

Soluble form of a receptor for advanced glycation end products (sRAGE) as a biomarker

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TABLE OF CONTENTS

1. Abstract
2. Introduction
3. sRAGE, diabetes and its vascular complications
4. sRAGE and CVD
5. sRAGE and hypertension (HT)
6. sRAGE and chronic kidney disease (CKD)
7. sRAGE and neurodegenerative disorders
8. sRAGE and inflammatory diseases
9. sRAGE and other disorders
10. Conclusion
11. Referemcs

1. ABSTRACT

There is a growing body of evidence that advanced glycation end products (AGEs) and their receptor (RAGE) system are implicated in various disorders, including vascular complications in diabetes, cardiovascular disease, neurodegenerative diseases, inflammatory and autoimmune disorders, and cancer growth and metastasis. Indeed, the engagement of RAGE with AGEs elicits oxidative stress generation and evokes inflammatory and thrombogenic responses, thus playing an important role in these devastating disorders. Moreover, since administration of a recombinant soluble form of RAGE (sRAGE), has been shown to block the AGE-RAGE signaling pathway in animal models, exogenously administered sRAGE may capture and eliminate circulating AGEs, thus protecting against the AGE-elicited tissue damage by acting as a decoy receptor for AGEs. Recently, sRAGE has been identified in humans. However, there are a few comprehensive papers about the regulation and role of sRAGE in humans. Therefore, in this paper, we review the kinetics, regulation and pathophysiological role of sRAGE in humans. We further discuss the potential clinical utility of measuring sRAGE in various disorders as a biomarker.

2. INTRODUCTION

Reactive derivatives from non-enzymatic glucose-protein condensation reactions, as well as lipids and nucleic acids exposed to reducing sugars, form a heterogeneous group of irreversible adducts called “AGEs (advanced glycation end products)”. AGEs were originally characterized by a yellow-brown fluorescent color and an ability to form cross-links with and between amino groups, but the term is now used for a broad range of advanced products of the glycation process (also called the “Maillard reaction”), including N-carboxymethyllysine (CML) and pyrraline, which show neither color nor fluorescence and do not cross-link proteins (1-5). The formation of AGEs *in vitro* and *in vivo* is dependent on the turnover rate of the chemically modified target, the time available, and the sugar concentration. The structures of the various cross-linked AGEs that are generated *in vivo* have not yet been completely determined. Because of their heterogeneity and the complexity of the chemical reactions involved, only some AGEs have been structurally characterized *in vivo* (1-5). The structural identity of AGEs that could mainly mediate their biological actions also remains unknown.

Under the hyperglycemic conditions elicited by diabetes, the glycation process begins with the conversion of reversible Schiff base adducts to more stable, covalently-bound Amadori rearrangement products. Over the course of days to weeks, these Amadori products undergo further rearrangement reactions to form the irreversibly-bound moieties known as AGEs. AGEs could contribute to the aging of proteins and to the pathological complications of diabetes (1-5). Further, there is accumulating evidence that AGEs and their receptor (RAGE) system are implicated in various disorders, including cardiovascular disease, neurodegenerative diseases, inflammatory and autoimmune disorders, and cancer growth and metastasis (6-14). Indeed, the AGE-RAGE interaction elicits oxidative stress generation and subsequently evokes inflammatory and thrombogenic reactions in a variety of cells, thus playing an important role in the development and progression of these devastating disorders (6-14).

Recently, administration of a recombinant soluble form of RAGE (sRAGE) consisting of the extracellular ligand-binding domain, has been shown to not only suppress the development of atherosclerosis but also to stabilize established atherosclerosis in diabetic apolipoprotein E-null mice (15,16). The blockade of the AGE-RAGE axis by administration of sRAGE also ameliorates neuronal dysfunction and reduces the development of acellular capillaries and pericyte ghosts in experimental diabetic retinopathy (17). Furthermore, recently, Kaji *et al.* have also shown that attenuation of the RAGE axis with injection of sRAGE inhibits retinal leukostasis and blood-retinal barrier breakdown in the diabetic RAGE-transgenic mice which are accompanied by decreased expression of vascular endothelial growth factor (VEGF) and intercellular cell adhesion molecule-1 (ICAM-1) in the retina (18). These observations suggest that exogenously administered sRAGE may capture and eliminate circulating AGEs, thus protecting against the AGE-elicited tissue damage by acting as a decoy for AGEs.

