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B7-H4 expression is upregulated by PKCδ activation and contributes to PKCδ-induced cell motility in colorectal cancer

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Abstract

Introduction: B7-H4 is overexpressed in colorectal cancer (CRC) and plays an important role in tumor growth and immunosuppression. However, the exact mechanism that regulates B7-H4 expression remains largely unknown. Here, we investigated whether protein kinase C δ (PKC δ) regulates the expression of B7-H4 in CRC.

Methods: By using immunohistochemical (IHC) and immunofluorescence (IF) staining, we analyzed the expression of B7-H4 and phospho-PKC δ (p-PKC δ) in 225 colorectal tumor samples and determined the clinical significance of the expression patterns. In vitro experiments were performed with the CRC cell lines HCT116 and SW620 to detect the effect of PKC δ activation on B7-H4 expression, and xenograft-bearing mice were treated with rottlerin to monitor the expression of B7-H4 and tumor metastasis.

Results: The B7-H4 expression level was significantly correlated with the p-PKC δ level (r = 0.378, P < 0.001) in tumor tissues. Coexpression of p-PKC δ and B7-H4 was significantly associated with moderate/poor differentiation (P = 0.024), lymph node metastasis (P = 0.001) and advanced Dukes' stage (P = 0.002). Western blot analysis showed that Phorbol-12-Myristate-13-Acetate (TPA) increased B7-H4 expression in a concentration-dependent manner and that rottlerin abrogated the TPA-induced increase in B7-H4 expression. The protein levels of B7-H4 and p-STAT3 were significantly reduced by a PKC δ -specific siRNA. Moreover, the STAT3 inhibitor cryptotanshinone significantly decreased the B7-H4 protein level in CRC cells. Knockdown of B7-H4 or PKC δ suppressed cell migration and motility. Rottlerin also inhibited B7-H4 expression and tumor metastasis in vivo.

Conclusion: The B7-H4 expression level is significantly correlated with the p-PKCδ level and tumor metastasis in CRC samples. B7-H4 expression is upregulated by STAT3 activation via PKCδ and plays roles in PKCδ-induced cancer cell motility and metastasis, suggesting that the PKCδ/STAT3/B7-H4 axis may be a potential therapeutic target for CRC.

Keywords: B7-H4, PKCδ, Regulation, Metastasis, Colorectal cancer

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Introduction

Colorectal cancer (CRC) is the third most common cancer worldwide, and approximately 30,000 new cases occur every year. CRC leads to 13,000 deaths annually and is the fourth leading cause of cancer-related mortality [1, 2]. Although advances in early diagnosis and therapeutic strategies have decreased the mortality of CRC,

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the survival rate is still poor [3, 4]. The high mortality rate is usually attributed to tumor recurrence and metastasis.

B7-H4 (VTCN1/B7x/B7S1) is a costimulatory molecule in the B7 family and is mainly expressed by antigen-presenting cells (APCs). The expansion of both neutrophil progenitors and T cells can be suppressed by B7-H4, resulting in immune suppression [5-7]. In numerous tumor tissues, B7-H4 is overexpressed and positively correlated with various clinicopathological features [8–13]. B7-H4 expression was found to be significantly higher in CRC tissues than in normal tissues and positively related to infiltration depth, lymph node metastasis and regulatory T cell (Treg) infiltration [14, 15]. Furthermore, soluble B7-H4 in serum has been shown to be a potential biomarker for diseases [16–19]. In addition to its role in immune suppression, B7-H4 also promotes tumor proliferation and metastasis, as revealed by many studies. Li et al. found that B7-H4 facilitates the proliferation and metastasis of CRC cells [20]. Zhang et al. revealed that B7-H4 also promotes lung cancer growth and metastatic progression [21]. Xie et al. showed that B7-H4 promotes tumor invasion and metastasis through activation of ERK1/2 signaling [22]. In summary, B7-H4 contributes to immune evasion, tumor growth and tumor metastasis, but the exact mechanism that regulates B7-H4 expression is not well elucidated.

The protein kinase C (PKC) family, comprising a series of serine/threonine kinases, regulates various cellular physiological processes [23, 24]. The PKCδ level is elevated in colorectal cancer tissue, suggesting a specific role for PKC δ in colon carcinogenesis [25–27]. As a unique novel PKC, PKC8 plays a significant role in diverse cancers and has different cell-specific effects [28– 30]. PKC δ is proposed to act mainly as a tumor suppressor due to its antiproliferative and proapoptotic activities. However, it was found that downregulation of PKC δ can induce cytotoxicity and inhibit the growth of cultured stem-like cells originating from human breast, pancreatic and prostate cancers[29, 31]. In colon cancer, PKC δ can inhibit cell growth and proliferation, and it also acts as a proapoptotic regulator [32, 33]. Furthermore, many studies have revealed that PKC δ is involved in colon cancer cell migration and invasion [26, 34-36].

In this study, we demonstrated a positive correlation between B7-H4 and p-PKC δ in clinical CRC samples. Furthermore, the p-PKC δ ⁺B7-H4⁺ phenotype was associated with tumor metastasis. To verify this finding in clinical samples, we investigated the effect of PKC δ on B7-H4 expression. Activation of PKC δ by TPA enhanced B7-H4 expression. However, interfering with PKC δ expression by transfection of specific small interfering RNA (siRNA) constructs downregulated B7-H4 in colorectal cancer cell lines. Furthermore, both PKC δ and B7-H4 contributed to CRC cell motility. Knockdown of B7-H4 abrogated PKC δ -induced cell metastasis, confirming that B7-H4 mediates PKC δ -induced cell metastasis and invasion.

Materials and methods

Clinical samples

The First Affiliated Hospital of Soochow University (Suzhou, China) approved our experimental protocols. A total of 225 colon cancer tissue specimens and 36 adjacent normal tissue specimens were obtained for immunohistochemical (IHC) analysis. All experimental procedures conformed to the tenets of the Declaration of Helsinki (as revised in 2013). Ethical approval of experiments involving clinical samples was given by the Institutional Review Board of Soochow University (No. 2014865082). In addition, informed consent was also obtained from the patients for the experimental use of their samples.

Immunohistochemical and immunofluorescence staining

For IHC analysis of p-PKC δ , a rabbit monoclonal antibody specific for human p-PKC δ (phospho-S299, ab133456) purchased from Abcam (Cambridge, MA, USA) was used at a 1:100 dilution. For B7-H4 IHC staining, a mouse monoclonal antibody (clone 3C8, Suzhou, China) specific for human B7-H4 was produced in our laboratory and used at a 1:200 dilution [37]. For IF analysis, Alexa Fluor 488-conjugated goat anti-rabbit IgG (1:100, A32731) and Alexa Fluor 594-conjugated goat anti-mouse IgG (1:200, A11005) were used as the secondary antibodies from Invitrogen (Carlsbad, CA, USA).

