

Measurement of endothelial dysfunction markers. Briefly, to 100µL of serum 50µL of internal standard containing solution (50µM d7-ADMA, d4-cystine, d8-homocystine, and d3-methionine; all in HPLC-grade water) and 40µL of 500mM DTT solution were added. For protein precipitation, 400µL of 0.05% trifluoric acid plus 0.1% formic acid containing acetonitrile solution were added to the sample. This solution also contained 500nM d3-SAM and d5-SAH as internal standards for adenosine, SAM and SAH analysis. The sample was finally vortexed for 5 minutes, centrifuged for 10 minutes at 13,000g and transferred into a HPLC vial.

20µL of the supernatant were injected onto a 4.6x12.5 mm guard/extraction column (Eclipse XDB-C8, 5µm, Agilent Technologies, Palo Alto, CA) inline with a 3.0x150 mm analytical column (RP-Amide, 3.5µm, Supelco, St. Louis, MI). For adenosine, SAM and SAH analysis, the starting mobile phase concentrations consisted of 5% acetonitrile and 95% 10 mM ammonium formate buffer (pH 3.4) with a flow of 0.6mL/min for the first minute. After one minute, the flow rate was increased to 0.8mL/min and a gradient from 5% to 95% acetonitrile within 2.5 min was run. Acetonitrile was then held at 95% for 0.5 minutes. The column was re-equilibrated for 1 min to starting conditions.

For all other compounds, the gradient started at 3% methanol and 97% 10mM ammonium formate buffer and was maintained at a flow of 0.8mL/min throughout the assay. At minute 4.5, the solvent gradient reached 25% methanol; after this the methanol content was raised to 98% and held for additional 1.5min. Hereafter the columns were re-equilibrated to the starting conditions for the remaining 2 minutes of the assay.

Prostaglandins (P) were extracted from 200 µL plasma following protein precipitation with 300 µL methanol/ 0.2M ZnSO₄ (70/30 v/v). Automated inline extraction was used for sample analysis. 425 µL of the extracted sample were injected onto a 4.6x12.5 mm Zorbax XDB-C8 guard/ extraction column (Agilent Technologies). The flow rate during the injection was 500µL/min and was increased to 3000 µL/min within one minute. The composition of solvents was 80% of aqueous 0.1% formic acid and 20% of methanol. After sample cleanup for one minute, the switching valve was activated and the analytes were back-flushed onto the 3x150mm 3.5 micron Zorbax XBD-C8 analytical column (Agilent Technologies). A solvent gradient of methanol and 0.1% aqueous formic acid was used for the chromatographic separation of study compounds within the next 11 minutes.

Bioactive lipid mediators. Briefly, to 200 µL of serum, 800 µL methanol/ ZnSO₄ (70:30, v/v)

protein precipitation solution containing the internal standards (2 ng/mL mixture of internal standards, see below) were added. Samples were vortexed for 10 minutes, centrifuged for 10 minutes at 13,000g and transferred into HPLC vials. Fifty μ L of the supernatants were injected onto a 4.6x12.5 mm guard column (Eclipse XDB-C18, 5 μ m, Agilent Technologies) and then back-flushed with 100% acetonitrile onto a 1.0x250 mm analytical column (Luna C18(2), 3 μ m, 100A, Phenomenex, Torrance, CA). For HPLC separation, the starting mobile phase consisted of 65% acetonitrile/0.1% formic acid (9:91 v/v) and 35% acetonitrile with a flow of 0.2 mL/min for the first minute. After 2 minutes, the gradient consisted of 82% acetonitrile and was increased to 95% acetonitrile within 8 minutes. Acetonitrile was then held at 95% for 2 minutes. The column was re-equilibrated for 2 min to starting conditions.

Krebs cycle intermediates and purine metabolites. Analytes were separated using a 150 x 3 mm Luna HILIC, 3 μ m column (Phenomenex) at an HPLC solvent flow rate of 450 μ L/min. The solvents were 0.1% aqueous formic acid (mobile phase A) and acetonitrile (mobile phase B). The gradient was: 0-1 min 5% acetonitrile, 1.0-3.5 min 5% acetonitrile to 15% acetonitrile, 3.5-4.5 min 100% acetonitrile. The column was then re-equilibrated to starting conditions (5% acetonitrile) between 4.6 and 5.5 min. The mass spectrometry parameters were identical for both positive (0 - 1.79 min) and negative (1.8 - 5.5 min) periods and were as follows; ion source gas one: 40, ion source gas two 45, source temperature 500°C, collision gas 10, curtain gas 20, and ion source voltage 4500V.