

**Supporting Information for**

**DBMLFF: Linear scaling machine learning force fields via electron density  
decomposition for molecular electrolytes**

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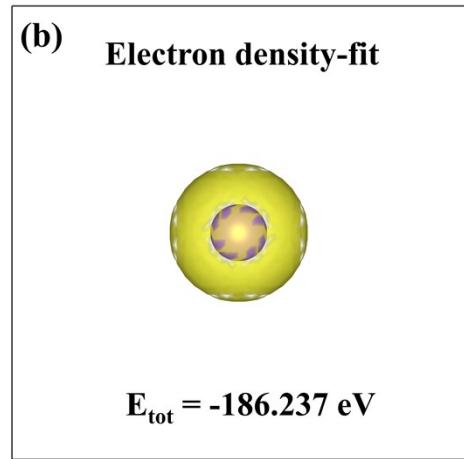
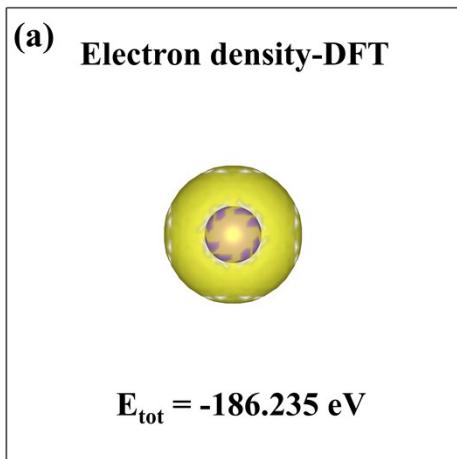
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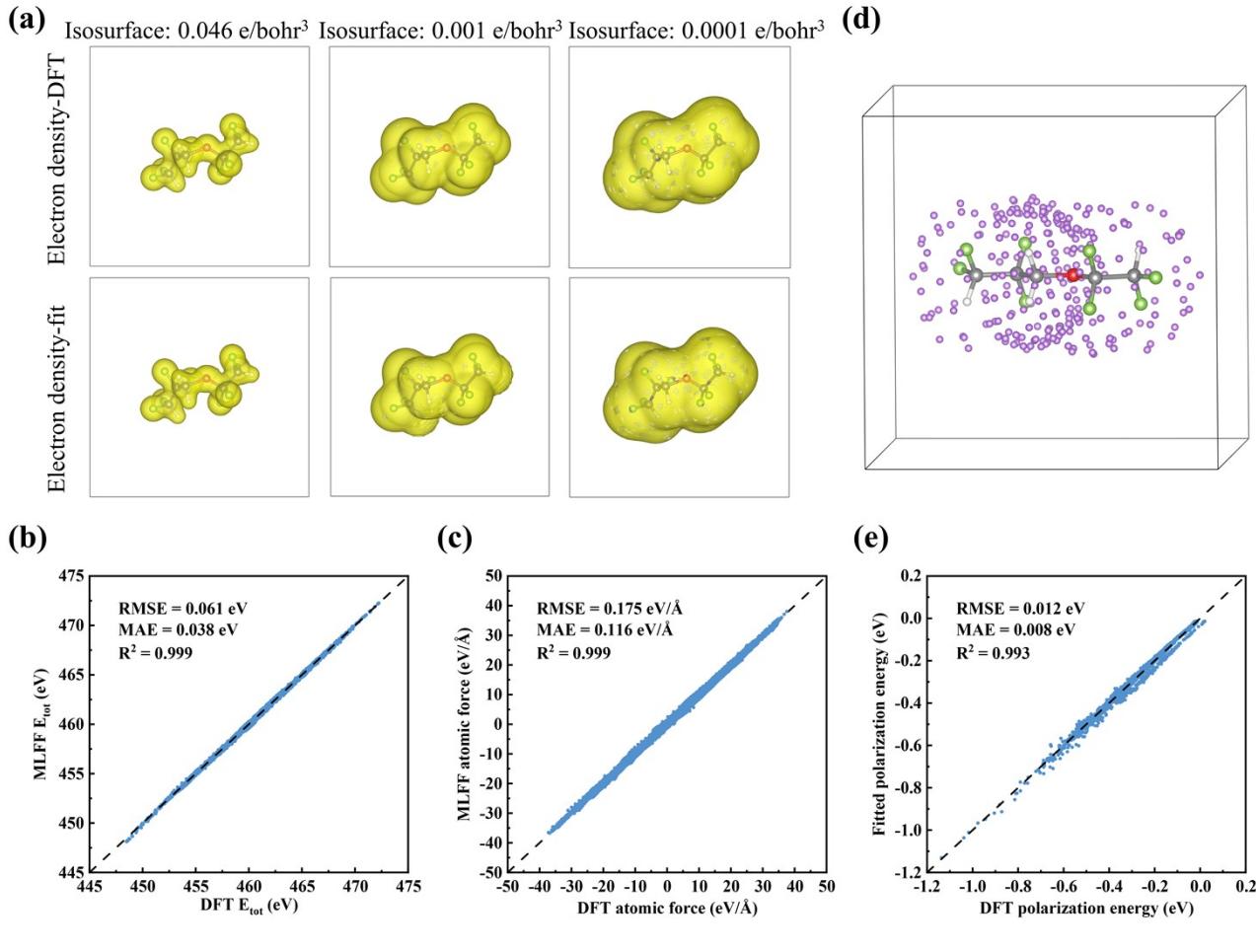
