## Supporting information

Structure-property relationships in organic battery anode materials: exploring redox reactions in crystalline Na- and Li-benzene diacrylate using combined crystallography and density functional theory calculations

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 Table S1 Space group of all crystal structures as theoretically predicted.

	Na <sub>2</sub> BDA	Li <sub>2</sub> BDA
Pristine	$P2_1/c$	$P2_1/c$
1 <sup>st</sup> ion insertion	P1	Pl
2 <sup>nd</sup> ion insertion	$P2_1/c$	PĪ

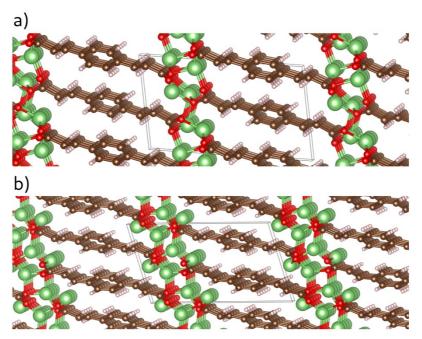
*Table S2* Lattice parameters of all crystal structures as theoretically predicted.

	Na <sub>2</sub> BDA	Na <sub>3</sub> BDA	Na <sub>4</sub> BDA	Li <sub>2</sub> BDA	Li <sub>3</sub> BDA	Li <sub>4</sub> BDA
a (Å)	13.77	15.02	14.03	11.11	13.03	13.33
b (Å)	5.34	5.39	5.11	5.36	5.13	5.28
c (Å)	6.89	6.72	8.09	8.28	7.57	7.59
$\alpha$ (deg)	90.01	90.01	90.05	89.97	86.84	86.40
$\beta$ (deg)	97.31	78.05	76.64	90.05	100.15	107.03
Y (deg)	89.95	96.09	90.05	87.15	86.80	101.05

*Table S3* Comparison of experimental data between the PXRD-Na<sub>2</sub>BDA and SCXRD-Li<sub>2</sub>BDA.

Parameter	Na <sub>2</sub> BDA	Li <sub>2</sub> BDA
Molar mass (g/mol)	262.02	230.07
Temp of measurement (K)	293	293
Space group	$P 2_{1}/c$	$P 2_{1}/c$
<i>a</i> axis (Å)	14.53	13.75
Cell volume (Å <sup>3</sup> )	526.35	1170.99
Z	2	4
Calc. density (g/cm <sup>3</sup> )	1.65429	1.39586
Crystal colour	white	white

The predicted lithiated phases for the Li<sub>2</sub>BDA compound are presented on the Figure S1.



*Figure S1* The predicted crystal structure for the lithiated phases a)  $Li_3BDA$  and b)  $Li_4BDA$ .

## **3D** electron diffraction

The sample was crushed in an agate mortar, dispersed in absolute ethanol. A droplet of the suspension was transferred onto a copper grid covered by a holey carbon film. The 3D electron diffraction data were collected using a JEOL JEM-2100 transmission electron microscope (TEM) operated at an accelerating voltage of 200 kV. The data were collected by continuously tilting the goniometer with a tilt speed of 0.28 °/s. During tilting the crystal was tracked by sequential defocusing of the intermediate lens using the software Instamatic.<sup>1</sup> The data were collected using X-ray Detector Software (XDS)<sup>2</sup>. The structure was solved using the software SHELXT<sup>3</sup>, the positions of all non-hydrogen atoms were directly obtained with reasonable geometries. A least-squares refinement was performed in SHELXL<sup>4</sup> using atomic scattering factors for electrons extracted from SIR2014<sup>5</sup>. The refinement converged with an R1 of 35% without the use of restraints. Details regarding the data and structure refinement can be found in Table S4.

Crystal system	Monoclinic
Space group	<i>P</i> 2 <sub>1</sub> / <i>c</i> (No. 14)
a, Å	14.49
b <i>,</i> Å	5.50
c, Å	7.21
α, °	90
β, °	101.1
γ, °	90
Volume, Å <sup>3</sup>	563.9
λ <i>,</i> Å	0.0251
Exposure time per frame (s)	0.6
Tilt range, °	-58.3 – +62.3
Resolution, Å	0.9
Completeness, %	77.2
R <sub>int</sub>	0.1142
R1	0.3535
No. of symmetry	617
independent reflections	
Restraints	0

Table S4. Details of the electron diffraction data and refinement.

## References

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