Clinicopathological and prognostic significance of PD-L1 expression in colorectal cancer: a systematic review and meta-analysis

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Abstract

Objective: To analyze the prognostic value of programmed death factor ligand 1 (PD-L1) in colorectal cancer.

Methods: Electronic databases, such as PubMed, Web of Science, Embase, and Cochrane library, were searched to identify studies evaluating the PD-L1 expression and overall survival (OS) in these patients. Afterwards, the relevant data were extracted to perform the meta-analysis.

Results: A total of 3481 patients were included in 10 studies. The combined hazard ratio (HR) was 1.22 (95%CI = 1.01–1.48, P = 0.04), indicating that high expression of PD-L1 was significantly correlated with poor prognosis of colorectal cancer. Apropos of clinicopathological features, the merged odds ratio (OR) exhibited that highly expressed PD-L1 was firmly related to lymphatic invasion (OR = 3.49, 95%CI = 1.54–7.90, P = 0.003) and advanced stage (OR = 1.77, 95%CI = 1.41–2.23, P < 0.00001), but not correlative with patients' gender, microsatellite instability, or tumor location.

Conclusion: The expression of PD-L1 can be utilized as an independent factor in judging the prognosis of colorectal cancer, and patients with advanced cancer or lymphatic invasion are more likely to express PD-L1. This conclusion may lay a theoretical foundation for the application of PD-1/PD-L1 immunoassay point inhibitors but still needs verifying by sizeable well-designed cohort studies.

Keywords: Colorectal cancer, PD-L1 expression, Prognosis, Clinicopathological features, Meta-analysis

Introduction

Among the most common cancers worldwide, colorectal cancer ranks third, accounting for 10% of all tumor cases [1]. In 2012, the disease engendered 1,400,000 new cases and nearly 700,000 deaths [2]. According to relevant research, 4.96% of the population born in the USA are suffering from colorectal cancer [3]. Even in Asia, where the incidence rate is reported to be the lowest [4], the threat posed by colorectal cancer cannot be underestimated. Taking China as an example, the incidence and mortality of colorectal cancer there have kept rising. China's cancer statistics manifest that the incidence and mortality of colorectal cancer ranked fifth among all malignant tumors in China, bringing about 380,000 new cases and 190,000 deaths annually. When they are seeking medical examination, most patients have already been in the advanced stage [5, 6]. Despite the continuous development of treatment technology, the 5-year survival rate of patients with metastatic disease is still less than 10% [7], which is probably due to the inability to diagnose early and the lack of specific markers to determine tumor development or patients' prognosis. Therefore, to enhance the prognosis of patients with colorectal cancer, it is indispensable to explore effective diagnostic and therapeutic methods.

Programmed cell death protein 1 (PD-1), a sort of inhibitory checkpoint molecule, was discovered and named by Japanese scholar Ishida in 1992 [8]. It belongs to the CD28 family and is expressed on the surface of activated T cells to regulate proliferation and activation [9]. PD-L1 (also known as B7-H1) is the dominant
ligand for PD-1 and expressed in activated T cells, B cells, dendritic cells, macrophages, endothelial cells, and a significant number of tumor cells [10]. In the healthy immune system, the activation of the PD-1/PD-L1 pathway can inhibit the immune function of T lymphocytes and promote the inhibitory function of regulatory T cells, which can reduce the immune response of the body to normal peripheral tissues. Consequently, it can inhibit autoimmune responses, prevent the development of autoimmune diseases, and maintain autoimmune tolerance in healthy individuals [11]. When cancer occurs, the tumor cells will reduce their immunogenicity by expressing PD-L1. Hence, they will not be recognized by the immune system and will evade immune attack [12]. In a variety of tumors, the PD-L1 expression is usually associated with poor prognosis [13, 14].

Current theories on the expression of PD-L1 in colorectal cancer and tumor prognosis are limited and controversial. Some studies have manifested the palpable connection between PD-L1 expression and overall survival rate of colorectal cancer patients [15–18], but the others utter the contradictory statement [19, 20]. So, we used meta-analysis to analyze the prognostic value of programmed death factor ligand 1 (PD-L1), which will also lay a theoretical foundation for the application of PD-1/PD-L1 immunoassay point inhibitors in colorectal cancer.