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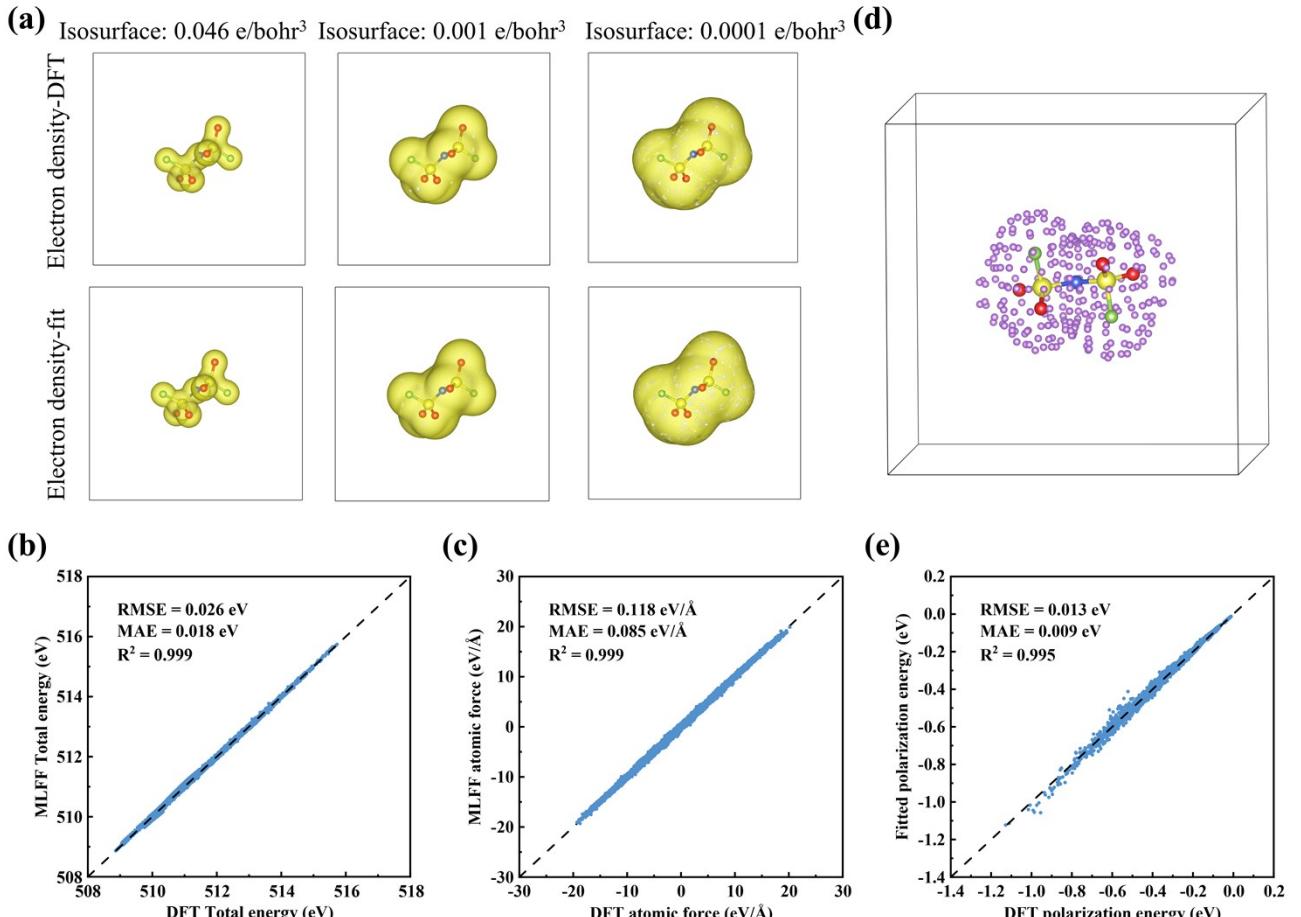
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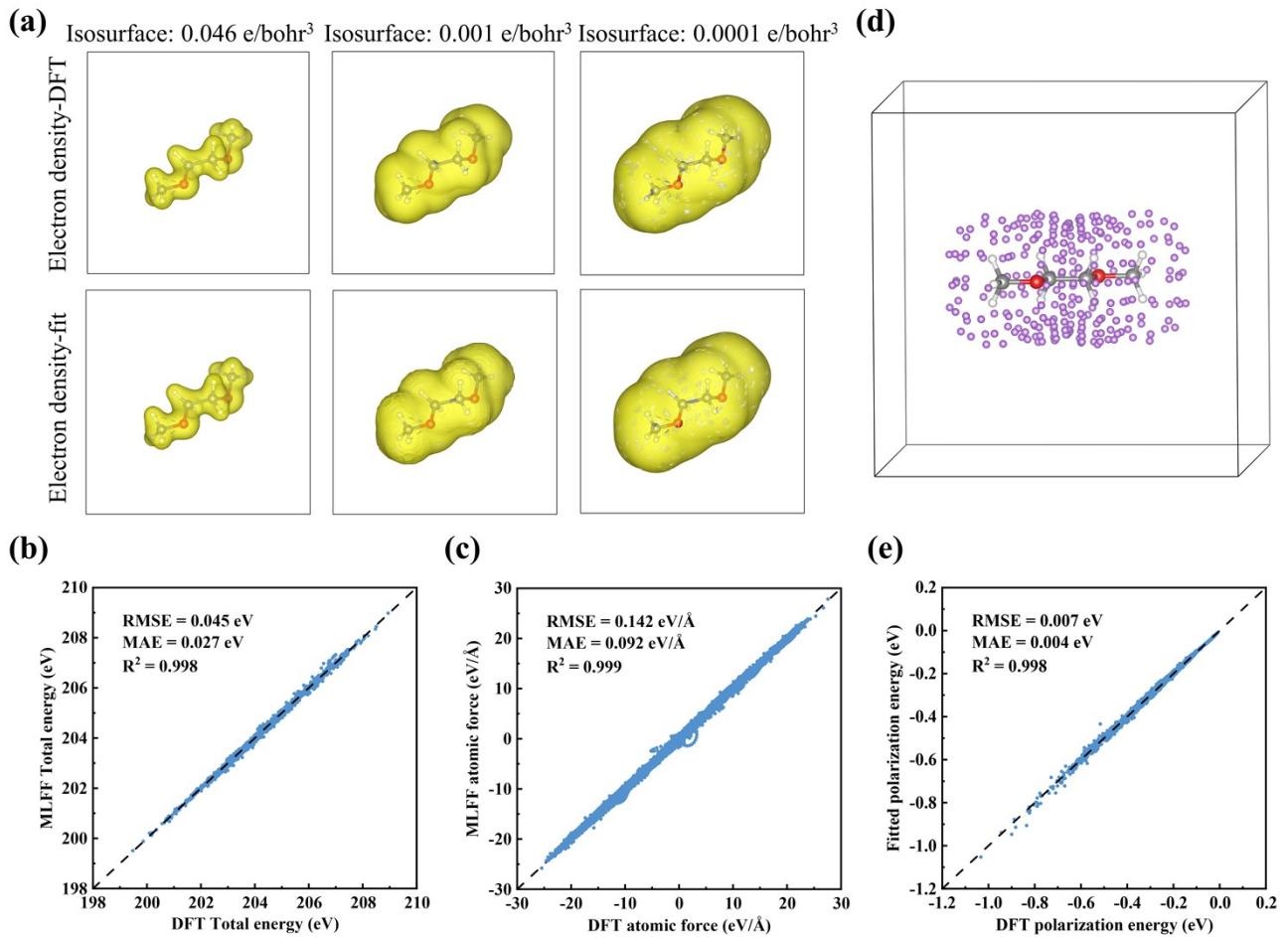
**Fig. S1** Isosurface plots and corresponding total energy values of (a) DFT-calculated and (b) spherically-fitted charge density distributions for  $\text{Li}^+$  ion with the isosurface value of  $0.00001 \text{ e/bohr}^3$ .



