# Daam1 activates RhoA to regulate Wnt5a-induced glioblastoma cell invasion

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Abstract. The signaling pathway of dishevelled-associated activator of morphogenesis 1 (Daam1) triggered by Wnt5a drives cell movement and migration during breast cancer metastasis. However, Wnt5a signaling in glioblastoma progression remains poorly defined. Wnt5a expression and activations of RhoA, Cdc42, and Rac1 were detected in human glioblastoma tissues by using ELISA assays and small G-protein activation assays, respectively. The cell invasion rate and Daam1 activation of glioblastoma U251 and T98MG cells were determined by cell invasion assays and pull-down assays, respectively. According to our experiments, Wnt5a expression and RhoA activation were upregulated in invasive glioblastoma tissues, with a significant positive correlation between them. Wnt5a activated Daam1 and RhoA, and subsequently promoted the invasion of glioblastoma U251 and T98MG cells. This process was abolished by secreted frizzled-related protein 2 (sFRP2), an antagonist that directly binds to Wnt5a. Specific small interfering RNA (siRNA) targeting Daam1 markedly inhibited Wnt5a-induced RhoA activation, stress fiber formation and glioblastoma cell invasion. CCG-1423, a RhoA inhibitor, decreased Wnt5a-induced stress fiber formation and glioblastoma cell invasion. Finally, siRNA targeting Daam1 or CCG-1423 treatment did not alter the cell proliferation of glioblastoma U251 and T98MG cells. We thus concluded that

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Wnt5a promoted glioblastoma cell invasion via Daam1/RhoA signaling pathway.

## Introduction

Glioblastoma, also known as glioblastoma multiforme (GBM), is the most aggressive primary tumor of the brain (1-3). Despite advances in surgical technology and adjuvant treatment for glioblastoma, this tumor remains the most lethal disease worldwide (2,3). The molecular mechanisms underlying the invasion and metastasis of glioblastoma are still unknown.

Formerly, Wnts have been divided into two classes: those that signal through canonical ( $\beta$ -catenin dependent) pathway and those that signal through non-canonical ( $\beta$ -catenin independent) pathway (4-6). Wnt signaling participates in the development of embryo and pathological processes, including tumorigenesis and metastasis (7). A homeobox transcription factor, MSX1 was able to inhibit the Wnt/ $\beta$ -catenin signaling pathway and suppress Wnt/ $\beta$ -catenin-induced migration and invasion of cultured glioblastoma cells (8). Inhibition of the Wnt/ $\beta$ -catenin pathway significantly abrogated the invasion effects of irradiation, indicating a pivotal role of the Wnt/ $\beta$ -catenin pathway in ionizing radiation-induced invasion of U87 cells (9).

Wnt5a is classified as a non-transforming Wnt family member that plays complex roles in cancer initiation and metastasis (10-12). Wnt5a activates small Rho-GTPases and regulates the cytoskeletal architecture and cellular polarity during development (13). Wnt5a signaling is a regulator in the proliferation of human glioma cells (14). However, very few studies have reported on the role of Wnt5a in glioblastoma cell invasion and migration (15,16). In the present study, we propose for the first time that Wnt5a promotes the invasion of glioblastoma cells and is upregulated in invasive glioblastoma tissues. Moreover, the mechanisms whereby the Wnt5a/Daam1/RhoA signaling pathway regulates glioblastoma cell invasion are described.

## Materials and methods

*Clinical samples*. Tissue samples of nine glioblastoma patients from the Jinan Fourth People's Hospital from 2015 to 2016 were recruited in the present study. All patients underwent surgical

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resection of the glioblastoma with the intention of maximally resecting the tumor. Glioblastoma tissues were frozen in liquid nitrogen or fixed in 10% formalin and embedded in paraffin wax. Slices were stained with hematoxylin and eosin (H&E) for light microscopy. The diagnosis of glioblastoma was based on the World Health Organization (WHO) 2007 and 2016 histopathologic criteria: the invasive phenotype showing an infiltrative astrocytic neoplasm with high proliferative activity and microvascular proliferation, necrosis or both (17,18). The non-invasive glioblastoma was thus called when the tumor at the anatomical site began to progress and further yielded a cancerous mass without the invasive phenotype aforementioned. All glioblastoma tissues with high tumor cell density were histopathologically confirmed by two pathologists prior to ELISA and small G-protein activation assay. Ethical approval of the study (no. LL-20140005) was granted by the Clinical Research Ethics Committee, Jinan Fourth People's Hospital. Written informed consent was obtained from each participant.

