CD163 as a marker of M2 macrophage, contribute to predict aggressiveness and prognosis of Kazakh esophageal squamous cell carcinoma

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ABSTRACT

M2 macrophages was domesticated by tumor microenvironment to produce some angiogenic molecules and protease, facilitating angiogenesis and matrix breakdown, promoting tumor invasive and metastasis. However, The function of M2 macrophages to progression of esophageal carcinoma, especially Kazakh esophageal carcinoma is still dimness. This study aims to investigate M2 macrophages correlated with matrix metalloproteinase-9 (MMP9) and microvessel density, and the role in the progression of Kazakh esophageal squamous cell carcinoma. CD163 and CD34 as the marker of M2 macrophages and endothelial cells, were used to identify the M2 macrophages density and microvessel density, respectively. Immunohistochemistry staining was evaluated the expression of MMP9. The number of infiltrated CD163-positive M2 macrophages in tumor islets and stroma was significantly higher than in cancer adjacent normal tissues. The increased of M2 macrophages and microvessel density were significantly correlated with more malignant phenotypes including lymph node metastasis and clinical stage progression. Meanwhile, the expression of MMP9 showed much higher level in esophageal squamous cell carcinoma than that in cancer adjacent normal tissues, and high expression of MMP9 in Kazakh esophageal squamous cell carcinoma was significantly associated with age, depth of tumor invasion, lymph node metastasis, and tumor clinical stage. The quantity of M2 macrophages in tumor stroma was positively associated with microvessel density and the expression of MMP9, and as an independent poorly prognostic factor for overall survival time of Kazakh esophageal squamous cell carcinoma. These findings suggest the increased number of M2 macrophages correlated with high expression of MMP9 and high microvessel density may contribute to the tumor aggressiveness and angiogenesis, promoting the progression of Kazakh esophageal squamous cell carcinoma.

INTRODUCTION

Esophageal carcinoma is one of most common malignant tumors in the world. According to its etiological and pathological characteristics, it is divided into two main forms: esophageal squamous cell carcinoma (ESCC) and esophageal adenocarcinoma. More than 90% of esophageal cancers are classified as ESCCs in China [1]. The Kazakh national minority (ethnic) living in Xinjiang (northwest of China) is a demographic with one of the highest rates of esophageal carcinoma incidence and mortality in China [2]. The 5-year survival rate of esophageal carcinoma is only 10%. Primary reasons for poor prognosis are associated with early stage cancer cell invasion and high metastasis [3].
The tumor microenvironment is important for cancer development and metastasis [4, 5]. It contains a range of inflammatory and immune cells. Macrophages are essential immune cells that play a critical role in carcinogenesis and tumor progression [6]. Mirroring T helper type 1, T helper type 2 (TH1, TH2) polarization, two distinct states of polarized activation for macrophages have been recognized: the classically activated (M1) macrophage phenotype and the alternatively activated (M2) macrophage phenotype [7, 8]. Bacterial moieties such as LPS and the interferon-γ (IFN-γ) polarize macrophages toward the M1 phenotype, showed more proinflammatory and antitumor activity, high expression of tumor necrosis factor alpha [TNF-α, interleukin [IL]-6, IL-12] and major histocompatibility complex (MHC-II) molecules. In contrast, IL-4, IL-10, and IL-13 polarize macrophages toward the M2 phenotype, showed more phagocytic activity and carcinogenesis function, high expression of IL-10 and arginase, but low expression of MHC class II and IL-12 [9]. Most M2 macrophages are considered to be tumor associated macrophages (TAMs) and function to promote tumor angiogenesis and metastasis [10]. One of the mechanism is M2 macrophages which could produce growth factors and proteases that degrade the tissue extracellular matrix (ECM), and facilitate the tumor cells escape from their constraining basement membranes [11]. In this process, matrix metalloproteinases (MMPs), particularly MMP9, which is delivered into the tumor microenvironment by M2 macrophages, favoring tumor stromal angiogenesis [12], and other study also showed M2 macrophages produces many proangiogenic factors and expresses high levels of MMP9, which promote cancer cell invasion, metastasis and angiogenesis [13]. However, the precise role of M2 macrophage in Kazakh esophageal carcinoma has yet to be elucidated. So we used CD163 and CD34 as marker of M2 macrophages and endothelial cells to assay the M2 macrophages density and microvessel density (MVD) separately, then we used immunohistochemistry (IHC) to evaluate the MMP9 expression. Ultimately, we aim to investigate whether the increased number of M2 macrophages was correlated with high expression of MMP9 and high MVD, which contributes to the tumor aggressiveness and angiogenesis, and facilitates the occurrence and progression of Kazakh ESCC.

