Supporting Information for

Understanding the Zinc-Doped Hydroxyapatite Structures Using the First-

Principles Calculations and Convolutional Neural Network Algorithm

Jing Wang¹, Ruihan Wang¹, Mingli Yang² and Dingguo Xu^{1,2*}

¹MOE Key Laboratory of Green Chemistry and Technology, College of Chemistry,

Sichuan University, Chengdu, Sichuan, 610064, PR China

²Research Center for Material Genome Engineering, Sichuan University, Chengdu,

Sichuan, 610065, PR China

* To whom correspondence should be addressed: dgxu@scu.edu.cn (D.X), Tel: 86-

28-85406156.

- 1. The number of structures for each doping concentration in the initial dataset
- 2. The number of the structures by the three substitution mechanisms in the lowest energy range.
- **3.** The two leading principal components of the feature extraction vector learned by 5 Zn atoms doped HAP.
- 4. Detailed display of three Zn²⁺ doped structures.
- 5. Substitution mechanisms when two Zn are doped in the same Ca2-V channel.
- 6. The volumes between the most stable and the most unstable structures.

Concentration (mol%)	1.25	2.5	3.75	5	6.25
all the structures	80	2800	56000	700000	5600000
Structures in the initial dataset	56	130	300	604	1370

Table S1. The number of structures for 5 doping concentrations in the initial dataset.

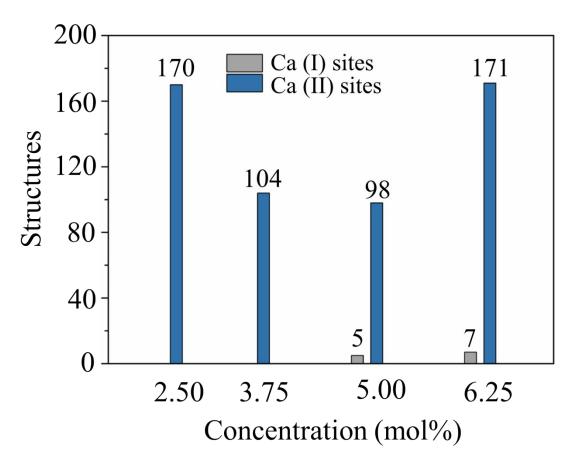


Figure S1. For the doping concentration of 2.5-6.25mol%, the number of the structures by the three substitution mechanisms in the lowest 5kcal/mol energy range.

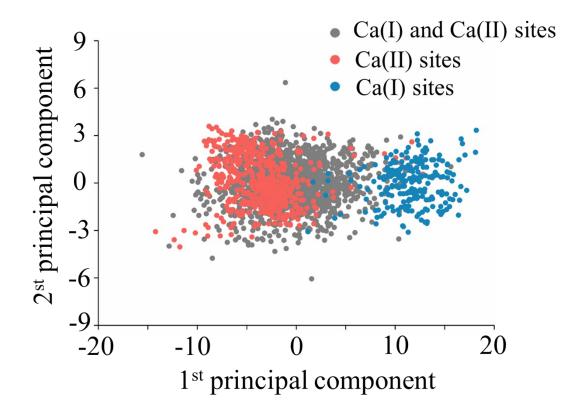


Figure S2. The two leading principal components of the feature extraction vector learned by Zn doped HAP with the concentration of 6.25 at.%.

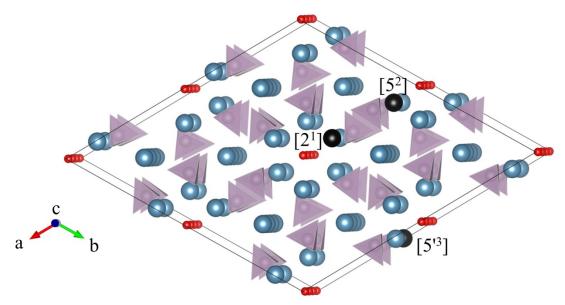


Figure S3. Detailed display of three Zn^{3+} doped structures.

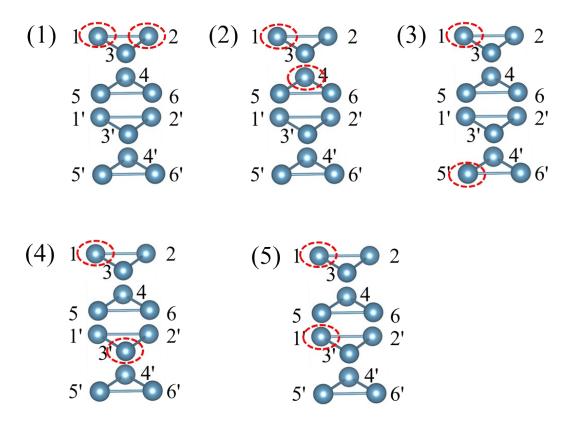


Figure S4. There are five different representative substitution mechanisms when two zinc atoms are doped in the same Ca2-V channel.

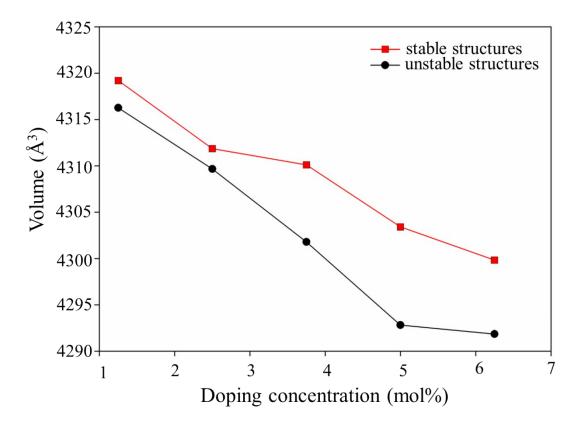


Figure S5. The comparison results of the volumes between the most stable and the most unstable structures.