ethyl-4-methyl-2-oxo-1,2-dihydropyrimidine-5carboxylate (34)

Nitric acid (50%, 25 ml) was cooled to -10° C. Subsequently, compound **33** (8.50 g, 40.1 mmol) was added in portions over a period of 5 min

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while maintaining the temperature at -10°C. After complete addition, the mixture was stirred additional 10 min at -10°C. Afterward, K₂CO₃ was added to adjust the pH to 7. The resulting aq. solution was extracted with ethyl acetate (10 × 100 ml). The combined organic phases were washed with brine, dried over Na₂SO₄, filtrated, and concentrated under reduced pressure to obtain **34** as an orange solid (7.30 g, 34.7 mmol, 87%). R_f = 0,45 (DCM/methanol 9:1); mp: 99°C; ¹H-NMR (400 MHz, DMSO-d₆): δ (ppm) = 12.14 (s, 1H), 4.27 (q, *J* = 7.1 Hz, 2H), 2.64 (q, *J* = 7.5 Hz, 2H), 2.33 (s, 3H), 1.29 (t, *J* = 7.1 Hz, 3H), 1.14 (t, *J* = 7.5 Hz, 3H); ¹³C-NMR (100 MHz, DMSO-d₆): δ (ppm) = 165.4, 155.2, 108.5, 61.0, 27.9, 20.8, 13.9, 12.5 (two ¹³C signals are apparently missing, possibly due to overlapping since the number of ¹³C signals of the predecessor compound **33** and the successor compound **35** is correct); IR (ATR): \tilde{v} = 2978 (w, v_{C-H}), 1709, 1645 (s, v_{C=O}), 1556 (m, δ_{N-H}).

Ethyl 2-chloro-6-ethyl-4-methyl-1,2-dihydropyrimidine-5carboxylate (35)

Compound **34** (1.10 g, 5.2 mmol) was dissolved in phosphorus oxychloride (3.20 ml, 18.4 mmol, 10.6 equiv.) and stirred for 1 h at 110°C. Subsequently, all volatiles were removed under reduced pressure. The residue was suspended in ice water (100 ml), and the resulting suspension was extracted with ethyl acetate (3 × 100 ml). The combined organic phases were washed with brine, dried over Na₂SO₄, filtrated, and concentrated under reduced pressure. The crude residue was purified by silica gel column chromatography (ethyl acetate/*n*-hexane 1:3), which yielded **35** as an orange liquid (540 mg, 2.36 mmol, 45%). $R_{\rm f} = 0.74$ (ethyl acetate/*n*-hexane 1:3); ¹H-NMR (400 MHz, DMSO-d₆): δ (ppm) = 4.41 (q, J = 7.1 Hz, 2H), 2.75 (q, J = 7.5 Hz, 2H), 2.48 (s, 3H), 1.33 (t, J = 7.1 Hz, 3H), 1.20 (t, J = 7.5 Hz, 3H); ¹³C-NMR (100 MHz, DMSO-d₆): δ (ppm) = 171.7, 167.7, 165.8, 159.5, 124.8, 62.1, 28.3, 22.2, 13.8, 12.4; IR (ATR): $\tilde{v} = 2981$ (w, v_{C-H}), 1726 (s, v_{C=O}).

2-Chloro-4-ethyl-6-methylpyrimidine-5-carboxylic acid (36)

Compound **35** (2.74 g, 12.0 mmol) was dissolved in THF (20 ml). The resulting solution was added to a solution of KOH (1.87 g, 47.9 mmol, 4.0 equiv.) in water (80 ml). The resulting mixture was stirred at room temperature for 12 h. Afterward, it was extracted with ethyl acetate (100 ml). The organic phase was discarded, and the aq. phase was acidified to pH 2–3 with conc. aq. HCl. Subsequently, it was extracted with ethyl acetate again (5 × 100 ml). The combined organic phases were washed with brine, dried over Na_2SO_4 , filtrated, and concentrated under reduced pressure. The resulting dark brown solid (1.74 g, 8.7 mmol, 72%) was used for the following reaction without characterization or any further purification.

2-Chloro-4-ethyl-N-(4-fluorobenzyl)-6-methylpyrimidine-5carboxamide (**37**)

Compound **36** (650 mg, 3.24 mmol), HATU (1.48 g, 3.9 mmol, 1.2 equiv.), DIPEA (1130 μ l, 6.48 mmol, 2.0 equiv.) and 4-fluorobenzylamine (555 μ l, 4.86 mmol, 1.5 equiv.) were successively dissolved in DMF (10 ml). The mixture was stirred at room temperature. After 8 h, the reaction was quenched by the addition of water (100 ml). The resulting suspension was

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extracted with ethyl acetate (100 ml). The organic phase was washed with water (2 × 100 ml) and brine, dried over Na₂SO₄, filtrated, and concentrated under reduced pressure. Finally, the crude residue was purified by silica gel column chromatography (ethyl acetate/*n*-hexane 1:1), which yielded **37** as a colorless oil (420 mg, 1.37 mmol, 42%). $R_{\rm f}$ = 0.51 (ethyl acetate/*n*-hexane 1:1); mp: 172°C; ¹H-NMR (400 MHz, DMSO-d₆): δ (ppm) = 9.14 (t, *J* = 5.9 Hz, 1H), 7.44–7.35 (m, 2H), 7.25–7.14 (m, 2H), 4.46 (d, *J* = 5.9 Hz, 2H), 2.63 (q, *J* = 7.5 Hz, 2H), 2.35 (s, 3H), 1.15 (t, *J* = 7.5 Hz, 3H); ¹³C-NMR (100 MHz, DMSO-d₆): δ (ppm) = 170.7, 166.6, 165.0, 161.3 (d, *J* = 242.8 Hz), 158.5, 134.8 (d, *J* = 3.0 Hz), 129.7 (d, *J* = 8.2 Hz), 128.9, 115.2 (d, *J* = 21.3 Hz), 41.9, 27.7, 21.5, 12.5; IR (ATR): \tilde{v} = 3259 (m, v_{N-H}), 3080, 2977 (w, v_{C-H}), 1637 (s, v_{C=O}), 1608 (m, δ_{N-H}).

