Intravital Imaging of Neocortical Heterotopia Reveals Aberrant Axonal Pathfinding and Myelination around Ectopic Neurons

Alice M. Li1,2,5, Robert A. Hill2,3 and Jaime Grutzendler1,2,4

1Interdepartmental Neuroscience Program, Yale School of Medicine, New Haven, CT 06510, USA, 2Department of Neurology, Yale School of Medicine, New Haven, CT 06510, USA, 3Department of Biological Sciences, Dartmouth College, Hanover, NH 03755, USA, 4Department of Neuroscience, Yale School of Medicine, New Haven, CT 06510, USA and 5Current address: Department of Pathology, Brigham and Women’s Hospital and Harvard Medical School, Boston, MA 02115, USA

Address correspondence to Robert Hill, Department of Biological Sciences, Dartmouth College, 78 College Street, Hanover, NH 03755, USA. Email: robert.hill@dartmouth.edu; Jaime Grutzendler, Department of Neurology, Yale School of Medicine, 300 George St., New Haven, CT 06511, USA. Email: jaime.grutzendler@yale.edu

Abstract

Neocortical heterotopia consist of ectopic neuronal clusters that are frequently found in individuals with cognitive disability and epilepsy. However, their pathogenesis remains poorly understood due in part to a lack of tractable animal models. We have developed an inducible model of focal cortical heterotopia that enables their precise spatiotemporal control and high-resolution optical imaging in live mice. Here, we report that heterotopia are associated with striking patterns of circumferentially projecting axons and increased myelination around neuronal clusters. Despite their aberrant axonal patterns, in vivo calcium imaging revealed that heterotopic neurons remain functionally connected to other brain regions, highlighting their potential to influence global neural networks. These aberrant patterns only form when heterotopia are induced during a critical embryonic temporal window, but not in early postnatal development. Our model provides a new way to investigate heterotopia formation in vivo and reveals features suggesting the existence of developmentally modulated, neuron-derived axon guidance and myelination factors.

Key words: axon pathfinding, cortical development, cortical malformation, heterotopia, myelination

Introduction

Up to one-third of routine postmortem examinations reveal the presence of neocortical heterotopia (Schulze and Braak 1978; Kaufmann and Galaburda 1989; Meencke and Veith 1992; Kasper et al. 1999), a heterogeneous group of focal cortical migration defects characterized by abnormally positioned clusters of neurons (Barkovich et al. 1992; Sisodiya 2004). Heterotopia have been linked to many neurological conditions including epilepsy, intellectual disability, and dyslexia (Galaburda and Kemper 1979; Hardiman et al. 1988; Palmiini et al. 1991; Farrell et al. 1992; Meencke and Veith 1992; Nordborg et al. 1999; Kakita et al. 2002). However, comprehensive exploration of their pathogenesis and pathophysiology has been limited by a lack of tools for their investigation in the live animal. Several genetic, traumatic, chemotoxic, and other heterotopia models have been described (Riggs et al. 1956; Dvorák et al. 1978; Sherman et al. 1987; Rosen et al. 1992; Ferrer et al. 1993; Amano et al. 1996; Brunstrom et al. 1997; Lee et al. 1997; Bai et al. 2003). However, these previous models have not been utilized for cellular intravital optical imaging analyses largely due to the lack of control over the position and timing of heterotopia induction and/or limited means for targeted cell labeling.
Here, we developed a methodology enabling the specific induction and visualization of focal heterotopia in the live mammalian neocortex. Our model uses in utero electroporation (Saito and Nakatsuji 2001; Tabata and Nakajima 2001; LoTurco et al. 2009) combined with in vivo optical imaging, to generate and track layer I heterotopia, a poorly understood developmental malformation characterized by misplaced neurons localized near the pial surface (Jacob 1940; Morel and Wildi 1952; Schulze and Braak 1978; Galaburda et al. 1985; Kaufmann and Galaburda 1989; Yamamoto et al. 1997; Sarkovich 1998). When visualized using label-free myelin imaging (Hill and Grutzendler 2014; Schain et al. 2014; Hill et al. 2018), we identify striking patterns of aberrantly projecting axons and focally increased myelination surrounding the heterotopic neurons. These distinct patterns emerge only when the heterotopia are induced during a critical embryonic period, suggesting the presence of locally derived, developmentally modulated signals that initiate the abnormal structural organization and myelination of the heterotopic axons. Identification and characterization of the neuronal and glial subtypes within embryonically induced heterotopia revealed many consistent features that are used to define spontaneously occurring heterotopia in both humans (Jacob 1940; Morel and Wildi 1952; Schulze and Braak 1978; Galaburda et al. 1985; Kaufmann and Galaburda 1989) and rodent models (Sherman et al. 1985, 1987, 1990; Ramos et al. 2008, 2014; Lipoff et al. 2011; Toia et al. 2017). Finally, using genetically encoded calcium biosensors, we demonstrate that heterotopic neurons display similar calcium transient frequencies as neighboring layer II/III neurons and respond to sensorimotor stimulation in behaving mice, opening new possibilities for the exploration of their influence on cortical function.

Materials and Methods

Animals

All experimental approaches and procedures were conducted in accordance with Yale University Institutional Animal Care and Use Committee regulations. Timed pregnant outbred CD1 mice were purchased from Charles River Laboratories, Inc, with the first 24 h of postnatal life designated as P0. We included both male and female mice, aged P30—P60, for this study. For some birth dating experiments as described in the text, pregnant dams were given a single intraperitoneal (i.p.) injection of 5-ethynyl-2'-deoxyuridine (EdU; 30 μg g⁻¹ body weight) at E11.5, prior to IUE surgery at E15. Litters were kept in individual ventilated cages until weaning age (P21), after which mice were housed in single-sex groups with 2–5 animals per unit. Cages were maintained in temperature-controlled facilities with 12-h light/12-h dark cycles.

In Utero Intracranial Injection, Electroporation, and Viral Infection

Embryonic cortical injections were performed as part of the IUE procedure during needle insertion into the lateral ventricle (Saito and Nakatsuji 2001; Saito 2006). Embryos were injected once unilaterally between embryonic day 14 and 17 (E14–E17) for all electroporations, as specified in the text. Intracranial injections were not performed at gestational ages below E14 or over E17 due to the technical challenges associated with maintaining embryo viability. To keep consistent experimental design parameters, however, all quantifications for in vivo and fixed-tissue analyses used only E15-injected embryos.

All in utero injections were targeted toward prospective somatosensory cortices. Injection solutions included the following plasmid and viral components for neuronal labeling: pCAG-tdTomato (based on Addgene plasmid 11150) and rAAV8-hSyn-eGFP (UNC Vector Core, Lot AV5075D; titer 3.9e12 GC mL⁻¹). Injection solutions contained either pCAG-tdTomato (1.5 μg μL⁻¹ final concentration) only, or both pCAG-tdTomato (1.5 μg μL⁻¹ final concentration) and AAV8-hSyn-eGFP (final titer 3.9e11 GC mL⁻¹) in saline solution. All injection solutions contained Fast Green FCF dye (0.2 mg mL⁻¹; TCI) to facilitate their visual tracking during the injection procedure. Procedures for in utero electroporation have been previously described (Saito and Nakatsuji 2001; Saito 2006).

