## **Supplementary Materials**

# Tilt grain boundaries in WS<sub>2</sub> from the low to high misorientation angles

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### 1. Other details of the coincidence site lattice (CSL) theory for WS<sub>2</sub>



**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]. The electron-ion interaction is represented by pseudopotentials in the norm-conserving method. The valence electrons of W 5d<sup>4</sup>6s<sup>2</sup> and S 3s<sup>2</sup>3p<sup>4</sup> 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] 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]. Formation energies from Siesta and FHI-aims code, shown in Figure S2, agree well with each other.





### 3. Relaxed structural models

3.1. The family of  $n_d = 1$ 



**Figure S3-1-1.** GB model of family  $n_d = 1$  and misorientation angle  $\theta = 1.297^{\circ}$ . The relaxed simulation cell sizes are a = 277.35 Å and b = 139.81 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 Å and b = 85.01 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 Å and b = 52.10 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 Å and b = 52.10 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 Å and b = 35.65 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 a = 98.92 Å and b = 24.74 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 a = 98.60 Å and b = 24.75 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 Å and b = 19.26 Å.



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



**Figure S3-1-8.** GB model of family  $n_d = 1$  and misorientation angle  $\theta = 21.787^{\circ}$ . The relaxed simulation cell sizes are a = 67.55 Å and b = 8.38 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 Å and b = 76.83 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 Å and b = 54.89 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 Å and b = 44.02 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 Å and b = 33.08 Å.



**Figure S3-2-5.** GB model of family  $n_d = 2$  and misorientation angle  $\theta = 16.426^{\circ}$ . The relaxed simulation cell sizes are a = 87.74 Å and b = 22.19 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 Å and b = 11.41 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 Å and b = 63.25 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 Å and b = 46.89 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 Å and b = 36.01 Å.



**Figure S3-3-4.** GB model of family  $n_d = 3$  and misorientation angle  $\theta = 17.897^{\circ}$ . The relaxed simulation cell sizes are a = 69.97 Å and b = 30.59 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 Å and b = 19.76 Å.



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

3.4. The family of  $n_d = 4$ 



**Figure S3-4-1.** GB model of family  $n_d = 4$  and misorientation angle  $\theta = 18.734^{\circ}$ . The relaxed simulation cell sizes are a = 76.79 Å and b = 38.60 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 Å and b = 28.13 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 Å and b = 17.62 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])meta-GGA, as established in our previous work in transition-metal mono-oxides [4]. Here, we use SCAN implemented in FHI-aims [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  $\theta = 7.341^{\circ}$ . (b) Family  $n_d = 4$  and the misorientation angle  $\theta = 42.103^{\circ}$ . The magnetic moments of W atoms are shown in the unit of  $\mu_{\text{B}}$ .

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

Carlsson et al. calculated the formation energies of graphene GBs [6], covering various misorientation angles in the range  $0^{\circ} < \theta \le 60^{\circ}$ . They found the Read-Shockley relation for low-angle GBs [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]. Following the same approach, we derive a critical angle of  $\theta_c \approx 14^{\circ}$  for the WS<sub>2</sub>-GBs [Figure S5(b)].



**Figure S5.** Agreement of the calculated data with the Read-Shockley relation. (a) Graphene results fitted up to the angle  $\theta = 18.73^{\circ}$ . (b) WS<sub>2</sub> results fitted up to  $\theta = 13.17^{\circ}$ .

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