Sections (4 µm thick) were sliced from paraffin-embedded samples with a Leica microtome (Wetzlar, Germany). Tissue microarrays were deparaffinized, rehydrated, rinsed, and stained for B7-H4 and p-PKC8 as previously described [38]. IHC staining was performed using the ${\rm ChemMate}^{{}^{\rm TM}}$ Envision/HRP technique (Gene Tech Company Limited). The sections were evaluated at low magnification $(100 \times)$ to identify positive staining. The samples were classified into four groups according to the percentage of cells positive for B7-H4 or p-PKCδ signals: 1, positive signals in less than 25% of cells; 2, in 26–50%; 3, in 51–75%; and 4, in more than 75%. The staining intensity was categorized based on the relative intensity as follows: 1, weak staining; 2, intermediate staining; and 3, strong staining. The total score was calculated as the positive percentage score X the staining intensity score. Cases with a Quickscore of > 4 were considered positive, and the others were regarded as negative. The stained slides were examined and evaluated by two independent investigators.

For IF analysis, serial sections were incubated with monoclonal anti-p-PKC δ , an anti-B7-H4 and/or isotype IgG antibodies for 1.5 h at room temperature, as previously described [38]. An Alexa Fluor 488-conjugated secondary antibody was used to detect p-PKC δ . An Alexa Fluor 594-conjugated secondary antibody was used to detect B7-H4. IF images were acquired under a Leica DM2500 microscope (Wetzlar, Germany).

Cell culture and transfection

The human CRC cell lines HCT116, SW620, SW480, RKO and NCM460 were purchased from Shanghai Cell Bank (Chinese Academy of Sciences, Shanghai, China). Cells were cultured in RPMI 1640 medium containing 10% fetal bovine serum (FBS) at 37 °C in a humidified atmosphere of 5% CO₂. RPMI 1640 medium and FBS were purchased from HyClone (Logan, UT, USA). The PKC δ activator TPA, PKC δ inhibitor rottlerin (Santa Cruz Biotechnology, CA, USA) and STAT3 inhibitor cryptotanshinone (Selleckchem, USA) were stored at – 80 °C.

Human PKC δ -specific siRNAs, a human B7-H4-specific siRNA and the corresponding control siRNAs (con siRNAs) were purchased from GenePharma Co. Ltd. (Shanghai, China). SiRNAs were transfected into HCT116 or SW620 cells using Lipofectamine 2000 reagent (Invitrogen, Carlsbad, CA, USA). RT–qPCR and Western blotting were performed to evaluate the transfection efficiency.

Total RNA isolation and RT-qPCR

Total RNA was isolated using TRIzol reagent (Invitrogen, Carlsbad, CA, USA), incubated for 20 min at 37 °C with RNase-Free DNase, purified with an RNeasy MinElute Cleaning Kit (74204, Qiagen, Hilden, Germany), and quantified with a spectrophotometer (BioDrop-µLite, UK). Total RNA was reverse transcribed into cDNA using PrimeScript[™] RT Master Mix (Takara Bio, Japan). A SYBR PrimeScript RT-qPCR Kit was used (Takara Bio, Japan) to examine individual genes. The thermal cycling procedure used for PCR was as follows: 95 °C for 2 min, followed by 45 cycles of denaturation at 95 °C for 10 s, annealing at 59 °C for 40 s and extension at 72 °C for 45 s. The levels of all genes examined were normalized to the GAPDH mRNA level. The primers for the individual genes used in RT-qPCR are listed in Additional file 7: Table S1.

Protein extraction and Western blot analysis

Human CRC cells were cultured in 6-well plates and were then lysed with RIPA lysis buffer (Beyotime, Shanghai, China). Protease inhibitor cocktail was added to the RIPA buffer. After determining the protein concentrations of all samples with BCA protein assay kits (Beyotime, Shanghai, China), samples containing equal amounts of total protein were loaded onto a gel and separated by electrophoresis. The protein bands on the gel were transferred to polyvinylidene difluoride (PVDF) membranes (Merck Millipore, Germany). The membranes were incubated with primary antibodies at 4 °C overnight. The primary antibodies used were as follows: mouse antihuman B7-H4 (3C8), rabbit anti-human PKCδ (#9616T, CST, Danvers, Massachusetts, USA), rabbit anti-human/ mouse p-PKC8 (ab133456, Abcam, USA), rabbit antihuman/mouse STAT3 (#12640, CST), rabbit anti-human/ mouse phospho-STAT3 (p-STAT3, Tyr705, #9145, CST), rabbit anti-human/mouse GAPDH (#5174, CST) and rabbit anti-human/mouse β -actin (#4970, CST). After three washes with PBST, the PVDF membranes were incubated with secondary antibodies at room temperature for 2 h. The secondary antibodies were as follows: HRP-conjugated goat anti-mouse/anti-rabbit IgG (H+L) and rabbit anti-goat IgG (H+L) (Thermo Fisher Scientific, Waltham, MA, USA). The membranes were washed with PBST five times. Then, the membranes were immersed in electrochemiluminescence (ECL) detection reagent (CST). Images were acquired with a Gel DocTM EZ System (Bio-Rad, USA). Image Lab 4.0.1 software (Bio-Rad, USA) was used to analyze band intensities.

Flow cytometric analysis and IF analysis

To examine intracellular B7-H4 expression, cells were permeabilized using Intracellular Fixation & Permeabilization Buffer (eBioscience, CA, USA). Then, the cells were stained with a PE-conjugated anti-B7-H4 antibody (#358104, Biolegend, CA, USA). PE-conjugated mouse IgG1 isotype control (eBioscience) was used as the control antibody. Flow cytometry was performed on a Beckman flow cytometer. Data were analyzed using FlowJo software (version 7.6, Tree Star Inc.).

For IF analysis of B7-H4 expression in cells, cultured cells were fixed with cold acetone for 10 min, washed in PBS buffer containing 1% FCS, and stained with a PE-conjugated anti-B7-H4 antibody.

Transwell invasion assay

After 16 h of transfection, cells were harvested by trypsin digestion and resuspended in culture medium containing 10% FBS. A total of 5×10^4 HCT116 or 8×10^4 SW620 cells were seeded in 24-well Transwell chambers containing membranes with a pore size of 8 µm (Falcon, USA). The membranes were precoated with 100 µl of a 1:16 dilution of Matrigel (Corning, New York, USA). After 4–6 h, the culture medium in each well was changed to culture medium containing 2% FBS and a drug or solvent. Then, 700 µl of medium containing 20% FBS was added

to the bottom compartment to act as a chemoattractant. After culture for 24 h, the cells that invaded to the underside of the Transwell membrane were fixed with methanol and stained with a 0.1% crystal violet solution. Then, the stained cells were observed under a Nikon TI-SR inverted microscope and imaged using a Nikon DS-Fi2 camera. ImageJ was used to determine the number of cells in three different fields of view.

Wound healing assay

For evaluation of migration by a wound healing assay, 12-well plates were seeded with cells at a density of 3×10^5 cells/well. After growth and attachment overnight, the surface of the cultured cells was scratched with 10-µl pipette tips. After another 24 h of culture, wound closure was observed and imaged using a Nikon DS-Fi2 camera. The wound closure percentage was calculated using ImageJ software.

