Anti-Stress Activity of Oviductus Ranae in Mice with Acute Stress Based on Network Pharmacology

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ABSTRACT

Stress is a state of threatened homeostasis that causes the body to release the hormone cortisol produced by the adrenal glands. Oviductus ranae (OR) is an animal-based raw material of folk medicine which plays a variety of activities. However, its anti-stress effects mechanism has not been fully revealed. In this work, based on network pharmacology, the potential targets of OR were screened, and a protein-protein interaction (PPI) network between the target of OR and anti-stress target was constructed using STRING database. Kyoto Encyclopedia of Genes and Genomes (KEGG) was used for analyzing the pathways of target gene. To further verify this, total 96 ICR mice were used, forced swim test and anoxic tolerance test were performed. The effect of OR on levels of monoamine neurotransmitters and phosphorylation of p38 which closely related to anti-stress were examined. The results showed that 203 potential OR targets and 126 stress-related gene targets were obtained, in which there were 15 common targets. Pathway enrichment analysis showed that there were 20 critical pathways. The results revealed that OR could increase the total swimming time, increase the survival time of enduring anoxia, and regulate monoamine neurotransmitters such as 5-hydroxytryptamine (5-HT), 5-hydroxyindoleacetic acid (5-HIAA), norepinephrine (NE) and dopamine (DA). Western blot analyses indicated that OR may decrease the phosphorylation of p38. In conclusion, the results revealed that OR may play the anti-stress effects by inhibiting mitogen-activated protein kinase (MAPK) pathway, thus promote the normalization of acute stress. This study revealed the possible mechanism of OR as a potential material for the treatment of acute stress-related problems, and laid a foundation for the further development and utilization of OR.

INTRODUCTION

Stress is regarded as the body’s nonspecific response to external stimuli, which threatens homeostasis that interferes with the normal physiological balance and endangers the survival of the individual (Mora et al., 2012; McEwen, 2007). Both acute and chronic stress stimuli can cause changes in daily behavior, regulate the function of hypothalamic-pituitary-adrenal axis (HPA axis) and autonomic nerve levels (Lupien et al., 2009). Stress response is affected by the duration and intensity of different stressors. Acute stress plays a vital part in various brain diseases and symptoms, including Parkinson’s disease, depression and neuroinflammation (Gispen-de Wied, 2000; Joëls et al., 2007). In addition to an important role against brain diseases, stress is also a risk factor for cardiovascular and skin diseases (Joëls and Baram, 2009). Oviductus ranae (OR), also called R. chensinensis oil, is the dry oviduct of Rana chensinensis. It has been for hundreds of years as a traditional medicine in China. It can be used to treat various diseases, such as frailty, night sweat, menopause syndrome, insomnia and neuroasthenia (Xiao and Jiang, 2010; Wang et al., 2010). In recent years, more and more studies have reported the active function of OR, and conducted in-depth studies on its mechanism of action in immune regulation, anti-oxidation, anti-fatigue, anti-
aging, estrogen-like, liver protection, lowering blood lipid, anti-osteoporosis, anti-depression, anti-tumor, anti-inflammation, anti-asthmatic and other aspects (Zhang et al., 2019; Li et al., 2019; Sheng et al., 2020). It has been shown that OR has good antioxidant and immunomodulatory effects, and can stimulate the activity of macrophages by regulating NF-κB pathway, and inhibit the ovarian cell apoptosis in rats induced by oxidative stress injury by declining the production of ROS and increasing mitochondrial membrane potential (Huang et al., 2014; Ling et al., 2019). These results demonstrate that OR has great potential and is worthy of further explore and development. In recent two decades based on the development of systems biology, the concept of network pharmacology began to rise. The methods of omics data integration and multi-target drug development and related databases were applied (Zhang et al., 2019; Yang et al., 2020), especially in the research of traditional Chinese medicine (Zhao et al., 2020; Li et al., 2021; Zuo et al., 2021).

In this study, the potential effect and mechanism of OR anti-stress were investigated by network pharmacology. The major activated components of OR were screened, and stress-related targets were gathered following by the intersecting and potential possible targets were predicted, and stress-related targets by constructing protein-protein interaction (PPI) networks. The analysis of PPI networks was performed with the Search Tool for Retrieval of interaction Genes/Proteins platform (STRING) (Chen et al., 2020), and the criterion was limited to Homo sapiens with a confidence level of 0.7.

**Acquisition of OR potential active ingredients and targets**

The chemical compounds of OR were screened from traditional Chinese medicine systems pharmacology database (TCMSP), with the active ingredients selected based on oral bioavailability (OB) ≥30% and drug similarity (DL) ≥0.18 (Zhang et al., 2016). Potential targets related to bioactive compounds of OR were further analyzed, and transformed to UniProt ID through UniProtKB database. Stress-related targets were collected using Drugbank and GeneCards database. On this basis, two sets of indicators were compiled: potential targets of OR and stress-related indicators. All the details of databases and bioinformatics tools were listed in Supplementary Table SI.

