

# Contrastive Learning of Cytoarchitecture using Spatial Information

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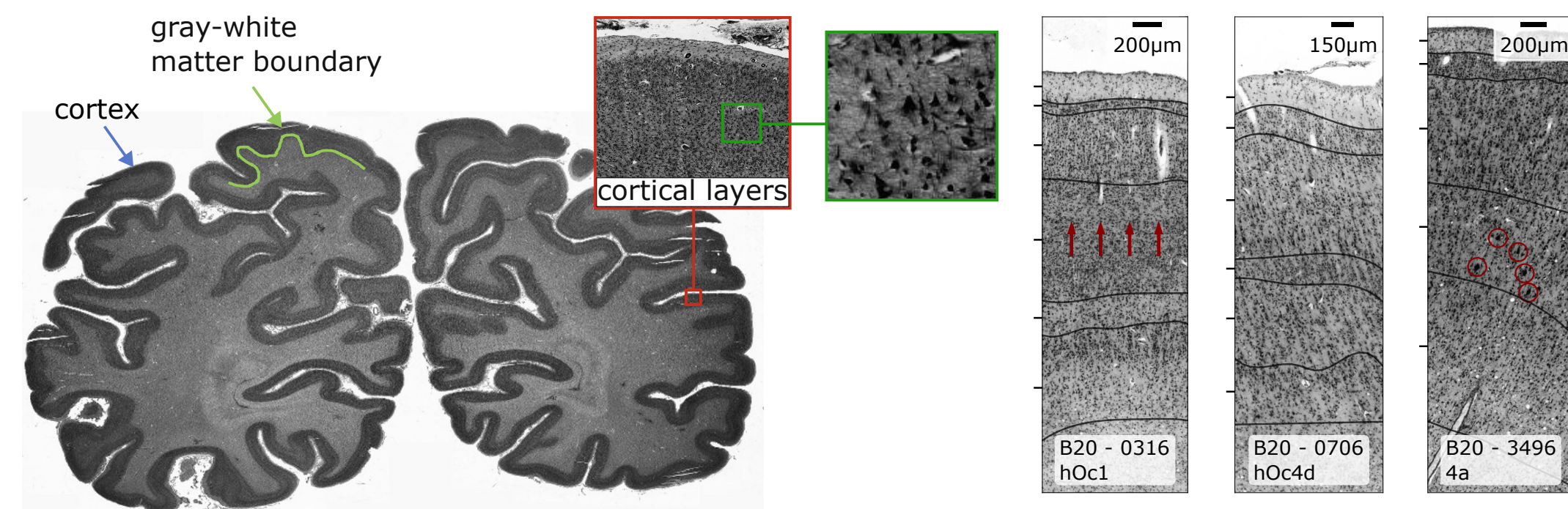