RESULTS

Distributions of CD163-positive M2 macrophages in Kazakh ESCCs and the correlation of M2 macrophages density with Kazakh ESCC clinicopathological parameters

We used CD163 as a marker to assess M2 macrophages distribution. Immunohistochemistry staining for CD163 revealed diffuse staining of M2 macrophages membranes and cytoplasm (Figure 1). We found that M2 macrophages were primarily located in the tumor stroma, but a small number of M2 macrophages reside in tumor islets. The density of M2 macrophages in Kazakh ESCC tumor islets (approximately 15/HPF, 0–45) and stroma (approximately 58/HPF, 9–139) were significantly higher compared to cancer adjacent normal (CAN) epithelia (approximately 2/HPF, 0–10) and stroma (approximately 19/HPF, 3–54) (all P < 0.001, Table 1). To explore the role of M2 macrophages in progression of Kazakh ESCCs, we divided the cases into high and low density of M2 macrophage groups according to the median value of CD163-positive macrophages (15 for tumor islet and 58 for tumor stroma), and evaluated possible correlations between M2 macrophages density and Kazakh clinicopathological parameters, including age, gender, tumor location, histological grade, invasion depth, nodal status, and clinical stages, found the increased number of M2 macrophages in tumor stroma were significantly correlated with more malignant phenotypes of Kazakh ESCCs, including lymph node metastasis and clinical stage progression (all P = 0.001). Similar, high density of M2 macrophages in tumor islet also were correlated with lymph node metastasis of ESCCs (P < 0.05). No significant correlations were found between M2 macrophages distribution and other parameters of Kazakh ESCCs (P > 0.05, Table 2).

Distributions of microvessel in Kazakh ESCCs and the correlation of microvessel density (MVD) with Kazakh ESCC clinicopathological parameters

CD34 staining was used to assess MVD in tumor tissues. Immunohistochemistry staining for CD34 revealed diffuse staining of endothelial cells membranes and cytoplasm (Figure 1). The mean value of MVD in Kazakh ESCCs was 17 per high-power field (range, 3–31), which was significantly higher compared to CAN (mean, 9/HPF; range, 2–19) (Supplementary Figure 1). Similar to classify of M2 macrophages density, we divided the cases into high and low density of microvessel groups according to the median value of MVD, and we found high MVD in Kazakh ESCCs was significantly associated with more malignant phenotypes including lymph node metastasis and clinical stage progression (all P < 0.05) (Table 3).

MMP9 expression in Kazakh ESCCs and CANs, and its relationship with ESCC clinicopathological parameters

M2 macrophages are reported to be involved in tumor growth and metastasis, and are suspected to be important for tumor progression [14]. MMP-9 is delivered into the tumor microenvironment by M2 macrophages, and subsequently sustains those distinct angiogenic
Figure 1: The distribution of CD163-positive M2 macrophages and microvessel in Kazakh esophageal squamous cell carcinoma (ESCC) and Cancer adjacent normal (CAN) tissues. (A–D) Immunohistochemical staining of CD163, which was used as a marker of M2 macrophages and to evaluate the density of M2 macrophages in ESCC and CAN tissues. (A) and (B) showed the distribution of M2 macrophages in ESCC tumor stromal and islet, CD163 revealed diffuse staining of membranes and cytoplasm of M2 macrophages, and showed the high density of M2 macrophages located in ESCC tissues (especially in tumor stroma). (C) and (D) showed the distribution of M2 macrophages in CAN stroma and epithelia. A small number of CD163-positive M2 macrophages appear in CAN tissues. (E and F) Immunohistochemical staining of CD34, which was used to mark endothelial cells and to evaluate the microvessel density (MVD) in ESCC and CAN tissues. (E) showed high MVD in Kazakh ESCC tissues. (F) showed low MVD in Kazakh ESCC tissues.
Table 1: The distribution of CD163-positive macrophages in Kazakh esophageal squamous cell carcinoma (ESCC) and Cancer adjacent normal (CAN) tissues

| Groups  | Cases (N) | Islets (count range) | Z/P   | Stroma (count range) | Z/P   |
|---------|-----------|----------------------|-------|----------------------|-------|
| ESCCs   | 100       | 15 (0–45)            | 11.559/0.000* | 58 (9–139)        | 10.081/0.000* |
| CANs    | 100       | 2 (0–10)             |       | 19 (3–54)           |       |

*P < 0.05.

