Abstract. Breast cancer is a leading cause of cancer-associated mortality globally amongst gynecologic tumors due to aggressive metastasis. A previous study reported that neurensin-2 (NRSN2) was implicated in human cancer cells, and that NRSN2 gene and protein expression levels were significantly upregulated in human breast cancer tissues compared with adjacent non-tumor tissues. The purpose of the present study was to analyze the role of NRSN2 in the metastasis of breast cancer cells and explore its potential mechanism. Reverse transcription-quantitative PCR, MTT, western blotting and immunohistochemistry was used to analyze the role of NRSN2 both in vitro and in vivo. The present study demonstrated that NRSN2 knockdown inhibited the proliferation, migration and invasion of breast cancer cells in vitro. NRSN2 upregulation promoted breast cancer cell proliferation and tissue growth in vitro and in vivo. In addition, the results demonstrated that the regulatory effects of NRSN2 on breast cancer cells were associated with PI3K/AKT/mTOR and NF-κB signaling pathways. Furthermore, NRSN2 overexpression in mice significantly promoted breast cancer cell proliferation. In conclusion, the results from the present study indicated that NRSN2 may be considered as a novel oncogenic protein and may represent a potential therapeutic target for breast cancer.

Introduction

Breast cancer is one of the most common gynecologic tumors, mainly occurring in the elderly (1). A previous review reported the most common genes involved in epigenetic modifications in patients with breast cancer (2). Most breast cancer-associated mortalities are caused by local breast cancer cell migration and distant metastasis (3). A review reported genetic analyses and inherited gene mutations in patients with breast cancer (4,5). Although advances in molecular diagnosis and medical treatments, including surgical techniques, radiation, chemotherapy and gene target therapy, have improved the 5-year survival rate of patients with breast cancer, the overall clinical outcomes remain poor (6-9). It is therefore essential to determine potential target proteins to inhibit breast cancer growth and metastasis.

Neurensin-2 (NRSN2) is a small neuronal membrane protein that is localized in small vesicles of neural cells (10). A previous study revealed that NRSN2 can promote non-small cell lung cancer cell proliferation via the PI3K/AKT/mTOR pathway (11). Tang et al (12) reported that NRSN2 overexpression was associated with malignant phenotypes in ovarian cancer, suggesting that it could be considered as a target for ovarian cancer treatment. However, Wang et al (13) demonstrated that NRSN2 upregulation inhibited cell proliferation and survival via the PI3K/AKT/mTOR pathway in hepatocellular carcinoma. These findings encouraged the present study to further investigate the role of NRSN2 in breast cancer cells.

The PI3K/AKT/mTOR, p65 and NF-κB signaling pathways contribute to breast cancer progression. Therefore, the present study hypothesized that NRSN2 may regulate breast cancer cell proliferation via the PI3K/AKT/mTOR and NF-κB pathways (14). The results from the present study demonstrated that NRSN2 overexpression significantly increased the proliferation, invasion and metastasis of breast cancer cells, suggesting that NRSN2 may be considered as a potential therapeutic target for breast cancer treatment via downregulation of the PI3K/AKT/mTOR and NF-κB signaling pathways.

Materials and methods

Ethical statements. This study was conducted in strict accordance with the recommendations of the Guide for the Care and Use of Laboratory Animals of the Tianjin Medical University. The protocol was approved by the Chinese Association for Laboratory Animal Operations. All surgery and euthanasia were performed under sodium pentobarbital anesthesia (intravenous, 35 mg/kg). Mice were sacrificed via cervical decapitation.

Patients and tissues. A total of 24 patients with breast cancer were recruited in Peking University between May 2015
and October 2016. Their average age was 54.5±24.5 years (range, 30-79 years). Breast cancer and adjacent noncancerous tissues were obtained from patients who underwent tumor resection and stored at -80°C prior to immunohistochemistry (IHC) and reverse transcription-quantitative polymerase chain reaction (RT-qPCR) analyses. Patients who had previously undergone radiotherapy, chemotherapy or administration of any other drug were excluded from this study. All patients provided written informed consent prior to any procedures of this study. The patient study was approved by the Ethics Committee of Peking University (approval no. PEK20150524).