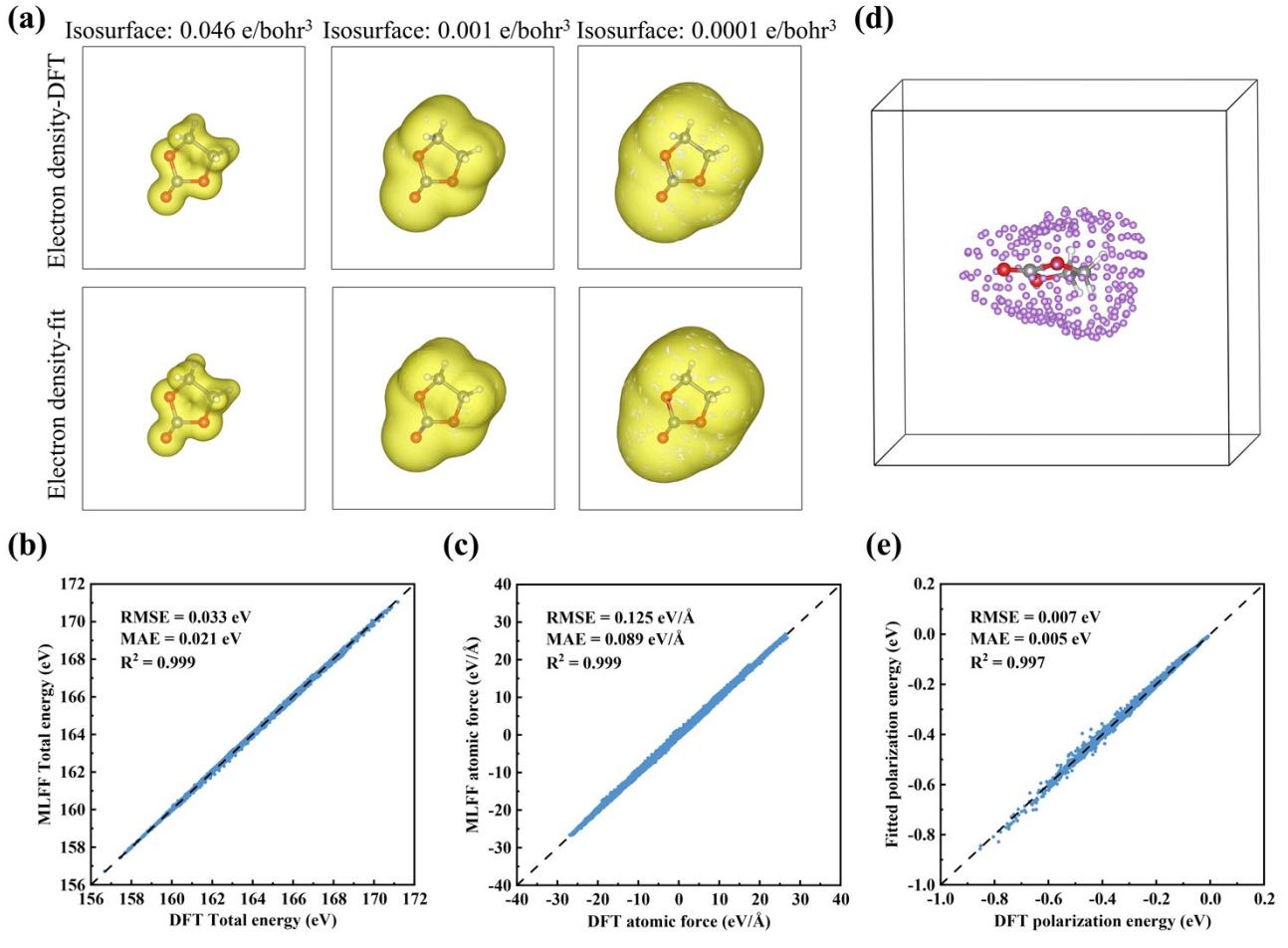
**Fig. S2** TTE molecule analysis: (a) DFT-calculated (upper) vs. spherically-fitted (lower) electron density isosurfaces at multiple values; (b) MLFF-predicted total energy and (c) atomic forces across configurations; (d) Probe charge position  $R_p$  distribution at a representative probe-molecule distance; (e) Correlation between DFT-calculated and fitted polarization energies under varied probe configurations.



