

# The expression status and prognostic significance of programmed cell death 1 ligand 1 in gastrointestinal tract cancer: a systematic review and meta-analysis

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**Background:** Programmed cell death 1 ligand 1 (PD-L1) expression has been observed in various malignancies. However, the association between PD-L1 expression and the survival of patients with gastrointestinal tract cancer remains controversial. Besides, the rate of PD-L1 positivity on tumor cells of digestive tract cancer is not clear. Thus, we performed a meta-analysis by incorporating all available evidence to evaluate the rate of PD-L1 positivity and the overall survival (OS) according to PD-L1 status in patients with gastrointestinal tract cancer.

**Methods:** Electronic databases were searched for eligible literature. Hazard ratios (HRs) for OS with 95% confidence intervals (CIs) according to the expression status of PD-L1 evaluated by immunohistochemistry were extracted. The outcomes were synthesized based on a random-effects model.

**Results:** Fifteen studies (only nine reported OS) that involved 2,993 gastrointestinal tract cancer patients stratified by PD-L1 status were eligible for inclusion in our study. We found the PD-L1-positive expression rate was 0.495 (95% CI 0.415–0.576) if 10% was taken as the cut-off value. When the H-score method was used to evaluate PD-L1 expression, it showed that the PD-L1 positive rate was 0.639 (95% CI 0.490–0.765) if the cut-off value was  $\leq 50$ , which was higher than when using  $> 50$  as the cut-off point (0.449, 95% CI 0.417–0.483). Additionally, PD-L1-positive gastrointestinal tract cancer patients were associated with significantly poorer OS when compared to negative ones (HR 1.61, 95% CI 1.10–2.35,  $P=0.014$ ). Subgroup analysis presented similar significant results in patients with esophageal cancer (HR 2.56, 95% CI 1.55–4.21,  $P<0.001$ ).

**Conclusion:** The positive expression rate of PD-L1 was nearly 50% no matter which method for immunohistochemistry evaluation we chose. Additionally, positive PD-L1 expression status in tumor cells is a risk factor for prognosis of gastrointestinal tract cancer, especially esophageal cancer.

**Keywords:** prognosis, esophageal cancer, immunohistochemistry, PD-L1-positive expression rate

## Introduction

Gastrointestinal tract cancer refers to malignant conditions of the gastrointestinal tract, including of the esophagus, stomach, small intestine, large intestine, rectum, and anus. Among those cancer patients, colorectal carcinoma, gastric cancer, and esophageal cancer are common types (the third, fourth, and sixth most-frequently diagnosed malignancies, respectively) globally.<sup>1–3</sup> Although recent advances in multidisciplinary therapies have improved treatment outcomes, the overall prognosis for gastrointestinal tract cancer remains poor. It is well known that the development

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and prognosis of malignant tumors are closely related to host immune functions.<sup>4</sup> Thus, novel therapeutic strategies, especially immunotherapy, are needed to be developed and established.<sup>4</sup>

It has been identified that immune escape plays an important role in tumor progression.<sup>5</sup> Improved understanding of the molecular mechanisms that govern the host response to tumors has led to the identification of checkpoint signaling pathways that limit the anticancer immune response.<sup>6</sup> A particularly important immune checkpoint that mediates tumor-induced immune suppression is the binding of programmed death 1 (PD-1) expressed on tumor-infiltrating lymphocytes and its ligand 1 (PD-L1) expressed on tumor cells.<sup>7</sup> PD-L1 has been reported to inhibit the proliferation of activated T-cells and induce the apoptosis of T-cells to form and maintain an immunosuppressive microenvironment since PD-L1 can recognize and bind the PD-1 on tumor-infiltrating lymphocytes.<sup>8</sup>

PD-L1 expression has been observed in various malignancies. Moreover, several meta-analyses have proved that PD-L1 overexpression indicates a poor prognosis for patients with non-small cell lung cancer.<sup>9–11</sup> However, the association between PD-L1 expression and the survival of patients with gastrointestinal tract cancer remains controversial. Besides, the rate of PD-L1 positivity on tumor cells of digestive tract cancer is not clear. Therefore, the expression status and prognostic significance of PD-L1 require further comprehensive study to clarify. Thus, we performed a meta-analysis by incorporating all available evidence to evaluate the expression rate of PD-L1 and the overall survival (OS) according to PD-L1 status in patients with gastrointestinal tract cancer.

