INTRODUCTION

Vascular dementia (VaD) is a syndrome of varying degrees of cognitive and memory impairment caused by cerebrovascular injury (Jia et al., 2018; Qian Wang, Yang, Zhang, Zhao, & Xu, 2020). VaD is a common form of dementia and affects millions of subjects around all the world. The World Health Organization (WHO) points out that there are currently about 35.6 million people with dementia worldwide, and the number is estimated to increase by 7.7 million annually (Xu et al., 2017). And among these dementia cases, VaD accounts for about 15%–20%, which is the second most common dementia subtype after Alzheimer’s disease (AD). With the increase in human life expectancy, the number of VaD patients and the cost of treatment are expected to increase exponentially, which is attracting more
and more attention (Llorens et al., 2020). Since currently VaD lacks specific treatments to slow down or prevent its progression, how to deal with the dementia caused by the aging population has become a public health problem that all human society must face.

Neuronal Pentraxin 2 (NPTX2), also named neuronal activity-regulated pentraxin, is a secreted glycoprotein characterized by a cyclic multimeric structure (Osera et al., 2012; Tang et al., 2019). As a member of the pentraxins family, NPTX2 is highly conservative in evolution and is mainly expressed in the brain, spinal cord, and dorsal root ganglia (Moreno-Rodríguez, Perez, Nadeem, Malek-Ahmad, & Mufson, 2020; Pribiag & Stellwagen, 2014). In vivo, NTPX2 exerts various neurological effects by combining with the transmembrane protein neuronal pentraxin receptor (NPTXR) (Chapman, Shanmugalingam, & Smith, 2019). NPTX2 was thought to play a vital role in transmitting neurotransmitters and maintaining synaptic plasticity (Kimoto, Zaki, Bazmi, & Lewis, 2015; Lee et al., 2017). Although the human NPTX2 gene was identified as early as 1995, little is known about its molecular spatial structure and biological function (Hsu & Perin, 1995).

In recent years, more and more studies on the correlation between NPTX2 and diverse neurological diseases have been reported. Till date, there are no data about the role of NPTX2 in VaD. Therefore, in our current study, we assume that NPTX2 is involved in the pathogenesis of VaD and its potential predicted value was investigated in VaD patients.

2 | METHODS

2.1 | Subjects

The cross-sectional study was conducted in Shanxian Central Hospital between March 2017 and February 2020. Locals older than 60 years in Heze City were included in the study. A total of 188 subjects including 112 VaD patients and 76 controls were recruited in the Shanxian Central Hospital from March 2017 to February 2020. The study included 112 VaD patients and 76 healthy controls at the Shanxian Central Hospital. This study complies with the Declaration of Helsinki and was approved by the Clinical Research Ethics Committee of Shanxian Central Hospital.

Subjects or their guardian signed an informed consent form. This study complies with the Declaration of Helsinki and was approved by the Clinical Research Ethics Committee of Shanxian Central Hospital.

2.2 | Cognitive assessment

Montreal cognitive assessment (MoCA) was a widely used scale for global cognitive screens. The total score of MoCA is 30 points, which contains seven aspects such as orientation, executive function, language abilities, visuospatial abilities, short-term and long-term memory, abstraction, and attention. The lower the MoCA score, the worse the cognitive function. Generally, 26 points are used as the cutoff point for MoCA to diagnose cognitive impairment (Xu et al., 2019). The MoCA scores were assessed by trained attending physicians, who were unaware of the grouping and the clinical baseline data of all the subjects.

2.3 | Laboratory assays

Venous blood was drawn early in the morning at least 8 hr after fasting. After centrifugation at 1200 g for 10 min at 4°C, the serum was separated immediately and frozen in a −80°C refrigerator, and the biochemical indicators were subsequently determined. Blood biochemical index including free triiodothyronine 3 (FT3), free triiodothyronine 4 (FT4), thyroid-stimulating hormone (TSH), fasting blood glucose (FBG), hemoglobin A1c (HbA1c), low-density lipoprotein cholesterol (LDL-C), triglycerides (TG), total cholesterol (TC), and high-density lipoprotein cholesterol (HDL-C) was measured by a blood automatic biochemical analyzer (VetScan HM5, Model No: 250735; M/s Abaxis, Pvt. Ltd). The serum concentrations of NPTX2 were determined using a commercial enzyme-linked immunosorbent assay (ELISA) reagent (RD, Inc.). All experimental protocols refer to reagent instructions and previous research reports (Zhang, Tang, Hu, Wang, & Xu, 2020).