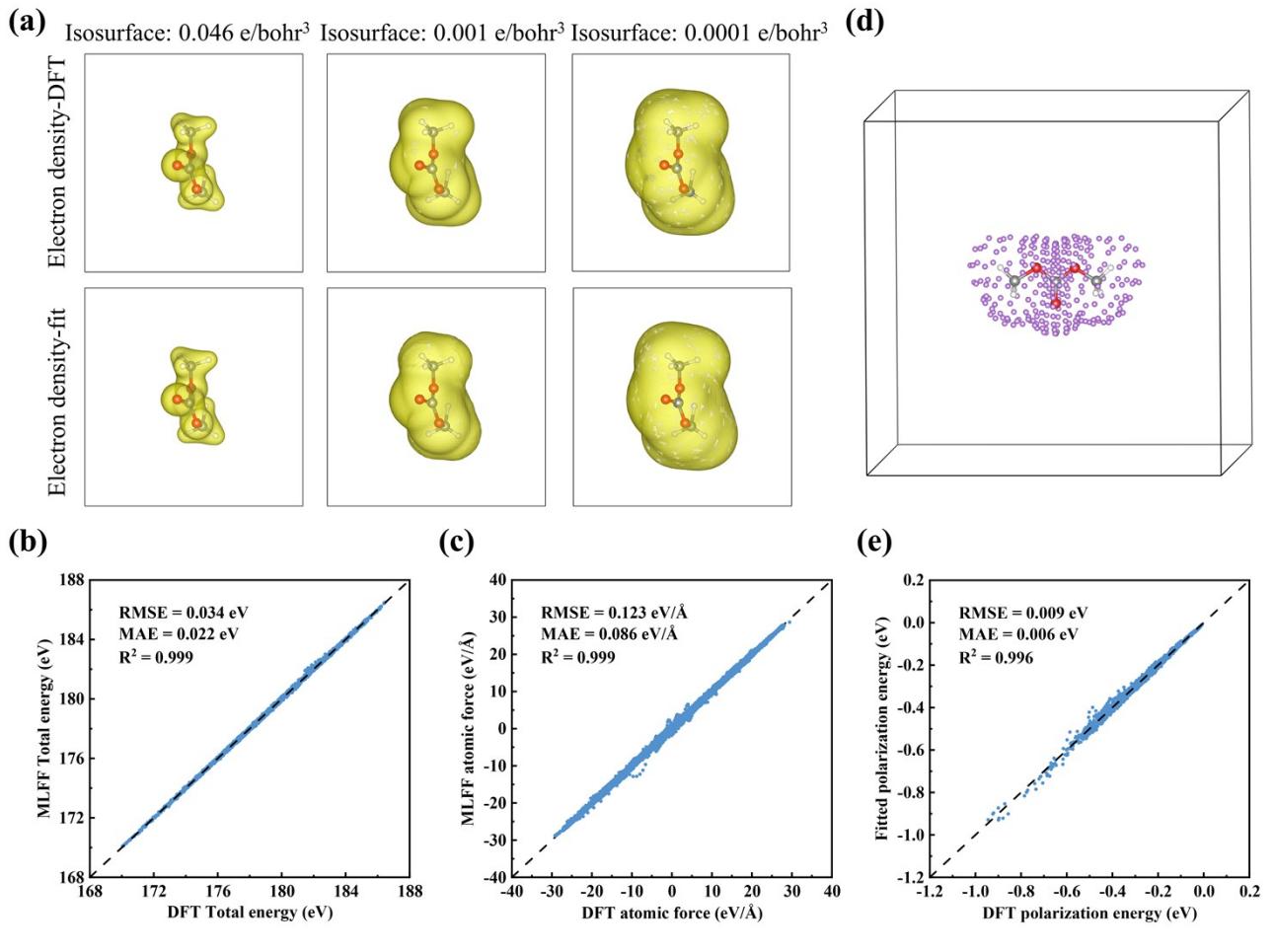
**Fig. S3** FSI<sup>-</sup> anion analysis: (a) DFT-calculated (upper) vs. spherically-fitted (lower) electron density isosurfaces at multiple values; (b) MLFF-predicted total energy and (c) atomic forces across configurations; (d) Probe charge position Rp distribution at a representative probe-molecule distance; (e) Correlation between DFT-calculated and fitted polarization energies under varied probe configurations.



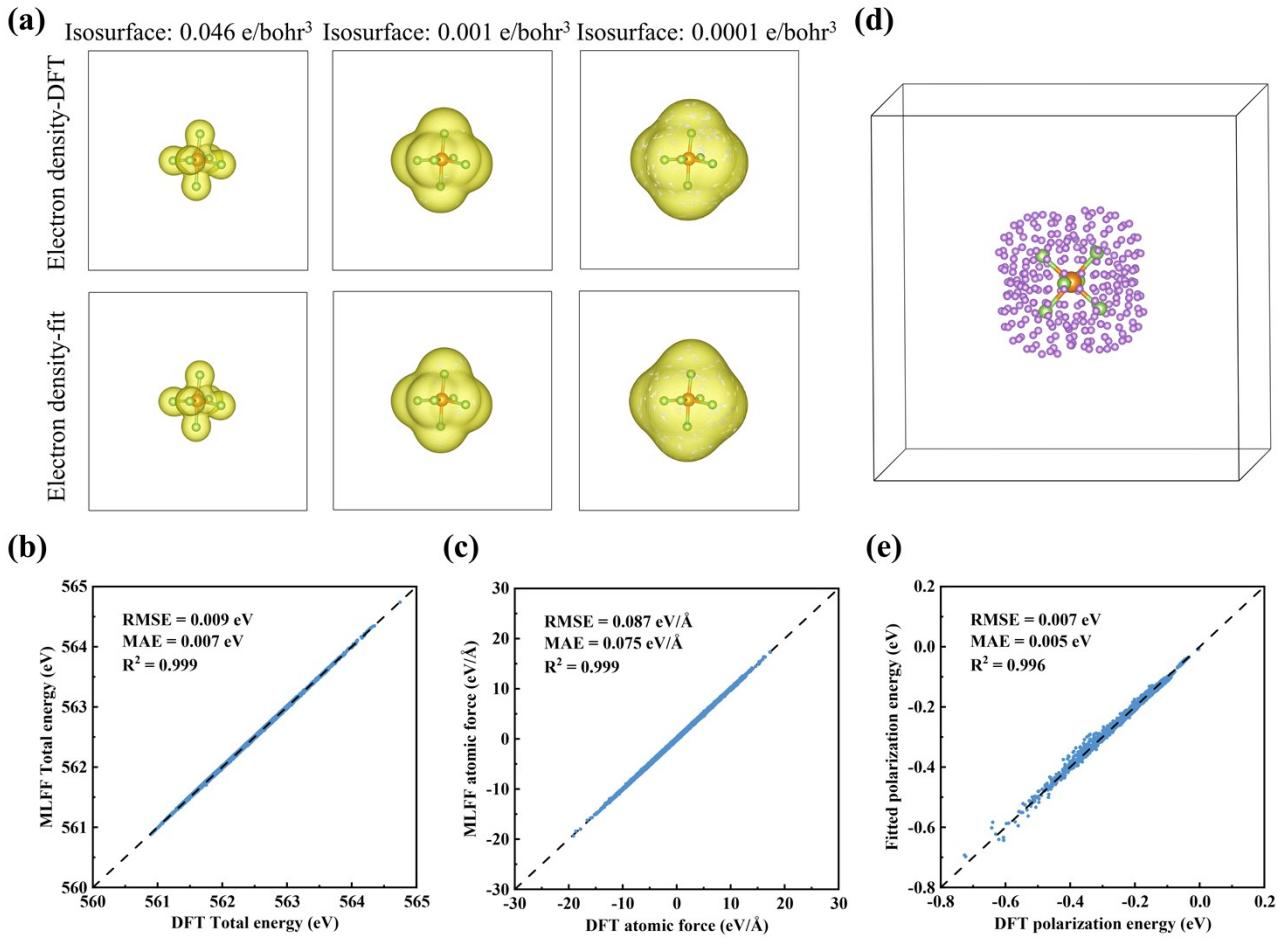
**Fig. S4** EGDME molecule analysis: (a) DFT-calculated (upper) vs. spherically-fitted (lower) electron density isosurfaces at multiple values; (b) MLFF-predicted total energy and (c) atomic forces across configurations; (d) Probe charge position  $R_p$  distribution at a representative probe-molecule distance; (e) Correlation between DFT-calculated and fitted polarization energies under varied probe configurations.



