Original Article

Assessing the role of IL-35 in colorectal cancer progression and prognosis

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Abstract: Despite the recent realization of Interleukin (IL)-35 in tumorigenesis, its exact impact on colorectal cancer (CRC) progression and prognosis, however, is yet to be elucidated clearly. We thus in the present report conducted comparative analysis of IL-35 levels between CRC patients and matched control subjects. IL-35 is highly expressed in all CRC tissues, which can be detected in vast majority of colorectal cancer cells. IL-35 levels in CRC lysates and serum samples are highly correlated to the severity of malignancy and the clinical stage of tumor. Particularly, a significant reduction for serum IL-35 was noted in patients after surgical resection, indicating that IL-35 promotes CRC progression associated with poor prognosis. Mechanistic study demonstrated a significant correlation between serum IL-35 levels and the number of peripheral regulatory T (Treg) cells in CRC patients, suggesting that IL-35 implicates in CRC pathogenesis probably by inducing Treg cells, while cancer cell-derived IL-35 may also recruit Treg cells into the tumor microenvironment in favor of tumor growth. Together, our data support that IL-35 could be a valuable biomarker for assessing CRC progression and prognosis in clinical settings.

Keywords: Interleukin-35, colorectal cancer, regulatory T cell, EBI3, IL-12p35

Introduction

Interleukin-35 (IL-35) is a heterodimeric cytokine composed of the p35 subunit of IL-12 and the Epstein-Barr Virus (EBV)-induced gene 3 (EBI3) subunit identified in B lymphocytes based on its induction following EBV infection [1-4]. Previous studies demonstrated that IL-35 is expressed in non-stimulated murine regulatory T cells (Tregs) [4] and in stimulated human Tregs [1, 4, 5]. Therefore, IL-35 is considered to be a characteristic factor for Treg cells [1, 4, 6-9], although its expression in non-stimulated human Tregs has yet to be detected [10]. Indeed, there is evidence supporting that IL-35 is effective to suppress the functionality of effector T cells such as Th1, Th2 and Th17 cells, and by which it attenuates the progression of inflammatory diseases and autoimmune diseases [1, 4, 11-14].

More recently, gene-expression analysis revealed that IL-35 may have a broader tissue distribution other than that observed in Tregs [15]. It was noted that EBI3 and IL-12p35 were upregulated in the placental trophoblasts [16], and EBI3 associates with p35 in the extracts of the trophoblastic components of human full-term normal placenta [2]. EBI3 has also been found expressed in Hodgkin lymphoma cells [17] and acute myeloid leukemia cells [18]. Immunohistochemical analysis further revealed that EBI3 and IL-12p35 are highly expressed in tumor tissues originated from lung cancer, colon cancer, esophageal carcinoma, hepatocellular carcinoma and cervical carcinoma [19].

Indeed, IL-35 can be induced in human cancer cell lines following TNF- α and IFN- γ stimulation [19]. Together, these data support a role for IL-35 in tumor development and metastasis.

In the tumor microenvironment, Foxp3+ Tregs and other Tregs are considered to be the primary IL-35 producer [20, 21]. Treg-derived IL-35 has been found to potently inhibit antitumor T cell responses [13]. In line with this notion, knockdown of EBI3 inhibited lung cancer cell proliferation, while upregulation of EBI3 promoted lung cancer cell growth [22]. Therefore, EBI3 expression is considered to be correlated to poor prognosis for human lung cancer [22]. Other than Treg cells, tumor-infiltrating dendritic cells (DCs) are also found to express EBI3 [17, 23], and tumor cells can produce EBI3 and IL-12p35 as well. Indeed, tumorderived IL-35 induces CD11b+Gr1+ myeloid cell accumulation in the tumor microenvironment, and thereby, promotes tumor angiogenesis [24]. While these discoveries are important and exciting, the exact impact of IL-35 on tumorigenesis, particularly in human colorectal cancer development and metastasis, is yet to be fully addressed. We herein, in the present report, conducted studies in colorectal cancer (CRC) patients to address this question. Our data revealed that IL-35 is highly expressed in all CRC tissues and is detected in vast majority of tumor cells. Particularly, much higher levels of p35 and EBI3 expressions are detected in poorly differentiated CRC patients than that of moderately and well differentiated CRC patients. Moreover, serum IL-35 levels are positively correlated to Treg numbers in the peripheral blood of CRC patients, and a significant reduction of serum IL-35 was noted in patients after tumor resection. Together, our data support that IL-35 could be a valuable biomarker for assessing CRC progression and prognosis in clinical settings.