Table 2: Tumor infiltrating CD163-positive macrophages in Kazakh esophageal squamous cell carcinoma (ESCC) islets and stroma, and their correlation with clinicopathological parameters

| Variable             | Cases (N) | CD163+ macrophage counts in tumor islets | CD163+ macrophage counts in tumor stroma |
|----------------------|-----------|-----------------------------------------|-----------------------------------------|
|                      |           | Low (< 15) | High (≥ 15) | P     | Low (< 58) | High (≥ 58) | P     |
| Age (y)              |           |            |             |       |            |             |       |
| ≤ Median (58 y)      | 53        | 34 (64.2%) | 19 (35.8%)  | 1.000 | 31 (58.5%) | 22 (41.5%)  | 0.164 |
| > Median             | 47        | 31 (66.0%) | 16 (34.0%)  |       | 20 (42.6%) | 27 (57.4%)  |       |
| Gender               |           |            |             |       |            |             |       |
| Male                 | 68        | 47 (69.1%) | 21 (30.9%)  | 0.301 | 35 (51.5%) | 33 (48.5%)  | 1.000 |
| Female               | 32        | 18 (56.3%) | 14 (43.8%)  |       | 16 (50.0%) | 16 (50.0%)  |       |
| Tumor location       |           |            |             |       |            |             |       |
| Upper                | 2         | 2 (100.0%) | 0 (0.00 %)  | 1.000 | 1 (50.0%)  | 1 (50.0%)   |       |
| Middle               | 70        | 47 (67.1%) | 23 (32.9%)  | 0.272 | 36 (51.4%) | 34 (48.6%)  | 0.991 |
| Lower                | 28        | 16 (57.1%) | 12 (42.9%)  |       | 14 (50.0%) | 14 (50.0%)  |       |
| Histologic grade     |           |            |             |       |            |             |       |
| Well                 | 29        | 21 (72.4%) | 8 (27.6%)   |       | 15 (51.7%) | 14 (48.3%)  |       |
| Moderate             | 47        | 30 (63.8%) | 17 (36.2%)  | 0.545 | 25 (53.2%) | 22 (46.8%)  | 0.838 |
| poor                 | 24        | 14 (58.3%) | 10 (41.7%)  |       | 11 (45.8%) | 13 (54.2%)  |       |
| Depth of invasion    |           |            |             |       |            |             |       |
| TI–T2                | 35        | 19 (54.3%) | 16 (45.7%)  | 0.153 | 20 (57.1%) | 15 (42.9%)  | 0.489 |
| T3–T4                | 65        | 46 (70.8%) | 19 (29.2%)  |       | 31 (47.7%) | 34 (52.3%)  |       |
| Nodal status         |           |            |             |       |            |             |       |
| pN−                  | 51        | 39 (76.5%) | 12 (23.5%)  | 0.025*| 35 (68.6%) | 16 (31.4%)  | 0.001*|
| pN+                  | 49        | 26 (53.1%) | 23 (46.9%)  |       | 16 (32.7%) | 33 (67.3%)  |       |
| Clinical stage       |           |            |             |       |            |             |       |
| I–II                 | 62        | 42 (67.7%) | 20 (32.3%)  | 0.604 | 40 (64.5%) | 22 (35.5%)  | 0.001*|
| III–IV               | 38        | 23 (60.5%) | 15 (39.5%)  |       | 11 (28.9%) | 27 (71.1%)  |       |

Abbreviations: pN−, no lymph node metastasis; pN+: node metastasis. *P < 0.05.
vessels capable, supporting tumor cell intravasation and metastatic dissemination [15]. So we decided to investigate whether MMP9 is the key molecule mediated by M2 macrophages to play an important role in the occurrence and progression of Kazakh ESCC. As shown in Figure 2, MMP9 staining was mainly observed in tumor stromal cells (located in cell membranes and cytoplasm), including macrophages, ESCC tumor cells and endothelial cells (ECs). We compared three categories of MMP9 positive staining combinations (1+, 2+/3+, and 1+/2+/3+) to MMP9 negative (0) staining (Table 4). The MMP9 positive rate (1+/2+/3+) in ESCC was higher than that in CAN tissues (90.0% vs 56.0%; *P* < 0.001). Differences in MMP9 positive rates from ESCC to CAN tissues were more prominent in strong staining cases (2+/3+; 55.0% vs 16.0%; *P* < 0.001).

In order to investigate the role of MMP9 in the progression of Kazakh ESCCs, we divided cases into two categories according to MMP9 expression level: low expression (−/1+) and high expression (2+/3+). The expression level of MMP9 was significantly higher in over-58-year old patients compared to under-58-years.

