1

1 C1q/CTRP1 exerts neuroprotective effects in TBI rats

2

by regulating inflammation and autophagy

- 3 Ming Pei^{1, 2, 3¶}, Chaoqun Wang⁴¶, Zhengdong Li^{2, 5}, Jianhua Zhang², Ping Huang²,
- 4 Jiawen Wang⁶, Jiang Huang⁶, Donghua Zou^{2,6*}, Yijiu Chen^{1, 2*}
- 5 1. West China School of Basic Medical Sciences & Forensic Medicine, Sichuan University, Chengdu,
- 6 Sichuan province, China
- 7 2. Shanghai Key laboratory of Forensic Medicine, Academy of Forensic Science, Ministry of Justice,
- 8 Shanghai, China
- 9 3. Institute of Forensic Science, Xuzhou Public Security Bureau, Xuzhou, Jiangsu province, China
- 10 4. Department of Pathology, Xuzhou Medical University, Xuzhou, Jiangsu province, China
- 11 5. Department of Forensic Medicine, School of Basic Medical Sciences, Fudan University, Shanghai,
- 12 China
- 13 6. School of Forensic Medicine, Guizhou Medical University, Guiyang, Guizhou province, China
- 14 *Corresponding author
- 15 E-mail: zoudh@ssfjd.cn (DZ)
- 16 E-mail: cyj1347@163.com (YC)
- 17 ¶ Ming Pei and Chaoqun Wang contributed equally to this work.
- 18

2

19 Abstract

20	Objective: C1q/CTRP1 is a newly discovered adiponectin protein, which is highly expressed in adipose
21	and heart tissues. Recent studies have revealed that C1q/CTRP1 can regulate metabolism and inhibit
22	inflammation. CTRP1 is also expressed in brain tissues and vascular cells of human and rat, and research
23	on cerebral hemorrhage and cerebral ischemia-reperfusion injury demonstrates that the CTRP family can
24	attenuate secondary brain injury and exert neuroprotective effects. Thus, this study was designed to
25	explore the role of CTRP1 in traumatic brain injury (TBI) and the underlying mechanism.
26	Main methods: Rats were assigned into rCTRP1 group, vehicle group, and sham group. Modified
27	Feeney's method was used to establish a closed traumatic brain injury model. Morris water maze was
28	used for directional navigation, reverse searching and space exploration tests in rats. In addition, Golgi-
29	Cox staining was utilized to visualize neurons, dendrites and dendritic spines. ELISA was conducted to
30	detect the levels of inflammatory factors (IL-6 and TNF- α). Finally, Western blot was adopted to detect
31	the relative expression of <i>p</i> -mTOR and autophagy-related proteins (Beclin-1 and LC3-II).
32	Results: CTRP1 improved the behavioral and histopathological outcomes, inhibited the inflammatory
33	response, activated mTOR and decreased autophagy-associated protein synthesis in TBI rats.
34	Conclusion: CTRP1 exerts neuroprotective effects in TBI rats by regulating inflammation and
35	autophagy and has potential therapeutic properties after TBI.
36	Keywords: Autophagy; CTRP1; Inflammation; Neuroprotective effect; Traumatic brain injury
37	Introduction
38	Traumatic brain injury (TBI) is the main cause of traumatic death and disability globally. The common
39	causes of TBI include violence, traffic accidents, tumble, falls, and sports accidents [1]. TBI is generally

40 divided into primary and secondary brain injury. To be specific, primary TBI is caused by direct external

41	force acting on the central nervous system. Primary TBI often leads to neurometabolic disorders,
42	hippocampal synapse damage, neuron, astrocytic degeneration (CA1/CA3 layer, dentate gyrus) and
43	glutamate excitotoxicity. In addition, with the subsequent destruction of the blood-brain barrier (BBB)
44	and the persistence of neurogenic inflammation [2], there are secondary changes in nerve tissues,
45	including a series of pathophysiological changes, energy metabolism disorders and inflammatory
46	response [3-5], which generally aggravate neuronal necrosis, dendrite and synapse damage [2]. Both
47	primary and secondary TBI can directly cause cognitive and behavioral dysfunction in patients, which
48	seriously affects the patient's quality of life [6,7]. Therefore, how to attenuate the damage of primary
49	and secondary TBI on the nerve tissue and to enhance the neuroprotective effects has become the present
50	research focus of neuroscience and trauma science.
51	C1q/TNF- related protein-1 (CTRP1) is a protein cytokine containing 281 amino acids, which belongs
52	to the CTRP family. And the CTRP family is mostly involved in regulating inflammation and metabolism
53	[8]. Similar to other CTRP family members, CTRP1 is expressed in adipose and heart tissues [9,10].
54	CTRP1-deficient mice show increased myocardial infarction area caused by ischemia reperfusion injury
55	(IRI), cardiomyocyte apoptosis and expression of pro-inflammatory genes, while the up-regulation of
56	CTRP1 protein expression can attenuate myocardial injury, indicating that CTRP1 can regulate
57	cardiomyocyte metabolism, inhibit inflammatory response and protect the damaged cardiomyocytes [10].
58	Additionally, inflammatory response and pro-inflammatory cytokines can increase the secretion of
59	CTRP1 [11-13]. Relevant studies have shown that the increased CTRP1 protein expression can suppress
60	the inflammatory response caused by cerebral IRI [14]. Moreover, CTRP1 can regulate the autophagy of
61	glial cells by activating the Akt/mTOR signaling pathway, thereby exerting neuroprotective effects [14].
62	Based on the above research, this study aimed to confirm the effectiveness of CTRP1 on neuroprotection,

4

63 and to investigate its effect on memory, cognitive and behavioral functions in TBI rats. In addition, we

64 further revealed the mechanism of the neuroprotective effects of CTRP1 in TBI rats.

