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Intratumoral interleukin-9 delineates a distinct immunogenic class of gastric cancer patients with better prognosis and adjuvant chemotherapeutic response

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ABSTRACT

Interleukin-9 (IL-9) is a T cell cytokine that is associated with inflammation and allergy, but the expression level of IL-9 in gastric cancer and its clinical significance are less well established. Our study aims to uncover the critical role of IL-9 in the progression of gastric cancer. Here, a total of 453 patients with gastric cancer undergoing curative resection were enrolled for immunohistochemical analyses, and Kaplan-Meier analysis was conducted to compare overall survival of patients in different subgroups. We further investigated the correlation between IL-9 expression and functional status of intratumoral CD8⁺ T cells by means of Flow cytometry. Moreover, in vitro study was preformed to further explore the influence of IL-9 on anti-tumor immunity. Results indicated that gastric cancer patients with high IL-9 expression showed improved overall survival and gained more benefit from 5-fluorouracil-based adjuvant chemotherapy (ACT). High IL-9 expression was associated with increased numbers and elevated function of intratumoral CD8⁺ T cells. In vitro study revealed that recombinant human IL-9 (rhIL-9) exhibit anti-tumor activity via enhancing the function of intratumoral CD8⁺ T cells. Moreover, we found rhIL-9 could augment the efficacy of Pembrolizumab in gastric cancer. In summary, these results suggest that IL-9 expression could act as an independent predictor for overall survival and ACT response and enhancing IL-9 signaling might represent an important therapeutic strategy in gastric cancer.

Introduction

Gastric cancer is one of the most common causes of cancerrelated death around the world. especially in Eastern Asia.^{1, 2} Owing to nonspecific symptoms in early stage, most patients are diagnosed at an advanced stage and showed a high rate of recurrence despite receiving standard radical gastrectomy. For these patients, a 5-fluorouracil-based adjuvant chemotherapy (ACT) regimen is commonly used as a first-line choice to reduce postoperative recurrent rate and prolong overall survival.^{3,4} However, large variations in clinical outcomes have been reported in patients with the same stage and similar treatment regimens. Hence, exploring the potential individual benefit of adjuvant chemotherapy is drawing more and more attention.

Recently, the clinical significance of tumor-infiltrating immune cells has been recognized in gastric cancer.^{5–7} As the bridge of interaction among immune cells, cytokines also play a critical role in tumor progression by orchestrating the entire immune microenvironment. Interleukin-9 (IL-9) is a newly identified cytokine in the late 1980s which was firstly described as a member of T cell growth factor.^{8,9} However, subsequent researches showed that IL-9 had pleiotropic functions with multiple target cells including T cells, mast cells, B cells as

well as nonimmune cells.⁸ So far, the main role of IL-9 has been described in the immune responses against parasites and pathogenesis of allergic diseases.¹⁰ However, the function of IL-9 in cancer biology is still dim and needs much more exploration. IL-9 was described as a pro-tumorigenic cytokine by promoting tumor cell growth in hepatocellular and breast cancer.^{11,12} While in melanoma, IL-9 exhibits anti-tumor activity by provoking CD8⁺ T cell-mediated antitumor immunity.¹³ These data revealed that IL-9 seemed to have pleiotropic effect depended on different tumor types. While in gastric cancer, the impact of IL-9 on anti-tumor immunity and its association with patients' prognosis remains unexplored.

Thus, in this study, we evaluated the intratumoral IL-9 expression in gastric cancer and further explored its prognostic value for overall survival and influence on anti-tumor immunity. We found that IL-9 expression exhibited outstanding prognostic value and predictive ability for overall survival benefit from ACT. Further analysis showed that IL-9 enhanced the anti-tumor immunity by intratumoral CD8⁺ T cells in gastric cancer. Moreover, we found that recombined human interleukin-9 (rhIL-9) could augment response to programmed cell death protein-1 (PD-1) blockade. All these results revealed

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that IL-9 expression could serve as a biomarker for prognosis and adjuvant response in gastric cancer patients and exogenous IL-9 might exert anti-tumor efficacy.

