Supporting Information:

Purine (N)-Methanocarba Nucleoside Derivatives Lacking an Exocyclic Amine as Selective A₃ Adenosine Receptor Agonists

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Scheme S1. A proposed mechanism for the unexpected formation of 6-Me purine derivatives.

Scheme S2. Reagents and Conditions: (i) isoamyl nitrite, iodine, CH₂I₂, CuI, THF, reflux; (ii) TFA, reflux.



Binding to off-target sites

K_i determinations and binding profiles in a broad screen of receptors and channels were generously provided by the National Institute of Mental Health's Psychoactive Drug Screening Program, Contract # HHSN-271-2008-00025-C (NIMH PDSP). The NIMH PDSP is Directed by Bryan L. Roth MD, PhD at the University of North Carolina at Chapel Hill and Project Officer Jamie Driscol at NIMH, Bethesda MD, USA. For experimental details please refer to the PDSP web site <u>http://pdsp.med.unc.edu/</u> and click on "Binding Assay" or "Functional Assay" on the menu bar.

Figure S1. Radiologand binding at (A) the human dopamine transporter (hDAT); compound 33256 =

Compound 15; (B) the human d-opioid receptor (hDOR); compound 40132 = Compound 19.





Figure S2. Per residue IS_{ele} (panel A) and IS_{vdW} (panel B) maps for the most energetically favorable docking poses of compounds with $K_i < 100$ nM. The maps have been computed for selected poses of the considered compound inside the putative binding site of hA₃ AR. Values are expressed in kcal/mol.

Figure S3. Representative structure for compounds 32a, 12, and 15 (shift data given in Tables S1, S2 and S3).



1 able S1. Compound 32a, 298K, CDCl ₃ (ρ H = $/.24$, ρ C $//.2$ ppn	Table S1. Com	oound 32a, 298K	, CDCl ₃ (δ H =7.24	, <i>δ</i> C 77.2	ppm
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#	δ H, multiplicity: <i>J</i> Hz	ðС	HMBC coupling to #
2		158.0	
4		148.0	
5		128.2	
6		145.5	
8	7.82	141.5	2, 4, 5
9	4.82	61.8	4, 8, 10, 11, 12, 13, 14
10	4.85, dd:7.1:1.4	89.6	12, 13
11	5.78, dd:7.1:0.8	83.9	9, 12, 14, 15, 29
12		42.1	

13	1.97, ddd:9.4:5.1:1.5	35.6	10, 11, 15
14	1.66, ddd: 9.4:5.1:1.4 1.28, t:5.2	15.1	9, 11, 12, 13, 15 9, 11, 12, 13, 15
15	NH=7.37, q:5.1	171.2	15, 16
16	2.81, d:4.8	27.5	12, 15
17		89.7	
18		85.2	
19		122.2	
20	7.66	132.7	18, 20, 22
21	7.41	129.1	19, 20
22	7.42	129.9	20
23	5.77	89.1	X(CH ₃), 2, 5, 6, 21, 25
24	NH=10.24, q:5.0 CH ₃ =3.0, d:5.4	163.6 32.6	23, 24(CH3), 25 24
25		136.9	
26	7.46	128.5	24, 26, 27, 28
27	7.42	128.8	25, 26
28	7.41	129.6	26
29		113.2	
30	1.28	24.8	29, 31
31	1.55	26.5	29, 30

Table S2. Compound 12, 298K, DMSO (*d*H =2.5, *d*C=39.5 ppm)

#	δ H, multiplicity: <i>J</i> Hz	∂ С(ОН), ∂ С(С=0)	HMBC coupling to #
2		154.5, 156.0	
4		148.8, 150.4	
5		126.4, 132.4	
6		141.1, 144.6	
8	8.53	143.6, 145.3	2, 4, 5
9	4.85	61.4, 61.5	4, 8, 10, 11, 12, 13, 14
10	4.03, bt		9, 12, 13
10	0H=5.46	76.3	9, 10, 11
11	5.01, t:6.3	70.9	9, 12, 14, 15
11	OH=4.86		10, 11, 12