Ethyl 4-ethyl-6-methyl-2-morpholinopyrimidine-5-carboxylate (38)

Compound **35** (4.72 g, 20.7 mmol) was dissolved in ethanol (100 ml). Morpholine (5.35 ml, 62.0 mmol, 3.0 equiv.) was added, and the reaction mixture was stirred at 80°C. After 2 h, volatiles were removed under reduced pressure. The resulting crude residue was purified by silica gel column chromatography (ethyl acetate/*n*-hexane 2:8) to obtain **38** as a yellow oil (5.20 g, 18.6 mmol, 90%). $R_f = 0.79$ (ethyl acetate/*n*-hexane 1:3); ¹H-NMR (400 MHz, DMSO-d_6): δ (ppm) = 4.29 (q, J = 7.1 Hz, 2H), 3.83–3.73 (m, 4H), 3.70–3.61 (m, 4H), 2.65 (q, J = 7.5 Hz, 2H), 2.34 (s, 3H), 1.30 (t, J = 7.1 Hz, 3H), 1.15 (t, J = 7.5 Hz, 3H); ¹³C-NMR (100 MHz, DMSO-d_6): δ (ppm) = 170.1, 167.4, 165.8, 159.7, 114.0, 65.9, 60.7, 43.7, 28.8, 23.3, 13.9, 12.6; (ATR): $\tilde{v} = 2972$ (w, v_{C-H}), 1711 (s, $v_{C=O}$).

4-Ethyl-6-methyl-2-morpholinopyrimidine-5-carboxylic acid (39)

Compound 38 (5.00 g. 17.9 mmol) was dissolved in ethanol (10 ml). A solution of KOH (4.93 g, 89.5 mmol) in water (20 ml) was added, and the reaction mixture was stirred at 80°C. After 4 h, all volatiles were removed under reduced pressure. The residue was dissolved in water (10 ml), and the resulting solution was acidified to pH 2-3 with conc. aq. HCl. Afterward, it was extracted with ethyl acetate (3 × 100 ml). The combined organic phases were dried over Na2SO4, filtrated, and concentrated under reduced pressure. The crude residue was purified by silica gel column chromatography (ethyl acetate/acetic acid 9:1) and successive recrystallization (toluene/n-hexane) to obtain 39 as a beigecolored solid (3.79 g, 15.1 mmol, 84%). R_f = 0,68 (ethyl acetate/toluene/ acetic acid 5:5:1); mp: 130°C; ¹H-NMR (400 MHz, DMSO-d₆): δ (ppm) = 13.02 (s, 1H), 3.86-3.70 (m, 4H), 3.70-3.58 (m, 4H), 2.69 (q, J = 7.5 Hz, 2H), 2.36 (s, 3H), 1.15 (t, J = 7.5 Hz, 3H); ¹³C-NMR (100 MHz, DMSO-d₆): δ (ppm) = 169.8, 169.1, 165.5, 159.7, 115.2, 66.0, 43.7, 28.8, 23.5, 12.6; (ATR): \tilde{v} = 2967 (w, v_{C-H}), 3300–2500 (b, v_{O-H}), 1675 (s, v_{C=O}).

4-Ethyl-N-(4-fluorobenzyl)-6-methyl-2-morpholinopyrimidine-5carboxamide (**40a**)

The synthesis was conducted from **37** (380 mg, 1.24 mmol) and morpholine in accordance with general procedure C. The crude residue was purified by silica gel column chromatography (ethyl acetate/*n*-hexane 1:1) and successive recrystallization (methanol/water), which yielded **40a** as a colorless solid (324 mg, 0.90 mmol, 73%). $R_{\rm f} = 0.66$

(ethyl acetate/*n*-hexane 3:2); mp: 187°C; ¹H-NMR (400 MHz, DMSO-d₆): δ (ppm) = 8.82 (t, *J* = 6.0 Hz, 1H), 7.42–7.32 (m, 2H), 7.23–7.12 (m, 2H), 4.40 (d, *J* = 5.9 Hz, 2H), 3.72–3.69 (m, 4H), 3.65–3.62 (m, 4H), 2.46 (q, *J* = 7.5 Hz, 2H, partially overlapped by DMSO signal), 2.19 (s, 3H), 1.11 (t, *J* = 7.5 Hz, 3H); ¹³C-NMR (100 MHz, DMSO-d₆): δ (ppm) = 167.7, 167.4, 163.3, 161.2 (d, *J* = 243.5 Hz), 160.0, 135.5 (d, *J* = 3.2 Hz), 129.5 (d, *J* = 8.1 Hz), 119.7, 115.0 (d, *J* = 21.3 Hz), 66.0, 43.8, 41.9, 27.8, 22.0, 12.5; IR (ATR): \tilde{v} = 3241 (m, v_{N-H}), 2975 (w, v_{C-H}), 1628 (s, v_{C=O}); ESI-HRAM-MS (*m*/*z*): calcd. for [C₁₉H₂₄N₄O₂F+H]⁺ 359.1878, found 359.1878; cpd purity (220 nm): 99.7%.

4-Ethyl-6-methyl-2-morpholino-N-[3-(trifluoromethyl)benzyl] pyrimidine-5-carboxamide (**40b**)

The synthesis was conducted from 39 (250 mg, 1.00 mmol) and (3-(trifluoromethyl)phenyl)methanamine (214 µl, 1.49 mmol, 1.5 equiv.) following general procedure D. The crude residue was purified by silica gel column chromatography (ethyl acetate/n-hexane 2:3) and successive recrystallization (methanol/water), which yielded 40b as a colorless solid (320 mg, 0.78 mmol, 79%). R_f = 0.50 (ethyl acetate/n-hexane 1:1); mp: 162°C; ¹H-NMR (400 MHz, DMSO-d₆): δ(ppm) = 8.94 (t, J = 6.1 Hz, 1H), 7.72-7.68 (m, 1H), 7.68-7.56 (m, 3H), 4.52 (d, J = 6.0 Hz, 2H), 3.77-3.68 (m, 4H), 3.68-3.56 (m, 4H), 2.48 (m, 2H, overlapped by DMSO signal), 2.21 (s, 3H), 1.12 (t, J = 7.5 Hz, 3H); ¹³C-NMR (100 MHz, DMSO-d₆): δ (ppm) = 167.7, 167.6, 163.3, 160.0, 140.8, 131.6, 129.4, 129.1 (q, J = 31.3 Hz), 124.3 (g, J = 270.7 Hz), 123.8 (g, J = 3.9 Hz), 123.6 (g, J = 4.0 Hz), 119.5, 66.0, 43.8, 42.1, 27.8, 21.9, 12.4; (ATR): v = 3251 (m, v_{N-H}), 2978 (w, v_{C-H}), 1634 (s, $v_{C=O}$), 1571 (m, δ_{N-H}); ESI-HRAM-MS (m/ z): calcd. for $[C_{20}H_{24}N_4O_2F_3+H]^+$ 409.1846, found 409.1845; cpd purity (220 nm): 99.7%.

