

Supplementary Information for:

***In Situ* Visualization of Hydrophilic Spatial Heterogeneity Inside
Microfluidic Chips by Fluorescence Microscopy**

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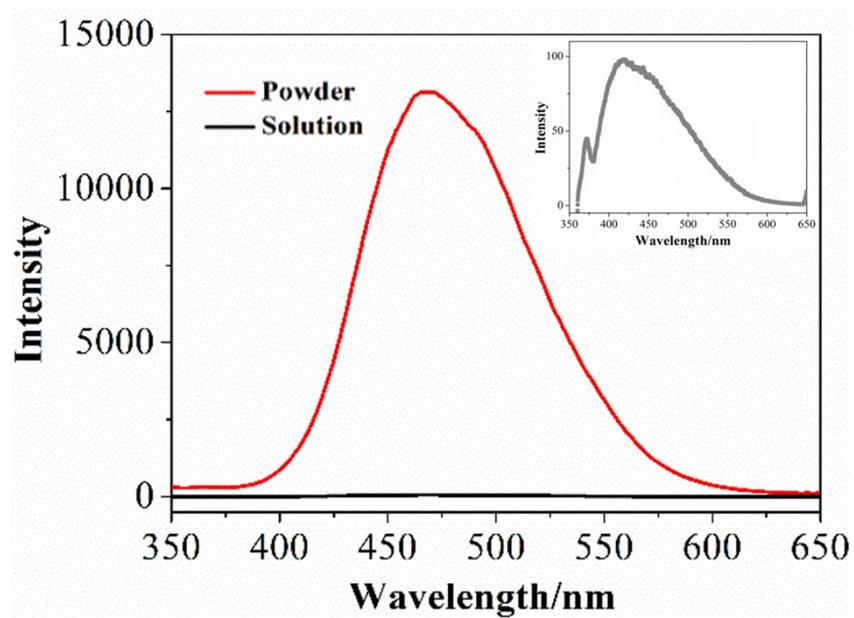


Fig. S1 Fluorescence emission spectra of TPEDB aqueous solution (50 μM) and powder under UV light irradiation (330 nm), the inset showed the enlarged spectra of TPEDB solution.

