Supplementary Materials

Tilt grain boundaries in WS2 from the low to high misorientation angles

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Contents

1. Other details of the coincidence site lattice (CSL) theory for WS2

Figure S1. (a) The CSL cell size as a function of the misorientation angles. **(b)** Periodicity length as a function of misorientation angles. The systems are grouped based on the GB families n_d .

2. First-principle simulation details

The first-principle simulations are mainly performed using the SIESTA code [\[1\]](#page-11-1). The electron-ion interaction is represented by pseudopotentials in the norm-conserving method. The valence electrons of W 5d⁴6s² and S 3s²3p⁴ are explicitly considered. For the basis set, the single-zeta (SZ) basis is used for the structural relaxation, and the "standard" choice of double-zeta plus polarization (DZP) is later adopted for calculating the electronic properties. Exchange-correlation functional is in the form of Perdew-Burke-Ernzerhof [\[2\]](#page-11-2) generalized gradient approximation. Structural optimization is the key to getting the low-energy motifs of the GB models. It is a difficult task and is performed carefully. The convergence criteria are 0.1 eV/Å for the force on atoms.

For cross-checking, calculations are also performed for a few selected systems using the all-electron FHI-aims code [\[3\]](#page-11-3). Formation energies from Siesta and FHI-aims code, shown in Figure S2, agree well with each other.

3. Relaxed structural models

Figure S3-1-1. GB model of family $n_d = 1$ and misorientation angle $\theta = 1.297$. The relaxed simulation cell sizes are $a = 277.35 \text{ Å}$ and $b = 139.81 \text{ Å}.$

Figure S3-1-2. GB model of family $n_d = 1$ and misorientation angle $\theta = 2.134^{\circ}$. The relaxed simulation cell sizes are $a = 169.08 \text{ Å}$ and $b = 85.01 \text{ Å}$.

Figure S3-1-3a. GB model of family $n_d = 1$ and misorientation angle $\theta = 3.481^{\circ}$. The distorted dislocations are 4|6 and 6|8 rings. The relaxed simulation cell sizes are $a = 103.75 \text{ Å}$ and $b = 52.10 \text{ Å}$.

Figure S3-1-3b. GB model of family $n_d = 1$ and misorientation angle $\theta = 3.481^{\circ}$. The distorted dislocations are 5|7 rings. The relaxed simulation cell sizes are $a = 104.15 \text{ Å}$ and $b = 52.10 \text{ Å}$.

Figure S3-1-4. GB model of family $n_d = 1$ and misorientation angle $\theta = 5.086^{\circ}$. The relaxed simulation cell sizes are $a = 142.75 \text{ Å}$ and $b = 35.65 \text{ Å}$.

Figure S3-1-5a. GB model of family $n_d = 1$ and misorientation angle $\theta = 7.341^{\circ}$. The distorted dislocations are 4|6 and 6|8 rings. The relaxed simulation cell sizes are $\alpha = 98.92 \text{ Å}$ and $b = 24.74 \text{ Å}$.

Figure S3-1-5b. GB model of family $n_d = 1$ and misorientation angle $\theta = 7.341^{\circ}$. The distorted dislocations are 5|7 rings. The relaxed simulation cell sizes are $\alpha = 98.60 \text{ Å}$ and $b = 24.75 \text{ Å}$.

Figure S3-1-6. GB model of family $n_d = 1$ and misorientation angle $\theta = 9.430^{\circ}$. The relaxed simulation cell sizes are $a = 76.83 \text{ Å}$ and $b = 19.26 \text{ Å}.$

Figure S3-1-7. GB model of family $n_d = 1$ and misorientation angle $\theta = 13.174$. The relaxed simulation cell sizes are $a = 82.58 \text{ Å}$ and $b = 13.81 \text{ Å}$.

Figure S3-1-8. GB model of family $n_d = 1$ and misorientation angle $\theta = 21.787$. The relaxed simulation cell sizes are $a = 67.55 \text{ Å}$ and $b = 8.38 \text{ Å}.$

3.2. The family of $n_d = 2$

Figure S3-2-1. GB model of family $n_d = 2$ and misorientation angle $\theta = 4.723^{\circ}$. The relaxed simulation cell sizes are $a = 152.44 \text{ Å}$ and $b = 76.83 \text{ Å}.$

Figure S3-2-2. GB model of family $n_d = 2$ and misorientation angle $\theta = 6.609^{\circ}$. The relaxed simulation cell sizes are $a = 109.10 \text{ Å}$ and $b = 54.89 \text{ Å}.$

Figure S3-2-3. GB model of family $n_d = 2$ and misorientation angle $\theta = 8.256^{\circ}$. The relaxed simulation cell sizes are $a = 86.45 \text{ Å}$ and $b = 44.02 \text{ Å}.$

Figure S3-2-4. GB model of family $n_d = 2$ and misorientation angle $\theta = 10.993^{\circ}$. The relaxed simulation cell sizes are $a = 131.18 \text{ Å}$ and $b = 33.08 \text{ Å}.$

Figure S3-2-5. GB model of family $n_d = 2$ and misorientation angle $\theta = 16.426$. The relaxed simulation cell sizes are $a = 87.74 \text{ Å}$ and $b = 22.19 \text{ Å}$.