N-(3,3-Dimethylbutyl)-4-ethyl-6-methyl-2-morpholinopyrimidine-5carboxamide (**40***c*)

The synthesis was conducted from 39 (250 mg, 1.00 mmol) and 3,3dimethylbutan-1-amine hydrochloride (274 mg, 1.99 mmol, 2.0 equiv.) following general procedure D. Deviating from general procedure D, triethylamine (416 µl, 2.99 mmol, 3.0 equiv.) was added and to the reaction mixture. The crude residue was purified by silica gel column chromatography (ethyl acetate/n-hexane 2:3) and successive recrystallization (methanol/water), which yielded 40c as a colorless solid (269 mg, 0.80 mmol, 81%). R_f = 0.59 (ethyl acetate/*n*-hexane 1:1); mp: 159°C; ¹H-NMR (400 MHz, DMSO-d₆): δ(ppm) = 8.22 (t, J = 5.6 Hz, 1H), 3.74-3.67 (m, 4H), 3.67-3.60 (m, 4H), 3.22 (m, 2H), 2.50 (m, 2H, overlapped by DMSO signal), 2.21 (s, 3H), 1.45-1.37 (m, 2H), 1.14 (t, J = 7.5 Hz, 3H), 0.92 (s, 9H); ¹³C-NMR (100 MHz, DMSO-d₆): δ (ppm) = 167.5, 167.0, 163.2, 159.9, 120.2, 66.0, 43.8, 42.5, 35.5, 29.6, 29.2, 27.8, 22.0, 12.5; (ATR): v = 3226 (w, v_{N-H}), 3067, 2952 (m, v_{C-H}), 1620 (m, $v_{C=0}$); ESI-HRAM-MS (m/z): calcd. for $[C_{18}H_{30}N_4O_2+H]^+$ 335.2442, found 335.2444; cpd purity (220 nm): 100.0%.

2-Chloro-4-methyl-6-morpholinonicotinonitrile (42)

2,6-Dichloro-4-methylnicotinonitrile (1.50 g, 8.0 mmol) was dissolved in methanol (15 ml). The solution was cooled to 0°C, and a solution of morpholine (1.75 ml, 20.1 mmol, 2.5 equiv.) in methanol (5 ml) was

added dropwise under stirring. After complete addition, the cooling was removed, and the reaction mixture was stirred for 16 h at room temperature. Subsequently, all volatiles were removed under reduced pressure, and the residue was dissolved in ethyl acetate (100 ml). The resulting solution was extracted with water (2 × 100 ml), washed with brine, dried over Na₂SO₄, filtrated, and concentrated under reduced pressure. The crude residue was purified by flash chromatography (mobile phase: ethyl acetate/*n*-hexane with 30%–70% ethyl acetate), which yielded **42** as a colorless solid (1.33 g, 5.6 mmol, 70%). R_f = 0.45 (ethyl acetate/*n*-hexane 1:1); mp: 156°C (lit mp: 158°C–160°C)^[63]; ¹H-NMR (400 MHz, DMSO-d₆): δ (ppm) = 6.88 (d, *J* = 0.8 Hz, 1H), 3.78–3.64 (m, 4H), 3.64–3.49 (m, 4H), 2.36 (d, *J* = 0.8 Hz, 3H); ¹³C-NMR (100 MHz, DMSO-d₆): δ (ppm) = 158.3, 153.5, 151.1, 115.9, 105.5, 95.9, 65.6, 44.5, 20.3; (ATR): \tilde{v} = 2982 (w, v_{C-H}), 2211 (m, v_{C=N}).

4-Methyl-6-morpholinoisoxazolo[5,4-b]pyridin-3-amine (43)

Acetohydroxamic acid (556 mg, 7.40 mmol, 2.2 equiv.) and potassium *tert*-butoxide (755 mg, 6.73 mmol, 2.0 equiv.) were set under an atmosphere of argon and dissolved in dry DMF (10 ml). The resulting mixture was stirred at room temperature for 30 min. Subsequently, a solution of **42** (800 mg, 3.37 mmol) in dry DMF (10 ml) was added. The reaction mixture was stirred at 50°C for 5 h. Afterward, it was concentrated under reduced pressure. The resulting residue was purified by flash chromatography (mobile phase: ethyl acetate/*n*-hexane with 50–90% ethyl acetate), which yielded **43** as a colorless solid (343 mg, 1.45 mmol, 43%). R_f = 0.43 (ethyl acetate/*n*-hexane 3:1); mp: 223°C (decomp.); ¹H-NMR (400 MHz, DMSO-d₆): δ (ppm) = 6.56 (d, *J* = 1.1 Hz, 1H), 5.80 (s, 2H), 3.72–3.65 (m, 4H), 3.59–3.52 (m, 4H), 2.49 (d, *J* = 0.8 Hz, 3H); ¹³C-NMR (100 MHz, DMSO-d₆): δ (ppm) = 169.5, 159.7, 158.8, 145.5, 103.4, 97.3, 65.8, 45.0, 18.4; (ATR): \tilde{v} = 3473, 3293 (m, v_{N-H}), 2961 (w, v_{C-H}).

