

# Protective effects of andrographolide against cerebral ischemia-reperfusion injury in mice

YAN LI<sup>1,2</sup>, LI-LING XIANG<sup>2</sup>, JIN-XIN MIAO<sup>2</sup>, MING-SAN MIAO<sup>2</sup> and CAN WANG<sup>2</sup>

<sup>1</sup>College of Traditional Chinese Medicine; <sup>2</sup>School of Pharmacy, Henan University of Chinese Medicine, Zhengzhou, Henan 450046, P.R. China

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Abstract. Ischemic stroke is one of the most common causes of mortality worldwide and is a primary cause of disability and mortality in adults. There is an unmet need for drugs that can effectively treat ischemic stroke. Hence, the present study explored the neuroprotective effects of andrographolide (Andro) in a mouse model of bilateral common carotid artery occlusion, and systematically evaluated the potential mechanisms underlying its effects. The effects of Andro on mouse brain tissue following cerebral ischemia-reperfusion injury (CIRI) were evaluated by histopathological (H&E and Nissl) and immunofluorescence [glial fibrillary acidic protein (GFAP) and neuronal nuclei (NeuN)] staining. A traditional Chinese medicine-based network pharmacology method was performed to establish and analyze compound-target-disease and function-pathway networks in order to elucidate the possible mechanisms responsible for the protective role of Andrographis paniculata in CIRI. In addition, western blot analysis and RT-qPCR was performed to evaluate the expression and activation of

*Correspondence to:* Professor Ming-San Miao or Dr Jin-Xin Miao, School of Pharmacy, Henan University of Chinese Medicine, 156 Jinshui East Road, Zhengzhou, Henan 450046, P.R. China E-mail: miaomingsan@163.com E-mail: jinxin.miao@yahoo.com

Abbreviations: Andro, andrographolide; BDNF, brain-derived neurotrophic factor; CCH, chronic cerebral hypoperfusion; CIRI, cerebral ischemia-reperfusion injury; DAPI, 4',6-diamidino-2-phenylindole; DL, drug-likeness; ELISA, enzyme-linked immunosorbent assay; GFAP, glial fibrillary acidic protein; GO, gene ontology; H&E, hematoxylin and eosin; IL-1β, interleukin 1β; IL-6, interleukin 6; KEGG, Kyoto Encyclopedia of Genes and Genomes; MCAO, middle cerebral artery occlusion; NeuN, neuronal nuclei; OB, oral bioavailability; PI3K, phosphoinositide 3-kinase; PVDF, polyvinylidene fluoride; RT-qPCR, reverse transcription-quantitative polymerase chain reaction; TBST, Tris-buffered saline/Tween-20; TNF-α, tumor necrosis factor-α

*Key words:* andrographolide, cerebral ischemia-reperfusion injury, TCM-based network pharmacology, PI3K/Akt

signaling proteins predicted to be involved in this mechanism. The amelioration of histopathological alterations was observed in mice pre-treated with Andro. Immunofluorescence staining revealed that Andro decreased the expression of GFAP and increased the expression of NeuN, and significantly decreased the levels of pro-inflammatory cytokines (IL-1 $\beta$ , IL-6 and  $TNF-\alpha$ ). Network pharmacology analysis revealed that neuroinflammatory response and apoptosis were associated with the effects of Andrographis paniculata on CIRI. Western blot analysis revealed that the mice pre-treated with Andro exhibited an upregulated protein expression of tropomyosin receptor kinase B (TrkB), p-PI3K and p-Akt, as well as a decrease in the expression of GFAP and an increase in the expression of NeuN. In addition, the data of RT-qPCR indicated that the mice pre-treated with Andro exhibited a significantly decreased expression of encoding IL-1 $\beta$  mRNA, IL-6 mRNA and TNF- $\alpha$ mRNA in the brain compared to the untreated mice following CIRI. On the whole, the findings of the present study suggest that pre-treatment with Andro exerts a protective effect against CIRI, which may be partly related to its potential to reduce neuroinflammatory response and apoptosis in patients with stroke.

## Introduction

Stroke is one of the leading causes of human mortality and long-term disability worldwide (1), and its annual incidence is gradually increasing. Approximately 80% of stroke cases involve ischemic stroke, which affects the daily life and work of patients, and places a heavy burden on family and society (2). Brain ischemia is caused by blood vessel obstruction (insufficiency) and brain damage is caused by vascular occlusion (3). The greater the decrease in cerebral blood flow, the more severe the brain damage (4). The pathophysiological mechanisms are complex and include inflammation, apoptosis, excitotoxicity and periinfarct depolarization (5). The reduction of inflammation and elimination of inflammatory genes exerts neuroprotective effects against brain damage following ischemic stroke (6). In addition, cerebral ischemia can affect the innate and adaptive immune systems, and can induce neuronal apoptosis (7). Although drugs have been developed to target inflammation, oxidative stress, apoptosis and other pathophysiological mechanisms, there remains a lack of effective neuroprotectors to prevent cerebral ischemia (8).

Thrombolytic therapy is one of the strategies recommended for the treatment of brain ischemia; however, its use is limited by a very narrow time window and adverse effects (9). Recombinant tissue plasminogen activator, a thrombolytic agent, is the only approved thrombolytic therapy, and it must be administered within a limited time frame in order for a clinical benefit to be achieved (10). Therefore, the identification of novel treatment strategies for stroke is essential. Notably, traditional Chinese medicine (TCM) has a long history in the treatment of ischemic stroke.

