

1 Assessing vesicular monoamine transport and
2 toxicity using fluorescent false neurotransmitters

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10 Abstract

11 Impairments in the vesicular packaging of dopamine result in an accumulation of dopamine in the
12 cytosol. Cytosolic dopamine is vulnerable to two metabolic processes - enzymatic catabolism and
13 enzymatic- or auto-oxidation - that form toxic metabolites and generate reactive oxygen species.
14 Alterations in the expression or activity of the vesicular monoamine transporter 2 (VMAT2),
15 which transports monoamines such as dopamine from the cytosol into the synaptic vesicle, result
16 in dysregulated dopamine packaging. Here, we developed a series of assays using the fluorescent
17 false neurotransmitter 206 (FFN206) to visualize VMAT2-mediated vesicular packaging at
18 baseline and following pharmacological and toxicological manipulations. As proof of principle,
19 we observed a significant reduction in vesicular FFN206 packaging after treatment with the
20 VMAT2 inhibitors reserpine (IC_{50} : 73.09 nM), tetrabenazine (IC_{50} : 30.41 nM), methamphetamine
21 (IC_{50} : 2.399 μ M), and methylphenidate (IC_{50} : 94.33 μ M). We then applied the assay to investigate
22 the consequences on vesicular packaging by environmental toxicants including the pesticides
23 paraquat, rotenone, and chlorpyrifos, as well as the halogenated compounds unichlor,
24 perfluorooctanesulfonic acid, Paroil, Aroclor 1260, and hexabromocyclododecane. Several of the
25 environmental toxicants showed minor impairment of vesicular FFN206 loading, suggesting that
26 the toxicants are weak VMAT2 inhibitors at the concentrations tested. The assay presented here
27 can be applied to investigate the effect of additional pharmacological compounds and
28 environmental toxicants on vesicular function, which will provide insight into how exposures to
29 such factors are involved in the pathogenesis of monoaminergic diseases such as Parkinson's
30 disease, and the assay can be used to identify pharmacological agents that influence VMAT2
31 activity.

32 Introduction

33 Proper vesicular dopamine storage is essential for the survival of dopaminergic neurons. When
34 dopamine is not properly packaged into the synaptic vesicle, it accumulates in the cytosol where
35 it is susceptible to oxidation and enzymatic deamination, which when occurring in excess result in
36 oxidative stress and subsequent cell death.¹⁻³ Cytosolic monoamines, including dopamine, are
37 packaged into synaptic vesicles by the vesicular monoamine transporter 2 (VMAT2; *SLC18A2*).
38 Given the toxic properties of cytosolic dopamine, this process is necessary to reduce potential
39 dopaminergic toxicity.^{2,4-16} In addition to sequestering monoamines, it has been shown that
40 VMAT2 transports the neurotoxicant 1-methyl-4-phenylpyridinium (MPP⁺), the toxic active
41 metabolite of 1-methyl-4-phenyl-1,2,3,6-tetrahydropyridine (MPTP), into synaptic vesicles. MPP⁺
42 is a potent inhibitor of complex I of the electron transport chain, and exposure to MPTP results in
43 immediate onset of parkinsonism.¹⁷ However, VMAT2-mediated sequestration of MPP⁺ prevents
44 its accumulation in the cytosol, thereby diminishing its neurotoxic effect and suggesting a
45 neuroprotective role for VMAT2.^{10,18-22} These data suggest a role of VMAT2 in regulating the
46 neurotoxic effects of both endogenous (e.g. dopamine) and exogenous (e.g. MPP⁺) toxicants.

47 While genetic deletion of VMAT2 is neonatal lethal, mice with 5% functional expression
48 (VMAT2-LO mice), 50% functional expression (VMAT2-HET), or 200% functional expression
49 (VMAT2-HI mice) are viable.^{6,9,11,13,20,23,24} Genetic reduction of VMAT2 expression results in
50 progressive nigrostriatal neurodegeneration as well as olfactory deficits, depressive behavior, and
51 altered sleep latency in mice—symptoms that mimic both the motor and non-motor symptoms of
52 Parkinson’s disease (PD).^{9,25-27} In addition, VMAT2-LO and VMAT2-HET mice exhibit increased
53 cell death upon exposure to the dopaminergic neurotoxicants methamphetamine and MPTP.<sup>4,8,9,11-
54 13</sup> Conversely, VMAT2-HI mice exhibit protection against the toxic consequences of

55 methamphetamine and MPTP.⁴⁻⁶ Collectively these data suggest that while losing VMAT2
56 expression or activity may facilitate neuronal toxicity, an increase in expression or activity may
57 allow VMAT2 to confer neuroprotection. For these reasons, it is essential to understand the factors
58 that modulate VMAT2 expression and activity.

59 Analyses from human studies further implicate VMAT2 as a necessary mediator of neuronal
60 health. Decreases both in the amount of, and activity of, VMAT2 have been detected in post-
61 mortem tissue from PD patients,^{28,29} and mutations in VMAT2 have been identified as causative
62 of infantile parkinsonism.³⁰ Recent work has identified low-activity variants in VMAT2 that may
63 increase the risk of PD, and gain of function polymorphisms in *SLC18A2* have been associated
64 with decreased risk for PD.³¹⁻³³ Furthermore, data shows a decrease in VMAT2 mRNA from
65 platelets of PD patients suggesting that a systemic deficiency in VMAT2 may be a pathologic
66 characteristic of the disease.³⁴ Genetic predisposition only accounts for a portion of PD risk, the
67 rest of which is explained by environmental exposures including manganese and the pesticides
68 rotenone and dieldrin.³⁵⁻³⁹ While the mechanisms by which these toxicants contribute to PD
69 pathogenesis remain unknown, it is possible that these toxicants exert their neurotoxicity in part
70 by affecting VMAT2 function.^{40,41}

71 Fluorescent false neurotransmitters (FFNs) were designed as specific substrates for VMAT2 as
72 a tool to visualize vesicular uptake. FFN206 is a fluorescent monoamine analog and substrate of
73 VMAT2 that was first reported in 2013.⁴² Here, we replicate the findings of the original study that
74 show FFN206 can be used to investigate VMAT2-mediated vesicular uptake, and report new data
75 at high resolution and in near real-time. Furthermore, we optimized a 96-well plate reader-based
76 screening assay to assess VMAT2 function and the dynamics of vesicle loading under: 1)
77 physiological conditions, 2) during treatment with pharmacological inhibitors of VMAT2, and 3)

78 during treatment with select pesticides and halogenated compounds. Our data show significant
79 reductions in vesicle packaging after treatment with the VMAT2 inhibitors tetrabenazine,
80 reserpine, methylphenidate, and methamphetamine, and modest reduction in vesicle packaging
81 after treatment with the toxicological compounds paraquat, rotenone, unichlor,
82 perfluorooctanesulfonic acid, Paroil, Aroclor 1260, and hexabromocyclododecane. The methods
83 of assessing vesicle function developed here can be used to further screen pharmacological and
84 toxicological factors that alter dopaminergic vesicular storage.