Figure S3-2-6. GB model of family $n_d = 2$ and misorientation angle $\theta = 32.204^{\circ}$. The relaxed simulation cell sizes are $a = 90.42 \text{ Å}$ and $b = 11.41 \text{ Å}.$

3.3. The family of $n_d = 3$

Figure S3-3-1. GB model of family $n_d = 3$ and misorientation angle $\theta = 8.613^{\circ}$. The relaxed simulation cell sizes are $a = 145.32 \text{ Å}$ and $b = 63.25 \text{ Å}$.

Figure S3-3-2. GB model of family $n_d = 3$ and misorientation angle $\theta = 11.635^{\circ}$. The relaxed simulation cell sizes are $a = 107.77 \text{ Å}$ and $b = 46.89 \text{ Å}.$

Figure S3-3-3. GB model of family $n_d = 3$ and misorientation angle $\theta = 15.178^{\circ}$. The relaxed simulation cell sizes are $a = 82.37 \text{ Å}$ and $b = 36.01 \text{ Å}$.

Figure S3-3-4. GB model of family $n_d = 3$ and misorientation angle $\theta = 17.897$. The relaxed simulation cell sizes are $a = 69.97 \text{ Å}$ and $b = 30.59 \text{ Å}$.

Figure S3-3-5. GB model of family $n_d = 3$ and misorientation angle $\theta = 27.796^{\circ}$. The relaxed simulation cell sizes are $a = 67.97 \text{ Å}$ and $b = 19.76 \text{ Å}.$

Figure S3-3-6. GB model of family $n_d = 3$ and misorientation angle $\theta = 38.213$. The relaxed simulation cell sizes are $a = 99.66 \text{ Å}$ and $b = 14.49 \text{ Å}.$

3.4. The family of $n_d = 4$

Figure S3-4-1. GB model of family $n_d = 4$ and misorientation angle $\theta = 18.734$. The relaxed simulation cell sizes are $a = 76.79 \text{ Å}$ and $b = 38.60 \text{ Å}$.

Figure S3-4-2. GB model of family $n_d = 4$ and misorientation angle $\theta = 26.008^{\circ}$. The relaxed simulation cell sizes are $a = 111.38 \text{ Å}$ and $b = 28.13 \text{ Å}$.

Figure S3-4-3. GB model of family $n_d = 4$ and misorientation angle $\theta = 42.103^{\circ}$. The relaxed simulation cell sizes are $a = 104.65 \text{ Å}$ and $b = 17.62 \text{ Å}$.

4. Weak magnetic instability

Magnetic moment could be strongly underestimated by the conventional LDA and GGA functional due to the well-known self-interaction error. This is significantly improved by the recently-developed SCAN (stands for *Strongly Constrained and Appropriately Normed* [\[5\]](#page-11-4))meta-GGA, as established in our previous work in transition-metal mono-oxides [\[4\]](#page-11-5). Here, we use SCAN implemented in FHI-aims [\[3\]](#page-11-3) to evaluate the magnetic moments of two systems.

Figure S4. Magnetic moments calculated using the SCAN meta-GGA for two GB models. (a) Family $n_d = 1$ and the misorientation angle $\theta = 7.341$ °. (b) Family $n_d = 4$ and the misorientation angle $\theta = 42.103$ °. The magnetic moments of W atoms are shown in the unit of μ_B .

5. Deriving the critical angle by fitting to the Read-Shockley relation

Carlsson et al. calculated the formation energies of graphene GBs [\[6\]](#page-11-6), covering various misorientation angles in the range $0 \le \theta \le 60$. They found the Read-Shockley relation for low-angle GBs [\[7\]](#page-11-7) is valid up to $\theta_c=12^\circ$. However, we argue that the critical angle could be extended to a higher value of $\theta_c\approx 20^\circ$, as shown in Figure S5(a). Too strict fitting criteria might have been used in Ref [\[6\]](#page-11-6). Following the same approach, we derive a critical angle of $\theta_c \approx 14^\circ$ for the WS₂-GBs [Figure S5(b)].

Figure S5. Agreement of the calculated data with the Read-Shockley relation. **(a)** Graphene results fitted up to the angle $\theta = 18.73$ ["]. (b) WS₂ results fitted up to $\theta = 13.17$ ["].

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