N-(4-Fluorobenzyl)-4-methyl-6-morpholinoisoxazolo[5,4-b]pyridin-3-amine (44)

Compound 43 (333 mg, 1.41 mmol) was suspended in dry DCM (30 ml). The suspension was set under an atmosphere of argon. 4-Fluorobenzaldehyde (181 µl, 1.69 mmol, 1.2 equiv.) was added, and the mixture was stirred at room temperature. After 1 h, triethylsilane (675 µl, 4.23 mmol, 3.0 equiv.) and TFA (324 µl, 4.23 mmol, 3.0 equiv.) were added, the temperature was raised to 60°C, and stirring was continued for 24 h. Afterward, 70 ml of DCM was added, and the resulting solution was extracted with a sat. aq. NaHCO₃ solution (100 ml). The organic phase was dried over Na₂SO₄, filtrated, and concentrated under reduced pressure. The resulting residue was purified by flash chromatography (mobile phase: ethyl acetate/n-hexane with 40%-70% ethyl acetate) and successive recrystallization (methanol/water), which yielded 44 as a colorless solid (386 mg, 1.12 mmol, 80%). Rf = 0.39 (ethyl acetate/nhexane 1:1); mp: 150°C; ¹H-NMR (400 MHz, DMSO-d₆): δ (ppm) = 7.49-7.39 (m, 2H), 7.19-7.09 (m, 2H), 6.65 (t, J = 6.1 Hz, 1H), 6.57 (d, J = 1.0 Hz, 1H), 4.36 (d, J = 6.0 Hz, 2H), 3.71-3.64 (m, 4H), 3.58–3.51 (m, 4H), 2.53 (d, J = 0.8 Hz, 3H); ¹³C-NMR (100 MHz, DMSOd₆): δ(ppm) = 169.5, 161.1 (d, J = 241.9 Hz), 159.6, 158.3, 145.3, 135.8 (d, J = 3.0 Hz), 129.4 (d, J = 8.2 Hz), 114.8 (d, J = 21.3 Hz), 103.4, 96.9, 65.8,

45.3, 45.0, 18.8; (ATR): \tilde{v} = 3470 (m, v_{N-H}), 2965 (w, v_{C-H}); ESI-HRAM-MS (*m/z*): calcd. for [C₁₈H₂₀N₄O₂F+H]⁺ 343.1565, found 343.1561; cpd purity (220 nm): 100.0%.

2-Methoxy-4-methyl-6-morpholinonicotinonitrile (45)

Compound 42 (768 mg, 3.23 mmol) was dissolved in dry methanol (30 ml). A 25% solution of sodium methoxide in methanol (2.96 ml, 12.92 mmol, 4.0 equiv.) was added, and the reaction mixture was stirred under reflux in an apparatus equipped with a CaCl₂ drying tube. After 24 h, it was cooled to room temperature and concentrated under reduced pressure. The residue was dissolved in ethyl acetate (100 ml), and the resulting solution was extracted with water (2 × 100 ml). The organic phase was washed with brine, dried over Na₂SO₄, filtrated, and concentrated under reduced pressure. Finally, the crude residue was purified by silica gel column chromatography (ethyl acetate/n-hexane 1:1), which yielded 45 as a colorless solid (678 mg, 2.91 mmol, 90%). $R_{\rm f}$ = 0.56 (ethyl acetate/*n*-hexane 1:1); mp: 150°C; ¹H-NMR (400 MHz, DMSO-d₆): δ (ppm) = 6.42 (d, J = 0.9 Hz, 1H), 3.88 (s, 3H), 3.71-3.64 (m, 4H), 3.64–3.57 (m, 4H), 2.29 (d, J = 0.8 Hz, 3H); ¹³C-NMR (100 MHz, DMSO-d₆): δ (ppm) = 164.1, 158.2, 153.8, 116.3, 99.5, 81.6, 65.7, 53.5, 44.4, 19.9; (ATR): \tilde{v} = 2972 (w, v_{C-H}), 2206 (m, v_{C=N}).

N'-Hydroxy-2-methoxy-4-methyl-6morpholinonicotinimidamide (46)

Compound **45** (644 mg, 2.76 mmol) was suspended in ethanol (50 ml). A 50% aq. solution of hydroxylamine (1.63 ml, 27.6 mmol, 10.0 equiv.) was added, and the reaction mixture was stirred in a closed vessel at 90°C for 4 d. After 24, 48, and 62 h, additional hydroxylamine solution (each time 1.63 ml, 27.6 mmol, 10.0 equiv.) was added. To terminate the reaction, it was cooled to room temperature, and all volatiles were removed under reduced pressure. The crude residue was purified by flash chromatography (mobile phase: DCM/methanol with 0%–10% methanol), which yielded **46** as a colorless solid (730 mg, 2.74 mmol, 99%). $R_f = 0.69$ (DCM/methanol 9:1); mp: 211°C; ¹H-NMR (400 MHz, DMSO-d₆): δ (ppm) = 9.04 (s, 1H), 6.20 (s, 1H), 5.45 (s, 2H), 3.75 (s, 3H), 3.72–3.65 (m, 4H), 3.45–3.38 (m, 4H), 2.14 (s, 3H); ¹³C-NMR (100 MHz, DMSO-d₆): δ (ppm) = 160.3, 157.1, 150.2, 148.3, 106.3, 99.2, 65.9, 52.5, 45.1, 19.0; (ATR): $\bar{v} = 3470$, 3323, 3229 (m, v_{N-H}/v_{O-H}), 2949 (w, v_{C-H}), 1713 (m, $v_{C=N}$).

