

# ADVANCED HEALTHCARE MATERIALS

## Supporting Information

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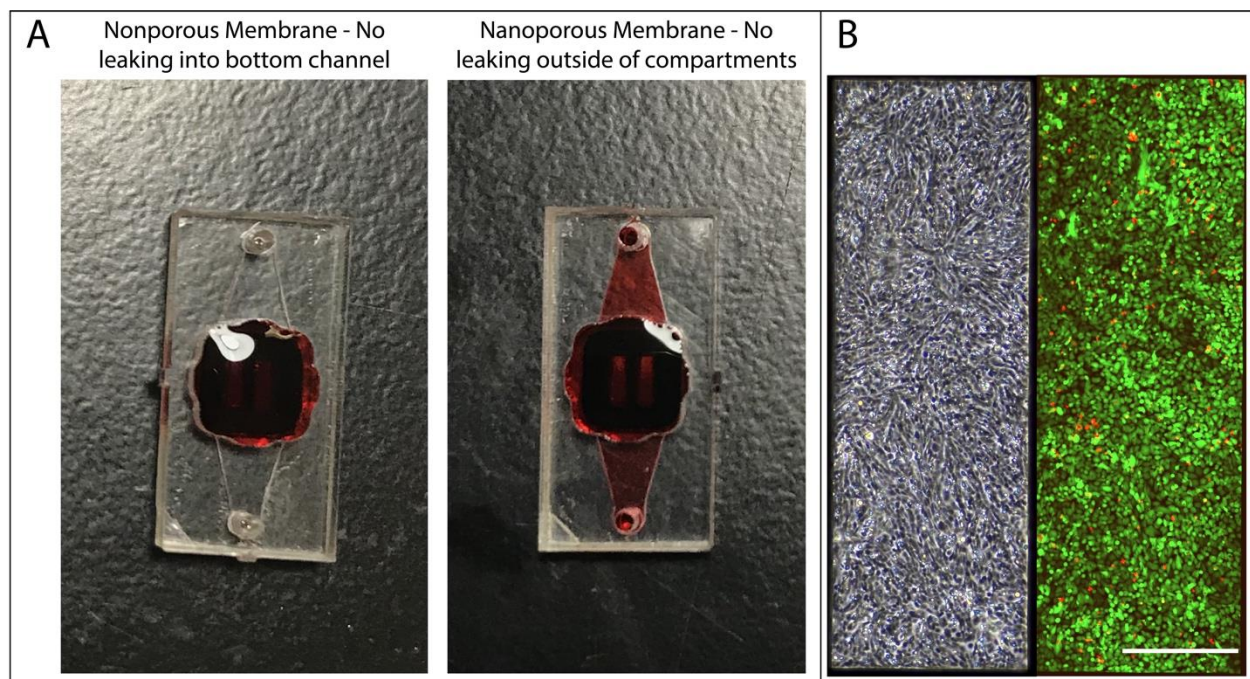
The Modular  $\mu$ SiM: A Mass Produced, Rapidly Assembled, and Reconfigurable Platform for the Study of Barrier Tissue Models In Vitro

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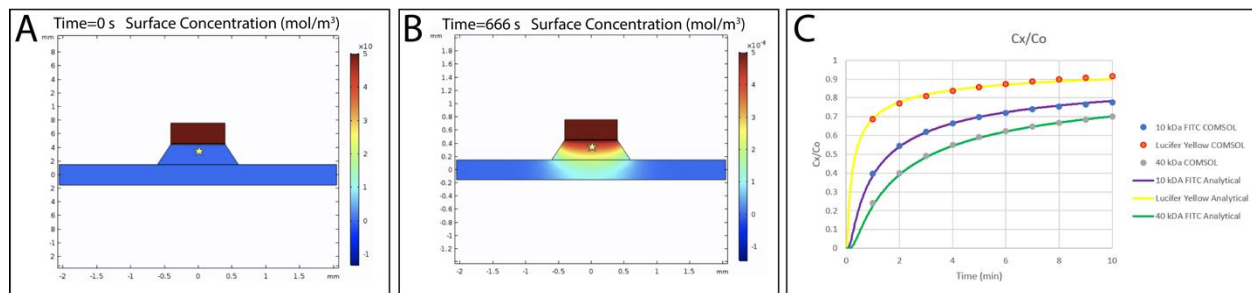
**Title:** The Modular  $\mu$ SiM: a Mass Produced, Rapidly Assembled, and Reconfigurable Platform for the Study of Barrier Tissue Models *In Vitro*

