Introduction

Stroke is widely recognized as a major disease that leads to death and disability in patients globally. Each year, approximately 2.4 million people suffer a stroke episode, with an estimated 1.1 million fatalities, consequently imposing a heavy burden on patients’ families and society. There are three primary types of strokes: ischemic, hemorrhagic, and transient. Ischemic stroke occurs when blood flow to the brain is obstructed by thrombosis or embolism and accounts for 87% of all strokes. Timely restoration of blood supply to the ischemic cerebral tissue by thrombolysis or mechanical recanalization of blood vessels is currently the optimal treatment for improving the prognosis of ischemic stroke. However, in several cases, the restoration of blood supply to ischemic tissue may exacerbate the brain injury, a condition known as “ischemia-reperfusion” (I/R) injury. Therefore, identifying an effective method to reduce the incidence of I/R injury is essential for treating ischemic stroke.

Background: Ferroptosis plays a crucial role in the occurrence and development of cerebral ischemia-reperfusion (I/R) injury and is regulated by mitogen-activated protein kinase 1/2 (ERK1/2). In China, Naodesheng Pills (NDSP) are prescribed to prevent and treat cerebrovascular disease and stroke. However, the protective effects and mechanism of action of NDSP against cerebral I/R-induced ferroptosis remain unclear. We investigated whether NDSP exerts its protective effects against I/R injury by regulating ferroptosis and aimed to elucidate the underlying mechanisms.

Methods: The efficacy of NDSP was evaluated using a Sprague-Dawley rat model of middle cerebral artery occlusion and an in vitro oxygen-glucose deprivation/reoxygenation (OGD/R) model. Brain injury was assessed using 2,3,5-triphenyltetrazolium chloride (TTC), hematoxylin and eosin staining, Nissl staining, and neurological scoring. Western blotting was performed to determine the expression levels of glutathione peroxidase 4 (GPX4), divalent metal-ion transporter-1 (DMT1), solute carrier family 7 member 11 (SLC7A11), and transferrin receptor 1 (TFR1). Iron levels, oxidative stress, and mitochondrial morphology were also evaluated. Network pharmacology was used to assess the associated mechanisms.

Results: NDSP (1.08 g/kg) significantly improved cerebral infarct area, cerebral water content, neurological scores, and cerebral tissue damage. Furthermore, NDSP inhibited I/R- and OGD/R-induced ferroptosis, as evidenced by the increased protein expression of GPX4 and SLC7A11, suppression of TFR1 and DMT1, and an overall reduction in oxidative stress and Fe levels. The protective effects of NDSP in vitro were abolished by the GPX4 inhibitor RSL3. Network pharmacology analysis revealed that ERK1/2 was the core target gene and that NDSP reduced the amount of phosphorylated ERK1/2.

Conclusion: NDSP exerts its protective effects against I/R by inhibiting cerebral I/R-induced ferroptosis, and this mechanism is associated with the regulation of ferroptosis via the ERK1/2 signaling pathway.

Keywords: Naodesheng pills, ferroptosis, cerebral ischemia/reperfusion, glutathione peroxidase 4, mitogen-activated protein kinase 1/2

Naodesheng Pills Ameliorate Cerebral Ischemia Reperfusion-Induced Ferroptosis via Inhibition of the ERK1/2 Signaling Pathway

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Ferroptosis is a distinct form of programmed cell death that differs from apoptosis, necrosis, and autophagy. It is characterized by the iron-dependent accumulation of lipid peroxides and is associated with various neurological pathologies, including cerebral I/R injury, Alzheimer’s disease, and Parkinson’s disease. Ferroptosis is typically regulated by the solute carrier family 7 member 11 (SLC7A11)/glutathione peroxidase 4 (GPX4) pathway, iron metabolism (transferrin receptor [TFR1]/divalent metal ion transporter-1 [DMT1] pathway), and lipid peroxidation. Iron chelation or restoration of GPX4 expression has been shown to attenuate I/R-induced brain injury.

The mitogen-activated protein kinase (MAPK)/extracellular signal-regulated kinase (ERK1/2) was reported to be activated and involved in cerebral I/R injury, and the inhibition of the ERK1/2 pathway reduces middle cerebral artery occlusion-induced infarct size. ERK1/2 plays a crucial role in modulating ferroptosis and the inhibition of ERK1/2 signaling has been reported to be essential for preventing ferroptosis. Therefore, inhibiting ERK1/2-mediated ferroptosis may be an effective strategy for preventing or treating cerebral I/R injury.

