ORIGINAL ARTICLE

Two new-type cannabimimetic quinolinyl carboxylates, QUPIC and QUCHIC, two new cannabimimetic carboxamide derivatives, ADB-FUBINACA and ADBICA, and five synthetic cannabinoids detected with a thiophene derivative α -PVT and an opioid receptor agonist AH-7921 identified in illegal products

Nahoko Uchiyama · Satoru Matsuda · Maiko Kawamura · Ruri Kikura-Hanajiri · Yukihiro Goda

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Abstract We identified two new-type cannabimimetic quinolinyl carboxylates, quinolin-8-yl 1-pentyl-(1H-indole)-3-carboxylate (QUPIC, 1) and quinolin-8-yl 1-(cyclohexylmethyl)-1*H*-indole-3-carboxylate (QUCHIC, 2); and two new cannabimimetic carboxamide derivatives, N-(1-amino-3,3dimethyl-1-oxobutan-2-yl)-1-(4-fluorobenzyl)-1H-indazole-3-carboxamide (ADB-FUBINACA, 3) and N-(1-amino-3,3dimethyl-1-oxobutan-2-yl)-1-pentyl-1H-indole-3-carboxamide (ADBICA, 4), as designer drugs in illegal products. Compound 3 was reported to have a potent affinity for cannabinoid CB₁ receptor by Pfizer in 2009, but this is the first report of its detection in illegal products. No chemical or pharmacological data for compounds 1, 2, and 4 have appeared until now, making this the first report on these compounds. We also detected synthetic cannabinoids, APICA N-(5-fluoropentyl) analog (5), APINACA N-(5-fluoropentyl) analog (6), UR-144 N-(5-chloropentyl) analog (7), JWH-122 N-(5-chloropentyl) analog (8), and AM-2201 4-methoxynaphthyl analog (4-MeO-AM-2201, 9) herein as newly distributed designer drugs in Japan. It is of interest that compounds 1 and 2 were detected with their synthetic component, 8-quinolinol (10). A stimulant thiophene analog, α -pyrrolidinovalerothiophenone (α-PVT, 11), and an opioid receptor agonist, 3,4-dichloro-N-([1-(dimethylamino)cyclohexyl]methyl)benzamide (AH-7921,

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N. Uchiyama · S. Matsuda · M. Kawamura · R. Kikura-Hanajiri · Y. Goda (⊠) National Institute of Health Sciences, 1-18-1 Kamiyoga, Setagaya-ku, Tokyo 158-8501, Japan e-mail: goda@nihs.go.jp **12**), were also detected as new types of designer drugs coexisting with several synthetic cannabinoids and cathinone derivatives in illegal products.

Keywords Quinolin-8-yl 1-pentyl-(1*H*-indole)-3carboxylate (QUPIC) · Quinolin-8-yl 1-(cyclohexylmethyl)-1*H*-indole-3-carboxylate (QUCHIC) · N-(1-Amino-3,3-dimethyl-1-oxobutan-2-yl)-1-(4fluorobenzyl)-1*H*-indazole-3-carboxamide (ADB-FUBINACA) · N-(1-Amino-3,3-dimethyl-1-oxobutan-2yl)-1-pentyl-1*H*-indole-3-carboxamide (ADBICA) · Synthetic cannabinoids · α -Pyrrolidinovalerothiophenone (α -PVT)

Introduction

The number of new psychotropic substances-not only synthetic cannabinoids but also other types of substances such as cathinone derivatives-has been increasing in Japan and in European countries year by year [1-10]. To prevent the abuse of these drugs, a total of 106 substances, including 35 synthetic cannabinoids, 17 cathinone derivatives, 26 phenethylamines, 13 tryptamines, 4 piperazines, 10 others, and 1 plant extract, were controlled as designated substances (Shitei-Yakubutsu) under the Pharmaceutical Affairs Law in Japan as of January 2013. Moreover, among them, 3 of the phenethylamines (2C-I, 2C-T-2, and 2C-T-4) have been strictly regulated as narcotic substances in Japan since January 2008. In August 2012, 2 synthetic cannabinoids (cannabicyclohexanol and JWH-018) and 2 cathinone derivatives (MDPV and mephedrone) were also classified as new narcotic substances.

We have been conducting an ongoing survey of designer drugs in the Japanese illegal market, and our



Fig. 1 Structures of newly detected (1-12, a), detected but known (b), and related compounds (c)



Fig. 2 Liquid chromatography–mass spectrometry (LC–MS) analysis of product A. Liquid chromatography–ultraviolet-photodiode array (LC–UV-PDA) chromatogram (**a**), total ion chromatogram (TIC) (**b**), and mass chromatogram at m/z 146 (**c**) using elution program (2).

survey has revealed nine synthetic cannabinoids (1-9), a substance (10) that is a synthetic component of 1 or 2, and two other psychotropic substances (11 and 12) newly detected in the present study (Fig. 1a). In this article, we describe our identification of these newly detected compounds in detail.

Ultraviolet (UV) and electrospray ionization (ESI) mass spectra of peaks 1 (d), 2 (e), 10 (f), 11 (h), α -PBP (i), pentedrone (j), and authentic 8-quinolinol (g) obtained by LC–MS

Materials and methods

Samples for analysis

The analyzed samples were purchased on the Internet between July 2012 and January 2013 as chemical-type or



Fig. 3 Gas chromatography-mass spectrometry (GC-MS) analysis of product A. TIC (a) and electron ionization (EI) mass spectra of peaks 1 (b), 2 (c), 10 (d), 11 (f), α -PBP (g), pentedrone (h), and authentic 8-quinolinol (e)

herbal-type products being sold in Japan. Each of the herbal-type products (A–J) contained about 3 g of mixed dried plants. The chemical product K, which was called "Fragrance Powder," consisted of about 400 mg of white powder.