Cells and small interfering RNA (siRNA) transfection. U87MG, U251 and T98MG are the most used cell lines for research on human glioblastoma. The original U87MG cell line was established in Uppsala University almost 50 years ago (19). However, Allen et al used short tandem repeat (STR) genotyping to screen out the DNA profile of U87MG. Different from that of the original cells, this friendly profile of U87MG is thought to have an unknown origin (20). Thus, two cell lines (U251 and T98MG) were used in this experiment. Human glioblastoma U251 or T98MG cell lines were purchased from the Cell Bank of Shanghai (Shanghai, China) and were grown in Eagle's Minimum Essential Medium (EMEM; HyClone, Thermo Scientific, Waltham, MA, USA) supplemented with 10% (v/v) fetal bovine serum (FBS), 2 mmol/l L-glutamine and 100 IU/ml penicillin, 100 µg streptomycin, 1 mmol/l sodium pyruvate and non-essential amino acids (HyClone) in a humidified incubator at 37°C with 5% CO<sub>2</sub> and 95% humidity. The cells were seeded in 6-well plates (Costar, Corning, NY, USA) and cultured to 80% confluence, and then transiently transfected with siRNA against Daam1 (21) using Lipofectamine 2000 reagent (Invitrogen, Carlsbad, CA, USA) in serum-free Opti-MEM according to the manufacturer's instructions. The cells were switched to fresh medium containing 10% FBS 6 h after the transfection and cultured for 48 h. The cells transfected with Daam1-siRNA were used for analyzing Rho activation and cell invasion.

*ELISA*. The glioblastoma tissues were grinded in liquid nitrogen. Equal weights of total tissue debris were dissolved in ice-cold phosphate-buffered saline (PBS) buffer. The experiments were then performed according to the manufacturer's protocol of the Wnt5a ELISA kit (CusaBio, Wuhan, China). The concentration of each glioblastoma tissue was calculated based on the concentration curve of the Wnt5a standard samples.

*Cell invasion assays.* Cell invasion was assessed in modified Boyden chambers (Costar). Two chambers were separated by a polycarbonate membrane (pore diameter, 8.0  $\mu$ m). Boyden chamber wells were coated with Matrigel (BD Biosciences,



Figure 1. Wnt5a is upregulated in invasive glioblastoma tissues. The expression level of Wnt5a was highly expressed in invasive glioblastoma tissues (n=5) when compared to that in non-invasive glioblastoma tissues (n=4). N0, non-invasive glioblastoma; N, invasive glioblastoma.

Franklin Lakes, NJ, USA) for 30 min at 37°C. U251 or T98MG cells treated with CCG-1423 (Selleck, Houston, TX, USA) were added to wells with a membrane placed in the bottom. Medium containing recombinant Wnt5a (rWnt5a) was added to the upper and lower compartment of the Boyden chamber. The cells were allowed to invade for 6 h at 37°C in this assay. Thereafter, the medium was discarded, stationary cells were removed with a cotton-tipped applicator, and the membranes were cut out of the chamber and stained with 0.5% crystal violet. The response was evaluated on a light microscope by counting the number of cells that had invaded into the Matrigel and membrane.

