## **Supplementary Information**

## Submicron-Thick Single Anion-Conducting Polymer Electrolytes

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Fig. S1 Contact-angle measurements for various iCVD-derived pDMAMS films.

Sample ID	R1	R2	Q <sub>1</sub>	Q <sub>1</sub> _a
Br_1	67	1128	1.0×10 <sup>-10</sup>	1
Br_2	52	1291	1.2×10 <sup>-10</sup>	1
Br_3	103	1070	0.7×10 <sup>-10</sup>	1
Br_4	22	1414	2.8×10 <sup>-10</sup>	1
Br_5	38	885	1.4×10 <sup>-10</sup>	1
Average	56	1158	1.4×10 <sup>-10</sup>	1
HCO <sub>3</sub> _1	62	97	4.5×10 <sup>-9</sup>	0.9
HCO <sub>3</sub> _2	93	11	5.3×10 <sup>-10</sup>	1
HCO <sub>3</sub> _3	87	56	1.2×10 <sup>-9</sup>	1
HCO <sub>3</sub> _4	119	78	2.2×10 <sup>-8</sup>	0.8
HCO <sub>3</sub> _5	58	145	6.7×10 <sup>-10</sup>	1
HCO <sub>3</sub> _6	115	48	1.1×10 <sup>-7</sup>	0.75
Average	84	77	5.8×10 <sup>-9</sup>	0.94
OH_1	76	56	1.7×10 <sup>-8</sup>	0.85
OH_2	76	37	2.5×10 <sup>-8</sup>	0.85
OH_3	67	66	1.7×10 <sup>-8</sup>	0.85
OH_4	75	42	1.1×10 <sup>-8</sup>	0.9
OH_5	74	56	7.6×10 <sup>-9</sup>	0.9
OH_6	58	89	6.0×10 <sup>-9</sup>	0.9
Average	74	51	1.6×10 <sup>-8</sup>	0.87

 Table S1 Fit parameters from dry Nyquist equivalent-circuit models.

**Table S2** Fit parameters from water-swelled Nyquist equivalent-circuit models.

Sample ID	R1	R2	Q <sub>1</sub>	Q <sub>1</sub> _a	Q <sub>2</sub>	Q <sub>2</sub> _a
Br_1	24	23	7.3×10 <sup>-9</sup>	1	5.4×10 <sup>-5</sup>	0.68
Br_2	15	68	4.7×10 <sup>-9</sup>	1	4.2×10 <sup>-5</sup>	0.70
Br_3	21	34	6.8×10 <sup>-9</sup>	1	6.2×10 <sup>-5</sup>	0.70
Br_4	29	23	8.7×10 <sup>-9</sup>	1	2.0×10 <sup>-4</sup>	0.59
average	22	37	6.9×10 <sup>-9</sup>	1	9.2×10 <sup>-5</sup>	0.67
HCO <sub>3</sub> 1	20	7	5.4×10 <sup>-7</sup>	0.95	9.2×10 <sup>-5</sup>	0.67
HCO <sub>3</sub> _2	19	11	1.8×10 <sup>-7</sup>	1	1.0×10 <sup>-4</sup>	0.66
HCO <sub>3</sub> _3	30	11	2.3×10 <sup>-7</sup>	0.97	1.0×10 <sup>-4</sup>	0.67
average	23	10	3.2×10 <sup>-7</sup>	0.97	9.8×10 <sup>-5</sup>	0.67
OH_1	19	26	7.3×10 <sup>-9</sup>	1	8.3×10 <sup>-5</sup>	0.72
OH_2	0	17	1.1×10 <sup>-5</sup>	0.45	1.9×10 <sup>-4</sup>	0.63
OH_3	21	14	6.5×10 <sup>-6</sup>	0.8	4.9×10 <sup>-5</sup>	0.71
OH_4	16	8	2.2×10 <sup>-6</sup>	0.91	5.5×10 <sup>-5</sup>	0.71
OH_5	13	9	5.5×10 <sup>-6</sup>	0.85	5.2×10 <sup>-5</sup>	0.74
average	14	15	5.0×10 <sup>-6</sup>	0.8	8.6×10 <sup>-5</sup>	0.70



**Fig. S2** Water uptake in hydration number ( $\lambda$ ) and percent mass (of the dry membrane mass) for pDMAMS<sup>+</sup>(Br<sup>-</sup>) equilibrated over P<sub>2</sub>O<sub>5</sub> (0% RH), various salt solutions (11–97% RH), and in liquid deionized water (100% RH).



**Fig. S3** EIS spectra of pDMAMS<sup>+</sup>(Br-) showing adequate electrode-electrolyte contact. Inset: current response of potentiostatic hold at 0.1 V.