Abstract. Considerable evidence has implied that α7 nicotinic receptor subtypes play an important role in chronic inflammatory and neuropathic pain signaling. The aim of the present study was to determine the role of endogenous α7nAChR signaling in tumor-associated macrophages (TAMs) in human colorectal cancer (CRC) metastasis and prognosis. α7nAChR expression in primary tumor cells and adjacent stroma cells especially in TAMs in 51 CRC patients was observed. Using a human monocyte THP-derived macrophages (TMs) with α7nAChR-siRNA knockdown (TMα7-/-) and a CRC cell Transwell co-culture model, the migration and invasion of two CRC cell lines, LoVo and SW620, were determined. Western blotting was carried out to investigate the expression of multiple molecules involved in the NF-κB, STAT3, PI3K signaling pathways in mimic TAMs, i.e., TMs exposed to in-direct LoVo cell stimulation. A nicotinic α7 receptor antagonist [α-bungarotoxin (α-Btx)] and three pharmaceutical inhibitors: AG490 (JAK2/STAT3 inhibitor), LY294002 (PI3K inhibitor) and Bay 11-7082 (NF-κB inhibitor) were applied to evaluate whether these signaling pathways were associated with the enhanced migration of CRC cells when co-cultured with α7nAChR knockdown TMs. The results revealed that the expression of α7nAChR in TAMs differed in patients. However, CRC patients who had a high incidence of hepatic metastasis showed no or low expression of α7nAChR in TAMs. TMs with α7nAChR-siRNA knockdown (TMα7-/-) significantly enhanced the migration and invasion of the two CRC cell lines LoVo and SW620. α7nAChR knockdown in TMs significantly downregulated phosphorylation of STAT3, PI3K p85 and NF-κB p65 after co-culturing with LoVo cells. Inhibition of JAK2/STAT3 prevented the TMα7-/--enhanced migration of LoVo cells. α7nAChR expressed in TAMs in human CRC patients plays an important role in preventing metastasis and could be a prognostic marker in CRCs, which may be regulated by the JAK2/STAT3 signaling pathway.

Introduction

Substantial evidence suggests that stromal cells including inflammatory cells and fibroblasts can interact with adjacent cancer cells (1). The inflammatory cells such as macrophages, neutrophils, and lymphocytes exist in the tumor microenvironment and play an indispensable role in cancer progression. Macrophages found in close proximity or within tumor masses are indicated as tumor-associated macrophages (TAMs) (2,3). Since pro-inflammatory macrophages have pronounced antitumor activity and are cytotoxic to tumor cells, TAMs are believed to play a pivotal role in preventing tumor development (4). Nevertheless, persistence of an inflammatory response appears to be detrimental and causes cancer initiation and/or progression probably through the generation of mutation-inducing reactive oxygen species and nitrogen-free radicals (5). Indeed, in some types of carcinomas, TAMs appear to not only fail to kill tumor cells but also contribute to tumor progression by promoting cell proliferation, tumor growth, angiogenesis and metastasis (6,7). Whether and how TAMs play an important role in colorectal cancer (CRC) remains controversial. According to an early study, TAMs may act as one line of defense against malignant cells in CRC patients (8). TAMs infiltrating at the invasive front in colon cancer are associated with improvement in both hepatic metastasis and overall survival (9). In contrast, Ishimaru et al found that TAMs induce the expression of osteopontin contributing to metachronous liver metastases in CRC (10). However, it has been reported that both CD169- and CD40-positive macrophages are associated with a favorable prognosis in colorectal carcinoma patients (10,11), indicating that TAMs
Table I. Clinical and pathological features of the examined patients.

| Patient characteristics         | All patients (n=51) | Patients with α7nAChR (TAM/stroma) | P-value |
|----------------------------------|---------------------|-----------------------------------|---------|
| Sex                              |                     |                                   |         |
| Male                             | 32                  | Grade ≤1 (n=33)                   |         |
| Female                           | 19                  | Grade ≥2 (n=18)                   |         |
| Mean (± SD) age at diagnosis (years) | 65±13.5            | 60±13.4                           | 57±14.0 | 0.4555 |
| Tumor metastasis (no.)³          |                     |                                   |         |
| Dukes’ A or B or C               | 37                  | 20                                | 17      | 0.0101 |
| Dukes’ D                         | 14                  | 13                                | 1       |        |
| 5-year survival (%)²             | 50.98               | 42.42                             | 77.77   | 0.0201 |

TAM, tumor-associated macrophage.

have protective potential in colon cancers and may serve as a novel therapeutic target.

