

## Supplementary Material

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## SUPPLEMENTARY MATERIAL 1: TWINNING CONFIGURATIONS

Fig. 1 shows the fully-relaxed atomic configuration of the twins (15,8), (19,8), (19,12) and (19,16), as reported in Arca *et al.*<sup>1</sup>.

## SUPPLEMENTARY MATERIAL 2: TRANSPORT CALCULATION METHODOLOGY

### Landauer-Büttiker formalism

For completeness and ease of reference, in this section we briefly summarize the basis of the Landauer-Büttiker formalism (cf. Datta<sup>2,3</sup>). The tight-binding Hamiltonian matrix (TBHM) of the model sketched in Fig. 2 can be written in terms of submatrices as

$$\mathbf{H} = \begin{pmatrix} \mathbf{H}_L & \mathbf{H}_{LC} & \mathbf{0} \\ \mathbf{H}_{CL} & \mathbf{H}_C & \mathbf{H}_{CR} \\ \mathbf{0} & \mathbf{H}_{RC} & \mathbf{H}_R \end{pmatrix}, \quad (1)$$

where  $\mathbf{H}_C$  corresponds to the central component,  $\mathbf{H}_{L(R)}$  corresponds to the isolated semi-infinite left (right) component and  $\mathbf{H}_{L(R)C}$  represents the coupling between the left (right) and central regions of the model. Within the TB formalism, the empirical TBHM follows as

$$H_{l\alpha,j\beta} = E_\alpha \delta_{lj} \delta_{\alpha\beta} + h_{l\alpha,j\beta} (1 - \delta_{lj} \delta_{\alpha\beta}), \quad (2)$$

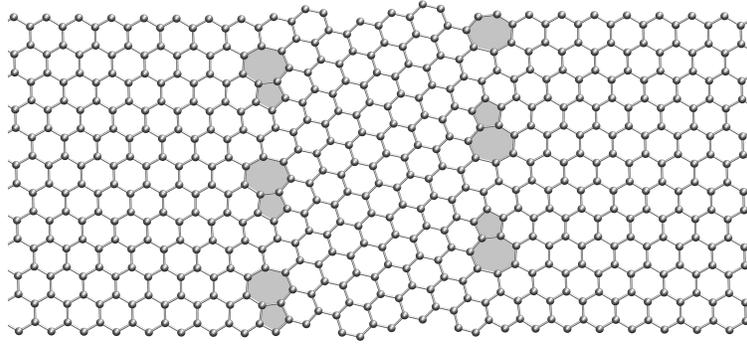
where  $l$  and  $j$  label atomic sites,  $\alpha$  and  $\beta$  label atomic orbitals, and  $E_\alpha$  and  $h_{l\alpha,j\beta}$  are the on-site parameters and hopping functions, respectively.

Bloch's theorem is applicable to periodic systems such as those considered in this work<sup>4</sup>. In  $\mathbf{k}$ -space, the sizes of the submatrices  $\mathbf{H}_C(\mathbf{k})$ ,  $\mathbf{H}_{L(R)}(\mathbf{k})$  and  $\mathbf{H}_{L(R)C}(\mathbf{k})$  are  $[N_c \times N_c]$ ,  $[\infty \times \infty]$  and  $[\infty \times N_c]$ , respectively, where  $N_c$  is the number of atoms in the central region.