Chemicals and reagents

Authentic APICA *N*-(5-fluoropentyl) analog (**5**), APINACA *N*-(5-fluoropentyl) analog (**6**), UR-144 *N*-(5-chloropentyl) analog (**7**), α -PBP, pentedrone, α -PVP, XLR11 (5FUR-144), MAM-2201, AH-7921 (**12**), and EAM-2201 were purchased from Cayman Chemical (Ann Arbor, MI, USA). 8-Quinolinol (**10**) was purchased from Tokyo Chemical Industry (Tokyo, Japan). Other compounds (**1**–**4**, **8**, and **9**) were isolated from herbal or chemical products. All other common chemicals and solvents were of analytical reagent grade or HPLC grade. As solvents for nuclear magnetic resonance (NMR) spectroscopy, CD₃OD (99.96 %), CD₃OH (99.8 %), CDCl₃ (99.96 %), benzene- d_6 (99.96 %), and dimethyl sulfoxide (DMSO)- d_6 (99.96 %) were purchased from the ISOTEC division of Sigma-Aldrich (St. Louis, MO, USA).

Preparation of sample solution

For qualitative analyses, 10 mg of each herbal-type product was crushed to a powder and extracted with 1 ml of methanol under ultrasonication for 10 min. A 2-mg portion of each powder-type product was extracted with 1 ml of methanol under ultrasonication for 10 min. After centrifugation (5 min, 3000 rpm) of each extract, the supernatant solution was passed through a centrifugal filter (Ultrafree-MC, 0.45 μ m filter unit; Millipore, Bedford, MA, USA) to serve as sample solution for analysis. If necessary, the solution was diluted with methanol to a suitable concentration before instrumental analysis.

Table 1 Nuclear magnetic resonance (NMR) data for QUPIC (1) and QUCHIC (2)

No.	QUPIC (1)		QUCHIC (2)		
	¹³ C	¹ H	¹³ C	¹ H	
1	165.2	_	165.2	_	
2'	137.7	8.33, 1H, s	138.2	8.28, 1H, s	
3'	106.3	_	106.2	_	
3′a	128.6	_	128.5	_	
4′	122.4	8.13, 1H, brd, $J = 7.9$ Hz	122.3	8.13, 1H, d, $J = 7.9$ Hz	
5'	123.1	7.24, 1H, ddd, $J = 7.9$, 6.9, 1.0 Hz	123.0	7.23, 1H, td, $J = 7.9$, 1.0 Hz	
6′	124.0	7.31, 1H, ddd, $J = 7.9$, 6.9, 1.0 Hz	124.0	7.30, 1H, td, $J = 7.9$, 1.0 Hz	
7′	111.7	7.57, 1H, brd, $J = 7.9$ Hz, overlapped	111.9	7.55, 1H, d, $J = 7.9$ Hz, overlapped	
7′a	138.3	_	138.6	_	
1″	48.0	4.33, 2H, t, <i>J</i> = 7.2 Hz	54.2	4.15, 2H, d, <i>J</i> = 7.2 Hz	
2"	30.9	1.95, 2H, q, $J = 7.2$ Hz	39.9	1.97, 1H, m	
3″	30.1	1.37, 2H, m, overlapped	_	_	
4″	23.4	1.40, 2H, m, overlapped	_	_	
5″	14.3	0.92, 3H, t, $J = 7.2$ Hz	27.4	1.69, 2H, m, overlapped	
3″/7″	_	_	31.9	1.66, 2H, m, overlapped	
				1.09, 2H, m	
4″/6″	_	_	26.8	1.28, 2H, m, overlapped	
				1.75, 2H, m	
1'''	_	_	_	_	
2'''	151.4	8.83, 1H, dd, J = 4.1, 1.7 Hz	151.4	8.83, 1H, dd, $J = 4.2$, 1.7 Hz	
3'''	123.1	7.56, 1H, m, overlapped	123.1	7.52, 1H, dd, $J = 8.3$, 4.2 Hz	
4'''	138.1	8.41, 1H, dd, J = 8.3, 1.4 Hz	138.1	8.41, 1H, dd, $J = 8.3$, 1.7 Hz	
4′′′a	131.2	_	131.2	_	
5'''	127.0	7.89, 1H, dd, $J = 7.6$, 1.4 Hz	127.0	7.89, 1H, brdd, $J = 7.6$, 1.4 Hz	
6'''	127.8	7.67, 1H, t, $J = 7.6$ Hz	127.8	7.66, 1H, t, $J = 7.6$ Hz	
7'''	123.7	7.63, 1H, dd, $J = 7.6$, 1.4 Hz	123.7	7.63, 1H, dd, $J = 7.6$, 1.4 Hz	
8'''	148.5	-	148.5	_	
8′′′a	142.8	-	142.8	_	

Recorded in CD₃OD at 600 MHz (¹H) or 150 MHz (¹³C); data in δ ppm



Fig. 4 Double quantum filtered correlation spectroscopy (DQF-COSY), selected heteronuclear multiple-bond correlation (HMBC), and selected rotating frame nuclear Overhauser effect (ROE)

Analytical conditions

Each sample solution was analyzed by ultra-performance liquid chromatography-electrospray ionization-mass spectrometry (UPLC-ESI-MS) and gas chromatographymass spectrometry (GC-MS) in the electron ionization (EI) mode according to our previous report [11]. Two elution programs were used in the LC-MS analysis. Programs (1) and (2) were used for synthetic cannabinoids and for the other compounds including cathinone derivatives, respectively [11]. The obtained GC mass spectra were compared to those of an EI-MS library [Mass Spectra of Designer Drugs 2012 (Wiley-VCH, Weinheim, Germany)]. In addition, our in-house EI-MS library of designer drugs obtained by our continuous survey of illegal products and commercially available reagents was also used for structural elucidation.