Nicotinic acetylcholine (ACh) receptors are a family of integral membrane proteins, which function as ligand-gated, pentameric ion channels responding to the binding of ACh, a neurotransmitter. In humans, 16 different subunits of nicotinic ACh receptors (α1-7, α9-10, β1-4, δ, ε, γ) are expressed on non-neuronal cells both within and outside the nervous system. The non-neuronal nAChRs have considerable implications for several diseases induced by tobacco usage such as cancers and cardiovascular disease through the non-neuronal nAChR signaling pathway (12). For example, AChRs, particularly, α7nAChR, can mediate nicotine-dependent upregulation of proliferative and survival genes that contribute to the growth and progression of lung cancer cell lines in vitro and in vivo (13-16). α7nAChR is an important factor in the regulation of cell signaling, which is expressed in many different non-neuronal cells such as vascular and brain endothelial cells, bronchial epithelial cells, keratinocytes, astrocytes, synoviocytes, thymocytes, lymphocytes, bone marrow cells, monocytes, macrophages and microglia (17-19). An α7nAChR unbalance may be involved in different diseases such as Alzheimer’s, Parkinson’s and cancer (20). α7nAChR is also an important component of the underlying mechanism that regulates the anti-inflammatory efficacy of the cholinergic anti-inflammatory pathway (21,22). It has also been reported that α7nAChR silencing inhibits lung cancer proliferation (23). Thus, α7nAChR offers rational bases to develop new drugs and therapeutic strategies for different diseases including cancers.

Recently, it was reported that nicotine increases cell proliferation and inhibits apoptosis through the activation of the α7nAChR/ERK/AKT pathway in human colon carcinoma cells (24). Our laboratory also found that nicotine enhances the invasion and metastasis of human CRC cells through the nicotinic ACh receptor downstream p38 MAPK signaling pathway (25). Inhibition of α7nAChR induces apoptosis in non-small cell lung carcinoma and may provide a feasible rationale for preventing the progression of head and neck squamous cell carcinoma (HNSCC) (26,27). Recently, Salaga et al reported that encenicline, an α7 nicotinic ACh receptor partial agonist, decreased macrophage infiltration in the colon and improved experimental colitis in mice (28). However, although α7nAChR has been proposed as a pharmacological target for inflammation in different human diseases including cancers, few studies on how α7nAChR is expressed in TAMs in human CRC have been reported. It is also unclear whether α7nAChR expressed in TAMs is involved in the prognosis of colorectal carcinoma. Here, we examined the expression of α7nAChR in TAMs in CRC patients and found that high expression of α7nAChR in TAMs was associated with low incidence of hepatic metastasis. We revealed that human monocyte THP-derived macrophages with α7nAChR knockdown significantly enhanced invasion and migration of CRC cells in a Transwell co-culture model, which probably acted through the JAK2/STAT3 signaling pathway.

Materials and methods

Patients. Fifty-one consecutive cases of colorectal carcinomas used for the study were recruited from the patients who had undergone curative resection at the First Affiliated Hospital of Zhejiang University (Zhejiang, China) between May 2005 and May 2011. Inclusion criteria were a histological diagnosis of colorectal adenocarcinoma without chemotherapy or radiation therapy and without blood transfusions before surgery. Follow-up examinations were performed in March of 2016 by the first author. The telephone follow-up included the quality of life, progression of disease and survival time. The study was approved by the Human Research Ethics Committee of the First Affiliated Hospital of Zhejiang University (Zhejiang, China) and formal consent was obtained from all the patients prior to participation.

