

## Supporting Information

### **Enhanced Biocompatibility and Osteogenic Differentiation of Mesenchymal Stem Cells of Titanium by Sr-Ga Clavate Double Hydroxides**

Maowen Chen<sup>a, 1</sup>, Bailong Tao<sup>b, 1</sup>, Yan Hu<sup>a\*</sup>, Menghuan Li<sup>c</sup>, Maohua Chen<sup>a</sup>, Lu Tan<sup>a</sup>,  
Zhong Luo<sup>c\*</sup>, Kaiyong Cai<sup>a\*</sup>

<sup>a</sup>Key Laboratory of Biorheological Science and Technology, Ministry of Education, College of Bioengineering, Chongqing University, Chongqing 400044, China.

<sup>b</sup>Laboratory Research Center, The First Affiliated Hospital of Chongqing Medical University, Chongqing, 400016, China.

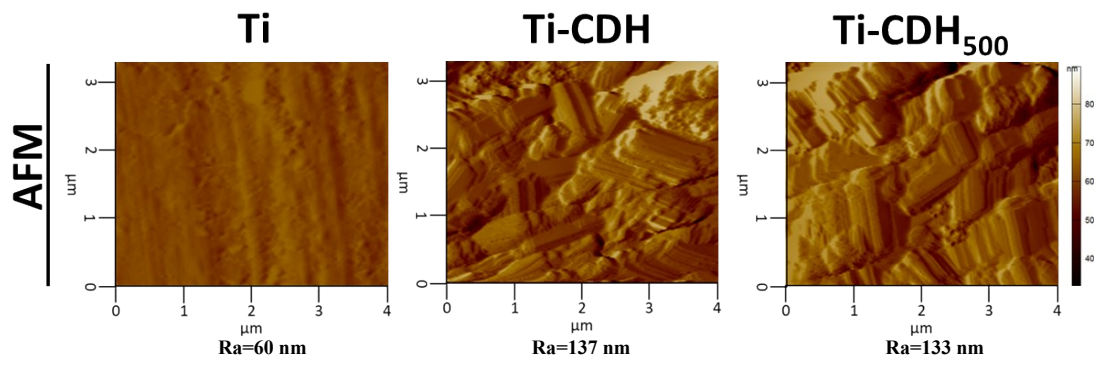
<sup>c</sup>School of Life Science, Chongqing University, Chongqing 400044, China

\*Corresponding author: Professor Yan Hu,

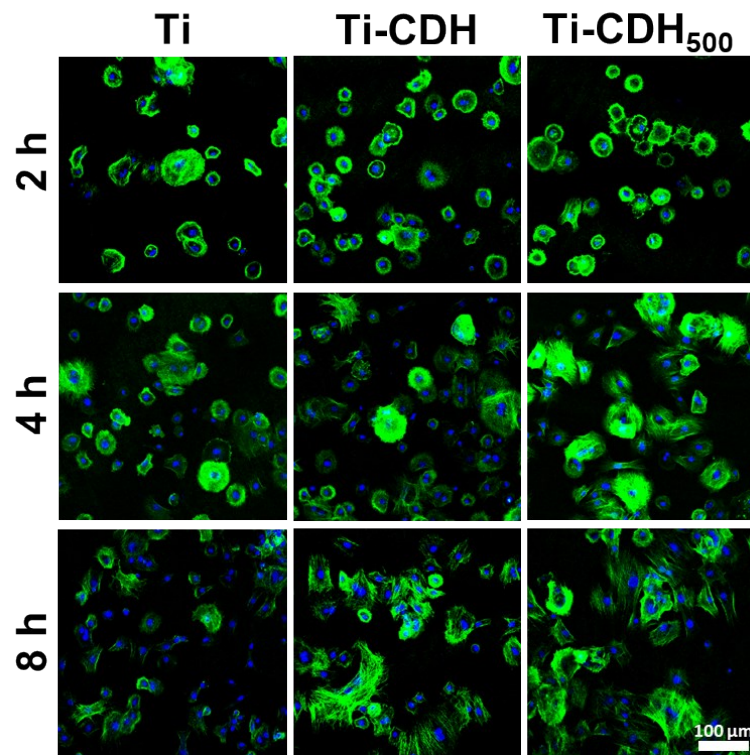
Professor Zhong Luo & Professor Kaiyong Cai

E-mail: huyan303@cqu.edu.cn; luozhong918@cqu.edu.cn; kaiyong\_cai@cqu.edu.cn.