Naadesheng pills (NDSP) are commonly prescribed traditional Chinese medicines consisting of Panax notoginseng (Burk.) F. H. Chen, Carthamus tinctorius L., Crataegus pinnatifida Bunge and Pueraria lobata (Wild.). Ohwi, and Ligusticum chuanxiong Hort. The major bioactive components of NDSP have been reported to include flavones (puerarin and daidzein), saponins (ginenosides Rg1, Rg2, Rb1, Rd, Rh1, and notoginsenoside R1), and phenylpropanoids (Safflower yellow A and ferulic acid). In clinical practice, NDSP has been used to treat ischemic stroke and cerebral hemorrhage. Animal experiments have shown that naadesheng (NDS) improves learning and memory function by inhibiting NF-κB and activation of cAMP/PKA/CREB signaling pathway in amyloid β-treated rats. The main components of NDSP, such as tetramethyl pyrazine, hydroxysafflor yellow A, and puerarin, could promote the recovery of neurological function after stroke. However, the therapeutic effects of NDS on ischemic brain injury have been inconsistent. In the FeCl₃-induced focal cerebral ischemic model, NDS improved the neurological deficit score and cerebral infarction. Li et al found that, after 26h of administration, NDS showed a decreasing trend in infarct size and neurological deficit score in a model of permanent cerebral ischemia. Seven days after administration, NDS significantly improved ischemia-induced nerve damage and superoxide dismutase. Further studies are required to evaluate the effects of NDSP on I/R-induced cerebral injury. Although recent metabolomic investigations have revealed that NDSP affects multiple metabolic pathways, such as energy and amino acid metabolism, oxidative stress, and inflammatory pathways, the therapeutic mechanism remains unclear. As the components of NDSP inhibit ERK1/2 activation and ferroptosis, we hypothesized that NDSP exerts a neuroprotective effect against I/R injury by regulating ferroptosis. Therefore, in this study, we investigated effects of NDSP and its mechanism of action on ferroptosis, both in vitro and in vivo.

Materials and Methods

Materials

NDSP was obtained from Hunan TianJi CaoTang Pharmaceutical Co., Ltd. (Changsha, China) and comprised Panax notoginseng (Burk.) F. H. Chen (11.73%), Carthamus tinctorius L. (13.68%), Crataegus pinnatifida Bunge (23.61%), Pueraria lobata (Wild.) Ohwi (39.25%), and Ligusticum chuanxiong Hort (11.73%). CoCl₂ was purchased from the Aladdin Company, Ltd. (Shanghai, China). Anti-SLC7A11 and iron assay kits were purchased from Abcam (Cambridge, UK). Anti-β-actin antibody was purchased by Boster Biological Technology Co., Ltd. (Wuhan, China). Anti-DMT1 and secondary antibodies were purchased from Boster Biological Technology Co. Ltd. (Wuhan, China). The anti-ERK1/2 antibody were purchased from Proteintech (Wuhan, China), and anti-phosphorylated ERK1-T202/Y204+ERK2-T185/Y187 antibodies were purchased from Abclonal (Wuhan, China). Anti-TFR1 antibodies were obtained from Affinity Biosciences (Jiangsu, China). In contrast, anti-GPX4 antibodies, superoxide dismutase (SOD), glutathione (GSH), hydrogen peroxide (H₂O₂), malondialdehyde (MDA), DCFH-DA, and Hoechst 33342/PI assay kits were purchased from the Beyotime Institute of Biotechnology (Jiangsu, China). The RSL3 was acquired from Selleck Co., Ltd. (Shanghai, China). A hematoxylin and eosin (H&E) staining kit, Nissl staining solution, and triphenyl tetrazolium chloride (TTC) solution (2%) were purchased from Solarbio Co., Ltd. (Beijing, China). Dulbecco’s Modified
Eagle’s medium (DMEM) and fetal bovine serum (FBS) were purchased from HyClone Laboratories, Inc. (Logan, UT, USA).

**Animals**

Male Sprague-Dawley rats aged eight weeks (200–250 g) were purchased from SiLaike JingDa (Changsha, China) and housed in cages with standard food and water provided ad libitum. All experimental protocols were designed according to the Guide for the Care and Use of Laboratory Animals (NIH Publication, 8th edition, 2011) and approved by the Animal Ethics and Welfare Committee of the Second Xiangya Hospital of Central South University (No. 2021955).