Chuanxinlian is the dry, aerial part of Andrographis paniculata, and one of its main components is andrographolide (Andro). In a previous study, a rat model of permanent middle cerebral artery occlusion (pMCAO) was established to investigate the potential therapeutic effects of Andro in cerebral ischemia; it was found that after Andro was administered intraperitoneally to rats with pMCAO, the neurological behavior scores and infarction area were decreased (11). Andro may exert therapeutic effects against hippocampal neuron injury and cognitive dysfunction induced by chronic cerebral hypoperfusion (CCH), and it also exerts a potential neuroprotective effect in cases of hippocampal neuronal damage and cognitive impairment from CCH due to the suppression of astrocyte activation (12). The present study aimed to further explore the effects of Andro on cerebral ischemia-reperfusion through both network analysis and in vivo experiments.

Network pharmacology is a novel drug research method that combines systems biology, multi-direction pharmacology, network analysis and computer technology. The association between drugs and disease has been investigated by combining drug component targets and disease targets from multiple perspectives (13,14). In the present study, the active components and disease targets of Chuanxinlian against ischemic stroke were summarized using the database of network pharmacology, and the association between the active components of Chuanxinlian, targets and pathways was explored through multi-directional systems biology. A TCM-based network pharmacology method was used to establish and analyze compound-target-disease and function-pathway networks in order to elucidate the possible mechanisms through which Andrographis paniculata exerts its protective effects against cerebral ischemia-reperfusion injury (CIRI). The classical pharmacology of Andro was also studied in vivo to verify the results of network pharmacology. Our study aims to provide a scientific basis for the clinical application of Andro in stroke. In addition, this study provides theoretical support for subsequent experimental studies.

# Materials and methods

Animals. A total of 90 male 6-week-old specific pathogen-free (SPF) Kunming mice weighing 18-22 g were obtained from Jinan Pengyue Experimental Animal Breeding Co., Ltd. (Laboratory animal certificate: SCXK 2014-0007). The animals were housed in cages with an ambient temperature of 23-25°C and a humidity of 35-45%, under a 12-h light/dark cycle beginning at 08:00; standard mouse chow and water were provided *ad libitum*. The present study was performed in accordance with the recommendations of the Ministry of Science and Technology of China's Guidance for the Care and



Figure 1. Chemical structure of andrographolide.

Use of Laboratory Animals. All procedures were approved by the Laboratory Animal Ethics Committee of Henan University of Chinese Medicine (the number of approval ethics code: DWLL201903026).

Drug administration. Using high-performance liquid chromatography, the purity of Andro provided by Shanghai Yuanye Bio-Technology Co., Ltd. (chemical structure presented in Fig. 1) was determined to be 98%. The maximum dose of Andro used in the present study (120 mg/kg; 12 mg/ml) was selected based on a previous study (15) and preliminary experiments (data not shown). A 200 mg/ml solution of Andro dissolved in dimethyl sulfoxide was prepared and diluted in distilled water to the appropriate concentration prior to use. The animals were randomly assigned to 5 groups as follows: The sham-operated group (n=10), repeated CIRI group (n=20) and 3 Andro groups (30, 60 and 120 mg/kg) (n=20). Following randomization, the corresponding drugs were administered orally, with the sham-operated group and CIRI group being treated with an equal volume of normal saline. At 1 h after the final administration on the 14th day, the mice in each group were subjected to repeated CIRI surgery.

Repeated CIRI in mice. Repeated CIRI was performed through bilateral common carotid artery occlusion (BCCAO) according to a previously published method (16). In brief, anesthesia was initialized with 4% isoflurane inhalation and then maintained with 1.5% isoflurane in 70% nitrous oxide  $(N_2O)/30\%$  oxygen  $(O_2)$  using a laboratory anesthesia system (Matrix VIP 3000; Midmark Corp.). The bilateral common carotid arteries (BCCAs) were exposed, the carotid artery was lifted with 4-0 silk sutures, and the BCCAs were ligated with artery clamps for 10 min. The artery clamps were then removed and the BCCAs were visually inspected for reperfusion. After the blood flow had been restored for 10 min, the BCCAs were again ligated for 10 min. The sham-operated mice received the same procedure, but without carotid artery ligation. Wounds were sutured with a 4-0 silk suture. During the surgery, the temperature was maintained at 37±0.5°C. To ameliorate the suffering of animals observed throughout the experimental period, the animals were euthanized with 4% isoflurane inhalation followed by the dislocation of the

Sense primer (5'-3')	Antisense primer (5'-3')		
GTGACGTTGACATCCGTAAAGA	GTAACAGTCCGCCTAGAAGCAC		
GCCCATGAAAGAAGTAAACGTCC	AGTGTCAGCCAGTGATGTCGTC		
TTCTTGGGACTGATGCTGGTG	GCCATTGCACAACTCTTTTCTC		
ACAGGCTCCGAGATGAACAAC	GTGGGTGTGCCGTCTTTCAT		
ACCCTCACACTCACAAACCA	ATAGCAAATCGGCTGACGGT		
	Sense primer (5'-3') GTGACGTTGACATCCGTAAAGA GCCCATGAAAGAAGTAAACGTCC TTCTTGGGACTGATGCTGGTG ACAGGCTCCGAGATGAACAAC ACCCTCACACTCACAAACCA		

Table I. Sequences of primers used for RT-qPCR.

