



Blood Selenium and Serum Glutathione Peroxidase Levels Were Associated with Serum β -Amyloid in Older Adults

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Abstract

Background Studies have established the association between blood β -amyloid ($A\beta$) levels and Alzheimer's disease, but population-based studies concerning the association between selenium (Se) and $A\beta$ levels in blood samples are very limited. Therefore, we explored the association in an elderly population with Se status and serum $A\beta$ measures.

Methods A cross-sectional study on 469 elderly individuals from four rural counties with diverse soil Se levels was carried out. Fasting blood Se, serum selenoprotein P (SELENOP), and glutathione peroxidase (GPX), serum $A\beta$ 42, and $A\beta$ 40 were measured. Quantile regression models were used to determine the associations of blood Se, serum GPX, and SELENOP with $A\beta$ levels.

Results Significant negative associations were observed between blood Se and serum $A\beta$ 42 and $A\beta$ 40 levels at all percentiles ($P < 0.05$). The associations were generally stronger at higher $A\beta$ 42 and $A\beta$ 40 percentiles than lower $A\beta$ 42 and $A\beta$ 40 percentiles. Blood Se was positively associated with serum $A\beta$ 42/ $A\beta$ 40 ratio at 25th, 50th, and 75th percentiles. Significant positive associations were observed between serum GPX and $A\beta$ 42 and $A\beta$ 40 levels at all percentiles ($P < 0.05$). The positive associations were generally stronger at higher $A\beta$ 42 and $A\beta$ 40 percentiles than at lower percentiles. Serum GPX was negatively associated with $A\beta$ 42/ $A\beta$ 40 ratio at 25th, 50th, 75th, and 95th percentiles. No associations with serum SELENOP and $A\beta$ levels were observed.

Conclusions Our results suggest that higher Se levels are associated with lower serum $A\beta$ 42 and $A\beta$ 40 levels and with higher $A\beta$ 42/ $A\beta$ 40 ratio, and the results are specific for different selenoproteins.

Keywords Selenium · Selenoprotein P · Glutathione peroxidase · β -Amyloid

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Introduction

Alzheimer's disease (AD) is a global public health threat in the context of population aging [1]. Identifying risk or protective factors associated with AD is highly critical. The deposition of neurotoxic β -amyloid ($A\beta$) in the brain is one of the key neuropathologic feature of AD development [2]. $A\beta$ 42 and $A\beta$ 40 are two predominant isoforms of $A\beta$ [3], which have been examined in an increasing number of research studies for AD pathogenesis and prognosis. Abnormal $A\beta$ status in cerebrospinal fluid (CSF) analysis has been incorporated in the diagnostic criteria for AD [4]. However, CSF acquisition may not be feasible in large observational studies of older adults. Previous studies have demonstrated that peripheral blood $A\beta$ levels were positively correlated with CSF $A\beta$ levels and could be regarded as predictive biomarkers for brain amyloidosis [5]. Moreover, higher $A\beta$ 42

and A β 40 levels in peripheral blood were associated with higher risk of AD [6–8], and higher A β 42/A β 40 ratio was typically associated with lower AD risk [9].

Selenium (Se) is an essential trace element for cerebral functions and the brain is the last Se-deficient organ in Se deficiency [10]. Changes in Se concentration and selenoprotein activity in blood and brain have been reported in AD and other brain diseases [11, 12]. Se had been found to inhibit the activity of γ -secretase, attenuate the alterations in APP expression and A β production, and have a protective effect on the neurotoxicity of A β [13, 14]. Selenoproteins are the major forms for Se to perform numerous biological functions [15]. Glutathione peroxidase (GPX) plays a unique role in protecting cells from free radical-induced oxidative damage [16], and GPX gene knockout mice has been reported to exhibit higher sensitivity to A β -induced oxidative stress [17]. Selenoprotein P (SELENOP) is an important Se transporter [18], which was found to be co-localized with A β plaques in the postmortem tissue from individuals with hallmark AD lesions [19]. Reduction of SELENOP gene expression rendered the N2A cell more sensitive to the toxicity of A β [20]. However, the function and mechanism of Se in AD development remain elusive, and there is a paucity of research in the association between Se and A β levels in bio-samples from the elderly population.

Considering the difference in Se nutrition status due to geologically uneven soil Se distribution [21] and the increasing concern on Se and AD [12], this study is focused on exploring the association between Se and A β levels in older adults with an extensive range of Se exposure.