85

86 Materials and Methods

87 *Cell culture.* Human embryonic kidney cells (HEK293, ATCC) were cultured at 37°C with 5%
88 CO₂ in media comprised of Dulbecco's Modified Eagle Medium (DMEM, Corning), 10% Fetal
89 Bovine Serum (FBS, Atlanta Biologicals), and 1% Penicillin-Streptomycin (Pen Strep, Corning).
90 All human VMAT2-containing constructs were made in pcDNA3.1 vectors (Life Technologies)
91 containing a zeocin resistance gene. Plasmids were transfected into HEK293 cells with
92 Lipofectamine 2000 using the manufacturer protocol and stable cell lines were generated by
93 repetitive rounds of limiting dilutions in selection media. HEK293 cells stably expressing human
94 VMAT2 (HEK+VMAT2) or mCherry-tagged human VMAT2 (HEK+mCherry-VMAT2) were
95 cultured at 37°C with 5% CO₂ in selection media comprised of DMEM (Corning), 10% FBS
96 (Atlanta Biologicals), 1% Pen Strep (Corning), and zeocin (100 µg/mL, InvivoGen). Experimental
97 media used to optimize the 96-well plate screening assay and to screen pharmacological inhibitors
98 of VMAT2 and environmental toxicants was comprised of DMEM without phenol red (Corning),
99 1% Pen Strep (Corning), and 1% L-glutamine (Gibco).

100

101 *Live-cell total internal reflection fluorescence (TIRF) microscopy.* HEK+VMAT2 cells were
102 seeded at 60,000 cells per well on laminin-coated glass-bottom 8-well chamber dishes (LabTek)
103 and maintained in selection media until they reached 60% confluence. Upon reaching confluence,
104 the selection media was aspirated and replaced with experimental media containing 1 μ M FFN206
105 (Abcam). Cells were incubated with FFN206 for 1 hour at 37°C with 5% CO₂ before the FFN206-
106 containing media was aspirated and replaced with experimental media. Cells were then imaged at
107 37°C with 5% CO₂ on the GE Delta Vision OMX total internal reflection fluorescence (TIRF)
108 microscope (FFN206 peak excitation = 369 nm; peak emission = 464 nm).

109
110 *Real-time uptake with confocal microscopy.* HEK+VMAT2 cells or HEK+mCherry-VMAT2
111 cells were seeded at 100,000 cells per plate in laminin-coated glass-bottom round 35 mm dishes
112 (ThermoFisher) and maintained in selection media until they reached 80% confluency. Upon
113 reaching confluency, selection media was aspirated and replaced with experimental media. Cells
114 were imaged on a Nikon A1R TE2000 confocal microscope at 37°C with 5% CO₂. Cells were
115 imaged for a 30 second baseline before the addition of FFN206 to a final concentration of 20 μ M.
116 Imaging lasted for a duration of 5 minutes or 1000 seconds (16.67 minutes; FFN206 peak
117 excitation = 369 nm; peak emission = 464 nm; mCherry peak excitation = 587 nm; peak emission
118 = 610 nm).

119
120 *96-well plate screening assay.* We adapted the protocol utilized by Hu and colleagues⁴² for use
121 in a 96-well plate reader. HEK+VMAT2 cells were seeded at 40,000 cells per well in half volume,
122 black-walled, laminin-coated 96-well plates (Grenier Bio One) and maintained in selection media
123 at 37°C with 5% CO₂ until 90-100% confluent (approximately 24 hours). Upon reaching

124 confluency, selection media was aspirated and replaced with 90 μ L of either experimental media
125 or experimental media containing the desired pharmacological compound or environmental
126 toxicant. Plates were incubated with the pharmacological compound and/or environmental
127 toxicant for 30 minutes at 37°C with 5% CO₂ before FFN206 (Abcam) was diluted in experimental
128 media and added to the appropriate wells to produce a final concentration of 1 μ M FFN206 per
129 well. Plates were incubated with FFN206 for 60 minutes at 37°C with 5% CO₂. Wells were then
130 washed with sterile phosphate-buffered saline (PBS, Gibco) and imaged in PBS on a BioTek
131 Synergy H1 multi-mode plate reader (FFN206 peak excitation = 369 nm; peak emission = 464
132 nm).

133

134 *Concentrations of pharmacological compounds and environmental toxicants.* Tetrabenazine
135 (Sigma) was diluted in experimental media to a final concentration of 10 μ M per well/plate for the
136 negative control group and to concentrations of 0.0001, 0.001, 0.01, 0.1, 1, and 10 μ M to produce
137 a dose response curve. Bafilomycin (InvivoGen) was diluted in experimental media to a final
138 concentration of 1 μ M per well/plate. Reserpine was diluted in experimental media to final
139 concentrations of 0.0001, 0.001, 0.01, 0.1, 1, and 10 μ M. Methamphetamine and methylphenidate
140 were diluted in experimental media to final concentrations of 0.01, 0.1, 1, 10, and 100 μ M. Stock
141 concentrations of rotenone, paraquat, chlorpyrifos, unichlor, PFOS, Paroil,
142 hexabromocyclododecane, and Aroclor 1260 were dissolved in dimethyl sulfoxide (DMSO,
143 Fisher) and then diluted in experimental media into a 200 μ M working stock. From this working
144 stock, each compound was diluted in experimental media to final concentrations of 0.01, 0.1, 1,
145 10 and 100 μ M.

