

1 Dual-color GRAB sensors for monitoring spatiotemporal 2 serotonin release *in vivo*

3 Fei Deng^{1,2,7}, Jinxia Wan^{1,2,7}, Guochuan Li^{1,2}, Hui Dong^{1,2}, Xiju Xia^{1,2,3}, Yipan Wang^{1,2},
4 Xuelin Li^{1,2}, Chaowei Zhuang⁴, Yu Zheng^{1,2,5}, Laixin Liu^{1,2,5}, Yuqi Yan^{1,2,5}, Jiesi Feng^{1,2},
5 Yulin Zhao^{1,2}, Hao Xie⁴, Yulong Li^{1,2,3,5,6*}

6

7 ¹State Key Laboratory of Membrane Biology, New Cornerstone Science Laboratory, School
8 of Life Sciences, Peking University, Beijing 100871, China.

9 ²PKU-IDG/McGovern Institute for Brain Research, Beijing 100871, China.

10 ³Peking University–Tsinghua University–National Institute of Biological Sciences Joint
11 Graduate Program, Academy for Advanced Interdisciplinary Studies, Peking University,
12 Beijing, China.

13 ⁴Department of Automation, Tsinghua University, Beijing 100084, China.

14 ⁵Peking-Tsinghua Center for Life Sciences, Academy for Advanced Interdisciplinary
15 Studies, Peking University, Beijing 100871, China.

16 ⁶Chinese Institute for Brain Research, Beijing 102206, China.

17 ⁷These authors contributed equally

18

19 *Manuscript correspondence: Yulong Li (yulongli@pku.edu.cn)

20

21 Abstract

22 (150 words)

23 The serotonergic system plays important roles in both physiological and pathological
24 processes, and is a widely used therapeutic target for many psychiatric disorders. Although
25 several genetically encoded GFP-based serotonin (5-HT) sensors were recently
26 developed, their sensitivities and spectral profiles are relatively limited. To overcome these
27 limitations, we optimized green fluorescent G-protein-coupled receptor (GPCR)-activation-
28 based 5-HT (GRAB_{5-HT}) sensors and developed a new red fluorescent GRAB_{5-HT} sensor.
29 These sensors have excellent cell surface trafficking, high specificity, sensitivity, and
30 spatiotemporal resolution, making them suitable for monitoring 5-HT dynamics *in vivo*.
31 Besides recording subcortical 5-HT release in freely moving mice, we observed both
32 uniform and gradient 5-HT release in the mouse dorsal cortex with mesoscopic imaging.
33 Finally, we performed dual-color imaging and observed seizure-induced waves of 5-HT
34 release throughout the cortex following calcium and endocannabinoid waves. In summary,
35 these 5-HT sensors can offer valuable insights regarding the serotonergic system in both
36 physiological and pathological states.

37

38 **Main**

39 **(3,735 words)**

40 **Introduction**

41 Serotonin (5-HT) is an important monoamine signaling molecule present virtually
42 throughout the body, widely regulating neural activity and other key biological processes¹.
43 In the central nervous system (CNS), 5-HT is an intensively studied neurotransmitter
44 involved in a wide range of neurobiological processes such as emotion, learning and
45 memory, reward, appetite, and the sleep-wake cycle¹⁻³. Moreover, impaired 5-HT
46 transmission is associated with a broad range of CNS disorders, including anxiety,
47 addiction, depression, and epilepsy⁴⁻⁶. As a consequence, many psychotropic and
48 psychedelic drugs have been developed to act on the serotonergic system in the CNS⁷.
49 The primary source of 5-HT in the CNS is serotonergic neurons in the brainstem, which
50 innervate most of the regions throughout the brain to drive various functions; moreover,
51 these neurons are highly heterogeneous with respect to their transcriptomics and
52 projection patterns⁸⁻¹². To encode these widespread 5-HT signals into specific downstream
53 signaling pathways, 15 different 5-HT receptor (5-HTR) subtypes have evolved¹, with half-
54 maximal effective concentration (EC₅₀) values ranging from nanomolar to micromolar¹³.
55 Understanding the serotonergic system in both physiological and pathological processes
56 requires the ability to directly monitor 5-HT dynamics in behaving animals in real time,
57 which in turn requires highly sensitive detection tools. However, given the anatomical and
58 functional complexities of the serotonergic system, classic detection methods such as
59 microdialysis and fast-scan cyclic voltammetry (FSCV) lack the simultaneous high
60 spatiotemporal resolution, specificity, sensitivity, and minimal invasiveness needed for the
61 *in vivo* detection of 5-HT¹⁴⁻¹⁶.

62 Recent advances in genetically encoded fluorescent 5-HT sensors have led to overall
63 optimal tools that surpass classic methods¹⁷⁻²⁰; however, these sensors have not yet hit
64 the proverbial “sweet spot” with respect to balancing apparent affinity with the magnitude
65 of the response. Specifically, sensors based on GPCRs, including GRAB_{5-HT1.0} (ref.¹⁷),
66 PsychLight2 (ref.¹⁸), and sDarken (ref.¹⁹), have high affinity for 5-HT but produce only a
67 modest change in fluorescence. On the other hand, the periplasmic binding protein (PBP)-
68 based sensor iSeroSnFR (ref.²⁰) has a relatively large response, but low affinity to 5-HT.
69 Thus, monitoring 5-HT dynamics *in vivo* requires a more sensitive, high-affinity sensor that
70 produces a sufficiently large response.

71 In the brain, the serotonergic system interacts with other neurotransmitters and
72 neuromodulators²¹; thus, simultaneously imaging 5-HT and other neurochemicals can
73 provide valuable information regarding the regulation of cognitive functions controlled by
74 these signaling processes. Unfortunately, most existing sensors for neurochemicals
75 contain a green fluorescent protein (GFP) as the fluorescent module, as do all genetically
76 encoded 5-HT sensors, precluding combined imaging due to spectral overlap. Although a
77 near-infrared 5-HT nanosensor based on single-wall carbon nanotubes has been
78 reported²², it may not be suitable for use in living animals due to limited sensitivity. On the

79 other hand, red-shifted sensors, such as a previously reported red calcium sensor^{23, 24}, are
80 compatible with other green fluorescent sensors and blue light-exitable actuators, with
81 intrinsically superior optical properties—including deeper tissue penetration, reduced
82 autofluorescence, and low phototoxicity—due to their longer excitation wavelengths. Thus,
83 red-shifted 5-HT sensors suitable for *in vivo* imaging, particularly multiplexed imaging, are
84 urgently needed.

85 Here, we report a series of green and red fluorescent 5-HT sensors generated by
86 transplanting the third intracellular loop (ICL3)—containing circular permuted enhanced
87 GFP (cpEGFP) or cpmApple—from existing green and red GRAB sensors into the 5-HTR4
88 subtype. The green fluorescent sensor, gGRAB_{5-HT3.0} (g5-HT3.0), produces a ~1,300%
89 increase in fluorescence in response to 5-HT, making it vastly superior to existing green
90 fluorescent 5-HT sensors both *in vitro* and *in vivo*. The red-shifted sensor, rGRAB_{5-HT1.0} (r5-
91 HT1.0), produces a >300% increase in fluorescence in response to 5-HT and is also
92 suitable for both *in vitro* and *in vivo* applications. Using mesoscopic imaging in mice
93 expressing g5-HT3.0, we found that 5-HT is released in a gradient along the anterior-to-
94 posterior axis in the mouse dorsal cortex upon optogenetic stimulation of serotonergic
95 neurons in the dorsal raphe nucleus (DRN), but is released with spatial homogeneity during
96 the sleep-wake cycle. Finally, dual-color mesoscopic imaging revealed cortex-wide 5-HT
97 waves that followed calcium and endocannabinoid (eCB) waves in a mouse seizure model.
98 Thus, these improved dual-color GRAB_{5-HT} sensors are powerful tools for monitoring
99 serotonin release *in vivo*, providing valuable new insights into the serotonergic system.

100 Results

101 Development and optimization of new GRAB_{5-HT} sensors

102 To expand the dynamic range and spectral profile of GRAB_{5-HT} sensors, we first
103 systematically searched for the most suitable GPCR scaffold by transplanting the ICL3
104 from existing green and red fluorescent GRAB sensors into a wide range of 5-HT receptor
105 subtypes (Fig. 1a). For green fluorescent sensors, we used green GRAB_{5-HT1.0} (g5-HT1.0)¹⁷
106 or GRAB_{NE1m} (ref.²⁵) as the ICL3 donor (including cpEGFP and surrounding linker
107 sequences). Possibly due to the conserved structures among the class A family of GPCRs,
108 we successfully obtained new candidate sensors that yielded a >50% increase in
109 fluorescence ($\Delta F/F_0$) in response to 10 μ M 5-HT following the screening of replacement
110 sites. One green fluorescent candidate based on 5-HTR4 had a higher response than the
111 original g5-HT1.0 sensor and was named g5-HT1.1. To develop red fluorescent 5-HT
112 sensors, we transplanted the ICL3 (including cpmApple and the linker sequences) from
113 rGRAB_{DA1m} (ref.²⁶) into the 5-HTR2C, 5-HTR4, and 5-HTR6 scaffolds—we chose these
114 three subtypes based on their high performance when developing the green fluorescent 5-
115 HT sensors. Once again, the top candidate was based on the 5-HTR4 subtype, with a ~40%
116 $\Delta F/F_0$; we named this sensor r5-HT0.1 (Fig. 1a).

117 To improve the sensors' sensitivity, then we optimized g5-HT1.1 and r5-HT0.1 by
118 performing saturation mutagenesis of critical residues believed to affect structural coupling,
119 fluorescence intensity²⁷, protein folding²⁸, and the 5-HT induced conformational change²⁹.

120 Iterative optimization by screening more than 4,500 candidates yielded two intermediate

121 green fluorescent sensors called g5-HT2h and g5-HT2m, followed by the final sensor g5-
122 HT3.0 (Fig. 1b,c and Extended Data Figs. 1a–c). When expressed in HEK293T cells, these
123 green fluorescent 5-HT sensors showed good trafficking to the cell surface and a robust
124 increase in fluorescence in response to 5-HT application (Fig. 1c,d and Extended Data Fig.
125 2a–c). In response to 100 μ M 5-HT, the change in fluorescence of g5-HT3.0 ($\Delta F/F_0$:
126 ~1,300%) was 6–20-fold larger than the response measured for previously developed 5-
127 HT sensors, g5-HT1.0, PsychLight2, and iSeroSnFR^{17, 18, 20}. Moreover, the g5-HT3.0
128 sensor is significantly brighter, with a higher signal-to-noise ratio (SNR) than the other
129 green fluorescent sensors (Fig. 1d). Dose-response curves showed that g5-HT3.0
130 produces a larger $\Delta F/F_0$ (i.e., is more sensitive to 5-HT) than previous sensors over a wide
131 range of concentrations, with an EC₅₀ value of ~150 nM (Fig. 1e). We also generated a 5-
132 HT insensitive version of the g5-HT3.0 sensor, named g5-HT3.0mut, by introducing the
133 D131^{3,32}F mutation^{30,31}; we confirmed that g5-HT3.0mut localizes to the plasma membrane
134 but does not respond to 5-HT even at 100 μ M (Fig. 1b–e and Extended Data Fig. 1a,c).

135 Similarly, we generated a red fluorescent 5-HT sensor called r5-HT1.0 by
136 screening >3,000 candidates, and we generated a 5-HT insensitive version, named r5-
137 HTmut, by introducing the D131^{3,32}Q and D149^{3,50}H mutations (Fig. 1b–e and Extended
138 Data Fig. 1d,f). Both r5-HT1.0 and r5-HTmut localized to the plasma membrane when
139 expressed in cultured HEK293T cells (Fig. 1c). Application of 10 μ M 5-HT to cells
140 expressing r5-HT1.0 elicited a ~330% increase in fluorescence, but had no effect on cells
141 expressing r5-HTmut (Fig. 1c,d). The EC₅₀ of r5-HT1.0 was ~790 nM (Fig. 1e). Moreover,
142 unlike the cpmApple-based calcium sensor jRGECO1a—in which blue light causes an
143 increase in fluorescence^{24, 32}—we found that blue light had no detectable effect on r5-
144 HT1.0 (Fig. 1f).

145 Characterization of GRAB_{5-HT} sensors in cultured cells

146 Next, we characterized the pharmacology, specificity, spectra, and kinetics of our new
147 5-HT sensors expressed in HEK293T cells. We found that both g5-HT3.0 and r5-HT1.0
148 inherited the pharmacological specificity of the parent 5-HTR4 receptor, as their 5-HT
149 induced responses were blocked by the 5-HTR4-specific antagonist RS 23597-190 (RS),
150 but not the 5-HTR2C-specific antagonist SB 242084 (SB); in addition, both sensors were
151 unaffected by application of a wide range of signaling molecules, including the 5-HT
152 precursor, 5-HT metabolite, and a variety of other neurotransmitters and neuromodulators
153 (Fig. 2a and Extended Data Fig. 3a,b). The emission spectra of the green fluorescent (with
154 a peak at 520 nm) and red fluorescent (with a peak at 595 nm) sensors are well-separated,
155 and we measured 1-photon/2-photon excitation peaks at 505/920 nm for g5-HT3.0 and
156 560/1050 nm for r5-HT1.0 (Fig. 2b and Extended Data Fig. 4a,b). We also measured a
157 425-nm isosbestic point for g5-HT3.0 under 1-photon excitation. With respect to the
158 sensors' kinetics, we measured the on rate (τ_{on}) by locally puffing 10 μ M 5-HT on the cells
159 and the off rate (τ_{off}) by puffing the 5-HTR4 antagonist RS in the continued presence of 10
160 μ M 5-HT (Fig. 2c,d), revealing sub-second τ_{on} rates and faster τ_{off} rates than our previously
161 reported g5-HT1.0 sensor¹⁷, with mean τ_{off} rates of 1.66 s, 1.90 s, 0.38 s, and 0.51 s for
162 g5-HT3.0, g5-HT2h, g5-HT2m, and r5-HT1.0, respectively (Fig. 2e and Extended Data Fig.
163 4c).

164 We then expressed the g5-HT3.0 and r5-HT1.0 sensors in cultured rat cortical neurons

165 and found that they trafficked well to the cell membrane and were distributed throughout
166 the soma, dendrites, and axon. Application of a saturating concentration of 5-HT induced
167 a fluorescence increase of ~2,700% for g5-HT3.0 and ~400% for r5-HT1.0, but had no
168 effect on neurons expressing g5-HT3.0mut or r5-HTmut (Fig. 2f,g,i,j). We also confirmed
169 the sensors' high specificity for 5-HT when expressed in cultured cortical neurons
170 (Extended Data Fig. 3c–f), and we measured EC₅₀ values of 80 nM, 70 nM, 2.4 μ M, and
171 600 nM for g5-HT3.0, g5-HT2h, g5-HT2m, and r5-HT1.0, respectively (Fig. 2h,k and
172 Extended Data Fig. 4d). Compared to previously reported GFP-based 5-HT sensors^{17, 18,}
173²⁰, g5-HT3.0 is brighter and has a greater fluorescence change and a higher SNR when
174 expressed in cultured neurons (Extended Data Fig. 5).

175 The steric hindrance of the bulky cpFP in sensors is likely to disturb the downstream
176 coupling of GPCR^{17, 25, 33}. To confirm that our 5-HT sensors do not couple to downstream
177 signaling pathways—and therefore do not likely affect cell activity—we used the luciferase
178 complement assay³⁴ and the Tango assay^{35, 36} to measure the GPCR-mediated G_s and β -
179 arrestin pathways, respectively. We found that g5-HT3.0, g5-HT2h, g5-HT2m, and r5-
180 HT1.0 had negligible downstream coupling; in contrast, the wild-type 5-HTR4 receptor had
181 high basal activity and robust, dose-dependent coupling (Fig. 2l and Extended Data Fig.
182 4e,f). In addition, our sensors do not undergo β -arrestin-mediated internalization or
183 desensitization when expressed in cultured neurons, as the 5-HT elicited increase in
184 fluorescence was stable for up to 2 hours in the continuous presence of 10 μ M 5-HT (Fig.
185 2m and Extended Data Fig. 4g).

186 **Measuring endogenous 5-HT release in freely moving mice**

187 To determine whether our newly developed red fluorescent sensor is suitable for *in vivo*
188 imaging in freely behaving mice, we expressed either r5-HT1.0 or r5-HTmut in the basal
189 forebrain (BF), which receives extensive DRN serotonergic projections³⁷, and expressed
190 the light-activated channel ChR2 (ref.^{38, 39}) in serotonergic neurons in the DRN of *Sert-Cre*
191 mice⁴⁰ (Fig. 3a). Optical stimulation of the DRN induced time-locked transient increases in
192 r5-HT1.0 fluorescence, the amplitude of which increased progressively with increasing
193 stimulation duration; moreover, the selective serotonin transporter blocker fluoxetine
194 further increased the amplitude of the response and prolonged the response's decay
195 kinetics (Fig. 3b–e). As expected, no response was measured for the 5-HT insensitive r5-
196 HTmut sensor (Fig. 3b–e).

197 To test whether r5-HT1.0 is compatible with green fluorescent sensors, we expressed
198 either r5-HT1.0 or r5-HTmut in the BF and expressed the axon-targeted green fluorescent
199 calcium sensor axon-GCaMP6s (ref.⁴¹) in DRN serotonergic neurons, which project to the
200 BF and regulate the sleep-wake cycle^{17, 42}. We then performed dual-color fiber photometry
201 recording in the BF while simultaneously recording the electroencephalography (EEG) and
202 electromyography (EMG) signals in order to track the animal's sleep-wake state (Fig. 3f).
203 We found that both the r5-HT1.0 and GCaMP6s signals were higher during both the wake
204 state and NREM (non-rapid eye movement) sleep than during REM sleep (Fig. 3g). In
205 addition, r5-HT1.0 revealed oscillations in 5-HT levels (Fig. 3h). Moreover, we found that
206 the r5-HT1.0 and GCaMP6s signals were temporally correlated, with no detectable lag,
207 revealing the rapid kinetics of 5-HT release and high consistency between 5-HT release
208 and the increase in presynaptic calcium (Fig. 3i). In contrast, the r5-HTmut signal was

209 largely unchanged throughout the sleep-wake cycle and was significantly smaller than the
210 r5-HT1.0 signal during the wake state and NREM sleep (Fig. 3j and Extended Data Fig. 6).
211 Finally, we found no significant difference in the GCaMP6s signal between mice co-
212 expressing r5-HTmut and mice co-expressing r5-HT1.0 (Fig. 3j).

213 To compare the performance of our optimized g5-HT3.0 sensor to previously reported 5-
214 HT sensors, we also performed bilateral recordings in the BF during the sleep-wake cycle
215 in mice expressing g5-HT3.0 in one hemisphere and g5-HT1.0, PsychLight2, or iSeroSnFR
216 in the other hemisphere (Extended Data Fig. 7a,d,g). Consistent with our *in vitro* results,
217 we found that the g5-HT3.0 sensor had significantly larger SNR during the wake state and
218 NREM sleep compared to the other three sensors, as well as more robust oscillations
219 measured during NREM sleep (Extended Data Fig. 7b,c,e,f,h,i).

220 **Mesoscopic imaging of 5-HT dynamics in the mouse dorsal cortex**

221 5-HT also plays important roles in the cerebral cortex, for example regulating cognition
222 and emotion in the prefrontal cortex^{12, 37, 43}. Given that the cortex generally receives sparser
223 serotonergic projections compared to subcortical regions such as the BF⁴⁴, measuring 5-
224 HT release in the cortex requires a highly sensitive 5-HT sensor. To image 5-HT dynamics
225 throughout the whole dorsal cortex, we expressed the g5-HT3.0 sensor by injecting AAV
226 in the transverse sinus⁴⁵. We then measured g5-HT3.0 fluorescence throughout the cortex
227 using mesoscopic imaging^{46, 47} in response to optogenetic stimulation of DRN serotonergic
228 neurons expressing ChrimsonR⁴⁸ (Fig. 4a). We found that light pulses induced transient
229 increases in g5-HT3.0 fluorescence, with increasing stimulation frequency causing
230 increasingly larger responses (Fig. 4b,c). As a negative control, no response was
231 measured when we expressed a membrane-tethered EGFP (memEGFP) (Fig. 4 b,c and
232 Extended Data Fig. 8g). Importantly, we found that treating mice with the SERT blocker
233 fluoxetine caused a gradual increase in baseline 5-HT levels and slowed the decay rate of
234 stimulation-induced transients; in contrast, the dopamine transporter (DAT) blocker GBR
235 12909 had no effect (Fig. 4d and Extended Data Fig. 8a-f).

236 Having shown that g5-HT3.0 can reliably detect 5-HT release in the cortex in response
237 to optogenetic stimulation, we then used this sensor to measure physiologically relevant
238 5-HT dynamics during the sleep-wake cycle using mesoscopic imaging combined with
239 simultaneous EEG and EMG recordings. Similar to our results measured in the subcortical
240 BF (Extended Data Fig. 7), we found that the g5-HT3.0 signal in the dorsal cortex was
241 highest during the wake state, followed by the NREM and REM states, with visible
242 oscillations during NREM sleep. In addition, we found no change in fluorescence through
243 the sleep-wake cycle using the 5-HT insensitive g5-HT3.0mut sensor (Fig. 4e,f).