Methods

Study Design and Participants

A cross-sectional study of 469 elderly people aged 60 and older from four rural counties in Enshi prefecture with diverse soil Se levels was conducted. Participants were enrolled between 2016 and 2017; biomarker measurements were conducted between 2018 and 2019. Data from 433 participants with serum Se status and A β measures was used for statistically analysis in this study. The four sites were from extremely high Se area (> 3.00 mg/kg), high Se area (0.40–3.00 mg/kg), Se-sufficient area (0.18–0.40 mg/kg), and Se-deficient area (< 0.18 mg/kg), respectively [21]. Residents were asked to enroll in the study if they met the following eligibility criteria: (a) had lived in the area for at least 30 years; (b) consumed local food and with no dietary supplement; (c) had no language communication problem; (d) agreed to complete a face-to-face interview and provide blood samples. All participants signed written informed consents before each interview. The study was approved by

the Institute for Environmental Health and Related Safety, Chinese Center for Disease Control and Prevention.

Blood Se Measures

Fasting blood samples were collected in the morning using 5-mL purple top (EDTA) Vacutainer® tubes. All samples were stored in polypropylene tubes at – 80 °C before laboratory analysis. Blood samples were digested with concentrated nitric acid for 1 h and then placed in a boiling water bath for 4 h until the samples were clear.

Se concentrations in whole blood were measured by inductively coupled plasma mass spectrometry (ICP-MS) (Agilent 7700x, USA). The instrument parameters were as follows: RF power, 1550 W; collision mode, helium flow 4 mL/min; carrier gas flow, 1.0 L/min; plasma gas flow, 15 L/min; and auxiliary gas flow, 0.15 L/min. External calibration curves were applied for quantification. And, two quality control samples (Seronorm™ Trace Elements Whole Blood L-1 and L-2, Norway) were used to ensure the accuracy of the Se measurement. The detection limit for Se was 0.54 μ g/L. The relative standard deviation (RSD) of replicate analysis of samples was less than 10%. In order to monitor the detection procedure, blank samples were analyzed for every 20 samples in all batches; 10% duplicated samples were also included; the samples were reanalyzed if the two results differed by > 10%.

Serum Selenoproteins Measures

Selenoproteins were measured by the enzyme-linked immunoassay (ELISA) method. Commercially available kits were selected according to pretest results. Serum GPX kits (Nanjing Jiancheng Bioengineering Institute, Nanjing, China) and SELENOP kits (Shanghai Enzyme-linked Biotechnology Co., Ltd., Shanghai, China) were used to determine the concentration according to the manufacturer's instructions. The absorbance of each well was read on ELISA reader using 412 nm as primary wavelength for serum GPX and 450 nm as primary wavelength for serum SELENOP. As per manufacturer, for GPX, the intra- and inter-assay CV were 3.56% and 6.80%. For SELENOP, the reported intra- and inter-assay CVs were all less than 10%. In addition to quality control samples, 10% parallel samples were measured in each batch for quality assurance.

Serum A β Peptides Measures

Serum A β levels were determined using ELISA kits (Wako Pure Chemical Industries, Ltd., Osaka, Japan) according to manufacturer's instructions. The absorbance was measured at a wavelength of 450 nm with a microplate reader (Bio-Tek, USA). In addition to quality control samples,

10% parallel samples were measured in each batch for quality assurance. Sensitivities were 0.12 pmol/L (assay range 1.0–100 pmol/L) in the A β 40 assay and 0.08 pmol/L (assay range 1.0–100 pmol/L) in the A β 42 assay. Intra- and inter-assay coefficients of variation were less than 10% in both assay systems.

Covariates

Information on socio-demographic characteristics (age, gender, and education) and lifestyle factors (alcohol consumption and smoking) were collected by standardized questionnaire from face-to-face interviews. Height and weight were also measured during the interview. The height-measuring instrument and electronic weighing scale were calibrated before measurement. The precision was 0.1 cm for height and 0.1 kg for weight. Body mass index (BMI, defined as body weight in kilograms divided by height in meters squared) was derived from height and weight measurements. The above variables will be used as covariates in the statistical analysis according to previous study [22].