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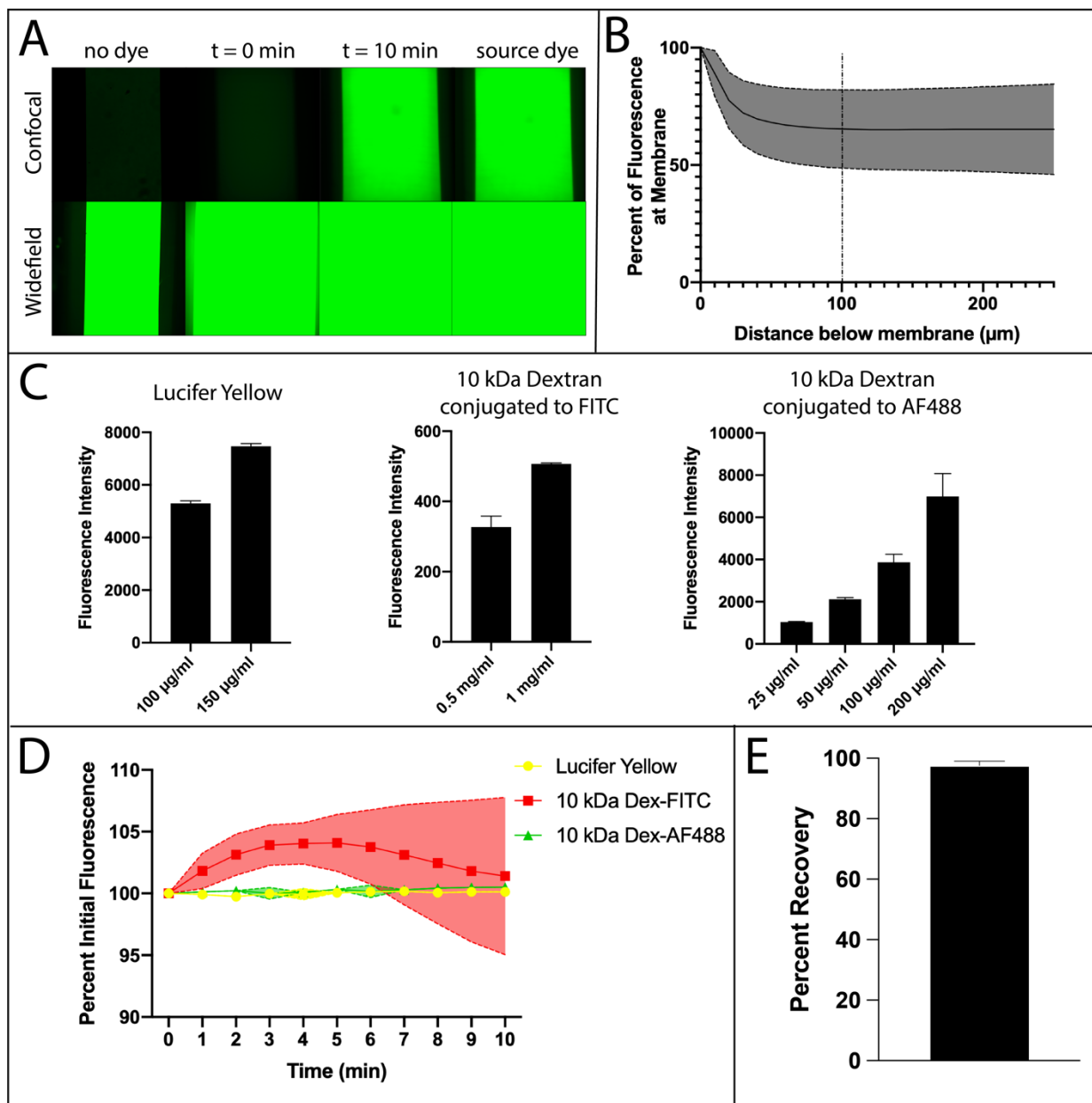


