

Figure S1. ADePT photo-stimulation rig, mapping DMD-to-CCD-to-2p images, and calibration of the DMD-modulated photo-stimulation patterns. Corresponding to Fig. 1. (a,b) Photostimulation and imaging rig pictures: CCD1/CCD2 – primary and secondary CCD cameras; DM, dichroic mirrors; DMD, digital micro-mirror device; I, iris diaphragm; L1-L12, lenses; PMT, photomultiplier tube; PS, periscope. Photo-stimulation (2D-light mask from a digital micro-mirror device illuminated with a solid-state CW 480 nm laser) and two photon imaging are confined to different optical z-planes that can be flexibly and independently adjusted by translating the diffuser and respectively the primary objective. (c) Fluorescence image of DMD-modulated light checkerboard projected on a 1 µm thin rhodamine B coated slide across the 1.2 x 1.5 mm field of stimulation (FOS) acquired using the secondary (inverted) microscope (CCD2). (d) DMD chipto-CCD camera-to-2p microscope registration. (i) two photon fluorescence image of pollen grains; (ii) widefield fluorescence image (full field illumination) of a larger field of view including the pollen grains from (i) marked by the square; (iii) ROIs selected from the two photon image were used to generate light masks on the DMD-chip that were further projected at the focal plane of the principal objective and imaged using the primary CCD camera (CCD 1); (iv) overlay of the DMD-chip generated photo-stimulation masks and widefield fluorescence image of pollen grains; fiduciary arrows mark indicate the location of same two pollen grains in the widefield fluorescence and two photon images. (e) Applying a correction mask for light power inhomogeneities arising due to the Gaussian profile of the photo-stimulation laser beam and aberrations along the optical path. A power distribution map (PDM) across the field of stimulation was obtained by sequentially projecting 25 µm spots in an 11 x 11 grid pattern covering 121-pixel coordinates across the DMDchip, and by linearly interpolating such as to estimate the power at the remaining positions. A power meter (Thorlabs) 50 mW sensor was placed in the focal plane of the primary objective with the projected DMD pattern in focus (Methods; Left/Right: uncorrected vs. corrected images). (f) The PMT output displayed a slowly-decaying (phosphorescence) signal upon illuminating a cover glass placed in the focal plane of the primary objective with the blue laser. By progressively blocking off different parts of the optical path, we determined that the main contributors to this signal were the objective and tube lens (Obj and L5 in Fig. 1). Replacing an N-BK7 glass tube lens with a fused silica lens with the same focal length, the signal dropped substantially. The artifact was not affected by the type of anti-reflective coating used.





Figure S2. Lateral resolution and tissue depth penetrance of ADePT photo-stimulation. Corresponding to Fig. 2. (a,b) Lateral confinement of 12 µm nominal diameter light spots projected onto a rhodamine B coated coverslip; (b) x-y FWHM = 17.1  $\mu$ m. (c) 50  $\mu$ m circular spot projected onto a rhodamine B coated coverslip through acute mouse brain slices of different thickness, from 0–150 µm. Note the difference in homogeneity of the light patterns in the presence/absence of the diffuser. (d) Dispersion of the pattern assessed by calculating the x-y FWHM increases with slice thickness, but is comparable in the presence (red) and absence the diffuser (blue). (e) Cartoon schematics depicting (ray diagram view) differences in pattern homogeneity through scattering tissue; in the absence of the diffuser, the example rays (drawn parallel to the optical axis) from three points within the pattern travel straight paths before propagating and scattering through the tissue and being further imaged on the secondary CCD camera; the diffuser diverges the example rays across multiple optical paths (spanning a solid angle cone) which further propagate through the tissue across a wider surface area, reducing the effect of variation in scattering properties along a particular path in the final image. (f) Lateral extent line profiles through the center of the spot for different slice thicknesses without (Left) and with (Right) the 5° holographic diffuser. (g) (Left) Normalized light intensity of 50  $\mu$ m circular spots decreased at comparable rates with and without the diffuser with increasing slice thickness. Measured values were normalized to signal obtained in the absence of the slice (0); (Right) Standard deviation (SD) of fluorescence within the spots as a function of slice thickness (0-250) $\mu$ m). Images taken with the diffuser in the photo-stimulation path have less fluctuation in brightness (fewer speckles).

