

Supplementary methods

Analysis of FOXC2 expression by real-time quantitative reverse transcription polymerase chain reaction (qRT-PCR)

Total RNA was extracted from LECs cultured on-chip and LECs cultured on a T75 flask (Thermo Fisher Scientific, Waltham, MA, USA) using TRIzol (Invitrogen, Waltham, MA, USA). At the end of the 4-day culture period, syringes were removed from the Luer connectors and culture medium was aspirated from the chips. To induce cell lysis, 100 µL of TRIzol were added to each Luer and 50 µL were added to the LEC channel (outlet) for each sample in the chip (3 per idenTx 3 device). Chips were incubated for 40 min on ice after which TRIzol was pipetted up-down before sample collection. Four samples were pooled to ensure sufficient RNA yield. For flask culture, at the end of the 4-day culture period cells were detached with TrypLE (Gibco, Life Technologies, Carlsbad, CA, USA) and 1 mL TRIzol was added to 0.5×10^6 cells for RNA isolation. RNA was extracted following manufacturer's instructions using 10 µg of glycogen (Invitrogen) per extraction to facilitate RNA precipitation. RNA was quantified by UV absorbance using a Nanodrop 8000 spectrophotometer (Thermo Fisher Scientific).

Reverse transcription was carried out using the High-Capacity cDNA Reverse Transcription (RT) kit (Applied Biosystems, Waltham, MA, USA) with 100 ng of total RNA per sample in a 20 µL final reaction volume. First, a 2X RT master mix was prepared following manufacturer's instructions containing 10X RT buffer, 25X dNTP Mix (100 nM), 10X RT random primers, MultiScribe™ Reverse Transcriptase and nuclease free water. Then, 10 µL of this master mix were mixed with 10 µL RNA sample and the RT was conducted with the following cycle: 10 min at 25 °C, 120 min at 37 °C and 5 min at 85 °C, after which samples were held at 4 °C.

The resulting cDNA (10 ng) was used to assemble qRT-PCR reactions in a final volume of 10 µL containing TaqMan Fast Advanced Master mix and TaqMan FAM-MGB gene expression assays Hs00270951_s1 (FOXC2), Hs02786624_g1 (GAPDH) and Hs01060665_g1 (ACTB) (Applied Biosystems). The thermal cycling was done in a QuantStudio™ 7 Flex Real-Time PCR system (Thermo Fisher Scientific) performing a 20 s hold at 95 °C followed by 40 cycles of 1 s at 95 °C and 20 s at 60 °C. No template controls were used in each reaction as negative control. FOXC2 expression values in LECs on-chip were normalized to internal controls (β -actin and GAPDH) and presented as fold change relative to LECs cultured on flask using the comparative Ct method.