The accurate mass numbers of the target compounds were measured by liquid chromatography-quadrupole

correlations (a) and $^1H\!-\!^{15}\!N$ HMBC (b) for compound 1 (QUPIC), and DQF-COSY, selected HMBC, and selected ROE correlations for compound 2 (QUCHIC, c) and compound 11 ($\alpha\text{-PVT}, d)$

time-of-flight-mass spectrometry (LC-QTOF-MS) system consisting of an Acquity UPLC and Xevo QTOFMS (Waters, Milford, MA, USA) with a photodiode array (PDA) detector (Waters). The sample solutions were separated with an ACQUITY UPLC HSS C18 column (150 mm \times 2.1 mm i.d., particle size 1.8 µm; Waters) at 50 °C. Each analysis was carried out with a binary mobile phase consisting of solvent A (10 mM ammonium formate in water, pH 3.0) and solvent B (0.1 % formic acid in acetonitrile). The elution program was: 87 % A/13 % B (0.5-min hold) to 50 % A/50 % B (0.5-10 min), and up to 10 % A/90 % B (10-15 min, 5-min hold) at a flow rate of 0.4 ml/min. The injection volume was 1 µl, and the wavelength of the PDA detector for screening was set from 210 to 400 nm. The MS conditions were: ion source, ESI in the positive mode; ion source temperature, 120 °C; desolvation gas, nitrogen at a flow rate of 800 l/h at 400 °C; capillary and cone voltages, 3000 and 30 V, respectively; collision energy, 27 V; mass spectral range,



Fig. 5 LC–MS analysis of hydrolysates of QUPIC (1). LC–UV-PDA chromatogram (a), TIC (b), and mass chromatograms at m/z 359 (c), 232 (d), and 146 (e) of the reaction mixture of QUPIC (1) after acid hydrolysis using elution program (2). LC–UV-PDA chromatograms

m/z 50–1000. Leucine enkephalin [m/z 278.1141 and 508.20783 ([M+H]⁺)] was used as a substance for lock mass ions during the measurements.

The NMR spectra were obtained on ECA-800 and 600 spectrometers (JEOL, Tokyo, Japan). Assignments were made via ¹H NMR, ¹³C NMR, heteronuclear multiple quantum coherence (HMQC), heteronuclear multiple-bond correlation (HMBC), ¹⁵N HMBC, double quantum filtered correlation spectroscopy (DQF-COSY), and rotating frame nuclear Overhauser effect (ROE) spectra.

Isolation of compound 1

A 3-g sample of mixed dried plants (product I) was extracted with 250 ml of chloroform by ultrasonication for 30 min. The extractions were repeated three times, and the supernatant fractions were combined and evaporated to dryness. The extract was placed on a preparative silica gel thin-layer chromatography (TLC) plate (Silica Gel 60,

of authentic QUPIC (1, f) and authentic 8-quinolinol (g). UV and ESI mass spectra of peaks 1 (h), hydrolysate-1 (i), and 10 (j). The putative hydrolysis mechanism of QUPIC (1) in acidic conditions (k)

 20×20 cm, 2 mm; Merck, Darmstadt, Germany), which was then developed using hexane/ethyl acetate (3:1). A portion of the silica gel containing a target compound in the TLC plate was detected under ultraviolet (UV) light (254 nm). It was then scraped from the plate and eluted with chloroform to give fraction 1, which was further purified by repeated preparative TLC with toluene/chloroform (1:1) and then recrystallized by methanol. Finally, compound **1** (130 mg) was obtained as a brown oil.

Isolation of compound 2

A 3-g sample of mixed dried plants (product A) was extracted with 250 ml of chloroform by ultrasonication for 30 min. The extractions were repeated three times, and the supernatant fractions were combined and evaporated to dryness. Separation of the extract by repeated preparative TLC [hexane/acetone (2:1) and toluene/ethyl acetate (10:1)] and silica gel column chromatography [toluene/

ethyl acetate (10:1, 9:1, 8:2, 7:3, 6:4)] gave compound **2** (13 mg) as a yellow oil.

Isolation of compounds 3 and 4

Each 3-g sample of mixed dried plants (products J and B) was extracted with 250 ml of chloroform by ultrasonication for 30 min. The extractions were repeated three times, and the supernatant fractions were combined and evaporated to dryness. Each extract was purified by preparative TLC [hexane/ethyl acetate (1:2)] to obtain compound **3** (65 mg) as a yellow solid and compound **4** (81 mg) as a pale yellow solid.

Isolation of compound 8

A 3-g sample of mixed dried plants (product F) was extracted by the same method as described above. The final separation of the extract by silica gel column chromatography [toluene/chloroform (5:5, 6:4)] resulted in the isolation of compound **8** (10 mg) as a yellow solid.

Isolation of compound 9

A 3-g sample of mixed dried plants (product G) was extracted by the same method described above. Separation of the extract by preparative TLC [hexane/ethyl acetate (3:1)] and recrystallization in methanol gave compound **9** (145 mg) as a pale yellow solid.

Hydrolysis of compounds 1 and 2

A 2-mg sample of each compound was dissolved in 20 % HCl aqueous solution and heated at 60 °C for 30 min. The reaction mixture was evaporated under a nitrogen stream, and the residue was redissolved in methanol. The solution was then analyzed by LC–MS and LC–QTOF–MS.

Results and discussion

Identification of unknown peaks 1, 2, 10, and 11

Four unknown peaks, **1**, **2**, **10**, and **11**, were detected along with known cathinone derivatives, α -PBP, and pentedrone in the LC–MS and GC–MS chromatograms for product A, as shown in Figs. 2a, b, 3a. In the LC–MS chromatograms using elution program (2) for the analysis of cathinones and others [11], two unknown peaks (1 and 2) at 31.1 and 32.0 min showed protonated molecular ion [M+H]⁺ signals at *m*/*z* 359 and 385, respectively (Fig. 2d, e). The UV spectra of both compounds showed the same absorbance maxima at 231 and 294 nm (Fig. 2d, e).

In addition, peaks 1 and 2 were detected at 10.2 and 12.2 min under elution program (1) for the analysis of cannabinoids (data not shown) [11]. The total ion chromatogram (TIC) by GC-MS showed peak 1 at 52.52 min and peak 2 at 56.57 min (Fig. 3a), which indicated putative molecular ion signals at m/z 358 and 384, respectively (Fig. 3b, c). Unknown peak 10 was presumed to be 8-quinolinol, based on the fragment patterns of the GC-MS analysis (Fig. 3d) and LC-MS analysis (Fig. 2c, f). Peak 10 was confirmed to be identical to 8-quinolinol by direct comparison of the data to those of the authentic compound (Figs. 2g, 3e). After isolation of compounds 1 and 2, their accurate mass spectra were measured by LC-QTOF-MS in the positive mode. The ion peaks observed at m/z 359.1764 and 385.1908 suggested that the protonated molecular formulae of compounds 1 and 2 were $C_{23}H_{23}N_2O_2$ (calcd. 359.1760) and C₂₅H₂₅N₂O₂ (calcd. 385.1916), respectively.