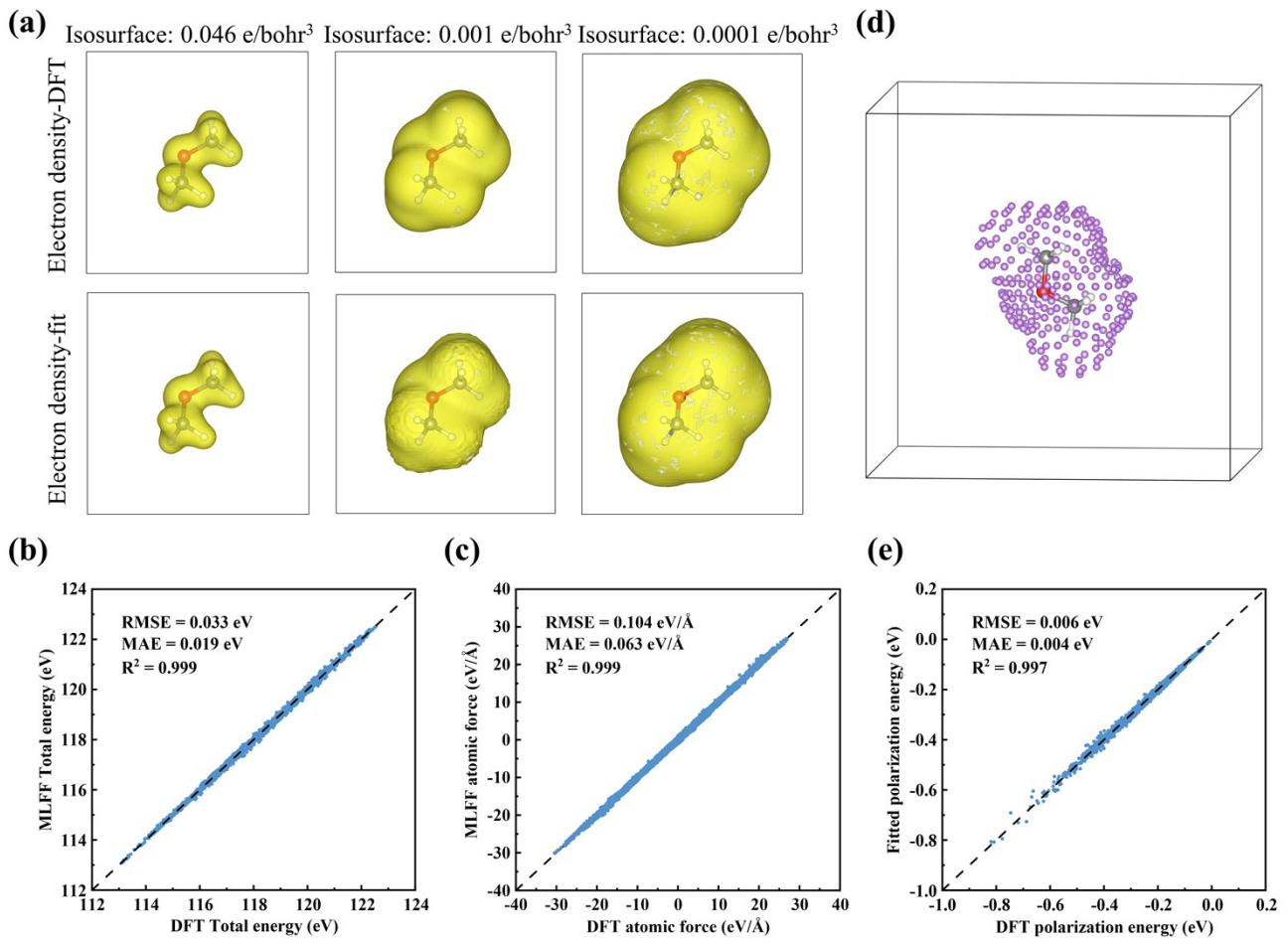
**Fig. S5** EC molecule analysis: (a) DFT-calculated (upper) vs. spherically-fitted (lower) electron density isosurfaces at multiple values; (b) MLFF-predicted total energy and (c) atomic forces across configurations; (d) Probe charge position  $R_p$  distribution at a representative probe-molecule distance; (e) Correlation between DFT-calculated and fitted polarization energies under varied probe configurations.