146

147 *Statistical analysis.* The data were analyzed by t-test, ANOVA, and z factor analysis as
148 appropriate using GraphPad Prism software. The z factor is commonly used in the design of
149 protocols for high-throughput screens and incorporates the positive control mean (μ_+), the positive
150 control standard deviation (σ_+), the negative control mean (μ_-), and the negative control standard
151 deviation (σ_-)⁴².

$$Z = 1 - \frac{(3\sigma_+ + 3\sigma_-)}{|\mu_+ - \mu_-|}$$

153

154 Results

155 *FFN206 packaging is dependent on VMAT2 function.* We first sought to confirm that FFN206
156 fluorescence was a reliable representation of VMAT2-mediated vesicular uptake. To this end,
157 HEK cells stably transfected with mCherry-tagged human VMAT2 (HEK+mCherry-VMAT2)
158 were treated with 1 μ M FFN206, and FFN206 fluorescence was recorded after one hour of
159 incubation. FFN206 fluorescence was observed to overlap with VMAT2 fluorescence (Figure 1A),
160 thus confirming that FFN206 was loaded into VMAT2-containing cells. Furthermore, analysis of
161 HEK+VMAT2 cells grown in a glass-bottom 8-chamber dish following incubation with 1 μ M
162 FFN206 using a GE Delta Vision OMX Blaze TIRF microscope demonstrated the localization of
163 FFN206 fluorescence within small vesicle-like compartments at high resolution (60x
164 magnification) within live cells (Figure 1B).

165 To further confirm that FFN206 was loaded via VMAT2 into vesicular compartments, HEK
166 cells stably transfected with human VMAT2 (HEK+VMAT2) were treated with 10 μ M
167 tetrabenazine (TBZ) – a pharmacological inhibitor of VMAT2. Compared to HEK cells lacking
168 VMAT2 that show no FFN206 uptake, treatment with tetrabenazine resulted in almost total loss
169 of FFN206 fluorescence in HEK + VMAT2 cells, indicating that VMAT2 function must be

170 maintained in order to observe FFN206 fluorescence (Figure 1C). Quantification of FFN206
171 fluorescence displayed as percent control demonstrated a significant increase in fluorescence in
172 HEK+VMAT2 cells treated with FFN206 than all other conditions (Figure 1D). As an additional
173 method of confirmation, cells were treated with bafilomycin, which inhibits the vesicular ATPase
174 that maintains the proton gradient present across the vesicular membrane that VMAT2 depends on
175 for sequestering dopamine. As expected, bafilomycin also depleted FFN206 fluorescence (data not
176 shown).

177

178 *Real-time VMAT2-mediated uptake of FFN206.* After examining FFN206 packaging in high-
179 resolution in a live cell, we sought to observe the dynamic packaging of FFN206 in living cells in
180 real-time. To that end, HEK+mCherry-VMAT2 cells were grown in glass-bottom round dishes
181 and recorded a baseline of mCherry and background fluorescence for 30 seconds. Cells were then
182 treated with 20 μ M FFN206 and fluorescence was recorded for 5 minutes. Image stills from 0.93s,
183 3.18s, 3.94s, and 5.94s demonstrate that over time, FFN206 fluorescence and its co-localization
184 with the VMAT2-containing vesicles is observed (Figure 2A). We then sought to examine how
185 perturbations in VMAT2 function and vesicle function affected uptake and retention of FFN206.
186 Uptake was recorded under four conditions: HEK cells without VMAT2 (HEK), HEK cells with
187 human VMAT2 (HEK+VMAT2), HEK+VMAT2 cells with VMAT2 inhibitor tetrabenazine
188 (HEK+VMAT2+TBZ), and HEK+VMAT2 cells with the proton gradient dissipater bafilomycin
189 (HEK+VMAT2+BAF). FFN206 fluorescence was recorded for 1000 seconds in each condition
190 (16.67 minutes), and quantification was performed on the fluorescence in the final 20 seconds of
191 the 1000 second uptake time-course. HEK+VMAT2 cells displayed significantly greater
192 fluorescence than HEK cells and HEK+VMAT2 cells treated with tetrabenazine or bafilomycin

193 (Figure 2B) (One-way ANOVA with Dunnett's multiple comparisons post-hoc test **** p
194 <0.0001 vs. control column HEK+VMAT2).

195

196 *Optimizing parameters for a 96-well plate assay.* We adapted the protocol utilized by Hu and
197 colleagues (2013) for use in our 96-well plate assay. In developing this protocol, we first
198 determined which concentration of FFN206 would produce a dynamic range of fluorescence in the
199 HEK+VMAT2 cell line. To this end, HEK+VMAT2 cells were treated with FFN206 at 0.0001,
200 0.001, 0.01, 0.1, 1, and 10 μ M concentrations. There was a positive relationship between
201 fluorescence and FFN206 concentration; however, variability in FFN206 fluorescence scaled with
202 the increase in signal strength (Figure 3A, C). Furthermore, the fluorescence emitted by 10 μ M
203 FFN206 approached the peak signal that can be detected by the 96-well plate reader, thus making
204 detection of subtle alterations in FFN206 fluorescence at this concentration unrealistic. Therefore,
205 a mid-level dose of 1 μ M FFN206 was chosen for use in further experiments to ensure the assay
206 had the dynamic range necessary to detect both potential decreases and increases in VMAT2
207 function.

208 The Hu (2013) protocol utilized tetrabenazine as a negative control; thus, we performed a dose
209 response of tetrabenazine-suppressed FFN206 fluorescence to determine the appropriate
210 tetrabenazine dose to utilize in our cell line. HEK+VMAT2 cells were treated with tetrabenazine
211 at 0, 0.0001, 0.001, 0.01, 0.1, 1, and 10 μ M concentrations (Figure 3B, D). The dose of
212 tetrabenazine was selected in order to achieve a good dynamic range with a high degree of FFN206
213 fluorescence suppression with the selected tetrabenazine concentration. With increasing
214 tetrabenazine concentration, FFN206 fluorescence decreased with the greatest suppression of
215 FFN206 fluorescence at 10 μ M tetrabenazine; thus, this dose was chosen for the assay.

216 High-throughput assays require a high degree of accuracy and sensitivity, and therefore demand
217 a wide dynamic range and minimal variability within the datasets. The z factor is commonly used
218 in the design of protocols for high-throughput screens and incorporates the positive control mean
219 (μ_+), the positive control standard deviation (σ_+), the negative control mean (μ_-), and the negative
220 control standard deviation (σ_-).⁴³ Z factor calculation ensures that assays with favorable z values
221 (as close to 1 as possible) will have a large band of separation between the distributions of the data
222 for the positive and negative control. A z factor above 0.5 represents a suitable assay. For the
223 FFN206 assay, the positive control was represented by HEK+VMAT2 cells treated with 1 μ M
224 FFN206 and 0 μ M tetrabenazine, while the negative control was represented by HEK+VMAT2
225 cells treated with 1 μ M FFN206 and 10 μ M tetrabenazine. After performing iterative experiments
226 to optimize cell density, incubation time, and reaction volume, a z factor of 0.76 was consistently
227 achieved, indicating that the protocol was suitable for high-throughput screening (Figure 3E).