Animal experiments

Ten female BALB/c nude mice aged 4-6 weeks were purchased from Shanghai Laboratory Animal Center (Shanghai, China). After 1 week of adaptation, HCT-116 cells (2×10^6) in PBS (100 µl) were injected intravenously into the nude mice. After 24 h, the mice were randomly allocated to two groups: one group of mice was orally administered 200 µl of solvent, while another was orally administered rottlerin (20 mg/kg) once every two days. The dosage of rottlerin was determined according to our preliminary experimental results and previous reports [39, 40]. All procedures in these experiments were approved by the Animal Protection and Use Committee of Soochow University. Fifty-four days after injection, the mice were sacrificed to assess tumor development, and the lungs were removed and embedded in paraffin for further analyses.

Bioinformatics analysis

Correlations between the expression of PKCs and that of B7-H4 in COAD patient samples was analyzed. The mRNA expression of PRKCD (encoding PKC\delta) and VTCN1 (encoding B7-H4) in COAD tissues and normal tissues from the TCGA colon adenocarcinoma (COAD) dataset was compared. These bioinformatic analyses were performed via the Gene Expression Profiling Interactive Analysis (GEPIA) website [41] (http://gepia.cancer-pku. cn). The correlation between the B7-H4 mRNA level and lymph node metastasis status in the TCGA COAD dataset was performed via the XENA website.

Statistical analysis

Statistical analyses of the experimental results were performed with GraphPad Prism version 5.0. Associations between p-PKC δ expression, B7-H4 expression and various clinicopathological parameters were evaluated by the χ^2 test. Correlations were evaluated by the Spearman rank correlation coefficient. Differences between groups were evaluated using two-tailed unpaired Student's t test. Replicate experiments were analyzed using paired Student's t test. All significance tests were two-tailed, and P < 0.05 was considered significant.

Results

B7-H4 and p-PKC δ were upregulated in CRC

First, we analyzed correlations between the expression of PKCs and that of B7-H4 in CRC patient samples based on the LinkedOmics and GEPIA databases and found that the expression of both PKRCA and PRKCD (PKCδ) was positively associated with B7-H4 expression, but only PKCδ was abnormally upregulated in colorectal cancer tissues (Additional file 1: Figure S1). Then, we evaluated the B7-H4 and p-PKC δ protein levels by IHC staining in 225 clinical colorectal tumor tissue specimens and 36 adjacent normal tissue specimens. Representative images of IHC staining are shown in Fig. 1A. Positive B7-H4 expression was detected in 132 of the CRC tissue specimens (132/225, 58.7%), and B7-H4 was expressed in the membrane, cytoplasm and nucleus of colorectal tumor cells (Fig. 1A). Positive staining for p-PKC δ was detected in 139 of the CRC tissue specimens (139/225, 61.8%), and p-PKC δ was expressed in the membrane and cytoplasm of colorectal tumor cells (Fig. 1A). Comparison of the tumor tissues and adjacent normal tissues revealed that the expression of both B7-H4 and p-PKCδ was significantly increased in the tumor tissue samples (Fig. 1B), consistent with previous reports [14, 25, 27, 40]. Furthermore, quantitative PCR analysis showed that the expression of both B7-H4 and PKC δ was significantly increased in the tumor tissues compared with the adjacent normal tissues (Fig. 1C).

The association between B7-H4 and p-PKCδ in CRC

Spearman correlation analysis of the IHC staining data showed that B7-H4 protein expression was significantly correlated with p-PKC δ protein expression (r=0.378, P<0.001). Analysis with the Pearson chi-square test also showed that B7-H4 protein expression was significantly correlated with p-PKC δ protein expression (P<0.001) (Table 1). Of the 139 p-PKC δ -positive tumor samples, 101 (72.7%, 101/139) showed positive B7-H4 expression, whereas of the 86 p-PKC δ -negative tumor samples, only 31 (36.0%, 31/86) showed positive B7-H4 expression. Analysis of the TCGA colon adenocarcinoma (COAD) dataset showed that the PRKCD mRNA level was also positively correlated with the VTCN1 (encoding B7-H4) mRNA level (Fig. 2A).

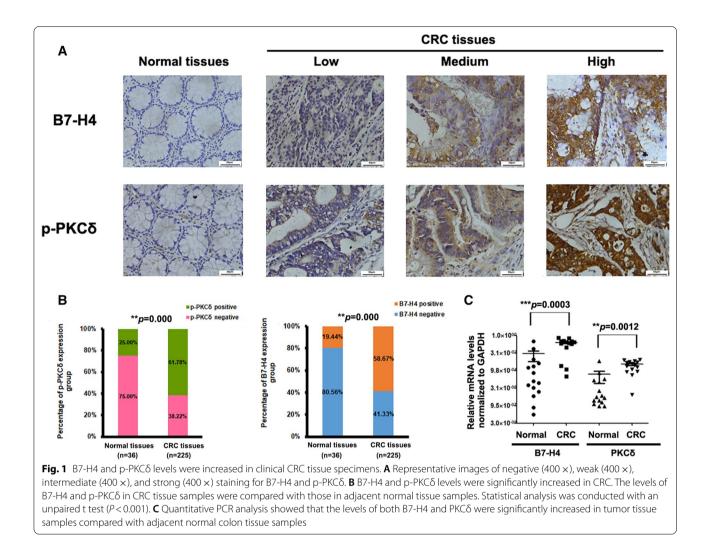


Table 1 The expression of B7-H4 and p-PKC δ in clinical CRC tissues

p-PKCδ expression	B7-H4 expression	P value ^a		
	Negative(n = 93)	Positive(n = 132)		
Negative (n = 86)	55 (64.0)	31 (36.0)	0.000 ^b	
Positive (n = 139)	38 (27.3)	101 (72.7)		

^a P value is obtained by Pearson chi-square, Asymp. Sig., two tailed

^b P < 0.05 was designated as significance

To verify the finding that the B7-H4 protein level is correlated with the p-PKC δ level in clinical samples, we examined B7-H4 and p-PKC δ levels in CRC cell lines. As shown in Fig. 2B and Additional file 2: Figure S2, the B7-H4 protein level was higher in the HCT116 and SW620 cell lines, intermediate in the SW480 and RKO cell lines, and lower in the NCM460 cell line. Consistent

with this pattern, the p-PKC δ level was higher in HCT116 and SW620 cells than in the other cells, verifying the correlation between the p-PKC δ and B7-H4 protein levels in CRC cell lines.

Collectively, the results obtained from analysis of clinical CRC samples and CRC cell lines showed that the B7-H4 level was correlated with the p-PKC δ level in CRC.

As we found that the B7-H4 protein level was significantly correlated with the p-PKC δ level, we further investigated whether B7-H4 and p-PKC δ are coexpressed in CRC tissue. Serial sections of p-PKC δ ⁺B7-H4⁺ CRC samples were subjected to a serial staining protocol as follows: IHC staining for p-PKC δ , IHC staining for B7-H4, and double IF staining for p-PKC δ and B7-H4. Representative images of stained sections are shown in Fig. 2C. IHC staining showed that the area of positive staining was similar for the two molecules. Merged images of double IF staining showed that the green and red fluorescence signals overlapped. Both the IHC and IF staining results demonstrated that p-PKC δ and B7-H4 were coexpressed in CRC tissue samples.