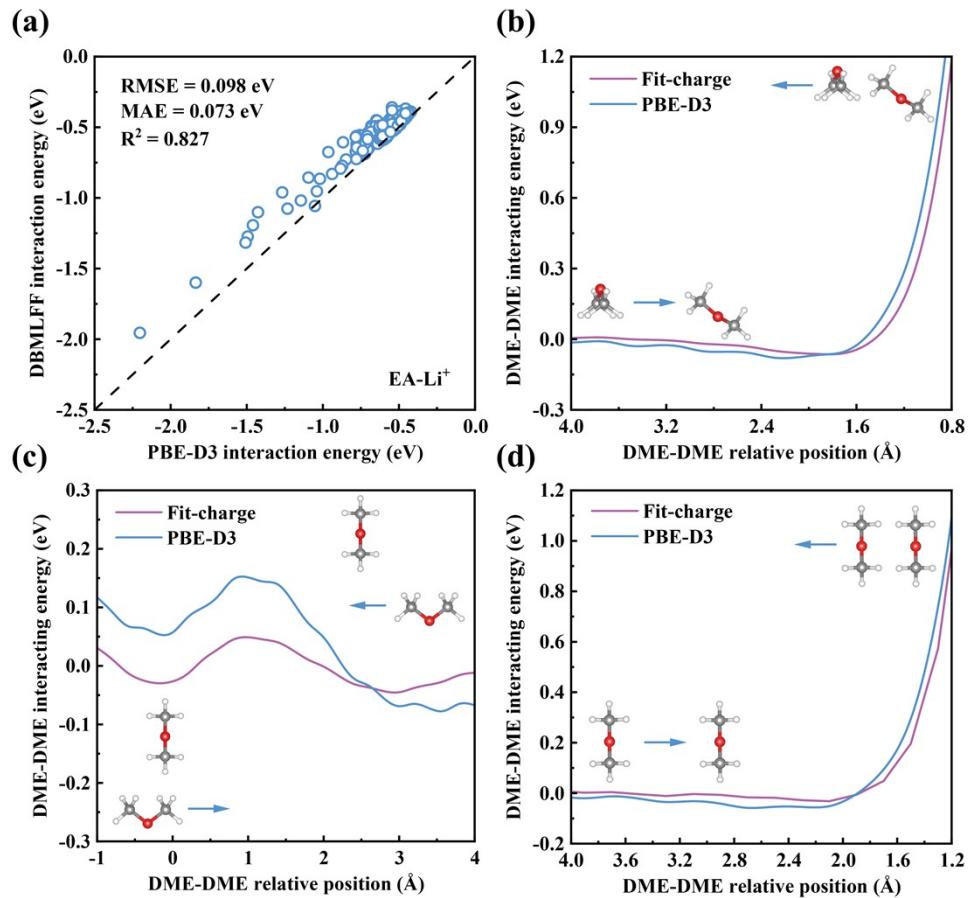
**Fig. S6** DMC molecule analysis: (a) DFT-calculated (upper) vs. spherically-fitted (lower) electron density isosurfaces at multiple values; (b) MLFF-predicted total energy and (c) atomic forces across configurations; (d) Probe charge position Rp distribution at a representative probe-molecule distance; (e) Correlation between DFT-calculated and fitted polarization energies under varied probe configurations.



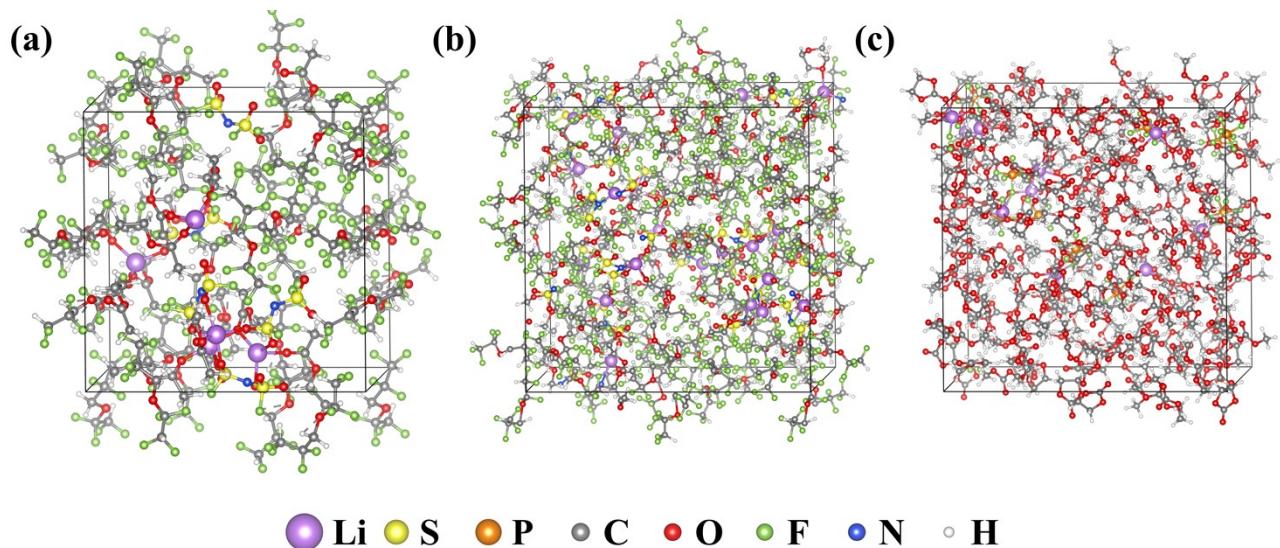
**Fig. S7**  $\text{PF}_6^-$  analysis: (a) DFT-calculated (upper) vs. spherically-fitted (lower) electron density isosurfaces at multiple values; (b) MLFF-predicted total energy and (c) atomic forces across configurations; (d) Probe charge position  $R_p$  distribution at a representative probe-molecule distance; (e) Correlation between DFT-calculated and fitted polarization energies under varied probe configurations.