228
229 *Pharmacological inhibitors of VMAT2.* After optimizing the 96-well plate protocol, the assay
230 was used to test a variety of pharmacological VMAT2 inhibitors to demonstrate the utility and
231 accuracy of the assay. Dose-dependent VMAT2 inhibition was observed in HEK+VMAT2 cells
232 treated with reserpine, tetrabenazine, methamphetamine, and methylphenidate from a range of
233 0.0001 to 10 μ M to determine the concentration at which VMAT2 was completely inhibited
234 (Figure 4). HEK+VMAT2 cells treated tetrabenazine yielded an IC₅₀ of 73.09 nM and showed
235 essentially complete inhibition at 1 μ M. HEK+VMAT2 cells treated with reserpine yielded an IC₅₀
236 of 30.41 nM and showed essentially complete inhibition at 0.1 μ M. HEK+VMAT2 cells treated
237 with methamphetamine yielded an IC₅₀ of 2.399 μ M and showed total inhibition by 100 μ M.
238 HEK+VMAT2 cells treated with methylphenidate yielded an IC₅₀ of 94.33 μ M.

239

240 *Environmental toxicants and VMAT2 function.* The optimized assay was then used to test a
241 variety of pesticides (Figure 5A) and halogenated environmental toxicants (Figure 5B) of interest.
242 All compounds were tested at concentrations of 0.01, 0.1, 1, 10 and 100 μM . The pesticides
243 rotenone and chlorpyrifos caused a minor decrease in FFN206 uptake at the highest concentration,
244 however, paraquat demonstrated a more significant effect on FFN206 fluorescence yielding an
245 IC_{50} of 12.41 μM (Figure 5A). The halogenated compounds unichlor, PFOS, and
246 hexabromocyclododecane did not show impairment of vesicular loading at the concentrations
247 tested (Figure 5B). However, Paroil and Arochlor 1260 showed modest effects on FFN206
248 fluorescence with IC_{50} s of 57.03 μM and 95.07 μM respectively (Figure 5B).

249

250 Discussion

251 Here, we demonstrate the utility of using false fluorescent neurotransmitters to investigate the
252 effect of pharmacological and environmental compounds on vesicular uptake in an *in vitro*
253 application. Importantly, we were able to reproduce key findings from Hu *et al.* 2013
254 demonstrating that FFN206 can be used to examine VMAT2 function and the dynamics of vesicle
255 packaging in HEK293 cells. We further went on to optimize a fluorescent 96-well plate assay with
256 a dynamic range that allows for detection of altered VMAT2-mediated vesicular uptake and is
257 amenable to high-throughput screening. We previously developed an assay (Bernstein *et al.* 2012)
258 to spatially resolve VMAT2-mediated packaging of dopamine utilizing high-content imaging with
259 a fluorescent dye and mCherry-tagged VMAT2, but this assay required time-intensive image
260 analysis to obtain suitable results.⁴⁴ The advent of FFN206 allowed us to adapt our assay to a
261 fluorescent plate reader format and to visualize monoamine transport with an ease and in a real-

262 time manner that was previously inaccessible. The methods presented here can be used in
263 experimental applications to understand how pharmacological and environmental manipulation
264 affects vesicle function and thus may contribute to monoaminergic neuron vulnerability.

265

266 *FFN206 as a tool to visualize VMAT2-dependent vesicular uptake*

267 Fluorescent false neurotransmitters were developed as specific substrates of VMAT2 for the
268 purpose of visualizing monoamine uptake and release.^{42,45-49} Previous work characterized vesicular
269 uptake of FFN206 in HEK cells stably transfected with rat VMAT2,⁴² and we first sought to
270 replicate these findings in HEK cells stably transfected to express human VMAT2. We
271 demonstrated that FFN206 fluorescence overlapped with mCherry-tagged human VMAT2
272 fluorescence, indicating that FFN206 is loaded into VMAT2-containing cells. Furthermore,
273 treatment with tetrabenazine resulted in almost total loss of FFN206 fluorescence, indicating that
274 VMAT2 function must be maintained in order to observe FFN206 fluorescence. As an additional
275 piece of evidence, TIRF-level resolution of FFN206 packaging revealed the localization of
276 FFN206 fluorescence to small vesicle-like compartments present in a single HEK+VMAT2 cell.

277

278 *Real-time visualization of FFN206 vesicular uptake*

279 We next sought to determine whether FFN206 could be used to measure real-time VMAT2-
280 mediated monoamine uptake. FFN206 uptake was recorded under four conditions: HEK cells not
281 expressing VMAT2, HEK cells expressing human VMAT2, HEK+VMAT2 cells incubated with
282 tetrabenazine, and HEK+VMAT2 cells incubated with bafilomycin. Quantification of FFN206
283 fluorescence demonstrated significant differences in HEK cells lacking VMAT2 and HEK cells
284 treated with tetrabenazine and bafilomycin compared to HEK cells expressing human VMAT2.

285 These results were as expected as tetrabenazine inhibits VMAT2 activity, thus preventing the
286 sequestration of FFN206. While cells treated with bafilomycin also exhibit decreased FFN206
287 fluorescence, this is caused by a different mechanism. Bafilomycin inhibits the vacuolar H⁺
288 ATPase present on vesicles, leading to the dissipation of the vesicular proton gradient present
289 across the vesicular membrane. As the proton gradient dissipates, the physiological state of the
290 vesicle is disrupted, and vesicular homeostasis cannot be maintained despite the continued normal
291 function of VMAT2. Thus, the decrease in FFN206 fluorescence of HEK+VMAT2 cells treated
292 with bafilomycin represent the necessity of proton motive force to sequester FFN206 within
293 vesicles.