Patients and materials

Patient collection

A total of 50 CRC patients undergone large bowel resection were collected at the Department of Surgery of Affiliated Hospital of Guangdong Medical College in Zhanjiang and the Second Clinical Medical School of Guangdong Medical College in Dongguan. The collections are composed of 28 male patients and 22 female patients with a mean age of 51.1 yr old (21~78 yr old). Both cancer and normal tissues (> 5 cm away from cancer tissues) were obtained from operative specimens and stored in -80°C until use. For normal controls, 50 healthy volunteers (HV) were organized by the Affiliated Hospital of Guangdong Medical Examination Center, which included 27 males and 23 females with a mean age of 48.1 yr old (24~78 yr old). Blood samples were collected before and after surgery (> 7 days after surgery). Serum aliquots (200 µl for each) were rapidly frozen and stored in a -80°C freezer until use. PBMCs were isolated from peripheral blood samples and then subjected to flow cytometry analysis immediately. Patients with recent blood transfusion or immunotherapy were excluded from the study. Informed consent was obtained from all study subjects, and the studies were approved by the Internal Review Board of Human Assurance Committee at the Guangdong Medical College.

Immunofluorescence and immunohistochemistry

Tissue sections (4-µM) were prepared from frozen tissue blocks and then subjected to Hematoxylin & Eosin (HE) staining as reported [25]. The results were assessed by two pathologists to demonstrate the presence of tumor and the proportion of tumor cells in each section. For immunostaining, tissue sections were subjected to antigen retrieval with sodium citrate (pH 6.0), followed by staining with antibodies against IL-12 p35 (sc-7925, Santa Cruz Biotechnology, Inc.) or EBI3 (sc-32868, Santa Cruz Biotechnology, Inc.) overnight at 4°C, respectively. After washes, the sections were stained with a goat anti-rabbit IgG labeled with either PE (bs-0295G-PE, Beijing Biosynthesis Biotechnology Co., Ltd) or FITC (bs-0295G-FITC, Beijing Biosynthesis Biotechnology Co., Ltd) [26]. The sections were then assessed for p35 and EBI3 expression under a fluorescent microscope (Nikon Ti-U). Immunohistochemistry (IHC) was next employed for quantitative analysis of p35 and EBI3 expression levels, which included 50 cancer sections and 50 normal sections. A rabbit polyclonal antibody against IL-12p35 (sc-7925, Santa Cruz Biotechnology, Inc.) or EBI3 (sc-32868, Santa Biotechnology, Inc.) was applied for the analysis as previously reported [27]. In brief, the slides were incubated in H₂O₂ solution for 10

