

Long-term exposure of human endothelial cells to metformin modulates miRNAs and isomiRs

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SUPPLEMENTARY FIGURES

Supplementary Figure S1. Pairwise sample-to-sample scatter plots of normalized counts from miR-seq analysis of n=3 SEN and n=3 SEN+M samples.

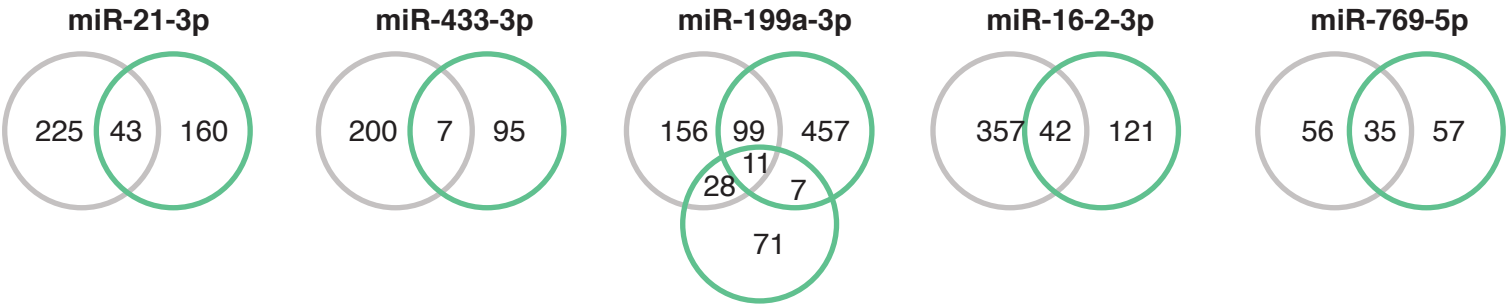
Supplementary Figure S2. Number of predicted target genes of canonical (gray) and 5' isomiRs (green) of miRNAs including at least one 5'isomiR showing a significant linear (group 1) or U-shaped/inverted U-shaped (group 2) trend in Young, SEN, and SEN+M samples.

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Group 1 (linear trend)



Group 2 (U-shaped / inverted U-shaped trend)