## Materials and methods

### Literature search

Our institutions Ethics Committee has exempted our study from Institutional Review Board approval as our study involves exclusively preexisting anonymous data. The preferred reporting items for systematic reviews and meta-analyses (PRISMA) statement for reporting systematic reviews recommended by the Cochrane Collaboration was followed for conducting this meta-analysis. Two authors independently carried out a comprehensive systematic search for published articles or abstracts by searching through PubMed, Embase, Scopus, and the Cochrane Library from inception to April 2015. Searches were limited to human studies, using a combination of the terms “PD-L1”, “CD274”, “B7-H1”, “programmed cell death 1 ligand 1”, “gastrointestinal”, “esophageal”, “gastric”, “colorectal”, “outcome”, “survival”, and “prognosis”. We also manually reviewed relevant reference lists and reviews. There were no language restrictions.

### Inclusion and exclusion criteria

Studies meeting the following criteria were eligible for the single-arm meta-analysis of PD-L1-positive expression rate on tumor cells: (1) cohort studies which investigated PD-L1 expression level in gastrointestinal tract cancer patients; (2) the expression level of PD-L1 was tested by immunohistochemistry (IHC) staining in the tissue specimens; (3) samples of PD-L1 positive and PD-L1 negative were available. Moreover, studies in the meta-analysis that focused on patients' survival according to PD-L1 status met the following criteria: (1) studies meeting the criteria of meta-analysis for PD-L1-positive expression rate; (2) survival data stratified by PD-L1 status was available. Studies that failed to meet the inclusion criteria were excluded.

### Outcome measures, data extraction, and quality assessment

The primary outcome for this meta-analysis was PD-L1-positive expression rate and OS. OS data were extracted in the form of hazard ratios (HRs) with the corresponding 95% confidence interval (CI). If the HR was not displayed directly, it was estimated according to the methods described by a previously published article.<sup>12</sup> The data collection and assessment of methodological quality followed the quality of reporting of meta-analyses (QUORUM) and the Cochrane Collaboration guidelines (<http://www.cochrane.de>). The following main items were abstracted from the included studies: author, year, tumor type, IHC evaluation method and cut-off value, sample size of total patients and PD-L1-positive group, and survival data. Two reviewers used the modified Newcastle–Ottawa scale to assess these studies.<sup>13</sup> All investigators discussed and resolved all discrepancies in the extracted data. All of the eligible studies were of high quality.

### Statistical analysis

All calculations for PD-L1-positive expression rate on tumor cells were performed using Meta-Analyst Beta (v 3.13; Tufts Medical Center, Boston, MA, USA). HRs for OS with 95% CIs according to the expression status of PD-L1 was pooled using STATA (v 11.0; StataCorp, College Station, TX, USA). Heterogeneity across the incorporated studies was assessed with a forest plot and the inconsistency statistic ( $I^2$ ). A random-effects model was employed in case of potential heterogeneity and to avoid underestimation of standard errors of pooled estimates in this meta-analysis. Subgroup analysis was conducted according to IHC evaluation method and cut-off value and tumor type. An HR that was greater than 1 reflected shorter OS for PD-L1-positive patients. All CIs had two-sided probability coverage of 95%. A statistical test with a  $P$ -value less than 0.05 was considered as significant.

## Publication bias

An extensive search strategy was made to minimize the potential publication bias. Graphical funnel plots were generated to visually assess publication bias. The statistical method to detect funnel plot asymmetry was Begg's test.<sup>14</sup>

## Results

### Eligible studies

In total, 574 records were identified according to the search strategy. Finally, we enrolled 15 studies<sup>15–29</sup> that involved 2,993 gastrointestinal tract cancer patients (1,461/1,532 cases for PD-L1 positive/negative) with available PD-L1 expression data stratified by PD-L1 status. Nine of the studies<sup>16–19,21,23,26–28</sup> reported HRs for OS as a clinical outcome. Figure 1 summarizes the flow chart of study selection. Our study covered three major types of gastrointestinal tract cancer, esophageal cancer, gastric cancer, and colorectal carcinoma. The H-score was one of the most common methods used to evaluate the expression of PD-L1 in the tumor cells among the included articles, as well as the percentage of positively stained cells. The characteristics of the enrolled studies are shown in Table 1.