Supplementary figures

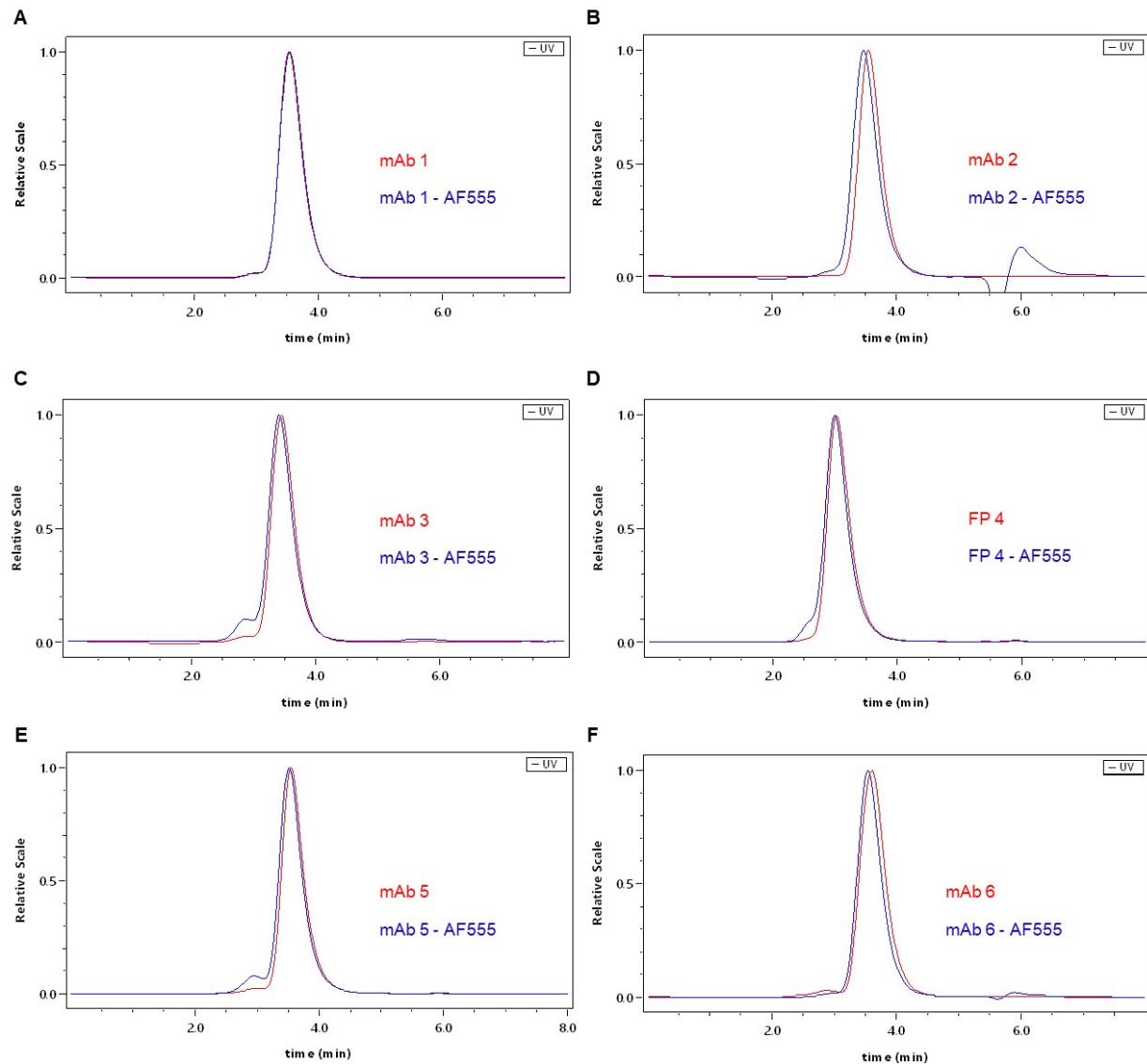


Fig. S1. Size exclusion chromatograms of monoclonal antibodies (mAb) and fusion protein (FP) before and after labeling. A-F: chromatograms of molecule list 1-6 showing the unconjugated and Alexa Fluor 555 (AF555) conjugated molecule in red and blue, respectively.

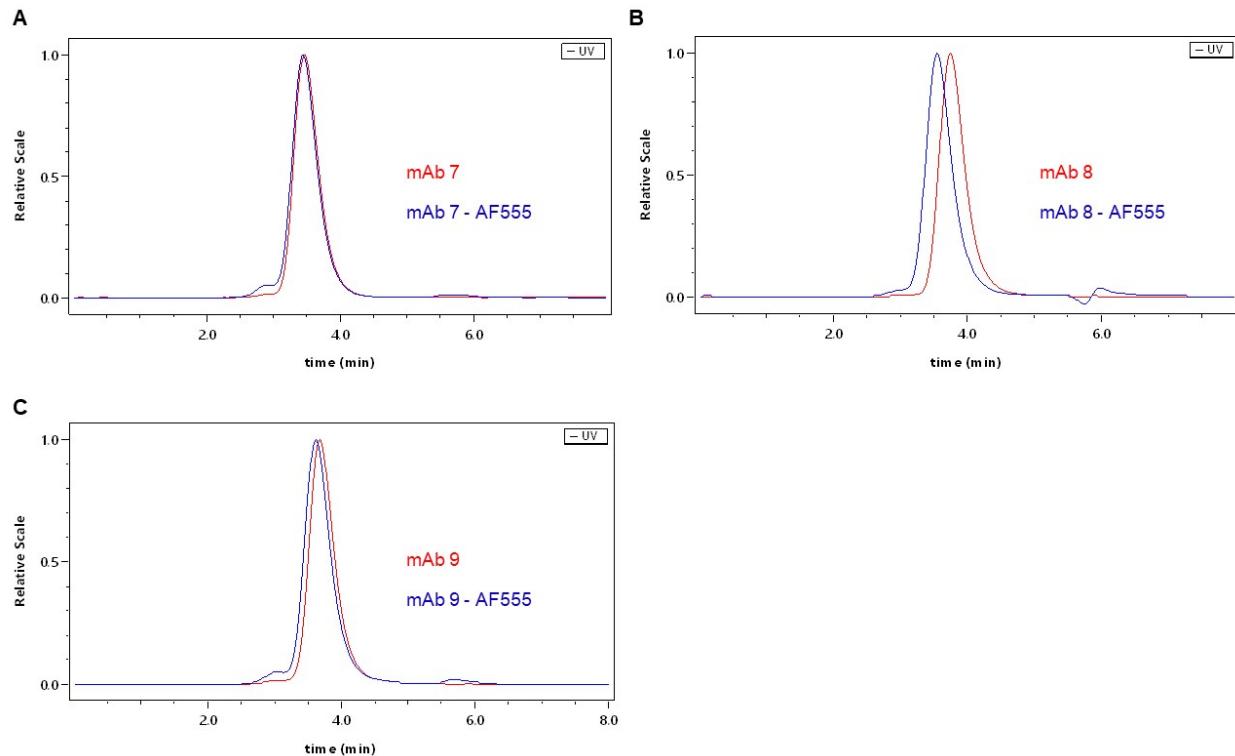


Fig. S2. Size exclusion chromatograms of monoclonal antibodies (mAb) and fusion protein (FP) before and after labeling. A-F: chromatograms of molecule list 7-9 showing the unconjugated and Alexa Fluor 555 (AF555) conjugated molecule in red and blue, respectively.

