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## 4 Supplementary Fig. 1 Differences in the expression and secretion of S100A9 in 5 macrophages and cardiomyocytes.

6 **a**, S100A9 expression in macrophages and cardiomyocytes. **b**, S100A9 expression in 7 THP-1 macrophages exposed to high glucose (33mM). **c**, S100A9 secretion in 8 macrophages and cardiomyocytes exposed to high glucose (33mM). **d**, schematic 9 diagram of co-cultured macrophages and cardiomyocytes. **e**, S100A9 level in 10 co-cultured supernatant. **f**, S100A9 expression in co-cultured macrophages and 11 cardiomyocytes. \*p < 0.05, \*\*p < 0.01, \*\*\*p < 0.001. All data are presented as 12 mean  $\pm$  SD. Statistical significance was determined by one-way ANOVA.

Supplementary Figure 2



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16 Supplementary Fig. 2 Blood glucose and lipid levels in diabetic mice.

a, OGTT (Oral Glucose Tolerance Test). b, ITT (Insulin Tolerance Test). c, random
blood glucose. d, fasting blood glucose. e, insulin levels. f, total triglyceride (TG). g,
total cholesterol (T-CHO). \*p < 0.05, \*\*p < 0.01, \*\*\*p < 0.001. All data are presented</li>

as mean  $\pm$  SD. Statistical significance was determined by unpaired student's t-test (2f,

21 2g), or one-way ANOVA (2c-2e).



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25 Supplementary Fig. 3 Quantification of heart morphology and fibrosis.

Quantification of body weight, heart weight (HW), HW/TL(tibial length), Masson-trichrome staining, and Sirius red staining in STZ/HFD-induced or db/db diabetic mice (**a-e**) and mice with cardiac-specific overexpression of S100A9 (**f-j**). \*p< 0.05, \*\*p < 0.01, \*\*\*p < 0.001. All data are presented as mean ± SD. Statistical

30 significance was determined by one-way ANOVA.



# Supplementary Fig. 4 S100A8 expression in diabetic mice and macrophage specific S100a9 knockout mice.

a, Serum S100A8 concentration in diabetic mice (n=7, STZ group; n=9, db/db group).
b, Serum S100A8 concentration in macrophage specific S100a9 knockout mice (n=7).
c, S100A8 expression in BMDM (n=5) and heart (n=6) of macrophage specific
S100a9 knockout mice. \*p < 0.05, \*\*p < 0.01, \*\*\*p < 0.001. All data are presented as</li>
mean ± SD. Statistical significance was determined by unpaired student's t-test

(4b-4c), or one-way ANOVA (4a).





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## Supplementary Fig. 5 Quantification of MitoSOX and JC-1 staining in AC16 cells.

AC16 cells exposed to rhS100A9 (2µg/mL), high glucose (33mM), and/or paquinimod (PAQ, 20µM). **a** and **c**, MitoSOX staining. **b** and **d**, JC-1 staining. \*p <0.05, \*\*p < 0.01, \*\*\*p < 0.001. All data are presented as mean ± SD. Statistical significance was determined by one-way ANOVA.

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