Abstract. Glioma is the most common primary brain malignant tumor. Receptor for activated C-kinase 1 (RACK1) is widely expressed in the central nervous system, and regulates multiple cellular processes including cell survival, proliferation, migration and metastasis. However, the role of RACK1 in glioma has never been revealed. The present study, for the first time, showed that RACK1 expression was significantly higher in glioma tissues and cell lines when compared with that in normal brain tissues, and was positively associated with the malignancy of glioma. siRNA-induced RACK1 downregulation significantly suppressed the proliferation and invasion of human glioma U87 and CHG-5 cells, while it promoted their apoptosis by upregulating Bax expression and reducing Bcl-2 expression. Furthermore, forced downregulation of RACK1 notably inhibited tumor xenograft growth in nude mice. These findings suggest that RACK1 plays a critical role in the development and progression of glioma in vitro and in vivo. Moreover, siRNA-induced RACK1 downregulation markedly reduced the activity of Src/Akt signaling pathway, which plays an important role in the growth and behavior of human malignancies, indicating that siRNA-mediated RACK1 downregulation inhibited glioma probably via suppressing Src/Akt signaling activity. The present study highlighted the role of RACK1 in glioma, and demonstrated that RACK1 is a novel promising therapeutic target for glioma treatment.

Introduction

Malignant glioma is the most common primary brain tumor, accounting for approximately 80% of malignant tumors in the central nervous system. It has a very poor prognosis, mainly due to its resistance to radiotherapy, chemotherapy and adjuvant therapies (1-3). Generally, patients diagnosed with glioblastoma, the most malignant form of glioma, live approximately only 1 year after diagnosis (4). Despite the marked developments in the therapy of other types of cancer, the median survival rate of malignant glioma has not improved in the past years (5). Therefore, a more effective therapeutic strategy is urgently required.

Receptor for activated C-kinase 1 (RACK1) is a member of the intracellular receptors for activated protein kinase C (PKC) (6). RACK1 is widely expressed in the central nervous system, and is involved in multiple crucial cellular processes including cell growth, proliferation, apoptosis and migration (7-9). In recent years, accumulating evidence has shown that RACK1 plays an important role in various types of cancer (10-12). Therefore, we hypothesized that RACK1 may act as an important regulator in the development and progression of malignant glioma.

In this study, we first determined the mRNA and protein expression levels of RACK1 in glioma tissues of different grades as well as in normal brain tissues. Then, the mRNA and protein expression of RACK1 was determined in two types of human glioma cell lines U87 and CHG-5. Using RACK1-specific siRNA, the effects as well as the molecular mechanism of RACK1 downregulation involved in proliferation, apoptosis and invasion of U87 and CHG-5 cells were further investigated in vitro and in vivo.

Materials and methods

Materials and reagents. The protocol used in the present study was approved by the Ethics Committee of Central South University. Written informed consent was obtained from each participant. Forty-five glioma tissues (10 cases of WHO I, 13 cases of WHO II, 12 cases of WHO III, and 10 cases of WHO IV) were obtained from patients who underwent surgery from October 2011 to October 2012 at the Department of Neurosurgery, Xiangya Hospital of Central South University (Hunan, China), and 10 normal brain tissues from patients with cerebral trauma were used as controls. The human glioma U87 (WHO IV) and CHG-5 (WHO II) cell lines were obtained from the Cell Bank of Central South University. Dulbecco's modified Eagle's medium (DMEM), opti-MEM, fetal bovine serum (FBS) and Lipofectamine 2000 transfection agent were purchased from Invitrogen Life Technologies (USA). MTT was

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obtained from Sigma (USA). The bicinchoninic acid (BCA) protein assay kit was purchased from Pierce Biotechnology, Inc. (USA). PVDF membrane was obtained from Pall (USA). All siRNAs and antibodies were obtained from Santa Cruz Biotechnology, Inc. (USA). The Annexin V-FITC Apoptosis Detection kit was purchased from Biovision (USA). Matrigel was purchased form BD Biosciences (USA).