### Table 3: Correlation between microvessel density (MVD) and clinicopathological parameters in Kazakh esophageal squamous cell carcinoma (ESCC) tumor stroma

| Variable                  | Cases (N) | MVD Low (≤ 17) | MVD High (≥ 17) | *P*  |
|---------------------------|-----------|----------------|-----------------|------|
| Age (y)                   |           |                |                 |      |
| ≤ Median (58y)            | 53        | 28 (52.8%)     | 25 (47.2%)      | 0.540|
| > Median                  | 47        | 21 (44.7%)     | 26 (55.3%)      |      |
| Gender                    |           |                |                 |      |
| Male                      | 68        | 31 (45.6%)     | 37 (54.4%)      | 0.435|
| Female                    | 32        | 18 (56.2%)     | 14 (43.8%)      |      |
| Tumor location            |           |                |                 |      |
| Upper                     | 2         | 0 (0.0%)       | 2 (100.0%)      | 0.255|
| Middle                    | 70        | 35 (50.0%)     | 35 (50.0%)      |      |
| Lower                     | 28        | 14 (50.0%)     | 14 (50.0%)      |      |
| Histologic grade          |           |                |                 |      |
| Well                      | 29        | 15 (51.7%)     | 14 (48.3%)      |      |
| Moderate                  | 47        | 26 (55.3%)     | 21 (44.7%)      | 0.197|
| poor                      | 24        | 8 (33.3%)      | 16 (66.7%)      |      |
| Depth of invasion         |           |                |                 |      |
| T1–T2                     | 35        | 19 (54.3%)     | 16 (45.7%)      | 0.571|
| T3–T4                     | 65        | 30 (46.2%)     | 35 (53.8%)      |      |
| Nodal status              |           |                |                 |      |
| pN−                       | 51        | 31 (60.8%)     | 20 (39.4%)      | 0.027*|
| pN+                       | 49        | 18 (36.7%)     | 31 (63.3%)      |      |
| Clinical stage            |           |                |                 |      |
| I–II                      | 62        | 36 (58.1%)     | 26 (41.9%)      | 0.035*|
| III–IV                    | 38        | 13 (34.2%)     | 25 (65.8%)      |      |

Abbreviations: pN−, no lymph node metastasis; pN+: node metastasis. *P* < 0.05.
patients (68.1% vs 43.4%; \( P < 0.05 \)). Cases with high MMP9 expression showed strong invasion (T3–T4 vs T1–T2 = 63.1% vs 40.0%; \( P < 0.05 \)) and metastasis (pN+ vs pN− =73.5% vs 37.7%; \( P = 0.001 \)), and were clearly present in advanced ESCC stages (III–IV vs I–II = 78.9% vs 40.3%; \( P < 0.001 \)) (Table 5).

**Correlation between the density of M2 macrophages, microvessel density (MVD) and MMP9 expression in Kazakh ESCCs**

To explore whether M2 macrophages promote the invasion, metastasis and angiogenesis of Kazakh ESCC by mediating MMP9, we investigated the relationship between the density of M2 macrophages, MVD and the expression of MMP9. Spearman’s rank analysis demonstrated positive correlation between MVD and the density of M2 macrophages in tumor stroma (\( r = 0.231; \ P < 0.05 \)). Similar, the expression of MMP9 and the density of M2 macrophages in tumor stroma (\( r = 0.455; \ P < 0.001 \)) and islets (\( r = 0.374; \ P < 0.001 \)) also showed positive correlation respectively (Figure 3C, 3D and 3E). Meanwhile, MVD showed positively association with the expression of MMP9 in Kazakh ESCCs (\( r = 0.292; \ P < 0.05 \); Figure 3F).

**Association of CD163-positive M2 macrophages with Kazakh ESCC Survival time**

To investigate the association of M2 macrophages with Kazakh ESCC prognosis, we divided Kazakh ESCC patients into two groups based on the median density of M2 macrophages, high and low density of M2 macrophages. Kaplan–Meier survival curves were used to investigate the influence of M2 macrophages to ESCC patients overall survival (OS) time. We found there was a significant negative correlation between the OS and the density of M2 macrophages in tumor stroma of Kazakh patients.

