Mapping brain networks in awake mice using combined optical neural control and fMRI

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¹McGovern Institute for Brain Research and Department of Brain and Cognitive Sciences, Massachusetts Institute of Technology (MIT), Cambridge; ²Biomedical Engineering Department, Tufts University, Medford; ³Howard Hughes Medical Institute and Center for Brain Science, Harvard University, Cambridge; ⁴The MIT Media Laboratory and MIT Department of Biological Engineering, MIT, Cambridge; and ⁵Center for Biodynamics and Departments of Mathematics and Statistics, Boston University, Boston, Massachusetts

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Desai M, Kahn I, Knoblich U, Bernstein J, Atallah H, Yang A, Kopell N, Buckner RL, Graybiel AM, Moore CI, Boyden ES. Mapping brain networks in awake mice using combined optical neural control and fMRI. J Neurophysiol 105: 1393-1405, 2011. First published December 15, 2010; doi:10.1152/jn.00828.2010. Behaviors and brain disorders involve neural circuits that are widely distributed in the brain. The ability to map the functional connectivity of distributed circuits, and to assess how this connectivity evolves over time, will be facilitated by methods for characterizing the network impact of activating a specific subcircuit, cell type, or projection pathway. We describe here an approach using high-resolution blood oxygenation level-dependent (BOLD) functional MRI (fMRI) of the awake mouse brain-to measure the distributed BOLD response evoked by optical activation of a local, defined cell class expressing the light-gated ion channel channelrhodopsin-2 (ChR2). The utility of this opto-fMRI approach was explored by identifying known cortical and subcortical targets of pyramidal cells of the primary somatosensory cortex (SI) and by analyzing how the set of regions recruited by optogenetically driven SI activity differs between the awake and anesthetized states. Results showed positive BOLD responses in a distributed network that included secondary somatosensory cortex (SII), primary motor cortex (MI), caudoputamen (CP), and contralateral SI (c-SI). Measures in awake compared with anesthetized mice (0.7% isoflurane) showed significantly increased BOLD response in the local region (SI) and indirectly stimulated regions (SII, MI, CP, and c-SI), as well as increased BOLD signal temporal correlations between pairs of regions. These collective results suggest opto-fMRI can provide a controlled means for characterizing the distributed network downstream of a defined cell class in the awake brain. Opto-fMRI may find use in examining causal links between defined circuit elements in diverse behaviors and pathologies.

Functional MRI (fMRI) based on the blood oxygenation leveldependent (BOLD) signal (Kwong et al. 1992; Ogawa et al. 1990, 1992) is widely used to indirectly measure neural activity in distributed brain networks in humans and nonhuman primates. A recent study has shown that optogenetic strategies, using activation of channelrhodopsin-2 (ChR2)-expressing neurons (Boyden et al. 2005), can be used with high-field MRI imaging to evoke BOLD signals in the anesthetized rodent (Lee et al. 2010). We present data obtained using optogenetic activation of a specific cell class in the awake animal, mapping and characterizing the distributed network responses that result. These opto-fMRI studies collectively open up a wide array of opportunities for exploring the relation between BOLD responses and neural activity and to use rodent models to assess the impact of causal manipulations on distributed brain networks.

The recent paper of Lee et al. (2010) illustrates the potential of optical neural control and fMRI in the anesthetized state, showing a BOLD response to light-driven epochs of stimulation in MI in anesthetized rats transfected with ChR2 in Ca²⁺/calmodulin-dependent protein kinase II α (CaMKII α)-expressing neurons. Like prior analyses of sensory-evoked responses in the human (Boynton et al. 1996), the measured BOLD response began a few seconds after stimulation and decreased ~6 s after stimulation offset. Lee et al. (2010) also showed downstream effects of optical stimulation by showing a robust BOLD response in the thalamus during local MI stimulation.

These results raise the possibility that opto-fMRI could serve as a tool to explore properties of distributed brain networks for example, enabling researchers to focus their electrophysiological or anatomical experiments on opto-fMRI-identified sets of brain regions or to examine how large-scale brain network properties are affected by molecular genetic manipulations. By providing a bridge between the increasingly widespread use of optical methods to probe causal neural circuit functions in animals and fMRI, which in humans is typically performed in the awake state, opto-fMRI in the awake mouse may play an important role in the translation of neural circuit insights derived from animal experimentation toward basic and clinical neuroscience in the human.

In this paper, we describe an initial exploration of an awake mouse model of optogenetic stimulation performed during fMRI, thus enabling measurement of large-scale circuit dynamics in a behaviorally relevant state. Using awake mice expressing ChR2 in pyramidal cells in the primary somatosensory (SI) barrel cortex as a model system, we show the detection, in individual mice, of cortical and striatal structures known to be connected to SI (Aronoff et al. 2010; Carvell and Simons 1986, 1987; Chakrabarti and Alloway 2006; Diamond et al. 2008; Ferezou et al. 2007; Megevand et al. 2008; White and DeAmicis 1977). We also show the use of opto-fMRI to assess how the administration of the common anesthetic iso-

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flurane modulates the set of recruited structures downstream of a cell type, as well as the connectivity between them.

METHODS

Animals and surgery

All procedures were conducted in accordance with National Institutes of Health guidelines and with the approval of the MIT Committee on Animal Care. Eleven wild-type mice (C57BL/6, purchased from Charles River) and 11 transgenic ChR2 mice [line 18, stock 007612, strain B6.Cg-Tg (Thy1-COP4/EYFP)18Gfng/J from Jackson Labs, Bar Harbor, ME; bred in-house with wild-type mice] were used. A summary of animals used for the different experiments throughout this study is presented in Supplemental Table S1.¹

For viral injection, eight wild-type mice were first anesthetized with isoflurane ($\sim 1-2\%$ mixed with oxygen), craniotomized (~ 0.5 mm wide), and injected with lentivirus encoding for ChR2-GFP under the CaMKII promoter [FCK-ChR2-GFP, 1 μ l, as used in Han et al. (2009), over a 30-min period, into the left SI barrel field]. Injections were performed using an injection pump (Quintessential Stereotaxic Injector, Stoelting Co., Wood Dale, IL) driving a 10 μ l Hamilton

¹ The online version of this article contains supplemental data.

syringe connected to a glass micropipette (100 μ m tip) via polyethylene tubing. The system was filled with mineral oil. The coordinates of viral injection relative to bregma were as follows: 1.0 mm posterior, 3 mm lateral, 0.7 mm ventral.

For opto-fMRI, animals were surgically implanted with a headpost (Fig. 1A) atop their skulls using dental cement (CandB-Metabond, Parkell). The headpost (weighing \sim 250 mg) and the headpost holder were custom made from Accura 55 plastic, an ABS-like plastic (Fig. 1A). For virally injected animals, the headpost surgery was done at least 3 wk after the surgery for viral injection. For Thy1-ChR2 mice, the kind of craniotomy performed depended on the experiment: in four Thy1-ChR2 mice used for controls for examining negative signal (Supplemental Fig. S2), a small craniotomy (~0.5 mm) was also drilled at this time through the skull over SI to allow positioning of the optical fiber above the cortical surface; in five other Thy1-ChR2 mice used for near-microscopic resolution opto-fMRI (Fig. 5) and for three control experiments (a blood pressure control, a yellow light control, and a paw stimulation control), instead of a craniotomy over SI, the skull was thinned to one half of its thickness to reduce distortion even further for high-resolution imaging (Fig. 5, A and B). At the end of each surgery, exposed skull (except for that which was thinned) was covered with a thin and uniform layer of dental cement to minimize echo-planar image (EPI) distortion caused by susceptibility mismatch



