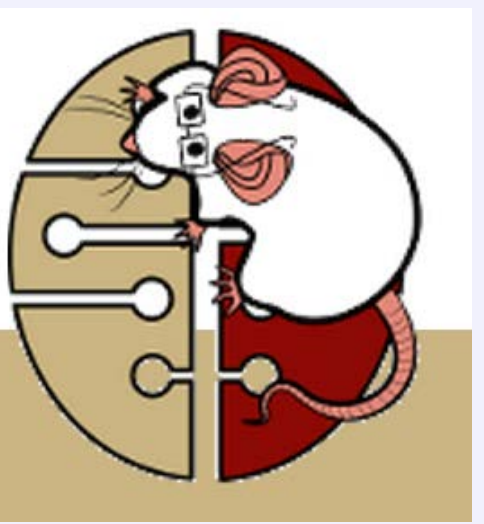




Cold Spring Harbor Laboratory  
Cold Spring Harbor, NY

# Combining high resolution MRI and DTI with dense whole-brain histology for mouse

V. Pinskiy  
A. Mukherjee  
A. Tolpygo  
A. Badea  
A.G. Johnson  
P. Mitra



Mouse Brain  
Architecture

## Overview

### Objective

With advances in digital imaging and computing technologies, there is increasing demand to cross-reference and freely share whole-brain image data sets of different modality. The Waxholm Space (WHS) digital atlas, now in its 3rd year and widely adopted, was developed to meet this demand by providing a common framework for whole mouse brain digital neuroanatomy.

The work we present is aimed at updating the histological component of the current WHS atlas with improved spatial resolution and image quality. We co-register the high-resolution whole-brain Nissl with simultaneously scanned MRI/DTI that represent significant advance in the quality of histological imaging for mouse brains. It will be made available online through the Mouse Brain Architecture Project website at <http://mouse.brainarchitecture.org>.

### Waxholm Space (WHS)

Ultra-high resolution MR atlas of the C57BL/6 mouse brain, co-registered with a whole-brain, segmented and annotated, Nissl-histology map of the same brain.

### Motivation

The existing Nissl map (obtained as horizontal stack) of the WHS can be improved upon by

- better preservation of tissue integrity
- better image quality
- higher digital spatial resolution – (current: 21  $\mu\text{m}$  section thickness; 9.9  $\mu\text{m}$  pxls)
- more accurate 3d-to-3d registration to MR atlas