BDNF, brain-derived neurotrophic factor; IL, interleukin; TNF- $\alpha$ , tumor necrosis factor  $\alpha$ .

cervical vertebra. Animal death was confirmed by respiratory and cardiac arrest, and no righting reflex. The brain tissues were removed (n=3), then fixed in 4% paraformaldehyde and embedded in paraffin prior to hematoxylin and eosin (H&E), Nissl and immunofluorescence staining. The other brain tissues of the mice were cryopreserved in liquid nitrogen for western blot analysis, reverse transcription-quantitative polymerase chain reaction (RT-qPCR) and biochemical detection.

*H&E staining*. Mouse brains were excised and fixed overnight in 4% paraformaldehyde at 4°C, embedded in paraffin and serial 5- $\mu$ m-thick sections were prepared. The sections were stained with H&E (C0105, Beyotime Institute of Biotechnology) to assess brain histology. Briefly, the sections were stained with hematoxylin for 5 min, then washed with running water for 5 min and differentiated in 1% acid alcohol for 10 sec at room temperature. The sections were then stained with 1% eosin for 1 min, then washed with running water and dehydrated in a decreasing concentration of alcohol (70, 80, 90 and 100%) and cleared in xylene. Finally, the sections were mounted with neutral resin and observed under a microscope (Olympus, BX63; Olympus Corporation).

*Nissl staining.* The brain tissue sections were stained with Nissl staining solution (C0117, Beyotime Institute of Biotechnology) for 30 min at 37°C. The brain sections were then washed with 95% ethyl alcohol and observed under a microscope (Olympus, BX63; Olympus Corporation). Large cell bodies with abundant cytoplasm and significant levels of Nissl bodies represent normal neurons. Cells with pyknosis or blurring of Nissl bodies represent damaged cells.

*Immunofluorescence staining*. Serial 5-µm-thick paraffinembedded sections were baked at 60°C for 30 min, deparaffinized with xylene and rehydrated. The sections were then submerged into the citrate antigen retrieval buffer, heated in a microwave for antigen retrieval, treated with 3% hydrogen peroxide to quench endogenous peroxidase activity and incubated with 1% bovine serum albumin to block non-specific binding. The sections were incubated with primary antibody [glial fibrillary acidic protein (GFAP), 1:500, GB12096, Wuhan Servicebio Technology Co., Ltd.; neuronal nuclei (NeuN), 1:500, GB11138, Wuhan Servicebio Technology Co., Ltd.] overnight at 4°C. After washing, the tissue sections were treated with fluorescence-conjugated secondary antibody (G1213-100UL, Wuhan Servicebio Technology Co., Ltd.) in the dark (37°C) for 2 h. After washing with PBS, the sections were immediately examined under a fluorescence microscope (Olympus, BX63; Olympus Corporation).

*Cytokine analysis.* The infarct sides of brain tissues following repeated CIRI surgery were used for assessing expression brain-derived neurotrophic factor (BDNF) and pro-inflammatory cytokines. The levels of BDNF and pro-inflammatory cytokines in the supernatant of the brain homogenates or serum were detected by enzyme-linked immunosorbent assay (ELISA).

Western blot analysis. The brain tissues of the mice were removed from the liquid nitrogen and placed in a pre-cooled mortar that had been cleaned and dried at high temperature in advance. The liquid nitrogen was poured into the mortar, and the tissues were ground into a powder in the liquid nitrogen. Approximately 100 mg of brain tissue powder were weighed and placed in a 5-ml EP tube, and 1 ml of RIPA lysis buffer containing 1% PMSF and 1% phosphatase inhibitor was added. Extraction with a syringe was performed until the tissue was completely fragmented completely and in full contact with the lysis buffer. The lysate was then placed on ice for 30 min. Lysates were centrifuged at 12,000 x g for 15 min in a high-speed centrifuge at 4°C, and the supernatant was collected to obtain total protein. The protein concentration was measured using a BCA protein assay kit (Beijing Solarbio Science & Technology Co., Ltd.). The samples were then separated on 10% SDS-PAGE gels and electro-transferred to polyvinylidene fluoride (PVDF) membranes. The membranes were blocked in Tris-buffered saline (TBS) with 5% non-fat milk for 2 h at room temperature, then incubated with primary antibodies [tropomyosin receptor kinase B (TrkB), 1:800, GB11295-1, Wuhan Servicebio Technology Co., Ltd.; GFAP, 1:1,000, GB12096, Wuhan Servicebio Technology Co., Ltd.; NeuN, 1:800, GB11138, Wuhan Servicebio Technology Co., Ltd.; p-PI3K, 1:1,000, 17366, Cell Signaling Technology, Inc.; PI3K, 1:800, 4257, Cell Signaling Technology, Inc.; Akt, 1:800, 4685, Cell Signaling Technology, Inc.; p-Akt, 1:1,000, 4058, Cell Signaling Technology, Inc.] at 4°C overnight. Subsequently, the membranes were washed with Tris-buffered saline/Tween-20 (TBST) and incubated with secondary antibodies, then washed with TBS. The secondary antibody used was IRDye800CW goat anti-rabbit antibody (1:20,000; LI-COR Biosciences). Finally, labeled proteins were detected using a near-infrared imaging system (LI-COR Biosciences) and analyzed using an image analyzer (Image Studio™

