

# Title; Sirt6 regulates postnatal growth plate differentiation and proliferation via Ihh signaling.

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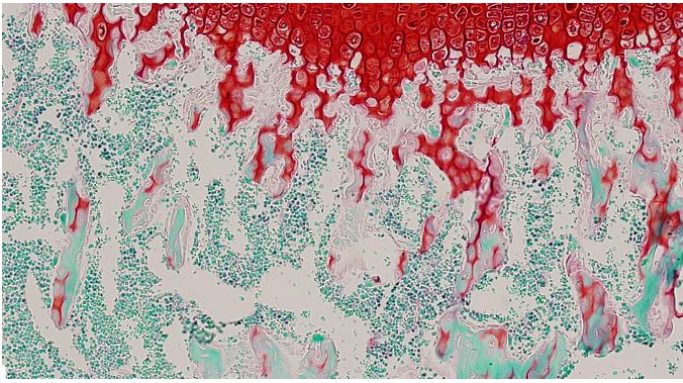
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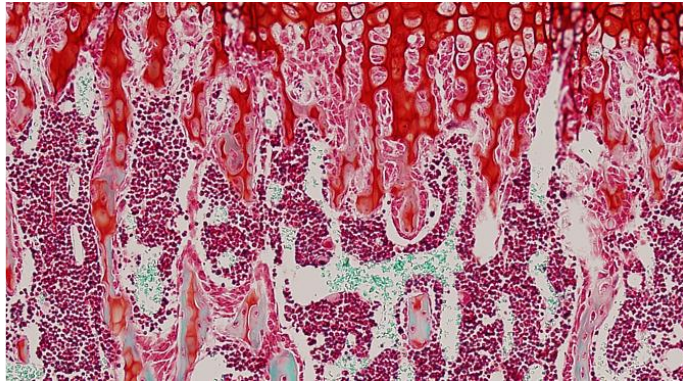
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WT



Sirt6<sup>-/-</sup>

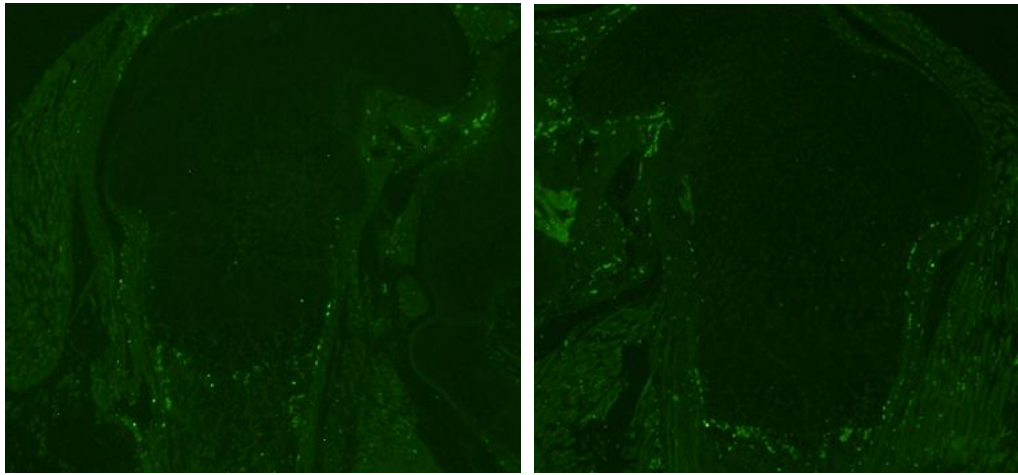


Supplementary Figure S1

Primary spongiosa from 3.5-week-old mice were stained with safranin O (X100).

Note the remarkable delay of ossification in Sirt6<sup>-/-</sup> mice.

a



WT

Sirt6-/-

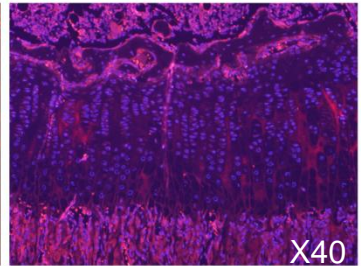
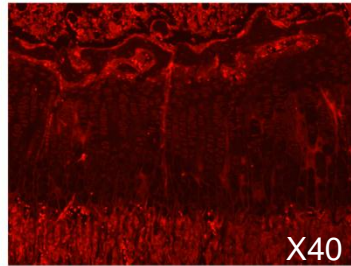
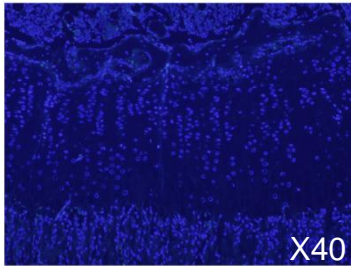
b