Statistical Analysis

The normality of the data was assessed by the Shapiro–Wilk test. Except for age and BMI, the distribution of other parameters was skewed in our study. The continuous variables of the normal distribution were presented as mean \pm standard deviation (SD), and continuous variables of skewed distribution were presented as median (interquartile range, IQR). Categorical variables were expressed as proportion (percentage). Comparisons of demographic characteristics among Se quintiles were performed by analysis of variance or Kruskal–Wallis rank-sum test for continuous variables and chi-square tests for categorical variables. Spearman rank correlation analysis was used to estimate associations among Se, SELENOP, and GPX.

Quantile regression models were used to examine the association between Se and A β levels. Quantile regression is a distribution-free method overcoming the limitations of traditional linear regression when the outcome variables fail to satisfy the normal distribution assumption [23, 24]. Conditional serum A β percentiles as functions of Se levels were estimated at the 5th, 25th, 50th, 75th, and 95th percentiles. The models were adjusted for age, gender, education, smoking, alcohol consumption, and BMI. Quantile regression coefficient graphs were used to present the changing trend of the association between Se levels and serum A β levels.

Statistical analyses were performed using STATA 16.0. All tests were two-tailed, and a *P*-value of less than 0.05 was considered statistically significant.

Results

Description of Participants

For the 433 participants included in this analysis, mean age was 72.38 ± 5.25 years, and 50.23% were females. Median and interquartile range (IQR) of blood Se, serum SELENOP, and GPX concentrations were 0.11 (0.08, 0.18) mg/L, 4.59 (3.62, 6.75) μ g/mL, and 0.20 (0.16, 0.25) mmol/L, respectively. Median and IQR serum of A β 42 and A β 40 concentrations were 2.48 (1.18, 4.08) pmol/L and 25.15 (8.56, 49.30) pmol/L, respectively.

Demographic characteristics of study participants by blood Se quintiles were presented in Table 1. There were significant differences in age, SELENOP, GPX, A β 42, A β 40, and A β 42/A β 40 ratio, while no significant difference was observed in BMI, gender, education, smoking, and alcohol consumption among five Se quintile groups. The blood Se level was negatively correlated with age ($r_s = -0.098$, $P = 0.044$). Non-smokers had higher SELENOP and GPX levels than that in smokers ($P < 0.001$). Participants who do not consume alcohol had higher serum SELENOP and GPX levels and lower Se levels than drinkers ($P < 0.05$). Blood Se was positively correlated with serum SELENOP ($r_s = 0.26$, $P < 0.001$) and GPX ($r_s = 0.11$, $P = 0.03$).

Blood Se and Serum A β

Regression coefficients using quantile regression were presented for the 5th, 25th, 50th, 75th, and 95th percentiles of A β distribution (Table 2). Significant negative associations were observed between blood Se and serum A β 42 and A β 40 levels at all percentiles ($P < 0.05$). There was an overall increasing trend of the absolute value of β -coefficients between blood Se and A β 42 and A β 40 percentiles, indicating that these associations were generally stronger at the higher A β 42 and A β 40 percentiles than lower A β 42 and A β 40 percentiles (Fig. 1A and B). For example, for every 1 mg/L increase in blood Se level, A β 42 decreased by 1.58 pg/mL at the lowest A β 42 percentile, and A β 42 decreased by 3.71 pg/mL at the highest A β 42 percentile. Importantly, Se was positively correlated with A β 42/A β 40 ratio at the 25th (β : 0.03, 95%CI: 0.01, 0.05), 50th (β : 0.06, 95%CI: 0.02, 0.10), and 75th (β : 0.07, 95%CI: 0.01, 0.14) percentiles. Interestingly, only at the highest and lowest percentiles for A β 42/A β 40 ratio, the association between blood Se and A β 42/A β 40 ratio was not significant.

Serum GPX and A β

Significant positive associations were observed between serum GPX, serum A β 42, and A β 40 levels at

Table 1 Demography characteristics of study participants by blood selenium quintile groups