4-[5-(5-Benzyl-1,2,4-oxadiazol-3-yl)-6-methoxy-4-methylpyridin-2yl]morpholine (**47**)

Compound **46** (250 mg, 0.94 mmol) was dissolved in 20 ml of DCM, and triethylamine (196 μ l, 1.41 mmol, 1.5 equiv.) was added in one portion. The reaction mixture was cooled to 0°C, and a solution of 2-phenylacetyl chloride (188 μ l, 1.41 mmol, 1.5 equiv.) in DCM (5 ml) was added dropwise. After complete addition, stirring at 0°C was continued for 1 h. Subsequently, all volatiles were removed under reduced pressure. The residue was dissolved in THF (10 ml) and treated with a 1 M solution of tetra-*n*-butylammonium fluoride in THF (939 μ l, 0.94 mmol, 1.0 equiv.). The reaction mixture was stirred at room temperature for 2 h. Afterward, it was dissolved in ethyl acetate (100 ml) and extracted with water (100 ml). The organic

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phase was washed with brine, dried over Na₂SO₄, filtrated, and concentrated under reduced pressure. The crude residue was purified by flash chromatography (mobile phase: ethyl acetate/*n*-hexane with 10%–50% ethyl acetate), which yielded **47** as a colorless solid (125 mg, 0.34 mmol, 36%). $R_f = 0.64$ (ethyl acetate/*n*-hexane 1:1); mp: 115°C; ¹H-NMR (400 MHz, DMSO-d₆): δ (ppm) = 7.40–7.35 (m, 4H), 7.31 (m, 1H), 6.36 (s, 1H), 4.40 (s, 2H), 3.75 (s, 3H), 3.77–3.66 (m, 4H), 3.55–3.48 (m, 4H), 2.09 (s, 3H); ¹³C-NMR (100 MHz, DMSO-d₆): δ (ppm) = 177.6, 165.1, 160.5, 158.0, 150.8, 134.3, 129.0, 128.7, 127.3, 99.7, 97.3, 65.8, 52.9, 44.7, 31.9, 19.7; (ATR): $\bar{v} = 2954$ (w, v_C-H); ESI-HRAM-MS (*m*/*z*): calcd. for $[C_{20}H_{23}N_4O_3+H]^+$ 367.1765, found 367.1763; cpd purity (220 nm): 99.4%.

(2-Methoxy-4-methyl-6-morpholinopyridin-3-yl)methanamine (48)

Compound 45 (600 mg, 2.57 mmol) was dissolved in a saturated solution of ammonia in methanol (50 ml). 1.00 g of a Raney nickel suspension in water (50%) was washed with methanol several times and added to the reaction mixture. The suspension was carefully set under a hydrogen atmosphere (balloon pressure) and stirred at 50°C. After 5 h, the catalyst was removed by filtration, and the filtrate was concentrated under reduced pressure. The resulting residue was dissolved in ethyl acetate and filtrated through a pad of silica gel, which was subsequently rinsed with an additional 250 ml of ethyl acetate. The obtained filtrate was discarded. Afterward, the silica gel pad was rinsed with 100 ml of a mixture of methanol and conc. aq. ammonia solution (9:1) to elute the desired product. The filtrate was concentrated under reduced pressure to obtain 48 as a colorless oil (434 mg, 1.83 mmol, 71%), used for the following reaction without any further purification and characterization.

4-Fluoro-N-[(2-methoxy-4-methyl-6-morpholinopyridin-3-yl)methyl] benzamide (49)

Compound 48 (434 mg, 1.83 mmol) was dissolved in DCM (20 ml). Triethylamine (507 µl, 3.66 mmol, 2.0 equiv.) was added in one portion. The reaction mixture was cooled to 0°C, and a solution of 4fluorobenzoyl chloride (325 µl, 2.75 mmol, 1.5 equiv.) in DCM (20 ml) was added dropwise over a period of 30 min. Cooling was removed, and the reaction mixture was stirred at room temperature. After 16 h, the reaction mixture was concentrated under reduced pressure. The crude residue was purified by flash chromatography (mobile phase: ethyl acetate/n-hexane with 30%-50% ethyl acetate) and successive recrystallization (methanol/water), which yielded 49 as a colorless solid (422 mg, 1.17 mmol, 64%). $R_f = 0.63$ (ethyl acetate/n-hexane 1:1); mp: 180°C; ¹H-NMR (400 MHz, DMSO-d₆): δ(ppm) = 8.31 (t, J = 4.7 Hz, 1H), 7.96-7.86 (m, 2H), 7.30-7.19 (m, 2H), 6.21 (s, 1H), 4.35 (d, J = 4.7 Hz, 2H), 3.81 (s, 3H), 3.73-3.66 (m, 4H), 3.44-3.37 (m, 4H), 2.25 (s, 3H); ¹³C-NMR (100 MHz, DMSO-d₆): δ(ppm) = 165.0, 163.7 (d, J = 249.0 Hz), 160.7, 156.6, 150.2, 130.9 (d, J = 3.0 Hz), 130.0 (d, J = 8.8 Hz), 114.9 (d, J = 21.6 Hz), 106.7, 100.1, 65.9, 52.7, $v_{C=O}$), 1599 (s, δ_{N-H}); ESI-HRAM-MS (m/z): calcd. for $[C_{19}H_{23}N_3O_4F]$ +H]⁺ 360.1718, found 360.1713; cpd purity (220 nm): 99.6%.

Methyl 6-chloro-2-methoxy-4-methylnicotinate (50)

Compound **15a** (2.25 g, 11.2 mmol) was dissolved in dry DMF (30 ml). K₂CO₃ (2.31 g, 16.7 mmol, 1.5 equiv.) was added, and the mixture was stirred for 30 min. Afterward, methyl iodide (1.04 ml, 16.7 mmol, 1.5 equiv.) was added dropwise, and the reaction mixture was stirred at room temperature for 8 h. Subsequently, the reaction was quenched by the addition of 150 ml of water. The resulting suspension was extracted with ethyl acetate (150 ml). The organic phase was successively washed with a sat. aq. solution of NaHCO₃ (2 × 100 ml) and brine, dried over Na₂SO₄, filtrated, and concentrated under reduced pressure to obtain **50** as a pale yellow solid (2.36 g, 10.9 mmol, 98%). R_f = 0.50 (ethyl acetate/*n*-hexane 1:9); mp: 70°C; ¹H-NMR (400 MHz, DMSO-d₆): δ (ppm) = 7.11 (d, J = 0.7 Hz, 1H), 3.85 (s, 3H), 3.82 (s, 3H), 2.24 (d, J = 0.6 Hz, 3H); ¹³C-NMR (100 MHz, DMSO-d₆): δ (ppm) = 165.7, 159.7, 150.8, 147.9, 118.1, 115.6, 54.4, 52.5, 18.4; (ATR): \tilde{v} = 3034, 2954 (w, v_{CH}), 1717 (s, v_{C=0}).