DAPI

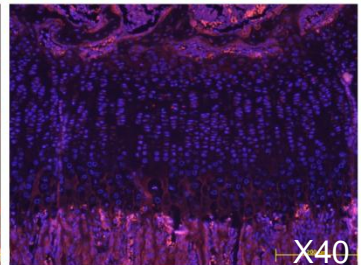
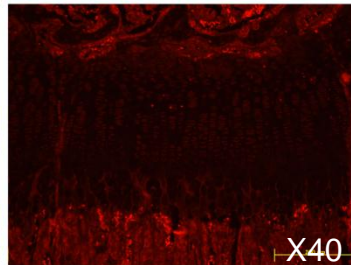
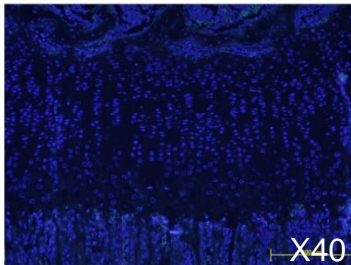
Caspase-3

Merge

WT



Sirt6 -/-



Supplementary Figure S2

(a) TUNEL staining of P1 WT and Sirt6<sup>-/-</sup> mouse femur sections.

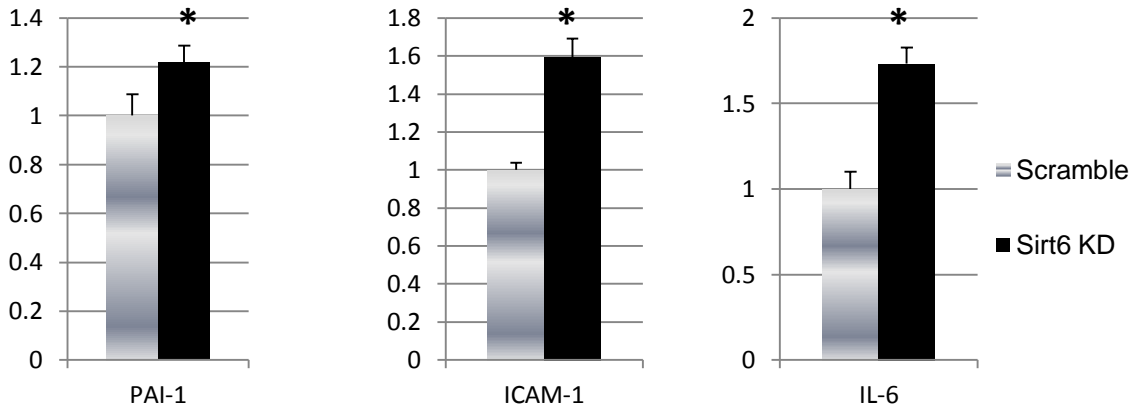
The number of apoptotic cells was similar between Sirt6<sup>-/-</sup> and WT.

(b) Immunohistological findings for activated caspase-3 in P14.

Caspase-3 was not activated in the proliferating zone of both WT and Sirt6<sup>-/-</sup> mice.