**Construction of PPI network**

To explore the interaction of OR therapeutic targets and stress-related targets by constructing protein-protein interaction (PPI) networks. The analysis of PPI networks was performed with the Search Tool for Retrieval of interaction Genes/Proteins platform (STRING) (Chen et al., 2020), and the criterion was limited to Homo sapiens with a confidence level of 0.7.

**KEGG pathway and gene ontology enrichment analyses**

To clarify the OR therapeutic targets at the system level role in the biological mechanisms of anti-stress, Gene Ontology (GO) and Kyoto Encyclopedia of Genomes (KEGG) pathway were executed with the ClueGO 2.5.4 plugin in Cytoscape (Li et al., 2021). The minimum number of enriched genes per GO or KEGG term was set to 3.

**Materials and experimental groups**

Total 96 ICR mice (6-week-old, half male and half female) were obtained from Changsheng Biotechnology Co. Ltd. (Shenyang, China), and were previously housed (12-h light and dark cycle, 22°C, 70% humidity) in pathogen-free conditions, 6 mice per cage, with food and water were freely available. Then, mice were divided into two same parts. In each part, total 48 mice were randomized into four groups (n=12): the control (20 mL/kg of distilled water), OR doze groups at 100, 200 and 400 mg/kg. Mice in each group were administered orally once a day for 14 consecutive days.

**Forced swimming test**

The forced swimming test was performed according to the method described by Aluko et al. (2015) and Yan and Hao (2016). In brief, one h after the last administration, each mice was forced to swim individually in a transparent tank (30, 45 and 40 cm in length, width, and height, respectively) filled with fresh water to a depth of 30 cm.
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and the water was maintained at 17±1°C. Exhaustion was determined by observing the loss of coordinated movements and failure to return to the surface within 10 s. The parameter measured were total time spent in active swimming, that is, the total duration during which the animal swims throughout the experimental period.

**Anoxic tolerance test**

The other part of 48 mice were used for anoxic tolerance test, according to the method described by Caillard *et al.* (1979) and Aluko *et al.* (2015). Before the experiment, 250 mL conical flasks were sealed with rubber cork. one hour after the last administration, the mice were placed in conical flask and the anoxic tolerance time was recorded. After that, the mice were removed immediately for recovery. The anoxic tolerance time was defined as the latency to the first appearance of anoxic convulsions.

**ELISA test**

After forced swimming test, mice brain tissues were taken from each group. The levels of monoamine neurotransmitters (5-hydroxyindoleacetic acid, 5-HIAA; 5-hydroxytryptamine, 5-HT; dopamine, DA and norepinephrine, NE) in brain tissues were measured by ELISA kit according to the kit instructions. The absorbance of each sample was measured at 450 nm using a multiskan spectrum (Thermo Fisher Scientific, USA). All experiments were performed in triplicate and repeated three times.

**Western blot analysis**

After forced swimming test, the proteins of brain tissues were extracted and the expression levels of p-p38 and p38 were analyzed by western blot. Brain tissue were reaped and lysed with RIPA buffer cracking 30 minutes on the ice, and the protein concentration was evaluated with the BCA assay kit. Those cerebrum tissue proteins (15 μL) were loaded onto 12% SDS-polyacrylamide gel. After electrophoresis, these gels were imprinted onto polyvinylidene fluoride (PVDF) membranes and sealed with 5% (W/V) skim milk for 1 h. Then, the transferred membrane and the primary antibody were incubated overnight at 4°C. The conjugation of primary antibody was detected by HRP coupled secondary antibody, and enhanced chemiluminescence (ECL) imaging was utilized.

**Statistical analysis**

All data were expressed as mean ± SD and SPSS 19.0 was used for statistical analysis. One-way analysis of variance (ANOVA) was used to evaluate the differences between groups. Differences at p <0.05 were considered statistically significant.

**RESULTS**

**Potential active ingredients and targets of OR**

All the active ingredients and related targets were searched in TCMSP database. In the result, a total of 14 active components were obtained from OR, corresponding to 203 target genes. Using “Stress” as the key word, 126 disease-related targets were obtained after screening, and 15 convergence targets then afterward the convergence of the two, which may be potential targets of OR anti-stress (Supplementary Fig. S1A, B).

**Construction of the PPI network**

The PPI network of OR interacting with anti-stress-related proteins was constructed in String database (Supplementary Fig. S1C). As shown in the result, there were SLC6A2, IL10, JUN, CAT, AKT1, VEGFA, CAV1, RELA, SOD1, IL1B, GSTP1, HSF1, APOA1, NR3C1 and FOS that to be considered as the key targets of OR against stress (Table I).