Characteristic [#]	Total (n=433)	Quintile groups of blood selenium levels (mg/L)					P values
		Q1 (n=88) (≤0.07)	Q2 (n=86) (0.07–0.09)	Q3 (n=86) (0.09–0.14)	Q4 (n=87) (0.14–0.20)	Q5 (n=86) (>0.20)	
Age (years)	72.38 ± 5.28	74.53 ± 5.92	71.48 ± 5.22 ^a	71.84 ± 5.06 ^a	72.02 ± 4.90 ^a	72.00 ± 4.78 ^a	0.001
BMI, kg/m ²	22.65 ± 3.49	22.07 ± 4.11	22.92 ± 3.78	22.59 ± 2.85	22.96 ± 3.46	22.72 ± 3.09	0.460
Female (%)	216 (50.23)	48 (55.81)	38 (44.19)	41 (47.67)	45 (52.33)	44 (50.23)	0.602
Attended school (%)	241 (56.04)	45 (52.33)	53 (61.63)	48 (55.81)	50 (58.14)	45 (52.33)	0.697
Alcohol consumer (%)	108 (25.12)	20 (23.26)	25 (29.07)	22 (25.58)	13 (15.12)	28 (32.56)	0.092
Smoking (%)	125 (29.07)	31 (36.05)	27 (31.40)	29 (33.72)	17 (19.77)	21 (24.42)	0.104
Selenium, mg/L	0.11 (0.08, 0.18)	0.06 (0.06, 0.07)	0.08 (0.08, 0.09)	0.11 (0.10, 0.13) ^{a,b}	0.16 (0.15, 0.18) ^{a,b,c}	0.27 (0.23, 0.39) ^{a,b,c,d}	<0.001
SELENOP, µg/mL	4.59 (3.62, 6.75)	3.93 (2.50, 7.50)	4.00 (3.23, 5.47) ^a	4.60 (3.32, 5.09) ^{a,b}	5.17 (3.61, 7.57) ^{a,b}	5.18 (3.90, 7.79) ^{a,b}	<0.001
GPX, mmol/L	0.20 (0.16, 0.25)	0.20 (0.16, 0.24)	0.21 (0.13, 0.25)	0.18 (0.15, 0.24)	0.24 (0.18, 0.32) ^{a,b,c}	0.18 (0.16, 0.26) ^d	<0.001
Aβ42, pmol/L	2.48 (1.18, 4.08)	2.65 (1.46, 4.13)	2.61 (1.57, 3.60)	1.75 (0.96, 2.89) ^{a,b}	3.83 (1.84, 5.17) ^{b,c}	2.15 (0.70, 3.45) ^{a,b,d}	<0.001
Aβ40, pmol/L	25.15 (8.56, 49.30)	29.40 (9.77, 48.60)	36.10 (13.07, 52.50)	13.20 (4.86, 29.70) ^{a,b}	38.90 (15.45, 61.40)	15.95 (3.65, 43.27) ^{a,b,d}	<0.001
Aβ42/Aβ40	0.10 (0.07, 0.14)	0.10 (0.08, 0.13)	0.09 (0.07, 0.11)	0.13 (0.08, 0.17) ^{a,b}	0.10 (0.08, 0.13)	0.10 (0.07, 0.15) ^c	0.001

BMI, body mass index; SELENOP, selenoprotein P; GPX, glutathione peroxidase; Aβ42, β-amyloid 42; Aβ40, β-amyloid 40; Aβ42/40, the ratio of β-amyloid 42 and β-amyloid 40

[#]Described as numbers and proportions, means ± standard deviations, or median and interquartile range

^aCompared with Q1, *P* < 0.05

^bCompared with Q2, *P* < 0.05

^cCompared with Q3, *P* < 0.05

^dCompared with Q4, *P* < 0.05

Table 2 Results of quantile regression models of the association between the blood selenium (mg/L) and serum β-amyloid (pmol/L) levels

	β-coefficient (95% confidence interval)				
	5th percentile	25th percentile	50th percentile	75th percentile	95th percentile
Aβ42					
Crude ^a	-1.51 (-2.01, -1.00)*	-1.66 (-2.69, -0.63)*	-1.56 (-3.30, -1.43)*	-2.58 (-5.04, -0.13)*	-4.77 (-5.95, -4.72)*
Adjusted ^b	-1.58 (-2.17, -0.10)*	-1.42 (-2.44, -0.39)*	-1.49 (-2.94, -0.03)*	-2.21 (-4.32, -0.11)*	-3.71 (-7.28, -0.13)*
Aβ40					
Crude ^a	-2.91 (-5.22, -0.60)*	-12.73 (-24.26, -1.21)*	-27.73 (-53.55, -1.91)*	-39.87 (-66.53, -13.21)*	-50.06 (-79.80, -20.32)*
Adjusted ^b	-2.52 (-4.58, -0.46)*	-12.35 (-22.46, -2.24)*	-22.70 (-35.16, -12.87)*	-26.91 (-50.60, -3.23)*	-48.48 (-79.74, -17.22)*
Aβ42/Aβ40					
Crude ^a	-0.01 (-0.04, 0.02)	0.04 (0.02, 0.06)*	0.06 (0.02, 0.10)*	0.07 (-0.00, 0.14)	-0.02 (-0.69, 0.65)
Adjusted ^b	-0.02 (-0.03, 0.00)	0.03 (0.01, 0.05)*	0.06 (0.02, 0.10)*	0.07 (0.01, 0.14)*	-0.08 (-0.61, 0.46)

