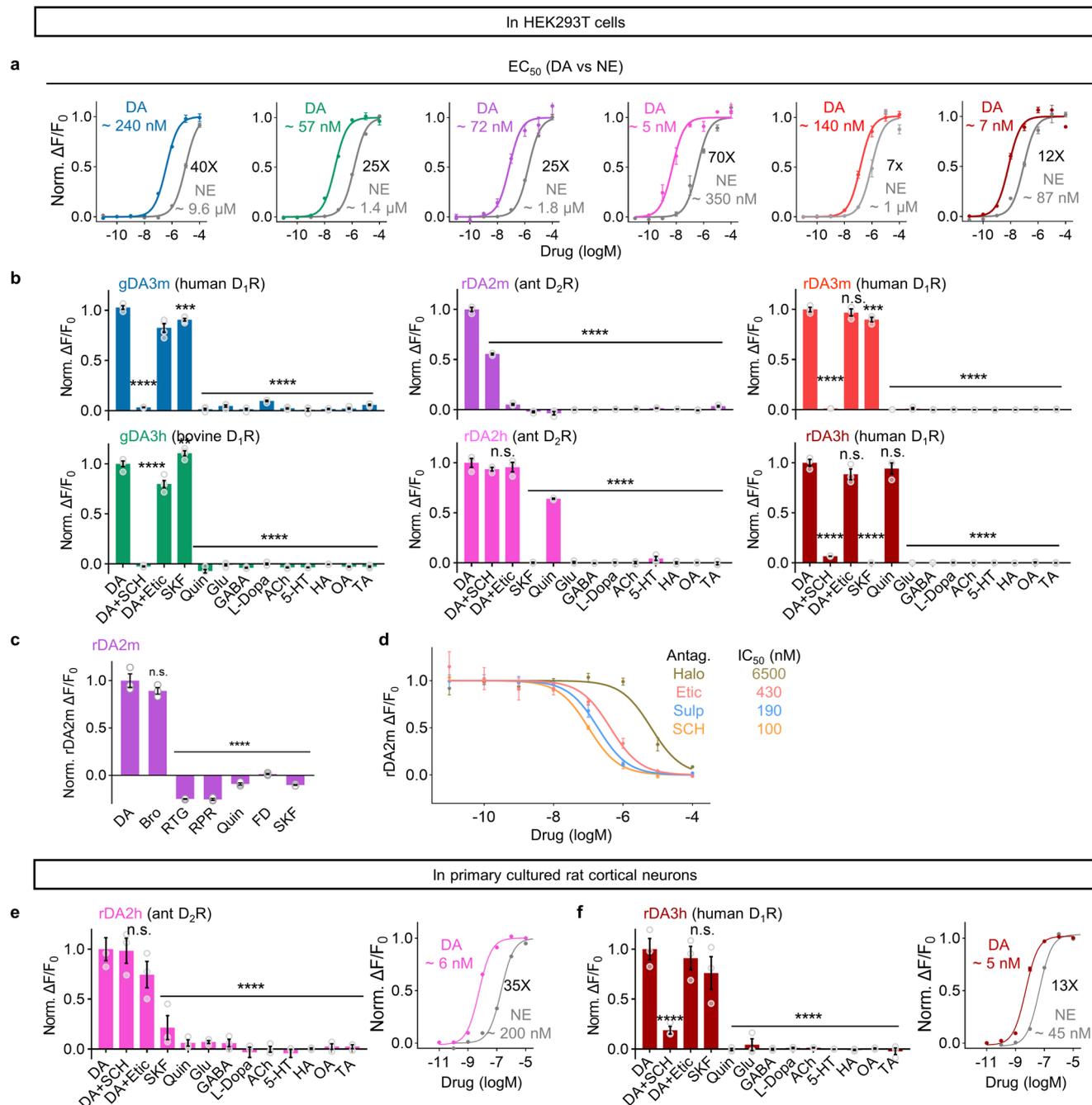


Extended Data Fig. 1 | Performance of DA-insensitive mutant sensors.

a, Representative images showing sensor expression (top) in HEK293T cells and fluorescence response to 100 μM DA (bottom) of indicated sensor variants. Scale bar, 20 μm . **b**, Group summary of maximal $\Delta F/F_0$ in response to 100 μM DA (left) and titration DA curves (right) of indicated sensors in HEK293T cells. Left, $n = 6, 6, 15, 12, 3$ wells for gDA3m, gDA3mut, rDA2m, rDA2mut, rDA3m and rDA3mut. Each well contains 400–500 cells. Two-tailed Student's *t*-test was performed.

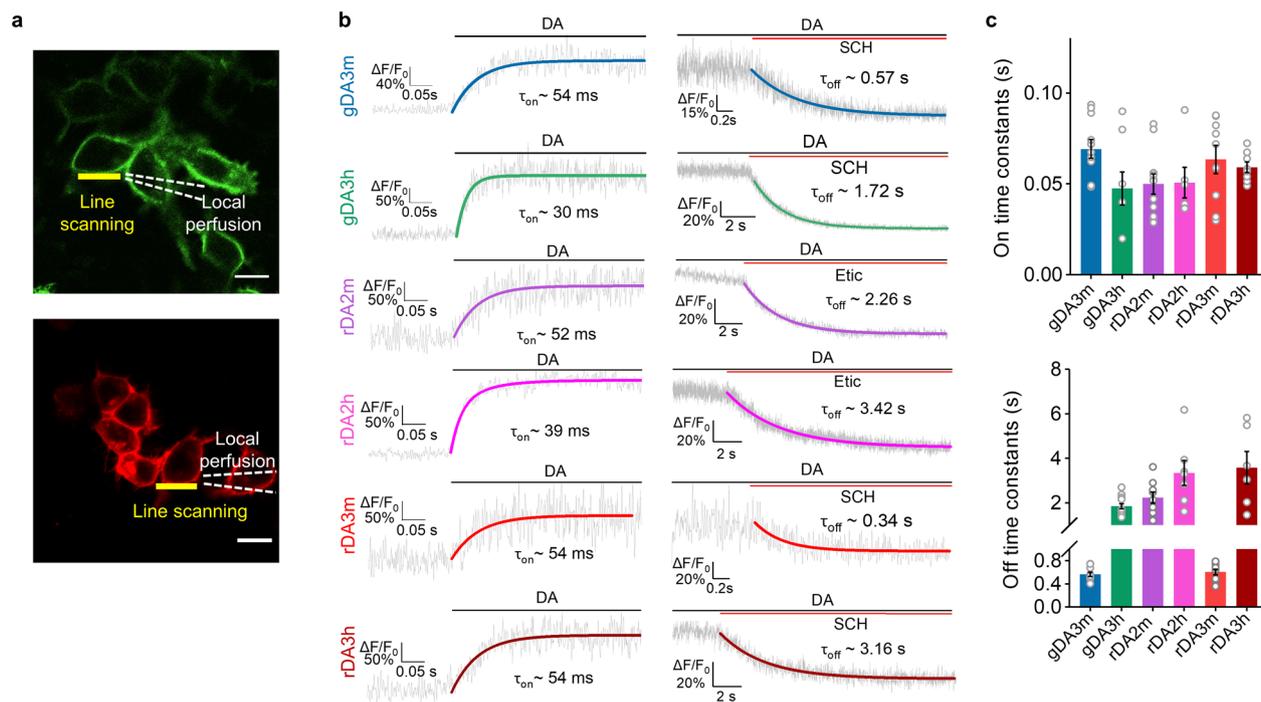
Right, $n = 3$ wells (with 400–500 cells per well) for each group. **c**, Representative images showing sensor expression (top) in cultured neurons and fluorescence response to 100 μM DA (bottom) of indicated sensor variants. Scale bar, 50 μm . **d**, Group summary of maximal $\Delta F/F_0$ of indicated sensors in response to 100 μM DA in cultured neurons. $n = 60$ neurons from 4 cultures for rDA2mut, $n = 30/2$ for others, mean \pm s.e.m. Two-tailed Student's *t*-test was performed.



