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ARTICLE OPEN



Elevated plasma sclerostin is associated with high brain amyloid- β load in cognitively normal older adults

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Osteoporosis and Alzheimer's disease (AD) mainly affect older individuals, and the possibility of an underlying link contributing to their shared epidemiological features has rarely been investigated. In the current study, we investigated the association between levels of plasma sclerostin (SOST), a protein primarily produced by bone, and brain amyloid-beta (A β) load, a pathological hallmark of AD. The study enrolled participants meeting a set of screening inclusion and exclusion criteria and were stratified into A β - ($n = 65$) and A β + ($n = 35$) according to their brain A β load assessed using A β -PET (positron emission tomography) imaging. Plasma SOST levels, apolipoprotein E gene (*APOE*) genotype and several putative AD blood-biomarkers including A β 40, A β 42, A β 42/A β 40, neurofilament light (NFL), glial fibrillary acidic protein (GFAP), total tau (t-tau) and phosphorylated tau (p-tau181 and p-tau231) were detected and compared. It was found that plasma SOST levels were significantly higher in the A β + group (71.49 ± 25.00 pmol/L) compared with the A β - group (56.51 ± 22.14 pmol/L) ($P < 0.01$). Moreover, Spearman's correlation analysis showed that plasma SOST concentrations were positively correlated with brain A β load ($\rho = 0.321$, $P = 0.001$). Importantly, plasma SOST combined with A β 42/A β 40 ratio significantly increased the area under the curve (AUC) when compared with using A β 42/A β 40 ratio alone (AUC = 0.768 vs 0.669, $P = 0.027$). In conclusion, plasma SOST levels are elevated in cognitively unimpaired older adults at high risk of AD and SOST could complement existing plasma biomarkers to assist in the detection of preclinical AD.

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INTRODUCTION

Increasing age is a major risk factor for both osteoporosis and Alzheimer's disease (AD), and clinical outcomes reveal underlying associations between the two disorders including the observation that patients with osteoporosis are prone to develop AD in their later life^{1,2}. Emerging evidence illustrates that bone can influence the function of other organs by secreting various proteins that enter the circulation³. In particular, osteocalcin which is exclusively produced by osteoblasts has been demonstrated to regulate brain development and play a role in age-related cognition decline⁴⁻⁶, and lower osteocalcin is indicated to be an independent predictor of worse cognitive performance⁷. Nevertheless, whether bone-derived factors are associated with AD pathological alterations is unknown.

We previously reviewed several bone-derived proteins with the potential to be involved in the pathogenesis of AD, and sclerostin (SOST) which is mainly produced by osteocytes, was one of the

candidate proteins discussed⁸. Sclerostin is encoded by the *Sost* gene and is one of the major antagonists of Wnt signaling⁹. Interestingly, circulating SOST protein levels increase with increasing age¹⁰ and it has been proposed as a candidate senescence marker¹¹. Moreover, dysfunction of Wnt/ β -catenin signaling has been implicated in AD pathological mechanisms of amyloid beta (A β) plaque formation. Specifically, inhibition of the Wnt signaling can promote the amyloidogenic processing of A β precursor protein (APP) and thereby leads to an enhanced production of the A β 42 peptide, resulting in a higher A β 42/A β 40 ratio and A β oligomer levels¹². In contrast, activation of Wnt signaling decreased A β 42 levels while increasing A β 40 levels, led to a reduced A β 42/A β 40 ratio. However, a direct involvement of SOST, as a Wnt signaling inhibitor, in AD pathogenesis has yet to be established¹².

Specific proteins in blood have been employed as biomarkers to predict the preclinical stage of AD (cognitively unimpaired while proteinopathies develop), such as: A β 42, A β 40, A β 42/A β 40, glial

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fibrillary acidic protein (GFAP), total tau (t-tau), phosphorylated tau (p-tau181, p-tau231), and neurofilament light (NFL)^{13–20}. Given the fact that SOST is a Wnt signaling inhibitor and its levels in the blood are elevated with age, we hypothesize that plasma SOST is associated with early AD pathological change, i.e., cerebral A β deposition, and could be a potential predictive blood biomarker for AD in individuals “at risk”. In the current study, we employed a cohort of cognitively healthy participants divided into A β – and A β + groups according to their brain A β load based on PET imaging assessment. To examine the association between plasma SOST and the presence of brain amyloidosis, plasma SOST concentrations were compared between A β – and A β + participants before and after stratification by sex and apolipoprotein E (*APOE*) gene ϵ 4 carriage. In addition, we evaluated the correlation between plasma SOST levels and brain A β load reflected by the mean standard uptake value ratio (SUVR) in different cortical regions. Furthermore, we conducted assessment of the correlation between plasma SOST and AD biomarkers including A β 40, A β 42, A β 42/A β 40 ratio, NFL, GFAP, t-tau, p-tau181 and p-tau231, as well as parallel comparisons of the diagnostic accuracy of plasma SOST with these blood biomarkers. Lastly, we evaluated the potential of plasma SOST as a predictor in distinguishing A β + from A β – individuals.