gredient code Ingredient name		OB (%)	DL	
MOL000173	Wogonin	30.68	0.23	
MOL002928	Oroxylin A	41.37	0.23	
MOL002932	Panicolin Panicolin		0.29	
MOL008203 Andrographolide		57.06	0.34	
MOL008206 Moslosooflavone		44.09	0.25	
MOL008218	1-Monoolein	34.13	0.3	
MOL008219	3-[2-[(1R,4aS,5R,8aS)-5,8a-dimethyl-2-methylene-5-	51.78	0.28	
	methylol-decalin-1-yl]ethyl]-5H-furan-2-one			
MOL008228	Andrographin	37.57	0.33	
MOL008229	IOL008229 Andrographin F		0.85	
MOL008230	IOL008230 Andrographidine F_qt		0.45	
MOL008232	(3Z,4S)-3-[2-[(1R,4aS,5R,6R,8aS)-6-hydroxy-5,8a-	46.96	0.36	
	dimethyl-2-methylene-5-methylol-decalin-1-yl]ethylidene]-4-			
	hydroxy-tetrahydrofuran-2-one			
MOL008238	3-[2-[(1S,4aR,5S,8aR)-5,8a-dimethyl-2-methylene-5-	63.54	0.28	
	methylol-decalin-1-yl]ethyl]-5H-furan-2-one			
MOL008239	Quercetin tetramethyl(3',4',5,7) ether	31.57	0.41	

Table II. Basic information on ingredients of Chuanxinlian.

OB, oral bioavailability; DL, drug-likeness.



Figure 2. Andrographis paniculata active ingredient/anti-ischemic target network map. The rose-red-colored circles represent the active ingredients of Andrographis paniculata, while the blue-colored squares represent the targets of these active ingredients against cerebral ischemia.





Figure 3. Interaction network of potential targets of *Andrographis paniculata* for the treatment of ischemic stroke. The 'node' represents the target protein, and the larger the degree value, the larger the node, and the larger the degree value corresponding to the color ranged from blue to red. The 'edge' represents the correlation between the targets, and the thickness of the edge is based on the combined score.

Software). The densitometric values were normalized to the corresponding GAPDH densitometry values.

*RT-qPCR*. Total RNA was extracted from the brain tissues using the TRIzol reagent (R0016, Beyotime Institute of Biotechnology). RNA concentration of the samples was determined by Natodrop 2000 (Thermo Fisher Scientific, Inc.). The total RNA reverse transcription was performed using the PrimeScript<sup>®</sup> RT Master Mix Perfect Real-Time kit (RR036A, Takara Bio, Inc.). In the following, cDNA was taken as the template for real-time PCR with TB Green<sup>®</sup> Premix Ex Taq (RR420A, Takara Bio, Inc.). The steps for qPCR included: Initial denaturation by incubating in 95°C for 30 sec; 40 qPCR cycles by 95°C for 5 sec and then 60°C for 30 sec in each cycle. Each sample was determined triply and relative expression levels were analyzed using the  $2^{-\Delta\Delta Cq}$  method (17).  $\beta$ -actin was used for normalization. The sequences of the primers are presented in Table I.

# *Network pharmacology analyses*. Network pharmacology analyses included the following:

i) *Establishment of the drug database*. Traditional Chinese Medicine Systems Pharmacology (TCMSP) was used to screen, predict and collect the active ingredients of Chuanxinlian using oral bioavailability (OB) and drug-likeness (DL) as the limiting conditions.

ii) Targets of CXL active ingredients. The simplified molecular-input line-entry system (SMILES) chemical formulas of all active ingredients of Chuanxinlian obtained in the current screen were downloaded from the PubChem database and the SMILES chemical formulas corresponding to the active ingredients were then entered into the similarity ensemble approach (SEA; http://sea.bkslab.org/) database to obtain therapeutic targets. Finally, the UniProtKB search function in the protein database (http://www.uniprot.org/uniprot/) was applied, and targets and UniProt numbers related to active ingredients were obtained after all retrieved targets were associated to their official symbol by entering the targets name and qualifying the species as human.

*iii) Potential disease targets.* Multiple databases (TTD, Drugbank and DisGeNET) were searched to collect therapeutic targets related to ischemic stroke using the terms 'Cerebral ischemic stroke', 'Cerebral infarction', 'Ischemic stroke' and 'Acute ischemic stroke'. The database of targets related to ischemic stroke was constructed by comparing and de-weighting the 3 databases.

iv) *Protein interaction*. The targets of the active ingredients of Chuanxinlian were intersected with the targets implicated in ischemic stroke to obtain the targets of Chuanxinlian potentially involved in treating ischemic stroke, referred to as Chuanxinlian-target-cerebral ischemia. The String database