244 Taking advantage of our ability to visualize the entire dorsal cortex using mesoscopic
245 imaging, we segmented the dorsal cortex into various brain regions based on the Allen
246 Common Coordinate Framework v3 (CCFv3) atlas⁴⁹ and analyzed the spatial distribution
247 of the 5-HT signals during optogenetic stimulation and during the sleep-wake cycle. We
248 found that the 5-HT signals measured in different brain regions were relatively spatially
249 homogenous and temporally synchronized during the sleep-wake cycle (Fig. 4g and
250 Extended Data Fig. 8h), reminiscent of our previous results recorded in subcortical regions,
251 including the orbital frontal cortex and the bed nucleus of the stria terminalis¹⁷. In contrast,
252 when we optogenetically stimulated DRN serotonergic neurons, the 5-HT signals had a

253 graded pattern, decreasing along the anterior-to-posterior axis (Fig. 4g,h); interestingly,
254 this pattern was consistent with the anatomically heterogeneous density of serotonergic
255 projections throughout the cortex⁴⁴.

256 Taken together, these results demonstrate that our next-generation g5-HT3.0 sensor is
257 sufficiently sensitive to monitor 5-HT release *in vivo* with high spatiotemporal resolution,
258 revealing key differences between specific brain regions.

259 **Dual-color *in vivo* imaging reveals cortex-wide neurochemical waves during seizure
260 activity**

261 The serotonergic system has been suggested to protect the CNS from epileptiform
262 activity⁵⁰⁻⁵⁴, which is characterized by excessive and hypersynchronous neuronal firing.
263 However, little is known regarding the spatiotemporal dynamics of 5-HT release during and
264 after seizure activity, let alone the relationship between 5-HT and other seizure-related
265 signals such as calcium^{55, 56} and endocannabinoid (eCB) levels⁵⁷. Therefore, we performed
266 dual-color mesoscopic imaging of g5-HT3.0 together with jRGECO1a (ref.²⁴) (to measure
267 both 5-HT and calcium) or r5-HT1.0 together with eCB2.0 (ref.⁵⁷) (to measure both 5-HT
268 and eCBs) in the mouse dorsal cortex, while simultaneously performing EEG recording to
269 identify seizures induced by an injection of the glutamate receptor agonist kainic acid
270 (KA)⁵⁸ (Fig. 5a). Similar to previous reports^{57, 59}, we observed an increase in Ca²⁺ during
271 the KA-induced seizure, followed by a spreading wave of Ca²⁺ with a larger magnitude. In
272 the same mouse, we observed a spreading wave of 5-HT, reported by g5-HT3.0 signals,
273 that closely followed the Ca²⁺ wave (Fig. 5b,c,f, Extended Data Fig. 9a,b and
274 Supplementary Video 1). The waves reported by g5-HT3.0 and jRGECO1a originated in
275 approximately the same location and propagated with similar speed (at ~76 μm/s and ~83
276 μm/s, respectively) and in the same direction, primarily from the lateral cortex to the medial
277 region (Fig. 5g,h). As a negative control, seizure activity had no effect on the signal
278 measured using g5-HT3.0mut (Fig. 5c,f and Extended Data Fig. 9a,b).

279 Finally, we obtained similar results in mice co-expressing r5-HT1.0 and eCB2.0, with a
280 propagating wave of 5-HT release following the eCB wave. Moreover, the waves reported
281 by r5-HT1.0 and eCB2.0 originated in approximately the same location, propagated at
282 similar speed (~83 μm/s and ~81 μm/s, respectively) and in the same direction (Fig. 5d,f–
283 h and Supplementary Video 2). As above, seizure activity had no effect on the signal
284 measured using r5-HTmut (Fig. 5e,f and Extended Data Fig. 9c,d).

285 Taken together, these results demonstrate that our g5-HT3.0 and r5-HT1.0 sensors can
286 reliably report 5-HT release *in vivo* with high sensitivity, specificity, and spatiotemporal
287 resolution both under physiological conditions and during seizure activity.

288 **Discussion**

289 Here, we report the development, optimization, characterization, and *in vivo* application
290 of a series of genetically encoded red and green fluorescent 5-HT sensors. These 5-HT
291 sensors have high specificity, sensitivity, and spatiotemporal resolution, as well as rapid
292 kinetics. More importantly, they do not couple to downstream signal pathways. Both the
293 g5-HT3.0 and r5-HT1.0 sensors reliably reported time-locked 5-HT release induced by
294 optogenetic activation of DRN serotonergic neurons, as well as changes in 5-HT levels

295 during the sleep-wake cycle. Moreover, using dual-color mesoscopic imaging, we found
296 that seizure activity triggers a cortex-wide 5-HT wave that follows a wave of Ca^{2+} and eCBs.

297 GRAB sensors consist of a ligand-binding module and a fluorescent reporting module⁶⁰.
298 Our ability to efficiently develop new GRAB_{5-HT} sensors was facilitated by our
299 transplantation strategy and our selection of the 5-HT₄ receptor as the scaffold. First, we
300 used the transplantation strategy to capitalize on previously optimized reporting modules
301 and linkers in existing sensors. Next, we chose the best-performing scaffold (5-HT₄) from
302 among the various 5-HT receptor subtypes as the ligand-binding module, thereby
303 accelerating the optimization processes. A similar strategy can be used in order to
304 accelerate the development of other GPCR-based sensors.

305 Our next-generation green fluorescent GRAB_{5-HT} sensors have three advantages over
306 previously reported genetically encoded fluorescent 5-HT sensors. First, g5-HT3.0 is
307 significantly more sensitive than other GPCR- and PBP-based sensors due to its larger
308 response and brightness, making it more suitable for both *in vitro* and *in vivo* applications.
309 Second, our high-affinity g5-HT3.0 (EC_{50} : ~150 nM) and medium-affinity g5-HT2m (EC_{50} :
310 ~1.1 μM) sensors fill the critical gap in measuring intermediate concentrations of 5-HT.
311 Thus, together with previous sensors such as g5-HT1.0 (EC_{50} : ~20 nM) and iSeroSnFR
312 (EC_{50} : >300 μM), we now possess a powerful toolbox covering a wide range of
313 physiological and pathological 5-HT concentrations⁶¹⁻⁶³. Third, these sensors have rapid
314 kinetics and are suitable for tracking transient 5-HT release in real time.

315 More importantly, our all-new red fluorescent 5-HT sensor expands the spectral profile
316 of genetically encoded 5-HT sensors and is suitable for dual-color imaging when combined
317 with green fluorescent sensors. Here, we simultaneously recorded red and green
318 fluorescence using r5-HT1.0 and axon-targeted GCaMP6s, respectively, in the mouse
319 basal forebrain during the sleep-wake cycle and found that the two signals were closely
320 correlated, consistent with Ca^{2+} -dependent rapid 5-HT release. In the future, further
321 expanding the spectra of 5-HT sensors (for example, to include the far-red and near-
322 infrared spectra) will provide the ability to simultaneously monitor multiple signals in
323 addition to 5-HT.

324 Using mesoscopic imaging, we found that 5-HT release in the mouse dorsal cortex
325 differs under different behavioral conditions. A previous study using a calcium sensor
326 suggested that 5-HT release might differ between the orbitofrontal cortex and central
327 amygdala in response to rewarding and aversive stimuli, but the 5-HT release was not
328 measured directly³⁷. Recently, using g5-HT1.0, we showed that 5-HT release is highly
329 synchronized between different brain regions during NREM sleep¹⁷. Here, we expanded
330 on this finding using mesoscopic imaging of g5-HT3.0 to image more brain regions in the
331 mouse dorsal cortex and found relatively homogenous release throughout the dorsal cortex
332 during the sleep-wake cycle. Moreover, we found a graded pattern of 5-HT release in the
333 mouse dorsal cortex evoked by optogenetic activation of the DRN, similar to the pattern of
334 serotonergic projections. These results suggest that although the potential 5-HT release is
335 spatially heterogeneous throughout the dorsal cortex, this release may be regulated on a
336 global scale during the sleep-wake cycle, leading to relatively homogenous 5-HT release
337 under these conditions. Furthermore, using a KA-induced seizure model, we found that 5-
338 HT release propagates as a wave across the cortex, a result made possible due to the

339 advantages of our new 5-HT sensors compared to traditional methods used to measure 5-
340 HT. Indeed, a previous study using microdialysis found increased 5-HT levels after seizures
341 in rats, but the results had relatively poor temporal resolution and lacked spatial resolution⁵⁴.
342 Interestingly, we further found that the seizure-induced 5-HT waves were spatially
343 correlated—but lagged behind—Ca²⁺ and eCB waves, consistent with the idea that
344 serotonergic activity may protect against neuronal hyperactivity^{51, 53, 64}.

345 In summary, our new 5-HT sensors can be used to monitor 5-HT release both *in vitro*
346 and *in vivo*, with high sensitivity and spatiotemporal resolution. Thus, when combined with
347 advanced imaging techniques, these new 5-HT sensors provide a robust toolbox to study
348 the serotonergic system in both health and disease.

349

350 **Methods**

351 **Molecular biology**

352 DNA fragments were amplified by PCR using primers (RuiBiotech) with 25–30-bp
353 overlap and used to generate plasmids via the Gibson assembly method⁶⁵ using T5
354 exonuclease (New England Biolabs), Phusion DNA polymerase (Thermo Fisher Scientific),
355 and Taq ligase (iCloning). Plasmid sequences were verified by Sanger sequencing
356 (RuiBiotech). For the replacement site screening, cDNAs encoding 12 different 5-HTR
357 subtypes (5-HTR_{1A}, 5-HTR_{1B}, 5-HTR_{1D}, 5-HTR_{1E}, 5-HTR_{1F}, 5-HTR_{2A}, 5-HTR_{2B}, 5-HTR_{2C},
358 5-HTR₄, 5-HTR_{5A}, 5-HTR₆, and 5-HTR₇) were cloned by PCR amplification of the full-length
359 human GPCR cDNA library (hORFeome database 8.1) or the PRESTO-Tango GPCR Kit³⁶
360 (Addgene Kit #1000000068). To optimize the 5-HT sensors, cDNAs encoding candidate
361 sensors were cloned into the pDisplay vector (Invitrogen) with an IgK leader sequence in
362 the sensor upstream, and either IRES-mCherry-CAAX (for green fluorescent 5-HT sensors)
363 or IRES-EGFP-CAAX (for red fluorescent 5-HT sensors) was fused downstream of the
364 sensor to calibrate the membrane signal. Site-directed mutagenesis was performed using
365 primers containing randomized NNB codons (48 codons in total, encoding 20 possible
366 amino acids) or defined codons at the target sites. For expression in cultured neurons, the
367 sensors were cloned into the pAAV vector under the control of the hSyn promoter. To
368 generate stable cell lines for measuring the excitation/emission spectra, sequences
369 encoding various 5-HT sensors were cloned into a vector called pPacific, containing a 3'
370 terminal repeat, IRES, the puromycin gene, and a 5' terminal repeat. Two mutations
371 (S103P and S509G) were introduced into pCS7-PiggyBAC to generate hyperactive
372 piggyBac transposase (ViewSolid Biotech)⁶⁶. To measure downstream coupling using the
373 Tango assay, DNA encoding various GRAB_{5-HT} sensors or wild-type 5-HTR4 was cloned
374 into the pTango vector³⁶. For the luciferase complementation assay, the β₂AR gene in the
375 β2AR-Smbit construct³⁴ was replaced with the indicated GRAB_{5-HT} sensors or wild-type 5-
376 HTR4; LgBit-mGs was a generous gift from Nevin A. Lambert (Augusta University).

377 **Cell lines**

378 HEK293T cells were purchased from ATCC (CRL-3216) and verified based on their
379 morphology and growth rate. Stable cell lines expressing different GRAB_{5-HT} sensors were
380 generated by co-transfected HEK293T cells with the pPacific plasmids encoding sensors
381 and the pCS7-PiggyBAC plasmid encoding the transposase⁶⁶. Cells expressing the
382 desired genes were selected using 2 µg/ml puromycin (Sigma). An HTLA cell line stably
383 expressing a tTA-dependent luciferase reporter and the β-arrestin2-TEV fusion gene used
384 in the Tango assay³⁶ was a generous gift from Bryan L. Roth (University of North Carolina
385 Chapel Hill). All cell lines were cultured at 37°C in 5% CO₂ in DMEM (Biological Industries)
386 supplemented with 10% (v/v) fetal bovine serum (GIBCO) and 1% penicillin-streptomycin
387 (GIBCO).

388 **Primary cultures**

389 Rat cortical neurons were prepared using postnatal day 0 (P0) Sprague-Dawley rat pups
390 (both sexes) purchased from Beijing Vital River. The cerebral cortex was dissected, and

391 neurons were dissociated using 0.25% trypsin-EDTA (GIBCO), plated onto 12-mm glass
392 coverslips coated with poly-D-lysine (Sigma-Aldrich), and cultured in neurobasal medium
393 (GIBCO) containing 2% B-27 supplement (GIBCO), 1% GlutaMax (GIBCO), and 1%
394 penicillin-streptomycin (GIBCO) at 37°C in humidified air containing 5% CO₂.

395 **Animals**

396 All procedures involving animals were performed using protocols approved by the Animal
397 Care and Use Committee at Peking University. *Sert-Cre* mice were generously provided
398 by Yi Rao at Peking University. All mice were group-housed or pair-housed in a
399 temperature-controlled room (18–23°C) with a 12-h/12-h light/dark cycle, with food and
400 water available ad libitum.

401 **Cell transfection and imaging**

402 HEK293T cells were plated either on 12-mm glass coverslips in 24-well plates or 96-well
403 plates without coverslips and grown to ~70% confluence for transfection with PEI (1 µg
404 plasmid and 3 µg PEI per well in 24-well plates or 300 ng plasmids and 900 ng PEI per
405 well in 96-well plates); the medium was replaced after 4–6 h, and the cells were used for
406 imaging 24–48 h after transfection. After 5–9 d of *in vitro* culture, rat cortical neurons were
407 infected with AAVs expressing the following 5-HT sensors: g5-HT3.0 (5.72×10¹² vg/ml,
408 BrainVTA), g5-HT2m (2.53×10¹² vg/ml, BrainVTA), g5-HT2h (2.67×10¹² vg/ml, BrainVTA),
409 g5-HT1.0 (AAV9-hSyn-tTA, 2.39×10¹³ vg/ml, and AAV9-TRE-5-HT1.0, 3.81×10¹³ vg/ml,
410 Vigene Biosciences), iSeroSnFR (2.76×10¹² vg/ml, BrainVTA), PsychLight2 (3.07×10¹²
411 vg/ml, BrainVTA), g5-HT3.0mut (6.16×10¹³ vg/ml, Vigene Biosciences), r5-HT1.0
412 (9.44×10¹³ vg/ml, Vigene Biosciences), or r5-HTmut (1.17×10¹⁴ vg/ml, Vigene
413 Biosciences). To compare the performance of various sensors, the volume of each AAV
414 was adjusted by titer to apply the same number of AAV particles.

415 Before imaging, the culture medium was replaced with Tyrode's solution consisting of (in
416 mM): 150 NaCl, 4 KCl, 2 MgCl₂, 2 CaCl₂, 10 HEPES, and 10 glucose (pH 7.4).

417 Cells were imaged using an inverted Ti-E A1 confocal microscope (Nikon) or an Opera
418 Phenix high-content screening system (PerkinElmer). The confocal microscope was
419 equipped with a 10x/0.45 NA (numerical aperture) objective, a 20x/0.75 NA objective, a
420 40x/1.35 NA oil-immersion objective, a 488-nm laser, and a 561-nm laser; the GFP signal
421 was collected using a 525/50-nm emission filter combined with the 488-nm laser, and the
422 RFP signal was collected using a 595/50-nm emission filter combined with the 561-nm
423 laser. The Opera Phenix system was equipped with a 20x/0.4 NA objective, a 40x/1.1 NA
424 water-immersion objective, a 488-nm laser, and a 561-nm laser; the GFP and RFP signals
425 were collected using a 525/50-nm and 600/30-nm emission filter, respectively. The
426 fluorescence signals produced by the green and red fluorescent GRAB_{5-HT} sensors were
427 calibrated using mCherry (the GFP/RFP ratio) or EGFP (the RFP/GFP ratio), respectively.

428 To measure the sensor responses induced by various chemicals, solutions containing
429 the indicated concentrations of 5-HT and/or other chemicals were administered to the cells
430 via a custom-made perfusion system or via bath application. To measure the sensors'
431 kinetics, a glass pipette was positioned in close proximity to cells expressing the sensors,

432 and the fluorescence signal was measured using confocal high-speed line scanning mode
433 with a scanning speed of 1024 Hz. To measure the on-rate constant (τ_{on}), 100 μ M 5-HT
434 was puffed from the pipette, and the increased trace in fluorescence was fitted with a
435 single-exponential function; to measure the off-rate constant (τ_{off}), 100 μ M RS 23597-190
436 was puffed on cells bathed in 10 μ M 5-HT, and the decreased trace in fluorescence was
437 fitted with a single-exponential function. To test for blue light-mediated photoactivation of
438 the red fluorescent sensors, a 488-nm laser lasting for 1 s was utilized (power: ~210 μ W,
439 intensity: ~0.4 W/cm²).

440 **Spectra measurements**

441 For 1-photon spectra, HEK293T cells stably expressing g5-HT3.0, g5-HT2m, g5-HT2h,
442 or r5-HT1.0 were harvested and transferred to a 384-well plate in the absence or presence
443 of 10 μ M 5-HT. Excitation and emission spectra were measured at 5-nm increments with a
444 20-nm bandwidth using a Safire2 multi-mode plate reader (Tecan). To obtain background
445 for subtraction, control cells (not expressing a sensor) were prepared to the same density
446 as the cells that expressed sensors, and were measured using the same protocol.

447 For 2-photon spectra, HEK293T cells expressing g5-HT3.0, g5-HT2m, g5-HT2h, or r5-
448 HT1.0) were cultured on 12-mm coverslips and placed in an imaging chamber under a 2-
449 photon microscope. For green fluorescent sensors, 2-photon excitation spectra were
450 measured at 10-nm increments ranging from 690 to 1030 nm using a 2-photon microscope
451 (Bruker) equipped with a 20x/1.0 NA water-immersion objective (Olympus) and an InSight
452 X3 tunable laser (Spectra-Physics). For the red fluorescent sensor, 2-photon excitation
453 spectra were measured at 10-nm increments ranging from 820 to 1300 nm using an A1R
454 MP+ multiphoton microscope (Nikon) equipped with a 25x/1.1 NA objective (Nikon) and a
455 Chameleon Discovery tunable laser (Coherent). Laser power was calibrated for various
456 wavelengths.

457 **Luciferase complementation assay**

458 The luciferase complementation assay was performed as previously described³⁴. In brief,
459 24–48 h after transfection, the cells were washed with PBS, dissociated using a cell scraper,
460 resuspended in PBS, transferred to opaque 96-well plates containing 5 μ M furimazine
461 (NanoLuc Luciferase Assay, Promega), and bathed in 5-HT at various concentrations
462 (ranging from 0.1 nM to 1 mM). After incubation for 10 minutes in the dark, luminescence
463 was measured using a VICTOR X5 multilabel plate reader (PerkinElmer).

464 **Tango assay**

465 A reporter cell line called HTLA, stably expressing a tTA-dependent luciferase reporter
466 and a β -arrestin2-TEV fusion gene, was transfected with pTango vectors to express
467 GRAB_{5-HT} sensors or wild-type 5-HTR4. After culturing for 24 h in 6-well plates, the cells
468 were transferred to 96-well plates and bathed with 5-HT at varying concentrations (ranging
469 from 0.01 nM to 100 μ M). The cells were then cultured for 12 h to allow the expression of
470 tTA-dependent luciferase. Bright-Glo reagent (Fluc Luciferase Assay System, Promega)
471 was added to a final concentration of 5 μ M, and luminescence was measured using a
472 VICTOR X5 multilabel plate reader (PerkinElmer).

473 **Fiber photometry recording of 5-HT release *in vivo***

474 To express the red fluorescent 5-HT sensors in the BF, adult *Sert-Cre* mice were
475 anesthetized with 1.5% isoflurane and placed on a stereotaxic frame (RWD Life Science).
476 AAV expressing hSyn-r5-HT1.0 (9.44×10^{13} vg/ml, Vigene Biosciences) or hSyn-r5-HTmut
477 (1.14×10^{14} vg/ml, Vigene Biosciences) was injected (400 nl per site) via a glass pipette
478 using a micro-syringe pump (Nanoliter 2000 Injector, World Precision Instruments) at the
479 following coordinates: AP: 0 mm relative to Bregma; ML: +1.5 mm; DV: 4.6 mm below the
480 dura. For optical activation of the DRN, 400 nl AAV9-EF1a-DIO-hChR2(H134R)-EYFP
481 (5.54×10^{13} vg/ml, BrainVTA) was injected into the DRN at the following coordinates: AP: -
482 4.1 mm relative to Bregma; ML: +1.1 mm; depth: 2.9 mm below the dura; at a 20° ML
483 angle). Two optical fiber cannulas (200 μ m, 0.37 NA, Inper, Zhejiang, China) were then
484 implanted; one cannula was implanted 0.1 mm above the virus injection site in the BF to
485 record the 5-HT sensors, and the other cannula was implanted 0.3 mm above the virus
486 injection site in the DRN for optically activating ChR2. The optical fibers were affixed to the
487 skull surface using dental cement. The fiber photometry system (Inper Tech, Zhejiang,
488 China) was used to record the fluorescence signals in freely moving mice. Yellow light-
489 emitting diode (LED) light was bandpass filtered (561/10 nm), reflected by a dichroic mirror
490 (495 nm), and then focused using a 20x objective lens (Olympus). An optical fiber was
491 used to guide the light between the commutator and the implanted optical fiber cannulas.
492 The excitation light emitted by the LED was adjusted to 20–30 μ W and delivered at 10 Hz
493 with a 20-ms pulse duration. The optical signals were then collected through the optical
494 fibers. Red fluorescence was bandpass filtered (520/20 nm and 595/30 nm) and collected
495 using an sCMOS camera. To induce ChR2-mediated 5-HT release, pulse trains (10-ms
496 pulses at 50 Hz for 1 s, 5 s, or 10 s) were delivered to the DRN using a 488-nm laser at 20
497 mW with a 5-min inter-stimulus interval. To test the effects of fluoxetine on the ChR2-
498 induced responses, three optical stimulation trains were applied at a 5-min interval. Then,
499 10 mg/kg fluoxetine was administered via i.p. injection; 30 min after injection, three optical
500 stimulation trains were applied.