Briefly, timed pregnant CD1 dams were anesthetized with a saline solution containing both ketamine (100–120 mg kg⁻¹) and xylazine (10–12 mg kg⁻¹), delivered via intraperitoneal injection. Following induction of deep surgical anesthesia, midline incisions (1½ inch) were made into the abdominal skin and muscle wall to access the underlying uterus. Pulled glass capillary needles (10 μL Drummond Scientific Glass Capillaries, Cat# 3-000-210-G; pulled to ∼50 μm diameter at tip and ∼125 μm diameter at 1 mm above tip) were then used to puncture the uterine wall and deliver (Picospritzer II, General Valve) 0.5 μL of injection solution into the lateral ventricle of individual embryos. Each embryo was injected only once and subsequently electroporated using BTX tweezer electrodes aimed at the somatosensory cortex of the injected hemisphere. All electroporations were conducted using four 50-ms, 50-V electrical pulses, delivered at 1-s intervals (BTX Harvard Apparatus 8300 pulse generator).

Postnatal Intracranial Viral Injection

Intracranial injections were performed on P0 pups from timed pregnant CD1 dams that were naïve to the in utero electroporation procedure. Injection solutions included one of the following viruses to mark injected regions: rAAV8-hSyn-eGFP (UNC Vector Core, Lot AV5075D; titer 3.9e12 GC mL⁻¹) and rAAV2-CaMKIIα-mCherry (UNC Vector Core, Lot AV4377d, 3.8e12 titer GC mL⁻¹). P0 intracranial injections were performed essentially as previously described (Kim et al. 2013). Briefly, neonates were cryoanesthetized (Phifer and Terry 1986) within 24 h following birth. After confirming loss of voluntary movement, neonates were placed in the prone position on a polymer cooling block. A pulled glass capillary needle (10 μL Drummond Scientific Glass Capillaries, Cat# 3-000-210-G; pulled to ∼75 μm diameter at tip and ∼325 μm diameter at 4 mm above tip), advanced at a 90° angle from the dorsal scalp through the lateral ventricle to a 4 mm depth, was then used to deliver into the brain parenchyma 1 μL of virus solution (diluted 1:10 in phosphate buffered saline [PBS] and 0.2 mg mL⁻¹ Fast Green FCF, from stock). Each neonate received only one unilateral intracranial injection targeted over somatosensory cortices. Injected neonates were then rewarmed on a heating pad and placed back with their biological mother. All P0-injected mice were sacrificed for analysis at P30, with quantifications and analyses performed in fixed tissue to circumvent the obstructive meningeal scarring and parenchymal adhesions associated with P0 injections on in vivo imaging.
Postnatal Subarachnoid Viral Injection

Injection solutions were prepared using AAV9-Syn-GCaMP6f-WPRE-SV40 (Penn Vector Core, Lot CS1001; titer 7.648e13 GC mL⁻¹) virus diluted 1:100 in PBS and Fast Green FCF (0.2 mg mL⁻¹). P21–P30 mice that had previously been electro-porated in utero were anesthetized by intraperitoneal injections of ketamine (100 mg kg⁻¹) and xylazine (10 mg kg⁻¹). The scalp was shaved, cleaned, and then incised to expose the underlying bone. A high-speed drill was next employed to introduce a small burr hole (~0.75 mm diameter) over the transfected hemisphere, taking care to avoid cortical regions suspected to have been directly punctured as part of the in utero electroporation procedure. The underlying dura was gently detached, and a 12-μL volume of injection solution (prepared as described above) was infused into the subarachnoid space to achieve viral transfection of both layer I heterotopic and layer II/III neurons via topical cortical application. The scalp incision was then closed with sutures. Cranial windows for in vivo imaging were prepared over injected cortical hemispheres 3–4 weeks following the subarachnoid space AAV infusion.

Cranial Window Surgery and In Vivo Imaging

All in vivo imaging was performed using cranial windows (Hill et al. 2018). Briefly, mice were anesthetized using ketamine (100 mg kg⁻¹) and xylazine (10 mg kg⁻¹), delivered via i.p. injection. The dorsal skull surgical field was shaved and cleaned, and an approximately 4-mm diameter circular region of skull and dura mater was excised from the injected hemisphere. A #0 transparent glass coverslip was then gently implanted on top of uncovered pial surface to serve as the cranial window. Glue and dental cement were applied to secure the window to the surrounding skull bone.

To detect neuronal cell bodies in vivo as described in the text, the fluorescent membrane-permeable probe NeuO (Neu-RoFluor, Stemcell Technologies Cat# 01801, diluted 1:25 in PBS) was applied to exposed cortex for 20 min followed by a 1–2-min PBS rinse, before the placement of the #0 glass coverslip during the cranial window surgery. In some cases, 100 μL Evans blue (TCI; 1 mg mL⁻¹) was injected intravenously after the cranial window surgery to label the cortical vasculature.

Except for GCaMP6f calcium imaging experiments, all in vivo imaging studies used mice that were anesthetized via i.p. ketamine and xylazine injection. In vivo imaging of anesthetized mice was performed at P30 immediately after cranial window surgery. All intravital GCaMP6f calcium imaging was carried out in P50–P60 awake head-fixed mice starting 4 h after arousal from cranial window surgical anesthesia.

Confocal in vivo images of previously injected cortical areas with or without the needle tract sites were acquired using a 20× water immersion objective (Leica, 1.0 NA) on a Leica SP5 upright laser scanning microscope. Spectral confocal reflectance (SCoRe) imaging to detect myelinated axon segments was performed as previously described (Schain et al. 2014; Hill et al. 2018) by capturing the simultaneously reflected light signals from 488, 561, and 633 nm multi-wavelength laser excitation outputs. Single-photon laser outputs were tuned to the following excitation wavelengths for fluorescence imaging: 488 nm for GFP and NeuO; 561 nm for tdTomato and mCherry; 633 nm for Evans blue. Sequential imaging was employed to minimize overlap between SCoRe reflection and individual fluorescence emission signals for all in vivo confocal imaging experiments.

Needle tracts from injections performed in utero were identified by the abrupt changes in orientations of labeled dendritic and axonal processes around breaks in the cortical surface in vivo. Since needle tracts identified in this manner always displayed ectopic neural clusters of similar expanse in layer I, the outer boundaries of “heterotopia” in vivo were defined as the layer I needle tract borders for all quantifications. Ipsilateral, noninjected cortical areas were defined as layer I regions at least 150 μm away from a discernable injection site border. Heterotopia and ipsilateral control regions were imaged using identical laser output and image acquisition configurations that were determined for each experimental data set. Confocal z-stacks were acquired at 1024 × 1024 pixel resolution starting from the pia through depths of up to 120 μm below the cortical pial surface. In some cases, time-lapse imaging of in vivo-induced heterotopia was also performed at 512 × 512 pixel resolution.

