© 2009 International Society of Nephrology

The kidney is the major site of S-adenosylhomocysteine disposal in humans

Giacomo Garibotto^{1,2}, Alessandro Valli^{1,2}, Björn Anderstam³, Monica Eriksson³, Mohamed E. Suliman³, Manrico Balbi^{1,2}, Daniela Rollando^{1,2}, Emanuela Vigo^{1,2} and Bengt Lindholm³

¹Department of Internal Medicine, University of Genoa, Genoa, Italy; ²Department of Cardionephrology, University of Genoa, Genoa, Italy and ³Divisions of Renal Medicine and Baxter Novum, Karolinska Institutet, Karolinska University Hospital at Huddinge, Stockholm, Sweden

S-adenosylhomocysteine (SAH), the metabolic precursor of homocysteine in the body, is a potent inhibitor of methylation reactions. Several methylation reactions play a major role in epigenetic regulation of protein expression, atherosclerosis, and cancer development. Here we studied the mechanisms responsible for the maintenance of circulating SAH levels by measurement of the arterio-venous differences across the kidney, splanchnic organs, and the lung in humans. The lungs did not remove or add any circulating SAH, whereas the liver released it into the hepatic veins. The kidney extracted 40% of SAH and the SAH arterio-venous difference across the kidney was directly and significantly related to its arterial levels. Thus, the kidney plays a major role in maintaining SAH levels and may, indirectly, control tissue transmethylation reactions. Our findings of a pivotal role for the human kidney in sulfur amino acid metabolism may also account for the increased plasma levels of SAH in patients with chronic kidney diseases.

Kidney International (2009) **76**, 293–296; doi:10.1038/ki.2009.117; published online 8 April 2009

KEYWORDS: atherosclerosis; epigenetics; homocysteine; methionine; methylation; S-adenosylhomocysteine

The biological effects of methionine transmethylation, whereby the methyl group of S-adenosylmethionine (SAM) is donated to a large variety of acceptor substrates, is of obvious importance for the biosynthesis of a wide range of compounds such as membrane phospholipids, neurotransmitters, proteins, creatine, and hormones.¹ Methylation processes also play a major role in the epigenetic regulation of protein expression and changes in human DNA methylation patterns are an important feature of many diseases, including atherosclerosis and cancer.^{2,3} S-Adenosylhomocysteine (SAH) is the by-product of methionine transmethylation and the metabolic precursor of homocysteine in all tissues (Figure 1). As SAH is a potent feedback inhibitor of most methyltransferases,⁴ including the methionine remethylation pathway, this compound plays an essential role in the control of the overall transmethylation rates.^{4,5} Thus, the efficiency of methyltransferase reactions is dependent on the efficient tissue removal of SAH.4,5

About 30 years ago, Wilcken and Wilcken⁶ observed that increased plasma sulfur amino acids levels were associated with atherosclerosis and cardiovascular damage. Among the sulfur compounds, homocysteine has been considered to be the culprit for this association. However, the recent homocysteinelowering intervention trials have cast into doubt the issue of causality.⁷ One potential explanation for these conflicting results is that hyperhomocysteinemia itself may not be the causative agent in vascular dysfunction, but instead may be a marker for another risk factor.^{8,9} Recent studies suggest an indirect mechanism for homocysteine toxicity, secondary to SAH accumulation.9 An increase in intracellular SAH is associated with DNA hypomethylation in endothelial cells.¹⁰ SAH levels are also associated with endothelial dysfunction in mice with a deficiency of the cystathionine β -synthase gene.¹¹ Recent experimental¹² and clinical evidence¹³⁻¹⁵ also suggests that the accumulation of SAH in body fluids, rather than increased homocysteine levels, is associated with vascular disease and tissue damage. Post-translational modification of proteins, associated with high SAH intracellular accumulation, has been described in patients with chronic kidney diseases.^{16,17}

The sites and mechanisms which prevent the body accumulation of SAH in the body are not well understood.⁹

Correspondence: Giacomo Garibotto, Division of Nephrology, Department of Internal Medicine, Viale Benedetto XV, 6, Genoa 16132, Italy. E-mail: gari@unige.it

Received 22 December 2008; revised 4 February 2009; accepted 25 February 2009; published online 8 April 2009