294

295 *Pharmacological and toxicological applications of high-throughput FFN206 assay*

296 We adapted the protocol utilized by Hu *et al.* for use in our 96-well plate assay for high-
297 throughput screening applications.⁴² We chose concentrations of 1 μ M FFN206 and 10 μ M
298 tetrabenazine to ensure the assay had the dynamic range necessary to detect both decreases and
299 increases in VMAT2 function. These concentrations were determined by our own dose-response
300 experiments and reproduce the optimal concentrations determined by Hu *et al.* 2013. Iterative
301 experiments were conducted to refine the protocol until a z score above 0.70 was consistently
302 achieved. We further verified the reliability of the assay by screening pharmacological compounds
303 known to inhibit VMAT2: tetrabenazine, methamphetamine, methylphenidate, and reserpine, and
304 observed dose-dependent VMAT2 inhibition that aligned with previously published results.⁴²

305 After confirming the reliability of the FFN206 assay, we treated HEK+VMAT2 cells with select
306 pesticides and halogenated environmental toxicants. There is an association between exposure to
307 environmental toxicants and PD and it has long been established that exposure to heavy metals

308 and pesticides contribute to PD risk.^{36-39,50-54} The extent to which altered VMAT2 function
309 mediates the toxicity of exposure to these environmental toxicants has not been extensively
310 studied. Thus, we tested representative pesticides and halogenated environmental toxicants
311 including rotenone, paraquat, chlorpyrifos, unichlor, PFOS, Paroil, HBCD, and Aroclor 1260 for
312 their effect on VMAT2 function. Of these compounds, exposures to rotenone and paraquat have
313 been most extensively associated with PD in both humans and animal models.^{52,53,55}

314 Rotenone is a known inhibitor of mitochondrial complex I and exerts its toxicity by oxidizing
315 mitochondrial proteins and causing oxidative stress that leads to cell death.^{56,57} Similarly, paraquat
316 exerts its toxicity predominantly through oxidative modification of cytosolic proteins, which
317 causes oxidative stress and leads to cell death.^{55,57,58} Here, we detected a reduction in VMAT2
318 function with administration of 100 μ M of rotenone and with 10 μ M and 100 μ M of paraquat,
319 though this reduced VMAT2 activity occurred at such high doses as to be physiologically
320 irrelevant. Previous studies examining rotenone and paraquat toxicity have observed 70-80% cell
321 death resulting from mitochondrial complex I inhibition with doses of 100 nM rotenone and 200
322 μ M paraquat.^{52,57,58} The present study differs from these studies in several respects, the first of
323 which being the length of exposure. We exposed cells to each toxicant for a total of 90 minutes
324 (30-minute incubation with the toxicant followed by a 60-minute incubation with FFN206), while
325 previous studies examined toxicity after 48 hours. We also screened for VMAT2 function and not
326 cell death. As the amount of cell death imposed by a 90-minute length of exposure is minimal, we
327 did not assess cell death in our assay. Finally, given the high IC_{50} value we determined for paraquat
328 and rotenone exposure, the mild inhibition of VMAT2 caused by exposures to these toxicants is
329 not likely to be mediated by mitochondrial complex I inhibition.

330 For the halogenated compounds, we observed a mild reduction in VMAT2 function at high
331 concentrations. While previous literature indicates exposure to PFOS⁵⁹, Aroclor 1260⁷, and
332 hexabromocyclododecane⁶⁰ exerts toxicity on the dopamine system, our results indicate that
333 unichlor, PFOS, Paroil, Aroclor 1260, and hexabromocyclododecane are weak inhibitors of
334 VMAT2 over the course of a short-term exposure, and therefore it is unlikely that these compounds
335 affect VMAT2 function at environmentally and physiologically relevant concentrations.

336

337 *Conclusions*

338 In conclusion, we have used FFN206 to investigate VMAT2-mediated vesicle uptake at high
339 resolution and in near real-time. Importantly, we were able to reproduce in HEK cells expressing
340 human VMAT2 the findings reported by Hu *et al.* 2013 performed in HEK cells expressing rat
341 VMAT2. We further optimized a 96-well plate assay that has a dynamic range and is amenable to
342 a high-throughput format, and we used this assay to assess VMAT2 function and the dynamics of
343 vesicle loading upon exposure to pharmacological and environmental compounds. We observed a
344 robust reduction in VMAT2 function after exposure to tetrabenazine, reserpine,
345 methamphetamine, and methylphenidate, and observed a modest reduction in VMAT2 function
346 after short-term exposure to high concentrations of environmental toxicants, though inhibition of
347 VMAT2 is unlikely to contribute to the mechanism of action of these compounds at
348 physiologically and environmentally relevant concentrations. The methods of assessing vesicle
349 function discussed here can be used to assess how other pharmacological and environmental
350 compounds exert their toxicity at the level of the vesicle. These assays can provide evidence of
351 potential mechanism by which exposures contribute to the progression of monoaminergic diseases
352 such as Parkinson's disease.

353

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358 **Author Contributions**

359 The manuscript was written through contributions of all authors. All authors have given approval
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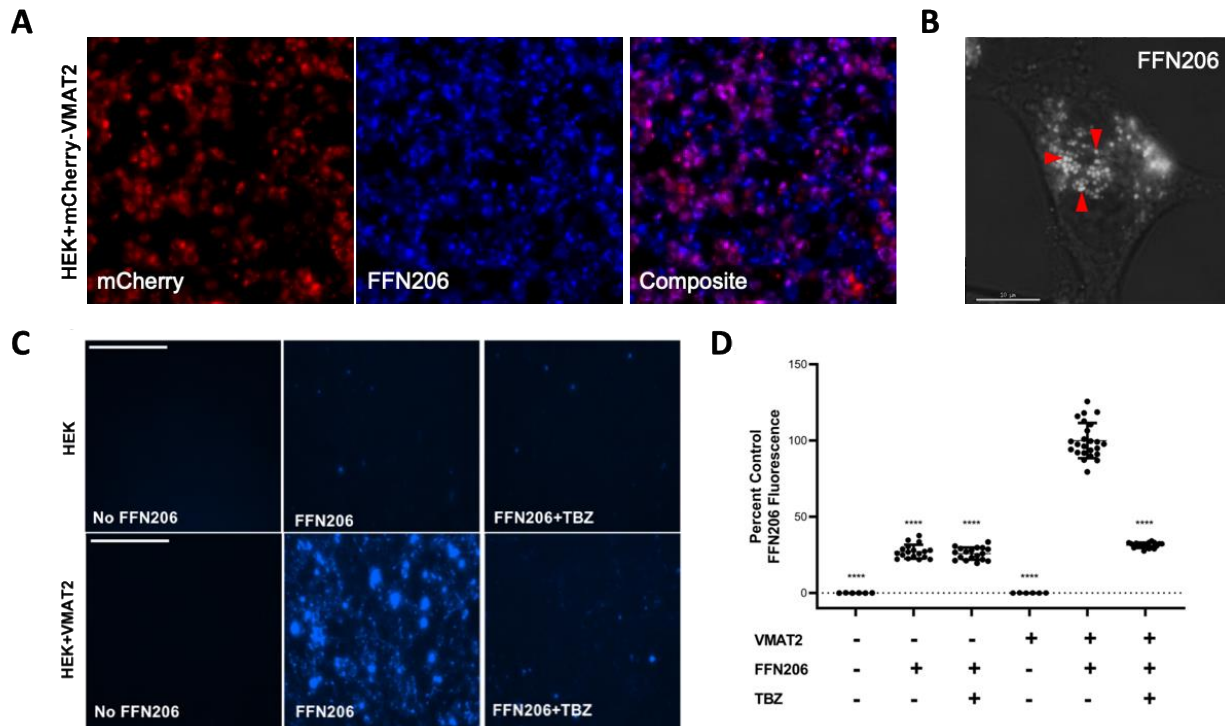
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368 **ABBREVIATIONS**