RESULTS

Cohort characteristics

Brain imaging and blood samples were obtained from 100 participants, with demographic and clinical characteristics summarized in Table 1. No significant difference was observed in terms of age, sex, MMSE and BMI between the A β – and A β + groups. As expected, there was a higher ratio of *APOE* ϵ 4 carriage in the A β + group (45.7%) compared with A β – group (7.7%; $P < 0.001$). By design, the SUVR values in the A β + group (1.71 \pm 0.26) were significantly higher than the A β – group (1.16 \pm 0.09; $P < 0.001$). Of the biomarkers assessed in the present

study, significant differences of the values of GFAP, p-tau181, p-tau231 and A β 42/A β 40 ratio were found between the groups (Table 1).

Associations of AD-related risk factors including age, sex and *APOE* allele status with plasma SOST

Plasma SOST levels were higher in males (76.66 \pm 27.67 pmol/L), compared with females (54.73 \pm 18.74 pmol/L; $P < 0.001$; Fig. 1a). Plasma SOST levels did not differ significantly in *APOE* ϵ 4 non-carriers and *APOE* ϵ 4 carriers (non-carriers = 62.45 \pm 25.76 pmol/L; carriers = 59.12 \pm 16.98 pmol/L; $P = 0.57$; Fig. 1b). Consistent with previous studies¹⁰, plasma SOST levels positively correlated with age ($\rho = 0.283$, $P = 0.004$; Fig. 1c).

Comparison of plasma SOST between A β – participants and A β + participants

Plasma SOST levels were significantly higher in the A β + group (71.49 \pm 25 pmol/L) compared with the A β – group (56.51 \pm 22.14 pmol/L), before and after adjusting for potential AD risk factors including age, sex and *APOE* ϵ 4 status ($P < 0.003$, $P^* = 0.008$; Table 2). On stratifying study participants by *APOE* ϵ 4 carriage status (ϵ 4 non-carriers: $n = 79$, and ϵ 4 carriers: $n = 21$), significantly higher plasma SOST levels were observed in the *APOE* ϵ 4 non-carrier A β + group ($n = 19$, 80.93 \pm 27.18 pmol/L) compared with the *APOE* ϵ 4 non-carrier A β – group ($n = 60$, 55.94 \pm 22.63 pmol/L), before and after adjusting for potential risk factors including age and sex ($P < 0.001$, $P^* = 0.001$; Table 2). Within the *APOE* ϵ 4 carriers, no significant difference in plasma SOST levels was observed between the A β + group ($n = 16$, 60.42 \pm 16.98 pmol/L) and the A β – group ($n = 5$, 63.42 \pm 15.06 pmol/L, $P = 0.728$, $P^* = 0.368$; Table 2). This observation could be attributed to the small sample size of the *APOE* ϵ 4 subset available within the present study.

When stratifying study participants by sex (male: $n = 32$; female: $n = 68$), there was a trend of higher plasma SOST levels in male A β + participants ($n = 13$, 87.07 \pm 24.28 pmol/L) compared with male A β – participants ($n = 19$, 69.55 \pm 28.18 pmol/L, $P = 0.078$, $P^* = 0.05$; Table 2). It is likely that an increased sample size of male participants would uncover a statistically significant effect. In females, a significantly higher plasma level of SOST was observed in A β + participants ($n = 22$, 62.28 \pm 20.87 pmol/L) compared with A β – participants ($n = 46$, 51.12 \pm 16.68 pmol/L), before and after adjusting for potential risk factors including age and *APOE* ϵ 4 status ($P = 0.02$, $P^* = 0.033$; Table 2).

Correlation of plasma SOST with selected AD biomarkers

We next assessed whether plasma SOST levels correlate with any AD-related biomarkers using Spearman’s correlation. We found a significant association between SOST levels and NFL, GFAP, phosphorylated versions of the tau protein (p-tau181 and p-tau231) and A β 40, while no significant association was found between SOST levels and other AD-related biomarkers (Table 3, unadjusted correlations are shown in Supplementary Fig. 1).

The cohort was then further stratified by brain A β deposition status. It was found that SOST levels were significantly correlated with NFL, p-tau181 and A β 40 while no correlation was observed with other biomarkers in A β – group. For individuals with A β + status, no significant association was found between SOST levels and all the selected biomarkers (Table 3).

Evaluation of plasma SOST as a predictor for brain A β status

Given the higher plasma SOST levels in the A β + group compared with the A β – group, we investigated the predictive value of plasma SOST for brain A β status. We first determined the correlation between plasma SOST levels and brain SUVR using Spearman’s correlation coefficient (ρ). Intriguingly, plasma SOST

Table 1. Cohort demographic characteristics.