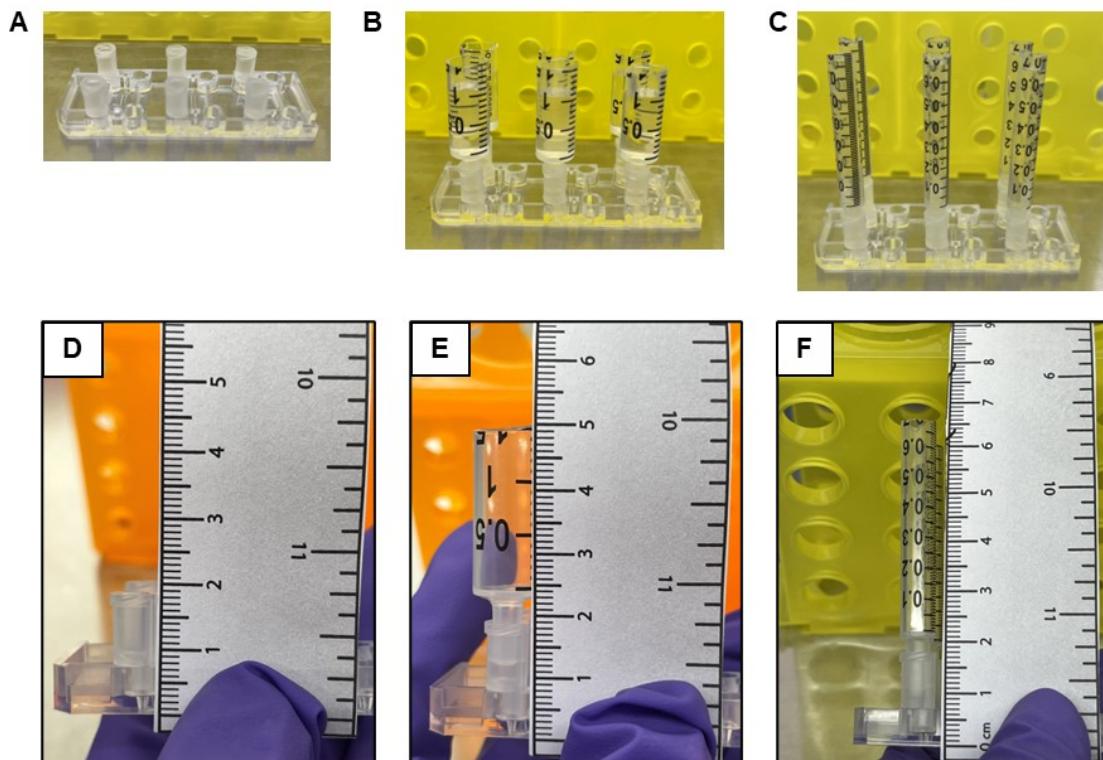


Fig. S3. Chip setups for hydrostatic pressure driven flow induction. A) Luer setup used for antibody transport measurements. B) 3 mL syringe setup used for lymphatic endothelial cell (LEC) growth. C) 1 mL syringe setup used for LEC growth. D) Pressure head in luer setup (~1.8 cm). E) Pressure head in 3 mL syringe setup (~4.2 cm). F) Pressure head in 1 mL syringe setup (~6.3 cm).

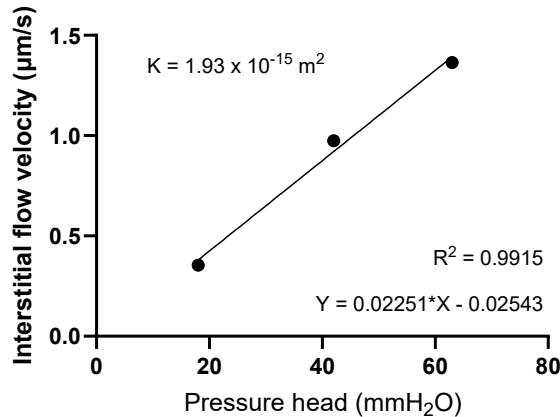


Fig. S4. Correlation analysis of pressure head and interstitial flow velocity. Linear regression analysis of the pressure head established by the different flow setups: luer (~18 mmH₂O), 3 mL syringe (~42 mmH₂O) and 1 mL syringe (~63 mmH₂O), and the resulting interstitial flow velocity within the extracellular matrix ($R^2 = 0.9915$). The resulting equation is used to calculate the hydraulic permeability at 18 mmH₂O as follows: Hydraulic permeability (K) = gel width (m) * viscosity (kg/m/s) * velocity (m/s) / pressure difference (kg/m/s²) $K = 1.3e-3 \text{ m} * 6.9e-4 \text{ kg/m/s} * 0.3797e-6 \text{ m/s} / 176.58 \text{ kg/m/s}^2 = 1.93e-15 \text{ m}^2$.

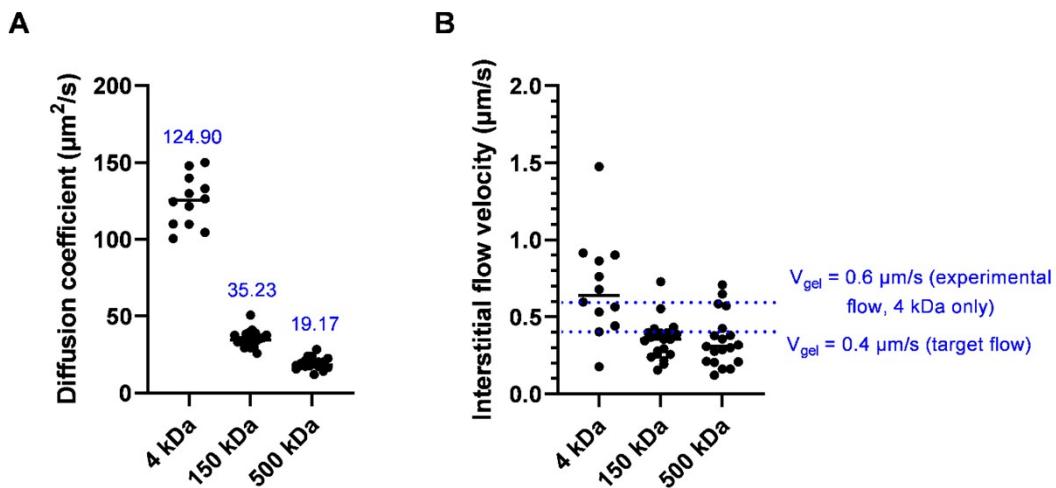


Fig. S5. Measured (black) vs simulated (blue) transport parameters for 4, 150 and 500 kDa FITC-dextran transport evaluations. A) Diffusion coefficients. B) Interstitial flow velocities. Experimental flow in simulations ($v_{gel}=0.6 \mu\text{m/s}$) was achieved by a 2-fold decrease in gel resistance as well as a 2-fold increase of the membrane conductivity compared to the conditions used for the simulation of target flow ($v_{gel}=0.4 \mu\text{m/s}$).