Cell culture. U87 and CHG-5 cell lines were cultured in DMEM containing 10% FBS, 100 U/ml penicillin and 100 U/ml streptomycin. Cells were incubated at 37°C in a humidified incubator of 95% air and 5% CO₂.

Real-time RT-PCR. The RACK1 mRNA expression in tissues or cell lines was detected by real-time RT-PCR (ABI 7500). Specific primers for different genes in this study were synthesized by BGI Company (Guangzhou, China). Specific primers for RACK1 were: sense, 5'-GAGCAAAATGACCCTCTGT-3' and antisense, 5'-GTAGTGCCCCGTTGTGAGA-3'. Specific primers for Bax were: sense, 5'-CCCAGAGGCTTTTTCCGAG-3' and antisense, 5'-CCAGCCCATGATGGTTCTGAT-3'. Specific primers for Bcl-2 were: sense, 5'-GATGCGGTCACTGTCATCATCC-3' and antisense, 5'-CGTTGAGGCTTCTGTCATGATGATGATGATGATGATG-3'. All amplifications were performed in 3 parallel samples.

Western blotting. Cells were lysed by cold RIPA lysis, and the protein concentrations were determined using BCA protein assay kit. Then, proteins of 20 µg/lane were loaded on 12% SDS-PAGE to separate, and then electrophoretically transferred to PVDF membranes. Proteins on the membranes were transferred using primary antibodies according to the supplier's protocol. Following incubation with secondary antibodies, results were visualized with peroxidase and an enhanced chemiluminescence system (Pierce Biotechnology, Inc.) and quantified by Quantity One software (Bio-Rad, USA).

siRNA interference. Human glioma U87 and CHG-5 cells were seeded at a density of 10³ cells/well in 6-well plates and cultured in DMEM containing 10% FBS. After incubating at 37°C, 5% CO₂ for 24 h, U87 and CHG-5 cells were transfected with siRNA and Lipofectamine 2000 according to the supplier's instruction. Briefly, 100 nmol siRNA and 5 µl Lipofectamine 2000 were diluted in opti-MEM to a final volume of 800 µl. After mixing for 20 min at room temperature, the siRNA/Lipofectamine 2000 mixture was added. Cells were incubated at 37°C 5% CO₂ for 6 h. Following incubation, the mixture was replaced with DMEM containing 10% FBS for 24 h.

Proliferation assay. Cell proliferation was determined by MTT assay. After 24 h post-transfection, the transfection medium in each well was replaced by DMEM medium containing 10% FBS used before, and was cultured for 12, 24, 36, 48 and 60 h. Then, the medium was replaced by 100 µl of fresh serum-free medium and cultured with 0.5 g/l MTT. Following incubation at 37°C for 4 h, the MTT medium was removed by aspiration and 50 µl of DMSO was added to each well. Following incubation at 37°C for a further 10 min, the A540 of each sample was measured using a plate reader.

Apoptosis analysis. Flow cytometry was used to determine the cell apoptosis with the Annexin V-FITC Apoptosis Detection kit. After 24 h post-transfection, cells were harvested and washed with cold PBS twice. Subsequently, 10⁶ cells were resuspended in 200 µl binding buffer supplemented with 10 µl Annexin V-FITC and 5 µl PI-PE, and incubated in the dark for 30 min. Then, 300 µl binding buffer was added followed by flow cytometry assay.

Transwell matrix penetration assay. Cells (10⁵) of different groups in 200 µl serum-free DMEM were plated on the upper chamber plated on the top side of polycarbonate Transwell filter coated with Matrigel and incubated at 37°C for 24 h. Subsequently, cells inside the upper chamber were removed. Cells on the lower membrane surface were fixed in 1% paraformaldehyde, stained with 0.1% crystal violet and counted.