Extended Data Fig. 2 | Pharmacological profiles of GRAB_{DA} sensors measured in cultured cells.

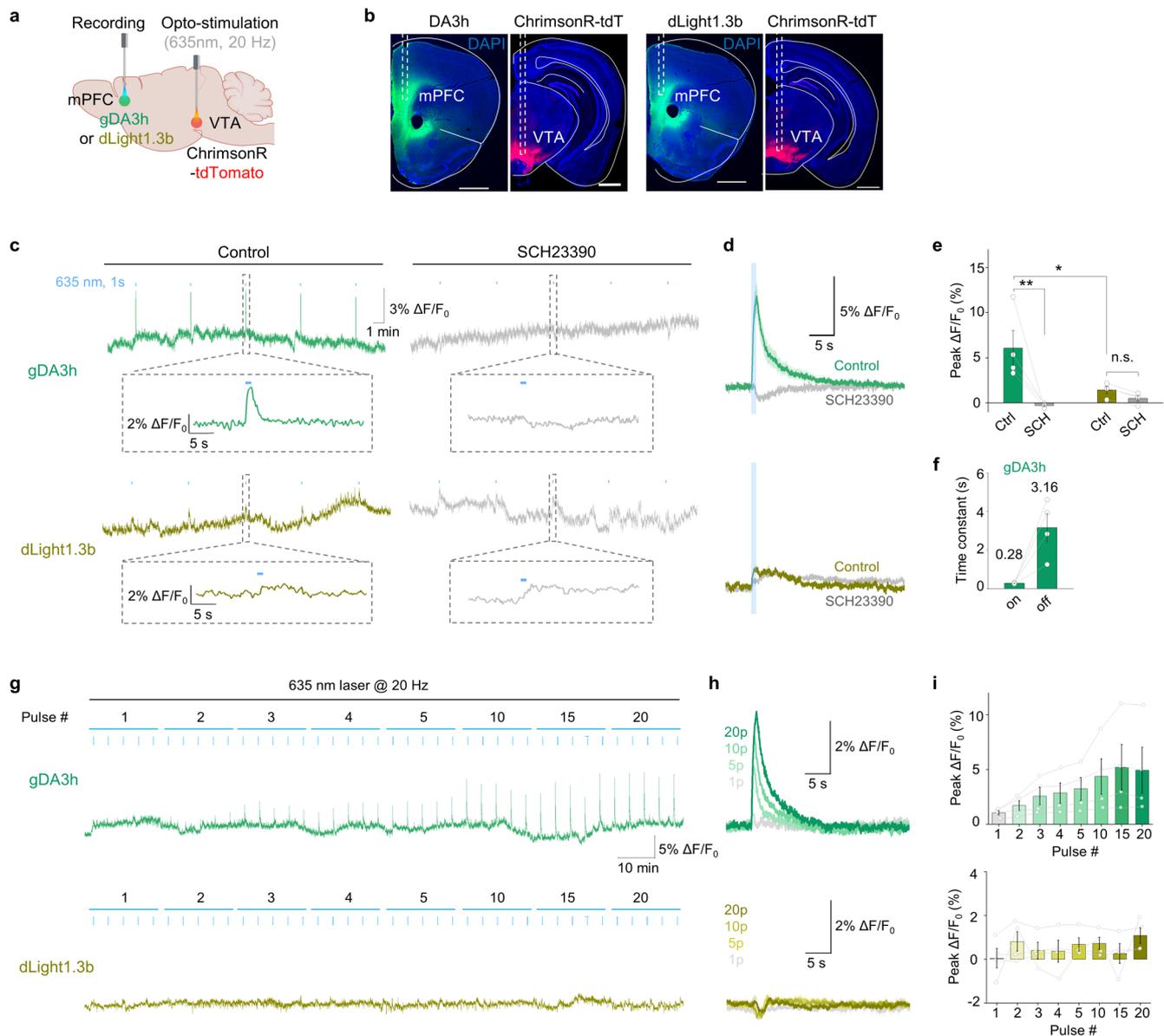
a, Titration curves of indicated sensors for the response to DA or NE in HEK293T cells. $n = 3$ cells with 400–500 cells per well, mean \pm s.e.m. **b**, The normalized $\Delta F/F_0$ in sensor-expressing HEK293T cells in response to the indicated compounds. Antagonists were applied at 10 μ M, others at 1 μ M. $n = 4$ wells for gDA3m and gDA3h, $n = 3$ wells for others, mean \pm s.e.m. One-way Anova, post hoc Dunnett's test: gDA3m, $p = 0.0002$, 8.9×10^{-10} between DA and SKF, or DA+Etic, $p = 1.3 \times 10^{-13}$ between DA and others; gDA3h, $p = 0.0020$, 4.3×10^{-9} between DA and SKF, or DA+Etic, $p = 1.3 \times 10^{-13}$ between DA and others; rDA2m, $p = 3.3 \times 10^{-14}$ between DA and others; rDA2h, $p = 0.6059$, 0.9530 between DA and DA + SCH, or DA+Etic, $p = 3.3 \times 10^{-14}$ between DA and others; rDA3m, $p = 0.9182$, 0.010 between DA and DA+Etic, or SKF, $p = 3.3 \times 10^{-14}$ between DA and others; rDA3h, $p = 0.0724$, 0.8723 between DA and DA+Etic, or Quin, $p = 3.3 \times 10^{-14}$ between DA and others. **c**, The normalized $\Delta F/F_0$ in rDA2m-expressing HEK293T cells in

