

Putative Synthetic Cannabinoids MEPIRAPIM, 5F-BEPIRAPIM (NNL-2), and Their Analogues Are T-Type Calcium Channel (Ca_v3) Inhibitors

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Cite This: *ACS Chem. Neurosci.* 2022, 13, 1395–1409

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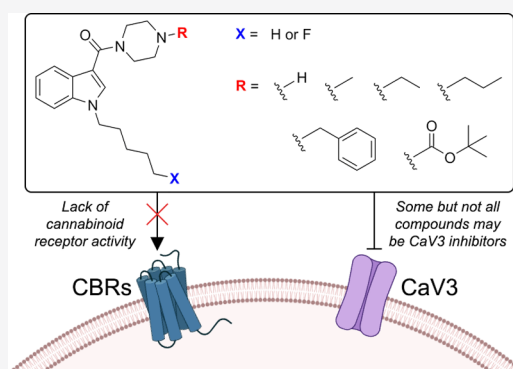
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ABSTRACT: Synthetic cannabinoid receptor agonists (SCRAs) are a large and growing class of new psychoactive substances (NPSs). Two recently identified compounds, MEPIRAPIM and 5F-BEPIRAPIM (NNL-2), have not been confirmed as agonists of either cannabinoid receptor subtype but share structural similarities with both SCRAs and a class of T-type calcium channel (Ca_v3) inhibitors under development as new treatments for epilepsy and pain. In this study, MEPIRAPIM and 5F-BEPIRAPIM and 10 systematic analogues were synthesized, analytically characterized, and pharmacologically evaluated using *in vitro* cannabinoid receptor and Ca_v3 assays. Several compounds showed micromolar affinities for CB₁ and/or CB₂, with several functioning as low potency agonists of CB₁ and CB₂ in a membrane potential assay. 5F-BEPIRAPIM and four other derivatives were identified as potential Ca_v3 inhibitors through a functional calcium flux assay (>70% inhibition), which was further confirmed using whole-cell patch-clamp electrophysiology. Additionally, MEPIRAPIM and 5F-BEPIRAPIM were evaluated *in vivo* using a cannabimimetic mouse model. Despite detections of MEPIRAPIM and 5F-BEPIRAPIM in the NPS market, only the highest MEPIRAPIM dose (30 mg/kg) elicited a mild hypothermic response in mice, with no hypothermia observed for 5F-BEPIRAPIM, suggesting minimal central CB₁ receptor activity.

KEYWORDS: cannabinoid, MEPIRAPIM, BEPIRAPIM, NNL, pharmacology, calcium channel



INTRODUCTION

Synthetic cannabinoid receptor agonists (SCRAs) are a diverse class of new psychoactive substances (NPSs) that exert intoxication via activation of cannabinoid type 1 receptors (CB₁). Most SCRAs function as agonists of both cannabinoid receptor subtypes, CB₁ and CB₂, with some compounds exhibiting potencies and efficacies equal to or greater than conventional “full agonists” like CP 55,940.^{1–3} The chemical evolution of SCRA NPS changes dynamically in response to local legislation,⁴ requiring the ongoing development of reference standards and detection methods for emerging compounds.^{5–11} Understanding structure–activity relationships for emergent SCRAs and their analogues facilitates more efficient allocation of resources to those SCRAs of greatest concern.

The earliest SCRAs were acylindoles such as JWH-018 (1; Figure 1) and XLR-11 (2), which possessed moderate potencies and efficacies at CB₁,¹² but as these compounds became prohibited in the United States and elsewhere, more potent and efficacious indole and indazole amides emerged in NPS markets

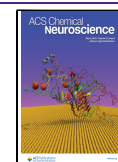
to replace them.¹³ More recent members of this class, such as amino acid-derived SCRAs AB-PINACA (3), ADB-CHMINACA (4), and AMB-FUBINACA (5), are frequently linked to mass intoxication events and serious adverse effects in the United States and other countries.^{14–20} Owing to their rapid proliferation, many recent SCRAs are unknown in the scientific literature at the time of initial detection in NPS markets; cannabimimetic activity remains to be confirmed *in vitro* and *in vivo*.

The present study focuses on the synthesis and characterization of a series of compounds (6–17; Figure 2) including and closely related to MEPIRAPIM (7) and 5F-BEPIRAPIM (NNL-

Received: December 12, 2021

Accepted: March 22, 2022

Published: April 20, 2022



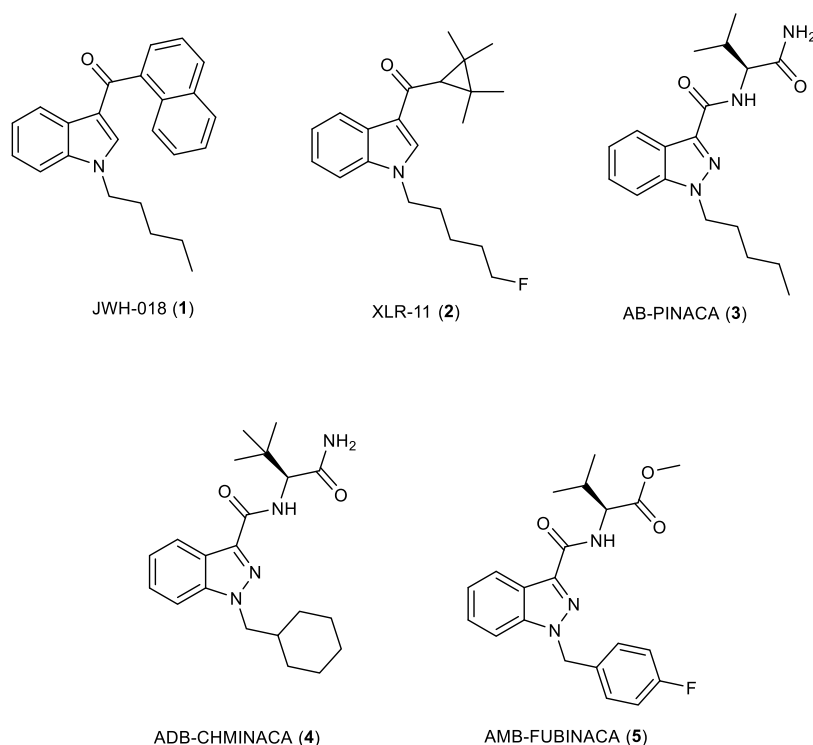


Figure 1. Synthetic cannabinoid receptor agonists (SCRAs) identified as new psychoactive substances.

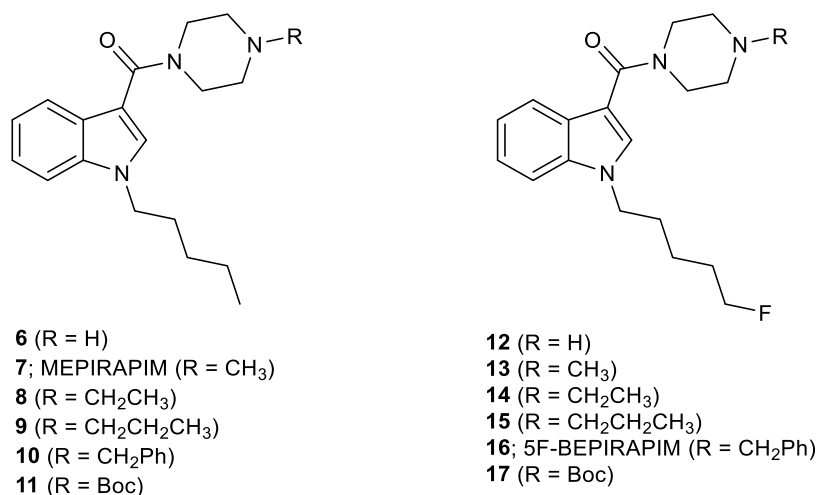


Figure 2. MEPIRAPIM, 5F-BEPIRAPIM (NNL-2), and their analogues.

2, 16). MEPIRAPIM (7) was first reported as an NPS in Japan in 2013²¹ and bears structural similarity to CB₁ and CB₂ agonists discovered by Organon International and later developed by Schering-Plough.^{22,23} MEPIRAPIM has since been reported in other regions, and a fluorinated benzylic analogue, 5F-BEPIRAPIM (NNL-2, 16), was recently identified in the seized material from a clandestine laboratory in China.^{24–26} Very little is known about the pharmacology of this class of SCRAs, however, and MEPIRAPIM and acetylfentanyl were detected in two cases of fatal overdose.²⁷ In these cases, MEPIRAPIM was detected at much higher urine concentrations compared to typical levels for other SCRAs, which may suggest unusual pharmacokinetics or metabolism.²⁸

MEPIRAPIM possesses low affinity for CB₁ ($K_i = 2650$ nM) and CB₂ ($K_i = 1850$ nM) receptors despite apparent design inspiration from optimized Organon International/Schering-

Plough CB₁/CB₂ agonists such as Org 28611 (also known as SCH-900,111, 18; Figure 3) and putative SCRA NPS like 5F-PY-PICA (19).^{10,22,29} Org 28611 possesses high affinities for both CB₁ ($K_i = 1.3$ nM) and CB₂ receptors ($K_i = 1.6$ nM) and functions as a potent, high efficacy CB₁ receptor agonist ($EC_{50} = 25$ nM). Org 28611 demonstrated adequate drug metabolism and pharmacokinetic (DMPK) profiles *in vitro* and *in vivo*, good aqueous solubility (129 mg·L⁻¹ at pH 6.9) and brain penetration, and was evaluated clinically as a potential intravenous analgesic agent.^{22,30} By contrast, 5F-PY-PICA shows negligible affinity for CB₁ ($K_i > 10,000$ nM) and micromolar affinity for CB₂ receptors ($K_i = 1737$ nM) and functions as a selective CB₂ partial agonist ($EC_{50} = 338$ nM; $E_{max} = 50\%$) with negligible effect on CB₁ ($E_{max} = 12\%$ at 10 μ M).²⁹

MEPIRAPIM also bears similarity to carbazoles designed to activate cannabinoid receptors and inhibit T-type calcium

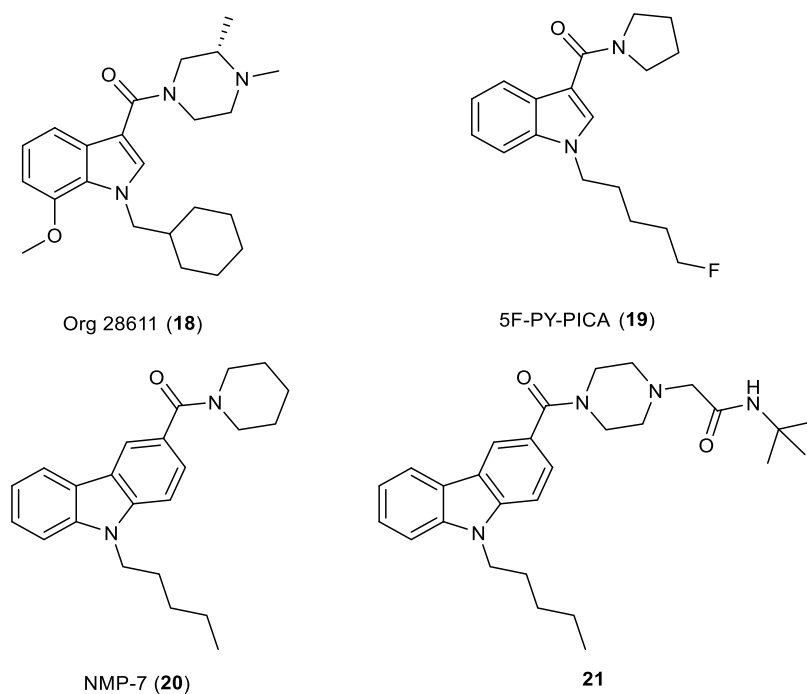


Figure 3. Analogues of MEPIRAPIM and 5F-BEPIRAPIM with diverse pharmacological profiles.