For some GCaMP6f calcium imaging experiments that involved visualization of deeper cortical regions in layers II/III as indicated in the text, time-lapse fluorescence images were acquired using a 20× water immersion objective (Zeiss, 1.0 NA) on a Prairie Technologies two-photon microscope fitted with a mode-locked, tunable Spectra Physics Mai-Tai laser. In these experiments, the two-photon laser was tuned to 920 nm for excitation of both GCaMP6f and tdTomato fluorophores. All two-photon time-lapse imaging was performed at 512 × 512 pixel resolution.

In Vivo Image Processing and Quantification

Except for GCaMP6f calcium imaging experiments, which were conducted using P50–P60 mice, all intravital imaging studies were performed using P30 mice, with quantifications carried out on those that had previously been injected at E15. All in vivo images were processed and quantified using ImageJ/FIJI.

For SCoRe density quantifications, we analyzed single z-sections located 10 μm deep to the cortical pial surface from both heterotopic and ipsilateral control regions. SCoRe density values were assessed using a custom-built, automated thresholding and binarization macro in ImageJ/FIJI, with Robust Automatic Threshold parameters set to noise = 25, lambda = 3, min = 31. Randomly selected, equally sized regions of interest (ROIs) were used to determine SCoRe density values within the centers and edges of the heterotopia versus control regions. A “heterotopion center” was defined as the circular region with a radius extending from the needle tract center to 1/3 of the radius of the needle tract. A “heterotopion edge” was defined as the needle tract concentric circular region just outside the “heterotopion center,” extending from the “heterotopion center” outer edge to the outermost border of the needle tract. Average SCoRe densities were determined for the heterotopion center, heterotopion edge, and control area for each mouse (n = 8 mice). Statistical analyses were carried out using Wilcoxon matched pairs, signed-rank nonparametric tests.

For NeuO dye-labeled cell body density quantifications, data were analyzed from the first 75 μm of cortex deep to the pial surface of both heterotopic and control regions in layer I. Equal-sized volumes (50 μm³) were randomly selected from the superficial cortical z-stacks captured from both heterotopic and control regions. The number of NeuO⁺ cell bodies was manually counted in each volume. Average NeuO⁺ cell body densities were then determined for the heterotopic and control region of each mouse (n = 6 mice). Statistical analyses were performed using Wilcoxon matched pairs, signed-rank nonparametric tests.
For quantification of neuronal calcium dynamics in heterotopia versus adjacent noninjected cortical areas, we used the two-photon microscope to capture 150 × 150 μm field of view (FOV) time-lapse images from both layer I heterotopia and noninjected ipsilateral layer II/III regions. Time-lapse imaging (512 × 512 pixel resolution; 2 Hz) was performed in awake, head-fixed mice during a 2-h time window that started 4 h after their arousal from cranial window surgical anesthesia. For heterotopic regions, FOVs were randomly selected from within the first 75 μm below the cortical surface in layer I. For layer II/III regions, FOVs were randomly selected between 120 and 175 μm below the cortical surface. Six to eight separate FOVs (exactly half from layer I heterotopia and half from ipsilateral layer II/III regions) were imaged in each mouse, with each FOV recorded for 120 s per trial for three trials. The order in which FOVs were acquired from heterotopia and layer II/III regions was alternated between mice. Time series analyses were performed using ImageJ/FIJI. Prior to quantifications, TurboReg plugin in ImageJ/FIJI was used to align all time series images in the XY plane.

To quantify GCaMP6f fluorescence changes in individual cells, ROIs were manually selected to encapsulate neuronal cell bodies. GCaMP6f+ cell bodies that exhibited tdTomato labeling from the IUE procedure were excluded from all analyses. Approximately 10–25 GCaMP6f+ cells were analyzed in each FOV. Baseline GCaMP6f fluorescence intensity values (F) for each cell were defined in each trial as the average of the lowest 20% of the recorded values per trial, with fluctuations from this baseline denoted as ΔF/F. A spike event was defined in each trial as a ΔF/F > 0.5. Spike event frequencies were averaged across three trials for each cell in both layer I heterotopia and layer II/III regions for n = 7 mice. Statistical analyses were performed using Wilcoxon matched pairs, signed-rank nonparametric tests.

For quantifications of the global synchronization index reflecting the relative coordination of GCaMP6f activity, analyses were performed using Fluorescence Single Neuron and Network Analysis Package (FluoroSNNAP) (Patel et al. 2015). Briefly, this semiautomated software implements a correlation matrix-based algorithm (Li et al. 2007, 2010; Patel et al. 2015) to compute the normalized global synchronization indices (ranging in value from 0 to 1) for time series calcium imaging data of neural populations. The highest value, 1, signifies entirely synchronized firing throughout an identified cluster of neurons, whereas 0 indicates the total absence of synchrony. Synchronization cluster analyses were performed using template-based calcium event detection parameters set to detection threshold = 0.85, minimum size of synchronization clusters = 2, and surrogate resampling = 20. A cluster of neurons was defined as the GCaMP6f+ cell population within one FOV. Global synchronization indices were averaged across three trials for each FOV in both layer I heterotopia and layer II/III regions for n = 7 mice. Statistical analyses were performed using Wilcoxon matched pairs, signed-rank nonparametric tests.

**Tissue Processing and Immunohistochemistry**

At P30, mice were deeply anesthetized using ketamine and xylazine and then perfused transcardially with 4% paraformaldehyde in PBS. Harvested brains were postfixed overnight using the same solution at 4 °C and then vibratome sectioned (coronal; 75 μm thickness) for fixed-tissue analysis.

To identify brain regions that had been injected as part of the IUE procedure, a fluorescence microscope was next used to examine the sections for visibly disrupted layer I regions showing transfected neurons and/or neuronal processes, similar to what has been previously described (Rosen et al. 2007; Wang et al. 2020). The disrupted regions often revealed cellular clusters containing eGFP+ neurons that protruded past the pial surface, which were never observed in intact cortical regions or in noninjected mice. Sections containing visibly disrupted layer I cytoarchitecture indicative of injection-associated trauma were then selected for further processing. Immunohistochemistry, performed as described below, was used to confirm the presence of layer I heterotopia at all identified IUE injection sites. Coronal sections from P0-injected mice were screened in a similar manner for injected areas, which were identified by virally transfected axonal fibers and neuronal cell bodies concentrated around visible needle tracts (data not shown).

