

Supplementary Information

Gd@C₈₂(OH)₂₂ harnesses inflammatory regeneration for osteogenesis of mesenchymal stem cells through JNK/STAT3 signaling pathway

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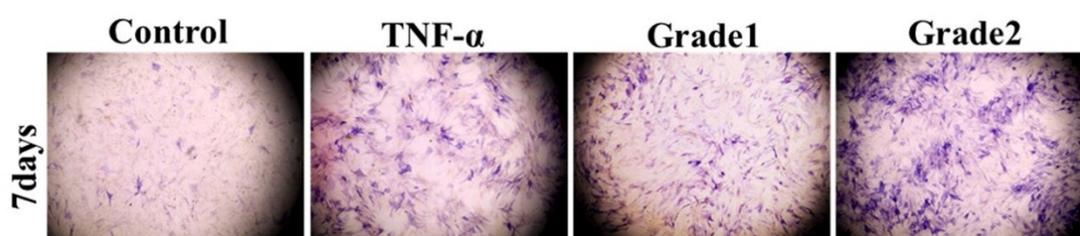


Figure S1. The effects of TNF- α , Grade 1 and Grade 2 on osteogenesis of hMSCs.

Alkaline phosphatase (ALP) staining was performed, after hMSCs were induced to osteoblasts with the treatment of 10 ng/ml TNF- α , Grade 1 and Grade 2 for 7 days.

Inflammatory microenvironment, Grade 1 and Grade 2.

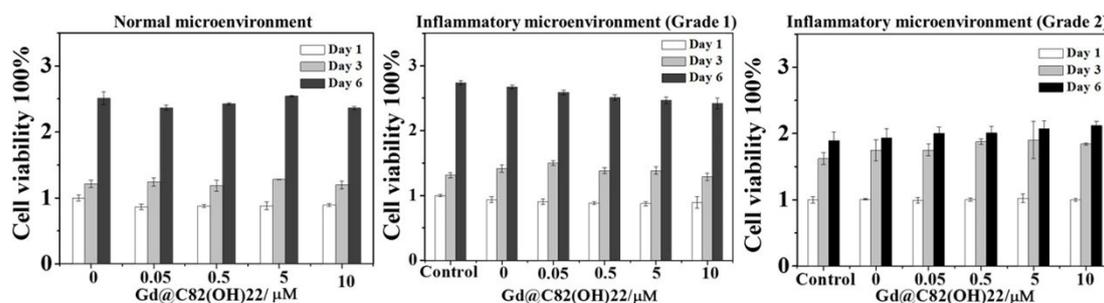


Figure S2. Gd@C₈₂(OH)₂₂ have no cytotoxicity of hMSCs in normal

microenvironment (growth medium) and inflammatory microenvironment (Grade 1, Grade 2). After treating hMSCs with 0.05, 0.5, 5, 10 μM $\text{Gd@C}_{82}(\text{OH})_{22}$ for 1, 3, 6 days, CCK-8 kits were used to measure the cell viability of hMSCs. Experiments were performed in triplicates. The data was presented as the mean \pm standard derivation.