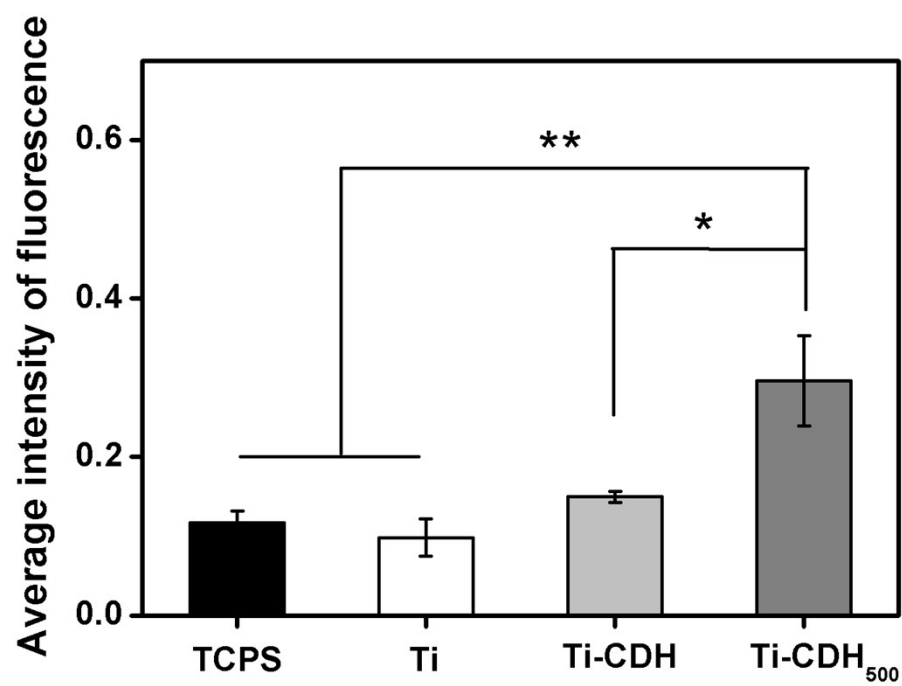
<sup>1</sup> Co-first authors: these authors contributed equally to this work.



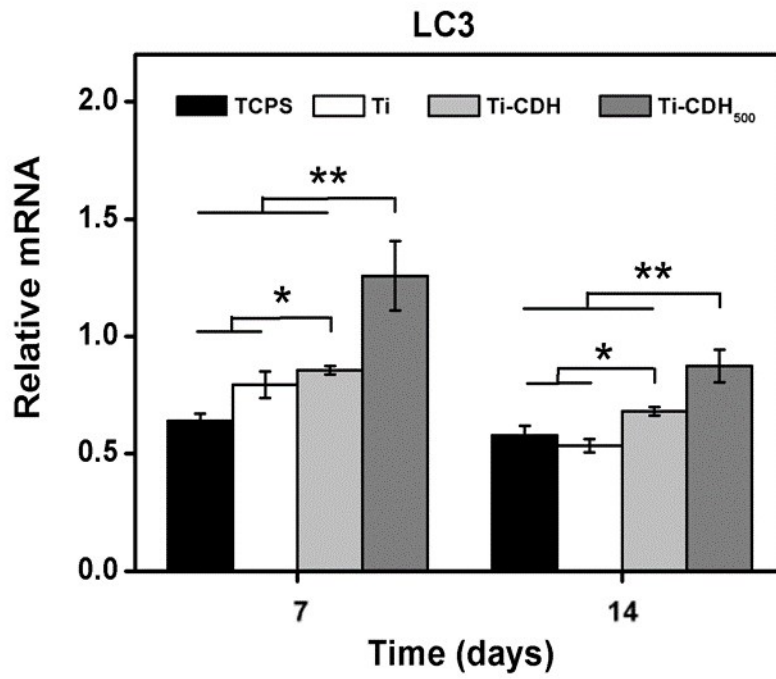
**Fig. S1.** AFM images of Ti, Ti-CDH and Ti-CDH<sub>500</sub> groups.



**Fig. S2.** (A) Initial MSCs adhesion on various Ti substrates for 2, 4, and 8 h with cytoskeleton staining.



**Fig. S3.** Average intensity of fluorescence statistics. Error bars represent means  $\pm$  SD,  $n=3$ ,  $*p < 0.05$ ,  $**p < 0.01$ .



**Fig. S4.** Relative mRNA expression of several autophagic related genes (LC3) in MSCs treated by various samples for 7 and 14 days, respectively. Error bars represent means ± SD, n=3, \*p < 0.05, \*\*p < 0.01.

**Table S1.** Statistics of chemical compositions on the surface of different substrates.

<i>Samples</i>	<i>Elemental content (wt%)</i>				
	Ti	O	C	Sr	Ga
Ti	88.95	5.79	4.40	—	—
CDH	—	26.55	10.21	34.77	20.62
CDH <sub>500</sub>	—	28.98	10.12	25.69	28.06

**Table S2. Real-time polymerase chain reaction primers used in this study.**

<i>Target gene</i>	<i>Primers</i>	<i>Product size (bp)</i>
ALP	AGCGACACGGACAAGAAGC GGCAAAGACCGCCACATC	183
OPG	GCCCAGACGAGATTGAGAG CAGACTGTGGGTGACGGTT	173
OPN	GACAGCAACGGGAAGACC CAGGCTGGCTTTGGAAC	216
BMP2	AGATTGTTGGGGCACAAGGT CCTTCAGCAGGGAAACCGAT	191
GAPDH	CATCCGTAAAGACCTCTATGCCAA C ATGGAGCCACCGATCCACA	223