Recently, sRAGE has been identified in mice and humans (19-23). Based on the data in sRAGE-administered animals, it is conceivable that sRAGE may also work as a decoy receptor for AGEs. However, since AGEs up-regulate RAGE expression levels in various tissues and that sRAGE could be mainly generated from proteolytic cleavage of membrane-bound RAGE by the actions of sheddase, a disintegrin and metalloproteinase 10 (ADAM 10) (6, 24-26), it is also possible that sRAGE may reflect tissue RAGE expression and the severity of target organ damage. Accordingly, whether sRAGE is a biomarker that could reflect tissue damage or a protective one against injury may be dependent on the disease types. As far as we know, since there are a few comprehensive papers about the regulation and role of sRAGE in humans (27-30), we review here the kinetics, regulation and pathophysiological role of sRAGE in humans. We further discuss the potential clinical utility of measuring sRAGE in various disorders as a biomarker.

3. sRAGE, DIABETES AND ITS VASCULAR COMPLICATIONS

Diabetic vascular complications are one of the leading causes of end-stage renal failure, acquired blindness, a variety of neuropathies and cardiovascular disease (CVD), which could account for disabilities and high mortality rates in patients with both type 1 and type 2 diabetes (31). Various hyperglycemia-induced metabolic and hemodynamic derangements, including increased formation of AGEs, enhanced reactive oxygen species (ROS) generation, activation of protein kinase C (PKC), stimulation of the polyol pathway and the renin-angiotensin system (RAS), are thought to contribute to the pathogenesis of vascular complications in diabetes (31). However, a clinical study, the Diabetes Control and Complications Trial-Epidemiology of Diabetes Interventions and Complications (DCCT-EDIC) Research, has shown that the reduction in the risk of progressive retinopathy and nephropathy resulting from intensive therapy in patients with type 1 diabetes persisted for at least several years, despite increasing hyperglycemia (32,33). In addition, intensive therapy during the DCCT resulted in decreased progression of intima-media thickness (IMT) and subsequently reduced the risk of nonfatal myocardial infarction, stroke, or death from cardiovascular disease by 57 percent 11 years after the end of the trials (34). Further, a recent follow-up study of United Kingdom Prospective Diabetes Study (UKPDS) also has shown that benefits of an intensive therapy in patients with type 2 diabetes are sustained after the cessation of the trial (35). In this study, despite an early loss of glycemic differences between intensive and conventional therapy, a continued reduction in microvascular risk and emergent risk reductions for myocardial infarction and death from any cause (legacy effects) were observed during 10 years of post-trial follow-up (35). These observations indicate that intensive therapy to control blood glucose has long-term beneficial effects on the risk of diabetic retinopathy, nephropathy, CVD and death in patients with both type 1 and type 2 diabetes, thus strongly suggesting that so-called 'metabolic memory' causes chronic damage in diabetic vessels that are not easily reversed, even by subsequent, relatively good control of blood glucose.

Although the exact pathophysiological mechanisms responsible for a legacy effect of the intensive glycemic control or metabolic memory are unclear, biochemical nature of AGEs, *that is*, gradual accumulation of AGEs under hyperglycemic conditions that are subsequently slowly degraded with intensive glycemic control, is compatible with the concept 'metabolic memory' (1-6). Further, AGEs and/or diabetic conditions can up-regulate RAGE expression in various cell types and subsequently induce sustained activation of transcriptional factor nuclear factor-kappaB (NF-kappaB) (6, 36-38). Under diabetic conditions, the AGE-RAGE-induced oxidative stress generation could further potentiate the formation and accumulation of AGEs (15). Therefore, these positive feedback loops between AGEs and RAGE-downstream pathways could make a vicious cycle, thus providing a mechanistic basis for understanding the

concept of ‘metabolic memory’ in vascular complications in diabetes.