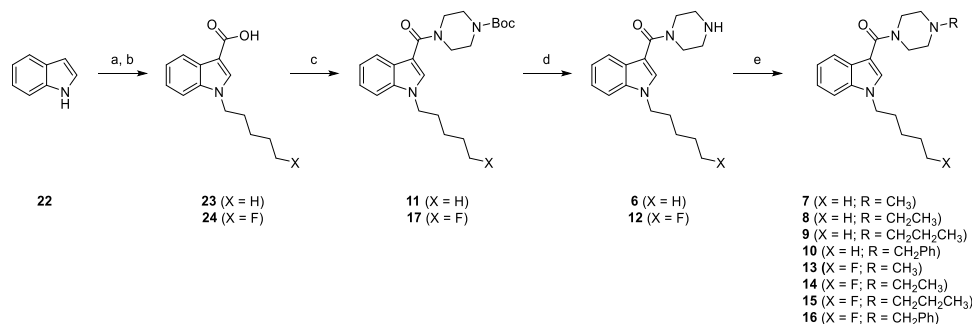


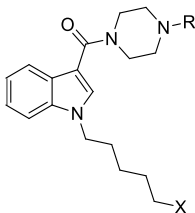
Figure 4. Reagents and conditions: (a) (i) NaH, X(CH₂)₅Br, *N,N*-dimethylformamide (DMF), 0 °C, rt, 1 h; (ii) (CF₃CO)₂O, 0 °C, rt, 1 h; (b) KOH, MeOH, PhMe, reflux, 2 h (**23**: 55%; **24**: 67%, over three steps); (c) Boc-piperazine, EDC·HCl, HOBT·H₂O, Et₃N, DMF, rt, 14 h (**11**: 92%; **17**: 89%); (d) 4 M HCl in dioxane, 0 °C, rt, 16 h (**6**-HCl: 91%; **12**-HCl: 84%); and (e) RCHO, NaBH(OAc)₃, ClCH₂CH₂Cl, rt, 2 h, 85–95%.

channels as dual mechanism analgesics, such as NMP-7 (**20**) and **21**.^{31,32} T-type calcium channels (Ca_v3) are a subfamily of low-voltage-gated Ca²⁺ channels, with three subtypes defined by their $\alpha 1$ subunit, namely, Ca_v3.1 ($\alpha 1G$), Ca_v3.2 ($\alpha 1H$), and Ca_v3.3 ($\alpha 1I$).^{33–36}

T-type calcium channels possess a number of unique biophysical properties, including small unitary conductance and rapid inactivation kinetics.³⁷ They also exhibit a hyperpolarized voltage dependence of activation and inactivation that give rise to a significant window current near typical neuronal resting membrane potentials.³⁸ These latter features allow T-type channels to efficiently regulate neuronal excitability by supporting both action potential initiation and rebound burst activity.³⁹ They also allow T-type channels to partake in the low threshold release of hormones and neurotransmitters.⁴⁰ Aberrant activation of T-type calcium channels has been linked to the development of conditions such as epilepsy and chronic pain, and genetic mutations in the three genes encoding the T-type channel family are associated with conditions such as epilepsy, neuromuscular diseases, primary aldosteronism, and autism.⁴¹ It is thus not surprising that T-type calcium channels

have emerged as pharmacological targets in the treatment of seizure disorders and are being explored as potential targets in the treatment of tremors (for review, see ref 42). Indeed, Ca_v3 inhibition is thought to be the key mechanism of action for the approved antiepileptic drugs zonisamide and ethosuximide, and Ca_v3 inhibitors Z944 and ACT-709,478 are currently in phase II clinical trials.^{43,44} Ca_v3.2 T-type calcium channels are also considered as potential targets for chronic pain,⁴⁵ although the inhibitor ABT-639 has failed in a phase II trial.⁴⁶ Finally, there is emerging evidence that Ca_v3.2 channels may be a potential target for treating pruritus.^{47,48} Hence, T-type channel inhibitors have considerable therapeutic potential, and it is important to identify novel pharmacophores that act on these channels for further optimization and development.⁴³

Activation of CB₁ and CB₂ receptors produces analgesia in many animal models of nociception, through actions at sites in the central and peripheral nervous systems, as well as the immune system.⁴⁹ It has also been reported that modest CB₁ agonism can contribute to the analgesic effects of moderately selective CB₂ agonists in the complete Freund's adjuvant (CFA) model of inflammatory nociception⁵⁰ and that low occupancy of

Table 1. Binding Affinities and Functional Activities of SCRA 6–17 at hCB₁ and hCB₂ Receptors^{a,b}

compound	X	R	calc cLogP	hCB ₁		hCB ₂	
				pK _i [M] ± SEM (K _i , μM) ^a	max ± SEM (% CP 55 940) ^b	pK _i [M] ± SEM (K _i , μM) ^a	max ± SEM (% CP 55 940) ^b
6	H	H	2.23	<5 (>10)	17.2 ± 13.7	<5 (>10)	9.8 ± 5.2
7	H	Me	2.60	<5 (>10)	40.0 ± 5.9	5.93 ± 0.02 (1.18)	16.9 ± 6.3
8	H	Et	2.94	<5 (>10)	37.0 ± 9.0	5.98 ± 0.03 (1.05)	13.1 ± 5.3
9	H	Pr	3.43	5.27 ± 0.08 (5.37)	9.8 ± 8.5	5.68 ± 0.08 (2.09)	15.9 ± 6.7
10	H	Bn	4.34	5.47 ± 0.07 (3.39)	17.7 ± 14.4	5.20 ± 0.13 (6.31)	15.5 ± 8.7
11	H	Boc	3.33	<5 (>10)	2.9 ± 1.0	<5 (>10)	3.8 ± 3.9
12	F	H	1.73	<5 (>10)	5.0 ± 8.2	<5 (>10)	16.7 ± 11.1
13	F	Me	2.11	<5 (>10)	44.3 ± 5.9	<5 (>10)	16.4 ± 4.6
14	F	Et	2.45	<5 (>10)	35.1 ± 7.6	5.53 ± 0.08 (2.95)	15.7 ± 3.8
15	F	Pr	2.94	<5 (>10)	12.3 ± 12.1	<5 (>10)	23.8 ± 6.6
16	F	Bn	3.84	5.65 ± 0.07 (2.24)	0.5 ± 2.6	5.34 ± 0.13 (4.57)	8.2 ± 2.7
17	F	Boc	2.84	<5 (>10)	0.2 ± 0.5	<5 (>10)	5.4 ± 4.3

^apK_i < 5 indicates less than 60% displacement of [³H]ligand at 10 μM in a binding assay. ^bMaximum response in the functional membrane potential assay at 10 μM compared to that at 1 μM CP 55,940 (100%). cLogP computed using ChemDraw (version 20).

CB₁ by an agonist with low intrinsic efficacy results in antinociception.⁵¹ Combined, these data suggest that compounds with low efficacy CB receptor agonism and potent T-type channel inhibition could be effective analgesics.

The activities of MEPIRAPIM, SF-BEPIRAPIM, and related compounds at CB receptors have not yet been defined *in vitro* or *in vivo* and, as indicated above, there seemed to be a possibility that they could also have activity at Ca_v3 channels. To address whether MEPIRAPIM, SF-BEPIRAPIM, and related compounds are SCRA and/or T-type calcium channel modulators, we devised a library of analogues based on the systematic modification of the pendant alkyl group to incorporate a homologous series from methyl to propyl substituents, as well as unsubstituted, *tert*-butyl carbamate-protected, and benzyl-substituted examples (see 6–17; Figure 2). Here, we report the synthesis, analytical chemistry, and *in vitro* cannabinoid and T-type calcium channel pharmacology of 6–17 and identify important structure–activity relationships (SARs) that could be used for further development and optimization of these pharmacophores. Additionally, MEPIRAPIM and SF-BEPIRAPIM, which have been detected in NPS products, were assessed for *in vivo* cannabinoid activity using radiotelemetry in mice.

RESULTS AND DISCUSSION

The synthesis of compounds 6–17 is shown in Figure 4. Indole (22) was alkylated with either bromopentane or 5-fluorobromopentane and treated with trifluoroacetic anhydride to give the corresponding trifluoroacetyl indoles, hydrolysis of which afforded carboxylic acids 23 or 24, respectively. Coupling of 23 or 24 with Boc-piperazine using 1-ethyl-3-(3'-dimethylaminopropyl)carbodiimide (EDC) gave 11 and 17, which were deprotected using hydrogen chloride in dioxane to give piperazines 6 and 12 as their hydrochloride salts. Reductive alkylation of 6 or 12 with either formaldehyde, acetaldehyde, propanal, or benzaldehyde in the presence of sodium

triacetoxyborohydride afforded 7–10 and 13–16 in excellent yield. The free base amines were purified by flash chromatography and characterized by NMR spectroscopy. The free base amines were then converted to their hydrochloride salts and purified by recrystallization and subjected to *in vitro* binding and functional activity assays at CB₁ and CB₂ receptors and *in vitro* functional activity assays at Ca_v3.1, Ca_v3.2, and Ca_v3.3 channels.

Ligands 6–17 were subjected to competition radioligand binding assays and fluorescence-based membrane potential assays at human CB₁ and CB₂ receptors (Table 1). Most compounds showed low or negligible affinities for CB₁ and CB₂ receptors, with K_i values generally not determined (i.e., pK_i < 5, <60% displacement of radioligand at 10 μM). CB₁ binding was greatest for derivatives featuring a pendant propyl (9: K_i = 5.37 μM) or benzyl group (10: K_i = 3.39 μM; 16: K_i = 2.24 μM) on the terminal piperazine nitrogen, suggesting a minimum steric requirement in this region for CB₁ binding. Fluorination had no major influence on CB₁ binding, with 10 showing a similar K_i value to the corresponding fluorinated compound, 16. CB₂ binding favored smaller piperazine substituents, with the highest affinities observed for the *N*-methyl (7: K_i = 1.18 μM) and *N*-ethyl analogues (8: K_i = 1.05 μM) in the desfluoro series and decreasing for the *N*-propyl (9: K_i = 2.09 μM) and *N*-benzyl (10: K_i = 6.31 μM) cases. In the corresponding fluorinated congeners, K_i values were only determinable for the ethyl (14) and benzyl cases (16) and not for the methyl (13) or propyl (15) derivatives. In both fluorinated and nonfluorinated series, neither an unsubstituted (6 and 12) nor a *tert*-butyl carbamate-protected (11 and 17) piperazine was tolerated for CB₁ or CB₂ binding.

In terms of functional activity at CB₁ and CB₂, all compounds demonstrated negligible or low potency agonist activity, which was not surprising given the low affinity for these receptors. It was not possible to determine EC₅₀ values; at 10 μM, all compounds produced only limited CB₁ or CB₂ receptor

activation compared to a maximally efficacious concentration of CP 55,940 (1 μ M; 100%). For CB₁, receptor activation ranged from approximately 0 to 44% of maximum response after normalization and approximately 4–49% of maximum for CB₂. Overall, several compounds appear to have some CB₁ and/or CB₂ agonist activity, albeit with apparent low potency compared to other recent generation SCARs.⁵² It was not possible to infer meaningful SARs for 6–17 given broadly similar low potency across all compounds.