369 VMAT2, vesicular monoamine transporter 2; FFN, fluorescent false neurotransmitter; MPTP, 1-
370 methyl-4-phenyl-1,2,3,6-tetrahydropyridine; MPP⁺, 1-methyl-4-phenylpyridinium; PD,
371 Parkinson's disease; HEK, human embryonic kidney; PFOS, perfluorooctanesulfonic acid.



372

373 **Figure 1. FFN206 packaging is dependent on VMAT2 function and maintenance of the**
 374 **vesicular proton gradient.**

375 A. Representative 10x image of HEK cells stably transfected with mCherry-tagged VMAT2

376 (HEK+mCherry-VMAT2, left image) treated with FFN206 (center image). FFN206 fluorescence

377 overlaps with VMAT2 fluorescence (purple, right image). B. Representative 60x TIRF microscope

378 image demonstrates FFN206 fluorescence localized within vesicle-like compartments in a stably

379 transfected HEK cell with VMAT2 (HEK+VMAT2), as denoted by red arrows. Scale bar = 10

380 μ M. C. Representative images of FFN206 fluorescence in control HEK cells with and without

381 VMAT2. In control HEK cells (top), there was no background fluorescence in the absence of

382 FFN206 (left). FFN206 treatment exhibited minimal fluorescence (middle), and FFN206

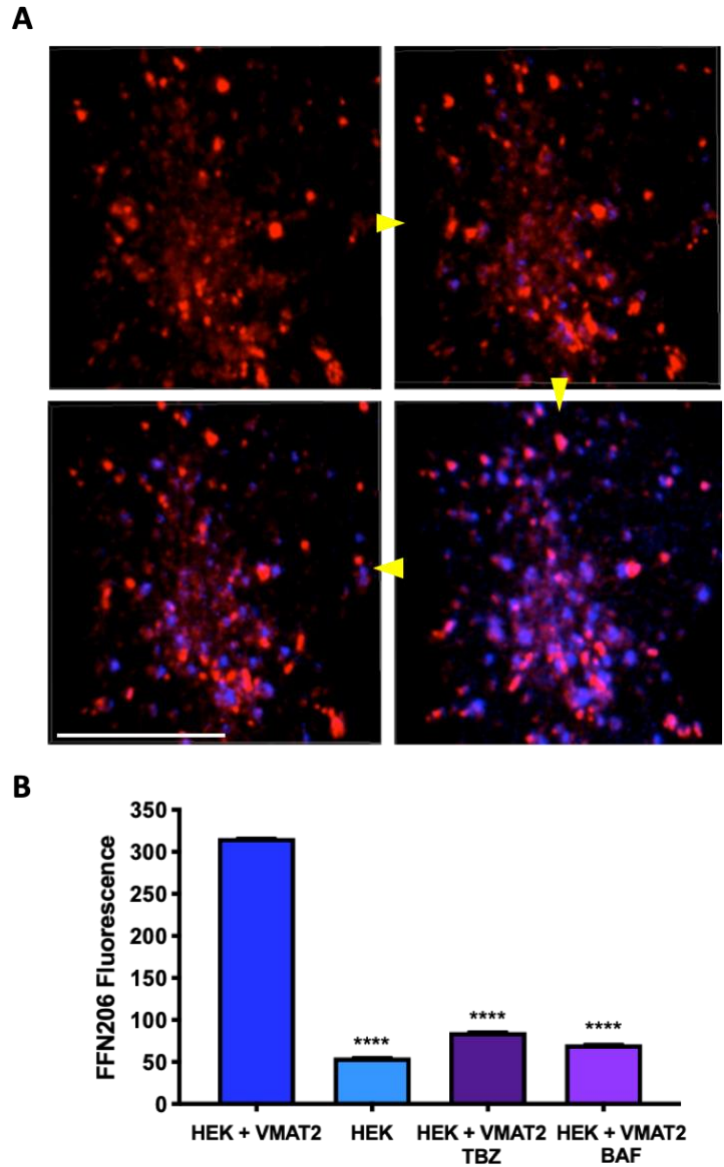
383 fluorescence was unchanged following treatment with the VMAT2 inhibitor tetrabenazine (10

384 μ M) (right). Representative images of FFN206 fluorescence in HEK+VMAT2 cells (bottom). In

385 the absence of FFN206, no background fluorescence is observed in HEK+VMAT2 cells (left).

386 HEK+VMAT2 cells treated with FFN206 exhibit robust fluorescence (middle) that was
387 diminished when VMAT2 function is inhibited by treatment with 10 μ M tetrabenazine (right). D.
388 Quantification of FFN206 fluorescence from panel C. Data displayed as percent control mean and
389 standard error of the mean, with the control group being HEK+VMAT2 cells incubated with
390 FFN206. Each point represents one well of cells from a 96-well plate. (One-way ANOVA with
391 Dunnett's multiple comparisons post-hoc test **** p <0.0001 vs. control column
392 HEK+VMAT2+FFN206).