2.4 | Statistical analysis

The clinical baseline data were analyzed with descriptive statistics. Data are expressed as n or mean ± SD. The comparison of clinical baseline data between groups was analyzed using Student’s t test or chi-square test. The Spearman correlation coefficient was used to correlate the MoCA score with clinical baseline data. A multivariate linear regression analysis was carried out for evaluation of NPTX2 with the MoCA scores. All statistical evaluations are statistically significant if the two-tailed p value is less than .05. SPSS version 23.0 software (SPSS Inc, Chicago, IL, USA) was used for all the analyses in our current study.

3 | RESULTS

3.1 | Clinical baseline data

The study included 112 VaD patients and 76 healthy controls at the Shanxian Central Hospital from March 2017 to February 2020. Clinical baseline data including demographic indicators and blood biochemical indicators of all subjects were recorded after admission, and they are summarized in Table 1. The differences in demographic indicators including age (73.2 ± 5.3 vs. 72.9 ± 6.0), gender (male/
female: 64/48 vs 45/31), and education years (8.0 ± 2.4 vs 8.1 ± 2.6) between the groups are not significant \((p > .05)\). There was also no significant difference between the two groups in the comparison of blood biochemical indexes including FT3, FT4, TSH, FBG, HbA1c, LDL-C, TC, TG, and HDL-C. However, compared with healthy controls, VaD patients had significantly lower serum NPTX2 levels (196.8 ± 16.5 vs. 242.6 ± 19.4, pg/ml) and MoCA scores (22.8 ± 2.4 vs. 27.8 ± 1.3), and the difference between them was statistically significant \((p < .05)\).

3.2 | Spearman's correlation analysis

The correlation analysis results of the MoCA score of VaD patients and clinical baseline data are presented in Table 2. The results of Spearman’s correlation analysis showed that the serum NPTX2 level of VaD patients was significantly positively correlated with the MoCA score \((r = 0.347, p = .042)\). However, other clinical baseline data of VaD patients were not significantly correlated with MoCA scores \((p > .05)\).

3.3 | Multivariate regression analysis

Multivariate regression analysis was used to evaluate the effect of clinical baseline data on VaD patients’ MoCA score (Table 3). The results showed that serum NPTX2 level was an independent risk factor for VaD patients’ cognitive function, even after adjusting for clinical baseline data including age, gender, education, FT3, FT4, TSH, FBG, HbA1c, LDL-C, TC, TG, and HDL-C, this predictive value still exists \((\beta = 0.346, p = .039)\).

### TABLE 1 Baseline characteristics of all subjects

|                        | VaD (n = 112) | Control (n = 76) | p   |
|------------------------|---------------|------------------|-----|
| Age, years             | 73.2 ± 5.3    | 72.9 ± 6.0       | .719|
| Gender, male/female    | 64/48         | 45/31            | .778|
| Education, years       | 8.0 ± 2.4     | 8.1 ± 2.6        | .787|
| FT3, pmol/L            | 4.28 ± 0.34   | 4.33 ± 0.38      | .347|
| FT4, pmol/L            | 12.65 ± 1.19  | 12.57 ± 1.23     | .656|
| TSH, mIU/L             | 0.45 ± 0.08   | 0.43 ± 0.09      | .112|
| FBG, mmol/L            | 5.26 ± 0.51   | 5.28 ± 0.60      | .806|
| HbA1c, mmol/L          | 5.44 ± 0.62   | 5.39 ± 0.67      | .600|
| LDL-C, mmol/L          | 2.51 ± 0.22   | 2.48 ± 0.25      | .387|
| TG, mmol/L             | 1.62 ± 0.17   | 1.60 ± 0.18      | .441|
| TC, mmol/L             | 4.85 ± 0.67   | 4.83 ± 0.71      | .845|
| HDL-C, mmol/L          | 1.33 ± 0.12   | 1.35 ± 0.11      | .248|
| NPTX2, pg/ml           | 196.8 ± 16.5  | 242.6 ± 19.4     | <.001|
| MoCA                   | 22.8 ± 2.4    | 27.8 ± 1.3       | <.001|