OMP-Cre x ReaChr-mCitrine x Thy1GCaMP6s



Targeting individual glomeruli by photo-stimulation of OSN terminals and Figure S3. monitoring glomerular responses of sister mitral and tufted cell dendrites. Corresponding to Fig. 4. (a) Confocal image of olfactory bulb sagittal slices showing expression of ReaChR-mCitrine in the OSNs (glomerular layer) and GcaMP6s in the mitral and tufted cells in OMP-Cre x ReaChR-Citrine x Thy1GCaMP6s.DKim4.3 mice. (b) (i) Example baseline fluorescence in a glomerular field of view (FOV) from an OMP-Cre x ReaChR-citrine x Thy1-GCaMP6s mouse; (ii) Response matrix of fluorescence changes (individual repeats in gray, and mean signal as thicker lines) corresponding to five glomerular regions of interest versus the five (matching) glomerular light masks (i) projected on the OB surface. Red traces mark average responses of the target glomerulus; (iii) Fluorescence changes (two dimensional glomerular light activation maps, 2DLAMs) evoked by photo-stimulating the five targeted glomeruli (i) across a range of four light intensities (2.5, 5.0 and 10.0 and 15.0 mW/mm<sup>2</sup>; 2 x 150 ms stimulus duration/frame); (iv) Statistically significant photo-stimulation responses shown as rectified 2DLAMs. Non-significant responses were thresholded to blue (Methods). All five glomeruli targeted for photo-stimulation were responsive. (c) Average lateral extent of glomerular fluorescence response while varying the light intensity (2.5 to 15 mW/mm<sup>2</sup>). (Left) Surface area of significantly responsive pixels; expected surface area, assuming average glomerulus diameter of 75 µm and no activation of cells surrounding the glomerulus, is ~4,400  $\mu$ m<sup>2</sup>. (Right) Responsive pixels area normalized using the ratio between the photo-stimulation mask and target glomerulus areas (Methods).



Figure S4. Identifying sister mitral and tufted cells via axially decoupled glomerular stimulation and two photon fluorescence imaging of cell body responses. Corresponding to Fig. 5. (a) anaesthetized, (**b**-g) awake. (a) (i) (Left) Specificity of enhanced responding mitral and tufted cells as a function of targeted glomerulus (N=5 mice; 16 responsive and specific glomeruli / 24 photostimulated target glomeruli; 357 mitral & tufted cells). Note that most cells were non-responsive, or responded specifically to the photo-stimulation of the target glomerulus (10.3 % of cells had off-target responses to more than one glomerulus; Methods); (Right) Same for suppressed responses. (ii) (Left) Number of observed sister cells (enhanced responses) per glomerulus photostimulation across fields of view (Left; Avg.  $\pm$  SD, 4.2  $\pm$  2.2; N = 6 FOVs; 5 mice); (Right) Number of cells whose baseline fluorescence was suppressed upon photo-stimulation of the same glomeruli (*Right*; Avg.  $\pm$  SD, 1.6  $\pm$  1.8; N = 6 FOVs; 5 mice). (b) Resting glomerular fluorescence (ReaChRcitrine and GCaMP6s), awake; (c) Fluorescence changes (raw, Top, and significant pixels glomerular 2DLAMs, Bottom) evoked by photo-stimulation of three target glomeruli (7.5 mW/mm<sup>2</sup>; 100 ms light on/trial); (d) Response matrix of fluorescence change traces (individual repeats in gray, and mean signal as thicker lines) corresponding to three glomerular regions of interest (#2,#3,#4) versus the 3 (matching) glomerular light masks projected on the OB surface. Red traces mark responses of the targeted glomerulus in a given trial; stimulation of glomeruli #1 and #5 resulted in non-specific responses. (e) Resting fluorescence (GCaMP6s) of mitral cell bodies at the same x-y coordinates, 180  $\mu$ m below the glomerular field of view shown in (b); (f) Significant cell body responses evoked by photo-stimulation of three target glomeruli (g,ii); (iii) Average fluorescence time traces corresponding to nine example mitral cells in each heat map, whose positions are indicated by numbers in (ii). Three example pairs of daughter sister cells (1, 35 with respect to parent glomerulus #2; 14, 33 with respect to parent glomerulus #3; and 25 and 49 with respect to parent glomerulus #4) are shown. In addition, glomerular photo-stimulation triggered spatially distributed suppressed responses (e.g. cells 13, 46).