**Figure S1. m- $\mu$ SiM assembly and cell culturing validation.** (A) Representative images of assembled m- $\mu$ SiM devices used for a leak test. Dye was added into Component 1's top well, and devices assembled with non-porous membranes (left) retained dye in the top well, whereas devices assembled with nanoporous membranes (right) showed diffusion into the channel after 2 hours. There was no leaking within the device. N = 4-5 devices per group. (B) Representative images of hCMEC/D3 growth in m- $\mu$ SiM. Cells were cultured for 5 days. A brightfield image was taken of the monolayer (left) and a LIVE/DEAD stain was performed (right), with  $98.2 \pm$

1.2% cell viability. Green indicates live cells, and red indicates dead cells.  $N = 4$ . Scalebar = 300  $\mu\text{m}$ .

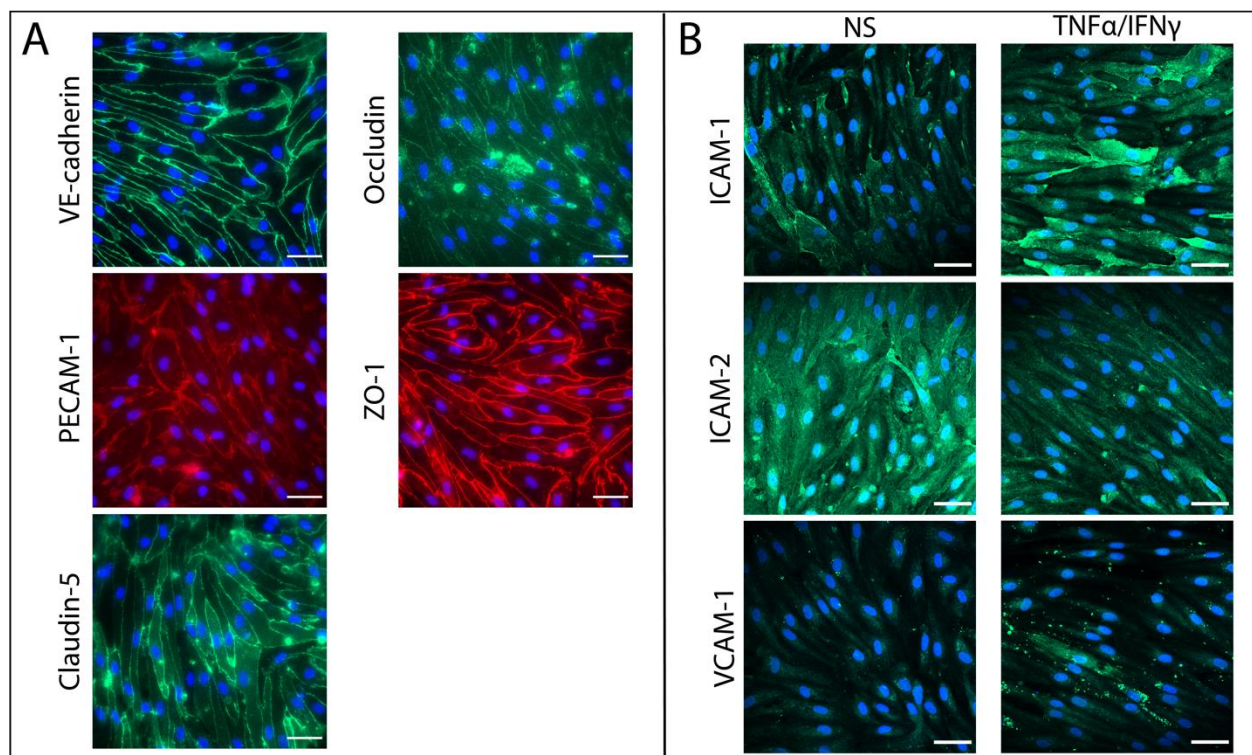


**Figure S2. Comparison between 1-D analytical model of diffusion and 2-D finite element model.** (A-B) COMSOL geometry representing membrane chip with trench, bottom chamber, and a ‘well’ of dye above the membrane at times zero (A) and 666 seconds (B). Concentration was measured at the center of the trench at a distance 100  $\mu\text{m}$  below the source bottom (yellow star). (C) Concentrations over time in the center of the trench at a position of 100  $\mu\text{m}$  below the membrane ( $C_x$ ) normalized to source ‘well’ concentration ( $C_0$ ). Agreement between the analytical (line) and computational (dot) models for free diffusion (Equation 1 in main text) for all molecular tracers indicates that a 1-D analytical solution can be used with the  $\mu\text{SiM}$  without error. Molecular tracers modeled were: 1) 10 kDa Dextran conjugated to FITC (purple line, blue dots), 2) lucifer yellow (yellow line, orange dots), and 3) 40 kDa Dextran conjugated to FITC (green line, gray dots).





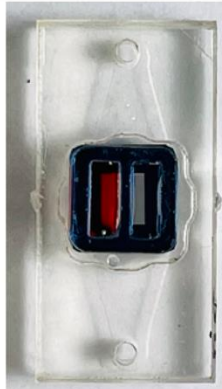
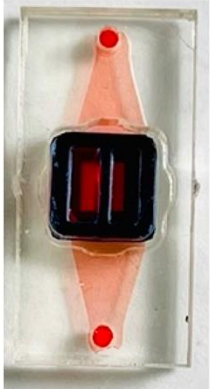