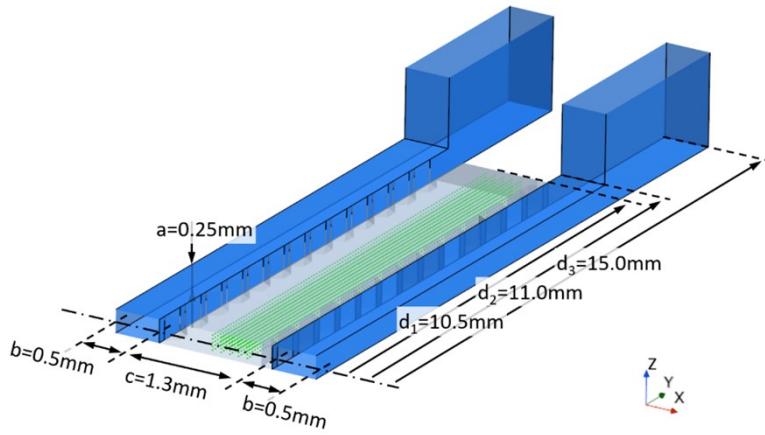


Fig. S6. Geometric dimensions of in- and outlet channel (blue) and gel matrix (grey) with lymph vessel locations (green). Only half-model is shown due to symmetry to the xz-plane (dashed dotted center line).

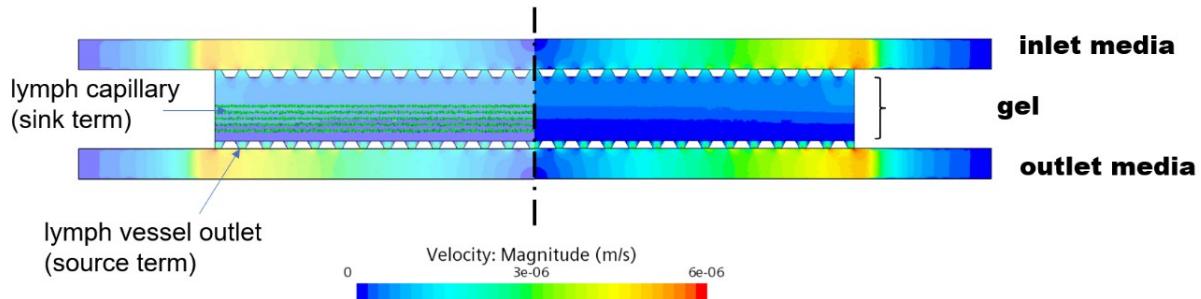


Fig. S7. Flow velocities in CFD simulation of the MPS with target gel flow velocity $v_{gel} = 0.4 \mu\text{m/s}$ showing the reduced velocity in the gel at the absorption sites and the high flows at the locations of re-administration into the outlet channel. For better visibility of the contour plot the elements used for lymph capillary absorption are not visualized on the right-hand side of the graphic. With the dimensions $a= 0.25 \text{ mm}$ and $b= 0.5 \text{ mm}$ from **Fig. S6** the Reynolds number for maximum flow velocity $6 \mu\text{m/s}$ is calculated to $Re = \rho * v * D_h / \mu = 0.003$ with $D_h = 2ab/(a+b)$.

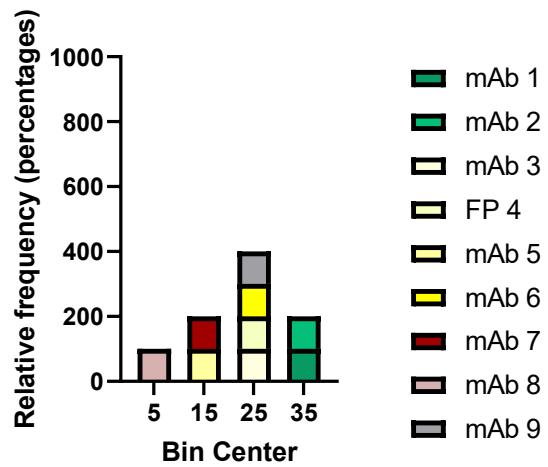


Fig. S8. Frequency distribution analysis of percentage monoclonal antibody (mAb) and fusion protein (FP) transport on-chip showing a 4-bin classification of the molecules. Bin width = 10; center of first bin = 5; center of last bin = 35.