501 The current output from the photomultiplier tube was converted to a voltage signal using
502 a model 1700 differential amplifier (A-M Systems) and passed through a low-pass filter.
503 The analog voltage signals were then digitized using an acquisition card (National
504 Instruments). To minimize autofluorescence of the optical fibers, the recording fibers were
505 photobleached using a high-power LED before recording. Background autofluorescence
506 was recorded and subtracted from the recorded signals in the subsequent analysis.

507 The photometry data were analyzed using a custom program written in MATLAB. To
508 calculate $\Delta F/F_0$ during the optogenetics experiments, a baseline was measured before
509 optical stimulation.

510 **Fiber photometry recording of 5-HT dynamics during the sleep-wake cycle**

511 Adult wild-type C57BL/6 mice and *Sert-Cre* mice were anesthetized with isoflurane and
512 placed on a stereotaxic frame (RWD Life Science). In Extended Data Fig. 7a–c, AAV
513 expressing CAG-g5-HT1.0 (3.16×10^{12} vg/ml) was injected into the BF in one hemisphere
514 and AAV expressing CAG-g5-HT3.0 (3.16×10^{12} vg/ml) was injected into the BF in another

515 hemisphere (400 nl per site) using the coordinates described above. In Extended Data Fig.
516 7d–f, AAV expressing hSyn-PsychLight2 (3.07×10^{12} vg/ml) and hSyn-g5-HT3.0 (5.72×10^{12}
517 vg/ml, diluted to 3.07×10^{12} vg/ml) were injected into the bilateral BF (400 nl per site),
518 respectively. In Extended Data Fig. 7g–i, AAV expressing CAG-iSeroSnFR (2.15×10^{12}
519 vg/ml) and CAG-5-HT3.0 (3.16×10^{12} vg/ml, diluted to 2.15×10^{12} vg/ml) were injected into
520 the bilateral BF (400 nl per site), respectively. In Extended Data Fig. 7, all AAVs were
521 produced by BrainVTA and mice were wild-type. In Fig. 3f–g, AAV expressing hSyn-r5-
522 HT1.0 (9.44×10^{13} vg/ml, Vigene Biosciences) or hSyn-r5-HTmut (1.14×10^{14} vg/ml, Vigene
523 Biosciences) was injected into the BF, and EF1α-DIO-axon-GCaMP6s (5.71×10^{12} vg/ml,
524 BrainVTA) was injected into the DRN (400 nl per site) of *Sert-Cre* mice. An optical fiber
525 cannula (200 μm, 0.37 NA, Inper, Zhejiang, China) was implanted 0.1 mm above the virus
526 injection sites in BF for recording the signals of 5-HT sensors and calcium sensor.

527 To record the animal's sleep-wake state, we attached and fixed custom-made EEG and
528 EMG electrodes to the skull via a microconnector. EEG electrodes were implanted into
529 craniotomy holes situated over the frontal cortex and visual cortex, and EMG wires were
530 placed bilaterally in the neck musculature. The microconnector was attached to the skull
531 using glue and a thick layer of dental cement. After surgery, the mice were allowed to
532 recover for at least 2 weeks.

533 The same fiber photometry system (Inper Tech, Zhejiang, China) was used to record the
534 fluorescence signals in freely moving mice during the sleep-wake cycle. In Extended Data
535 Fig. 7, a 10-Hz 470/10-nm filtered light (20–30 μW) was used to excite green fluorescent
536 5-HT sensors, and a 520/20-nm and 595/30 nm dual-band bandpass filter was used to
537 collect the fluorescence signals. In Fig. 3f,g, a 10-Hz 470/10-nm filtered light (20–30 μW)
538 was used to excite the green fluorescent calcium sensor, and a 561/10-nm filtered light
539 (20–30 μW) was used to excite the red fluorescent 5-HT sensors. Fluorescence signals
540 were collected using a dual-band bandpass filter (520/20 nm and 595/30 nm), with
541 excitation light delivered as 20-ms pulses at 10 Hz.

542 Photometry data were analyzed using a custom MATLAB program. To calculate the z-
543 score during the sleep-wake cycle, baseline values were measured during a period of REM
544 sleep in which no apparent fluctuations were observed.

545 **Polysomnographic recording and analysis**

546 The animal's sleep-wake state was determined using the EEG and EMG recordings. The
547 EEG and EMG signals were amplified (NL104A, Digitimer), filtered (NL125/6, Digitimer) at
548 0.5–100 Hz (EEG) and 30–500 Hz (EMG), and then digitized using a Power1401 digitizer
549 (Cambridge Electronic Design Ltd.). The Spike2 software program (Cambridge Electronic
550 Design Ltd.) was used for recording with a sampling rate of 1000 Hz. The animal's sleep-
551 wake state was classified semi-automatically in 4-s epochs using AccuSleep⁶⁷ and then
552 validated manually using a custom-made MATLAB GUI. The wake state was defined as
553 desynchronized EEG activity combined with high EMG activity. NREM sleep was defined
554 as synchronized EEG activity with high-amplitude delta activity (0.5–4 Hz) and low EMG
555 activity. REM sleep was defined as high-power theta frequencies (6–9 Hz) combined with
556 low EMG activity.

557 **Mesoscopic *in vivo* imaging**

558 To express the sensors throughout the cortex, we injected the indicated AAVs into the
559 transverse sinus as described previously⁴⁵. In detail, P0–P1 C57BL/6 mouse pups were
560 removed from their home cages, placed on a warm pad, anesthetized on ice for 2–3 min,
561 and fixed on an ice cooled metal plate. Two small incisions were then made over the
562 transverse sinuses for AAV injection using a glass pipette. The pups were injected
563 bilaterally with the following pairs of AAVs (6 μ l total volume): AAV9-hSyn-5-HT3.0
564 (8.67 \times 10¹³ vg/ml, Vigene Biosciences) and AAV9-hSyn-GAP43-jRGECO1a (1.47 \times 10¹³
565 vg/ml, OBiO); AAV9-hSyn-r5-HT1.0 (9.44 \times 10¹³ vg/ml, Vigene Biosciences) and AAV9-
566 hSyn-eCB2.0 (9.11 \times 10¹³ vg/ml, Vigene Biosciences); AAV9-hSyn-5-HT3.0mut (6.16 \times 10¹³
567 vg/ml, Vigene Biosciences) and AAV9-hSyn-GAP43-jRGECO1a; or AAV9-hSyn-r5-HTmut
568 (1.17 \times 10¹⁴ vg/ml, Vigene Biosciences) and AAV9-hSyn-eCB2.0. The AAVs were injected
569 at a rate of 1.2 μ l/min, and the pipette was left in the sinus for at least 30 s. After injection,
570 the incisions were sealed with Vetbond glue (3M Animal Care Products) and the pups were
571 placed on a warm pad for recovery. After recovery, the pups were gently rubbed with
572 bedding and returned to their home cage.

573 About eight weeks after AAV injection, surgery was performed for implanting the imaging
574 window and the EEG and EMG electrodes. Anesthesia was induced with an i.p. injection
575 of 2,2,2-tribromoethanol (Avertin, 500 mg/kg, Sigma-Aldrich) and maintained using
576 inhalation with 1% isoflurane. The mouse was fixed in a stereotaxic frame, 2% lidocaine
577 hydrochloride was injected under the scalp, and the eyes were covered with erythromycin
578 ophthalmic ointment for protection. Part of the scalp above the skull and the underlying
579 muscles were removed and cleaned carefully to expose the skull. Most of the skull above
580 the dorsal cortex was then carefully removed and replaced with a flat custom-made
581 coverslip (D-shape, ~8 mm \times 8 mm) to create an optical window. EEG and EMG electrodes
582 were implanted and fixed as described above. After surgery, the mice were returned to
583 their home cage for at least 7 d to recover, and then fixed to the base for over 3 d to
584 habituate before imaging until the mouse can fall asleep (especially REM sleep) within the
585 first 3 h. To optically activate the DRN, we used *Sert-Cre* mice as described above, except
586 before the surgery, 300 nl AAV9-EF1a-DIO-ChrimsonR-iP2A-Halotag9-V5 (6.81 \times 10¹³
587 vg/ml, Vigene Biosciences) was injected into the DRN using the following coordinates: AP:
588 –6.1 mm relative to Bregma; ML: 0 mm relative to Bregma; depth: 3 mm below the dura;
589 at a 32° AP angle (to avoid the imaging window, the fiber is inserted forward and down
590 from the back of the interparietal bone). An optical fiber cannula (200 μ m, 0.37 NA, Inper,
591 Zhejiang, China) was then implanted 0.2 mm above the virus injection site and affixed to
592 the skull surface using dental cement.

593 Mesoscopic imaging was performed using a custom-made dual-color macroscope
594 equipped with a 2x/0.5 NA objective lens (Olympus, MVPLAPO2XC), two 1x/0.25 NA tube
595 lenses (Olympus, MVPLAPO1X), and two sCMOS cameras (Andor, Zyla 4.2 Plus,
596 2,048 \times 2,048 pixels, 16-bit)^{68, 69}. Three excitation wavelengths (405 nm, 488 nm, and 561
597 nm) were generated using a multi-line fiber coupled laser system (Changchun New
598 Industries Optoelectronics Tech. Co., Ltd., RGB-405/488/561/642nm-220mW-CC32594).
599 The emission light was passed through a 567-nm cut-on longpass dichroic mirror (Thorlabs,

600 DMLP567L), then through either a 525/36-nm or 609/34-nm emission filter (Chroma), and
601 finally into the sCMOS cameras (one for the green channel and one for the red channel).
602 Both the excitation laser and the camera imaging were triggered by an Arduino board (Uno)
603 with custom-written programs. For green sensor imaging, a 488-nm laser was used and
604 interleaved with a 405-nm laser; while for dual-color imaging, 488-nm and 561-nm lasers
605 were simultaneously generated and interleaved with a 405-nm laser. Images were acquired
606 using Micro-Manager 2.0 at a resolution of 512×512 pixels after $4\times$ pixel binning, and each
607 channel was acquired at either 1 Hz or 5 Hz with 40-ms exposure.

608 During imaging, the mice were head-fixed to the base and could freely run on a
609 treadmill⁷⁰. An infrared camera equipped with LEDs was placed around the mouse's head
610 to capture its face and eyes for recording behavioral data. To induce ChrimsonR-mediated
611 5-HT release in the optogenetics experiment, pulse trains of light (10-ms pulses for 10 s at
612 5, 10, 20, 30, 40, or 50 Hz) were applied to the DRN using a 635-nm laser at 10 mW with
613 a 3-minute interval, and three trials were performed at each frequency. To test the effects
614 of fluoxetine and GBR 12909 on ChrimsonR-evoked 5-HT release, three trains of optical
615 stimulation (10-ms pulses for 1 s at 20 Hz, 10-ms pulses for 10 s at 20 Hz, or 10-ms pulses
616 for 10 s at 50 Hz) were applied with a 5-min interval, and three trials were conducted for
617 each parameter. Then, 10 mg/kg GBR 12909 was injected i.p., and the same optical
618 stimulation procedure was performed 30 min later. Next, 20 mg/kg fluoxetine was injected
619 i.p., followed by the same optical stimulation procedure 30 min later (see Extended Data
620 Fig. 8a). For the seizure experiments, another infrared camera was hung above the back
621 of the mouse to record its behavioral data. After recording ~ 1 h of baseline data, 10 mg/kg
622 KA was injected i.p. to induce seizures. All recordings, including mesoscopic imaging, EEG
623 and EMG recording, optical stimulation trains, and the infrared cameras, were
624 synchronized using a Power1401 acquisition board (Cambridge Electronic Design Ltd.).

625 **Analysis of mesoscopic imaging data**

626 **Preprocessing.** Raw images acquired by each camera were calibrated for the uniformity
627 of the imaging system, and movement-related artifacts were corrected using the motion-
628 correction algorithm NoRMCorre (ref.⁷¹). The corrected image stack with a size of
629 512×512 pixels was downsampled by a factor of 0.7 to 359×359 pixels for subsequent
630 analysis. For dual-color imaging, the averaged red-channel image was registered to the
631 averaged green-channel image via automatic transformation using the MATLAB function
632 “imregtform” with the “similarity” mode. The same geometric transformation was applied to
633 all red-channel images to register to corresponding green-channel images. The image
634 stack was saved as a binary file to accelerate the input and output of large files (typically >8
635 GB). To remove pixels belonging to the background and blood vessels (particularly large
636 veins), we generated a mask for further analysis of the pixels. Specifically, the outline of
637 the entire dorsal cortex in the field of view was generated manually, and pixels outside the
638 outline were set as background and excluded from further analysis. The blood vessels
639 were then removed from the image using the machine learning-based ImageJ plugin
640 Trainable Weka Segmentation⁷² (v3.3.2) in order to minimize artifacts caused by the
641 constriction and dilation of blood vessels. The final mask without the background and blood
642 vessel pixels was applied to the image stack for further analysis.

643 **Spectral unmixing.** For simultaneous dual-color imaging, the bleed-through of
644 fluorescence intensity for each pixel between the green and red channels was removed
645 using linear unmixing based on the spectra of the various sensors and the setup of the
646 microscope system.

647 **Hemodynamic correction and response calculation.** Hemodynamic changes can affect
648 the absorption of light, resulting in changes in fluorescence^{73, 74}. According to the spectra
649 of the sensors, when excited with 405-nm light, the g5-HT3.0, eCB2.0, r5-HT1.0, and
650 jRGECO1a sensors are ligand-insensitive, which can reflect hemodynamic absorption. To
651 correct for hemodynamic artifacts, we performed a pixel-by-pixel correction based on linear
652 regression⁷⁵ of the ligand-dependent signals (excited by 488 nm or 561 nm) against the
653 ligand-independent signals (excited by 405 nm). The baseline images were spatially
654 smoothed using a Gaussian filter ($\sigma=2$) and used for the linear regression. Then, for each
655 pixel, the baseline fluorescence intensity of the 405-nm excited channel was regressed
656 onto the 488-nm or 561-nm signal, obtaining regression coefficients for rescaling the 405-
657 nm channel. The rescaled 405-nm signal was subtracted from the 488-nm or 561-nm signal
658 to generate a corrected signal for each pixel. To avoid the corrected signal becoming close
659 to zero or even less than zero, the corrected signal was added to the average rescaled
660 405-nm channel signal as the final corrected signal for the response calculation. The
661 response of each pixel was calculated using the equation $\Delta F/F_0 = (F-F_0)/F_0$, where F_0 is
662 defined as the average baseline fluorescence intensity. When analyzing the data obtained
663 during the sleep-wake cycle, the baseline was defined as the REM sleep state, in which
664 the signal had low fluctuations.

665 **Parcellation of cortical areas.** Based on previous studies^{76, 77}, we rigidly registered the
666 average fluorescence image to a 2D projection of the Allen Common Coordinate
667 Framework v3 (CCFv3) using four manually labeled anatomical landmarks, namely the left,
668 center, and right points in the boundary between the anterior cortex and the olfactory bulbs,
669 and the medial point at the base of the retrosplenial cortex. To analyze the time series
670 response in an individual brain region, we averaged the $\Delta F/F_0$ value for all available pixels
671 within that brain region. To obtain the average response map from multiple mice (see Fig.
672 4g), we developed a custom code that first register the response image for each individual
673 mouse to the Allen CCFv3, and then averaged the images, keeping only the intersection
674 among all mice. For the analysis of serotonergic projection in the mouse dorsal cortex, the
675 serotonergic projection map was modified from Allen Mouse Brain Connectivity Atlas,
676 connectivity.brain-map.org/projection/experiment/cortical_map/480074702.

677 **Analysis of propagating waves**

678 **Peak response calculation.** The time series of the images obtained before (~30 s before
679 the wave originated), during, and after (~30 s after the wave disappeared) wave
680 propagation was extracted as an event for further analysis. Images taken during the first
681 20 s (20 frames) were set as the event baseline. The event response image was spatially
682 filtered, and each pixel during the event was then corrected to set the average response
683 of the event baseline to zero. The peak response site was automatically found by a circle
684 with a 500- μ m diameter across the event, and its average value was defined as the peak

685 response of the event.

686 **Identification of wave directions using optical flow analysis.** To determine the direction
687 of the waves, we adopted an optical flow method for automatically detecting the wave
688 directions based on the NeuroPatt toolbox⁷⁸. In detail, the corrected response image stack
689 was smoothed over time, downsampled in size by a factor of 0.2, and normalized to the
690 maximum response for each pixel. The phase velocity fields were then calculated using
691 the “opticalFlowHS” MATLAB function (smoothness parameter $\alpha = 0.05$). For each frame,
692 velocity fields were ignored in pixels with a low response, defined as smaller than 3-fold
693 standard deviation (SD) of the baseline. Finally, we obtained the frequency distribution of
694 these wave directions in each event and the average distribution of all events (see Fig. 5h).

695 **Calculation of wave speed.** The velocity fields calculated using the optical flow method
696 depend on two parameters⁷⁸ and tend to be underestimated; therefore, we used a different
697 method to calculate the speed of waves (see Extended Data Fig. 9e). In detail, the time (T)
698 to peak response for each pixel was determined, and the pixel with the shortest time (T_0)
699 to reach the peak response was defined as the origin. The wave-propagating region was
700 then divided by fans centered at the origin with 0.5° intervals, and the relative distance (ΔS)
701 between the distal pixel and the origin was calculated. The speed (v) in each direction was
702 then calculated using the equation $v = \Delta S / (T - T_0)$. Finally, we obtained the frequency
703 distribution for speed in each event and the average distribution of all events (see Fig. 5h).

704 **Quantification and statistical analysis**

705 Where appropriate, cells or animals were randomly assigned to either the control or
706 experimental group. Imaging data were processed using ImageJ (1.53q) software (NIH)
707 and custom-written MATLAB (R2020b) programs. Data were plotted using OriginPro 2020b
708 (Originlab) or Adobe Illustrator CC. Except where indicated otherwise, all summary data
709 are reported as the mean \pm SEM. The SNR was calculated as the peak response divided
710 by the SD of the baseline fluorescence fluctuation. All data were assumed to be distributed
711 normally, and equal variances were formally tested. Differences were analyzed using the
712 two-tailed Student’s *t*-test or one-way ANOVA; * $P < 0.05$, ** $P < 0.01$, *** $P < 0.001$, and n.s.,
713 not significant ($P \geq 0.05$). Some cartoons in Fig. 3a,f, 4a, 5a and Extended Data Fig. 7a,d,g,
714 8a were created with BioRender.com.

715 **Data availability**

716 The plasmids used to express the sensors in this study and the related sequences are
717 available from Addgene. The human GPCR cDNA library was obtained from the
718 hORFeome database 8.1 (<http://horfdb.dfci.harvard.edu/index.php?page=home>). Source
719 data are provided with this paper.

720 **Code availability**

721 The custom-written MATLAB, Arduino, and ImageJ programs will be provided upon
722 request.

723 Acknowledgments

724 This work was supported by the National Key R&D Program of China (2022YFC3300905
725 to H.D.); the National Basic Research Program of China (2019YFE011781), the National
726 Natural Science Foundation of China (31925017), the Beijing Municipal Science &
727 Technology Commission (Z220009), the NIH BRAIN Initiative (1U01NS113358 and
728 1U01NS120824), grants from the Peking-Tsinghua Center for Life Sciences and the State
729 Key Laboratory of Membrane Biology at Peking University School of Life Sciences, the
730 Feng Foundation of Biomedical Research, the Clement and Xinxin Foundation, and the
731 New Cornerstone Science Foundation (to Y.L.); the National Major Project of China
732 Science and Technology Innovation 2030 for Brain Science and Brain-Inspired Technology
733 (2022ZD0205600), the Postdoctoral Science Foundation (2022M720258), the Peking
734 University Boya Postdoctoral Fellowship (to J.W.). We thank Y. Rao for sharing *Sert-Cre*
735 mice and X. Lei at PKU-CLS and the National Center for Protein Sciences at Peking
736 University for support and assistance with the Opera Phenix high-content screening
737 system. We thank P. Gong at the University of Sydney and M. Mohajerani at University of
738 Lethbridge for their help with the optical flow analysis of waves.

