

Notoginsenoside R2 induces colonic microvascular injuries via regulating the Rap1GAP/PI3K/Akt signaling pathway

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Background: Notoginsenoside R2 (NGR2) is an important active saponin molecule of *Panax Notoginseng* (PN), but its effect on regulating angiogenesis is unclear. Here, we investigated the role of NGR2 in angiogenesis *in vivo* and *in vitro*.

Methods: NGR2 was administered to rats by intragastric administration for 7 days. The colonic histopathology and microvessel density (MVD) were observed and evaluated under an inverted microscope. The colonic mucosal permeability (MP) and vascular permeability (VP) were evaluated by measuring the transmittance of FD-4 and the vascular leakage of Evans blue, respectively. The serum IL-2, TNF- a, IL-4, IL-10, VEGFA165, and VEGFA121 levels were detected with ELISA. *In vitro*, pHUVECs were cultured and treated with NGR2. The mechanism of NGR2 in angiogenesis was assessed by VEGFA165, LY294002, silencing, and overexpression plasmids of *Rap1GAP in vitro*. The cell viability, cell proliferation, tube formation, and concentration of intracellular pyruvate and lactic acid of pHUVECs were measured after 24 h. The expression of proteins was detected with western blotting.

Results: NGR2 could significantly induce inflammatory injuries in the colonic mucosa and microvessels. Both MP and VP in rats treated with NGR2 increased in proportion to the serum VEGFA165 and VEGFA121 level, the ratio of VEGFA165/VEGFA121, and the concentration of intracellular pyruvate and lactic acid. *In vitro*, NGR2 reduced cell viability, proliferation, and tube formation, and enhanced the intracellular glycolysis of pHUVECs. Furthermore, the cell viability, proliferation, and tube formation of pHUVECs were inhibited by NGR2 via blocking the Rap1GAP/PI3K/Akt signaling pathway.

Conclusions: NGR2 could induce colonic mucosal microvascular injuries and promoted the intracellular glycolysis of pHUVECs via blocking the Rap1GAP/PI3K/Akt signaling pathway.

Keywords: Notoginsenoside R2 (NGR2); glycolysis; angiogenesis; PI3K/Akt signaling pathway

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Introduction

Panax Notoginseng (PN) has been used traditionally for the treatment of traumatic injuries in East Asia for more than 2,000 years. The formation of granulation tissue during the repair of traumatic injuries is the result of the controllable regulation of angiogenesis, including promoting angiogenesis and anti-angiogenesis (1), which have been shown to be regulated by PN (2). Our previous research found that PN could significantly promote angiogenesis and reduce vascular inflammatory injuries (3,4). Indeed, some saponins in PN, which are important active chemical components including ginsenosides and notoginsenosides, have appeared to lack toxicity in both animals and humans (5). For example, notoginsenoside, such as NGR1, NGFc, and NGRg1, had the effect of promoting angiogenesis, protecting the blood vessels from injuries, and inhibiting inflammation (6,7). However, the antiangiogenesis effect and active ingredients of PN have been seriously neglected in previous studies.

Notoginsenoside R2 (NGR2) is an active saponin molecule extracted from the root of *PN*. Previous studies showed that it reduced A25-35-induced neuronal apoptosis and inflammation (8), and stimulated the growth of bone marrow nucleated cells and hematopoietic stem cells, as well as burst-forming unit-erythroid and colony-forming unit granulocyte-monocyte colony expansion (9). However, the role of NGR2 in angiogenesis remains unexplored, which is precisely the purpose of this study.

The Ras-related protein 1-GTPase-activating protein 1 (Rap1GAP)/PI3K/Akt signaling pathway is an important signal pathway regulating the cell proliferation of vascular endothelial cells (VECs), which can be activated by vascular endothelial growth factor A (VEGFA). Activating the VEGFA/Rap1GAP/PI3K/Akt signaling pathway could significantly promote the cell proliferation of human umbilical vein endothelial cells (HUVECs) and reduce inflammation (10,11). Our previous studies also found that the VEGF/Rap1GAP signaling pathway had a critical role in regulating the cell proliferation of VECs, as blocking the VEGF/Rap1GAP signaling pathway could significantly increase colonic mucosal inflammatory injuries in rats via up-regulating the expression of thrombospondin 1 (TSP1) (12). TSP1 is a critical protein involved in increasing vascular injuries and promoting vascular remodeling (13,14). Our results in previous studies

showed that PN could significantly regulate the secretion of VEGFA165 and VEGFA121 and down-regulate the expression of TSP1 protein in colonic mucosa (3). However, the regulatory effect of NGR2 on the Rap1GAP/PI3K/Akt signaling pathway is unclear. Here, we investigated the role of NGR2 in angiogenesis *in vivo* and *in vitro* to further understand the anti-angiogenesis effect of PN. The findings suggest that PN may have potential vascular toxicity except for promoting angiogenesis.