**Drug Treatment and Focal Cerebral I/R Rat Model**

The Resource Equation Method was used to estimate the sample size of animal experiments. According to the formula, $n = \frac{10}{k} + 1$ (k = number of groups, n = number of subjects per group), the minimum number of animals in each group is three rats, and the total sample size is at least 18 rats. In order to obtain sufficient experimental data, a total of 48 rats were used for the experiments and randomly divided into six groups of eight individuals each: control, sham, I/R, low-dose NDSP (NDSP-L), high-dose NDSP (NDSP-H), and edaravone group (positive control). The daily dose of NDSP administered to adults weighing 60 kg was 0.54 g/kg. This dose was used as the low-dose group in this experiment, and twice the dose was employed as the high-dose group. The NDSP was dissolved in 0.25% CMC-Na and intragastrically administered at doses of 0.54 g/kg or 1.08 g/kg once daily for seven days. The rats in the I/R, normal control, and sham groups were administered equal volumes of CMC-Na. After seven days of drug treatment, rats in the I/R, NDSP-L, edaravone, and NDSP-H groups were subjected to I/R. A rat cerebral I/R model was established. Briefly, the rats were anesthetized with an intraperitoneal injection of 2% pentobarbital sodium (30 mg/kg), and a midline incision was made in the neck. The left common carotid artery (CCA), external carotid artery (ECA), and internal carotid artery (ICA) were isolated. The CCA and ECA were ligated, and the ICA was temporarily clipped using an artery clamp. A nylon suture was then inserted externally into the ECA. After blocking the blood flow for 2 h, reperfusion was performed for 24 h. Rats in the sham group underwent the same procedure without nylon suture insertion.

**Assessment of Neurological Function**

Neurological deficits were assessed based on a five-point neurological deficit score, as previously described (0 = no observable symptoms of neurological deficits, 1 = failure of fully extending the right forepaw, 2 = circling to the right, 3 = falling to the right, 4 = unable to walk spontaneously, and loss of consciousness). Scoring was performed independently by two researchers who were blinded to the experimental groups.

**Assessment of Cerebral Infarct Area**

After 24 h of I/R, the rats were injected intraperitoneally with 2% sodium pentobarbital (150 mg/kg) and sacrificed by decapitation. Infarct area following cerebral I/R injury in rats was determined by 2,3,5-triphenyltetrazolium chloride (TTC) staining. The brain was removed quickly and cut into five 2-mm slices using a rat brain fixator and blade. The sections were incubated with 2% TTC for 25 min at 37 °C and fixed in 4% formaldehyde solution for 24 h. Next, the slices were captured as a digital image and analyzed using ImageJ software (1.37c). The infarct area ratio (%) was calculated as follows:

\[
\text{Infarct area (\%) = infarct area/total cross sectional area} \times 100\%
\]

**Assessment of Cerebral Water Content**

Following I/R, the brain tissues were weighed before being placed in a constant-temperature drying oven at 110 °C for 48 h. The dry brain tissue was weighed again, and the cerebral water content was calculated as follows:

\[
\text{Cerebral water content (\%) = (wet mass} - \text{dry mass)/wet mass} \times 100\%
\]
Histopathological Evaluation
Rat brain tissue was fixed in 4% paraformaldehyde. Then, the tissue block was dehydrated in 70%, 80%, 90%, and 95% ethanol for 10 min each, embedded in paraffin, and sliced into five μm-thick slices. The slices were deparaffinized and stained with hematoxylin, eosin, or Nissl stain to observe histological changes.

Immunohistochemistry
The sections were then deparaffinized and rehydrated. Following antigen retrieval, the sections were treated with 3% H$_2$O$_2$ to block endogenous peroxidase activity and were subsequently blocked with 3% sheep serum at 37 °C for 1 h. Then, sections were incubated with the primary anti-p-ERK1/2 antibody at 4 °C overnight. After washing with PBS, the sections were incubated with HRP-conjugated secondary antibodies and stained with DAB, followed by hematoxylin counterstaining. The stained sections were observed under a light microscope.

Biochemical Analysis
The levels of MDA, GSH, H$_2$O$_2$, SOD, released lactate dehydrogenase (LDH), and iron in the cerebral tissue or SH-SY5Y cells were determined according to the manufacturer’s instructions.

Western Blot Analysis
Total protein from the brain tissue or SH-SY5Y cells was extracted in RIPA lysis buffer (Beyotime, P0013C), and the protein concentration was quantified using a BCA kit (Beyotime, P0010S). SDS-PAGE (10%) separated forty micrograms of protein from each sample; the samples were subsequently transferred from the gel to a nitrocellulose membrane. Membranes were blocked for 1 h with 5% skimmed milk at room temperature, followed by incubation with GPX4 (1:500), SLC7A11 (1:250), TFR1 (1:500), DMT1 (1:500), ERK1/2 (1:500), and p-ERK1/2 (1:500) at 4 °C overnight. The membrane was washed three times in TBST buffer and incubated with secondary antibodies at room temperature for 1 h. Blots were developed using enhanced chemiluminescence reagents. Band intensity was employed as a readout for the relative expression level of each target protein, and this was normalized to the expression of β-actin in each sample.