The calcium channel activity of 6–17 was investigated using a fluorometric imaging plate-reader (FLIPR) calcium flux assay in Ca_v3.1-, Ca_v3.2-, and Ca_v3.3-transfected cells, and the data are shown in Table 2 and plotted in Figure 5, with representative

Table 2. Mean Calcium Fluorescence Inhibition at Ca_v3.1, Ca_v3.2, and Ca_v3.3 by Compounds 6–17 and NNC Using a Functional Calcium Flux Assay

compound	X	R	% fluorescence inhibition \pm SEM		
			Ca _v 3.1	Ca _v 3.2	Ca _v 3.3
NNC 55-0396			78.5 \pm 5.1	85.2 \pm 3.1	94.2 \pm 1.9
6	H	H	28.3 \pm 12.3	38.7 \pm 11.8	28.4 \pm 6.6
7	H	Me	40.9 \pm 8.1	42.6 \pm 9.9	29.1 \pm 5.8
8	H	Et	48.0 \pm 7.1	45.7 \pm 4.6	66.9 \pm 8.3
9	H	Pr	78.9 \pm 5.9	67.6 \pm 5.2	76.0 \pm 7.4
10	H	Bn	79.5 \pm 7.6	74.0 \pm 8.2	55.6 \pm 7.4
11	H	Boc	92.7 \pm 2.3	88.2 \pm 4.7	5.1 \pm 0.8
12	F	H	31.2 \pm 6.5	21.4 \pm 7.1	2.3 \pm 4.2
13	F	Me	36.1 \pm 7.7	29.6 \pm 11.5	14.2 \pm 5.8
14	F	Et	1.8 \pm 3.9	4.5 \pm 5.5	14.1 \pm 6.9
15	F	Pr	43.7 \pm 8.4	21.4 \pm 7.1	6.6 \pm 4.5
16	F	Bn	92.2 \pm 2.5	74.5 \pm 6.6	87.5 \pm 6.6
17	F	Boc	96.0 \pm 1.3	95.5 \pm 1.1	26.8 \pm 10.6

traces included in Supporting Figure S20. Figure 5 shows results of SCARs screened at 10 μ M in the calcium flux assay on stably expressed human T-type calcium channels in HEK293T cells (Ca_v3.1, Ca_v3.2, and Ca_v3.3). Inhibition induced by the nonselective Ca_v3 inhibitor NNC 55-0396 (NNC; 10 μ M) was included as a positive control. The functional calcium flux assay identified five compounds that potently inhibited (>70%, italic values in Table 2) the calcium response for one or more Ca_v3 subtype: 9, 10, 11, 16, and 17. Appreciable Ca_v3.1/3.2 inhibition was achieved for compounds with R = Bn (i.e., 10 and 16) or R = Boc (i.e., 11 and 17), regardless of compounds possessing a pentyl chain (X = H) or a 5-fluoropentyl group (X = F), suggesting that terminal fluorination does not influence

Ca_v3.1/3.2 inhibition with these pendant R-groups. Conversely, comparing 8 and 14 (R = Et) or 9 and 15 (R = Pr), terminal fluorination substantially reduces activity across all Ca_v subtypes.

Notably, both 11 and its corresponding fluorinated derivative 17 feature the *tert*-butyl carbamate moiety on the piperazine ring and showed a similar profile, with selective inhibition of Ca_v3.1 (93% and 96%, respectively) and Ca_v3.2 (88% and 96%, respectively) but not Ca_v3.3. Strong Ca_v3.2 inhibition was also achieved by carbazole analogues,³² indicating that the choice of carbazole or indole core is not a critical determinant of the Ca_v3.2 channel blockade. Only two compounds, 9 (X = H, R = Pr) and 16 (X = F, R = Bn), produced strong (>70%) inhibition of Ca_v3.3, suggesting that a more complex interaction between terminal fluorination and R-group is necessary for strong inhibition of this channel subtype in the calcium flux assay.

To confirm Ca_v3 inhibition by the more active compounds (9, 10, 11, 16, and 17), follow-up electrophysiology experiments were carried out (Figure 6; Supporting Figure S21). As

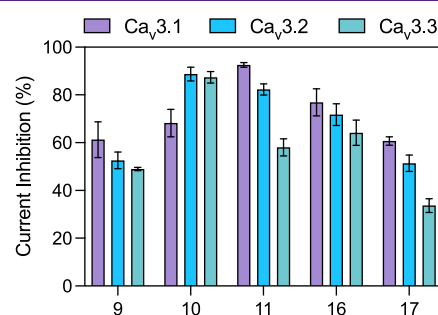


Figure 6. Mean current inhibition at Ca_v3.1, Ca_v3.2, and Ca_v3.3 by compounds 9–11, 16, and 17 using patch-clamp electrophysiology. Data are % current inhibition at 10 μ M and represent the mean \pm SEM of six independent whole-cell patch-clamp recordings per compound on each stably expressed ion channel.

compounds with little inhibitory activity display similarly low activity in electrophysiology experiments,⁵³ compounds 6, 7, 8, 12, 13, 14, and 15 were not tested in this platform. Compounds 9, 10, 11, 16, and 17 showed the inhibition of current for each of Ca_v3.1, Ca_v3.2, and Ca_v3.3, with, generally, slightly reduced inhibitory values compared to the calcium assay data. The greatest inhibition differences between the two assays were observed for Ca_v3.3, suggesting that the predictive utility of the calcium assay system is less robust for this subtype. For example, compared to the fluorescence inhibition in the calcium flux assay experiments, current inhibition at Ca_v3.3 decreased slightly for 9

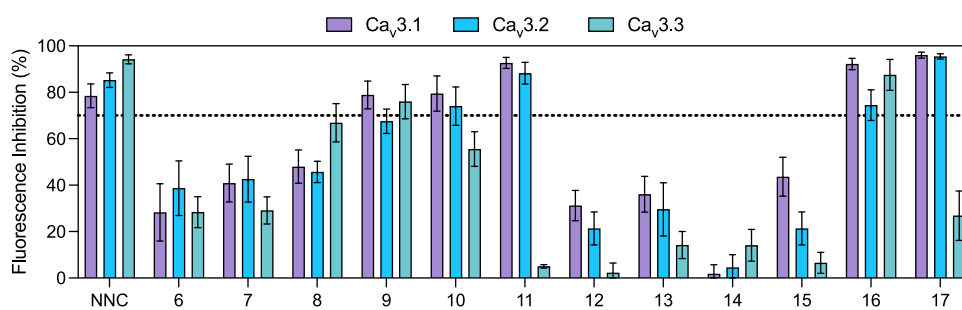


Figure 5. Mean fluorescence inhibition at Ca_v3.1, Ca_v3.2, and Ca_v3.3 by compounds 6–17 and positive control NNC 55-0396 (NNC; Supporting Figure S22) using a functional calcium flux assay. Data are % fluorescence inhibition at 10 μ M and represent the mean \pm standard error of the mean (SEM) of six independent experiments per compound.

and 16, increased for 10 and 11, and was consistent/unchanged for 17. This had the effect of abolishing any observed Ca_v3 subtype selectivity suggested by the calcium flux assay data.

Overall, 9, 10, 11, 16, and 17 are efficacious inhibitors of $\text{Ca}_v3.1$, $\text{Ca}_v3.2$, and $\text{Ca}_v3.3$, with little selectivity between the subtypes. Several of these compounds, in particular, 11 and 17, are structurally similar to cannabinoids shown to mediate potent analgesic responses *in vivo* via $\text{Ca}_v3.2$,^{31,32} whereas 9, 10, and 16 have important steric and electronic differences.

The observed differences between the results of the electrophysiology and functional assays on $\text{Ca}_v3.3$ could be due to the much slower kinetics of this subtype versus $\text{Ca}_v3.1$ and $\text{Ca}_v3.2$. Given the depolarized resting membrane potential (V_m) of HEK cells (approximately -60 mV), the functional assay may detect more inactivated state inhibition compared to the electrophysiology assay that holds cells artificially at -100 mV. Therefore, any putative inactivated state blocking component could be diminished and lead to changes in both apparent affinity and subtype dependence. To help mitigate this, low potassium HBSS was used in the calcium flux assay to keep the V_m of the cells at more negative potentials. In addition, depolarization of the membrane in the calcium flux assay was created by the addition of 10 mM Ca^{2+} , versus a voltage step protocol to -30 mV, to open the expressed channel subtype. Determination of whether the addition of the relatively high concentration of Ca^{2+} , versus 5 mM Ba^{2+} in the electrophysiology assay, contributes to the observed differences in $\text{Ca}_v3.3$ inhibition in the two assays would require detailed kinetic experiments that are beyond the scope of the current study.

MEPIRAPIM (7) and 5F-BEPIRAPIM (16) have been detected in the NPS market as putative SCRA based on structural motifs common in other SCRA. It was therefore hypothesized that these two compounds might possess cannabimimetic activities *in vivo*. In rodents, centrally active CB_1 receptor agonists produce hypothermia, hypolocomotion, and bradycardia that are typically concordant with the central cannabinoid activity in humans,^{11,54–57} and these effects may be used as a proxy for potential psychoactivity. To explore the potential cannabimimetic effects of these SCRA *in vivo*, the effect of MEPIRAPIM and 5F-BEPIRAPIM on core body temperature was evaluated in mice using radiotelemetry (Figure 7). No hypothermic effects were observed except for a modest and brief effect after the administration of 30 mg/kg MEPIRAPIM (AUC: $t(3) = 3.82$, $p = .02$). Compared to other recent generation SCRA tested using this model, which produce pronounced hypothermic effects (exceeding -5 °C from baseline) at doses as low as 0.3 mg/kg,^{1,3,11,54} MEPIRAPIM and 5F-BEPIRAPIM appear to possess relatively poor cannabimimetic efficacy in mice. However, this is entirely consistent with their *in vitro* pharmacological profiles at CB_1 (and CB_2) and mirrors our observations for other putative SCRA NPSs that possess comparably poor binding/agonist efficacy at CB_1 , e.g., 5F-PY-PICA or AB-001.^{29,58}

To gain a better understanding of the SARs underlying cannabinoid receptor binding, compounds 7, 9, 10, 14, 16, and Org 28611 were selected for induced fit docking in CB_1 (PDB: 6N4B) and CB_2 (PDB: 6PT0) receptors. The compounds share almost identical binding poses at both receptors, which is not surprising considering that the agonist-binding pockets of CB_1 and CB_2 receptors are very similar in nature.⁵⁹ In CB_1 receptors, the indole core makes π - π interactions with Phe268 and/or Phe200. The carbonyl group makes the H-bond interaction with

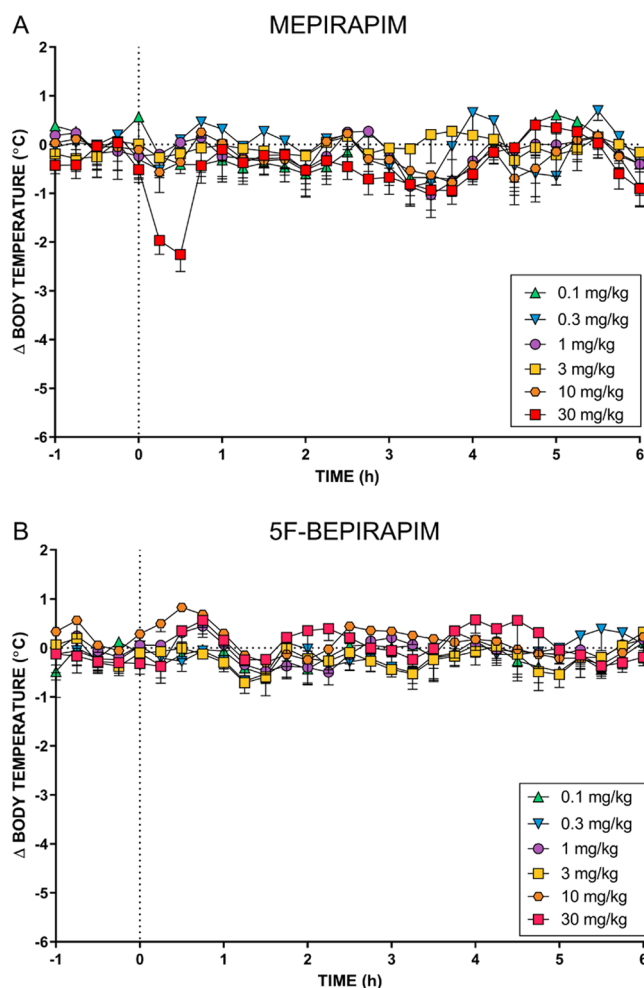


Figure 7. Change in body temperature following the intraperitoneal injection of (A) MEPIRAPIM and (B) 5F-BEPIRAPIM relative to a baseline vehicle injection. The vertical dashed line denotes the time of injection, and each data point represents the mean (\pm SEM) change in body temperature of four mice.