## MR Methods & Data Sets

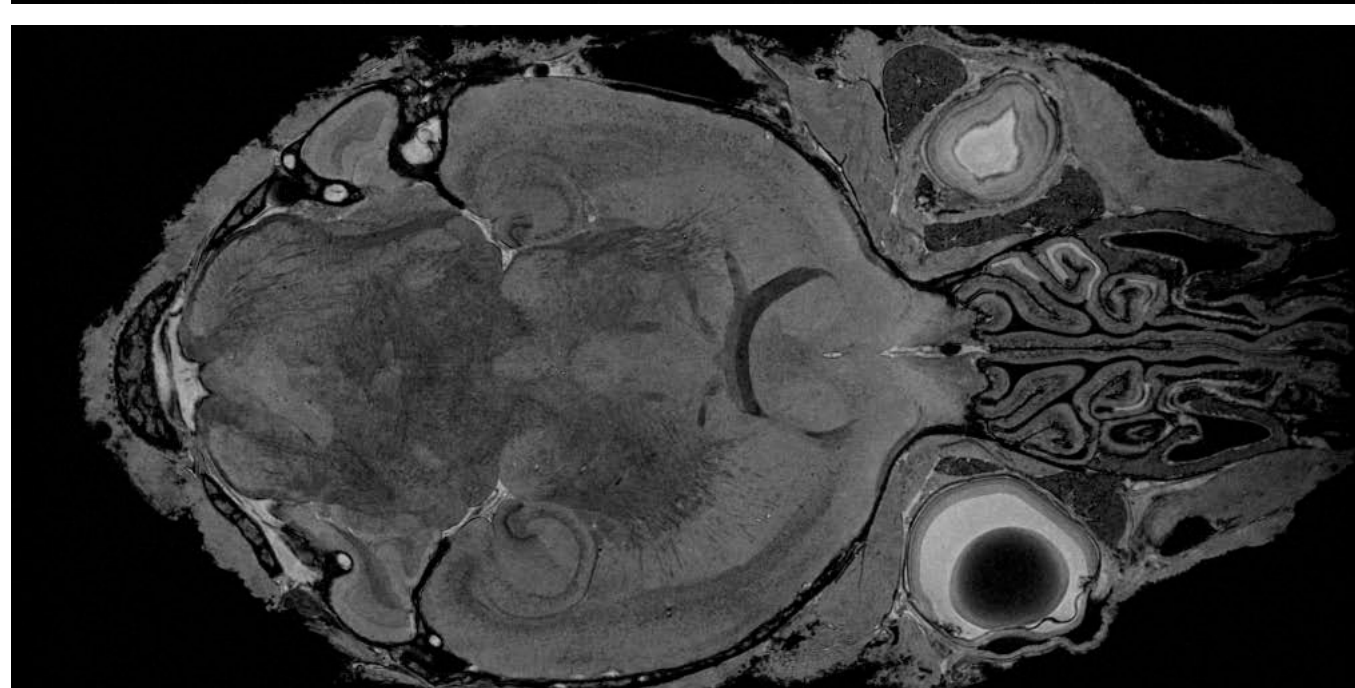