 Table 1. Demographic and clinical characteristics for the enrolled subjects

		·'		CRC patients		Healthy Volunteers		
NO.	Gender	Age	Tumor size	Histological grade	TNM (stage)	NO.	Gender	Age
1	F	27	< 5 cm	Poorly	I	1	F	32
2	F	64	> 5 cm	Poorly	IV	2	M	46
3	F	50	< 5 cm	Well	III	3	F	48
4	M	64	< 5 cm	Moderately	1	4	M	51
5	F	43	> 5 cm	Poorly	III	5	F	34
6	M	29	< 5 cm	Adenocarcinoma	III	6	M	38
7	M	47	> 5 cm	Moderately	III	7	F	46
8	M	49	< 5 cm	Moderately	IV	8	F	42
9	F	61	< 5 cm	Well	III	9	F	51
10	M	56	> 5 cm	Well	III	10	M	29
11	M	53	> 5 cm	Moderately	II	11	M	31
12	F	12	> 5 cm	Adenocarcinoma	IV	12	M	35
13	M	51	< 5 cm	Moderately	IV	13	M	64
14	M	65	< 5 cm	Well	III	14	F	62
15	F	56	< 5 cm	Moderately	II	15	M	45
16	F	78	< 5 cm	Moderately	III	16	F	58
17	M	54	< 5 cm	Well	1	17	M	67
18	F	55	< 5 cm	Well	1	18	M	38
19	F	57	< 5 cm	Moderately	II	19	M	42
20	M	50	< 5 cm	Poorly	IV	20	F	46
21	M	58	> 5 cm	Adenocarcinoma	III	21	M	49
22	M	46	< 5 cm	Adenocarcinoma	1	22	F	38
23	M	42	> 5 cm	Adenocarcinoma	II	23	F	65
24	M	34	< 5 cm	Well	II	24	M	64
25	M	71	< 5 cm	Adenocarcinoma	III	25	F	63
26	M	67	< 5 cm	Well	1	26	M	71
27	M	41	> 5 cm	Adenocarcinoma	II	27	M	70
28	F	45	< 5 cm	Well	1	28	M	42
29	F	59	< 5 cm	Moderately	1	29	M	41
30	F	55	< 5 cm	Well	1	30	F	36
31	M	47	> 5 cm	Adenocarcinoma	II	31	M	37
32	M	63	< 5 cm	Well	1	32	M	56
33	M	37	< 5 cm	Well	II	33	M	54
34	F	56	> 5 cm	Moderately	III	34	F	29
35	F	48	> 5 cm	Poorly	IV	35	F	36
36	M	46	< 5 cm	Moderately	1	36	F	33
37	F	48	> 5 cm	Adenocarcinoma	II	37	M	46
38	M	64	> 5 cm	Poorly	III	38	F	44
39	F	35	< 5 cm	Moderately	1	39	M	37
40	M	66	< 5 cm	Well	II	40	F	53
41	F	64	< 5 cm	Well	1	41	M	64
42	M	21	< 5 cm	Moderately	1	42	M	55
43	F	39	< 5 cm	Poorly	II	43	F	37
44	F	53	< 5 cm	Moderately	1	44	M	46
45	M	46	> 5 cm	Poorly	IV	45	F	47
46	M	55	< 5 cm	Adenocarcinoma	III	46	М	65

47	F	43	< 5 cm	Poorly	II	47	F	66
48	F	70	< 5 cm	Adenocarcinoma	I	48	M	38
49	M	68	> 5 cm	Poorly	III	49	F	67
50	М	46	> 5 cm	Adenocarcinoma	II	50	F	52

TNM: Tumor-node-metastasis.

min at room temperature to block the endogenous peroxidase activity. Antigenic epitopes were next retrieved by heating for 5 min in 10 mmol/L citrate buffer (pH 6.0). The slides were then first incubated with the above indicated primary antibodies for 30 min at room temperature, followed by a secondary peroxidase antirabbit IgG at 1:500 dilutions.

Score of immunohistochemical sections

All sections were assessed by two pathologists in a blinded fashion to the clinical status of the patients. The immunoreactive area for EBI3 and p35 was scored as 1 (0~25%), 2 (25~50%), 3 (50~75%) or 4 (> 75%), as previously reported by Michalski and colleagues [28]. Based on the intensity of staining, 0 scored with absent of staining, 1 scored as weak staining, and 2 scored as strong staining. The final score was established by the average scores from two pathologists. Cases with high variations between two pathologists were evaluated again without the knowledge of previous results.

ELISA analysis

Frozen tissues (1 gm) were homogenized in a glass homogenizer in 200 µl PBS, supernatants were collected after centrifugation. A human IL-35 ELISA kit (BlueGene, Shanghai, China) was employed for analysis of IL-35 levels in the tissue supernatants and serum samples using the established techniques [29].

Flow cytometry analysis

Isolated PBMCs were resuspended in FACS buffer (PBS with 1% BSA and 0.1% sodium azide), and then subjected to staining with FITC labeled mouse anti-human CD4 (BD Pharmingen, San Jose, CA), APC labeled mouse anti-human CD25 (BD Pharmingen, San Jose, CA), and PE labeled mouse anti-human Foxp3 (BD Pharmingen, San Jose, CA). After washes, the stained cells (2 x 10^6) were resuspended in 400 μ L FACS buffer containing 2% paraformal-dehyde and then analyzed via three-color flow

cytometry (Epics Altre II, Beckman Coulter) as reported [30]. A mouse IgG isotype was served as negative controls. All data were analyzed using the Flowjo.7.6.1 software (Treestar, Ashland, OR, USA).