	A β –	A β +	<i>P</i>
Age (mean \pm SD)	77.62 \pm 5.56	79.23 \pm 5.38	0.165
Sex (M/F)	19/46	13/22	0.419
<i>APOE</i> ϵ 4 carriers (n (%))	5 (7.7)	16 (45.7)	<0.001
SUVR (mean \pm SD)	1.16 \pm 0.09	1.71 \pm 0.26	<0.001
MMSE (mean \pm SD)	28.51 \pm 1.16	28.80 \pm 1.11	0.219
BMI (mean \pm SD)	27.39 \pm 4.48	28.05 \pm 4.74	0.494
GFAP (pg/mL, mean \pm SD)	145.5 \pm 49.2	210.5 \pm 83.9	<0.001
NFL (pg/mL, mean \pm SD)	19.36 \pm 9.00	21.02 \pm 10.09	0.416
t-tau (pg/mL, mean \pm SD)	1.19 \pm 0.37	1.34 \pm 0.39	0.068
p-tau181 (pg/mL, mean \pm SD)	13.55 \pm 5.69	17.27 \pm 5.58	0.003
p-tau231 (pg/mL, mean \pm SD)	11.39 \pm 6.45	19.08 \pm 7.16	<0.001
A β 42 (pg/mL, mean \pm SD)	21.96 \pm 4.58	19.54 \pm 5.74	0.041
A β 40 (pg/mL, mean \pm SD)	95.39 \pm 14.75	98.37 \pm 16.88	0.395
A β 42/40 (mean \pm SD)	0.23 \pm 0.04	0.20 \pm 0.05	0.003

Baseline characteristics including sex, age, body mass index (BMI), *APOE* ϵ 4 status, Mini-mental State Examination (MMSE) scores, brain amyloid-beta (A β) load represented by the standard uptake value ratio (SUVR) of ligand 18F-Florbetaben (FBB) in the neocortical region normalized with that in cerebellum, and blood-based biomarkers (GFAP, NFL, t-tau, p-tau181, p-tau231, A β 42, A β 40 and A β 42/40) have been compared between A β – (SUVR < 1.35, $n = 65$) and A β + (SUVR \geq 1.35, $n = 35$) participants. Student’s *t* test or Chi-square tests were employed as appropriate. Data are presented in mean \pm SD, and *P* values in bold font were considered as significant ($P < 0.05$). ($N = 100$, A β – $n = 65$, A β + $n = 35$).
GFAP glial fibrillary filament protein, NFL neurofilament light.