Like plasma homocysteine, plasma SAH levels are inversely related to glomerular filtration rate.^{18,19} However, plasma SAH appears to be more sensitive than plasma homocysteine to small decrements in glomerular filtration rate.¹⁹ In patients with chronic kidney diseases a 45-fold increase in SAH as compared with a sixfold increase in SAM and fourfold increase in plasma homocysteine is observed.¹³ These changes are not likely dependent on different urine excretion of these compounds, as the fractional excretion of SAM and SAH compared with that of creatinine has been reported to be 93% for SAM and 39% for SAH, which may explain the high ratio of SAM to SAH in normal urine.²⁰ Stable-isotope studies in non-diabetic^{21,22} and diabetic chronic kidney disease patients²³ have shown impaired metabolic clearance of homocysteine by both the transsulfuration and the remethylation pathways. It is interesting to note that Stam et al.²² observed that in chronic kidney disease patients the methionine remethylation and transmethylation rates are inversely related to plasma SAH, suggesting a strong inhibitory action of SAH on methionine-dependent remethylation pathway.

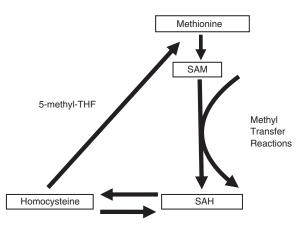


Figure 1 | The metabolic conversion of methionine to homocysteine. SAM = S-adenosylmethionine, SAH = S-adenosylhomocysteine, 5-methyl-THF = 5-methyl-tetrahydrofolate.

In this study, to explore the sites and mechanisms underlying the regulation of circulating SAH levels, we measured plasma SAH across the kidney, splanchnic bed and lung in humans. Our results show for the first time that the human kidney plays a unique role in the removal of SAH from circulation indicating that the kidney may have an important role in the control of body transmethylation reactions.

RESULTS

Individual arterial and venous levels of SAH, as well as their arterio-venous difference across the kidney and splanchnic organs are reported in Table 1. Renal vein SAH concentrations were remarkably lower (by ~40%; P<0.001) than the corresponding arterial values measured in the same subjects, demonstrating that plasma SAH decreases substantially after a single pass across the kidney. Fractional extraction of SAH was twofold greater than that of creatinine (P<0.01). The SAH arterio-venous difference across the kidney was directly related to SAH arterial levels (r=0.72; P<0.02) (Figure 2).

SAH levels in the hepatic veins were $\sim 20\%$ greater than those in the artery (P < 0.02), indicating that splanchnic organs are a major source for SAH into the circulation. No significant A–V gradient of SAH was observed across the lung.

Mean homocysteine levels in the renal and in the liver veins were not statistically different from those in the artery (Femoral artery 10.4 ± 1 , Renal vein 10.3 ± 1 , Liver vein $10.1 \pm 1 \,\mu\text{mol}\,l^{-1}$; P = NS). Similarly, no significant arteriovenous homocysteine gradient was observed across the lung (Femoral artery 10.4 ± 1 , Lung artery $10.1 \pm 1 \,\mu\text{mol}\,l^{-1}$).

DISCUSSION

We observed for the first time that the kidney is a major site for the disposal of plasma SAH in humans. The SAH arteriovenous difference across the kidney represents $\sim 40\%$ of the SAH arterial plasma concentration, positioning the human kidney as a major tissue in the metabolic disposal of plasma SAH. It is interesting to note that the arterio-venous

Table 1 SAH pla	asma levels and fractiona	l extractions across the kidney,	splanchnic organs, and lung
-------------------	---------------------------	----------------------------------	-----------------------------