Methyl 2-methoxy-4-methyl-6-morpholinonicotinate (51)

Compound **50** (800 mg, 3.71 mmol) was dissolved in NMP (10 ml). Triethylamine (1030 µl, 7.42 mmol, 2.0 equiv.) and morpholine (388 µl, 4.45 mmol, 1.2 equiv.) were added successively, and the resulting solution was stirred at 90°C for 2 d. Afterward, the reaction mixture was dissolved in ethyl acetate (100 ml) and extracted with water (3 × 100 ml). The organic phase was washed with brine, dried over Na₂SO₄, filtrated, and concentrated under reduced pressure. The crude residue was purified by flash chromatography (mobile phase: ethyl acetate/*n*-hexane with 30%–40% ethyl acetate), which yielded **51** as a colorless solid (386 mg, 1.45 mmol, 39%). R_f = 0.64 (ethyl acetate/*n*-hexane 1:1); mp: 123°C; ¹H-NMR (400 MHz, DMSO-d₆): δ (ppm) = 6.26 (d, *J* = 0.8 Hz, 1H), 3.79 (s, 3H), 3.73 (s, 3H), 3.71–3.64 (m, 4H), 3.56–3.45 (m, 4H), 2.20 (d, *J* = 0.6 Hz, 3H); ¹³C-NMR (100 MHz, DMSO-d₆): δ (ppm) = 166.9, 160.1, 157.5, 150.0, 103.9, 99.6, 65.8, 52.9, 51.5, 44.6, 19.9; (ATR): \tilde{v} = 2918 (w, v_{C-H}), 1747 (s, v_{C=O}).

2-Methoxy-4-methyl-6-morpholinonicotinohydrazide (52)

Compound **51** (250 mg, 0.94 mmol) was suspended in an 80% aq. solution of hydrazine hydrate (20.0 ml, 303 mmol, 323 equiv.). The suspension was heated to reflux, and methanol was added until complete dissolution of methyl 2-methoxy-4-methyl-6-morpholinonicotinate. Subsequently, stirring under reflux was continued. After 24 h, all volatiles were removed under reduced pressure. The crude residue was purified by flash chromatography (mobile phase: DCM/methanol with 0%–10% methanol), which yielded **52** as a beige-colored solid (168 mg, 0.63 mmol, 67%). $R_f = 0.73$ (DCM/methanol 9:1); mp: 158°C; ¹H-NMR (400 MHz, DMSO-d₆): δ (ppm) = 9.08 (s, 1H), 6.20 (s, 1H), 4.55 (s, 2H), 3.75 (s, 3H), 3.72–3.65 (m, 4H), 3.48–3.40 (m, 4H), 2.13 (s, 3H); ¹³C-NMR (100 MHz, DMSO-d₆): δ (ppm) = 165.4, 159.0, 157.1, 148.9, 108.3, 99.4, 65.9, 52.6, 45.0, 19.1; (ATR): $\tilde{v} = 3308$ (m, v_{N+H}), 2977 (w, v_{C+H}), 1593 (s, $v_{C=O}$).

2-(4-Fluorophenyl)-N'-(2-methoxy-4-methyl-6-

morpholinonicotinoyl)acetohydrazonamide (53)

Compound **56** (169 mg, 0.83 mmol, 1.4 equiv.) was added in portions to a solution of sodium methoxide (45 mg, 0.83 mmol, 1.4 equiv.) in dry methanol (20 ml) at 0°C. After complete addition, the cooling was

removed, and the reaction mixture was stirred at room temperature. Subsequently, compound **52** (158 mg, 0.59 mmol) was added, and stirring at room temperature was continued for 24 h. Afterward, volatiles were removed under reduced pressure, and the residue was purified by silica gel column chromatography (DCM/methanol 9:1), which yielded **53** as a colorless solid (111 mg, 0.28 mmol, 47%). $R_f = 0.57$ (DCM/methanol 9:1); mp: 219°C; ¹H-NMR (400 MHz, DMSO-d_6): δ (ppm) = 9.45 (s, 1H), 7.47–7.34 (m, 2H), 7.28–7.09 (m, 2H), 6.23 (s, 1H), 6.08 (s, 2H), 3.76 (s, 3H), 3.73–3.59 (m, 4H), 3.54–3.41 (m, 4H), 3.37 (s, 2H), 2.15 (s, 3H); ¹³C-NMR (100 MHz, DMSO-d_6): δ (ppm) = 161.1, 161.0 (d, J = 240.0 Hz), 159.1, 157.0, 152.3, 149.0, 134.0, 130.6 (d, J = 8.0 Hz), 114.9 (d, J = 21.1 Hz), 99.5, 65.9, 52.7, 45.1, 39.1, 19.1; (ATR): $\bar{v} = 3389$, 3220 (m, v_{N-H}), 3047, 2986 (w, v_{C-H}), 1653 (m, $v_{C=O}$).

4-{5-[5-(4-Fluorobenzyl)-1H-1,2,4-triazol-3-yl]-6-methoxy-4methylpyridin-2-yl}morpholine (54)

Compound 52 (133 mg, 0.50 mmol), K₂CO₃ (35 mg, 0.25 mmol, 0.5 equiv.), and 2-(4-fluorophenyl)acetonitrile (180 µl, 1.50 mmol, 3.0 equiv.) were dissolved in n-butanol (2 ml) and stirred in a closed vessel at 150°C in a microwave reactor. After 4 h, the reaction was cooled to room temperature, and all volatiles were removed under reduced pressure. The residue was dissolved in ethyl acetate (100 ml) and extracted with water (100 ml). The organic phase was washed with brine, dried over Na₂SO₄, filtrated, and concentrated under reduced pressure. The crude residue was purified by silica gel column chromatography (ethyl acetate/n-hexane 1:1), which yielded 54 as a colorless solid (90 mg, 0.23 mmol, 47%). $R_f = 0.41$ (ethyl acetate/n-hexane 1:1); mp: 164°C; ¹H-NMR (400 MHz, DMSO-d₆): δ(ppm) = 11.35 (s, 1H), 7.43-7.32 (m, 2H), 7.06-6.95 (m, 2H), 6.16 (s. 1H), 4.13 (s. 2H), 4.01 (s. 3H), 3.87-3.80 (m. 4H), 3.62-3.55 (m. 4H), 2.70 (s, 3H); 13 C-NMR (100 MHz, DMSO-d₆): δ (ppm) = 162.3, 161.8 (d, J = 242.0 Hz), 160.6, 157.1, 152.7, 152.5, 134.4, 130.6 (d, J = 7.9 Hz), 115.3 (d, J = 21.2 Hz), 101.9, 98.7, 66.8, 53.9, 45.3, 34.2, 23.4; (ATR): v = 3220 (m, v_{N-H}), 2977 (w, v_{C-H}), ESI-HRAM-MS (m/z): calcd. for [C₂₀H₂₃N₅O₂F+H]⁺ 384.1830, found 384.1833; cpd purity (220 nm): 98.6%.