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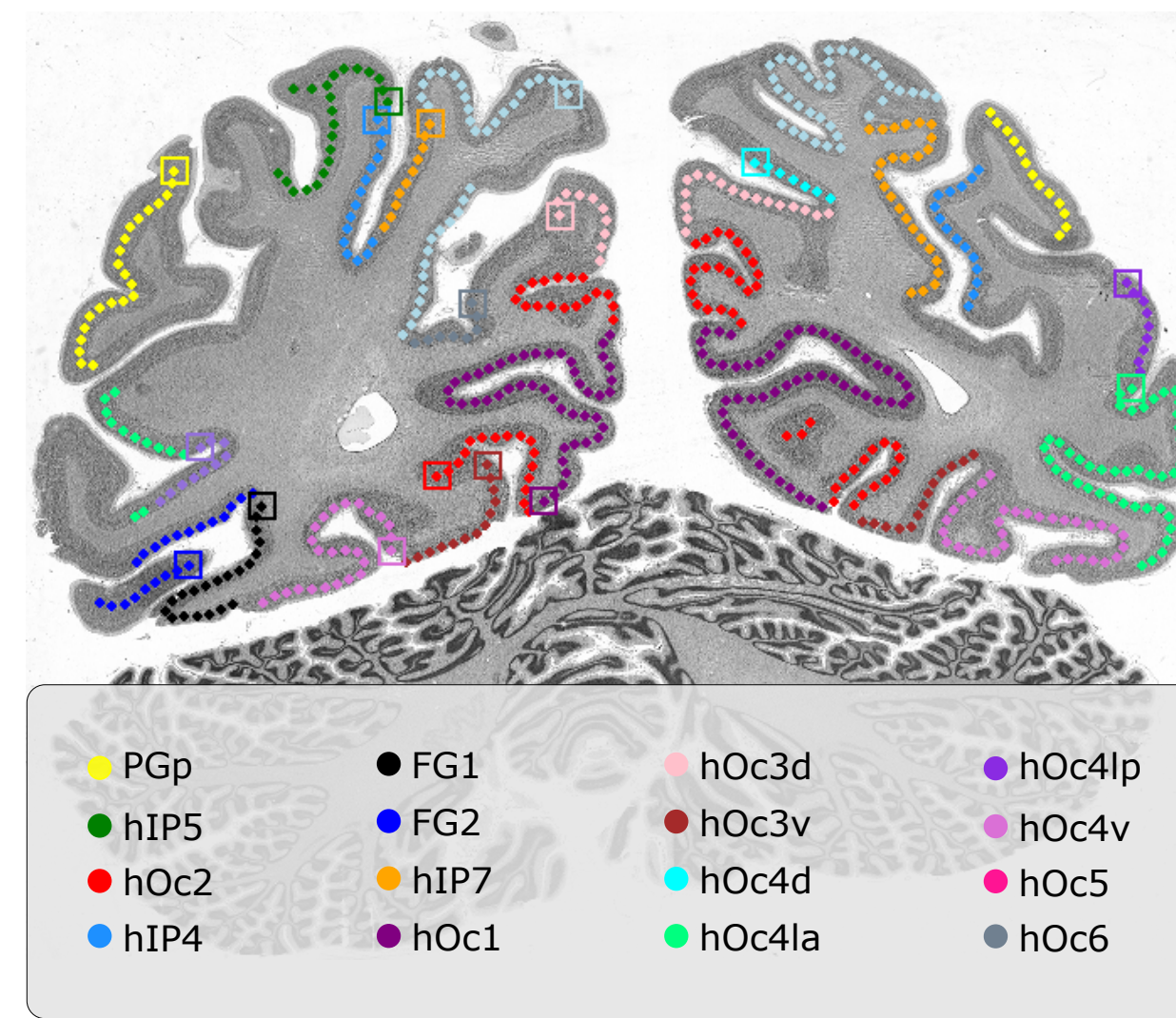
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## Mapping of Cytoarchitectonic Areas in Histological Sections