**Figure S3. *In situ* and sampling-based small molecule permeability assay optimizations.** (A) Demonstration that confocal microscopy is necessary for *in situ* permeability assay. Images are acquired on the same microscope (Andor Spinning Disc Confocal Microscope) in confocal and widefield modalities. Background fluorescence in widefield overwhelms the field of view, whereas in confocal, background is minimal and can be subtracted. (B) Optimization of confocal plane for *in situ* assay. Dye is added into the top well of a nonporous device and fluorescence intensity is measured starting at the membrane and shifting the objective down. Distance below membrane refers to distance the objective moves. At approximately 100  $\mu\text{m}$ , background fluorescence from dye in the well is minimized.  $N = 5$  devices. (C) Optimization of dye

concentration for three dyes of interest in *in situ* assay. The concentration selected for each is within the linear range of fluorescence intensities to appropriately assume correlation between concentration and fluorescence. (D) Photo bleach test for three dyes of interest for *in situ* assay. 10-25% of the optimized source dye concentration is added into the channel of a nonporous device. Fluorescence intensity is measured once every minute for ten minutes. Fluorescence intensity does not diminish over the course of the experiment. (E) Experimentally tested percent recovery for sampling assay. Dye was added into the top well of uncoated nanoporous devices and allowed to diffuse for one hour. After one hour, dye was removed from the channel using a 50  $\mu\text{L}$  reservoir. A second 50  $\mu\text{L}$  flush was done to remove remaining dye. A fluorescent plate reader was used to measure fluorescence intensity and percent recovery was calculated as concentration in flush one divided by the sum of the concentration from both flushes.  $N = 10$ .