Materials and methods

Bibliographic search

Two authors independently searched PubMed, Web of Science, Embase, and Cochrane Library for published literature on PD-L1 and colorectal cancer. Publication time of the included articles ranges from the time when the database was established until August 2018. All publications are in English. Search strategies are ("colorectal neoplasms" OR "colorectal cancer" OR "colorectal carcinoma" OR "colorectal cancers" OR "colonic neoplasms" OR "rectal neoplasms") AND ("PD-1" OR "PD-L1" OR "programmed death 1" OR "programmed death ligand 1" OR "programmed cell death ligand 1" OR "programmed death 1 ligand 1" OR "B7-H1" OR "CD274").

Inclusion criteria

The following are the inclusion criteria:

1. The clinical and pathological data of all cases are complete, and all were diagnosed as colorectal cancer by pathological examination;
2. Detecting the PD-L1 expression in colorectal cancer tissues by immunohistochemical staining;
3. The literature provides the relationship between PD-L1 expression and overall survival (OS) in patients with colorectal cancer;
4. The literature provides the relationship between PD-L1 expression and clinicopathological features, such as primary tumor size, clinical stage, and differentiation;
5. The literature provides sufficient information to estimate the hazard ratio (HR).

Exclusion criteria

The following are the exclusion criteria:

1. The included literature is not an original study;
2. The data contained in the research is wrong, or the quality of the incorporated literature is low;
3. The included literature is based on animal or cell experiments;
4. Cannot use the data provided in the literature to calculate the hazard ratio (HR) associated with PD-L1;
5. The included literature did not analyze the expression of PD-L1 in tumor cells.

Data extraction and quality evaluation

Two authors independently screened and extracted data found on inclusion and exclusion criteria and discussed together or adjudicated by third parties in case of disagreement. For the lack of information, we contacted the original author as much as possible. Extracted contents included author, publication year, country, positive threshold, follow-up period, baseline and clinicopathological information of patients, hazard ratio (HR), and 95% confidence interval (95%CI) related to PD-L1 expression.

Methodological quality assessment of the included data was carried out using the Newcastle-Ottawa Scale (NOS). Scores of the NOS are split into three aspects: object selection, inter-group comparability, and outcome measurement. The highest rating is 9 points, and the study with more than 6 points is considered as a high-quality one [21].

Statistical analysis

We implemented the meta-analysis based on the Preferred Reporting Items for Systematic Review and Meta-Analysis (PRISMA) Checklist (Additional file 1: Table S1). The hazard ratio (HR) and 95%CI of each study were combined to assess the relationship between the expression of PD-L1 and the prognosis, but the correlation between the PD-L1 expression and clinicopathological features was calculated using the pooled odds ratio (OR) and 95%CI. The heterogeneity test was performed using the $\chi^2$ test, and then we utilized the
fixed-effect model or the random-effect model according to heterogeneity. All the analyses above were presented by Revman 5.3 software.

All the studies are retrospective cohort studies, whose heterogeneity is often inevitable. Therefore, based on the analyses above, we carried out publication bias test and sensitivity analysis with the Stata 12.0 software to explore the sources of heterogeneity. And according to Begg’s or Egger’s test, \( P > 0.05 \) manifested that there was no publication bias in the study.

**Results**

**Data collection and characteristics**

A total of 1013 related articles were initially retrieved. After the layer-by-layer screening, 10 items were ultimately included, totaling 3481 cases (Fig. 1). The basic characteristics of the included studies were presented in Table 1. The NOS was used to estimate the quality of the included studies, and all were proved to be high-quality ones (Table 2).

**Relationship between PD-L1 expression and prognosis of colorectal cancer**

The relationship between the overexpression of PD-L1 and the poor prognosis of colorectal cancer patients was evaluated, and the consequence displayed a significant correlation (HR = 1.22, 95%CI = 1.01–1.48, \( P = 0.04 \), random effect) (Fig. 2).

**Relationship between PD-L1 expression and clinicopathological features of colorectal cancer**

Apropos of clinicopathological features, the merged odds ratio (OR) exhibited that highly expressed PD-L1 was firmly related to lymphatic invasion (OR = 3.49, 95%CI = 1.54–7.90, \( P = 0.003 \)) and advanced stage (OR = 1.77, 95%CI = 1.41–2.23, \( P < 0.00001 \)), but not correlative with patients’ gender, microsatellite instability, or tumor location (Table 3) (Additional file 2: Figure S1).