Prior to immunostaining, all free-floating sections were heated to 95 °C for 30 min in 50-mM sodium citrate buffer (0.05% Tween-20, pH 6.0) for antigen retrieval and elimination of endogenous eGFP expression. After rinsing the sections in PBS at room temperature, EdU labeling was next carried out in some experiments as specified by the Click-iT EdU Alexa Fluor-647 Imaging Kit protocol (Cat# C10340). Before proceeding to antibody staining, all tissues were preincubated for 1–2 hr in 0.1% Triton X-100 and 5% Normal Goat Serum (NGS; Jackson Immunoresearch, Cat# 005-000-121) in PBS at room temperature. Slices were then incubated with primary antibody for 1.5 h–2 days as needed in 0.1% Triton X-100 and 5% NGS in PBS at 4 °C. The primary antibodies used were rabbit anti-NeuN (Abcam, Cat# ab177487, 1:3000), mouse anti-NeuN (Abcam, Cat# ab104224, 1:1000), mouse antimyelin CNPase (clone SMI 91, Biologend, Cat# 836404, 1:1000), rabbit anti-Cux1 (Novus Biologicals, Cat# NBP2-13883, 1:100), mouse anti-TLE4 (E-10) Alexa Fluor 647 (Santa Cruz Biotechnology, Cat# sc-365406 AF647, 1:100), mouse anti-GAD-67 (EMD Millipore, Cat# MAB5406, 1:1000), rabbit anti-Iba1 (Wako, Cat# 019-19741, 1:600), mouse anti-Aldh1l1 (clone N103/39, NeuroMab, Cat# 75-140, 1:500), chicken antineurofilament NF-H (EnCor, Cat# CPCA-NF-H, 1:500), chicken anti-GFP (Abcam, Cat# ab13970, 1:500), mouse anti-myelin CNPase (clone RX-20, Leica objectives. All fixed-tissue images of layer I heterotopia at all identified IUE injection sites. Coronal sections from P0-injected mice were screened in a similar manner for injected areas, which were identified by virally transfected axonal fibers and neuronal cell bodies concentrated around visible needle tracts (data not shown).

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**Fixed-Tissue Imaging and Quantification**

Images of stained sections were collected using a Leica SP5 upright confocal laser scanning microscope. Lower magnification views of analyzed sections for presentation were acquired using ×10 and ×20 Leica objectives. All fixed-tissue images for quantification were captured through a ×40 Leica water immersion objective. For immunohistochemical analyses of in utero–injected mice, “heterotopia” were defined as the layer I cortical regions demarcated by ectopic NeuN+ cell body clusters identified at IUE needle tract sites. For P0-injected mice, because the heterotopia associated with needle tracts were more variable in size, ranging from several cells to 350 μm in diameter (data...
not shown), data for these injected areas were obtained from 300 μm × 90 μm FOVs centered on visible needle tracts that were within the first 100 μm below the pial surface. Control, noninjected regions were specified on corresponding contralateral cortices at similar mediolateral distances from the midline. Identical laser output and image acquisition configurations were used to capture images of both heterotopia and contralateral control regions throughout each immunostaining set.

All data for Aikd111, Iba1, CNPase, and GAD-67 cell density quantifications were acquired from the most superficial 100 μm of cortical tissue of both heterotopic and contralateral homologous control regions in layer I. The number of cell bodies confirmed by DAPI labeling that were positive for each marker was manually counted in randomly selected volumes within z-stacks acquired from both regions. The volumes were kept at identical dimensions between heterotopia and contralateral control regions for each experiment. For GAD-67 immunostaining, the presence of neuronal cell bodies was also confirmed by NeuN labeling. Two coronal sections containing heterotopia were analyzed per animal, with each immunostaining set comprising n = 6 animals. Statistical analyses were performed using Wilcoxon matched pairs, signed-rank nonparametric tests.

Data for CNPase density quantifications were acquired from the most superficial 100 μm of cortical tissue in both heterotopia and contralateral control regions. CNPase intensity was assessed using an automated thresholding and binarization plugin in ImageJ (Robust Automatic Threshold parameters set to noise = 1, lambda = 2, min = 208) for equally sized ROIs that were randomly selected from within heterotopic and contralateral control z-projections (15 μm thickness). CNPase density values for each section were determined by subtracting the automated measurements made in additional background regions from those made in assessed heterotopic and control regions. Two coronal sections containing heterotopia were analyzed per animal (n = 6 animals). Statistical analyses were performed using Wilcoxon matched pairs, signed-rank nonparametric tests.

To examine the neuronal composition of the embryonically induced heterotopia, coronal sections were stained for NeuN, DAPI, and one of the other following stains: Cux1, Tle4, or EdU, as indicated in the text. Data were acquired from within the most superficial 100 μm of the neocortex for layer I heterotopia and across the entire cortical wall for contralateral homologous control regions. For quantification of layer I heterotopia, equally sized volumes of interest (VOIs) were randomly selected from within the first 100 μm below the pial surface. For quantification of control regions on the contralateral hemisphere, the cortical wall was partitioned into 11 equal volume bins extending from the pial surface to white matter, with Bin1 encompassing layer 1 and Bin 11 adjoining white matter at its lower boundary. For each heterotopic VOI and control bin, the 1) number of NeuN+ cells and 2) proportion of NeuN+ cells that were also Cux1, Tle4, or EdU positive were manually quantified in ImageJ. Cells were identified as EdU positive if >50% of their nuclear volume, defined by DAPI, was occupied by EdU. Two sections were analyzed per mouse, with sample sizes for each group denoted in the text and figure legends.

The sites of injection analyzed in this study were positioned in somatosensory cortices and immediately adjacent areas in both male and female mice. Given the minimal variation in the morphological appearances of needle tracts between male and female mice and across different cortical areas, we pooled together these variables in our analyses. No animal subjects or experimental data points were excluded from analysis, and the nature of our study did not require subject randomization or experimenter blinding. No statistical methods were used to predetermine sample sizes, although our sample sizes are comparable to those published and generally accepted in the field. GraphPad Prism 7 was utilized for all statistical analyses, and Wilcoxon matched pairs signed-rank nonparametric tests were used to determine statistical significance (declared for P values below 0.05; two-tailed) because a normal distribution of differences between the paired data could not be assumed. Data are reported as mean ± SEM, unless otherwise noted.

