# MKL-1 suppresses ferroptosis by activating system Xc- and increasing glutathione synthesis

#### **Supplementary materials**

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Figure S1



Figure S2











Figure S7





#### **Figure legends**

**Figure S1.** Construction of ferroptosis stress model. A. CCK-8 assay was used to detect cell viability with different concentrations of ferroptosis inducer Erastin. B. CCK-8 assay was used to detect cell viability with different concentrations of ferroptosis inducer Erastin and ferroptosis inhibitor Fer-1. C. After the addition of ferroptosis inducer Erastin and ferroptosis inhibitor Fer-1, the Lipid ROS level of cells was detected by C11 BODIPY 581/591 probe and FCM. D. After the addition of ferroptosis inducer Erastin and ferroptosis inhibitor Fer-1, the level of the membrane lipid peroxidation product MDA was detected by the MDA lipid oxidation level detection kit. E. Typical images of mitochondria in MGC-803 and HGC-27 cells were observed by transmission electron microscopy at 80KV. The t-test was used to determine the statistical significance for both groups of analysis. The CCK-8 assay, Lipid ROS assay and MDA assay results represented the average values obtained from three independent experiments. The results presented were Mean  $\pm$  SD. \* p < 0.05, \*\* p < 0.01, \*\*\* p< 0.001, ns = p > 0.05.

**Figure S2.** Knockdown efficiency of MKL-1. A. The protein expression of MKL-1 in cells was detected by Western Blot. B. Image J software was used for the grayscale analysis of Western Blot results. C. The mRNA expression of MKL-1 was detected by qRT-PCR. D. The protein expression of MKL-1 in human gastric cancer cell lines AGS, MKN-45, HGC-27 and SGC-7901 were detected by Western Blot. The t-test was used to determine the statistical significance for both groups of analysis. The Western blot and qRT-PCR results represented the average values obtained from three independent

experiments. The results shown are Mean  $\pm$  SD. \*\* p < 0.01.

**Figure S3.** Overexpression of MKL-1 promoted cystine uptake through System Xc-. A. Cystine (U-<sup>13</sup>C<sup>6</sup>, U-<sup>15</sup>N<sup>2</sup>) fed cells were used for analysis and GSH biosynthesis pathway by LC/MS. B. The isotope tracks the metabolism of cystine in the cell. The t-test was used to determine the statistical significance for both groups of analysis. The results shown are Mean  $\pm$  SD. \* p < 0.05, \*\* p < 0.01, \*\*\* p < 0.001, ns = p > 0.05.

**Figure S4.** Overexpression of MKL-1 reduces ferroptosis induced by Erastin in both in vitro and in vivo. A. The level of GSH in cells was detected by GSH detection kit. B. The mRNA expression of SLC3A2 and SLC7A11 in cells was detected by qRT-PCR. C. Typical images of the subcutaneous tumor tissue of nude mice after Erastin treatment (n = 3-5 tumors/group). D. The mass of subcutaneous tumor tissue in nude mice after Erastin treatment. E. The volume of subcutaneous tumor tissue in nude mice after Erastin treatment. E. The volume of subcutaneous tumor tissue in nude mice after Erastin treatment. F. The level of Lipid ROS in subcutaneous tumor tissue of nude mice was detected by C11 BODIPY 581/591 probe and FCM. G. The level of GSH in subcutaneous tumor tissue of nude mice was detected by GSH detection kit. H. Pearson correlation analysis of SLC3A2, SLC7A11 and MKL-1 mRNA expression in subcutaneous tumor tissue of nude mice. The t-test was used to determine the statistical significance for both groups of analysis. The GSH assay and qRT-PCR results represented the average values obtained from three independent experiments. The results shown are Mean  $\pm$  SD. \* p < 0.05, \*\* p < 0.01, \*\*\* p < 0.001, ns = p > 0.05.

Figure S5. Identification of MSCs. A. Typical images of MSCs cells under an inverted microscope. B. Typical images of alizarin red staining after 14 days of osteogenic

induction. C. Typical images of oil red O staining after 14 days of adipogenic induction. D. The expression of MSCs markers CD90, CD105, CD34 and CD45 was detected by FCM. The FCM result represented the average values obtained from three independent experiments. The results presented were Mean  $\pm$  SD.

**Figure S6.** Extraction and identification of MSCs exosmoes. A. Typical images of exosomes observed by transmission electron microscopy at a voltage of 80KV. B. NTA analysis of exosome particle size. C. Western Blot detection of CD9, CD63 and Calnexin in MSCs supernatant and MSCs Exo Expression. D. Image J software for grayscale analysis of Western Blot results. The t-test was used to determine the statistical significance for both groups of analysis. The Western blot results represented the average values obtained from three independent experiments. The results presented were Mean  $\pm$  SD in three independent trials. The results presented were Mean  $\pm$  SD. \*\*\* p < 0.001.

**Figure S7.** MSCs Exo as a carrier to insert miR-149-5p and exogenous MKL-1 siRNA. A. Typical images of MSCs after lentivirus infection. B. The expression of miR-149-5p in cells and its Exo was detected by qRT-PCR. C. After transfection of MKL-1 siRNA exosome treatment 72 hours, the protein expression of MKL-1 in cells was detected by Western Blot. D. After transfection of MKL-1 siRNA exosome treatment, the mRNA expression of MKL-1 in cells was detected by qRT-PCR. The t-test was used to determine the statistical significance for both groups of analysis. The qRT-PCR result represented the average values obtained from three independent experiments. The results presented were Mean  $\pm$  SD. \*\* p < 0.01, \*\*\* p < 0.001. Figure S8. Therapeutic impact of exosomes on xenograft tumor models in nude mice.

A. The volume of subcutaneous tumors formed in nude mice (n = 4-8 tumors/group). Multiple t-tests (followed by post-hoc test Holm-Sidak) was used to determine the statistical significance against control group. B. The Kaplan-Meier survival curve of nude mice (n = 10 tumors/group). Log-rank was used to determine the statistical significance. ns = p > 0.05.