Materials and methods

Patients and specimens

We enrolled 453 gastric cancer patients who received total or partial gastrectomy by the same surgical team in Zhongshan Hospital, Fudan University (Shanghai, China) during August 2007 and December 2008 for survival analysis. Initially, 496 patients were recruited in this study. 28 patients were excluded owing to lack of chemotherapy information, clinicopathological data, or survival outcomes for complete analysis. In this study, we utilized the tissue microarrays (TMAs) constructed by Shanghai Outdo Biotech Co, Itd, and 7 dots were detached after immunohistochemistry. Additionally, we excluded 8 patients of TNM stage IV. Finally, 453 patients were included in our study and we randomly divided them into two independent data sets, the Discovery set (n = 207) and the Validation set (n = 246). There were totally of 260 patients (57.4%) received fluorouracil-based ACT according to practical needs and patients' will after their total or partial gastrectomy. Clinicopathological characteristics and their association with intratumoral IL-9 expression were shown in Table 1. Additionally, fresh gastric cancer samples from 44 patients who underwent gastrectomy in Zhongshan from August 2018 to October 2020 were obtained. Written

informed consents of all patients in this study were provided and the Clinical Research Ethics Committee of Zhongshan Hospital, Fudan University had approved our study.

Assay methods

The protocol of immunohistochemistry (IHC) and flow cytometry (FCM) were listed in Supplementary Methods, and antibodies involved were summarized in Supplementary Table S1. TMA slides were scanned on Leica DM6000 B (Leica Microsystems, Wetzlar, Germany). In our study, two pathologists (Dr. Peipei Zhang and Dr. Lingli Chen) who were blinded to the clinical data evaluated all samples separately. Both of them scored independently according to the intensity of cellular staining and the proportion of stained cells.¹⁴ The mean score of their evaluation was adopted. Variations in IL-9 IHC score, exceeding 10, were reevaluated separately by both pathologists to acquire consensus. In brief, the IHC cellular staining intensity was stratified as 0 (negative staining), 1 (weak staining, light yellow), 2 (moderate staining, yellow brown) and 3 (strong staining, brown), while the proportion was defined as the percentage of positive cells (0-100%). The staining intensity and proportion were multiplied to generate an overall IL-9 IHC score ranging from 0 to 300. The density of immune cells was recorded as the average number of cells with positive staining marker per high power field (HPF). Determined by X-tile software using minimum *p*-value method in the discovery set, the cutoff point was set as 20 to define the intratumoral

	Discovery set			Validation set			Combined set		
Characteristics	Low	High	Р	Low	High	Р	Low	High	Р
All patients	97	110		105	141		202	251	
Age(years)			0.851			0.907			0.864
Median	61	58		63	59		61.5	59	
IQR	(55.0–71.5)	(50.8–67.0)		(53.0–70.5)	(50.5–68.0)		(54.0-71.0)	(51.0-68.0)	
Gender			0.724			0.225			0.238
Male	63	74		74	109		137	183	
Female	34	36		31	32		65	68	
Localization			0.324			0.460			0.648
Proximal	19	24		32	37		51	61	
Middle	23	17		8	17		31	34	
Distal	55	69		65	87		120	156	
Tumor size(cm)			0.172			0.441			0.147
Median	3.5	3.5		3.5	3.5		3.5	3.5	
IQR	(2.0-5.0)	(2.0-4.6)		(2.0-5.3)	(2.0-5.0)		(2.0-5.0)	(2.0-5.0)	
Differentiation			0.940			0.569			0.718
Well+Moderately	26	30		31	37		57	67	
Poorly	71	80		74	104		145	184	
Lauren classification			0.889			0.786			0.901
Intestinal type	60	67		66	91		126	158	
Diffuse type	37	43		39	50		76	93	
T stage			0.883			0.083			0.265
T1	20	23		16	28		36	51	
T2	15	17		8	24		23	41	
T3	18	16		25	25		43	41	
T4	44	54		56	64		100	118	
N stage			0.744			0.155			0.329
NO	45	51		30	50		75	101	
N1	11	9		11	21		22	30	
N2	18	18		19	29		37	47	
N3	23	32		45	41		68	73	
TNM stage			0.573			0.068			0.217
1	30	28		18	38		48	66	
II	23	32		19	32		42	64	
III	44	50		68	71		112	121	

Table 1. Relationship between intratumoral IL-9 expression expression and baseline characteristics of patients.