65 1. Materials and methods

66 1.1 Animals

67 The use of animals and the animal procedures were conducted following the guidelines approved and 68 formulated by the Animal Care and Use Committee of the Xuzhou Medical University. A total of 80 69 healthy male specific pathogen free (SPF) Sprague-Dawley (SD) rats (10 to 12 weeks old, 220 to 250 g 70 of weight) were purchased from the Animal Experimental Center of Xuzhou Medical University. Rats 71 were maintained under SPF condition, with the temperature of 25 ± 1 °C, the relative humidity of 40%-60% 72 and the light/dark cycle of 12 h/12 h. Rats were freely accessible to food and water. The rats were 73 adaptively fed for one week before the experiment. All surgery was performed under sodium 74 pentobarbital anesthesia, and all efforts were made to minimize suffering.

75 **1.2 Construction of the TBI model**

76 The modified Feeney's method was used to establish the closed TBI model¹⁶. In brief, rats were fasted 77 and deprived of water for 12 h before operation. The head was fixed on the stereotaxic apparatus, 78 followed by shearing and disinfection of the operation area. Afterwards, the scalp was cut along the 79 midline sagittal of the head to expose the right parietal skull. To penetrate the skull and open a circular 80 window 5 mm in diameter while preventing endocranial injury, a dental drill was used on the skull at a 81 distance of 3.5 mm to the right of the skull midline and about 0.2 mm from the bregma. Subsequently, 82 a 40 g skull batting stick was released from a height of 20 cm of the stereotaxic apparatus vertically 83 fixed to the cannula, so as to control the subsidence depth of 0.2 cm and the diameter of striking end of 84 approximately 0.2 cm, causing contusion of the right hemisphere. Finally, the skull defect was sealed

85	with bone wax	, and the scal	p was intermittently	y sutured. Rats	in the sham gr	oup were subjected	to skull
----	---------------	----------------	----------------------	-----------------	----------------	--------------------	----------

- 86 window exposure without further hitting. Symptoms such as limb twitch, urinary incontinence, nasal
- 87 bleeding and a few seconds of apnea were shown in rats after the hit. Based on the neurological
- 88 severity score (NSS) at 0.5 h and 1 h after injury, the rats with common TBI were screened, excluding
- 89 eight rats that died or did not present moderate TBI.

90 1.3 Grouping and drug administration

- 91 The remaining 72 rats were randomly grouped into three categories matching the NSS results,
- 92 including the rCTRP1 group (TBI + rCTRP1 recombinant protein), which was given an acute
- 93 intracerebroventricular injection of 80 µg/kg mouse-derived rCTRP1 recombinant protein
- 94 (ANNORON, China) every 24 h for up to 7d from half an hour after TBI; the vehicle group (TBI
- +vehicle), which was administered with the same amount of normal saline at equal frequency to that of
- 96 the rCTRP1 group after craniocerebral trauma; and the sham group, which received a craniotomy
- 97 without TBI. Three time points after TBI, namely, 24 h, 72 h and 1 w, were chosen, and eight rats in
- 98 each group were examined at each time point.

99 1.4 Morris water maze

100 The Morris water maze consisted of a round dark metallic pool 160 cm in diameter and 60 cm in depth

- 101 that was filled with water $(22\pm0.5^{\circ}C)$. Water was made opaque by the addition of a dark nontoxic
- 102 water-based paint to a depth of 50 cm and surrounded by a dark curtain. The pool was virtually divided
- 103 into four quadrants. An escape platform (12 ×12 cm) was submerged at 1 cm below the opaque water
- surface and located in the center of one quadrant of the maze, approximately 30 cm from the edge of
- 105 pool. Randomized to one of the four quadrants, rats were allowed to search for the hidden platform for
- 106 1 min. The rats were placed on the platform manually for 15 s if they exceeded the allotted time. Rats

107 were trained for five consecutive days before the experiment. Then, at 24 h, 72 h and 1 w after TBI,

- 108 rats were allowed to seek the platform in the 1-min test. After the water maze, rats were sacrificed.
- 109 Hippocampal tissue samples were immediately removed and stored at -80°C.

110 **1.5 Golgi-Cox staining and microscopy procedures**

111 One side of the hippocampus of rats in each group was transferred into the impregnation solution at room 112 temperature based on the instructions of the Golgi-Cox OptimStain Prekit (HiTO, USA). The 113 impregnation solution was placed and stored at room temperature in dark for two weeks. The tissue was 114 subsequently transferred into solution-3 for 12 h at 4°C. Then solution-3 was replaced and stored for 48 115 h at 4°C. The temperature of isopentane was attenuated to -70 °C using dry ice. The hippocampal tissue 116 was then immersed in the isopentane and cooled down for approximately 40 s, and the absorbent paper 117 was used to remove excess isopentane from the tissue. The freezing microtome was pre-cooled to -19°C, 118 which was used to slowly cut the tissue into sections 120-um in thickness. The sections were transferred 119 to a gelatin-coated slide and dried at room temperature in darkness overnight. Thereafter, the slides were 120 rinsed in distilled water twice for 3 min each time. Later, the slides were placed in the staining mixing 121 solution for 10 min and then in renewed distilled water twice for 4 min each time. The slides were then 122 dehydrated in 50%, 75% and 95% ethanol (5 min each) and in 100% ethanol thrice (5 min each). 123 Afterwards, the slides were cleaned in xylene twice (5 min each), sealed with a coverslip, and viewed by 124 light microscopy. At least 10 neurons were randomly selected in each slice (magnification 200 times) by 125 a trained observer blind to the experimental condition. Equi-distant (10 µm) concentric rings were placed 126 over the tracings of the dendritic tree. The total dendritic arborization and dendritic length were measured 127 by the amount of ring intersections with the dendritic tree. Over 10 primary dendritic branches at a length 128 of $\ge 20\mu m$ were traced (at 1000×). The amount of dendritic spine was computed utilizing the analysis

129 system of Image pro-plus 6.0.