**Subgroup analysis of heterogeneity sources**

As for subgroup analysis of heterogeneity sources, the heterogeneity of each subgroup decreased in varying degrees (Additional file 3: Figure S2). Among them, the non-Asian group had the minimum heterogeneity (\( I^2 = 12\% < 50\% \)). In addition, the results of the Asian group (HR = 1.73, 95%CI = 1.10–2.73, \( P = 0.02 \), \( I^2 = 60\% \)), the non-Asian group (HR = 0.93, 95%CI = 0.89–0.97, \( P = 0.001 \), \( I^2 = 12\% \)), and the tumor stages I–IV group (HR = 1.32, 95%CI = 1.06–1.63, \( P = 0.01 \), \( I^2 = 52\% \)) were still statistically significant, but other subgroup analyses failed to arrive at such a statistically significant conclusion (Table 4).

**Publication bias analysis and sensitivity analysis**

The funnel plot is a conventional method to evaluate whether there is a “publication bias” in the meta-analysis, but as a qualitative judgment, its subjectivity makes different observers come to different conclusions [22]. Given
this, Begg’s test [23] and Egger’s test [24] were created to evaluate “publication bias” quantitatively. In this meta-analysis, according to Begg’s test (P = 0.428 > 0.05), there was no publication bias in the included literature involving PD-L1 and OS (Fig. 3). The detection results of publication bias in subgroup analyses are shown in Table 2 and Additional file 4: Figure S3. Sensitivity analysis pointed out that the conclusions were generally stable (Fig. 4 and Additional file 5: Figure S4).

**Discussion**

PD-1 and PD-L1 are inhibitory costimulatory molecules. When tumor cells express PD-L1 to combine with the PD-1 provided by tumor-infiltrating lymphocytes, the immune effect of T cells in the tumorous microenvironment is inhibited, which mediates the occurrence of tumor immune escape and promotes the progress of cancer [25]. At present, a lot of research has been done on this pair of inhibitory costimulatory molecules, but the regulatory mechanism of the signaling pathway in colorectal carcinoma has not been clarified, and theories in many fields are controversial. Although some systematic reviews focused on the prognostic value of PD-L1 in all types of solid tumors also mentioned the relationship between PD-L1 and prognosis of colorectal cancer in passing [26–28], yet their results in this regard had limitations because of the lack of in-depth research. Wu et al. suggested that PD-L1 overexpression was positively correlated with 5-year OS deterioration in colorectal cancer, but they only included two papers and used OR value to evaluate the results, meaning the existence of great bias [28]. Different from the above conclusion, Pyo et al. applied HR involving PD-L1 to assess the relevance between PD-L1 expression and prognosis of colorectal cancer and then concluded that there was no connection between them. However, they merely contained four retrospective studies, which lacked persuasiveness [27]. Xiang et al. also argued that PD-L1 could not be used as a prognostic indicator of colorectal cancer, but they misapplied risk ratio (RR), a specific measure for evaluating prospective studies, to the analysis of retrospective studies, so the findings should be treated cautiously [26]. Therefore, to resolve the controversy and deficiency mentioned above, this meta-analysis comprehensively collected relevant literature based on the inclusion criteria and adopted the hazard ratio (HR) associated with PD-L1 to estimate the
prognostic value of PD-L1 in colorectal cancer. In addition to that, we also explored the relationship between the expression of PD-L1 and the clinicopathological characteristics of colorectal cancer to make the outcome more convincing.

This meta-analysis demonstrated that PD-L1 expression could be utilized as an independent factor in judging the prognosis of colorectal cancer (HR = 1.22, 95%CI = 1.01–1.48, \( P = 0.04 \), random effect). Nevertheless, there was inevitable heterogeneity among the retrospective studies included in this meta-analysis (\( P = 0.0002 \), \( I^2 = 67% \)). In order to make the conclusion more persuasive and scientific, we adopted the subgroup analysis to explore the heterogeneity sources. And as the findings suggested, the heterogeneity of each subgroup decreased in varying degrees, indicating that these factors have certain degrees of influence. Among them, the non-Asian group had the minimum heterogeneity (\( I^2 = 12% < 50% \)), which implied the essential role the regional or ethnic differences played on engendering heterogeneity.