### TABLE 2 Correlation coefficients between MoCA and baseline characteristics in patients with VaD

|          | r   | p     |
|----------|-----|-------|
| Age      | -0.214 | .087  |
| Gender, male | .352 | .130  |
| Education | .407 | .274  |
| FT3      | .318 | .572  |
| FT4      | .389 | .415  |
| TSH      | .436 | .383  |
| FBG      | .311 | .256  |
| HbA1c    | .394 | .428  |
| LDL-C    | .323 | .149  |
| TG       | .265 | .601  |
| TC       | .450 | .513  |
| HDL-C    | -.276 | .395  |
| NPTX2    | .347 | .042  |

### TABLE 3 Multivariable analysis between MoCA and baseline characteristics in patients with VaD

| Regression coefficient | p   | 95% CI          |
|------------------------|-----|----------------|
| Age                    | 0.363 | .119  | 0.736–1.062 |
| Gender, male           | 0.284 | .162  | 0.583–1.197 |
| Education              | 0.327 | .254  | 0.865–1.058 |
| FT3                    | 0.451 | .387  | 0.914–1.206 |
| FT4                    | 0.382 | .435  | 0.832–1.145 |
| TSH                    | 0.238 | .206  | 0.879–1.156 |
| FBG                    | 0.174 | .293  | 0.631–1.039 |
| HbA1c                  | 0.229 | .158  | 0.710–1.104 |
| LDL-C                  | 0.315 | .427  | 0.826–1.217 |
| TG                     | 0.341 | .370  | 0.739–1.060 |
| TC                     | 0.143 | .608  | 0.927–1.149 |
| HDL-C                  | 0.212 | .275  | 0.883–1.241 |
| NPTX2                  | 0.346 | .039  | 0.562–0.913 |

### Abbreviations

- FBG, fasting blood glucose; FT3, free triiodothyronine 3; FT4, free triiodothyronine 4; HbA1c, hemoglobin A1c; HDL-C, high-density lipoprotein cholesterol; LDL-C, low-density lipoprotein cholesterol; MoCA, Montreal cognitive assessment; NPTX2, neuronal pentraxin 2; TC, total cholesterol; TG, triglycerides; TSH, thyroid-stimulating hormone; VaD, vascular dementia.
The main finding of this study is that VaD patients have lower serum NPTX2 levels than normal controls, and this serum NPTX2 levels are also positively correlated with VaD patients’ MoCA scores. The association is independent of the effects of age, gender, education, FT3, FT4, TSH, FBG, HbA1c, LDL-C, TC, TG, and HDL-C. As far as I know, there has been no research report on the correlation between serum NPTX2 level and the cognitive function of VaD patients. The findings of this study can be extended to patients with the same characteristics.

The relationship between NPTX2 and a number of neurological diseases has been widely reported. One study had shown that the levels of NPTX2 in the inferior colliculus of audiogenic seizures (AGS)-susceptible P77PMC rats were significantly increased after AGS, suggesting that NPTX2 is involved in the pathogenesis of AGS (Li, Xu, & Jia, 2003). Not only in epilepsy, Lang and his colleagues found that NPTX2 plays an important role in Parkinson’s disease. Their results show that HOX transcript antisense intergenic RNA (HOTAIR) can increase the expression of NPTX2 in the substantia nigra through microRNA-221-3, thereby triggering excessive autophagy of dopamine neurons (Lang et al., 2020). This further enriches the mechanisms by which NPTX2 participates in the pathogenesis of Parkinson’s disease. Interestingly, studies have also shown that high levels of NPTX2 expression may be a biomarker for poor prognosis in human neuroblastoma (Bartolini et al., 2015). However, our results indicate that serum NPTX2 is decreased in VaD, which expands the spectrum of neurological diseases that NPTX2 may affect.