### Single-arm meta-analysis of positive expression rate of PD-L1

After stratifying the enrolled studies according to IHC evaluation method and cut-off value, we found the PD-L1-positive expression rate was 0.495 (95% CI 0.415–0.576) if 10% was taken as the cut-off value. When the H-score method

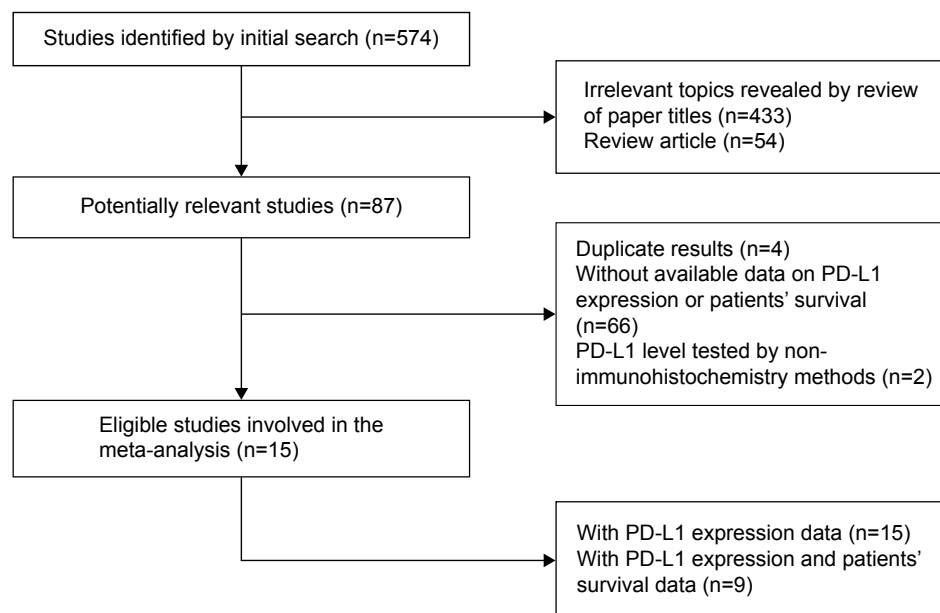
was used to evaluate PD-L1 expression, it showed that the PD-L1 positive rate was 0.639 (95% CI 0.490–0.765) if the cut-off value was  $<50$ , which was higher than when  $>50$  was used as the cut-off point (0.449, 95% CI 0.417–0.483) (Figure 2).

### Meta-analysis of PD-L1 positive versus PD-L1 negative in terms of OS

Overall, positive expression of PD-L1 in the tumor cells of gastrointestinal tract cancer patients was associated with a significantly poorer OS than negative expression (HR 1.61, 95% CI 1.10–2.35,  $P=0.014$ ) (Figure 3).

### Subgroup analyses, sensitivity analyses, and publication bias

The H-score evaluation method showed inferior survival in the PD-L1-positive group if we took  $<50$  as the cut-off value (HR 1.75, 95% CI 1.33–2.31,  $P<0.001$ ). The results were similar when using  $>50$  as the cut-off value (HR 1.79, 95% CI 0.69–4.62,  $P=0.231$ ). However, an only slightly numerical adverse prognostic effect of PD-L1-positive expression was presented if 10% was taken as the cut-off value using the percentage method (HR 1.05, 95% CI 0.42–2.61,  $P=0.919$ ) (Figure 4). Additionally, when stratifying studies according to tumor type, we observed that meta-analysis presented significant poorer OS in the PD-L1-positive group of patients with esophageal cancer (HR 2.56, 95% CI 1.55–4.21,  $P<0.001$ ). Similar numerical survival benefits were found in the PD-L1 negative