393



394

395 **Figure 2. Real-time uptake of FFN206 in a single cell.**

396 A. Sequential stills taken from a video of FFN206 uptake (blue fluorescence) into the vesicle-like

397 compartments of a single HEK cell stably transfected with mCherry-tagged VMAT2 (red

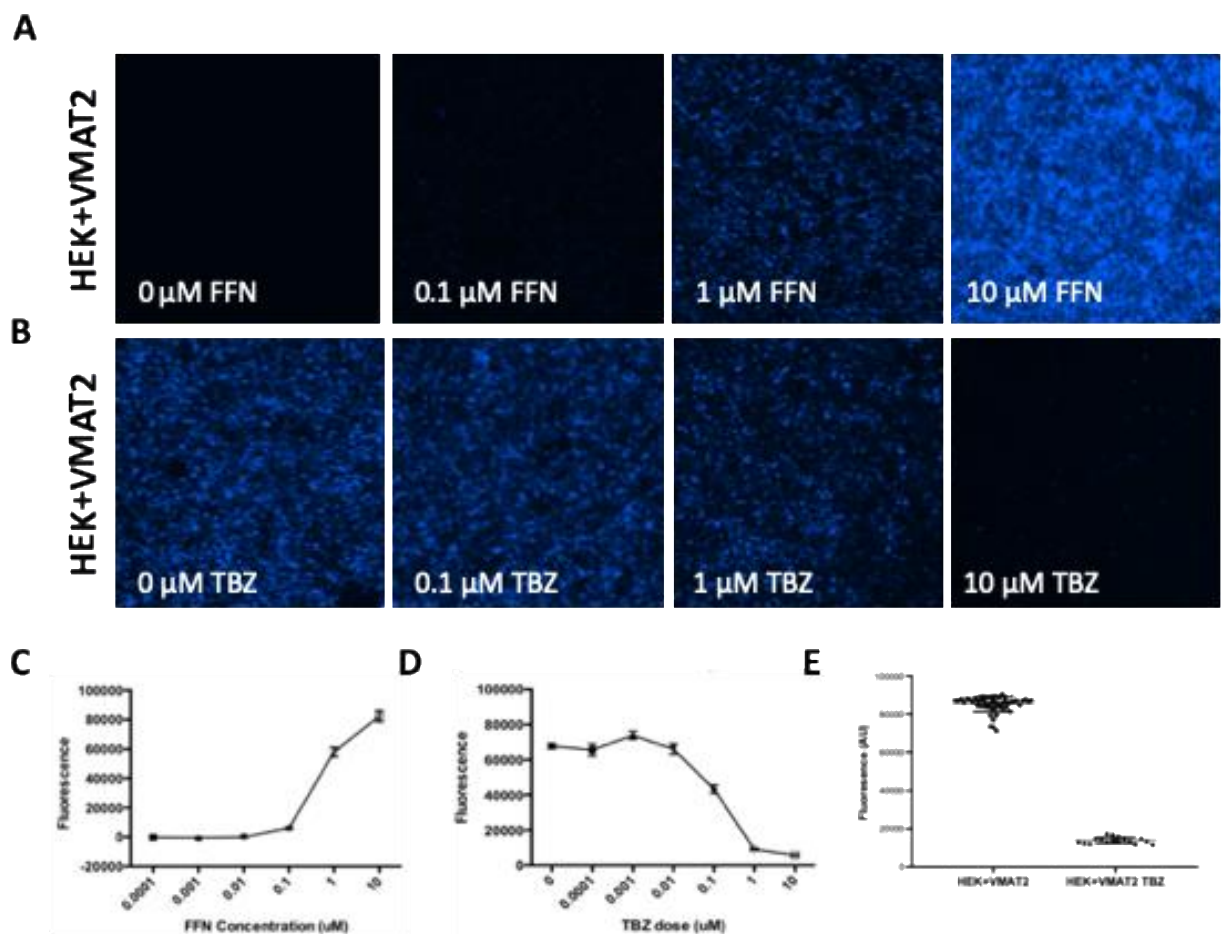
398 fluorescence). Still photos taken at 0.93s, 3.18s, 3.94s, and 5.94s following FFN206 application.

399 Scale bar = 10 μ M. Yellow arrows indicate image progression. B. Comparison of peak FFN206

400 fluorescence in the final 20 seconds of 1000 total seconds of uptake. HEK+VMAT2 cells display

401 significantly greater fluorescence than HEK cells and HEK+VMAT2 cells treated with

402 tetrabenazine or bafilomycin. Bars are displayed as mean and standard error of the mean. (One-
403 way ANOVA with Dunnett's multiple comparisons post-hoc test **** p <0.0001 vs. control
404 column HEK+VMAT2).



405

406 **Figure 3. Determining optimal parameters for a FFN206 96-well plate assay.**

407 A. Representative images of HEK+VMAT2 cells treated with 0, 0.1, 1, or 10 μM of FFN206. Cells

408 imaged at 10x magnification. B. Representative images of HEK+VMAT2 cells treated with 0, 0.1,

409 1, and 10 μM of tetrabenazine (TBZ) and 1 μM FFN206. Cells imaged by EVOS system at 10x

410 magnification. Scale bar = 400 μM. C. Quantification of fluorescence in 3A demonstrating

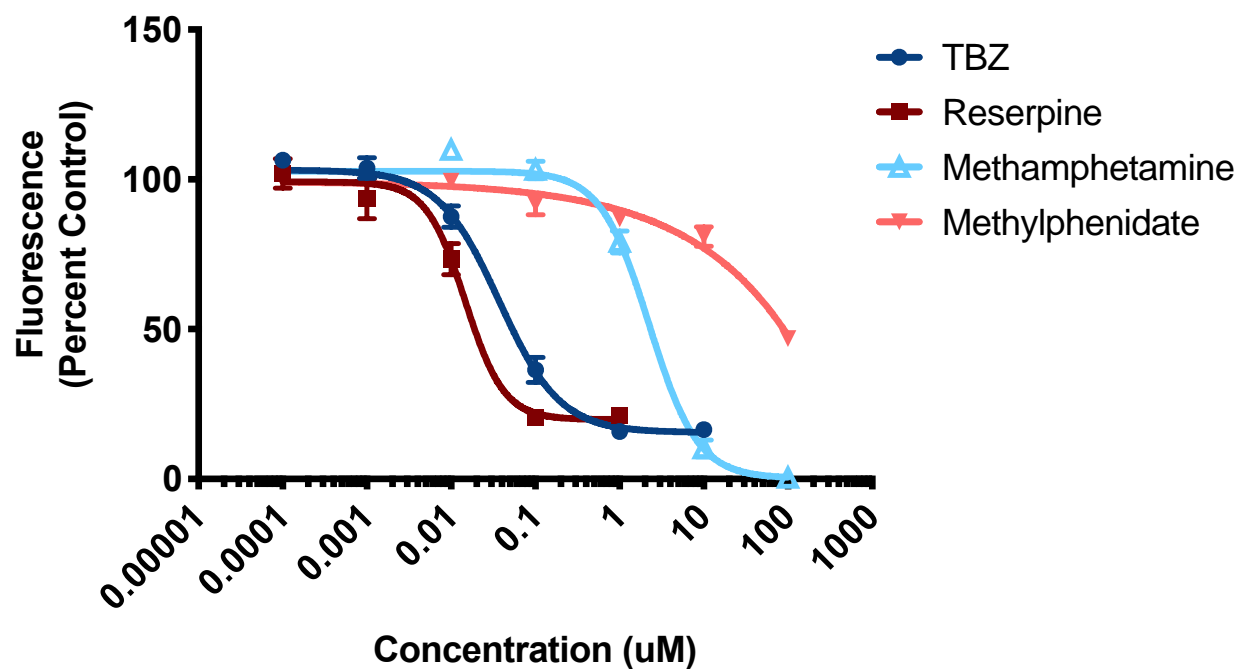
411 FFN206 fluorescence increases as FFN206 dose increases. D. Quantification of fluorescence in