**Cytoarchitecture** can be analyzed in high-resolution microscopic scans of cell-body stained histological sections. It encompasses the distribution and size of neuronal cell bodies, their shape, orientation, type, as well as their columnar and laminar organization.



Postmortem human brains were obtained from the body donor programs of the anatomical institutes of the universities of Düsseldorf, Rostock and Aachen.

Histological processing of a postmortem brain [1]

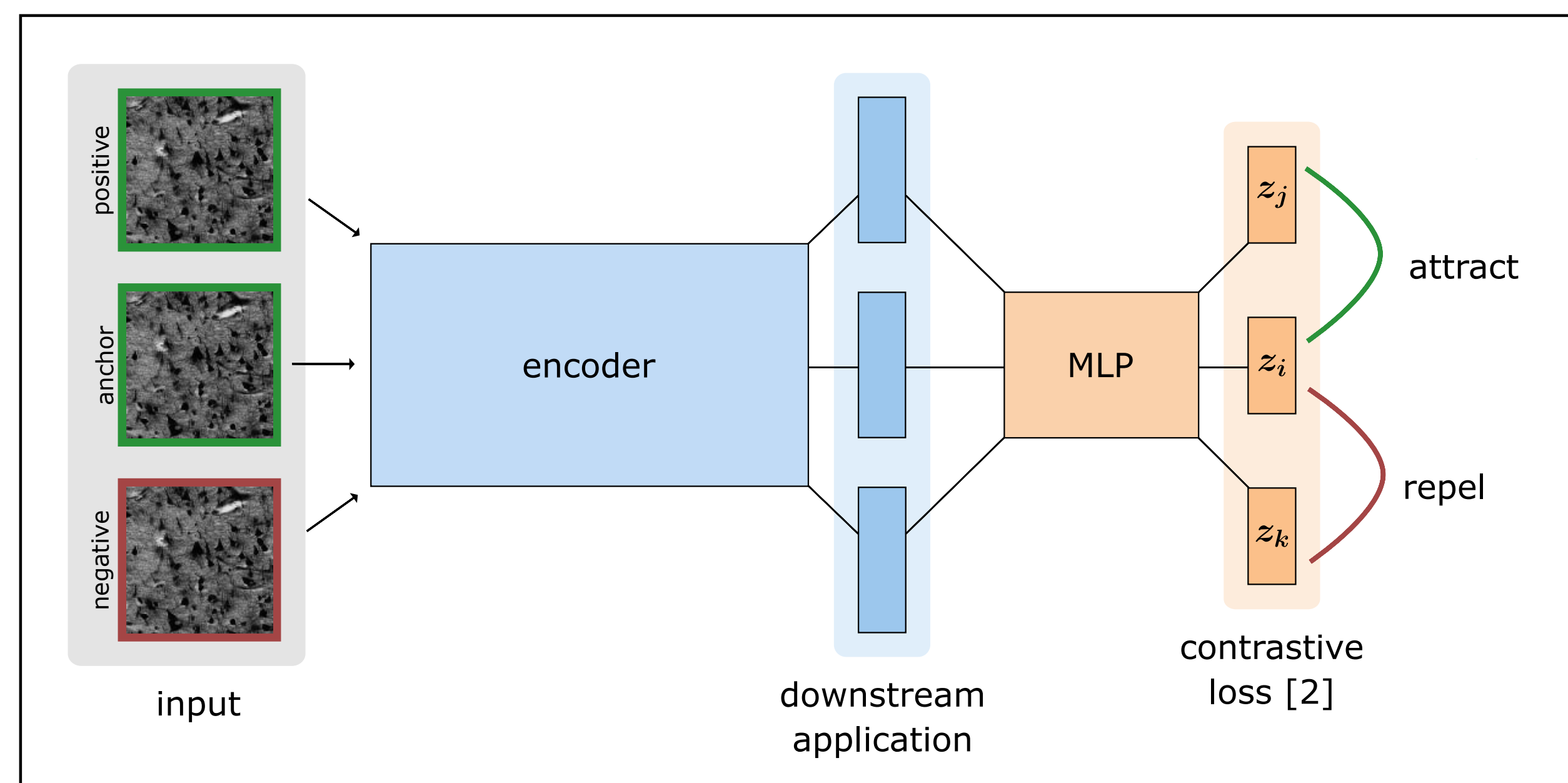
- Cutting brain into 6000-7500 thin histological slices
- Staining for cell bodies
- Scanning with high-throughput light microscopic scanners at a resolution of 1µm/px

### Aim: Cytoarchitectonic Brain Mapping

We train deep neural networks to predict brain areas from high-resolution image patches sampled along the midline of the cerebral cortex.

## Method: Distance-Based Representation Learning for Brain Mapping

### Representation Learning: Distance-Based Contrastive Learning



**Assumption:** Image patches sampled from *spatially close* locations are more similar than those extracted from distant locations.

