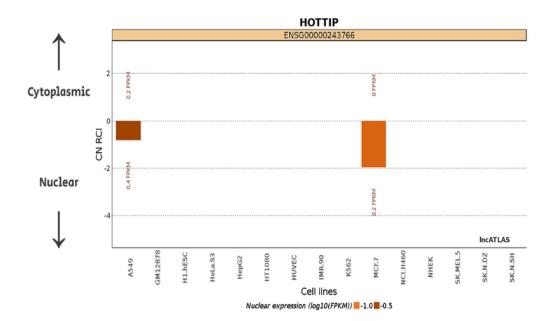
Supplemental Information

Silencing of Long Non-coding RNA HOTTIP

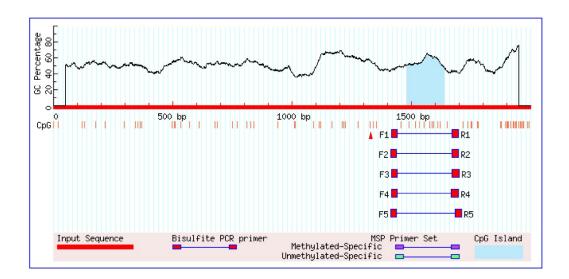
Reduces Inflammation in Rheumatoid

Arthritis by Demethylation of SFRP1

Xumin Hu, Jianhua Tang, Xuyun Hu, Peng Bao, Weixi Deng, Jionglin Wu, Yuwei Liang, Zhipeng Chen, Liangbin Gao, and Yong Tang



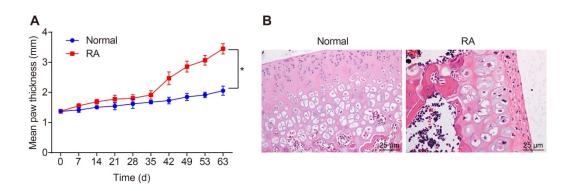
Supplementary Figure 1. HOTTIP was mainly located in the nucleus of RASFs predicted by the lncATLAS website.



Supplementary Figure 2. The CpG islands in the promoter region of SFRP1 were analyzed by the Methprimer website.

序号▼	StartInGenome 🔽	EndInGenome 🔽	MeanStability 🔽	MeanIdentidy (%) 🔽	Score -	TFO_sequence 🔻	TTS_sequence 🔽
1	41310643	41310703	2.27	65.08	1.67	TTTTGACCCCTATTATAATTT CATCTTCAGTGTTTTATTAT CCACTTCCTCTCTCTAT CTT	CACTAGGGCAGACGCCAAG AGGACGGTATCAGCACAAC AGCTCGGCCCCCAGGAAC AGAAG
2	41309649	41309702	1.89	62.96	1.44	ATTATTTTTAAAAAAGAATGA ATAAAGATGTTCTGGTTTCT TTTGTTTT	CCTTCTTTTTCTCCCCTTG TCTCTTTCCTCCTCCCCCT TTTATTTATGTATTTT
3	41309658	41309745	1.85	60.23	0.88	CTAAAAATTACATATGAAAGA GGAAGATTTATGTTACTTTTT TATATGAGAGGAATCGTCCTT TAATAGAAAAATTTCTATTG	CGTCCGCCCTGGTCTCTC TCCACCCCACGCCGTGAT CCATTCCCCTTCTTTTTCT CCCCTTGTCTCTTTATTT
4	41309650	41309701	2.08	64.15	1.55	TCATCTTTCTAAAAATTACAT ATGAAAGAGGAAGATTTATG TTACTTTTTTAT	
5	41311229	41311283	1.25	63.16	1.26	CAAGGCCAGCTCCACATTC TTCCCTCCCCCTCCCACTT CACCGTAGCCCCGAACCC	
6	41311134	41311196	1.1	63.64	1.36	GAGTAGGGTTCTAGGCCCC TGTTCCTGGGGACTTGAAG GCGGTTTTACATACTGGTCA GACACGGC	GTGCACACAGGAGGCTTCT

Supplementary Figure 3. The binding sites of HOTTIP and SFRP1 were predicted by the website Long Target.



Supplementary Figure 4. The rat model of RA was identified using paw swelling scores of rats and pathological changes in the lesions induced after modeling. (A) The paw thickness of rats after modeling. (B) The pathological changes in the lesion after modeling detected by HE staining (× 100; scale bar = 200 μ m). The normal group refers to rats without treatment. *, p < 0.05. The results of paw thickness were expressed as mean \pm standard deviation. Comparisons among multiple groups were analyzed by one-way ANOVA. n = 15.