Statistical analysis

The normality test was first performed to determine whether our dataset was well-modeled by a normal distribution. Student's t-test was employed for analysis of two-sample and two-tailed comparisons. Pearson correlation was used to measure the degree of dependency between variables by GraphPad Prism version 5.0 (GraphPad Software Inc., San Diego, CA, USA). In all cases, p < 0.05 was considered with statistical significance.

Results

Demographics and clinical data

According to the WHO grading system, tumors in 14 patients (28%) were defined as well differentiated, 14 patients (28%) fell into the category of moderately differentiated, 10 patients (20%) were poorly differentiated, and 12 patients (24%) were diagnosed as mucinous adenocarcinoma. Based on the standard for tumor-node-metastasis (TNM) stage, patients (32%) were in stage I, 13 patients (26%) were found in stage II, 14 patients (28%) were characterized in stage III, and 7 patients (14%) were defined in stage IV. Among all patients, 18 patients (36%) were noted with tumor size > 5 cm. The demographic and clinical characteristics of the selected subjects were summarized in **Table 1**. There was no significant difference in age and gender between CRC patients and control subjects.

p35 and EBI3 are highly expressed in colorectal cancer tissues

We first conducted immunostaining for detection of IL-35 expression in colorectal cancer sections using a FITC-conjugated anti-p35 anti-

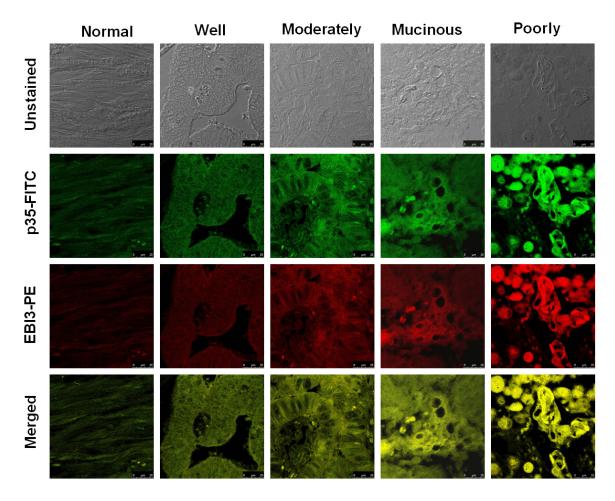


Figure 1. Detection of IL-35 expression in colorectal cancer tissues by immunostaining of EBI3 and p35. CRC sections were stained with FITC-conjugated anti-p35 antibody and PE-conjugated anti-EBI3 antibody, respectively, and the sections were then assessed under a fluorescent microscope by two pathologists. A perfect co-localization for the immunofluorescence of EBI3 and p35 was noted in all studied sections, indicating the expression of IL-35 in colorectal cancer cells. All images were taken under a confocal laser scanning microscope with 2000 x amplifications.

body and a PE-conjugated anti-EBI3 antibody, respectively. It was found that both p35 and EBI3 were expressed in all CRC sections. Particularly, much higher levels of p35 and EBI3 were detected in poorly differentiated CRC samples as compared with that of well or moderately differentiated samples (Figure 1). Of note, p35 and EBI3 showed perfect co-localization in the tissue sections, demonstrating the expression of IL-35 in colorectal cancer cells.

For quantitative analysis of IL-35 expression levels in different types of colorectal cancer tissues, we did immunohistochemical analysis of p35 and EBI3 as described. In line with the above results, cancer sections from all patients displayed positive reactivity for p35 and EBI3

as shown in Figure 2A. It was interestingly noted that the expression levels for p35 and EBI3 in tumors were correlated to the extent of cell differentiation, in which the highest expression was detected in poorly differentiated tumor samples, while well differentiated tumor samples showed the lowest expression, and the expression levels for moderately differentiated tumor samples were in the between of these two types of tumors (Figure 2B). However, no correlation was detected in terms of clinical stage, age and gender, and tumor size with p35 and EBI3 expressions (data not shown). Of note, high levels of p35 and EBI3 expressions were also detected in patients with mucinous adenocarcinoma (Figure 2B). The sections were next scored for immunoreactive area as