*Small G-protein activation assay.* For RhoA, Cdc42 and Rac1 activation assays, the glioblastoma tissues were grinded in liquid nitrogen. Equal weights of total tissue debris were dissolved in ice-cold PBS buffer. Glioblastoma cells were seeded into 6-well plates and transfected with Daam1-siRNA or treated with sFRP2 (R&D Systems, Minneapolis, MN, USA). The experiments were then performed according to the manufacturer's protocol (Cytoskeleton Inc., Denver, CO, USA). The activation of RhoA, Cdc42 and Rac1 was normalized to the NC control group.

Western blotting. Subconfluent cells were washed twice with PBS, and then lysed with ice-cold RIPA lysis buffer (Beyotime Biotechnology, Nantong, China). The lysates were then clarified by centrifugation at 12,000 x g for 20 min at 4°C. The protein extracts were separated by 8% sodium dodecyl sulfate-polyacrylamide gel electrophoresis (SDS-PAGE). The immunoblotting procedure was performed as previously described (22), and the following antibodies were used: anti-GAPDH (Sigma, St. Louis, MO, USA), anti-Daam1 (Santa Cruz Biotechnology, Santa Cruz, CA, USA) antibodies. Protein bands were detected by incubation with horseradish peroxidase-conjugated antibodies and visualized with enhanced chemiluminescence (ECL) reagent (Thermo Scientific, Rockford, IL, USA).

*Pull-down assays.* For the detection of active Daam1, GST-RhoA beads were incubated with 0.1 mmol/l GTP $\gamma$ S (Sigma) at 30°C for 15 min with constant agitation. Equal volumes of total cellular protein were incubated with GST-RhoA beads captured on MagneGST Glutathione



Figure 2. RhoA is activated and positively correlated with Wnt5a in invasive glioblastoma tissues. (A-C) The activation of (A) RhoA and (B) Cdc42 was highly expressed in invasive glioblastoma tissues (n=5) when compared to that in non-invasive glioblastoma tissues (n=4). (C) The activations of Rac1 exhibited no significant difference between invasive glioblastoma and non-invasive glioblastoma tissues. N0, non-invasive glioblastoma; N, invasive glioblastoma; ns, no significance. (D-F) The correlation analysis in glioblastoma tissues was performed between the expression of Wnt5a and activation of RhoA, Cdc42 and Rac1. (D) The activation of RhoA was positively correlated with Wnt5a expression. Correlation coefficients ( $R^2$ ) are shown in the panels.

Particles (Promega, Madison, WI, USA) at 4°C with constant rotation for 90 min. The beads were washed three times with washing buffer (4.2 mmol/l Na<sub>2</sub>HPO<sub>4</sub>, 2 mmol/l KH<sub>2</sub>PO<sub>4</sub>, 140 mmol/l NaCl and 10 mmol/l KCl, pH 7.2). At the end of this period, the beads were captured using a magnet on a magnetic stand. After being washed three times with ice-cold buffer, the beads were resuspended in Laemmli buffer, boiled, and subjected to western blot analysis. SDS-PAGE and western blotting were performed using standard methods.

Actin cytoskeleton staining and immunofluorescence. Transfected cells were fixed in 4% paraformaldehyde in PBS for 20 min, permeabilized in 0.2% Triton X-100 and blocked in PBS containing 1% BSA for 1 h at room temperature. F-actin was stained with FITC-labeled phalloidin (5 mg/ml) (Beyotime Biotechnology) for 40 min at room temperature. After being washed with PBS, the coverslips were mounted on glass slides with DAPI Fluoromount-G (Southern Biotech, Birmingham, AL, USA). The images were acquired with a fluorescence microscope (Zeiss, LSM 710 system; Carl Zeiss, Jena, Germany). Statistical analysis. The data were analyzed using Student's t-test with the SPSS statistical software package. All the results were expressed as the mean  $\pm$  SD. For all analyses a two-sided P-value of <0.05 was deemed statistically significant.

