

1 Supporting Information of
2 Computational Investigation of Geometrical Effect in 2D Boron
3 Nitride Nanopores for DNA Detection

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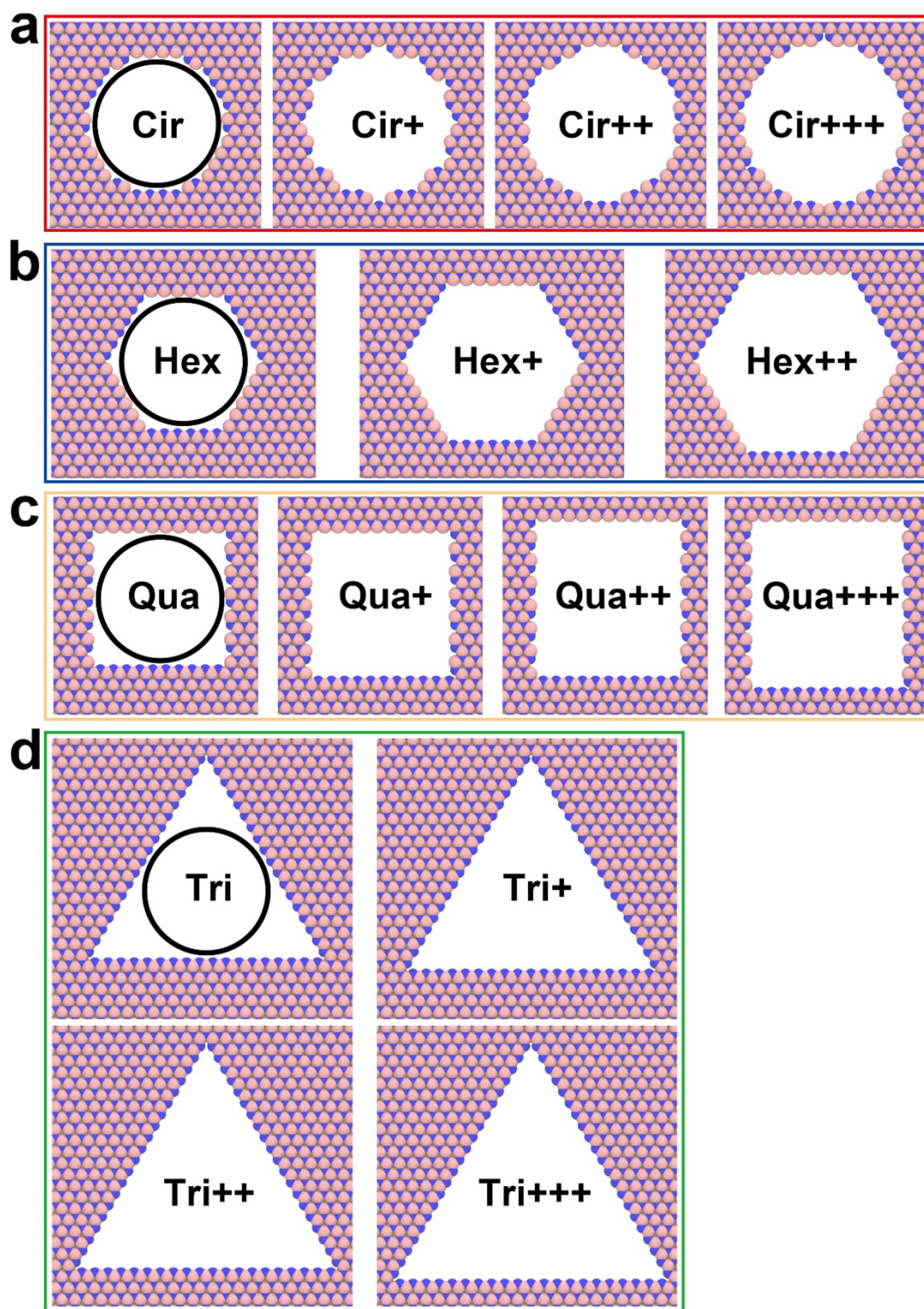
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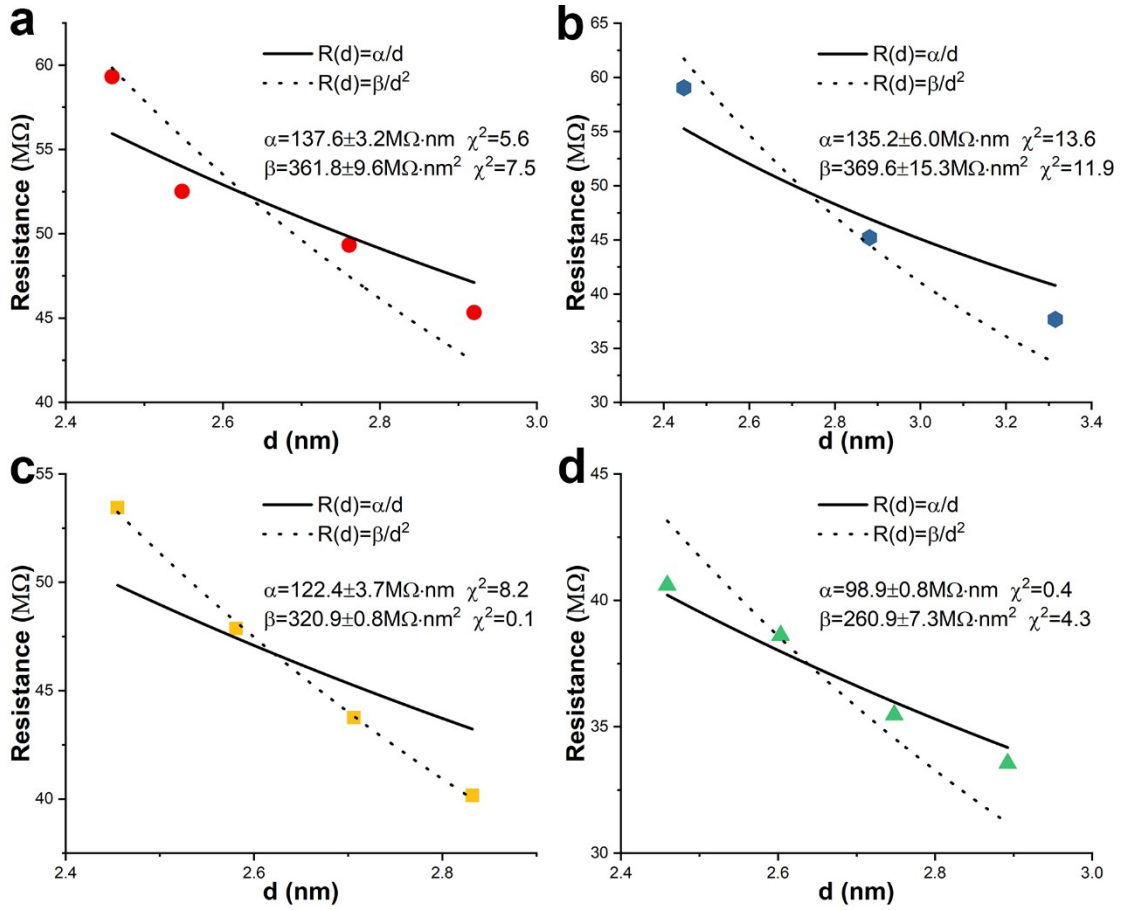