Ser383, and the tail group sits in the side pocket formed by Leu193, Thr197, Thr275, Ile271, Trp276, and Met363. The alkyl substituent at the piperazine ring makes hydrophobic interactions with Lys192, Phe177, and Phe189, whereas compounds 10 and 16, with a benzene substituent, also form π - π interaction with Phe177 and Phe189.

Similarly, in the CB_2 receptor, the indole ring makes π - π interactions with Phe117 and/or Phe183, and the carbonyl group makes H-bond interaction with Ser285. The tail group now sits in the hydrophobic pocket that is surrounded by Thr114, Phe183, Tyr190, Leu191, Leu193, Trp194, and Met265. The substituents at the piperazine ring interact with hydrophobic residues Phe94, Phe106, and Ile110, and the benzene ring of compounds 10 and 16 also makes π - π interactions with Phe94.

Sitemap^{60,61} analysis of the CB_1 and CB_2 orthosteric binding sites shows that the CB_2 receptor has a much smaller binding site volume (284.69 \AA^3) than the CB_1 receptor (502.50 \AA^3).

Compound 7 (MEPIRAPIM) and Org 28611 were directly compared in the CB_1 receptor to identify why MEPIRAPIM is relatively inactive at CB_1 compared to Org 28611 (18) despite a number of structural similarities. Compared to 7, Org 28611 has a methyl substitution at the 3-position of the piperazine ring and

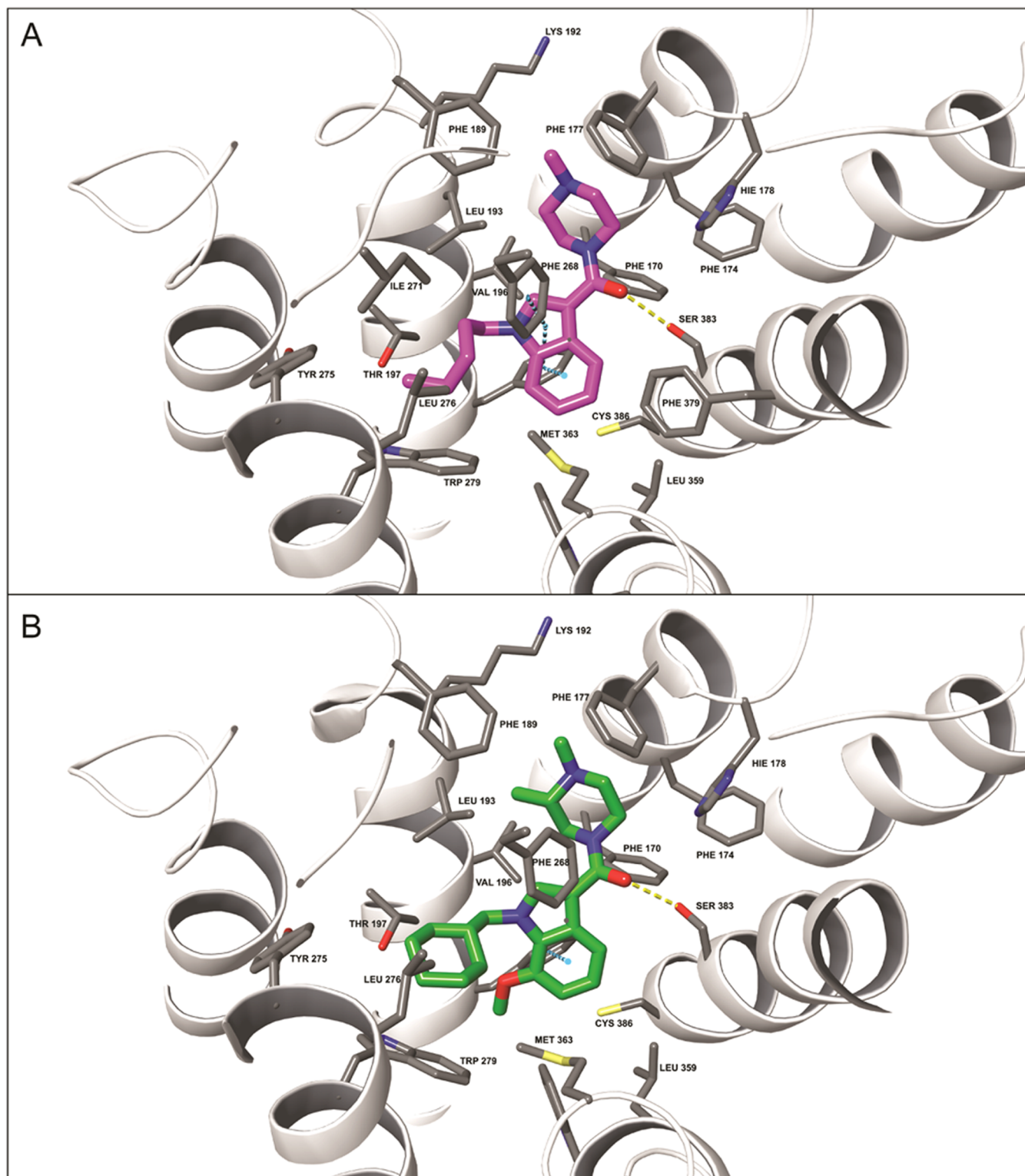


Figure 8. Docking simulation at CB₁ receptors for (A) MEPIRAPIM (7) and (B) Org 28611. Hydrophobic contacts between Leu193/Val196 and the methyl substituent of Org 28611 (18) appear to be a key determinant of the CB₁ receptor binding strength.

a methoxy substituent at the 6-position of the indole ring. No interactions between the methoxy group and the binding site were observed in the docking analysis. The methyl substituent of the piperazine ring of Org 28611 made hydrophobic contacts with Leu193 and Val196 and appeared to be responsible for the difference in potency between the two compounds (Figure 8). A methyl group at the 3- or 5-position of the piperazine ring is

reported to improve potency,⁶² and our previous work has also shown that SCRAAs that interact with these two residues may give rise to stronger binding affinity to the receptor.⁶³ The IFD score, which indicates binding affinity, shows that Org 28611 has a stronger binding affinity to both the CB₁ receptors compared to compound 7 (−9744.10 and −9726.62 kcal/mol, respectively).

CONCLUSIONS

Based on recent detections of MEPIRAPIM and 5F-BEPIRAPIM in drug markets and due to their structural similarities to known Ca_v3 inhibitors, we tested a library of MEPIRAPIM and 5F-BEPIRAPIM analogues for cannabinoid receptor and Ca_v3 activity. All tested compounds possessed low or negligible potency and efficacy at CB_1 and CB_2 receptors *in vitro*. A subset of compounds (**9**, **10**, **11**, **16**, and **17**) showed broad inhibition of Ca_v3 channels, with terminal fluorination having minimal impact on Ca_v3 activity or subtype selectivity, as measured by patch-clamp electrophysiology. Interestingly, Bladen et al. (2015) demonstrated that the related carbazole compound DX332 (Supporting Figure S23) had no activity on CB_1 and CB_2 and other carbazole derivatives in this series also did not interact strongly with CB_1 and CB_2 . However, in the series of carbazole derivatives described by You et al. (2011), CB receptor interactions were observed. Together with the present study, these reports indicate that both indole and carbazole derivatives can have mixed Ca_v3/CB receptor activity and that the substituents in the R position (Figure 2) are more critical for determining whether a compound acts on Ca_v3 , CBRs, or both. Further exploration and optimization of these compound classes may be warranted to develop an effective combined CBR agonist and Ca_v3 inhibitor.

The poor *in vivo* potency and efficacy of MEPIRAPIM and 5F-BEPIRAPIM, as compared to recent generation carboxamide-type SCRA, suggest limited psychoactivity in humans, which appears to be at odds with the detection of these compounds in the NPS market. However, this is not the first time that apparently nonpsychoactive SCRA have been detected. We previously reported that 5F-PY-PICA and 5F-PY-PINACA, detected in drug markets in 2015, possessed low binding affinity and efficacy at CB_1 receptors and that they failed to produce hypothermic effects in mice.²⁹ A potential explanation for this phenomenon is poor or absent pharmacological assessment in illicit/ clandestine SCRA development production. Since clandestine NPS producers are unlikely to be systematically testing new scaffold-hopping compounds for CB_1 activity *in vitro* or *in vivo*, from time to time, unexpectedly low efficacy compounds may be introduced into the NPS market. However, we cannot exclude the possibility that these compounds could exert psychoactive effects by noncannabinoid-mediated mechanisms.

METHODS

General Chemical Synthesis Details. All reactions were performed under an atmosphere of nitrogen unless otherwise specified. All reagents, reactants, and solvents were obtained from Sigma-Aldrich (St. Louis, MO) and used as purchased. Analytical thin-layer chromatography was performed using Merck aluminum-backed silica gel 60 F254 (0.2 mm) plates (Merck, Darmstadt, Germany), which were visualized using shortwave (254 nm) UV fluorescence. Flash chromatography was performed using a Biotage Isolera Spektra One and Biotage SNAP KP-Sil silica cartridges (Uppsala, Sweden), with gradient elution terminating at the solvent combination indicated for each compound (*vide infra*). Melting point ranges (m.p.) were measured in open capillaries using a Stuart SMP50 Automated melting point apparatus (Cole-Palmer, Staffordshire, U.K.) and are uncorrected. Nuclear magnetic resonance spectra were recorded at 298 K using an Agilent 400 MHz spectrometer (Santa Clara, CA). The data are reported as chemical shift (δ ppm) relative to the residual protonated solvent resonance, multiplicity (s = singlet, br s = broad singlet, d = doublet, br d = broad doublet, t = triplet, q = quartet, quin. = quintet, m = multiplet), coupling constants (*J* Hz), relative integral, and

assignment. High-resolution mass spectrometry (HRMS) data were collected using an Agilent LC 1260-QTOF-MS 6550 (Santa Clara, CA) or a Thermo Scientific Q Exactive HF-X hybrid quadrupole-orbitrap mass spectrometer (Waltham, MA). A methanolic extract of each pure standard was run using an electrospray ionization source in an automated MS/MS (information-dependent acquisition) mode. Accurate mass for the parent ion and its corresponding mass error expressed in parts per million (ppm) is reported.

General Procedure for the Synthesis of (4-Alkylpiperazin-1-yl)(1-alkyl-1H-indol-3-yl)methanones (7–10 and 13–16). To a solution of the free base of the appropriate (piperazin-1-yl)(1-alkyl-1H-indol-3-yl)methanone (**6** or **12**, 1 mmol) in 1,2-dichloroethane (3.5 mL) was added the appropriate aldehyde (1 mmol, 1.0 equiv), followed by sodium triacetoxymethylborohydride (297 mg, 1.4 mmol, 1.4 equiv) in a single portion and the mixture was stirred at ambient temperature for 2 h. The reaction was quenched by the addition of 1 M aq. NaOH (3.5 mL), and the layers were separated. The aqueous phase was extracted with CH_2Cl_2 (3 \times 3.5 mL), and the combined organic layers were washed with brine (10 mL), dried (Na_2SO_4), and the solvent was evaporated under reduced pressure. The crude products were purified by flash chromatography, converted to their hydrochloride salts, and recrystallized for pharmacological evaluation. Analytical data were obtained for each compound as the hydrochloride salt (M.p., HRMS) or as the free base liberated from the corresponding salt by dissolution in H_2O , adjusting the pH to 14 (1 M aq. NaOH), extracting with CH_2Cl_2 (and drying the extracts over Na_2SO_4), evaporating the solvent under reduced pressure, and drying the resultant oil under high vacuum (R_f , 1H , and ^{13}C NMR).

