Identification of juvenility-associated genes in the mouse hepatocytes and cardiomyocytes.

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Supplementary Information

Identification of juvenility-associated genes in the mouse hepatocytes and cardiomyocytes.

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Supplementary Figures

Supplementary Figure S1. Appearance of the isolated hepatocytes and cardiomyocytes.

(A) The image of hepatocytes isolated from the P1 mouse taken one day after the isolation. Scale bar, 50 μ m.

(B) The image of hepatocytes isolated from the P7 mouse taken one day after the isolation. Scale bar, 50 μ m.

(C) The image of hepatocytes isolated from the P56 mouse taken one day after the isolation. Scale bar, 50 μ m.

(D) The image of cardiomyocytes isolated from the P1 mouse taken one day after the isolation. Scale bar, 100 μ m.

(E) The image of cardiomyocytes isolated from the P7 mouse taken one day after the isolation. Scale bar, 100 μm.

(F) The image of cardiomyocytes isolated from the P56 mouse taken on the day of isolation. Scale bar, 100 μ m.

Supplementary Figure S2. Validation of RNA-seq results with quantitative PCR analyses.

(A) The quantitative PCR (qPCR) analysis of *Igfbp2*, a hepato-JAG, in the hepatocytes and cardiomyocytes. Data were normalized by *Polr2a*.

(B) The qPCR analysis of *Pleiotrophin*, a cardio-JAG, in the hepatocytes and cardiomyocytes. Data were normalized by *Polr2a*.

(C) The qPCR analysis of common JAGs in the hepatocytes and cardiomyocytes.

Data were normalized by Polr2a.

*p < 0.05, **p < 0.01, Student's t test. Data are represented as mean ± SEM.

Supplementary Figure S3. Association of the common JAGs to the human diseases.

The rates for the successful annotation of the common JAGs and all the genes to any human disease.

Supplementary Table S1. Sequences of the qPCR primers used in this study.

Gene	Forward	Reverse
Polr2a	5'-GAGTCCAGAACGAGTGCATGA-3'	5'-ACAGGCAACACTGTGACAATC-3'
lgfbp2	5'-CAGACCTCGGGTGAGAAAAG-3'	5'-CTGCTACCACCTCCCAACAT-3'
Pleiotrophin	5'-TTTTCATCTTGGCAGCTGTG-3'	5'-ACACTCCACTGCCATTCTCC-3'
Ezh2	5'-CCTGTTCCCACTGAGGATGT-3'	5'-GAGCCGTCCTTTTTCAGTTG-3'
Gpc3	5'-ACGGGATGGTGAAAGTGAAG-3'	5'-AGGTGGTGATCTCGTTGTCC-3'
Uhrf1	5'-ACGGTGCCTACTCATTGGTC-3'	5'-GCTTCTGGTCAGAGGACTGG-3'
Postn	5'-TGGTCACTTCACGCTCTTTG-3'	5'-GCCACTTTGTCTCCCATGAT-3'
Map4k4	5'-CTGGGTCCATCACAGACCTT-3'	5'-TCGGTGAATAACGTGGTGAA-3'
Prmt1	5'-GCCTGCAAGTGAAGAGGAAC-3'	5'-CTCAGGACTGGTGGAGAAGC-3'
Tia1	5'-AGATGCCCGTGTGGTAAAAG-3'	5'-TTCTGCATCCCATTTGTTGA-3'
Sirt6	5'-CCTGTAGAGGGGAGCTGAGA-3'	5'-GAGGTACCCAGGGTGACAGA-3'

Supplementary Figure S1. Appearance of the isolated hepatocytes and cardiomyocytes.

A P1 B P7 C P56

Images of the isolated hepatocytes

Images of the isolated cardiomyocytes



A qPCR with a hepato-JAG **B** qPCR with a cardio-JAG Pleiotrophin lgfbp2 2.0 2.0 Hepatocytes P1 P7 P56 1.8 1.8 8.0 Kelative expression 8.0 expression 8.0 A.1.0 8.0 A.1 1.6 1.6 Relative expression 1.4 1.2 Cardiomyocytes 1.0 P1 P7 P56 0.8 0.6 0.4 0.2 0.2 0 0 **C** qPCR with common JAGs 6.0 5.0 4.0 3.0 2.0 Relative expression 1.4 1.2 1.0 0.8 0.6 0.4 0.2 0

Postn

Map4k4

Prmt1

Tia1

Sirt6

Ezh2

Gpc3

Urhf1

Supplementary Figure S2. Validation of RNA-seq results with quantitative PCR analyses.

Supplementary Figure S3. Association of common JAGs with human diseases.