### Results

Wnt5a and RhoA are upregulated in invasive glioblastoma tissues. In order to evaluate Wnt5a expression in invasive glioblastoma tissues, we assessed the expression of Wnt5a using ELISA assays in nine samples of glioblastoma. Wnt5a expression was higher in invasive glioblastoma tissues compared to that in non-invasive glioblastoma tissues, with the highest expression at 43.7 ng/ml (Fig. 1). We also assessed the activations of Rho GTPases in the same glioblastoma tissues and found that the activation of RhoA and Cdc42 were higher in invasive glioblastoma tissues (Fig. 2A and B). However, Rac1 activity had an insignificant increase in invasive glioblastoma tissues (Fig. 2C). Moreover, the activation of RhoA was positively correlated



Figure 3. RhoA activation is required for Wnt5a-induced glioblastoma cell invasion. (A) U251 glioblastoma cells were treated with 20 ng/ml Wnt5a and allowed to invade for 6 h. The rates of cell invasion were determined by Boyden chamber assays. The left panel shows the glioblastoma cells that had invaded onto the bottom surface. Magnification, x200. (B and C) RhoA activation, but not Rac1 and Cdc42 activation, was induced by Wnt5a, but blocked by sFRP2 treatment. Serum-starved U251 and T98MG cell monolayers were incubated with 20 ng/ml rWnt5a for 1 h, or pre-treated with 100 ng/ml sfrp2 for 1 h and then incubated with 20 ng/ml recombinant Wnt5a (rWnt5a) for 1 h. Cell lysates were assayed for small G-protein activation assay. ns, no significance. (D and E) Wnt5a-induced cell invasion was abolished by CCG-1423. U251 and T98MG cells were pre-incubated with RhoA-specific inhibitor CCG-1423 (10 ng/ml) for 1 h, and then allowed to invade in Matrigels with 20 ng/ml rWnt5a treatment for 6 h. The cell invasion rate was determined by Boyden chamber assays.

with the expression of Wnt5a in glioblastoma tissues (Fig. 2D), while there was a weak correlation between the activation of Cdc42 or Rac1 and Wnt5a expression (Fig. 2E and F). These results indicated that Wnt5a and RhoA had tumor-promoting roles in glioblastoma invasion.

*Wnt5a stimulates glioblastoma cell invasion in vitro*. To assess the effect of Wnt5a on glioblastoma cell invasion, we treated U251 glioblastoma cells with 20 ng/ml rWnt5a, and assessed the invasion rate using the Boyden chamber assay. An approximately threefold increase of cell invasion was observed in U251 cells treated with 20 ng/ml rWnt5a, which indicated that Wnt5a had a potent stimulatory effect on glioblastoma cell invasion (Fig. 3A). *RhoA activation participates in Wnt5a-induced glioblastoma cell invasion.* We investigated whether RhoA activation was induced by Wnt5a in glioblastoma cells. Small G-protein assays revealed a significant increase of active RhoA after 20 ng/ml rWnt5a treatment (Fig. 3B). Pre-incubation of secreted frizzled-related protein 2 (sFRP2), an antagonist that directly binds to Wnt5a (21), abolished rWnt5a-RhoA activity in U251 and T98MG cells (Fig. 3B). However, Wnt5a and/or sFRP2 treatment did not alter the activation of Rac1 and Cdc42 in U251 cells (Fig. 3C). To assess the effect of RhoA on glioblastoma cell invasion, we treated glioblastoma cells with 10 ng/ml RhoA-specific inhibitor CCG-1423 and 20 ng/ml rWnt5a, and assessed the invasion rate using the Boyden chamber assay. We



Figure 4. Dishevelled-associated activator of morphogenesis 1 (Daam1) is upstream of RhoA and its activation is essential for Wnt5a-induced glioblastoma cell invasion. (A) RhoA was a downstream target of Daam1. U251 and T98MG cells were transfected with Daam1 siRNA or scramble siRNA, and then incubated with 20 ng/ml recombinant Wnt5a (rWnt5a) for 1 h. Cell lysates were assayed for small G-protein activation assay. (B and C) Wnt5ainduced cell invasion was abolished by Daam1 siRNA. U251 and T98MG cells were transfected with Daam1 siRNA or scramble siRNA, and then allowed to invade in Matrigels with 20 ng/ml rWnt5a treatment for 6 h. The cell invasion rate was determined by Boyden chamber assays.

found that CCG-1423 blocked Wnt5a-induced glioblastoma cell invasion (Fig. 3D and E), indicating that RhoA activation was indispensable for Wnt5a-induced glioblastoma cell invasion.