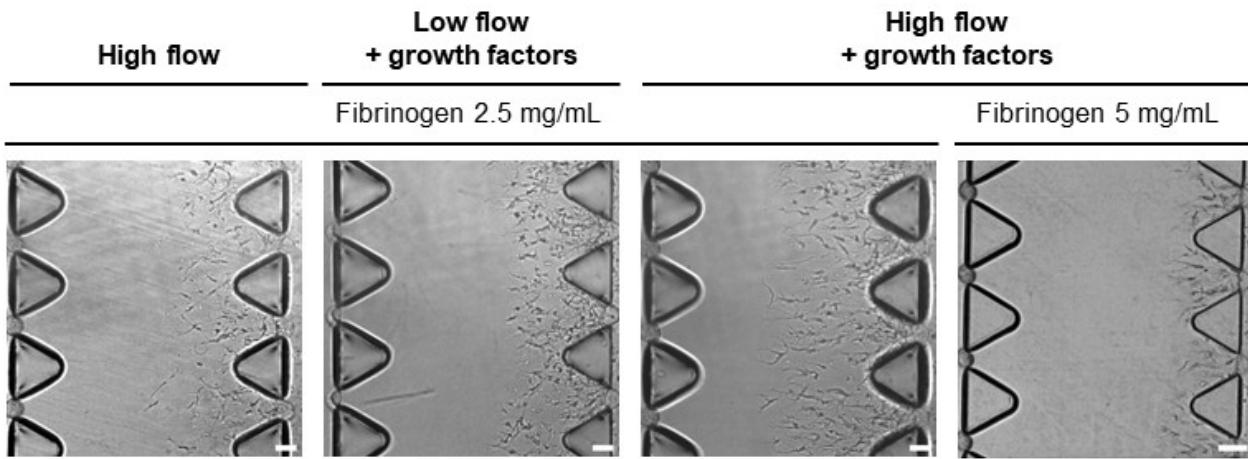


Fig. S9. Bright field images showing the morphology of lymphatic capillaries grown under high and low flow, in 2.5 and 5 mg/mL fibrin EMCs, with and without growth factor supplementation. Scale bar = 100 μ m.

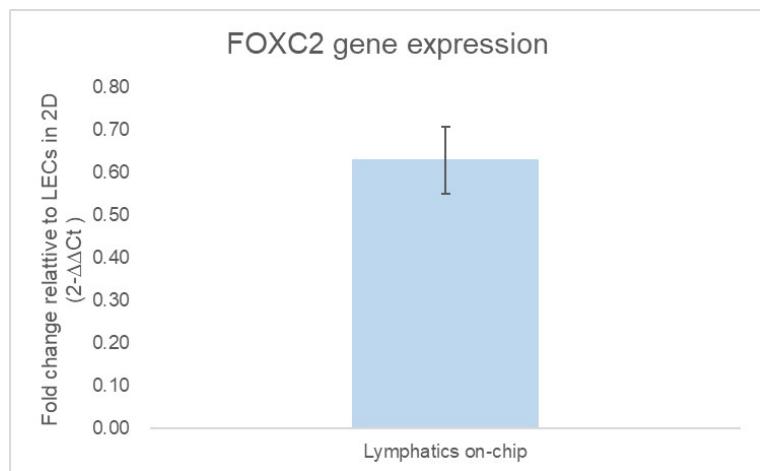


Fig. S10. FOXC2 mRNA expression in lymphatics on-chip. Expression values in LECs on-chip were normalized to internal controls (β -actin and GAPDH) and presented as fold change relative to LECs cultured on 2D flask using the comparative Ct method.

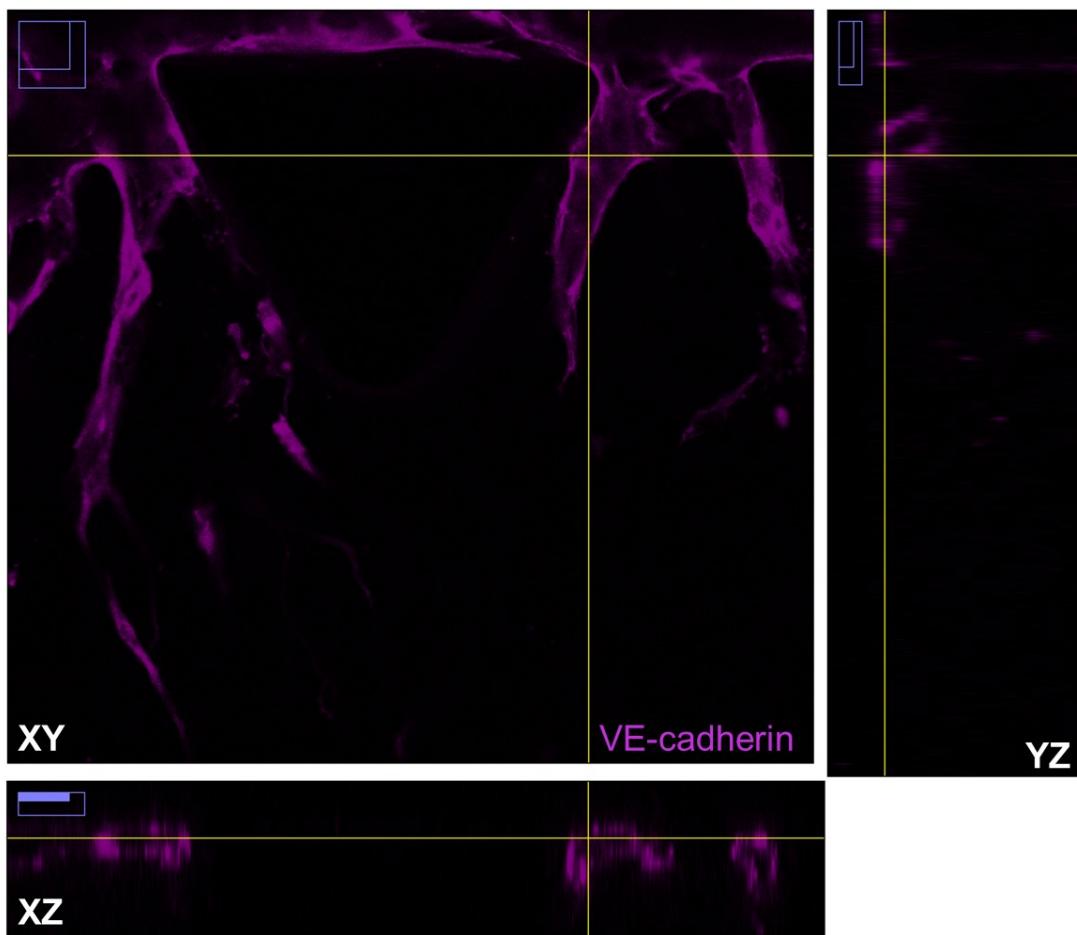


Fig. S11. Confocal cross-sections of VE-cadherin immunostaining showing lymphatic capillary lumens. XY image shows the morphology of lymphatic capillaries growing from the lymphatic tubule in the