The results of serial CRC sample staining further indicated that B7-H4 expression in CRC tissues was correlated with PKC δ activation.

Positivity for B7-H4 and p-PKC δ was associated with tumor metastasis in CRC samples

Next, we examined the associations of B7-H4 and p-PKCδ with clinical parameters. As shown in Table 2, in the 225 CRC samples, positive B7-H4 expression was correlated with moderate/poor differentiation ($\chi 2 = 8.992$, P = 0.003), lymph node metastasis ($\chi^2 = 8.919$, P = 0.003) and advanced Dukes' stage ($\chi^2\!=\!5.427,\ P\!=\!0.02)$ and that positivity for p-PKC\delta was correlated with advanced Dukes' stage ($\chi^2 = 4.118$, P = 0.042). Notably, coexpression of B7-H4 and p-PKC8 was significantly associated with moderate/poor differentiation ($\chi^2 = 5.072$, P = 0.024), lymph node metastasis ($\chi^2 = 10.909$, P = 0.001) and advanced Dukes' stage ($\chi^2 = 10.017$, P = 0.002). TCGA COAD dataset analysis via the XENA website showed that the B7-H4 mRNA level was also positively correlated with lymph node metastasis (Fig. 2D). These results suggested that B7-H4 and p-PKC8 were associated with tumor metastasis.

Activation of PKC δ induced B7-H4 expression in CRC cell lines

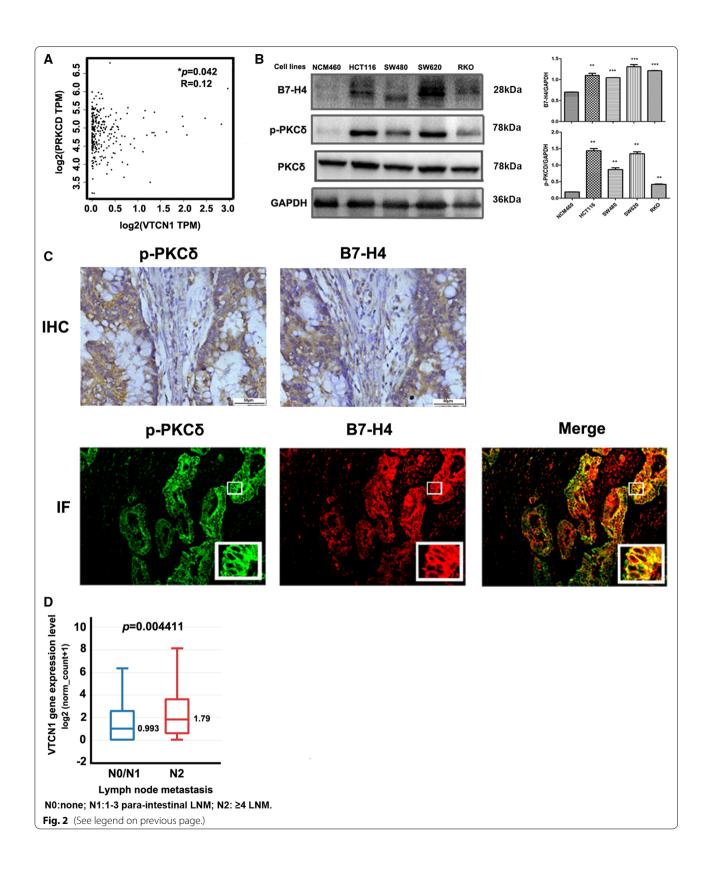
Since TPA can induce strong PKC δ localization mainly to the plasma membrane [26], we examined the effect of TPA on B7-H4 expression in CRC cell lines. HCT116 and SW620 cells were treated with various concentrations of TPA for 20 h, and B7-H4 protein levels were determined by Western blotting. As shown in Fig. 3A and Additional file 3: Figure S3A, treatment with 10 and 50 nM TPA for 20 h effectively increased the p-PKC δ level, whereas only treatment with 50 nM TPA effectively increased the B7-H4 level. Treatment with 100 nM TPA for 20 h began to deactivate PKC δ , possibly due to the effect of chronic treatment with TPA [42, 43]. Similarly, time point at which B7-H4 expression began to decrease was delayed, and the required concentration increased to 200 nM. The explanation for the delay between these decreases in B7-H4 expression and PKCS deactivation might be due to the time needed for signal transduction downstream of PKCS. Therefore, the delay strongly confirmed that B7-H4 was regulated by the PKC δ signaling pathway. B7-H4 and nuclei in HCT116 cells were stained with a PE-conjugated anti-B7-H4 antibody and DAPI, respectively, and the cells were then observed under a confocal microscope (Fig. 3B). The results showed that TPA treatment significantly increased the percentage of B7-H4 positive cells. Furthermore, cell lines were treated with rottlerin, a PKC δ inhibitor. Rottlerin inhibited PKCα, PKCβ, PKCγ, PKCδ, PKCη, CKII and PKA and preferentially inhibited PKC δ activity at low concentrations but inhibited the other PKC isoforms at high concentrations^[44]; thus, we used rottlerin at a concentration of 1-2 µM (low concentration) to investigate the inhibitory effect of PKCδ on B7-H4. The results showed that rottlerin effectively decreased the B7-H4 protein level in a concentration-dependent manner in the HCT116 and SW620 cell lines (Fig. 3C and Additional file 3: Figure S3B). Flow cytometric analysis also demonstrated that TPA increased B7-H4 expression and that rottlerin decreased B7-H4 expression in the HCT116 and SW620 cell lines (Fig. 3D). Then, the HCT116 and SW620 cell lines were treated first with $1 \,\mu M$ rottlerin and then with 100 nM TPA. This experiment showed that rottlerin effectively abrogated the TPA-induced increase in B7-H4 expression (Fig. 3E and Additional file 3: Figure S3C).

$PKC\delta$ activation upregulated B7-H4 expression in CRC cell lines

After using a PKC δ activator and a PKC δ inhibitor, we knocked down PKC δ expression and examined the effect on B7-H4 expression in CRC cell lines. PKC δ mRNA expression was knocked down in HCT116 and SW620 cells as described in the Materials and Methods section. The PKC δ mRNA level was decreased dramatically in the treated group compared with the mock group (Fig. 4A). Western blot analysis showed that the PKC δ and B7-H4 protein levels were significantly decreased in the PKC δ siRNA/HCT116 and PKC δ siRNA/SW620 cell lines (Fig. 4B and Additional file 4: Figure S4A). The con siRNA and PKC δ siRNA cell lines were treated with TPA, and PKC δ knockdown was found to abrogate the

Fig. 2 Correlation between B7-H4 and p-PKC δ levels. **A** TCGA dataset analysis was performed on the GEPIA website (http://gepia.cancer-pku.cn/). This analysis showed that PRKCD (PKC δ) mRNA expression was positively correlated with VTCN1 (encoding B7-H4) mRNA expression. **B** The protein levels of B7-H4 and p-PKC δ in the NCM460, SW480, HCT116, SW620 and RKO cell lines were determined by Western blot analysis. The data are expressed as the mean \pm SD values; n = 3. **C** Serial p-PKC δ^+ tumor sections were examined for B7-H4 expression, and positive staining (400 ×) of B7-H4 and p-PKC δ in CRC samples is shown. Double IF staining of CRC samples was performed. CRC tissue samples were stained for p-PKC δ (green, 200 ×) and B7-H4 (red, 200 ×). B7-H4⁺/p- PKC δ^+ cells were identified in CRC tissue specimens (dark yellow; original magnification). **D** TCGA COAD dataset analysis was performed on the XENA website. This analysis showed that the B7-H4 mRNA level was positively correlated with lymph node metastasis

⁽See figure on next page.)