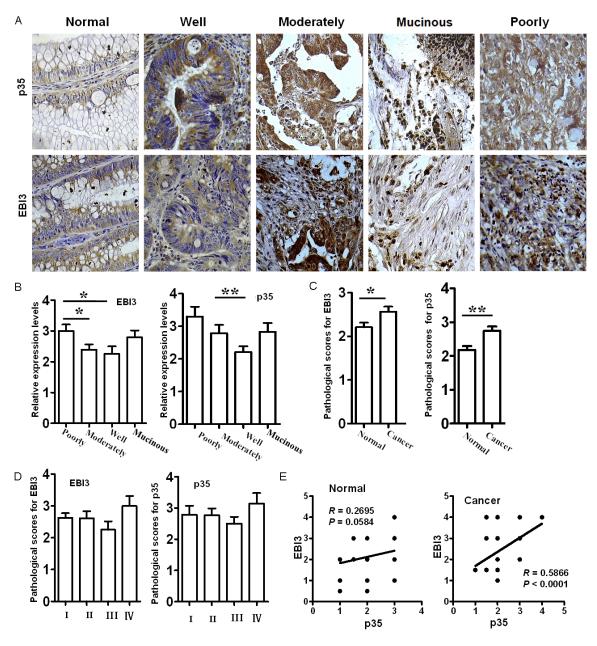


Figure 2. Quantitative analysis of EBI3 and p35 expression in colorectal cancer tissues by immunohistochemistry. Immunohistochemical analysis was employed for quantitative analysis of p35 and EBI3 expression in CRC tissues. A. Representative images for the immunohistochemical staining of EBI3 and p35 in different types of CRC tissues. All images were taken under a light microscope with 400 x amplifications. B. Bar graphic figures showing the relative expression levels of EBI3 and p35 assessed in all sections. C. Pathological scores assessed based on the intensity of immunoreactive area of all sections analyzed. D. Bar graphic figures showing the differences of EBI3 and p35 expression in different stages of CRC sections. E. Results for correlation analysis of EBI3 and p35 in normal and CRC sections. *, p < 0.05; **, p < 0.01; ***, p < 0.001.

detailed earlier based on the intensity of staining. In consistent with the expression data, the patients showed significantly higher scores than that of controls for both p35 and EBI3 (Figure 2C). Next, we examined p35 and EBI3 expression in different stages of tumor sam-

ples, and much higher levels of p35 and EBI3 expressions were noted in stage IV tumor samples (Figure 2D). Furthermore, p35 was noted to be positively correlated to EBI3 in colorectal cancer tissues, but not in normal tissues after Pearson correlation analysis (Figure 2E).

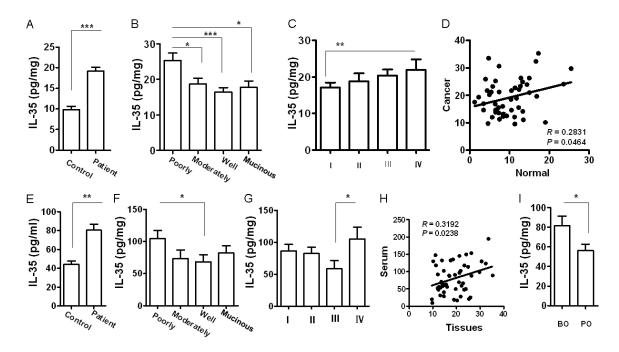


Figure 3. ELISA analysis of IL-35 levels in colorectal cancer lysates and serum samples. A. A bar graphic figure showing the difference of IL-35 levels between control and CRC lysates. B. IL-35 levels in different types of CRC lysates. C. IL-35 levels in different stages of CRC samples. D. Results for correlation analysis of IL-35 levels between CRC tissues and normal tissues. E. Serum IL-35 levels in controls and CRC patients. F. Serum IL-35 levels in patients with different types of colorectal cancer. G. Comparison of serum IL-35 levels between patients with different stages of colorectal cancer. H. Results for correlation analysis of IL-35 levels between serum samples and CRC lysates. I. Comparison of serum IL-35 levels in patients before and after surgical resection of tumors. BO, before operation; PO, post operation. *, p < 0.05; **, p < 0.01; ***, p < 0.001.