Cell Culture and the Oxygen-Glucose Deprivation Model
The SH-SY5Y cells were provided by the Chinese Academy of Sciences (Shanghai, China) and were cultured in DMEM supplemented with 10% (v/v) FBS at 37 °C and 5% CO$_2$. An oxygen-glucose deprivation/reperfusion (OGD/R) model was established to mimic I/R injury. SH-SY5Y cells were incubated in glucose- and serum-free DMEM containing CoCl$_2$ (400 μM) for 6 h to simulate ischemia. The culture medium was then replaced with fresh DMEM supplemented with 10% fetal bovine serum (FBS) for 24 h to represent reperfusion. Control cells were incubated in DMEM supplemented with 10% FBS throughout the whole process.

Assessment of Cell Viability
The MTT assay measured the effect of NDSP on SH-SY5Y cellular viability. Briefly, SH-SY5Y cells were grown in a 96-well plate and divided into the control and NDSP groups. NDSP was dissolved in PBS. The cells in the NDSP group were treated with varying concentrations of NDSP (6.25, 12.5, 25, 50, and 100 μg/mL) for 24 h. Cells in the control group were treated with the same volume of PBS. Next, the cells in each well were incubated with MTT (5 mg/mL) at 37 °C for 3 h. Formazan was dissolved in DMSO, and the optical density (OD) value was determined at 570 nm using a microplate reader. Cell viability was calculated as the percentage of the OD value relative to that of the control group (representing 100% viability).

To evaluate the protective effect of NDSP against OGD/R-induced cell injury, the cells were divided into control, OGD/R, and NDSP groups. The cells in the NDSP group were pre-incubated with different concentrations of NDSP (3.125, 6.25, 12.5, and 25 μg/mL) for 6 h. OGD/R and NDSP cells were subjected to OGD/R. Afterward, the cell viability was measured by the MTT assay.
**Hoechst 33342-PI Staining**
The cell death was detected using Hoechst 33342/PI following the manufacturer’s protocol. Briefly, cells were divided into control, NDSP, OGD/R, and +NDSP groups. Cells in the NDSP and +NDSP groups were pretreated with NDSP (6.25 μg/mL) for 6 h. Cells in the OGD/R and +NDSP groups were subjected to OGD/R. After treatment, the culture medium was removed, and the cells were incubated with a working Hoechst 33342/PI solution for 30 min in the dark at 37°C. The fluorescence intensity of each well was recorded using a fluorescence microscope (Zeiss AG, Oberkochen, Germany).

**DCFH-DA Staining**
Reactive oxygen species (ROS) were detected using a DCFH-DA assay kit following the manufacturer’s protocol. Briefly, the cells were divided into the control, OGD/R, OGD/R+NDSP, and OGD/R+NDSP+RSL3 group. Cells in the OGD/R+NDSP+RSL3 group were preincubated with NDSP (6.25 μg/mL) and RSL3 (30 nM) for 6 h. Cells in the OGD/R+NDSP group were pre-incubated with NDSP for 6 h. Cells in the control and OGD/R groups were treated with the same volume of DMSO (0.1%) for 6 h. Cells in the OGD/R+NDSP and OGD/R+NDSP+RSL3 groups were subjected to OGD/R. After cell treatment, the culture medium was removed, and the cells were incubated with a working solution of DCFH-DA for 30 min in the dark at room temperature. The fluorescence intensity of each well was recorded using a fluorescence microscope (Zeiss AG, Oberkochen, Germany).

**Transmission Electron Microscopy**
The SH-SY5Y Cells were fixed in 4% glutaraldehyde and 1% osmic tetroxide at 4 °C. The cells were then dehydrated with 50%, 70%, 80%, and 90% ethanol for 15 min each, followed by dehydration with 100% ethanol three times for 30 min each, and embedded in Eponate 12 resin. Ultrathin sections (50–70 nm thick) were obtained and stained with uranyl acetate and lead citrate solutions for transmission electron microscopy (TEM). Images were captured using a JEOL 1200EX-II electron microscope (Akishima, Japan).

**Screening for Active Ingredients and Targets in NDSP**
The candidate ingredients of NDSP (Panax notoginseng [Burk] F.H. Chen, Carthamus tinctorius L., Crataegus pinnatifida Bunge, Pueraria lobata [Willd.] Ohwi, and Ligusticum chuanxiong Hort) were obtained from the Traditional Chinese Medicine Systems Pharmacology Database and Analysis Platform (TCMSP, [http://tcmspw.com/tcmsp.php](http://tcmspw.com/tcmsp.php)), Encyclopedia of Traditional Chinese Medicine (ETCM; [http://www.tcmip.cn/ETCM/index.php](http://www.tcmip.cn/ETCM/index.php)), and Batman databases. The candidate compounds were first screened through the TCMSP database, using the following screening criteria: Oral bioavailability ≥ 30% and Drug Likeness ≥ 0.18. If no absorption, distribution, metabolism, or excretion (ADME) parameters existed for the candidate compound in the TCMSP database, we used the Swiss ADME database to obtain the compound’s ADME parameters. The screening criteria met the requirements for the gastrointestinal absorption Glabsort to be “High” and with at least two “Yes” hits in Drug Likeness (Lipinski, Ghose, Veber, Egan, Muegge). In addition, based on literature reports, puerarin, ursolic acid, hydroxy safflower yellow A, and notoginseng saponin R1 are potential supplementary active ingredients of NDSP. Targets of the active compounds were obtained from the Swiss TargetPrediction database; targets with scores greater than 0 were classified as possible targets.