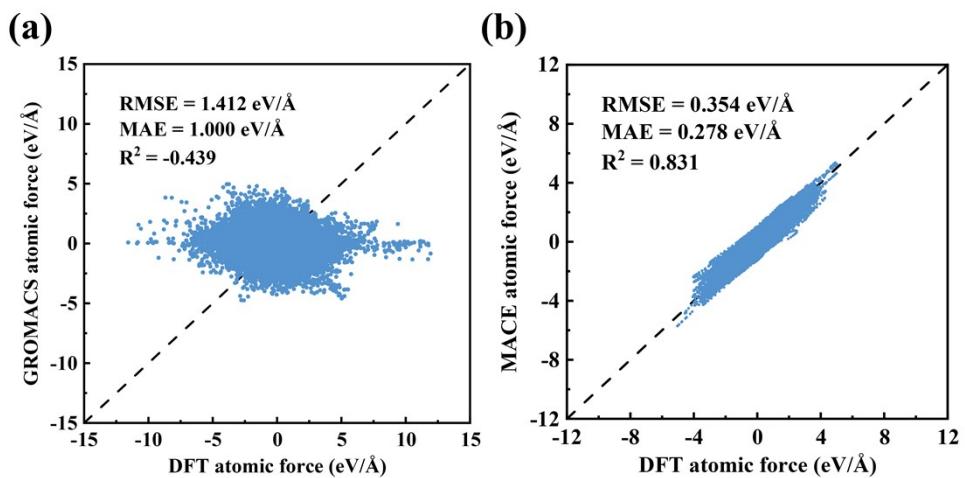
**Fig. S8** DME analysis: (a) DFT-calculated (upper) vs. spherically-fitted (lower) electron density isosurfaces at multiple values; (b) MLFF-predicted total energy and (c) atomic forces across configurations; (d) Probe charge position Rp distribution at a representative probe-molecule distance; (e) Correlation between DFT-calculated and fitted polarization energies under varied probe configurations.



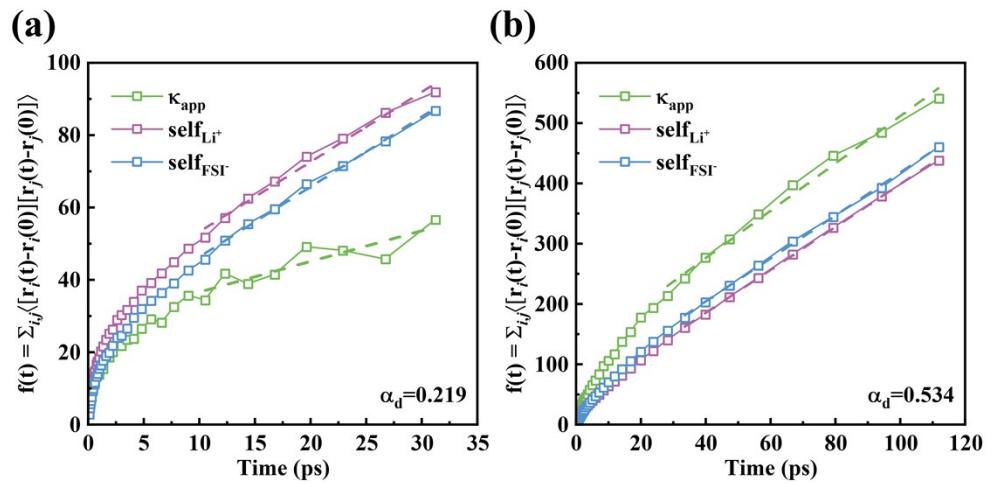
**Fig. S9** (a) Comparison of intermolecular interaction energies for the EA- $\text{Li}^+$  system between the DBMLFF model and DFT (PBE-D3) calculations, with random molecular displacements and orientations. (c)-(d) Comparison of DME-DME intermolecular interaction energies between the DBMLFF model and DFT (PBE-D3) for three representative configurations.



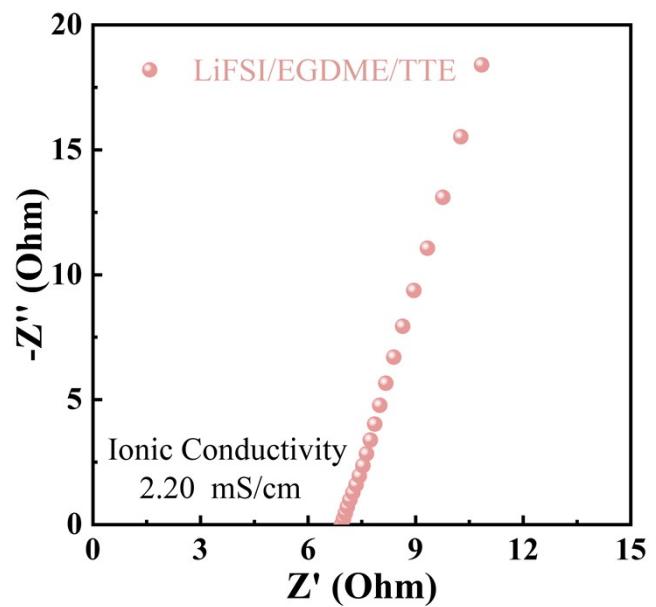
**Fig. S10** Structural snapshots of (a) the LiFSI/EA/TTE system, (b) the LiFSI/EGDME/TTE system, and (c) the LiPF<sub>6</sub>/EC/DMC system.



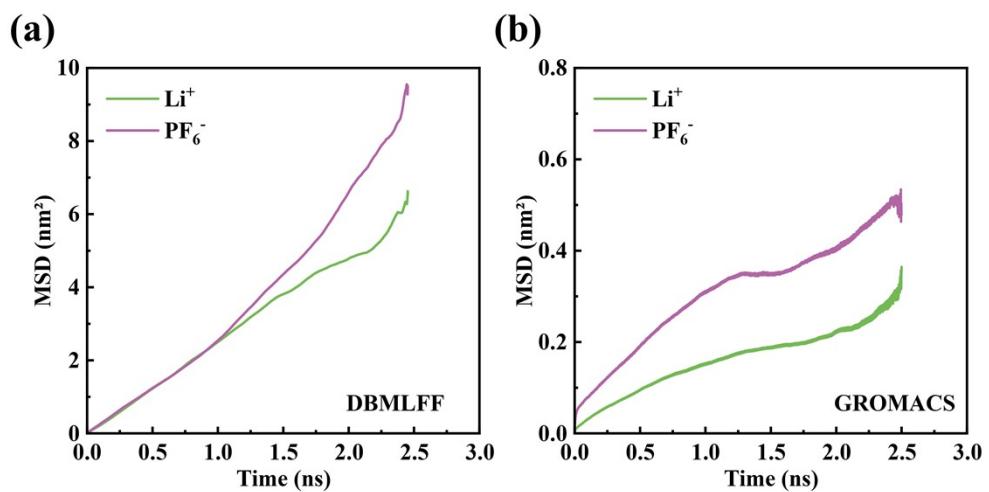
**Fig. S11** Benchmarking of simulated atomic forces against DFT for a 460-atom LiFSI/EA/TTE electrolyte system. (a) GROMACS. (b) MACE.