Fig. 1. Opto-functional MRI (fMRI): mapping of distributed neural circuits causally downstream of a defined cell class in the awake mouse brain. A: left: schematic of the opto-fMRI setup for measuring distributed neural activation patterns in the awake mouse, downstream of optical perturbation of an opsin-expressing cell type. Shown is a mouse bearing an MRI-compatible head post, attached to a head post holder, and surrounded by a body restraint tube, which is in turn attached to a cradle with radio frequency (RF) coil attached (surrounding the head post; not shown for clarity; see Supplemental Fig. S1A for photograph of the RF coil). An optical fiber holder orients an optical fiber (200 µm wide) into the craniotomy, aimed at the cortical surface. The cradle is inserted into a 9.4-T MRI scanner. Top Right: boxcar protocol for delivery of a series of 15 s periods (indicated by blue bars, in this and subsequent figures) of 40 Hz pulse trains of 8 ms blue laser pulses (473 nm, 5 mW unless otherwise indicated), driven by a PC running MATLAB. Bottom right: images [echo planar images (EPIs)] are motion-corrected, slice-timing corrected, detrended, converted into percent signal change [expressed as a percent difference from the blood oxygenation level-dependent (BOLD) signal during "laser off" periods], and averaged across scans. This percent signal change is correlated, voxelwise, to the boxcar pattern of the protocol, shifted forward by 2.5 s to compensate for the BOLD signal delay to determine the voxels that are significantly changing. Significance was judged with respect to a Monte Carlo simulation at a multiple comparisons corrected P value of 0.05 for the boxcar correlation (uncorrected P < 0.005; r > 0.83). Bi: channelrhodopsin-2 (ChR2)-GFP expression, overlaid on a mouse atlas (Paxinos and Franklin 2001) in a representative mouse injected with FCK-ChR2-GFP lentivirus in left hemisphere somatosensory cortex (SI) (bregma AP -1.0 mm, ML 3 mm, DV -0.7 mm). Bii: voxels with significant increases in BOLD signal are indicated for the representative mouse (same as Bi) undergoing opto-fMRI. The image in Bii shows the voxels with significant increases in signal; voxel color indicates the boxcar correlation of the voxel; the data are shown for each of 4 0.5 mm thick EPI slices ($150 \times 150 \times 500 \ \mu m$ voxel resolution), overlaid over corresponding 0.5 mm thick single slice T1 anatomical images (these slices are shown from posterior to anterior, shown from left to right). Scale bar in the figure and subsequent figures is 1 mm. Blue triangle indicates illumination site. The time course of SI activation for the representative mouse in Bii is plotted in Biii (red trace), averaged across all 4 scans of an opto-fMRI session. The black trace in Biii is the time course of SI activation (same set of contiguous voxels as for the red trace), when the laser was not on.

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in fMRI. The reason for this final dental cement step was to reduce distortion: Supplemental Fig. S1*B* shows a pair of images (each maximum-intensity projected across 4 coronal slices) from a mouse, which shows the effect of a uniform layer of Metabond on the EPI distortion artifact. Specifically, this headposted wild-type mouse was imaged [spin-echo EPI (SE-EPI)] without any Metabond (Supplemental Fig. S1*Bi*) and reimaged after applying a thin and uniform layer of Metabond (Supplemental Fig. S1*Bii*), showing in the latter case a reduction in artifact.

Awake and anesthetized opto-fMRI of mouse: experimental setup and protocol

Opto-fMRI data were acquired on a 9.4-T (Bruker BioSpin MRI, Ettlingen, Germany), 20 cm ID, horizontal bore magnet. Custom-built radio frequency (RF) transmit-receive surface coils, specifically designed for mice, were used for imaging (Supplemental Fig. S1A). Functional data were acquired using an SE-EPI in the coronal orientation; we compared this protocol to gradient-echo EPI (GRE-EPI) in Supplemental Fig. S1C, in an acute (0.5% isoflurane anesthetized) experiment on a wild-type mouse, and found GRE-EPI to be more susceptible to distortion artifacts than SE-EPI.

Opto-fMRI experiments were performed on eight virally injected wild-type mice, nine Thy1-ChR2 mice, and two wild-type mice. Of the eight virally injected mice, five underwent awake imaging followed by 0.7% isoflurane anesthetized imaging, and three were imaged at 0.5% isoflurane or 1% isoflurane for laser power and negative BOLD controls (Fig. 2*B*; Supplemental Fig. S2*B*). Of the nine Thy1-ChR2 mice, three were imaged at 0.5% isoflurane for near-microscopic imaging (Fig. 5), four were imaged at 0.5% isoflurane for negative BOLD controls (Supplemental Fig. S2), and two were imaged at 0.7% used for the blood pressure control, a yellow light control, and a paw stimulation control.

For animals that underwent awake imaging, 3 consecutive days of acclimation to head restraint and scanner noise were first performed, followed by fMRI data collection starting on the fourth day. Past experiments in awake head-restrained mice have shown that, after a few days of such acclimation, overt stress is greatly reduced, as observed behaviorally (e.g., little or no struggling, eye secretions indicative of stress in mice, or excessive vocalizations or other signs of stress) (Boyden and Raymond 2003; Boyden et al. 2006). Each session was performed at approximately the same time of day, around 2:00 PM to minimize the influence of circadian rhythm. The headposted animal was inserted into the custom built G-10 fiberglass MRI cradle and restrained for \sim 20 min to acclimate the animal to this head restrained position. The animal was given chocolate sprinkles while restrained on the MRI cradle. After 20 min of head restraint acclimation, the cradle was inserted into the 9.4-T MRI scanner. The cradle was locked to the MRI scanner stage to position the head of the animal



Fig. 2. Positive and negative signals shown by opto-fMRI in the presence and absence of ChR2. *A*: voxels with significant increase in signal, colored according to boxcar correlation, for a wild-type mouse (non-ChR2 bearing) undergoing opto-fMRI under 0.5% isoflurane and a laser power of 10 mW, superimposed on the corresponding single-slice T1-anatomical image. Images were collected at $200 \times 200 \times 500 \ \mu m$ voxel resolution. *B*: the proportion of voxels throughout the imaged brain volume (*y*-axis) that possess uncorrected boxcar correlation *P* values below a given value (*x*-axis) are plotted; *x*- and *y*-axes are plotted on a log scale. Vertical dashed line indicates the threshold value of the multiple comparisons corrected *P* value of 0.05 (that is, uncorrected *P* = 0.005). *C* and *D*: voxels with significant decreases in signal, colored according to boxcar correlation, for a wild-type mouse (non-ChR2 bearing; the same mouse as in *A*) and for an FCK-ChR2-GFP lentivirus-injected mouse undergoing opto-fMRI ($150 \times 150 \times 500 \ \mu m$; 5 mW laser power), respectively (see Supplemental Fig. S2 for further supporting data). *E*: as in *B* and *C*, but for an MRI phantom and a laser power of 15 mW, in this case a 2-cm-diam plastic centrifuge tube (50 ml) filled with agarose and a LEGO brick. Shown here is a single coronal slice collected at a voxel resolution of 200 × 200 × 500 μm . Scale bar in *A* and *C*-*E* is 1 mm. Blue triangle in *A* and *C*-*E* indicates illumination site.

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in the center of the magnet bore. An SE-EPI, similar to the one used for fMRI data collection, was run for 40 min to acclimate the animal to scanner noise. The animal was taken out of the scanner and given two more chocolate sprinkles before finally being taken out of the body restraint tube and the headpost holder.

fMRI data collected during all three acclimation sessions were used to estimate head motion of the animal being scanned. Animal head motion was computed using the 3dvolreg program in AFNI. The 3dvolreg program assumes a rigid body transformation and minimizes a least squares difference between each source sub-brick volume and the base. We find the animal head motion reduced significantly [Supplemental Fig. S1D; F(1,48) = 261.82, P < 0.0001, main effect of acclimation in a 2-way ANOVA of acclimation × orientation of motion; F(5,48) = 59.38, P < 0.0001, main effect of orientation of motion in a 2-way ANOVA of acclimation \times orientation of motion; F(5,48) = 34.122, P < 0.001, interaction between acclimation and orientation of motion] following 3 days of acclimation compared with unacclimated animal (1st day of acclimation). Changes in signal-tonoise (SNR) ratio for EPI scans were also computed for all 3 days of acclimation. Temporal SNR (tSNR) was computed from the scans without optical illumination from 100 voxels in the SI barrel field region (contralateral to the optical illumination). tSNR was determined from the mean voxel value across 200 time points, divided by temporal SD of that voxel's signal. The change in average tSNR after 3 days of acclimation compared with the first day of acclimation is plotted in Supplemental Fig. S3E. There was a trend toward an increase in tSNR following 3 days of acclimation (paired *t*-test, n =5 mice; P = 0.12). During awake opto-fMRI, an RF transmit-receive coil (single copper loop, milled out from a copper plated 1/16-in epoxy material (FR-4, T-Tech, Norcross, GA), shown in Supplemental Fig. S1A) was positioned over the animal's head, surrounding the headpost. The animal was attached by its headpost to the headpost holder, which was in turn attached to the MRI cradle. The animal was positioned in a body restraint tube-a 4.5 cm diam, 2 mm wall thickness, plastic tube with an enclosed end at the tail end of the tube, padded with an absorbent material-that lightly fits around the animal's body and restricts motion during opto-fMRI experiments. The restraint tube was bolted to the MRI cradle. A 200 μ m optical fiber (Ocean Optics, Dunedin, FL) attached to an adjustable optical fiber holder was also, at this time, positioned directly on the target (i.e., into the craniotomy, right above the brain), passing through the RF coil. A nose-cone for isoflurane delivery was positioned around the animals' snout. The MRI cradle was slid into the magnet bore. The cradle was locked to the MRI scanner stage to position the head of the animal in the center of the magnet bore. Animals were anesthesia induced with isoflurane right after the awake functional imaging experiment. The anesthesia level for these animals was maintained, as indicated by end-tidal isoflurane level at 0.7%.