Cell line, chemicals and reagents. The breast tumor cell lines MDA-MB-231 and BT549, and the normal breast cell line MCF-10A were purchased from the American Type Culture Collection. All cells were cultured in Dulbecco's modified Eagle's medium (DMEM; Invitrogen; Thermo Fisher Scientific, Inc.) containing 10% fetal bovine serum (Invitrogen; Thermo Fisher Scientific, Inc.) containing 10% fetal bovine serum (Invitrogen; Thermo Fisher Scientific, Inc.) or PBS as control for NRSN2 expression levels measured in an iCycler thermal reaction (RT-qPCR) analyses. The thermocycling conditions were: 95°C for 120 sec; followed by 45 cycles at 95°C for 120 sec; 56°C for 20 sec and 65°C for 30 sec. The RT-qPCR kit (Invitrogen; Thermo Fisher Scientific, Inc.) was used for RT-qPCR. MDA-MB-231 and BT549, and the normal breast cell line MCF-10A were purchased from the American Type Culture Collection. All cells were cultured in Dulbecco's modified Eagle's medium (DMEM; Invitrogen; Thermo Fisher Scientific, Inc.) containing 10% fetal bovine serum (Invitrogen; Thermo Fisher Scientific, Inc.) and placed at 37°C in a humidified incubator containing 5% CO₂. Cells were treated with the NF-κB inhibitor caffeic acid phenethyl ester (CAPE; 70 mM; Apex Biotechnology Corp.), PI3K inhibitor (cat. no. P0126; Beyotime Institute of Biotechnology) according to manufacturer's protocol. The efficiency of siR-NRSN2 transfection was verified via western blotting 12 h following transfection, prior to subsequent experiments. siR-NRSN2 and siR-NC transfection was verified via western blotting 12 h following transfection, prior to subsequent experiments.

Small interfering RNA (siRNA) transfection. All siRNAs (si-R-NRSN2, 5′-CAA TCT TCT GTG CAG ACT ATC-3′; si-R-NRSN2, 5′-CAATCTTCTGTCGACTATC-3′; si-R-NRSN2, 5′-CAATCTTCTGTCGACTATC-3′; si-R-NRSN2, 5′-CAATCTTCTGTCGACTATC-3′) were synthesized by Invitrogen; Thermo Fisher Scientific, Inc. MDA-MB-231 cells (1x10⁴ cells/well) were seeded in six-well plates and transfected with 150 µM si-R-NRSN2 or si-control using a Cell Line Nucleofector kit (cat. no. VCA-1003; Lonza Group, Ltd.) according to manufacturer's protocol. The efficiency of si-R-NRSN2 transfection was verified via western blotting at 72 h following transfection, prior to subsequent experiments.

NRSN2 overexpression. An expression plasmid (pRK5-hNRSN2) with a Flag tag at the C-terminus was constructed by Invitrogen (Thermo Fisher Scientific, Inc.). MDA-MB-231 cells (1x10⁴) were seeded in six-well plates (Corning Inc.) and transiently transfected with pRK5-hNRSN2 (2 µg) or pRK5-control (pControl) (2 µg) using Lipofectamine™ 2000 (cat. no. 11668-027; Invitrogen; Thermo Fisher Scientific, Inc.) according to the manufacturer's protocol. The efficiency of NRSN2 overexpression was verified by western blotting at 72 h following transfection, prior to subsequent experiments.

RT-qPCR. Total RNA was isolated from tissues or cells by using an RNAeasy Mini kit (Qiangen, Inc.). The expression of NRSN2 in tissues and cells was measured using a Hairpin-it™ RT-qPCR kit (Invitrogen; Thermo Fisher Scientific, Inc.). NRSN2 expression levels were measured in an iCycler thermal cycler (Bio-Rad Laboratories, Inc.) using SYBR Green Supermix (Bio-Rad Laboratories, Inc.). The thermocycling conditions were: 95°C for 120 sec; followed by 45 cycles at 95°C for 20 sec and 65°C for 30 sec. The primers were designed as follows: NRSN2, forward 5′-GGG AGACGCAGGTCCAGAGGAT-3′; reverse 5′-TATGCATACTGTATTTAGGAAGG-3′; and β-actin, forward 5′-GTG GGGCAGGAGCCACCA-3′ and reverse 5′-CTCCCTAAT GTCACGGCAGATT-3′. Relative mRNA expression changes were calculated using the 2-ΔΔCq method (16). The results are expressed compared to β-actin expression.