response to indicated DA agonists. Bromocriptine (Bro), Rotigotine (RTG), D₂/D₁R agonists; Ropinirole (RPR), Quin, D₂R-specific agonists; Fenodopam (FD), SKF, D₁R-specific agonist. All chemicals were bath-applied in 100 μ M. $n = 3$ wells, mean \pm s.e.m. One-way Anova, post hoc Dunnett's: $p = 0.1074$ between DA and Bro, $p = 8.0 \times 10^{-12}$ between DA and others. **d**, Titration curves of indicated dopamine receptor antagonists. The fluorescence intensity in the presence with 10 μ M DA was set as F_0 and the relative fluorescence changes under indicated compound concentration were plotted. $n = 3$ wells with 400–500 cells per well, mean \pm s.e.m. **e, f**, Pharmacological specificity (left) and titration curves of indicated sensors for the response to DA or NE (right) in cultured neurons. Left, antagonists at 10 μ M, others at 1 μ M. $n = 3$ wells, mean \pm s.e.m. One-way Anova, post hoc Dunnett's test: rDA2h, $p = 0.9998$, 0.1458 between DA and DA + SCH, or DA+Etic, $p = 1.5 \times 10^{-7}$ between DA and others; rDA3h, $p = 0.9591$, 0.1309 between DA and DA+Etic, or SKF, $p = 4.0 \times 10^{-9}$ between DA and others.



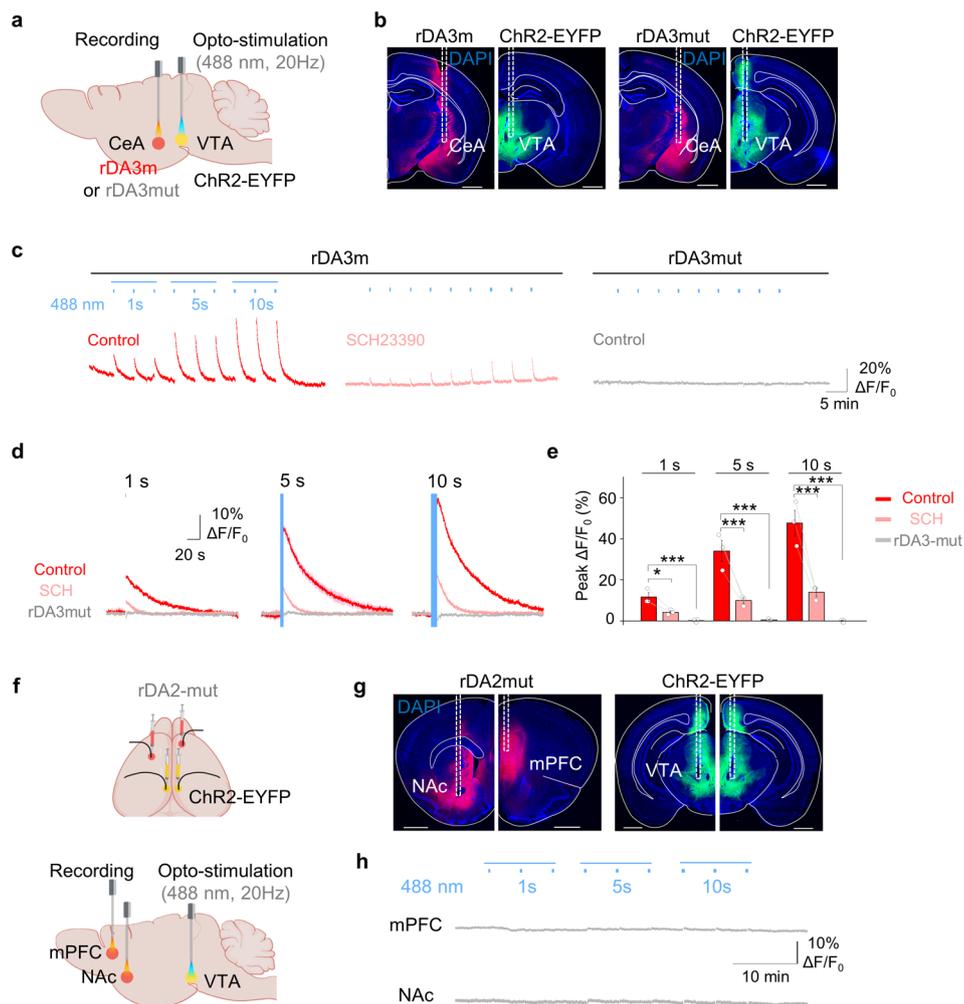
Extended Data Fig. 3 | Kinetics measurement of GRAB_{DA} sensors in HEK293T cells. **a**, Schematic illustration showing the local perfusion system using a glass pipette containing 100 μM DA and/or receptor-specific antagonist positioned above the sensor-expressing cell. The yellow line indicates the area for line scanning. The dash lines indicate the pipette. Scale bar, 20 μm. **b**, Representative traces showing the response measured using line-scanning; when indicated, DA

and receptor-specific antagonist were puffed onto the cell. The trace were the average of 3 different ROIs on the scanning line. Data are shown as mean ± SD. Each trace was fitted with a single-exponential function to determine the τ_{on} (left) and τ_{off} (right). **c**, Group summary of τ_{on} and τ_{off}. τ_{on}, $n = 11, 8, 11, 6, 9, 8$ cells for gDA3m, gDA3h, rDA2m, rDA2h, rDA3m, rDA3h; τ_{off}, $n = 10, 14, 9, 7, 10, 6$ cells for gDA3m, gDA3h, rDA2m, rDA2h, rDA3m, rDA3h, mean ± s.e.m.