For purely anesthetized imaging, animals were anesthetized with isoflurane (2–3% in oxygen) before positioning them on the MRI cradle. The MRI cradle was slid into the magnet bore. The cradle was locked to the MRI scanner stage to position the head of the animal in the center of the magnet bore. Breathing rate (Small Animal Monitoring 1025, SA Instruments, Stony Brook, NY) and end-tidal expired isoflurane (V9004 Capnograph Series, Surgivet, Waukesha, WI) were continuously monitored during awake and anesthetized imaging experiments. The anesthesia level was set at 0.5 or 1.0%, as indicated by end-tidal isoflurane level.

Functional images were collected at voxel resolution $150 \times 150 \times 500 \ \mu\text{m}$ (5 slices; mouse images in Figs. 1*B*, 2*D*, 3*A*, and 4*A*; Supplemental Figs. S3, *A* and *B*, S4, and S5*A*), $100 \times 100 \times 500 \ \mu\text{m}$ (5 slices; mouse images in Fig. 5, *A* and *B*; Supplemental Fig. S2*A*), or $200 \times 200 \times 500 \ \mu\text{m}$ (10 slices; mouse images in Fig. 2, *A* and *C*; Supplemental Figs. S1, *B* and *C*, and S3, *B* and *C*). For simplicity, a summary of fMRI parameters used for different experiments throughout this study is presented in Supplemental Table S1. For the high-resolution imaging sessions (that is, $<150 \times 150 \times 500 \ \mu\text{m}$ in voxel

size), we took advantage of the speed and strength of our gradient coils to frame the volume imaged with four saturation slices (e.g., Fig. 5A), thus avoiding wrap-around artifacts (Wang et al. 2004; Wilm et al. 2007). Functional data were acquired using a SE-EPI; 2.5 s repetition time (T_R) and 25 ms echo time (T_E). High-resolution T1-weighted anatomical images ($78 \times 78 \times 500 \ \mu$ m) were acquired using a rapid acquisition process with relaxation enhancement (RARE) sequence in the coronal orientation, after physiological data acquisition was complete. Statistical maps of the correlation of BOLD percent signal change to the boxcar pattern of light delivery (aka boxcar correlation maps) were aligned to the high-resolution T1-weighted structural volumes using AFNI's align_epi_anat.py program.

The optical protocol used for opto-fMRI in this paper was a 10 s baseline period of darkness followed by 16 repetitions of 15 s on, 15 s off periods of 40 Hz trains of 8 ms laser pulses (the boxcar pattern shown in Fig. 1*A*, *top right*), applied to SI. Light was delivered with a 473 nm blue laser (Shanghai Dream Lasers, Shanghai, China), placed outside the magnet room and coupled to a 200 μ m optical fiber (~5 m in length), which was passed inside the magnet room through a small duct. A MATLAB program and a USB Data Acquisition Module (Cole-Parmer, Vernon Hills, IL) were used to control the laser to deliver the protocol boxcar pattern (Fig. 1*A*, *top right*).

For mapping experiments in the five awake mice (and the 0.7% isoflurane states) and for the near-microscopic opto-fMRI experiments, the laser power out the fiber was 5-10 mW (fiber tip irradiance, $150-300 \text{ mW/mm}^2$). For control experiments, three virally injected mice, three Thy1-ChR2 mice, and two wild-type mice, we used 10-15 mW laser power; these last two wild-type mice were exposed to 25-30 mW as controls.

Control experiments

Three control experiments were performed on two 0.7% isofluraneanesthetized Thy1-ChR2 mice to examine 1) blood pressure dependence on SI illumination, 2) fMRI response to yellow light, and 3) fMRI response to paw stimulation. For the first of these control mice, blood pressure, heart rate, body temperature, and respiration rate were continuously monitored (sampling rate of 5 s; Advisor, Vital Signs Monitor, Surgivet, Waukesha, WI) throughout the length of an optofMRI experiment, on one mouse, to observe the effect of laser stimulation on the animal's physiology. The femoral artery was cannulated with polyethylene tubing (ID 0.28 mm, OD 0.61 mm) to measure blood pressure and heart rate. The mean arterial blood pressure (mabp), mean heart rate (hr), mean temperature (t), and mean breathing rates (br) were 89.4 \pm 8.7 mmHg, 577.4 \pm 12.8 beats/min, 35.8 ± 1.5 °C, and 98.7 ± 3.7 breaths/min, respectively, during the on periods (mean \pm SE; taken across 64 on periods, in 4 scans of 16 on periods each) and 89.3 \pm 8.5 mmHg, 576.5 \pm 15.7 beats/min, 35.8 \pm 1.0° C, and 98.9 ± 3.4 breaths/min during the off periods. On and off were not significantly different for any of these measures (paired *t*-test, n = 64 on periods, each followed by an off period; $P_{\text{mabp}} >$ 0.90; $P_{\rm hr} > 0.90; P_{\rm t} > 0.55; P_{\rm br} > 0.85$). For the second mouse, the effect of paw stimulation was measured (15 s, 40 Hz, 3 ms pulse duration, 3 mA-amplitude pulse current delivery) on the BOLD signal, finding the percent change in SI to be, at its temporal peak, a 2.2% increase (at 7.5 s after stimulus onset), with a shape that matches the hemodynamic response function (HRF) with P < 0.0005 (least squares linear regression analysis, comparing the canonical HRF with each of the 4 scans within the session; $R^2 = 0.9627$, F = 32.93). In this same mouse, we measured the effect of yellow light (593 nm, 5 mW) on the fMRI signal and found no induced BOLD signal changes (positive or negative).

Control experiments investigating putative negative signals were conducted using an MRI phantom of 3% agarose in saline containing a LEGO brick (Fig. 2*E*).



Rank ordering of the ROIs at left, by reliability of observation across animals (with key ROIs labeled, with their respective atlas names)

Fig. 3. Unbiased algorithmic mapping of neural targets recruited by SI pyramidal cell activation in the awake mouse brain. *A*: population data for 5 mice undergoing awake opto-fMRI. The image in *A* shows the voxels with significant increases in signal in any animal. Voxel color indicates the median of the boxcar correlation, taken across all animals for which that voxel exhibits a statistically significant increase, for the voxel. The data are shown for each of 4 0.5 mm thick EPI slices $(150 \times 150 \times 500 \ \mu m$ voxel resolution), overlaid over corresponding 0.5 mm thick single slice T1 anatomical images (these slices are shown from posterior to anterior, shown from *left* to *right*). Scale bar in the figure is 1 mm. Blue triangle indicates illumination site. For completeness, response maps for all 5 animals injected with FCK-ChR2-GFP lentivirus and undergoing opto-fMRI are shown in Supplemental Fig. S4, and statistical robustness is explored in Supplemental Fig. S5. *Bi*: plot of regions of interest that are consistently activated across mice during SI pyramidal cell activation, generated by an unsupervised k-means clustering of all sets of contiguous significantly activated voxels obtained during the opto-fMRI experiment performed in *A*. Each filled circle indicates the centroid of a k-means derived cluster and is connected to the sets of contiguous significantly activated voxels. The shape of the open symbol, localized to the location of the peak correlation voxel for that set of contiguous significantly activated voxels. The shape of the open symbol indicates which animal the set of contiguous significantly activated voxels is from. K-means derived clusters were considered as regions of interest (ROIs), with key ROIs (annotated based on what they correspond to in the atlas) labeled as SI (SI barrel field), c-SI (contralateral SI), SII (secondary sensory cortex), MI (motor cortex), and CP (caudoputame). *Bii*: percentage of mice that exhibit opto-fMRI activations for each of the 11 ROIs generated in the

Opto-fMRI of mouse: data analysis

The analysis pipeline is outlined in the *bottom right* of Fig. 1A. Boxcar correlation maps for EPI functional data were generated using AFNI (Cox 1996; Nelson et al. 2006) (National Institutes of Health, http://afni.nimh.nih.gov/afni, Bethesda, MD) and MAT-LAB (The Mathworks, Natick, MA). EPI functional data were motion corrected in all three spatial dimensions, but no spatial smoothing or undistortion was performed (to preserve full resolution). The images were slice-time corrected and detrended, as is standard for EPI scanning (Lindquist 2008; Smith et al. 1999). Percent signal change was computed by subtracting, from each voxel's BOLD time series, the temporal average of all off periods for all scans in a session and dividing by the average of the signal across all off periods. We performed four scans per session, and percent signal change was averaged across scans in a session unless otherwise indicated.

0 mm

To determine which voxels had significant increases or decreases in BOLD signal, a voxelwise time series correlation of the percent signal change was performed, as determined above, with the protocol boxcar (Fig. 1*A, top right*), delayed by one repetition time (T_R ; 2.5 s) to compensate for the BOLD response time lag (Cox

1996; Nelson et al. 2006); we label this analysis "boxcar correlation" in the text. Activation in a region was deemed significant if a cluster (here used in the fMRI sense of the word) of at least six contiguous voxels had correlation P values at an uncorrected threshold of P < 0.005. This cluster size of 6 and this uncorrected P value threshold of 0.005 were objectively chosen via Monte Carlo simulations performed in AFNI (family-wise error correction using AlphaSim program), to result in a multiple comparisons corrected type I error of 5%, appropriate for the performance of statistics for individual voxels taken throughout the entire imaging volume (Forman et al. 1995; Xiong et al. 1995). The parameters used for the four Monte Carlo simulations (1 for each scanning resolution used) were as follows: number of voxels, $100 \times 100 \times$ number of slices; voxel size (functional resolution); size of smoothing filter (same as voxel size); number of Monte Carlo iterations. 10.000.