**Figure 1** Flow chart of study selection.

**Abbreviation:** PD-L1, programmed cell death 1 ligand 1.

Table 1 Characteristics of included studies for meta-analyses

Study, year	Tumor type	IHC evaluation <sup>a</sup>	Cut-off value for PD-L1 (+)	Patients, N	Age ≥60 years (%)	Male, n (%)	Country	PD-L1 (+) patients (%)	HR <sup>b</sup> for OS	Lower limit of 95% CI	Upper limit of 95% CI
Ohigashi et al <sup>15</sup> , 2005	EC	Percentage	≥10%	31	NA	NA	Japan	13 (41.94)	NA	NA	NA
Loos et al <sup>16</sup> , 2011	EC	H-score	≥66	101	NA	NA	Germany	37 (36.63)	2.92	1.5	5.66
Chen et al <sup>17</sup> , 2014	EC	H-score	>0	99	48 (48.48)	76 (76.77)	People's Republic of China	82 (82.83)	2.157	1.017	4.577
Wu et al <sup>18</sup> , 2006	GC	NA	NA	102	64 (62.75)	75 (73.53)	People's Republic of China	43 (42.16)	2.74	1.23	6.1
Geng et al <sup>19</sup> , 2014	GC	H-score	≥22	100	65 (65.00)	61 (61.00)	People's Republic of China	65 (65.00)	1.65	1.16	2.73
Hou et al <sup>20</sup> , 2014	GC	Percentage	≥10%	111	56 (50.45)*	75 (67.57)	People's Republic of China	70 (63.06)	NA	NA	NA
Kim et al <sup>21</sup> , 2014	GC	Percentage	≥10%	243	53 (21.81) <sup>#</sup>	152 (62.55)	Korea	106 (43.62)	0.651	0.416	1.017
Jiang et al <sup>22</sup> , 2014	GC	H-score	≥60	205	128 (62.44)	151 (73.66)	People's Republic of China	88 (42.93)	NA	NA	NA
Qing et al <sup>23</sup> , 2015	GC	Percentage	≥10%	107	62 (57.94)	72 (67.29)	People's Republic of China	54 (50.47)	1.649	1.173	2.318
Hua et al <sup>24</sup> , 2012	CRC	H-score	≥200	33	NA	NA	People's Republic of China	15 (45.45)	NA	NA	NA
Zhang et al <sup>25</sup> , 2012	CRC	Percentage	≥10%	57	36 (63.16)	35 (61.40)	People's Republic of China	26 (45.61)	NA	NA	NA
Droeser et al <sup>26</sup> , 2013	CRC	H-score	≥200	1,420	NA	673 (47.6)	Switzerland	669 (47.11)	0.85	0.81	0.89
Shi et al <sup>27</sup> , 2013	CRC	H-score	≥200	143	69 (48.25)	61 (42.66)	People's Republic of China	64 (44.76)	2.611	1.008	3.576
Liang et al <sup>28</sup> , 2014	CRC	H-score	≥20	185	114 (61.62)	87 (47.03)	People's Republic of China	102 (55.14)	1.74	1.195	2.713
Zhao et al <sup>29</sup> , 2014	CRC	H-score	≥20	56	30 (53.57)	35 (62.50)	People's Republic of China	27 (48.21)	NA	NA	NA

Notes: <sup>a</sup>H-score = SI (staining intensity) × PP (percentage of positive cells). SI was determined as: 0, negative; 1, weak; 2, moderate; and 3, strong. PP was defined as: 0, negative; 1–100, 1–100% positive cells. <sup>b</sup>Age ≥58 years; <sup>#</sup>age ≥70 years. <sup>c</sup>HR represents HR PD-L1 positive/PD-L1 negative.

Abbreviations: CI, confidence interval; CRC, colorectal cancer; EC, esophageal cancer; GC, gastric cancer; HR, hazard ratio; IHC, immunohistochemistry; NA, not available; OS, overall survival; PD-L1, programmed cell death ligand 1; PD-L1 (+), PD-L1 positive.

groups of patients with gastric cancer (HR 1.42, 95% CI 0.84–2.41,  $P=0.194$ ) and colorectal carcinoma (HR 1.49, 95% CI 0.74–3.00,  $P=0.260$ ) (Figure 4). Funnel plots with Begg's test are shown in Figure S1.

## Discussion

For patients with gastrointestinal tract malignancies, the association of the expression of PD-L1 and their prognosis remains unclear. In addition, the positive expression rate of PD-L1 in the tumor cells of digestive tract cancer is still unknown. A meta-analysis incorporating all available data from correlative studies is a reasonable method by which to address these questions. We conducted this study and found nearly 50% of gastrointestinal tract cancer patients showed positive expression of PD-L1. Furthermore, patients with positive expression of PD-L1 had significantly poorer survival than with those with negative expression.

It is notable that the PD-L1 expression rate was delineated less when stricter criteria for positive PD-L1 expression were applied. We could see that studies using a cut-off value >50 for H-score<sup>16,22,24,26,27</sup> showed a lower positive PD-L1 expression rate than those using <50.<sup>17,19,28,29</sup> A similar result was found if 10% was taken as the cut-off value when using the percentage method. Since IHC evaluation methods and cut-off values were not consistent for each gastrointestinal tract cancer type, subgroup meta-analysis was not possible when studies were stratified by tumor type. We could also see that studies using a cut-off value >50 for H-score<sup>16,26,27</sup> showed a greater difference in OS between PD-L1 positive and negative groups than those using <50,<sup>17,19,28</sup> although no significant survival benefits were obtained in the former due to limited sample sizes. Moreover, the unfavorable prognostic value of PD-L1 was significantly seen in patients with esophageal cancer.<sup>16,17</sup> Despite some trends observed, we currently cannot draw a valid conclusion that PD-L1 status is a predictor of prognosis for patients with gastric cancer or colorectal carcinoma. Considering the consistent trend of the prognostic significance of PD-L1 when using different kinds of anti-PD-L1 antibodies to stain tumor cells has been proven in another meta-analysis,<sup>30</sup> subgroup analyses regarding source and type of antibody are not necessary.