Wnt5a induces glioblastoma cell invasion via Daaml activation. We examined whether Wnt5a-induced RhoA activation was regulated by Daam1 in glioblastoma cells. Specific siRNA against Daam1 blocked Wnt5a-induced RhoA activation (Fig. 4A). Moreover, Daam1 siRNA was fully capable of retarding the invasion of U251 and T98MG



Figure 5. Dishevelled-associated activator of morphogenesis 1 (Daam1) activation is triggered by Wnt5a in glioblastoma cells. (A) Serum-starved U251 and (B) T98MG cell monolayers were pre-treated with 100 ng/ml sFRP2 for 1 h, and then incubated with 20 ng/ml recombinant Wnt5a (rWnt5a) for 30 min. Cell lysates were assayed for pull-down assays and western blotting. The ratios of active Daam1 and total Daam1 are displayed in the bottom panels. GAPDH was used as the internal control.

cells (Fig. 4B and C). To analyze the role of Wnt5a on Daam1 activation (a state allowing its interaction with RhoA), we blocked Wnt5a signaling with sFRP2 treatment. SFRP2 significantly inhibited Wnt5a-induced Daam1 activity in U251 and T98MG cells (Fig. 5A and B). These experiments demonstrated that RhoA was a downstream target of Wnt5a/Daam1 signaling in U251 and T98MG cells.

Wnt5a/Daam1/RhoA signaling sustains the formation of stress fibers. We performed fluorescent phalloidin staining to investigate the distribution pattern of filamentous actin (F-actin) in glioblastoma cells. Wnt5a treatment sustained the formation/maintenance of actin stress fibers in U251 cells (Fig. 6). In contrast, Daam1-siRNA or CCG-1423 treatment disrupted the formation of actin stress fibers in U251 cells (Fig. 6). Finally, neither Daam1-siRNA nor CCG-1423 treatment altered the cell proliferation of U251 and T98MG cells (Fig. 7A and B). Thus, the findings from the clinical and cellular biological assays indicated that RhoA activation required Daam1 activity to mediate Wnt5a-induced glioblastoma cell invasion (Fig. 7C).

## Discussion

Wnt5a acts both as a suppressor and an inducer in different types of tumors (12,23-26). The Wnt/planar cell polarity (PCP) pathway triggered by Wnt5a activates small Rho-GTPases and reassembles the cytoskeletal architecture and cellular polarity during embryo development and pathological processes (27-29). In our previous studies, we found that Wnt5a stimulated the migration of breast cancer cells via Rho signaling pathways (21,30). In gastric cancer, Wnt5a also enhanced the ability of cell mobility via RhoA signaling pathways (31). In the present study, we found that Wnt5a induced the invasion of U251 and T98MG glioblastoma cells. Moreover,



Figure 6. Daam1 and RhoA participates in the rearrangement of stress fibers in glioblastoma cells. Daam1 siRNA and CCG-1423 disrupted the formation of actin stress fibers in U251 cells. U251 cells were grown on coverslips, then transiently transfected with Daam1 siRNA or pre-treated with 10 ng/ml CCG-1423 for 1 h. U251 cells were then incubated in culture medium containing 20 ng/ml recombinant Wnt5a (rWnt5a) for 4 h. Subsequently, the cells were fixed and F-actin organization was analyzed by phalloidin staining. Magnification, x400.