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**Figure 2: Immunohistochemical staining of MMP9 in Kazakh esophageal squamous cell carcinoma (ESCC) and Cancer adjacent normal (CAN) tissues.** MMP9 staining is primarily observed in tumor stroma (cell membranes and cytoplasm), some ESCC cells also show staining. (A) Negative MMP9 staining is shown in CAN tissues (scored as 0). (B) Weak MMP9 staining is shown in ESCC tissues (scored as 1). (C) and (D) show moderate and strong MMP9 staining in Kazakh ESCC tissues, respectively (scored as 2 and 3, respectively).
### Table 4: The expression of MMP9 in Kazakh esophageal squamous cell carcinoma (ESCC) and cancer adjacent normal (CAN) tissues

| Characteristics | N   | Negative | Positive combination |
|-----------------|-----|----------|----------------------|
|                 |     | 0 (%)    | 1+ (%) | X²   | P   | 2+3+ (%) | X²   | P   | 1+2+3+ | X²   | P   |
| ESCCs           | 100 | 10 (10.0%) | 35 (35.0%) | 9.393 | 0.002* | 55 (55.0%) | 40.369 | 0.000* | 90 (90.0%) | 27.625 | 0.000* |
| CANs            | 100 | 44 (44.0%) | 41 (41.0%) | 5.178 | 0.023* | 56 (56.0%) | 11.819 | 0.001* | 56 (56.0%) | 12.684 | 0.000* |

*p < 0.05.

### Table 5: Correlation between expression of MMP9 and clinicopathological parameters in Kazakh esophageal squamous cell carcinoma (ESCC) tissues

| Variable | Cases (N) | MMP9 low expression | MMP9 high expression | X² | P   |
|----------|-----------|----------------------|----------------------|----|-----|
|          |           | 0/1+(%)              | 2+/3+(%)             |    |     |
| Age (y)  |           | 53                   | 30 (56.6%)           | 23 (43.4%) | 5.178 | 0.023* |
|          | ≤ Median (58y) | 53                   | 30 (56.6%)           | 23 (43.4%) | 5.178 | 0.023* |
|          | > Median   | 47                   | 15 (31.9%)           | 32 (68.1%) | 0.002 | 0.966  |
| Gender   |           | 68                   | 30 (44.1%)           | 38 (55.9%) | 2.597 | 0.273  |
|          | Male       | 68                   | 30 (44.1%)           | 38 (55.9%) | 2.597 | 0.273  |
|          | Female     | 32                   | 15 (46.9%)           | 17 (53.1%) | 0.002 | 0.966  |
| Tumor location |       | 70                   | 30 (42.9%)           | 40 (57.1%) | 2.699 | 0.259  |
|          | Upper      | 2                    | 2 (100.0%)           | 0 (0.0%) | 0.002 | 0.966  |
|          | Middle     | 70                   | 30 (42.9%)           | 40 (57.1%) | 2.699 | 0.259  |
|          | Lower      | 28                   | 13 (46.4%)           | 15 (53.6%) | 0.002 | 0.966  |
| Histologic grade |     | 29                   | 10 (34.5%)           | 19 (65.5%) | 11.819 | 0.001* |
|          | Well       | 29                   | 10 (34.5%)           | 19 (65.5%) | 11.819 | 0.001* |
|          | Moderate   | 47                   | 25 (53.2%)           | 22 (46.8%) | 2.699 | 0.259  |
|          | poor       | 24                   | 10 (41.7%)           | 14 (58.3%) | 0.002 | 0.966  |
| Depth of invasion |   | 35                   | 21 (60.0%)           | 14 (40.0%) | 4.007 | 0.045* |
|          | TI–T2      | 35                   | 21 (60.0%)           | 14 (40.0%) | 4.007 | 0.045* |
|          | T3–T4      | 65                   | 24 (36.9%)           | 41 (63.1%) | 4.007 | 0.045* |
| Nodal status |       | 51                   | 32 (62.7%)           | 19 (37.7%) | 11.819 | 0.001* |
|          | pN−        | 51                   | 32 (62.7%)           | 19 (37.7%) | 11.819 | 0.001* |
|          | pN+        | 49                   | 13 (26.5%)           | 36 (73.5%) | 11.819 | 0.001* |
| Clinical stage |     | 62                   | 37 (59.7%)           | 25 (40.3%) | 12.684 | 0.000* |
|          | I–II       | 62                   | 37 (59.7%)           | 25 (40.3%) | 12.684 | 0.000* |
|          | III–IV     | 38                   | 8 (21.8%)            | 30 (78.9%) | 12.684 | 0.000* |

Abbreviations: pN−, no lymph node metastasis; pN+: node metastasis. *P < 0.05.
Figure 3: Cross correlation analyses reveal strong relationships among the density of M2 macrophage in tumor islet, tumor stroma and expression of MMP9, microvessel density (MVD) in Kazakh esophageal squamous cell carcinoma (ESCC). (A) Significant correlation was observed between the density of M2 macrophages in Kazakh ESCCs tumor islet and stroma ($r = 0.422; P < 0.001$). (B) and (C) Significant correlation was observed between M2 macrophages in Kazakh ESCCs tumor islet, stroma and the expression of MMP9 ($r = 0.374$, $r = 0.455; P < 0.001$). (D) and (E) Significant correlation between the density of M2 macrophages in Kazakh ESCCs tumor stroma and MVD ($r = 0.231; P < 0.05$), but not between the density of M2 macrophages in Kazakh ESCCs tumor islets and MVD ($r = 0.101; P > 0.05$). (F) Significant correlation between the density of the expression of MMP9 and MVD in Kazakh ESCCs.
ESCC patients \((P < 0.001)\), but not with the density of M2 macrophages in tumor islet \((P > 0.05, \text{Figure 4})\).