412 3B demonstrating FFN206 fluorescence decreases as tetrabenazine dose increases. E.

413 Quantification of FFN206 fluorescence in HEK+VMAT2 cells treated with 1 μM FFN206 and 0

414 μM TBZ compared to cells treated with 1 μM FFN206 and 10 μM TBZ.

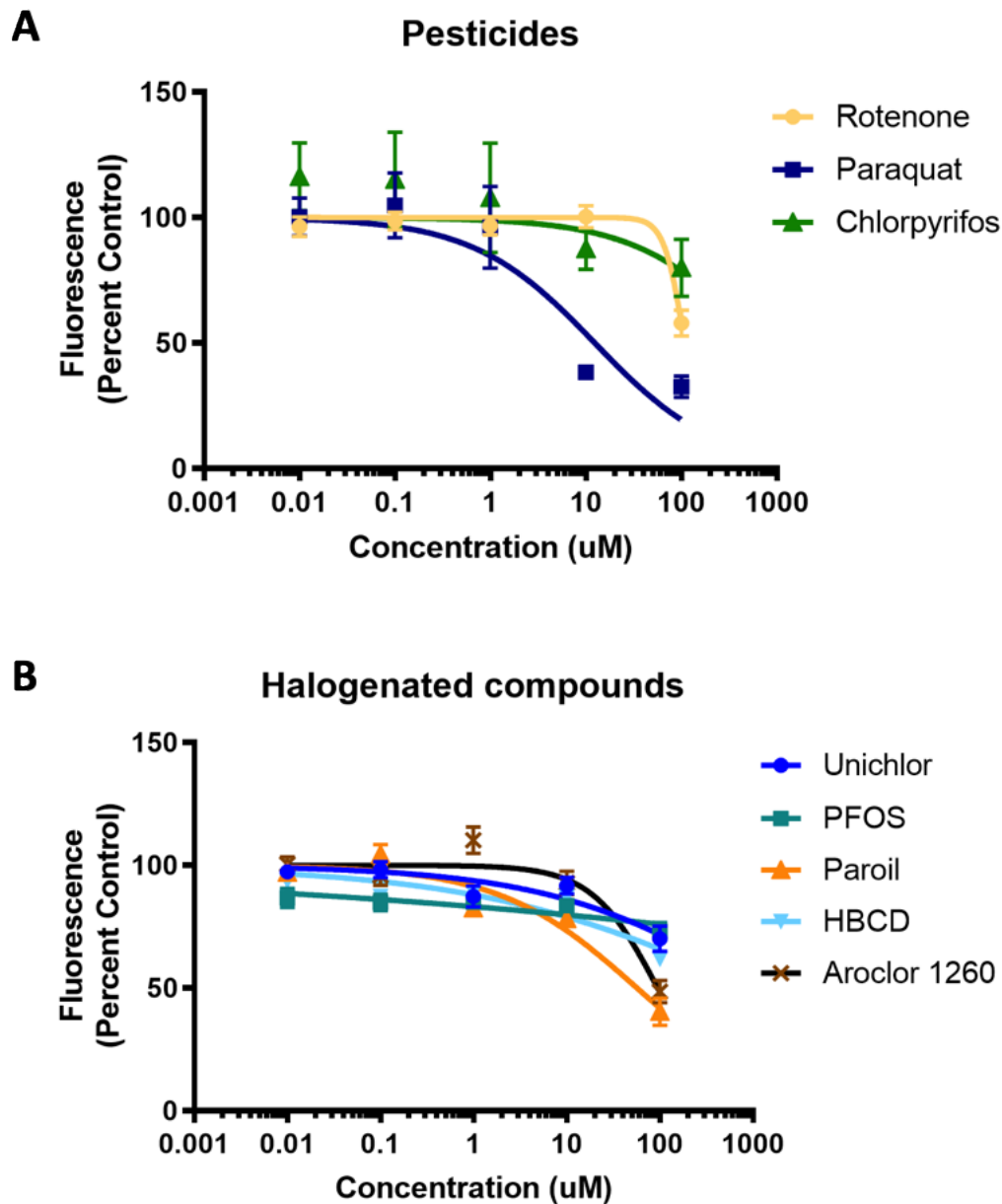
VMAT2 inhibitors



415

416 **Figure 4. Screening pharmacological inhibitors of VMAT2.**

417 HEK+VMAT2 cells were treated with 0.0001, 0.001, 0.01, 0.1, 1, and 10 µM of tetrabenazine
418 (TBZ); 0.0001, 0.001, 0.01, 0.1, 1, and 10 µM of reserpine; 0.01, 0.1, 1, 10, and 100 µM of
419 methamphetamine; or 0.01, 0.1, 1, 10, and 100 µM of methylphenidate. FFN206 fluorescence
420 represented as percent of control (HEK+VMAT2 cells treated with 0 µM drug). Graphs depict
421 mean and standard error of the mean and curves represent non-linear regressions.



422

423 **Figure 5. Screening environmental toxicants for influence on VMAT2 function.**

424 A. HEK+VMAT2 cells were treated with 0.01, 0.1, 1, 10 and 100 μ M of the pesticides rotenone,

425 paraquat, or chlorpyrifos. B. HEK+VMAT2 cells were treated with 0.01, 0.1, 1, 10 and 100 μ M

426 of the halogenated compounds unichlor, PFOS, Paroil, hexabromocyclododecane, or Aroclor

427 1260. FFN206 fluorescence represented as percent of control (HEK+VMAT2 cells treated with 0

428 μM drug). Graphs depict mean and standard error of the mean and curves represent non-linear
429 regressions.

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