**Collection of Disease-Related Targets**
Disease targets were obtained from the OMIM ([https://omim.org/](https://omim.org/)), Genecards ([http://www.genecards.org/](http://www.genecards.org/)), Drugbank ([https://www.drugbank.com](https://www.drugbank.com)), and TTD databases ([http://db.idrblab.net/tdt/](http://db.idrblab.net/tdt/)) using the terms “cerebral ischemia-reperfusion injury” and “ischemic stroke”.

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Drug-Ingredient-Disease-Target Network Construction and Vital Active Ingredients Analysis
Common targets of the drugs and diseases were identified using Venny2.1 (https://bioinfogp.cnb.csic.es/tools/). The Drug-Ingredient-Disease-Target network was constructed using Cytoscape 3.9.2. The key active ingredients were determined using a NetworkAnalyzer. The degree value represents the number of links between ingredients and their targets. The higher the degree value, the more critical the component.

Protein-Protein Interaction Network Construction and Core Target Analysis
The protein-protein interaction (PPI) network of common targets was constructed using the STRING database (https://string-db.org/), with the species limited to “Homo sapiens” and a minimum interaction threshold of 0.9. At the same time, the PPI network interaction data was obtained, and the topology analysis of the common targets was carried out using the NetworkAnalyzer of Cytoscape 3.9.2 software. The top 20 targets were selected based on the degree value, and the core target diagram was drawn using R 4.2.1.

GO Enrichment and KEGG Pathway Analysis
GO enrichment analysis and KEGG pathway annotation of common targets were performed using the online tool DAVID (https://david.ncifcrf.gov).

Molecular Docking
The top six compounds were selected for molecular docking analysis with ERK1 or ERK2 based on the degree values. The predicted structures of the active ingredients were downloaded from PubChem (https://pubchem.ncbi.nlm.nih.gov/), and those of the target proteins were downloaded from the Protein Data Bank database (https://www.rcsb.org/). The Schrödinger Maestro software was used to perform molecular docking, and the binding activity of the target protein to the compound was ranked based on the docking score. Discovery Studio 3.6 software was used to visualize the docking results and identify the binding residues. Molecular dynamics simulations were performed with AMBER 18 program.

Statistical Analysis
All data are expressed as the mean ± standard error of the mean (SEM), and data were analyzed using SPSS 29.0 software or GraphPad Prism 7.03. For n ≥ 5, use the Q-Q plot and Levene's test to test the normality and homogeneity of variance of the data, respectively. If the data showed normality and equal variation, the significance of the difference between multiple groups was analyzed by one-way analysis of variance (ANOVA) followed by Tukey’s post-hoc test. If the data showed non-normal distribution or biased variation, statistical significance was analyzed by Kruskal Wallis test with a two-stage linear step-up procedure from the Benjamini, Krieger, and Yekutieli test for multiple comparisons using GraphPad Prism. For n=3, the bootstrap method was used to analyze the data. A P-value < 0.05 was considered statistically significant.

Results
NDSP Attenuates OGD/R-Induced Ferroptosis in SH-SY5Y Cells
The SH-SY5Y cells were exposed to different concentrations of NDSP (6.25, 12.5, 25, 50, and 100 μg/mL) for 24 h. We observed that NDSP did not exert any toxic effects on SH-SY5Y cells at concentrations below 100 μg/mL (Figure 1A). To evaluate the protective effects of NDSP against cell death, an in vitro CoCl₂-induced OGD/R model was established. As depicted in Figure 1B, treatment with NDSP (6.25–25 ng/mL) significantly enhanced cell viability compared to that in the OGD/R group. Moreover, SH-SY5Y cells treated with NDSP (6.25 μg/mL) exhibited an inhibition of OGD/R-induced cell death (P=0.039) (Figure 1C) along with increased levels of Fe²⁺ (P=0.035) and MDA (P=0.038) (Figure 1D and E). Additionally, treating SH-SY5Y cells with NDSP mitigated the decline in GPX4 expression (P=0.011) mediated by OGD/R and promoted the expression of TFR1 (P=0.032) and DMT1 (P=0.043) (Figure 1F).
NDSP Reduces OGD/R-Induced Cell Injury via Ferroptosis

RSL3 is an inhibitor of GPX4 as it sequesters and inactivates the enzyme, thereby promoting ferroptosis. Incubation with RSL3 (30 nM) abolished the protective effects of NDSP against the OGD/R-induced decline in cell viability ($P=0.001$) (Figure 2A) and LDH release ($P=0.002$) (Figure 2B). Similarly, the NDSP-treated group showed downregulated levels of MDA and ROS, accompanied by the upregulation of GSH.