Region of interest identification and cross-registration

To identify regions showing positive BOLD responses, boxcar correlation maps (that is, the correlation of BOLD percent signal change to the boxcar model, as described above) were aligned to the

Innovative Methodology



Fig. 4. Opto-fMRI analysis of brain state modulation of causal network dynamics: application to characterization of anesthesia effects on SI network recruitment. A: overlay of the sets of contiguous significantly activated voxels, color-coded by ROI membership, onto T1 anatomical images, in the awake (*top*) and 0.7% isoflurane anesthetized (*bottom*) state (see Supplemental Fig. S3 for raw datasets). For each set of contiguous significantly activated voxels, a colored point is shown, localized to the location of the peak correlation voxel for that set of contiguous significantly activated voxels. Scale bar, 1 mm. *B*: percent change in the BOLD signal, plotted over time, for each key ROI indicated by a label in Fig. 3*Bi*. The BOLD signal time course was averaged across all pulse trains within a session (consisting of 4 scans of 16 pulse trains each) and across all the voxels within the contiguous significantly-activated voxels associated with each key ROI. Solid lines indicate the average of the percent signal change, taken across animals, for awake (red) and anesthetized (blue) states; shaded areas indicate the SE of percent signal change (n = 5 mice). The dashed curves in *B* represent scaled canonical hemodynamic response functions (HRFs) fitted to the BOLD signal time course for each of the ROIs, in either the awake (red) or anesthetized (blue) state. *C*: peak percent change in the BOLD signal for the experiments plotted in Fig. 4*B*. ***P* < 0.0001, factor of anesthesia level (2-way ANOVA of anesthesia level × region). *D*: normalized cross-correlations of the time series of percent change in the BOLD signal, averaged across animals, and plotted on a continuous color scale. ***P* < 0.0001, factor of anesthesia level (2-way ANOVA of anesthesia level × region). *D*: normalized ross-correlations of the time series of percent change in the BOLD signal, averaged across animals, and plotted on a continuous color scale. ***P* < 0.0001, factor of anesthesia level (2-way ANOVA of anesthesia level × region). f

high-resolution T1-weighted structural volumes using AFNI's align_epi_anat.py program. High-resolution T1-weighted structural images were cross-registered between mice using AFNI's 3dAllineate program; this alignment was used to bring the boxcar correlation maps into a standard coordinate space.

For each imaging session, the coordinate location of the voxel with peak boxcar correlation, in each contiguous set of significantly activated voxels, was identified. Using k-means clustering we clustered (note: the term cluster is now being used in the k-means sense, not in the fMRI preprocessing sense used in the previous section of the METHODS) the coordinate locations of the peak-correlation voxels thus identified. We adapted an algorithm (Duda et al. 2000) to perform the k-means analysis in an unsupervised fashion, following the steps listed below, starting with C = 1:

- performing k-means clustering with C clusters;
- computing for each resultant cluster the within-cluster sumof-squares position variance – that is, the sum, over all peak-correlation voxels in a cluster, of the squared distance l(coordinate of the peak-correlation voxel) – (mean coordinate of all peak-correlation voxels within the cluster)²|;
- comparing the C sum-of-squares variances thus computed, using a *t*-test, to those obtained when k-means was run with C + 1 clusters.

We iterated this process, increasing C by 1 each time until the P value of this t-test was no longer significant (P < 0.05); then, the number of clusters used for k-means was set to C.

Key regions of interest (ROIs) were named overlaying T1 anatomical images onto corresponding atlas plates from the Paxinos and Franklin (2001) atlas, by matching their anteroposterior coordinates and scaling atlas plates along the *x*- and *y*-axes until the borders of the T1 anatomical images and the atlas plate corresponded.

As a test of robustness, the k-means clustering analysis was repeated on correlation maps obtained for two different values of boxcar correlation P value threshold (5 times higher and 5 times lower than the 1 objectively chosen via Monte Carlo simulation to result in a multiple comparisons corrected type I error of 5%; Supplemental Fig. S5).

ROI time series extraction and analysis

Extracted time series from each key ROI were signal averaged across the contiguous significantly activated voxels associated with each key ROI and averaged across pulse trains within a session (consisting of 4 scans, of 16 pulse trains each) to generate an average time series of percent change in the BOLD signal for each key ROI (Fig. 4*B*). The peak percent change in the BOLD signal, across time, was extracted from these thus-averaged traces (Fig. 4*C*).



fig. 5. Opto-fMRI applied to transgenic mice. A: voxels with significant increases in BOLD signal are indicated, color coded by boxcar correlation, for a representative Thy1-ChR2 mouse undergoing high-resolution $(100 \times 100 \times 500 \ \mu\text{m})$ opto-fMRI (here performed under 0.5% isoflurane anesthesia; blue laser pulse power, 10 mW). The data are shown for a single EPI slice, overlaid over a corresponding 0.5 mm thick single slice T1 anatomical image. Scale bar in the figure is 1 mm. Blue triangle indicates illumination site. *B*: voxels in SI with significant increase in BOLD signal, for 3 Thy1-ChR2 mice undergoing high-resolution opto-fMRI; the leftmost of the 3 images is a zoomed-in subpicture of Fig. 5A. C: peak percent change in the BOLD signal (*x*-axis) is plotted as a function of cortical depth (*y*-axis, measured relative to the pia). Solid line indicates the average of the percent signal change taken across the mice shown in *B*; shaded areas indicate the SE of percent signal change (across n = 3 Thy1-ChR2 mice). *Di*: epifluorescence image of a section (50 μ m thick) of the brain of a Thy1-ChR2 is fused with YFP. Multiunit activity (MUA) recorded across neocortical layers was measured using a laminar silicon electrode array (contacts spaced, 100 μ m). A representative episode of MUA recorded from an electrode contact in layer V of cortex during 1 s pulse train (40 Hz, 8 ms pulses) illumination is plotted in *Dii*. MUA firing rate (*E*) and local field potential (LFP) power (*F*) during the illumination period (*x*-axis; LFP power is normalized to the power in the 1 s preceding the illumination period) are plotted as a function of cortical depth (0–1,200 μ m; *y*-axis). Solid line indicates the average of the MUA firing rate and the LFP power; shaded areas indicate SE (n = 2 Thy1-ChR2 mice).

We assessed the extent to which BOLD signal time series could be fit with a human-derived canonical HRF. We produced a goodnessof-fit (R^2) measure and probability that the fit is statistically significant, by applying a least squares linear regression for each key ROI, in each of the awake and anesthetized (0.7% isoflurane level) states, each time taking into account data from each of the five animals to compare against the canonical HRF. In a separate analysis, we sought to test whether the goodness-of-fits themselves differed between the awake and anesthetized states or between regions. For this analysis, the goodness-of-fit (R^2) for the default value of delay of response (6) s, relative to onset) was first computed on each animal's individual time series (averaged across scans within a session) and then through Fisher's z-transform computed the z-score for the goodness-of-fit to the HRF. The resultant set of z-scores was subjected to a two-way ANOVA with factors of brain state and ROI. In a second analysis, we varied the delay of response parameter from 1 to 12 s. The delay of response for which the R^2 value was maximum was computed for each animal independently, and the resultant set of delay of responses that resulted in peak R^2 values was subjected to a two-way ANOVA with factors of brain state and ROI.

Normalized cross-correlations of BOLD percent signal change time series were computed between pairs of key ROIs (Fig. 4D) (Haralick and Shapiro 1992). We computed, for each key ROI, the average of all

the normalized cross correlations between that ROI and all the other key ROIs; this average was termed the "total connectivity" of that region.

BOLD signal responses as a function of cortical depth were computed by drawing a line perpendicular to the cortical surface and choosing sets of three voxels (the 1 closest to the line and the 2 horizontally flanking ones) for each cortical depth. Peak percent BOLD signal change was averaged across each of these sets of three voxels and across animals (n = 3) for each depth below the surface of the cortex (Fig. 5*C*).

We used *t*-test and ANOVA (StatView, SAS Institute, Cary, NC) throughout, e.g., to analyze the dependence of BOLD signal change or significantly activated brain volume on anesthesia and brain region (Fig. 4*C*, etc.).