outlet channel (top) into the extracellular matrix in the central channel (bottom). Cross-sections of the capillary lumens are shown in the XZ and YZ images, marked by the yellow lines. Image dimensions are XY: 708.49x708.49 μm ; XZ: 708.49x253.29 μm and YZ: 253.29x708.49 μm .

Table S1. Material parameters for fluids, gel and membrane used for lymphatic chip simulations with computational fluid dynamics software.

Fluid Properties

| | |
|----------------------------------------------------------------------------|--------|
| Density (ρ , kg/m ³) | 1000.0 |
| Viscosity (μ , Pa*s) | 6.9e-4 |
| Diffusion coefficient for 4 kDa (D_4 , $\mu\text{m}^2/\text{s}$) | 124.9 |
| Diffusion coefficient for 150 kDa (D_{150} , $\mu\text{m}^2/\text{s}$) | 35.2 |

Gel Properties

| | |
|---------------------------------------------------------------------------|--------|
| Porosity (χ , --) | 0.9 |
| Porous viscous resistance for target flow (P_v , kg/m ³ /s) | 1e11 |
| Porous viscous resistance for exp. flow (P_v , kg/m ³ /s) | 0.5e11 |

Membrane Properties

| | |
|-----------------------------------------------------------------|---------|
| Thickness (Δx , μm) | 1.0 |
| Hydraulic conductivity for target flow (K_L , m^2) | 1.0e-18 |
| Hydraulic conductivity for exp. flow (K_L , m^2) | 2.0e-18 |

Table S2. Fluorescence intensity values detected by confocal microscopy in the outlet and inlet (reference) chip channels and corresponding drainage rate values calculated for fluorescein isothiocyanate-dextran of 4 kDa.

| Time (min) | Dextran 4 kDa | | | Drainage rate (1/min) (n=3) | | |
|------------------|--------------------------------------|----------|-----------|-----------------------------|----------|----------|
| | Mean fluorescence intensity (n=3) | | | | | |
| 0 | 308.923 | 376.636 | 332.402 | | | |
| 2 | 285.783 | 353.295 | 340.518 | -0.00042 | -0.00043 | 0.000211 |
| 4 | 310.554 | 543.301 | 358.64 | 0.000452 | 0.003534 | 0.00047 |
| 6 | 456.827 | 1114.475 | 421.485 | 0.00267 | 0.010622 | 0.00163 |
| 8 | 817.793 | 2090.094 | 526.261 | 0.006589 | 0.018144 | 0.002718 |
| 10 | 1380.343 | 3215.411 | 738.919 | 0.010269 | 0.020928 | 0.005516 |
| 12 | 2210.958 | 4479.696 | 1083.197 | 0.015162 | 0.023512 | 0.00893 |
| 14 | 3312.04 | 5792.161 | 1562.312 | 0.020099 | 0.024408 | 0.012427 |
| 16 | 4499.643 | 7177.83 | 2277.098 | 0.021678 | 0.02577 | 0.01854 |
| 18 | 5666.84 | 8603.799 | 3175.466 | 0.021306 | 0.026519 | 0.023301 |
| 20 | 6741.724 | 9890.584 | 4054.231 | 0.019621 | 0.023931 | 0.022793 |
| 22 | 7806.882 | 11196.65 | 5117.516 | 0.019443 | 0.024289 | 0.027579 |
| 24 | 8902.579 | 12533.11 | 6370.354 | 0.020001 | 0.024855 | 0.032496 |
| 26 | 9929.354 | 13678.28 | 7565.425 | 0.018743 | 0.021297 | 0.030997 |
| 28 | 10940.692 | 14697.2 | 8650.904 | 0.018461 | 0.018949 | 0.028155 |
| 30 | 11980.47 | 15713.42 | 9911.693 | 0.01898 | 0.018899 | 0.032702 |
| Reference | 27391.444 | 26885.64 | 19277.067 | | | |

Table S3. Fluorescence intensity values detected by confocal microscopy in the outlet and inlet (reference) chip channels and corresponding drainage rate values calculated for fluorescein isothiocyanate-dextran of 150 kDa.