IL-9 high/low expression. In fresh tumor samples, intratumoral IL-9 expression was assessed by mean fluorescence intensity (MFI) with flow cytometry. The cutoff value for the classification of IL-9 high/low subgroup in the 44 fresh tumor samples was the median value of MFI. All flow cytometry data were obtained using BD FACS Celesta and analyzed by FlowJo v10.0 (Treestar). The gating strategy of flow cytometry is provided in Supplementary Figure 1.

Elisa (Enzyme linked immunosorbent assay)

IFN-γ, GZMB, and PRF1 were measured by ELISA in tumor tissue supernatant. ELISA kit used was the human IFN-γ, GZMB and PRF1 kits obtained from RUIXIN BIOTECH (RUIXIN BIOTECH, Cat.# 10296, Cat.#16262, Cat.#10448).

In vitro therapeutic assay

The fresh tumor tissue was dissociated into single-cell suspensions as above. The digests were cultured in assay medium (RPMI-1640 with 100 U/mL penicillin, 100 µg/mL streptomycin and 10% fetal bovine serum) with anti-CD3 antibody (0.5 µg/ml, OKT3; Biolegend) and anti-CD28 antibody (2 µg/ml, clone 28.2; eBioscience). In CD8⁺ T cells-deprived system, CD8⁺ T cells were depleted from single cell suspensions by negative selection using CD8 MicroBeads (Life Technologies). Recombinant human IL-9 (0.3 ng/ml, R&D Systems), anti-IL-9 antibody (5 µg/mL, R&D Systems) or PBS were added to the culture system to study the effect of IL-9 on anti-tumor immunity. Pembrolizumab (10ug/ ml; Pembrolizumab, Selleck) were used to explore the synergistic effect of PD-1 blockade and rhIL-9. After 24 hours of incubation, we collected cells for flow cytometry as indicated above.

Statistical analysis

Pearson χ^2 test or Student's t test was respectively applied to compare categorical variables and continuous variables. Overall survival of the two patient data sets was calculated by Kaplan–Meier method and analyzed by log rank test. Cox proportional hazard regression model was used to perform multivariate analyses. Spearman correlation analysis was conducted to assess the correlation between IL-9 expression and other immune cells infiltration. All analyses were performed using IBM SPSS Statistics v20.0, MedCalc 15.6.1 and GraphPad Prism v6.02. All statistical tests were two–tailed and only P < .05 were considered as statistically significant.

Results

Intratumoral IL-9 expression and its association with patients' clinicopathological characteristics in gastric cancer

By means of immunohistochemistry, we found a widely distribution of IL-9 in tumor tissues and the staining intensity of IL-9 varied greatly among different specimens. The representative pictures of IL-9 high and low expression are shown in Supplementary Figure 2. By flow cytometry, we assessed the surface markers expressed by tumor-infiltrating IL-9⁺ cells in freshly isolated clinical specimens. Firstly, we found that the majority of IL-9 was produced by CD45⁺ immune cells in gastric cancer tissues. Further analysis showed that nearly half of CD45⁺ IL-9⁺ cells expressed CD3 and CD4, which indicated that CD4⁺ Th cells might be one of the main producer of IL-9 in gastric cancer (Supplementary Figure 3). We further assessed the association between IL-9 expression and clinicopathologic characteristics, but no significant association was observed (Table 1).

