

Supplementary Figure 1. miR-4500 mediates $\mathrm{H}_{2} \mathrm{O}_{2}$-induced endothelial cell aging and dysfunction by targeting ARG2. HUVECs were treated with $100 \mathrm{H}_{2} \mathrm{O}_{2}$ for 24 h , then transduced with miR4500 mimics or miR-4500-MUT mimics harboring mutated ARG2-binding sites for 24 h , the cells were subjected to (A) qPCR analysis of ARG2 mRNA expression. (B) SA- $\beta$-gal staining for detecting senescent cells. Quantification is presented below the images. (C) DCFH-DA and DAF2DA staining for the detection of ROS and NO. (D) Quantification of the ROS and NO signals in (C). Scale bar $=100 \mu \mathrm{~m} . \mathrm{n}=4,{ }^{* *} \mathrm{p}<0.01, * * * \mathrm{p}<0.001$. n.s.: not significant.


Supplementary Figure 2. The IncRNA OIP5-AS1/miR-4500 axis targeting ARG2 modulates $\mathrm{H}_{2} \mathrm{O}_{2}$-induced premature cellular senescence and endothelial dysfunction HAECs. HAECs were treated with $100 \mathrm{H}_{2} \mathrm{O}_{2}$ for 24 h , then transduced with miR4500 mimics or OIP5-AS1 shRNA for 24 h . The cells were subjected to (A) qPCR analysis of ARG2 mRNA expression. (B) SA- $\beta$-gal staining for detecting senescent cells. Quantification is presented on the right panel. (C) DCFH-DA and DAF-2DA staining for the detection of ROS and NO. Quantification is presented on the right panel. Scale bar $=100 \mu \mathrm{~m} . \mathrm{n}=4,{ }^{* *} \mathrm{p}<0.01, * * * \mathrm{p}<0.001$.

A
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B


Supplementary Figure 3. The expression levels of miR-4500 and OIP5-AS1 in blood serum samples from young and elder people. The qPCR analysis of (A) miR-4500 expression, and (B) OIP5-AS1 expression in the blood samples of young and old people with atherosclerosis. ARG2 mRNA was not detectable in the serum from either group. $\mathrm{n}=8,{ }^{* *} \mathrm{p}<0.01,{ }^{* * *} \mathrm{p}<0.001$.


Supplementary Figure 4. OIP5-AS1 promotes endothelial cell senescence via sponging miR-4500. HUVECs were transfected with OIP5-AS1-WT or OIP5-AS1-MUT harboring mutated miR-4500binding sites. The cells were subjected to (A) qPCR analysis of ARG2 mRNA expression. (B) SA- $\beta$ gal staining for detecting senescent cells. Quantification is presented below the images. (C) DCFH-DA and DAF-2DA staining for the detection of ROS and NO. (D) Quantification of the ROS and NO signals in (C). Scale bar $=100 \mu \mathrm{~m} . \mathrm{n}=4,{ }^{*} \mathrm{p}<0.05,{ }^{* *} \mathrm{p}<0.01,{ }^{* * *} \mathrm{p}<0.001$.