130 **1.6 Detection of IL-6 and TNF-α by ELISA**

- 131 Hippocampal tissue was harvested immediately and homogenized in lysis buffer, followed by
- 132 centrifugation at 8000g for 10 min at 4°C to collect the supernatants. The levels of IL-6 and TNF- α
- **133** were determined by ELISA kit purchased from Abcam Company (ab100785 and ab100772, Abcam,
- 134 America) according to the manufacturer's instruction.

135 **1.7 Western blot analysis**

136 Hippocampal tissue was centrifuged at 12000 rpm for 15 min and homogenized in ice-cold tissue lysis 137 buffer (50 mm Tris, PH 7.5, 0.15 mm NaCl, 2% NP-40, 0.5% sodium deoxycholate, 4% SDS, and 138 protease and phosphatase inhibitor cocktails) for 15 min. In every supernatant fraction, the BAC protein 139 assay kit was used to measure the total protein concentration (Pierce, Rockford, IL, USA), and 15 µg 140 Beclin-1, LC3-II and p-mTOR were separated by electrophoresis and then transferred onto the 141 nitrocellulose membranes (Bio-Rad; Trans-Blot Turbo Transfer System). After several washes with 142 TBST buffer, the membranes were blocked for 2 h with the blocking buffer (LI-COR Biosciences, 143 Lincoln, Nebraska, USA) and later incubated with rabbit anti-Beclin-1 (1:4000; Abcam, America), rabbit 144 anti-LC3-II (1:2000; Abcam, America), polyclonal rabbit Anti-mTOR (1:5000, Abcam, America), and 145 rabbit anti-GAPDH (1:10,000; Abcam, America) in TBST at 4°C overnight. After four washes, the HRP-146 linked secondary antibodies (1:5,000, Boster Bioengineering, China) were incubated in dark for 1 h. The 147 Image J software was used to quantify the bands (NIH, Bethesda, MD, USA).

148 **1.8 Statistical analysis**

149	The description of continuous	variables in our dataset wa	as expressed as mean \pm standard de	eviation. The
-----	-------------------------------	-----------------------------	--	---------------

- 150 repeated measurement ANOVA was employed to analyze the behavioral statistics during the training
- 151 period. The one-way ANOVA was utilized to examine the spine density, dendritic length and
- branching, as well as the ELISA and Western blot data. Further multiple comparisons were undertaken
- 153 via the LSD method if the overall ANOVA analysis was significant. All the statistical analyses in the
- 154 present study were implemented by the SPSS software (version 19.0 for Windows, SPSS Inc., USA)
- and the significance level was set to be 0.05 throughout our study.

156 **2 Results**

157 **2.1 The expression of CTRP1 in hippocampus**

- 158 The expression of CTRP1 in rat hippocampus was detected at 24 h after TBI. The results showed that
- the expression of CTRP1 increased in rCTRP1 group and vehicle group compared with sham group (p
- 160 <0.001). In addition, the expression of CTRP1 was significantly higher in rCTRP1 group than that in
- 161 vehicle group (p<0.001) (Fig 1A B).
- **162** Fig 1. Expression of CTRP1 in hippocampus at 24h after TBL. *** p < 0.001 vs. sham group, ###p
- 163 < 0.001 vs vehicle group.
- 164 **2.2 Results of memory and spatial learning**

165 **2.2.1 Spatial orientation ability**

166 The escape latency is defined as the time from entering water to boarding the platform of rats, which is

- 167 one of the commonly indicators of spatial orientation ability. In this study, the escape latency time was
- detected in rats of three different groups at 24 h, 72 h, and 1 w after craniocerebral trauma. As a result,
- the escape latency was significantly longer in rCTRP1 group and vehicle group than that in sham group
- 170 at the first two time points (p<0.005), and there was no significant difference in the escape latency

171 between rCTRP1 group and sham group at 1 w after craniocerebral trauma. However, the time was

172 significantly shorter in rCTRP1 group than that in vehicle group at 1 w after TBI (p<0.05) (Fig 2A).

173 Fig 2. The results of two Morris water maze tests in three groups at 24h, 72h and 1w after TBI. (A)

- 174 Escape latency to locate the target platform. (B) Escape latency of reverse searching to locate the target
- 175 platform. (C) After the platform was removed, the number of entries into the target quadrant (TA). *p<
- 176 0.05 vs. sham group, ** p < 0.005 vs. sham group, ***p < 0.001 vs. sham group, # p < 0.05 vs vehicle
- 177 group.
- 178 **2.2.2 Reverse searching ability**

179 The second quadrant of the platform was changed to the fourth quadrant, followed by re-examination of

180 the escape latency of rats, which was used as an indicator of working memory ability. The escape latency

181 of reserve searching was significantly prolonged in rCTRP1 group and vehicle group than that in sham

182 group at 24 h and 72 h after craniocerebral trauma (p<0.05), and there was no significant difference in

183 the time between rCTRP1 group and sham group at 1 w after craniocerebral trauma. However, the time

184 was significantly shorter in rCTRP1 group than that in vehicle group at 1 w after TBI (p<0.05) (Fig 2B).