Aβ42, β-amyloid 42; Aβ40, β-amyloid 40; Aβ42/40, the ratio of β-amyloid 42 and β-amyloid 40

^aCrude model was not adjusted for any covariates

^bAdjusted for age, gender, education, BMI, tobacco smoking, and alcohol consumption

* *P* values less than 0.05

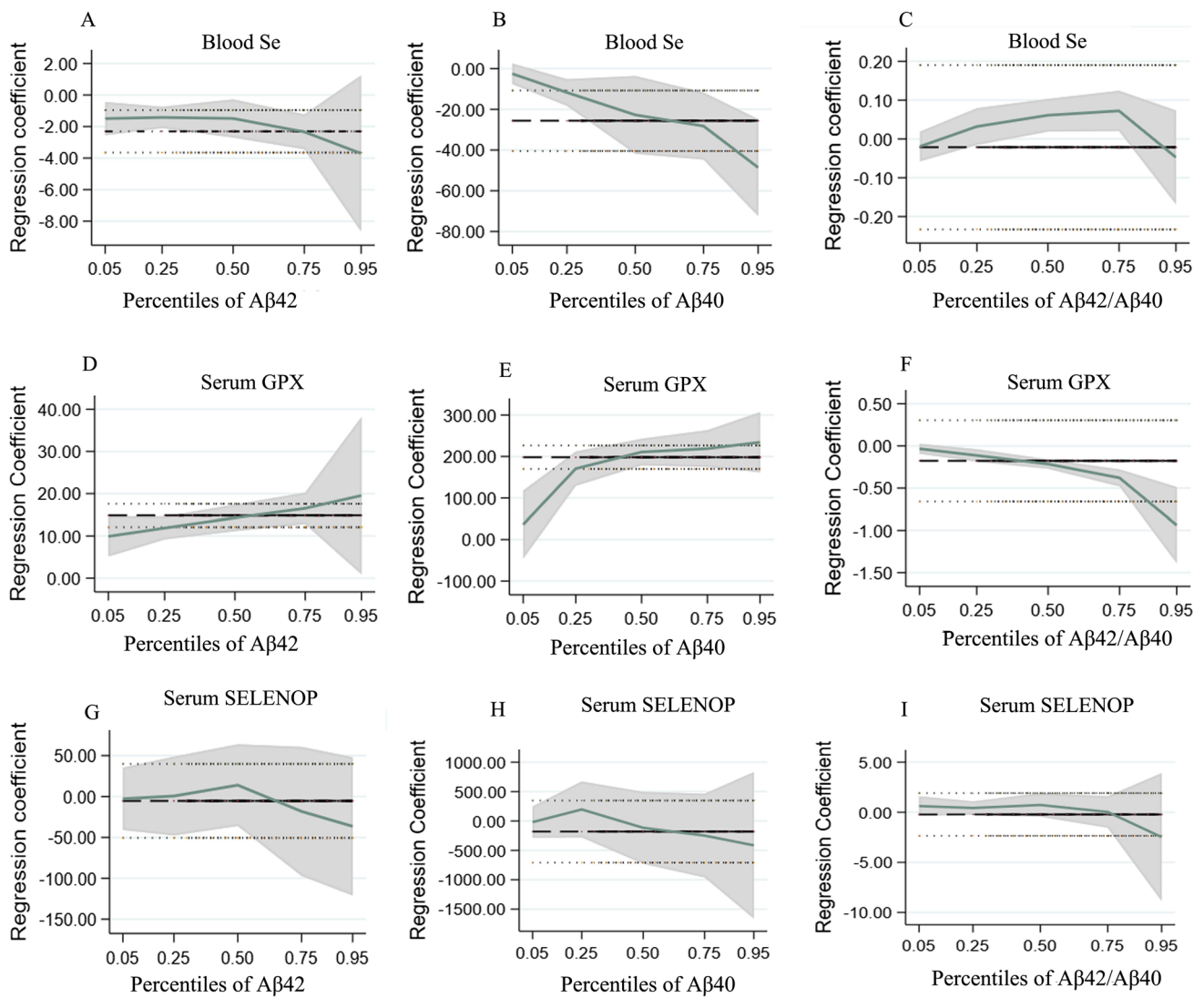


Fig. 1 a–i Graphics of quantile regression coefficient between serum A β and blood Se, serum GPX and SELENOP levels. The green solid line represented the quantile regression estimated coefficients and the gray area represented 95%CI of the corresponding parameters. The black dashed line represented the ordinary least squares (OLS) esti-

mate regression of the corresponding explanatory variable, and the area between the two black dashed lines represented the 95%CI of the OLS regression value. All estimations were adjusted for age, gender, education, BMI, tobacco smoking, and alcohol consumption.

all percentiles (Table 3). There was an overall increasing trend of the β -coefficients between serum GPX and A β 42 and A β 40 percentiles, indicating that the positive associations were generally stronger at the higher A β 42 and A β 40 percentiles than lower percentiles (Fig. 1D and E). For example, for every 1 mmol/L increase in serum GPX level, A β 42 increased by 11.21 pg/mL in the lowest A β 42 percentile, and A β 42 increased by 23.51 pg/mL in the highest A β 42 percentile. In addition, serum GPX was negatively associated with A β 42/A β 40 ratio at 25th (β : -0.11, 95%CI: -0.17, -0.06),

50th (β : -0.31, 95%CI: -0.31, -0.15), 75th (β : -0.38, 95%CI: -0.50, 0.26), and 95th (β : -0.88, 95%CI: -1.75, -0.02) percentiles, but there was no association at 5th percentile (Table 3). The negative associations were generally stronger at the higher A β 42/A β 40 percentiles than lower percentiles (Fig. 1F).