Variables	B7-H4 (%)			ρ-ΡΚϹδ (%)			p-PKCδ ⁺ B7-H4 ⁺ (%)		
	Positive (n = 132)	Negative (n=93)	P value ^a	Positive (n = 139)	Negative (n=86)	P value ^a	Positive (n = 101)	Others (n = 124)	P value ^a
Gender									
Male (n = 122)	72 (59.0%)	50 (41.0%)	0.908	78 (63.9%)	44 (36.1%)	0.469	53 (43.4%)	69 (56.6%)	0.635
Female (n = 103)	60 (58.3%)	43 (41.7%)		61 (59.2%)	42 (40.8%)		48 (46.6%)	55 (53.4%)	
Histology									
Well (n = 36)	13 (36.1%)	23 (63.9%)	0.003 ^b	22 (61.1%)	14 (38.9%)	0.928	10 (27.8%)	26 (72.2%)	0.024 ^b
Moder- ate/poor (n = 189)	119 (63.0%)	70 (37.0%)		117 (61.9%)	72 (38.1%)		91 (48.1%)	98 (51.9%)	
Depth of tumor									
T1 + T2 (n = 51)	26 (51%)	25 (49%)	0.205	31 (52.6%)	20 (47.4%)	0.868	19 (15.8%)	32 (84.2%)	0.213
T3 + T4 (n = 174)	106 (60.9%)	68 (39.1%)		108 (69.1%)	66 (30.9%)		82 (47.1%)	92 (52.9%)	
Lymph node me	etastasis								
Negative (n = 136)	69 (50.7%)	67 (49.3%)	0.003 ^b	79 (58.1%)	57 (41.9%)	0.159	49 (36.0%)	87 (64.0%)	0.001 ^b
Positive (n = 89)	63 (70.8%)	26 (29.2%)		60 (67.4%)	29 (32.6%)		52 (58.4%)	37 (41.6%)	
Dukes' stage									
A + B (n = 122)	63 (51.6%)	59 (48.4%)	0.020 ^b	68 (55.7%)	54 (44.3%)	0.042 ^b	43 (35.2%)	79 (64.8%)	0.002 ^b
C + D (n = 103)	69 (67.0%)	34 (33.0%)		71 (68.9%)	32 (31.1%)		58 (56.3%)	45 (43.7%)	

^a P value is obtained by Pearson chi-square, Asymp. Sig., two tailed

 $^{\rm b}\it P\,{<}\,0.05$ was designated as significance

TPA-induced increase in B7-H4 expression (Fig. 4C and Additional file 4: Figure S4B). Taken together, these data suggested that PKC δ knockdown downregulated B7-H4 expression in CRC cells.

STAT3 mediated PKC δ -induced B7-H4 upregulation in CRC cell lines

We next explored the signaling pathway by which PKC δ mediates the expression of B7-H4. A previous study showed that activated STAT3 can bind to the B7-H4 promoter and enhance the expression of the B7-H4 protein in microglial cells [45]. In addition, PKC δ is a primary regulator of STAT3 phosphorylation in keratinocytes and luteal cells [46, 47]. Therefore, we hypothesized that activated PKC δ can induce B7-H4 expression by increasing the phosphorylation of STAT3 in CRC cells. Western blot analysis showed that the levels of B7-H4 and p-STAT3 were decreased in PKC δ siRNA CRC cells (Fig. 5A and Additional file 5: Figure S5A). Moreover, we observed that cryptotanshinone, a STAT3 phosphorylation inhibitor, significantly decreased the B7-H4 protein level in a concentration-dependent manner in HCT116 and

SW620 cells (Fig. 5B and Additional file 5: Figure S5B). These results suggested that PKC δ could regulate the expression of B7-H4 via the STAT3 signaling pathway in CRC cells.

The PKCδ/B7-H4 axis promoted CRC cell motility

As analysis of clinical samples and datasets showed that B7-H4 and p-PKCδ were associated with CRC metastasis, we further examined whether CRC cell invasion and migration are mediated via the PKC8/B7-H4 axis. Previously, the PKC δ siRNA/HCT116 and PKC δ siRNA/ SW620 cell lines were established. Here, we established the B7-H4 siRNA/HCT116 and B7-H4 siRNA/SW620 cell lines (Fig. 6A and Additional file 6: Figure S6A). A Transwell assay showed that knockdown of B7-H4 or PKC δ expression inhibited the constitutive invasion of CRC cells (Fig. 6B and Additional file 6: Figure S6B), suggesting that B7-H4 and PKC8 promoted cell motility. In addition, we found that pharmacological induction of PKC δ expression by 24 h of TPA treatment effectively enhanced cell invasion (Fig. 6C). However, knockdown of B7-H4 prevented the increase in invasion (Fig. 6C),