Subject	Arterial SAH (nmol/l)	Renal vein SAH (nmol/l)	SAH kidney A–V (nmol/l)	SAH kidney FE (%)	Creatinine kidney FE (%)	Liver vein SAH (nmol/l)	SAH splanchnic AV (nmol/l)	SAH splanchnic FE (%)	Lung artery SAH (nmol/l)	SAH lung V–A (nmol/l)
1	26.0	25.0	1.0	+3.8	+22	40.0	-14	-53.8	21.0	-5
2	24.9	19.9	5.0	+20	+21.2	34.0	-9.1	-36.0	24.9	0
3	42.0	20.0	22.0	+52.4	+19	44.0	-2	-4.8	38.0	-4
4	22.6	14.6	7.9	+34.8	+18	25.9	-3.3	-15.0	42.0	19
5	29.9	10.1	19.8	+66.7	+28	33.1	-3.2	-10.5	17.0	-13
6	28.9	14.0	14.9	+51.6	+17	34.6	-5.7	-19.7	34.9	5.2
7	17.8	8.0	9.8	+55.1	+18	19.6	-1.8	-10.1	17.9	0.2
8	17.6	10.4	7.2	+40.9	+23	25.0	-7.4	-42.8	20.4	2.8
9	18.1	11.9	6.2	+34.2	+18	20.0	-1.9	-10.5	21.2	3.1
10	12.2	6.65	5.6	+45.7	+23	13.0	-0.8	-6.1	12.6	0.4
Mean \pm s.e.m.	24.1 ± 2.7	14.1 ± 1.9	10.0 ± 2.2**	40.5 ± 5.8***	21 ± 1	28.9 ± 3.1	$-4.9 \pm 1.12^{*}$	-21 ± 5.0	24.9 ± 3.1	-0.8 ± 2.6

A-V, arterio-venous; FE, fractional extraction (negative extraction means production); SAH, S-adenosylhomocysteine, V-A, venous-arterial.

*P<0.02 and **P<0.001 or less Artery vs Vein; ***P<0.01 vs creatinine FE across the kidney. 'Splanchnic' indicates the metabolic activity of the liver + intestine.

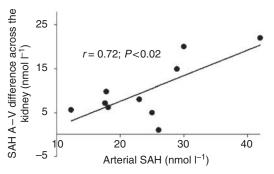


Figure 2 | Relationship between the arterio-venous (A-V) difference of *S*-adenosylhomocysteine (SAH) across the kidney and arterial SAH levels.

difference of SAH is directly related to the arterial levels of the same metabolite, implying that the regulatory role of the kidney on plasma SAH occurs over a wide range of SAH plasma levels. SAH has a molecular mass of 384 D,¹ which is within the filtration range of normal glomeruli. Although SAH may be filtered and lost with the urine, its urinary clearance is remarkably lower than creatinine clearance.²⁰ In our study, the fractional extraction of SAH across the kidney was ~2-fold greater than that of creatinine. This finding strongly indicates that a large part of SAH is removed by kidney metabolic clearance.

As a new finding, we observed that SAH is released by the splanchnic organs. SAH levels in the hepatic veins were markedly greater ($\sim 21\%$) than those in the artery, indicating that splanchnic organs are a major source for SAH entering into the circulation. In contrast, no significant SAH gradient was observed across the lung.

The efficiency of methyltransferase reactions is dependent on the efficient removal of SAH. This is effectively accomplished by SAH hydrolase, an enzyme that appears to act in close proximity to the methyltransferases. SAH hydrolase is highly expressed in the rat kidney and is homogeneously localized in proximal and distal tubule cells as well as in glomeruli.^{24,25} This distribution is likely involved in maintaining efficient transmethylation reactions and thereby low intracellular kidney SAH levels. Considering an average kidney plasma flow of 600 ml/min²⁶ the removal of SAH by the kidney would according to the results of our study amount to $\sim 8.6 \,\mu mol/day$. This figure clearly matches the estimated release of SAH by splanchnic organs (about $\sim 6 \,\mu mol/day$, considering a plasma flow of about 800 ml/ min).²⁶ According to these findings, extracellular SAH pools $(\sim 350 \text{ nmol}, \text{ considering a plasma SAH concentration})$ 25 nmol/l and extracellular fluid 141 in a 70 kg man)²⁷ are composed of only a small fraction (about 4%) of the estimated kidney daily removal, suggesting that the extracellular SAH pool is renewed by tissue SAH generation and SAH kidney removal several times a day.

In conclusion, this study is the first to show that the human kidney is the major site for the metabolic disposition of plasma SAH in the body. These data display a new role for the kidney in the metabolic regulation of sulfur amino acids and, indirectly, of tissue methyl transfer reactions in humans.

