Electronic Supplementary Material (ESI) for Journal of Materials Chemistry A. This journal is © The Royal Society of Chemistry 2022

Towards higher electrochemical stability of electrolytes: lithium salt design through *in silico* screening

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Electronic Supplementary Information

 Table S1 – List of abbreviations for anion structures investigated in this study.

Abbreviation		Chemical Name
R_1	TFSI	bis((trifluoromethyl)sulfonyl)imide
R_2	FSI	bis(fluorosulfonyl)imide
R_3	FTFSI	(fluorosulfonyl)((trifluoromethyl)sulfonyl)imide
S_1	MSI	bis(methylsulfonyl)imide
S_2	ESI	bis(ethylsulfonyl)imide
S_3	BETI	bis((perfluoroethyl)sulfonyl)imide
S_4	TFESI	bis((2,2,2-trifluoroethyl)sulfonyl)imide
S_5	TbSI	bis(tert-butylsulfonyl)imide
S_6	NFSI	bis((1,1,1,3,3,3-hexafluoro-2-(trifluoromethyl)propan-2-yl)sulfonyl)imide
S ₇	CSI	bis(cyanosulfonyl)imide
S ₈	CmSI	bis(cyanomethyl)sulfonyl)imide
S ₉	MOSI	bis(methoxysulfonyl)imide
A_1	HMSI	(hydrosulfonyl)(methylsulfonyl)imide
A ₂	FMSI	(fluorosulfonyl)(methylsulfonyl)imide
A ₃	HESI	(ethylsulfonyl)(hydrosulfonyl)imide
A_4	MESI	(ethylsulfonyl)(methylsulfonyl)imide
A_5	FPFESI	(fluorosulfonyl)((perfluoroethyl)sulfonyl)imide
A ₆	MTbSI	(tert-butylsulfonyl)(methylsulfonyl)imide
A ₇	TFNFSI	((1,1,1,3,3,3-hexafluoro-2-(trifluoromethyl)propan-2-yl)sulfonyl)((trifluoromethyl)sulfonyl)imide
A ₈	FCSI	(cyanosulfonyl)(fluorosulfonyl)imide
A9	TFCSI	(cyanosulfonyl)((trifluoromethyl)sulfonyl)imide

		N	S ₁	S ₂	01	O ₂	O ₃	O ₄	R'	R''
						(<i>e</i>)				
R_1	LiTFSI	-1.26	2.27	2.27	-0.97	-0.97	-0.96	-0.96	-0.22	-0.22
R ₂	LiFSI	-1.23	2.56	2.56	-0.96	-0.96	-0.95	-0.95	-0.53	-0.53
R ₃	LIFTFSI	-1.24	2.27	2.56	-0.96	-0.97	-0.95	-0.96	-0.53	-0.22
S_1	LiMSI	-1.26	2.32	2.32	-1.03	-1.03	-1.01	-1.01	-0.15	-0.15
S ₂	LiESI	-1.26	2.33	2.33	-1.01	-1.01	-1.03	-1.04	-0.15	-0.15
S ₃	LiBETI	-1.25	2.30	2.30	-0.96	-0.96	-0.96	-0.96	-0.26	-0.26
S ₄	LITFESI	-1.26	2.34	2.34	-0.98	-0.98	-1.01	-1.01	-0.22	-0.22
S ₅	LiTbSI	-1.30	2.35	2.35	-1.01	-1.01	-1.02	-1.02	-0.17	-0.17
S ₆	LiNFSI	-1.28	2.40	2.40	-0.96	-0.96	-0.94	-0.94	-0.36	-0.36
S ₇	LiCSI	-1.22	2.33	2.33	-0.94	-0.94	-0.94	-0.94	-0.34	-0.34
S ₈	LiCmSI	-1.25	2.35	2.35	-0.98	-0.98	-1.01	-1.01	-0.24	-0.24
S ₉	LiMOSI	-1.27	2.54	2.55	-0.98	-0.99	-0.99	-0.98	-0.44	-0.44
A ₁	Lihmsi	-1.26	2.32	2.19	-1.01	-1.00	-1.03	-1.02	-0.04	-0.15
A ₂	LiFMSI	-1.25	2.55	2.32	-0.99	-1.00	-0.97	-1.00	-0.53	-0.14
A ₃	LiHESI	-1.26	2.19	2.32	-1.00	-1.03	-1.02	-1.01	-0.04	-0.15
A ₄	LiMESI	-1.26	2.33	2.32	-1.01	-1.01	-1.03	-1.03	-0.15	-0.15
A ₅	LIFPFESI	-1.24	2.56	2.30	-0.95	-0.96	-0.95	-0.96	-0.53	-0.26
A ₆	LiMTbSI	-1.28	2.34	2.33	-1.01	-1.03	-1.02	-1.02	-0.15	-0.17
A ₇	LITFNFSI	-1.27	2.39	2.28	-0.95	-0.95	-0.96	-0.96	-0.21	-0.38
A ₈	LiFCSI	-1.22	2.33	2.56	-0.95	-0.95	-0.95	-0.94	-0.53	-0.35
A ₉	LiTFCSI	-1.23	2.33	2.26	-0.96	-0.95	-0.96	-0.94	-0.20	-0.35