(4-Methylpiperazin-1-yl)(1-pentyl-1H-indol-3-yl)methanone Hydrochloride (7-HCl, MEPIRAPIM-HCl). Subjecting **6** (300 mg, 1.00 mmol) and 37% aqueous formaldehyde (80 μ L, 1.00 mmol, 1.0 equiv) to the procedure described above with additional sodium triacetoxymethylborohydride (848 mg, 4.00 mmol, 4.0 equiv)⁶⁴ gave, following purification by flash chromatography (CH_2Cl_2 -MeOH, 90:10), **7** as a colorless oil (278 mg, 89%). The free base was converted to the hydrochloride and recrystallized from EtOAc to give a white solid. M.p. (HCl) 184–186 °C; R_f (free base) 0.14 (CH_2Cl_2 -MeOH, 95:5); 1H NMR (400 MHz, $CDCl_3$, free base): δ 7.71–7.68 (m, 1H), 7.46 (s, 1H), 7.35–7.70 (m, 1H), 7.18–7.27 (m, 2H), 4.12 (t, *J* = 7.2 Hz, 2H, NCH_2), 3.75 (t, *J* = 4.8 Hz, 4H, 2 \times NCH_2), 2.45 (t, *J* = 4.8 Hz, 4H, 2 \times NCH_2), 2.33 (s, 3H, NCH_3), 1.86 (quin., *J* = 7.2 Hz, 2H, CH_2), 1.30–1.38 (m, 4H, 2 \times CH_2), 0.89 (t, *J* = 7.2 Hz, 3H, CH_3); ^{13}C NMR (100 MHz, $CDCl_3$, free base): δ 166.9 (CO), 135.9 (quat.), 130.8 (CH), 126.3 (quat.), 122.4 (CH), 120.9 (CH), 120.8 (CH), 110.5 (quat.), 110.1 (CH), 55.4 (2 \times NCH_2), 46.9 (NCH_2), 46.3 (NCH_3), 45.4 (br, 2 \times NCH_2), 30.0 (CH_2), 29.2 (CH_2), 22.4 (CH_2), 14.1 (CH_3); HRMS (ESI) $C_{19}H_{27}N_3O$ exact mass 313.2154, accurate mass 313.2158 (mass error 1.09 ppm).

(4-Ethylpiperazin-1-yl)(1-pentyl-1H-indol-3-yl)methanone Hydrochloride (8-HCl). Subjecting **6** (300 mg, 1.00 mmol) and acetaldehyde (56 μ L, 1.00 mmol, 1.0 equiv) to the general procedure gave, following purification by flash chromatography (CH_2Cl_2 -MeOH, 90:10), **8** as a colorless oil (304 mg, 93%). The free base was converted to the hydrochloride and recrystallized from EtOAc to give a white solid. M.p. (HCl) 165–167 °C; R_f (free base) 0.15 (CH_2Cl_2 -MeOH, 95:5); 1H NMR (400 MHz, $CDCl_3$, free base): δ 7.46 (s, 1H), 7.34–7.37 (m, 1H), 7.23–7.27 (m, 1H), 7.20 (m, 1H), 4.12 (t, *J* = 7.2 Hz, 2H, NCH_2), 3.76 (t, *J* = 4.8 Hz, 4H, 2 \times CH_2), 2.44–2.51 (m, 6H, 3 \times CH_3), 1.87 (quin., *J* = 7.2 Hz, 2H, CH_2), 1.03–1.38 (m, 4H, 2 \times CH_2), 1.11 (t, *J* = 7.2 Hz, 3H, CH_3), 0.89 (t, *J* = 7.2 Hz, 3H, CH_3); ^{13}C NMR (100 MHz, $CDCl_3$, free base): δ 166.8 (CO), 136.0 (quat.), 130.8 (CH), 126.3 (quat.), 122.3 (CH), 120.8(8) (CH), 120.8(6) (CH), 110.6 (quat.), 110.1 (CH), 53.3 (2 \times NCH_2), 52.5 (NCH_2), 46.9 (NCH_2), 29.9 (CH_2), 29.2 (CH_2), 22.4 (CH_2), 14.1 (CH_3), 12.1 (CH_3) (one missing or overlapping); HRMS (ESI) $C_{20}H_{29}N_3O$ exact mass 327.2311, accurate mass 327.2309 (mass error –0.40 ppm).

(4-Propylpiperazin-1-yl)(1-pentyl-1H-indol-3-yl)methanone Hydrochloride (9-HCl). Subjecting **6** (300 mg, 1.00 mmol) and propanal (72 μ L, 1.00 mmol, 1.0 equiv) to the general procedure gave, following purification by flash chromatography (CH_2Cl_2 -MeOH, 95:5), **9** as a

colorless oil (290 mg, 85%). The free base was converted to the hydrochloride and recrystallized from EtOAc to give a white solid. M.p. (HCl) 102–105 °C; R_f (free base) 0.20 (CH₂Cl₂-MeOH, 95:5); ¹H NMR (400 MHz, DMSO-*d*₆, free base): δ 7.79 (s, 1H), 7.73–7.75 (m, 1H), 7.55–7.57 (m, 1H), 7.23 (ddd, *J* = 8.3, 7.0, 1.2 Hz, 1H), 7.15 (ddd, *J* = 8.0, 7.0, 1.0 Hz, 1H), 4.37 (br d, *J* = 14.0 Hz, 2H, CH₂), 4.22 (t, *J* = 7.2 Hz, 2H, NCH₂), 3.46–3.51 (m, 4H, 2 × CH₂), 3.02–3.09 (m, 4H, 2 × CH₂), 1.67–1.82 (m, 4H, 2 × CH₂), 1.18–1.35 (m, 4H, 2 × CH₂); ¹³C NMR (100 MHz, DMSO-*d*₆, free base): δ 165.8 (CO), 136.1 (quat.), 132.3 (CH), 126.9 (quat.), 122.6 (CH), 121.1 (CH), 120.9 (CH), 111.0 (CH), 108.0 (quat.), 57.5 (NCH₂), 51.2 (2 × NCH₂), 46.3 (NCH₂), 41.6 (2 × NCH₂), 29.8 (CH₂), 28.8 (CH₂), 22.2 (CH₂), 17.1 (CH₂), 14.3 (CH₃), 11.4 (CH₃); HRMS (ESI) C₂₁H₃₁N₃O exact mass 341.2467, accurate mass 341.2467 (mass error –0.14 ppm).

(4-Benzylpiperazin-1-yl)(1-pentyl-1H-indol-3-yl)methanone Hydrochloride (10-HCl). Subjecting **6** (300 mg, 1.00 mmol) and benzaldehyde (102 μL, 1.00 mmol, 1.0 equiv) to the general procedure gave, following purification by flash chromatography (CH₂Cl₂-MeOH, 95:5), **10** as a colorless oil (370 mg, 95%). The free base was converted to the hydrochloride and recrystallized from MeOH-Et₂O to give a white solid. M.p. (HCl) 219–221 °C; R_f (free base) 0.38 (CH₂Cl₂-MeOH, 95:5); ¹H NMR (400 MHz, CDCl₃, free base): δ 7.69–7.72 (m, 1H), 7.44 (s, 1H), 7.30–7.36 (m, 5H), 7.17–7.28 (m, 3H), 4.11 (t, *J* = 7.2 Hz, 2H, NCH₂), 3.74 (t, *J* = 4.8 Hz, 4H, 2 × CH₂), 3.55 (s, 2H, CH₂Ph), 2.50 (t, *J* = 4.8 Hz, 4H, 2 × CH₂), 1.86 (quin., *J* = 7.2 Hz, 2H, CH₂), 1.26–1.41 (m, 4H, 2 × CH₂), 0.89 (t, *J* = 6.8 Hz, 3H, CH₃); ¹³C NMR (100 MHz, CDCl₃, free base): δ 166.8 (CO), 137.9 (quat.), 135.9 (quat.), 130.7 (CH), 129.3 (2 × CH), 128.4 (2 × CH), 127.3 (CH), 126.3 (quat.), 122.3 (CH), 120.9 (CH), 120.8 (CH), 110.6 (quat.), 110.0 (CH), 63.1 (CH₂Ph), 53.5 (2 × NCH₂), 46.8 (NCH₂), 45.5 (2 × NCH₂), 29.9 (CH₂), 29.2 (CH₂), 22.4 (CH₂), 14.1 (CH₃); HRMS (ESI) C₂₅H₃₁N₃O exact mass 389.2467, accurate mass 389.2467 (mass error 0.40 ppm).

(4-Methylpiperazin-1-yl)(1-(5-fluoropentyl)-1H-indol-3-yl)methanone Hydrochloride (13-HCl). Subjecting **12** (317 mg, 1.00 mmol) and 37% aqueous formaldehyde (80 μL, 1.00 mmol, 1.0 equiv) to the procedure described above with additional sodium triacetoxymethylborohydride (848 mg, 4.00 mmol, 4.0 equiv)⁶⁴ gave, following purification by flash chromatography (CH₂Cl₂-MeOH, 90:10), **13** as a colorless oil (298 mg, 90%). The free base was converted to the hydrochloride and recrystallized from *i*-PrOH-Et₂O to give a white solid. M.p. (HCl) 157–159 °C; R_f (free base) 0.14 (CH₂Cl₂-MeOH, 95:5); ¹H NMR (400 MHz, CDCl₃, free base): δ 7.69–7.71 (m, 1H), 7.45 (s, 1H), 7.34–7.36 (m, 1H), 7.24–7.28 (m, 1H), 7.18–7.23 (m, 1H), 4.42 (dt, ²*J*_{HF} = 47.6 Hz, ³*J*_{HH} = 6.0 Hz, 2H, CH₂F), 4.15 (t, *J* = 7.2 Hz, 2H, NCH₂), 3.74 (t, *J* = 4.8 Hz, 4H, 2 × CH₂), 2.45 (t, *J* = 4.8 Hz, 4H, 2 × CH₂), 2.33 (s, 3H, NCH₃), 1.92 (quin., *J* = 7.2 Hz, 2H, CH₂), 1.67–1.77 (m, 2H, CH₂), 1.43–1.51 (m, 2H, CH₂); ¹³C NMR (100 MHz, CDCl₃, free base): δ 166.7 (CO), 135.9 (quat.), 130.7 (CH), 126.3 (quat.), 122.5 (CH), 121.0 (CH), 120.9 (CH), 110.8 (quat.), 110.0 (CH), 83.8 (d, ¹*J*_{CF} = 164.0 Hz, CH₂F), 55.5 (2 × NCH₂), 46.7 (NCH₂), 46.3 (NCH₃), 45.2 (2 × NCH₂), 30.1 (d, ²*J*_{CF} = 20.0 Hz, CH₂), 29.9 (CH₃), 23.0 (d, ³*J*_{CF} = 5.0 Hz, CH₂); HRMS (ESI) C₁₉H₂₆FN₃O exact mass 331.2060, accurate mass 331.2063 (mass error 0.93 ppm).