The clinical and pathological data of the patients included in the study are shown in Table I. For histopathological examination with hematoxylin and eosin, all surgical specimens were freshly cut and tissue samples were obtained from the primary tumors from the area of maximal invasion in paraffin wax. We recorded the tumor size, Dukes’ classification (UICC, 2002), number of lymph nodes involved and the stage of the disease. The histological grade of the tumors consisting of high-grade...
10% FBS, 2 mM glutamine, 50 U/ml penicillin, 50 µg/ml TIB-202™) were cultured in RPMI-1640 medium containing from American Type Culture Collection (ATCC, catalog no. mycin. Human monocytic leukemia cell line THP-1 cells containing 10% heat-inactivated fetal bovine serum (FBS),

Cell culture. LoVo cells (ATCC CCL-229™) and SW620 cells (ATCC CCL-227™), both derived from human colon adenocarcinoma metastases were grown in RPMI-1640 medium containing 10% heat-inactivated fetal bovine serum (FBS), 2 mM glutamine, 50 U/ml penicillin and 50 µg/ml streptomycin. Human monocytic cell line THP-1 cells from American Type Culture Collection (ATCC, catalog no. TIB-202™) were cultured in RPMI-1640 medium containing 10% FBS, 2 mM glutamine, 50 U/ml penicillin, 50 µg/ml streptomycin and 0.05 mM β-mercaptoethanol (β-ME) at 2-8x10^5 cells/ml. The cells were maintained at 37°C, 95% humidity and 5% CO₂. Cell viability was determined by trypan blue exclusion assay. All the cell culture reagents were purchased from Invitrogen (Shanghai, China).

Establishment of human monocyte THP-1-derived macrophage (TM) cell lines. Stable α7nAChR knockout human monocyte THP-1 cell lines were generated by transfection of the cells with α7nAChR-siRNA expression plasmid Lenti-X followed by puromycin selection, which was provided by Jinsirui Biological Technology Ltd. (Nanjing, China). The following shRNA sequence was used for knockdown of human α7nAChR: 5′-ggatccGATCACTATTTACAGTGGAA TTTCAGAGAATTCCACTGTGTAATATGTATCTTTTTT CATGGTctgta-3′. THP-1 cells stably bearing a mock α7nAChR expression plasmid (Jinsirui Biological Technology Ltd.) were used as control cells. Both RT-qPCR and western blot analysis confirmed that the two THP-1 cell lines were very stable without or with expression of α7nAChR at both mRNA and protein levels (Fig. 1).

The two THP-1 cell lines grown in RPMI-1640 medium were stimulated with 20 ng/ml phorbol 12-myristate 13-acetate (PMA; Sigma-Aldrich, St. Louis, MO, USA) for 48 h to induce the cells to differentiate into macrophages. After removing non-attached cells by washing three times with serum-free RPMI-1640 medium, the adherent macrophages were consid-

Figure 1. α7nAChR is knocked down in THP-1 derived macrophages (TM). The (A) mRNA and (B) protein expression level of α7nAChR in TM was determined by RT-qPCR and western blotting, respectively.

Cancer cell invasion and migration assay. Cell invasion and migration assays were carried out using modified Boyden chambers consisting of Transwell (8-µm pore size; Corning Costar Corp., Cambridge, MA, USA) membrane filter inserts in 24-well tissue culture plates. For the invasion assay, the upper surfaces of the membranes were coated with 100 µl of Matrigel (BD Biosciences, San Jose, CA, USA) at 37°C for 6 h and then placed into 24-well tissue culture plates which were placed with TMα7- and TM cells (2x10^5) in 500 µl medium containing 10% FBS, SW620 or LoVo cells (5.0x10^3) in 200 µl serum-free RPMI-1640 medium were added to each Transwell chamber and allowed to invade toward the underside of the membrane for 36 h for LoVo cells and 24 h for SW620 cells. Non-invasive cells were removed by wiping the upper side of the membrane with a cotton swab and the invaded cells were fixed with ice-cold methanol. Then the inserts were stained with 0.5% crystal violet in 20% ethanol for 30 min. The cells that invaded to the underside of the membrane were quantitated by cell counting under a light microscope in four predetermined fields at a magnification of x400. For the migration assay, the cell cultures and treatments were the same as those of the invasion assay except that there was no Matrigel in the upper surfaces of the membranes.