We present the following article in accordance with the ARRIVE reporting checklist (available at https://dx.doi.org/10.21037/atm-21-5898).

Methods

Cell culture and animals

Primary human umbilical vein endothelial cells (pHUVECs) were obtained from ATCC (Maryland, USA) and maintained in vascular cell basal medium (ATCC) at 37 °C in a humidified atmosphere of 95% air and 5% CO₂.

Sprague Dawley (SD) rats (120–140 g) were purchased from Shanghai SLAC Laboratory Animal Co. Ltd. (Shanghai, China). The care and treatment of mice were in accordance with institutional guidelines. All animals used in the present study were housed in a pathogen-free environment, and animal experiments were approved by the Institutional Animal Care and Use Committee of Shanghai University of TCM (No. SZY201612006), in compliance with the Institutional Animal Care and Use Committee of Shanghai University of TCM's guidelines for the care and use of animals. After 1 week of adaptive feeding, the study begun when the rats behaved normally.

Reagents

Formamide (purity was above 99%) and Evans Blue (no: 295876, dye content was above 75%) were purchased from Sigma Chemical. Co. (Sigma-Aldrich, MO, USA). NGR2 was obtained from Shanghai R&D Center for Standardization of Chinese Medicines (Shanghai, China). Enzyme-linked immunosorbent assay (ELISA) kits (IL-10, IL-4, IL-6, and TNF-a) were from R&D Systems (R&D, USA). VEGFA121 (no: SEB851Ra) and VEGFA (no: SEA143Ra) ELISA kits were purchased from Cloud-Clone Co. Anti-VEGFR2 antibody (ab11939), anti-VEGFR2

(phospho Y1214) antibody (ab5475), and anti-Rap1GAP antibody (ab32373) were purchased from Abcam Co. (London, United Kingdom). Anti-VEGFR2 (phospho Y1175) antibody (AF4426) was from Affinity Biosciences (USA). Human VEGFA165 protein (#8065), LY294002 (PI3K inhibitor, #9901), anti-PI3K antibody (#4249), phospho-PI3K (#4228), Akt (#4691), phospho-Akt (#13038 and #4060) antibodies, goat anti-mouse antibody, and goat anti-rabbit antibody were from Cell Signaling Technology (USA). Anti-CD31 antibody (sc-376764) and anti-TSP1 antibody (sc-393504) were from Santa Cruz Biotechnology Co. (Santa Cruz, CA, USA). Human Rap1GAP gene silencing plasmid (designed by targeting the NM_001145658 gene) and human Rap1GAP gene overexpression plasmid (targeting the NM_001145658 gene) were constructed and synthesized by GK Gene (Shanghai, China). Lipofectamine 2000 Transfection Reagent (#11668019) was purchased from Life Technologies Co. (Waltham, MA).

NGR2 treatment in vivo

NGR2 (5.0 10.0, and 20.0 μ M) was administered to normal rats once a day by intragastric administration (i.g., n=8). In the normal control group (n=8), normal saline was administered to the rats i.g. once a day. After 7 days, the rats were anesthetized with 180 μ L 3% sodium pentobarbital per 100 g. Blood was collected from the abdominal aorta and the colon tissues were obtained. The colon tissues were formaldehyde-fixed, paraffin-embedded, and stained with hematoxylin and eosin (HE). A protocol was prepared before the study without registration.

pHUVECs treated with LY294002, VEGFA165, and NGR2

pHUVECs were treated with VEGFA165 (1.0 nM), LY294002 (50 μ M), and NGR2 (5.0, 10.0, and 20.0 nM) for 24 h or in combination. The medium was removed and the adherent cells were washed 3 times with phosphate-buffered saline (PBS) to eliminate any residual LY294002 and NGR2. Fresh DMEM without fetal bovine serum (FBS) was added to the cells for 3 h. This conditioned medium (CM) was centrifuged for 15 min at 5,000 ×g, passed through a 0.2- μ m filter, and incubated for 2 h at 37 °C with

the remaining medium removed.

Histopathology, microvessel density (MVD), vascular permeability (VP), mucosal permeability (MP), cell viability assay, cell proliferation assay, tube formation assay, plasmid construction and cell transfection, and western blotting

Histopathology, MVD, cell viability assay, cell proliferation assay, tube formation assay, plasmid construction and cell transfection, and western blotting were carried out as previously described (12). The VP and MP in the colon were detected as previously described (3,15). All assays were performed at least 3 times.

Metabolic substrate assay

After treatment, the content of pyruvate and lactate in cell lysates was measured using the NADH optical test as previously described (16). For determining the intracellular pyruvate and intracellular lactate levels, 100 μL of lysate from 1×10⁶ trypsinized cells was prepared with a hypotonic buffer (20 mM HEPES, 0.5 mM CaCl₂, 1 mM MgCl₂). Following centrifugation at 13,000 ×g, 100 µL of supernatant was collected and adjusted to pH 6.5-8.5 by adding 64.6 µL of ice-cold 2 M KOH for 20 min. The resulting precipitate was centrifuged and the supernatant was collected. After incubating for 30 min at room temperature, the change in fluorescence was measured at the excitation and emission wavelengths of 355 and 460 nm, respectively (Victor 3, Perkin Elmer). Pyruvate and lactate concentrations were calculated from a reference curve. Subsequently, the intracellular lactate concentration was calculated based on the concentration of protein in the cell lysate (1 μg/μL).