Intratumoral IL-9 expression indicates better prognosis in gastric cancer patients

In order to investigate the prognostic value of IL-9 expression, we performed Kaplan-Meier survival analysis according IL-9 expression level. Patients with high IL-9 expression had an obvious better overall survival in the discovery (P = .002, Figure 1a), validation (P < .001, Figure 1d) and combined set (P < .001, Figure 1g). We further classified patients into an early-stage (TNM stage I) disease subgroup or an advancedstage (TNM stage II and III) disease subgroup. IL-9 high expression could still stratify patients' survival in advancedstage disease group (P < .001, P < .001 and P < .001, respectively; Figure 1c, f and i). . However, overall survival didn't show significant difference between IL-9 low and high expression subgroup in stage I patients (P = .566, P = .729 and P = .357, respectively; Figure 1b, e and h). Overall, intratumoral IL-9 expression was associated with overall survival in the patients with gastric cancer, especially for advanced-stage disease.

Intratumoral IL-9 expression acts as an independent prognostic factor

Multivariate cox regression model including IL-9 expression, adjuvant chemotherapy, TNM classification, localization, size, differentiation, Lauren classification, sex, and age were constructed to demonstrate the independent prognostic value of IL-9 expression. In multivariate cox regression model, IL-9 high expression could still indicate longer overall survival patients in discovery (HR 0.495, 95%CI 0.321–0.764, P = .001), validation (HR 0.561, 95% CI 0.382–0.823, P = .003) and combined set (HR 0.553, 95%CI 0.418–0.732, P < .001) (Table S2). These results indicated that intratumoral IL-9 expression could act as an independent prognostic factor.

Intratumoral IL-9 expression associates with response to adjuvant chemotherapy

Gastric cancer patients with Stage II/III who received postoperative ACT showed better overall survival compared with those who received surgery only (Figure 2a), which had been identified by the MAGIC and CLASSIC trails.^{15,16} To evaluate whether intratumoral IL-9 expression has an impact on response to adjuvant chemotherapy, we investigated the association between IL-9 expression and overall survival among stage II/III patients who either did or did not receive adjuvant chemotherapy. The results presented that treatment with ACT was associated with better OS only in patients with high intratumoral IL-9 expression (P < .001, Figure 2c), but not in patients with low intratumoral IL-9



Figure 1. Interleukin-9 high expression predicts better overall survival in gastric cancer. (a, d and g). Kaplan-Meier analyses comparing OS of all patients according to IL-9 expression in Discovery set (n = 207, P = .002), Validation set (n = 246, P < .001) and Combined set (n = 453, P < .001). (b, e and h). Kaplan-Meier analyses comparing OS of stage I patients according to IL-9 expression in Discovery set (n = 114, P = .357). (c, f and i). Kaplan-Meier analyses comparing OS of stage II/III patients according to IL-9 expression in Discovery set (n = 149, P < .001), Validation set (n = 190, P < 0.001) and Combined set (n = 339, P < 0.001). P-value was calculated by log-rank t test. OS = overall survival.

expression (P = .087, Figure 2b). A test for the interaction between IL-9 expression and the treatment revealed that the benefit observed in IL-9 high subgroup was significantly superior to that observed in IL-9 low subgroup (P for interaction = 0.034). Multivariate analysis suggested that association between IL-9 expression and benefit from adjuvant chemotherapy was not confounded by risk factors that are known to affect the survival rates among patients with stage II and stage III disease (Figure 2d). All these results demonstrated that patients with high intratumoral IL-9 expression might benefit more from adjuvant chemotherapy.