The structure of compound **1** was elucidated by NMR analysis (Table 1; Fig. 4a, b). The ¹H and ¹³C NMR spectra of compound **1** suggested the existence of 22 protons and 23 carbons as shown in Table 1. The analyses by DQF-COSY, HMQC, HMBC, and one-dimensional (1D) ROE spectra for compound **1** revealed the presence of an *N*-(1-pentyl)-1*H*-indole-3-carbonyl moiety (Fig. 4a). In addition, the NMR spectra of the remaining C_9H_6NO unit

Table 2 NMR data for α -PVT (11)

No.	¹³ C	¹ H
1	188.7	_
2	63.1	4.89, 1H, t, $J = 4.8$ Hz
3	32.9	2.31, 1H, m
		2.03, 1H, m, overlapped
4	19.8	1.47, 1H, m
		1.30, 1H, m
5	13.9	0.93, 3H, t, $J = 7.2$ Hz
1′	-	_
2′	143.4	_
3′	134.5	7.88, 1H, d, $J = 4.8$ Hz
4′	129.4	7.23, 1H, t, $J = 4.8$ Hz
5′	137.7	7.85, 1H, d, $J = 4.8$ Hz
2″	52.9	3.77, 1H, m
		2.82, 1H, m
3″	23.9	2.15, 1H, m, overlapped
		1.98, 1H, m, overlapped
4″	23.6	2.18, 1H, m, overlapped
		2.08, 1H, m, overlapped
5″	48.6	3.89, 1H, m
		3.63, 1H, m

Recorded in CDCl₃ at 600 MHz (¹H) or 150 MHz (¹³C); data in δ ppm



Fig. 6 LC–MS and GC–MS analyses of product B. LC–UV-PDA chromatogram (a) and TIC (b) using elution program (1) obtained by LC–MS. UV and ESI mass spectra of peaks 3 (c) and 4 (d). TIC (e) and EI mass spectra of peaks 3 (f) and 4 (g) obtained by GC–MS

suggested the presence of a quinolinol group, and the observed ¹⁵N HMBC correlations (Fig. 4b) and the fragment ions at m/z 214 and 144 of peak **1** revealed by the GC–MS analysis (Fig. 3b) supported the existence of *N*-(1-pentyl)-1*H*-indole-3-carbonyl and quinolinol moieties.

On the basis of the two-dimensional (2D) NMR correlations and the quaternary carbon signals at $\delta_{\rm C}$ 165.2 (C-1) and $\delta_{\rm C}$ 148.5 (C-8'''), we concluded that the *N*-(1-pentyl)-1*H*-indole moiety was attached to the 8-quinolinol moiety by an ester linkage at position-1. However, the ester bond between the *N*-(1-pentyl)-1*H*-indole-3-carbonyl and quinolinol moieties was not clear. To determine the structure of compound **1**, we hydrolyzed it under acidic conditions, and then analyzed the reaction mixture by LC–MS (Fig. 5). The peak at 3.0 min was identified as 8-quinolinol (**10**) by direct comparison of its spectral data with those of the purchased compound (Fig. 5e, g, j). The other peak at 27.6 min showed the ion signal at m/z 232 (Fig. 5d, i). Its LC–QTOF–MS analysis showing the ion signal at 232.1353 suggested the protonated molecular formula of C₁₄H₁₈NO₂ (calcd. 232.1338) for the expected hydrolysate, *N*-(1-pentyl)-1*H*-indole-3-carboxylic acid (Fig. 5k). Thus, the structure of compound **1** was determined as quinolin-8-yl 1-pentyl-(1*H*-indole)-3-carboxylate and was named QUPIC (**1**).

Given the estimated protonated molecular formulae of compounds **2** and **1** of $C_{25}H_{25}N_2O_2$ and $C_{23}H_{23}N_2O_2$, respectively, the elemental difference between them was C_2H_2 . The ¹H and ¹³C NMR spectra of compound **2** were very similar to those of compound **1** except for the

Table 3 NMR data for ADB-PINACA (3) and ADBICA (4)

No.	AB-FUBINACA ^{a,c}	ADB-PINACA (3) ^a		ADBICA (4) ^b	
	¹³ C	¹³ C	¹ H	¹³ C	¹ H
1	161.2	160.6	-	165.0	_
2'	_	_	_	131.5	7.70, 1H, s
3'	137.1	136.9	_	110.3	-
3′a	122.3	122.2	_	125.4	-
4′	121.8	121.8	8.17, 1H, d, J = 7.9 Hz	120.2	7.99, 1H, m
5'	122.8	122.8	7.29, 1H, t, $J = 7.9$ Hz	121.6	7.24, 1H, m, overlapped
6'	127.0	127.1	7.46, 1H, brt, $J = 7.9$ Hz	122.5	7.26, 1H, m, overlapped
7′	110.6	110.7	7.79, 1H, d, $J = 7.9$ Hz	110.3	7.36, 1H, m
7′a	140.6	140.6	_	136.6	-
1″	51.6	51.6	5.78, 2H, s	46.9	4.10, 2H, t, $J = 7.2$ Hz
2"	133.0, <i>J</i> = 2.9 Hz	133.0, <i>J</i> = 2.9 Hz	_	29.7	1.84, 2H, q, J = 7.2 Hz
3″/7″	129.5, $J = 8.7$ Hz	129.5, $J = 8.7$ Hz	7.31, 2H, dd, $J = 7.2$, 1.3 Hz, overlapped	-	_
4″/6″	115.5, $J = 21.7$ Hz	115.6, $J = 21.7$ Hz	7.16, 1H, brd, $J = 8.6$ Hz, 7.15, 1H, brd, $J = 8.6$ Hz	_	_
3″	-	_	_	29.0	1.29, 2H, m, overlapped
4″	-	_	_	22.3	1.32, 2H, m, overlapped
5″	161.6, d, <i>J</i> = 242.8 Hz	160.9, d, J = 244.2 Hz	_	13.9	0.87, 3H, t, J = 6.9 Hz
1'''	172.6	171.7	_	173.2	-
2'''	56.9	58.7	4.45, 1H, d, $J = 9.6$ Hz	59.7	4.68, 1H, d, J = 9.3 Hz
3'''	31.2	34.6	_	34.7	-
4'''	19.4	_	_	-	-
5'''	18.1	_	_	-	-
4'''/5'''/6'''	_	26.6	0.98, 9H, s	26.8	1.13, 9H, s
1-CONH	_	_	7.60, 1H, d, $J = 9.6$ Hz	-	6.7, 1H, d, <i>J</i> = 8.9 Hz
1'''-CONH _{2a}	_	_	7.27, 1H, brs, overlapped	-	6.41, 1H, brs
1 ^{'''} -CON <u>H_{2b}</u>	-	-	7.72, 1H, brs	-	5.63, 1H, brs