185 **2.2.3 Spatial exploration ability**

186 The number of entries into the target quadrant (TA) after removing the underwater platform is one of the

187 commonly used indicators to detect the spatial exploration ability of rats. In this study, the number of

- 188 entries into the TA was significantly less in rCTRP1 group and vehicle group than that in sham group at
- 189 24 h and 72 h after TBI (p<0.05), and there was no significant difference in the number between rCTRP1
- 190 group and sham group at 1 w after craniocerebral trauma. While the number significantly increased in
- 191 rCTRP1 group compared to vehicle group at 1 w after TBI (p<0.05) (Fig 2C).

192 2.3 Changes in dendrites and dendritic spines

193	The alterations in dendrites and dendritic spines were observed at 24 h, 72 h and 1 w after TBI,
194	respectively. The length and branches of dendrites significantly decreased in rCTRP1 group and
195	vehicle group than those in sham group at 72 h and 1 w after TBI (p<0.05), and there was no
196	significant difference in those between rCTRP1 group and sham group at 24 h after craniocerebral
197	trauma. Moreover, the length and branches of dendrites significantly increased in rCTRP1 group
198	compared to those in vehicle group at 1 w after TBI (p<0.05) (Fig 3ABC, Fig 4A). The dendritic spines
199	decreased in rCTRP1 group and vehicle group than those in sham group at 24 h and 72 h after TBI
200	(p<0.005), and there was no significant difference in those between rCTRP1 group and sham group at 1
201	w after craniocerebral trauma. However, the dendritic spines significantly increased in rCTRP1 group
202	compared to those in vehicle group at 72 h and 1 w after TBI (p<0.05) (Fig 3 DEF, Fig 4B).
203	Fig 3. Changes in dendrites and dendritic spines at 72h after TBI. (A) (the black bar is 50 μ m) and
204	(D) (the black bar is 10 μ m), sham group; (B) and (E), vehicle group; (C) and (F), rCTRP1 group.
205	Fig 4. The histogram of changes in dendrites and dendritic spines at 24h, 72h and 1w after TBL
206	(A) The number of ring intersections of the dendritic arborization. (B) The density of the dendritic
207	spines. * p < 0.05 vs. sham group, **p<0.005 vs. sham group, *** p<0.001 vs. sham group, # p <
208	0.05 vs vehicle group.
209	2.4 Effects of CTRP1 protein on neuroinflammation
210	As presented in Fig. 5, after TBI, enhanced concentrations of TNF- α in rCTRP1 group and vehicle group

211 were observed at 24 h, 72 h and 1 w after TBI compared to sham group (p < 0.05); typically, the

- 212 concentrations rapidly increased within 24 h after TBI and then slowly recovered to normal levels. The
- 213 change trend of IL-6 concentrations in the three groups was similar to that of TNF- α at the three time
- points. The IL-6 and TNF-α levels in rCTRP1 group significantly decreased compared to those in vehicle

215	group at 24 h and 72 h afte	: TBI (p<0.001)	. The results suggested that the	e inflammatory response was
-----	-----------------------------	-----------------	----------------------------------	-----------------------------

- 216 relatively significant within 1 w after craniocerebral trauma and CTRP1 exerted anti-inflammatory
- effects in the early stage of injury.

218 Fig 5. The changes of TNF-α and IL-6 levers in each group at different time points after

- **219** craniocerebral trauma. * p < 0.05 vs. sham group, ** p < 0.005 vs. sham group, *** p < 0.001 vs.
- 220 sham group, ### p < 0.001 vs vehicle group.

221 **2.5 Effects of CTRP1 protein on autophagy**

222 Western blot was used to detect the expression of Beclin-1, LC3-II and *p*-mTOR in the hippocampus

at 24 h, 72 h and 7 d after TBI and to demonstrate the changes in autophagy after TBI and the regulatory

roles of CTRP1 on autophagy (Fig 6ABCD). The expression of the autophagy-related proteins Beclin-1

and LC3-II in hippocampus of vehicle group was significantly higher than that in sham group at 24 h,

226 72 h, and 1 w after craniocerebral trauma (p<0.05). The properly increased level of autophagy can

227 attenuate cellular functions and decrease energy consumption, which is one of the self-protection

228 mechanisms of damaged cells. However, intense autophagy can also cause cell death. The expression of

- Beclin-1 and LC3- II in hippocampus tissue in rCTRP1 group was significantly lower than that in
- vehicle group at three time points (p<0.001), indicating that CTRP1 attenuated autophagic injury after
- craniocerebral trauma. The expression of *p*-mTOR was significantly lower in vehicle group than that in
- sham group at 24 h, 72 h and 7 d after TBI (p < 0.001). Additionally, the expression of *p*-mTOR was
- 233 significantly higher in rCTRP1 group than that in vehicle group at three time points (p<0.001), suggesting

that the CTRP1 recombinant protein was involved in the regulation of the mTOR pathway.

Fig 6. Western-blot analysis was used to detect the expression of *p*-mTOR, Beclin-1 and LC3-II

in the hippocampus of each group after craniocerebral trauma. (A) Relative expression of *p*-mTOR,

Beclin-1 and LC3- II in each group at three time points. (B-D) The histogram of the expression of pmTOR, Beclin-1 and LC3- II in each group at 24h, 72h and 1w after craniocerebral trauma. *p < 0.05 vs. sham group, **p < 0.005 vs. sham group, ***p < 0.001 vs. sham group, ###p < 0.001 vs vehicle group.