No.	UniProt ID	Gene name	Protein name	Degree	Betweenness
1	O14920	IKBKB	Inhibitor of nuclear factor kappa-B kinase subunit beta	18	0.00467253
2	P00734	F2	Coagulation factor II, thrombin	11	0.00838562
3	P00747	PLG	Plasminogen	15	0.05091082
4	P01375	TNF	Tumor necrosis factor	23	0.19812513
5	P03372	ESR1	Estrogen receptor 1	6	0.01264776
6	P05121	SERPINE1	Serpin family E member 1	8	0.00336983
7	P07550	ADRB2	Adrenoceptor beta 2	7	0.01043808
8	P08069	IGF1R	Insulin-like growth factor 1 receptor	5	0.00441784
9	P08172	CHRM2	Muscarinic acetylcholine receptor M2	6	0.00606293
10	P08253	MMP2	Matrix metallopeptidase 2	7	0.01000518
11	P11712	CYP2C9	Cytochrome P450 family 2 subfamily C member 9	7	0.04508112
12	P13726	F3	Coagulation factor III, tissue factor	5	0.00583017
13	P14780	MMP9	Matrix metallopeptidase 9	9	0.04962157
14	P25116	F2R	Coagulation factor II thrombin receptor	11	0.00838562
15	P28482	MAPK1	Mitogen-activated protein kinase 1	19	0.21744713
16	P30556	AGTR1	Angiotensin II receptor type 1	16	0.0567444
17	P33261	CYP2C19	Cytochrome P450 family 2 subfamily C member 19	6	0.0237056
18	P35354	PTGS2	Prostaglandin-endoperoxide synthase 2	8	0.12466034
19	P37231	PPARG	Peroxisome proliferator-activated receptor gamma	5	0.04858176
20	P42574	CASP3	Caspase-3	9	0.00262056
21	Q07869	PPARA	Peroxisome proliferator-activated receptor alpha	7	0.02624386
22	Q9HBW0	LPAR2	Lysophosphatidic acid receptor 2	13	0.01747176

Table III. Information on potential targets and topological properties of Chuanxinlian against cerebral ischemia.

(https://string-db.org/) (17), a database containing known and predicted protein-protein interactions, was used.

The protein targets of Chuanxinlian were imported into the String database to obtain their interaction associations, qualifying the species as human. The results were saved in TSV format, and the nodes (node 1, node 2) and combined score information in the file were reserved, then imported into Cytoscape 3.2.1 to draw the interaction network, and the network results were analyzed and saved. Using the 'generate style from statistics' tool in Cytoscape, the node size and color were set to reflect the number of components connected to the target (degree), and the thickness of edge setting was set to reflect the size of the combined core. Finally, the ultimate protein interaction network was obtained.

v) Construction of active ingredient-target network diagram. The active ingredients and targets of Cuanxinlian were imported into Cytoscape 3.2.1 software and the ingredient-target network diagram of Chuanxinlian was constructed to predict the main targets and active ingredients of Chuanxinlian involved in treating cerebral ischemia.

vi) Analysis of biological processes and pathways. The targets of Chuanxinlian were input into the DAVID database (18), with the selected Identifier set to the OFFICIAL GENE SYMBOL, List Type set to the GENE List, and species limited to human. Gene ontology (GO) functional enrichment analysis and Kyoto Encyclopedia of Genes and Genomes (KEGG) pathway enrichment analysis were carried out for the targets of Andro. Its mechanism was investigated and the results were saved. The top biological processes and signaling pathways were obtained by setting P<0.05.

Statistical analysis. All data are expressed as the means  $\pm$  standard deviation and analyzed using SPSS 17.0 (SPSS, Inc.). One-way analysis of variance and the Bonferroni post hoc test were used to evaluate statistical significance. Results were considered statistically significant when P<0.05.

#### Results

Components of Chuanxinlian and their corresponding target proteins. Through the TCMSP database and literature investigation, 49 chemical components of Chuanxinlian were obtained. Using OB  $\geq$  30% and DL  $\geq$  0.18 as the filter criteria, 13 active ingredients were found to meet the requirements. Basic information on the ingredients is presented in Table II.

*Target prediction*. All targets were obtained from the TCMSP, SEA and SIA databases, and the duplicates and false positives were removed. Subsequently, 428 targets of Chuanxinlian were integrated; 421 ischemic stroke symptom targets were obtained from the DisGeNET, Drugbank and TTD databases; in addition, 68 potential targets were obtained from the intersection of Chuanxinlian targets and ischemic stroke targets (Table II).

Active ingredient-target network. Cytoscape 3.2.1 was used to construct the Andrographis paniculata active





Figure 4. Interaction network of potential targets of Chuanxinlian for the treatment of ischemic stroke. (A) Enriched gene ontology terms for biological processes, (B) molecular functions, (C) cell components, and (D) KEGG pathway enrichment analysis of potential targets of the main active ingredients of Chuanxinlian.

ingredient-anti-ischemic target map (Fig. 2), which contained 81 nodes and 249 edges. The rose-red-colored circles represent the active ingredients of Chuanxinlian, while the blue-colored

squares represent the targets of these active ingredients against cerebral ischemia, and the edge represents the interaction between the active ingredient and anti-cerebral ischemia targets.



Figure 5. Effect of andrographolide pre-treatment on pathological changes of the (A) brain cortex and (B) hippocampal CA1 region following CIRI (H&E staining; magnification, x200; scale bar, 50  $\mu$ m; n=3). Pre-treatment with 30, 60 and 120 mg/kg andrographolide reduced the histopathological alteration following CIRI to a certain degree. Sham, sham-operated group; CIRI, cerebral ischemia-reperfusion injury; Andro, andrographolide.



Figure 6. Effect of andrographolide pre-treatment on changes in Nissl staining in the brain cortex following CIRI (Nissl staining; magnification, x200; scale bar, 50  $\mu$ m; n=3). Sham, sham-operated group; CIRI, cerebral ischemia-reperfusion injury; Andro, andrographolide.