Statistical analysis. Data are presented as means ± standard deviation (SD). SPSS 17.0 software was used to perform statistical analyses using a two-tailed Student's t-test or one-way ANOVA. P<0.05 was considered to indicate statistically significant differences. All experiments were repeated at least 3 times.

Results

Upregulation of RACK1 in human glioma tissues and glioma cell lines. To preliminarily investigate the relationship between RACK1 expression and glioma, we first examined the mRNA and protein expression levels of RACK1 in normal brain tissues, gliomas of different grades and 2 glioma cell lines. As shown in Fig. 1A and B, real-time PCR and western blotting results showed that the expression of RACK1 showed an increasing tendency with the malignancy of glioma. We further determined the expression of RACK1 in 2 human glioma cell lines, U87 (grade IV) and CHG-5 (grade II). As shown in Fig. 1C and D, real-time PCR and western blotting results demonstrated that RACK1 expression was also upregulated in U87 and CHG-5 cells compared to normal brain tissues (P<0.01). However, there was no difference of RACK1 expres-
sion between these 2 glioma cell lines (P>0.05). Accordingly, these data above indicate that the increased expression of RACK1 may be associated with the malignancy of glioma.

siRNA-induced RACK1 downregulation in U87 and CHG-5 cells. Due to the increased expression of RACK1 in glioma tissues and cell lines, we used RACK1-specific siRNA to downregulate the expression of RACK1 in U87 and CHG-5 cells to further investigate the role of RACK1 in glioma. After transfection, we first determined the mRNA and protein expression of RACK1 in U87 and CHG-5 cells using real-time RT-PCR and western blotting, respectively. As shown in Fig. 2, after transfection with RACK1-specific siRNA, the mRNA and protein levels of RACK1 in U87 and CHG-5 cells were effectively downregulated (P<0.01), when compared with those in the untreated (control) and the non-specific siRNA group (NC).

siRNA-induced RACK1 downregulation suppresses proliferation of glioma U87 and CHG-5 cells. To further investigate the role of RACK1 in glioma cells in vitro, MTT assay was performed to determine the effect of siRNA-induced RACK1 downregulation on proliferation of U87 and CHG-5 cells. As shown in Fig. 3, MTT assay demonstrated that in RACK1-downregulated U87 and CHG-5 cells, the cell proliferation rate was lower when compared with controls (P<0.01). These results suggest that RACK1 promotes proliferation of human glioma cells.

siRNA-induced RACK1 downregulation enhances apoptosis of glioma U87 and CHG-5 cells. We further investigated the effect of RACK1 downregulation on apoptosis of glioma cells. As shown in Fig. 4, in RACK1-downregulated U87 and CHG-5 cells, the cell apoptosis rate was higher when compared with controls (P<0.01). These results suggest that RACK1 promotes apoptosis of human glioma cells.

Figure 1. The mRNA and protein expression of RACK1 in glioma tissues and cells. (A) Real-time RT-PCR was performed to determine the relative mRNA expression of RACK1 in gliomas of different grades. (B) Western blotting was applied to determine the protein expression of RACK1 in gliomas of different grades. (C) Real-time RT-PCR was performed to determine the relative mRNA expression of RACK1 in 2 glioma cell lines, U87 and CHG-5. (D) Western blotting was applied to determine the protein expression of RACK1 in 2 glioma cell lines, U87 and CHG-5. All data in the experiment groups were compared with the normal group. *P<0.05; **P<0.01. RACK1, receptor for activated C-kinase 1.