Figure S5. Optogenetic control of DAT+ (superficial short axon) cells within individual glomeruli. Corresponding to Fig. 6. (a) Confocal images of an olfactory bulb slice of GCaMP6f (Left), ChR2-mCherry (Center) and merged signals (Right) expression in DAT+ neurons. (b) Schematic of in vivo experiment (Top) and baseline 2p fluorescence of three adjacent glomeruli in the field of view (Bottom); GABAergic/dopaminergic superficial short axon cells (DAT+) coexpressing GCaMP6f and ChR2-mCherry; circles and numbers indicate the location of projected light masks. (c) (i) Fluorescence (two-dimensional light activity maps) evoked by photostimulating DAT+ cells in glomeruli (1-3) and one (no apparent opsin expression) control site (4) marked in (b) using light masks equivalent to the dotted blue circles. Color scale in heatmap indicates  $\Delta F/F$ , comparing fluorescence after light stimulation to baseline. (ii) Fluorescence traces (average, thick line, and four individual repeats, thin lines) corresponding to a region of interest (ROI) in each heatmap, whose position is indicated by matching color squares in (i). As in the case of OSN terminals photo-stimulation experiments, we flashed single 100 ms subglomerular light masks (20 µm diameter) between the imaging frames and recorded calcium transients in the DAT+ cells via multiphoton imaging; for analysis, the field of view was divided into a 20 by 20 grid (15 x 15  $\mu$ m grid units), and the response was defined as the average change in fluorescence in each unit of the grid during the two frames (10 Hz) immediately following the light stimulus. (d) For each ROI, the evoked  $\Delta F/F$  response is plotted against the distance of that ROI to the center of each stimulation pattern across repeats. For all stimuli except the control (4), the relationship was modeled as an exponential decay. (e) Width of the exponential fits in (d)against spots of same size and varying light intensity. Dots correspond to individual ROIs. We tested a range of photo-stimulation light intensities ranging from 0.5 to 10.0 mW/mm<sup>2</sup>. Within the range of light intensity used, the characteristic width constant varied from 25 µm for the lowest to 62 µm for the highest.