(4-Ethylpiperazin-1-yl)(1-(5-fluoropentyl)-1H-indol-3-yl)methanone Hydrochloride (14-HCl). Subjecting **12** (317 mg, 1.00 mmol) and acetaldehyde (56 μL, 1.00 mmol, 1.0 equiv) to the general procedure gave, following purification by flash chromatography (CH₂Cl₂-MeOH, 90:10), **14** as a colorless oil (324 mg, 94%). The free base was converted to the hydrochloride and recrystallized from *i*-PrOH-Et₂O to give a white solid. M.p. (HCl) 175–177 °C; R_f (free base) 0.17 (CH₂Cl₂-MeOH, 95:5); ¹H NMR (400 MHz, CDCl₃, free base): δ 7.69–7.71 (m, 1H), 7.44 (s, 1H), 7.33–7.36 (m, 1H), 7.17–7.27 (m, 2H), 4.41 (dt, ²*J*_{HF} = 47.2 Hz, ³*J*_{HH} = 6.0 Hz, 2H, CH₂F), 4.14 (t, *J* = 7.2 Hz, 2H, NCH₂), 3.75 (t, *J* = 4.8 Hz, 4H, 2 × NCH₂), 2.43–2.50 (m, 6H, 2 × NCH₂ and 1 × CH₂), 1.91 (quin., *J* = 7.2 Hz, 2H, CH₂), 1.64–1.76 (m, 2H, CH₂), 1.42–1.50 (m, 2H, CH₂), 1.10 (t, *J* = 7.2 Hz, 3H, CH₃); ¹³C NMR (free base, CDCl₃, 100 MHz): δ 166.6

(CO), 135.8 (quat.), 130.6 (CH), 126.3 (quat.), 122.4 (CH), 120.92 (CH), 120.87 (CH), 110.7 (quat.), 110.0 (CH), 83.8 (d, ¹*J*_{CF} = 164.0 Hz, CH₂F), 53.2 (2 × NCH₂), 52.4 (NCH₂), 46.7 (NCH₂), 45.4 (2 × CH₂), 30.0 (d, ²*J*_{CF} = 20.0 Hz, CH₂), 29.8 (CH₂), 23.0 (d, ³*J*_{CF} = 5.0 Hz, CH₂), 12.1 (CH₃); HRMS (ESI) C₂₀H₂₈FN₃O exact mass 345.2216, accurate mass 345.2217 (mass error 0.17 ppm).

(4-Propylpiperazin-1-yl)(1-(5-fluoropentyl)-1H-indol-3-yl)methanone Hydrochloride (15-HCl). Subjecting **12** (317 mg, 1.00 mmol) and propanal (72 μL, 1.00 mmol, 1.0 equiv) to the general procedure gave, following purification by flash chromatography (CH₂Cl₂-MeOH, 95:5), **15** as a colorless oil (342 mg, 95%). The free base was converted to the hydrochloride and recrystallized from EtOAc to give a white solid. M.p. (HCl) 105–108 °C; R_f (free base) 0.29 (CH₂Cl₂-MeOH, 95:5); ¹H NMR (400 MHz, CDCl₃, free base): δ 7.69–7.71 (m, 1H), 7.44 (s, 1H), 7.34–7.36 (m, 1H), 7.18–7.28 (m, 2H), 4.42 (dt, ²*J*_{HF} = 47.2 Hz, ³*J*_{HH} = 6.0 Hz, 2H, CH₂F), 4.14 (t, *J* = 7.2 Hz, 2H, NCH₂), 3.74 (t, *J* = 4.8 Hz, 4H, 2 × NCH₂), 2.48 (t, *J* = 4.8 Hz, 4H, 2 × NCH₂), 2.32–2.36 (m, 2H, CH₂), 1.91 (quin., *J* = 7.2 Hz, 2H, CH₂), 1.66–1.77 (m, 2H, CH₂), 1.43–1.57 (m, 4H, 2 × CH₂), 0.91 (t, *J* = 7.2 Hz, 3H, CH₃); ¹³C NMR (free base, CDCl₃, 100 MHz): δ 166.6 (CO), 135.9 (quat.), 130.6 (CH), 126.3 (quat.), 122.4 (CH), 120.94, (CH), 120.91 (CH), 110.8 (quat.), 110.0 (CH), 83.8 (d, ¹*J*_{CF} = 164.0 Hz, CH₂F), 60.7 (NCH₂), 53.6 (2 × NCH₂), 46.7 (NCH₂), 45.3 (2 × NCH₂), 30.1 (d, ²*J*_{CF} = 19.0 Hz, CH₂), 29.9 (CH₂), 23.0 (d, ³*J*_{CF} = 5.0 Hz, CH₂), 20.1 (CH₂), 12.0 (CH₃); HRMS (ESI) C₂₁H₃₀FN₃O exact mass 359.2373, accurate mass 359.2375 (mass error 0.46 ppm).

(4-Benzylpiperazin-1-yl)(1-(5-fluoropentyl)-1H-indol-3-yl)methanone Hydrochloride (16-HCl). Subjecting **12** (317 mg, 1.00 mmol) and benzaldehyde (102 μL, 1.00 mmol, 1.0 equiv) to the general procedure gave, following purification by flash chromatography (CH₂Cl₂-MeOH, 95:5), **16** as a colorless oil (383 mg, 94%). The free base was converted to the hydrochloride and recrystallized from MeOH-Et₂O to give a white solid. M.p. (HCl) 199–201 °C; R_f (free base) 0.34 (CH₂Cl₂-MeOH, 95:5); ¹H NMR (400 MHz, CDCl₃, free base): δ 7.69–7.72 (m, 1H), 7.44 (s, 1H), 7.30–7.36 (m, 5H), 7.17–7.28 (m, 3H), 4.42 (dt, ²*J*_{HF} = 47.6 Hz, CH₂F, 2H), 4.14 (t, *J* = 7.2 Hz, 2H, NCH₂), 3.74 (t, *J* = 4.8 Hz, 4H, 2 × NCH₂), 3.55 (s, 2H, CH₂Ph), 2.50 (t, *J* = 4.8 Hz, 4H, 2 × NCH₂), 1.91 (quin., *J* = 7.6 Hz, 2H, CH₂), 1.65–1.78 (m, 2H, CH₂), 1.43–1.51 (m, 2H, CH₂); ¹³C NMR (100 MHz, CDCl₃, free base): δ 166.7 (CO), 137.9 (quat.), 135.9 (quat.), 130.6 (CH), 129.3 (2 × CH), 128.4 (2 × CH₂), 127.3 (CH), 126.3 (quat.), 122.4 (CH), 120.9 (CH), 110.8 (quat.), 110.0 (CH), 83.8 (d, ¹*J*_{CF} = 164.0 Hz, CH₂F), 63.1 (CH₂Ph), 53.5 (2 × NCH₂), 46.7 (NCH₂), 45.5 (2 × NCH₂), 30.1 (d, ²*J*_{CF} = 19.0 Hz, CH₂), 29.85 (CH₂), 23.0 (d, ³*J*_{CF} = 5.0 Hz, CH₂); HRMS (ESI) C₂₅H₃₀FN₃O exact mass 407.2373, accurate mass 407.2374 (mass error 0.20 ppm).

Synthesis of Piperazin-1-yl(1-alkyl-1H-indol-3-yl)methanone Hydrochlorides (6-HCl and 12-HCl). (Piperazin-1-yl)(1-pentyl-1H-indol-3-yl)methanone Hydrochloride (**6-HCl**). *tert*-Butyl 4-(1-pentyl-1H-indole-3-carbonyl)piperazine-1-carboxylate (**11**, 639 mg, 1.6 mmol) was dissolved in 1,4-dioxane (12 mL) and added dropwise to a cooled (0 °C) solution of 4 M hydrogen chloride in 1,4-dioxane (4 mL, 16 mmol, 10 equiv). The solution was stirred for 16 h, and the reaction mixture was evaporated under reduced pressure to afford the crude hydrochloride salt, which was washed with ice-cold anhydrous Et₂O (3 × 10 mL) and recrystallized from MeOH-Et₂O to afford **6-HCl** as a white solid (489 mg, 91%). M.p. (HCl) 161–163 °C; ¹H NMR (400 MHz, CDCl₃, free base): δ 7.68–7.70 (m, 1H), 7.45 (s, 1H), 7.34–7.37 (m, 1H), 7.17–2.27 (m, 2H), 4.12 (t, *J* = 7.2 Hz, 2H, NCH₂), 3.71 (t, *J* = 4.8 Hz, 4H, 2 × CH₂), 2.91 (t, *J* = 4.8 Hz, 4H, 2 × CH₂), 1.86 (quin., *J* = 7.2 Hz, 2H, CH₂), 1.28–1.39 (m, 4H, 2 × CH₂), 0.89 (t, *J* = 7.2 Hz, 3H, CH₃); ¹³C NMR (100 MHz, CDCl₃, free base): δ 166.9 (CO), 135.9 (quat.), 130.7 (CH), 126.2 (quat.), 122.4 (CH), 120.9 (CH), 120.8 (CH), 110.5 (quat.), 110.1 (CH), 46.9 (NCH₂), 46.5(2) (2 × NCH₂), 46.5(0) (br overlapping, 2 × NCH₂), 29.9, (CH₂), 29.2 (CH₂), 22.4 (CH₂), 14.1 (CH₃); HRMS (ESI) C₁₈H₂₅N₃O exact mass 299.1998, accurate mass 299.2001 (mass error 0.98 ppm).

(Piperazin-1-yl)(1-(5-fluoropentyl)-1H-indol-3-yl)methanone Hydrochloride (**12-HCl**). Subjecting *tert*-butyl 4-(1-(5-fluoropentyl)-1H-

indole-3-carbonyl)piperazine-1-carboxylate (**17**, 668 mg, 1.6 mmol) to the procedure described for **6•HCl** gave **12•HCl**, which was recrystallized from MeOH-Et₂O to furnish a white solid (476 mg, 84%). M.p. (HCl) 139–142 °C; ¹H NMR (400 MHz, CDCl₃, free base): δ 7.67–7.70 (m, 1H), 7.46 (s, 1H), 7.34–7.36 (m, 1H), 7.28–7.18 (m, 2H), 4.42 (dt, ²J_{HF} = 47.2 Hz, ³J_{HH} = 6.0 Hz, 2H, CH₂F), 4.15 (t, J = 7.2 Hz, 2H, NCH₂), 3.72 (t, J = 5.2 Hz, 4H, 2 × NCH₂), 2.93 (t, J = 5.2 Hz, 4H, 2 × NCH₂), 1.92 (quin., J = 7.6 Hz, 2H, CH₂), 1.65–1.78 (m, 2H, 2 × CH₂), 1.43–1.51 (m, 2H, 2 × CH₂); ¹³C NMR (100 MHz, CDCl₃, free base): δ 166.9 (CO), 135.9 (quat.), 130.7 (CH), 126.2 (quat.), 122.5 (CH), 121.0 (CH), 120.8 (CH), 110.6 (quat.), 110.0 (CH), 83.8 (s, ¹J_{CF} = 164.0 Hz, CH₂F), 46.7 (NCH₂), 46.3 (2 × NCH₂), 46.2 (br overlapping, 2 × NCH₂), 30.1 (d, ²J_{CF} = 20.0 Hz, CH₂), 29.8 (CH₃), 23.0 (d, ³J_{CF} = 5.0 Hz, CH₂); HRMS (ESI) C₁₈H₂₄FN₃O exact mass 317.1903, accurate mass 317.1908 (mass error 1.32 ppm).