Univariate Cox proportional hazard analysis revealed lymph node metastasis \((HR = 15.629)\), later clinical stage \((HR = 22.352)\) and the high density of M2 macrophages in tumor stroma \((HR = 18.521)\) were poor prognostic factors for the patients of Kazakh ESCC \((\text{all } P < 0.001, \text{Table 6})\). Moreover, after Multivariate Cox proportional hazard analysis, we found high-density of M2 macrophages in tumor stroma could serve as an independent prognostic factor for the patients of Kazakh ESCCs \((HR = 5.464, \ P < 0.05, \text{Table 6})\).

**DISCUSSION**

M2 macrophages are prominent stromal cells that play an important role in tumor growth, angiogenesis and metastasis. Several studies also showed higher density of M2 macrophages in tumor islets and stroma have close relationship with tumor progression and poor prognosis \([11, 14]\). Nevertheless, the precise role of M2 macrophages in occurrence and progression of Kazakh esophageal carcinoma has yet to be elucidated.

CD163 is confirmed to be a phenotypic marker of M2 macrophages that can be used to distinguish M2 and M1 macrophages. In this study, we found that CD163-positive M2 macrophages were primarily located in the tumor stroma, a few also distributed in the tumor islet of Kazakh ESCCs. The density of M2 macrophages in the tumor islet and stroma was significantly higher than that in corresponding CAN tissues, and the increased number of M2 macrophages in stroma was positively associated with more malignant phenotypes including lymph node metastasis and clinical stage progression of Kazakh ESCCs. Such results are similar to those reported in gastric cancers, oral carcinoma and endometrial adenocarcinoma \([16–19]\). Further study we found high-density of M2 macrophages infiltrated in tumor stroma could be an independent poorly prognostic factor for Kazakh ESCC evaluation. He and Clear’s studies also showed that a high-density of macrophages was associated with worse prognosis in oral cancer and Follicular lymphoma \([20, 21]\). Prompt increased numbers of M2 macrophages, especially in tumor stroma, were closely related to the occurrence and progression of Kazakh ESCCs, and could be served as an independent poorly prognostic factor for Kazakh ESCCs.

Widespread acceptance, angiogenic vessels developing in the primary tumor are structurally abnormal and functionally immature \([22]\), which provides not only the nutrients and oxygen for tumor growing, but also formed the vascular networks to sustain active intravasation of tumor cells and conduits for tumor cells metastasis. In human cancer tissues, the most common method for semiquantitative evaluation of angiogenesis is to measure microvessel density (MVD) using endothelial markers \([23]\), and which is recognized as one of the most useful prognostic markers for tumor progression and survival in multitudinous cancer patients \([24–26]\). In this study, CD34 as marker of endothelial cells was used to assess MVD in ESCC tissues. We found the mean value

![Figure 4: Kaplan–Meier overall survival curves of Kazakh esophageal squamous cell carcinoma (ESCC) patients stratified by the density of CD163 positive M2 macrophages (MØ) in tumor islet and stroma based on the median number. (A) No significant difference for overall survival between Kazakh patients with high and low density of CD163 positive M2 macrophages (MØ) in tumor islets \((P > 0.05)\). (B) Patients with high density of CD163 positive M2 macrophages (MØ) in tumor stroma had a poor overall survival \((P < 0.001)\).](https://www.impactjournals.com/oncotarget)
of MVD in tumor was significantly higher compared to corresponding CAN tissues, similar to classify of M2 macrophages density, high MVD in tumor was significantly associated with more malignant phenotypes including lymph node metastasis and clinical stage progression. Then we analyzed the correlation between MVD and the density M2 macrophages, and found the increased MVD was closely related to the increased density of M2 macrophages in tumor stroma. The results are similar to Manabu’s in ESCCs study [27]. Bailey also recently identified that increased MVD correlated with higher-density of tumor associated macrophages, play an important role in colorectal cancer progression [28]. These studies may suggest that M2 macrophages are critical for tumor angiogenesis, promoting metastasis and progression of tumor in Kazakh ESCCs.