Local field potential and multiunit activity recordings

Two Thy1-ChR2 mice were anesthetized with isoflurane (0.5%) and headposted for electrophysiology, followed by the opening of a small craniotomy over the barrel cortex. A laminar silicon linear electrode array (Neuronexus Technologies) was connected to a Cheetah32 data acquisition system (Neuralynx). The array consisted of 16 contacts along the probe shank, each with a diameter of 15 μ m and

spaced 100 μ m apart. The probe shank was inserted such that ≥ 12 contacts covered the cortical laminae from immediately below the surface to a depth of ~1,200 μ m continuously (depicted in Fig. 5*D*). Recordings were performed with minimal filtering (band-pass 0.1–9,000 Hz) and separated into local field potential (LFP) and multiunit activity (MUA) signals in software using MATLAB. Specifically, MUA was determined by filtering the original signal between 600 and 6,000 Hz and thresholding at 4 SD above background noise; LFP was determined by filtering the original signal between 7 and 200 Hz. MUA and LFP plots in Fig. 5, *E* and *F*, were averaged across two animals, for 10 trials of 1 s of light stimulation (40 Hz, 8 ms pulses) each time, and plotted as a function of cortical depth. LFP power was normalized to the 1 s baseline before light onset.

Histology

After opto-fMRI experiments were concluded, we verified the extent of ChR2 expression. Animals were transcardially perfused with 100 mM PBS followed by 4% formaldehyde in PBS. Brains were postfixed overnight at 4°C and cryoprotected in 30% sucrose for 24–48 h before sectioning. Free-floating sections (50 μ m) were cut using a cryostat (Leica 3050S, Leica Microsystems, Wetzlar, Germany), mounted on glass slides with Vectashield mounting medium with DAPI (Vector Laboratories, Burlingame, CA), and coverslipped. Spread and labeling efficiency were estimated by examination of 50 μ m coronal sections near the site of injection for the presence of GFP. For the five animals that underwent the core awake opto-fMRI experiments, four were histologically examined; the fifth mouse (CaMKII-ChR2-D) was not examined because of potential health issues unrelated to the opto-fMRI experiment. Examination of the entire brain between V1 and frontal cortex showed no GFP beyond the site of injection. To estimate the volume of the cortex containing ChR2-GFP expressing cells, we measured the area in each slice by measuring image intensity of GFP (field of view 1.9×1.9 mm, $4 \times$ objective, BX51, Olympus, Tokyo, Japan). Each injection of $\sim 1 \ \mu l$ labeled an ellipsoid volume with a diameter of 700–1,000 μ m along the anterior-posterior aspect, 700-1,000 along the medial-lateral aspect, and 800-1,100 mm along the dorsal-ventral aspect. Next, we collected confocal image stacks ($21 \times 21 \times 45 \mu m$; FV10i, Olympus) sampling two regions along the dorsal-ventral axis of each GFPpositive area, on each of three equally spaced slices sampled from each animal. For each of these regions, we counted the number of GFP-positive and DAPI-positive cell bodies. By this measure, the local density of cells expressing ChR2 would be estimated as $314,938 \pm 110,485/\text{mm}^3$ (SE) over the regions identified (n = 4 mice).

RESULTS

Optimization of opto-fMRI methods for awake mice

A diagram of the opto-fMRI hardware for awake mouse imaging is shown in Fig. 1A. This equipment consists of a blue laser coupled to a long optical fiber, a custom-engineered MRI cradle for holding an awake mouse (CAD design available on request), a body restraint tube within the cradle for minimizing body motion, an RF coil (Supplemental Fig. S1A), and an optical fiber and holder (which passes through the open loop of the RF coil) over the desired neural target. To facilitate placement of the animal in the MRI cradle, the animal is surgically prepared in advance with a plastic head post that is bolted into the cradle at the beginning of an experiment. The animal is also prepared with a craniotomy or thinned skull over the area of interest (in this case, the SI barrel cortex, which has been transgenically or virally labeled to express ChR2 in pyramidal neurons). The experimental opto-fMRI protocol (Fig. 1*A*, *right*) used EPI in conjunction with a boxcar protocol of light delivery (15 s periods of 40 Hz delivery of 8 ms laser pulses, separated by 15 s periods of no illumination). Significantly activated voxels in images were identified by correlating the percent signal change of the BOLD response with a boxcar function that corresponded to the periods of light delivery and selecting the voxels whose correlation coefficients were significant at a familywise multiple comparisons-corrected *P* value level of <0.05 (equivalent to an uncorrected *P* value of 0.005) and had five or more significant neighbors (i.e., a cluster threshold of 6). The uncorrected *P* value of 0.005 and the cluster threshold of 6 were objectively determined using the Monte Carlo simulation program of AFNI (Forman et al. 1995; Xiong et al. 1995).

Our method was optimized along a number of experimental axes. First, to enable fMRI in awake mice, it was critical to acclimate mice to head restraint and scanner noise over a multi-day period under conditions similar to actual opto-fMRI experimentation. In addition, the body restraint tube was important for reducing body motion. Although the body was not in the fMRI field of view, motion can cause susceptibility artifacts in brain images by changing the magnetic environment. Several other methodological choices were important for preventing magnetic susceptibility artifacts. First, although GRE-EPI is commonly used because of its excellent signal-tonoise properties, we found that SE-EPI minimized distortion caused by magnetic susceptibility mismatch between tissue and air (Bandettini et al. 1994). This sequence approach (SE-EPI) also provides for higher-resolution imaging (Kim et al. 2000; Lee et al. 1999; Moon et al. 2007) (Supplemental Fig. S1C). Second, adding a thin layer of dental cement homogeneously across the surface of the skull reduced magnetic susceptibility artifacts caused by the mismatch between intact tissue and air (Supplemental Fig. S1B). Third, we adapted a strategy from diffusion tensor imaging (Wang et al. 2004; Wilm et al. 2007), namely, the use of saturation slices, to eliminate signals from nearby tissue outside the field of view.

BOLD signal changes downstream of SI pyramidal cell activation

Opto-fMRI was performed on awake mice that had undergone viral infusion of the excitatory neuron-targeting lentivirus FCK-ChR2-GFP (Han et al. 2009) into the SI barrel cortex. These mice expressed ChR2-GFP in pyramidal neurons within a spherical volume between 700 and 1,000 μ m in diameter, containing neurons distributed in layers 2/3 and layer 5 (Fig. 1Bi). On illumination of SI pyramidal cells, we observed robust, distributed BOLD responses in the brain ipsilateral and contralateral to the fiber illumination site, time locked to the illumination period. Figure 1Bii shows the results from a representative awake mouse. The time series of evoked BOLD signals in SI is plotted for this mouse in Fig. 1Biii (red trace). When the laser was not on, there was no visible change in the SI BOLD signal (Fig. 1Biii, black trace). We did not detect positive BOLD responses in control mice not expressing ChR2 undergoing opto-fMRI (Fig. 2A; n = 2 mice; representative mouse shown), even with laser powers increased to levels threefold to sixfold higher than used in the experiments here described (Fig. 2B). Few voxels were significantly activated in

control even at an uncorrected P value of 0.15 (Fig. 2*B*). This control experiment mitigates the possibility that positive BOLD responses were produced by non-ChR2-related effects of cortical laser illumination, such as any potential temperature changes induced by illumination.

A putative negative BOLD response was observed in the brain during opto-fMRI in the SI barrel cortex immediately under the optical fiber tip (Fig. 2D). This negative BOLD response was similar in ChR2-negative mice (Fig. 2C). To probe the mechanism of negative signal, we carried out a specific set of experiments to compare the negative signal under the fiber tip in three ChR2-virally transduced mice, two non-ChR2-expressing mice, and four Thy1-ChR2 mice and found negative responses in all three sets of mice of comparable magnitude [Fig. 2*F*; P > 0.80, F(2,6) = 0.225 for peak reduction in signal, factor of mouse type, 1-way ANOVA]. The fact that lack of ChR2 did not alter the signal reduction, despite its elimination of the signal increases observed (e.g., Fig. 2, A and B) led us to hypothesize that the signal reduction observed under illumination was not related to ChR2 but instead reflected a direct and local effect of light on the sample that was observable using MRI. In support of this hypothesis, we observed small but significant signal reductions even in plain agarose (Fig. 2E), on illumination. This local effect was at least in part unrelated to optical activation of ChR2-expressing neurons and may therefore reflect temperature-related modulation of the fMRI signal (Yablonskiy et al. 2000) or other nonspecific local effects of illumination (see Supplemental Results and Supplemental Fig. S2 for more analysis of the negative signal, i.e., analysis for Thy1-ChR2 mice). We confine the rest of our analysis to observation of positive BOLD signals, because negative signal was not consistently observed except for this nonspecific local response.

To characterize the consistency of the opto-fMRI map across different mice, we performed opto-fMRI on five awake mice, each injected with 1 μ l FCK-ChR2-GFP lentivirus in SI of the left hemisphere. To facilitate this comparison, the resultant functional scans were aligned for each mouse to its respective T1 anatomical scan, and these datasets were cross-registered across all mice. All mice exhibited BOLD activations in the illuminated SI and also displayed activations in nearby cortical regions, subcortical regions, and the contralateral cortex. Voxels that were significant in any of the five mice are color-coded according to the median (taken across mice) boxcar correlation value in Fig. 3*A* (raw datasets for all 5 mice are shown in Supplemental Fig. S4).