As the subgroup analysis of PD-L1 and clinicopathological features indicated, PD-L1 overexpression in colorectal cancer cells was associated with lymphatic invasion. Previous experimental studies have shown that they are indeed relevant. Epithelial-to-mesenchymal transition (EMT) leads to lymphatic invasion [29], and the expression of PD-L1 in tumor cells facilitates immunosuppression, both of which contribute to tumor progression and metastasis. MiR-200/Zinc finger E-box-binding homeobox 1 (ZEB1) was initially known as EMT regulatory axis, and the bidirectional negative feedback regulation mechanism between ZEB1 and miR-200 makes the corresponding cells actualize EMT [30]. But recently, the mechanism that miR-200/ZEB1 axis can also regulate PD-L1 to facilitate immunosuppression has been proved by experiments [31]. Therefore, the relationship between lymphatic invasion and PD-L1 overexpression can be considered to be mutually “parallel.” Also, consistent with another inference of this meta-analysis that the inhibition of the PD-1/PD-L1 signaling pathway in advanced colorectal cancer could achieve

| Study or Subgroup | Log(Hazard Ratio) | SE | Weight | Hazard Ratio N, Random, 95% CI | Hazard Ratio N, Random, 95% CI |
|-------------------|-------------------|----|--------|-------------------------------|-------------------------------|
| Hailong Zhu, 2015 | -0.3682           | 0.4672 | 3.4%   | 0.69 [0.28, 1.73]             |                              |
| Kyu Sang Lee, 2018 | 1.331            | 0.4906 | 3.2%   | 3.78 [1.45, 9.95]             |                              |
| Ming Liang, 2014  | 0.5539           | 0.1917 | 9.6%   | 1.74 [1.20, 2.53]             |                              |
| Raoul A. Droeser, 2014 | -0.0033         | 0.0222 | 15.7%  | 0.92 [0.89, 0.98]             |                              |
| Suzumori Saigusa, 2016 | 0.8969         | 0.4081 | 4.2%   | 2.45 [1.10, 5.48]             |                              |
| T.J. Miller, 2017  | 0.4937           | 0.31%  | 1.99   | 0.38 [2.63]                   |                              |
| Tsuyoshi Hamada, 2017 | 0.1823         | 0.1641 | 10.9%  | 1.20 [0.87, 1.68]             |                              |
| Tumenjin Enkhbat, 2018 | 1.354          | 0.6907 | 2.3%   | 3.87 [1.19, 11.57]            |                              |
| Yao Li, 2018       | 0.0469           | 0.2524 | 7.7%   | 1.95 [0.84, 2.72]             |                              |
| Yohpei Masugi-1, 2016 | 0.1169        | 0.1735 | 10.6%  | 1.12 [0.80, 1.59]             |                              |
| Yohpei Masugi-2, 2016 | -0.0202       | 0.1722 | 10.6%  | 0.98 [0.70, 1.37]             |                              |
| Yohpei Masugi-3, 2016 | 0.0441        | 0.1705 | 10.7%  | 1.04 [0.75, 1.48]             |                              |
| Yohpei Masugi-4, 2016 | 0.3149        | 0.2394 | 7.8%   | 1.37 [0.84, 2.23]             |                              |
| Total (95% CI)     | 0.100%           | 1.22 [1.01, 1.48] | 0.06    | 0.0002; \( \chi^2 = 36.90, df = 12; P = 0.0002 \); \( \phi = 67% \) | 0.04 |

*The article scored one point in the project
**The article scored two points in the project

As the subgroup analysis of PD-L1 and clinicopathological features indicated, PD-L1 overexpression in colorectal cancer cells was associated with lymphatic invasion. Previous experimental studies have shown that they are indeed relevant. Epithelial-to-mesenchymal transition (EMT) leads to lymphatic invasion [29], and the expression of PD-L1 in tumor cells facilitates immunosuppression, both of which contribute to tumor progression and metastasis. MiR-200/Zinc finger E-box-binding homeobox 1 (ZEB1) was initially known as EMT regulatory axis, and the bidirectional negative feedback regulation mechanism between ZEB1 and miR-200 makes the corresponding cells actualize EMT [30]. But recently, the mechanism that miR-200/ZEB1 axis can also regulate PD-L1 to facilitate immunosuppression has been proved by experiments [31]. Therefore, the relationship between lymphatic invasion and PD-L1 overexpression can be considered to be mutually “parallel.” Also, consistent with another inference of this meta-analysis that the inhibition of the PD-1/PD-L1 signaling pathway in advanced colorectal cancer could achieve
remarkable results, existing clinical trials have exhibited the high security and activity of the treatment with PD-1/PD-L1 immunocheckpoint inhibitors [32].

