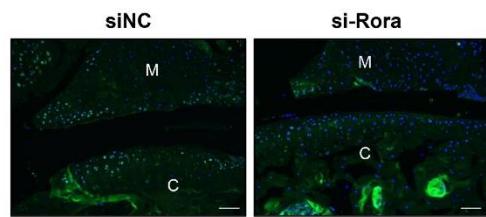
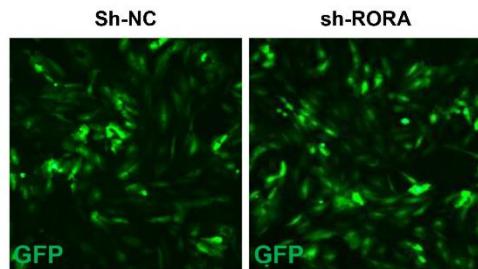


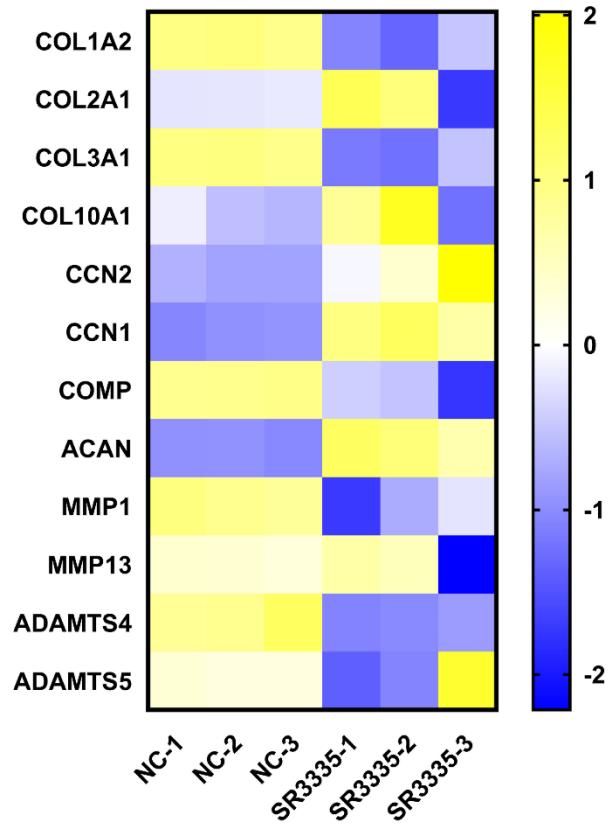
A



B

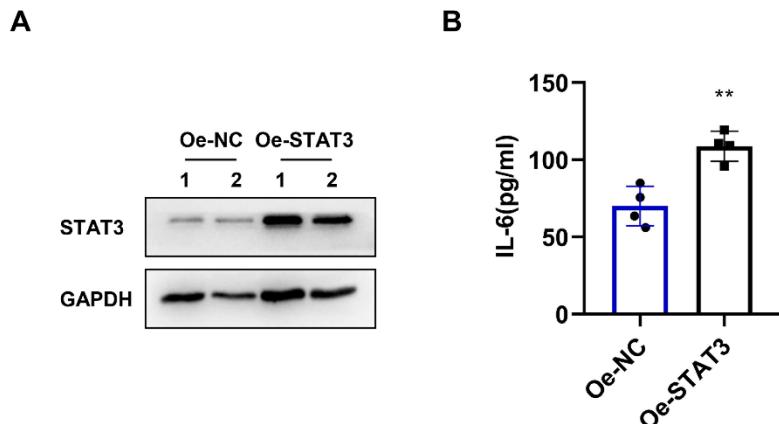


Supplementary Figure 1 The infection efficiency in vivo and in vitro. (A) The immunofluorescence image of ROR α (green) and DAPI (blue) 5 days after intra-articular injection of si -Rora or si-NC; M, meniscus; AC, articular cartilage. Scale bars, 100 μ m. **(B)** Fluorescence of GFP in human chondrocyte infected with sh-RORA or sh-NC lentivirus at 80 MOI separately.



Supplementary Figure 2 The heat map for certain genes after SR3335 treatment.

The relative expression of COL1A2, COL2A1, COL3A1, COL10A1, CCN2, CCN1, COMP, ACAN, MMP1, MMP13, ADAMTS4, and ADAMTS5 was shown.



Supplementary Figure 3 The concentration of IL-6 after overexpressing STAT3.