| Dextran 150 kDa | | | | | | | | |
|------------------|-----------------------------------|----------|-----------|----------|-----------------------------|----------|----------|----------|
| Time (min) | Mean fluorescence intensity (n=4) | | | | Drainage rate (1/min) (n=4) | | | |
| 0 | 314.455 | 313.407 | 252.072 | 281.143 | | | | |
| 2 | 284.053 | 310.21 | 286.237 | 229.626 | -0.00048 | -0.0001 | 0.000751 | -0.00118 |
| 4 | 289.455 | 339.219 | 244.337 | 228.278 | 8.54E-05 | 0.000912 | -0.00092 | -3.1E-05 |
| 6 | 341.125 | 383.712 | 260.07 | 239.074 | 0.000817 | 0.001399 | 0.000346 | 0.000246 |
| 8 | 512.779 | 464.268 | 289.967 | 294.381 | 0.002714 | 0.002533 | 0.000658 | 0.001262 |
| 10 | 923.163 | 620.215 | 366.659 | 480.345 | 0.006489 | 0.004904 | 0.001687 | 0.004244 |
| 12 | 1596.636 | 858.59 | 512.393 | 823.961 | 0.01065 | 0.007497 | 0.003205 | 0.007842 |
| 14 | 2500.848 | 1167.642 | 745.138 | 1374.949 | 0.014298 | 0.009719 | 0.005119 | 0.012574 |
| 16 | 3647.629 | 1609.417 | 1050.702 | 2089.123 | 0.018134 | 0.013893 | 0.00672 | 0.016298 |
| 18 | 5002.178 | 2162.725 | 1464.786 | 2985.897 | 0.021419 | 0.017401 | 0.009107 | 0.020465 |
| 20 | 6440.882 | 2752.866 | 1947.709 | 3962.381 | 0.02275 | 0.018559 | 0.010621 | 0.022284 |
| 22 | 7902.854 | 3413.98 | 2475.545 | 4925.954 | 0.023118 | 0.020791 | 0.011609 | 0.02199 |
| 24 | 9556.24 | 4174.561 | 3051.693 | 5902.047 | 0.026145 | 0.02392 | 0.012672 | 0.022275 |
| 26 | 11172.412 | 4957.157 | 3654.356 | 6978.572 | 0.025556 | 0.024612 | 0.013255 | 0.024567 |
| 28 | 12748.981 | 5716.462 | 4297.961 | 7941.059 | 0.02493 | 0.023879 | 0.014155 | 0.021965 |
| 30 | 14230.691 | 6457.991 | 4916.189 | 8677.694 | 0.02343 | 0.02332 | 0.013597 | 0.016811 |
| Reference | 31619.66 | 15898.75 | 22733.898 | 21909.6 | | | | |

Table S4. Fluorescence intensity values detected by confocal microscopy in the outlet and inlet (reference) chip channels and corresponding drainage rate values calculated for fluorescein isothiocyanate-dextran of 500 kDa.

| Dextran 500 kDa | | | | | | |
|-----------------|-----------------------------------|----------|----------|-----------------------------|----------|----------|
| Time (min) | Mean fluorescence intensity (n=3) | | | Drainage rate (1/min) (n=3) | | |
| 0 | 266.232 | 297.724 | 319.84 | -0.00039 | -0.00025 | 0.001068 |
| 2 | 251.402 | 288.825 | 360.479 | 0.000399 | 5.36E-05 | -0.00064 |
| 4 | 266.729 | 290.746 | 336.184 | -5.9E-05 | -0.00018 | 0.000494 |
| 6 | 264.467 | 284.306 | 354.973 | -3.6E-05 | 5.3E-05 | -4.6E-05 |
| 8 | 263.09 | 286.205 | 353.221 | 0.000115 | 0.00014 | 0.000903 |
| 10 | 267.515 | 291.222 | 387.6 | 0.00031 | 0.00064 | 0.001932 |
| 12 | 279.424 | 314.163 | 461.134 | 0.000784 | 0.002175 | 0.004525 |
| 14 | 309.504 | 392.082 | 633.377 | 0.001758 | 0.003983 | 0.008546 |
| 16 | 376.98 | 534.782 | 958.711 | 0.003177 | 0.008114 | 0.012788 |
| 18 | 498.914 | 825.465 | 1445.515 | 0.004496 | 0.011673 | 0.014196 |
| 20 | 671.489 | 1243.647 | 1985.942 | 0.005977 | 0.013831 | 0.01732 |
| 22 | 900.876 | 1739.121 | 2645.272 | 0.008473 | 0.015677 | 0.022227 |
| 24 | 1226.066 | 2300.725 | 3491.423 | 0.010029 | 0.022346 | 0.020903 |

| | | | | | | |
|------------------|-----------|----------|-----------|----------|----------|----------|
| 26 | 1610.988 | 3101.225 | 4287.169 | 0.012277 | 0.025731 | 0.021153 |
| 28 | 2082.189 | 4023.012 | 5092.42 | 0.013949 | 0.027514 | 0.023066 |
| 30 | 2617.546 | 5008.66 | 5970.518 | | | |
| Reference | 19190.375 | 17911.76 | 19034.228 | | | |

Table S5. Fluorescence intensity values detected by confocal microscopy in the outlet and inlet (reference) chip channels and corresponding drainage rate values calculated for IgG-DyLight650.

Table S6. Fluorescence intensity values detected by confocal microscopy in the outlet channel for fluorescein isothiocyanate-dextran of 4 kDa normalized to the values detected in the inlet (reference) channel.

| Dextran 4 kDa | | | |
|---------------|-----------------------------------------------------------|----------|-------------|
| Time (min) | Mean fluorescence intensity normalized to reference (n=3) | | |
| 0 | 0.011278 | 0.014009 | 0.017243391 |
| 2 | 0.010433 | 0.013141 | 0.017664409 |
| 4 | 0.011338 | 0.020208 | 0.01860449 |
| 6 | 0.016678 | 0.041452 | 0.021864581 |
| 8 | 0.029856 | 0.07774 | 0.027299848 |
| 10 | 0.050393 | 0.119596 | 0.038331506 |
| 12 | 0.080717 | 0.16662 | 0.056190965 |
| 14 | 0.120915 | 0.215437 | 0.081045109 |
| 16 | 0.164272 | 0.266976 | 0.118124713 |
| 18 | 0.206884 | 0.320015 | 0.164727653 |
| 20 | 0.246125 | 0.367876 | 0.210313685 |
| 22 | 0.285012 | 0.416455 | 0.265471713 |
| 24 | 0.325013 | 0.466164 | 0.330462824 |
| 26 | 0.362498 | 0.508758 | 0.392457265 |
| 28 | 0.39942 | 0.546656 | 0.44876661 |
| 30 | 0.43738 | 0.584454 | 0.51417018 |