Serum SELENOP and A β

No statistically significant associations were observed between serum SELENOP, A β 42, A β 40, and A β 42/A β 40 ratio (Table 4 and Fig. 1).

Table 3 Results of quantile regression models of the association between serum glutathione peroxidase (mmol/L) and serum β -amyloid (pmol/L) levels

	β -coefficient (95% confidence interval)				
	5th percentile	25th percentile	50th percentile	75th percentile	95th percentile
Aβ42					
Crude ^a	11.44 (9.19, 13.70)*	12.62 (10.47, 14.78)*	14.34 (11.75, 16.92)*	17.33 (12.77, 21.88)*	30.09 (18.4, 41.78)*
Adjusted ^b	11.21 (9.23, 13.20)*	12.06 (9.78, 14.33)*	14.33 (11.69, 16.98)*	16.99 (13.07, 20.92)*	23.51 (17.46, 29.56)*
Aβ40					
Crude ^a	45.68 (34.53, 56.84)*	179.39 (151.05, 207.73)*	214.13 (177.62, 250.65)*	246.09 (200.38, 291.80)*	212.15 (121.92, 302.37)*
Adjusted ^b	37.83 (32.05, 43.60)*	169.29 (142.11, 196.46)*	211.7 (176.02, 247.38)*	223.4 (180.82, 265.98)*	237.36 (159.14, 315.58)*
Aβ42/Aβ40					
Crude ^a	-0.01 (-0.07, 0.06)	-0.11 (-0.16, -0.05)*	-0.21 (-0.29, -0.14)*	-0.41 (-0.54, -0.28)*	-0.94 (-1.42, -0.46)*
Adjusted ^b	-0.03 (-0.06, 0.01)	-0.11 (-0.17, -0.06)*	-0.23 (-0.31, -0.15)*	-0.38 (-0.50, -0.26)*	-0.88 (-1.75, -0.02)*

A β 42, β -amyloid 42; A β 40, β -amyloid 40; A β 42/40, the ratio of β -amyloid 42 and β -amyloid 40

^aCrude model was not adjusted for any covariates

^bAdjusted for age, gender, education, BMI, tobacco smoking and alcohol consumption

* *P* values less than 0.05

Table 4 Results of quantile regression models of the association between serum selenoprotein P (mg/mL) and serum β -amyloid (pmol/L) levels