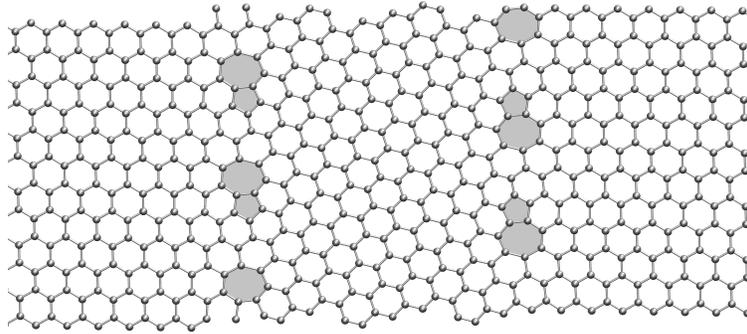
Following Datta<sup>2,3</sup>, the transmission function  $T$  across the central region at energy  $E$  can be written in the form

$$T(E) = \frac{1}{N} \sum_{\mathbf{k}} Tr[\Gamma_L(E, \mathbf{k}) \mathbf{G}_C^\dagger(E, \mathbf{k}) \Gamma_R(E, \mathbf{k}) \mathbf{G}_C(E, \mathbf{k})], \quad (3)$$

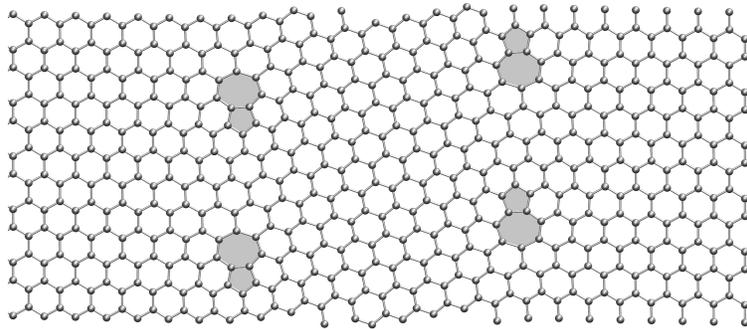
where  $\Gamma_L(E, \mathbf{k})$  and  $\Gamma_R(E, \mathbf{k})$  are the coupling  $[N_c \times N_c]$  matrices, accounting for the coupling between the central and the left (right) regions, and  $\mathbf{G}_C$  is Green's matrix of the central region. We recall that the transmission function provides the rate at which electrons are transmitted across the



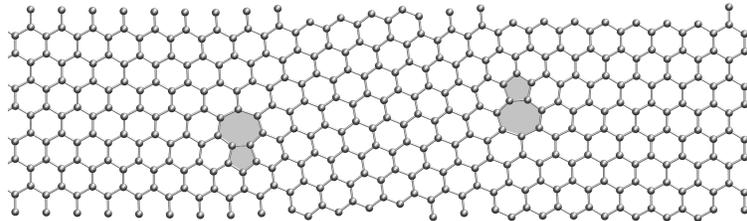
(a)



(b)



(c)



(d)

FIG. 1: Relaxed atomic configurations of the dislocation structures considered in this work: (a) (15,8), (b) (19,8), (c) (19,12), and (d) (19,16).

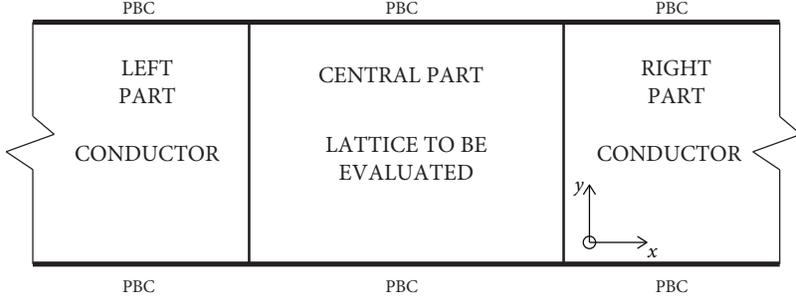


FIG. 2: Geometry of the model used to compute charge carrier transmission across a lattice.