The protective effects of NDSP were negated when the cells were treated with RSL3 ($P<0.001$) (Figure 2C–F). Regarding morphological changes, TEM analysis revealed that NDSP treatment attenuated mitochondrial swelling and reduced mitochondrial density in SH-SY5Y cells; however, co-incubation with RSL3 abolished these changes (Figure 2G). Collectively, these results indicated that NDSP effectively inhibited OGD/R-induced ferroptosis in vitro.
NDSP Decreases I/R-Induced Cerebral Injury in vivo

There was no significant difference in the infarct area between the sham and control groups. However, the infarct area, neurological score, and cerebral water content in the I/R-treated rats were significantly higher than in the sham group \((P < 0.001)\). Rats treated with NDSP-H exhibited smaller cerebral infarct area, reduced neurological damage, and lower cerebral water content than the rats in the I/R group \((P = 0.007, P = 0.025, \text{and } P < 0.001)\) \((\text{Figure 3A–D})\). Histological examination of these rats’ cerebral cortex and hippocampus revealed that NDSP attenuated the shrunken and pyknotic nuclei and the swollen tissue induced by I/R, as depicted in \(\text{Figure 3E}\). In addition, Nissl staining showed fewer Nissl bodies in the I/R group than in the sham group. However, NDSP-H-treated rats had higher Nissl body than the I/R group \((\text{Figure 3F})\). These data suggested that NDSP reduced cerebral I/R injury.

NDSP Alleviates Cerebral I/R-Induced Ferroptosis

\(\text{Figure 4A–C}\) demonstrates that NDSP administration in rats significantly inhibited the I/R-induced increases in MDA, \(\text{H}_2\text{O}_2\), and \(\text{Fe}^{2+}\) levels \((P < 0.001, P = 0.018, \text{and } P = 0.005)\). Furthermore, although the protein levels of GSH and SOD decreased following I/R, they were elevated in rats treated with NDSP relative to those in the I/R group \((P = 0.034 \text{ and } P = 0.002)\) \((\text{Figure 4D and E})\). Western blot analysis revealed that NDSP-treated rats exhibited higher protein levels of SLC7A11 and GPX4 \((P = 0.003 \text{ and } P = 0.028)\) and lower protein expression of TFR1 and DMT1 \((P = 0.049 \text{ and } P = 0.025)\).
than I/R-treated rats (Figure 4F and G). These findings suggest that NDSP mitigates I/R-induced ferroptosis in the cerebral tissue.

Ferroptosis-Regulating Genes as Candidate Targets of NDSP Activity
We employed a computational approach based on network pharmacology to delineate further the mechanism of action of NDSP in cerebral I/R-mediated ferroptosis. Some 135 active ingredients were identified (Supplementary Table S1), and 1042 drug targets were selected from the Swiss TargetPrediction database (Supplementary Table S2). Furthermore, 669 disease-related genes were included in the I/R dataset (termed “disease dataset”), after querying the OMIM, Genecards, DrugBbank, and TTD databases (Supplementary Table S3). Next, we constructed an overlapping set of 203 common targets whose expression was affected by either NDSP or I/R (Supplementary Table S4). The percentages of targets in the NDSP, I/R, and their overlapping set are shown in Figure 5A. The Drug-Ingredient-Disease-Target dataset used 135 potential active ingredients and 203 common drug-disease targets (Figure 5B). Our analysis of the key active ingredients showed that officinalic acid, baicalein, cnidium lactone, senkyunone, lignan, and (-)-dicentrine were the top candidates...
that may be primarily responsible for the protective effects of NDSP (Supplementary Table S5). Results of GO-biological process analysis of 203 common targets indicated that the targets of NDSP mainly included “response to a drug”, “response to hypoxia”, “inflammatory response”, and “positive regulation of the ERK1 and ERK2 cascade”. Cellular components include the plasma membrane, cell surface, and membrane rafts. Molecular functions were predominantly related to identical proteins, enzymes, and protein binding (Figure 5C). The results of the KEGG enrichment analysis indicated that lipids and atherosclerosis, the MAPK signaling pathway, and the hypoxia-inducible factor signaling pathway may mediate the effects of NDSP in NDSP-regulated cerebral injury (Figure 5D).