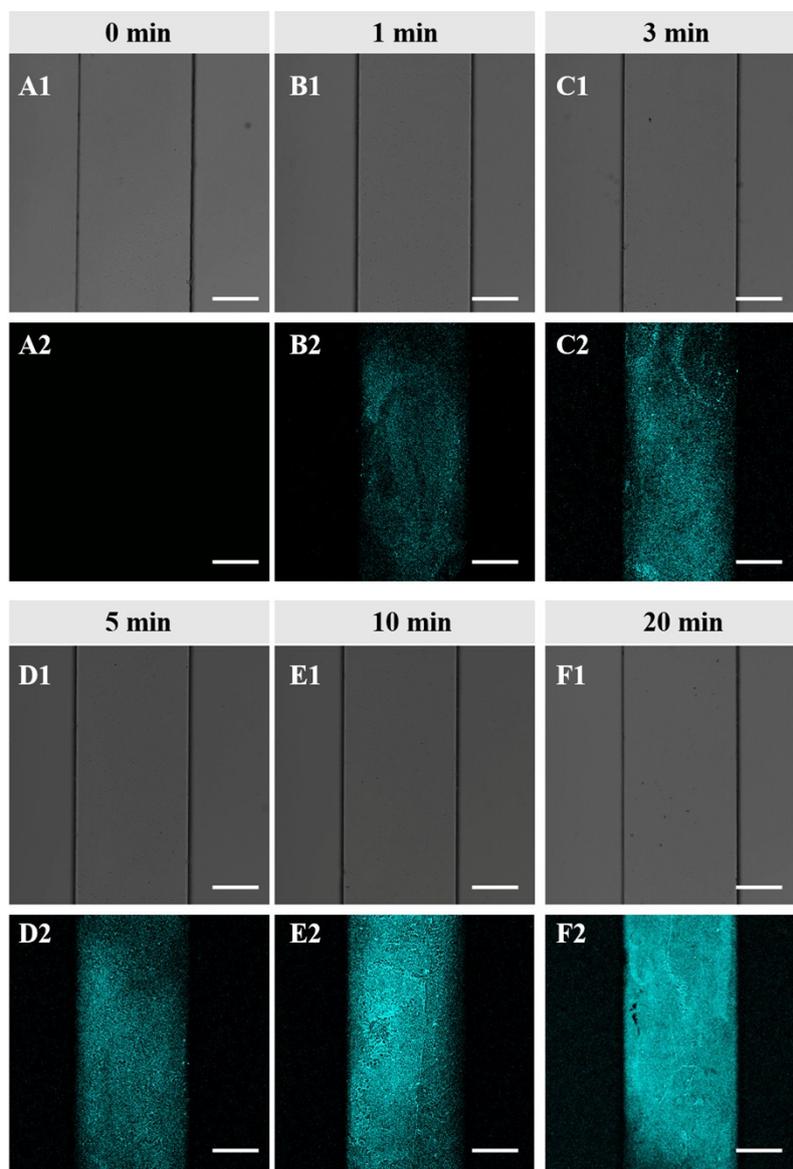


Fig. S2 Bright-field and dark-field images of confocal microscopy ($1200 \times 1200 \mu\text{m}^2$) for the PDMS microfluidic channels after hydrophilization treatment in PVA for different time, from A to F: treatment time is prolonged from 0 min, 1 min, 3 min, 5 min, 10 min to 20 min. The scale bars represented for 200 μm .

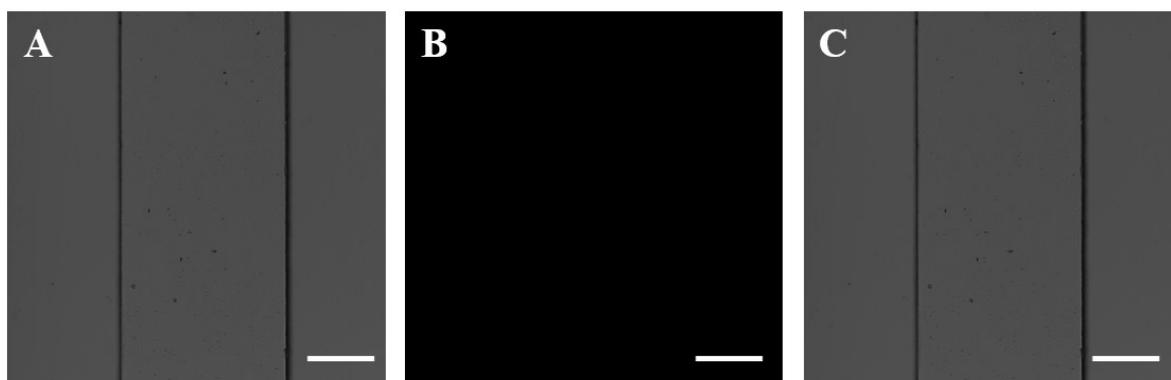


Fig. S3 Confocal microscopy images of (A) bright-field, (B) dark-field and (C) merged images ($1200 \times 1200 \mu\text{m}^2$) for the PDMS microfluidic channels after the dissociation of TPEDB molecules. The scale bars represented for $200 \mu\text{m}$.