*Protein interaction network diagram.* Target interactions were obtained from the String database, using Cytoscape 3.2.1 to draw the network diagram (Fig. 3). The 'edge' represents the correlation between the targets, and the thickness of the edge corresponds to the combined score: The thicker the edge, the greater the value of the combined score and the greater the degree of association. The node represents the target, and the value of 'degree' represents the effect intensity. The larger the degree value, the larger the node, and the larger the degree value corresponding to the color ranged from blue to red. There were 98 nodes and 1,099 edges in the figure. The average node degree was 20.4 and the average local clustering coefficient was 0.624. Finally, nodes with degree ≥5 and betweenness ≥0.00262056 were selected as the main nodes; thus, there were a total of 22 main nodes, as listed in Table III.

# *Gene function and pathway analysis.* Gene function and pathway analysis revealed the following results:

i) GO enrichment analysis. The GO enrichment function and the KEGG pathway in DAVID were used to analyze the 22 selected targets. GO enrichment analysis includes 3 branches: Biological processes, molecular function and cellular component. At the P<0.05 setting, 298 biological processes or pathways were obtained. Biological processes or pathways with a higher P-value were screened, and GraphPad Prism7.0 was used for mapping.

As shown in Fig. 4A, the main biological processes of the targets were the regulation of response to an external stimulus,

regulation of apoptosis and the regulation of programmed cell death. As shown in Fig. 4B, the main molecular functions of the targets were drug binding, endopeptidase activity, lipid binding and peptidase activity. Plasminogen (PLG) is a single-chain glycoprotein that plays an important role in thrombolytic therapy for acute cardiogenic cerebral infarction. When stimulated, vascular endothelial cells secrete a large amount of tissue-type plasminogen activator, which converts PLG into fibrinase and this activates the fibrinolytic system *in vivo*. Some autosomal genes associated with ischemic heart disease, including the gene encoding PLG, have been identified (19). PLG is also a key molecule involved in the thrombolytic treatment of cerebral ischemia. As shown in Fig. 4C, the cell components of the targets included cell fraction, plasma membrane and neuron projection.

ii) *Pathway analysis*. The results of KEGG pathway enrichment analysis are shown in Fig. 4D. The targets identified were associated with the NOD-like receptor, mitogen-activated protein kinase (MAPK) and phosphoinositide 3-kinase (PI3K)/Akt and Toll-like receptor signaling pathways. The results revealed that the main active ingredients of Chuanxinlian were largely distributed into different metabolic pathways. Thus, the mechanisms through which Chuanxinlian protects against damage following ischemic stroke may involve multiple ingredients, multiple targets and multiple pathways.

Andro attenuates the pathological abnormalities caused by CIRI as shown by H&E staining. The brain sections of the







Figure 7. Influence of Andro on protein expression levels of (A) GFAP and (B) NeuN in brains of mice subjected to CIRI based on immunofluorescence staining. DAPI was used to stain the nuclei blue, GFAP-positive cells are indicated in red, and NeuN-positive cells are indicated in green. (C) Quantification of GFAP-positive cells. (D) Quantification of NeuN-positive cells. Data are presented as the means  $\pm$  standard deviation (n=3). <sup>#</sup>P<0.01 vs. sham group; <sup>\*\*</sup>P<0.01 vs. CIRI group. Scale bar, 50  $\mu$ m. GFAP, glial fibrillary acidic protein; NeuN, neuronal nuclei; sham, sham-operated group; CIRI, cerebral ischemia-reperfusion injury; Andro, andrographolide.

mice were stained with H&E (Fig. 5), and the results revealed that neurons in the cerebral cortex and hippocampal CA1 region in the sham group were normal. In these mice, brain tissue was not damaged and no obvious vacuolar space was observed. The nerve cells were densely and evenly arranged, with normal structure and morphology, clear cell contour, complete normal morphology and clearly visible nucleoli. By contrast, in the CIRI group, ischemic necrosis occurred in the cerebral cortex, with the shrinkage of neuronal nuclei; some cells exhibited spot-like necrosis, with evident infiltration of microglial cells, neuronal disorder and tissue edema in the hippocampus.

Andro increases Nissl positive staining. The mouse brain sections were stained with Nissl (Fig. 6), and the results revealed that, compared with the sham group, the numbers of Nissl-positive cells in the CIRI group were markedly decreased. Following pre-treatment with Andro, the numbers of Nissl-positive cells were significantly increased compared to the CIRI group.

Andro decreases GFAP and increases NeuN expression, as shown by immunofluorescence staining. The results of immunofluorescence staining illustrated that, compared with the sham group, the expression of GFAP was significantly increased (Fig. 7A and C) and the expression of NeuN was significantly decreased (Fig. 7B and D) in the brains of the mice subjected to CIRI. Following pre-treatment with Andro, the expression of GFAP significantly decreased, while the expression of NeuN significantly increased.