To investigate the effects of signaling pathway inhibitors on TMα7-/-enhanced migration of LoVo cells: TM cells (2x10^5) in 500 µl of medium in 24-well tissue culture plates were treated with or without α-bungarotoxin (α-Btx, a potent nicotinicα7 receptor antagonist; Sigma-Aldrich) at 100 nM for 6 h. After being washed with PBS three times, the TM cells were further treated with or without three signaling pathway inhibitors: AG490 (JAK2/STAT3 inhibitor, 20 µM), LY294002 (PI3K inhibitor, 5 µM) and Bay 11-7082 (NF-κB inhibitor, 2 µg/ml) for 6 h. After being washed with PBS three times, 500 µl of medium containing 10% FBS was added into the lower chamber and 5.0x10^5 LoVo cells in 200 µl of serum-free RPMI-1640 medium were added to each Transwell chamber for 36 h. The cell migration assay was the same as the procedure aforementioned.
**Indirect co-culture of LoVo cells and TMs.** To obtain TAMs, the indirect co-culture of TMac7−/− and TM with CRC LoVo cells was achieved using a Transwell co-culture system (0.4-µm pore size; Corning) in 6-well tissue culture plates. In brief, 2x10⁵ of LoVo cells in 1.6 ml of culture medium were seeded in the upper side of the membrane of the Transwell. Transwell inserts were then placed into 6-well plates cultured with 1x10⁶ of TMac7−/− or TM cells in 2.5 ml of culture medium. The polycarbonate Transwell membrane with 0.4-µm pores allows the exchange of the medium but limits the cell-cell interactions between the cancer cells and macrophages.

**Western blotting.** Cells were lysed in 1X RIPA buffer from Cell Signaling Technology (CST; Danvers, MA, USA) containing 1 mM phenylmethylsulfonyl fluoride (PMSF) and a protease inhibitor cocktail. The protein concentration was determined by Bradford protein assay. Total proteins (40 µg) were separated by SDS-PAGE and blotted onto PVDF membranes. The blots were first probed by antibodies against different phosphorylated proteins including phospho-STAT3, phospho-PI3 kinase p85, phospho-NF-κB p65, and β-actin. All the antibodies were bought from Cell Signaling Technology (CST). Then the membranes were probed with a horseradish peroxidase-conjugated goat anti-rabbit secondary antibody (Lianke, Hangzhou, China) followed by the detection of signals with FluorChem E system (Protein Simple, Santa Clara, CA, USA).

**Statistical analysis.** All assays were performed in triplicate. Data were analyzed using Stata 9.1 (Stata Corp., College Station, TX, USA) and presented as the mean ± SD. Significance of differences between clinical groups was determined by Chi-squared test and the invasion/migration data were determined by Student’s t-test. A p-value <0.05 was considered to be statistically significant.