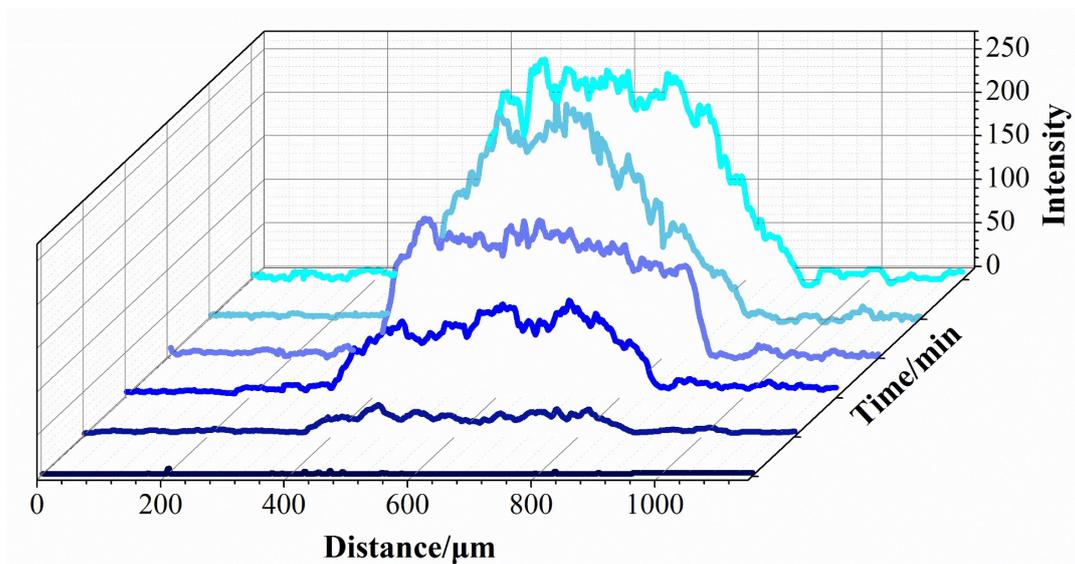


Fig. S4 Fluorescent intensity analysis for the PDMS microfluidic channels after treatment in PVA for different time (0 min, 1 min, 3 min, 5 min, 10 min and 20 min).

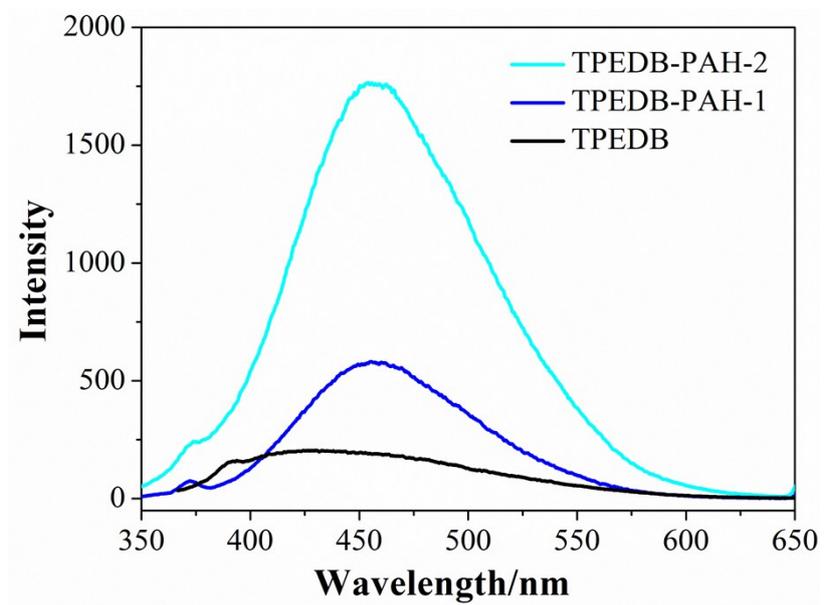


Fig. S5 Fluorescent emission spectra for TPEDB, TPEDB-PAH-1 and TPEDB-PAH-2 composites with the excitation of 330 nm.