Cell proliferation assay. Cell proliferation was determined using a Cell Counting Kit-8 assay (CCK-8; Dojindo Molecular Technologies, Inc.) according to the manufacturer's instructions. Briefly, pRK5-hNRSN2 or SiR-NRSN2-transfected MDA-MB-231 cells and their controls were seeded in 96-well plates at a density of 1x10⁴/well and cultured at 37°C in a 5% CO₂ atmosphere for 48 h. CCK-8 solution (10 µl) was added to each well for 2 h. The cell proliferation was monitored by measuring the absorbance at 450 nm using a microplate reader (Thermo Fisher Scientific, Inc.).

Colonies formation assay. For the colony formation assay, pRK5-hNRSN2 or SiR-NRSN2-transfected MDA-MB-231 cells were seeded in 6-well plates at a density of 1x10⁴/well and cultured at 37°C in a 5% CO₂ atmosphere for 7 days until visible colonies were formed. Cells were stained with 5% crystal violet for 10 min at room temperature. The numbers of colonies were then counted using a light microscope at x20 magnification.

Cell migration and invasion assay. MDA-MB-231 cells were transfected with pRK5-hNRSN2 or si-R-NRSN2. Matrigel-coated and -coated migration inserts (8-µm pore size; Corning Inc.) were used for migration and invasion assays, respectively.

For the migration assay, MDA-MB-231 cells (1x10⁴) in DMEM were plated into the upper chamber with the non-coated membrane. For the invasion assay, pRK5-hNRSN2 or si-R-NRSN2-transfected cells were prepared at a density of 1x10⁵ cells in 500 µl serum-free DMEM in the upper chamber and the lower chamber contained DMEM with 5% FBS. Cells were seeded in the upper chamber of a BD BioCoat Matrigel Invasion Chamber (BD Biosciences) according to the manufacturer's instructions. Following incubation for 48 h, cells were fixed with 4% paraformaldehyde for 1 h at room temperature and stained with 0.1% crystal violet (Sigma-Aldrich; Merck KGaA) for 10 min at 37°C. The membranes were mounted onto a glass slide with antifade mounting medium (cat. no. P0126; Beyotime Institute of Biotechnology), and the number of migrating and invading tumor cells was counted in at least three randomly selected fields under a light microscope (Olympus Corporation) at x200 magnification.

Western blotting. MDA-MB-231 cells were transfected with pRK5-hNRSN2 or si-R-NRSN2. Following transfection, cells were lysed using radioimmunoprecipitation assay buffer (Sigma-Aldrich; Merck KGaA) containing protease-inhibitor (Sigma-Aldrich; Merck KGaA) and were centrifuged at 12,000 x g, at 4°C for 10 min. Breast cancer tissues (10 mg) were lysed using radioimmunoprecipitation assay buffer containing protease inhibitor and were centrifuged at 1,000 x g at 4°C for 10 min. Supernatant was collected for protein analysis. The concentrations of protein were measured using a BCA Protein Concentration Assay kit (Beyotime Institute of Biotechnology). Protein samples (50 µg per lane) were separated by sodium...
dodecyl sulfate-polyacrylamide electrophoresis (SDS-PAGE) on a 10% gel and transferred to polyvinylidene difluoride (PVDF; Bio-Rad Laboratories, Inc.) membrane. Membranes were incubated with rabbit anti-human primary antibodies against NRSN2 (1:1,000; cat. no. ab237739; Abcam), PI3K (1:1,000; cat. no. ab32089; Abcam), phosphorylated (p)-PI3K (1:1,000; cat. no. ab154598; Abcam), p65 (1:1,000; cat. no. ab16502; Abcam), p-p65 (1:1,000; cat. no. ab86299; Abcam), IκBα (1:1,000; cat. no. ab7217; Abcam) and p-IκBα (1:1,000; cat. no. ab133462; Abcam), β-actin (1:5,000; cat. no. 20536-1-AP; ProteinTech Group, Inc.), AKT (1:1,000; cat. no. 51077 -1-AP; ProteinTech Group, Inc.), p-mTOR (1:1,000; cat. no. ab2731; Abcam), and mTOR (1:1,000; cat. no. ab109268; Abcam), overnight at 4°C. The membranes were subsequently incubated with HRP-conjugated anti-rabbit IgG secondary antibodies (diluted 1:5,000; cat. no. A9169; Sigma-Aldrich; Merck KGaA) for 24 h at 4°C. Enhanced chemiluminescence reagent (Millipore; Merck KGaA) was used to visualize the bands. Quantitation of the signal intensities were analyzed using the Quantity One software package (version 2.0; Bio-Rad Laboratories, Inc.).