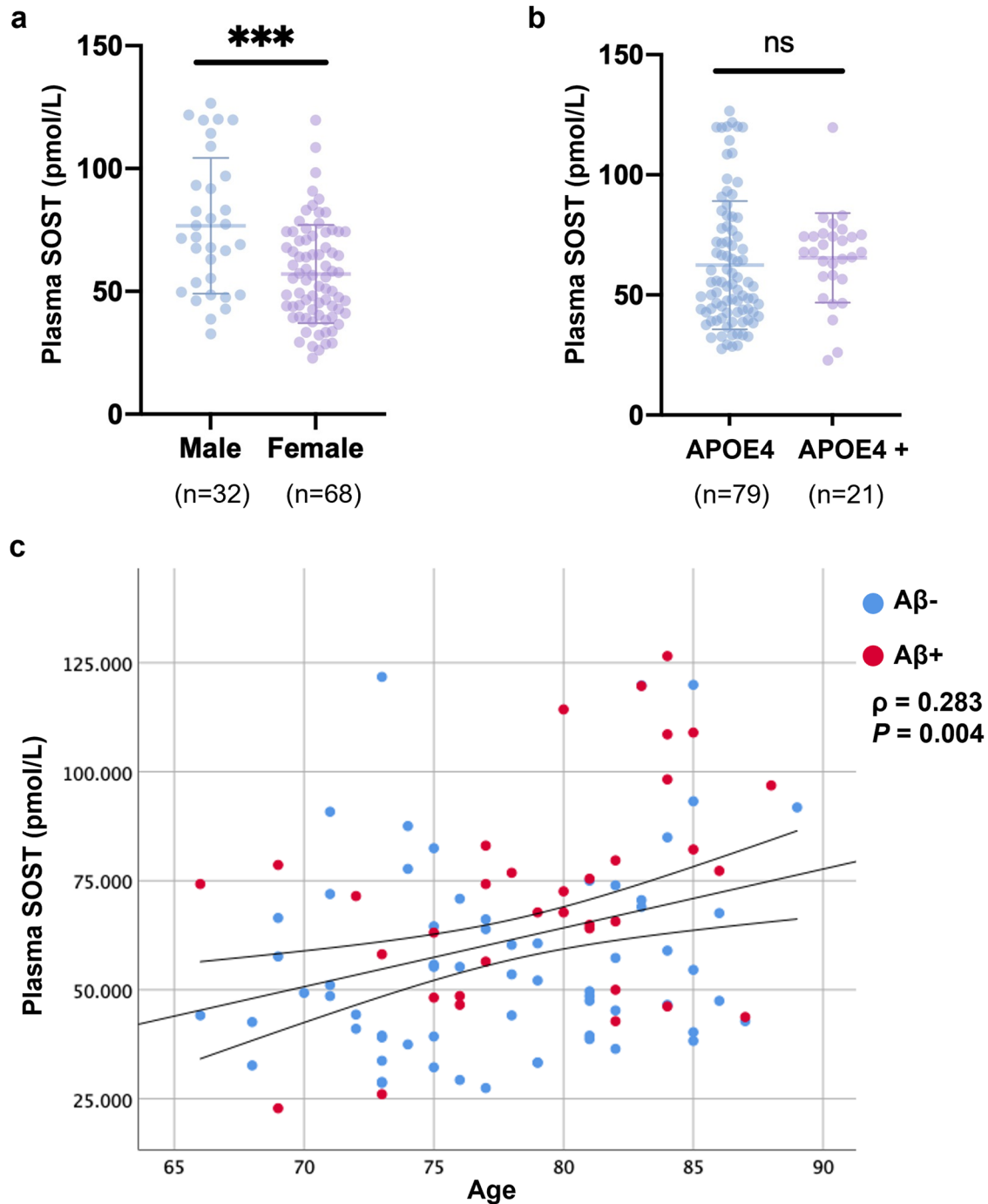


Fig. 1 Associations of AD-related risk factors, age, sex and *APOE4* allele status with plasma sclerostin. **a** Plasma sclerostin levels were compared between male and female participants. **b** Plasma sclerostin levels were compared between *APOE4*⁻ and *APOE4*⁺ participants. **c** Correlations between plasma sclerostin levels and age. The line segment within each jitter plot represents the median of the data and error bars in the graphs represent the interquartile range for the groups compared. *** $P \leq 0.001$, ns no significant difference. Correlation coefficients and P values were calculated using Spearman's correlation coefficient (ρ).

levels were found to have a significant positive correlation with SUVR ($\rho = 0.321$, $P = 0.001$; Supplementary Fig. 2). AUC analysis was then performed to determine if SOST levels improved the diagnosis of individuals with ongoing brain deposition, either alone or in combination with other known AD-related biomarkers (GFAP, t-tau, p-tau181, p-tau231, A β 40, A β 42 and A β 42/A β 40 ratio). A base model (BM) including age, sex and *APOE* ϵ 4 status was included in the analysis. The AUC of the base model was 0.787

(95% CI = 0.693–0.882), however, when combining the base model with the plasma SOST model, higher accuracy (AUC = 0.818, 95% CI = 0.733–0.903) was obtained compared with the base model alone, albeit not to a significant level ($P = 0.228$; Fig. 2a). The addition of SOST to the models including the BM plus known AD-related biomarkers (p-tau181, p-tau231, GFAP and A β 42/A β 40 ratio) also reached a higher discriminative accuracy between individuals with A β ⁺ status from those with A β ⁻ status,

Table 2. Comparison of plasma sclerostin within all A β - and A β + participants and subgroups stratified by APOE ϵ 4 status and sex, respectively.

	A β -	A β +	<i>P</i>	<i>P</i> *
SOST in all participants	56.51 \pm 22.14	71.49 \pm 25.00	0.003	0.008
SOST in APOE ϵ 4 non-carriers	55.94 \pm 22.63	80.93 \pm 27.18	<0.001	0.001
SOST in APOE ϵ 4 carriers	63.42 \pm 15.06	60.42 \pm 16.98	0.728	0.368
SOST in males	69.55 \pm 28.18	87.07 \pm 24.28	0.078	0.050
SOST in females	51.12 \pm 16.68	62.28 \pm 20.87	0.020	0.033

Student's *t* test or Chi-square tests were employed as appropriate. Data are presented in mean \pm SD in pmol/L, and *P* values in bold font were considered as significant (*P* < 0.05). *P** represents *P* values adjusted for age, sex and APOE ϵ 4 status.

Table 3. Correlations between plasma sclerostin levels and AD biomarkers among all participants and subgroups stratified by brain A β status.

	All participants		A β -		A β +	
	ρ	<i>P</i>	ρ	<i>P</i>	ρ	<i>P</i>
NFL	0.337	<0.001	0.445	<0.001	0.095	0.587
GFAP	0.314	0.001	0.218	0.081	0.067	0.703
t-tau	-0.017	0.865	-0.018	0.889	-0.267	0.121
p-tau181	0.393	<0.001	0.279	0.025	0.143	0.434
p-tau231	0.316	0.002	0.241	0.053	0.114	0.541
A β 42	0.097	0.348	0.217	0.090	0.061	0.737
A β 40	0.283	0.005	0.336	0.007	0.153	0.395
A β 42/40	-0.155	0.133	-0.076	0.559	-0.128	0.479

Unadjusted correlation coefficients (ρ) and *P* values between plasma sclerostin and AD-related biomarkers were calculated using Spearman's correlation analysis.