739 Author contributions

740 Y.L. conceived and supervised the project. F.D., G.L., J.W. and Yu Zheng developed and
741 optimized the sensors. F.D., J.W. and G.L. performed the experiments related to
742 characterizing the sensors with help from X.X., Y.W., X.L. and Y.Y. J.W. performed the *in*
743 *vivo* fiber photometry recordings of r5-HT sensors during optogenetic stimulation and sleep
744 wake cycles. J.W., H.D., and L.L. performed the fiber photometry recordings of green 5-
745 HT sensors for *in vivo* comparison during sleep wake cycles. F.D. performed the
746 mesoscopic imaging in head-fixed mice. H.X., F.D., C.Z. and J.F. built the mesoscopic
747 imaging system. All authors contributed to the data interpretation and analysis. F.D. and
748 Y.L. wrote the manuscript with input from all other authors, especially the review and editing
749 from Yulin Zhao.

750 Competing interests

751 J.W., J.F. and Y. L have filed patent applications whose value might be affected by this
752 publication.

753 References

- 754 1. Berger, M., Gray, J.A. & Roth, B.L. The expanded biology of serotonin. *Annual Review
755 of Medicine* **60**, 355-366 (2009).
- 756 2. Li, Y. et al. Serotonin neurons in the dorsal raphe nucleus encode reward signals. *Nat
757 Commun.* **7**, 10503 (2016).
- 758 3. Portas, C.M. et al. On-line detection of extracellular levels of serotonin in dorsal raphe
759 nucleus and frontal cortex over the sleep/wake cycle in the freely moving rat.

760 *Neuroscience* **83**, 807-814 (1998).

761 4. Lesch, K.P. et al. Association of anxiety-related traits with a polymorphism in the
762 serotonin transporter gene regulatory region. *Science* **274**, 1527-1531 (1996).

763 5. Theodore, W.H., Juhasz, C., Savic, V. & Drevets, W. Serotonin, depression, and epilepsy.
764 *Epilepsia* **46**, 3-3 (2005).

765 6. Li, Y. et al. Synaptic mechanism underlying serotonin modulation of transition to cocaine
766 addiction. *Science* **373**, 1252-1256 (2021).

767 7. Vaswani, M., Linda, F.K. & Ramesh, S. Role of selective serotonin reuptake inhibitors in
768 psychiatric disorders: a comprehensive review. *Prog Neuropsychopharmacol Biol
769 Psychiatry* **27**, 85-102 (2003).

770 8. Steinbusch, H.W.M. Distribution of serotonin-immunoreactivity in the central nervous
771 system of the rat—Cell bodies and terminals. *Neuroscience* **6**, 557-618 (1981).

772 9. Huang, K.W. et al. Molecular and anatomical organization of the dorsal raphe nucleus.
773 *Elife* **8** (2019).

774 10. Ren, J. et al. Single-cell transcriptomes and whole-brain projections of serotonin neurons
775 in the mouse dorsal and median raphe nuclei. *Elife* **8** (2019).

776 11. Okaty, B.W. et al. A single-cell transcriptomic and anatomic atlas of mouse dorsal raphe
777 Pet1 neurons. *Elife* **9** (2020).

778 12. Hensler, J.G. Serotonergic modulation of the limbic system. *Neuroscience &
779 Biobehavioral Reviews* **30**, 203-214 (2006).

780 13. Hannon, J. & Hoyer, D. Molecular biology of 5-HT receptors. *Behavioural Brain Research*
781 **195**, 198-213 (2008).

782 14. Fuller, R.W. Uptake inhibitors increase extracellular serotonin concentration measured
783 by brain microdialysis. *Life Sci* **55**, 163-167 (1994).

784 15. Lama, R.D., Charlson, K., Anantharam, A. & Hashemi, P. Ultrafast detection and
785 quantification of brain signaling molecules with carbon fiber microelectrodes. *Analytical
786 Chemistry* **84**, 8096-8101 (2012).

787 16. Candelario, J. & Chachisvilis, M. Mechanical stress stimulates conformational changes
788 in 5-hydroxytryptamine receptor 1B in bone cells. *Cellular and Molecular Bioengineering*
789 **5**, 277-286 (2012).

790 17. Wan, J. et al. A genetically encoded sensor for measuring serotonin dynamics. *Nat
791 Neurosci* **24**, 746-752 (2021).

792 18. Dong, C. et al. Psychedelic-inspired drug discovery using an engineered biosensor. *Cell*
793 **184**, 2779-2792.e2718 (2021).

794 19. Kubitschke, M. et al. Next generation genetically encoded fluorescent sensors for
795 serotonin. *Nat Commun* **13**, 7525 (2022).

796 20. Unger, E.K. et al. Directed evolution of a selective and sensitive serotonin sensor via
797 machine learning. *Cell* **183**, 1986-2002.e1926 (2020).

798 21. Guiard, B.P., El Mansari, M., Merali, Z. & Blier, P. Functional interactions between
799 dopamine, serotonin and norepinephrine neurons: an in-vivo electrophysiological study
800 in rats with monoaminergic lesions. *International Journal of Neuropsychopharmacology*
801 **11**, 625-639 (2008).

802 22. Jeong, S. et al. High-throughput evolution of near-infrared serotonin nanosensors.
803 *Science Advances* **5**, eaay3771 (2019).

804 23. Zhao, Y. et al. An expanded palette of genetically encoded Ca²⁺ indicators. *Science* **333**,
805 1888-1891 (2011).

806 24. Dana, H. et al. Sensitive red protein calcium indicators for imaging neural activity. *Elife* **5**,
807 e12727 (2016).

808 25. Feng, J. et al. A genetically encoded fluorescent sensor for rapid and specific in vivo
809 detection of norepinephrine. *Neuron* **102**, 745-761.e748 (2019).

810 26. Sun, F. et al. Next-generation GRAB sensors for monitoring dopaminergic activity in vivo.
811 *Nat Methods* **17**, 1156-1166 (2020).

812 27. Bajar, B.T. et al. Improving brightness and photostability of green and red fluorescent
813 proteins for live cell imaging and FRET reporting. *Sci Rep-Uk* **6**, 20889 (2016).

814 28. Pédelacq, J.-D., Cabantous, S., Tran, T., Terwilliger, T.C. & Waldo, G.S. Engineering and
815 characterization of a superfolder green fluorescent protein. *Nat Biotechnol* **24**, 79-88
816 (2006).

817 29. Peng, Y. et al. 5-HT_{2C} receptor structures reveal the structural basis of GPCR
818 polypharmacology. *Cell* **172**, 719-730.e714 (2018).

819 30. Ballesteros, J.A. & Weinstein, H. in *Methods in Neurosciences*, Vol. 25. (ed. S.C. Sealfon)
820 366-428 (Academic Press, 1995).

821 31. Peng, Y. et al. 5-HT_{2C} receptor structures reveal the structural basis of GPCR
822 polypharmacology. *Cell* **172**, 719-730 e714 (2018).

823 32. Wu, J. et al. Improved orange and red Ca²⁺ indicators and photophysical considerations
824 for optogenetic applications. *ACS Chemical Neuroscience* **4**, 963-972 (2013).

825 33. Sun, F. et al. A genetically encoded fluorescent sensor enables rapid and specific
826 detection of dopamine in flies, fish, and mice. *Cell* **174**, 481-496.e419 (2018).

827 34. Wan, Q. et al. Mini G protein probes for active G protein-coupled receptors (GPCRs) in
828 live cells. *The Journal of Biological Chemistry* **293**, 7466-7473 (2018).

829 35. Barnea, G. et al. The genetic design of signaling cascades to record receptor activation.
830 *Proceedings of the National Academy of Sciences* **105**, 64-69 (2008).

831 36. Kroeze, W.K. et al. PRESTO-Tango as an open-source resource for interrogation of the
832 druggable human GPCRome. *Nature Structural & Molecular Biology* **22**, 362-369 (2015).

833 37. Ren, J. et al. Anatomically defined and functionally distinct dorsal raphe serotonin sub-
834 systems. *Cell* **175**, 472-487.e420 (2018).

835 38. Nagel, G. et al. Channelrhodopsin-2, a directly light-gated cation-selective membrane
836 channel. *Proceedings of the National Academy of Sciences* **100**, 13940-13945 (2003).

837 39. Nagel, G. et al. Light activation of channelrhodopsin-2 in excitable cells of *Caenorhabditis*
838 *elegans* triggers rapid behavioral responses. *Current Biology* **15**, 2279-2284 (2005).

839 40. Zhuang, X., Masson, J., Gingrich, J.A., Rayport, S. & Hen, R. Targeted gene expression
840 in dopamine and serotonin neurons of the mouse brain. *Journal of Neuroscience*
841 *Methods* **143**, 27-32 (2005).

842 41. Broussard, G.J. et al. In vivo measurement of afferent activity with axon-specific calcium
843 imaging. *Nat Neurosci* **21**, 1272-1280 (2018).

844 42. Xu, M. et al. Basal forebrain circuit for sleep-wake control. *Nat Neurosci* **18**, 1641-1647
845 (2015).

846 43. Celada, P., Puig, M.V. & Artigas, F. Serotonin modulation of cortical neurons and
847 networks. *Frontiers in Integrative Neuroscience* **7** (2013).

848 44. Oh, S.W. et al. A mesoscale connectome of the mouse brain. *Nature* **508**, 207-214 (2014).

849 45. Hamodi, A.S., Sabino, A.M., Fitzgerald, N.D., Moschou, D. & Crair, M.C. Transverse
850 sinus injections drive robust whole-brain expression of transgenes. *Elife* **9** (2020).

851 46. Ferezou, I. et al. Spatiotemporal dynamics of cortical sensorimotor integration in
852 behaving mice. *Neuron* **56**, 907-923 (2007).

853 47. Cardin, J.A., Crair, M.C. & Higley, M.J. Mesoscopic imaging: Shining a wide light on large-
854 scale neural dynamics. *Neuron* **108**, 33-43 (2020).

855 48. Klapoetke, N.C. et al. Independent optical excitation of distinct neural populations. *Nat
856 Methods* **11**, 338-346 (2014).

857 49. Wang, Q.X. et al. The Allen Mouse Brain Common Coordinate Framework: A 3D
858 reference atlas. *Cell* **181**, 936-953.e920 (2020).

859 50. Prendiville, S. & Gale, K. Anticonvulsant effect of fluoxetine on focally evoked limbic
860 motor seizures in rats. *Epilepsia* **34**, 381-384 (1993).

861 51. Pasini, A., Tortorella, A. & Gale, K. The anticonvulsant action of fluoxetine in substantia
862 nigra is dependent upon endogenous serotonin. *Brain Research* **724**, 84-88 (1996).

863 52. Tecott, L.H. et al. Eating disorder and epilepsy in mice lacking 5-HT2C serotonin
864 receptors. *Nature* **374**, 542-546 (1995).

865 53. Cheng, H.-M., Gao, C.-S., Lou, Q.-W., Chen, Z. & Wang, Y. The diverse role of the raphe
866 5-HTergic systems in epilepsy. *Acta Pharmacologica Sinica* **43**, 2777–2788 (2022).

867 54. Lin, W.-h. et al. Seizure-induced 5-HT release and chronic impairment of serotonergic
868 function in rats. *Neuroscience Letters* **534**, 1-6 (2013).

869 55. Wenzel, M., Hamm, J.P., Peterka, D.S. & Yuste, R. Reliable and elastic propagation of
870 cortical seizures in vivo. *Cell Rep* **19**, 2681-2693 (2017).

871 56. Rossi, L.F., Wykes, R.C., Kullmann, D.M. & Carandini, M. Focal cortical seizures start as
872 standing waves and propagate respecting homotopic connectivity. *Nat Commun* **8**, 217
873 (2017).

874 57. Dong, A. et al. A fluorescent sensor for spatiotemporally resolved imaging of
875 endocannabinoid dynamics in vivo. *Nat Biotechnol*, 787–798 (2021).

876 58. Ben-Ari, Y., Lagowska, J., Tremblay, E. & Le Gal La Salle, G. A new model of focal status
877 epilepticus: intra-amygdaloid application of kainic acid elicits repetitive secondarily
878 generalized convulsive seizures. *Brain Research* **163**, 176-179 (1979).

879 59. Farrell, J.S. et al. In vivo assessment of mechanisms underlying the neurovascular basis
880 of postictal amnesia. *Sci Rep-Uk* **10** (2020).

881 60. Wu, Z., Lin, D. & Li, Y. Pushing the frontiers: tools for monitoring neurotransmitters and
882 neuromodulators. *Nature Reviews Neuroscience* **23**, 257-274 (2022).

883 61. Bunin, M.A. & Wightman, R.M. Quantitative evaluation of 5-hydroxytryptamine (serotonin)
884 neuronal release and uptake: An investigation of extrasynaptic transmission. *The Journal
885 of Neuroscience* **18**, 4854-4860 (1998).

886 62. Thorré, K. et al. Differential effects of restraint stress on hippocampal 5-HT metabolism
887 and extracellular levels of 5-HT in streptozotocin-diabetic rats. *Brain Research* **772**, 209-
888 216 (1997).

889 63. Hashemi, P., Dankoski, E.C., Petrovic, J., Keithley, R.B. & Wightman, R.M. Voltammetric
890 detection of 5-hydroxytryptamine release in the rat brain. *Analytical Chemistry* **81**, 9462-
891 9471 (2009).

892 64. Lazarova, M., Bendotti, C. & Samanin, R. Studies on the role of serotonin in different
893 regions of the rat central nervous system of pentylenetetrazol-induced seizures and the
894 effect of di-n-propylacetate. *Naunyn-Schmiedeberg's Archives of Pharmacology* **322**,
895 147-152 (1983).

896 65. Gibson, D.G. et al. Enzymatic assembly of DNA molecules up to several hundred
897 kilobases. *Nat Methods* **6**, 343-345 (2009).

898 66. Yusa, K., Zhou, L., Li, M.A., Bradley, A. & Craig, N.L. A hyperactive piggyBac transposase
899 for mammalian applications. *Proceedings of the National Academy of Sciences* **108**,
900 1531-1536 (2011).

901 67. Barger, Z., Frye, C.G., Liu, D.Q., Dan, Y. & Bouchard, K.E. Robust, automated sleep
902 scoring by a compact neural network with distributional shift correction. *Plos One* **14**
903 (2019).

904 68. Werley, C.A., Chien, M.P. & Cohen, A.E. Ultrawidefield microscope for high-speed
905 fluorescence imaging and targeted optogenetic stimulation. *Biomed Opt Express* **8**,
906 5794-5813 (2017).

907 69. Zhuang, C. et al. Real-time brain-wide multi-planar microscopy for simultaneous cortex
908 and hippocampus imaging at the cellular resolution in mice. *Biomed Opt Express* **12**,
909 1858-1868 (2021).

910 70. Jackson, J., Karnani, M.M., Zemelman, B.V., Burdakov, D. & Lee, A.K. Inhibitory control
911 of prefrontal cortex by the claustrum. *Neuron* **99**, 1029-1039.e1024 (2018).

912 71. Pnevmatikakis, E.A. & Giovannucci, A. NoRMCorre: An online algorithm for piecewise
913 rigid motion correction of calcium imaging data. *Journal of Neuroscience Methods* **291**,
914 83-94 (2017).

915 72. Arganda-Carreras, I. et al. Trainable Weka Segmentation: a machine learning tool for
916 microscopy pixel classification. *Bioinformatics* **33**, 2424-2426 (2017).

917 73. Ma, Y. et al. Wide-field optical mapping of neural activity and brain haemodynamics:
918 considerations and novel approaches. *Philosophical Transactions of the Royal Society
919 B: Biological Sciences* **371**, 20150360 (2016).

920 74. Valley, M.T. et al. Separation of hemodynamic signals from GCaMP fluorescence
921 measured with wide-field imaging. *Journal of Neurophysiology* **123**, 356-366 (2020).

922 75. Vesuna, S. et al. Deep posteromedial cortical rhythm in dissociation. *Nature* **586**, 87-94
923 (2020).

924 76. Musall, S., Kaufman, M.T., Juavinett, A.L., Gluf, S. & Churchland, A.K. Single-trial neural
925 dynamics are dominated by richly varied movements. *Nat Neurosci* **22**, 1677-1686
926 (2019).

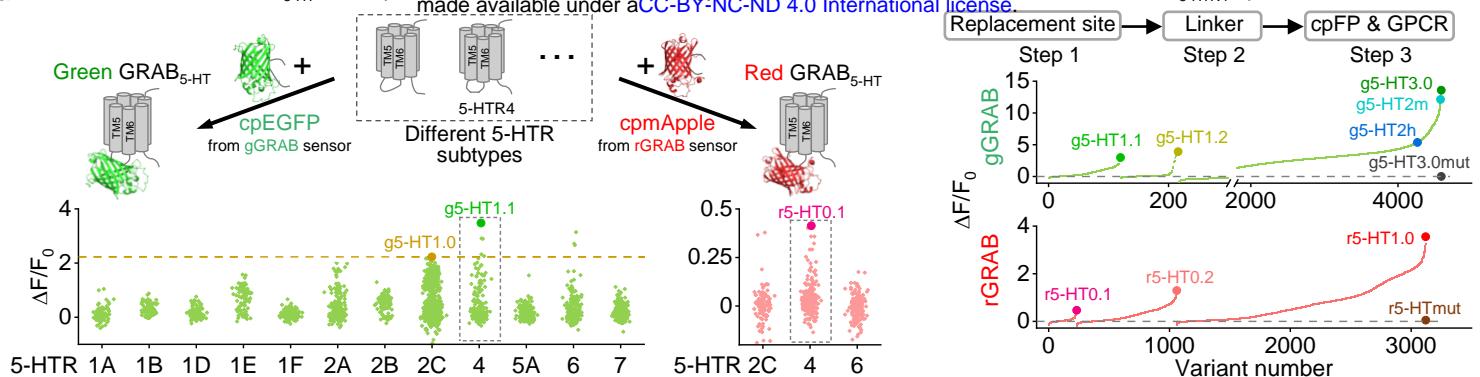
927 77. Saxena, S. et al. Localized semi-nonnegative matrix factorization (LocaNMF) of widefield
928 calcium imaging data. *PLOS Computational Biology* **16**, e1007791 (2020).

929 78. Townsend, R.G. & Gong, P. Detection and analysis of spatiotemporal patterns in brain
930 activity. *PLOS Computational Biology* **14**, e1006643 (2018).

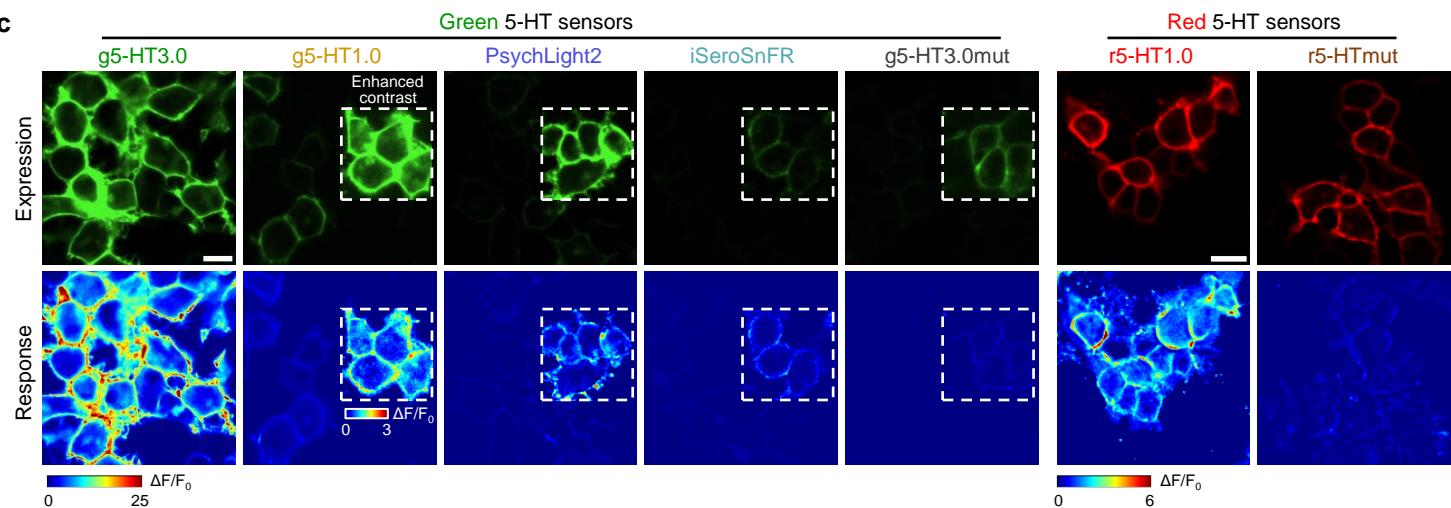
931 79. Ballesteros, J.A. & Weinstein, H. [19] Integrated methods for the construction of three-
932 dimensional models and computational probing of structure-function relations in G
933 protein-coupled receptors. *Methods in Neurosciences* **25**, 366-428 (1995).