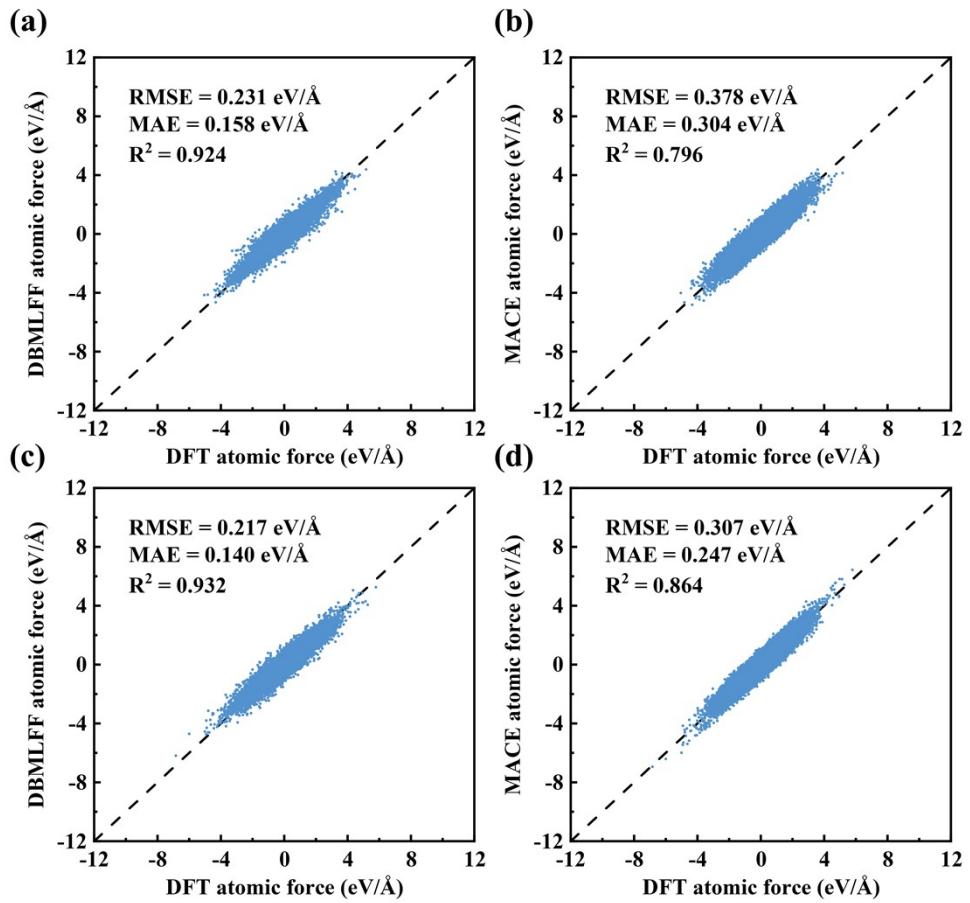
**Fig. S12** Compute  $\alpha(t)$  using the initial 4% segment of the MD trajectory for (a) LiFSI/EGDME/TTE and (b) LiPF<sub>6</sub>/EC/DMC electrolyte systems. Here,  $\kappa_{app}$  is the time correlation function for collective ion conduction;  $self_{Li^+}$  and  $self_{FSI^-}$  denote the self-diffusion correlation functions for Li<sup>+</sup> and FSI<sup>-</sup>, respectively.



**Fig. S13** Nyquist plot of the LiFSI/EGDME/TTE electrolyte system, showing an ionic conductivity of 2.20 mS/cm.



**Fig. S14** Mean square displacement (MSD) of  $\text{Li}^+$  and  $\text{PF}_6^-$  in the  $\text{LiPF}_6/\text{EC/DMC}$  system from (a) DBMLFF and (b) GROMACS simulations.



**Fig. S15** Benchmarking of ML-predicted atomic forces against reference DFT calculations. (a, b) Forces in the LiFSI/EGDME/TTE system predicted by (a) the DBMLFF and (b) the MACE model. (c, d) Forces in the LiPF<sub>6</sub>/EC/DMC system predicted by (c) the DBMLFF and (d) the MACE model.

Table S1. Diffusion coefficients and ionic conductivities ( $\kappa$ ) in the LiFSI/EGDME/TTE system, determined using various techniques.

Methods	$D_{\text{Li}^+}$ ( $10^{-10} \text{ m}^2 \text{ s}^{-1}$ )	$D_{\text{FSI}^-}$ ( $10^{-10} \text{ m}^2 \text{ s}^{-1}$ )	$\kappa$ ( $\text{mS cm}^{-1}$ )
GROMACS	0.486	0.704	1.511
DBMLFF	0.849	1.030	2.386
Experiments	1.377	1.180	2.200

Table S2. Diffusion coefficients and ionic conductivities ( $\kappa$ ) in the LiPF<sub>6</sub>/EC/DMC system, determined using various techniques.

Methods	$D_{\text{Li}^+}$ ( $10^{-10} \text{ m}^2 \text{ s}^{-1}$ )	$D_{\text{PF}_6^-}$ ( $10^{-10} \text{ m}^2 \text{ s}^{-1}$ )	$\kappa$ ( $\text{mS cm}^{-1}$ )
GROMACS	0.211	0.422	1.393
DBMLFF	3.581	4.205	13.800
Experiments	1.8	2.7	11.2

Table S3. Enthalpies of vaporization of different substances at 298 K and 1 atm.

Molecule	GROMACS (kJ·mol <sup>-1</sup> )		DBMLFF (kJ·mol <sup>-1</sup> )		Experiments (kJ·mol <sup>-1</sup> )
	Mean	SE	Mean	SE	
DME	26.37	0.01	23.40	0.03	21.2 <sup>1</sup>
EA	53.87	0.03	38.91	0.02	35.69 <sup>2</sup>
TTE	44.79	0.05	36.21	0.09	40.2 <sup>3</sup>
EGDME	50.08	0.39	38.36	0.06	36.76 <sup>4</sup>
DMC	62.92	0.02	43.75	0.09	37.70 <sup>5</sup>

## References

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