	β -coefficient (95% confidence interval)				
	5th percentile	25th percentile	50th percentile	75th percentile	95th percentile
Aβ42					
Crude ^a	-3.21 (-39.49, 33.06)	8.84 (-35.73, 53.42)	17.83 (-36.18, 71.84)	20.56 (-60.34, 101.46)	-49.62 (-193.36, 94.13)
Adjusted ^b	-2.74 (-40.2, 34.72)	0.61 (-47.26, 48.48)	13.94 (-36.51, 64.38)	-18.18 (-103.34, 66.98)	-36.44 (-135.55, 62.68)
Aβ40					
Crude ^a	10.29 (-249.51, 270.09)	42.68 (-367.19, 452.56)	150.5 (-807.16, 1108.16)	-95.92 (-956.07, 764.22)	-446.74 (-1397.82, 504.35)
Adjusted ^b	-18.88 (-262.93, 225.18)	196.13 (-277.6, 669.86)	-115.95 (-898.33, 666.43)	-248.47 (-1211.47, 714.52)	-414.01 (-1453.18, 625.16)
Aβ42/Aβ40					
Crude ^a	-0.11 (-1.38, 1.17)	0.52 (-0.45, 1.49)	0.54 (-0.59, 1.66)	-0.71 (-2.8, 1.38)	-3.89 (-13.78, 6.00)
Adjusted ^b	0.62 (-0.23, 1.47)	0.43 (-0.6, 1.46)	0.73 (-0.57, 2.03)	0.03 (-1.92, 1.98)	-2.48 (-9.59, 4.64)

A β 42, β -amyloid 42; A β 40, β -amyloid 40; A β 42/40, the ratio of β -amyloid 42 and β -amyloid 40

^aCrude model was not adjusted for any covariates

^bAdjusted for age, gender, education, BMI, tobacco smoking and alcohol consumption

Discussion

In this cross-sectional study, we observed that blood Se was positively associated with serum SELENOP and GPX levels. Higher Se levels were significantly associated with lower serum A β 42 and A β 40 levels and associated with higher A β 42/A β 40 ratio. Higher serum GPX levels were significantly associated with higher serum A β 42 and A β 40 levels and associated with lower A β 42/A β 40 ratio. No association with serum SELENOP and A β levels was observed.

Total Se and A β

Our results were consistent with a cohort study in Italy involving 71 subjects, showing that higher inorganic Se concentrations in baseline CSF samples were inversely associated with lower CSF A β concentration in older adults [25]. However, a case-control study involving 30 AD patients, 35 vascular dementia patients, and 40 healthy controls in Indian reported that there was no association between plasma Se and A β 42 [26]. Nevertheless, the results of some in vitro

or in vivo studies could also support our results. A cellular experiment showed that selenite could reduce A β 40 generation in APP695 stably transfected Chinese hamster ovary cells [27]. An animal study reported that Se combined with fish oil could inhibit the activity of serum β - and γ -secretase in a mouse model of aging [28]. Another in vitro study also observed that selenite could reduce A β 40, A β 42, and sAPP beta production by reducing A β producing β -secretase and γ -secretase activities [29]. In addition, Se could attenuate lead exposure-induced changes in APP expression and A β production in Wistar rat hippocampus [30]. These findings suggest that blood Se may affect the APP processing by regulating the activity of β - and γ -secretase and further influence the serum A β levels.

Metabolic disturbances of A β homeostasis are the initial culprit in the pathogenesis of AD [31]. Recent reports suggested that high-accuracy assays of plasma A β 42/A β 40 ratio strongly predicted brain amyloidosis, and higher plasma A β 42/A β 40 ratios were correlated with lower A β deposition in brain tissue [5, 32]. However, no observational studies had focused on the association between Se and A β 42/A β 40 ratio before. As far as we know, our study is the first report showing that higher blood Se levels were associated with higher A β 42/A β 40 ratio, and we could further infer that higher blood Se levels could be associated with higher cognitive performance. Actually, this inference has been confirmed by our previous observational study in China [33] and another cross-sectional study in the USA [34]. Therefore, we speculate that Se could affect cognitive function by affecting A β metabolism; higher Se could reduce the generation of A β and increase A β 42/A β 40 ratio, which may delay the onset of cognitive impairment.