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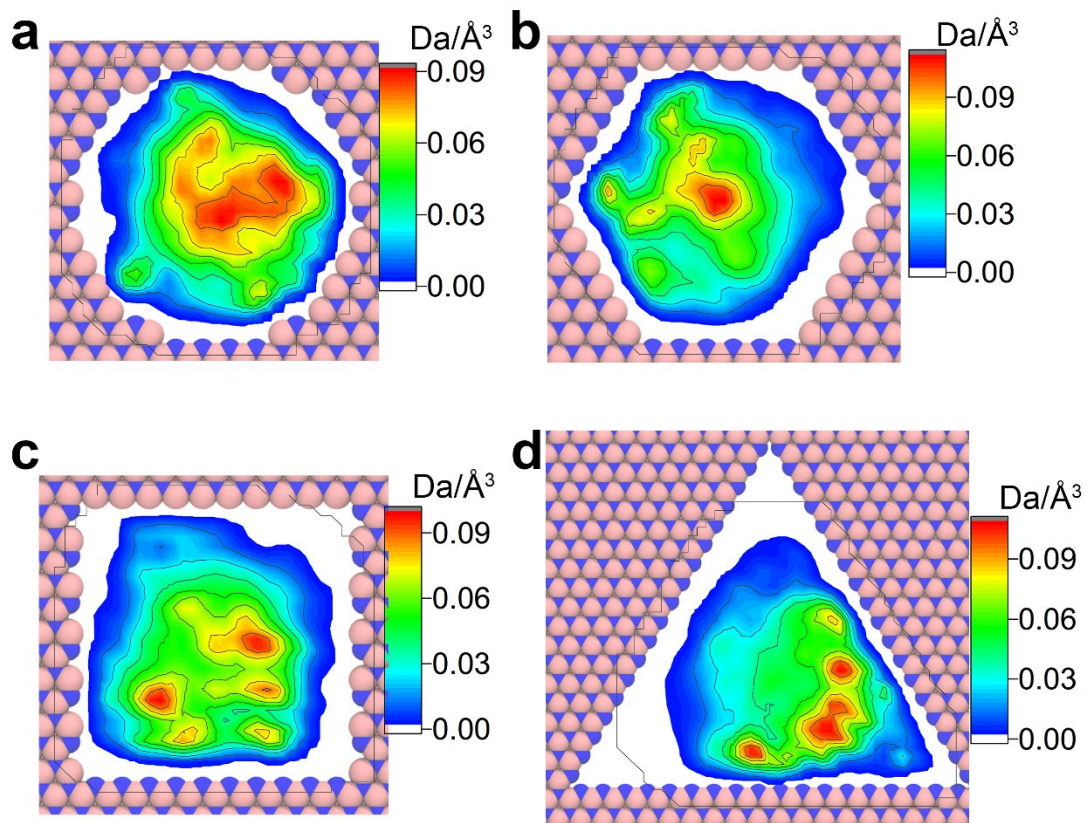
2 **Figure S1. Atomic models of BN nanopore.** Atomic models of BN nanopore with circular (a),
 3 hexagonal (b), quadrangular (c) and triangular (d) shapes investigated in this study. The solid
 4 circles in the Cir, Hex, Qua and Tri systems indicate the minimum inscribed circles with same
 5 2.46 nm effective diameter, which is defined as the distance between the pore atoms at the diameter
 6 minus the van der Waals radius of boron and nitrogen (~ 0.3 nm). More geometrical information
 7 of the nanopore is list in table S1.