934

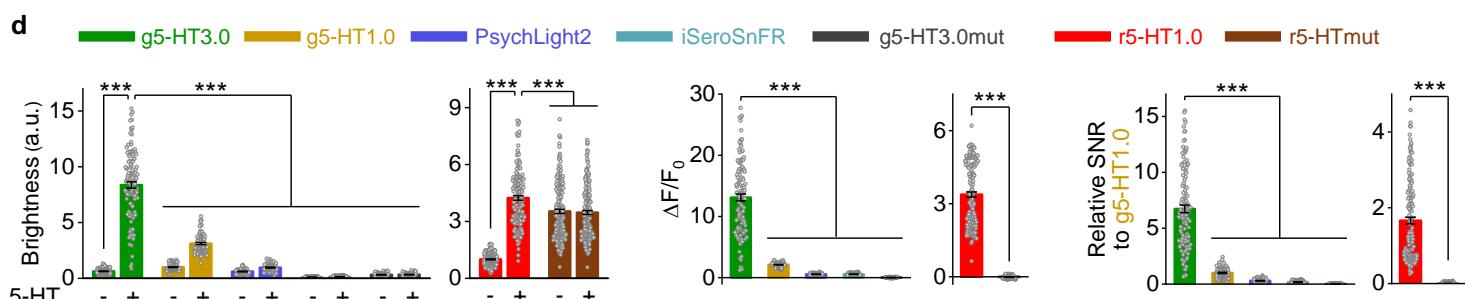
a



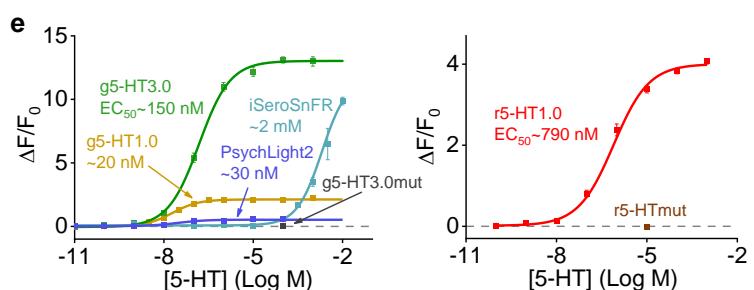
c



d



e



f

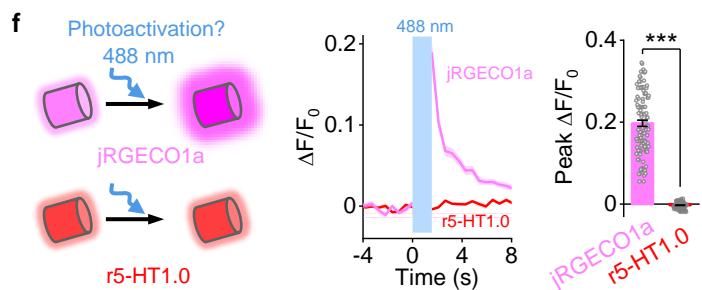


Fig. 1 | Development of improved green fluorescent 5-HT sensors and new red 5-HT sensors.

935 **Fig. 1 | Development of improved green fluorescent 5-HT sensors and new red 5-HT sensors.**

936 **a**, Schematic illustrating the strategy for developing new GRAB_{5-HT} sensors (top). Performance of
937 sensor candidates based on different receptor subtypes for green (bottom left) and red 5-HT sensors
938 (bottom right). The dashed horizontal line represents g5-HT1.0 response (bottom left), and the best
939 candidates in green and red sensors are denoted by enlarged green and pink dots, respectively.

940 **b**, Optimization of the replacement site, linker, cpFP and GPCR. Responses to 10 μ M 5-HT of various
941 candidates are presented, and different versions are indicated with enlarged dots.

942 **c**, Representative images of sensors' expression (top, with 5-HT) and response (bottom) to 5-HT in
943 HEK293T cells. Insets with white dashed outlines in images have either enhanced contrast (top) or
944 different pseudocolor scales (bottom). 100 μ M 5-HT for green sensors and 10 μ M 5-HT for red sensors.
945 Scale bar, 20 μ m.

946 **d**, Group summary of the brightness (left), peak $\Delta F/F_0$ (middle) and SNR (right) of different 5-HT
947 sensors. The SNR of all sensors were relative to g5-HT1.0; a.u., arbitrary units, the basal brightness
948 of g5-HT1.0 is set as 1. $n = 119$ cells from 3 coverslips (hereafter denoted as 119/3) for g5-HT3.0, 82/3
949 for g5-HT1.0, 64/3 for PsychLight2, 139/3 for iSeroSnFR, 92/3 for g5-H3.0mut, 159/5 for r5-HT1.0 and
950 191/5 for r5-HTmut; 100 μ M 5-HT for green sensors and 10 μ M 5-HT for red sensors. (One-way
951 ANOVA followed by Tukey's multiple-comparison tests for green sensors; for brightness, $F_{9,982} = 600.2$,
952 $P = 0$, post hoc test: $P < 10^{-8}$ for g5-HT3.0 with 5-HT versus g5-HT3.0 without 5-HT and other sensors
953 with or without 5-HT; for peak $\Delta F/F_0$, $F_{4,491} = 387.1$, $P = 2.76 \times 10^{-150}$, post hoc test: $P < 10^{-8}$ for g5-HT3.0
954 versus other sensors; for relative SNR, $F_{4,491} = 285.7$, $P = 1.13 \times 10^{-126}$, post hoc test: $P < 10^{-6}$ for g5-
955 HT3.0 versus other sensors. One-way ANOVA followed by Tukey's multiple-comparison tests for
956 brightness of red sensors, $F_{3,696} = 178.3$, $P = 9.26 \times 10^{-86}$, post hoc test: $P < 10^{-5}$ for r5-HT1.0 with 5-HT
957 versus r5-HT1.0 without 5-HT and r5-HTmut with or without 5-HT. Two-tailed Student's *t*-test for r5-
958 HT1.0 and r5-HTmut; for peak $\Delta F/F_0$, $P = 3.13 \times 10^{-72}$; for relative SNR, $P = 2.67 \times 10^{-43}$.)

959 **e**, Dose-response curves of different 5-HT sensors. $n = 3$ wells for each sensor with 300–500 cells per
960 well.

961 **f**, Schematic illustrates the photoactivation properties of jRGECO1a and r5-HT1.0 (left), representative
962 traces (middle) and group summary of peak $\Delta F/F_0$ (right) in response to blue light (488 nm, without
963 imaging) in cells expressing jRGECO1a or r5-HT1.0. $n = 105/4$ for jRGECO1a and 88/4 for r5-HT1.0.
964 (Two-tailed Student's *t*-test, $P = 2.07 \times 10^{-48}$).

965 Data are shown as mean \pm SEM in **d–f**, with the error bars or shaded regions indicating the SEM,
966 *** $P < 0.001$.

967

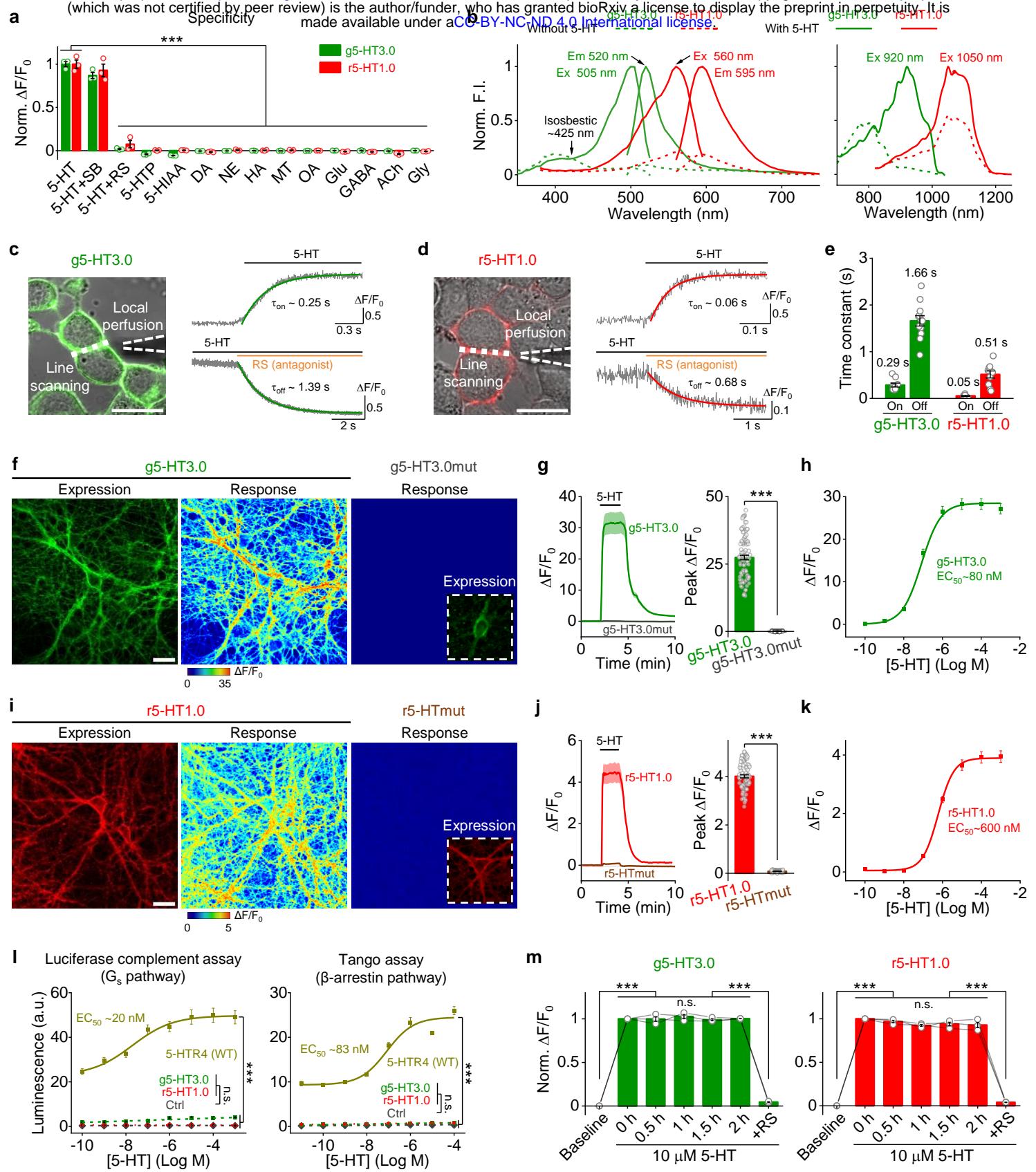


Fig. 2 | Characterization of 5-HT sensors in HEK293T cells and cultured rat cortical neurons.

968 **Fig. 2 | Characterization of 5-HT sensors in HEK293T cells and cultured rat cortical neurons.**

969 **a**, Normalized $\Delta F/F_0$ of g5-HT3.0 and r5-HT1.0 in response to different compounds (each at 10 μM
970 except RS at 100 μM). 5-HTP, 5-hydroxytryptophan; 5-HIAA, 5-hydroxyindole acetic acid; DA,
971 dopamine; NE, norepinephrine; HA, histamine; MT, melatonin; OA, octopamine; Glu, glutamate; GABA,
972 gamma-aminobutyric acid; ACh, acetylcholine; Gly, glycine. Norm., normalized. $n = 3$ wells per group,
973 200–500 cells per well. (For g5-HT3.0, $F_{13,28} = 745.7$, $P = 5.74 \times 10^{-32}$, post hoc test: $P = 0$ for 5-HT
974 versus 5-HT and RS, and other compounds; for r5-HT1.0, $F_{13,28} = 180.6$, $P = 2.02 \times 10^{-23}$, post hoc test:
975 $P = 0$ for 5-HT versus 5-HT and RS, and other compounds.)

976 **b**, One-photon excitation (Ex) and emission (Em) spectra and two-photon excitation spectra of g5-
977 HT3.0 and r5-HT1.0 in the absence (dashed line) and presence of 10 μM 5-HT (solid line). F.I.,
978 fluorescence intensity.

979 **c–e**, Kinetic of g5-HT3.0 and r5-HT1.0 in cultured HEK293T cells. Illustration of the local puffing
980 system (**c,d**, left). Representative traces of sensor fluorescence increase to 5-HT puffing (**c,d**, top
981 right) and decrease to RS puffing (**c,d**, bottom right). Group summary of on and off kinetics (**e**). $n = 10$
982 cells from 3 coverslips (short for 10/3) for g5-HT3.0 on kinetics, 12/4 for g5-HT3.0 off kinetics, 9/3 for
983 r5-HT1.0 on kinetics, 12/4 for r5-HT1.0 off kinetics.

984 **f**, Representative images showing the expression and responses of g5-HT3.0 and g5-HT3.0mut to
985 100 μM 5-HT in cultured rat cortical neurons. The inset in the g5-HT3.0mut response image shows
986 the contrast-enhanced expression image.

987 **g**, Representative traces and peak response summary of g5-HT3.0 and g5-HT3.0mut in response to
988 100 μM 5-HT. $n = 96$ ROIs from 5 coverslips (short for 96/5) for g5-HT3.0 and 92/5 for g5-HT3.0mut.
989 ($P = 1.40 \times 10^{-53}$ for g5-HT3.0 versus g5-HT3.0mut.)

990 **h**, The dose-response curve of g5-HT3.0. $n = 76/4$.

991 **i**, Representative images showing the expression and responses of r5-HT1.0 and r5-HTmut to 10 μM
992 5-HT.

993 **j**, Representative traces and peak response summary of r5-HT1.0 and r5-HTmut in response to 10
994 μM 5-HT. $n = 80/4$ for r5-HT1.0 and 60/3 for r5-HTmut. ($P = 4.46 \times 10^{-70}$ for r5-HT1.0 versus r5-HTmut.)

995 **k**, The dose-response curve of r5-HT1.0. $n = 80/4$.

996 **l**, Downstream coupling tests. WT, wild type; Ctrl, control, without expression of wild type 5-HTR4 or
997 sensors; a.u., arbitrary units. $n = 3$ wells per group, 200–500 cells per well. (For luciferase complement
998 assay, $F_{3,8} = 256$, $P = 2.77 \times 10^{-8}$, post hoc test: $P = 0$ and 0.37 for g5-HT3.0 versus 5-HTR4 (WT) and
999 Ctrl in 1 mM 5-HT, respectively, $P = 0$ and 1 for r5-HT1.0 versus 5-HTR4 (WT) and Ctrl, respectively;
1000 for Tango assay, $F_{3,8} = 766.4$, $P = 3.55 \times 10^{-10}$, post hoc test: $P = 0$ and 0.89 for g5-HT3.0 versus 5-
1001 HTR4 (WT) and Ctrl in 100 μM 5-HT, respectively, $P = 0$ and 0.86 for r5-HT1.0 versus 5-HTR4 (WT)
1002 and Ctrl, respectively.)

1003 **m**, Normalized $\Delta F/F_0$ of g5-HT3.0 and r5-HT1.0 in response to the 2-h application of 10 μM 5-HT,
1004 followed by 100 μM RS. $n = 3$ wells for each sensor. (For g5-HT3.0, $F = 359.8$, $P = 0.034$, post hoc
1005 test: $P = 1.29 \times 10^{-6}$ for baseline versus 0 h, $P = 1.76 \times 10^{-6}$ for 2.0 h versus RS, $P = 1, 0.77, 1, 1$ for 0 h
1006 versus 0.5 h, 1 h, 1.5 h or 2.0 h, respectively; for r5-HT1.0, $F = 250.9$, $P = 0.04$, post hoc test: $P =$
1007 2.85×10^{-6} for baseline versus 0 h, $P = 5.82 \times 10^{-6}$ for 2.0 h versus RS, $P = 0.95, 0.44, 0.66, 0.64$ for 0 h
1008 versus 0.5 h, 1 h, 1.5 h or 2.0 h, respectively.)

1009 **a–e** and **l** tested in HEK293T cells; **f–k** and **m** tested in cultured rat cortical neurons.

1010 All scale bar, 20 μM . Data are shown as mean \pm SEM in **a,e,g,h,j–m**, with the error bars or shaded
1011 regions indicating the SEM. One-way ANOVA (in **a,l**) and one-way repeated measures ANOVA (in **m**)
1012 followed by Tukey's multiple-comparison tests; two-tailed Student's *t*-test in **g,j**; *** $P < 0.001$, n.s., not
1013 significant.

a

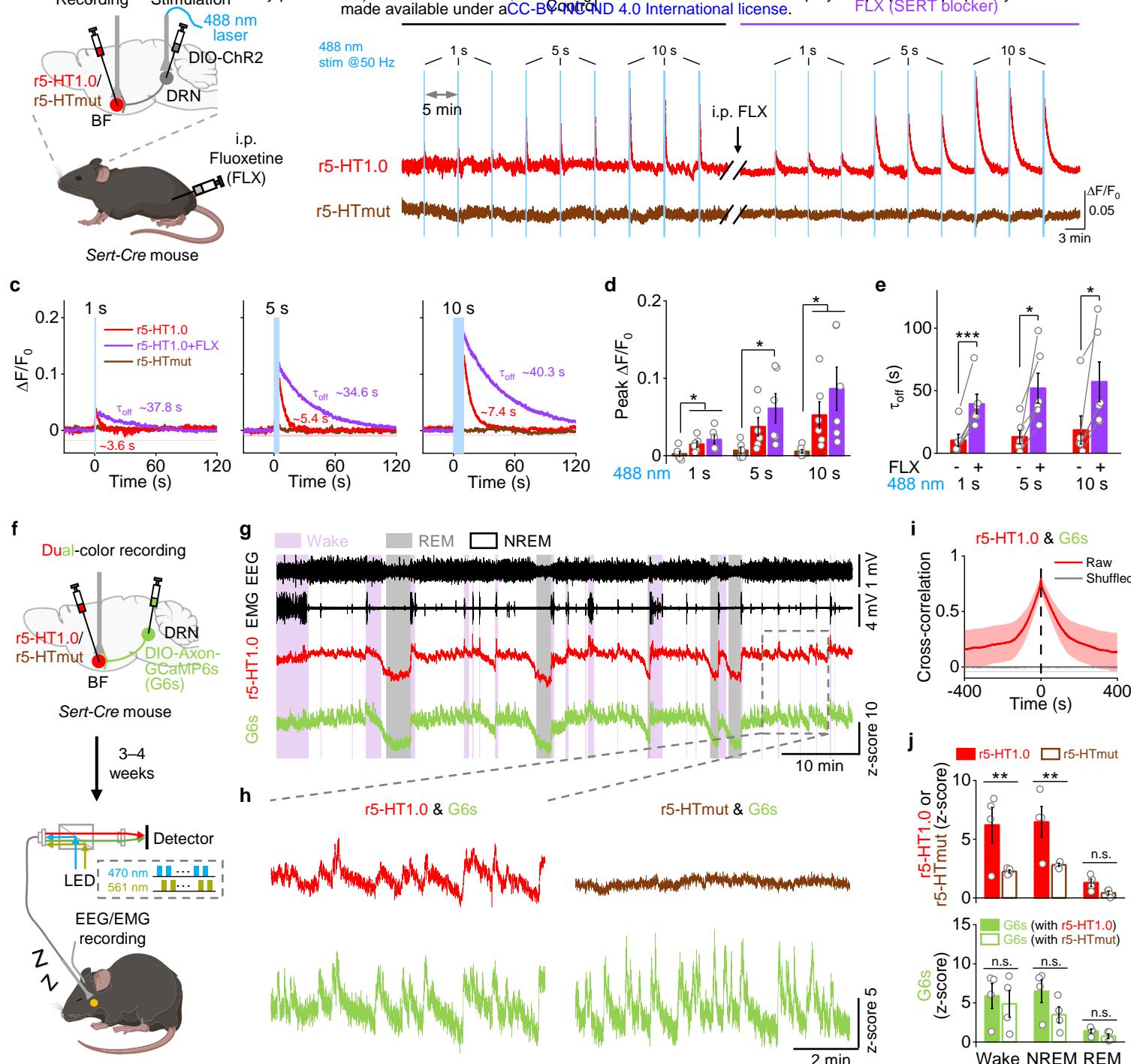


Fig. 3 | Red GRAB_{5-HT} sensor can monitor endogenous 5-HT release in freely moving mice.

1015 **Fig. 3 | Red GRAB_{5-HT} sensor can monitor endogenous 5-HT release in freely moving mice.**

1016 **a**, Schematic depicting the fiber-photometry recording setup using red 5-HT sensors with optogenetic
1017 activation of DRN in *Sert-Cre* mice.

1018 **b**, Representative $\Delta F/F_0$ traces of r5-HT1.0 and r5-HTmut in response to optical stimulation in the DRN
1019 under different stimulation durations before or after fluoxetine (FLX) application. Blue shading, period
1020 for 488-nm stimulation.

1021 **c**, Averaged $\Delta F/F_0$ traces of r5-HT1.0 and r5-HTmut under different stimulation durations in an
1022 example mouse.

1023 **d**, Summarized peak responses of r5-HT1.0 and r5-HTmut under different stimulation durations. $n =$
1024 6 mice for each treatment. (Two-tailed Student's *t*-tests, for r5-HTmut versus r5-HT1.0, $P = 0.030$,
1025 0.052, 0.041 under 1 s, 5 s, 10 s stimulation, respectively; for r5-HTmut versus r5-HT1.0+FLX, $P =$
1026 0.016, 0.034, 0.033 under 1 s, 5 s, 10 s stimulation, respectively.)

1027 **e**, Summarized decay kinetics of r5-HT1.0 with or without FLX application under different stimulation
1028 durations. $n = 6$ mice for each treatment. (Paired *t*-tests for r5-HT1.0 and r5-HT1.0 + FLX, $P = 4.44 \times 10^{-4}$,
1029 1.44×10^{-2} , 3.19×10^{-2} for 1 s, 5 s and 10 s stimulation, respectively.)

1030 **f**, Schematic showing the setup for dual-color recording of r5-HT1.0 or r5-HTmut and GCaMP6s (G6s)
1031 during the sleep-wake cycle.