**Results**

α7nAChR in TAMs prevents hepatic metastasis in CRC patients. We first determined the expression of CD68 in primary colonic adenocarcinoma of 51 colorectal patients using IHC staining as this intracellular glycoprotein is tightly associated with TAMs (31,32). As shown in Fig. 2, CD68 was highly expressed in tumors stroma, especially, in invasive cells of 2 representative colonic adenocarcinomas (Fig. 2C and D). We then determined whether α7nAChR was expressed in TAMs in 51 CRC patients using IHC staining. Expression of α7nAChR paralleled with CD68 in primary colonic adenocarcinoma of the 2 representative patients (Fig. 2A and B). We classified the α7nAChR expression in TAMs into 5 grades (0–4) based on the α7nAChR intensity detected. Grade 0 indicated no expression of α7nAChR, grade 1 only had weak α7nAChR expression while grade 2 to 4 exhibited medium to strong expression of α7nAChR. The results revealed that 33 CRC patients (64.7%) had grade 0 or 1 α7nAChR expression. Fourteen patients
(42.4%) had a 5-year survival. Eighteen patients (35.3%) had grade 2 to 4 expression (Table I). Notably, the other 18 patients who exhibited grade 2-4 expression of $\alpha_7nAChR$ had a 5-year survival at 77.8%, which is statistically significant. We observed that 14 of the 51 CRC patients (27.5%) with Dukes' D classification had hepatic metastasis. Among these 14 patients, 13 had grade 0 or 1 $\alpha_7nAChR$ expression and only 1 had grade 2 $\alpha_7nAChR$ expression. The other 37 patients (72.5%) with Dukes' A to C classification did not have hepatic metastasis. In this 37-patient group, 20 had grade 0 or 1 $\alpha_7nAChR$ expression and 17 had grade 2 to 4, respectively. Our results indicated that most of the hepatic metastatic CRC patients had low expression of $\alpha_7nAChR$ (grade 0 to 1) while non-metastatic patients exhibited strong expression of $\alpha_7nAChR$. Statistically the difference between the patient groups was significant, suggesting that $\alpha_7nAChR$ may play a functional role in the prevention of CRC metastasis.

Knockdown of $\alpha_7nAChR$ in TMs promotes CRC cell migration and invasion in vitro. To demonstrate our clinical observations that $\alpha_7nAChR$ expressed in TAMs may play a role in preventing CRC metastasis, we investigated the migration and invasion of CRC cells in a CRC/TM co-culture model. Two CRC cell lines LoVo and SW620 were co-cultured with THP-1 derived macrophages with or without $\alpha_7nAChR$ knockdown (TM and TM$\alpha_7^{-/-}$), respectively. The results revealed that both CRC cell lines co-cultured with TMs only had a few migrated or invaded cells after a certain period (Figs. 3 and 4). In contrast, the number of migrated or invaded CRC cells were significantly increased in the co-cultured with TM$\alpha_7^{-/-}$ CRC cell lines (Figs. 3 and 4), demonstrating that $\alpha_7nAChR$ may play an important role in cancer cell migration and invasion in vitro.

$\alpha_7nAChR$ regulates the expression of phosphorylated STAT3, PI3K p85 and NF-$\kappa$B p65 in TAMs. Next, we determined whether $\alpha_7nAChR$ had any regulatory roles in the expression and functions of key signaling molecules in both TAM and TAM$\alpha_7^{-/-}$. TM and TAM$\alpha_7^{-/-}$ cells were indirectly co-cultured with LoVo cells and then expressions of three key signaling...
molecules [phosphorylated-STAT3 (p-STAT3), phosphorylated-PI3K p85 (p-PI3K p85) and phosphorylated-NF-κB p65 (p-NF-κB p65)] were detected by western blotting. As shown in Fig. 5, results revealed that knockdown of α7nAChR in TMs significantly inhibited the expression of p-STAT3, p-PI3K and p-NF-κB in macrophages, which are involved in the metastasis of cancer cells.

Inhibition of phosphorylated-STAT3 in TMs prevents cancer cell migration. To determine whether the expression of p-STAT3, p-PI3K and p-NF-κB regulated by α7nAChR in TMs may mediate CRC cell migration, we first treated TMs with or without a potent nicotinic α7 receptor antagonist: α-Btx for 6 h and then with three inhibitors: AG490 (JAK2/STAT3 inhibitor), LY294002 (PI3K inhibitor) and Bay 11-7082 (NF-κB inhibitor) for 1 h, respectively. The inhibitor-treated TMs after being washed with PBS were co-cultured with CRC cells (LoVo cells) for 36 h to determine the migration of LoVo cells. The results clearly revealed that a very low number of LoVo cells had migrated when they were co-cultured with TMs under the condition that TMs were treated only with AG490, LY294002 and Bay 11-7082 [Fig. 6A (upper panel) and B]. In contrast, the number of migrated LoVo cells was significantly increased when TMs were only treated α-Btx (Fig. 6).