IL-9 high expression was associated with an anti-tumor microenvironment

In order to figure out a possible mechanism for the different OS between IL-9 high and low expression patients, we decided to

look into the differences of immune cell infiltration between two groups of patients. We found the number of intratumoral CD8⁺ T cells, NK cells, CD4⁺ T cells, macrophages and B cells were significantly higher in patients with IL-9 high expression (Supplementary Figure 4). Then, we focused on CD8⁺ T cells owing to their central role in anti-tumor immunity.¹⁷ Positive correlation was found between intratumoral IL-9 expression and number of CD8⁺ T cells by immunohistochemistry (Figure 3a). In order to detect the infiltration and exact functional state of CD8⁺ T cells in IL-9 high/low tumors, flow cytometry analysis were conducted with 44 additional patients and showed higher proportion of CD8⁺ T cells among CD45⁺ leukocytes (Figure 3b). Then, we compared the functional state of CD8⁺ T cells between two groups of patients. It was found that the percentage of CD8⁺ T cells expressing proliferative marker Ki-67 and effector molecules including interferon gamma (IFN-y), granzyme B (GZMB)



Figure 2. Interleukin-9 high expression is associated with better benefit from adjuvant chemotherapy in stage II/III gastric cancer patients. (a-c). Kaplan-Meier analyses comparing OS according to treatment in all stage II/III patients (left panel), IL-9 low expression subgroup (middle panel) and IL-9 high expression subgroup (right panel). Subgroup interaction analysis showed IL-9 expression could distinguish the benefit from ACT (P = .034). (d). Multivariate analysis adjusted for TNM stage, localization, tumor size, differentiation, Lauren classification Sex, age and IL-9 expression. Interaction *p*-value for association between clinical variable and adjuvant chemotherapy benefit are shown.

and perforin (PRF1) increased significantly in IL-9 high tumors, while CD107a didn't show significant difference (Figure 3c and d). Consistently, IL-9 expression was positively correlated with intratumoral expression of effector molecules including INF- γ , GZMB, and PRF1 by immunohistochemistry (Supplementary Figure 5). Furthermore, we found a decreased expression of immune checkpoint receptors including PD-1, CTLA-4, and TIM-3 (Figure 3e), which was reported to be associated with T cell dysfunction in tumor tissues. Together, we found elevated intratumoral IL-9 expression correlated with an enhanced antitumor immunity in gastric cancer.

Recombined human IL-9 enhanced anti-tumor immune response by CD8⁺ T cells in vitro

With the positive correlation between IL-9 expression and antitumor immunity by CD8⁺ T cells, we detected the expression of IL-9 receptor (IL-9 R) on CD8⁺ T cells in gastric cancer tissues and found that about 59.8% CD8⁺ T cells showed positive expression of IL-9 R (Supplementary Figure 6). Furthermore, in vitro experiments were conducted to validate the impact of IL-9 on intratumoral CD8⁺ T cells. Flow cytometry analysis indicated that adding recombined human IL-9 (rhIL-9) significantly increased the expression of effector molecules including IFN-γ, GZMB and PRF1 on CD8⁺ T cells (Figure 4a–c), which was validated by Elisa (Supplementary Figure 7). We also detected the proliferative and apoptotic status of tumor cells to evaluate the anti-tumor effect of rhIL-9. It was shown that rhIL-9 caused decreased frequency of Ki67⁺ but increased frequency of Annexin V⁺ tumor cells (Figure 4d and e) Interestingly, the anti-tumor activity of rhIL-9 was abolished when CD8⁺ T cells were depleted from single-cell suspension (Figure 4d and e), which indicated that the anti-tumor activity of IL-9 might be mediated by CD8⁺ T cells. Taken together, these findings confirmed that IL-9 enhanced the anti-tumor immunity by CD8⁺ T cells in gastric cancer.

Recombined human IL-9 augmented responses to PD-1 blockade

As immune checkpoint inhibitors (ICPI) such as Pembrolizumab exert therapeutic effect also by motivating cytotoxicity by CD8⁺ T cells, we asked whether rhIL-9 was synergistic with ICPI treatment. Thus, we performed a combined treatment utilizing rhIL-9 and Pembrolizumab, an anti-PD-1 antibody. It was shown that combination of rhIL-9 and Pembrolizumab led to further functional restoration of intratumoral CD8⁺ T cells (Figure 5a–c and