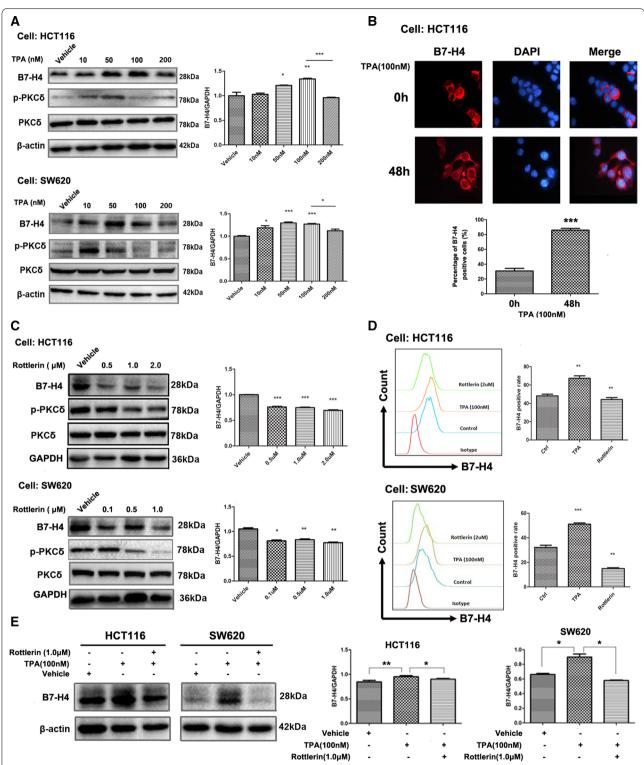
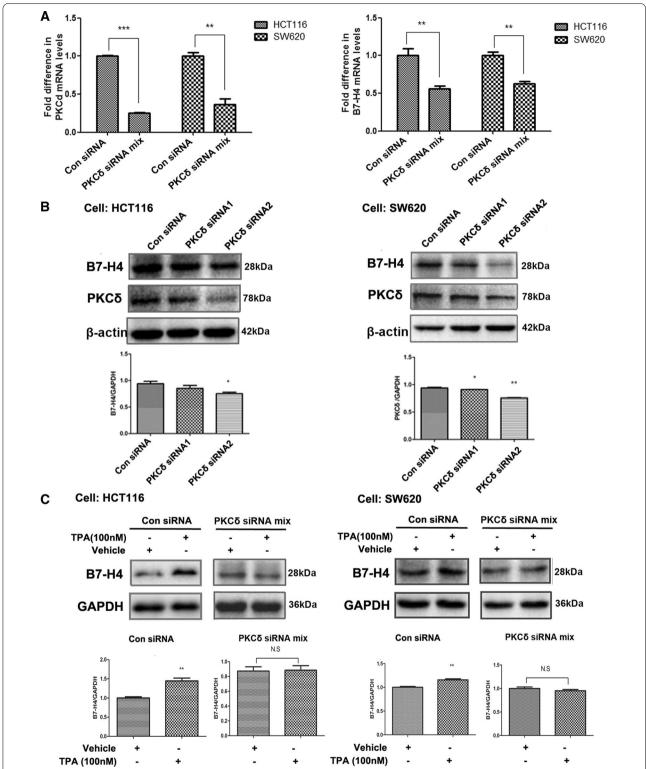
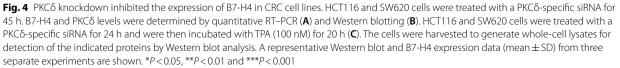
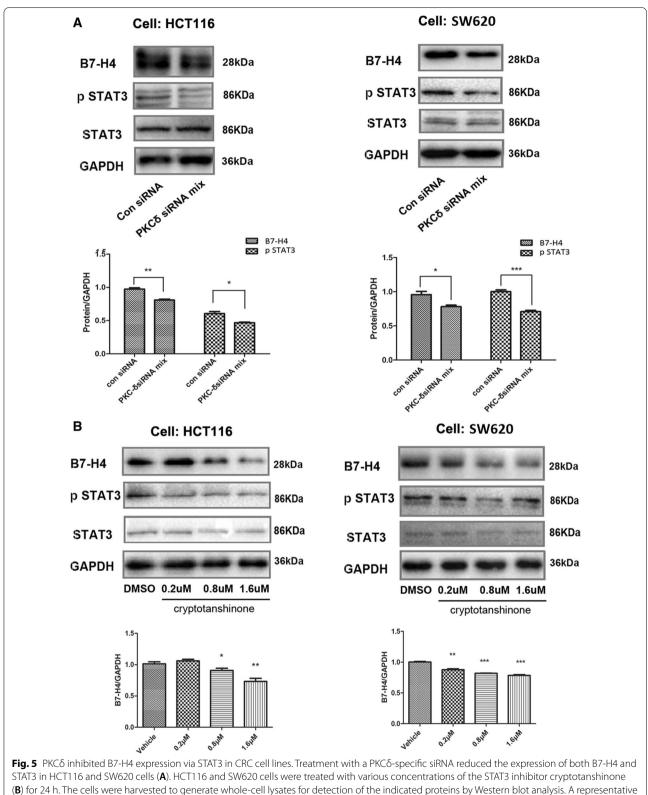


Fig. 3 PKCS mediates B7-H4 upregulation in CRC cell lines. HCT116 and SW620 cells were treated with various concentrations of TPA (**A**) or rottlerin (**C**) for 20 h. **B** B7-H4 and nuclei in HCT116 cells were stained with a PE-conjugated anti-B7-H4 antibody and DAPI, respectively. **D** Flow cytometric analysis was used to detect changes in B7-H4 expression in TPA- or rottlerin-treated HCT116 and SW620 cells. CRC cells were treated with TPA or rottlerin for 24 h, and intracellular B7-H4 expression was examined by flow cytometry. **E** The HCT116 and SW620 cell lines were treated with 1 μ M rottlerin and 100 nM TPA for 24 h, and B7-H4 levels were determined by Western blotting. A representative Western blot image is shown in the left panel. Statistical analysis of B7-H4 expression (mean ± SD) from three separate experiments is shown in the right panel. **P* < 0.05, ***P* < 0.01 and ****P* < 0.001







(B) for 24 h. The cells were harvested to generate whole-cell lysates for detection of the indicated proteins by Western blot analysis. A re Western blot and B7-H4 expression data (mean \pm SD) from three separate experiments are shown. *P < 0.05, **P < 0.01 and ***P < 0.001

suggesting that B7-H4 plays a role in PKCδ activationinduced cell invasion.

To evaluate the effects of PKCδ and B7-H4 on cell migration, we performed a wound healing assay. Images showing wound closure are shown in Fig. 6D and Additional file 6: Figure S6B. Knockdown of PKCδ or B7-H4 expression effectively inhibited cell motility. TPA treatment effectively enhanced the motility of con siRNA/HCT116 cells, but the effect was obviously reduced in B7-H4 siRNA/HCT116 and PKCδ siRNA/HCT116 cells. Cell viability was also assessed by a wound healing assay. There were no differences between the TPA treatment groups and the control groups (data not shown).

Rottlerin inhibited B7-H4 expression and tumor metastasis in mice

To validate the effect of the PKC δ /B7-H4 axis on tumor metastasis in vivo, we injected rottlerin into HCT116 tumor-bearing nude mice. Rottlerin treatment significantly reduced lung metastasis (Fig. 7A and B). The IHC results showed that the levels of human B7-H4 and p-PKC δ were decreased in the lung tissues of rottlerin-treated mice (Fig. 7C; lung tissues from three mice per group). These results show that rottlerin treatment inhibited colon cancer metastasis compared with that in the control group via the PKC δ /B7-H4 axis.

Discussion

CRC cell migration and invasion are related to the occurrence of postsurgical metastasis and poor survival in CRC patients; thus, we need to find new CRC therapies to block metastasis. As a novel member of the PKC family, PKC δ can be activated independent of Ca²⁺ and phospholipids and has multiple functions associated with cancer progression, including functions in the proliferation, survival, apoptosis and motility of cancer cells [30]. PKC δ is involved in colon epithelial cell migration via the IGF-I signaling pathway [34]. Although some studies found that PKC δ mRNA expression was decreased in most primary CRC tumors and some CRC cell lines, the p-PKC δ level was increased in a subset of colorectal cancers, and this increase can also enhance the migration and invasiveness of colon carcinoma cells by enhancing KITENIN expression and phosphorylation of HuR and Trop-2 [26, 27, 36]. Knockdown of B7-H4 effectively inhibits the proliferation, invasion, and migration of CRC cells, gastric cancer cells and lung cancer cells via various signaling pathways [20, 21, 48]. Both PKC δ and B7-H4 contribute to tumor metastasis; therefore, we sought to investigate the correlation among PKC δ , B7-H4 and metastasis in CRC.