(A) The chondrocytes were transfected with STAT3 overexpression plasmid and control plasmid for 48 h. The expression level of STAT3 protein was detected with Western blot. **(B)** ELISA assay to detect IL-6 protein expression after transfected with STAT3 overexpression or control plasmid for 48 h. n = 4 independent experiments. Data in (B) are presented as mean \pm SD. The statistical data in (B) was analyzed with Student's t test. ** P < 0.01. All data shown above are presented as mean \pm SD.

Supplementary Table 1 Primer sequence used in this study

Gene	Forward/ Reverse	Primer sequence (5'-3')
<i>RORA</i>	Forward	ACTCCTGTCCTCGTCAGAAGA
	Reverse	CATCCCTACGGCAAGGCATT
<i>COL2A1</i>	Forward	GGCAATAGCAGGTTCACGTACA
	Reverse	CGATAACAGTCTGCCCACTT
<i>ACAN</i>	Forward	TGCATTCCACGAAGCTAACCTT
	Reverse	GACGCCTCGCCTTCTTGAA
<i>COL10A1</i>	Forward	CAAGGCACCATCTCCAGGAA
	Reverse	AAAGGGTATTGTGGCAGCATATT
<i>SOX9</i>	Forward	AGCGAACGCACATCAAGAC
	Reverse	GCTGTAGTGTGGGAGGTTGAA
<i>MMP13</i>	Forward	CCAGACTTCACGATGGCATTG
	Reverse	GGCATCTCCTCCATAATTGGC
<i>ADAMTS4</i>	Forward	GAGGAGGAGATCGTGTTC
	Reverse	CCAGCTCTAGTAGCAGCGTC
<i>ADAMTS5</i>	Forward	GAACATCGACCAACTCTACTCCG
	Reverse	CAATGCCACCAGAACCATCT
<i>IL6</i>	Forward	ACTCACCTCTTCAGAACGAATTG
	Reverse	CCATTTGGAAGGTTCAGGTTG
<i>IL6R</i>	Forward	CCCCTCAGCAATGTTGTTGT
	Reverse	CTCCGGGACTGCTAACTGG
<i>STAT3</i>	Forward	CAGCAGCTTGACACACGGTA
	Reverse	AAACACCAAAGTGGCATGTGA
<i>SOCS3</i>	Forward	CCTGCGCCTCAAGACCTTC
	Reverse	GTCACTGCGCTCCAGTAGAA
<i>JAK2</i>	Forward	TCTGGGGAGTATGTTGCAGAA
	Reverse	AGACATGGTGGGTGGATACC
<i>IL1B</i>	Forward	AGCTACGAATCTCGACCAC
	Reverse	CGTTATCCCATGTGTCGAAGAA
<i>IL17A</i>	Forward	TCCCCACGAAATCCAGGATGC
	Reverse	GGATGTTCAGGTTGACCATCAC
<i>TNF</i>	Forward	CCTCTCTCTAATCAGCCCTCTG
	Reverse	GAGGACCTGGGAGTAGATGAG
<i>GAPDH</i>	Forward	AGAAAAAACCTGCCAAATATGATGAC

Reverse TGGGTGTCGCTGTTGAAGTC

Supplementary Table 2 Oligo sequence used in this study

Oligo	Forward/ Reverse	Primer sequence (5'-3')
Si-Rora-1	Forward	CCGGGCGCAGGCAGAGCTATGCGAGCTCG AGCTCGCATAGCTCTGCCTGCGCTTTTG
	Reverse	AATTCAAAAAGCGCAGGCAGAGCTATGCG AGCTCGAGCTCGCATAGCTCTGCCTGCGC
Si-Rora-2	Forward	CCGGCAGGCAGAGCTATGCGAGCTCCTCG AGGAGCTCGCATAGCTCTGCCTTTTG
	Reverse	AATTCAAAAACAGGCAGAGCTATGCGAGC TCCTCGAGGAGCTCGCATAGCTCTGCCTG
Si-Rora-3	Forward	CCGGGCTATGCGAGCTCCAGCCGAGCTCG AGCTCGGCTGGAGCTCGCATAGCTTTTG
	Reverse	AATTCAAAAAGCTATGCGAGCTCCAGCCG AGCTCGAGCTCGGCTGGAGCTCGCATAGC
Sh-RORA-1	/	GGAGAAGTCAGCAAAGCAATGCTCGAGC ATTGCTTGCTGACTCTCC
Sh-RORA-2	/	GCAGAGAGACAGCTTGTATGCCTCGAGGC ATACAAGCTGTCTCTGC
Sh-RORA-3	/	AAACGTGACACGTTGGAGAACGAATTCT CCGAACGTGTCACGTTT