MATERIALS AND METHODS Patients and procedures

Ten patients (seven men, three women, mean age 67, range 56-74 years, BMI $24 \pm 1 \text{ kg/m}^2$) who were scheduled for elective cardiac catheterization for hemodynamic evaluation or the assessment of coronary heart disease were eligible for enrollment in this protocol. The patients were enrolled in the study on a consecutive basis if they met the following exclusion criteria: NYHA Class III congestive heart failure, a recent myocardial infarction, pregnancy, or unstable renal function. Their estimated²⁸ glomerular filtration rate was 83 ± 5 ml/min per 1.73 m². No subject had evidence of liver disease or diabetes mellitus. The study was approved by the Ethical Committee of the Department of Internal Medicine of the University of Genoa. Subjects were informed about the purposes, procedures, and possible risks of the study, before their informed consent was obtained. The procedures were in accordance with the Helsinki declaration. Three sets of blood samples were obtained at \sim 20-min intervals from the femoral artery as well as from the renal veins, the hepatic veins, and the pulmonary artery, in the postabsorptive state. Plasma SAH and homocysteine levels were determined in triplicate by HPLC.¹⁵ This method measures both SAH and SAM in body fluids. However, although the intra-assay coefficient of variation for SAH is remarkably low (4.7%), coefficient of variation for SAM analysis is greater ($\sim 10\%$), which makes unlikely the possibility to detect small SAM arterio-venous differences.¹⁵ Therefore, data regarding SAM are not reported. All samples from one individual were always run in the same batch. Blood acid-base, glucose, lactate, and creatinine were measured with an ABL800 Flex apparatus (Radiometer, Copenhagen, Denmark).

Calculations and statistical methods

The arterio-venous difference of SAH and homocysteine across the splanchnic organs, lung, and kidney was calculated as: [A]-[V], where [A] and [V] are the metabolite concentrations in arterial and venous plasma. Fractional extractions were calculated as $100 \times ([A]-[V]/[A])$. Statistical analysis was performed using analysis of variance for repeated measures to compare arterial data with venous data (Statview Statistical Package, Abacus, Berkeley, CA, USA). Linear regression and correlation were employed to evaluate the relation between two variables. A *P*-value of <0.05 was considered statistically significant. All data are expressed as means \pm s.e.

DISCLOSURE

Bengt Lindholm is employed by Baxter Healthcare.

ACKNOWLEDGMENTS

This study was supported by grants from the Ministero dell'Università e della Ricerca Scientifica e Tecnologica (PRIN), the ISN/Baxter Extramural Grant Program, and the University of Genoa, Italy.

REFERENCES

- Finkelstein JD. Regulation of homocysteine metabolism. In: Carmel R, Jacobsen DW (eds). *Homocysteine in health and disease*. Cambridge University Press: Cambridge, UK, 2001, pp 92–99.
- Wilson AS, Power BE, Moloy PL. DNA hypomethylation and human diseases. *Biochim Biophys Acta* 2000; **1775**: 138–162.
- Lee ME, Wang H. Homocysteine and hypomethylation: a novel link to vascular disease. *Trends Cardiovasc Med* 1999; 9: 49–54.