Statistical analysis

Data were presented as the mean ± SEM. One-way ANOVA or general linear model with repeated measures was used to analyze data sets with 3 or more groups, and the least significant difference *post hoc* test was used for multiple comparisons. Student's *t*-test was used to analyze data sets with two groups. P<0.05 was considered significant.

Results

NGR2 induced microvascular and mucosal injuries in the rat colons

SD rats were divided into the control group (NS, iv), low-dose group (5.0 µM NGR2, iv), medium-dose group (10.0 µM NGR2, iv), and high-dose group (20.0 µM NGR2, iv). After 7 days of treatment, obvious colonic mucosal inflammation and vascular injuries were found in the colonic mucosa of the high-, medium-, and lowdose groups compared with the control group (Figure 1A). Meanwhile, VP was increased greatly in both the highdose group and medium-dose group, along with increased MP (Figure 1B, 1C). In addition, the serum contents of pro-inflammatory cytokines (IL-6 and TNF-α) were significantly higher, and serum IL-4 and IL-10 were significantly lower in the high-dose group and mediumdose group than those in the control group (Figure 1D-1G). The results showed that NGR2 could lead to vascular and mucosal inflammatory injuries, with increased VP in rat colons.

NGR2 increased hypoxia in the colon mucosa with an increased ratio of serum VEGFA165/VEGFA121 and increased protein expression of Rap1GAP and TSP1

With the colonic mucosal vascular injuries increased, the protein expression levels of Rap1GAP and TSP1 in the colon mucosa were significantly higher in the high-dose group and medium-dose group than in the control group. However, there were no significant differences in the protein expression of VEGFR2, VEGFR2 (p1175), and VEGFR2 (p1214) between the control group and treatment groups (*Figure 2A*). Meanwhile, the serum contents of VEGFA165 and VEGFA121 and the ratio of VEGFA165/ VEGFA121 were significantly higher in the high-dose group and medium-dose group than in the control group (*Figure 2B,2C*).

In addition, the protein expression of HIF-1 α and the concentration of intracellular pyruvate and lactic acid were significantly increased in the colon mucosa in the high-dose group and medium-dose group (*Figure 2A*,2*D*,2*E*).

The results suggested that the microvascular injuries of the colon mucosa in rats treated with NGR2 aggravated the hypoxia in the colon mucosa, and might be related to the increased serum ratio of VEGFA165/VEGFA121 and the protein expression of Rap1GAP and TSP1.

NGR2 promoted intracellular glycolysis and inhibited the cell proliferation and cell viability of pHUVECs

To clarify the mechanism of NGR2 in inducing vascular injury in colon tissues, the effect of NGR2 on regulating the cell viability of pHUVECs was measured at 24 and 48 h. NGR2 (from 0 to 50 nM) inhibited the cell viability in a concentration-dependent manner (*Figure 3A*). The IC50 of NGR2 was 25.94 nM at 24 h. Therefore, in the following experiments, the concentrations used to treat pHUVECs were 20.0, 10.0, and 5.0 nM. Compared with the control group, the cell viability and the cell proliferation of the groups treated with NGR2 (10.0 and 20.0 μM) were significantly inhibited at 24 h (*Figure 3B,3C*).

In addition, the concentrations of intracellular pyruvate and lactic acid were significantly increased in pHUVECs in the high-dose group and medium-dose group (*Figure 3D,3E*).

The results showed that NGR2 could inhibit the cell viability and proliferation of pHUVECs and promote intracellular glycolysis.

NGR2 inhibited the cell proliferation and tube formation of pHUVECs induced by VEGFA165

VEGFA165, one of the VEGFA isoforms that promotes the cell proliferation of pHUVECs, was used to explore the mechanism of NGR2 in inhibiting angiogenesis. As shown in *Figure 4A* and *Figure 4B*, cell viability and proliferation were increased by VEGFA165 (1.0 nM). When pHUVECs were co-treated with VEGFA165 and NGR2 (10.0 nM), cell viability and proliferation were significantly inhibited. In addition, the number of tube formations also obviously decreased in the VEGFA165 + NGR2 group compared to the VEGFA165 group. However, there was no significant difference in the tube formation between the NGR2 group and the VEGFA165 + NGR2 group (*Figure 4C,4D*).

The results demonstrated that NGR2 may counteract the role of VEGF in the regulation of cell viability and proliferation.