In general, MMP-9-positive tumor-associated macrophages (TAMs) concomitant with the “restoration” of tumor angiogenesis and metastasis in genetically-deficient tumor bearers transplanted with wild-type hematopoietic cells [29], imply that MMP9 may be critical regulatory factor for macrophages to promote angiogenesis and sustain active of tumor invasion and metastasis. So, we evaluated the expression of MMP9 in ESCCs, found that MMP9 was mainly expressed in stroma cells, only a small section of tissue showed tumor cells staining, and MMP9 expression was much higher in ESCCs than in CANs, particularly in the strong MMP9 staining positive cases. Furthermore, high expression of MMP9 was significantly associated with invasion depth, lymph node metastasis and later clinical stage of Kazakh ESCCs. These results were consistent with previous esophageal carcinoma studies [30–32]. To explore whether M2 macrophages promote the invasion, metastasis and angiogenesis of Kazakh ESCCs by mediating of MMP9, we investigated the relationship between the density of M2 macrophages, MVD and the expression of MMP9, and we found significant correlation between MMP9 expression and the density of M2 macrophages either in the ESCC tumor islets or tumor stroma. We also found increased MVD was correlated with higher expression of MMP9 in Kazakh ESCC tumor tissues. Similarly, El-Kenawy’s study also showed MMP9 positively correlated with MVD in esophageal cancer [30]. These results suggest the M2 macrophages may promote angiogenesis by means of MMP9, and high density M2 macrophages and high expression of MMP9 reflect strong aggressiveness and activity of cancer lesions. Moreover, the results were also similar to some tumor vitro studies, genetic ablation of MMP9 in tumor recipients, resulting in decreased MVD in developing tumors and even preventing the angiogenic switch during cancer progression, provided original evidence for the functional involvement of host MMP9 in tumor angiogenesis [12]. Specific ablation of MMP9 positive TAMs with zoledronic acid resulted in reduced tumor angiogenesis, leading to a conclusion that TAMs deliver angiogenesis-inducing MMP9 are implicated in invasion-promoting processes such as flicking of the angiogenic switch [33, 34]. Combining the research above with our study, we prompted that M2 macrophages induced MMP9 secretion and promoted tumor aggressiveness and angiogenesis, which is closely associated with the occurrence and progression of Kazakh ESCCs.

### MATERIALS AND METHODS

#### Ethics statement

All participants were recruited from the Yili Friendship Hospital in Xinjiang, China. Each participant provided written, informed consent before enrolling in this study.
study. Protocols were approved by the institutional ethics committee of Yili Friendship Hospital in accordance with Helsinki Declaration ethical guidelines.

Studies population

A total of 200 surgically resected and paraffin-embedded human tissues were collected, including 100 Kazakh ESCC tissues and 100 Kazakh cancer adjacent normal tissues (CANs), from the Department of Pathology at Yili Friendship Hospital in Xinjiang, China (collected from 2008 to 2014). The patients consisted of 68 men and 32 women with age range of 33–76 years old and mean age of 58 years old, all had been diagnosed with ESCC, but none had received radiotherapy or chemotherapy before surgery. CAN group participants consisted of 68 men and 32 women with age range of 33–73 years old and mean age of 58 years old. All specimens were sectioned into 5 μm slices and subjected to conventional hematoxylin and eosin staining. A diagnosis of ESCC was confirmed by two pathologists following the World Health Organization histological tumor classification criteria [35]. There were 29 cases of well-differentiated ESCC, 47 cases of moderately differentiated ESCC, and 24 cases of poorly differentiated ESCC. From these, 35 and 65 cases exhibited invasion depths of T1–T2 and T3–T4, respectively. There were 49 cases with lymph node metastasis, 62 cases in clinical stages I–II, and 38 cases in clinical stages III–IV. CAN specimens, which were sampled more than 5 cm away from the cancer region, were confirmed to be free of cancer tissue. The survival status of all patients was followed up by telephone contact until December 2015. The median follow-up for living patients was 30 months (range, 1–84 months). Overall survival (OS) was defined as the interval between surgery and death or between surgery and the last follow-up for surviving patients. Among the 100 patients who were recruited, 43 (43.0%) died, and 57 (57.0%) remained alive during the follow-up period.