241 **3 Discussion**

242 CTRP1 is a secretory glycoprotein that is expressed in the heart, liver, kidney, placenta, brain and 243 interstitial vascular cells (including macrophages) in human and rat [15-18]. Previous studies have shown 244 that the level of circulating CTRP1 elevates in patients with type 2 diabetes [19-20], coronary artery 245 disease [21] and hypertension [22], and CTRP1 may be involved in the vasculitis and coagulation process 246 in the acute phase of Kawasaki disease [23]. Studies have also demonstrated that CTRP1 possesses 247 insulin-sensitizing effects [24,25]. Recent studies have revealed that the expression of CTRP1 248 significantly increases in the serum of stroke patients, and is positively correlated with the high-249 sensitivity C-reactive protein, suggesting that CTRP1 may exert a neuroprotective effect after ischemic 250 stroke [14]. In our study, we found that the expression of CTRP1 increased in the hippocampus of rats 251 at 24 h after craniocerebral trauma and CTRP1 improved the behavioral and histopathological outcomes, 252 indicating that CTRP1 might exert a neuroprotective effect after craniocerebral trauma, which was 253 consistent with the above studies concerning stroke.

A variety of studies have reported that CTPR1 can inhibit inflammation. For instance, CTPR1 protects the heart from IRI by attenuating myocardial cell apoptosis and inflammation [26]. CTRP1 is associated with ischemic heart disease. The myocardial infarction area, cardiomyocyte apoptosis and proinflammatory gene expression following IRI are up-regulated in CTPR1 knockout mice, compared with wild-type (WT) mice. In contrast, the up-regulation of CTRP1 protein can attenuate myocardial damage

259	after IRI in WT mice. Treatment of cardiomyocytes with CTRP1 can reduce the hypoxia-reoxygenation-
260	induced apoptosis and the LSP-stimulated expression of proinflammatory cytokines, which can be
261	reversed by inhibiting the sphingosine-1-phosphate (S1P) signaling [26]. Therefore, CTRP1 can be used
262	as an endogenous cardioprotective factor with anti-apoptotic and anti-inflammatory effects. In addition,
263	the detection of CTRP1 expression in the plasma of patients with different types of stroke reveals that
264	the expression level of CTRP1 in atherosclerosis-related strokes is significantly lower than that in other
265	types of strokes, indicating that CTRP1 may be involved in the occurrence and development of stroke
266	via atherosclerosis and the mechanism is associated with the suppression of inflammatory response of
267	macrophages, thereby protecting brain function [27]. In our study, we found that rCTRP1 decreased the
268	expression of inflammatory factors (TNF- α and IL-6) in the hippocampus of rats after TBI.
269	Autophagy exerts a dual role in TBI. On the one hand, enhanced autophagy can remove the damaged
270	proteins and organelles, and reduce mitochondrial energy consumption, which are beneficial to maintain
271	cell homeostasis. On the other hand, the enhanced autophagy can simultaneously destroy functional
272	macromolecular substances and organelles in cells to aggravate cell damage [28]. Autophagy is regulated
273	by a variety of molecular mechanisms, and the Akt/mTOR signaling pathway plays an important role in
274	regulating autophagy [29]. In addition, the Akt/mTOR pathway attenuates neuronal apoptosis by
275	suppressing calcification-dependent pathways, which has a protective effect in neurological diseases
276	[30,31]. CTRP1 can specifically activate the Akt (protein kinase B) [32] and MAPK signaling pathways
277	in myotubes of the differentiated mice [25]. Huilin Wang et al. [14] found that activation of the
278	Akt/mTOR pathway by IGF-1 significantly inhibited the promoting role of si-CTRP1 in microglial
279	autophagy, while A6730 inhibited the Akt and reversed the inhibitory effect of CTRP1 recombinant
280	protein on microglial autophagy. Consistent with previous reports, we discovered that rCTRP1 reduced

- the expression of hippocampal autophagy-related proteins LC3-II and Beclin-1 by activating the mTOR
- signaling pathway in rats, and exerted an inhibitory effect on autophagy in hippocampus tissue after TBI,
- thereby decreasing autophagic injury caused by craniocerebral trauma. These findings suggest that
- rCTRP1 plays a neuroprotective effect by activating the mTOR signaling pathway.

285 4. Conclusion

- 286 In this study, we find that the CTRP1 recombinant protein can improve the behavioral and
- 287 histopathological outcomes, inhibit inflammatory response, activate mTOR and decrease autophagy-
- associated protein synthesis in TBI rats. Therefore, CTRP1 exerts neuroprotective effects in TBI rats by
- regulating inflammation and autophagy and has potential therapeutic properties after TBI.

290

291 CONFLICT OF INTEREST

292 The authors declare that there are no conflicts of interest in the authorship or publication of the

293 contribution.

294

295 **REFERENCES**

DeKosky ST, Asken BM. Injury cascades in TBI-related neurodegeneration. Brain Inj. 2017;31(9):1177–
 82.
 Shandra O, Winemiller A, Heithof B, Munoz-Ballester C, George, Kijana, et al. Repetitive diffuse mild
 traumatic brain injury causes an atypical astrocyte response and spontaneous recurrent seizures. J
 Neurosci. 2019;39(10):1944–63.