**Table I. The key targets of OR anti-stress targets by PPI network.**

<table>
<thead>
<tr>
<th>Items</th>
<th>Details</th>
</tr>
</thead>
<tbody>
<tr>
<td>SLC6A2</td>
<td>Solute carrier family 6 member 2</td>
</tr>
<tr>
<td>IL10</td>
<td>Interleukin 10</td>
</tr>
<tr>
<td>JUN</td>
<td>Jjun proto-oncogene</td>
</tr>
<tr>
<td>CAT</td>
<td>Catenin</td>
</tr>
<tr>
<td>AKT1</td>
<td>Serine/threonine kinase 1</td>
</tr>
<tr>
<td>VEGFA</td>
<td>Vascular endothelial growth factor A</td>
</tr>
<tr>
<td>CAV1</td>
<td>Caveolin-1</td>
</tr>
<tr>
<td>RELA</td>
<td>v-rel avian reticuloendotheliosis viral oncogene homolog A</td>
</tr>
<tr>
<td>SOD1</td>
<td>Superoxide dismutase 1</td>
</tr>
<tr>
<td>IL1B</td>
<td>Interleukin 1B</td>
</tr>
<tr>
<td>GSTP1</td>
<td>Glutathione S-transferaz pi</td>
</tr>
<tr>
<td>HSF1</td>
<td>Heat shock transcription factor 1</td>
</tr>
<tr>
<td>APOA1</td>
<td>Apolipoprotein A1</td>
</tr>
<tr>
<td>NR3C1</td>
<td>Nuclear receptor subfamily 3 group C member 1</td>
</tr>
<tr>
<td>FOS</td>
<td>Fructooligosaccharides</td>
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</tbody>
</table>

**GO function enrichment analysis and KEGG pathway**

Total 141 proteins involved in the target network of named components were analyzed. In GO enrichment analysis, 55 GO items and 20 biological process related items were identified, the results revealed that the critical targets of effective compounds in OR are enriched significantly and the main biological functions were transcription factor binding, antioxidant activity and steroid binding in Figure 1A.
The KEGG pathway enrichment analysis showed that, OR mainly involved in the signaling pathways of toll-like receptor, C-type lectin receptor, T cell receptor, TNF, Relaxin, MAPK and B cell receptor; the involved 20 critical pathways were shown in Figure 1B. The results suggest that, the higher the order of enrichment results, the more likely the OR is to exert its antipyretic effect through these pathways. The top five enriched KEGG pathways include Fluid shear stress and atherosclerosis, Chagas disease, Yersinia infection, lipid and atherosclerosis and MAPK signaling pathway.

Effect of OR on the forced swim endurance and anoxic tolerance tests

The total swimming time of 400 mg/kg OR group significantly increased, the 100 mg/kg OR group also increased compared with the control group, but there was no significant difference (Fig. 2A), indicating that OR could enhance the endurance of mice by prolonging swimming time, thus improving their stress ability. The convulsion latency (400 mg/kg OR group) increased significantly comparing to the control group, the 200 and 100 mg/kg OR group also increased but no significant difference (Fig. 2B), revealing that OR could enhance mice tolerance to hypoxia by prolonging the convulsion latency, thus improving their anti-stress ability.

Effects of OR on the monoamine neurotransmitters

The expressions of 5-HT, 5-HIAA, NE and DA in the brain tissues were detected. The results showed that, the levels of 5-HIAA and DA in 400 mg/kg OR group increased markedly \( p < 0.05 \), compared with control group. The contents of 5-HT and NE in 100, 200 and 400 mg/kg OR groups also increased significantly \( p < 0.05 \) or \( p < 0.01 \); while the levels of 5-HIAA, 5-HT, DA and NE in other groups all have an increasing trend with no significant difference (Fig. 3). It revealed that OR could improve the anti-stress ability of mice by increasing the content of various monoamine transmitters in brain.

Fig. 2. Effect of OR on the forced swimming endurance test (A) and anoxic tolerance test (B) in mice. * \( p < 0.05 \) compared with the control.

Fig. 3. The effect of OR on the changes of monoamine neurotransmitters in brain tissue under stress state. 5-HIAA, 5-hydroxyindoleacetic acid (A); 5-HT, 5-hydroxytryptamine (B); DA, dopamine (C); NE, norepinephrine (D). * \( p < 0.05 \) or ** \( p < 0.01 \) compared with the control.
**Effects of OR on phosphorylation of p38**

The mitogen-activated protein kinases (MAPK) mediate a variety of cellular behaviors that in response to extracellular stimuli. The p38 group, as a main sub-groups of MAPK, serves as a junction for signal transduction, and play a crucial role in many biological processes (Zarubin and Han, 2005). The expression of p-p38 and p38 from the brain tissues in 400 mg/kg OR group was clearly increased (Fig. 4) comparing to the control group. It indicated that OR could increase phosphorylation of p38 in mice brains.