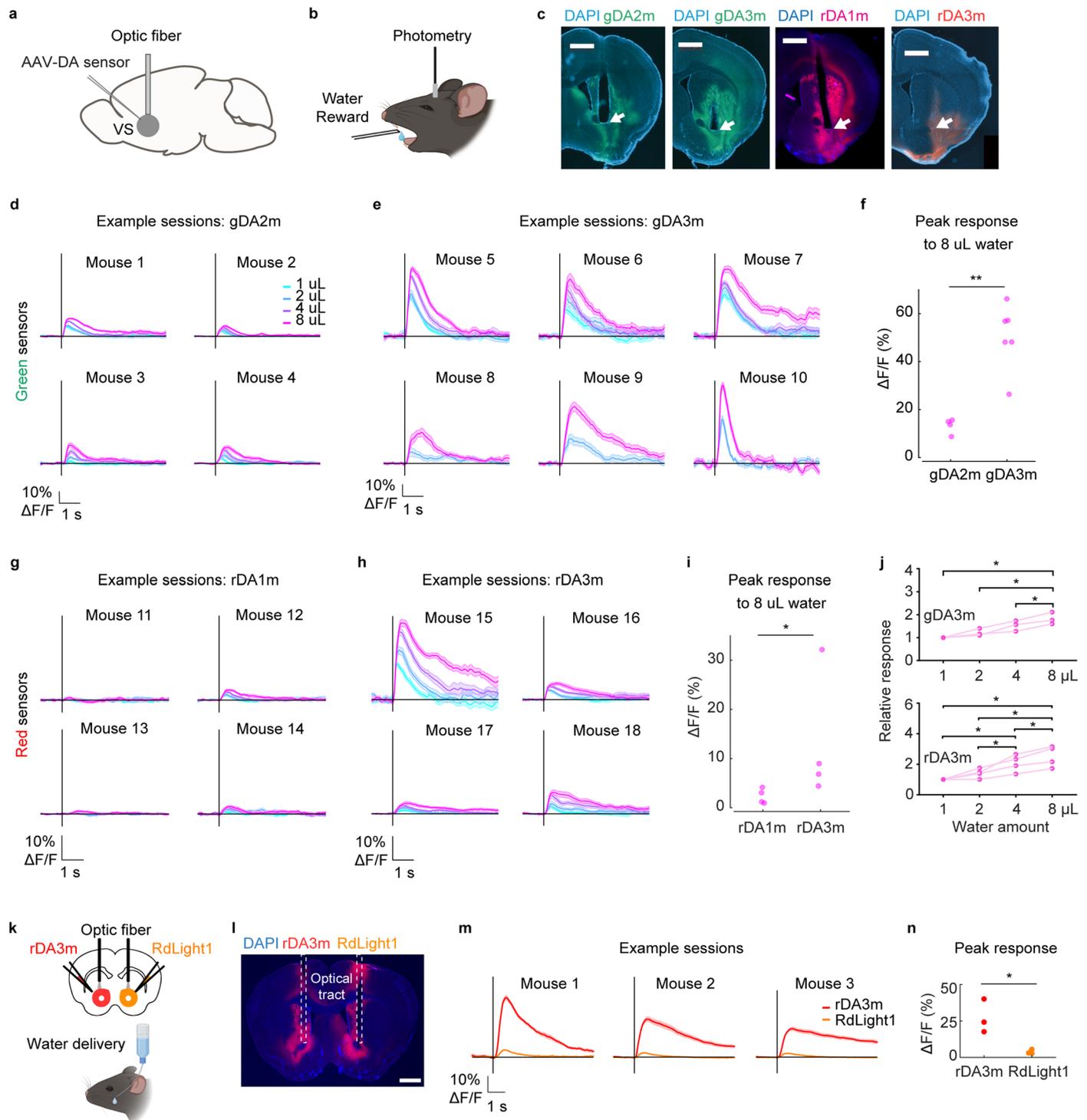
Extended Data Fig. 4 | gGRAB_{DA3h} sensors report optogenetically-elicited DA release in the mouse mPFC. **a**, Schematic illustration depicting the experimental design for panel **b-i**. **b**, Histological verification of indicated sensor expression in mPFC and ChrimsonR expression in VTA. Dashed boxes indicate the location of optical tract. Scale bar, 1 mm. **c**, Representative fluorescence changes and zoom-in view (indicated by dashed box) of indicated sensors during optogenetic stimulations under control condition or in the presence of SCH-23390 (SCH). **d**, Average traces of the change in gDA3h (top) or dLight1.3b (bottom) fluorescence from a mouse. Data are shown as mean±s.d. **e**, Group summary of $\Delta F/F_0$ for the indicated sensors. $n = 4$ mice for gDA3h and dLight1.3b,

respectively, mean±s.e.m. One-way ANOVA, post hoc Tukey's test was performed. ** $p = 0.0035$ for gDA3h; n.s. $p = 0.9122$ for dLight1.3b; * $p = 0.0295$ between gDA3h and dLight1.3b. **f**, Group summary of the rise and decay time constant of the gDA3h signals in response to optogenetic stimulations. $n = 4$ mice, mean±s.e.m. **g, h**, Example fluorescence response (**g**) and corresponding average traces (**e**) of gDA3h (top) or dLight1.3b (bottom) to indicated optogenetic stimulation. The average traces are shown as mean±s.d. **i**, Group summary of peak $\Delta F/F_0$ of gDA3h or dLight1.3b in response to indicated optogenetic stimulation. $n = 4$ mice for gDA3h and dLight1.3b, mean±s.e.m.