**Results**

**Spatiotemporally Precise Layer I Heterotopia Generation and Visualization in the Live Mouse Brain**

In utero electroporation (IUE) is a powerful technique that permits labeling and genetic manipulation of targeted cortical neurons (Saito and Nakatsuji 2001; Tabata and Nakajima 2001; Lo Turco et al. 2009). As part of the IUE procedure, a thin glass microcapillary needle is used to deliver genetic material into the lateral ventricle of embryonic stage animals for the electropropagation of neuronal progenitors (Fig. 1a). While implementing IUE to label and image cortical axons in vivo, we unexpectedly found that cortical sites that had been injected during the procedure developed marked accumulations of labeled neuronal cell bodies in layer I of adult mice (Fig. 1c, top row, and Supplementary Fig. 1a). Electroporated pyramidal cells are normally found in layers II/III of cortical regions; thus, their distinct presence at injected layer I sites stood out in contrast to noninjected surrounding cortical areas (Fig. 1c, top row, and Supplementary Fig. 1a). In vivo labeling of neurons with a fluorescent dye called NeuO (Er et al. 2015) further revealed that the IUE-labeled neuronal cell bodies constituted only a small fraction of the total neuronal cell bodies ectopically positioned (Fig. 1b and c; for detailed statistics, see Supplementary Table 1; mean diameter of NeuO+ ectopic cell clusters 474 ± 174 μm SD in n = 23 mice). Abnormal superficial clustering of neuronal cell bodies is a defining feature of layer I heterotopia (Jacob 1940; Morel and Wildi 1952; Schulze and Braak 1978; Kaufmann and Galaburda 1989; Meencke and Veith 1992; Yamamoto et al. 1997), a subtype of neocortical heterotopia that has been linked to learning impairments in humans (Galaburda and Kemper 1979; Galaburda et al. 1985; Humphreys et al. 1990). Their presence demonstrates that direct injections into the cortex performed as part of the IUE procedure can induce neocortical heterotopia amenable to precise molecular and cellular studies in vivo.

**Aberrant Axonal and Myelin Patterns Characterize Layer I Heterotopia Formed during a Critical Embryonic Time Window**

Using this methodology, we set out to examine the cellular composition of the layer I heterotopia. Interestingly, we observed markedly aberrant patterns of myelinated axons exclusively associated with the ectopic neuronal clusters (Figs 2a and 3a, Supplementary Fig. 1a,b, and Supplementary Videos 1 and 2), as visualized in vivo by SCoRe microscopy, a technique that enables high-resolution label-free imaging of myelinated axons (Schain et al. 2014; Hill et al. 2018). This was further corroborated by fixed-tissue immunohistochemistry of myelin markers, showing increased oligodendrocyte and myelin densities at the sites of injection (Figs 2b–d and
To investigate whether the marked axonal and myelin changes were dependent on induction of heterotopia at specific developmental ages, we performed IUE at various time points during cortical development between embryonic days 14 (E14) to E17 and postnatal injections at P0. We found that whereas embryonic injections always resulted in similar nest-like patterns of densely myelinated fiber bundles around heterotopic neurons (100% frequency in n > 30 mice; Figs 4 and 5b), P0 injections did not result in robust axon guidance or myelinization abnormalities, despite the presence of ectopic neural clusters (Fig. 5a). Although postnatally induced heterotopia tended to be smaller and more variable in size (data not shown), the axonal and myelin abnormalities were never observed even around larger P0-induced clusters (Fig. 5a). Consistent with this, we found no significant difference in the myelin or oligodendrocyte cell body densities (Fig. 5; for detailed statistics, see Supplementary Table 1) between heterotopia and contralateral control regions of P0-injected mice. Likewise, immunolabeling for neurofilament heavy chain (NF-H) did not reveal aberrant accumulations of axonal fibers (Fig. 5a). Together, our data indicate previously unrecognized differences in the cellular composition and organization of neocortical heterotopia based on the timing of their induction.

Astrocyte and Microglia Cell Densities Are Not Altered in Induced Heterotopia

Given the striking abnormalities in oligodendrocyte density, myelination and axon pathfinding observed in the embryonically induced heterotopia, we next wondered whether other glial cell types such as astrocytes and microglia also exhibited altered morphology or density. Immunolabeled cortical brain sections against Aldh1l1 revealed no significant difference in the astrocyte cell density between layer 1 heterotopia and corresponding contralateral control regions (Fig. 6a–c; for detailed statistics, see Supplementary Table 1). Similarly, using Iba1 immunolabeling, we found no regional differences in microglia density or morphology (Fig. 6d–f; for detailed statistics, see Supplementary Table 1). Moreover, there was no evidence of pronounced astrocytic or microglial accumulation or altered cellular morphology at the borders of the heterotopia suggesting that a glial scar had not formed due to the embryonic injection. These data suggest that while heterotopia induce marked oligodendrocyte generation, astrocytes and microglia are not significantly influenced by potential local factors derived from ectopic neuronal clusters.

Heterotopia Contain a Variety of Excitatory and Inhibitory Neurons Born at Different Embryonic Ages

To characterize the neural composition of the embryonically induced heterotopia, we examined the expression of neuronal subtype markers Cux1, Tle4, and GAD-67 (Fig. 7). Cux1 is a transcription factor predominantly expressed in callosal projection neurons in layers II–IV (Molyneaux et al. 2009), whereas Tle4 is primarily restricted to deeper corticothalamic projection neurons of layers V and VI (Molyneaux et al. 2015; Sorensen et al. 2015). We found both Cux1+ and Tle4+ neurons present to varying degrees in all layer I heterotopia, consistent with a mixed population of excitatory projection neurons from different cortical layers (Fig. 7bc and Supplementary Fig. 3). Further analyses revealed the presence of GAD-67+ interneurons, which...
Figure 2. Heterotopia contain an abundance of myelinated axons. (a) In vivo combined fluorescence and label-free SCoRe myelin images of a representative induced heterotopia (right; same as in Fig. 1c) and neighboring ipsilateral, noninjected control region (left) in a P30 mouse. The tightly packed heterotopic neuronal cell bodies (NeuO) are surrounded by a circumscribed abundance of aberrantly projecting, myelinated axon segments (SCoRe). (b, c) Low magnification (b) and high magnification (c) immunostainings of an induced heterotopia and its corresponding contralateral control region taken from a P30 mouse, showing increased oligodendrocyte CNPase expression associated with the heterotopia, as defined by the ectopically positioned NeuN+ neuronal cell bodies in layer I (b, white arrowheads). Notice the more horizontal orientation of myelin segments located closer to the pial surface of the heterotopia (c, blue arrowhead). Deeper myelin segments, in contrast, are organized in a more radial fashion (c, blue arrow) and fasciculate into densely myelinated fiber bundles that project through lower cortical layers (b, white arrow). (d) Quantifications of SCoRe in vivo (n = 8 mice, top) and CNPase in fixed tissue (n = 6 mice, middle and bottom), showing increased myelination and oligodendrocyte cell body densities in induced layer I heterotopia compared with noninjected contralateral control regions. Each point represents the layer I heterotopion or control region from one animal. Horizontal lines and error bars denote mean and SEM, respectively (Wilcoxon matched-pairs signed rank nonparametric test; *P < 0.05, **P < 0.01). HTP, Heterotopia. Descriptive statistics are given in Supplementary Table 1. Scale bars, 50 μm (a), 200 μm in (b), and 50 μm (c).
Figure 3. Myelinated and unmyelinated axons follow aberrantly looping and concentric paths. (a–d) Intravital imaging of a nest-like heterotopion at P30 in (a) using SCoRe and confocal fluorescence microscopy showing b, labeled neuronal cell bodies (tdTomato; green) dispersed throughout the aberrantly oriented myelinated fibers (SCoRe; magenta) and c, winding axonal projections that occasionally fasciculate into bundles at the edges of the heterotopion. (d) Myelinated (arrowheads) and unmyelinated axon segments (arrows) are observed within the heterotopion. (e–g) Low-magnification (e) and high-magnification (f, g) immunostainings of an embryonically induced heterotopion (g), confirming the presence of both myelinated (g, arrowhead) and unmyelinated (g, arrow) axon segments (f and g show high-magnification images of areas indicated by arrow and arrowhead in e, respectively). NF-H, neurofilament heavy chain. Images are representative of experiments performed in at least six animals. Scale bars, 100 μm (a), 15 μm (b–d), 100 μm (e), and 25 μm (f, g).
occurred at similar densities within heterotopia as in corresponding layer I control regions (Fig. 7d–f; for detailed statistics, see Supplementary Table 1). In addition, layer I heterotopia always contained GAD-67+ puncta, consistent with inhibitory synapses (Fig. 7d,e). Together, our findings suggest that embryonically induced layer I heterotopic neural clusters comprise a diverse cohort of glutamergic and GABAergic neurons, in line with the heterogeneous population of neuronal subtypes previously described in those occurring spontaneously (Gabel and LoTurco 2001; Ramos et al. 2008, 2014; Gabel 2011).