**Prerequisite:** Registration of each brain in an anatomical reference space to obtain canonical spatial coordinates.

**Idea:** Learn latent space so that representations of similar inputs are closer than those of negative pairs

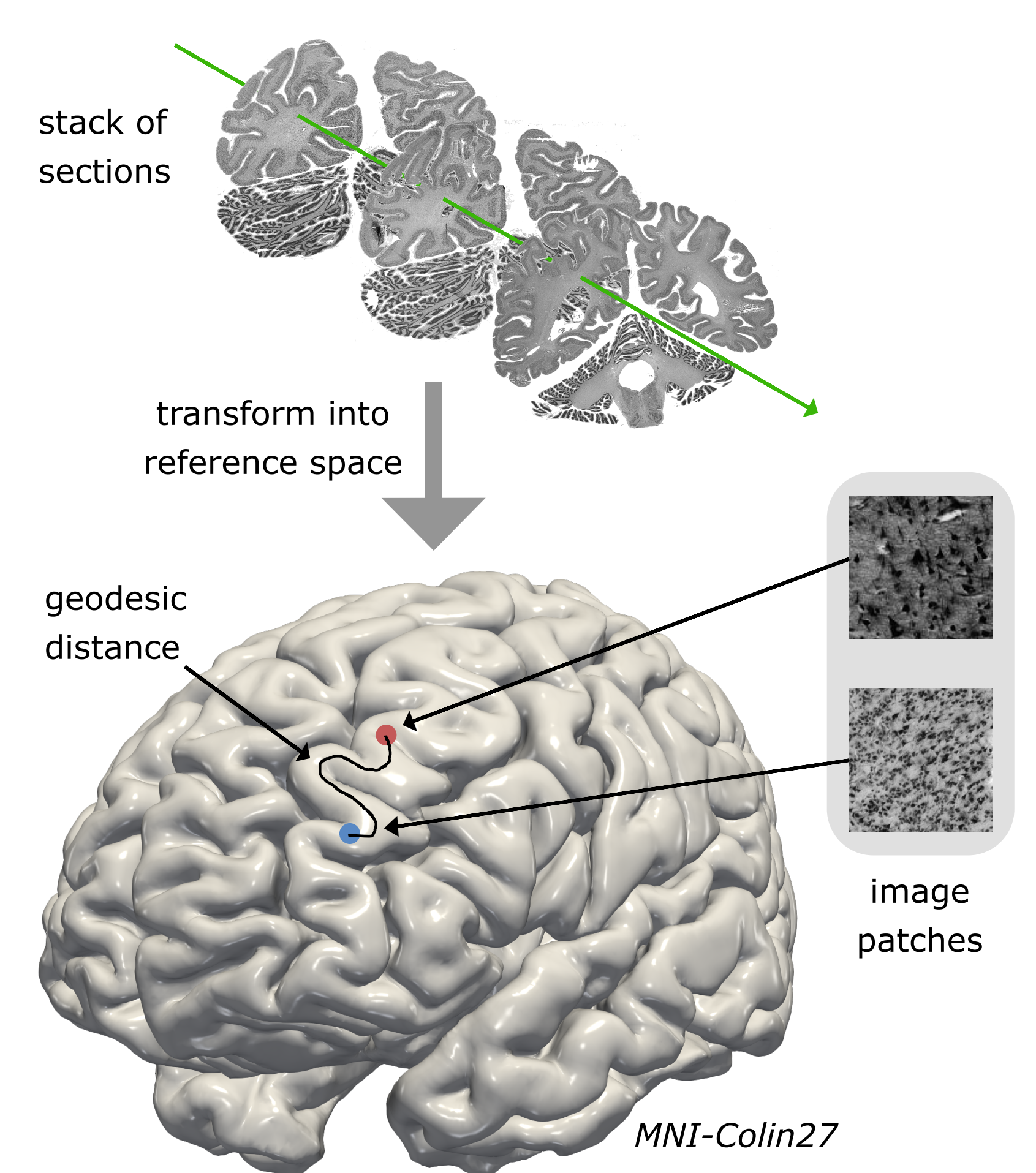
#### Geodesic distance:

Length of the shortest path along the surface of an object.  
→ similarity measure used to weight pairs of patches

#### Data sampling:

- Uniform sampling of coordinates within cerebral cortex
- Patches are centered around these coordinates
- Image patches of size 2048 x 2048 px at 2µm resolution