Figure 2. The mRNA and protein expression of RACK1 in U87 and CHG-5 cells after siRNA-mediated interference. Control, cells without any treatment. NC, cells transfected with non-specific siRNA. RACK1 siRNA, cells transfected with RACK1-specific siRNA. (A) Real-time RT-PCR was performed to determine the relative mRNA expression of RACK1 in U87 and CHG-5 cells after siRNA-mediated interference. (B) Western blotting was applied to determine the protein expression of RACK1 in U87 and CHG-5 cells after siRNA-mediated interference. All data in the experiment groups were compared with the control group. *P<0.01. RACK1, receptor for activated C-kinase 1.
cells, the cell apoptosis rate was much higher when compared with controls (P<0.01), suggesting that RACK1 may play an inhibitory role in the regulation of apoptosis of human glioma cells. We then examined the expression of apoptotic-related genes in each group. Real-time PCR assay showed that the mRNA expression of pro-apoptotic gene Bax was upregulated.
while the mRNA expression of anti-apoptotic gene Bcl-2 was reduced, in RACK1-downregulated glioma U87 and CHG-5 cells, when compared with those in controls. These results indicated that RACK1 may suppress apoptosis of glioma cells in vitro through directly or indirectly regulating the expression of apoptosis-related genes, Bax and Bcl-2.

**siRNA-induced RACK1 downregulation inhibits invasion of glioma U87 and CHG-5 cells.** The alternations of cell invasion ability of human glioma U87 and CHG-5 cells were examined after transfection with RACK1-specific siRNA. As shown in Fig. 5, RACK1-downregulated cells showed decreased invasion ability when compared with controls (P<0.05). These data indicate that RACK1 promotes invasion ability of human glioma cells in vitro.

**siRNA-induced RACK1 downregulation suppresses survival, apoptosis, migration, and proliferation relative signaling pathways in U87 and CHG-5 cells.** Bax is a pro-apoptotic gene, while Bcl-2 has an anti-apoptotic function. It has been well established that the ratio of Bax/Bcl-2 protein plays a crucial role in regulating cell apoptosis. Hence, we applied real-time RT-PCR to determine the mRNA expression of Bax and Bcl-2 in U87 and CHG-5 cells of each group. As shown in Fig. 6A, the relative mRNA level of Bax was significantly upregulated in U87 and CHG-5 cells after transfection with RACK1-specific siRNA, when compared with that in the control group (P<0.01). However, the relative mRNA level of Bcl-2 was decreased in U87 and CHG-5 cells after transfection with RACK1-specific siRNA, compared with that in the control group (P<0.01). These data partly explain the above findings that forced downregulation of RACK1 promotes the apoptosis of U87 and CHG-5 cells.

Src/Akt signaling pathway plays a crucial role in the regulation of survival, proliferation and migration of multiple types of cells. It has been reported that RACK1 induces colon cell apoptosis, partly by suppressing Src activity in Akt pathway. However, whether a similar molecular mechanism exists in glioma cells remains unknown. Thus, we determined the activity of Src/Akt signaling pathway in RACK1-downregulated U87 and CHG-5 cells. As shown in Fig. 6B, the phosphorylation levels of Src and Akt were much lower in U87 and CHG-5 cells after transfection with RACK1-specific siRNA, when compared with those in controls (P<0.05). These results indicate that the effects of RACK1 on the survival, proliferation, invasion of glioma cells may be involved in the regulation of the Src/Akt signal pathway.

**Forced downregulation of RACK1 suppresses tumor xenograft growth in nude mice.** To further investigate the role of RACK1 in vivo, a tumor xenograft animal model was conducted using U87 cells, in which RACK1 was successfully knocked down by lentivirus infection. After all animals were subcutaneously implanted with the infected tumor cells, the size of tumors in the RACK1-specific shRNA group was significantly smaller than that in the control and NC group (Fig. 7A). As shown in Fig. 7B, the average tumor weight in the RACK1-specific shRNA group was markedly lower than that in the control and
Figure 6. The activity of Src/Akt signaling pathway in U87 and CHG-5 cells after siRNA-mediated RACK1 downregulation. Western blotting was applied to determine the phosphorylation levels of Bax and Bcl-2 proteins in U87 and CHG-5 cells after siRNA-mediated RACK1 downregulation. Control, cells without any treatment. NC, cells transfected with non-specific siRNA. RACK1 siRNA, cells transfected with RACK1-specific siRNA. All data in the experiment groups were compared with the control group. *P<0.01. RACK1, receptor for activated C-kinase 1.