Animal study. Pathogen-free female Balb/c (8-week-old; 20-25 g body weight) nude mice were purchased from Slack...
Figure 2. NRSN2 knockdown suppresses and NRSN2 overexpression increases the proliferation of breast cancer cells in vitro. (A) NRSN2 knockdown decreased NRSN2 expression in MDA-MB-231 cells. (B) NRSN2 overexpression increased NRSN2 expression in MDA-MB-231 cells. (C) NRSN2 knockdown inhibited MDA-MB-231 cell proliferation. (D) NRSN2 overexpression promoted MDA-MB-231 cell proliferation. (E) NRSN2 knockdown inhibited the colony formation of MDA-MB-231 cells. (F) NRSN2 overexpression promoted the colony formation of MDA-MB-231 cells. NC, negative control; NRSN2, neurensin-2; ns, not significant; si, small interfering (RNA); siR-NRSN2, siRNA against NRSN2; pRK5-NRSN2, NRSN2 overexpression vector. *P<0.01.
All mice were treated in accordance with the China Legislation on the Protection of Animals and the Guide for the Care and Use of Laboratory Animals. The study was approved by the ethics committees of Shanxi Traditional Chinese Medical University. Animals were housed in a temperature-controlled facility at 23±1˚C with 50±5% humidity under a 12-h light/dark cycle. All mice had free access to food and water. The siR-NRSN2-transfected MDA-MB-231, pRK5-control vector-transfected MDA-MB-231 cells or MDA-MB-231 cells (1x10^7) were subcutaneously injected into the flanks of female Balb/c mice. The tumor volume was calculated every 3 days and calculated as Volume = (D x d^2)/2 (D represents the maximal diameter, d represents the minimal one). The mice were sacrificed on day 30 following anesthesia.

**Human breast tissues or mouse cancer tissues were analyzed for NRSN2 expression using IHC, as previously described (17). Briefly, tumor tissue samples were fixed in 10% formalin for 12 h at room temperature and embedded in paraffin blocks for 8 h at room temperature. Thin tissue sections (4-µm thick) were de-waxed and rehydrated. Tissue sections were immersed for 15 min at room temperature in 0.3% H_2O_2 (diluted with 100% methanol) to block endogenous peroxidase activity. The tumor sections were incubated with specific primary antibodies: Rabbit anti-human or mouse antibodies against rabbit anti-human/mouse NRSN2 (1:1,000; cat. no. ab237739; Abcam) overnight at 4˚C. Tumor tissues were then incubated with horseradish peroxidase-conjugated goat anti-rabbit immunoglobulin G secondary antibody (1:5,000; cat. no. PV-6001; OriGene Technologies, Inc.) for 2 h at room temperature. Finally, tissue sections were stained with 3-3'-diaminobenzidine tetrahydrochloride (Sigma-Aldrich; Merck KGaA) for 1 h at room temperature and lightly counter-stained by hematoxylin for 30 min at room temperature. A Ventana Benchmark automated staining system was used to detect NRSN2 protein expression in tumor tissues (Olympus BX51; Olympus Corporation) at x200 magnification. The quantification of NRSN2 density was analyzed using Image J software (version 4.6; National Institutes of Health).

**Statistical analysis.** All data were expressed as the mean ± standard deviation. Experiments were repeated at least three times. Statistical analyses were performed using Student's t-test or one-way ANOVA followed by Tukey's honestly significant difference post hoc test. Data were analyzed using SPSS Statistics 19.0 software (IBM Corp.) and GraphPad Prism 5.0 (GraphPad Software, Inc.). P<0.05 was considered to indicate a statistically significant difference.
Results

NRSN2 expression is elevated in breast cancer tissues and cell lines. The mRNA and protein levels of NRSN2 in breast cancer tissues and cell lines were determined. As presented in Fig. 1A and B, mRNA and protein levels of NRSN2 were significantly increased in breast cancer tissues compared with adjacent tissues. Furthermore, the results from IHC demonstrated that NRSN2 was highly expressed in breast cancer tissues compared with adjacent tissues (Fig. 1C). In addition, as presented in Fig. 1D and E, the mRNA and protein levels of NRSN2 were significantly increased in the breast cancer cell lines BT549 and MDA-MB-231 compared with the MCF-10A cell line. These results demonstrated that NRSN2 was highly expressed in breast cancer tissues and cell lines, which suggested that NRSN2 may have a role in breast cancer progression.