Limitations
To mention first, all the included articles were retrospective studies, whose bias could not be eliminated, so the consequences were generally stable. Also, it should be noted that all the articles included are in English, meaning the lack of research especially those negative studies in non-English speaking countries, which leads to the absence of representativeness and the production of bias.

Secondly, it has been reported that the expression of PD-L1 in tumor cells is a critical factor in making the monoclonal antibody against PD-1/PD-L1 effective. However, in every article, the threshold of PD-L1 positive is different and brings about a tremendous impact

Table 3 The relationship between PD-L1 expression and clinicopathological characteristics in subgroup analysis

| Subgroup analysis                        | No. of studies | No. of patients | Experimental group positive/total | Control group positive/total | OR       | 95%CI     | P value | Heterogeneity (I²), % | Begg’s test (P value) | Egger’s test (P value) |
|------------------------------------------|----------------|----------------|-----------------------------------|------------------------------|----------|-----------|---------|------------------------|------------------------|------------------------|
| Gender                                   | 9              | 3706           | Male 898/1852                     | Female 1018/1854             | 0.92     | 0.79–1.07 | 0.29    | 0                      | 0.917                  | 0.639                  |
| The situation of primary tumor           | 7              | 2809           | T3+T4 1018/2176                   | T1+T2 347/633                | 1.15     | 0.68–1.95 | 0.60    | 70                     | 0.764                  | 0.115                  |
| The involvement of regional lymph nodes | 6              | 2652           | N1+N2 558/1225                    | N0 729/1427                  | 1.32     | 0.84–2.09 | 0.23    | 74                     | 0.060                  | 0.012                  |
| Stage                                    | 6              | 2064           | III+IV 594/936                    | I+II 659/1128                | 1.77     | 1.41–2.23 | P < 0.0001 | 23     | 0.452                  | 0.512                  |
| Vascular invasion                        | 6              | 2052           | Positive 201/564                  | Negative 491/1488            | 1.12     | 0.72–1.75 | 0.62    | 63                     | 0.133                  | 0.090                  |
| Tumor location                           | 6              | 3133           | Left 916/1873                     | Right 711/1260               | 0.86     | 0.53–1.40 | 0.55    | 82                     | 0.452                  | 0.283                  |
| Microsatellite instability               | 4              | 2012           | MSI-H 223/362                     | MSI-L+MS 1012/1650           | 0.95     | 0.46–1.97 | 0.89    | 84                     | 1.000                  | 0.347                  |
| Lymphatic invasion                       | 4              | 723            | Positive 122/280                  | Negative 83/443              | 3.49     | 1.54–7.90 | 0.003   | 73                     | 1.000                  | 0.764                  |
| Tumor differentiation                    | 4              | 1862           | Poor 108/156                      | Well to moderate 1057/1706   | 1.90     | 0.55–6.63 | 0.31    | 88                     | 0.734                  | 0.292                  |
| Mucinous properties                      | 3              | 1563           | Mucinous 34/108                   | Other 573/1455               | 0.94     | 0.42–2.10 | 0.88    | 56                     | –                      | –                      |
| Grade                                    | 3              | 1562           | III 86/239                        | I+II 569/1323                | 0.66     | 0.42–1.03 | 0.07    | 52                     | –                      | –                      |

Table 4 Subgroup analysis of heterogeneity sources

| Subgroup analysis                        | No. of studies | HR       | 95%CI     | P value | Heterogeneity (I²), % |
|------------------------------------------|----------------|----------|-----------|---------|------------------------|
| Country                                  | Asian          | 6        | 1.73      | 1.10–2.73 | 0.02                  | 60           |
|                                          | Non-Asian      | 4        | 0.93      | 0.89–0.97 | 0.001                 | 12           |
| Stages                                   | I–IV           | 5        | 1.32      | 1.06–1.63 | 0.01                  | 52           |
| Follow-up period                         | ≥ 5 years      | 4        | 1.12      | 0.94–1.34 | 0.20                  | 65           |
|                                          | < 5 years      | 6        | 1.62      | 0.93–2.82 | 0.09                  | 61           |
| Postoperative adjuvant chemotherapy      |                | 4        | 1.61      | 0.88–2.94 | 0.12                  | 54           |
| Sample size                              | ≥ 200          | 5        | 1.09      | 0.92–1.28 | 0.32                  | 54           |
|                                          | < 200          | 5        | 1.61      | 0.99–2.60 | 0.05                  | 47           |
on the experimental results. According to published articles, patients with PD-L1 positive can obtain a better outcome in the treatment with immunocheckpoint inhibitors as the threshold increases [33]. So, uniformly applying the most suitable PD-L1 positive threshold to the following research should be a top priority. Besides that, using of different immunohistochemical antibodies in various studies leads to specific errors.