We next used birthdating techniques to determine whether neurons within the induced heterotopia originate at similar or different time points during development. Edu pulse labeling at E11.5 revealed small but distinct populations of Edu+NeuN+ cells in all embryonically-generated layer I heterotopia (Supplementary Fig. 4), suggesting that heterotopia contain neurons born during early corticogenesis (Angevine and Sidman 1961; Raedler and Raedler 1978). We also observed cells "birthdate-labeled" via IUE at E15 in heterotopia (~90% heterotopia in n > 30 mice; Fig. 1c), suggesting the additional presence of later-born neurons (Lo Turco et al. 2009). These data indicate that induced heterotopia contain neurons born at various stages of cortical development.

Heterotopic Neurons Exhibit Similar Calcium Dynamics as Neighboring Normotopic Neurons

Despite their linkage to a wide spectrum of neurological disorders, the cellular dynamics and mechanisms by which individual heterotopic cells contribute to disrupted neural circuit functioning are poorly understood. We examined the calcium activity of single cortical neurons in embryonically induced heterotopia and in surrounding, noninjected cortical layer II/III regions of awake head-fixed mice using the genetically encoded calcium indicator, GCaMP6f. Surprisingly, although we found highly variable calcium spike patterns of individual cells within heterotopia (Fig. 8a–c and Supplementary Video 3), there were no significant differences in their overall calcium spike event frequency, variance, or synchrony compared with surrounding nonheterotopic layer II/III regions (Fig. 8d; for detailed statistics, see Supplementary Table 1). Furthermore, we did not identify any epileptiform activity in our imaging sessions, although we cannot rule out that continuous recordings could have revealed sporadic aberrant activity (Fig. 8a–c and Supplementary Video 3). Interestingly, we found that mice that had been started with brief whisker stimulation consistently responded via neuronal calcium spikes within layer 1 heterotopia (Fig. 8ef and Supplementary Video 4). This result indicates that heterotopic neurons are connected to other brain areas and could thus influence network function.

Discussion

Here, we describe an inducible model that allows the visualization of focal heterotopia at cellular resolution in the live animal. Building upon previous models, this system employs in utero cortical microinjections to induce layer 1 heterotopia that closely recapitulate those seen in humans (Jacob 1940; Schulze and Braak 1978; Galaburda et al. 1985; Kaufmann and Galaburda 1989; Humphreys et al. 1990; Meencke and Veith 1992; e.g., see Fig. 7 in Jacob; Fig. 1b in Humphreys et al.; Fig. 3 in Meencke et al.) and in spontaneous animal models (Sherman et al. 1985; Sherman et al. 1987; Ramos et al. 2008; Lipoff et al. 2011; Toia et al. 2017; e.g., see Fig. 1 in Sherman et al. 1987; Fig. 1 in Ramos et al.; Fig. 1 in Lipoff et al.). Pairing this methodology with intravital fluorescence and label-free (SCoRe) microscopy of neuronal cell bodies, axons and myelin, as well as time-lapse axonal and calcium imaging (Saito and Nakatsuji 2001; Lo Turco et al. 2009; Schain et al. 2014; Er et al. 2015; Hill et al. 2018), enables detailed studies of heterotopia in the intact brain.

Many animal models of focal heterotopia have been described; however, most involve subcortical or intrahippocampal heterotopia (Riggs et al. 1956; Singh 1977; Amano et al. 1996; Lee et al. 1997; Bai et al. 2003), which are not easily amenable to intravital optical imaging due to their distance from the cortical surface. Moreover, other animal models of more superficially occurring heterotopia (Sherman et al. 1987; Rosen et al. 1992, 2007; Ferrer et al. 1993; Brunstrom et al. 1997) have not been imaged successfully at high resolution in vivo mainly as a result of their temporally and spatially unpredictable nature and/or lack of visible cell transfection. Our methodology improves upon these limitations by employing minimally invasive microinjections to both induce layer I heterotopia and deliver plasmid DNA and/or viral vectors for cellular genetic manipulation in the live mouse.

Using our model, we revealed several striking characteristics of neocortical heterotopia. First, we discovered that axonal projections within layer 1 heterotopia formed swirled and concentric morphologies specifically heterotopic neuronal
clusters. These aberrantly projecting swirling axons were associated with radially oriented fibers extending through the base of the heterotopia, consistent with previous fixed-tissue studies of spontaneously occurring heterotopia (Sherman et al. 1990; Ramos et al. 2008). Interestingly, there was no evidence of gliotic scarring or mechanical tissue barriers surrounding the heterotopia, suggesting that the directional changes in axon pathfinding are instead likely mediated by disruptions in the precise balance of repulsive and attractive gradients of local axon guidance cues (Métin et al. 1997; Richards et al. 1997; Bagnard et al. 1998; Gao et al. 1998; Polleux et al. 1998). Future studies of the individual neuronal identities of the aberrantly projecting axons, which could derive from various cortical and subcortical sources (Jenner et al. 2000; Dwyer et al. 2011; Gabel 2011; Ramos et al. 2014), may shed light on the possible molecular factors leading to the disrupted axon pathfinding.

The axon guidance abnormalities led us to investigate whether the developmental timing of heterotopia induction could modulate their formation. We found that whereas embryonically induced heterotopia always displayed distinct axon guidance abnormalities, similar defects were not observed in postnatally induced heterotopia. One potential explanation for this surprising finding is the presence of a critical developmental time period during which cortical axon pathfinding is uniquely sensitive to local environmental disturbances. The distinct responsiveness of axons during this critical period could be mediated by the enhanced expression of axon growth cone receptors that increase the ability of axons to extend toward guidance cues (Shewan et al. 2002; Baudet et al. 2012). However, developmentally modulated changes in the signaling gradients of guidance cues themselves could also play a role (Skaliora et al. 1998). An alternative explanation is the induction of subtle glial alterations that could mechanically impair axon outgrowth, although the lack of glial changes in our images does not favor this hypothesis.