NRSN2 knockdown inhibits the proliferation of breast cancer cells in vitro. The role of NRSN2 in the breast cancer cell line MDA-MB-231 was further investigated. As presented in Fig. 2A and B, NRSN2 knockdown (siR-NRSN2) and overexpression (pRK5-NRSN2) significantly decreased and increased, respectively, NRSN2 expression in MDA-MB-231 cells. Furthermore, NRSN2 knockdown inhibited MDA-MB-231 cell proliferation (Fig. 2C), whereas NRSN2 overexpression promoted MDA-MB-231 cell proliferation (Fig. 2D). Colony formation assays demonstrated that NRSN2 overexpression increased the numbers of MDA-MB-231 cell colonies, whereas NRSN2 knockdown decreased the numbers of MDA-MB-231 cells colonies formed (Fig. 2E and F). These results demonstrated that NRSN2 knockdown could inhibit breast cancer cell proliferation in vitro.

NRSN2 promotes tumor cell migration and invasion in vitro. The effects of NRSN2 on cell motility were analyzed using...
migration and invasion assays. The results demonstrated that NRSN2 overexpression promoted the migration and invasion of MDA-MB-231 cells (Fig. 3A and B), whereas NRSN2 knockdown inhibited the migration and invasion of MDA-MB-231 cells (Fig. 3C and D).

NRSN2 promotes proliferation, migration and invasion of breast cancer cells by activating PI3K/AKT/mTOR and NF-κB signaling pathways. The potential molecular mechanisms mediated by NRSN2 were investigated in MDA-MB-231 cells. The results demonstrated that NRSN2 knockdown significantly inhibited the phosphorylation of PI3K, AKT and mTOR in MDA-MB-231 cells (Fig. 4A). Furthermore, NRSN2 overexpression significantly increased the phosphorylation of PI3K, AKT and mTOR in MDA-MB-231 cells (Fig. 4B). NRSN2 overexpression induced opposing effects (Fig. 4C and D). In addition, the NF-κB inhibitor CAPE (NF-κBIR) inhibited the pro-proliferation effects of NRSN2 (NF-κBIR) in MDA-MB-231 cells (Fig. 4E). Furthermore, NF-κBIR suppressed the NRSN2 overexpression-induced (NF-κBIR-NRSN2) migration and invasion of MDA-MB-231 cells (Fig. 4F and G). Treatment of NRSN2 overexpressing cells with PI3K inhibitor (PI3KIR-NRSN2) inhibited NRSN2 overexpression-induced proliferation, migration and invasion of MDA-MB-231 cells (Fig. 4H-J). However, PI3K inhibitor or NF-κB inhibitor induced no effects on the proliferation, migration or invasion of MDA-MB-231 cells. These results suggested that NRSN2 may regulate the proliferation and aggressiveness of MDA-MB-231 cells through PI3K/AKT/mTOR and NF-κB signaling pathways.

NRSN2 overexpression promotes tumor growth in vivo. The role of NRSN2 in tumor growth was further investigated in vivo in subcutaneous breast cancer xenograft nude mice. The results demonstrated that tumor growth was faster in nude mice injected with pRK5-NRSN2 plasmid-transfected MDA-MB-231 cells than in nude mice injected with pRK5-control vector (control; Fig. 5A and B). However,
injection of siRNA-NRSN2 plasmid-transfected MDA-MB-231 cells induced smaller tumor volume and weight compared with in nude mice injected with cells transfected pRK5-control vector (Fig. 5A and B). The results from western blotting and IHC demonstrated that NRSN2 protein expression was increased in pRK5-NRSN2 plasmid-transfected tumor tissues compared with in siRNA-NRSN2 plasmid-transfected tumor tissues (Fig. 5C and D). These results suggested that NRSN2 may promote tumor growth in vivo.