Neural targets recruited by SI pyramidal cell activation can be automatically identified

To identify, in an unbiased manner, the brain regions downstream of SI activation (Figs. 1*Bii* and 3*A*), we developed an unsupervised algorithm that first clusters sets of contiguous significantly activated voxels into ROIs and then localizes the centroid of each ROI to a reference mouse atlas. The clustering algorithm is based on an automated variant of the k-means clustering algorithm (see METHODS for details), which adjusts the number of clusters according to objective statistical criteria, and showed 11 ROIs for the five opto-fMRI experiments conducted (Fig. 3*Bi*). Of these 11 ROIs, we designated 5 as key ROIs, which were robustly activated in all five mice (Fig. 3Bii). The centroid-atlas alignment showed these key ROIs as ipsilateral SI, contralateral SI (c-SI), ipsilateral primary motor cortex (MI), ipsilateral secondary somatosensory cortex (SII), and the caudoputamen (CP), which are regions known to be targets of SI pyramidal neurons (Aronoff et al. 2010; Carvell and Simons 1986, 1987; Chakrabarti and Alloway 2006; Diamond et al. 2008; Ferezou et al. 2007; Megevand et al. 2008; White and DeAmicis 1977). The algorithm also reported ROIs that were less consistent across individual mice (Fig. 3Bii). These ROIs might represent additional activated circuits, for which the reliability of observation was not as high as the selected five key ROIs. For example, in a subset of the five opto-fMRI experiments, clusters emerged in regions that were atlas-aligned to thalamus and contralateral SII (Fig. 3Bi; these activated regions were also visible in the raw datasets shown in 11 ROIs; Supplemental Fig. S4). This variability could result from natural variations in the strength of connectivity of circuits from one brain to another, from subtle variations in the location of virally labeled neurons or of the placement of the optical fiber across mice, or from the fact that, in our algorithm, the choice of a boxcar correlation statistical threshold could admit significant voxels for some mice but reject others, leading to a perceived heterogeneity across mice.

To assess statistical robustness, we repeated the opto-fMRI algorithm on boxcar correlation maps obtained by varying the uncorrected boxcar correlation P value to values 5 times higher and 5 times lower than that used in Fig. 3. We found that varying the uncorrected boxcar correlation P value to 0.001 or 0.025 resulted in few qualitative changes in the appearance of the correlation maps (representative mouse shown in Supplemental Fig. S5Ai; median boxcar correlation map shown in Supplemental Fig. S5Aii). When we ran the unsupervised neural target identification algorithm on the opto-fMRI data, few differences emerged in the cluster map, with key ROIs preserved (Supplemental Figs. S5, B and C) and with most clusters remaining centered at the same location but expanding or contracting as the P value was relaxed or made stricter. Our algorithm thus exhibits robustness to variations in the specific *P* value chosen for the initial boxcar correlation threshold, indicating its utility in comparing neural maps across animals and across experiments.

Opto-fMRI evokes a greater BOLD response in awake than anesthetized animals

We anticipate a major use of opto-fMRI will be in awake animals, given the potential for comparing animal opto-fMRI data to human fMRI data (which is predominantly collected in the awake state) and for understanding behavior in head-posted rodents, an approach that has recently been optimized for a variety of imaging modalities (Andermann et al. 2010). To show the use of opto-fMRI in analyzing how light-activated neural networks can be modulated, we compared, for the five mice analyzed in Fig. 3, how administration of 0.7% isoflurane varied the network impact of SI illumination. Isoflurane was chosen because it is commonly used in small animal fMRI to prevent motion artifacts and immobilization stress (King et al. 2005; Lahti et al. 1998; Sicard et al. 2003), and it was also the anesthetic used in an earlier opto-fMRI study (Lee et al. 2010). The role that isoflurane anesthesia (or any anesthesia) has on local versus global network dynamics,

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and on the coupling of these neural signals to blood flow, is controversial (Corfield et al. 2001; Matta et al. 1999; Tsurugizawa et al. 2010).

To visualize the effects of anesthesia on the set of regions causally recruited by SI, we plot in Fig. 4A the clusters or ROIs identified by the algorithm described above, for the awake (top) and anesthetized (bottom) states. Raw datasets leading up to these ROI plots, akin to those shown for the awake state in Fig. 3, are provided in Supplemental Fig. S3. In the anesthetized state, only 6 ROIs were activated compared with the 11 ROIs in the awake state. Of the five key ROIs identified in the awake state (SI, c-SI, SII, MI, and CP), four were represented in the anesthetized state, but the striatal response was less prominent. This finding is consistent with previous studies using electrophysiology (Detsch et al. 2002; West 1998) reporting that anesthesia suppresses somatosensory-evoked information transfer to subcortical regions. In the isoflurane anesthetized condition, there was also a decrease in the BOLD response for the voxels in all five key ROIs identified in the awake state [Fig. 4, B and C; F(1,40) = 172.63, P < 0.0001, main effect of anesthesia level in a 2-way ANOVA of anesthesia level × region; F(4,40) = 7.830, P < 0.0001, main effect of region in a 2-way ANOVA of anesthesia level \times region; no interaction, F(4,40) = 0.558, P = 0.6943]. These contrasts show the potential utility of opto-fMRI for probing dynamics in network activation.

To address the question of how the time course of the BOLD response is altered by anesthesia administration, we examined the goodness-of-fit of BOLD responses in different regions under different levels of anesthesia to the canonical HRF. The BOLD response in SI was well fit with the canonical HRF in the awake state [Fig. 4B, dotted line superimposed on red trace in left plot; $R^2 = 0.94$, F(1,4) = 21.37, P = 0.0002, n = 5mice; least square linear regression, allowing only the amplitude scaling of the HRF to vary, as well as the amplitude offset, but not allowing for scaling or shifting of the HRF along the time axis]. Indeed, all of the awake responses were well fit with the canonical HRF [R^2 ranging from 0.78 to 0.94, F(1,4)] ranging from 7.28 to 21.37, P values ranging from 0.0002 to 0.009; dotted lines superimposed on red trace in each plot of Fig. 4B]. The anesthetized responses were less well fit by the HRF [R^2 ranging from 0.15 to 0.90, F(1,4) ranging from 1.57 to 17.57, P value ranging from 0.0008 to 0.27].

To apply a direct statistical comparison of the goodness-offit to the HRF for the awake and anesthetized BOLD signal changes, we computed an R^2 for every opto-fMRI session BOLD signal versus the HRF. In our first analysis, we computed this R^2 using the default value of delay in HRF (6 s relative to onset) and then through Fisher's z-transform computed the z-score for the goodness-of-fit to the HRF. We ran an ANOVA to test how these z-scores varied across anesthesia level and region. We found that anesthesia significantly lowered the z-score for the goodness-of-fit to the HRF of the BOLD signal relative to the awake state [F(1,40) = 167.62], P < 0.0001, main effect of anesthesia level in a 2-way ANOVA of anesthesia level \times region], and regions also differed in their match to the HRF [F(4,40) = 34.42, P < 0.0001, main effectof region in a 2-way ANOVA of anesthesia level \times region]. There was a significant interaction between anesthesia level and region [F(4,40) = 7.28, P = 0.0002]. Specifically, local SI activity was least affected in its BOLD signal response shape

by anesthesia (z-score for the goodness-of-fit to the HRF decreased from 2.07 to 1.59), whereas the striatum was most affected (z-score for the goodness-of-fit to the HRF decreased from 1.48 to 0.35). In a second analysis, we varied the delay parameter for the HRF from 1 to 12 s and recomputed the R^2 for each fit to determine the delay that would result in the highest R^2 value. We found that anesthesia significantly increased the delay in HRF that resulted in the peak R^2 , relative to the best-fit delay seen for the awake state [F(1,40) = 98.33], P < 0.0001, main effect of anesthesia level in a 2-way ANOVA of anesthesia level \times region], and regions also differed in their delay in HRF that would result in peak R^2 [F(4,40) = 11.04, P < 0.0001, main effect of region in a 2-wayANOVA of anesthesia level \times region]. There was a significant interaction between anesthesia level and region [F(4,40) =18.96, P < 0.0001] in this delay computation.

Anesthesia affects BOLD functional connectivity in the network recruited by pyramidal cell activation

We applied opto-fMRI to probe the functional connectivity of the brain network downstream of SI as a function of brain state, again using isoflurane anesthesia as a model. We performed normalized cross-correlation analyses of the BOLD signal changes in different ROIs during opto-fMRI. This technique provides a measure of temporal correlation across regions while nullifying the effect of absolute changes in signal amplitude (as reported above), allowing focus on how anesthesia level modulates the coordination between regions. We found that the anesthetized state had generally lower normalized cross-correlations between pairs of regions than did pairs of regions in the awake state [Fig. 4D; F(1,80) = 71.67, P < 1000.0001, main effect of anesthesia level in a 2-way ANOVA of anesthesia level \times region pair]. This measure of correlation also varied between different pairs of regions [F(9,80) = 4.76]. P < 0.0001, main effect of region in a 2-way ANOVA of anesthesia level \times region pair, and no interaction between the two factors, F(9,80) = 1.22, P = 0.29].