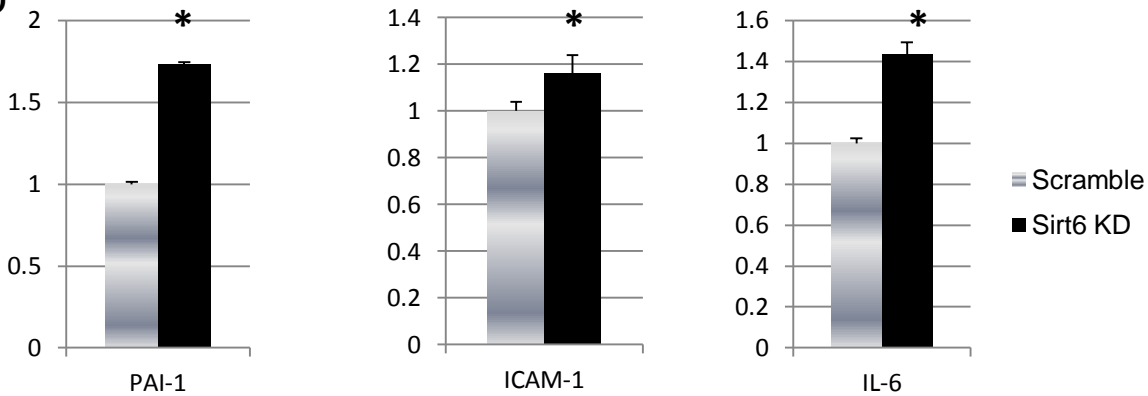
# Primary chondrocyte

a



# ATDC5 cells

b



## Supplementary Figure S3

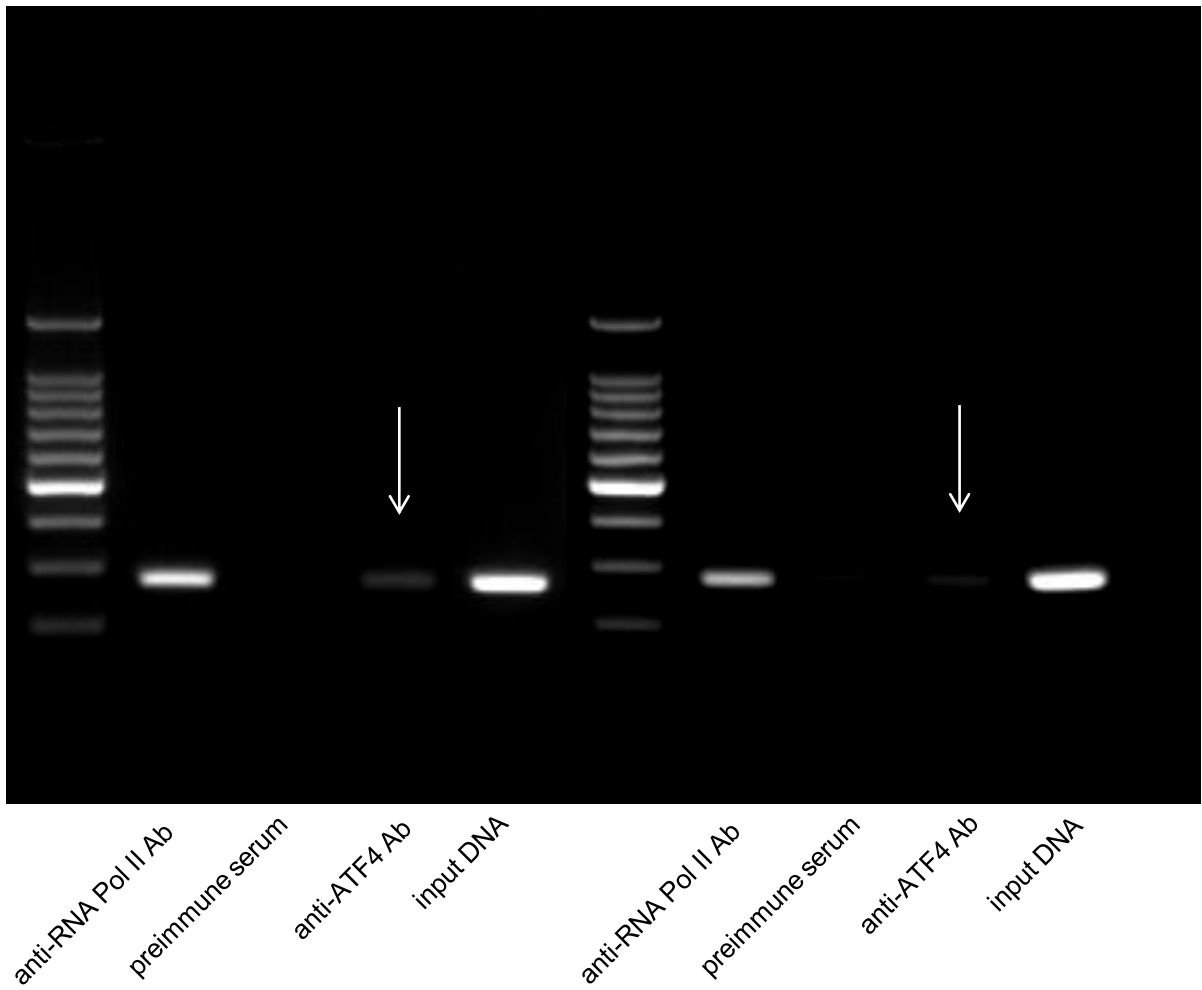
mRNA expression of PAI-1, ICAM-1 and IL-6 was enhanced by Sirt6 knockdown in primary chondrocyte (a) and ATDC5 cells (b).

The graph shows relative levels of gene expression.

Values represent the mean  $\pm$  SD of 3 samples per group. \*; p < 0.05.

scrambled siRNA

Sirt6 siRNA



Supplementary Figure S4

Representative CHIP of *Ihh* with anti-ATF-4 antibody (Ab) and qRT-PCR of CHIP-ed in primary chondrocytes treated with scrambled siRNA (left panel) and Sirt6 siRNA (right panel).