Synthesis of tert-Butyl 4-(1-Alkyl-1H-indole-3-carbonyl)piperazine-1-carboxylates (11 and 17). tert-Butyl 4-(1-Pentyl-1H-indole-3-carbonyl)piperazine-1-carboxylate (**11**). To a solution of 1-pentyl-1H-indole-3-carboxylic acid (**23**, 463 mg, 2.0 mmol) in DMF (10 mL) was added HOBt•H₂O (337 mg, 2.2 mmol, 1.1 equiv), EDC•HCl (498 mg, 2.6 mmol, 1.3 equiv), 1-Boc-piperazine (373 mg, 2.0 mmol, 1.0 equiv), and Et₃N (833 μL, 6.0 mmol, 3.0 equiv), and the mixture was stirred at ambient temperature for 14 h. The mixture was poured onto H₂O (400 mL) and extracted with EtOAc (3 × 100 mL), and the combined organic phases were washed with H₂O (2 × 100 mL) and brine (100 mL), dried (MgSO₄), and the solvent was evaporated under reduced pressure. The crude material was purified using flash chromatography (CH₂Cl₂-MeOH, 98:2) to give **11** as a colorless solid (734 mg, 92%). M.p. 82–84 °C; R_f 0.38 (CH₂Cl₂-MeOH, 95:5); ¹H NMR (400 MHz, CDCl₃): δ 7.68 (d, J = 7.4 Hz, 1H), 7.47 (s, 1H), 7.37 (d, J = 7.9 Hz, 1H), 7.17–7.29 (m, 2H), 4.12 (t, J = 7.2 Hz, 2H, NCH₂), 3.63–3.75 (m, 4H, 2 × NCH₂), 3.45–3.55 (m, 4H, 2 × NCH₂), 1.87 (quin., J = 7.2 Hz, 2H, CH₂), 1.47 (s, 9H, 3 × CH₃), 1.28–1.41 (m, 4H, 2 × CH₂), 0.89 (t, J = 6.8 Hz, 3H, CH₃); ¹³C NMR (100 MHz, CDCl₃): δ 167.1 (CO), 154.9 (NCO₂tBu), 135.9 (quat.), 130.9 (CH), 126.1 (quat.), 122.5 (CH), 121.0 (CH), 120.7 (CH), 110.6 (CH), 110.1 (quat.), 80.3 (quat.), 46.9 (NCH₂), 45.4 (br, 2 × NCH₂), 44.0 (br, 2 × NCH₂), 29.8 (CH₂), 29.2 (CH₂), 28.5 (3 × CH₃), 22.4 (CH₂), 14.0 (CH₃); HRMS (ESI) C₂₃H₃₃N₃O₃ exact mass 399.2522, accurate mass 399.2523 (mass error 0.23 ppm).

tert-Butyl 4-(1-(5-Fluoropentyl)-1H-indole-3-carbonyl)piperazine-1-carboxylate (**17**). Subjecting 1-(5-fluoropentyl)-1H-indole-3-carboxylic acid (**24**, 463 mg, 2.0 mmol) to the procedure described for **11** afforded, following purification by flash chromatography (CH₂Cl₂-MeOH, 95:5), **17** as a colorless resin (741 mg, 89%). R_f 0.29 (CH₂Cl₂-MeOH, 95:5); ¹H NMR (400 MHz, CDCl₃): δ 7.68 (d, J = 7.5 Hz, 1H), 7.46 (s, 1H), 7.36 (d, 1H), 7.17–7.30 (m, 2H), 4.43 (dt, ²J_{HF} = 47.3, ³J_{HH} = 5.9 Hz, 2H, CH₂F), 4.16 (t, J = 7.2 Hz, 2H, NCH₂), 3.63–3.75 (m, 4H, 2 × NCH₂), 3.46–3.53 (m, 4H, 2 × NCH₂), 1.92 (quin., J = 7.2 Hz, 2H, CH₂), 1.63–1.83 (m, 2H, CH₂), 1.47 (s overlapping, 9H, 3 × CH₃), 1.44–1.53 (m overlapping, 2H, CH₂); ¹³C NMR (100 MHz, CDCl₃): δ 167.0 (CO), 154.8 (NCO₂tBu), 135.9 (quat.), 130.8 (CH), 126.2 (quat.), 122.6 (CH), 121.1 (CH), 120.8 (CH), 110.4 (quat.), 110.1 (CH), 83.76 (d, ¹J_{CF} = 165.0 Hz, CH₂), 80.3 (quat.), 46.7 (NCH₂), 45.3 (br, 2 × NCH₂), 44.6 (br, 2 × NCH₂), 30.7 (d, ²J_{CF} = 19.8 Hz, CH₂), 29.8 (CH₂), 28.5 (3 × CH₃), 23.0 (d, ²J_{CF} = 4.9 Hz, CH₂); HRMS (ESI) C₂₃H₃₂FN₃O₃ exact mass 417.2428, accurate mass 417.2433 (mass error 1.28 ppm).

General Procedure for Synthesis of 1-Alkyl-1H-indole-3-carboxylic Acids (23 and 24). To a cooled (0 °C) suspension of sodium hydride (60% dispersion in mineral oil, 1.20 g, 30.0 mmol, 2.0 equiv) in DMF (15 mL) was added dropwise a solution of indole (1.76 g, 15.0 mmol) in DMF (2 mL) and the mixture was allowed to stir at ambient temperature for 10 min. The mixture was cooled (0 °C), treated dropwise with the appropriate alkyl bromide (15.8 mmol, 1.05 equiv), and stirred at ambient temperature for 1 h. The mixture was cooled (0 °C), treated portionwise with trifluoroacetic anhydride (5.20 mL, 37.5 mmol, 2.5 equiv), and stirred at ambient temperature for 1 h. The solution was poured portionwise onto vigorously stirred ice water (900

mL) until precipitation was complete, and the formed red-pink solid was filtered and air-dried overnight.

To a refluxing solution of potassium hydroxide (2.78 g, 49.5 mmol, 3.3 equiv) in methanol (5 mL) was added portionwise a solution of the crude 1-alkyl-3-trifluoroacetyl-1H-indole in toluene (15 mL) and the solution was heated at reflux for 2 h. The solution was cooled to ambient temperature and partitioned between 1 M aq. NaOH (400 mL) and Et₂O (50 mL). The layers were separated, and the aqueous layer was adjusted to pH 1 with 10 M aq. HCl. The aqueous phase was extracted with Et₂O (3 × 100 mL), dried (Na₂SO₄), and solvent-evaporated under reduced pressure. The crude product was recrystallized from *i*-PrOH.

1-Pentyl-1H-indole-3-carboxylic Acid (23). Subjecting 1-bromopentane (1.95 mL, 15.8 mmol) to the general procedure above gave **23** as a colorless crystalline solid (1.92 g, 55% over two steps). ¹H NMR (400 MHz, CDCl₃): δ 8.24–8.27 (m, 1H), 7.93 (s, 1H), 7.38–7.40 (m, 1H), 7.29–7.33 (m, 2H), 4.17 (t, 2H, J = 7.2 Hz, NCH₂), 1.90 (quin, 2H, J = 7.2 Hz, CH₂), 1.32–1.39 (m, 4H, 2 × CH₂), 0.91 (t, 3H, J = 7.0 Hz, CH₃). All physical and spectral properties matched those reported previously.⁵⁸

1-(5-Fluoropentyl)-1H-indole-3-carboxylic Acid (24). Subjecting 1-bromo-5-fluoropentane (0.195 mL, 1.58 mmol) to the general procedure gave **23** as a colorless crystalline solid (0.249 g, 67% over two steps). ¹H NMR (400 MHz, CDCl₃): δ 8.21–8.29 (m, 1H, CH), 7.93 (s, 1H, CH), 7.36–7.42 (m, 1H, CH), 7.29–7.34 (m, 2H, CH), 4.43 (dt, ²J_{CF} = 47.2, ³J_{HH} = 5.9 Hz, 2H, CH₂F), 4.20 (t, J = 7.1 Hz, 2H, NCH₂), 1.96 (quin., J = 7.3 Hz, 2H, CH₂), 1.65–1.83 (m, 2H, CH₂), 1.43–1.53 (m, 2H, CH₂). All physical and spectral properties matched those reported previously.¹²

In Vitro CB₁ and CB₂ Binding Experiments. Receptor affinity was determined as previously described.¹¹ Membranes containing either triple-hemagglutinated, human CB₁ with preprolactin signal sequence, 3HA-hCB1⁶⁵ or CB₂ 3HA-hCB2,⁶⁶ were isolated from HEK293 cells (ATCC #CRL-1573, Manassas, VA) stably expressing each receptor (both in pEF4a). Cells were cultivated until semiconfluent in 175 cm² polystyrene culture flasks in Dulbecco's modified Eagle's medium supplemented with 10% fetal bovine serum (FBS, *v/v*, ThermoFisher Scientific, Waltham MA) and 250 μg/ml zeocin. Cells were suspended with 5 mM ice-cold EDTA in phosphate-buffered saline (PBS) and pelleted at 300g for 5 min. The supernatant was discarded, and the pellet was snap-frozen at –80 °C. To purify membranes, pellets were thawed on ice with tris–sucrose buffer (50 mM tris–HCl pH 7.4, 200 mM sucrose, 5 mM MgCl₂, 2.5 mM EDTA) and homogenized manually using a glass homogenizer. The homogenate was centrifuged at 1000g for 10 min at 4 °C. The supernatant was retained and further centrifuged at 27,000 rcf for 30 min at 4 °C. The pellet was suspended in a minimal volume of tris–sucrose buffer, aliquoted and stored at –80 °C. The protein concentration was quantified using a DC protein assay (Bio-Rad, Hercules, CA) adhering to the manufacturer's protocol.

CB₁- or CB₂-containing membranes (7.5 μg/point) were, respectively, incubated with 1 nM radioactively labeled [³H]-SR141716A (specific activity 55 Ci/mmol, PerkinElmer, Waltham MA) or [³H]CP 55,940 (specific activity 175 Ci/mmol, PerkinElmer) and either 10 μM (preliminary screen) or a range from 0.1 nM of 10 μM (for K_i determination) of the unlabeled test compound for 1 h at 30 °C. All components were diluted to a final assay volume of 200 μL in binding buffer (50 mM HEPES pH 7.4, 1 mM MgCl₂, 1 mM CaCl₂, 0.2% *w/v* bovine serum albumin (BSA, MP Biomedicals, Auckland, NZ)). Harvest plates (96-well GF/C, PerkinElmer) were simultaneously soaked for 1 h at room temperature in 0.1% (*w/v*) branched polyethyleneimine (Sigma-Aldrich) in H₂O. Harvest plates were applied to a vacuum manifold (Pall Corp., New York, NY) at 5–10 mmHg and washed with 200 μL wash buffer (50 mM HEPES pH 7.4, 500 mM NaCl, 0.1% (*w/v*) BSA). Test solutions were filtered through and thrice washed with 200 μL wash buffer. Harvest plates were dried overnight at room temperature. After sealing the base of the plate with a clear plate sealer, scintillation fluid (IRGASAFE Plus, PerkinElmer) was applied to each well (50 μL/well) and incubated in dark conditions for 30 min prior to detection using a Wallac TriLux MicroBeta2 scintillation counter for 2 min/well (PerkinElmer). Data were exported

and analyzed in GraphPad PRISM (version 8.0., GraphPad Software Inc. San Diego, CA) and expressed as receptor affinity (pK_i), and the maximum displacement of the reference compound (%) from at least three individual experiments was performed in duplicate. Receptor affinity was estimated using a “one-site fit K_i ” model, with radioligand K_d previously determined to be 1 nM (for both [^3H]SR141716A and [^3H]CP 55,940 for CB₁ and CB₂, respectively).

In Vitro CB₁ and CB₂ and Ca_v3.x Functional Activity Experiments. Transfection and Cell Culture. All Ca_v3.x experiments were performed using HEK293 FlpIn T-Rex cells stably transfected with pcDNA5 constructs encoding human Ca_v3.x cDNA.⁶³ Ca_v3.x expression was induced 24 h before membrane potential assays or electrophysiology experiments by adding 2 $\mu\text{g}/\text{mL}$ tetracycline. Human CB₁ or CB₂ was stably transfected into AtT-20 FlpIn and maintained using 80 $\mu\text{g}/\text{mL}$ hygromycin within the culture media. Cells were maintained and passaged at an 80% confluency in 75 cm² flasks and kept at 37 °C/5% CO₂ and used for up to 30 passages. Cells for calcium flux and membrane potential assays were grown in the same conditions but used at >90% confluence.