The striatum showed lower correlations with other regions than did other pairings. The 17 post hoc tests of differences between the awake and anesthetized state that showed P values using Fisher's protected least significant difference (PLSD) test significant with respect to a 0.05 significance level, all involved the striatum. When we computed, for each region, the average normalized cross-correlation between the striatum and all the other regions, we found that this "total connectivity" was lowered by anesthesia [F(1,40) = 86.50, P < 0.0001, maineffect of anesthesia level in a 2-way ANOVA of anesthesia level \times region pair]. Different regions also exhibited different total levels of connectivity with the rest of the network [F(4,40) = 3.39, P < 0.02, main effect of region in 2-way]ANOVA of anesthesia level \times region pair; no interaction, F(4,40) = 0.76, P > 0.55], with the striatum exhibiting less total connectivity than the others (P < 0.02, Fisher's PLSD, comparison with all other regions).

Opto-fMRI applied to transgenic mice expressing ChR2 in defined cortical layers

Opto-fMRI can be applied to other models of potentially significant interest, and we explored these signals in transgenic mice. These preparations provide disease models and have

been important in a variety of basic science studies. We applied opto-fMRI to Thy1-ChR2 transgenic mice that express ChR2-YFP in the neocortex in layer 5 pyramidal neurons (Arenkiel et al. 2007) (Fig. 5*Di*). In this mouse strain, multiunit neural activity appeared predominantly in layer 5 (Fig. 5, *Dii* and *E*); LFP power was broadly distributed in these mice, perhaps because of the extensive ChR2-positive dendrites (Fig. 5*F*). When Thy1-ChR2 mice received light isoflurane (0.5%) and underwent opto-fMRI, positive responses could be observed in SI sensory cortex (Fig. 5*B*), including prominent activity in layer 5. The ChR2-independent negative signal in Thy1-ChR2 mice (Supplemental Fig. S2) also was observed in the most superficial layers of the cortex (Fig. 5*C*). These findings extend the application of opto-fMRI to transgenic mice, suggesting that application to other lines will be successful.

DISCUSSION

Optogenetic strategies using ChR2 expression have previously been shown to evoke a BOLD response in the anesthetized rat (Lee et al. 2010). Here we extend this work is three ways. First, robust positive BOLD responses were shown in the awake mouse. The ability to perform opto-fMRI in the awake mouse opens up translational links between causal circuit studies in animals and behaviorally and clinically important human studies. Second, using photoactivation of cortical pyramidal cells of the mouse SI barrel cortex, which has wellcharacterized targets throughout the brain (Aronoff et al. 2010; Carvell and Simons 1986, 1987; Chakrabarti and Alloway 2006; Diamond et al. 2008; Ferezou et al. 2007; Megevand et al. 2008; White and DeAmicis 1977), we showed that distributed regions of the network could be evoked by light stimulation and measured using an unbiased, automated analysis. The network response included regions in the striatum: the MI, c-SI, and SII. Third, functional connectivity measures showed the BOLD response was correlated between regions within the network consistent with observations from evoked and intrinsic BOLD fluctuations in the human (Fox and Raichle 2007; Friston 1994; Hampson et al. 2006; Pezawas et al. 2005; Ramnani et al. 2004; Stephan et al. 2008; Van Dijk et al. 2010). The absolute magnitude of BOLD response in each region of the network and the functional correlation strengths between regions were attenuated by anesthesia. These collective results suggest that opto-fMRI provides an approach to map circuits in the awake mouse and thus complements fMRI approaches used in vivo in humans.

This study also provides a detailed account, intended to facilitate the implementation of this method by future researchers. A number of technical challenges were overcome to achieve the combination of optical neural activation and fMRI in the awake mouse. Specialized hardware for awake mouse imaging was implemented (Fig. 1) that enables positioning of optical fibers and immobilization without incurring significant motion artifacts, using a combination of head posts and restraint tubes. Surgical and pulse sequence strategies were devised for reducing magnetic susceptibility artifacts while preserving high resolution. For the majority of the experiments in this study, we chose to use 150 mW/mm² irradiance, well within the range used for repeated stimulation (Ayling et al. 2009; Cardin et al. 2009, 2010; Kravitz et al. 2010; Zhang et al. 2010) and at the

lower limit of what is needed to evoke EMG responses when targeting MI (Ayling et al. 2009); however, it will generally be useful to select the least dosage of light necessary for achieving a given scientific goal. We also performed a number of critical control experiments, for example, showing that negative signal changes under the optical fiber are nonspecific effects of opto-fMRI, unrelated to the activity of ChR2-expressing neurons. Future studies will be required to differentiate these nonspecific signals from negative BOLD responses.

Opto-fMRI provides a strategy for characterizing the network downstream of a target in a fashion that is amenable to repeated assessment, potentially over extended time periods suitable to longitudinal study of learning and plasticity, and disease progression. Thus this work complements recent studies by Lee et al. (2010) in the rat discussed above and Logothetis et al. (2010) in the monkey that used electrical stimulation to measure the BOLD response in distributed regions across the visual system. In the future, the addition of new technologies such as optical neural silencing reagents (Chow et al. 2010; Han and Boyden 2007; Zhang et al. 2007; Zhao et al. 2008) or multisite illuminators, the targeting of different neuron types, and further improvements of fMRI, will augment the performance of opto-fMRI. Awake mouse optofMRI also opens up the possibility of implementing a psychophysical paradigm and assessing brain-wide effects of changing the psychometric threshold curves through optogenetically manipulating a specific subcircuit (Cardin et al. 2009).

There are also clear limitations to opto-fMRI. Weak, modulatory functional connections or transient connections may not be detectable because of the slow temporal response of the BOLD signal and the nature of neural-to-hemodynamic coupling. This limitation of poor temporal resolution is in part offset by the utility of this approach for exploring widespread distributed regions and the specificity of the target cell types and location that can be achieved. Another limitation of optofMRI method is the need for a head-fixed animal. Head-fixed animals have been shown to have sparse firing rate in the sensory cortex compared with freely moving animals (de Kock and Sakmann 2009; Sakata and Harris 2009).The higher firing rate is thought be important for information processing in various ways, in a behaviorally relevant state (Vijayan et al. 2010).

Other approaches for mapping neural circuits in a causal fashion using fMRI exist. Electrical microstimulation has proven to be an important tool in combination with fMRI (Canals et al. 2008; Ekstrom et al. 2009; Logothetis 2003; Logothetis et al. 2010; Sultan et al. 2007; Tolias et al. 2005) for driving local circuits in the context of whole-brain mapping, but electrical activation can recruit a diversity of cell types within a microcircuit, as well as fibers of passage, smooth muscle, and other brain circuit elements (Ranck 1975). However, the passive spread of current can confound the analysis of functional connections (local and distal), as determined by fMRI, requiring indirect methods to characterize the current spread such as behavioral controls (Tehovnik et al. 2006). Future work that directly compares stimulation methods will be valuable for understanding tradeoffs between methods and resolve the uses of these tools in different spheres of neuroscience.

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DISCLOSURES

No conflicts of interest, financial or otherwise, are declared by the authors.

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Mouse coil



В

А



С









Thy1-ChR2 mouse, high-resolution scan (decreases in MRI signal)



% change in the MRI signal, peak over time, for Thy1-ChR2 mice, averaged across all voxels with significant % signal change in negative direction



В

A





В

А

Median opto-fMRI activation, taken across all mice (0.7% isoflurane anesthesia)



С

i

Regions of interest (ROIs) downstream of SI pyramidal cell activation, revealed via unsupervised clustering of opto-fMRI activations across animals



△ CaMKII-ChR2-A
□ CaMKII-ChR2-B
○ CaMKII-ChR2-C
☆ CaMKII-ChR2-C
◇ CaMKII-ChR2-E
8 sono constant
8 sono constant
8 sono constant

ii



Percentage of mice which exhibit opto-fMRI activations for each ROI



Rank ordering of the ROIs at left, by reliability of observation across animals (with key ROIs labeled, with their respective atlas names)





С

В





Percentage of mice which exhibit opto-fMRI activations for each ROI, for uncorrected boxcar correlation p-value to values 5x higher and 5x lower than that used in Fig. 1 and Fig. 3



Rank ordering of the ROIs at left, by reliability of observation across animals (with the five key ROIs labeled, with their respective atlas names)

ii

Supplementary Table 1.