Furthermore, the latest experiments indicate that cancer patients [34] with intestinal flora disorders have a worse prognosis in the treatment with anti-PD-1/PD-L1 antibodies, which can be concluded that intestinal flora balance plays a vital role in the efficacy of PD-1/PD-L1 immunocheckpoint inhibitors. Therefore, the following study in this field should take intestinal flora into consideration.

**Conclusions**

This study analyzed all the available interrelated information in the published literature and exhibited that the expression of PD-L1 was significantly correlated with the overall survival rate of colorectal cancer. The more the PD-L1 was expressed, the worse prognosis the colorectal cancer patients would undergo. Concerning clinicopathological features, the expression of PD-L1 was bound up with lymphatic invasion and tumor stage, but not gender,

![Fig. 3 Begg's funnel plot for publication bias test including PD-L1 expression and prognosis in colorectal cancer](image1)

![Fig. 4 Sensitivity analysis including PD-L1 expression and prognosis in colorectal cancer](image2)
microsatellite instability, or tumor differentiation. In other words, the expression of PD-L1 could be utilized as an independent factor in judging the prognosis of colorectal cancer, and patients with advanced cancer or lymphatic invasion were more likely to express PD-L1. This conclusion may lay a theoretical foundation for the application of PD-1/PD-L1 immunoassay point inhibitors but still need to be verified by sizeable well-designed cohort studies.

Additional files

Additional file 1: Table S1. PRISMA 2009 Checklist. (DOC 33 kb)
Additional file 2: Figure S1. Forest plots assessing the relationship between PD-L1 and clinicopathological characteristics: (a) gender; (b) grade; (c) lymphatic invasion; (d) microsatellite instability; (e) mucinous properties; (f) stage; (g) the involvement of regional lymph nodes; (h) tumor differentiation; (i) tumor location; (j) the situation of primary tumor; (k) vascular invasion. (PNG 220 kb)
Additional file 3: Figure S2. Subgroup analysis of heterogeneity sources: (a) Asian; (b) non-Asian (c) stages I-IV; (d) follow-up more than 5 years; (e) follow-up less than 5 years (f) postoperative adjuvant chemotherapy; (g) sample size ≥ 200; (h) sample size < 200. (PNG 287 kb)
Additional file 4: Figure S3. Detection of publication bias in subgroup analysis: (a) gender; (b) lymphatic invasion; (c) microsatellite instability; (d) stage; (e) the involvement of regional lymph nodes; (f) the situation of primary tumor; (g) tumor location; (h) tumor differentiation; (i) vascular invasion. (PNG 99 kb)
Additional file 5: Figure S4. Sensitivity analysis of subgroup analysis: (a) gender; (b) lymphatic invasion; (c) microsatellite instability; (d) stage; (e) the involvement of regional lymph nodes; (f) the situation of primary tumor; (g) tumor location; (h) tumor differentiation; (i) vascular invasion. (PNG 147 kb)

Abbreviations
CE: Confidence interval; EMT: Epithelial-to-mesenchymal transition; HR: Hazard ratio; NOS: Newcastle-Ottawa Scale; OR: Odds ratio; OS: Overall survival; PD-1: Programmed cell death protein 1; PD-L1: Programmed death factor ligand 1; PRISMA: Preferred Reporting Items for Systematic Review and Meta-Analysis; ZEB1: Zinc finger E-box-binding homeobox 1

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Availability of data and materials
The datasets supporting the conclusions of this article are included within the article and its additional files.

Authors’ contributions
ZFS designed the research process. LHG and MMC searched the database for corresponding research. DYM and ZFS extracted the useful information from the articles above. ZFS and RJU used the statistical software for analysis. ZFS and LHG drafted the meta-analysis. The final draft came into being after the careful examination by all the authors. All authors read and approved the final manuscript.

Ethics approval and consent to participate
Not applicable

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Competing interests
The authors declare that they have no competing interests.

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