Recorded at 600 MHz (¹H) or 150 MHz (¹³C); data in δ ppm

^a Recorded in dimethyl sulfoxide (DMSO)-d₆

^b Recorded in CDCl₃

^c From Uchiyama et al. [15]





Fig. 7 DQF-COSY, selected HMBC, and selected ROE correlations (a) and $^{1}H^{-15}N$ HMBC correlations (b) for compound 3 (ADB-FUBINACA); DQF-COSY, selected HMBC, and selected ROE

N-cycloalkyl moiety (position-1" to 7") as shown in Table 1. The observed DQF-COSY, HMQC, HMBC, 1D ROE correlations and the quarternary carbon signals at $\delta_{\rm C}$ 165.2 (C-1) and $\delta_{\rm C}$ 148.5 (C-8"") for compound **2** suggested the presence of the *N*-(1-cyclohexylmethyl)-1*H*-indole-3-carbonyl and quinolinol moieties, and an ester linkage between these two moieties at position-1, similar to compound **1** (Fig. 4c; Table 1). Compound **2** was similarly hydrolyzed under acidic conditions to determine the structure (Supplementary material, Fig. S1). The structure of compound **2** was deduced to be quinolin-8-yl 1-(cyclohexylmethyl)-1*H*-indole-3-carboxylate and was named QUCHIC (**2**).

The chemical and pharmacological data on compounds **1** and **2** have not been reported, although the quinoline derivatives have been synthesized as cannabinoid receptor ligands [12]. Even though compounds **1** and **2** are being sold under the names "PB-22" and "BB-22," respectively, on illegal drug markets on the Internet, we named these compounds QUPIC and QUCHIC, respectively, taking into

correlations for compound **4** (ADBICA, **d**), and deuterium-induced isotope shifts of NH protons for the ¹³C NMR signals of compound **3** in CD_3OD (**c**)

account the IUPAC (International Union of Pure and Applied Chemistry) naming system and regulatory action.

Unknown peak 11 was detected together with two other peaks by LC-MS and GC-MS analyses of the same product A (Figs. 2h-j, 3f-h). The latter two peaks were readily found to be identical to α -PBP and pentedrone by direct comparison of the data with those of the purchased authentic compounds (data not shown). Both compounds were detected in European countries in 2011 [9, 13]. The LC-MS chromatogram demonstrated that unknown peak **11** at 11.7 min showed a protonated ion signal $([M+H]^+)$ at m/z 238 and absorbance maxima at 270 and 299 nm in the UV spectrum (Fig. 2h). The accurate mass spectrum was measured by LC-QTOF-MS in the positive mode. The ion peak observed at m/z 238.1252 suggested the protonated molecular formula of compound 11 to be $C_{13}H_{20}NOS$ (calcd. 238.1266). The LC-MS and GC-MS analyses revealed that product K mainly contained compound 11. Therefore, product K was directly dissolved in CDCl₃ and analyzed by NMR spectroscopy. The ¹H and ¹³C NMR



(e) JWH-122 *N*-(5-chloropentyl) analog (8) (11.1 min)



Fig. 8 GC-MS and LC-MS analyses of product F. TIC (a) and EI mass spectra of peak 8 (b) obtained by GC-MS analysis. LC-UV-PDA chromatogram (c) and TIC (d) using elution program (1) obtained by LC-MS. UV and ESI mass spectra of peak 8 (e)

spectra of compound **11** suggested the existence of 19 protons and 13 carbons (Table 2). The analyses by DQF-COSY, HMQC, HMBC, and 1D-ROE spectra for compound **11** suggested the presence of a 2-(pyrrolidin-1-yl)pentanoyl moiety (Fig. 4d). The ¹H, ¹³C NMR, and 2D NMR spectra of the remaining C_4H_3S unit suggested the existence of a 2-substituted thiophene moiety (position-1' to 5') as shown in Fig. 4d. The connection of the thiophene

moiety to the carbonyl group was shown by HMBC correlations to be from the aromatic proton (H-3') to the carbonyl carbon (C-1) (Fig. 4d). Additionally, the major fragment ions at m/z 126 and 111 of peak 11 in GC–MS spectra suggested the presence of 1-butylpyrrolidine and thiophene-2-carbonyl moieties, respectively (Fig. 3f). Therefore, the structure of 11 was determined as α -pyrrolidinovalerothiophenone [α -PVT, IUPAC: 2-(pyrrolidin-1yl)-1-(thiophen-2-yl)pentan-1-one]. Compound 11 (α -PVT) is a novel designer drug, and its chemical and pharmacological data have not been reported. However, chloro- or methyl-substituted (position-5' in Fig. 4d) thiophene analogs of α -PVT have been reported as monoamine uptake inhibitors [14]. It is thus possible that compound 11 has similar inhibitory activity.

Identification of unknown peaks 3 and 4

Two unknown peaks 3 and 4 were detected together with the known synthetic cannabinoid XLR-11 (Fig. 1b, [11]) in the GC-MS and LC-MS chromatograms of product B (Fig. 6a, b, e). In the LC-MS chromatogram, two unknown peaks 3 and 4 at 5.8 and 6.1 min showed protonated ion signals at m/z 383 and 344 and absorbance maxima at 302 and 291 nm in UV spectra, respectively (Fig. 6c, d). In the GC-MS chromatogram, peaks 3 and 4 at 50.1 and 49.5 min showed putative molecular ion signals at m/z 382 and 343, respectively (Fig. 6f, g). After the isolation of compounds 3 and 4, the accurate mass spectra were measured by LC-OTOF-MS in the positive mode. The observed ion peaks at m/z 383.1891 and 344.2347 suggested the protonated molecular formulae of compounds 3 and 4 to be $C_{21}H_{24}FN_4O_2$ (calcd. 383.1883) and $C_{20}H_{30}N_3O_2$ (calcd. 344.2338), respectively.