Table S2 – Anion natural charge for each atom of the sulfonyl core, and sum of the charges for each functional group,calculated from the M06-2X/6-311⁺⁺G(2d,p) geometry.

Table S3 – Lithium salt natural charge for each atom of the sulfonyl core, and sum of the charges for each functional group,calculated from the M06-2X/6-311⁺⁺G(2d,p) geometry.

		N	S.	S.,	0.	0.	0.	0.	Li	R'	R"
		(e)									
R ₁	LiTFSI	-1.24	2.28	2.28	-0.90	-0.90	-1.09	-1.09	0.95	-0.14	-0.14
R_2	LiFSI	-1.21	2.59	2.59	-0.89	-0.89	-1.08	-1.08	0.96	-0.50	-0.50
R_3	LIFTFSI	-1.22	2.27	2.59	-0.89	-0.90	-1.08	-1.10	0.95	-0.49	-0.14
S_1	LiMSI	-1.25	2.33	2.30	-0.95	-0.98	-1.13	-1.12	0.95	-0.08	-0.07
S ₂	LiESI	-1.25	2.33	2.30	-0.95	-0.99	-1.13	-1.12	0.95	-0.08	-0.06
S ₃	LiBETI	-1.21	2.30	2.30	-0.89	-0.90	-1.08	-1.08	0.94	-0.19	-0.17
S_4	LITFESI	-1.22	2.34	2.34	-0.93	-0.93	-1.09	-1.09	0.91	-0.17	-0.17
S ₅	LiTbSI	-1.28	2.34	2.36	-0.98	-0.97	-1.13	-1.14	0.95	-0.07	-0.08
S_6	LiNFSI	-1.26	2.41	2.41	-0.90	-0.90	-1.09	-1.08	0.95	-0.26	-0.27
S ₇	LiCSI	-1.20	2.34	2.34	-0.88	-0.88	-1.07	-1.07	0.96	-0.27	-0.27
S ₈	LiCmSI	-1.26	2.36	2.37	-0.93	-0.92	-1.08	-1.07	0.93	-0.18	-0.21
S9	LiMOSI	-1.23	2.53	2.56	-0.91	-0.93	-1.05	-1.08	0.94	-0.45	-0.39
A ₁	Lihmsi	-1.24	2.30	2.20	-0.97	-0.93	-1.12	-1.12	0.95	0.00	-0.06
A ₂	LiFMSI	-1.23	2.58	2.33	-0.89	-0.94	-1.08	-1.12	0.95	-0.53	-0.07
A ₃	LiHESI	-1.24	2.20	2.30	-0.93	-0.98	-1.12	-1.12	0.95	0.00	-0.06
A ₄	LiMESI	-1.25	2.30	2.33	-0.95	-0.99	-1.13	-1.12	0.95	-0.08	-0.06
A ₅	LIFPFESI	-1.20	2.59	2.29	-0.88	-0.89	-1.07	-1.08	0.94	-0.50	-0.19
A ₆	LiMTbSI	-1.25	2.32	2.34	-0.95	-0.99	-1.13	-1.12	0.95	-0.09	-0.06
A ₇	LITFNFSI	-1.25	2.39	2.28	-0.90	-0.90	-1.08	-1.07	0.94	-0.13	-0.29
A ₈	LiFCSI	-1.20	2.34	2.59	-0.88	-0.88	-1.08	-1.08	0.96	-0.50	-0.27
A ₉	Litfcsi	-1.22	2.34	2.27	-0.89	-0.88	-1.09	-1.08	0.95	-0.13	-0.28