12		37.9	
13	1.94, dd:9.0:4.6	27.5	10, 11, 12, 14, 15
14	1.35, ddd: 9.0:4.6:1.5	13.9	9, 11, 12, 13, 14, 15
17	1.65, t:4.6	15.7	9, 11, 12, 13, 14, 15
15	NH=7.59	171.1	NH=15, 16
16	2.68	26.1	12, 15
17		86.89, 88.9	
18		86.94, 85.2	
19		120.2, 120.8	
20	7.76, dm:6.7	132.2, 132.0	18, 20, 21, 22
21	7.53	128.9	20, 21
22	7.55	130.2, 129.8	20, 22
23	6.76	88.0, 42.8	2, 6, 24, 25, 26
24	OH: 4.93	170.3, 194.5	OH: 2, 5, 24
25		135.0, 136.2	
26	7.99	126.0, 128.4	24, 25, 26, 27, 28
27	7.53, dm:7.4	128.8	
28	7.56	131.0, 133.6	

Table S3. Compound 15, 298K, MeOD (*d*H =3.31, *d*C=49.2 ppm)

#	δH , multiplicity: <i>J</i> Hz	ðС	HMBC coupling to #
2		160.8	
4		151.8	
5		133.5	
6		147.2	
8	8.51	146.5	2, 4, 5, 6, 9, 17
9	4.99	64.2	4, 8, 10, 11, 12, 13, 14
10	4.10, dt:6.6:0.9	77.9	9, 12, 13, 14
11	5.14, dd:6.3:1.3	73.5	9, 10, 12, 14, 15
12		39.9	
13	2.14, ddd:9.0:4.9:1.3	28.9	10, 11, 12, 14, 15
14	1.40, ddd: 9.0:5.1:1.7 1.87, t:4.9	15.5	9, 11, 12, 13, 15 9, 11, 12, 13, 15

15		174.9	
16	2.84	27.3	12, 15
17		89.3	
18		87.6	
19		123.1	
20	7.68	133.5	18, 19, 20, 21, 22
21	7.46	130.1	19, 20, 21
22	7.47	131.1	20, 21
23	2.81	19.3	2, 6, 17, 20



Most protonated carbon and ¹H-spin system assignments were relatively straightforward via HSQC and COSY/TOCSY, respectively, for **32a** and **15**, with ¹³C-coupled and HMBC experiments providing additional information for overlapping resonances such as 15's #5 and #20 carbons. HMBC also provided the remaining pertinent information for the quaternary carbons. Notably, the purine #8 proton of **12** coupled to each ¹³C in its ring and to that of the

#17 acetylene carbon. The structurally important couplings are shown with arrows in Figure S4, with all couplings listed in Tables S1-S3.

Compound **12** was a keto/enol mixture that was not exchanging on the chemical shift time scale. The ¹³C-shift column in Table 2 shows the enol resonances first (major component, 60%) and then the keto resonances if they were assignable. Every aromatic ¹³C and ¹H resonance as well as #9 shifted with tautomerization. Figures S5A and S5B show the ¹H and ¹³C DMSO spectra of **12**, with examples of the enol/keto shifts in the insets.



Figure S5A. Compound 12 ¹H spectra in DMSO-d6



Figure S5B. Compound 12 ¹³C spectra in DMSO-*d*6.

Selected NMR and mass spectra



Monoisotopic Mass, Even Electron lons 376 formula(e) evaluated with 6 results within limits (up to 50 best isotopic matches for each mass) Elements Used: C: 0-500 H: 0-1000 N: 0-6 O: 0-8 09-Mey-2013 13:19:40 TOF MS ES+

dkt-ix-25-up 80 (1.361) AM (Top,6, Ht,10000.0,0.00,1.00); Sm (SG, 2x3.00) 2.04e+002



Compound 12







Monoisotopic Mass, Even Electron Ions 66 formula(e) evaluated with 7 results within limits (up to 19 closest results for each mass) Elements Used:

Elements Used: C: 0-100 H: 0-200 N: 5-5 O: 0-30 35Cl: 1-1 32S: 1-1 18-Jun-2015 dkt-18jun15-xiii-18 75 (1.276) Cn (Cen,7, 50.00, Ar); Sm (SG, 1x2.00); Sb (12,5.00)





Compound 19

TOFMSES+







Compound 21





Compound 22