**NDSP Regulates Ferroptosis via ERK1/2**

A PPI network comprising 203 common targets was constructed (Figure 6A). According to the topological analysis of the constructed PPI network, the top 20 core targets ranked according to the P-values are shown in Supplementary Table S6. MAPK3, SRC, MAPK1, STAT3, TP53, AKT1, JUN, NFKB1, MAPK14, and PIK3CA were the top ten core targets implicated in the effects of NDSP (Figure 6B). Among these, ERK1/2 (MAPK3 and MAPK1) emerged as crucial genes involved in ferroptosis regulation and served as a core target, ranking first and third in the network pharmacology approach. Therefore, we investigated the effects of NDSP on the expression of ERK1/2. As shown in Figure 6C, NDSP markedly reduced the ratio of phosphorylated ERK1/2 (p-ERK1/2) to total ERK1/2 in SH-SY5Y cells exposed to OGD/R (P=0.019). Similarly, immunohistochemical staining of the rat brain tissue sections demonstrated that NDSP-H treatment attenuated the increased expression of p-ERK1/2 after I/R injury (Figure 6D). To further explore the molecular interactions, we performed molecular docking of the six most important active ingredients, namely, officinalic acid, baicalein, cnidium lactone, senkyunone, lignan, and (-)-dicontrine, with ERK1 and ERK2. These results indicate that baicalein exhibits a strong binding affinity for ERK1 and ERK2. Several residues, including LYS-52 and Glu-69 of ERK2 and Asp-123 and Met-125 of ERK1, formed hydrogen bonds with baicalein (Figure 6E). The binding score are presented in Supplementary Table S7. Next, molecular dynamics simulations were performed for the baicalein-ERK1/2 complex to explore potential interaction mechanisms. The root mean square deviation (RMSD) for the ERK1-baicalein and ERK2-baicalein systems ranged from 0.10–2.50 Å and 0.15–2.25 Å (Figure 6F). The root mean square fluctuation
RMSF results indicate that the ERK1/2 protein has good structural rigidity (Figure 6G). The number of hydrogen bonds in the Erk2-baicalein complex remained at approximately three during the molecular dynamics simulations, whereas that in the Erk1-baicalein complex was approximately two (Figure 6H). Thus, the molecular dynamics simulation results were consistent with the molecular docking results showing that ERK1/2-baicalein had a good binding capacity. These findings suggested that NDSP inhibited I/R-induced ferroptosis by modulating the activity of ERK1/2.

Discussion
Several mechanisms, such as oxidative stress, inflammation, glutamate-mediated neuroexcitatory toxicity, and mitochondrial dysfunction, are involved in cerebral I/R injury. Neuronal death is the main cause of neurological and cognitive impairment. Ferroptosis is a type of programmed cell death that plays an important role in pathological conditions, including cancer, cardiovascular disease, diabetes and its associated complications, I/R organ injury, as well as numerous neurodegenerative diseases. Increasing evidence shows that ferroptosis is closely related to ischemic stroke. In the present study, we assessed the protective effects of NDSP against I/R-induced cerebral ferroptosis.
and explored its underlying mechanisms. We demonstrated that pretreatment of rats with NDSP 1.08 g/kg alleviated I/R-induced ferroptosis, decreased infarct area, and improved overall neurological scores.

The regulatory mechanism of ferroptosis involves perturbations in iron transport, amino acid metabolism, and lipid peroxidation. Iron is crucial for normal neuronal function at baseline, and an overabundance of iron is associated with the exacerbation of several neurodegenerative diseases. Regarding the modulation of iron levels, TFR1 and DMT1 are involved in intracellular iron regulation, maintaining iron metabolism and homeostasis. TFR1 facilitates the cellular uptake of Fe$^{3+}$ via endocytosis of transferrin, which is subsequently reduced to Fe$^{2+}$ by the six-transmembrane epithelial antigen protein 3. DMT1 and ZIP8/14 transport Fe$^{2+}$ into the cytoplasm. Aberrant expression of TFR1 and DMT1 serves as a marker of ferroptosis. Here, we investigated the effect of NDSP on iron metabolic pathways and found that NDSP attenuated the imbalance in iron metabolism by decreasing TFR1 and DMT1 expression.

The accumulation of iron-dependent lipid peroxidation products is a characteristic feature of ferroptosis. Excessive lipid ROS levels induce oxidative stress, damaging proteins, lipids, nucleic acids, membrane rupture, and subsequent cell death. In this context, the inhibition of ferroptosis presents a potential strategy for ameliorating I/R injury. We observed that in brain tissue samples of I/R-treated rats or OGD/R-treated SH-SY5Y cells, there was a significant surge in the levels of MDA, Fe$^{2+}$, and H$_2$O$_2$ in brain tissue samples from I/R-treated rats. The NDSP treatment reversed these
changes. The SLC7A11/GPX4 signaling axis is a critical pathway that regulates glutathione production.\textsuperscript{51} Downregulation of SLC7A11 leads to reduced intracellular cystine levels, depleting glutathione biosynthesis, and indirectly suppressing GPX4 activity, consequently promoting ferroptosis. NDSP treatment restored the expression of SLC7A11 and GPX4, thereby exerting a protective effect against cerebral I/R injury. In contrast, incubation with RSL3, a GPX4 inhibitor, abrogated the protective effects of NDSP against OGD/R-induced ROS generation. These findings suggested that NDSP protects against I/R-induced cerebral injury by inhibiting ferroptosis.