IL-35 levels are correlated to colorectal cancer progression and prognosis

The above expression data prompted us to examine IL-35 levels in cancer tissues. For this purpose, we conducted ELISA analysis of IL-35 using CRC lysates. In line with the expression data, IL-35 levels were almost 1-fold higher in colorectal cancer tissues than that of control tissues (Figure 3A), and a similar correlation for IL-35 levels and the extent of differentiation of cancer cells (Figure 3B), as well as the clinical stages of tumors (Figure 3C) was noted. Particularly, higher levels of IL-35 were highly correlated to patients with colorectal cancer (Figure 3D). Next, we analyzed the differences of serum IL-35 levels between cancer patients and normal controls. Although serum IL-35 was likely less significantly as compared with that in cancer tissues, a similar trend was also noted (Figure 3E-G), and serum IL-35 was positively correlated to that in cancer tissues (Figure 3H). More importantly, a significant reduction for serum IL-35 was noted in all patients after tumor resection (**Figure 3I**). All together, those data suggest that IL-35 could be a valuable biomarker for assessing colorectal cancer progression and prognosis.

Serum IL-35 is associated with higher peripheral Tregs in colorectal cancer patients

Given that increased number for Treg cells is a characteristic feature for the extent of colorectal cancer malignancy and progression, we therefore next, examined the correlation between IL-35 levels and Treg numbers in the peripheral blood for dissecting the mechanisms underlying IL-35 promotion of CRC progression. To this end, we first gated CD4+CD25+ cells and then analyzed Foxp3 expression in the gated cells as shown in Figure 4A. As expected, CRC patients showed almost 1-fold higher Foxp3+ Treg cells than that of control subjects (Figure 4B). Particularly, significantly higher proportion of Foxp3+ Treg cells was noted in patients with poorly differentiated tumors as compared with that of patients with

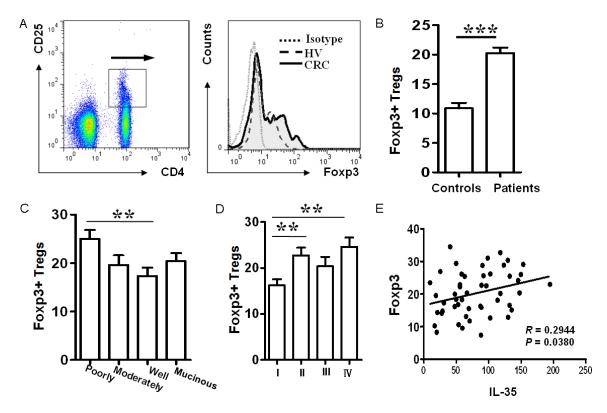


Figure 4. Serum IL-35 levels associate with the number of peripheral Treg cells in colorectal cancer patients. A. A paradigm for flow cytometry analysis of the peripheral Foxp3+ Treg cells. B. Comparison of peripheral Foxp3+ Treg cells between normal controls and CRC patients. A total of 50 patients and 50 matched controls were analyzed. C. The distribution of peripheral Treg cells in patients with different types of colorectal cancer. D. Comparison for the number of peripheral Treg cells between patients with different stages of colorectal cancer. E. Serum IL-35 levels are highly correlated to the number of peripheral Treg cells in CRC patients. **, p < 0.01; ***, p < 0.001.

moderately or well differentiated tumors (Figure **4C**), and patients in stage I displayed the lowest number for Treg cells, while highest number of Treg cells was detected in patients in stage IV (Figure 4D). Next, we conducted correlation analysis between serum IL-35 levels and peripheral Treg numbers, Remarkably, serum IL-35 levels were positively correlated to higher number of peripheral Treg cells in colorectal cancer patients (Figure 4E). Together, our data suggest that IL-35 could be a potent factor for induction of peripheral Treg cells in colorectal cancer patients, and through which it promotes CRC progression associated prognosis.