### Registration of Sections to MNI-Colin27



$$\ell_{dist}(i) = \frac{1}{\sum_{j=1}^b w_{ij}} \sum_{j=1}^b \mathbb{I}_{i \neq j} w_{ij} \log \frac{\mathbb{I} \exp(\text{sim}(z_i, z_j)/\tau)}{\sum_{k=1}^b \mathbb{I}_{k \neq i} \exp(\text{sim}(z_i, z_k)/\tau)}$$

Number of samples in a batch

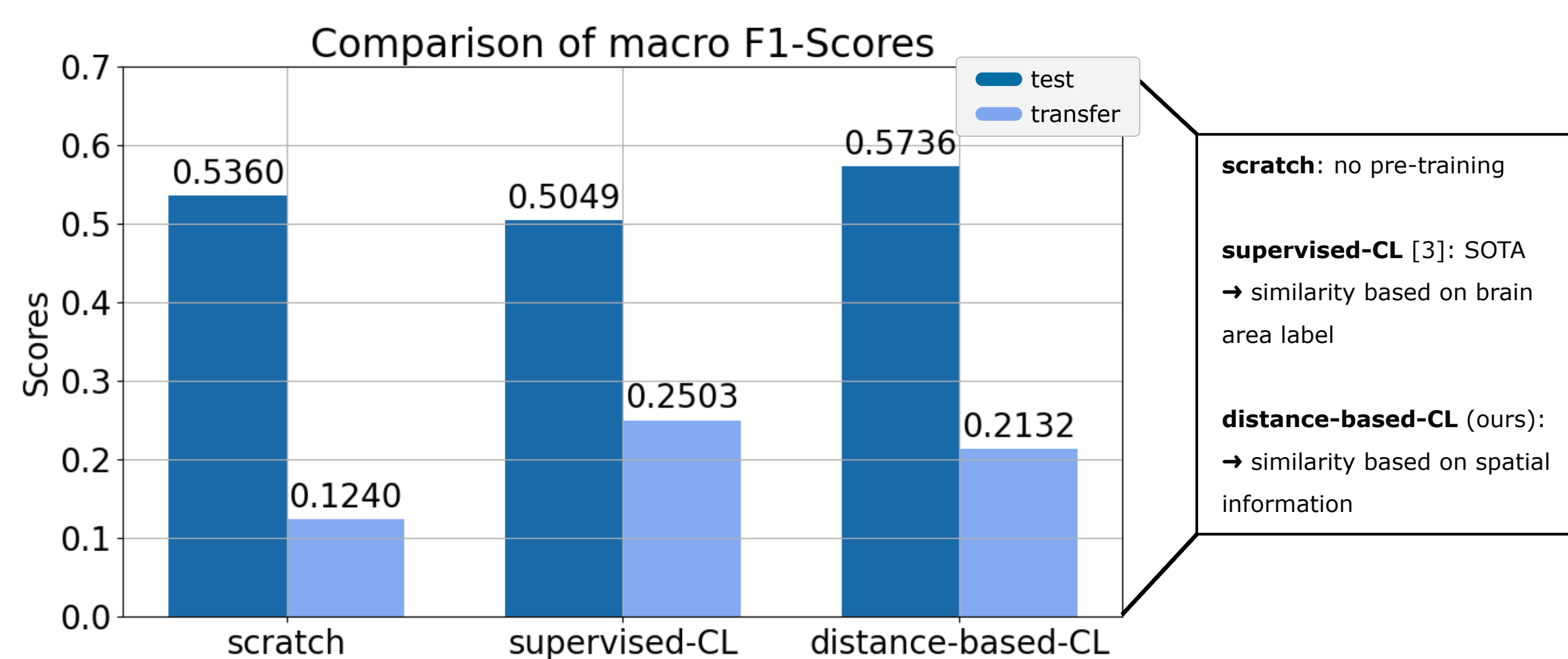
Cosine similarity

Sample in the batch at index  $i$

Weighting of pairs based on geodesic distance along the MNI-Colin27 brain

Feature vector computed by the neural network

## Results

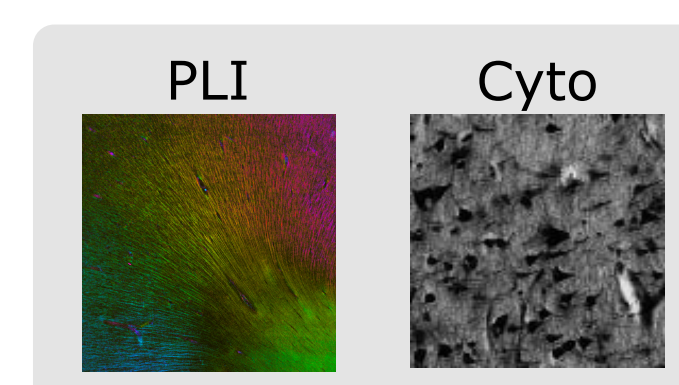


The proposed method enables learning of cytoarchitectonic features and outperforms the supervised contrastive learning method on the test dataset. However, generalization to unseen brains is a challenging task that remains unsolved.

Compared to the sota method [3], the proposed method does not use annotations for pre-training, has **lower inductive bias with respect to predefined areas**, and is thus applicable to a wider range of applications.

## Next Steps: Multi-Modal Learning

Training deep neural networks on image patches derived from various imaging techniques [1,6], with the goal of learning representations to **identify and extract both shared and distinct features**.



Sections derived from various imaging modalities contain shared and **complementary information**:

- Cell-body staining: cellular architecture
- Polarized light imaging [6]: connectivity and fiber tracts

#### Motivation:

- Spatial anchoring: attaching patches to specific locations in a 3D brain volume
- Cross-modality image generation: predict one image modality from the other
- Identifying common and shared information in both modalities

**Idea:** Our goal is to develop a CLIP-inspired [4] neural network architecture that integrates image patches from different modalities and learns meaningful representations from them. As our work has shown promising results, future work will further investigate the capabilities of learning from spatial information.

[1] K. Amunts, K. Zilles (2020) *Jülich-Brain: A 3D Probabilistic Atlas of the Human Brain's Cytoarchitecture*. Science, 369, no. 6506, 988, 2020.  