Extended Data Fig. 5 | rGRAB sensors report optogenetically-elicited DA release in multiple brain regions *in vivo*. **a**, Schematic illustration depicting the experimental design for panel **b-e**. **b**, Histological verification of indicated sensor expression in CeA and ChrimsonR expression in VTA. Dashed boxes indicate the location of optical tract. Scale bar, 1 mm. **c**, Representative traces of rDA3m or rDA3mut signals during optogenetic stimulations. rDA3m signals were measured before and after SCH-23390 (SCH) administration. **d**, Average traces of the change in sensor fluorescence to 1-, 5- or 10-s opto-stimulation from a mouse. Data are shown as mean \pm SD. The blue shaded area indicates the application of opto-stimulation. **e**, Group summary of peak response of

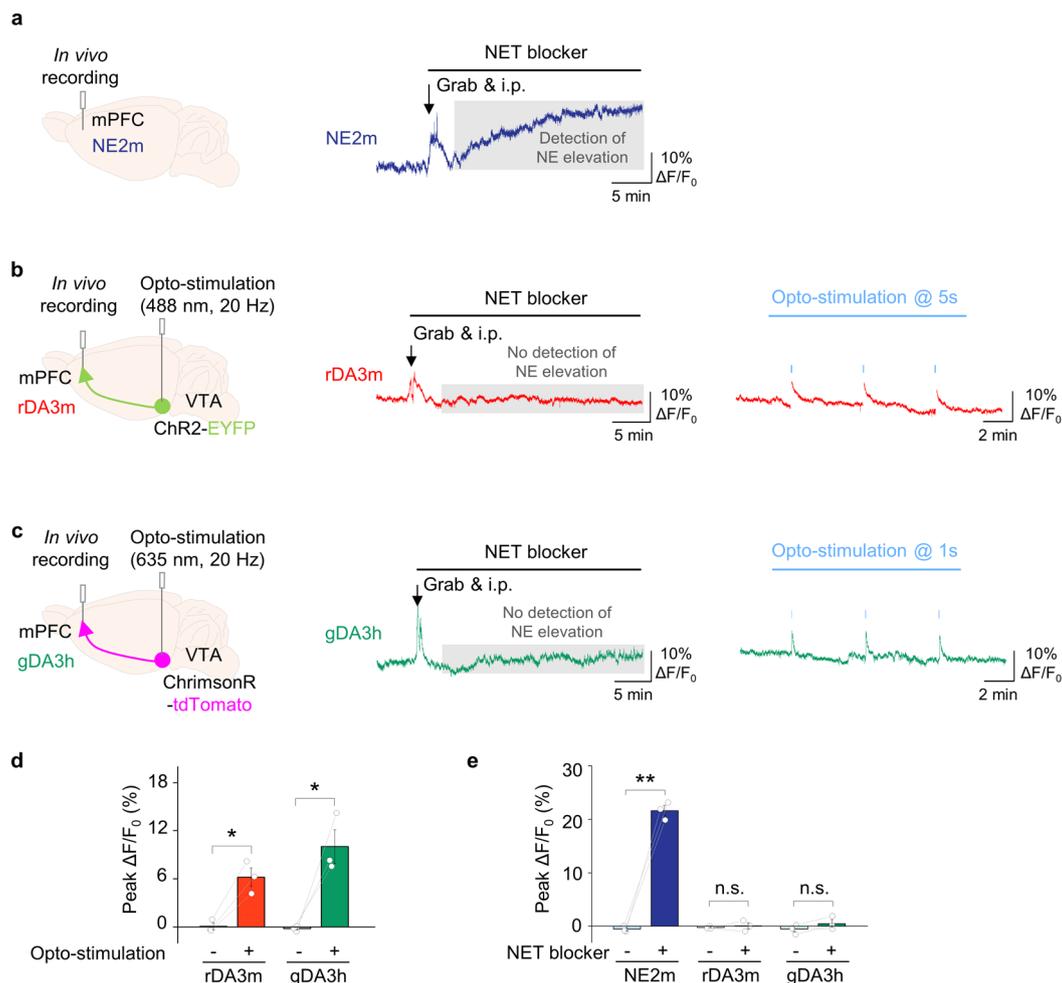
rDA3m or rDA3mut to indicated optogenetic stimulation. $n = 3$ mice for rDA3m and $n = 5$ for rDA3mut, mean \pm s.e.m. Two-tailed Student's *t*-test was performed. $p = 0.0278, 0.0101, 0.0068$ between control and SCH to 1-, 5-, 10-s opto-stimulation. $p = 0.0003, 0.0001, 0.00004$ between rDA3m and rDA3mut to 1-, 5-, 10-s opto-stimulation. **f**, Schematic illustration depicting the experimental design for panel **g**. **g**, Histological verification of rDA2mut expression in mPFC and NAc, and ChrimsonR expression in VTA. Dashed boxes indicate the location of optical tract. Scale bar, 1 mm. **h**, Representative traces of rDA2mut signals simultaneously recorded in the mPFC (top) and NAc (bottom) during optogenetic stimulations.



Extended Data Fig. 6 | *In vivo* comparison of the third-generation DA sensors versus previous variants in water-restricted mice receiving water rewards.