301 3. Adams JH, Doyle D, Graham DI, LAWRENCE AE, MCLELLAN DR, GENNARELLI TA, et al. The contusion
 302 index: A reappraisal in human and experimental non-missile injury. Neuropathol Appl Neurobiol.
 303 1985;11(4):299-308.

- 4. Alessandrini A, Namura S, Moskowitz MA, Bonventre JV. MEK1 protein kinase inhibition protects
 against damage resulting from focal cerebral ischemia. Proc Natl Acad Sci USA. 1999; 96(22): 1286612869.
- 307 5. Artuso M, Esteve A, Bresil H, Vuillaume M, Hall J, et al. The role of the Ataxia telangiectasia gene in
 308 the p53, WAF1/CIP1(p21)- and GADD45-mediatedresponse to DNA damage produced by ionizing
 309 radiation. Oncogene. 1995; 11(8):1427-1435.

15

6. Kraus JF, McArthur DL. Epidemiologic aspects of brain injury. Neurol Clin. 1996; 14(2):435-50.
7. Raymont V, Salazar AM, Krueger F, Grafman J. "Studying injured minds": the- Vietnam head injury
study and 40 years of brain injury research. Front Neurol. 2011; 2:1-13.
8. Weng H, Pei Q, Yang M, Zhang J, Yi Q. Hypomethylation of C1q/tumor necrosis factor-related protein-
1 promoter region in whole blood and risks for coronary artery aneurysms in Kawasaki disease. Int J
Cardiol. 2020; 307:159-163.
9. Yang Y, Liu S, Zhang RY, Luo H, Chen L, He WF, et al. Association between c1q/tnf-relatedprotein-
1levelsinhumanplasmaand epicardial adipose tissues and congestive heart failure. Cell. Physiol.
Biochem.2017; 42 (5) :2130–2143.
10. Yuasa D,Ohashi K,Shibata R,Ouchi N, Kataoka Y, Kambara T, et al. C1q/TNF-related protein-1
functions to protect against acute ischemic injury in the heart. FASEB J. 2016;30 (3) :1065–1075.
11. Lu L, Zhang RY, Wang XQ, Liu ZH, Shen Y, Ding FH, et al. C1q/TNF-related protein-1: an adipokine
marking and promoting atherosclerosis. Eur Heart J. 2016;37(22):1762–71.
12. Muendlein A, Leiherer A, Saely C, Ebner J, Geiger K, Brandtner EM, et al. The novel adipokine CTRP1
is significantly associated with the incidence of major adverse cardiovascular events. Atherosclerosis.
2019; 286:1–6.
13. Kim D, Park SY. C1q and TNF related protein 1 regulates expression of inflammatory genes in
vascular smooth muscle cells. Genes Genomics. 2019;41(4):397–406.
14. Wang HL, Liu QH, Zhang XJ. C1q/tumor necrosis factor-related protein-1 attenuates microglia
autophagy and inflammatory response by regulating the Akt/mTOR pathway. Life
Sciences.2020;256:117992.
15. Si Y, Fan W, Sun L. A Review of the Relationship Between CTRP Family and Coronary Artery Disease.
Curr Atheroscler Rep.2020;22(6) :22.
16. Wong GW, Krawczyk SA, Kitidis-Mitrokostas C, Revett T, Gimeno R, Lodish HF. Molecular,
biochemical and functional characterizations of C1q/ TNF family members: adipose-tissue-selective
expression patterns, regulation by PPAR-gamma agonist, cysteine-mediated oligomerizations,
combinatorial associations and metabolic functions. Biochem. J. 2008; 416 (2) :161-77.
17. Schaffler A, Buechler C. CTRP family: linking immunity to metabolism. Trends Endocrinol. Metab.
2012; 23(4):194-204.
18. Kim KY, Kim HY, Kim JH, Yang Y. Tumor necrosis factor-alpha and interleukin-1beta increases CTRP1
expression in adipose tissue. FEBS Lett. 2006; 580(16): 3953-3960.
19. PanX, Lu T, Wu F, Jin L, Zhang Y, Shi L, et al. Circulating complementC1q TNF-related protein 1 levels
are increased in patients with type 2 diabetes and are associated with insulin sensitivity in Chinese
subjects. PLoS One. 2014; 9 (5) :94478.
20. Xin Y, Lyu X, Wang C, Fu Y, Zhang S, Tian C, et al. Elevated circulating levels of CTRP1, a novel
adipokine, in diabetic patients. Endocr. J.2014; 61 (9) :841–847.
21. Shen Y, Lu Y, Liu ZH, Wu F, Zhu JZ, Shen WF, et al. Increased serum level of CTRP1 is associated with
low coronary collateralization in stable angina patients with chronic total occlusion. Int. J. Cardiol. 2014;

- **348** 174 (1) :203–206.
- 349 22. Jeon JH, Kim KY, Kim JH, Baek A, Yang Y, Lee YH, et al. A novel adipokine CTRP1 stimulates
 350 aldosterone production. FASEB J.2008; 22 (5):1502–1511.
- 351 23. Feng S, Su Y, Luo L, Jing F, Yi Q. Serum levels of C1q/tumor necrosis factor related protein-1 in
 352 children with Kawasaki disease. Pediatr. Res. 2018; 83 (5): 999–1003.
- 353 24. Wong GW, Krawczyk SA, Kitidis-Mitrokostas C, Revett T, Gimeno R, Lodish HF. Molecular,