Methyl 2-(4-fluorophenyl)acetimidate hydrochloride (56)

2-(4-Fluorophenyl)acetonitrile (3.00 ml, 25.0 mmol) was dissolved in dry methanol (20.0 ml, 49 mmol, 2.0 equiv.). The reaction mixture was set under an atmosphere of argon and cooled to 0°C. HCl gas was passed through the solution for 3 h under stirring. Afterward, all volatiles were removed under reduced pressure. The residue was suspended in dry THF, filtered off, and rinsed with additional dry THF to obtain **56** as a colorless solid (2.94 g, 14.5 mmol, 58%). Mp: 157°C; ¹H-NMR (400 MHz, CDCl₃): δ (ppm) = 12.91 (s, 1H), 11.83 (s, 1H), 7.48–7.38 (m, 2H), 7.10–6.99 (m, 2H), 4.28 (s, 3H), 4.06 (s, 2H); ¹³C-NMR (100 MHz, CDCl₃): δ (ppm) = 178.2, 162.9 (d, *J* = 247.9 Hz), 131.6 (d, *J* = 8.2 Hz), 126.9 (d, *J* = 3.3 Hz), 116.4 (d, *J* = 21.8 Hz), 61.3, 38.4; (ATR): \tilde{v} = 3100–2600 (b, v_{N-H}), 1652 (m, v_{C=N}).

1-Amino-2-(4-fluorophenyl)ethan-1-iminium chloride (57)

Compound **56** (2.20 g, 10.8 mmol) was dissolved in dry methanol (25 ml). Ammonia gas was passed through the solution for 30 min. Subsequently, the reaction vessel was closed, and the mixture was stirred at room temperature. After 24 h, all volatiles were removed under reduced pressure. The resulting mucilaginous substance was treated with ethyl acetate to obtain **57** as a colorless solid, which was filtered off (1.25 g, 6.6 mmol, 61%). Mp: 134°C; ¹H-NMR (400 MHz, MeOH-d₄): δ (ppm) = 7.75–7.29 (m, 2H), 7.29–6.99 (m, 2H), 3.82 (s, 2H); ¹³C-NMR (100 MHz, MeOH-d₄): δ (ppm) = 171.6, 164.1 (d, *J* = 245.5 Hz), 132.2 (d, *J* = 8.2 Hz), 130.5 (d, *J* = 3.3 Hz), 117.1 (d, *J* = 22.0 Hz), 38.6; (ATR): \tilde{v} = 3300–2600 (b, v_{N-H}), 1669 (s, v_{C=N}).

4.1.3 | LogD_{7.4} estimation

The logD_{7,4} estimation was carried out as previously reported by employing a standard HPLC-based method.^[23] Briefly, the capacity factors of seven reference substances with known logD7,4 values were determined from their retention times (acetophenone, benzene, ethyl benzoate, benzophenone, phenyl benzoate, diphenyl ether, bibenzyl) with uracil used as a dead-time marker. The logarithm of the capacity factors was then plotted against the corresponding logD7.4 values to obtain a calibration function, which was used to calculate the logD7.4 values for each compound of interest. For HPLC analysis, a Phenomenex Luna 5 µm Phenyl-Hexyl 100 Å column (150 × 4.6 mm) was used with a mixture of methanol (75%) and 10 mM Tris/HCl buffer (25%) at pH 7.4 as mobile phase at a flow rate of 1.0 ml/min. The compounds of interest were dissolved in methanol (2 mg/ml), and the retention time was determined as the mean value from two measurements. The reference substances were injected as a mixture, which was prepared by dissolving 2 mg of a reference substance in 1 ml of methanol and subsequently combining 50 µl aliquots of each reference solution. The reference mixture was measured before and after the compounds of interest. Then, the mean retention time from both measurements was used to calculate the calibration function.

4.2 | Biology

4.2.1 | K_V7.2/3 channel opening activity

The FLIPR Potassium Assay Kit (Molecular Devices) was used to determine the K_V7.2/3 channel opening activity of the synthesized analogs according to the protocol of the assay kit. HEK-293 cells transfected with KCNQ2/3 were obtained from SB Drug Discovery. The cell culture and the data analysis were carried out as previously described elsewhere.^[24,64] Briefly, the cells were grown in minimum essential medium with non-essential amino acids (Thermo Fisher Scientific), supplemented with 10% heat-inactivated fetal bovine serum, 2 mM L-glutamine, 4 μ g/ml blasticidin S-HCl, 1% penicillin/streptomycin and 0.78 mg/ml G418 sulfate. The cells had been seeded at densities of 60,000 cells/well in 100 μ l of cell culture media using black-walled 96-well plates with a clear bottom (4titude Vision Plates from Azenta Life Sciences) suitable for fluorimetric measurements. After incubation for 24 h, 100 μ l of loading buffer containing