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2 **Figure S2. Ionic resistance for circular (a), hexagonal (b), quadrangular (c) and triangular**
 3 **(d) shape nanopores of various size, fitted according to two functional dependences**
 4 **$R(d)=\alpha/d$ and $R(d)=\beta/d^2$. The effective diameter (d) of the minimum inscribed circle is used for**
 5 **the hexagonal, quadrangular and triangular shape nanopores.**

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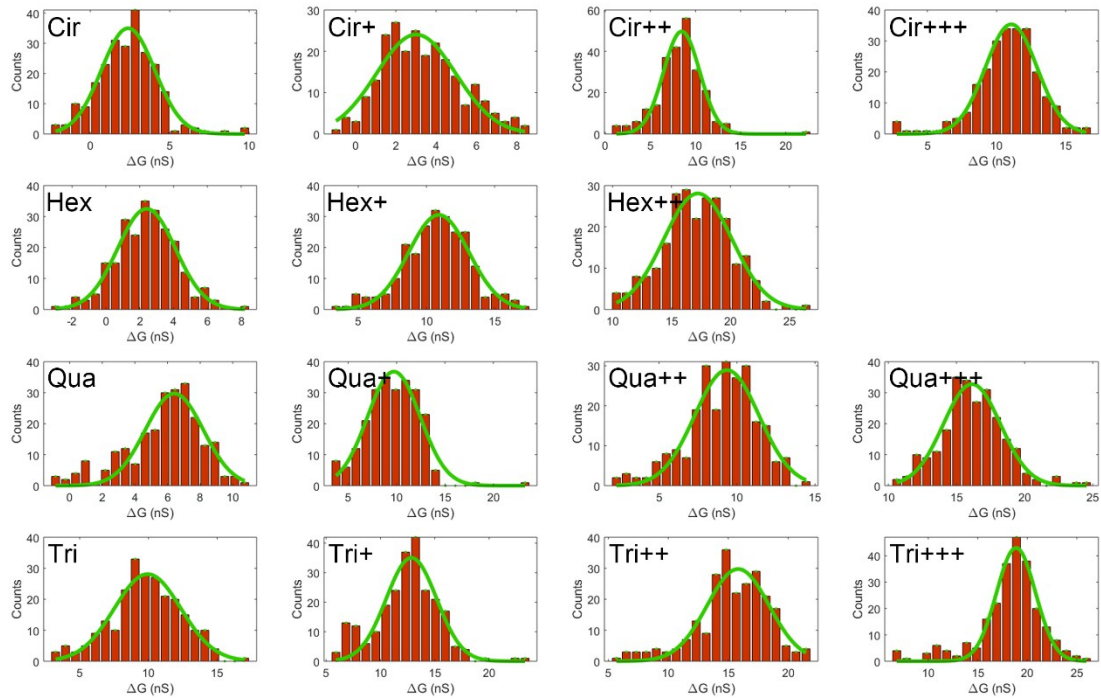
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3 **Figure S3. Mass density of DNA for the Cir (a), Hex (b), Qua (c) and Tri (d) systems.**

4 Heavy atoms of DNA molecule within 6 Å of the BN membrane are counted.

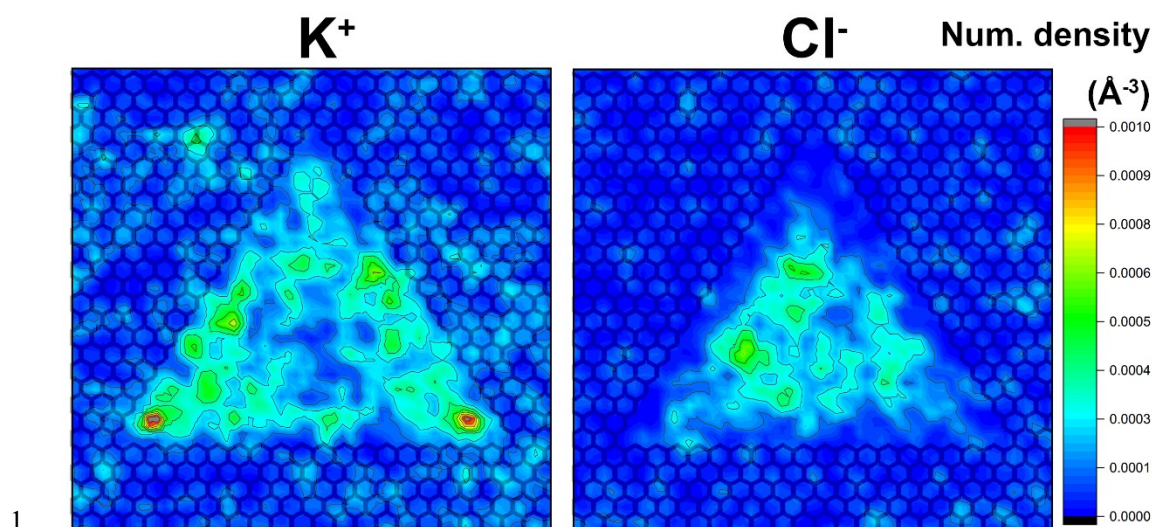
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2 **Figure S4. Histograms of conductance blockades for all the systems with dsDNA**
 3 **translocating through the nanopores. Gaussian fit (Green solid line) was performed to obtain**
 4 **the expectation of every distribution.**

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Table S1. Geometrical information of BN nanopores