Andro increases the levels of BDNF. BDNF is involved in the development of the nervous system, and CIRI promotes the release of BDNF. Compared with the sham group, the BDNF



Figure 8. Influence of Andro on BDNF, TrkB, GFAP, NeuN, p-PI3K and p-Akt in mouse brains following CIRI was examined. Mice were treated with Andro (30, 60, or 120 mg/kg) for 14 days prior to CIRI, and repeated CIRI was then performed. BDNF was detected by (A) enzyme-linked immunosorbent assay and (B) RT-qPCR. (C-I) Protein levels of TrkB, p-PI3K, PI3K, p-Akt, Akt, GFAP and NeuN in brains of CIRI mice were assessed by western blot analysis. <sup>#</sup>P<0.05, <sup>##</sup>P<0.01 vs. sham group; <sup>\*</sup>P<0.05, <sup>\*\*</sup>P<0.01 vs. CIRI group BDNF, brain-derived neurotrophic factor; TrkB, tropomyosin receptor kinase B; GFAP, glial fibrillary acidic protein; NeuN, neuronal nuclei; p-PI3K, phosphorylated phosphoinositide 3-kinase; Sham, sham-operated group; CIRI, cerebral ischemia-reperfusion injury; Andro, andrographolide.

content in the serum of mice in the CIRI group was significantly increased (P<0.05), and the *BDNF* mRNA level in the brain tissues also exhibited an increasing tendency (Fig. 8A and B). This indicates that cerebral ischemia-reperfusion can stimulate the release of BDNF in a short period of time. Compared with the CIRI group, the serum BDNF levels were significantly increased in mice pre-treated with Andro (P<0.05), and the *BDNF* mRNA levels in the brain tissues of mice receiving CIRI also exhibited an increasing trend. In addition, Andro pre-treatment enhanced the transcription of BDNF in mice subjected to CIRI.

Andro increases the levels of TrkB, and downstream proteins. TrkB is a primary receptor of BDNF. Compared with the sham group, TrkB protein expression in the brain tissues of mice in the CIRI group was significantly decreased (P<0.01), and compared with the CIRI group, the mice that received Andro (120 mg/kg) pre-administration exhibited a significantly upregulated TrkB protein expression in the brain tissue (P<0.01; Fig. 8C and D). The levels of proteins downstream of TrkB [including phosphorylated PI3K (p-PI3K) and phosphorylated Akt (p-Akt) were reduced in the mice in the CIRI group, while Andro pre-treatment markedly increased the phosphorylation levels of these proteins compared to the CIRI group (Fig. 8C, E and F).

GFAP is a specific marker of activated astrocytes, and NeuN is a specific marker of mature neurons. CIRI increased the expression of GFAP and inhibited the expression of NeuN, while Andro administration markedly reduced the expression of GFAP and enhanced the expression of NeuN (Fig. 8G-I).

Andro reduces the levels of pro-inflammatory cytokines. ELISA was used to determine the effects of Andro on the levels of the inflammatory cytokines,  $IL-1\beta$ , IL-6 and  $TNF-\alpha$ in serum. The serum levels of  $IL-1\beta$  and  $TNF-\alpha$  were significantly increased in the CIRI group compared to the sham group. However, compared with the CIRI group, the serum levels of  $IL-1\beta$ , IL-6 and  $TNF-\alpha$  were markedly decreased in the Andro groups (Fig. 9A-C).



Figure 9. Influence of Andro on inflammatory cytokine levels in mouse serum and brain following CIRI was examined. Enzyme-linked immunosorbent assay was used to examine serum levels of inflammatory cytokines (A) IL-1 $\beta$ , (B) IL-6, and (C) TNF- $\alpha$  in mice subjected to CIRI with or without Andro pre-treatment. (D-F) RT-q-PCR was used to measure the mRNA expression levels of (D) IL-1 $\beta$ , (E) IL-6, and (F) TNF- $\alpha$  in the brain tissues of mice subjected to CIRI. Data are presented as the means  $\pm$  standard deviation (n=3). <sup>#</sup>P<0.01 vs. sham group; <sup>\*\*</sup>P<0.01 vs. CIRI group. IL, interleukin; TNF- $\alpha$ , tumor necrosis factor  $\alpha$ ; Sham, sham-operated group; CIRI, cerebral ischemia-reperfusion injury; Andro, andrographolide.

In addition, the *IL-1* $\beta$  mRNA, *IL-6* mRNA and *TNF-a* mRNA levels in brain samples were analyzed by RT-qPCR (Fig. 9D-F). Compared with the sham group, the *IL-1* $\beta$  mRNA, *IL-6* mRNA and *TNF-a* mRNA levels were significantly increased in the CIRI group. In addition, pre-treatment with Andro significantly downregulated the *IL-1* $\beta$  mRNA, *IL-6* mRNA and *TNF-a* mRNA levels (Fig. 9D-F).

## Discussion

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Stroke is the third-leading cause of mortality in several developed countries and is associated with high morbidity, disability and mortality (20). Inflammation and apoptosis, as the critical factors involved in CIRI, can interact with each other and further create a destructive cascade effect (21). When brain tissue is damaged, the levels of pro-inflammatory cytokines (*IL-1β*, *IL-6* and *TNF-α*) increase. Due to the high incidence and severe consequences of ischemic stroke, new drugs are urgently required for the treatment of ischemic stroke (22).

The present study mainly aimed to predict and analyze the target and pathway of *Andrographis paniculata* through network pharmacology. From previous studies (6,11,12), Andro has been demonstrated to possess potent neuroprotective action in the treatment of neurological injury and cognitive dysfunction. Andro is one of the main components of *Andrographis paniculata*. Therefore, Andro was used to further verify the results, and other components could also be selected for subsequent experiments.