1032 **g**, Representative traces of simultaneous EEG, EMG, r5-HT1.0, and G6s recording during the sleep-
1033 wake cycle in freely behaving mice. Pink shading, wake state; gray shading, REM sleep.

1034 **h**, Zoom-in traces of r5-HT1.0 and G6s (from **g**) or r5-HTmut and G6s (mainly during the NREM sleep).

1035 **i**, The averaged cross-correlation between r5-HT1.0 and G6s signals during the sleep-wake cycle.

1036 **j**, Averaged responses of 5-HT sensors (red channel, r5-HT1.0 or r5-HTmut) and G6s (green channel).
1037 $n = 4$ mice for each group. (Two-way repeated measures ANOVA followed by Tukey's multiple-
1038 comparison tests; for r5-HT1.0 versus r5-HTmut, post hoc test: $P = 5.65 \times 10^{-3}$, 9.22×10^{-3} and 0.47
1039 during wake, NREM and REM sleep state, respectively; for G6s (with r5-HT1.0) versus G6s (with r5-
1040 HTmut), post hoc test: $P = 0.56$, 0.11 and 0.71 during wake, NREM and REM sleep state, respectively.)

1041 Data are shown as mean \pm SEM in **c-e,i,j**, with the error bars or shaded regions indicating the SEM,
1042 $*P < 0.05$, $**P < 0.01$, $***P < 0.001$, n.s., not significant.

1043

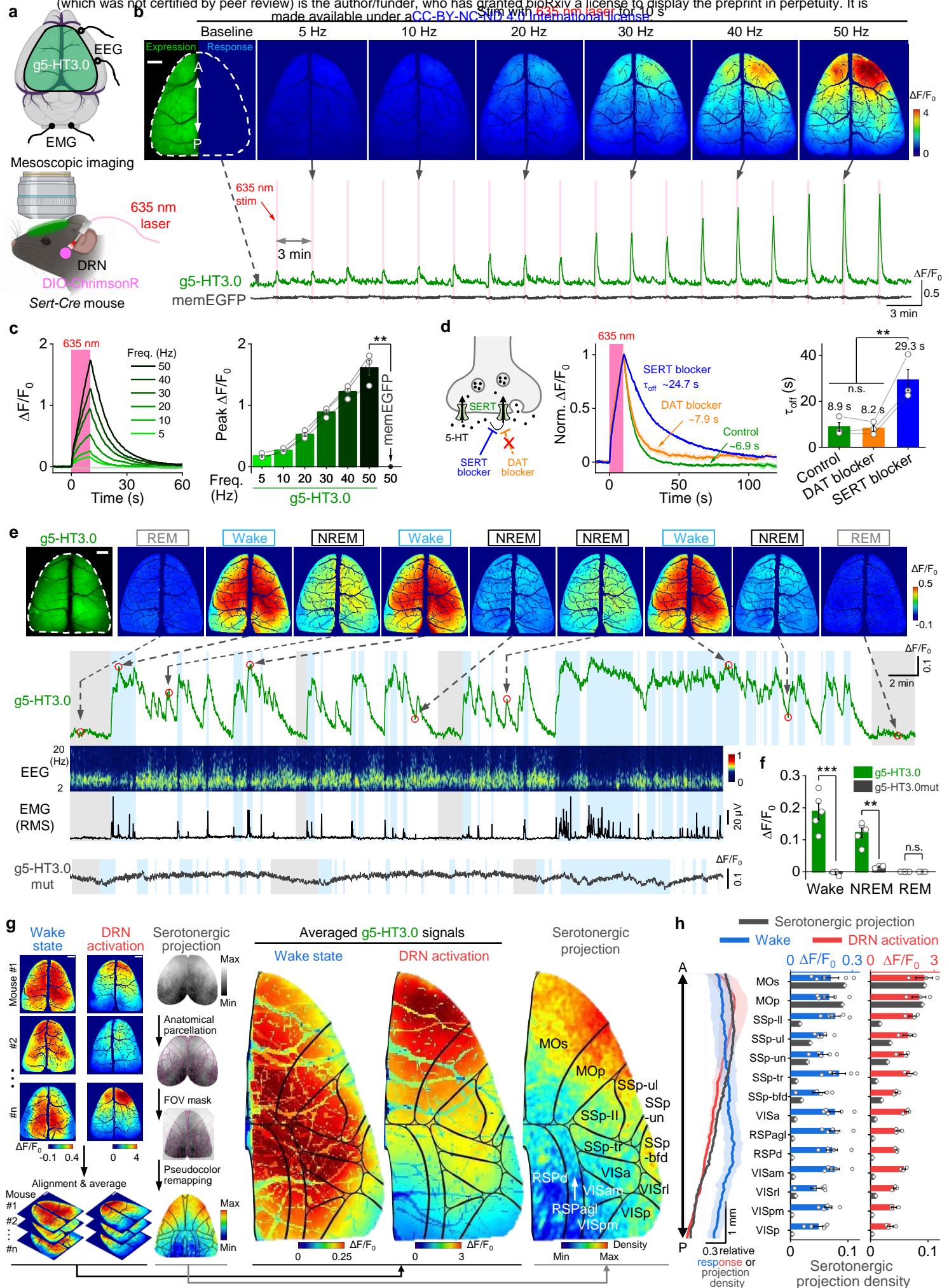


Fig. 4 | gGRAB_{5-HT3.0} reveals 5-HT release in mouse dorsal cortex *in vivo* by mesoscopic imaging.

1044 **Fig. 4 | gGRAB_{5-HT3.0} reveals 5-HT release in mouse dorsal cortex *in vivo* by mesoscopic**
1045 **imaging.**

1046 **a**, Schematic depicting the setup of mesoscopic imaging.

1047 **b**, Representative images showing the cortical g5-HT3.0 expression and response to optical
1048 stimulation in the DRN with incremental frequencies (top). Representative traces of g5-HT3.0 and a
1049 negative control memEGFP (bottom). The dashed white outline indicates the ROI.

1050 **c**, Representative $\Delta F/F_0$ traces of g5-HT3.0 (left) and group data of peak response (right) with
1051 increased frequencies of 635 nm laser. $n = 3$ mice for each group. (Two-tailed Student's *t*-tests, $P =$
1052 8.48×10^{-3} for g5-HT3.0 versus memEGFP under 50 Hz stimulation.)

1053 **d**, Schematic illustrating the effect of SERT blocker and DAT blocker on extracellular 5-HT level (left).
1054 Representative $\Delta F/F_0$ traces of g5-HT3.0 (middle) and summary data of decay kinetics (right) during
1055 50 Hz 10 s stimulation after treatment with indicated compounds. (One-way repeated measures
1056 ANOVA followed by Tukey's multiple-comparison tests, $F = 28.9$, $P = 4.18 \times 10^{-3}$, post hoc test: $P = 0.98$
1057 for DAT blocker versus control, 6.45×10^{-3} for SERT blocker versus control and 5.72×10^{-3} for SERT
1058 blocker versus DAT blocker.)

1059 **e**, Representative fluorescence and pseudocolor images of g5-HT3.0 during the sleep-wake cycle
1060 (top). Representative traces of g5-HT3.0 response, EEG, EMG (by root mean square, RMS) and g5-
1061 HT3.0mut response in the dorsal cortex during the sleep-wake cycle (bottom). The dashed white
1062 outline in the top left image indicates the ROI. Dashed arrows and red circles indicate the timepoint of
1063 frames shown at the top. Gray shading, REM sleep; light blue shading, wake state.

1064 **f**, Group data of g5-HT3.0 and g5-HT3.0mut responses in mice during the awake state, NREM and
1065 REM sleep. $n = 5$ mice for g5-HT3.0 and 3 mice for g5-HT3.0mut. (Two-way repeated measures
1066 ANOVA followed by Tukey's multiple-comparison tests for g5-HT3.0 and g5-HT3.0mut, $P = 5.77 \times 10^{-6}$,
1067 1.89×10^{-3} and 1 during the wake, NREM and REM sleep state, respectively.)

1068 **g**, Snapshots of g5-HT3.0 responses in different mice in the awake state and DRN activation, and
1069 serotonergic projection map modified from Allen Brain (left). Averaged pseudocolor images of g5-
1070 HT3.0 responses under indicated conditions (middle) and serotonergic projection map overlaid with
1071 black outlines aligned to the Allen Mouse Brain CCF (right). $n = 5$ and 3 mice for the awake state and
1072 DRN activation group, respectively.

1073 **h**, Averaged relative responses of g5-HT3.0 and serotonergic projection density along the anterior-to-
1074 posterior (AP) axis (left) and summary of g5-HT3.0 signals or serotonergic projection density in
1075 different cortex regions.

1076 All scale bar, 1 mm. Data are shown as mean \pm SEM in **c,d,f,h**, with the error bars or shaded regions
1077 indicating the SEM, ** $P < 0.01$, *** $P < 0.001$, n.s., not significant.

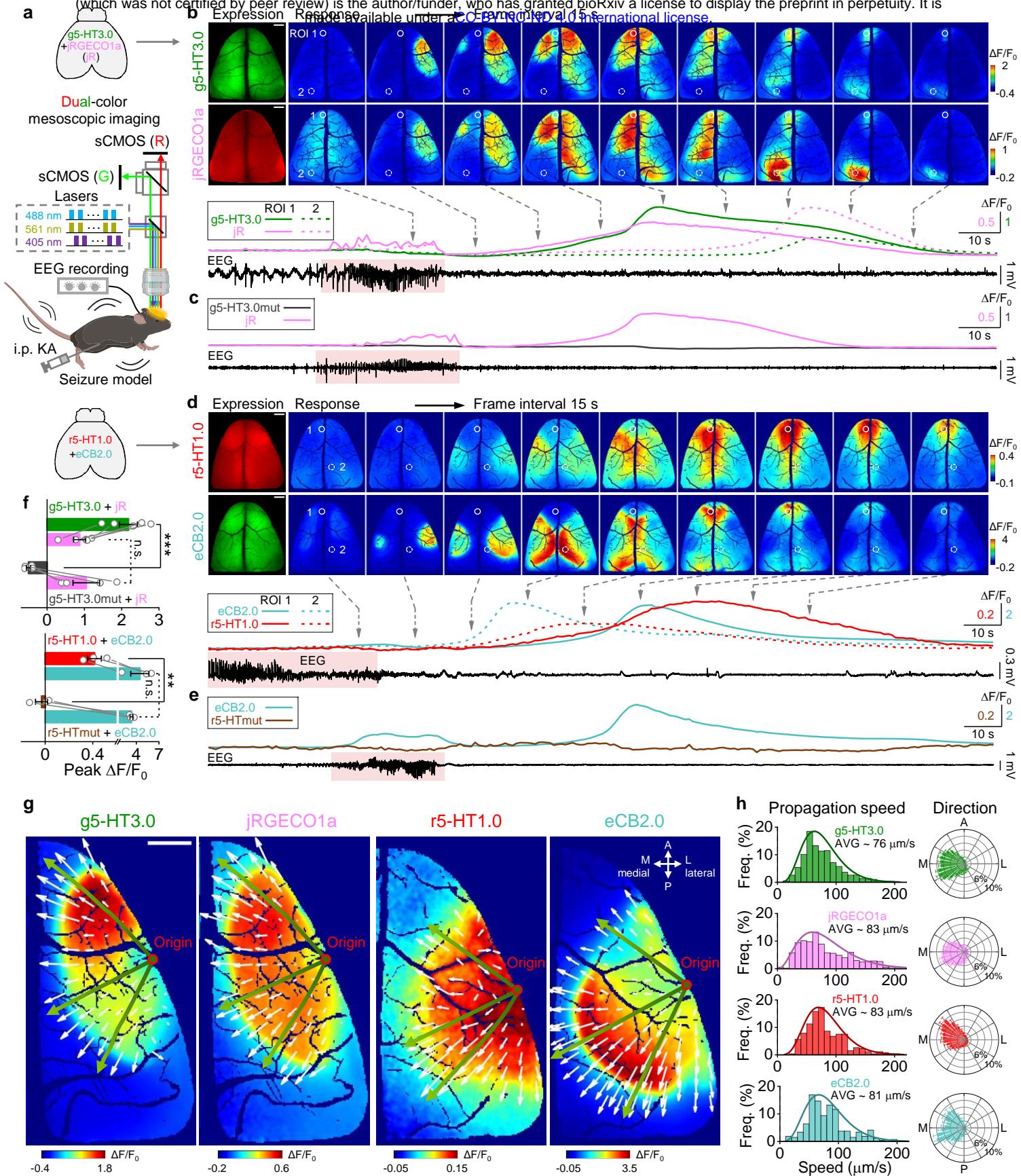


Fig. 5 | Dual-color imaging of cortex-wide neurochemical waves during seizures.

1079 **Fig. 5 | Dual-color imaging of cortex-wide neurochemical waves during seizures.**

1080 **a**, Schematic depicting the setup of dual-color mesoscopic imaging in a KA-induced seizure model.

1081 **b**, Representative images and $\Delta F/F_0$ traces of g5-HT3.0 and jRGECO1a during seizures. Two ROIs

1082 (500 μm in diameter) are labeled; ROI 1 (the white circle) and ROI 2 (the white dashed circle) show

1083 the maximum response regions of g5-HT3.0 and jRGECO1a, respectively. The solid and dashed lines

1084 in traces correspond to ROI 1 and ROI 2, respectively. The red shading in the EEG trace indicates the

1085 epileptic discharges.

1086 **c**, Representative $\Delta F/F_0$ traces of g5-HT3.0mut and jRGECO1a during seizures, similar to **b**, and

1087 images are showed in Extended Data Fig. 9b.

1088 **d**, Representative images and $\Delta F/F_0$ traces of r5-HT1.0 and eCB2.0 during seizures, similar to **b**,

1089 except that ROI 1 (the white circle) and ROI 2 (the white dashed circle) show the maximum response

1090 regions of r5-HT1.0 and eCB2.0, respectively.

1091 **e**, Representative traces of r5-HTmut and eCB2.0 signals during seizures, similar to **d**, and images

1092 are showed in Extended Data Fig. 9d.

1093 **f**, Group summary of different sensors' peak responses. $n = 5$ mice for the group co-expressing g5-

1094 HT3.0 and jRGECO1a, $n = 4$ for g5-HT3.0mut and jRGECO1a, $n = 3$ for r5-HT1.0 and eCB2.0, $n = 3$

1095 for r5-HTmut and eCB2.0. (Two-tailed Student's *t*-tests, $P = 2.36 \times 10^{-4}$ for g5-HT3.0 versus g5-

1096 HT3.0mut, $P = 0.64$ for jRGECO1a between two groups; $P = 4.41 \times 10^{-3}$ for r5-HT1.0 versus r5-HTmut,

1097 $P = 0.45$ for eCB2.0 between two groups.)

1098 **g**, Representative images showing the wave propagation detected by indicated sensors. The red circle

1099 indicates the origin of waves; small white arrows indicate the wave-propagating velocity vector; green

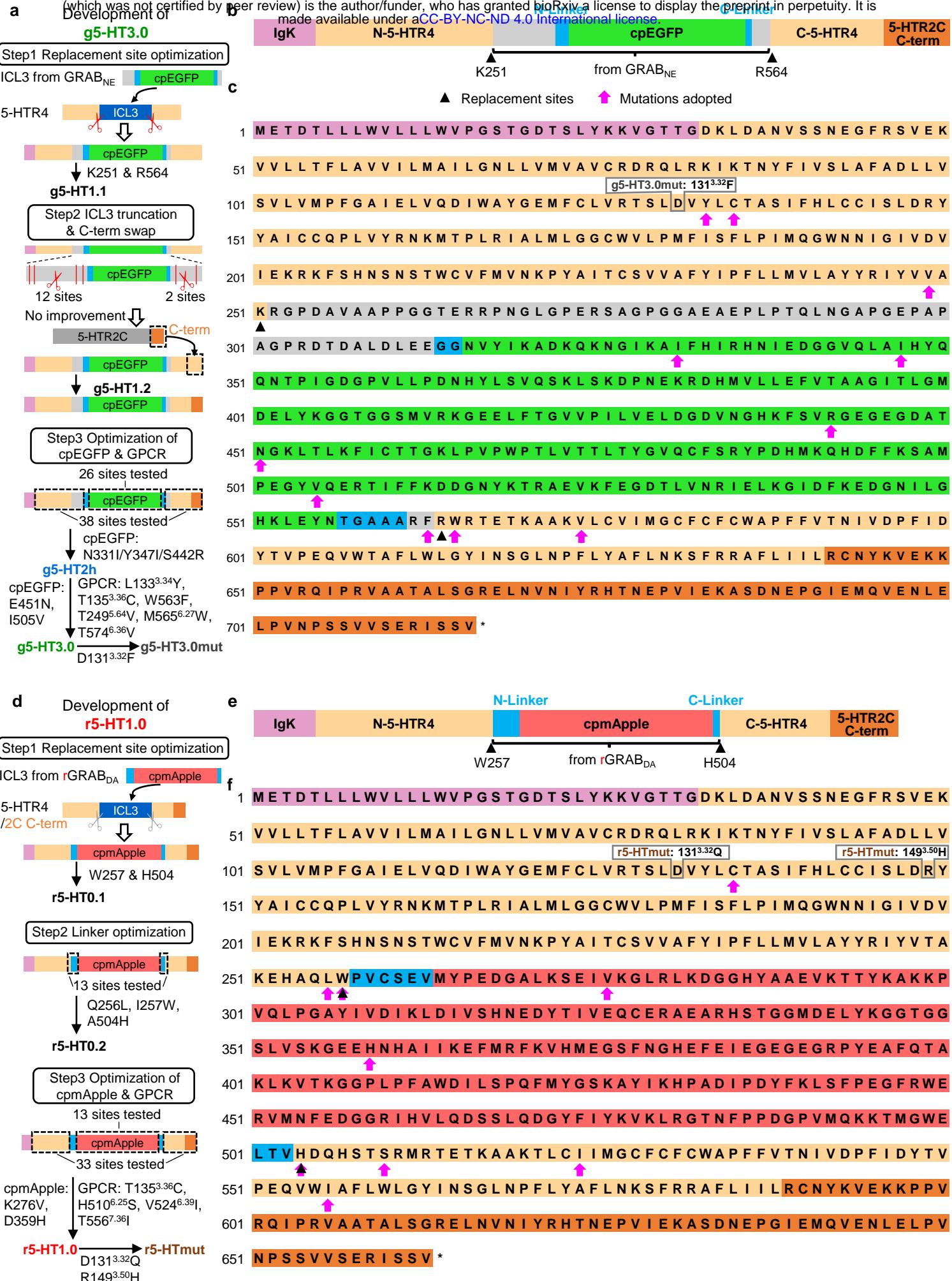
1100 lines with arrow indicate example propagating trajectories. L, lateral, M, medial, A, anterior, P, posterior.

1101 **h**, Probability distributions of wave-propagating speed and direction calculated by indicated sensors.

1102 Scale bar in all images, 1 mm. Data are shown as mean \pm SEM in **f,h**, with the error bars indicating

1103 the SEM, ** $P < 0.01$, *** $P < 0.001$, n.s., not significant.

1104



Extended Data Fig. 1 | Development and sequence of GRAB_{5-HT} sensors.

1105 **Extended Data Fig. 1 | Development and sequence of GRAB_{5-HT} sensors.**

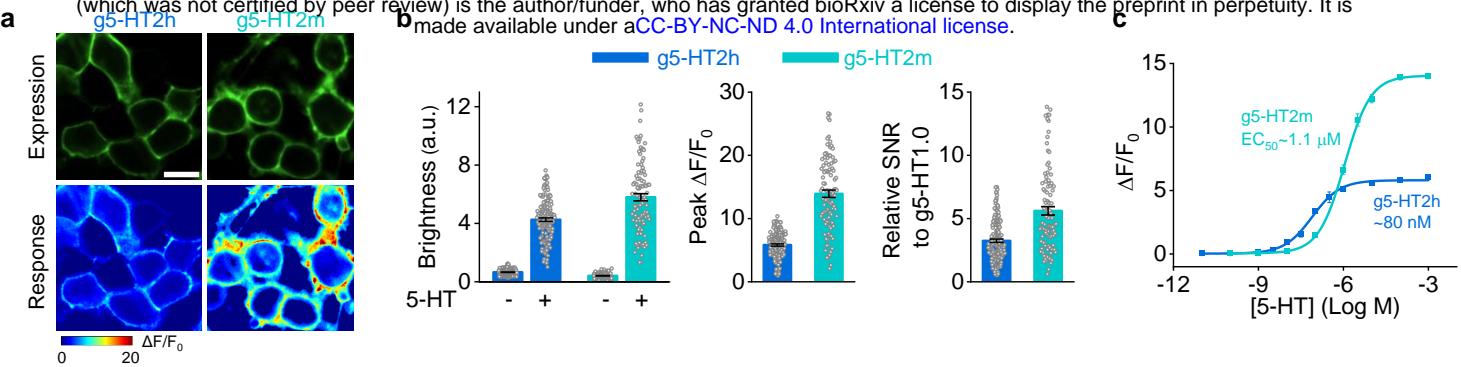
1106 **a**, A flowchart depicting the development of the g5-HT3.0 sensor, including replacement site
1107 optimization, C-terminal (C-term) swap with 5-HTR2C, linker, cpEGFP and GPCR optimization.
1108 Mutations adopted in each step are noted.