P values in bold font were considered as significant (*P* < 0.05).

while not to a significant level, neither (AUC = 0.806 vs 0.836, *P* = 0.27 for BM + p-tau181 vs BM + p-tau181 + SOST; AUC = 0.838 vs 0.852, *P* = 0.47 for BM + p-tau231 vs BM + p-tau231 + SOST; AUC = 0.916 vs 0.918, *P* = 0.69 for BM + GFAP vs BM + GFAP + SOST; AUC = 0.833 vs 0.857, *P* = 0.27 for BM + A β 42/A β 40 vs BM + A β 42/A β 40 + SOST; Fig. 2c–e). Furthermore, the addition of SOST to the models including only AD-related biomarkers (not including age, sex, APOE carriage) was also assessed and did not significantly increase the diagnostic potential for brain amyloid deposition in p-tau181 vs p-tau181 + SOST (AUC = 0.712 vs 0.759, *P* = 0.23; Fig. 2f), p-tau231 vs p-tau231 + SOST (AUC = 0.792 vs 0.816, *P* = 0.36, Fig. 2g) and GFAP vs GFAP + SOST (AUC = 0.762 vs 0.789, *P* = 0.25; Fig. 2h). Interestingly, there was a significant increase of accuracy in diagnosing brain amyloidosis upon the addition of SOST to A β 42/A β 40 (AUC = 0.669 vs 0.768, *P* = 0.027, A β 42/A β 40 vs A β 42/A β 40 + SOST; Fig. 2i).

DISCUSSION

The present study has demonstrated that the plasma level of the bone-derived protein, SOST, is positively correlated with brain A β load in a cognitively unimpaired older population. To our knowledge, this is the first study demonstrating the association

between plasma SOST levels and preclinical AD pathological alterations related to A β accumulation in the brain. Importantly, the positive correlation between plasma SOST and age also highlights a possible pathophysiological relationship between osteoporosis and AD with ageing.

It is demonstrated that SOST participates in bone turnover, and for this reason has been identified as a therapeutic target for the treatment of osteoporosis²¹. In addition to its role in bone, SOST is likely to have functions in other organs beyond the skeletal system. For example, blood SOST levels correlate inversely with glomerular filtration rate after correction for age and sex in patients with chronic kidney disease²². In addition, higher SOST blood levels have been observed in type 2 diabetes mellitus patients compared to healthy individuals^{23,24}, and there is a positive correlation between blood SOST levels and insulin resistance²⁵. These findings implicate a cross-talk between bone and other organs mediated by SOST.

To further investigate the correlation of SOST plasma levels and brain A β load, this study also investigated the association between blood SOST and several major AD risk factors by stratifying the participants based on sex and APOE ϵ 4 status. It was found that in both male and female subgroups, participants with an A β + status had higher blood SOST levels compared with participants with an A β - status. This indicates that the association between blood SOST and brain A β load is independent of sex. Interestingly, we also observed high blood levels of SOST in A β + individuals who did not carry the APOE ϵ 4 allele, whereas there was no difference in APOE ϵ 4 carriers. This observation could be attributable to the modest sample size of the APOE ϵ 4 carriers, or more likely, due to the greater influence of the ApoE 4 protein compared with SOST in the development of AD. When evaluating the contribution of plasma SOST in diagnosing brain amyloidosis, it was found that in the presence of a base model including age, sex and APOE ϵ 4, the addition of plasma SOST improved, albeit not to a significant level, the diagnostic accuracy of brain amyloidosis. Based on the results showing that plasma SOST levels were significantly different in A β - compared to A β + in APOE ϵ 4 non-carriers, but were similar in APOE ϵ 4 carriers, we assume that the influence of the APOE ϵ 4 allele overrides the influence of plasma SOST in terms of A β deposition.

We also evaluated the potential of combining plasma SOST with a number of AD-related biomarkers in diagnosing A β + status without using the base model including age, sex and APOE ϵ 4 status. Interestingly, the addition of SOST increased the validity of these biomarkers, and in particular, resulting in a significant improvement in diagnostic value for amyloid deposition when combined with the A β 42/A β 40 ratio alone. Further, Spearman's correlation analysis in the whole cohort showed that plasma SOST levels were positively correlated with A β 40 levels in the plasma. In addition, it was found that with increasing plasma SOST levels, there was an increasing trend of A β 42 levels accompanied by a decreasing trend of the A β 42/A β 40 ratio in the plasma, despite such correlations not reaching statistical significance. These findings re-enforce the underlying association between plasma levels of SOST and A β 40, A β 42 and A β 42/A β 40 and amyloid deposition.