			Ν	S ₁	S ₂	01	O ₂	O ₃	O ₄	Li	R'	R''
							(e)					
Ref.	R_1	LiTFSI	0.017	0.012	0.012	0.064	0.064	-0.138	-0.138	-0.046	0.076	0.076
	R_2	LiFSI	0.024	0.031	0.031	0.073	0.073	-0.129	-0.129	-0.044	0.034	0.034
	R_3	LiFTFSI	0.025	0.003	0.030	0.068	0.069	-0.132	-0.134	-0.046	0.036	0.081
	S_1	LiMSI	0.015	0.010	-0.020	0.082	0.051	-0.123	-0.107	-0.052	0.064	0.081
	S_2	LiESI	0.016	0.008	-0.021	0.059	0.028	-0.102	-0.089	-0.053	0.068	0.086
a	S_3	LiBETI	0.039	-0.006	0.000	0.067	0.058	-0.122	-0.126	-0.062	0.070	0.083
tric	S_4	Litfesi	0.035	-0.002	-0.002	0.058	0.058	-0.084	-0.084	-0.086	0.052	0.052
.əu	S_5	LiTbSI	0.026	-0.013	0.007	0.028	0.046	-0.111	-0.118	-0.052	0.100	0.086
Symr	S_6	LiNFSI	0.018	0.009	0.009	0.051	0.052	-0.144	-0.143	-0.052	0.103	0.097
	S_7	LiCSI	0.020	0.015	0.015	0.059	0.059	-0.131	-0.131	-0.044	0.070	0.070
	S_8	LiCmSI	-0.009	0.017	0.021	0.048	0.054	-0.077	-0.061	-0.072	0.054	0.026
	S ₉	LiMOSI	0.039	-0.009	0.016	0.068	0.063	-0.061	-0.095	-0.063	-0.010	0.054
	A_1	Lihmsi	0.018	-0.019	0.012	0.036	0.067	-0.087	-0.099	-0.053	0.038	0.087
	A_2	LiFMSI	0.019	0.023	0.009	0.094	0.059	-0.113	-0.121	-0.047	0.008	0.068
a	A_3	LiHESI	0.020	0.012	-0.021	0.066	0.055	-0.100	-0.110	-0.054	0.037	0.095
tric	A_4	LiMESI	0.017	-0.022	0.009	0.060	0.028	-0.103	-0.088	-0.053	0.063	0.089
me	A_5	LiFPFESI	0.038	0.025	-0.005	0.071	0.072	-0.124	-0.117	-0.061	0.030	0.071
хл	A_6	LiMTbSI	0.024	-0.023	0.004	0.060	0.041	-0.117	-0.106	-0.053	0.062	0.107
As	A ₇	LitfnfSI	0.021	0.001	-0.002	0.045	0.043	-0.120	-0.111	-0.058	0.089	0.091
	A ₈	LiFCSI	0.020	0.011	0.031	0.071	0.065	-0.126	-0.135	-0.045	0.030	0.078
	A9	LiTFCSI	0.019	0.012	0.009	0.064	0.061	-0.130	-0.136	-0.045	0.074	0.073

Table S4 – Change in net charges (Δq) between the most stable anion and Li salt structure.