Figure 3. Interleukin-9 expression positively correlates with the numbers and functional status of CD8⁺ T cells in gastric cancer. (a). Scatter diagram showing the positive correlation between IL-9 expression and CD8⁺ T cells infiltration by means of immunohistochemistry. (b). Proportion of CD8⁺ T cells among CD45⁺ leukocytes in IL-9 high and low expression subgroup by flow cytometry (n = 37). (c). Expression of proliferation marker Ki-67 on CD8⁺ T cells in IL-9 high and low expression subgroup (n = 12). (d). Expression of effector molecules (IFN- γ , GZMB, PRF1 and CD107a; n = 37,37,37,35, respectively) on CD8⁺ T cells in IL-9 high and low expression subgroup. (e). Expression of co-inhibitory receptors (PD-1, CTLA-4, TIM-3 and LAG-3; n = 37,37,37, respectively) on CD8⁺ T cells in IL-9 high and low expression subgroup. Each symbol represents an individual patient. IFN- γ , interferon- γ ; GZMB, granzyme B; PRF1, perforin-1; PD-1, programmed cell death-1; CTLA-4, cytotoxic T-lymphocyte-associated protein 4; TIM-3, T-cell immunoglobulin and mucin-domain containing-3; LAG-3, lymphocyte-activation gene-3. *P < .05, **P < .01 and *** *P* < .001.



Supplementary Figure 7). Likewise, we uncovered the synergistic effect of rhIL-9 and Pembrolizumab on triggering the apoptosis and limiting proliferation of tumor cells (Figure 5d and e). Taken together, these findings indicate that rhIL-9 may elicit a synergistic effect with Pembrolizumab to facilitate functional restoration of CD8⁺ T cells and the clearance of tumor cells.

Discussion

In this study, we explored the clinical significance and impact on anti-tumor immunity of intratumoral IL-9 in gastric cancer. It was shown that intratumoral IL-9 expression had strong prognostic value and indicated better adjuvant chemotherapy benefits in gastric cancer patients. In vitro experiments showed that rhIL-9 could enhance anti-tumor immunity by CD8⁺ T cells and raise the efficacy of PD-1 blockade on CD8⁺ T cells. These findings suggested that efforts to increase intratumoral IL-9 level might be new immunotherapeutic strategy in gastric cancer.

Prognostic assessment is crucial for formation of appropriate treatment choices. Recently, the association between immune contexture and clinical outcome has been drawing more and more attention owing to their critical role in tumorigenesis and progression.¹⁸ Our previous studies have demonstrated the clinical significance of tumor-infiltrating immune cells including macrophages, neutrophils, mast cells and IL-17-producing cells in gastric cancer,^{6,7,19,20} here we concluded that the immune regulatory cytokine, IL-9 could also serve as an independent prognostic factor for overall survival in gastric cancer patients.

Remarkably, further analysis showed IL-9 high expression also indicated better benefit from 5-fluorouracil-based adjuvant chemotherapy in gastric cancer patients with stage II/III disease. Although 5-fluorouracil-based ACT has been recommended as a first-line adjuvant therapy for stage II/III patients with gastric cancer, not everyone was responsive to chemotherapy and the criterion for selection of candidates is still controversial.²¹ Thus, our findings provided more evidence for better selection of patients who should receive adjuvant chemotherapy.

We showed that IL-9 exhibited anti-tumor role in gastric cancer, which was consistent with studies on melanoma¹³ and colon cancer,²² but contradictory with those on breast cancer,¹² non-small cell lung cancer²³ and hepatocellular cancer.¹¹ Clearly, IL-9 has a complicated dual function depending on tumor type, tumor microenvironment, or other experimental conditions. For example, it was found that IL-9 was involved in proliferation, cell growth, and migration process by triggering JAK/STAT signaling of tumor cells.²⁴ However, multiple studies demonstrated that IL-9 also had anti-tumoral functions by provoking robust anti-tumor immunity.²⁵ Our study discovered that IL-9 enhanced the cytotoxicity by CD8⁺ T cells in gastric cancer, which gave possible explanations for the better prognosis and chemotherapy sensitivity of patients with high IL-9 expression. But indeed, multiple cell types like mast cells and dendritic cells might also be target cells by IL-9 in tumor microenvironment.26,27