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the fluorescent dye and probenecid as an inhibitor for the anionexchange protein were added. Afterward, the plates were incubated for 1 h at room temperature under the exclusion of light. The test compounds were serially diluted in DMSO and added to the wells to obtain a final DMSO concentration of 1% (V/V). Equally treated wells containing loading buffer and 1% DMSO without test compound were deemed as the vehicle control. After an additional 30 min of incubation, fluorimetric measurements were performed at extinction/ emission wavelengths of 485 nm and 535 nm with an Infinite F200 Pro plate reader (Tecan). The baseline fluorescence signal was recorded for 20 s. Subsequently, a stimulus buffer (25 mM K⁺, 15 mM TI⁺) was added to each well, and measuring the fluorescence intensity was continued for 2.5 min. The resulting data were processed by normalizing the measured fluorescence intensity with the average baseline signal (F/F_0) . A correction was then performed by calculating the difference between the normalized vehicle control signal and the normalized baseline signal and subtracting the result from the F/F_0 value. To obtain a concentration-activity curve, the maximal corr. $\Delta F/$ F₀ values were plotted against the logarithmic compound concentration. Relative EC₅₀ values were calculated with GraphPad Prism 6 by determining the inflection point of the resulting sigmoidal curves. The corresponding E_{max} values as a measure for the intrinsic activity were determined by relating the maximum corr. $\Delta F/F_0$ value of a specific compound to the maximum corr. $\Delta F/F_0$ value of flupirtine, which was defined as 100%. All results are means of at least three independent experiments ± standard deviation (SD).

4.2.2 | Hepatic cell viability

The culturing of the TAMH and HEP-G2 cells and the MTT cell viability assay were carried out as previously described.^[24,64] Briefly, TAMH mouse liver cells (School of Pharmacy, University of Washington, Seattle, WA, USA) were grown in serum-free DMEM/ F12 medium supplemented with 5% PANEXIN NTA, 10 mM nicotinamide, and 10 µg/ml gentamicin. HEP-G2 human liver cancer cells (DSMZ, Braunschweig, Germany) were cultured in RPMI 1640 (PAN Biotech), supplemented with 10% heat-inactivated fetal bovine serum and 1% penicillin/streptomycin. Both cell lines were seeded into 96-well plates with 20,000 cells per well for TAMH and 15,000 cells per well for HEP-G2 and incubated at 37°C in a 5% CO2 atmosphere. After 24 h, the medium was replaced with fresh culture medium containing the test compounds at defined concentrations. For this purpose, the compounds were dissolved in DMSO and serially diluted in the corresponding culture medium to achieve a final concentration of 1% DMSO (V/V). Equally treated wells containing 1% DMSO without test compound were deemed as the vehicle control. Additional wells without cells were treated analogously to the control wells to determine the background optical density (OD). After 24 h of incubation with the test compounds, the medium was replaced with fresh medium containing 10% (V/V) of a 2.5 mg/ml solution of 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide (MTT). After an additional 4 h of incubation, the culture medium

in each well was removed, and DMSO (50 μ I) was added to dissolve the formazan crystals. Afterward, ODs were determined at 570 nm by using a SpectraMax 190 microplate reader. A correction of the OD values of the test compound (T) and control (C) wells by subtracting the bank value followed. Subsequently, the $T/C_{\rm corr.}$ -ratios were calculated and plotted against the logarithmic compound concentration to obtain a dose-response curve. After interpolation of the sigmoidal standard curve with GraphPad Prism 6, the LD₅₀ and LD₂₅ values were determined as the concentrations that reduced cell viability to 50% and 75%, respectively. If no reduction in cell viability to 75% was observed at the maximum concentration possible without precipitating the compounds, the LD₂₅ value was reported as higher than the highest concentration tested. All results are the means of at least three independent experiments ± SD.

4.3 | Molecular modeling

If not stated otherwise, all calculations were performed using the Schrodinger software suite release 2022-1 (Schrödinger, LLC, 2022).

4.3.1 | Structure preparation

The heterotetrameric protein structure of the KCNQ2/3 potassium channel was taken from previous studies^[23] and is based on the cryo-EM structure of KCNQ2 with bound retigabine (PDB 7CR2),^[36] while two opposing chains were used as a template for homology modeling within the Multiple Sequence Viewer in Maestro. Finally, the obtained model was prepared by the Protein Preparation Wizard^[65] to optimize the protonation states and the hydrogen bond network, followed by a restraint minimization with OPLS4 force field parameters.^[66]

4.3.2 | Molecular docking

All ligands were prepared using the Ligand Preparation application in Maestro and then docked into the heterotetrameric KCNQ2/3 model using an induced fit approach^[67] in Glide (version 94137)^[68] with standard sampling protocol of protein sidechains and an implicit membrane representation. The resulting 20 docking poses per compound were visually analyzed and selected. The compounds **14** and **18a** were re-docked into the same KCNQ2/3 retigabine binding site after molecular dynamics simulations using the Glide XP scoring function^[69] for direct comparison.

4.3.3 | Molecular dynamics simulations

The previously generated protein structure was prepared for molecular dynamics simulation using Desmond System Builder. The POPC bilayer membrane^[70] position was placed on the

transmembrane helices, and the system was solvated in an orthorhombic box with a 1 nm buffer around the protein with explicit TIP3P^[71] water molecules and 0.15 M NaCl salt concentration. OPLS4 force field parameters^[66] were applied in all simulations. Since the binding pockets are highly conserved, four different ligands (retigabine, **18a**, **18c**, **49**) were used in a single molecular dynamics run after molecular docking. The molecular system was minimized for 500 ps and equilibrated using the standard protocol (distributed with Desmond v6.9^[72]) for membrane systems at 300 K and NpγT ensemble.^[73] The production run was performed for 50 ns with a timestep of 2 fs. The temperature and pressure were maintained by a Langevin thermostat and a Langevin barostat, respectively. The snapshots were saved to the trajectory every 20 ps and analyzed using the Simulation Interaction Diagram in Maestro.

ACKNOWLEDGMENTS

Konrad W. Wurm and Frieda-Marie Bartz are funded by grants DFG LI 765/7-2 and DFG BE 1287/6-2 awarded to Andreas Link and Patrick J. Bednarski by the Deutsche Forschungsgemeinschaft (DFG–German Research Foundation). We thank Ms. Anne Schüttler and Ms. Maria Hühr for excellent technical assistance. Open Access funding enabled and organized by Projekt DEAL.

CONFLICT OF INTEREST

The authors declare no conflict of interest.

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How to cite this article: K. W. Wurm, F.-M. Bartz, L. Schulig, A. Bodtke, P. J. Bednarski, A. Link, Arch. Pharm.
2023;356:e2200473. https://doi.org/10.1002/ardp.202200473