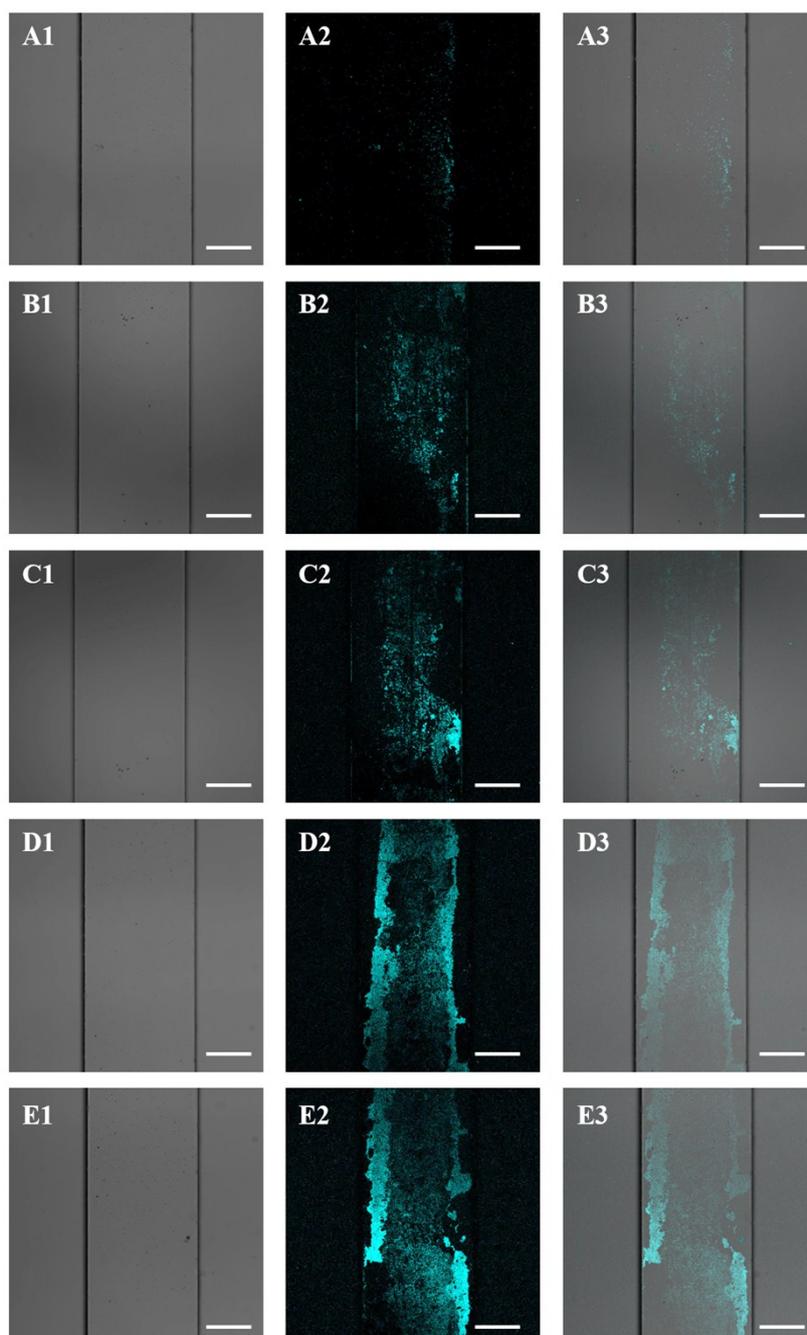


Fig. S6 Bright-field, dark-field and merged images of confocal microscopy ($1200 \times 1200 \mu\text{m}^2$) for the PDMS microfluidic channels after hydrophilization treatment in PAH for different time, from A to E: treatment time is prolonged from 1 min, 3 min, 5 min, 10 min to 20 min. The scale bars represented for $200 \mu\text{m}$.