While the strengths of this study include the utilization of data from the highly characterized KARVIAH aging cohort which has been reported in previous studies^{26,27}, it needs to be acknowledged that the current study has some limitations, such as the modest sample size and cross-sectional nature of the study design. Another limitation is that SOST levels in the blood may be influenced by other conditions such as chronic kidney disease and type 2 diabetes and ageing pathophysiology¹¹, therefore additional inclusion/exclusion criteria taking into account other comorbidities of the participants will need to be considered in future studies. Further studies are also required to validate these observations through longitudinal monitoring of SOST levels in

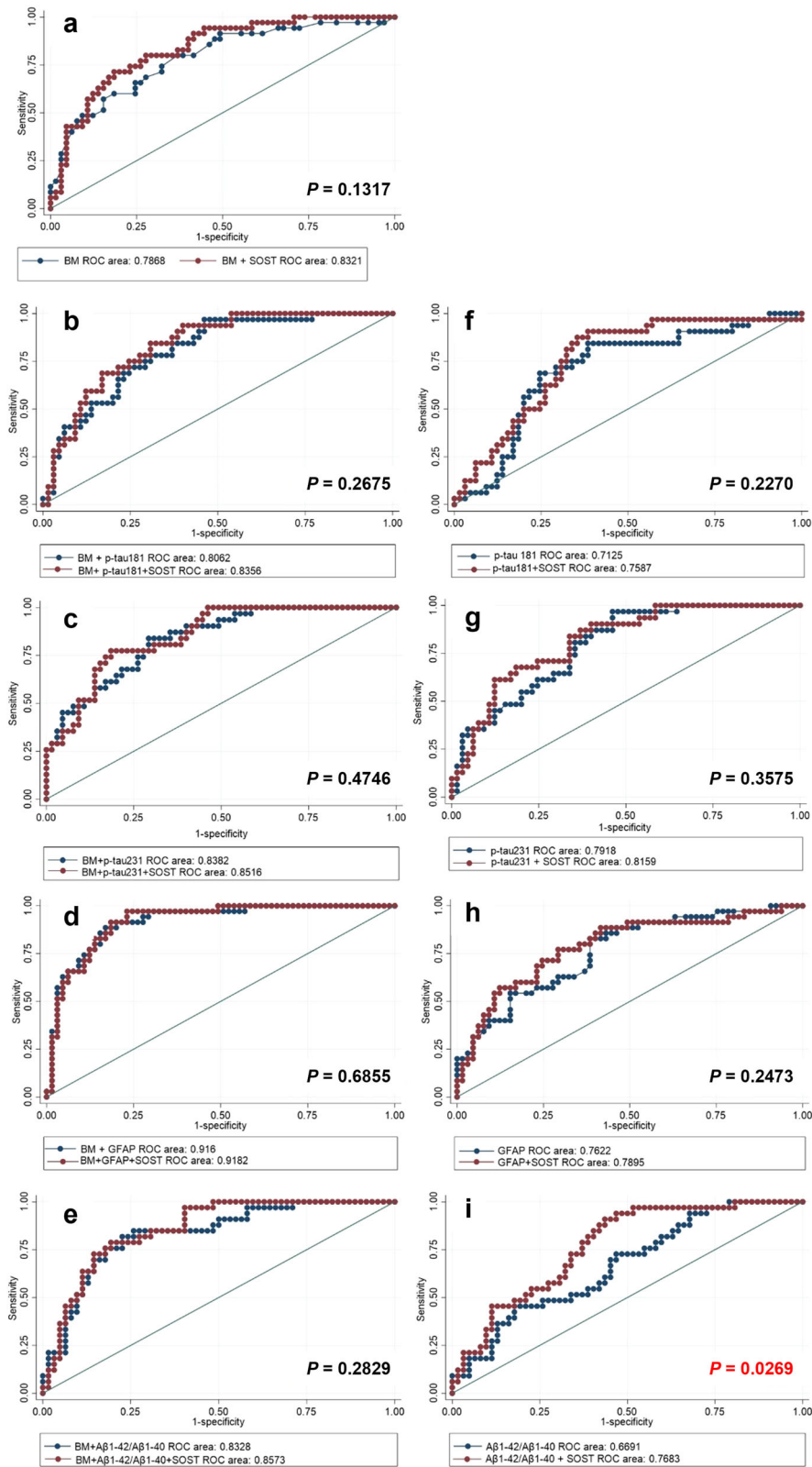


Fig. 2 Receiver operating characteristic curves for distinguishing $A\beta^+$ individuals from $A\beta^-$ individuals using plasma sclerostin in combination with the base model (BM) and/or AD-related biomarkers. A base model (BM) including age, sex and *APOE* $\epsilon 4$ carriage status was included in the analysis. Receiver operating characteristic (ROC) curves are presented with the BM (a–e) or without (f–i). **a** (BM vs BM + SOST), **b** (BM + p-tau181 vs BM + p-tau181 + SOST), **c** (BM + p-tau231 vs BM + p-tau231 + SOST), **d** (BM + GFAP vs BM + GFAP + SOST), **e** (BM + A β 1-42/A β 1-40 vs BM + A β 1-42/A β 1-40 + SOST), **f** (p-tau181 vs p-tau181 + SOST), **g** (p-tau231 vs p-tau231 + SOST), **h** (GFAP vs GFAP + SOST) and **i** (A β 1-42/A β 1-40 vs A β 1-42/A β 1-40 + SOST).

an independent patient cohort. In addition, the comparison of blood SOST levels between healthy individuals and those with mild cognitive impairment or established AD would further strengthen the usefulness of SOST as an AD biomarker, and its role in AD progression. Moreover, if bone-derived SOST can have a biological effect in the brain through the Wnt/ β -catenin signaling pathway, an outstanding question which has yet to be addressed relates to the extent to which it is able to cross the blood-brain barrier (BBB).

This study has demonstrated that plasma SOST levels are increased in cognitively preserved older adults at high risks of AD. A higher degree of accuracy in differentiating $A\beta^-$ and $A\beta^+$ individuals was achieved when combining SOST blood measurements with other major AD risk factors and several putative plasma AD biomarkers, indicating that plasma SOST is a promising biomarker to assist in the detection of preclinical AD. Importantly, the association between bone-derived SOST and brain $A\beta$ load status has drawn further attention to the importance of the 'bone-brain axis' and has indicated a possible common link in the pathogenesis of osteoporosis and AD.