-0.5 0.0 0.5 Correlation coefficient

1.0

0.00 -1.0 0.0 0.5 -0.5

Correlation coefficient

1.0

-1.0 -0.5 0.0 0.5 1.0

Odor response correlation

Figure S6. Optogenetic control of DAT+ (superficial short axon) cells to determine their impact on mitral cell responses. Corresponding to Fig. 6a-e. (a) (i) Cartoon schematics: in vivo photostimulation of opsin expressing DAT+ cells while monitoring mitral cell activity (GCaMP6f) by axial decoupling of the photo-stimulation and imaging planes. (ii) Confocal image of olfactory bulb sagittal slices showing expression of ChrimsonR-tdtomato in the glomerular DAT+ cells (superficial short axon cells) and GCaMP6f in the mitral and tufted cells in DAT-Cre x Thy1GCaMP6f (5.11) mice. (iii) td-tomato intensity across glomeruli as a function of glomerular size (area). (b) (Left) Correlation distribution of mitral cell ensemble responses across different pairs of DAT+ cell glomerular photo-stimulation masks (G1-G6) per field of view; Avg. ± SD, data:  $0.42 \pm 0.23$ ; shuffled:  $0.29 \pm 0.19$ ; self:  $0.81 \pm 0.10$ . (Right) Same across pairs of odors in the panel (1-11); data:  $0.25 \pm 0.35$ ; shuffled:  $0.06 \pm 0.19$ ; self:  $0.87 \pm 0.10$ ; p < 0.001, Wilcoxon signed-rank test for all pairs of comparisons. Shuffled mitral cell index control distributions are shown in red; 'self' control quantifying variability in response across repeats of the same photostimulation mask is shown in blue. (c) Pairwise correlation of mitral cell responses across stimuli in an example field of view (correlation between all pairs of cells, corresponding to Fig. 6e). (Left) Distribution of pairwise correlations of mitral cells responses across DAT+ cell glomerular photostimulation masks (G1-G6); data: 0.52  $\pm$  0.30; shuffled: 0.37  $\pm$  0.30. (Right) Distribution of pairwise correlation of same cells (N=49 mitral cells) in response to different odors in the panel (1-11). data:  $0.17 \pm 0.33$ ; shuffled:  $0.02 \pm 0.31$ . Shuffled mitral cell index control distributions are shown in red. (d) Pairwise correlation of mitral cell responses to different DAT+ cell glomerular photo-stimulation masks (G1-G6) vs. pairwise correlation of same mitral cell responses to odors in the panel (1-11). Each dot corresponds to a pair of mitral cells. Responses across light stimuli appeared more similar than across odors at the level of mitral cell pairs. However, no apparent trend emerged when analyzing the response correlation of a given pair of mitral cells responded to light stimulation as a function of their response similarity across the odors in the panel.



d ADE 2-HEX 2-HEX + Light ADE+ Light PEN PEN + Light G1 G2 G3 G4 G1G2G3G4 dF/F Tufted cells 61 61

f



Pairwise correlations Same odor across different glomerular light masks



Pairwise correlations Same glomerular light mask across different odor responses



Figure S7. Specificity of modulation of mitral and tufted odor responses by glomerular photoactivation of DAT+ interneurons. Corresponding to Fig. 6f-h. (a) (i) Schematic of the experiment: in vivo photo-stimulation of ChrimsonR-tdtomato expressing DAT+ cells, while monitoring its impact on mitral and tufted cell odor responses (GCaMP6f) using ADePT; (ii) baseline 2p fluorescence of an example field of stimulation in glomerular layer, and of two optical z-planes sampling tufted (iii) and mitral cells (iv) across different depths (100 µm vs. 275 µm from surface); circles and numbers indicate the location of projected light masks (ii) and example tufted and mitral cell bodies (iii,iv). (b) (i) Fluorescence change  $(\Delta F/F)$  elicited by odor presentation in one example tufted cell, and its subsequent modulation by projecting of a glomerular light mask to optogenetically boost DAT+ cell activity (500 ms, 10 mW/mm<sup>2</sup>) 2s from odor onset. (ii, iii) Modulation of example tufted (ii) and mitral cell (iii) odor responses by photo-stimulating DAT+ interneurons across different glomeruli (G1-G4); (c) Odor response modulation ( $\Delta F/F$ ; two dimensional light suppression/activation maps) of tufted (i) and mitral cell (ii) odor responses (ADE, PEN, 2-HEX) evoked by glomerular photo-stimulation of DAT+ cells across 4 individual glomeruli (G1-G4); each cell is color coded (blue-to-read) to indicate its response amplitude to light/odor stimulation; non-significant responses were thresholded to white (Methods). (d) Tufted cell peak responses across the field of view (a); shown are odor responses (ADE, PEN, 2-HEX) and their modulation by 4 DAT+ cell glomerular light masks. (e) Distribution of pairwise correlations of tufted cell response modulation: different glomerular DAT+ cells photostimulation masks across same odor (0.44  $\pm$  0.33, 23 pairs, 2 FOVs, 64 TC cells); (f) Distribution of pairwise correlation of response modulation for different odors across the same DAT+ cells glomerular photo-stimulation mask, aggregated across 4 masks (G1-G4; TC: 0.41  $\pm$  0.32, Avg.  $\pm$ SD; 14 pairs, 2 FOVs, 64 TCs; MC: 0.13±0.34; 14 pairs 2 FOVs, 39 MC cells). Avg. ± SD for shuffled cell index control distributions are shown in red.