NGR2 inhibited the cell proliferation of pHUVECs via targeting Rap1GAP

To explore the role of *Rap1GAP* in regulating angiogenesis, a *Rap1GAP* interference plasmid was used. The expression of the *Rap1GAP* gene in pHUVECs was up-regulated by

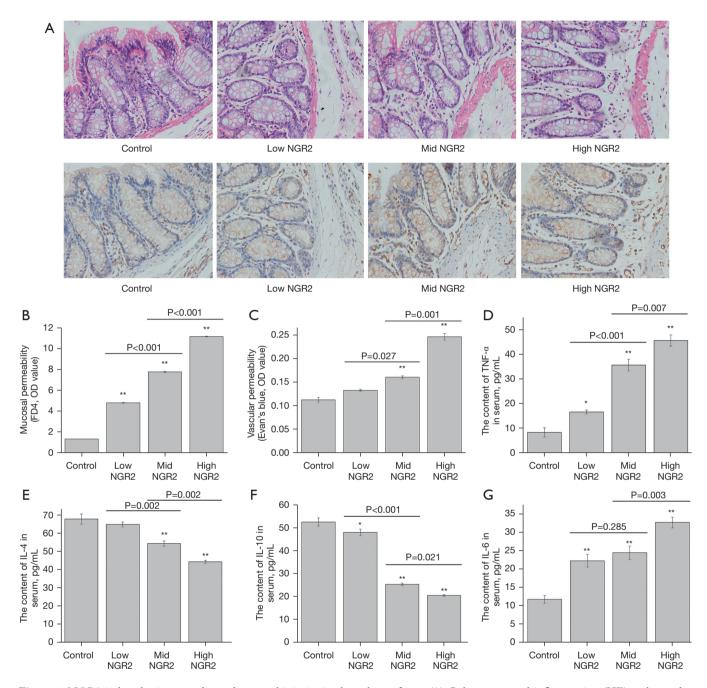


Figure 1 NGR2 induced microvascular and mucosal injuries in the colons of rats. (A) Colonic mucosal inflammation (HE) and vascular injury (400×); (B,C) Vascular permeability and mucosal permeability in the colonic mucosa; (D-G) the serum contents of IL-6, TNF- α , IL-4, and IL-10. The data are means \pm SEM of 3 independent experiments performed in triplicate. One-way ANOVA and Student's *t*-test (*P<0.05, **P<0.001 *vs.* control).

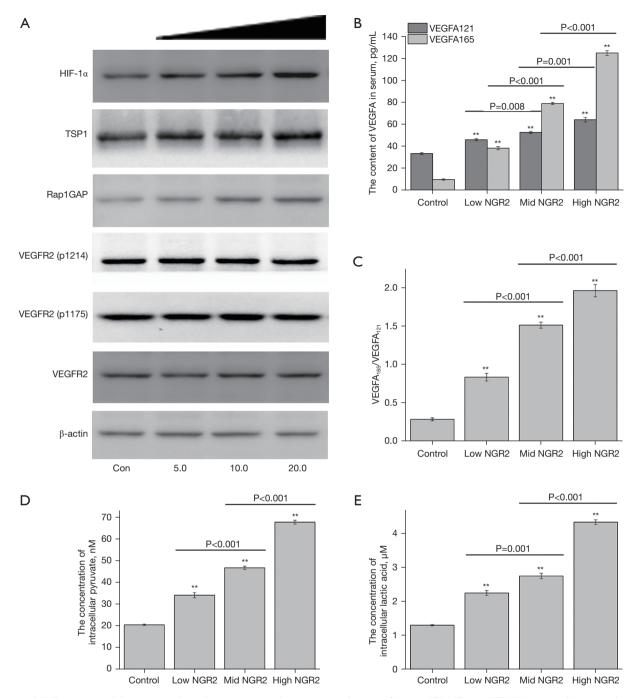


Figure 2 NGR2 increased hypoxia in the colon mucosa with an increased ratio of serum VEGFA165/VEGFA121 and increased protein expression of Rap1GAP and TSP1. (A) The protein expression levels of VEGFR2, VEGFR2 (p1175), VEGFR2 (p1214), Rap1GAP, TSP1, and HIF-1α in the colon mucosa; (B,C) the serum contents of VEGFA165 and VEGFA121 and the ratio of VEGFA165/VEGFA121; (D,E) the concentration of intracellular pyruvate and lactic acid in the colonic mucosa. The data are means ± SEM of 3 independent experiments performed in triplicate. One-way ANOVA and Student's *t*-test (**, P<0.001 *vs.* control).

