

## YOUNG INVESTIGATOR AWARD

# Optoacoustic imaging of the glymphatic system

Sarah Shaykevich,<sup>1,2,3</sup> Russell W. Chan,<sup>1,2,4</sup> Chandni Rana,<sup>1,2</sup>

Mohamed Eltaeb,<sup>1,2</sup> Justin P. Little,<sup>1,2</sup> Daniel Razansky,<sup>5,6</sup> Kevin C. Chan,<sup>1,3,4</sup> Shy Shoham,<sup>1,2,4</sup>

<sup>1</sup>Neuroscience Institute, <sup>2</sup>Tech4Health Institute, <sup>3</sup>Department of Radiology, <sup>4</sup>Department of Ophthalmology, NYU Grossman School of Medicine, New York, USA; <sup>5</sup>Institute for Biomedical Engineering, <sup>6</sup>Institute of Pharmacology and Toxicology, University of Zurich and ETH Zurich, Zurich, Switzerland

### Background

The glymphatic system is a brain waste clearance system mediated via cerebrospinal fluid (CSF) flow,<sup>1</sup> with implications for influence on neurodegenerative diseases.<sup>2</sup> Most preclinical glymphatic studies employ fluorescence imaging, which provides higher specificity, but a smaller field-of-view (FOV), or magnetic resonance imaging (MRI), which provides brain-wide FOV, but lower specificity. Functional optoacoustic neuro-tomography<sup>3</sup> (FONT) offers a larger FOV compared to classical optical methods, and higher specificity compared to MRI. However, FONT has not yet been applied to probe the glymphatic system. In this study, we used fluorescence and optoacoustic imaging of a near-infrared dye, Janelia Fluor 669 (JF669), to track CSF and multimodal CSFhemodynamic flows in mice.

## Methods

After observing strong fluorescence and optoacoustic signatures of JF669 in phantom experiments, we performed a series of *in vivo* experiments in isoflurane-anesthetized C57BL/6 mice (n=3 fluorescence and n=4 FONT experiments, respectively) (Figure 1A). Lumbar injection was applied to deliver JF669 at a rate of 2  $\mu$ L/min for 30 minutes. A polyethylene tube was placed intrathecally at the lumbar region (L4-L5). The scalp was removed. Fluorescence or FONT images were obtained every 5 minutes after injection.

## Results

## Fluorescence imaging and FONT probe CSF flow

Images (Figure 1B) and time traces (Figure 1C) revealed time-dependent anatomical routes of paravascular influx, including the transport along the olfactory artery (OFA), superior cerebellar artery (SCA), and bilateral middle cerebral artery (MCA). For FONT imaging, since the OFA

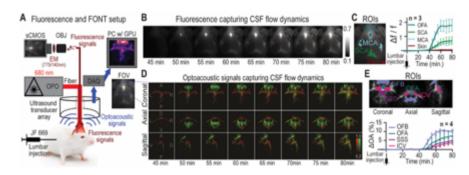


Figure 1. Fluorescence imaging and functional optoacoustic neuro-tomography (FONT) visualize and characterize anatomical routes of the glymphatic system. (A) Experimental setup. (B) Fluorescence images captured CSF flow dynamics. (C) ROI-based fluorescence traces revealed time-dependent routes of paravascular influx. (D) Maximum intensity projections of FONT volumes captured static blood vessels and CSF flow dynamics. Blood (red) and JF669 (green) areas were determined by unmixing multispectral data. (E) ROI-based FONT traces revealed time-dependent routes of paravascular influx, CSF-ISF exchange and paravascular efflux.

Correspondence: Sarah Shaykevich, 433 1st Ave, Fl 10 Rm 1026, New York, 10010, USA. Tel. (570) 730 6102. E-mail: sarah.shaykevich@nyulangone.org

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showed strong fluorescence (Figure 1B), we positioned the ultrasound transducer array at the anterior of the mouse brain with a FOV of 5x5 mm<sup>2</sup> (Figure 1A). Standard filtered back-projection reconstruction was applied. Besides the OFA route of the paravascular influx, the dynamic images (Figure 1D) and time-traces (Figure 1E) also revealed time-dependent anatomical routes of CSF-interstitial fluid (ISF) exchange in the olfactory bulb (OFB) and paravascular efflux in the superior sagittal sinus (SSS) and the bilateral inferior cerebral vein (ICV).

Next, we studied the aquaporin-4 (AQP4) dependence of glymphatic flow by subcutaneously injecting AQP4 inhibitor TGN020, in addition to the prior procedures. Under fluorescence imaging and FONT, we observed that TGN020 significantly decreased and spatially restricted the spread of JF669 in the brain.

## FONT spectral unmixing separates CSF and blood

We swept the OPO through 680 nm to 750 nm, with 10 nm steps at 10 Hz in the phantom and in each animal. The multispectral reconstructions were unmixed using the known absorption spectra of hemoglobin and the JF669 OA spectrum obtained from the phantom.<sup>4</sup> This enabled the separation of blood and JF669 signal (Figure 1D).

### Conclusions

We characterized anatomical routes of



paravascular influx (OFA), CSF-ISF exchange (OFB) and paravascular efflux using FONT, while fluorescence imaging only enabled visualization of the paravascular influx. We thus demonstrated the feasibility and advantages of FONT for probing the glymphatic system. These findings present great opportunities for monitoring and understanding the glymphatic system during development, aging, disease, pharmacological interventions and genetic modifications.

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