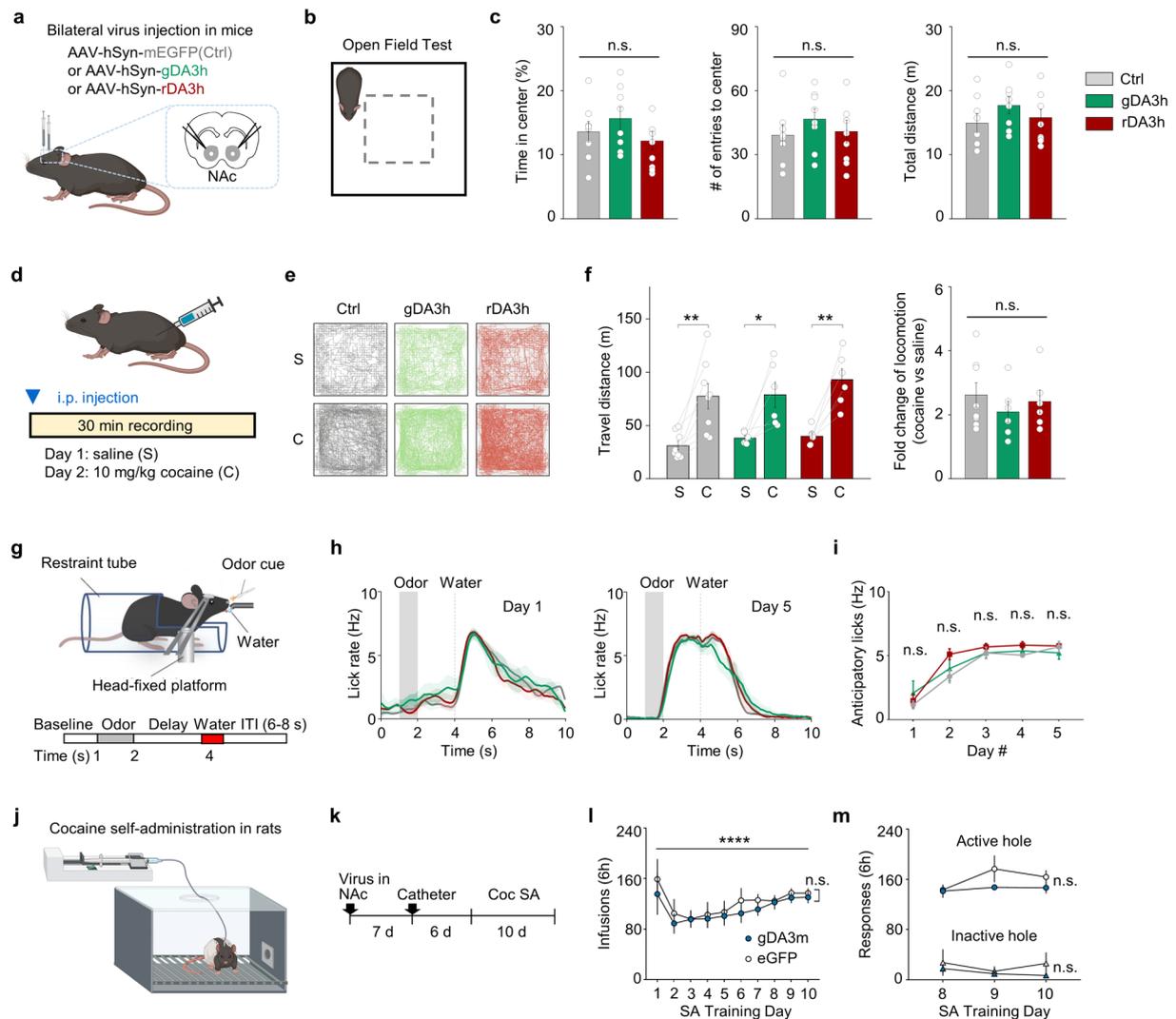
a, Diagram of mouse surgical procedure. AAVs carrying gGRAB_{DA2m}, gGRAB_{DA3m}, rGRAB_{DA1m}, or rGRAB_{DA3m} were injected unilaterally into NAc. An optic fiber was implanted above the injection site. **b**, Illustration of behavioral experiment. **c**, Histological verification of indicated sensor expression in NAc. White arrows indicate the location of fiber tips. Scale bar, 1 mm. **d**, Recording sessions from gDA2m mice, mean \pm s.e.m. Vertical black bars indicate water delivery. Colors indicate water volume. **e**, Recording sessions from gDA3m mice, mean \pm s.e.m. Vertical black bars indicate water delivery. Colors indicate water volume. **f**, Peak response to 8 μ L water for the sessions shown in **d** and **e**. $n = 4$ mice for gDA2m, $n = 6$ for gDA3m. $p = 0.0095$, Two-tailed Mann-Whitney U test. **g**, Recording sessions from rDA1m mice, mean \pm s.e.m. Vertical black bars indicate water delivery. Colors indicate water volume. **h**, Recording sessions from rDA3m mice, mean \pm s.e.m. Vertical black bars indicate water delivery. Colors indicate water volume. **i**, Peak response to 8 μ L water for the sessions shown in

g and **h**. $n = 4$ for rDA1m, $n = 4$ for rDA3m. $p = 0.0286$, Two-tailed Mann-Whitney U test. **j**, Group summary of sensor responses to each water amount. The response of each mouse was relative to that to 1 μ L water reward. $n = 3$ mice for gDA3m, $n = 4$ mice for rDA3m. Two-tailed Student's t-test was performed between groups. For gDA3m, $p = 0.032$, 0.0185 and 0.0312 between 1, 2, 4 vs 8 μ L, respectively; For rDA3m, $p = 0.0217$, 0.017 and 0.0179 between 1, 2, 4 vs 8 μ L, and $p = 0.032$ and 0.0378 between 1, 2 vs 4 μ L respectively. **k**, Schematic illustration depicting the mouse surgical procedure and the experimental design for panel **l-n**. **l**, Histological verification of rDA3m (left side) and RdLight1 (right side) in NAc. Dashed boxes indicate the location of optical tract. Scale bar, 1 mm. **m**, Recording sessions from 3 mice. Vertical black bars indicate water delivery. Colors indicate sensor version. **n**, Peak response of rDA3m and RdLight1 for the sessions shown in **m**. $n = 3$ mice for rDA3m and RdLight1. $p = 0.0249$, Two-tailed Student's t-test.



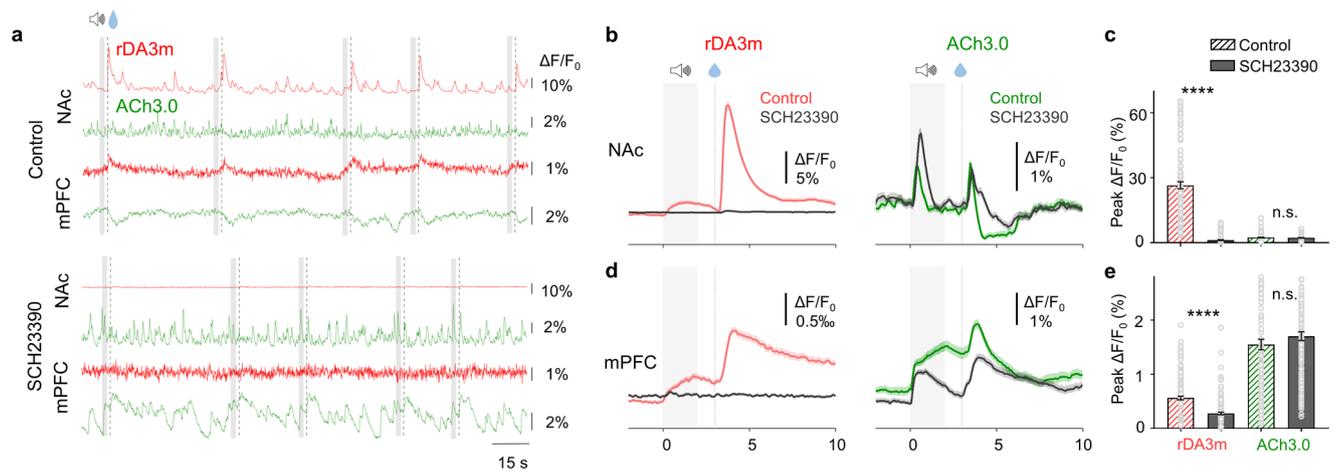
Extended Data Fig. 7 | GRAB_{DA} sensors show minimal signal changes towards endogenous NE elevation. **a**, Schematic illustration (left) depicting the experimental design and representative fluorescence changes of NE2m towards systematic administration of NET blocker (3 mg/kg desipramine). **b**, Schematic illustration (left) depicting the experimental design and representative fluorescence changes of rDA3m towards systematic administration of NET blocker (middle) and upon optogenetic stimulations (right). **c**, Schematic illustration (left) depicting the experimental design and representative fluorescence changes (right) of gDA3h towards systematic administration of NET