Functional Assays. Changes in membrane potential and calcium flux of cells were measured using the FLIPR membrane potential assay (MPA) or calcium 5 assay (Ca5) kits (Molecular Devices, Sunnyvale, CA), as previously described.^{53,67} Briefly, 24 h prior to the assay, AtT-20-CB₁/CB₂ or HEK293-Ca_v3.x cells were detached from their flasks using trypsin/EDTA (Sigma-Aldrich) and resuspended in 10 mL Leibovitz's L-15 media supplemented with 1% FBS, 100 μg penicillin and streptomycin per mL, and 15 mM glucose (and 2 $\mu\text{g}/\text{mL}$ tetracycline for HEK293-Ca_v3.x). Cells were plated in a volume of 90 μL per well in black-walled, clear-bottomed 96-well microplates (Corning, Castle Hill, Australia) and incubated overnight at 37 °C in ambient CO₂. MPA or Ca5 dye was reconstituted with the assay buffer (HBSS) containing the following compounds in mM: NaCl 145, HEPES 22, Na₂HPO₄ 0.338, NaHCO₃ 4.17, KH₂PO₄ 0.441, MgSO₄ 0.407, MgCl₂ 0.493, CaCl₂ 1.26, and glucose 5.56, pH 7.4, osmolarity 315 mOSM at half the manufacturer's recommended concentration. For Ca5 experiments, the HBSS was supplemented with probenecid (2.5 mM, Biotium, Scoresby, VIC, Australia) to minimize dye loss from the cells. Cells were loaded with 90 μL per well of the dye solution without removal of the L-15 and incubated at 37 °C for 60 min in ambient CO₂. Fluorescence was measured using a FlexStation 3 microplate reader (Molecular Devices, λ_{ex} 530 nm/ λ_{em} 565 nm for MPA or λ_{ex} 485 nm/ λ_{em} 525 nm for Ca5) using SoftMax Pro 7 (Molecular Devices). Baseline readings were taken every 2 s for at least 2 min, at which time the diluted drug was added in a volume of 20 μL . For the MPA assay of CB₁/CB₂ activity, changes in fluorescence elicited by the addition of drug were expressed as a percentage of baseline fluorescence after subtraction of the changes produced by vehicle addition, normalized to the maximal effective response elicited from 1 μM CP 55,940 within each column, as previously described.⁶⁷ For measuring the modulation of Ca_v3.x activity, CaCl₂ was added to cells (final concentration 10 mM) 5 min after the drug and the changes in Ca5 fluorescence were expressed as a change relative to 10 mM Ca alone (area under the curve). The final concentration of the vehicle (dimethyl sulfoxide (DMSO)) was not more than 0.1% in both assays, and Ca²⁺ and K⁺ concentrations in the CaV assay buffer solution were approximately 10 and 2.5 mM, respectively. Data from both assays were analyzed with Graphpad PRISM and are expressed as the mean \pm SEM of at least five independent determinations performed in duplicate unless otherwise stated.

Electrophysiology. Whole-cell voltage-clamp recordings from HEK293-Ca_v3.x cells were performed at room temperature. At least 24 h prior to experiments, cells were detached from flasks using trypsin/EDTA and plated into 10 cm sterile tissue culture dishes containing 10 mL of supplemented DMEM and 10–15 glass coverslips (12 mm diameter, ProScitech, QLD, Australia). Culture dishes were then kept overnight in the same conditions as flasks to allow cells to adhere to coverslips. They were then transferred to a 30 °C/5% CO₂ incubator to inhibit cell proliferation until ready to be used for electrophysiology experiments.

Recording Solutions. External recording solutions contained (in mM) 114 CsCl, 5 BaCl₂, 1 MgCl₂, 10 HEPES, 10 glucose, adjusted to pH 7.4 with CsOH. The internal patch pipette solution contained (in mM) 126.5 CsMeSO₄, 2 MgCl₂, 11 EGTA, and 10 HEPES adjusted to pH 7.3 with CsOH. The internal solution was supplemented with 0.6 mM GTP and 2 mM ATP and mixed thoroughly just prior to use. Liquid junction potentials for the above solutions were calculated prior to experiments using pClamp 10 software and corrected for during experiments. Compounds were prepared daily from 10 mM DMSO stocks and diluted into the external solution just prior to use. Compounds were then applied rapidly and locally to the cells using a custom-built gravity-driven microperfusion system.⁶⁸ Initial vehicle experiments were performed to ensure that 0.1% DMSO had no effect on current amplitudes or channel kinetics (data not shown), and all subsequent experiments contained 0.1% DMSO in control external solutions. Currents were elicited from a holding potential of -100 mV and were measured by conventional whole-cell patch-clamp techniques using an Axopatch 200B amplifier in combination with Clampex 9.2 software (Molecular Devices, Sunnyvale). After establishing whole-cell configuration, the cellular capacitance was minimized using the amplifier's built-in analogue compensation. Series resistance was kept to <10 M Ω and was compensated to at least 85% in all experiments. All data were digitized at 10 kHz with a Digidata 1320 interface (Molecular Devices) and filtered at 1 kHz (8-pole Bessel filter). Raw and online leak-subtracted data were both collected simultaneously, and P/N4 leak subtraction was performed using opposite polarity and after the protocol sweep. For tonic inhibition of T-type current, membrane potential was stepped from -100 to -30 mV for 200 ms and then allowed to recover for 12 s (one sweep). A minimum of 10 sweeps were collected under control external perfusion to allow for control peak current to equilibrate. The drug was then continuously perfused, and sweeps were recorded until no further inhibition is seen (minimum of three sweeps with the same amplitude). All electrophysiology data were acquired and analyzed using Clampfit 9.2 (Molecular Devices) and are expressed as a percentage of means \pm standard errors of at least six experiments per compound on each calcium channel.

In Vivo Radiotelemetry. Radiotelemetric assessment of MEPIRAPIM and 5F-BEPIRAPIM was carried out according to previously reported procedures.^{11,54,69} Briefly, TA-F10 radiotelemetry probes (Data Sciences International, St. Paul) were intraperitoneally implanted according to the manufacturer's instructions into male C57BL/6J mice. The mice were aged 8 weeks at the time of surgery, weighed between 20 and 25 g, and were allowed 10 days of recovery time before data collection. All work involving animals was carried out in accordance with the Australian Code of Practice for the Care and Use of Animals for Scientific Purposes and approved by The University of Sydney Animal Ethics Committee.

Four mice were used per drug, and the first received a “baseline” vehicle injection of a 5:5:90 mixture of ethanol, polysorbate 80, and saline, respectively. Mice then received an ascending dose sequence of 0.1, 0.3, 1, 3, 10, and 30 mg/kg, with two drug-free washout days between each dose to minimize the development of drug tolerance. The drugs were administered via intraperitoneal injection with an injection volume of 10 $\mu\text{L}/\text{g}$.

Raw body temperature data were gathered continuously at 1000 Hz and averaged into 15 min bins using Dataquest A.R.T. software (version 4.33, Data Sciences International). Using Prism (version 8.4.3), baseline (vehicle injection) data for each animal were subtracted to yield the change in body temperature data. Area under baseline curves (AUCs) for the first 2.5 h were calculated for each drug dose using R (version 4.0.5) as a measure of the total drug effect. For the 30 mg/kg MEPIRAPIM drug dose, the mean AUC was compared to zero (i.e., no change from baseline) using a one-sample *t*-test.

Molecular Docking Simulations. Protein Preparation. The cryo-EM structures of the CB₁ receptor (PDB: 6N4B)⁷⁰ and CB₂ receptor (PDB: 6TP0)⁷¹ were retrieved from RCSB PDB (<https://www.rcsb.org/>). The structure was prepared using Protein Preparation Wizard.⁷² The G proteins and cholesterol molecule were removed, leaving only the receptor and ligand in the active site. The preparation process involved assigning bond orders, adding hydrogens, creating zero-order

bonds to metals, generating disulfide bonds, filling in missing side chains and loops using Prime, generating het states using Epik⁷³ at pH 7.0 ± 2.0, and deleting water molecules beyond 5 Å from het groups. The hydrogen bond network was optimized, and water orientations were sampled. The pK_a values of the protein were predicted using PROPKA,⁷⁴ and the target pH value was set at 7.0. Lastly, the protein structure was minimized using the OPLS_2005 force field⁷⁵ where the RMSD of the atom displacement for terminating the minimization was set as 0.3 Å.

Ligand Preparation. Ligands were first prepared using LigPrep⁷⁶ to generate energy-minimized three-dimensional (3D) structures. OPLS3e force field was used for minimization. Epik was used to generate all possible ionized states at pH 7.0 ± 2.0. The desalt setting was used to remove any counterions or water molecules. Tautomer and stereoisomers were generated (at most 32 per ligand), where specified chiralities were retained.

Induced Fit Docking. To generate docking poses, prepared ligands were docked against the prepared CB1 and CB2 structures using induced fit docking.⁷⁷ Extended sampling protocol and OPLS3e force field were used. The ligand in the binding site was used to define the receptor. The core constraint was applied to restrict docking to the ligand with a tolerance of 1.0 Å; core atoms were determined based on the maximum common structure. Ring conformation sampling was performed with an energy window set as 2.5 kcal/mol; nonplanar conformation was penalized for amide bonds. Residues within 5.0 Å of ligand poses were refined using Prime.

■ ASSOCIATED CONTENT

SI Supporting Information

The Supporting Information is available free of charge at <https://pubs.acs.org/doi/10.1021/acschemneuro.1c00822>.

¹H and ¹³C NMR spectra for all novel compounds and representative fluorescence inhibition and electrophysiology traces for selected compounds at T-type calcium channels (PDF)

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A.M., J.L.L., and W.J. synthesized and analytically characterized all compounds under the supervision of S.D.B. J.M. performed the *in vitro* binding assays under the supervision of M.G. R.B. performed the *in vitro* CB₁ and CB₂ receptor membrane potential assays, and S.M. performed the Ca_v calcium flux and electrophysiology assays, under the supervision of M.C. and C.B. R.C.K. performed radiotelemetry experiments under the supervision of I.S.M. R.E. conducted high-resolution mass spectrometry under the supervision of R.R.G. F.L. conducted molecular docking simulations under the supervision of D.E.H. R.C.K., M.G., M.C., C.B., and S.D.B. conceived the experiments, and R.C.K., E.A.C., A.A., C.B., G.W.Z., and S.D.B. wrote the manuscript. All authors reviewed and edited drafts of the manuscript and approved the final version.

Funding

R.C.K., R.B., S.M., E.A.C., A.A., J.L.L., I.S.M., and S.D.B. gratefully acknowledge support from The Lambert Initiative for

Cannabinoid Therapeutics, a philanthropic research program based at the Brain and Mind Centre, The University of Sydney. Research by R.C.K., E.A.C., A.A., J.L.L., and S.D.B. was partially supported by the National Health and Medical Research Council Project Grant 1161571 and a Brain and Mind Centre Research Development Grant from The University of Sydney. W.J. is supported by a National Health and Medical Research Council Peter Doherty Early Career Research Fellow. Research by J.M. and M.G. was supported by a contract from the Health Research Council of New Zealand.

Notes

The authors declare no competing financial interest. All work involving animals was carried out in accordance with the Australian Code of Practice for the Care and Use of Animals for Scientific Purposes and approved by The University of Sydney Animal Ethics Committee.

ABBREVIATIONS

ANOVA, analysis of variance; CB₁, cannabinoid receptor 1; CB₂, cannabinoid receptor 2; EDC, 1-ethyl-3-(3-dimethylaminopropyl)carbodiimide; EMCDDA, European Centre for Drugs and Drug Addiction; FBS, fetal bovine serum; FLIPR, fluorometric imaging plate reader; GTPγS, guanosine 5'-O-[gamma-thio]triphosphate; HEK, human embryonic kidney; HOBt, 1-hydroxybenzotriazole; NMR, nuclear magnetic resonance; NPS, new psychoactive substances; QTOF-MS, quadrupole time-of-flight mass spectrometry; SAR, structure–activity relationship; SCRA, synthetic cannabinoid receptor agonist; SEM, standard error of mean; THC, Δ⁹-tetrahydrocannabinol

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