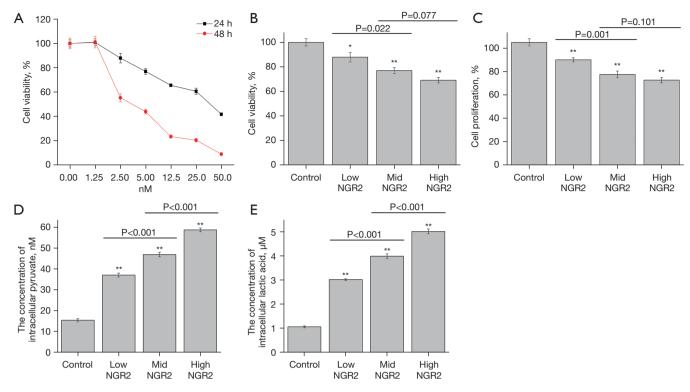


Figure 3 NGR2 promoted intracellular glycolysis and inhibited the cell proliferation and cell viability of pHUVECs. (A) The effect of NGR2 on regulating the cell viability of pHUVECs was measured at 24 and 48 h (IC50 of NGR2 was 25.94 nM at 24 h); (B,C) the cell viability and cell proliferation of pHUVECs treated with NGR2; (D,E) the concentration of intracellular pyruvate and lactic acid in pHUVECs. The data are means ± SEM of 3 independent experiments performed in triplicate. One-way ANOVA and Student's *t*-test (*P<0.05, **P<0.001 *vs.* control).

the *Rap1GAP* overexpression plasmid (*Rap1GAP*^{ver} plasmid, *Rap1GAP-ov*) (*Figure 5A*). Furthermore, VEGFA165 could down-regulate the protein expression of Rap1GAP in pHUVECs (*Figure 5B*). Meanwhile, the cell viability and cell proliferation of pHUVECs increased in the VEGFA165 group compared with the control group (*Figure 4A*,4*B*).

The protein expression of Rap1GAP in pHUVECs in the groups treated with NGR2 (10.0 nM), VEGFA165 (1.0 nM) + NGR2, VEGFA165 (1.0 nM) + Rap1GAP-ov, and Rap1GAP-ov + NGR2 + VEGFA165 was significantly up-regulated compared with the control and VEGFA165 group, and the expression of TSP1 was also up-regulated. Additionally, the expression of TSP1 in the group treated with VEGFA165 was also down-regulated compared with the control group (Figure 5B).

Meanwhile, cell viability and cell proliferation in the groups treated with NGR2 (10.0 nM), VEGFA165 (1.0 nM) + NGR2, VEGFA165 (1.0 nM) + Rap1GAP-ov, Rap1GAP-ov + NGR2 + VEGFA165, and Rap1GAP-ov

were inhibited compared with the control and VEGFA165 treatment group, with increased concentrations of intracellular pyruvate and lactic acid. There were no significant differences in cell viability, cell proliferation, and the concentrations of intracellular pyruvate and lactic acid between the groups treated with VEGFA165 + Rap1GAP-ov + NGR2 and Rap1GAP-ov + VEGFA165 or between the groups treated with VEGFA165 + NGR2 and NGR2 (Figure 5C-5F).

The *Rap1GAP* silencing plasmid (*Rap1GAP*^{si} plasmid, *Rap1GAP-si*) was used to silence the expression of *Rap1GAP* in pHUVECs to explore the mechanism of NGR2 in regulating angiogenesis. We found that *Rap1GAP-si* 3 could silence *Rap1GAP* expression (*Figure 5G*). In the following experiments, *Rap1GAP-si* 3 was chosen as the silencing plasmid.

Furthermore, the protein expression levels of Rap1GAP and TSP1 in pHUVECs in the groups treated with Rap1GAP-si, NGR2 + Rap1GAP-si, and VEGFA165 +

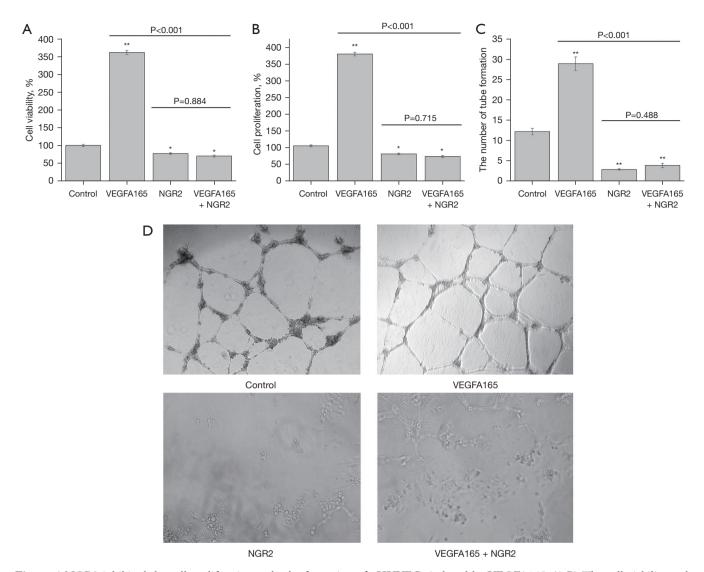


Figure 4 NGR2 inhibited the cell proliferation and tube formation of pHUVECs induced by VEGFA165. (A,B) The cell viability and proliferation of pHUVECs; (C,D) the tube formation of pHUVECs (100x). The data are means ± SEM of 3 independent experiments performed in triplicate. One-way ANOVA and Student's *t*-test (*P<0.05, **P<0.001 *vs*. control).