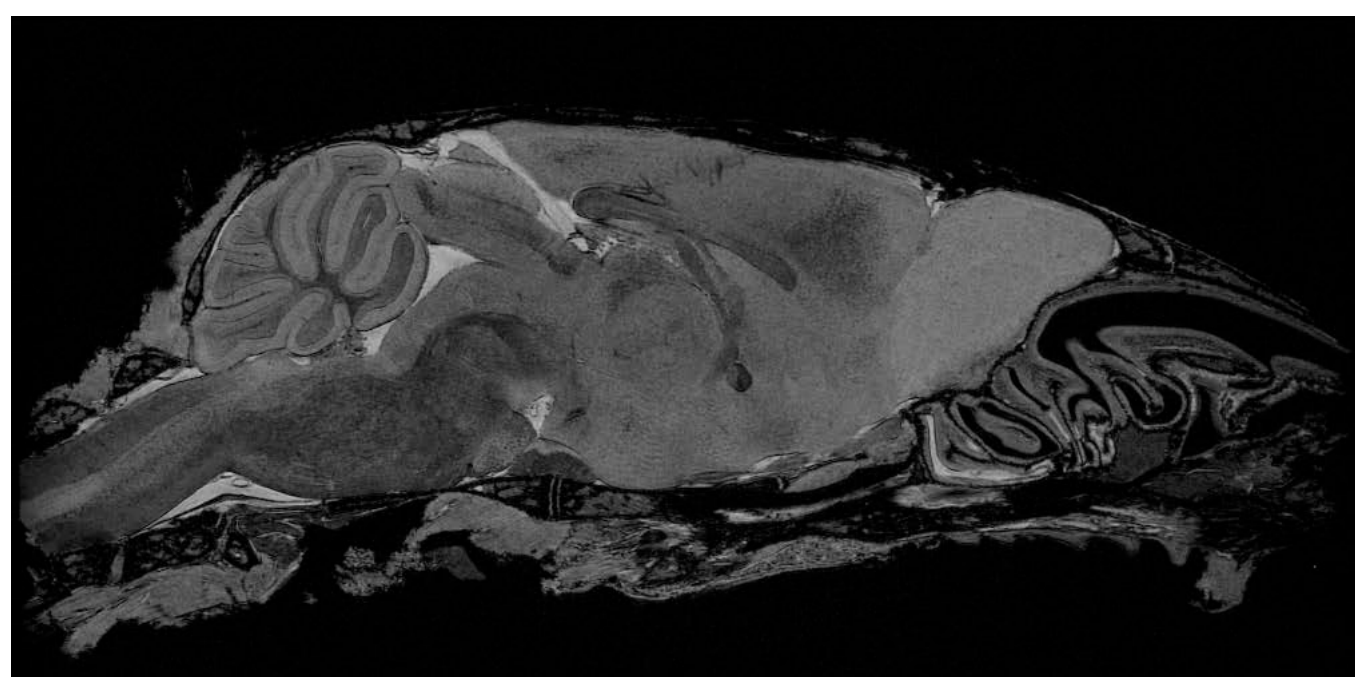
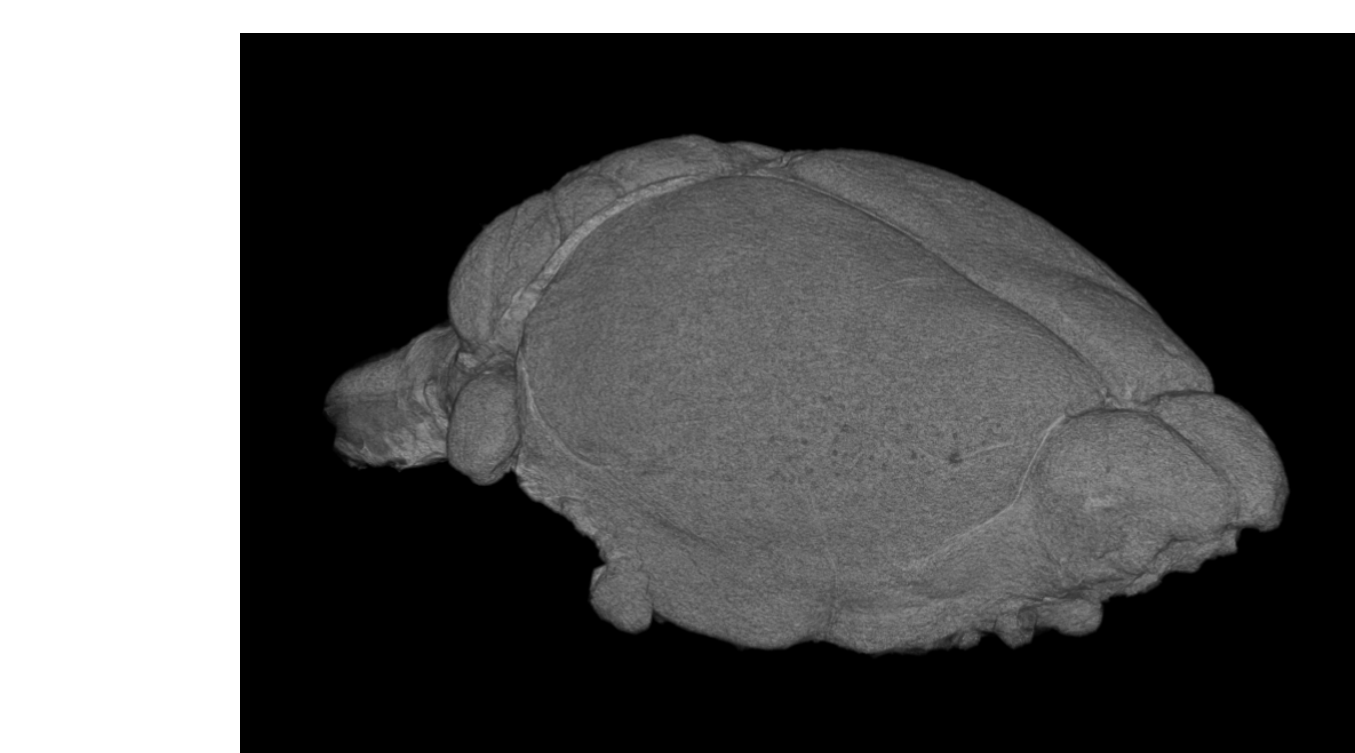
### Experiment