Andrographis paniculata, an agent used in TCM, has the effect of clearing heat and removing toxicity, and clearing heat

and removing toxicity is one of the 8 treatment methods of TCM. Andro is a terpene that has been used extensively in China for the treatment of a range of diseases. Previous studies have illustrated that Andro significantly inhibits free radical formation, blood-brain barrier destruction and cerebral infarction in rats subjected to MCAO, and the heme oxygenase 1 (HO-1) inhibitor zinc protoporphyrin IX reverses these effects, indicating that Andro increases Nrf2-HO-1 expression through the regulation of p38 MAPK, confirming that it protects against MCAO-induced brain injury (23). Moreover, Andro reduces the expression of NADPH oxidase 2 (NOX2) and inducible nitric oxide synthase (iNOS), possibly by impairing PI3K/Akt-dependent NF-kB and hypoxia-inducible factor (HIF)-1a activation. This affects microglial activation, which in turn mediates the protective effects of Andro in mice subjected to CIRI (24).

The present study aimed to explore the neuroprotective effects of Andro on CIRI. The data demonstrated that the administration of Andro attenuated pathological changes and increased the number of Nissl-positive cells. Moreover, immunofluorescence staining confirmed that Andro effectively reduced the expression of GFAP and increased the expression of NeuN in mice following CIRI. Nissl bodies, which exist in the cytoplasm of neurons, are characteristic structures of neurons, and their role is to synthesize structural proteins, enzymes, and nerve modulators for neurons. When neurons undergo pathological changes, Nissl bodies begin to dissolve and become dust-like particles that spread out from the nucleus until they disappear (25). NeuN is a specific marker of mature neuronsand is expressed only after the neuron matures, with its expression increasing gradually. Therefore, the expression of NeuN can reflect the maturity of neurons (26). GFAP is a specific marker of activated astrocytes (27). Astrocytes, which are a substantial component of nerve cells, are less sensitive to ischemic hypoxia (28). They are activated when cerebral ischemia and other injuries occur (28).

The results of the present study illustrate that Andro protects the brain from CIRI through a reduction in the levels of apoptosis-related proteins by activating the BDNF-TrkB/PI3K/Akt signaling pathway. It has been demonstrated that BDNF promotes post-CIRI neurological recovery, prevents neuron death and supports the survival of multiple neurons (29). BDNF is mainly synthesized and secreted by presynaptic and postsynaptic neurons and may mediate axonal growth and brain plasticity (30). BDNF activates downstream signaling pathways such as the PI3K/Akt pathway through binding to its cell surface receptor TrkB after being secreted into the extracellular space (31). It has been confirmed that the PI3K/Akt signaling pathway, a significant cell signal transduction pathway, plays an essential role in regulating cell survival, apoptosis, necrosis and other pathophysiological processes (32). The suppression of Akt activity has been associated with neuronal cell death following I/R injury (33), and a previous study demonstrated that Akt was activated by Ser473 phosphorylation (34). The results of the present study demonstrated that following CIRI, the phosphorylation of both PI3K and Akt was increased, indicating that this signal transduction pathway was activated.

The findings of the present study provide novel insight into a potential method of protection against cerebral ischemia, as well as the mechanism through which Andro exerts its neuroprotective effects. Multiple factors and pathways may be responsible for the therapeutic effects of Andro, including inhibiting inflammation and mediating cell apoptosis. Future studies are required to focus on other pathways that are likely involved in this process. It is difficult to identify the true target proteins of Andro and to measure binding affinity. Future studies may focus on verifying the effects reported herein, particularly through the use of knockout mice. Moreover, proteomics and transcriptomics techniques can also be used to find differential targets. The PI3K/Akt pathway is an important neuroprotective and anti-apoptotic pathway. These findings provide novel insight into a potential method of protection against cerebral infarction, as well as the mechanisms through which Andro exerts its neuroprotective effects.

In the present study, the protective effects of Andro were assessed in a mouse model of CIRI, and as shown in the pathological section, the CIRI of neurons was significantly attenuated. Pre-treatment with Andro upregulated the protein expression of TrkB, p-PI3K and p-Akt. It also decreased the expression of GFAP and increased the expression of NeuN. Furthermore, Andro attenuated inflammation significantly by reducing the serum levels of IL- $I\beta$ , IL-6 and TNF- $\alpha$ .

In conclusion, the present study demonstrates that Andro can protect brain tissue from ischemia-reperfusion injury in mice by reducing the inflammatory response and apoptosis of brain tissue cells. It was found that Andro exerts this neuroprotective effect by regulating the expression of important proteins in the PI3K/Akt pathway, indicating that this signaling pathway may be the mechanism behind this protective effect. The results of the present study suggest that Andro may be an effective agent for the treatment of CIRI. Moreover, the findings presented herein provide a potential explanation for the clinical anti-apoptotic and neuroprotective effects of Andro.

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#### Availability of data and materials

All data generated or analyzed during this study are included in this published article or are available from the corresponding author on reasonable request.

#### Authors' contributions

YL, LLX, JXM, CW and MSM conceived and designed the all experiments. YL, LLX and JXM performed the experiments. YL, LLX and CW analyzed the data. YL wrote and edited the manuscript. JXM and MSM revised the manuscript. YL and LLX confirm the authenticity of all the raw data. All authors have read and approved the final manuscript.

#### Ethics approval and consent to participate

The present study was performed in accordance with the recommendations of the Ministry of Science and Technology of China's Guidance for the Care and Use of Laboratory Animals. All procedures were approved by the Laboratory Animal Ethics Committee of Henan University of Chinese Medicine. (the number of approval ethics code: DWLL201903026).

## Patient consent for publication

Not applicable.

# **Competing interests**

The authors declare that they have no competing interests.

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