In addition to the aberrant axonal organization, we used label-free SCoRe in vivo imaging to discover focally increased myelination specifically within the heterotopia. This striking change was due to an increased density of local myelinating oligodendrocytes that deposited myelin sheaths primarily along

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Figure 5. Postnatally induced heterotopia do not develop aberrant axons or myelination. (a) Confocal images of an immunolabeled coronal section from P30 mouse cortex, showing no changes in myelination (CNPase) or axonal pathfinding (NF-H) associated with a P0-induced heterotopion (NeuN; arrowheads), compared with its contralateral control region (left). (b) In contrast, neuronal heterotopia induced in utero at E15 (NeuN; arrowheads) show dramatic changes in local layer I myelin (CNPase) expression and patterning. (c) Quantifications of CNPase expression, showing no significant differences in the myelin or oligodendrocyte cell body densities (n = 6 animals) between postnatally induced heterotopia and control regions. Each point represents the P0-induced layer I heterotopion or control region from one animal, with horizontal lines and error bars denoting mean and SEM, respectively. Wilcoxon matched-pairs signed rank nonparametric test was used to determine significance; NS, no significance. NF-H, neurofilament heavy chain. Descriptive statistics are given in Supplementary Table 1. Scale bars, 100 μm (a, b).
Figure 6. Astrocytes and microglia retain a normal density despite aberrant axonal and myelin distribution. (a, b) Low-magnification (a) and high-magnification (b) images of Aldh111-positive astrocytes in P30 mouse cortex, showing similar cell densities in layer I heterotopia (right images, arrowheads) compared with corresponding contralateral control regions (left images). (c) Quantifications using P30 mouse forebrain tissue showing no significant difference in astrocyte cell density (Wilcoxon nonparametric matched-pairs, signed rank test, n = 6 animals; ns, no significance). (d, e) Low-magnification (d) and high-magnification (e) images of Iba1-immunolabeled P30 mouse forebrain, showing no difference in microglia cell densities between heterotopia (right images, arrowheads) and contralateral control regions (left images). (f) Quantifications using P30 mouse forebrain tissue demonstrating no significant difference in microglia cell density between layer I heterotopia and corresponding contralateral control regions (Wilcoxon matched-pairs, signed rank nonparametric test, n = 6 animals; ns, no significance). For graphs in (c) and (f), each point represents an embryonically induced layer I heterotopia or control region from one animal, with horizontal lines and accompanying error bars denoting mean and SEM, respectively. Descriptive statistics are given in Supplementary Table 1. HTP, heterotopia. Scale bars, 100 μm (a), 20 μm (b), 100 μm (d), and 20 μm (e).
Heterotopia contain a mixed population of excitatory and inhibitory neurons. (a) Approach for quantifying layer-specific neuronal marker expression (symbolized by blue and green dots) in layer I heterotopia and across corresponding contralateral control cortices in P30 mice. Contralateral control cortices are subdivided into 11 equally sized bins that together span the entire thickness of the cortex. (b, c) Both Cux1+ (b, cyan) and Tle4+ (c, green) neuronal cell bodies occur within layer I heterotopia (right images, arrowheads). In contralateral control regions (left images), Cux1 is concentrated in more superficial cortical areas corresponding to layers II/III, whereas deeper areas corresponding to layers V/VI encapsulate the majority of Tle4 labeling. Quantifications (far right) show the percentages of NeuN+ cell bodies that also express Cux1 (top) or Tle4 (bottom) within heterotopia and across corresponding contralateral control bins. Each dot corresponds to the layer I heterotopia or control bin of a single animal, with all dots of the same color belonging to the same animal (n=4 animals for Cux1; n=5 animals for Tle4). The black line denotes the mean. WM, white matter.

(d, e) Low- (d) and high- (e) magnification z-projections of GAD-67-immunostained P30 mouse forebrain, showing the presence of GAD-67-labeled cells and puncta in both layer I heterotopia (right images, arrowheads) and corresponding contralateral control regions (left images). Quantifications from P30 mouse tissue showing no significant difference in GAD-67+ neuronal cell density between layer I heterotopia and contralateral control regions (Wilcoxon matched-pairs, signed rank nonparametric test, n=6 animals; ns, no significance). Each point represents the layer I heterotopia or contralateral control region from one animal, with horizontal lines and accompanying error bars denoting mean and SEM, respectively. HTP, heterotopia. Descriptive statistics are given in Supplementary Table 1. Scale bars 200 μm (b, c), 100 μm (d), and 10 μm (e).

Figure 7. Heterotopia contain a mixed population of excitatory and inhibitory neurons.
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Figure 8. Heterotopic neurons display calcium dynamics similar to adjacent layer II/III neurons and respond to sensorimotor input. (a–c) In vivo time-lapse imaging of layer I heterotopic neurons expressing the calcium sensor GCaMP6f in an awake, head-fixed mouse. Dotted lines in (a) indicate an analyzed heterotopic region that is displayed in (b). (d) Example traces of spontaneous neuronal calcium transients from layer I heterotopic (right) and ipsilateral layer II/III nonheterotopic (left) regions displayed in (b). (e) Quantifications showing no significant difference in calcium spike event frequency (Hz), variance (SD), or global synchronization index between layer I heterotopia and adjacent layer II/III nonheterotopic regions. Each dot corresponds to the average value for each measure in a layer I heterotopic or nonheterotopic layer II/III region from a single animal. Data represent 493 layer I heterotopic neurons and 594 layer II/III neurons from n = 7 mice. Wilcoxon matched-pairs, signed rank nonparametric test was used for all three quantifications; ns, no significance; HTP, heterotopia. Descriptive statistics are given in Supplementary Table 1. (e) In vivo z-projection of the same GCaMP6f-transfected heterotopia displayed in (a), captured at a different depth, showing neurons that were imaged during brief applications of whisker stimulation. (f) Baseline calcium dynamics (left column) and calcium fluctuations (middle and right columns) of individual heterotopic neurons in response to stimuli (denoted by solid blue line in two trials) obtained from the awake, head-fixed mouse imaged in (e). Experimental observations in (e,f) were replicated in three mice. All data and images from (a–f) were obtained from P50 to P60 mice with heterotopia induced at E15 and virally transfected with AAV-GCaMP6f at P21–P30. Some neuronal cell body and axonal labeling by pCAG-TdTomato (red) is the result of E15 IUE-mediated transfection. Scale bars, 50 μm (a), 25 μm (b), and 50 μm (e).