![Fig. 4. Effects of OR on phosphorylation levels of p38 in the brain tissues by western blot.](image)

**DISCUSSION**

The method of network pharmacology has been widely used in the research of traditional Chinese medicine (Zuo et al., 2021; Zhao et al., 2021; Yin et al., 2021). In this study, we constructed a comprehensive PPI network based on OR-related and stress-related networks, the result showed total 15 core targets involved in the anti-stress pharmacological action of OR were identified. Subsequent analysis of GO and KEGG signaling pathway enrichment of these key targets were performed. It is suggested that the anti-stress effect of OR may be caused by regulating key signaling pathways of oxidative stress, inflammation, and cell proliferation. To our knowledge, there are no similar studies base on network pharmacology for OR. Notably, over the top oxidative stress produces oxidative radicals, for example, ROS, that might animate MAPK course phosphorylation (Subramaniam and Unsicker, 2010; Madrigal et al., 2003). Some evidence suggests that acute stress is associated with the release of oxidative free radicals that promote phosphorylation of MAPK cascade in the hippocampus of stressed rats (Chen et al., 2018). And Inhibition of oxidative-stress-induced MAPK phosphorylation relieves the stress response in a rat stress model (Madrigal et al., 2001; Sasaguri et al., 2005). In fact, in this study, the western blot showed that brain tissue p-p38 levels were significantly reduced after the OR treatment, which was similar to an earlier study (Madrigal et al., 2003), and also the predictions of network pharmacology were preliminarily validated.

The regulation of monoamine transmitters is an important mechanism of drug resistance to stress (Ahmad et al., 2012). Exposure to stress causes certain regions of the brain (mainly the hippocampus, amygdala, prefrontal cortex, and nucleus accumbens) to rapidly release monoamines such as dopamine, norepinephrine, and serotonin (Ramadan and Alahamrani, 2015; Adamec et al., 2008; Joëls and Baram, 2009). Stress-induced release of monoamines typically occurs within minutes of the onset of stress and may persist throughout the duration of stress (Aston-Jones and Cohen, 2005). Our study also found that OR significantly modulates monoamine transmitters in the brain tissue of stressed mice during anti-stress. It showed that the content of 5-HIAA, 5-HT, DA and NE in the brain was significantly increased.

At present, studies on the biological activities of OR mainly focus on antioxidant, anti-inflammatory, immune regulation and bone metabolism (Huang et al., 2014; Li et al., 2019). Among them, some researchers have preliminarily investigated its anti-oxidative stress effect at the cellular level, and found that serum containing OR can down-regulate negative regulatory factors of cell proliferation, activate ERK1/2, inhibit JNK and p38 activities, reduce ROS production, improve mitochondrial membrane potential, and inhibit H2O2-induced apoptosis of rat ovarian cells (Ling et al., 2019). Our recent study also proved that OR has effect on improving cognitive disorder (Li et al., 2022). In this work, the anti-stress effect of OR was confirmed by network pharmacology prediction and animal experimental verification, which further confirmed the aforementioned reports and provided a reference for its in-depth research and development.

**CONCLUSION**

In conclusion, a total of 14 active components in OR corresponding to 15 anti-stress related targets were found, and through forced swimming endurance and anoxic tolerance test, it suggested that the regulation of monoamine neurotransmitters and the phosphorylation of p38 play a crucial part in the anti-stress process. This study aims to provide a reference for the basic function research.
of OR, and the development of its related biological products.

ACKNOWLEDGMENTS

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Ethical compliance

Research experiments conducted in this article with animals were approved by the Animal Experiment Ethics Committee of Changchun University of Chinese Medicine (Ethic code 2020217) following all guidelines, regulations, legal, and ethical standards as required for humans or animals.

Supplementary material

There is supplementary material associated with this article. Access the material online at: https://dx.doi.org/10.17582/journal.pjz/20210818020819

Statement of conflict of interest

The authors have declared no conflict of interest.

REFERENCES


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Supplementary Material

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Supplementary Table SI. Databases and bioinformatics tools.

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0030-9923/2022/0001-0001 $ 9.00/0

Supplementary Fig. S1. The therapeutic targets of OR against stress and the bioactive compound-therapeutic target network. (A) The Venn diagram of the overlapping targets between the putative targets of OR and stress-related targets. (B) The bioactive compound-therapeutic target network of OR against stress. (C) The PPI network of the therapeutic targets of OR against stress.