These results further demonstrated that α7nAChR expressed in TMs prevented CRC cell migration as shown in Fig. 4. Furthermore, the number of migrated LoVo cells was also markedly increased when TMs were dually treated with α-Btx and either LY294002 or Bay 11-7082, suggesting that the PI3K and NF-κB signaling pathway is not involved in α7nAChR-mediated migration of CRC cells. Conversely, dual treatment of α-Btx and AG 490 substantially prevented the migration of CRC cells (Fig. 6), suggesting that α7nAChR
expressed in TMs inhibits the migration of CRC cells through the JAK2/STAT3 signaling pathway.

Discussion

Metastasis is a multistep process by which cancer cells disseminate from primary tumors and establish secondary lesions in distant organs (33). The liver is one of the most significant targets for organ-specific metastasis in various cancer types and is commonly the sole site of metastasis for CRC (34). Hepatic metastasis (e.g., liver metastasis) is an ominous event in the natural history and progression of CRC (35), which is a major cause of mortality in CRC patients. To date, the underlying mechanisms of the hepatic metastasis in CRC patients have not been fully elucidated. Also, few studies have been published which demonstrate that TAMs play a role in preventing hepatic metastasis. In the present study we first observed that TAMs with α7nAChR expression are associated with less incidents of hepatic metastasis in CRC patients. Furthermore we observed that THP-derived macrophages with α7nAChR knockdown (TMα7-/−) significantly promoted the invasion and migration of two CRC cell lines in a two-cell Transwell co-culture model, similar to a previously published study which demonstrated that siRNA-knockdown of androgen receptor promotes...

Figure 5. α7nAChR regulates the expression of phosphorylated STAT3, PI3Kp85 and NF-κB p65 in tumor-associated macrophages (TAMs). THP-1 derived macrophages (TMs) and TMα7-/− at 1x10^6 cells were indirectly co-cultured with LoVo cells for 24 h. TMs were then harvested for western blotting to determine the expression of phosphorylated STAT3, PI3Kp85 and NF-κB p65. Results are representative of three independent experiments.

Figure 6. α7nAChR expressed in THP-1 derived macrophages (TMs) inhibits the migration of CRC cells through the JAK2/STAT3 signaling pathway. TMs were treated with or without α-bungarotoxin (α-Btx) for 6 h and then with DMSO (control), AG490, LY294002 and Bay 11-7082 for 1 h, respectively. Subsequently, LoVo cells were co-cultured with TM in 24-well Transwell plates with 8 µm polycarbonate membrane filters for 36 h for the migration assay. (A) Upper panel: Representative images show the migration of LoVo cells under the condition that the TMs did not have the α-Btx treatment and were treated with AG490, LY294002 and Bay 11-7082 only, respectively. Lower panel: Representative images show the migration of LoVo cells under the condition that the TMs were treated with α-Btx only, or dually treated with AG490, LY294002 and Bay 11-7082, respectively. (B) Number of migrated LoVo cells are presented as the mean ± SD of six images (n=6) of three separate experiments. Significance of differences between groups was determined by Student's t-test. P-value <0.05 indicates that there was a significant difference between treatments.
prostate cancer metastasis through enhanced macrophage recruitment (36). Thus, our data from both in vivo and in vitro experiments revealed that α7nAChR expressed in TAMs may play an important role in preventing metastasis and could be a prognosis marker in human CRCs.

Significant progress has been made in identifying the molecular basis for both macrophage phenotypes and their actions in promoting specific aspects of tumor behavior. A published study revealed that NF-κB activation is extremely crucial for mediating inflammation-induced tumor growth and progression, and modulating tumor surveillance and rejection (37).

It appeared that STAT3 activation was much more important for CRC growth and progression, and its surveillance and rejection in the present study.