central region. For economy of notation, henceforth we drop the  $k$ -dependence of the submatrices. The central region Green's matrix,  $\mathbf{G}_C$ , follows readily as

$$\mathbf{G}_C(E) = [E^+ \mathbb{I} - \mathbf{H}_C - \Sigma_L(E) - \Sigma_R(E)]^{-1} \approx [E \mathbb{I} - \mathbf{H}_C - \Sigma_L(E) - \Sigma_R(E)]^{-1}, \quad (4)$$

where  $E^+ = E + i\eta$ , with  $\eta \rightarrow 0^+$ ,  $\Sigma_L$  and  $\Sigma_R$  are the self-energy  $[N_c \times N_c]$  matrices corresponding to the left and right regions, respectively, which describe the effect of these regions on the electronic structure of the central region. We have,

$$\Sigma_l(E) = \mathbf{H}_{Cl} \mathbf{G}_l(E) \mathbf{H}_{lC}, \quad l = L, R, \quad (5)$$

where

$$\mathbf{G}_l(E) = (E^+ \mathbb{I} - \mathbf{H}_l)^{-1}, \quad l = L, R \quad (6)$$

are the semi-infinite left and right regions Green's functions, respectively. It can be shown that the self-energy matrices,  $\Sigma_L(E)$  and  $\Sigma_R(E)$ , can be further reduced to

$$\Sigma_l(E) = \mathbf{H}_{Cl} \mathbf{G}_l^0(E) \mathbf{H}_{lC}, \quad l = L, R, \quad (7)$$

where  $\mathbf{G}_L^0(E)$  and  $\mathbf{G}_R^0(E)$  are the surface Green's matrices of the left and right regions, respectively. In order to compute these matrices, we follow the recursive method of Li and Lu<sup>5</sup>, John and Pulfrey<sup>6</sup> and Sancho *et al.*<sup>7</sup>. Finally, the coupling matrices follow as

$$\Gamma_l(E) = \imath(\Sigma_l(E) - \Sigma_l^\dagger(E)), \quad l = L, R \quad (8)$$

i. e., as the difference between the retarded and the advanced self-energy matrices.

## Tight-binding model

The TBH matrix has the structure

$$\begin{matrix} & 1 & \dots & j & \dots & N \\ \begin{matrix} 1 \\ \vdots \\ i \\ \vdots \\ N \end{matrix} & \begin{pmatrix} \mathbf{H}_{1\alpha,1\beta} & \dots & \dots & \dots & \mathbf{H}_{1\alpha,N\beta} \\ \vdots & \vdots & \vdots & \vdots & \vdots \\ \dots & \dots & \mathbf{H}_{i\alpha,j\beta} & \dots & \dots \\ \vdots & \vdots & \vdots & \vdots & \vdots \\ \mathbf{H}_{N\alpha,1\beta} & \dots & \dots & \dots & \mathbf{H}_{N\alpha,N\beta} \end{pmatrix} & \cdot & \end{matrix} \quad (9)$$

When  $i \neq j$ , the sub-matrices are called hopping sub-matrices and have the form

$$\mathbf{H}_{i\alpha,j\beta} = \begin{pmatrix} h_{is,js} & h_{is,jp_x} & h_{is,jp_y} & h_{is,jp_z} \\ h_{ip_x,js} & h_{ip_x,jp_x} & h_{ip_x,jp_y} & h_{ip_x,jp_z} \\ h_{ip_y,js} & h_{ip_y,jp_x} & h_{ip_y,jp_y} & h_{ip_y,jp_z} \\ h_{ip_z,js} & h_{ip_z,jp_x} & h_{ip_z,jp_y} & h_{ip_z,jp_z} \end{pmatrix} = \begin{pmatrix} h_{ss\sigma} & d_x h_{sp\sigma} & d_y h_{sp\sigma} & d_z h_{sp\sigma} \\ -d_x h_{sp\sigma} & d_x^2 h_{pp\sigma} + d_y^2 h_{pp\pi} & d_x d_y (h_{pp\sigma} - h_{pp\pi}) & d_x d_z (h_{pp\sigma} - h_{pp\pi}) \\ -d_y h_{sp\sigma} & d_x d_y (h_{pp\sigma} - h_{pp\pi}) & d_x^2 h_{pp\sigma} + (1 - d_y^2) h_{pp\pi} & d_y d_z (h_{pp\sigma} - h_{pp\pi}) \\ -d_z h_{sp\sigma} & d_z d_x (h_{pp\sigma} - h_{pp\pi}) & d_z d_y (h_{pp\sigma} - h_{pp\pi}) & d_z^2 h_{pp\sigma} + (1 - d_z^2) h_{pp\pi} \end{pmatrix}, \quad (10)$$