16

354	biochemical and functional characterizations of C1q/TNF family members: adiposetissue-selective
355	expression patterns, regulation by PPAR-gamma agonist, cysteine-mediated oligomerizations,
356	combinatorial associations and metabolic functions. Biochem. J. 2008; 416:161–177.
357	25. Peterson JM, Aja S, Wei Z, Wong GW. CTRP1 protein enhances fatty acid oxidation via AMP-
358	activated protein kinase (AMPK) activation and acetyl-CoA carboxylase (ACC) inhibition. J. Biol.
359	Chem.2012; 287: 1576–158.
360	26. Yuasa D, Ohashi K, Shibata R, Mizutani N, Kataoka Y, Kambara T, et al. C1q/TNF-related protein-1
361	functions to protect against acute ischemic injury in the heart. FASEB J. 2016;30 (3):1065–1075.
362	27. Li SP, Xie Y, Xu F, Liang Y, He BM, Sun HB. The expression of C1q/TNF-related protein-1 in patients
363	with acute ischemic stroke and its predictive value for the severity of neurological deficits. The Journal
364	of Practical Medicine. 2018; 34(10).
365	28. Xing S, Zhang Y, Li J, Zhang J, Li Y, Dang C, et al. Beclin 1 knockdown inhibits autophagic activation
366	and prevents the secondary neurodegenerative damage in the ipsilateral thalamus following focal
367	cerebral infarction. Autophagy. 2012; 8:63-76.
368	29. Shinojima N, Yokoyama T, Kondo Y, Kondo S. Roles of the Akt/mTOR/p70S6K and ERK1/2 signaling
369	pathways in curcumin-induced autophagy. Autophagy. 2007; 3 (6):635–637.
370	30. Zhang R, Zhu Y, Dong X, Liu B, Zhang N, Wang X, et al. Celastrol Attenuates Cadmium-Induced
371	Neuronal Apoptosis via Inhibiting Ca 2+ -CaMKII-Dependent Akt/mTOR Pathway: Celastrol attenuates
372	Cd neurotoxicity via Ca 2+ signaling. J Cell Physiol. 2017; 232(8):2145-2157.
373	31. Hou Y, Wang K, Wan W, Cheng Y, Pu X, Ye X. Resveratrol provides neuroprotection by regulating the
374	JAK2/STAT3/PI3K/AKT/mTOR pathway after stroke in rats. Genes Dis. 2018; 5 (3):245–255.
375	32. Chen H, Gao L, Huang Z, Liu Y, Guo S, Xing J, et al. C1qTNF-related protein 1 attenuates doxorubicin-
376	induced cardiac injury via activation of AKT. Life Sci. 2018; 207:492–498.