sRAGE can generate from cleavage of cell surface full-length RAGE by sheddase or novel splice variants of RAGE as a C-truncated splice isoform of endogenous secretory RAGE (esRAGE) (19-23). Most of the sRAGE (*ca.* 80%) is derived from the cleavage of membrane-bound RAGE, and esRAGE only occupies about 20% of sRAGE (19-23,26). We, along with others, have shown that sRAGE levels are elevated in patients with type 1 and type 2 diabetes and that serum sRAGE levels are positively, rather than inversely, associated with AGE levels in both non-diabetic and diabetic subjects (39-44). Age-, sex- and body mass index-adjusted AGE levels are significantly increased in proportion to the increasing levels of sRAGE in non-diabetic subjects as well (44). These findings suggest that sRAGE could not efficiently capture and eliminate circulating AGEs *in vivo* by working as a decoy receptor for AGEs. Since AGEs are positive regulators of cell expression of RAGE (6,36,37), our present observations suggest that sRAGE levels may reflect tissue RAGE expression and be elevated in parallel with serum AGE levels as a counter-system against the AGE-elicited tissue damage, even in a non-diabetic general population without apparent cardiovascular disease or renal disease. The concept that sRAGE levels may be elevated in response to serum AGE levels and reflect tissue RAGE expression in diabetes is supported by the following observations; (1) RAGE belongs to the same immunoglobulin superfamily as ICAM-1 and vascular cell adhesion molecule-1 (VCAM-1) and that serum levels of soluble forms of ICAM-1 (sICAM-1) and VCAM-1 (sVCAM-1) are elevated in patients with diabetes as biomarkers of endothelial injury by reflecting overexpression of these adhesion molecules in endothelial cells (ECs) (45), (2) Angiotensin II (Ang II) increases RAGE mRNA levels in ECs and subsequently stimulates sRAGE formation *in vitro* (25). Treatment with telmisartan, an Ang II type 1 receptor blocker (ARB), not only inhibits the Ang II-elicited sRAGE generation by ECs, but also decreases serum levels of sRAGE in patients with essential hypertension (25), (3) AGEs up-regulate RAGE expression in various types of cells and that RAGE is up-regulated in atherosclerotic plaques in diabetes, diabetic nephropathy and retinopathy (6,36,37,46-48), (4) Vitreous levels of sRAGE are increased in proliferative retinal diseases by reflecting enhanced RAGE expression in epiretinal membranes of the eyes (48), and (5) Although exogenously administered sRAGE was shown to block the harmful effects of AGEs in animals by acting as a decoy receptor (15-18), it is questionable that endogenous sRAGE could also exert the same biological effect because serum levels of sRAGE in humans are 1000 times lower than needed for the binding AGEs (29,49,50).

The findings that sRAGE levels are positively associated with circulating AGEs and could reflect tissue RAGE expression led us to examine whether sRAGE levels may be a novel biomarker of vascular injury and/or target organ damage in patients with diabetes. To address these issues, we examined whether circulating sRAGE levels

were elevated in type 2 diabetic patients, compared with non-diabetic healthy subjects, and explored the association between sRAGE levels and coronary artery disease (CAD) in diabetic subjects. As a result, we found that serum sRAGE levels were significantly higher in type 2 diabetic patients than in non-diabetic age- and sex-matched control subjects and that the presence of diabetes was a sole independent determinant of sRAGE in our subjects (41). Furthermore, serum sRAGE levels were significantly elevated in diabetic patients with CAD, compared to those without CAD, and one of the independent determinants of CAD in patients with diabetes. In addition, we have recently found that serum levels of sRAGE are positively correlated with inflammatory biomarkers such as tumor necrosis factor-alpha (TNF-alpha), sVCAM-1 and monocyte chemoattractant protein-1 (MCP-1) in type 2 diabetic patients (51-53). Moreover, circulating levels of AGEs and sRAGE were the independent determinants of serum MCP-1 levels in patients with type 2 diabetes. Taken together, these observations suggest the possibility that circulating sRAGE level is a novel biomarker of vascular inflammation and target organ damage in both type 1 and type 2 diabetic patients. In support of this, Tan *et al.* reported that serum sRAGE levels and circulating AGEs were correlated with each other and positively associated with the severity of nephropathy in type 2 diabetes (42). Yan *et al.* also reported that sRAGE levels were significantly higher in type 2 diabetic patients with CAD, compared with those without CAD (54). Further, EURODIAB Prospective Complications Study revealed that patients of type 1 diabetes with CVD had higher levels of sRAGE than those without CVD or any microvascular complications and that endothelial damage, renal dysfunction and low-grade inflammation could partially account for the associations between sRAGE and prevalent CVD (55). In this study, sRAGE levels tended to be higher in the presence and across the levels of severity of albuminuria and retinopathy as well (55). In addition, a recent 12-year follow-up study published in the meeting of EASD 2009 showed that higher plasma sRAGE levels were associated with incident cardiovascular morbidity and mortality as well as all-cause mortality in type 1 diabetes (56).