Experiment type	Mice			fMRI parameters		
	Animal type	Number of animals	Animal IDs	Voxel resolution	Number of slices	Referenced in
Opto-fMRI						
Awake followed by 0.7% isoflurane	Virally injected wild types	5	CamKII-ChR2-A thru E	150 μm × 150 μm × 500 μm	5	Fig. 1B, Fig. 2B & D, Fig. 3, Fig. 4, Supp. Fig. 1D & E, Supp. Fig. 3, Supp. Fig. 4 and Supp. Fig. 5.
0.5% isoflurane	Virally injected wild types	3	CamKII-ChR2-F thru H	200 μm × 200 μm × 500 μm	10	Fig. 2B & F
	Thy1-ChR2 mice	3	Thy1-ChR2-A thru C	100 μm × 100 μm × 500 μm	5	Fig. 5A , B & C
	Un-injected wild types	2	WT-noChR2-A and B	200 μm × 200 μm × 500 μm	10	Fig. 2A, B, C & F
0.5% and 1.0% isoflurane	Thy1-ChR2 mice	4	Thy1-ChR2-D thru G	200 μm × 200 μm × 500 μm	10	Fig. 2F and Supp. Fig. 2A & B
Electrophysiology experiment						
LFP and MUA recordings for optogenetic stimulation	Thy1-ChR2 mice	2	Thy1-ChR2-H and I	NA	NA	Fig. 5D & E
Control fMRI						
Head post and pulse sequence EPI distortion	Un-injected wild types	1	WT-noChR2-C	200 μm × 200 μm × 500 μm	10	Supp. Fig. 1B & C
Yellow light optical and electrical paw stimulation	Thy1-ChR2 mice	1	Thy1-ChR2-K	200 μm × 200 μm × 500 μm	10	methods section
physiological measurements	Thy1-ChR2 mice	1	Thy1-ChR2-J	200 μm × 200 μm × 500 μm	10	methods section

Supplementary Figure 1. RF coil design, cement usage, and EPI distortion elimination. A. Custom-built RF transmit-receive surface coils used for opto-fMRI. B, Single-slice images obtained via SE-EPI, in an anesthetized animal, without (i) and with (ii) a thin and uniform layer of Metabond coating the skull. Images were collected at 200 µm x 200 µm x 500 µm voxel resolution, and maximum-intensity projected across 4 coronal slices (2 mm). Scale bar in **B** and **C** is 1 mm. **C**, Single-scan images of the same headposted mouse shown in **B**, with cement already applied, obtained via GRE-EPI (i) and SE-EPI (ii). Each image was acquired at 200 μ m x $200 \,\mu\text{m} \ge 500 \,\mu\text{m}$. White open triangle in **B** and **C** indicates position of the optical fiber tip; the laser was left unilluminated for these experiments. **D**, Average (n = 5 mice) measurements of motion (estimated from EPI scans) while the animals are in the fMRI scanner. The measurements are from the first day of acclimation (grey bars) and the day opto-fMRI data was collected, following 3 days of acclimation sessions (black bars). Motion (μm) is measured in three orientations (Superior-Inferior (SI), Left-Right (LR) and Anterior-Posterior (AP)) and three rotational directions (roll, pitch, yaw). **, p < 0.0001, factor of acclimation (two-way ANOVA) of acclimation \times orientation of motion). E, Average (n=5 mice) SNR measurements from the first day of acclimation (grey bar) and the day opto-fMRI data were collected (following 3 days of acclimation sessions). Temporal SNR (tSNR) was computed from the scans without optical illumination and from 100 voxels in the S1BF region (contralateral to the optical illumination). tSNR was determined from the mean pixel value across the 200 time points divided by its temporal standard deviation.

Supplementary Figure 2. Negative BOLD signal in Thy1-ChR2 mice. A, Voxels with significant decreases in BOLD signal, colored according to boxcar correlation, for a Thy1-ChR2 transgenic mouse undergoing opto-fMRI under 0.5% isoflurane at a near-microscopic resolution (100 μm x 100 μm x 500 μm; 10 mW laser power). Scale bar is 1 mm. Blue triangle indicates

illumination site. **B**, Negative BOLD signal, (at the peak, after timecourses of BOLD signal change were averaged across all significantly-decreased voxels within SI), for four Thy1-ChR2 mice for which isoflurane was varied (0.5% and 1.0%).

Supplementary Figure 3. Unbiased algorithmic mapping of neural targets recruited by SI pyramidal cell activation in the anesthetised mouse brain. A. Voxels with significant increases in BOLD signal are indicated for the representative mouse (same as in Fig. 1B) that is anesthetized (0.7% isoflurane concentration) and undergoing opto-fMRI. Voxel color in A indicates the boxcar correlation of the voxel. **B**. Population data for 5 anesthestised (0.7%)isoflurane concentration) mice undergoing opto-fMRI (same mice as in Fig. 3). The image in B shows the voxels with significant increases in signal in any animal. Voxel color indicates the median of the boxcar correlation, taken across all animals for which that voxel exhibits a statistically significant increase, for the voxel. The data are shown for each of four 0.5 mm thick EPI slices (150 μ m × 150 μ m × 500 μ m voxel resolution), overlaid over corresponding 0.5 mm thick single slice T1 anatomical images (these slices are shown from posterior to anterior, displayed from left to right). Scale bar in the figure is 1 mm. Blue triangle indicates illumination site. C, i, Plot of regions of interest that are consistently activated across mice during SI pyramidal cell activation, generated by an unsupervised k-means clustering of all sets of contiguous significantly activated voxels obtained during the opto-fMRI experiment performed in **Supp. Fig. 3B**. Each filled circle indicates the centroid of a k-means derived cluster, and is connected to the sets of contiguous significantly activated voxels that make up this cluster; each of these sets of contiguous significantly activated voxels is marked by an open symbol, localized to the location of the peak correlation voxel for that set of contiguous significantly activated voxels. The shape of the open symbol indicates which animal the set of contiguous significantly activated voxels is from. K-means derived clusters were considered as regions of interest (ROIs), with key ROIs (annotated based upon what they correspond to in the atlas) labeled as SI (SI barrel field), c-SI (contralateral SI), SII (secondary sensory cortex), MI (motor cortex), and CP (caudate/putamen) respectively. **Cii**, Percentage of mice which exhibit opto-fMRI activations for each of the 11 ROIs generated in the k-means clustering of **Supp. Fig. 3Ci**, rank ordered, from left to right, by reliability of observation across animals.

Supplementary Figure 4. Raw opto-fMRI data for all 5 individual mice expressing ChR2 in SI and imaged awake. Four boxcar correlation+T1 anatomy images, plotted as in Fig. 1Bii, are shown, for each mouse (shown from posterior to anterior, displayed from left to right), for each of the five FCK-ChR2-GFP lentivirus injected mice. Functional images are collected at a resolution of 150 μ m x 150 μ m x 500 μ m (5 mW laser power). Scale bar in the figure is 1 mm. Blue triangle indicates illumination site.

Supplementary Figure 5. Robustness of the algorithmic component of the opto-fMRI mapping process. A, i, voxels with significant increases in BOLD signal are indicated for the representative mouse displayed in Fig. 1Bii, but varying the boxcar correlation *p*-value to different levels. Shown in Aii is the population data (n = 5 mice) similar to Fig. 3A, but varying the boxcar correlation *p*-value to different levels. Scale bar in the figure is 1 mm. Blue triangle indicates illumination site. B, Regions of interest (ROIs) downstream of SI pyramidal cell activation, revealed via unsupervised clustering of opto-fMRI activations across animals, for uncorrected boxcar correlation *p*-value to values 5x higher (0.025) and 5x lower (0.001) than that used in Fig. 3Bi. C, Percentage of mice which exhibit opto-fMRI activations for each of the ROIs generated in the k-means clustering for the two uncorrected *p*-values in B, rank ordered, from left to right, by reliability of observation across animals.

Supplementary Table 1. Summary of animal usage and fMRI parameters for different experiments throughout this manuscript. The table presents a summary of all the animals that are used throughout this study and the kind of experiments they are used for. The summary also presents the fMRI parameters (resolution and number of slices) employed for a specific experiment and the figure it represents in this manuscript.

Supplementary Results

Negative BOLD responses

BOLD signal reductions were observed in opto-fMRI in Thy1-ChR2 mice (**Supp. Fig. 2A**), localized to the region of the brain just under the fiber tip. The negative BOLD response in Thy1-ChR2 mice (n = 4) was similar to the negative BOLD response observed in 3 ChR2-virally transduced mice and 2 non ChR2-expressing mice (**Fig. 2C, Fig. 2D** and **Fig. 2F**). We varied the isoflurane concentration from 0.5% to 1.0% while illuminating SI in ChR2-expressing Thy1 mice. We saw no effect of isoflurane level on the peak percent change in the BOLD signal for voxels that significantly decreased in BOLD signal (**Supp. Fig. 2B**; p > 0.90, paired t-test; n = 4 mice) in Thy1-ChR2 mice where isoflurane is changed from 0.5% to 1.0%. Thus, multiple independent lines of experiments support the idea that BOLD signal reductions observed during light delivery to the brains of ChR2-expressing mice are not due to ChR2-mediated neural activation, but instead reflect a direct effect of light on the sample.

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