The ${}^{13}C$ NMR spectra of compound **3** was very similar to that of a known synthetic cannabinoid, AB-FUBINACA, except for a dimethylpropyl moiety (position-2''' to 6''') as shown in Table 3 and Fig. 1a, c [15]. The difference between the molecular formula of compound 3 and that of AB-FU-BINACA $(C_{20}H_{21}FN_4O_2)$ is the additional CH₂. The observed DQF-COSY, HMQC, HMBC, 15N HMBC, and 1D ROE correlations of compound 3 suggested the presence of 1-(4-fluorobenzyl)-1H-indazole and (1-amino-3,3-dimethyl-1-oxobutan-2-yl)-carboxamide moieties (Fig. 7a, b). That is, compound 3 may have an additional methyl group at the 3^{'''}-position in the structure of AB-FUBINACA. However, no HMBC correlation between the two moieties was observed. We therefore measured the deuterium isotope effect of the NH amide proton on the ¹³C chemical shift to determine the connection between the two moieties. The ${}^{13}C$ NMR spectrum of compound 3, measured in CD₃OH, was compared with that in CD₃OD. The isotope shift values for the ¹³C NMR signals of this compound are shown in Fig. 7c. The first, second, and third largest deuterium shifts (0.15, 0.14, and 0.05 ppm) were observed at the C-1^{'''} and C-2^{'''} positions of the 1-amino-3,3-dimethyl-1-oxobutane moiety and the C-1 position of the carboxamide moiety, respectively. The fourth largest shift of 0.02 ppm was attributed to the three-bond deuterium isotope effects of the NH amide proton on the indazole carbon (C-3') and the 1-amino-3,3-dimethyl-1-oxobutane carbon (C-3'''). These results strongly indicate that the 1-(4-fluorobenzyl)-1*H*-indazole moiety is connected at the 3'-position of the indazole to the carboxyamide (1-CONH). In addition, the major fragment ion signals at m/z 109, 253, and 338 revealed by the GC–MS analyses (Fig. 6f) supported the presumed structure of compound **3**. Therefore, compound **3** was identified as *N*-(1-amino-3,3-dimethyl-1oxobutan-2-yl)-1-(4-fluorobenzyl)-1*H*-indazole-3-carbox-amide (Fig. 1a).

The S-form of compound **3** has been reported to have an affinity for the cannabinoid CB₁ receptor ($K_i = 0.36$ nM) that is 25- and 2.5-fold more potent than those of JWH-018 ($K_i = 9.0$ nM) and AB-FUBINACA ($K_i = 0.9$ nM), respectively [16, 17]. Considering its general properties, we propose a new name for this compound, ADB-FU-BINACA (**3**), with agreement from Pfizer. This is the first case in which compound **3** has been detected in an illegal product.

The major fragment ion signals at m/z 214 and 144 of peak 4 (Fig. 6g) shown by the GC–MS analysis suggested the presence of a 1-pentyl-1*H*-indole-3-carbonyl moiety from comparison of mass fragment patterns of known

Table 4 NMR data for JWH-122 N-(5-chloropentyl) analog (8) and AM-2201 4-methoxynaphthyl analog (4-MeO-AM-2201) (9)

No.	JWH-122 ^a ¹³ C	JWH-122 N-(5-Chloropentyl) analog (8) ^b		4-MeO-AM-2201 (9) ^c	
		¹³ C	¹ H	¹³ C	¹ H
1	192.2	191.5	_	190.5	_
2'	137.8	137.1	6.96, 1H, s, overlapped	138.9	7.83, 1H, s
3'	117.7	118.6	_	116.2	_
3′a	127.0	127.0	_	126.6	_
4′	122.9	123.9	9.13, 1H, d, $J = 7.9$ Hz	121.7	8.30, 1H, d, $J = 7.6$ Hz
5'	122.7	123.2	7.34, 1H, t, $J = 7.9$ Hz	122.3	7.29, 1H, t, $J = 7.6$ Hz, overlapped
6'	123.5	123.7	7.22, 1H, brt, $J = 7.9$ Hz	123.2	7.32, 1H, t, $J = 7.6$ Hz, overlapped
7′	109.9	109.9	6.95, 1H, d, $J = 7.9$ Hz	110.9	7.64, 1H, d, $J = 7.6$ Hz
7′a	137.0	137.2	_	136.7	_
1″	47.1	46.2	3.05, 2H, t, $J = 7.2$ Hz	46.0	4.23, 2H, t, $J = 6.9$ Hz
2"	29.5	28.8	0.93, 2H, m	29.0	1.76, 2H, quintet, $J = 6.2$ Hz
3″	28.9	23.9	0.71, 2H, m	21.9, d, J = 5.8 Hz	1.28, 2H, quintet, $J = 6.2$ Hz
4″	22.2	31.9	1.04, 2H, m	29.3, d, <i>J</i> = 18.8 Hz	1.63, 1H, quintet, $J = 6.2$ Hz, 1.58, 1H, quintet, $J = 6.2$ Hz
5″	13.9	44.3	2.85, 2H, t, $J = 6.5$ Hz	83.6, d, <i>J</i> = 161.8 Hz	4.40, 1H, t, <i>J</i> = 6.2 Hz, 4.32, 1H, t, <i>J</i> = 6.2 Hz
1'''	137.5	138.9	_	125.0	_
2'''	125.8	125.8	7.54, 1H, d, $J = 6.9$ Hz	127.9	7.70, 1H, d, $J = 7.9$ Hz
3'''	125.2	125.6	7.09, 1H, d, $J = 6.9$ Hz	103.1	7.06, 1H, d, $J = 7.9$ Hz
4'''	136.6	136.3	_	156.1	_
4′′′a	132.8	133.3	_	130.8	_
5'''	124.2	124.5	7.83, 1H, brd, $J = 8.4$ Hz	121.7	8.25, 1H, m
6'''	126.1	126.4	7.29, 1H, m	125.7	7.55, 1H, m, overlapped
7'''	126.4	126.7	7.26, 1H, m	127.2	7.54, 1H, m, overlapped
8'''	126.6	127.4	8.61, 1H, brd, $J = 7.6$ Hz	125.4	8.12, 1H, m
8′′′a	130.9	131.7	_	131.5	_
4'''-Me	19.8	19.6	2.42, 3H, s	_	_
4'''-OMe	_	-	_	55.9	4.06, 3H, s