RAGE mediates the AGE-signaling to MCP-1 and TNF-alpha expression in various types of cells including ECs (57-59). Further, there are several papers to show the positive correlation between serum levels of MCP-1 and TNF-alpha and circulating AGE levels (60,61). These observations suggest that enhancement of the AGE-RAGE axis under diabetic conditions could account for the positive association between sRAGE levels and these inflammatory markers in our patients. Recently, Basta *et al.* reported that plasma sRAGE (measured by the same commercial enzyme-linked immunosorbent assay kit as ours) was inversely associated with the inflammatory marker C-reactive protein (CRP) and hemoglobin A1c which were also independently associated with sRAGE on stepwise regression analysis in their subjects (diabetic patients and controls) (62). However, in their study, CRP was not identified as independent predictors of plasma sRAGE in diabetic patients alone. Therefore, our data to

show the positive association between sRAGE and some inflammatory markers in diabetics are not necessarily discrepant against their findings. Further, recently, sRAGE, but not esRAGE, has been shown to be independently correlated with albuminuria in type 2 diabetic patients (49). sRAGE itself might be capable of triggering inflammatory reactions via binding of Mac-1 and subsequent activation of NF-kappaB, thus being involved in diabetic vascular complications (63).

Although number of patients was very small (one included 30 subjects, and the other 50), there were two papers to show that sRAGE levels were decreased in type 2 diabetic patients (62,64). Devangelio *et al.* reported that sRAGE levels were significantly decreased in type 2 diabetic patients and that HbA1c was inversely and independently associated with sRAGE (64). Basta *et al.* also showed that decreased sRAGE levels in type 2 diabetic patients were associated with the increased CVD risk (62). In addition, Grossin *et al.* reported that although serum sRAGE levels in type 2 diabetic patients and healthy controls were similar, sRAGE levels were significantly decreased in the diabetic patients with renal and retinal complications compared with those without complications (65). At present, we do not know the reasons for the opposite results in type 2 diabetic patients. The difference of subject population (age and duration of diabetes) and ethnicity could account for the discrepancies. Anyway, we should present the possibility of the other side of story, *that is*, sRAGE levels could be inversely associated with the risk of vascular injury in type 2 diabetic patients. However, it is unlikely that decreased *rather than* increased sRAGE level could be a biomarker of vascular complications in diabetes, because as far as we know, there are no papers to show that (1) sRAGE efficiently captures and eliminates circulating AGEs in diabetic patients by working as a decoy receptor for AGEs, (2) sRAGE levels are inversely associated with serum levels of AGEs in diabetics, and (3) sRAGE levels are decreased in type 1 diabetic patients compared with controls. Since interaction of RAGE with the ligand promotes the RAGE shedding (26), sRAGE levels could correlate with high levels of ongoing inflammation in diabetes.

There is still some controversy over the therapeutic modulation of sRAGE. Forbes *et al.* reported that treatment of ramipril, an angiotensin-converting enzyme inhibitor (ACEI) restored the decreased plasma levels of sRAGE in diabetic rats (66). Similarly, they also found that there was a significant increase in plasma sRAGE levels in patients who had type 1 diabetes and were treated with an ACEI perindopril (66). As described above, Nakamura *et al.* reported that telmisartan, an ARB, inhibited sRAGE generation by Ang II-exposed ECs and decreased serum levels of sRAGE in patients with essential hypertension (25). Although atorvastatin or rosiglitazone has been shown to increase serum sRAGE levels (67,68), whether modulation of circulating sRAGE levels has a beneficial effect on the progression of diabetic vascular complications is unknown.