blocker (middle) and upon optogenetic stimulations (right). **d**, Group summary of $\Delta F/F_0$ for the indicated sensors upon opto-stimulation of VTA neurons. $n = 3$ mice for rDA3m and gDA3h, mean \pm s.e.m. Paired two-tailed Student's *t*-test was performed within group. * $p = 0.0167$ for rDA3m and * $p = 0.0463$ for gDA3h. **e**, Group summary of $\Delta F/F_0$ for the indicated sensors towards systematic administration of NET blocker. $n = 3$ mice for NE2m, rDA3m and gDA3h, respectively, mean \pm s.e.m. Paired two-tailed Student's *t*-test was performed within group. $p = 0.6827$, 0.2155 and 0.0012 for rDA3m, gDA3h and NE2m.



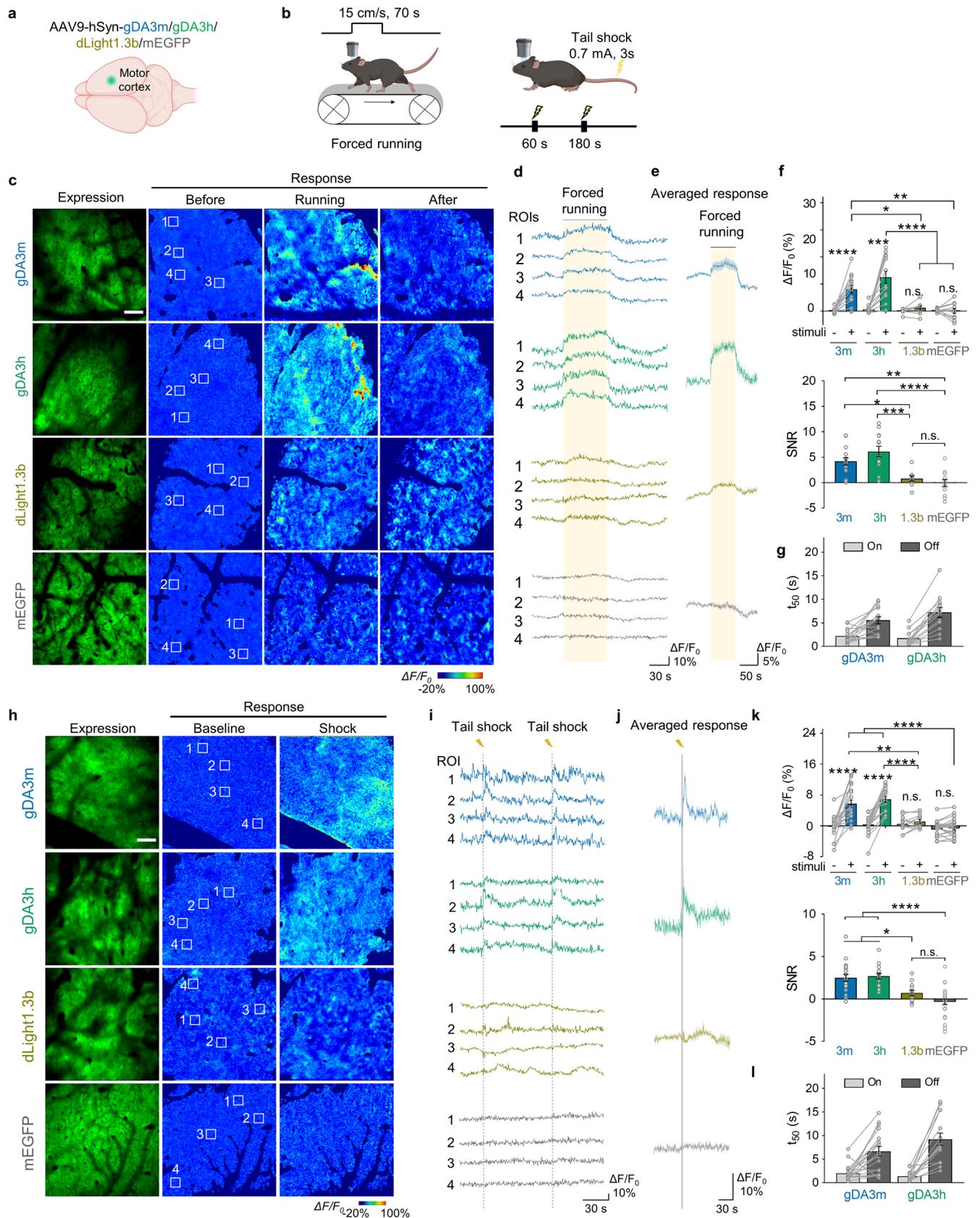
Extended Data Fig. 8 | GRAB_{DA} expression in NAc has minimal effects on DA-related animal behaviors. **a**, Schematic representation of viral injections in the bilateral NAc. **b**, Schematic illustration showing the open field test (OFT). **c**, Quantification of behavioral parameters in the OFT. $n = 8, 8$ and 10 mice for the control, gDA3h and rDA3h group, respectively, mean \pm s.e.m. One-way ANOVA was performed. $p = 0.3118, 0.5870$ and 0.3736 . **d**, Schematic illustration depicting the experimental designs for panel **e-f**. **e**, Representative track of control, gDA3h and rDA3h animals. **f**, Quantification of behavioral parameters during the experiments. $n = 8, 6$ and 6 mice for the control, gDA3h and rDA3h group, respectively, mean \pm s.e.m. Two-tailed Student's t tests were performed within groups: $p = 0.0026, 0.0196$ and 0.0039 ; One-way ANOVA was performed among groups: $p = 0.6016$. **g**, Schematic illustration showing the odor-reward associative learning task. **h**, Mean lick rate of Ctrl, gDA3h and rDA3h mice on day 1 and day 5 conditioning. $n = 5$ mice for each, mean \pm s.e.m. **i**, Quantification