In recent years, NPTX2 has been confirmed to be closely related to the pathogenesis of AD. Galasko and his colleagues found that the levels of cerebrospinal fluid (CSF) synaptic protein NPTX2 in AD patients significantly decreased, and NPTX2 can be used as a biomarker for the progression of cognitive and global decline (Galasko et al., 2019). Further research shows that the predictive effect of NPTX2 is superior to the traditional biomarkers Aβ1-42 and Tau. Iowa State University’s research also shows that high baseline levels of NPTX2 in AD may have neuroprotective effects and can predict lower degrees of medial temporal lobe atrophy and cognitive decline (Swanson & Willette, 2016). Neuropathologists found that the expression of NPTX2 decreased significantly in the cerebral cortex of autopsy in AD patients (Hendrickson et al., 2015). All the above studies show that low levels of NPTX2 are involved in the pathogenesis of AD.

In addition to AD, NPTX2 is also associated with other forms of cognitive disorders. A newly published study points out that NPTX2 is significantly lower in genetic frontotemporal dementia (FTD) and is a novel synaptic marker to predict the progression of FTD disease (van der Ende et al., 2020). The results of BIOCARD Research Team show that the expression level of NPTX2 in mild cognitive impairment (MCI) is significantly lower than that of normal people, and NPTX2 may participate in the regulation of cognitive tasks in MCI (Soldan et al., 2019). Similar to the above studies, we found that serum NPTX2 levels of VaD patients were significantly lower than those of normal controls. Therefore, decreased NPTX2 may be associated with a variety of cognitive disorders.

Although some studies have observed fluctuations in the level of NPTX2 in various neurological diseases, the role of NPTX2 in it is largely unknown. NPTX2 can enhance the expression of brain-derived neurotrophic factor (BDNF), while BDNF with neuroprotective effect can also promote the expression of NPTX2 (Mariga et al., 2015). The protective effect of NPTX2 on cognition depends on the synapses (Gu et al., 2013; Miskimon et al., 2014). Under normal circumstances, NPTX2 is secreted into the synapse and cannot adhere to the cell surface. The perineuronal net (PNN), a proteoglycan network structure on cell surface, can capture NPTX2 to the presynaptic membrane and postsynaptic membrane to regulate synaptic homeostasis and plasticity (Van’t Spijker et al., 2019). However, the specific mechanism by which NPTX2 regulates synapses has not been fully elucidated.

There are several limitations in our study. First of all, we did not record the duration of VaD, so it is impossible to clarify the effect of different duration of VaD on the expression of NPTX2. Second, we did not dynamically observe the NPTX2 expression levels and MoCA scores of all the subjects. Third, there may be a mixed type of dementia among VaD subjects. Finally, it is a single-center clinical study, and the findings may not apply to people in other regions or ethnic groups.

The conclusion of this study is that VaD patients have lower serum NPTX2 levels than normal controls, and serum NPTX2 levels are positively correlated with cognitive function. Moreover, the correlation between serum NPTX2 and cognitive function in VaD patients is independent of age, gender, education, FT3, FT4, TSH, FBG, HbA1c, LDL-C, TC, TG, and HDL-C. Therefore, NPTX2 may be a novel serum biomarker of VaD. It is expected that future research will further reveal the pathogenesis of NPTX2 in VaD, which will have important clinical significance.

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### Conflict of Interests
The authors declare no conflict of interests.

### Author Contribution
Keke Shao and Cuihua Ma designed the experiments. Keke Shao, Shiqin Shan, and Wenwen Ru participated in the cognitive
assessments and ELISA experiments. Keke Shao drafted the manuscript and analyzed the data. Cuihua Ma edited the manuscript.

DATA AVAILABILITY STATEMENT
The data used to support the findings of this study are available from the corresponding author upon request.

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