Another study revealed that STAT3 is a point of convergence for numerous oncogenic signaling pathways, which is constitutively activated in both tumor cells and immune cells in the tumor microenvironment (38). The constitutive activation of STAT3 inhibits the expression of mediators necessary for immune activation against tumor cells and promotes the production of immunosuppressive factors leading the tumor to escape from host immune surveillance (38). Also, the invasive capacity of tumor cells can increase in the presence of inflammatory cytokines such as IL-1β and IL-6 (39). IL-6 is crucial for inducing and maintaining a cancer-promoting inflammatory environment and macrophage derived IL-6 promotes colorectal tumor cell survival and proliferation (40,41). Kaler et al (42) reported that TAMs, through IL-1β production, promote Wnt signaling in colon cancer cells, supporting tumor growth. It has been demonstrated that in macrophages, STAT3 is crucial for regulating the expression of cytokines (43-47). Based on the aforementioned evidence, we hypothesized that IL-6 and IL-1β may be the potential downstream molecules of the α7nAChR/JAK2/STAT3 pathway in macrophages, which regulate CRC metastasis. In future studies, we will explore the molecular mechanism of this phenomenon.

In addition, Mantovani et al (48) reported that PI3K activation may be a universal requirement for macrophage M2 differentiation and activation which may play the pro-tumoral role. Our laboratory tried to find a potential signaling pathway that correlated macrophage α7nAChR and CRC cell metastasis.

Activation of nAChRs in non-neuronal cells elicits the non-ionic signaling events that regulate protein phosphorylation and dephosphorylation, which is a novel function of nAChR subunit proteins in non-excitable cells. Activation of α7nAChR can activate p38 MAPK, AKT, RAS/RAF/MEK/ERK and JAK2 (49,50). In the present study, we determined the expression of phosphorylated STAT3, NF-κB p65 and PI3K p85 in THP-derived macrophages with or without α7nAChR knockdown (TMα7c and TM). The results revealed that phosphorylation of STAT3 was undetectable in TMα7c cells and NF-κB p65 and PI3K p85 were faint after co-culturing with LoVo cells, suggesting that the STAT3, NF-κB and PI3K signaling pathways may be related with α7nAChR expression. Furthermore, TAMs treated with the α7nAChR inhibitor α-Btx increased CRC cell invasion/migration that was prevented by treatment of the JAK2/STAT3 inhibitor AG490. These results further demonstrated that α7 nicotinic ACh receptor in TAMs inhibits CRC metastasis through the JAK2/STAT3 signaling pathway.

Previously, de Jonge et al (51) reported that the anti-inflammatory action of the activated nicotinic receptor in peritoneal macrophages was associated with the activation of the transcription factor STAT3. It has been well established that STAT3 is a key transcription factor in both immunity and inflammatory pathways. Ablation in myeloid cells of Stat3, a transcription factor whose function suppresses inflammatory responses since it is a major target of immunosuppressive cytokine IL-10 (52), causes inflammation in the colon. In the present study, α7nAChR knockdown substantially inhibited the expression of phosphorylated STAT3 in macrophages, providing evidence that phosphorylation of STAT3 is essential for α7nAChR to prevent functional invasion and migration of CRC cells although STAT3 is considered to be an oncogene that is frequently activated in a variety of human cancer cells and promotes the growth and survival of tumor cells (53,54).

In the present study, we clinically observed that the expression of α7nAChR in TAMs was associated with a low incidence of hepatic metastasis in CRC patients. We found that α7nAChR knockdown in THP-derived macrophages (TMs) significantly enhanced the migration and invasion of CRC cells, which was regulated through the JAK2/STAT3 signaling pathway in in vitro experiments. Our results revealed that α7nAChR expressed in TAMs may play an important role in preventing metastasis and could be a prognosis marker in CRCs. If validated in larger study sets, these findings may have potential clinical relevance since the detected cytokines and signaling proteins are current targets for molecularly-targeted therapeutics, leading to liver metastasis specific molecular therapies for the CRC.

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