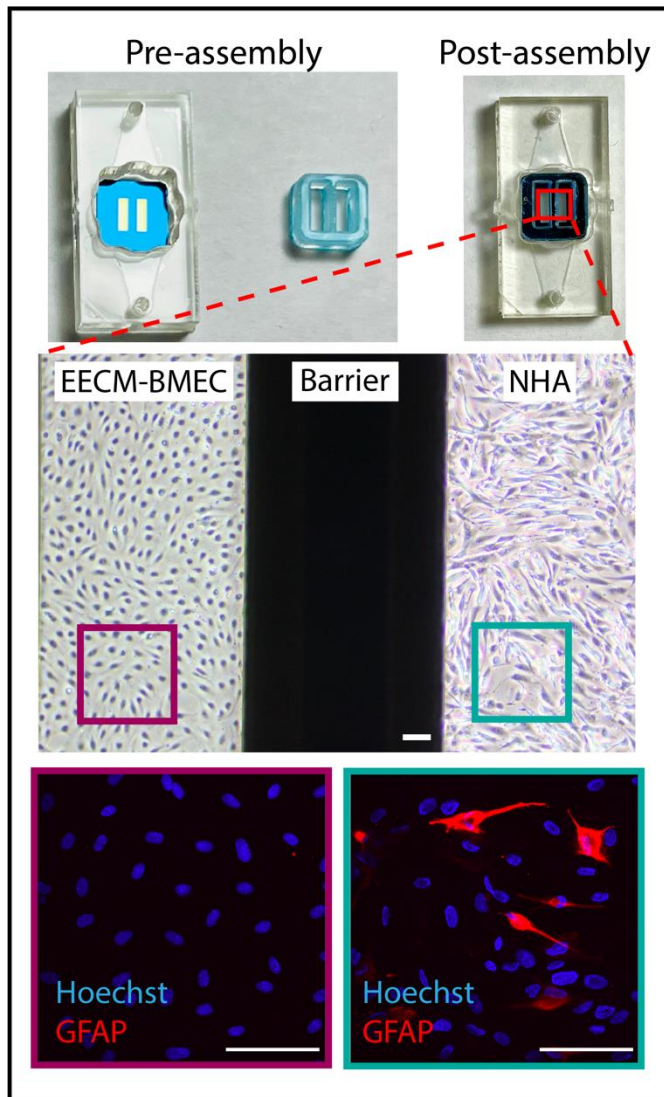


**Figure S4. Reproducibility of EECM-BMEC-like cell differentiation and culture on m- $\mu\text{SiM}$  at UR.** (A) EECM-BMEC-like cells differentiated at UR and cultured on the m- $\mu\text{SiM}$  express key molecules of junctional complexes, similar to those produced by UniBe (see Main Text). (B) EECM-BMEC-like cells express key cell adhesion molecules upon exposure to proinflammatory stimuli (0.1 ng/ml TNF $\alpha$  + 2 IU/ml IFN $\gamma$ ) when differentiated at UR and

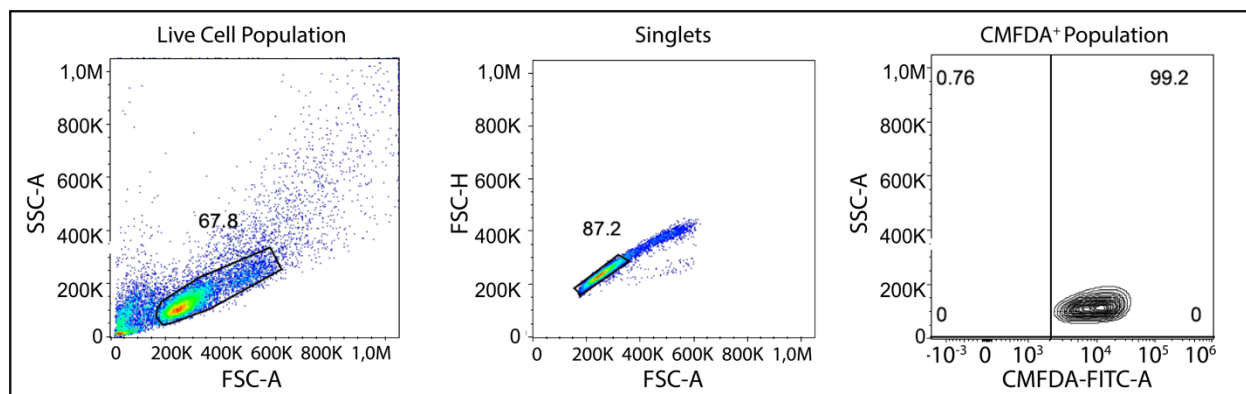
cultured on m- $\mu$ SiM devices. Images were acquired on an Andor Spinning Disc Confocal Microscope using a long-working distance 40X objective. Scalebar = 50  $\mu$ m.