Immune checkpoint inhibitors have made a clinically significant impact on the treatment of several different types of



Figure 5. Recombined human IL-9 augmented responses to PD-1 blockade by CD8⁺ T cells in gastric cancer. (a, b and c). Effector molecule expression (GZMB, IFN- γ , PRF-1) on CD8⁺ T cells treated with isotype, rhlL-9, Pembrolizumab or combination of rhlL-9 and Pembrolizumab (n = 10,10,5, respectively). (d). Ki-67⁺ tumor cells frequencies of tumor cells in gastric cancer tissue samples from patients after treated isotype, rhlL-9, Pembrolizumab or combination of rhlL-9 and Pembrolizumab (n = 10). *P < .05, **P < .01 and ***P < .001.

cancer, of which PD-1 inhibitors were the most successful immunotherapy drugs. However, PD-1 inhibition is not capable of reversing all immunosuppressive mechanisms, and a large proportion of patients do not respond adequately to anti-PD-1 immunotherapies.²⁸ Thus, synergistic combinations of checkpoint with other strategies have been proposed to further enhance the antitumor efficacy of anti-PD-1 individual treatments.²⁹ A previous study revealed that peripheral blood Th9 cells, which produce IL-9, play an important role in the successful treatment of melanoma patients with PD-1 blockade.³⁰ While in our study, we further discovered that rhIL-9 and Pembrolizumab had a synergistic role in reactivated intratumoral CD8⁺ T cells. These results provided the preclinical rational for combination rhIL-9 and PD-1 blockade to restore anti-tumor immune response.

Several mechanisms involved in IL-9-mediated tumor suppression has been discovered. Purwar R et al found that IL-9 could activate the antitumorigenic effect of innate immune cells like mast cells.²⁵ Moreover, IL-9 was shown to involve CCL20-mediated leukocyte recruitment into the tumor and the regulation of dendritic cell function in a CCR6-dependent manner.¹³ Apart from the stimulation of the innate immune system, IL-9 could also activate adaptive immune cells. For example, IL-9 is responsible for the expansion and IFN-γ production of CD8⁺ T cells.³¹ However, there are also studies demonstrated that IL-9 also had pro-tumoral functions in several hematopoietic malignancies as well as solid tumors like hepatocellular cancer and lung cancer. Most studies indicated that IL-9 could drive JAK/STAT3 signaling activity and thus trigger tumor cell proliferation and migration.^{11,24} All this evidence suggested that, as a pleiotropic cytokine, IL-9 is not limited to any one function and has been shown to induce various physiological responses in tumor.

A mainly limitation of our study was the lack of in vivo validation for our findings because no appropriate mouse-derived cells line for gastric cancer could be acquired for building immunecompetent mice model. Furthermore, as the intervention experiments were based on bulks of cells dissociated from tumor tissues, it remains to be explored whether the effect of rhIL-9 on CD8⁺ T cells was direct or indirect.

In summary, we reported for the first time that intratumoral IL-9 expression could serve as a prognostic and predictive indicator in gastric cancer. Enhancing IL-9 signaling might be an effective therapeutic approach in gastric cancer, either alone or in combination with immune checkpoint inhibitors.

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Authors' contributions

H. Fang, R. Li and Y. Gu for acquisition of data, analysis and interpretation of data, statistical analysis and drafting of the manuscript; Y. Fei, K. Jin, Y. Chen, Y. Cao, X. Liu, K. Lv, J. Wang, K. Yu, C. Lin, H. Liu, H. Li and H. He for technical and material support; W. Zhang, H. Zhang, and Z. Shen for study concept and design, analysis and interpretation of data, drafting of the manuscript, obtained funding and study supervision. All authors read and approved the final manuscript.

Disclosure of potential conflicts of interest

No potential conflicts of interest were disclosed.

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