The aberrantly projecting and concentric axons. Interestingly, the abundant myelin phenotype was only observed in embryonically induced but never in postnatally induced heterotopia. Similar to the critical period for disrupted axonal patterning, these data suggest that the excess myelination seen in embryonically induced heterotopia is instructed by developmentally regulated molecules or biophysical cues, which could accelerate the local production of myelin by stimulating oligodendrocyte precursor cell recruitment and/or differentiation into mature oligodendrocytes. Myelin formation is influenced by axon caliber (Sturrock 1980) and genetic manipulations increasing axon caliber can trigger oligodendrocyte precursor cell proliferation and the ensheathment of classically unmyelinated fibers (Goebbels et al. 2017). Thus, axon caliber could serve as one mechanism that differs between the aberrantly projecting axons found in embryonically induced heterotopia compared with adjacent nonmyelinated fibers. Other potential causes for the increased myelin deposition around heterotopic axons could include cell surface markers or molecular identity. Specific subpopulations of excitatory (Tomassy et al. 2014) and inhibitory (Micheva et al. 2016) neurons exhibit variable myelination patterns, seemingly not directly linked to axon caliber, but instead potentially due to their differential expression of adhesion molecules or altered patterns of neuronal activity.
Our methodology now provides a means to test these and other possibilities in precise, real-time live animal studies.

Despite the striking axon and myelin abnormalities associated with the induced heterotopia, we found no evidence of spontaneous epileptiform activity occurring in heterotopic regions or nearby nonheterotopic cortices of awake animals, consistent with previous studies (Roper et al. 1995; Gabel and LoTurco 2001; Ishii et al. 2015). Moreover, spontaneous calcium fluctuations and the relative synchrony of firing between neurons in layer I heterotopia and neighboring layer IV/III cortex were similar. This lack of observable difference in activity may reflect the similar neural and glial composition of both brain areas, which could enable a similar balance of excitatory and inhibitory inputs to these regions. Indeed, we identified microglia, astrocytes, Cux1+ cells, GABAergic neurons, and GABAergic puncta in layer I heterotopia that are also prevalent in layers II/III (Chmielowska et al. 1988; Nieto et al. 2004; Molyneaux et al. 2009; Meyer et al. 2011; Farhy-Tselnicker and Allen 2018; Jara et al. 2019). A second, nonmutually exclusive possibility is that more metabolically demanding conditions are required to trigger aberrant activity in mice with induced heterotopia. Consistent with this hypothesis, the application of normally subthreshold doses of convulsant drugs can provoke epileptiform activity in mice with heterotopia in vitro (Gabel and LoTurco 2002) and in vivo (Gabel and LoTurco 2002; Manent et al. 2009; Ishii et al. 2015). Other animal models of heterotopia, such as the tish mutant (Lee et al. 1997) and Ihara’s genetically epileptic rat (Amano et al. 1996) have documented spontaneous epileptiform activity in vivo. Reasons for this difference from our findings remain unclear; however, they may be related to disparities in the number, location, or size of heterotopia, or unknown off-target effects of the inherited mutations themselves in the genetic models.

There is little known about the functional dynamics of individual heterotopic neurons during behavioral stimulation in the live animal. Using time-lapse calcium imaging, we revealed that heterotopic neurons display robust responses following brief whisker stimulation in awake mice, consistent with the functional connectivity implicated by previous anatomical, electrophysiological, and behavioral studies (Lee et al. 1997; Chevassus-Au-Louis, Congar, et al. 1998a; Chevassus-Au-Louis, Rafiki, et al. 1998b; Jenner et al. 2000; Gabel and LoTurco 2001; Schottler et al. 2001; Ishii et al. 2015). This functional connectivity suggests that, even though we did not detect aberrant spontaneous activity within heterotopia, these neurons are integrated into the local neural network and are potentially capable of altering neural network function at baseline or under metabolically demanding conditions. There remains considerable debate about the exact role for layer I heterotopia in neurological disorders (Galaburda and Kemper 1979; Galaburda et al. 1985; Kaufmann and Galaburda 1989; Humphreys et al. 1990; Meencke and Veith 1992; Barkovich 1998). They may provoke more subtle cognitive and behavioral disturbances, as implicated by their widely distributed axon projections throughout both ipsilateral and contralateral cortical layers (Jenner et al. 2000; Gabel 2011; Ramos et al. 2014), and subcortical structures (Jenner et al. 2000; Ramos et al. 2014). Indeed, their presence has been linked to reading disability in humans (Galaburda and Kemper 1979; Galaburda et al. 1985; Humphreys et al. 1990) and changes in perceptual and mnemonic abilities in rodents (Waters et al. 1997; Clark et al. 2000; Frenkel et al. 2000; Hyde et al. 2000, 2001; Peiffer et al. 2001). There is evidence that the functional outcomes associated with individual heterotopia may vary with respect to their intracortical positioning (Hyde et al. 2001), which could be related to underlying differences in neuronal connectivity (Jenner et al. 2000; Ramos et al. 2014) or composition (Ramos et al. 2008).

Previous genetic and neuropathological studies (De Bernabé et al. 2002; Halfter et al. 2002; Moore et al. 2002; Devisme et al. 2012; Myshral et al. 2012; Radner et al. 2013) have suggested that layer I heterotopia arise from acquired defects in the integrity of pial basement membranes during corticogenesis (Nakano et al. 1996; Yamamoto et al. 1997; Romero et al. 2018). Consistently, we found that layer I heterotopia form only at pial surface sites that are previously punctured in utero, and never in noninjected cortical regions of CD1 mice, in accordance with the findings of previous studies (Rosen et al. 1993; Ramos et al. 2008). A breach in the pial basement membrane during corticogenesis could trigger the inappropriate migration of diverse neuronal subtypes into layer I by causing the displacement of local anchoring radial glial foot processes, which normally serve as scaffolds for radially migrating neuroblasts during cortical development (Rakic 1972; Nadarajah and Parnavelas 2002). This hypothesis is supported by histopathological studies that have revealed the protrusion of radial glial fibers through breaks in the pial basement membrane that coincide with the location of layer I heterotopia in both spontaneous and mutant animal models (Sherman et al. 1992; Myshral et al. 2012). Spontaneously occurring heterotopia in rodents (Sherman et al. 1990; Ramos et al. 2008; Lipoff et al. 2011) and in humans (Jacob 1940; Morel and Wildi 1952) display aberrant accumulations of superficial myelinated axons and radially oriented axon bundles that resemble those resulting from embryonic cortical injections (e.g., see Fig. 3 in Jacob; Fig. 8 in Morel et al.; Fig. 8 in Ramos et al. in comparison to induced heterotopia shown here, Fig. 2 and Supplementary Fig. 2). Future studies using our model, including at earlier developmental time points, may provide additional information about the cellular and molecular mechanisms by which layer I heterotopia form and lead to aberrant axon and myelin patterns, as well as their functional implications.

Supplementary Material
Supplementary material can be found at Cerebral Cortex online.

Author Contributions
A.M.L. and J.G. conceived the initial project. A.M.L., R.A.H., and J.G. designed the experiments. A.M.L. performed all experiments, analyzed the data, and prepared the figures. A.M.L., R.A.H., and J.G. wrote the manuscript.

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Notes
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