Condition 1	Condition 2	Condition 3	Condition 4	Condition 4
Backside sealed chip	Backside sealed chip	No seal	No seal	Post-diffusion: Channel flushed with water
Air on backside Water other chamber	Water on backside and other chamber	Air on backside Water other chamber	Water on backside and other chamber	
				
No leaking into other chamber	No leaking into other chamber	No leaking into other chamber	Dye diffuses through channel	Dye diffused to other chamber

**Figure S5. Two-slot NPN insert dye leak validation.** Representative images of m- $\mu$ SiM with 2-slot culture chamber insert used for a dye leak tests. Devices with the backside sealed retained dye in one chamber, with no leaking into the second chamber (Conditions 1 and 2). Devices with the backside unsealed but air in the channel (Condition 3) retained dye one chamber. When dye is added to one chamber with water in the channel and other chamber (Condition 4), dye diffuses through the channel and into the second chamber. After flushing the channel with water (Condition 4, Post-diffusion), it is clear dye does diffuse into the second chamber. This indicates communication between chambers occurs solely through the channel and not between compartments. N = 4-5 devices each group.



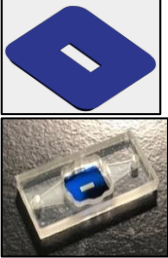
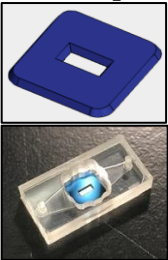
**Figure S6. Two-slot NPN insert for immunofluorescence staining of astrocytes.** EECM-BMEC-like cells were cultured in one chamber and human astrocytes (NHA) in the other chamber. Cells were stained for astrocyte marker (GFAP) and nuclear marker Hoechst (blue). Images show there is no cross-contamination of cells between culture chambers. It should be noted that GFAP is a marker of reactive astrocytes, so while it is clear through phase contrast imaging that astrocytes are growing in their respective chamber, only through staining can we see that only a handful of astrocytes are expressing GFAP in co-culture. Phase images were acquired on a Nikon Eclipse Ts2 phase contrast microscope and fluorescence images were acquired on an Andor Spinning Disc Confocal Microscope using a long-working distance 40X objective. Scalebar = 100  $\mu\text{m}$ .



**Figure S7. Flow cytometry gating strategy for CMFDA-labelled T-cells.** Plots are initially gated to remove dead cell debris (left), then gated to remove doublets (middle). Final gate includes only CMFDA<sup>+</sup> T-cells (right). Plots above are representative gating using our input T-cell population.