METHODS

Cohort

The study enrolled participants who have met a set of screening inclusion and exclusion criteria from the Kerr Anglican Retirement Village Initiative in Aging Health (KARVIAH) cohort^{26,27}. The inclusion criteria comprised an age range of 65–90 years, good general health, no known significant cerebrovascular disease, fluent in English, adequate/corrected vision and hearing to enable testing, and no dementia or pathological cognitive impairment as screened by a Montreal Cognitive Assessment (MoCA) score ≥ 26 . MoCA scores lying between 18–25 were assessed on a case-by-case basis by the study neuropsychologist following stratification of scores according to age and education. The exclusion criteria comprised the previous diagnosis of dementia based on the revised criteria from the National Institute on Aging-Alzheimer's Association, the presence of an acute functional psychiatric disorder (including lifetime history of schizophrenia or bipolar disorder), severe or extremely severe depression (based on the Depression, Anxiety, Stress Scales; DASS), a history of stroke, and uncontrolled hypertension (systolic blood pressure [BP] > 170 mm Hg or diastolic BP > 100 mm Hg).

In total, 134 volunteers met the inclusion and exclusion criteria; of these, 105 underwent neuroimaging, neuropsychometric evaluation, and blood collection; the remainder declined to undergo neuroimaging or withdrew from the study. Within these 105 participants, 100 participants were considered to have normal global cognition based on their Mini-Mental State Examination (MMSE; scores can range from 0 to 30, with higher scores indicating better cognitive function) wherein, a cut-off score < 26 was employed to screen out individuals with potential early dementia. All volunteers provided written informed consent prior to participation, and the Bellberry and the Macquarie University Human Research Ethics Committees provided approval for the study.

Brain amyloid- β load evaluation via PET

Neuroimaging was conducted within 3 months of blood collection at Macquarie Medical Imaging in Sydney. Positron emission tomography (PET) studies were conducted over a 20-min static scan (4 \times 5 min frames) that was acquired 50 min after an intravenous bolus of 18F-florbetaben (FBB) administered slowly over 30 s. Brain $A\beta$ load was calculated as the mean standard uptake value ratio (SUVR) of the frontal, superior parietal, lateral temporal, lateral occipital, and anterior and posterior cingulate regions using image processing software, CapAIBL. Participants

with an SUVR ≥ 1.35 were considered to have a high brain $A\beta$ load ($A\beta^+$), while those with an SUVR < 1.35 were considered to have a low $A\beta$ load ($A\beta^-$)²⁶.