NGR2 + Rap1GAP-si were significantly down-regulated compared with the control group (Figure 6A). In addition, cell viability and cell proliferation in the groups treated with Rap1GAP-si, NGR2 + Rap1GAP-si, and VEGFA165 + NGR2 + Rap1GAP-si were significantly increased compared to the corresponding group untreated with Rap1GAP-si, while the concentrations of intracellular pyruvate and lactic acid decreased (Figure 6B-6E). There were no significant differences in cell viability, cell proliferation, and the concentrations of intracellular pyruvate and lactic acid between the two groups treated with Rap1GAP-si (Figure 6B-6E).

The results demonstrated that NGR2 inhibited the cell proliferation of pHUVECs via targeting Rap1GAP.

NGR2 blocked the Rap1GAP/PI3K/Akt signaling pathway

The *Rap1GAP* silencing plasmid and overexpression plasmid were transfected into pHUVECs to explore whether NGR2 regulated angiogenesis via the Rap1GAP and PI3K/Akt signaling pathway. Compared with the control group, Rap1GAP proteins down-regulated by VEGFA165 were up-regulated in the NGR2 (10.0 nM) group, *Rap1GAP-ov* group, and *Rap1GAP-ov* + NGR2 group. Meanwhile, the

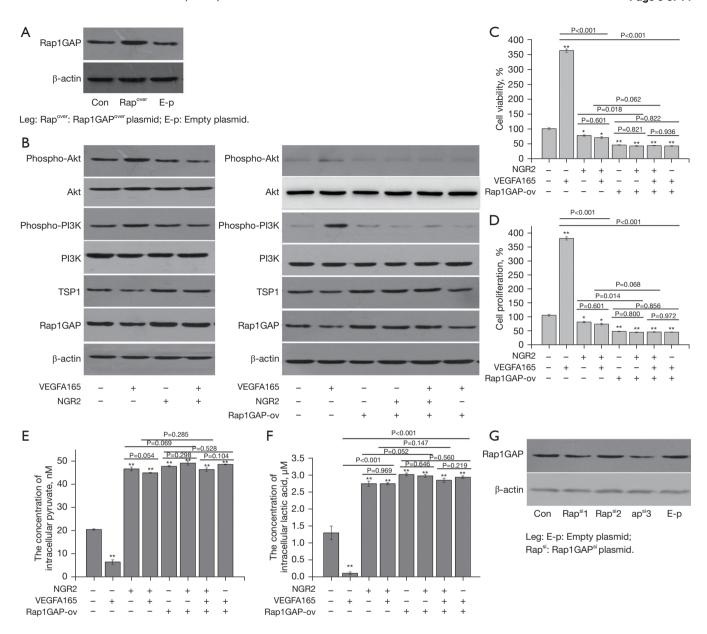


Figure 5 Overexpression of Rap1GAP and NGR2 inhibited the cell proliferation of pHUVECs. (A) The gene expression of Rap1GAP in pHUVECs was up-regulated by Rap1GAPover plasmid; (B) the protein expression of Rap1GAP, TSP1, PI3K, phospho-PI3K, Akt, and phospho-Akt in pHUVECs treated with VEGFA165, NGR2, and Rap1GAPover plasmid; (C-F) the cell viability, cell proliferation, and concentration of intracellular pyruvate and lactic acid in pHUVECs treated with VEGFA165, NGR2, and Rap1GAPover plasmid; (G) the expression of Rap1GAP in pHUVECs was down-regulated by Rap1GAPsi plasmid; The data are means ± SEM of 3 independent experiments performed in triplicate. One-way ANOVA and Student's *t*-test (*P<0.05, **P<0.001 *vs*. control).