**Table S1. Microscopy objectives and working distances for different chip configurations.**

Chip Orientation	Distance from bottom of $\mu$ SiM to Membrane [ $\mu$ m]	Reasons to Select Orientation	Objectives Used in Orientation	Numerical Aperture (NA)	Working Distance (WD) [mm]
<b>Trench-down</b> 	615	Endothelial cells cultured in Component 1's well grow on flat surface	4X: Nikon, dry, Model MRH20041	0.13	16.4
			10X: Nikon, air, Model MRD00105	0.45	4
			10X: Nikon, air, Model MRH00105	0.30	16.0
			LWD 40X: Model, water, Part MRD77410	1.15	0.61–0.59
			LWD 40X: Nikon, air, Model MRP46402	0.55	2.1
<b>Trench-up</b> 	305	Shorter working distance to cells and endothelial cells cultured bottom channel grow on flat surface	20X: Nikon, multi-immersion, Model MRH07241	0.75	0.51–0.33
			LWD 40X: Nikon, water, Model MRD77410	1.15	0.61–0.59
			60X: Nikon, water, Model MRD07602	1.2	0.31–0.28
			60X: Olympus UPlanSApo, water, Model N1480800	1.2	0.28

**Table S2. Antibody list for immunofluorescence staining at UniBe and UR**

Antibodies	Fixative	Clone	Source	Cat. N.	Secondary Antibody
<b>VE-cadherin</b>	MeOH	UniBe: F-8	UniBe: Santa Cruz	UniBe: sc-9989	UniBe: Cy3 AffiniPure F(ab') <sub>2</sub> Fragment Goat Anti-Mouse IgG
		UR: 123413	UR: R&D Systems	UR: MAB9381	UR: Goat Anti-Mouse IgG (H+L) Cross-Adsorbed, Alexa Fluor 488
<b>PECAM-1</b>	MeOH	UniBe: MEM-05	UniBe: Invitrogen	UniBe: 37-0700	UniBe: Cy3 AffiniPure F(ab') <sub>2</sub> Fragment Goat Anti-Mouse IgG
			UR: Invitrogen		

		UR: polyclonal		UR: PA5-32321	UR: Goat Anti-Rabbit IgG (H+L) Cross-Adsorbed, Alexa Fluor 568
<b>Claudin-5</b>	MeOH	4C3C2	Invitrogen	35-2500	UniBe: Cy3 AffiniPure F(ab') <sub>2</sub> Fragment Goat Anti-Mouse IgG  UR: Goat Anti-Mouse IgG (H+L) Cross-Adsorbed, Alexa Fluor 488
<b>Occludin</b>	MeOH	OC-3F10	Invitrogen	33-1500	UniBe: Cy3 AffiniPure F(ab') <sub>2</sub> Fragment Goat Anti-Mouse IgG  UR: Goat Anti-Mouse IgG (H+L) Cross-Adsorbed, Alexa Fluor 488
<b>ZO-1</b>	MeOH	polyclonal	Invitrogen	40-2200	UniBe: Cy3 AffiniPure F(ab') <sub>2</sub> Fragment Donkey Anti-Rabbit IgG  UR: Goat Anti-Rabbit IgG (H+L) Cross-Adsorbed, Alexa Fluor 568
<b>ICAM-1</b>	live	HA58	Biolegend	353102	UniBe: Goat Anti-Mouse IgG (H+L) Highly Cross-Adsorbed, Alexa Fluor 488  UR: Goat Anti-Mouse IgG (H+L) Cross-Adsorbed, Alexa Fluor 488
<b>ICAM-2</b>	live	CBR-IC2/2	FITZGERALD	10R-7606	UniBe: Goat Anti-Mouse IgG (H+L) Highly Cross-Adsorbed, Alexa Fluor 488  UR: Goat Anti-Mouse IgG (H+L) Cross-Adsorbed, Alexa Fluor 488
<b>VCAM-1</b>	live	51-10C9	BD Biosciences	555645	UniBe: Goat Anti-Mouse IgG (H+L)

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					Highly Cross-Adsorbed, Alexa Fluor 488
					UR: Goat Anti-Mouse IgG (H+L) Cross-Adsorbed, Alexa Fluor 488
<b>GFAP</b>	MeOH	EP672Y	abcam	ab33922	UR: Goat Anti-Rabbit IgG (H+L) Cross-Adsorbed, Alexa Fluor 568

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