Equation S1 - Calculation of lithium metal theoretical capacity.

$$Q = \frac{(1e \times 96,485.3321 \ C.mol^{-1})}{6.94 \ g.mol^{-1} \times 3.6 \ C.mAh^{-1}} = 3,861.87 \approx 3,860 \ mAh.g^{-1}$$



Figure S1 – Conformer scan and lithiation site screening of R_1 (TFSI⁻), depicting the 5 calculated unique adsorption sites and binding energies. The most stable site is outlined in green.



Figure S2 – Conformer scan and lithiation site screening of R_2 (FSI⁻), depicting the 5 calculated unique adsorption sites and binding energies. The most stable site is highlighted in green.



Figure S3 – Conformer scan and lithiation site screening of R₃ (FTFSI⁻), depicting the 9 calculated unique adsorption sites and binding energies. The most stable site is outlined in green.



Figure S4 – Conformer scan and lithiation site screening of S_1 (MSI⁻), depicting the 5 calculated unique adsorption sites and binding energies. The most stable site is outlined in green.



Figure S5 – Conformer scan and lithiation site screening of S_2 (ESI⁻), depicting the 11 calculated unique adsorption sites and binding energies. The most stable site is outlined in green.



Figure S6 – Conformer scan and lithiation site screening of S_3 (BETI⁻), depicting the 8 calculated unique adsorption sites and binding energies. The most stable site is outlined in green.



Figure S7 – Conformer scan and lithiation site screening of S_4 (TFESI⁻), depicting the 5 calculated unique adsorption sites and binding energies. The most stable site is highlighted in green.



Figure S8 – Conformer scan and lithiation site screening of S_5 (TbSI⁻), depicting the 5 calculated unique adsorption sites and binding energies. The most stable site is highlighted in green.



Figure S9 – Conformer scan and lithiation site screening of S₆ (NFSI⁻), depicting the 9 calculated unique adsorption sites and binding energies. The most stable site is outlined in green.



Figure S10 – Conformer scan and lithiation site screening of S₇ (CSI⁻), depicting the 6 calculated unique adsorption sites and binding energies. The most stable site is outlined in green.



Figure S11 – Conformer scan and lithiation site screening of S₈ (CmSI⁻), depicting the 9 calculated unique adsorption sites and binding energies. The most stable site is outlined in green.



Figure S12 – Conformer scan and lithiation site screening of S_9 (MOSI⁻), depicting the 11 calculated unique adsorption sites and binding energies. The most stable site is outlined in green.



Figure S13 – Conformer scan and lithiation site screening of A₁ (HMSI⁻), depicting the 9 calculated unique adsorption sites and binding energies. The most stable site is outlined in green.



Figure S14 – Conformer scan and lithiation site screening of A_2 (FMSI⁻), depicting the 9 calculated unique adsorption sites and binding energies. The most stable site is highlighted in green.



Figure S15 – Conformer scan and lithiation site screening of A_3 (HESI⁻), depicting the 10 calculated unique adsorption sites and binding energies. The most stable site is outlined in green.



Figure S16 – Conformer scan and lithiation site screening of A4 (MESI-), depicting the 9 calculated unique adsorption sites and binding energies. The most stable site is outlined in green.



Figure S17 – Conformer scan and lithiation site screening of A_5 (FPFESI⁻), depicting the 13 calculated unique adsorption sites and binding energies. The most stable site is outlined in green.



Figure S18 – Conformer scan and lithiation site screening of A_6 (MTbSI⁻), depicting the 10 calculated unique adsorption sites and binding energies. The most stable site is outlined in green.



Figure S19 – Conformer scan and lithiation site screening of A₇ (TFNFSI⁻), depicting the 14 calculated unique adsorption sites and binding energies. The most stable site is outlined in green.



Figure S20 – Conformer scan and lithiation site screening of A8 (FCSI-), depicting the 10 calculated unique adsorption sites and binding energies. The most stable site is outlined in green.



Figure S21 – Conformer scan and lithiation site screening of A_9 (TFCSI⁻), depicting the 11 calculated unique adsorption sites and binding energies. The most stable site is outlined in green.



Figure S22 – Calculated electrochemical windows for the reference, symmetric and asymmetric salts.