species C57BL/6 mice  
age / sex adult (P56) male  
brain preparation post-fixed intracranial  
fixative 10% buffered formalin (+ contrast agent)

### MR data acquisition

protocol T2\* -weighted  
sequence gradient-recalled echo  
TR 50.0 ms  
TE 4.3 ms  
field strength 9.4 T  
coil 12 mm  
gradient 860 mT/m  
contrast enhancement high-proton density active stain

(gadopentetate dimeglumine)  
acquisition matrix 768 x 512 x 512  
display matrix 1024 x 512 x 512  
image volume 3mm isotropic cube  
voxel resolution (MRI) 21.4  $\mu\text{m}$  isotropic  
voxel resolution (DTI) 43  $\mu\text{m}$  isotropic



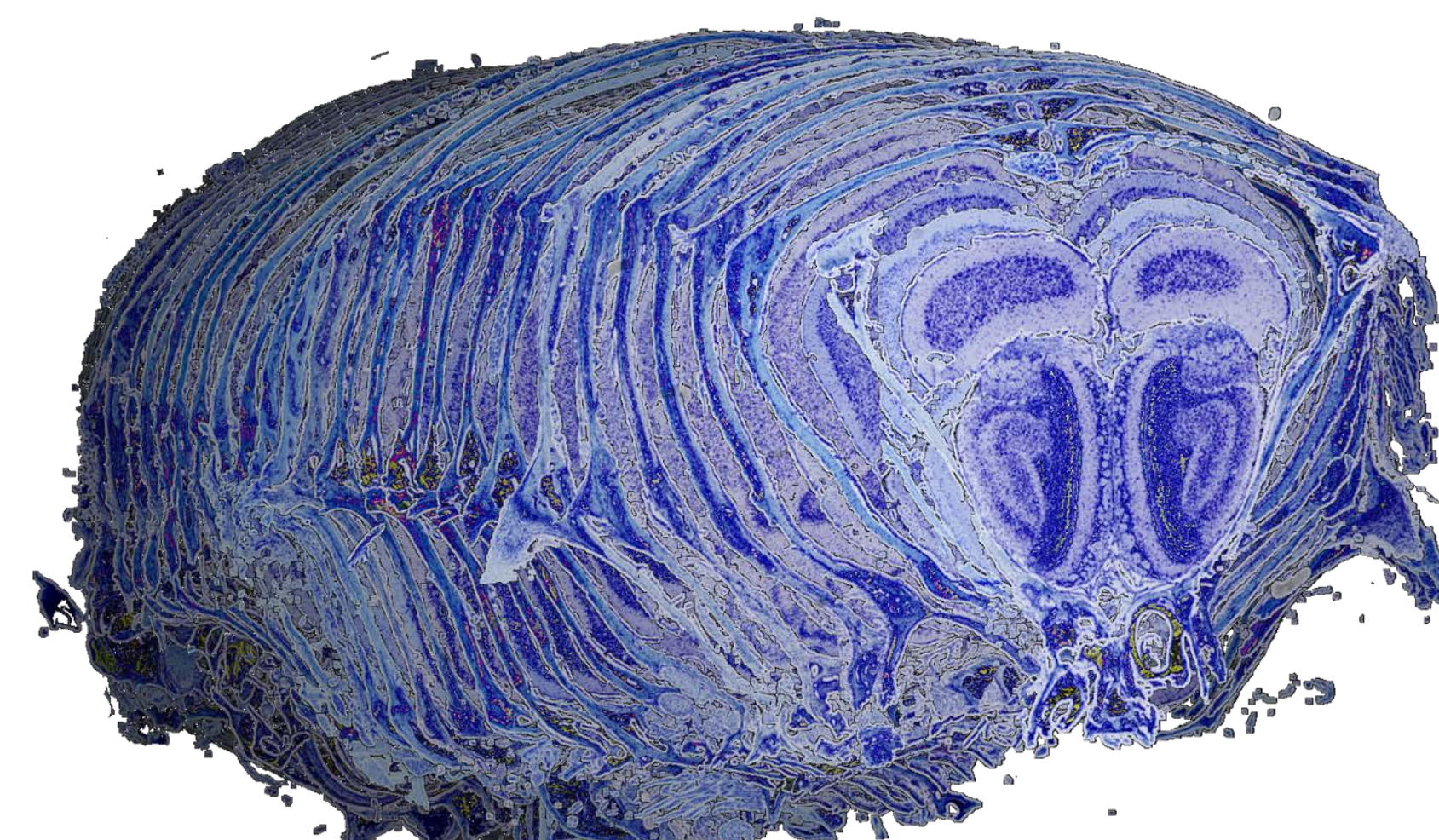
## MBA Pipeline

After the MR imaging, the perfusion-fixed brains of the same animals were histologically processed using the automated Mouse Brain Architecture Project (MBA) histological pipeline. The histological data set consists of a thin-sectioned, Nissl-stained, whole-brain, coronal stacks.