377

378 Supporting information

379 S1 Dataset. Raw data.

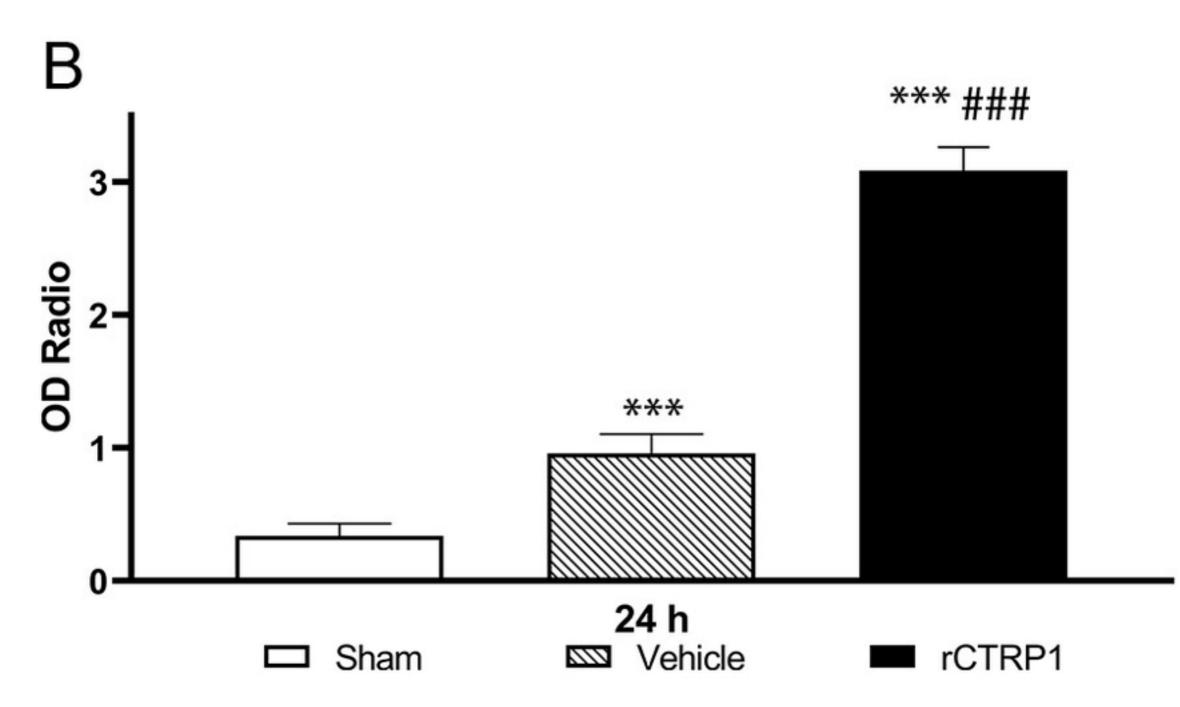


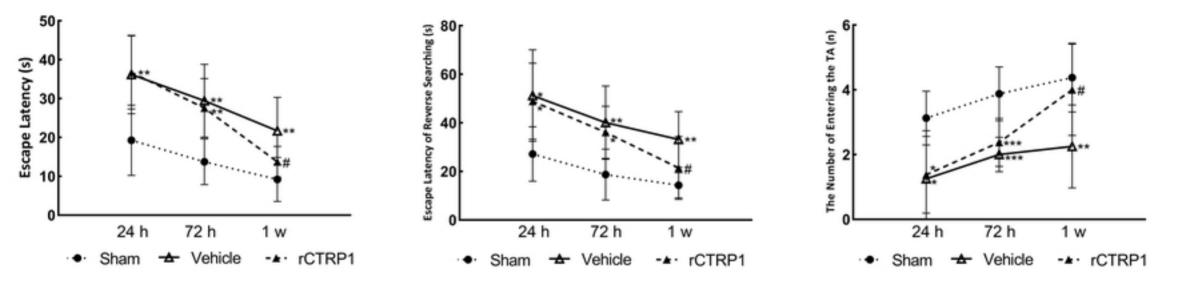
Sham Vehicle

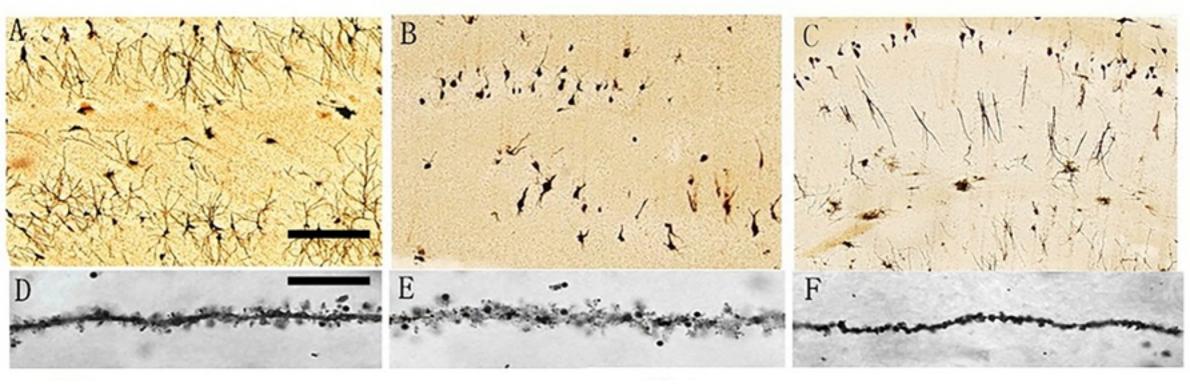


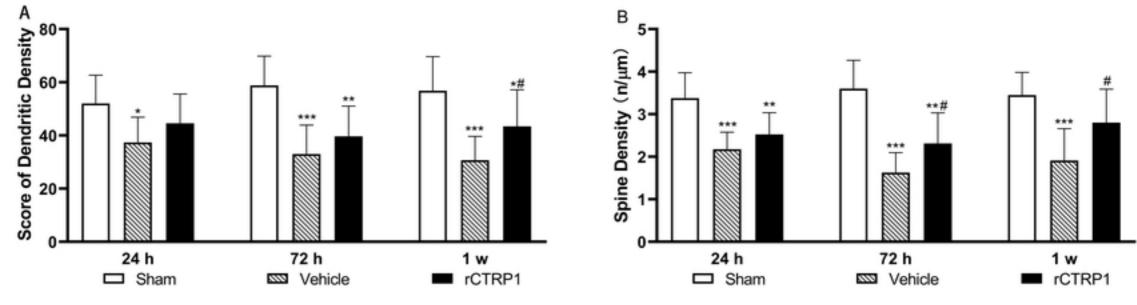
TRP1 bioRxiv preprint doi: https://doi.org/10.1101/2020.12.29.424653; this version posted December 29, 2020. The copyright holder for this preprint (which was not certified by peer review) is the author/funder, who has granted bioRxiv a license to display the preprint in perpetuity. It is made available under aCC-BY 4.0 International license.

GAPDH

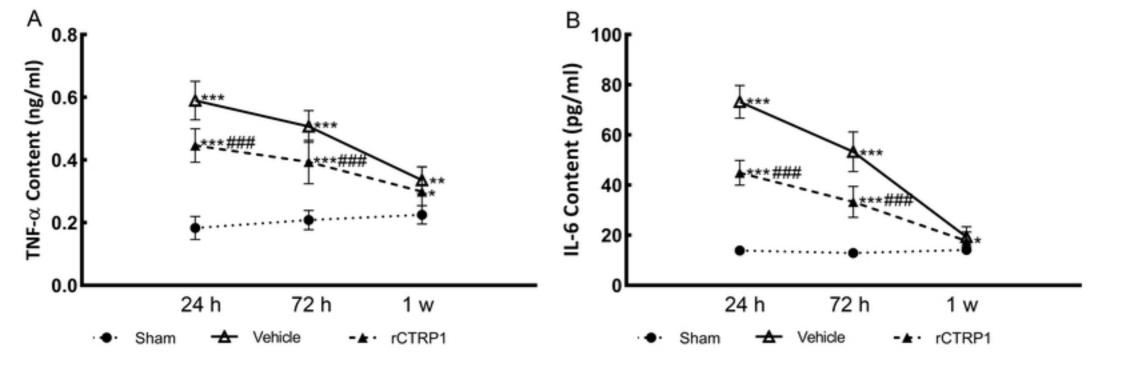








Figure



Figure