Figure 7. Daam1 and RhoA do not regulate the proliferation of glioblastoma cells. (A and B) U251 and T98MG cells were transfected with (A) Daam1 siRNA or (B) pre-treated with 10 ng/ml CCG-1423 for 1 h, and then subjected to the cell proliferation assays. Daam1 siRNA and CCG-1423 did not alter the proliferation of glioblastoma cells; ns, no significance. (C) A schematic illustration of the regulation of RhoA in Wnt5a/Daam1 signaling to mediate cell glioblastoma invasion (see text for details).

Wnt5a was highly expressed in invasive glioblastoma tissues compared to that in non-invasive glioblastoma tissues. These results revealed that Wnt5a may accelerate glioblastoma invasion *in vitro* and *in vivo* and that blocking Wnt5a signaling may prevent glioblastoma metastasis in clinic. In addition, the overexpression of Wnt5a increased the proliferation of glioblastoma GBM-05 and U87MG cells, indicating that Wnt5a is a regulator in the proliferation of human glioblastoma (14). This evidence demonstrated that Wnt5a acts as an inducer and promoter in glioblastoma tumorigenesis and metastasis.

With the help of small G-protein promoting cellular migration, Wnt5a can remodel the cellular cytoskeleton of melanoma cells (11). In the present study, Wnt5a promoted glioblastoma cell invasion by activating the Daam1/RhoA signaling pathway. In clinical samples, Wnt5a and RhoA are upregulated in invasive glioblastoma tissues, with a significant positive correlation between them. Studies with much larger samples may provide statistical power to validate the role of Wnt5a and RhoA in glioblastoma. Our previous study revealed that Rac1 activation was increased by Wnt5a and stimulated the cell migration of MCF-7 breast cancer cells (30). Unfortunately, Wnt5a did not alter the activation of Rac1 and Cdc42 in U251 glioblastoma cells in the present study. These results reveal the specificity of elevated RhoA activation in glioblastoma invasion.

Containing multiple regulatory domains, Daam1 exists in an auto-inhibited state through intramolecular interaction in unstimulated cells (10,32). Our results showed that sFRP2, an antagonist that directly binds to Wnt5a, markedly blocked the activity of Wnt5a-induced Daam1 in glioblastoma cells. Knockdown of Daam1 expression via siRNA transfection inhibited RhoA activation and the cell invasion stimulated by Wnt5a in U251 and T98MG cells. Furthermore, we tested the reestablishment of stress fibers in Daam1-knockdown or RhoA-blocked cells. The decrease of RhoA activity, formation of stress fibers and cell invasion in Daam1-knockdown cells indicated that Daam1 may be required for the activation of RhoA after Wnt5a treatment in glioblastoma. Thus, these results clearly demonstrated that Daam1/RhoA signaling under Wnt5a stimulation participated in the invasion of glioblastoma and that retarding them may prevent glioblastoma metastasis in clinic.

Soluble frizzled-related proteins (sFRPs) function as modulators of Wnt signaling through direct interaction with Wnts (33). We reported in our previous study that Wnt5ainduced RhoA activation and cell migration could be abolished by sFRP2 pretreatment in breast cancer cells (21). In the present study, we also found that sFRP2 blocked the Wnt5a-induced RhoA activation in glioblastoma cells. These results demonstrated that sFRP2 acted as an antagonist of Wnt5a mediating the progression of glioblastoma and breast cancer. A recent study in immunohistochemistry revealed that the expression level of soluble frizzled-related protein 3 (sFRP3) was decreased in the nucleus in higher grade astrocytoma, indicating the antagonistic ability of Wnt signaling (34). Further studies are needed to decipher whether sFRP2 and sFRP3 function in a common pathway or in parallel pathways to block Wnt5a signaling.

In conclusion, the present study partially clarified the associations between Wnt5a/RhoA signaling and glioblastoma progression. It is the first to demonstrate that Wnt5a may regulate the invasion of glioblastoma cells, at least in part via the Daam1/RhoA signaling pathway. Therefore, the Wnt5a/Daam1/RhoA signaling pathway is a candidate accelerator in glioblastoma and may be a potential clinical classification marker and therapeutic target for human glioblastoma.

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