<i>Nanopore geometry</i>	Diameter d (nm)	Area A (nm ²)
<i>Cir</i>	2.46	4.75
<i>Cir+</i>	2.55	5.10
<i>Cir++</i>	2.76	5.99
<i>Cir+++</i>	2.92	6.70
<i>Hex</i>	2.46*	5.19
<i>Hex+</i>	2.88*	7.19
<i>Hex++</i>	3.32*	9.52
<i>Qua</i>	2.46*	6.36
<i>Qua+</i>	2.58*	7.06
<i>Qua++</i>	2.71*	7.80
<i>Qua+++</i>	2.83*	8.98
<i>Tri</i>	2.46*	7.85
<i>Tri+</i>	2.60*	8.80
<i>Tri++</i>	2.75*	9.81
<i>Tri+++</i>	2.89*	10.87

2 * List is the effective diameter of the minimum inscribed circle.

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Table S2. Information of systems simulated

<i>Nanopore geometry</i>	<i>DNA (bp)</i>	<i>No. of atoms</i>	<i>Simulation duration (ns)</i>
<i>Cir</i>	n/a	193815	27
<i>Cir</i>	[A-T] ⁴⁵	208847	60*
<i>Cir</i>	[C-G] ⁴⁵	208994	75*
<i>Cir+</i>	n/a	205491	27
<i>Cir+</i>	[A-T] ⁴⁵	204680	90*
<i>Cir+</i>	[C-G] ⁴⁵	204992	65*
<i>Cir++</i>	n/a	205482	27
<i>Cir++</i>	[A-T] ⁴⁵	204635	70*
<i>Cir++</i>	[C-G] ⁴⁵	204935	60*
<i>Cir+++</i>	n/a	205482	27
<i>Cir+++</i>	[A-T] ⁴⁵	204647	70*
<i>Cir+++</i>	[C-G] ⁴⁵	204761	65*
<i>Hex</i>	n/a	193809	27
<i>Hex</i>	[A-T] ⁴⁵	208787	70*
<i>Hex</i>	[C-G] ⁴⁵	209012	70*
<i>Hex+</i>	n/a	205473	27
<i>Hex+</i>	[A-T] ⁴⁵	204677	70*
<i>Hex+</i>	[C-G] ⁴⁵	208790	75*
<i>Hex++</i>	n/a	205479	27
<i>Hex++</i>	[A-T] ⁴⁵	208787	60*
<i>Hex++</i>	[C-G] ⁴⁵	208796	70*
<i>Qua</i>	n/a	193815	27
<i>Qua</i>	[A-T] ⁴⁵	208718	75*
<i>Qua</i>	[C-G] ⁴⁵	208904	65*
<i>Qua+</i>	n/a	209501	27
<i>Qua+</i>	[A-T] ⁴⁵	208663	75*
<i>Qua+</i>	[C-G] ⁴⁵	208828	75*

<i>Qua++</i>	n/a	209510	27
<i>Qua++</i>	[A-T] ⁴⁵	208732	60*
<i>Qua++</i>	[C-G] ⁴⁵	208744	60*
<i>Qua+++</i>	n/a	209508	27
<i>Qua+++</i>	[A-T] ⁴⁵	208721	60*
<i>Qua+++</i>	[C-G] ⁴⁵	208739	75*
<i>Tri</i>	n/a	205488	27
<i>Tri</i>	[A-T] ⁴⁵	208718	90*
<i>Tri</i>	[C-G] ⁴⁵	208748	60*
<i>Tri+</i>	n/a	205499	27
<i>Tri+</i>	[A-T] ⁴⁵	208777	65*
<i>Tri+</i>	[C-G] ⁴⁵	208795	70*
<i>Tri++</i>	n/a	205472	27
<i>Tri++</i>	[A-T] ⁴⁵	208834	80*
<i>Tri++</i>	[C-G] ⁴⁵	208816	75*
<i>Tri+++</i>	n/a	205500	27
<i>Tri+++</i>	[A-T] ⁴⁵	208871	80*
<i>Tri+++</i>	[C-G] ⁴⁵	208757	60*

1 *Simulations with three replicas (refer to simulation S1, S2 and S3).

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