of anticipatory lick rate across five conditioning days. $n = 5$ mice for each, mean \pm s.e.m. Two-way ANOVA was performed among groups, $p = 0.1076$. **j**, Schematic diagram illustrating intravenous cocaine self-administration in rats. **k**, Timeline describing intravenous cocaine self-administration experiments. **l**, Cocaine infusions over 10 days of SA training did not differ between rats expressing gDA3m and eGFP virus bilaterally in the NAc core. $n = 8$ rats for each, mean \pm s.e.m. Two-way ANOVA mixed-effects model (Day \times Virus): Day, $F(9,126) = 4.50, p = 0.00004$; Virus, $F(1,14) = 0.35, p = 0.56$; Day \times Virus, $F(9,126) = 0.21, p = 0.99$. **m**, Nose-pokes in the active and inactive ports over the last 3 days of SA training did not differ between virus groups. Two-way ANOVA mixed-effects models (Day \times Virus). $n = 8$ rats for each, mean \pm s.e.m. Active port: Day, $F(2,28) = 3.21, p = 0.06$; Virus, $F(1,14) = 1.35, p = 0.27$; Day \times Virus, $F(2,28) = 1.48, p = 0.24$. Inactive port: Day, $F(2,28) = 1.97, p = 0.16$; Virus, $F(1,14) = 0.48, p = 0.50$; Day \times Virus, $F(2,28) = 0.92, p = 0.41$.



Extended Data Fig. 9 | The signals in the mouse NAc and mPFC during Pavlovian conditioning. **a**, Representative fluorescence signals recorded during consecutive water trials pre (top, control) and post SCH-23390 (bottom, SCH-23390) treatment. The audio and water delivery are indicated above. **b**, Averaged traces of rDA3m (left) and ACh3.0 (right) fluorescence measured in the NAc from a mouse under control condition or in the presence of SCH-23390 in one mouse, mean ± s.e.m. The grey shaded area indicates the application of audio. The dashed

line indicates the delivery of water. **c**, Group summary of the peak fluorescence change of rDA3m and ACh3.0 signals in the NAc under the indicated condition. $n = 155$ trials from 3 mice for each group, mean ± s.e.m. Two-tailed Student's *t*-test was performed between control and SCH-23390 group. $p = 0.2624$ for ACh3.0. **d**, **e**, same as (b, c) with simultaneously recorded rDA3m and ACh3.0 signals in the mPFC. Two-tailed Student's *t*-test was performed between control and SCH-23390 group. $p = 0.2274$ for ACh3.0.



Extended Data Fig. 10 | See next page for caption.

Extended Data Fig. 10 | *In vivo* two-photon imaging of cortical DA dynamics in mice. **a, b**, Schematic illustration depicting the experimental design for panel **c–j**. **c–e**, Representative expression and pseudocolored response images (**c**), representative traces measured at the indicated ROIs (**d**), and average traces per forced running (**e**) measured in the motor cortex expressing indicated sensors. Scale bar, 100 μm . **f**, Group summary of the peak response (top) and SNR (bottom) of indicated sensors measured during forced running. $n = 14/4$ (14 trials from 4 mice), 13/4, 9/3 and 12/4 for gDA3m, gDA3h, dLight1.3b and mEGFP, respectively, mean \pm s.e.m. Paired two-tailed Student's *t*-test was performed for response: $p = 6 \times 10^{-5}$, 0.0002, 0.0683, 0.6275 for gDA3m, gDA3h, dLight1.3b and mEGFP. One-way ANOVA, post hoc Tukey's test was performed across groups: response, $p = 9 \times 10^{-5}$, 4×10^{-6} , 0.0214, 0.0022, 0.1611 and 0.9577 between gDA3h and dLight1.3b, gDA3h and EGFP, gDA3m and dLight1.3b, gDA3m and mEGFP, gDA3m and gDA3h, and dLight1.3b and mEGFP, respectively; SNR, $p = 9 \times 10^{-6}$, 0.0004, 0.0016, 0.0337 and 0.8812 between gDA3h and mEGFP, gDA3h and dLight1.3b, gDA3m and mEGFP, gDA3m and dLight1.3b, and dLight1.3b and

mEGFP, respectively. **g**, Summary of the rise and decay t_{50} values of indicated sensors to forced running. $n = 14/4$ for gDA3m, $n = 13/4$ for gDA3h, mean \pm s.e.m. **h–j**, Same as (**c–e**) except mice were subjected to tail shock. **k**, Group summary of the response (top) and SNR (bottom) of indicated sensors measured upon tail shock. $n = 19/4$ for gDA3m, 16/4 for gDA3h, 12/3 for dLight1.3b, 26/4 for mEGFP, mean \pm s.e.m. Paired two-tailed Student's *t*-test was performed for response: $p = 3 \times 10^{-5}$, 4×10^{-7} , 0.1774 and 0.2554 for gDA3m, gDA3h, dLight1.3b and mEGFP. One-way ANOVA, post hoc Tukey's test was performed across groups: response, $p = 8 \times 10^{-5}$, 1×10^{-8} , 0.0013, 4×10^{-8} , 0.7169 and 0.3714 between gDA3h and dLight1.3b, gDA3h and EGFP, gDA3m and dLight1.3b, gDA3m and mEGFP, gDA3m and gDA3h, and dLight1.3b and mEGFP, respectively; SNR, $p = 1 \times 10^{-7}$, 0.0104, 1×10^{-6} , 0.0186 and 0.2607 between gDA3h and mEGFP, gDA3h and dLight1.3b, gDA3m and mEGFP, gDA3m and dLight1.3b, and dLight1.3b and mEGFP, respectively. **l**, Summary of the rise and decay t_{50} values of indicated sensors to tail shock. mEGFP data replotted from Fig. 6f. $n = 18/4$ for gDA3m, $n = 15/4$ for gDA3h, mean \pm s.e.m.