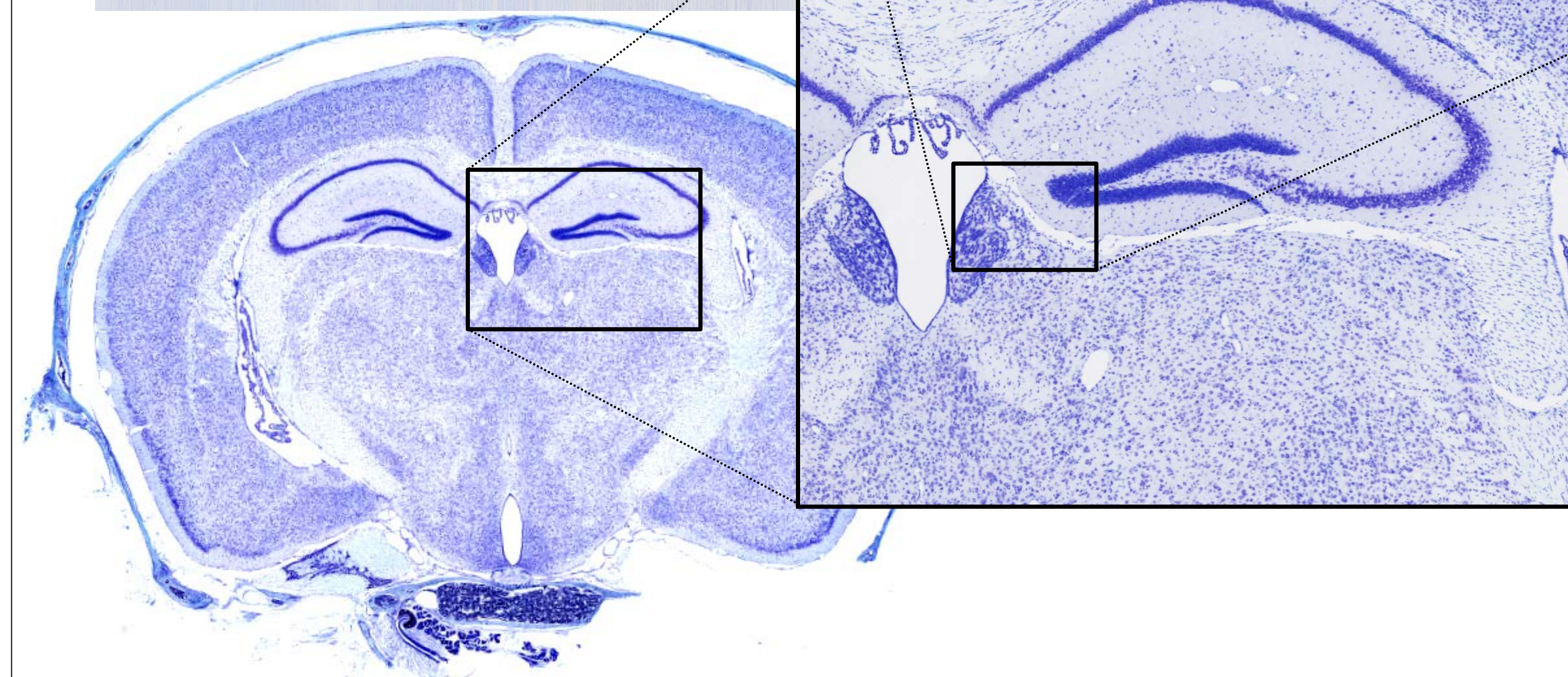
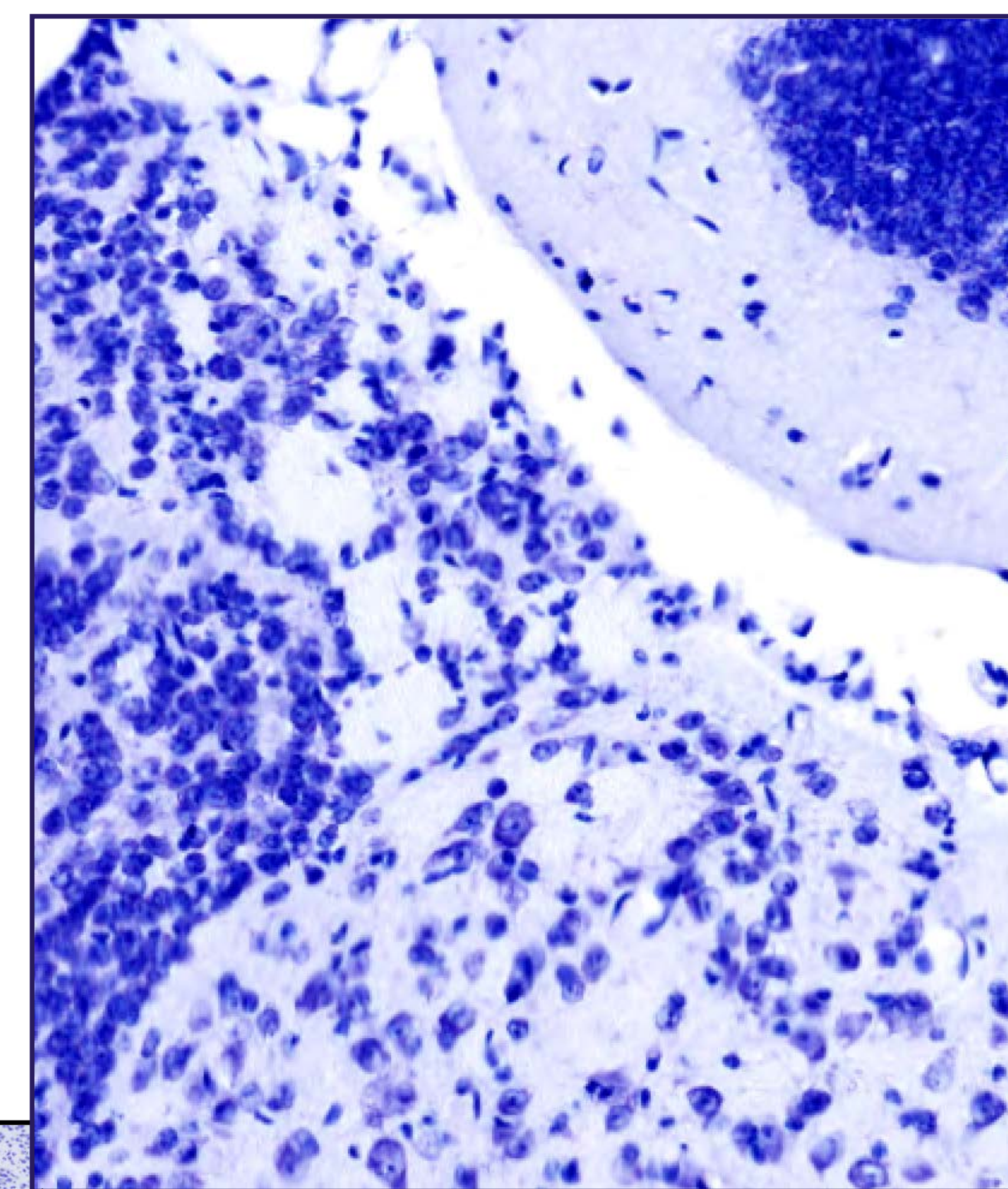
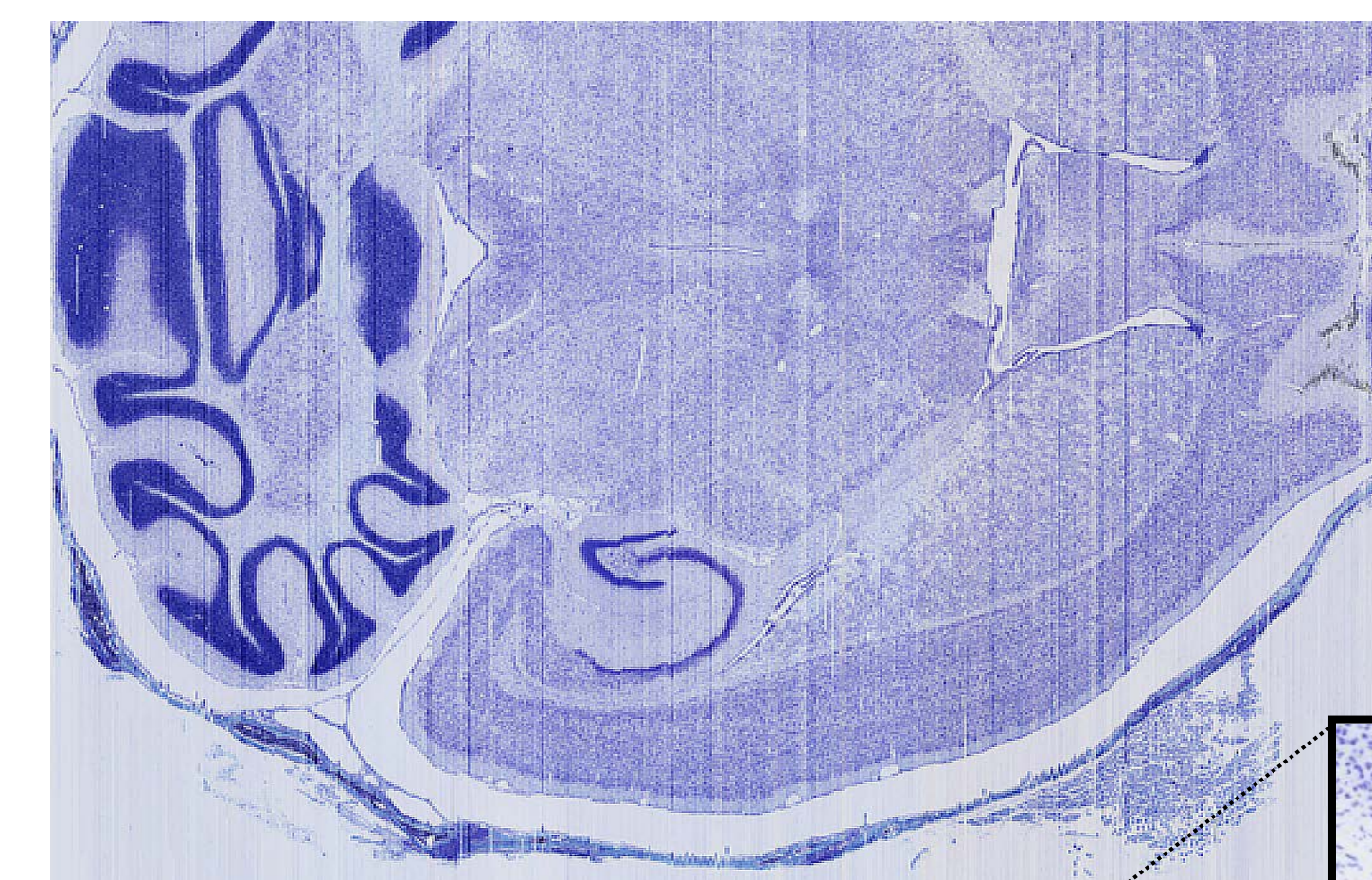
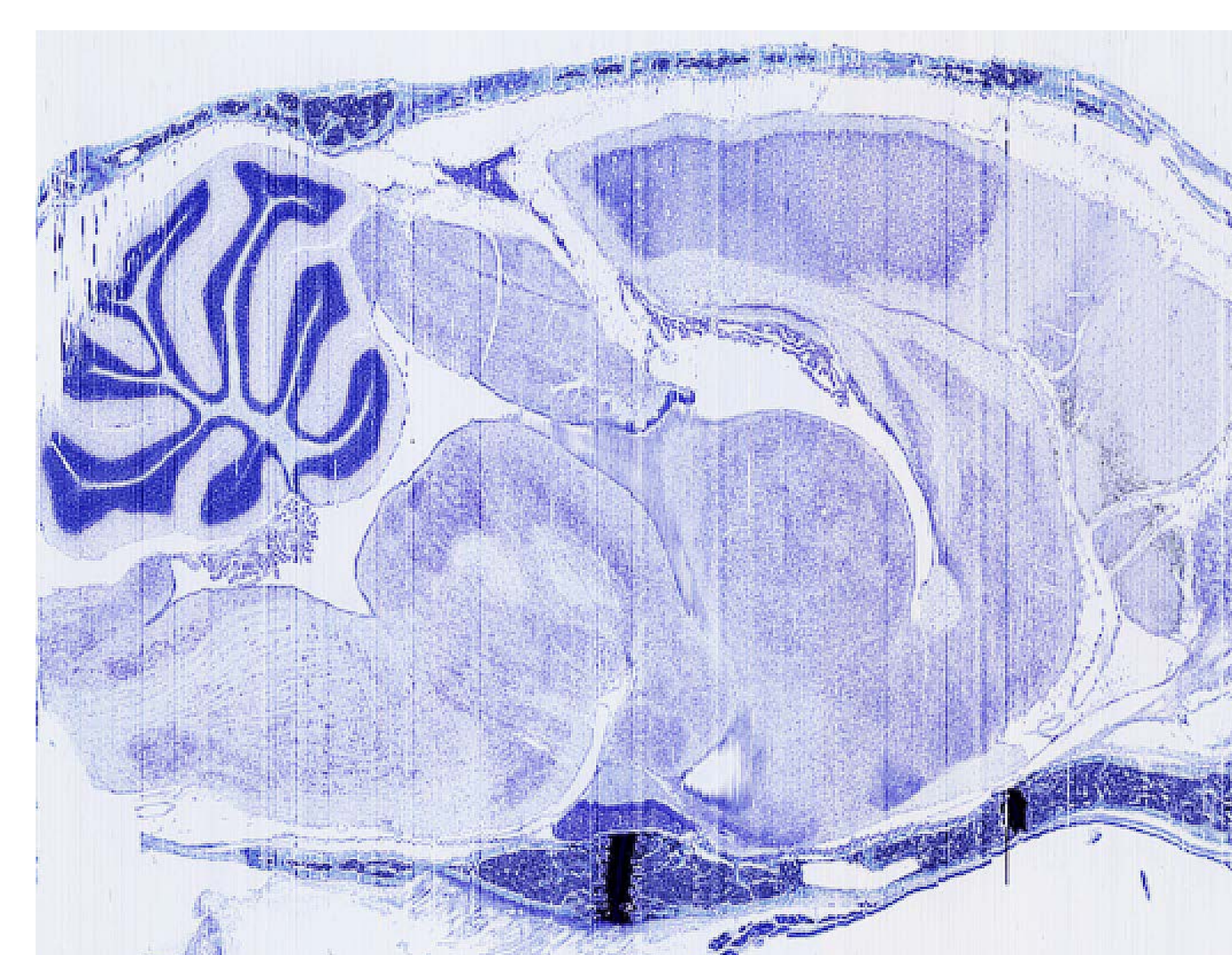
- brain inside the lightly decalcified skull cryo-sectioned (10  $\mu\text{m}$  thickness)
- sections (skull+brain) transferred to slide using the tape-transfer system
- sections were Nissl stained on slide.
- whole slides imaged using the NanoZoomer 2.0HT (0.46  $\mu\text{m}$ /pixel in plane)