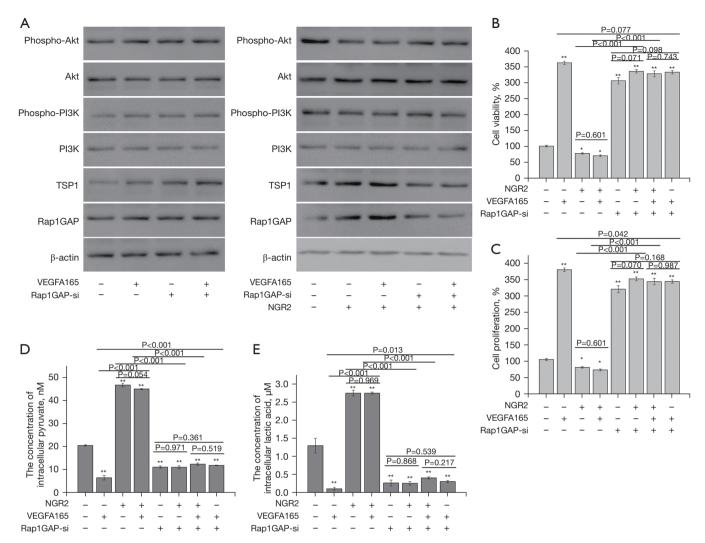


Figure 6 Silencing Rap1GAP promoted the cell proliferation of pHUVECs. (A) The protein expression of Rap1GAP, TSP1, PI3K, phospho-PI3K, Akt, and phospho-Akt in pHUVECs treated with VEGFA165, NGR2, and Rap1GAPsi plasmid; (B-E) the cell viability, cell proliferation, and concentration of intracellular pyruvate and lactic acid in pHUVECs treated with VEGFA165, NGR2, and Rap1GAPsi plasmid. The data are means ± SEM of 3 independent experiments performed in triplicate. One-way ANOVA and Student's *t*-test (*P<0.05, **P<0.001 *vs.* control).

protein expression levels of phospho-PI3K and phospho-Akt were significantly down-regulated by Rap1GAP1-ov compared with the control group and the empty plasmid (E-p) group, and were up-regulated by Rap1GAP-si (Figure 5B and Figure 6A). However, there were no differences in the protein expression of total PI3K and Akt compared with the control group. In addition, the protein expression of phospho-Akt could be down-regulated in the NGR2, NGR2 + LY294002, and LY294002 groups. Meanwhile, there were no differences in the protein expression of phospho-Akt found between the

NGR2 + LY294002 group and LY294002 group. The cell proliferation of pHUVECs was inhibited with the down-regulation of phospho-Akt protein expression (*Figure 7A*). Furthermore, to explore the regulatory effect of Rap1GAP on the PI3K/Akt signaling pathway, we found that the upregulated phospho-PI3K and phospho-Akt proteins in the *Rap1GAP-si* group and VEGFA165 group were significantly decreased by LY294002. In addition, there was no difference in the protein expression of phospho-PI3K and phospho-Akt between the LY294002 group and LY294002 + *Rap1GAP-ov* group (*Figure 7B*).

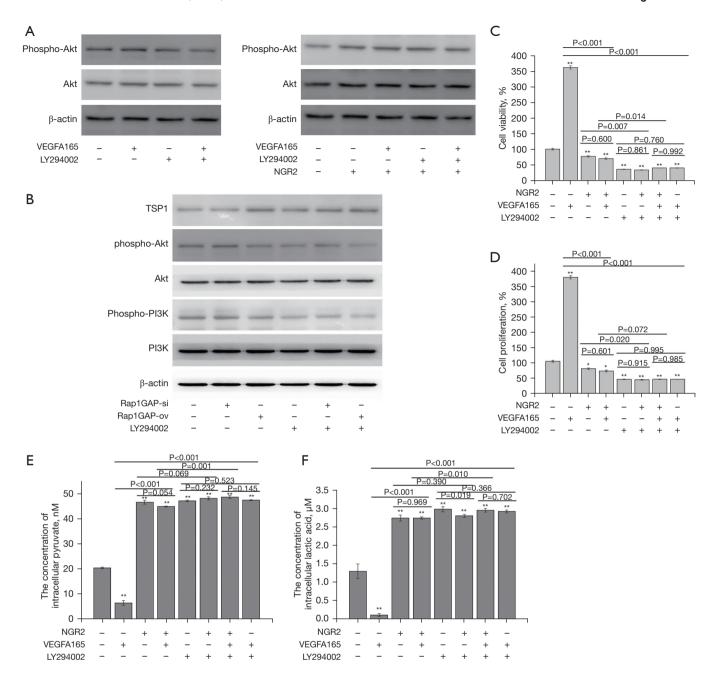


Figure 7 NGR2 blocked the Rap1GAP/PI3K/Akt signaling pathway. (A) The protein expression of phospho-Akt and Akt in the NGR2, NGR2 + LY294002, and LY294002 groups; (B) the expression of proteins in the PI3K/Akt signaling pathway could be regulated by Rap1GAPover plasmid and Rap1GAPsi plasmid, and could also be blocked by LY294002; (C-F) the cell viability, cell proliferation, and concentration of intracellular pyruvate and lactic acid of pHUVECs. The data are means ± SEM of 3 independent experiments performed in triplicate. One-way ANOVA and Student's *t*-test (*P<0.05, **P<0.001 *vs.* control).

The cell viability and cell proliferation of pHUVECs in the *Rap1GAP-si* group and VEGFA165 group were significantly inhibited by LY294002, while the concentration of intracellular pyruvate and lactic acid increased (*Figure 7C-7F*).

Discussion

PN has been shown to promote angiogenesis, regulate metabolism, and reduce the inflammatory response (17). Our previous research also found that PN could significantly promote angiogenesis and reduce vascular inflammatory injuries in colitis (3,4). Indeed, some saponins in PN, which are important active chemical components including ginsenosides and notoginsenosides, have appeared to lack toxicity in both animals and humans (5). Therefore, PN is regarded as a health-preserving drug that can protect the structure and function of blood vessels in daily life. PN is a traditional medicine that has the effect of promoting angiogenesis and antiangiogenesis. However, the anti-angiogenesis effect and active ingredients of PN have been seriously neglected in studies.