Blood collection, measurement of plasma SOST and APOE genotyping

All study participants fasted for a minimum of 10 h overnight prior to venesection employing standard serological methods and processing. Following blood sample processing, plasma fractions were stored at -80°C until further testing. Plasma SOST concentration was measured using the human SOST ELISA kit (BI-20492, Biomedica, Wien, Austria) according to the manufacturer's instructions. In brief, Assay buffer was added into pre-coated plates followed with addition of plasma samples in duplicate. Samples were then incubated in wells simultaneously with biotinylated antibody overnight (18–24 h) at room temperature (18–24 $^\circ\text{C}$) and washed 5 \times prior to incubation with conjugation buffer for 1 h at room temperature in the dark. The conjugation buffer was aspirated and wells washed 5 \times prior to a 30-min incubation with substrate buffer in the dark. At the end of the incubation, stop solution was added into each well and the absorbance measured immediately at 450 nm. The average signal coefficient of variation (%) of all data is 5.57%. A standard curve was constructed from the absorbance values of the standards. Plasma SOST concentrations of the samples were obtained from the standard curve. Apolipoprotein E (APOE) genotype was determined from purified genomic DNA extracted from 0.5 mL whole blood. Each sample was genotyped for the presence of the three APOE variants ($\epsilon 2$, $\epsilon 3$ and $\epsilon 4$) based on TaqMan SNP genotyping assays for rs7412 (C 904973) and rs429358 (C3084793) as per the manufacturer's instructions (AB Applied Biosystems by Life Technologies, Scoresby, VIC, Australia). Five percent of the samples were genotyped in duplicate and 100% inter- and intra-assay concordance was observed.

Blood biomarkers analysis

The single molecule array (Simoa) platform was used to measure protein biomarker concentrations in EDTA plasma specimens. GFAP, t-tau and NFL concentrations were measured using the Neurology 4-Plex A kit (QTX-102153, Quanterix, MA, USA), while p-tau181 and p-tau231 concentrations were measured with in-house assays developed at the University of Gothenburg, Sweden^{15,17}. Calibrators and samples were run in duplicates in all assays. Two internal controls provided with the Simoa kits were included in duplicate at the beginning of each experiment. Plasma $A\beta$ concentrations were measured using the Amyblood test that was developed at Amsterdam University Medical Center in collaboration with ADx NeuroSciences (Ghent, Belgium), on the Simoa platform (HDx instrument, Quanterix)^{27,28}.

Statistical analyses

Student's *t* tests or Chi-square tests were employed to compare descriptive statistics including means and standard deviations calculated for $A\beta^-$ and $A\beta^+$ groups. Continuous variables between $A\beta^-$ and $A\beta^+$ groups were compared by employing linear models after correcting for covariates age, sex, and APOE $\epsilon 4$ carrier status. To better approximate normality and variance homogeneity, dependent variables were transformed to natural log as required. Correlations between continuous variables were investigated using the Spearman correlation coefficient (ρ). Predictive models and receiver-operating characteristic (ROC) curves constructed from the logistic scores were evaluated by using logistic regression with $A\beta^-/+$ as response. All data analyses and visualization were carried out using IBM® SPSS (v27) or Stata MP (v17.0) and GraphPad Prism (v8), respectively.

Reporting summary

Further information on research design is available in the Nature Research Reporting Summary linked to this article.

DATA AVAILABILITY

All data generated or analyzed during this study are included in this paper or the supplementary materials. Additional data are available from the authors upon reasonable request.

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AUTHOR CONTRIBUTIONS

M.Z. and R.N.M. designed the project and the main conceptual ideas. J.Y., R.T. and J.D. conducted the statistical analyses. J.Y. and S.P. wrote the manuscript. S.P., C.E.T., I.M.W.V., E.S., H.V., N.A., H.Z. and K.B. measured plasma biomarkers. P.C., H.R.S., B.P.M., C.M., S.R.S. K.G., A.C.P.T., F.M. and C.I. contributed to editing this manuscript and interpreting the results. J.G. and D.L. contribute to figure preparation process. All authors critically reviewed the manuscript. J.Y. and S.P. are co-first authors. M.Z. and R.N.M. both supervised the project.

COMPETING INTERESTS

K.B. has served as a consultant, at advisory boards, or at data monitoring committees for Abcam, Axon, BioArctic, Biogen, JOMDD/Shimadzu, Julius Clinical, Lilly, MagQu, Novartis, Ono Pharma, Pharmatrophix, Prothena, Roche Diagnostics, and Siemens Healthineers, and is a co-founder of Brain Biomarker Solutions in Gothenburg AB (BBS), which is a part of the GU Ventures Incubator Program, outside the work presented in this paper. H.Z. has served at scientific advisory boards and/or as a consultant for Abbvie, Acumen, Alector, ALZPath, Annexon, Apellis, Artery Therapeutics, AZTherapies, CogRx, Denali, Eisai, Nervgen, Novo Nordisk, Passage Bio, Pinteon Therapeutics, Red Abbey Labs, reMYND, Roche, Samumed, Siemens Healthineers, Triplet Therapeutics, and Wave, has given lectures in symposia sponsored by Cellectricon, Fujirebio, Alzecure, Biogen, and Roche, and is a co-founder of Brain Biomarker Solutions in Gothenburg AB (BBS), which is a part of the GU Ventures Incubator Program (outside submitted work). C.T. has a collaboration

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ADDITIONAL INFORMATION

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