This study used network pharmacology to analyze how NDSP inhibited ferroptosis induced by cerebral ischemia-reperfusion. We identified 203 common targets and performed GO and KEGG for these 203 common targets. GO enrichment analysis suggested that NDSP treatment for I/R-induced injury involves positive regulation of the ERK1 and ERK2 cascades in biological processes, and KEGG enrichment analysis indicated that common targets are involved in the MAPK signaling pathway. Although the MAPK signaling pathway did not rank high in the GO and KEGG analyses, the results of the topology analysis of the PPI network suggested that genes such as MAPK3, SRC, MAPK1, STAT3, TP53, and AKT1 are the core targets of NDSP. These results suggest that ERK1/2 may be involved in the effects of NDSP on I/R-induced ferroptosis.

The MAPK signaling cascade is essential for regulating ferroptosis, as activated ERK1/2 regulates the expression of ferroptosis-related genes. ERK1/2 increases the sensitivity of human breast epithelial cells to ferroptosis, and ERK inhibition upregulated the expression of GPX4.\textsuperscript{15} In pancreatic ductal adenocarcinoma, ERK1/2 is required for elastin-induced ferroptosis.\textsuperscript{52} In macrophages, ERK1/2-mediated ferritinophagy contributes to ferroptosis induced by cadmium telluride quantum dots.\textsuperscript{53} Zille et al demonstrated that the ERK1/2 inhibitor U0126 abrogated hemin-induced ferroptosis in primary cortical neurons.\textsuperscript{54} In a spinal cord ischemia/reperfusion injury model, the ERK1/2/c-Myc/TFR1 signaling axis participates in the Eph receptor A4 regulation of motor neuron ferroptosis.\textsuperscript{55} Our observations indicated that NDSP alleviated the increase in ERK1/2 phosphorylation mediated by I/R and OGD/R. These results are consistent with those reported by pharmacologists. Based on our molecular docking results, baicalein displayed the most satisfactory binding affinity for ERK1/2 compared to the other identified active ingredients. Furthermore, the molecular dynamics simulation results of baicalein-ERK1/2 supported those of molecular docking. However, further research is needed to explore how the active ingredients of NDSP regulate ERK1/2 phosphorylation.

**Conclusion**

In summary, we have demonstrated that NDSP exerts a protective effect against ferroptosis during cerebral I/R injury. Furthermore, our findings indicated that this protective mechanism is associated with the inhibition of ERK1/2 phosphorylation. Our study offers valuable insights for potentially improving therapeutic approaches for the treatment of stroke.

**Abbreviations**

ADME, absorption, distribution, metabolism and excretion; CCA, common carotid artery; DMT1, divalent metal-ion transporter-1; ECA, external carotid artery; ERK1/2, mitogen-activated protein kinase 1/2; GPX4, glutathione peroxidase 4; GSH, glutathione; H\textsubscript{2}O\textsubscript{2}, hydrogen peroxide; ICA, internal carotid artery; I/R, ischemia/reperfusion; MDA, malondialdehyde; NDSP, Naodesheng pill; NDSP-L, low-dose NDSP; NDSP-H, high-dose NDSP; OGD/R, oxygen glucose deprivation/reoxygenation; OD, optical density; ROS, reactive oxygen species; SEM, standard error of the mean; SLC7A11, solute carrier family 7 member 11; SOD, superoxide dismutase; TC MSP, Traditional Chinese Medicine Systems Pharmacology Database and Analysis Platform; TEM, transmission electron microscopy; TFR1, transferrin receptor 1; TTC, triphenyltetrazolium chloride.

**Data Sharing Statement**

The datasets used and/or analyzed in the current study are available from the corresponding author (Yong-Yu Yang) upon reasonable request. Uncropped gels and blots are provided in Supplementary Figure 1.
Ethics Approval and Consent to Participate
All experimental protocols were designed according to the Guide for the Care and Use of Laboratory Animals (NIH Publication, 8th edition, 2011) and approved by the Animal Ethics and Welfare Committee of the Second Xiangya Hospital of Central South University (No. 2021955).

Author Contributions
All authors made a significant contribution to the work reported, whether that is in the conception, study design, execution, acquisition of data, analysis, and interpretation, or in all these areas; took part in drafting, revising, or critically reviewing the article; gave final approval of the version to be published; agreed on the journal to which the article has been submitted; and agreed to be accountable for all aspects of the work.

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Disclosure
The authors have no competing interest to declare in this work.

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