- 4. Cantoni GL. Experimental and clinical roles of S-Adenosylmethionine. In: Borchardt RT, Creveling CR, Ueland PM (eds). *Biological Methylation and Drug Design*. Humana Press: Clifton, 1986, pp 227–238.
- Clarke S, Banfield K. S-adenosylmethionine-dependent methyltransferases. In: Carmel R, Jacobsen DW (eds). *Homocysteine in health and diseases*. Cambridge University Press: Cambridge, UK, 2001, pp 63–78.
- Wilcken DE, Wilcken B. The pathogenesis of coronary artery disease. A possible role for methionine metabolism. J Clin Invest 1976; 57: 1079–1082.
- Maron BA, Loscalzo J. The Treatment of Hyperhomocysteinemia. Annu Rev Med 2008; 60: 39–54.
- Weiss N, Keller C, Hoffmann U *et al.* Endothelial dysfunction and atherothrombosis in mild hyperhomocysteinemia. *Vasc Med* 2002; 7: 227–239.
- Wagner C, Koury MJ. S-Adenosylhomocysteine—a better indicator of vascular disease than homocysteine? Am J Clin Nutr 2007; 86: 1581–1585.
- Castro R, Rivera I, Martins C *et al.* Intracellular S-adenosylhomocysteine increased levels are associated with DNA hypomethylation in HUVEC. *J Mol Med* 2005; **83**: 831–836.
- Dayal S, Bottiglieri T, Arning E *et al.* Endothelial dysfunction and elevation of S-Adenosylhomocysteine in cystathionine ß-Synthase–Deficient Mice. *Circ Res* 2001; 88: 1203–1212.
- Liu C, Wang Q, Guo H *et al.* Plasma S-adenosylhomocysteine is a better biomarker of atherosclerosis than homocysteine in apolipoprotein Edeficient mice fed high dietary methionine. *J Nutr* 2008; **138**: 311–315.
- Loehrer FM, Angst CP, Brunner FP *et al.* Evidence for disturbed Sadenosylmethionine:S adenosylHcy ratio in patients with end-stage renal failure: a cause for disturbed methylation reactions? *Nephrol Dial Transplant* 1998; **13**: 656–661.
- Kerins DM, Koury MJ, Capdevila A et al. Plasma S-adenosyl-Homocysteine is a more sensitive indicator of cardiovascular disease than plasma Homocysteine. Am J Clin Nutr 2001; 74: 723–729.
- 15. Valli A, Carrero JJ, Qureshi AR *et al.* The increase of serum Sadenosylhomocysteine, but not homocysteine, is associated with cardiovascular disease in chronic kidney disease patients. *Clin Chim Acta* 2008; **395**: 106–110.
- Perna AF, Ingrosso D, Zappia V *et al.* Enzymatic methyl esterification of erythrocyte membrane proteins is impaired in chronic renal failure. Evidence for high levels of the natural inhibitor S-adenosyl-Homocysteine. J Clin Invest 1993; **91**: 2497–2503.

- Ingrosso D, Cimmino A, Perna AF *et al.* Folate treatment and unbalanced methylation and changes of allelic expression induced by hyperhomocysteinaemia in patients with uraemia. *Lancet* 2003; **17**: 1693–1694.
- Wollesen F, Brattstrom L, Refsum H *et al.* Plasma total Homocysteine and cysteine in relation to glomerular filtration rate in diabetes mellitus. *Kidney Int* 1999; 55: 1028–1035.
- Jabs K, Koury MJ, Dupont WD *et al.* Relationship between plasma S-adenosyl-homocysteine concentration and glomerular filtration rate in children. *Metabolism* 2006; 55: 252–257.
- Stabler SP, Allen RH. Quantification of serum and urinary S-Adenosylmethionine and S-Adenosylhomocysteine by stable-isotope-dilution liquid chromatography-mass spectrometry. *Clin Chem* 2004; **50**: 365–372.
- Van Guldener C, Kulik C, Berger R *et al.* Homocysteine and methionine metabolism in ESRD: a stable isotope study. *Kidney Int* 1999; 56: 1064–1071.
- 22. Stam F, van Guldener C, ter Wee PM *et al.* Homocysteine clearance and methylation flux rates in health and end-stage renal disease: association with S-adenosylhomocysteine. *Am J Physiol Renal Physiol* 2004; **287**: F215–F223.
- Tessari P, Coracina A, Kiwanuka E et al. Effects of insulin on methionine and homocysteine kinetics in type 2 diabetes with nephropathy. *Diabetes* 2005; 54: 2968–2976.
- Helland S, Ueland PM. S-adenosylhomocysteine and Sadenosylhomocysteine hydrolase in various tissues of mice given injections of 9-beta-D-arabinofuranosyladenine. *Cancer Res* 1983; 43: 1847–1850.
- 25. Kloor D, Stumvoll W, Schmid H *et al.* Localization of S adenosylhomocysteine Hydrolase in the Rat Kidney. *J Histochem Cytochem* 2000; **48**: 211–218.
- Tessari P, Garibotto G, Inchiostro S *et al*. Kidney, splanchnic, and leg protein turnover in humans. Insight from leucine and phenylalanine kinetics. J Clin Invest 1996; **98**: 1481–1492.
- Defronzo RA, Felig P, Ferrannini E *et al.* Effect of graded doses of insulin on splanchnic and peripheral potassium metabolism in man. *Am J Physiol* 1980; **238**: E421–E427.
- Levey AS, Bosch JP, Lewis JB *et al.* A more accurate method to estimate glomerular filtration rate from serum creatinine: a new prediction equation. Modification of diet in renal disease study group. *Ann Intern Med* 1999; **130**: 461–470.