Serum levels of esRAGE are also positively

correlated with circulating AGEs such as pentosidine and CML in both type 1 and type 2 diabetes (69,70). esRAGE levels were increased levels in type 2 diabetic patients with decreased renal function, and serum creatinine was one of the independent determinants of esRAGE levels in these patients (69). However, in contrast to the case of sRAGE, esRAGE levels are decreased, rather than increased, in both type 1 and type 2 diabetic patients with microalbuminuria or retinopathy, compared with those without microvascular complications (71-73). Further, esRAGE levels were inversely associated with carotid IMT in both type 1 and type 2 diabetes, and one of the independent risk factors for the progression of carotid atherosclerosis in patients with type 1 diabetes (74,75). In addition, Koyama *et al.* reported that esRAGE levels were decreased in Japanese type 2 diabetic patients compared with non-diabetic subjects and that its low levels were associated with the components of the metabolic syndrome and carotid atherosclerosis (76). Decreased esRAGE levels were associated with inflammation, arterial stiffness and severity of CAD in patients with type 2 diabetes (77,78). These observations were contrary to the above-mentioned findings that sRAGE levels were associated with conventional coronary risk factors, including inflammatory biomarkers, and one of the independent determinants of CAD in diabetic patients. Therefore, the kinetics and pathophysiological role of sRAGE and esRAGE could differ in diabetes. Decreased levels of esRAGE may be associated with co-morbidity such as diabetic microangiopathy and accelerated atherosclerosis in unknown mechanisms other than working as a decoy receptor for AGEs because (1) as mentioned above, esRAGE levels are approximately 3~5-fold lower than sRAGE and are not sufficient to efficiently eliminate circulating AGEs in humans (29,49,50) and (2) there are no reports to show the inverse associations between esRAGE and AGE levels. We should also note that there was a controversy about the associations between decreased esRAGE levels and vascular complications in diabetes. esRAGE levels were reported not to be associated with albuminuria, carotid IMT, peripheral and autonomic neuropathy in type 2 diabetic patients (49,79). Sakurai *et al.* reported that there was no significant difference of esRAGE levels between type 1 diabetic patients and healthy controls (73).

4. sRAGE AND CVD

Plasma CML and sRAGE levels were elevated after percutaneous coronary interventions with stenting, and positively associated with the extent of vascular injury (80). Further, Basta *et al.* reported that plasma sRAGE levels were higher in non-diabetic patients with symptomatic carotid atherosclerosis than those without symptom (81). They also showed that in symptomatic patients, plasma levels of sRAGE were positively associated with CML, and CML and RAGE levels in carotid plaques were correlated with each other (81). These observations suggest that higher levels of sRAGE may be a biomarker of a high degree of vascular inflammation in non-diabetic patients with CAD as well. It is interesting that, as described before, the same authors previously reported that sRAGE levels were inversely associated with

coronary risk factors in type 2 diabetes (62). These findings suggest that the regulation of sRAGE and its association with vascular disease may differ between patients with and without diabetes.

Koyama *et al.* have shown that serum sRAGE levels are increased with advancing New York Heart Association functional class, and higher in patients with cardiac events than in event free patients (82). In their study, sRAGE and serum pentosidine levels were independent risk factors for cardiac events in the multivariate Cox proportional hazard analysis, thus suggesting that serum sRAGE level is an independent prognostic factor for heart failure (82). In addition, elevated plasma levels of sRAGE were positively associated with primary graft dysfunction after lung transplantation as well (83).