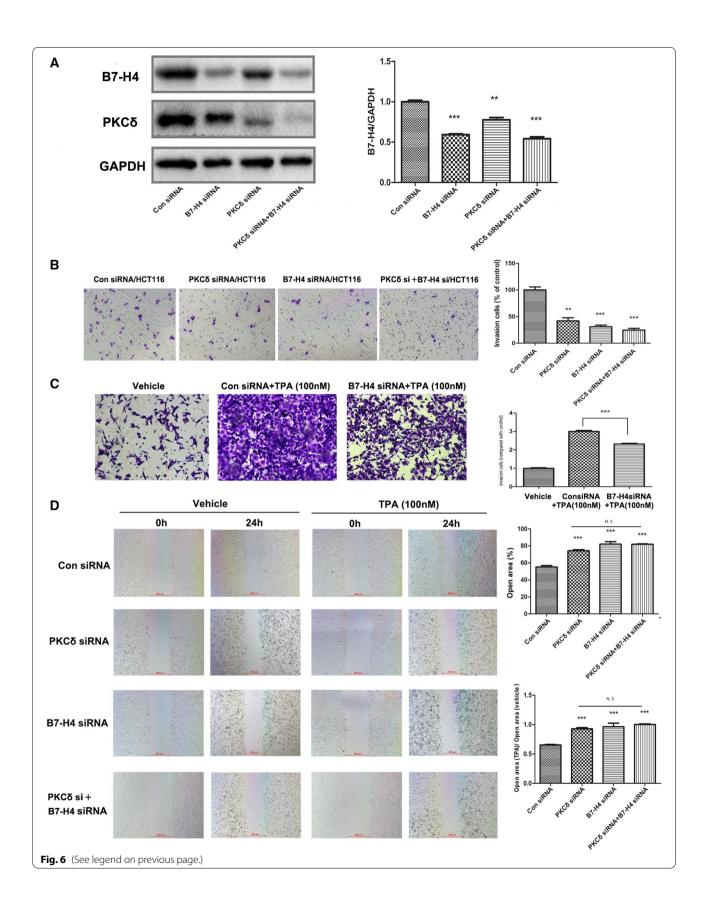
In this study, we evaluated the protein levels of p-PKC δ and B7-H4 in tumor cells and found increased levels of p-PKC δ in some colorectal tumor samples (139/225, 61.8%). IHC analysis of serial sections from identical tissues showed that 101 of the p-PKC δ^+ samples also expressed B7-H4 (72.7%, 101/139), and the IF staining results further confirmed this finding. The IHC results showed that a p-PKC δ^+ B7-H4⁺ phenotype in colorectal tumor samples was significantly associated with moderate/poor differentiation, lymph node metastasis and advanced Dukes' stage. Thus, we speculated that the magnitude of PKC δ activation is related to the B7-H4 level and plays an important role in cancer progression.

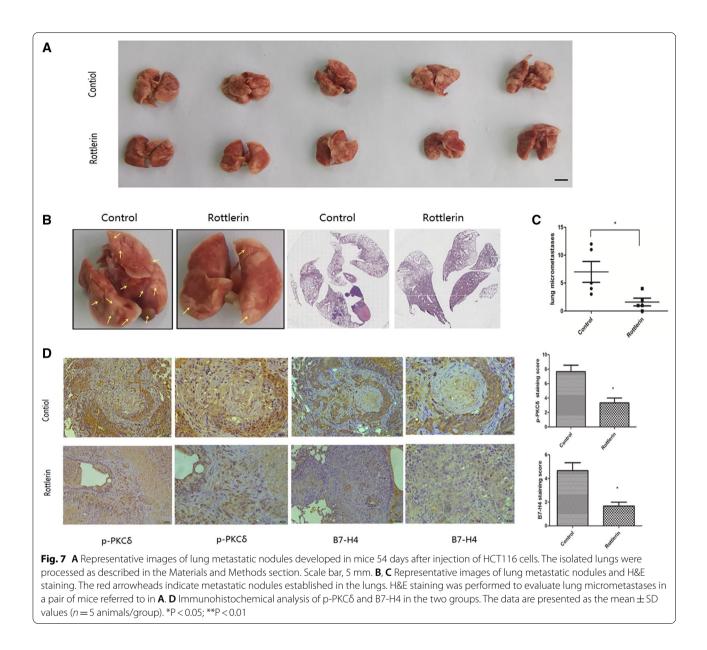
It has been reported that B7-H4 expression can be upregulated by many inflammatory mediators. In a renal cell carcinoma cell line, IFN-a, IL-2, and IFN-y were found to upregulate B7-H4 expression [49]. In human ovarian cancer and glioma cancer, tumor-associated Tregs trigger macrophages to secrete IL-10 and IL-6, which activate STAT3 and induce B7-H4 transcription [50, 51]. In human lung cancer, tumor-associated macrophages secrete TNF- α , IL-10, and IFN- γ , which induce B7-H4 expression in lung cancer cells [52]. Our previous study revealed that B7-H4 can be upregulated by IGF1R activation through the MEK/ERK1/2 signaling pathway in lung cancer [53]. In multiple myeloma, hypoxia-inducible factor-1 α (HIF-1 α) can bind to the B7-H4 promoter and induce B7-H4 expression [54]. In addition, the NF-κB (P65) pathway could increase PD-L1 and B7-H4 levels in hepatocellular carcinoma (HCC) tissues [55]. A recent study demonstrated that TGF-\u03b31-driven SMAD3/4 signaling can increase B7-H4 expression in CRC [56]. Most of these studies identified the triggering factors contributing to B7-H4 expression, but the exact signaling pathways involved in the regulation of B7-H4 expression still require further elucidation.