## Nissl (in Skull) Data



### Image Stack

Sections: coronal (10  $\mu\text{m}$  thickness)  
Volume:  $\sim$ 1000 sections per brain  
pxl resolution: 0.46  $\mu\text{m}$



## MR-to-Nissl Co-Registration & Atlas Segmentation

### Data preprocessing

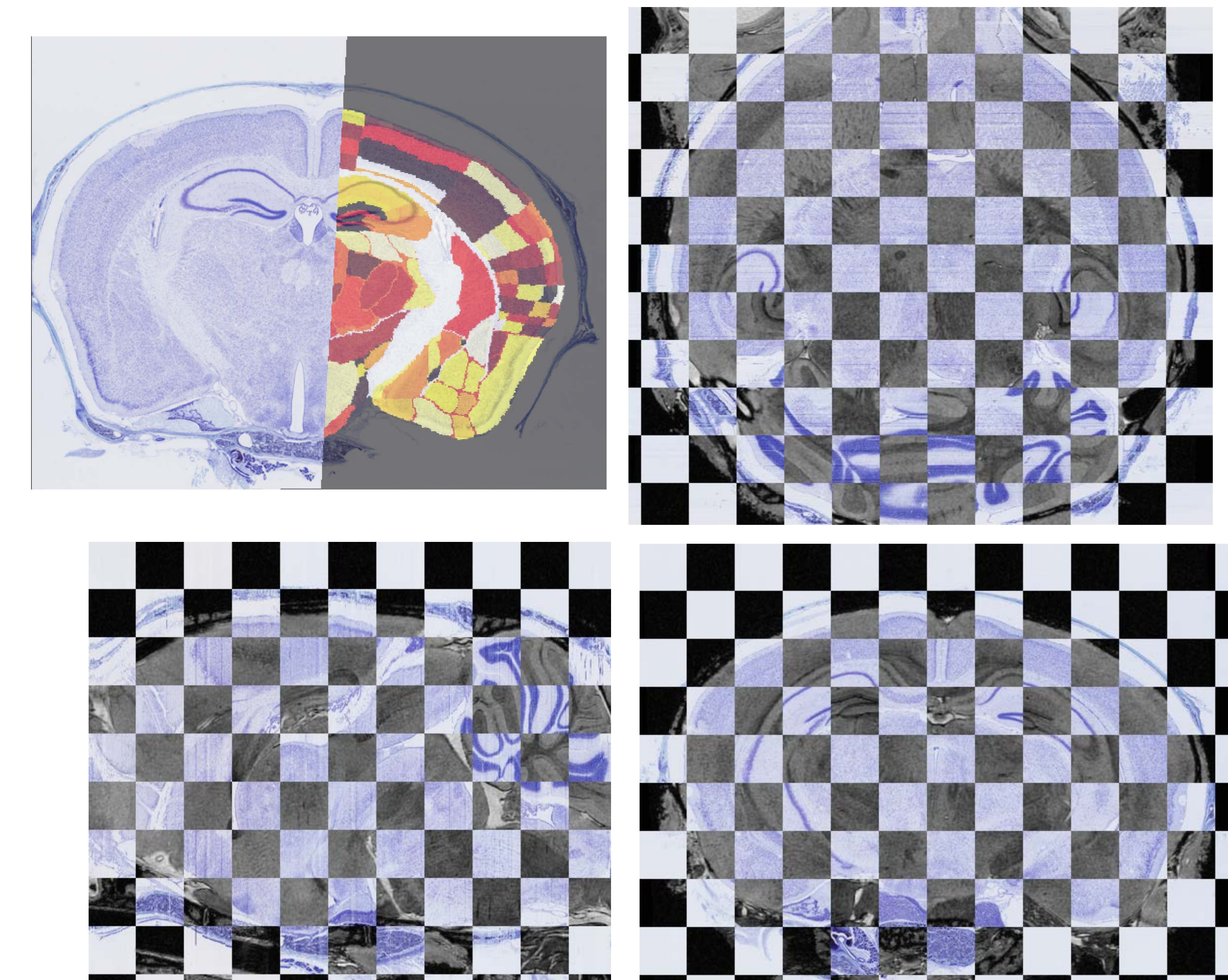
- Nissl image stack 2d-aligned (rigid transformation of adjacent sections)
- Nissl images subsampled to 20  $\mu\text{m}$  (to approximate MRI voxels size of 21.4  $\mu\text{m}$ )
- Nissl stack converted to inverted gray-scale (to compare with MR modality)
- MRI & Nissl stacks skull stripped by morphological operations

### Multimodal deformable registration [1]

- Step 1: Affine transformation
- Step 2: Free-form deformation estimated on a 3D grid
- 3D Grid-size: 240  $\mu\text{m}$  cube
- Resampling: cubic-splines
- Multi resolution: 5 levels
- Objective function: maximized normalized mutual information

### Atlas segmentation

- mouse brain reconstructed with high resolution from serial block-face tomography & annotated by co-registration with Allen Reference Atlas (P. Osten Lab; CSCHL)
- Annotated mouse brain morphed to MR-Nissl atlas via deformable registration



Reference  
[1] S. Klein, M. Staring, K. Murphy, M.A. Viergever, J.P.W. Pluim, "elastix: a toolbox for intensity based medical image registration," IEEE Transactions on Medical Imaging, vol. 29, no. 1, pp. 196 - 205, January 2010.

## Future Work

- Repeat procedure on multiple animals
- Deformable registration to be aided by salient anatomical landmarks
- Improve rostro-caudal resolution via optical re-slicing
- Assemble and publish high-resolution Nissl component of WHS

## Acknowledgements

We gratefully acknowledge funding support from the

- NIH (Directors Transformative Award MH087988, DA036400)
- NSF (Inspire Award PHY-1344069)
- Mathers Foundation
- Crick-Clay Professorship to Dr. Mitra
- CSHL internal funding.