Although the above-mentioned observations suggest that increased sRAGE levels is a biomarker of CVD, opposite scenario-*that is*, decreased sRAGE level could be related with CVD, was also presented by some researchers. Indeed, increased CRP and decreased sRAGE levels were observed in non-diabetic patients with angiographically proven CAD, compared with those with coronary risk factors and without CAD (84). In addition, Falcone *et al.* reported that, independent of established vascular risk factors and lipid parameters, the odds ratio for the presence of CAD in non-diabetic male subjects was 6.7 when the lowest quartile of the sRAGE levels (below 766 pg/ml) compared with the highest quartile (85). Further, plasma levels of sRAGE were positively associated with endothelial function and major cardiovascular event-free survival in non-diabetic subjects (86). Meanwhile, we have recently found that sRAGE levels are independently and inversely associated with high mobility group box 1 (HMGB1) levels in an apparently healthy population (87). HMGB1 is one of the ligands for RAGE, and HMGB1 and RAGE interaction promotes chemotaxis and maturation of immune cells, enhances the expression of adhesion molecules, and stimulates the production of cytokines by various types of cells (88). Moreover, sRAGE is absent and HMGB1 levels are higher in diabetic RAGE^{-/-}/apoE^{-/-} mice, compared with diabetic apoE^{-/-} mice (89). These observations suggest that sRAGE may protect against CVD by working as a decoy receptor for circulating HMGB1. Binding affinity of HMGB1 to RAGE is 10-times higher than that of AGEs (50, 90), whereas serum concentration of HMGB1 is 1000-times less than that of AGEs (87,91), thus supporting the concept that circulating HMGB1 *but not* AGEs is a molecular target for sRAGE. Therefore, the differences of kinetics and serum concentrations of the RAGE ligands (ex. AGEs and HMGB1) in various disorders could account for the discrepant results between sRAGE levels and the disease status.

5. sRAGE AND HYPERTENSION (HT)

Plasma levels of sRAGE were decreased in patients with essential HT, compared with age- and sex-matched normotensive subjects (92). Further, sRAGE levels were inversely associated with pulse pressure, thus

suggesting that sRAGE may be a marker of arterial stiffness in essential HT (92). When we analyzed anthropometric, metabolic and inflammatory variables, including sRAGE in 271 consecutive non-diabetic outpatients with essential HT, 24-hour creatinine clearance ($p<0.0001$, inversely), gamma-glutamyltranspeptidase ($p<0.001$), body mass index ($p=0.007$, inversely), and TNF- α ($p=0.024$) were independently associated with sRAGE levels (93). These findings suggest that anthropometric and inflammatory variables and liver and renal function may be the determinants of sRAGE levels in non-diabetic hypertensive patients. The exact reasons for the discrepant results between the two papers, *that is*, an inverse correlation between sRAGE and pulse pressure in one hand and a positive correlation between body mass index and TNF- α with sRAGE in the other, are unclear. Difference of patient population, disease activity, and concentrations of serum levels of AGEs and HMGB1 between the two studies could affect the results.

6. sRAGE AND CHRONIC KIDNEY DISEASE (CKD)

Serum AGE, HMGB1, sRAGE, and an endogenous inhibitor of nitric oxide synthase, asymmetric dimethylarginine (ADMA) levels in CKD patients were significantly higher than those in control subjects (94). In addition, AGE and sRAGE levels were correlated with each other, and AGEs and HMGB1 were independently associated with ADMA in non-diabetic CKD patients (94). These findings suggest that elevation of the RAGE ligands may enhance ADMA levels, suggesting the active involvement of AGE/HMGB1-RAGE-ADMA axis in CKD patients.

sRAGE levels were increased in patients with decreased renal function, especially those with end-stage renal failure (95). There was a significant inverse relationship between estimated glomerular filtration rate (eGFR) and serum sRAGE levels (96). However, significant inverse relationships were also found for sRAGE to IMT and plaque number in patients with CKD (96). Further, Gross *et al.* reported that low sRAGE levels were associated with a 2-3 times higher risk for mortality in renal transplant recipients especially after correction for creatinine clearance (97). In addition, low circulating sRAGE is reported to predict cardiovascular mortality in end-stage renal disease patients (98). The pathophysiological significance of these correlations between decreased sRAGE/esRAGE and cardiovascular events in CKD patients will have to await more mechanistic studies.

7. sRAGE AND NEURODEGENERATIVE DISORDERS

Emanuele *et al.* reported that levels of sRAGE were significantly reduced in the plasma of patients with Alzheimer disease compared with that for those with either vascular dementia ($p<0.05$) or with controls ($p<0.001$) (99). Further, circulating sRAGE levels were decreased in patients with mild cognitive impairment (MCI), and could be a marker of extreme longevity in humans (100,101).

