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Supporting Information

Dimensionally stable hexamethylenetetramine functionalized

polysulfone anion exchange membranes

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Fig. S1. RDFs and CNs of different atoms in the dry PSF-ImOH membrane.



Fig. S2. RDFs and CNs of different atoms in the dry PSF-QuOH membrane.



Fig. S3. RDFs (solid lines) and CNs (dashed lines) of N⁺-N in the dry and hydrated (a) PSF-QuOH and (b) PSF-ImOH membranes. (WU: water uptake)

AEMs	N ⁺ -O ₁		N ⁺ -O ₂		N ⁺ -O(S)		N-S		N ⁺ -N	
	r ^a (Å)	$CN(O_1)$	r ^a (Å)	CN(O ₂)	r ^a (Å)	CN(O(S))	r ^a (Å)	CN(S)	r ^a (Å)	CN(N)
PSF-ImOH	_b	0	_ b	0	4.9	0.8	7.1	0.8	9.0	2.7°
PSF-QuOH	_ b	0	_ b	0	5.0	0.5	7.9	2.0	9.6	9.1 ^d

Table S1 Coordination Numbers of different atoms.

^a Upper boundary for integration of the first shell.

^bThe first peaks of the RDFs (except for the first strong peaks (<4 Å)) are very weak. This indicates the atoms are uniform distribution in the AEMs and the corresponding CNs are close to zero.

^{c, d}CN(N)=CN(total)-CN(one repeat unit), CN(total) is the integration of the RDFs; CN(one repeat unit) is the numbers of N atoms around N^+ ion in one repeat unit, and equal to 1 for PSF-ImOH and 3 for PSF-QuOH, respectively.



Fig. S4. TGA and DTG curves of the PSF-QuOH 1.67 membrane.



Fig. S5. FTIR of the PSF-QuOH 1.94 membrane before and after 1 M KOH immersion at 60 °C for 168 h.



Fig. S6. Alkaline stability of the PSF-QuOH 1.39 after 4 M NaOH immersion at 80 °C for 264 h.



Fig. S7. ¹H NMR spectra of BHMTA degradation in 1 M NaOH CD₃OD/D₂O (3:1).



Fig. S8. Chemical shift correlated spectroscopy (COSY) of BHMTA degradation products