Immunohistochemistry

Paraffin-embedded tissue samples were cut into 4 μm-thick sections and mounted on polylysine-coated slides. Samples were dewaxed in xylene and rehydrated using a graded series of ethanol solutions. After deparaffinization, endogenous peroxidase activity was blocked by incubation in a 3% peroxide-methanol solution at room temperature (RT) for 10 mins, and then antigen retrieval was performed at 100°C in an autoclave for 7 mins. Samples were then incubated at RT for 30 mins. Afterwards, sections were washed with phosphate-buffered saline (PBS) three times for 5 mins each time. They were then incubated with mouse anti-human CD163 antigen monoclonal antibody (clone 10D6, Zhongshan Goldenbridge Biotechnology Co., LTD., Beijing, China), mouse anti-human MMP9 antigen monoclonal antibody (clone Sc-2c3, Santa Cruz Biotechnology, Santa Cruz, CA, USA) and mouse anti-human CD34 antigen monoclonal antibody (clone QBEnd/10, Zhongshan Goldenbridge Biotechnology Co., LTD., Beijing, China). Thoroughly washing with PBS was then performed, and primary antibody binding was visualized using a DAKO EnVision kit (DAKO, Glostrup, Denmark) following the manufacturer’s instructions. Finally, sections were faintly counterstained with hematoxylin and mounted with glycerol gelatin.

Immunoreactivity evaluation

The numbers of CD163-positive macrophages were analyzed as described previously [36]. Slides were examined by light microscopy and the five most representative hot spots were selected from low-power fields (LPFs, 100×) per slide using an Olympus BX51TF microscope (Olympus, Japan). Tumor islet and stroma areas were defined and the numbers of CD163-positive macrophages were counted in high-power fields (HPFs, 400×) by two pathologists. When cell counts differed by more than 10 cells per HPF, samples would be counted again a week later until recording differences were below 10 counts. The mean number of macrophages per HPF across five hot spots for every sample (tumor islet and tumor stroma) was defined as the M2 macrophages density.

Microvessel was measured by the average number of CD34 positive vessels. Similar to macrophages counts, we choose five most representative hot spots to observe and use a 400× magnification field to count the number of microvessels. Determined microvessel density (MVD) was expressed as the number of stained microvessels per optical field. The vessel was composed by any CD34 positive cell or cell cluster, and the diameter less than eight red blood cells was counted as a microvessel, as described in the Weidner method [37]. When microvessel counts differed by more than 10 per HPF, samples would be counted again a week later until recording differences were below 10 counts. The mean number of microvessel per HPF across five hot spots for every sample (tumor islet and tumor stroma) was defined as the MVD.

MMP9 immunohistochemistry (IHC) reactivity was evaluated following previously described methods [38, 39]. Positive IHC stains were defined as yellow-brown color following the manufacturer’s guidelines (Figure 3). IHC staining slides were scored as positive or negative by percentage and intensity of positive cells, where the scoring percentage of positively stained cells was as follows: 0 = < 5%, 1 = 6% – 25%, 2 = 26% – 50%, 3 = 51% – 75%, and 4 = 76% – 100%; staining intensity scoring was: 0 = absent, 1 = weak, 2 = moderate, and 3 = strong. A final score was based on multiplying both scores from individual slides (Supplementary Table 1), where: 0–1 was negative (−), 2–3 was weak positive (1+),
4–6 was moderate positive (2+), and 8–12 was strong positive (3+).

**Statistical analysis**

SPSS version 13.0 software (IBM, Endicott, NY) was used for all statistical analyses. The Mann-Whitney nonparametric test was used to compare the density of M2 macrophages and MVD between ESCC and CAN groups. For categorical analysis, the median value of M2 macrophages density and MVD were used as a cut-off point to dichotomize the continuous variables. The X² test was adopted for analysis of correlations between the expression of MMP9 and clinicopathological parameters of ESCCs. Spearman’s rank correlation method was used to evaluate the correlations between the density of M2 macrophages, MVD and MMP9 expression. Survival curves were estimated by the Kaplan Meier method. The correlation between the density of M2 macrophages, clinicopathological parameters and overall survival (OS) rates of Kazakh ESCCs was evaluated in univariate analysis, and was further analyzed with multivariate analysis by using Cox proportional hazard regression model. P-values were calculated by using the Epi-Info program, and P-values < 0.05 were considered significant.

**Abbreviations**

MØ, macrophages; ESCC, esophageal squamous cell carcinoma; CAN, cancer adjacent normal; MMPs, matrix metalloproteinases; MVD, microvessel density; interferon-γ, IFN-γ; tumor necrosis factor alpha, TNF-α; histocompatibility complex, MHC; interleukin-6, IL-6; TAM, tumor associated macrophage; IHC, immunohistochemistry.

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**CONFLICTS OF INTEREST**

The authors declare that they have no competing interests.

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