Further, sRAGE levels were significantly decreased in serum of the patients with amyotrophic lateral sclerosis comparing to the control group, although the correlation between the serum sRAGE levels and clinical parameters of the disease was not significant (102). Sternberg *et al.* reported that serum levels of sRAGE were significantly lower in multiple sclerosis (MS) patients compared to levels in healthy controls (103). They also showed the association of sRAGE with disease activity and rate of clinical relapse. These observations suggest that sRAGE may protect against various types of neurodegenerative disorders and that decreased sRAGE level could be a biomarker of these degenerative diseases.

8. sRAGE AND INFLAMMATORY DISEASES

sRAGE levels at baseline significantly correlated with bone/cartilage turnover markers including C-terminal propeptide of type I procollagen, carboxyterminal telopeptide of type I collagen and cartilage oligomeric matrix protein in rheumatoid arthritis (RA) patients, and the decrease of sRAGE levels following hormone replacement therapy paralleled with diminished concentration of these molecules (104). Bone mineral density (BMD) in hip and femoral neck and progression of Larsen score at 1 year were associated with baseline sRAGE levels. The same authors reported that sRAGE levels were decreased in RA patients and that its synovial fluid levels were higher in patients treated with methotrexate, compared without disease-modifying anti-RA treatment (105). sRAGE levels in RA patients were associated with CRP levels as well (106). These observations suggest serum sRAGE was associated with BMD and markers of bone/cartilage metabolism in RA.

Yoshizaki *et al.* reported that serum HMGB1 and sRAGE levels in systemic sclerosis SSc were higher than those in controls (107). Since SSc patients with elevated HMGB1 and sRAGE levels had more frequent involvement of several organs and immunological abnormalities compared to those with normal levels, increased sRAGE level may be a biomarker that could reflect disease severity and immunological abnormalities in SSc. In contrast the case with SSc, sRAGE levels were decreased in patients with Kawasaki disease and primary Sjögren's syndrome (108,109).

We, along with others, have recently found that sRAGE levels are elevated in septic shock patients, and that nonsurvivors after 28 days have had higher plasma sRAGE concentrations than survivors (2302 \pm 189 versus 1326 \pm 112 pg/ml) (110,111). Receiver operating characteristic curve analysis of plasma sRAGE concentrations of septic patients showed a specificity of 75% and a sensitivity of 84.6% with 1596 pg/ml as cutoff.

Taken together, for different autoimmune diseases, sRAGE showed both a positive correlation and a negative correlation. The correlation between sRAGE levels and disease activity may be dependent on autoimmune disorder types and/or serum levels of RAGE ligands such as AGEs and HMGB1.

9. sRAGE AND OTHER DISORDERS

Plasma levels of sRAGE were significantly decreased in patients with nonalcoholic steatohepatitis, a hepatic manifestation of insulin resistance compared with controls (112).

Atherosclerotic mechanism may be involved in the pathogenesis of aortic valve stenosis (AVS) (113). sRAGE levels were significantly lower in AVS patients than in controls, and inversely correlated with coronary calcification in AVS patients (113).

sRAGE levels may be altered in patients with cancers. Despite higher serum levels of AGEs, sRAGE levels were significantly decreased in patients with breast cancer compared with healthy controls (114). Further, patients with better outcome (low grade histology and estrogen receptor-positive breast cancer) are reported to have higher sRAGE levels (114). Overall survival rate of lung cancer patients with low expression of cytoplasmic esRAGE was significantly lower than that of patients with normal expression (115).

It would be interesting to introduce some information on sRAGE in pregnancy. sRAGE levels in women with preterm labor are decreased and correlated negatively with the leukocyte count (116). Further, in women with preeclampsia, sRAGE is elevated and correlated with serum creatinine concentration and with uric acid concentration (116). In the first trimester, pregnant diabetic (DP) patients showed lower sRAGE and higher AGEs compared to healthy pregnant. In the DP group, significant negative correlations were seen between TNF- α and lipopolysaccharide-stimulated interleukin-6 in the first trimester and sRAGE in the third trimester (117).

10. CONCLUSION

We review here the kinetics, regulation and pathophysiological role of sRAGE and esRAGE in diabetes, CVD, HT, CKD, neurodegenerative disorders, inflammatory diseases and others. Further basic and clinical investigations are needed to clarify whether sRAGE and/or esRAGE levels are one of the most convenient biomarkers for evaluating disease activity and target organ damage in these devastating disorders.

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