Discussion

IL-35, a member of the IL-12 family of cytokines, is well documented as an anti-inflammatory cytokine implicated in the regulation of autoimmune diseases and allograft rejection. For example, previous studies revealed that

IL-35 produced by Treg cells is potent to suppress the functionality of Th1, Th17 and Th2 cells [13]. More recently, IL-35 has been further recognized to play a pivotal role in the pathogenesis of tumor development, progression and prognosis. In general, IL-35 expression is considered with immunosuppression along with tumor progression and poor prognosis. However, contradictory results have also been reported; for example, it has been noted that ectopic IL-35 expression in human cancer cells suppresses cell growth by inducing cell cycle arrest at the G1 phase along with enhanced apoptosis [19]. Therefore, the exact impact of IL-35 on tumorigenesis, particularly on colorectal cancer (CRC) progression and prognosis, is yet to be fully addressed.

In the present report, we conducted expression analysis of IL-35 in 50 CRC patients and 50 age-matched healthy controls. We demonstrated experimental evidence indicating that IL-35 is highly expressed in vast majority of colorec-

tal cancer cells. We next analyzed IL-35 expression levels by immunohistochemical analysis and ELISA analysis of cancer tissue samples. Our data have consistently revealed that the expression levels for IL-35 are associated with the extent of CRC malignancy and clinical stage, in which much higher levels of IL-35 were detected in poorly differentiated CRC tissues, while well differentiated CRC tissues manifested the lowest IL-35 expression as compared with that of poorly or moderately differentiated CRC tissues. Correlation analysis suggested that IL-35 is positively correlated to CRC malignancy and progression. To further confirm the results in CRC tissues, we analyzed serum IL-35 levels. Similar as the data obtained from cancer tissues, CRC patients manifested 1-fold higher serum IL-35 than that of control subjects, and higher levels of serum IL-35 were associated with the extent of CRC differentiation and clinical stage. A noteworthy finding is that patients underwent a significant reduction for serum IL-35 after surgical resection of tumor tissues. All together, our data support that IL-35 could be a valuable biomarker essential for assessing CRC progression and prognosis after surgical resection and chemotherapy in clinical settings.

To address the mechanisms underlying IL-35 promotion of CRC progression associated with poor prognosis, we performed correlation analysis between serum IL-35 levels and peripheral Treg production. Remarkably, serum IL-35 levels were highly correlated to the number of peripheral Treg cells in CRC patients. Of interestingly note, IL-35 has been found to mediate potent induction of Treg cells [13]. Given the role of Treg cells played in tumorigenesis and cancer progression, we thus assume that IL-35 enhances CRC progression at least partly by inducing Treg production.

As aforementioned, other than Treg cells, colorectal cancer cells also express high levels of IL-35. IL-35 derived from CRC cells may recruit Treg cells or other immune cells with suppressive function into the cancerous milieu, and through which it promotes CRC progression. In line with this assumption, tumor-derived IL-35 has been found to increase CD11b+Gr1+ myeloid cell accumulation in the tumor microenvironment and, thereby, promotes tumor angiogenesis [24]. Also, Olson and colleagues characterized a population of CD8+CTLA-4+

IL-35-secreting tumor Ag-specific Tregs in prostate cancer patients, which prevents Ag-specific effector responses by an IL-35-dependent mechanism [31].

In consistent with published data, we noted that IL-35 levels in the tumor tissues and peripheral blood are negatively correlated to colorectal cancer cell differentiation. Unexpectedly, IL-35 levels have also been found to be positively correlated to the clinical stage of CRC patients. Given the role of IL-35 played in protection of tumor cells against immunity, this observation in fact further supports our original hypothesis. Indeed, as the disease evolves, higher levels of IL-35 in the tumor microenvironment are likely to be a prerequisite to ensure CRC progression and metastasis.

In summary, we demonstrated evidence that IL-35 is highly expressed in colorectal cancer cells. High levels of IL-35 in CRC tissues and peripheral blood are correlated to the extent of CRC malignancy and clinical stage. Particularly, CRC patients undergo a significant reduction of serum IL-35 after surgical resection of tumors. Mechanistic study revealed that serum IL-35 is correlated to the production of peripheral Treg cells, while IL-35 derived from colorectal cancer cells may recruit Treg cells and other immunosuppressive cells into the tumor microenvironment, and through which, IL-35 promotes CRC progression associated with poor prognosis. All together, our data support that IL-35 could be a valuable biomarker for assessing CRC progression and prognosis in clinical settings.

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Disclosure of conflict of interest

The authors declare that there are no conflicts of interest.

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