where  $h_{ss\sigma}$ ,  $h_{sp\sigma}$ ,  $h_{pp\sigma}$  and  $h_{pp\pi}$  are the hopping parameters between atoms  $i$  and  $j$ , and  $d_x$ ,  $d_y$  and  $d_z$  are the directional cosines defined as

$$d_x = \frac{\mathbf{r}_{ij} \cdot \mathbf{x}}{r_{ij}}, \quad (11)$$

$$d_y = \frac{\mathbf{r}_{ij} \cdot \mathbf{y}}{r_{ij}}, \quad (12)$$

$$d_z = \frac{\mathbf{r}_{ij} \cdot \mathbf{z}}{r_{ij}}, \quad (13)$$

where  $\mathbf{x}$ ,  $\mathbf{y}$  and  $\mathbf{z}$  represent the unit vectors of a cartesian coordinate system

$$\mathbf{x} = \begin{pmatrix} 1 \\ 0 \\ 0 \end{pmatrix}, \quad \mathbf{y} = \begin{pmatrix} 0 \\ 1 \\ 0 \end{pmatrix}, \quad \mathbf{z} = \begin{pmatrix} 0 \\ 0 \\ 1 \end{pmatrix}, \quad (14)$$

and  $\mathbf{r}_{ij}$  is the vector that goes from atom  $i$  to atom  $j$ , and  $r_{ij}$  is the norm of the vector. Normally,  $h_{ss\sigma}$ ,  $h_{sp\sigma}$ ,  $h_{pp\sigma}$  and  $h_{pp\pi}$  are functions of the interatomic distance between atoms  $i$  and  $j$ .

When  $i = j$ , the sub-matrices are called on-site sub-matrices and have the form

$$\mathbf{H}_{ii} = \begin{pmatrix} h_{is,is} & 0 & 0 & 0 \\ 0 & h_{ip_x,ip_x} & 0 & 0 \\ 0 & 0 & h_{ip_y,ip_y} & 0 \\ 0 & 0 & 0 & h_{ip_z,ip_z} \end{pmatrix} = \begin{pmatrix} E_s & 0 & 0 & 0 \\ 0 & E_{p_x} & 0 & 0 \\ 0 & 0 & E_{p_y} & 0 \\ 0 & 0 & 0 & E_{p_z} \end{pmatrix}. \quad (15)$$

### Definition of the tight binding Hamiltonian matrix in $k$ -space

The Hamiltonian matrix in  $k$ -space for pristine graphene lattice is defined by the sub-matrices

$$\hat{\mathbf{H}} = \begin{pmatrix} \hat{\mathbf{H}}_{11}(\mathbf{k}) & \hat{\mathbf{H}}_{12}(\mathbf{k}) \\ \hat{\mathbf{H}}_{21}(\mathbf{k}) & \hat{\mathbf{H}}_{22}(\mathbf{k}) \end{pmatrix}. \quad (16)$$

Since the unit cell is composed of two atoms, labeled as atom type 1 and 2 in our model, the TBH matrix is built from 4 sub-matrices.