Recorded at 600 MHz (¹H) or 150 MHz (¹³C); data in δ ppm

^a Recorded in CDCl₃ at 150 MHz (¹³C)

^b Recorded in benzene-d₆

^c Recorded in DMSO-*d*₆



Fig. 9 DQF-COSY, selected HMBC, and selected ROE correlations for compound 8 [JWH-122 *N*-(5-chloropentyl) analog, **a**] and compound 9 [AM-2201 4-methoxynaphthyl analog (4-MeO-AM-2201), **b**]

synthetic cannabinoids that have the same moiety, such as JWH-018 and JWH-122 [18, 19].

The ¹H and ¹³C NMR spectra of compound **4** suggested the existence of 29 protons and 20 carbons as shown in Table 3. The fragment ions at m/z 214 and 299 of peak 4 by GC-MS analysis (Fig. 6g) and the observed DQF-COSY, HMQC, HMBC, and 1D ROE spectra of compound 4 suggested the presence of 1-pentyl-1H-indole and (1amino-3,3-dimethyl-1-oxobutan-2-yl)-carboxamide moieties (Fig. 7d; Table 3). In addition, the HMBC correlations from the amide proton (1-CONH) and the indole proton (H-2') to the carboxyamide carbon (C-1) suggested that the carboxyamide carbon (C-1) in the (1-amino-3,3-dimethyl-1-oxobutan-2-yl)carboxamide moiety was attached to the carbon at the 3'-position of the 1-pentyl-1H-indole moiety (Fig. 7d). On the basis of the mass spectral and NMR data, the structure of compound 4 was determined to be N-(1amino-3,3-dimethyl-1-oxobutan-2-yl)-1-pentyl-1H-indole-3-carboxamide. We named the compound ADBICA (Fig. 1a). This is the first study in which compound 4 has been detected in an illegal product. Compound 4 is a novel cannabimimetic substance and its chemical and pharmacological data have not been reported, although its structure is similar to that of a known indazole derivative, AB-PINACA (Fig. 1c, [15]).

Identification of the unknown peaks 5-7

GC–MS and LC–MS analyses were performed to identify the unknown peaks **5**, **6**, and **7** in products C (Suppl., Fig. S2), D (Fig. S3), and E (Fig. S4), respectively. Based on the GC–MS and LC–MS data, the three peaks were identified as APICA *N*-(5-fluoropentyl) analog (Fig. S2a, b, d–f), APINACA *N*-(5-fluoropentyl) analog (Fig. S3a, b, d–f), and UR-144 *N*-(5-chloropentyl) analog (Fig. S4a, b, d–f) by direct comparison of the data with those of the purchased authentic compounds, respectively (Fig. S2c, g; Fig. S3c, g; Fig. S4c, g). In addition, compound **7** [UR-144 *N*-(5-chloropentyl) analog] was detected along with XLR-11 and a cathinone derivative, α -PVP, which are controlled as designated substances (Shitei-Yakubutsu) in Japan, in product E (Fig. 1b; Fig. S4a, d, e).

Compounds 5, 6, and 7 were detected as newly distributed designer drugs in Japan. These compounds are analogs of known cannabimimetic substances, APICA, APINACA, and UR-144, respectively, which have been controlled as designated substances (Shitei-Yakubutsu) in Japan since 2012 [5, 11, 20].

Identification of unknown peaks 8 and 9

Unknown peak **8** was detected together with known synthetic cannabinoid MAM-2201 (Fig. 1b) in the GC–MS and LC–MS chromatograms of product F (Fig. 8a, c, d). The proposed fragment patterns and presumed structure of peak **8** obtained by GC–MS analysis are shown in Fig. 8b. The LC–MS data revealed that peak **8** showed absorbance maxima at 226 and 314 nm in the UV spectrum, a protonated ion signal at m/z 390 ([M+H]⁺), and an isotopic ion signal at m/z 392 ([M+2+H]⁺) due to the presence of chlorine atom (Fig. 8e). After the isolation of compound **8**, the accurate mass spectrum obtained by LC–QTOF–MS gave an ion peak at m/z 390.1632, suggesting the protonated molecular formula of C₂₅H₂₅ClNO (calcd. 390.1625).

The ${}^{13}C$ NMR spectrum of compound **8** was very similar to that of JWH-122 (Fig. 1c) except for the chlorine-



Fig. 10 GC-MS and LC-MS analyses of product G. TIC (a) and EI mass spectra of peak 9 (b) obtained by GC-MS analysis. LC-UV-PDA chromatogram (c) and TIC (d) using elution program (1) obtained by LC-MS. UV and ESI mass spectra of peak 9 (e)

substituted moiety (C-5") of compound **8**, as shown in Table 4. On the basis of the mass spectra and the observed DQF-COSY, HMBC, and 1D ROE correlations shown in Fig. 9a, the structure of compound **8** was identified as JWH-122 *N*-(5-chloropentyl) analog (IUPAC: [1-(5-chloropentyl)-1*H*-indol-3-yl](4-methylnaphthalen-1-yl)methanone) (Fig. 1a).

In the GC–MS and LC–MS chromatograms of product G, unknown peak 9 was detected along with the peak of QUPIC (1) (Fig. 10a, c, d). Peak 9 at 55.00 min showed a putative molecular ion signal at m/z 389 by GC–MS

analysis (Fig. 10b). The proposed fragment patterns and presumed structure of compound **9** by GC–MS analysis are also shown in Fig. 10b. The LC–MS data revealed that peak **9** showed absorbance maxima at 232 and 316 nm in the UV spectrum and a protonated ion signal at m/z 390 ([M+H]⁺) (Fig. 10e). After isolation of compound **9**, the accurate mass spectrum obtained by LC–QTOF–MS showed an ion peak at m/z 390.1859, suggesting the protonated molecular formula of C₂₅H₂₅FNO₂ (calcd. 390.1869).