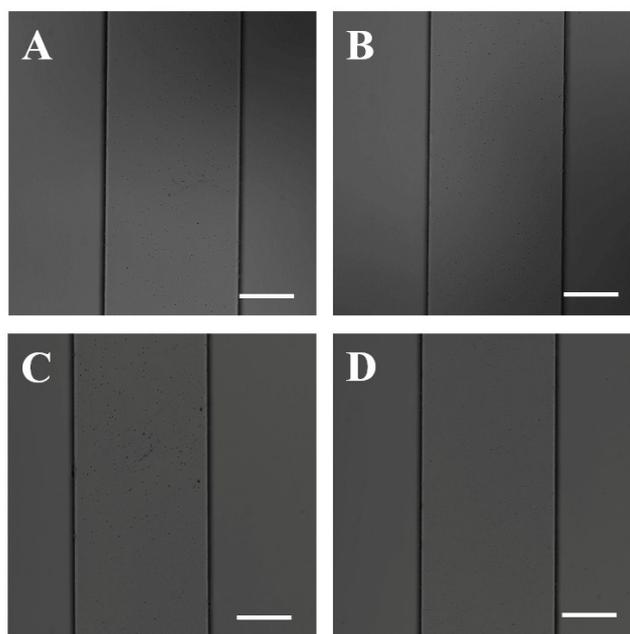


Fig. S7 Bright field images of confocal microscopy ($1200 \times 1200 \mu\text{m}^2$) for the PDMS microfluidic channels after different treatment: (A) HCl and H₂O₂, (B) 3% glycerol, (C) 3% PEG and (D) 3% PVA. The scale bars represented for 200 μm .

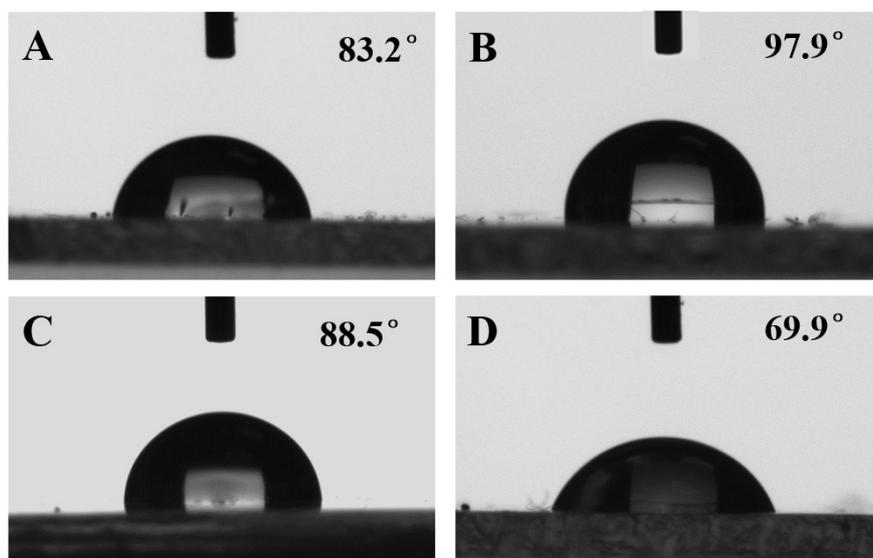


Fig. S8 Contact angles of PDMS flat substrates under different hydrophilization treatment: (A) HCl-H₂O₂ mixture, (B) 3% glycerol, (C) 3% PEG and (D) 3% PVA.

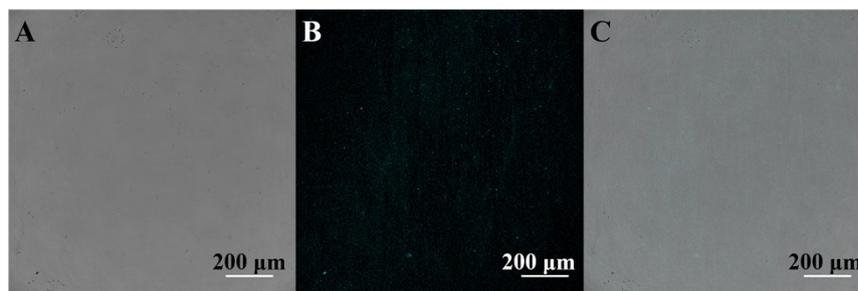


Fig. S9 (A) Bright-field, (B) dark-field and (C) merged images of confocal microscopy ($1500 \times 1500 \mu\text{m}^2$) for the glass substrate labelled by the TPEDB molecules.

