

Supporting Information

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SI Methods

Animals. We obtained estimates of neuron and other cell numbers across the cortical expanse from four species of primates. All brain tissue for these experiments was obtained from ongoing experiments of other investigators at Vanderbilt University or was purchased, so tissue preparation methods varied across cases but not in any way that would affect our cell counts. The analyzed material came from two flat hemispheres from two different prosimian galagos (*Otolemur garnetti*), one flat hemisphere from a New World owl monkey (*Aotus nancymae*), one hemisphere from an Old World macaque monkey (*Macaca mulatta*), and one flat hemisphere from an Old World baboon (*Papio cynocephalus anubis*). The baboon brain was purchased from the tissue program at the University of Washington National Primate Research Center (<http://www.wanprc.org/primate-resources/pathology-tissue-program/>).

Dissection of Cortex. All brains were perfused with PBS followed by 2–4% paraformaldehyde fixative. The baboon brain was perfused with PBS only and then was fixed by immersion in 4% paraformaldehyde after flattening and dissection. The cortical hemisphere from each galago, owl monkey, and baboon was separated from subcortical structures, the pia was removed, and the sulci were opened to flatten the cortex, as described elsewhere (1–3). In the owl monkey and in galago #2 (case 08–07), identifiable visual, somatosensory, auditory, and motor areas or regions were dissected after the flat cortex was viewed on a light box, so that myelin-dense sensory areas appeared dark relative to surrounding cortex (Fig. 1A). Dissected areas were processed individually. The flattened hemispheres from galago #1 (case 07–104) and the baboon (case 09–27) were dissected into a grid of 5 × 5 mm² tissue pieces (pieces along the margins of the hemisphere were smaller and irregular). Each piece was weighed, and, where possible, pieces were assigned to a cortical area identified by transmitted light. Some pieces were processed individually, and some were combined with small neighboring pieces.

Tissue Processing and Cell Counting. The isotropic fractionator method (4) was used to determine the number of neurons and

nonneuron cells in each sample. Details of processing steps are detailed in previous publications (e.g., ref. 5). Briefly, tissue samples from each case were homogenized in a dissociation solution of Triton X-100 and sodium citrate in distilled water, using glass Pyrex Tenbroeck tissue grinders (Fisher Scientific). The resulting suspensions contained cellular nuclei, with very few cell membranes remaining intact. Because tissue samples from the present experiment were immersed in fixative solution for at least 4 wk before dissociation and processing, all went through an epitope retrieval step, which consisted of 60 min in 0.2 M boric acid solution in an oven set at 70 °C, before processing for immunocytochemistry. Samples were spun down in a centrifuge and resuspended in a mixture of PBS and DAPI. All suspension volumes were between 3 and 15 mL and were determined based on approximate cellular density. DAPI-stained nuclei were counted on a fluorescence microscope (Nikon E-800, Nikon Instruments, Inc.) using a Neubauer counting chamber; then a sample was taken from the main suspension and stained for NeuN (Millipore, Inc.), an antibody that stains neuronal nuclear antigen expressed only in the cytoplasm and nuclei of neurons. A fluorescent secondary antibody, goat anti-mouse IgG-Alexa Fluor 594 (Invitrogen), was used to visualize stained nuclei, and the ratio of neurons to nonneuron cells was determined using a fluorescence microscope. For each sample, a minimum of 500 DAPI-positive nuclei were evaluated for labeling with NeuN/AF594.

Data Analysis. Percentages of neurons relative to nonneuron cells and neuron densities were calculated for all tissue samples. For the galago cortex (07–104) dissected into a grid of samples, surface areas were measured using ImageJ software (National Institutes of Health), and *x* and *y* centroids were determined for each tissue piece in the cortical grid (Fig. 1B). This process allowed us to produce contour plots of the cellular and neuron density data depicted in Fig. 1C and D, respectively, using the “Akima” package (6) of R statistical software (<http://www.r-project.org>). (See plot of neuronal density in the galago cortex, case 07–104; Fig. 1D.) Larger samples from the macaque monkey cortex were related to the estimated locations of primary sensory and motor areas as well as to several other previously described subdivisions of cortex.

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Other Supporting Information Files

[Dataset S1 \(XLS\)](#)