The ¹H and ¹³C NMR spectra of compound 9 suggested the presence of N-(5-fluoropentyl)-3-carbonyl indole and 4-methoxynaphthyl moieties as shown in Fig. 9b. The fragment ions at m/z 232 and 157 of compound 9 in the GC-MS spectrum also supported the presence of these moieties (Fig. 10b). The connection of the two moieties was revealed by HMBC correlations from the indole proton (H-2') and the naphthyl proton (H-2") to the carbonyl carbon (C-1) as shown in Fig. 9b. Therefore, the structure of compound 9 was identified as AM-2201 4-methoxynaphthyl analog (4-MeO-AM-2201, IUPAC: [1-(5-fluoropentyl)-1H-indol-3-yl](4-methoxynaphthalen-1-yl)methanone) (Fig. 1a). Compounds 8 and 9, which have no reported pharmacological data, are analogs of the known synthetic cannabinoids JWH-122 and AM-2201, respectively.

Identification of unknown peaks 12 and 13

In the GC–MS and LC–MS analyses using elution program (2) [11], unknown peaks 12 and 13 were detected together with α -PBP and known synthetic cannabinoid EAM-2201 (Fig. 1b, [11]) in product H (Fig. 11a, e, f). Based on the GC–MS and LC–MS data, peak 12 was identified as opioid receptor agonist AH-7921 [3,4-dichloro-*N*-([1-(dimethyl-amino)cyclohexyl]methyl)benzamide] (Figs. 1a, 11b, g) by direct comparison of the data with those of the purchased authentic compound (Fig. 11c, h). AH-7921 (12) has been classified as an opioid analgesic with high addictive liability and has been reported to act as a selective μ -opioid receptor agonist [21, 22]. The present study is the first reported case in which AH-7921 has been detected in an illegal product.

Unknown peak **13** was presumed to be AB-001 *N*-(5-fluoropentyl) analog, from the proposed fragment patterns obtained by GC–MS analysis (Fig. 11d). The LC–MS chromatograms showed that peak **13** exhibited protonated ion signals ($[M+H]^+$) at m/z 368 and showed absorbance maxima at 220, 246, and 303 nm (Fig. 11i). Peak **13** was detected at 12.8 min under elution program (1) (data not shown [11]). The accurate mass spectrum was measured by LC–QTOF–MS. The observed ion peak at m/z 368.2389 suggested that the protonated molecular formula of



Fig. 11 GC–MS and LC–MS analyses of product H. TIC (a) and EI mass spectra of peaks 12 (b) and 13 (d) and authentic AH-7921 (c) obtained by GC–MS. LC–UV-PDA chromatogram (e) and TIC

spectra of peaks 12 (g) and 13 (i) and authentic AH-7921 (h)

(f) using elution program (2) obtained by LC-MS. UV and ESI mass

compound **13** was $C_{24}H_{31}FNO$ (calcd. 368.2389). A product ion peak at m/z 135.1169 indicating the presence of an adamantyl group $[C_{10}H_{15}$ (calcd. 135.1174)] in the structure of compound **13** was also detected by LC–QTOF–MS-MS analysis (data not shown). The above results supported the putative structure of compound **13** as AB-

001 N-(5-fluoropentyl) analog. A strict confirmation analysis of the structure using NMR spectroscopy is now in progress.

To the best of our knowledge, the chemical and/or pharmacological data of most of the new detected compounds (except compounds 3 and 12) have not been

reported. We are, therefore, conducting the following two examinations: (1) the affinities of the abused synthetic cannabinoids for cannabinoid CB_1/CB_2 receptors as described in our previous reports [11, 23], and (2) the inhibitory activities of the cathinones and their related derivatives on the neuronal uptake of the monoamines. The results will be reported in the near future (Kikura-Hanajiri et al., in preparation).

Of the new designer drugs distributed since late 2011 in Japan, the new synthetic cannabinoids belong to chemically diverse families, such as the naphthoylpyrroles (to which JWH-307 and JWH-030 belong [11]), the adamantyl-indoles/indazoles (to which APICA and APINACA belong [5]), and the dicarboxamide-indazoles (to which AB-PINACA and AB-FUBINACA belong [15]). With the marked increase in the detection of new cathinone derivatives in illegal products, other substances belonging to an expanding range of chemical families that are derivatives of controlled drugs such as aminoindanes (5-IAI), tryptamines (4-OH-DET), and arylcyclohexylamines (methoxetamine) have begun to appear in illegal drug markets [7– 11, 24].

Our ongoing survey of designer drugs in the illegal market in Japan has revealed that a recent trend is the supply of a mixture of different designer drugs, such as cathinones (stimulants) and tryptamines (hallucinogens) with synthetic cannabinoids in one illegal product [11]. More recently, several potent hallucinogenic *N*-(2-methoxy)benzyl phenethylamine derivatives, 25I-NBOMe, 2C-C-NBOMe, and 25H-NBOMe [25, 26], were detected together with synthetic cannabinoids, AM-2201 and MAM-2201, in an illegal herbal product distributed in Japan (unpublished observation).

Conclusions

In this study, we detected two new-type cannabimimetic quinolinyl carboxylates, QUPIC (1) and QUCHIC (2); two new cannabimimetic carboxamide derivatives, ADB-FU-BINACA (3) and ADBICA (4); and five new distributed synthetic cannabinoids, APICA N-(5-fluoropentyl) analog (5), APINACA N-(5-fluoropentyl) analog (6), UR-144 N-(5-chloropentyl) analog (7), JWH-122 N-(5-chloropentyl) analog (8), and AM-2201 4-methoxynaphthyl analog (4-MeO-AM-2201, 9) in illegal products in Japan. In addition, 8-quinolinol (10), a synthetic component of compound 1 or 2 was detected. Moreover, a stimulant thiophene analog, α -PVT (11), and an opioid receptor agonist, AH-7921 (12), were detected as new types of designer drugs together with several synthetic cannabinoids and cathinone derivatives in illegal products. The types of designer drugs and their combinations in illegal products seem to be diversifying,

and more serious health risks will be associated with their use than ever before. Therefore, continuous monitoring and rapid identification of newly distributed designer drugs are essential to prevent their abuse.

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