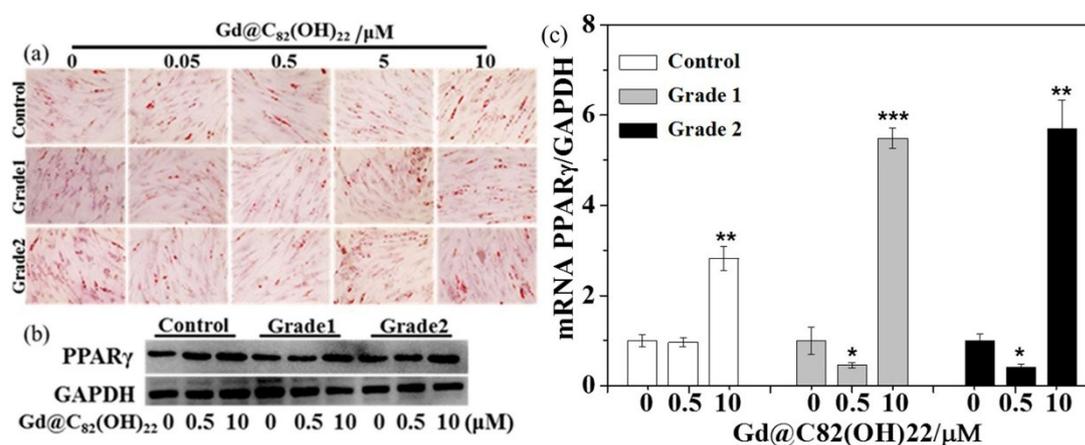


Figure S3. The effect of $\text{Gd@C}_{82}(\text{OH})_{22}$ on adipogenesis of hMSCs in Control (adipogenic induction medium, AIM) and inflammatory microenvironment (dilute CM with AIM 50 (Grade 1) and 20 (Grade 2) folds). **(a)** hMSCs were induced to adipocytes with treatment of 0, 0.05, 0.5, 5 and 10 μM $\text{Gd@C}_{82}(\text{OH})_{22}$ for 10 days, Oil droplets were stained (first row: AIM; second row: Grade 1; third row: Grade 2). **(b)** PPAR γ expression was measured by western blotting after adipogenic differentiation 5 days in different microenvironment. **(c)** PPAR γ expression was measured by RT-PCR after hMSCs were induced to adipocytes with treatment of 0, 0.5, 10 μM $\text{Gd@C}_{82}(\text{OH})_{22}$ for 5 days. * $P < 0.05$; ** $P < 0.01$; *** $P < 0.001$.

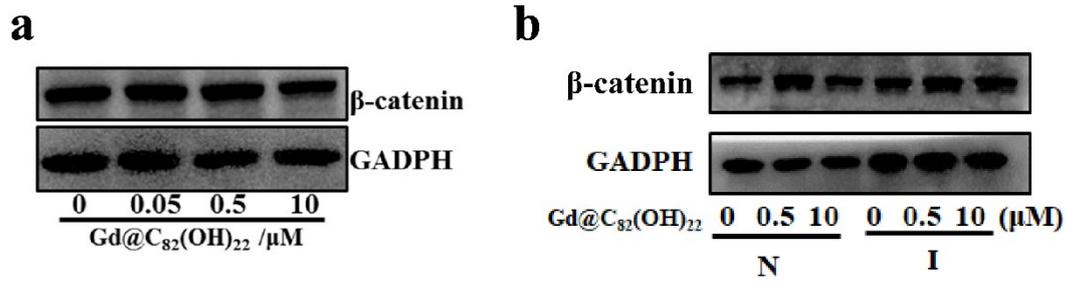


Figure S4. Gd@C₈₂(OH)₂₂ regulates osteogenesis by the β -catenin-independent signaling pathway. (a) After 0.05, 0.5 and 10 μ M Gd@C₈₂(OH)₂₂ treatment in normal microenvironment, the expression of β -catenin of hMSCs were analyzed by western blotting. (b) The role of inflammatory cytokines on the β -catenin expression of hMSCs. N: Normal microenvironment; I: inflammatory microenvironment.

Table S1. Concentration of main inflammatory factors in inflammatory microenvironment

Inflammatory factors	Inflammatory microenvironment(pg/ml)	
	Grade 1	Grade 2
IL-6	1.3±0.1	3.2±0.2
TNF- α	1.1±0.1	2.6±0.1
IL-1 β	1.1±0.1	2.8±0.1
IL-10	0.3±0.1	0.9±0.1

Table S2. Primer of specific osteogenic and adipogenic gene

Gene	Upstream primer	Downstream primer
<i>GAPDH</i>	5'-CCTGCACCACCAACTGCTTA-3'	5'- GGCCATCCACAGTCTTCTGAG-3'
<i>PPARγ</i>	5'-GCGAGGGCGATCTTGACAGG-3'	5'- TGATCACCTGCAGTAGCTGCAC-3'
<i>OCN</i>	5'-GGCTCACCTCCATCACTC-3'	5'- TCCAGCACTGTTTATACCCTCT-3'

OPN

5'-CAGATGCTGTGGCCACATGG-3'

5'-GGTGAGACTCATCAGACTGGTGAG-3'