GPX and A β

Higher serum GPX was associated with higher serum A β and lower A β 42/A β 40 ratio in our study, which was opposite to the findings from total blood Se. This difference could be partly explained by the feedback effect of oxidative stress. As a group of antioxidant enzymes, the main biological role of GPX is to protect the organism from oxidative damage, for example, GPX4 could protect cortical neurons from oxidative injury and amyloid toxicity [35], and neurons of GPX1 gene knockout mice exhibited higher sensitivity to A β -induced oxidative stress [17]. Some studies suggested that the increase in antioxidant enzymes may represent a compensatory upregulation in response to increased oxidative stress [36, 37]. Interestingly, several studies have confirmed that oxidative stress could increase the expression of APP and the intracellular and secreted A β levels [38], and the increase of A β levels can aggravate oxidative stress in turn, forming a vicious cycle [39, 40]. This feedback effect could be found in other population-based studies; higher

plasma GPX activity was observed in patients with AD and in those with vascular dementia compared with healthy controls [41, 42].

In addition, other selenoproteins may be responsible for the opposite finding in our study. Except for GPX and SELENOP, there are several more selenoproteins which could be associated with AD. In vitro studies showed that selenoprotein S could prevent A β production by degrading C99 which could be cleaved to A β [43], and overexpression of selenoprotein M could inhibit the α/γ -secretase activity related to the protection of A β 42 production [44]. The thioredoxin reductase (TrxR) protein family may also have a protective roll in AD [10]. Therefore, we hypothesized that other selenoproteins or Se compounds might have covered up the inverse association of GPX compared with total Se.

SELENOP and A β

In the current study, we did not find significant association between serum SELENOP and A β levels. However, as the main Se transport protein, SELENOP has been supposed to be responsible for maintaining Se homeostasis and be vital for proper brain development and function [11, 45]. A postmortem study did observe an association that SELENOP was co-localized with A β plaques after examining postmortem brain tissue from individuals with the hallmark lesions of AD and individuals without these lesions [19]. In the triple-transgenic AD (3 \times Tg-AD) model of mice, overexpression of SELENOP can inhibit the aggregation of A β in the brain of mice [46]. Meanwhile, some studies found consistent results as ours. An animal experiment found no effect on the number of amyloid plaque nor intracellular A β -containing neurons [47]. And, a cross-sectional study reported that CSF SELENOP levels were not associated with CSF A β levels [25]. The conflicting findings suggested that SELENOP may not directly affect the formation and deposition of A β ; the function of SELENOP in brain health needs to be verified in future studies.

Strengths and Limitations

Our study has several strengths. Firstly, this is the first cross-sectional study with a relatively large sample to explore the relationship between peripheral blood Se levels and A β levels in the general population. Compared with CSF sample collection in other studies, blood sample collection in our study is less invasive. Therefore, blood A β measures are more suitable for epidemiological studies on general population. Secondly, our study design ensures an extensive range of Se exposure as subjects came from areas with different soil Se levels. Finally, the rural elderly Chinese participants

were mostly lifelong residents consuming local food and without Se supplementation; hence, the ascertained Se levels could be inferred to be lifelong Se exposure without the influence of supplements.

There are also some limitations. On the one hand, we did not determine oxidative stress biomarkers to explore the relationship between Se and A β . On the other hand, with the cross-sectional study design, we cannot make causal inferences on the association between Se and A β levels. Therefore, more longitudinal studies are necessary to confirm our results.

Conclusion

Our results suggest that higher Se levels are associated with lower serum A β 42 and A β 40 levels and with higher serum A β 42/A β 40 ratio, and the results are specific for different selenoproteins. More longitudinal evidence is needed to confirm our findings, and further studies are needed to elucidate the underline mechanisms.

Author Contribution Jiao Luo: data curation; software; formal analysis; writing—original draft; Liqin Su: conceptualization, funding acquisition, project administration, writing—review and editing; Xiaohong He: investigation, data curation, resources; Yegang Du: methodology, data curation, resources; Ning Xu: investigation, supervision; Rangpeng Wu: investigation, data curation; Yunfeng Zhu: investigation, data curation; Ting Wang: data curation, validation; Ranqi Shao: data curation, software; Frederick W. Unverzagt: resources, supervision; Ann M. Hake: resources, supervision; Yinlong Jin: resources, funding acquisition, supervision; Sujuan Gao: resources, funding acquisition, supervision.

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Declarations

Conflict of Interest The authors declare no competing interests.

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