Discussion

Breast cancer is the most common female cancer worldwide. It is commonly diagnosed at advanced stages and exhibits
rising incidence and mortality rates (18). Previous studies have indicated that NRSN2 is highly expressed in numerous human cancer cells, including lung cancer and hepatocellular carcinoma, and may therefore be considered as a potential target for human cancer treatment (10,11,13). The results from the present study suggested that NRSN2 may serve an important role in the carcinogenesis and progression of breast cancer in vitro and in vivo. The results demonstrated that NRSN2 was highly expressed in breast cancer tissues and cells, and may therefore stimulate the progression of breast cancer and promote the proliferation of breast cancer cells. These findings also indicated that NRSN2 knockdown may inhibit the migration and invasion of breast cancer cells in vitro. Notably, NRSN2 knockdown potentially regulated human breast cancer proliferation via PI3K/AKT/mTOR and NF-κB signaling pathway inactivation.

Tumor markers have been widely used for the diagnosis of early stage breast cancer in patients (19-21). The present study demonstrated that NRSN2 was significantly upregulated in breast cancer tissues compared with adjacent noncancerous tissues. Although a previous study reported that NRSN2 downregulation promoted cell proliferation and survival via the PI3K/AKT/mTOR pathway in hepatocellular carcinoma (13), the results from the present study demonstrated that NRSN2 downregulation significantly inhibited breast cancer cell proliferation and aggressiveness. Notably, NRSN2 controlled human breast cancer proliferation via the regulation of PI3K/AKT/mTOR and NF-κB signaling pathways; however, NF-κB inhibitor or PI3K inhibitor had no effects on the proliferation, migration and invasion of MDA-MB-231 cells. These results suggested that NRSN2 may regulate the proliferation and aggressiveness of breast cancer cells via PI3K/AKT/mTOR and NF-κB signaling pathways.

Currently, the PI3K/AKT/mTOR signaling pathway is considered as a potential target for breast cancer therapy (22). A recent study reported that intermittent hypoxia induces the overexpression of prometastatic genes in breast cancer cells via NF-κB, such as tenascin-C (an essential factor of the metastatic niche) and matrix metalloproteinase 9, and induces pro-inflammatory processes, via cyclooxygenase-2 (COX-2) for example (23). According to previous studies (24-26), alterations of the PI3K/AKT/mTOR and NF-κB signaling pathways were investigated in breast cancer cell lines following NRSN2 overexpression or knockdown.

Shen et al (27) reported that the tumor volume in 6-week-old female NOD/SCID mice injected with wild-type MDA-MB-231 cells (1x10^6 cells in 30 µl PBS) was ~200 mm³ at day 22. In the present study, specific pathogen-free female Balb/c nude mice were injected with MDA-MB-231 cells transfected with pRK5-NRSN2- or pRK5-vector (control; 1x10^7 cells) or MDA-MB-231 cells to analyze the role of NRSN2 in breast cancer growth. The tumor volume of wild-type MDA-MB-231-injected mice at day 27 was ~2,100 mm³. The difference in the tumor volumes between the present study and the study by Shen et al (27) may be due to the types of mice used and the number of cells injected. The results from the present study demonstrated that NRSN2 stimulated PI3K/AKT/mTOR and NF-κB phosphorylation in MDA-MB-231 cells, which further promoted breast cancer cells growth both in vitro and in vivo. However, only 8 mice were used in this study to explore the inhibitory effects of NRSN2.
Further investigation using a larger sample size would therefore be needed to determine other signaling pathways associated with NRSN2. In addition, the present study did not use ideal controls when investigating the effects of NF-κB inhibitor or PI3K inhibitor on the proliferation, migration and invasion of breast cancer cells, as an overexpression-only group was not included. The effects of NF-κB inhibitor or PI3K inhibitor on NRSN2 expression will also be further analyzed in the future.

In conclusion, the results from the present study demonstrated that NRSN2 was overexpressed in breast cancer tissues and cells, and that it significantly promoted the proliferation and aggressiveness of breast cancer by activating PI3K/AKT/mTOR and NF-κB signaling pathways. These findings suggested that NRSN2 may be considered as a potential therapeutic target for breast cancer.

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Availability of data and materials

The datasets used and/or analyzed during the current study are available from the corresponding author on reasonable request.

Authors’ contributions

FR and WZ performed all experiments. SL and HR prepared for the experiments, analyzed and collected data. YG designed the study and wrote the manuscript. All authors read and approved the final manuscript.

Ethics approval and consent to participate

The animal study was approved by the Chinese Association for Laboratory Animal Operations. The patient study was approved by the Ethic Committee of Peking University (approval no. PEK20150524). All patients provided written informed consent prior to the study.

Patient consent for publication

Not applicable.

Competing interests

The authors declare that they have no competing interests.

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