Taking into account interactions up to the first-nearest neighbors for the hopping parameters, we have

$$\hat{\mathbf{H}}_{11}(\mathbf{k}) = \hat{\mathbf{H}}_{22}(\mathbf{k}) = \begin{pmatrix} E_s & 0 & 0 & 0 \\ 0 & E_p & 0 & 0 \\ 0 & 0 & E_p & 0 \\ 0 & 0 & 0 & E_p \end{pmatrix}, \quad (17)$$

$$\hat{\mathbf{H}}_{12}(\mathbf{k}) = \sum_j e^{i\mathbf{k}\cdot\mathbf{r}_{ij}} \begin{pmatrix} h_{ss\sigma} & d_x \cdot h_{sp\sigma} & d_y \cdot h_{sp\sigma} & d_z \cdot h_{sp\sigma} \\ -d_x \cdot h_{sp\sigma} & d_x^2 \cdot h_{pp\sigma} + d_y^2 \cdot h_{pp\pi} & d_x \cdot d_y \cdot (h_{pp\sigma} - h_{pp\pi}) & d_x \cdot d_z \cdot (h_{pp\sigma} - h_{pp\pi}) \\ -d_y \cdot h_{sp\sigma} & d_x \cdot d_y \cdot (h_{pp\sigma} - h_{pp\pi}) & d_x^2 \cdot h_{pp\sigma} + (1 - d_y^2) \cdot h_{pp\pi} & d_y \cdot d_z \cdot (h_{pp\sigma} - h_{pp\pi}) \\ -d_z \cdot h_{sp\sigma} & d_z \cdot d_x \cdot (h_{pp\sigma} - h_{pp\pi}) & d_z \cdot d_y \cdot (h_{pp\sigma} - h_{pp\pi}) & d_z^2 \cdot h_{pp\sigma} + (1 - d_z^2) \cdot h_{pp\pi} \end{pmatrix}, \quad (18)$$

$$\hat{\mathbf{H}}_{21}(\mathbf{k}) = \hat{\mathbf{H}}_{12}^\dagger(\mathbf{k}), \quad (19)$$

where  $j$  runs over the first-nearest neighbors of atom  $i$ .

### SUPPLEMENTARY MATERIAL 3: VALIDATION OF THE DEFORMATION

For verification purposes, we have also computed the deformed configuration of the dislocation structures (15,8) using the DFT SIESTA code<sup>8</sup>. Fig. 3 shows the resulting structure for  $\epsilon_x = 0.18$

using molecular dynamics (MD) and the AIREBO potential, and DFT-based MD calculations. As is evident from the figure, in general both configurations show good agreement for the atomic structure, although we observe small discrepancies in the  $y$ -dimension of the cell due to the difference in the lattice constant predicted by AIREBO, 1.397 Å, and SIESTA, 1.42 Å.

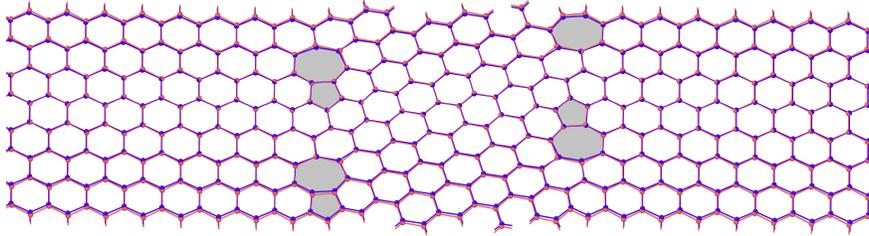


FIG. 3: Comparison of the (15,8) configuration under a uniaxial strain of  $\epsilon_x = 0.18$ , calculated with LAMMPS-AIREBO (blue) and with SIESTA (red).

This calculation was carried out employing the PBE exchange–correlation functional within the generalized gradient approximation (GGA)<sup>9</sup>, the double- $\xi$  with polarization basis set<sup>10</sup>, and smooth norm-conserving pseudopotentials<sup>11</sup>. We apply an increasing force in the  $x$ -direction until the target deformation is reached and conjugate gradients (CG) energy-minimization in every step to obtain the equilibrium structure. We use a grid of 80  $k$ -points and a mesh cut-off of 300 Ry.

## SUPPLEMENTARY REFERENCES

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