 [2] T. Chen (2020) *A Simple Framework for Contrastive Learning of Visual Representations*. International Conference on Machine Learning (ICML 2020), pp. 1597–1607.  
 [3] C. Schiffer (2021) *Contrastive Representation Learning For Whole Brain Cytoarchitectonic Mapping In Histological Human Brain Sections*. 2021 IEEE 18th International Symposium on Biomedical Imaging (ISBI 2021), pp. 603–606.  
 [4] Radford, A., Kim, J. W., Hallacy, C., Ramesh, A., Goh, G., Agarwal, S., Sastry, G., Askell, A., Mishkin, P., Clark, J., Krueger, G., & Sutskever, I. (2021) *Learning Transferable Visual Models From Natural Language Supervision*. arXiv preprint arXiv:2103.00020  
 [5] Kipf, T. N., & Welling, M. (2017) *Semi-Supervised Classification with Graph Convolutional Networks*. International Conference on Learning Representations (ICLR 2017).  
 [6] Axer, M., Graessel, D., Kleiner, M., Dammers, J., Dickscheid, T., Reckfort, J., Huetz, T., Eiben, B., Pietrzyk, U., Zilles, K., Amunts, K., 2011. *High-Resolution Fiber Tract Reconstruction in the Human Brain by Means of Three-Dimensional Polarized Light Imaging*. Frontiers in Neuroinformatics 5.

This project received funding from the European Union's Horizon 2020 Research and Innovation Programme, grant agreement 101147319 (EBRAINS 2.0 Project), and the Helmholtz Association port-folio theme "Supercomputing and Modeling for the Human Brain", and the Helmholtz Association's Initiative and Networking Fund through the Helmholtz International BigBrain Analytics and Learning Laboratory (HIBALL) under the Helmholtz International Lab grant agreement InterLabs-0015, and by HELMHOLTZ IMAGING, a platform of the Helmholtz Information & Data Science Incubator [X-BRAIN, grant number: ZT-I-PF-4-061].  
 Computing time was granted through JARA on the supercomputer JURECA-DC at Jülich Supercomputing Centre (JSC).