Figure 7. Forced downregulation of RACK1 suppresses tumor xenograft growth in nude mice. The tumor xenograft animal model was conducted using U87 cells, in which RACK1 was successfully knocked down by lentivirus infection. Control, U87 cells without any treatment. NC-LV, U87 cells infected with non-specific shRNA lentivirus. RACK1 shRNA-LV, U87 cells infected with RACK1-specific shRNA lentivirus. (A) The average tumor volume in each group was determined at day 10, 15, 20, 25 and 30 after implantation. (B) The relative tumor weight in each group after excision. (C) The image of typical tumor in each group after excision. All data in the experiment groups were compared with the control group. *P<0.01.
Ribosomal RACK1 promotes the proliferation, survival, and migration (27,28). Accumulating evidence has revealed that Src could be recruited by RACK1, and acts in downstream of Src and plays an important role in the regulation of cell survival, proliferation and migration partly via suppressing Src activity, in colon cancer cells, indicating that this regulatory mechanism may exist in multiple types of cancer cells.

In conclusion, the present study indicated that the upregulation of RACK1 may suppress the development of glioma, and hence may become a promising therapeutic target for this cancer.

Based on these clinical findings, we speculated that forced downregulation of RACK1 may suppress the development of gliomas. To test this hypothesis, RACK1-specific siRNA was applied to successfully downregulate the RACK1 expression in human glioma U87 and CHG-5 cells. As expected, RACK1 downregulation significantly inhibited the proliferation and invasion in U87 and CHG-5 cells. The Bcl-2 family plays an essential role in the regulation of cell survival and apoptosis, including both pro-apoptotic members such as Bax, as well as anti-apoptotic members such as Bcl-2. The balance between Bax and Bcl-2 determines the susceptibility of cells to the apoptotic signal (20). We found that siRNA-induced RACK1 downregulation notably promoted cell apoptosis of U87 and CHG-5 cells, partly through suppressing Bax expression and upregulating Bcl-2 expression, and hence breaking this balance for maintaining cell survival.

Based on these findings in vitro, we further applied tumor xenograft animal models to test whether RACK1 downregulation has an inhibitory effect on tumor growth in vivo, and found that stable downregulation of RACK1 in human glioma U87 cells markedly suppressed the tumor growth in vivo. These findings suggest that RACK1 may play a crucial role in tumorigenesis and progression of glioma in vitro and in vivo.

Src is a protein tyrosine kinase and participates in the regulation of multiple cellular processes including cell survival, proliferation and migration (21,22). Accumulating evidence has revealed that Src could be recruited by RACK1, and acts as an oncogene in some types of cancer including glioma (23-26). Akt, also known as protein kinase B (PKB), acts in downstream of Src and plays an important role in the regulation of cell survival, proliferation and migration (27,28). However, whether Src/Akt signaling activity is involved in RACK1-mediated glioma development has yet to be investigated. The present study demonstrated that forced RACK1 downregulation suppressed the activity of Src/Akt signaling pathway in U87 and CHG-5 cells. These findings suggest that siRNA-induced RACK1 downregulation inhibits glioma development partly via suppressing Src/Akt signaling activity. Our results are consistent with the findings of Mamidipudi and Cartwright (29) that RACK1 promotes mitochondrial cell death and blocked Akt-mediated cell survival, partly via suppressing Src activity, in colon cancer cells, indicating that this regulatory mechanism may exist in multiple types of cancer cells.

In conclusion, the present study indicated that the upregulation of RACK1 is a common event in glioma, and that RACK1 plays a critical role for glioma development and progression in vitro and in vivo. Moreover, the underlying mechanism involves RACK1-mediated Src/Akt signaling activity. Thus, the present study suggests that RACK1 may be a novel promising therapeutic target for glioma treatment.

References