Notoginsenoside is the main active component of PN which regulates angiogenesis and inflammation in vivo and vitro (18,19). Among them, NGR2 is an important active component in PN (20). Hu et al. found that NGR2 could prevent neuronal apoptosis and inflammation via miR-27a/ SOX8/-catenin axis in Alzheimer's disease (8). However, the effect of NGR2 on angiogenesis is still unclear. Our results showed that NGR2 could inhibit the cell viability, cell proliferation, and tubulogenesis of pHUVECs and induce colonic mucosal and microvascular injuries in the colons of SD rats. Meanwhile, the protein expression of HIF-1α and the content of intracellular pyruvate and lactic acid were significantly increased in the colonic mucosa of rats treated with NGR2. Therefore, NGR2 could be a negative regulator of angiogenesis and enhance hypoxia in the colonic mucosa.

Angiogenesis is a complex physiological and pathological process. It is well known that VEGFA, an important subtype of VEGF, plays an important role in regulating the cell proliferation and tube formation of VECs (21). In addition, VEGFA consists of many isoforms, such as VEGFA121 and VEGFA165, among others (13). The results of previous studies showed that the abnormal ratio of VEGFA165/VEGFA121 is closely correlated with vascular injuries in a variety of diseases, such as

ulcerative colitis and cancers (12,13). VEGFR2 is the major receptor implicated in process of regulating cell viability, cell proliferation, cell migration, and VP stimulated by VEGFA (22). Our previous study found that the abnormal ratio of VEGFA165/VEGFA121 was closely correlated with colonic mucosal microvascular injuries in UC rats, and the activation of different phosphorylation sites of VEGFR2 led to VEGFA121 and VEGFA165 exerting opposite effects on angiogenesis (12). Besides, VEGFA also plays an important role in regulating glucose metabolism in VECs. VEGFA could reduce intracellular glycolysis and the protein expression of HIF-1 α in HUVECs (23). In the present study, after being treated with NGR2 for 7 days, the contents of VEGFA165 and VEGFA121 increased significantly, accompanied by the abnormal ratio of VEGFA165/VEGFA121 in rats treated with NGR2, and colonic mucosal and microvascular injuries also increased as hypoxia increased. It was also found that NGR2 could significantly inhibit the effects of VEGFA on regulating the cell proliferation and tubulogenesis of HUVECs and increase the intracellular glycolysis, which could cause microvascular injuries and hypoxia in tissue.

The PI3K/Akt signaling pathway is an important signaling pathway which has been shown to regulate cell proliferation, metastasis, and differentiation, among other processes (24). In addition, related studies found that VEGFA could promote angiogenesis via activating the PI3K/Akt signaling pathway (25). In this study, LY294002 was used as a PI3K/Akt signaling pathway inhibitor to investigate the mechanism of NGR2 in inhibiting angiogenesis. We found that NGR2 was an analog of the PI3K inhibitor which inhibited protein expression of phospho-Akt, and was similar to LY294002. Meanwhile, the cell viability and cell proliferation of pHUVECs were inhibited by LY294002 and NGR2. In addition, the signal transduction of the PI3K/Akt signaling pathway could be regulated by multiple signaling molecules, such as VEGFR2, PTEN, and Rap1GAP, among others (26,27). Of these, overexpression of Rap1GAP1, a GTPase-activating protein inhibiting Rap1 expression, effectively regulated angiogenesis via inhibiting cell proliferation, cell migration, tubulogenesis, and integrin-dependent adhesion (12,28). Besides, TSP1, a major protein driving endothelial cell senescence, could also inhibit angiogenesis directly by interacting with VEGF or indirectly by combining with TSP1 receptors on endothelial cells (29). The present study found that NGR2 could significantly up-regulate the protein expression of Rap1GAP and TSP1 in the colonic

mucosa of rats and pHUVECs, and enhanced intracellular glycolysis and hypoxia. Furthermore, the effect of NGR2 on anti-angiogenesis could be recovered via down-regulating the expression of Rap1GAP in pHUVECs, which could inhibit the PI3K/Akt signaling pathway as well as LY294002. Overall, NGR2 inhibited angiogenesis and enhanced intracellular glycolysis via activating the Rap1GAP/PI3K/Akt signaling pathway.

In summary, NGR2 was an analog of the Rap1GAP inhibitor. It could inhibit the cell proliferation, tube formation, and glucose aerobic oxidation of HUVECs via regulating the Rap1GAP/PI3K/Akt signaling pathway. However, there are still some deficiencies in this study. For example, there is a lack of *in vivo* verification of the regulatory effects of NGR2 on angiogenesis using RapGAP gene intervention animal models. Additionally, although NGR2 could promote the intracellular glycolysis of HUVECs, the mechanism remains unclear.

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