In this study, we found that the protein levels of p-PKC δ and B7-H4 were higher in CRC cell lines than in a normal cell line. The PKC activator TPA increased the B7-H4 level in HCT116 and SW620 cells in a concentration-dependent manner. To confirm that B7-H4 is

Fig. 6 The PKC δ /B7-H4 axis promoted the migration of HCT116 cells. HCT116 cells were treated with a PKC δ -specific siRNA and/or a B7-H4-specific siRNA for 45 h, and B7-H4 protein levels were then determined by Western blot analysis (**A**). A Transwell assay was performed to examine the constitutive invasion of B7-H4 siRNA/HCT116, PKC δ siRNA/HCT116, PKC δ siRNA + B7-H4 siRNA/HCT116 and con siRNA/HCT116 cells (**B**). The invasion of B7-H4 siRNA/HCT116 cells was evaluated after the cells were treated with 100 nM TPA for 24 h (**C**). A wound healing assay was performed to evaluate the effects of PKC δ and B7-H4 on cell migration (**D**). Experiments were performed in triplicate. **P* < 0.05, ***P* < 0.01 and ****P* < 0.001

⁽See figure on next page.)





regulated by activation of PKC δ , the specific PKC δ inhibitor rottlerin and a PKC δ -specific siRNA were used. Initially, rottlerin was found to inhibit PKC partially due to competition with ATP for the ATP-binding site in PKC, and the structure of rottlerin is much more selective for the PKC δ isozyme [44]. However, a series of subsequent studies showed that rottlerin can inhibit agonist-induced PKC δ translocation, thereby inhibiting PKC δ activity [57, 58]. Regardless of the exact mechanism, all of these studies have shown that rottlerin can selectively inhibit PKC δ activation. Therefore, in this study, we used rottlerin as a specific PKC δ inhibitor. The results showed that rottlerin decreased B7-H4 expression and abrogated the TPA-induced increase in B7-H4 expression. Furthermore, treatment with the PKC δ -specific siRNA also effectively decreased the B7-H4 level and abrogated the TPA-induced increase in B7-H4 expression. Collectively, these results verified the findings reported for clinical samples and confirmed that B7-H4 expression can be upregulated by PKC δ activation. Moreover, PKC δ siRNA treatment also reduced the activity of STAT3. Treatment with the STAT3 inhibitor cryptotanshinone significantly decreased the B7-H4 protein level in a concentrationdependent manner in CRC cell lines. These results suggested that PKC δ could regulate the expression of B7-H4 via STAT3. We also investigated whether PKC δ promotes

B7-H4 expression through the MEK/ERK1/2 pathway in CRC cells, but we did not find a related change (data not shown). Previous studies revealed that PKC8 plays an important role in colon cancer cell migration and invasion. B7-H4 could increase cell migration and invasion by targeting the angiogenic factors MMP2, MMP9 and VEGF. B7-H4 overexpression activates a variety of signaling pathways, such as the NF-KB, ERK1/2, AKT/ STAT3 and PI3K/AKT/mTOR pathways, to promote epithelial-mesenchymal transition (EMT) and invasion of cancer cells [20, 22, 59, 60]. Here, we also performed Transwell invasion and wound healing assays to evaluate the motility of CRC cells. We found that knockdown of B7-H4 or PKC δ suppressed cell motility and suppressed the enhancing effect of TPA on cell invasion and migration. We also found that rottlerin treatment significantly inhibited B7-H4 expression and tumor metastasis in vivo. The rottlerin dose (20 mg/kg) used in the present study was lower than the safe dose established in a previous study.

In conclusion, our results identify a canonical PKC δ / STAT3/B7-H4 signaling pathway that is constitutively active in colorectal carcinoma cells. B7-H4 expression was upregulated by PKC δ activation and contributed to PKC δ -induced cell motility, which plays a role in the immune escape of CSCs. This result suggests that B7-H4 and PKC δ may be therapeutic targets in tumor metastasis. It might be important to consider the effect on B7-H4 expression when a PKC δ inhibitor is used clinically.

Abbreviations

CRC: Colorectal cancer; PKC: Protein kinase C; TPA: Phorbol-12-Myristate-13-Acetate; APCs: Antigen-presenting cells; Treg: Regulatory T cells; siRNA: Small interfering RNA; IHC: Immunohistochemistry; IF: Immunofluorescence; FBS: Fetal bovine serum; SDS-PAGE: Sodium dodecyl sulfate–polyacrylamide gel; PVDF: Polyvinylidene difluoride; FCS: Flow cytometry analysis; HIF-10: Hypoxiainducible factor-10; HCC: Hepatocellular carcinoma.

Supplementary Information

The online version contains supplementary material available at https://doi.org/10.1186/s12935-022-02567-1.

Additional file 1: Figure S1. Correlation analysis of the expression of PKCs and B7-H4 in CRC based on the LinkedOmics and GEPIA databases. A Correlation analysis between PKCs and B7-H4; B comparison of PKRCA and PRKCD expression in cancer and adjacent tissues.

Additional file 2: Figure S2. Western blot analysis was performed to detect the expression of B7-H4 in CRC cell lines. The protein levels of B7-H4 and p-PKC δ in the NCM460, SW480, HCT116, SW620 and RKO cell lines were determined.

Additional file 3: Figure S3. PKC δ mediated B7-H4 upregulation in CRC cell lines. HCT116 and SW620 cells were treated with various concentrations of TPA (**A**) or rottlerin (**B**) for 20 h. **C** The HCT116 and SW620 cell lines were treated with 1 μ M rottlerin and 100 nM TPA for 24 h, and B7-H4 levels were determined by Western blotting.

Additional file 4: Figure S4. PKC& knockdown inhibited the expression of B7-H4 in CRC cell lines. HCT116 and SW620 cells were treated with a PKC&-specific siRNA for 45 h. B7-H4 and PKC& levels were determined by Western blotting (**A**). HCT116 and SW620 cells were treated with a PKC&-specific siRNA for 24 h and were then incubated with TPA (100 nM) for 20 h (**B**, **C**). The cells were harvested to generate whole-cell lysates for detection of the indicated proteins by Western blot analysis.

Additional file 5: Figure S5. PKC δ inhibited B7-H4 expression via STAT3 in CRC cell lines. Treatment with a PKC δ -specific siRNA reduced the expression of both B7-H4 and STAT3 in HCT116 and SW620 cells (A). HCT116 and SW620 cells were treated with various concentrations of the STAT3 inhibitor cryptotanshinone (B) for 24 h. The cells were harvested to generate whole-cell lysates for detection of the indicated proteins by Western blot analysis.

Additional file 6: Figure S6. The PKC\delta/B7-H4 axis promoted the migration of SW620 cells. HCT116 and SW620 cells were treated with a PKCδ-specific siRNA and/or a B7-H4-specific siRNA for 45 h, and B7-H4 protein levels were then determined by Western blot analysis (**A**). A Transwell assay was performed to examine the constitutive invasion of B7-H4 siRNA/SW620, PKCδ siRNA/SW620, PKCδ siRNA + B7-H4 siRNA/SW620 and con siRNA/SW620 cells (**B**). A wound healing assay was performed to evaluate the effects of PKCδ and B7-H4 on cell migration (**C**). The viability of HCT116 cells in different groups was assessed by a CCK-8 assay (**D**). Experiments were performed in triplicate. *P < 0.05, **P < 0.01 and ***P < 0.001.

Additional file 7: Table S1. The primers of real-time PCR and the siRNAs.

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Author contributions

XZ and LZ designed research; BZ, YL, ZZ and HW performed research; BZ, TS and WC contributed to data analysis and interpretation; BZ and LZ wrote the paper. All authors read and approved the final manuscript.

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Availability of data and materials

The datasets generated during the current study can be discovered in online repositories, further inquiries can be available from the corresponding author.

Declarations

Ethics approval and consent to participate

This work obtained ethics approval by The First Affiliated Hospital of Soochow University, Reference Number 2014865082. All patients included were consented to participate in the study and to use their materials in research.

Consent for publication

Not applicable.

Competing interests

The authors declare that they have no competing interests.

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