1109 **b**, Schematic showing components of the g5-HT3.0 sensor. The cpEGFP and linkers were
1110 transplanted from GRAB_{NE}.

1111 **c**, The amino acid sequence of g5-HT3.0, in which replacement sites for ICL3 loop from GRAB_{NE} are
1112 denoted by black arrowhead and mutated amino acids are indicated by pink arrow. The numbering of
1113 amino acid corresponds to the start of the IgK leader in the sensor and superscripts in the insets of
1114 **a,c** are based on the Ballesteros-Weinstein numbering system⁷⁹.

1115 **d-f**, The development (**d**), components (**e**) and sequence (**f**) of the r5-HT1.0 sensor. Similar to **a-c**,
1116 except that the C-term was swapped with 5-HTR2C C-term in the first step (replacement site
1117 optimization).

1118



Extended Data Fig. 2 | Performance of g5-HT2h and g5-HT2m in HEK293T cells.

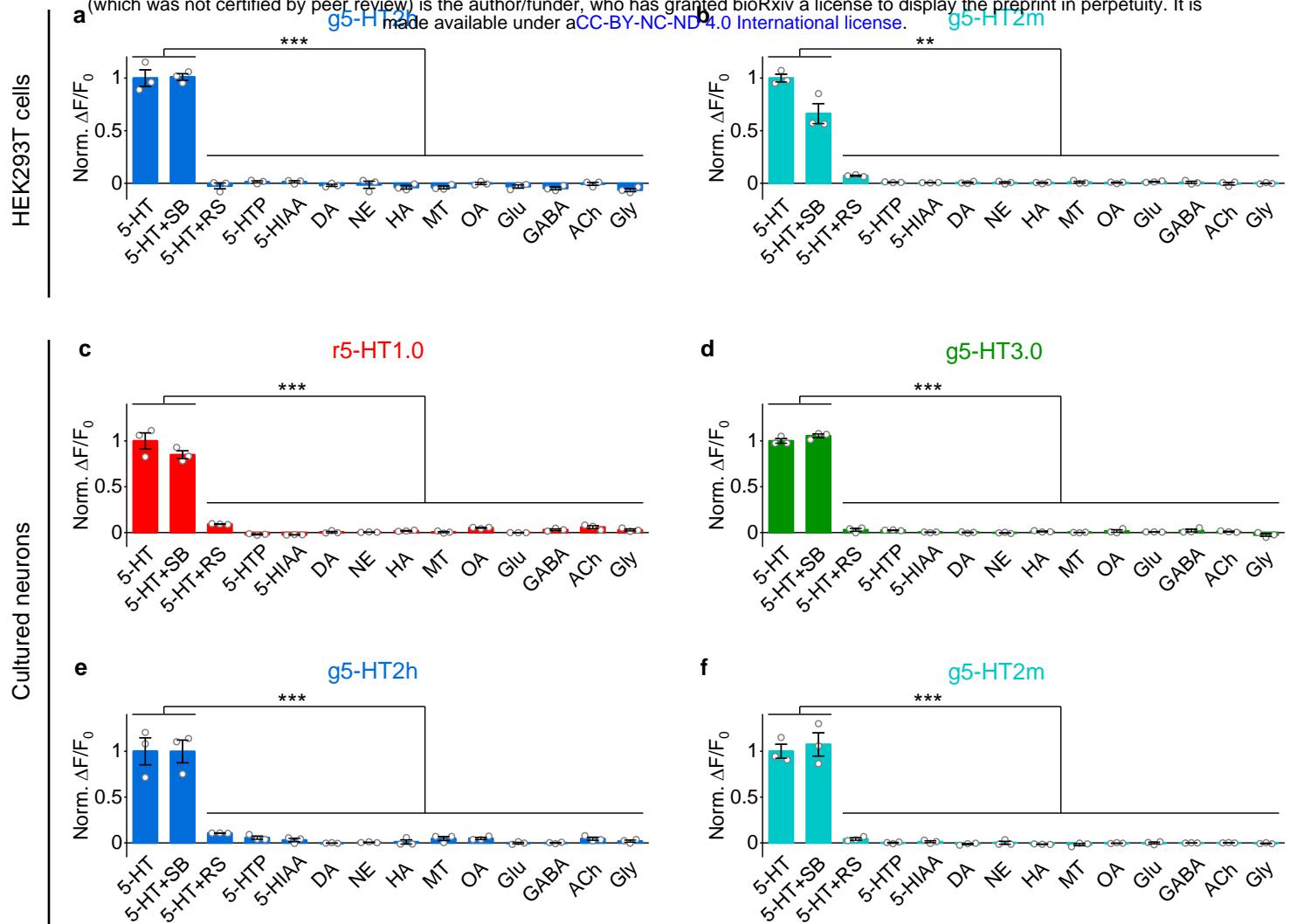
1119 **Extended Data Fig. 2 | Performance of g5-HT2h and g5-HT2m in HEK293T cells.**

1120 **a**, Representative images showing the expression (top, with 5-HT) and responses (bottom) to 100 μ M
1121 5-HT for g5-HT2h (left) and g5-HT2m (right). Scale bar, 20 μ m.
1122 **b**, The group summary of the brightness (left), peak $\Delta F/F_0$ (middle) and SNR (right) of g5-HT2h and
1123 g5-HT2m. The SNR is relative to g5-HT1.0; a.u., arbitrary units. $n = 154$ cells from 3 coverslips (154/3)
1124 for g5-HT2h, 98/3 for g5-HT2m.

1125 **c**, Dose-dependent curves of g5-HT2h and g5-HT2m. $n = 3$ wells for each sensor with 300–500 cells
1126 per well.

1127 Data are shown as mean \pm SEM in **b,c**, with the error bars indicating the SEM.

1128

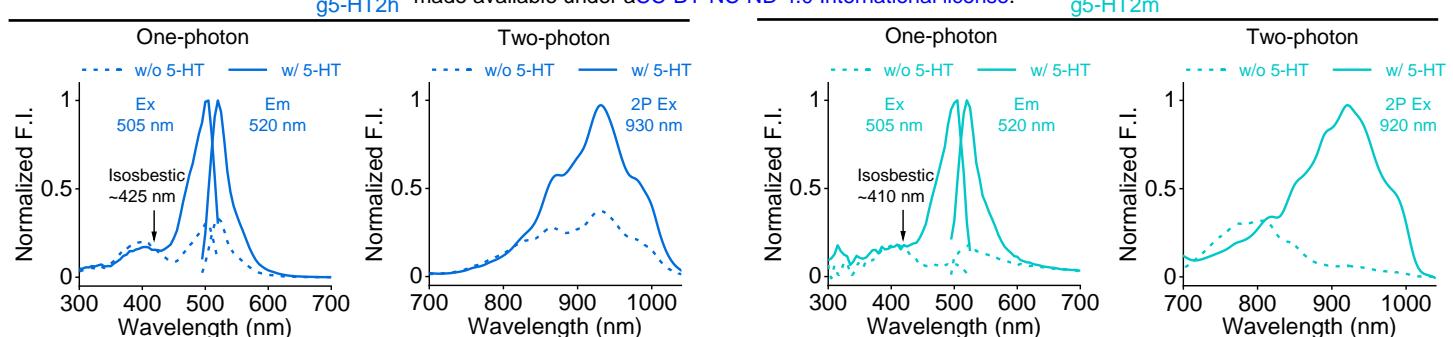


Extended Data Fig. 3 | Specificity of 5-HT sensors.

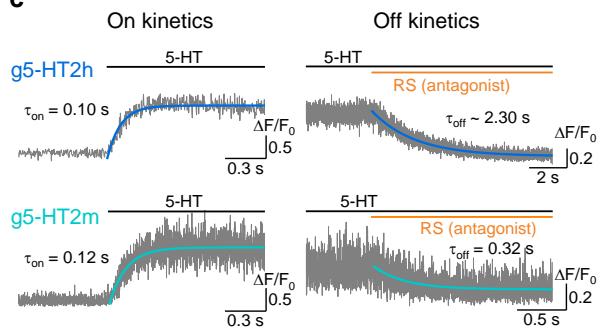
1129 **Extended Data Fig. 3 | Specificity of 5-HT sensors.**

1130 Specificity test of indicated sensors in HEK293T cells (**a, b**) or cultured rat cortical neurons (**c–f**) to 5-
1131 HT alone, 5-HT together with SB, 5-HT together with RS, and 5-HT precursor, 5-HT metabolites, as
1132 well as other neurotransmitters and neuromodulators (all compounds at 10 μ M except RS at 100 μ M).
1133 5-HTP, 5-hydroxytryptophan; 5-HIAA, 5-hydroxyindole acetic acid; DA, dopamine; NE, norepinephrine;
1134 HA, histamine; MT, melatonin; OA, octopamine; Glu, glutamate; GABA, gamma-aminobutyric acid;
1135 ACh, acetylcholine; Gly, glycine. Norm., normalized. n = 3 wells for each group with 200–500 cells per
1136 well. (One-way ANOVA followed by Tukey's multiple-comparison tests were performed, ** P < 0.01,
1137 *** P < 0.001, n.s., not significant.)
1138 Data are shown as mean \pm SEM, with the error bars indicating the SEM.
1139

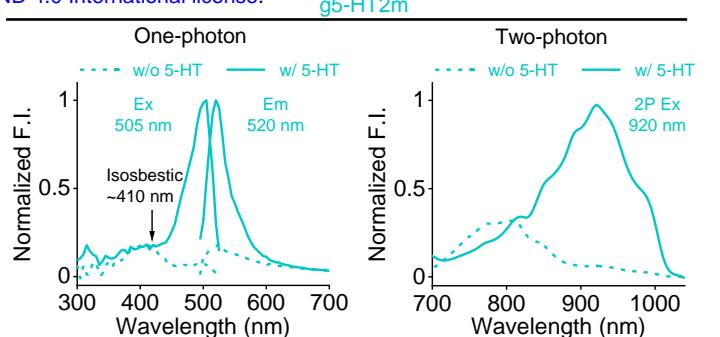
a



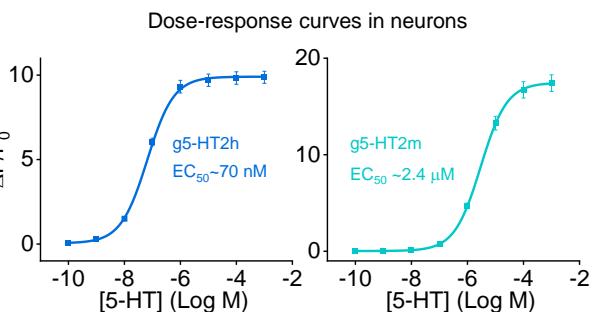
c



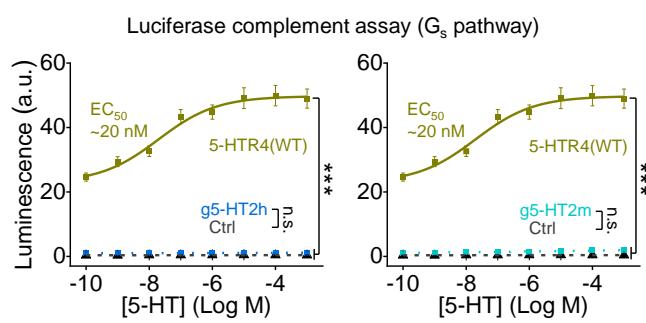
b



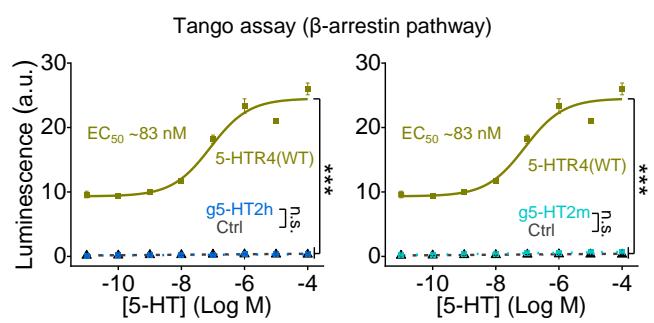
d



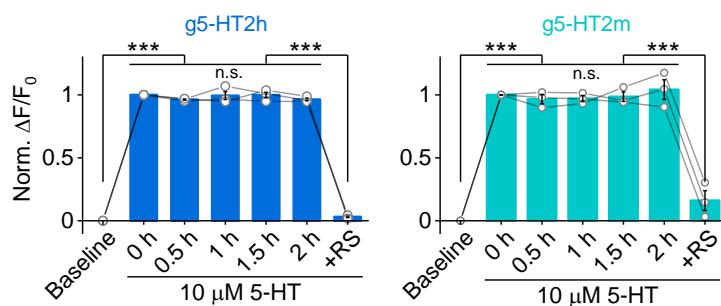
e



f



g



Extended Data Fig. 4 | Characterization of g5-HT2h and g5-HT2m in HEK293T cells and cultured rat cortical neurons.

1140 **Extended Data Fig. 4 | Characterization of g5-HT2h and g5-HT2m in HEK293T cells and**
1141 **cultured rat cortical neurons.**

1142 **a–b**, Excitation (Ex) and emission (Em) spectra of g5-HT2h (**a**) and g5-HT2m (**b**) in the absence (dash
1143 line) and presence of 10 μ M 5-HT (solid line) under one-photon (left), and two-photon excitation (right).
1144 w/o, without; w/, with.

1145 **c**, Representative traces of sensor fluorescence increase to 5-HT puffing and decrease to RS puffing
1146 (left). Group summary of on and off kinetics (right). $n = 16$ cells from 4 coverslips (short for 16/4) for
1147 g5-HT2h on kinetics, 10/3 for g5-HT2h off kinetics, 11/3 for g5-HT2m on kinetics, 9/3 for g5-HT2m off
1148 kinetics.

1149 **d**, Dose-response curves of g5-HT2h (left) and g5-HT2m (right) in cultured rat cortical neurons. $n =$
1150 60 ROIs from 3 coverslips for g5-HT2h and g5-HT2m.

1151 **e–f**, Downstream coupling tests of g5-HT2h and g5-HT2m by the luciferase complement assay for G_s
1152 coupling (**e**) and the Tango assay for β -arrestin coupling (**f**), respectively. WT, wild type; Ctrl, control,
1153 without expression of wild type 5-HTR4 or sensors; a.u., arbitrary units. Data of WT and Ctrl groups
1154 were replotted from Fig. 2I. $n = 3$ wells per group with 200–500 cells per well. (For luciferase
1155 complement assay, post hoc test in 1 mM 5-HT: $P = 2.65 \times 10^{-6}$ and 0.96 for g5-HT2h versus WT and
1156 Ctrl, respectively, $P = 2.93 \times 10^{-6}$ and 0.82 for g5-HT2m versus WT and Ctrl, respectively; for Tango
1157 assay, post hoc test: $P = 4.94 \times 10^{-8}$ and 1 for g5-HT2h versus WT and Ctrl, respectively, $P = 5.96 \times 10^{-8}$
1158 and 0.88 for g5-HT2m versus WT and Ctrl, respectively.)

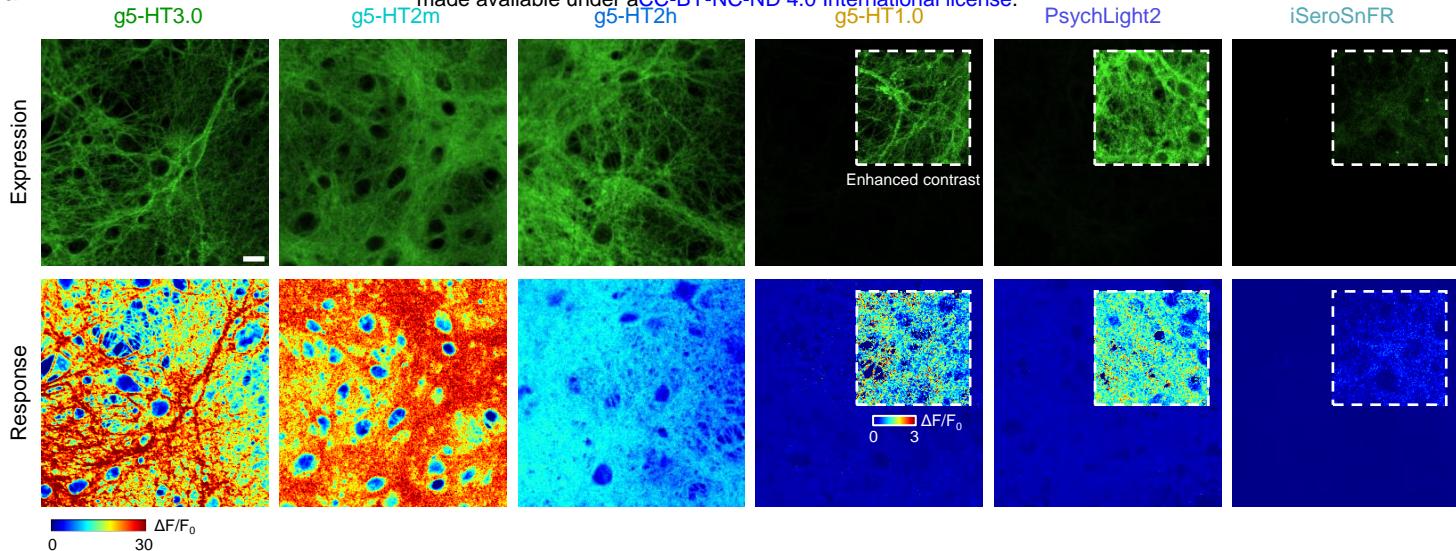
1159 **g**, The fluorescence of g5-HT2h (left) and g5-HT2m (right) expressed in cultured rat cortical neurons
1160 in response to a 2-h application of 5-HT, followed by 5-HTR4 antagonist RS. (For g5-HT2h, $F = 670$,
1161 $P = 2.83 \times 10^{-5}$, post hoc test: $P = 0$ for baseline versus 0 h, $P = 0$ for 2.0 h versus RS, $P = 0.76$, 1, 1,
1162 0.80 for 0 h versus 0.5 h, 1 h, 1.5 h or 2.0 h, respectively; for 5-HT2m, $F = 100.3$, $P = 0.006$, post hoc
1163 test: $P = 1.13 \times 10^{-6}$ for baseline versus 0 h, $P = 1.77 \times 10^{-7}$ for 2.0 h versus RS, $P = 1, 1, 1, 0.99$ for 0 h
1164 versus 0.5 h, 1 h, 1.5 h or 2.0 h, respectively.)

1165 $n = 3$ wells for each sensor. One-way ANOVA (in **e,f**) and one-way repeated measures ANOVA (in **g**)
1166 followed by Tukey's multiple-comparison tests, *** $P < 0.001$, n.s., not significant.

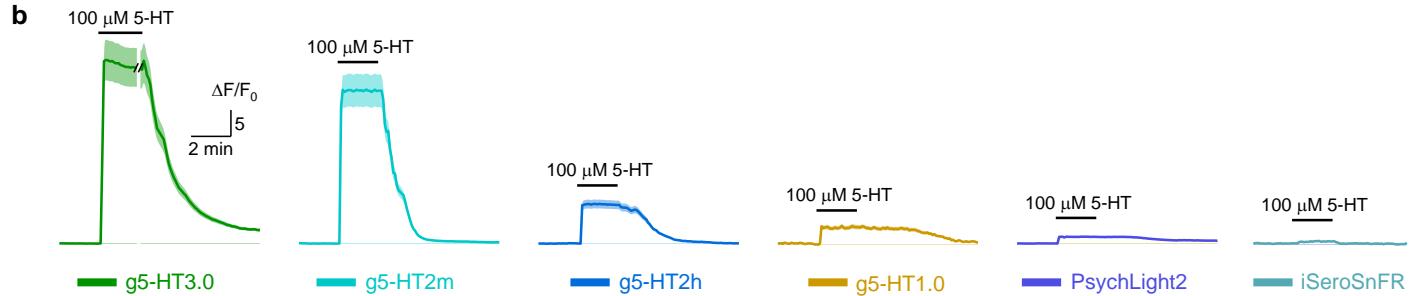
1167 Data are shown as mean \pm SEM in **c–g**, with the error bars indicating the SEM.

1168

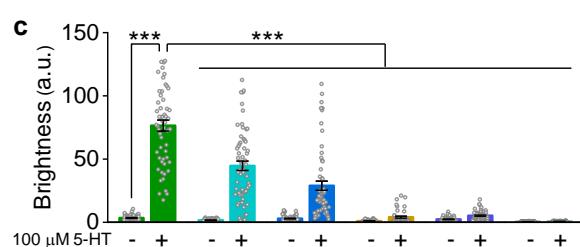
a



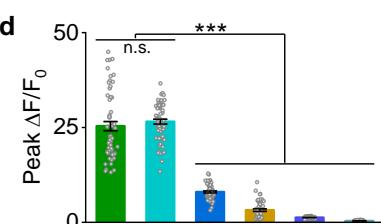
b



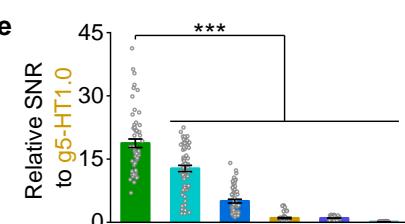
c



d



e



Extended Data Fig. 5 | Comparison of single GFP-based 5-HT sensors in cultured rat cortical neurons.