Table S7. Fluorescence intensity values detected by confocal microscopy in the outlet channel for fluorescein isothiocyanate-dextran of 150 kDa normalized to the values detected in the inlet (reference) channel.

| Dextran 150 kDa | | | | |
|-----------------|-----------------------------------------------------------|----------|-------------|----------|
| Time (min) | Mean fluorescence intensity normalized to reference (n=4) | | | |
| 0 | 0.009945 | 0.019713 | 0.011087936 | 0.012832 |
| 2 | 0.008983 | 0.019512 | 0.012590758 | 0.010481 |
| 4 | 0.009154 | 0.021336 | 0.010747695 | 0.010419 |
| 6 | 0.010788 | 0.024135 | 0.011439745 | 0.010912 |
| 8 | 0.016217 | 0.029202 | 0.01275483 | 0.013436 |
| 10 | 0.029196 | 0.03901 | 0.016128294 | 0.021924 |
| 12 | 0.050495 | 0.054004 | 0.022538722 | 0.037607 |
| 14 | 0.079092 | 0.073442 | 0.032776517 | 0.062756 |
| 16 | 0.11536 | 0.101229 | 0.046217415 | 0.095352 |
| 18 | 0.158198 | 0.136031 | 0.064431801 | 0.136283 |
| 20 | 0.203699 | 0.17315 | 0.085674221 | 0.180851 |
| 22 | 0.249935 | 0.214733 | 0.108892237 | 0.224831 |
| 24 | 0.302225 | 0.262572 | 0.134235361 | 0.269382 |
| 26 | 0.353338 | 0.311795 | 0.160744805 | 0.318517 |
| 28 | 0.403198 | 0.359554 | 0.189055172 | 0.362447 |
| 30 | 0.450058 | 0.406195 | 0.216249277 | 0.396068 |

Table S8. Normalized fluorescence intensity values detected by confocal microscopy for fluorescein isothiocyanate-dextran of 4 kDa (**Table S6**) transformed to concentration.

| Dextran 4 kDa | | | |
|---------------|------------------------------------------------------------------------|----------|----------|
| Time (min) | Mean fluorescence intensity transformed to concentration (mg/mL) (n=3) | | |
| 0 | 0.000112781 | 0.00014 | 0.000172 |
| 2 | 0.000104333 | 0.000131 | 0.000177 |
| 4 | 0.000113376 | 0.000202 | 0.000186 |
| 6 | 0.000166777 | 0.000415 | 0.000219 |
| 8 | 0.000298558 | 0.000777 | 0.000273 |
| 10 | 0.000503932 | 0.001196 | 0.000383 |
| 12 | 0.000807171 | 0.001666 | 0.000562 |
| 14 | 0.001209151 | 0.002154 | 0.00081 |
| 16 | 0.001642718 | 0.00267 | 0.001181 |
| 18 | 0.002068836 | 0.0032 | 0.001647 |
| 20 | 0.002461252 | 0.003679 | 0.002103 |
| 22 | 0.002850117 | 0.004165 | 0.002655 |
| 24 | 0.003250131 | 0.004662 | 0.003305 |
| 26 | 0.003624984 | 0.005088 | 0.003925 |
| 28 | 0.003994201 | 0.005467 | 0.004488 |
| 30 | 0.0043738 | 0.005845 | 0.005142 |

Table S9. Normalized fluorescence intensity values detected by confocal microscopy for fluorescein isothiocyanate-dextran of 150 kDa (**Table S7**) transformed to concentration.

| Dextran 150 kDa | | | | |
|-----------------|------------------------------------------------------------------------|----------|----------|----------|
| Time (min) | Mean fluorescence intensity transformed to concentration (mg/mL) (n=4) | | | |
| 0 | 9.94492E-05 | 0.000197 | 0.000111 | 0.000128 |
| 2 | 8.98343E-05 | 0.000195 | 0.000126 | 0.000105 |
| 4 | 9.15427E-05 | 0.000213 | 0.000107 | 0.000104 |
| 6 | 0.000107884 | 0.000241 | 0.000114 | 0.000109 |
| 8 | 0.000162171 | 0.000292 | 0.000128 | 0.000134 |
| 10 | 0.000291959 | 0.00039 | 0.000161 | 0.000219 |
| 12 | 0.00050495 | 0.00054 | 0.000225 | 0.000376 |
| 14 | 0.000790916 | 0.000734 | 0.000328 | 0.000628 |
| 16 | 0.001153595 | 0.001012 | 0.000462 | 0.000954 |
| 18 | 0.001581983 | 0.00136 | 0.000644 | 0.001363 |
| 20 | 0.002036986 | 0.001731 | 0.000857 | 0.001809 |
| 22 | 0.002499348 | 0.002147 | 0.001089 | 0.002248 |
| 24 | 0.003022246 | 0.002626 | 0.001342 | 0.002694 |
| 26 | 0.003533375 | 0.003118 | 0.001607 | 0.003185 |
| 28 | 0.004031979 | 0.003596 | 0.001891 | 0.003624 |
| 30 | 0.004500583 | 0.004062 | 0.002162 | 0.003961 |