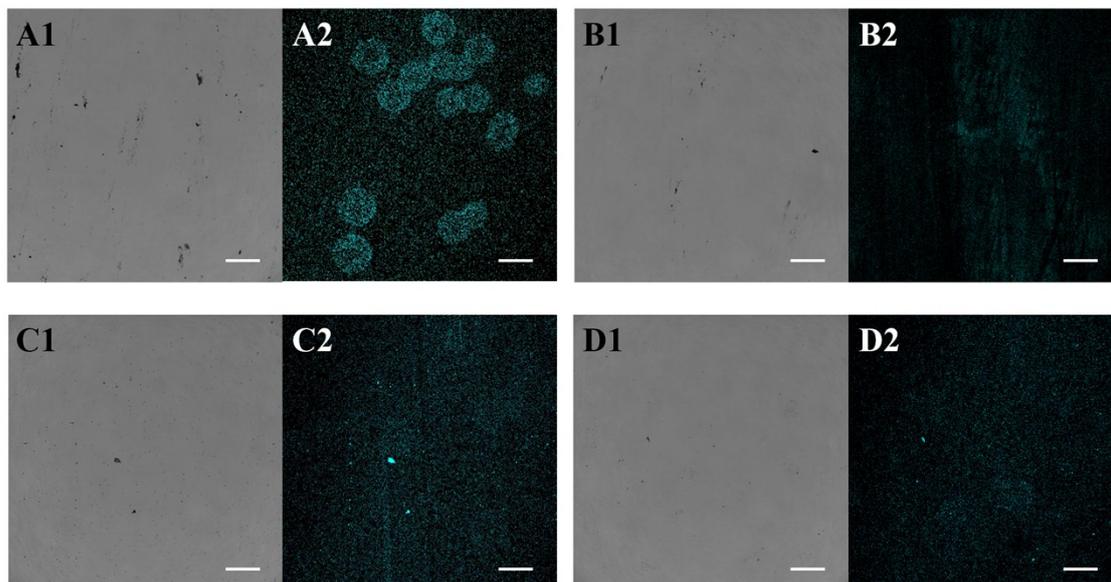


Fig. S10 Bright-field and dark-field images of confocal microscopy ($1500 \times 1500 \mu\text{m}^2$) for the glass substrate treated by (A) HCl-H₂O₂ mixture, (B) 3% glycerol, (C) 3% PEG and (D) 3% PVA, followed by the fluorescent labelling of the TPEDB molecules. Scale bars represented for 200 μm .

Table S1. Fluorescent imaging analysis of PDMS microfluidic channels after treatment in PVA solution for different time.

Time treated in PVA solution/min	Mean values/pixel
0	0.54
1	13.44
3	22.30
5	33.69
10	45.14
20	49.99

Table S2. Fluorescent imaging analysis of PDMS microfluidic channels after treatment in HCl-H₂O₂ mixture, 3% glycerol, 3% PEG and 3% PVA for 10 min.

Treatment	Mean values/pixel
HCl-H ₂ O ₂	23.37
3% glycerol	16.68
3% PEG	13.93
3% PVA	44.05

Table S3. Contact angle measurements of glass substrate after treatment in HCl-H₂O₂ mixture, 3% glycerol, 3% PEG and 3% PVA for 10 min.

Treatment	Contact angle/degree	Mean values/pixel
Blank	55.6	4.89
HCl-H ₂ O ₂	32.4	19.32
3% glycerol	50.9	7.65
3% PEG	45.5	9.35
3% PVA	49.2	9.03