1169 **Extended Data Fig. 5 | Comparison of single GFP-based 5-HT sensors in cultured rat cortical
1170 neurons.**

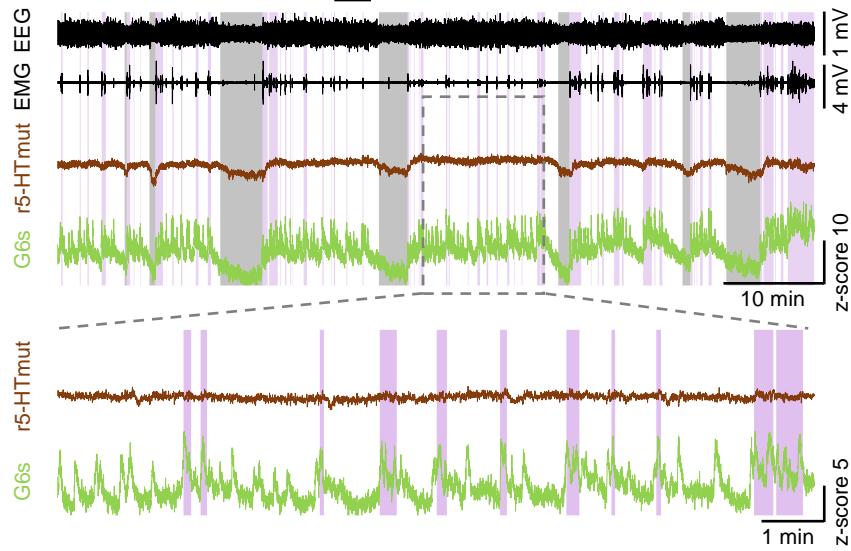
1171 **a**, Representative images showing the fluorescence expression (top) and responses (bottom) to 100
1172 μM 5-HT for different sensors as indicated. Insets with white dashed outlines in images have either
1173 enhanced contrast (top) or different pseudocolor scales (bottom). Scale bar, 20 μm .

1174 **b**, Representative traces in response to 100 μM 5-HT for different sensors as indicated.

1175 **c–e**, Group summary of the brightness (**c**), peak $\Delta F/F_0$ (**d**) and SNR (**e**). The SNR of all sensors is
1176 relative to the SNR of g5-HT1.0; a.u., arbitrary units, the basal brightness of g5-HT1.0 is set as 1. $n =$
1177 56 ROIs from 3 coverslip (short for 56/3) for g5-HT3.0, 60/3 for g5-HT2m, 60/3 for g5-HT2h, 48/3 for
1178 g5-HT1.0, 60/3 for PsychLight2 and 60/3 for iSeroSnFR. (One-way ANOVA followed by Tukey's
1179 multiple-comparison tests for **c–e**; for brightness in **c**, $F_{11,676} = 141.4$, $P = 4.97 \times 10^{-167}$, post hoc test: P
1180 $<10^{-7}$ for g5-HT3.0 with 5-HT versus g5-HT3.0 without 5-HT and other sensors with or without 5-HT;
1181 for peak $\Delta F/F_0$ in **d**, $F_{5,338} = 446.9$, $P = 1.46 \times 10^{-146}$, post hoc test: $P = 0.696$ for g5-HT3.0 versus g5-
1182 HT2m, $P < 10^{-7}$ for g5-HT3.0 and g5-HT2m versus other sensors; for relative SNR in **e**, $F_{5,338} = 195.1$,
1183 $P = 2.46 \times 10^{-97}$, post hoc test: $P < 10^{-7}$ for g5-HT3.0 versus other sensors, *** $P < 0.001$, n.s., not
1184 significant.)

1185 Data are shown as mean \pm SEM in **b–e**, with the error bars or shaded regions indicating the SEM.

1186

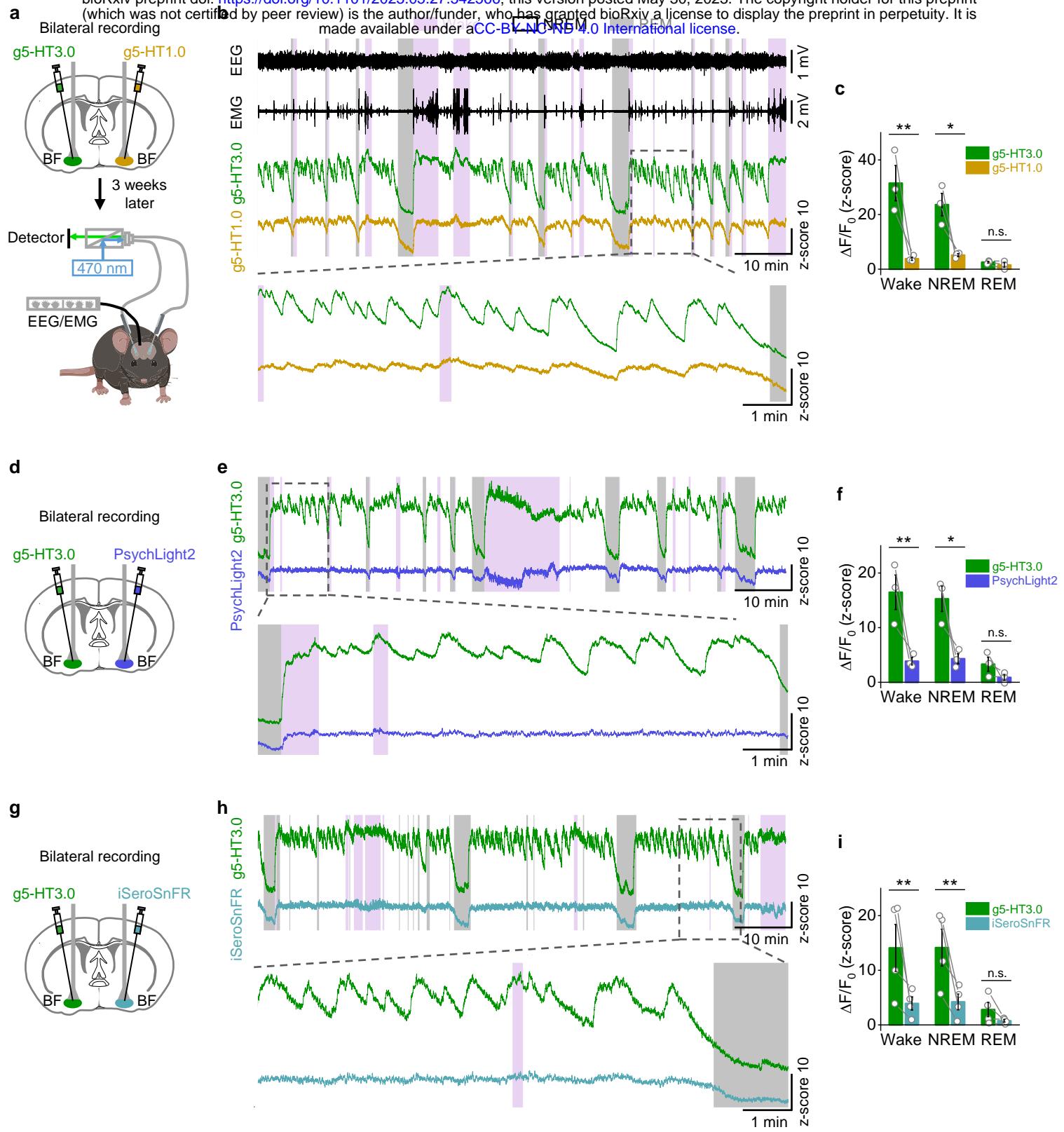


Extended Data Fig. 6 | Representative r5-HTmut and GCaMP6s signals during the sleep-wake cycle in freely moving mice.

1187 **Extended Data Fig. 6 | Representative r5-HTmut and GCaMP6s signals during the sleep-wake**
1188 **cycle in freely moving mice.**

1189 Representative r5-HTmut and GCaMP6s (G6s) traces in the mouse basal forebrain (BF) along with
1190 EEG and EMG recording during the spontaneous sleep-wake cycle.

1191



Extended Data Fig. 7 | Comparison of gGRAB_{5-HT3.0} and other green 5-HT sensors during the sleep-wake cycle in freely moving mice.

1192 **Extended Data Fig. 7 | Comparison of gGRAB_{5-HT3.0} and other green 5-HT sensors during the**
1193 **sleep-wake cycle in freely moving mice.**

1194 **a**, Schematic showing the setup of bilateral fiber-photometry recording of g5-HT3.0 and g5-HT1.0
1195 during the sleep-wake cycle in mice.

1196 **b**, Representative traces of simultaneous EEG, EMG, g5-HT3.0 and g5-HT1.0 recording during the
1197 sleep-wake cycle in freely behaving mice. Pink shading, wake state; gray shading, REM sleep.

1198 **c**, Summary of averaged g5-HT3.0 and g5-HT1.0 signals in indicated sleep-wake states. $n = 3$ mice.
1199 ($P = 0.0034$, 0.014 and 0.83 during wake, NREM and REM sleep state, respectively.)

1200 **d-f**, Similar to **a-c**, except bilateral recording of g5-HT3.0 and PsychLight2, $n = 3$ mice in **f**. ($P =$
1201 0.0066, 0.011 and 0.38 during wake, NREM and REM sleep state, respectively.)

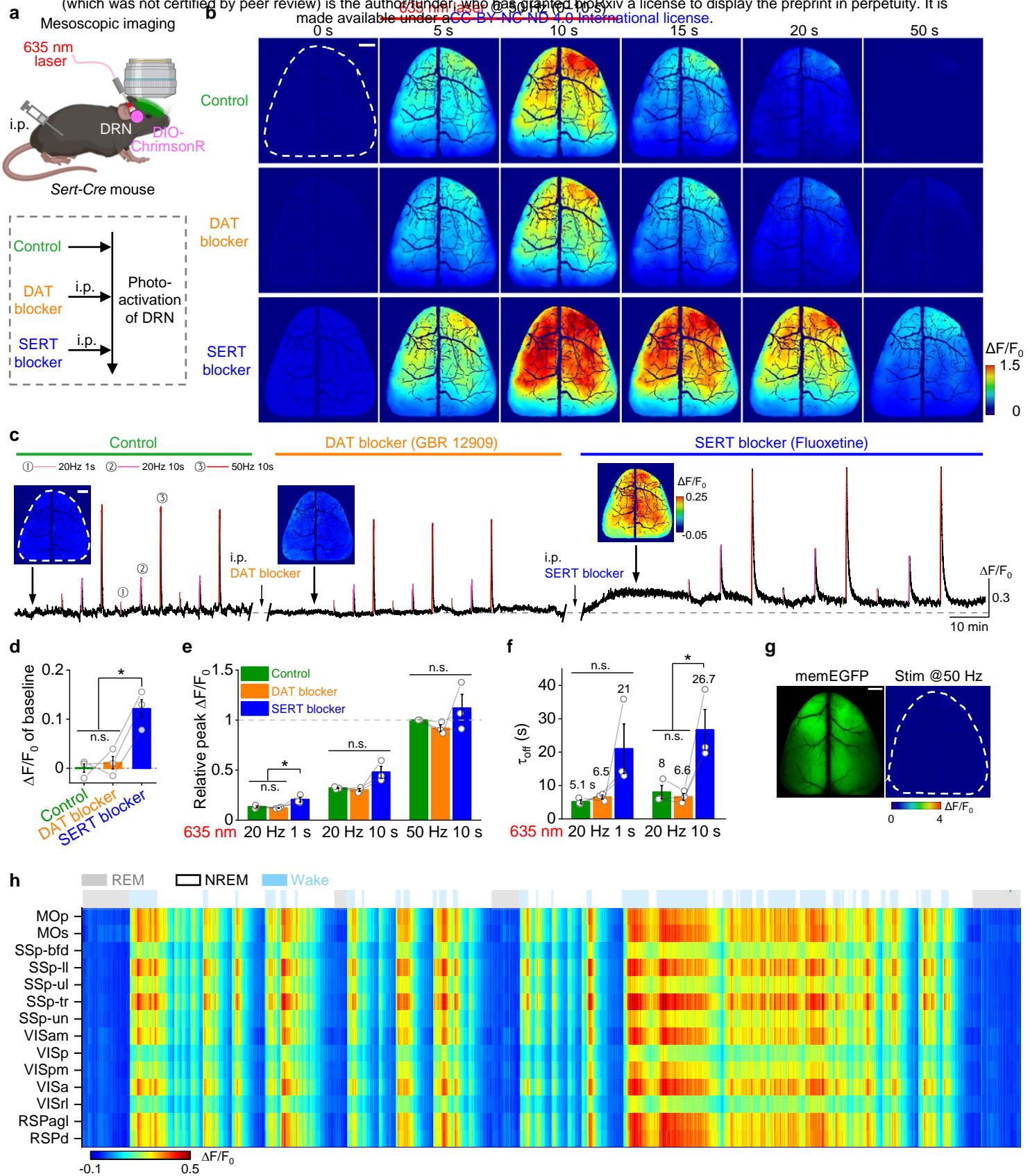
1202 **g-i**, Similar to **a-c**, except bilateral recording of g5-HT3.0 and iSeroSnFR, $n = 4$ mice in **i**. ($P = 0.0086$,
1203 0.0095 and 0.47 during wake, NREM and REM sleep state, respectively.)

1204 Data are shown as mean \pm SEM in **c,f,i**, with the error bars or shaded regions indicating the SEM.

1205 Two-way repeated measures ANOVA followed by Tukey's multiple-comparison tests in **c,f,i**, $^*P < 0.05$,

1206 $^{**}P < 0.01$, n.s., not significant.

1207



Extended Data Fig. 8 | gGRAB_{5-HT3.0} reveals 5-HT dynamics in mouse dorsal cortex *in vivo*.

1208 **Extended Data Fig. 8 | gGRAB_{5-HT3.0} reveals 5-HT dynamics in mouse dorsal cortex *in vivo*.**

1209 **a**, Schematic depicting the protocol for mesoscopic imaging along with optogenetic activation of DRN
1210 with different drug treatments.

1211 **b**, Representative pseudocolor images in response to the 50 Hz 10 s optical stimulation of DRN with
1212 indicated treatments.

1213 **c**, Representative trace of g5-HT3.0 with indicated treatments, including the application of different
1214 drugs and activation of DRN using a 635 nm laser with different frequencies and durations. Insets
1215 above the trace are averaged images in the indicated baseline timepoint (by the black arrow) of
1216 different stages.

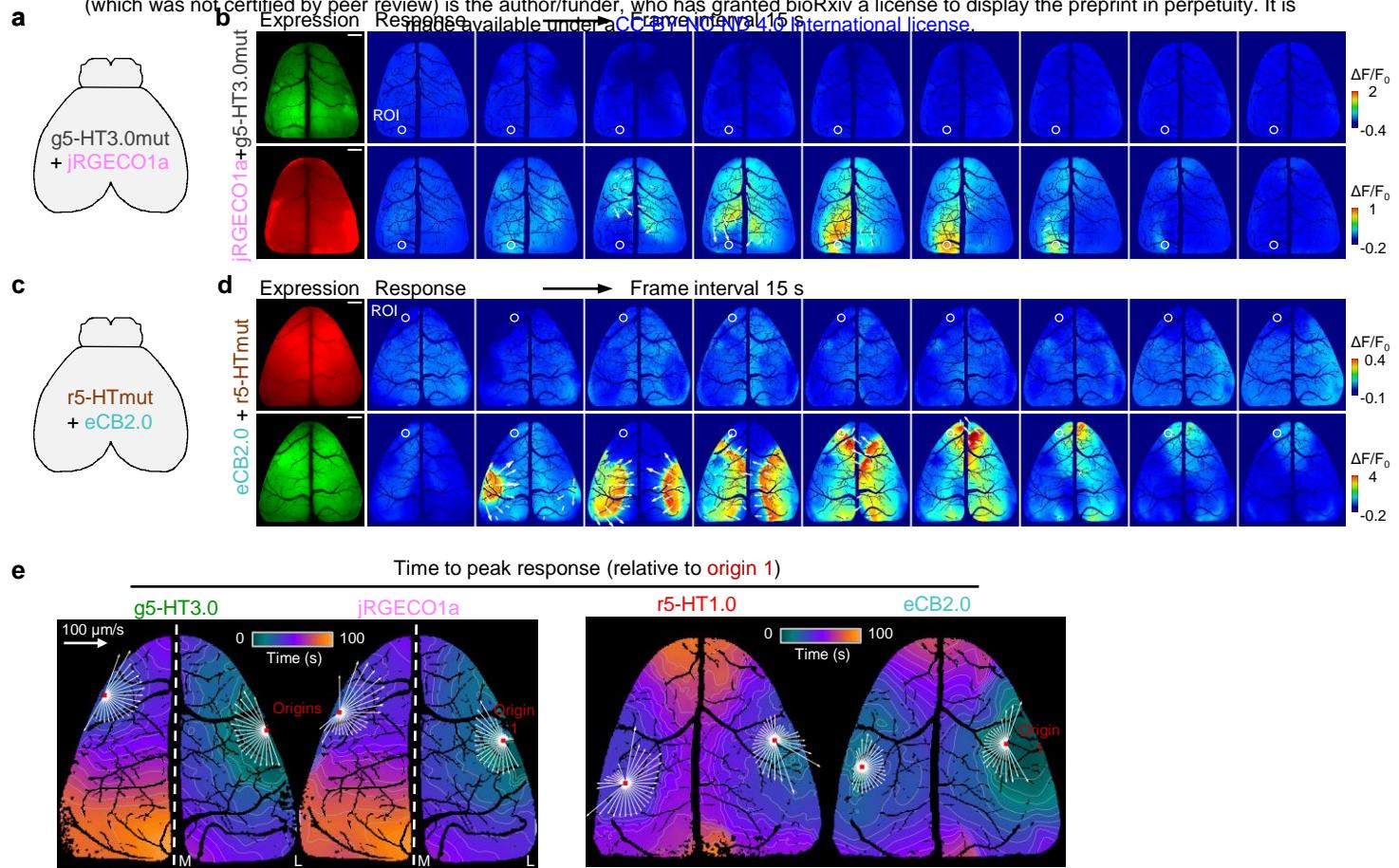
1217 **d**, Group data of averaged g5-HT3.0 baseline fluorescence changes under indicated treatments.
1218 (One-way repeated measures ANOVA followed by Tukey's multiple-comparison tests, $F = 19.9$, $P =$
1219 0.047, post hoc test: $P = 0.896$ for control versus DAT blocker, 0.016 for SERT blocker versus control
1220 and 0.022 for SERT blocker versus DAT blocker.)

1221 **e–f**, Group summary of optical stimulation evoked peak response (**e**) and decay kinetics (**f**). $n = 3$ mice
1222 in **d–f**. (One-way repeated measures ANOVA followed by Tukey's multiple-comparison tests. For
1223 relative peak $\Delta F/F_0$ in **e**, under 20 Hz 1 s stimulation, $F = 11.1$, $P = 0.023$, post hoc test: $P = 0.81$ for
1224 control versus DAT blocker, 0.043 for SERT blocker versus control and 0.026 for SERT blocker versus
1225 DAT blocker; under 20 Hz 10 s stimulation, $F = 6.67$, $P = 0.053$; under 50 Hz 10 s stimulation, $F = 1.39$,
1226 $P = 0.348$. For decay kinetics τ_{off} in **f**, under 20 Hz 1 s stimulation, $F = 4.06$, $P = 0.182$; under 20 Hz
1227 10 s stimulation, $F = 16.78$, $P = 0.011$, post hoc test: $P = 0.932$ for control versus DAT blocker, 0.018
1228 for SERT blocker versus control and 0.014 for SERT blocker versus DAT blocker.)

1229 **g**, Representative images showing the memEGFP expression and response to the 50 Hz 10s optical
1230 activation.

1231 **h**, Representative heatmap showing changes of g5-HT3.0 fluorescence in different brain regions
1232 during the sleep-wake cycle. Gray shading, REM sleep; light blue shading, wake state.
1233 The dashed white outlines **b,c,g** indicate the ROI. All scale bar, 1 mm. Data are shown as mean \pm SEM
1234 in **d–f**, with the error bars indicating the SEM, $*P < 0.05$, n.s., not significant.

1235



Extended Data Fig. 9 | Mesoscopic imaging of 5-HT, Ca^{2+} and eCB waves during seizures.

1236 **Extended Data Fig. 9 | Mesoscopic imaging of 5-HT, Ca²⁺ and eCB waves during seizures.**

1237 **a**, Schematic showing the co-expression of g5-HT3.0mut and jRGECO1a in the mouse dorsal cortex.
1238 **b**, Representative images show fluorescence changes of g5-HT3.0mut (top) and jRGECO1a (bottom)
1239 during seizures. A ROI labeled with the white circle (500 μ m in diameter) shows the maximum
1240 response regions of jRGECO1a, which corresponds to the trace in Fig. 5c. White arrows indicate the
1241 direction of wave propagation and the length of arrows indicates relative magnitudes of velocities.
1242 Scale bar, 1 mm.

1243 **c-d**, Similar to **a-b**, but co-expressing r5-HTmut and eCB2.0. The ROI shows the maximum response
1244 regions of eCB2.0 and corresponds to the trace in Fig. 5e.

1245 **e**, Representative time to peak response maps of waves relative to the origin 1, monitored by different
1246 sensors. Red dots indicate origin locations of waves; white arrows indicate velocity vectors calculated
1247 based on the propagation distance and duration along the corresponding direction; L, lateral, M,
1248 medial; scale bar of speed, 100 μ m/s.