## Limitations and future directions

Efforts were made to conduct a comprehensive analysis to answer the question of expression rate of PD-L1 and the impact of PD-L1 status on patient prognosis in gastrointestinal tract cancer, but some limitations still should be acknowledged. First of all, we calculated or estimated

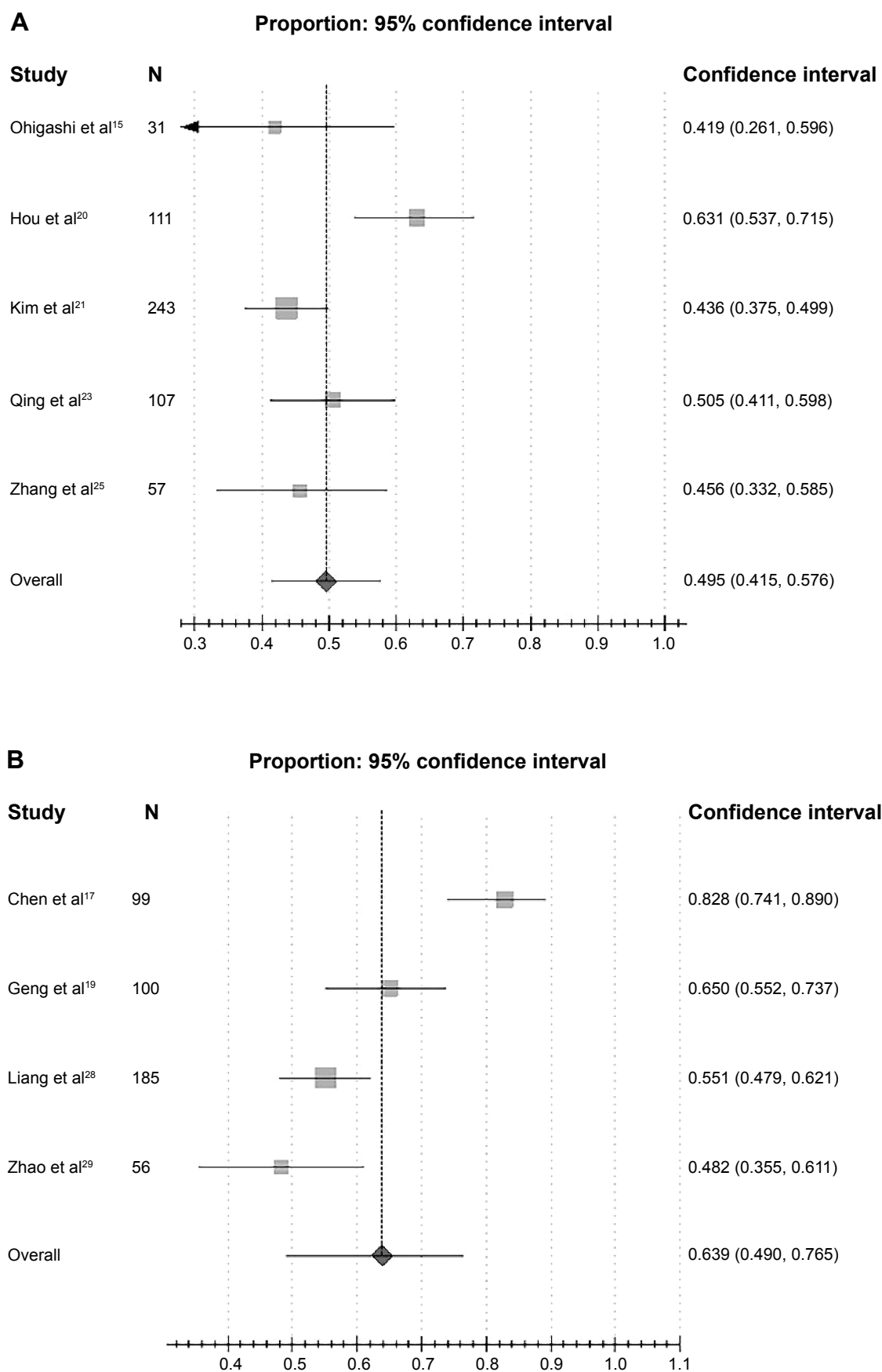
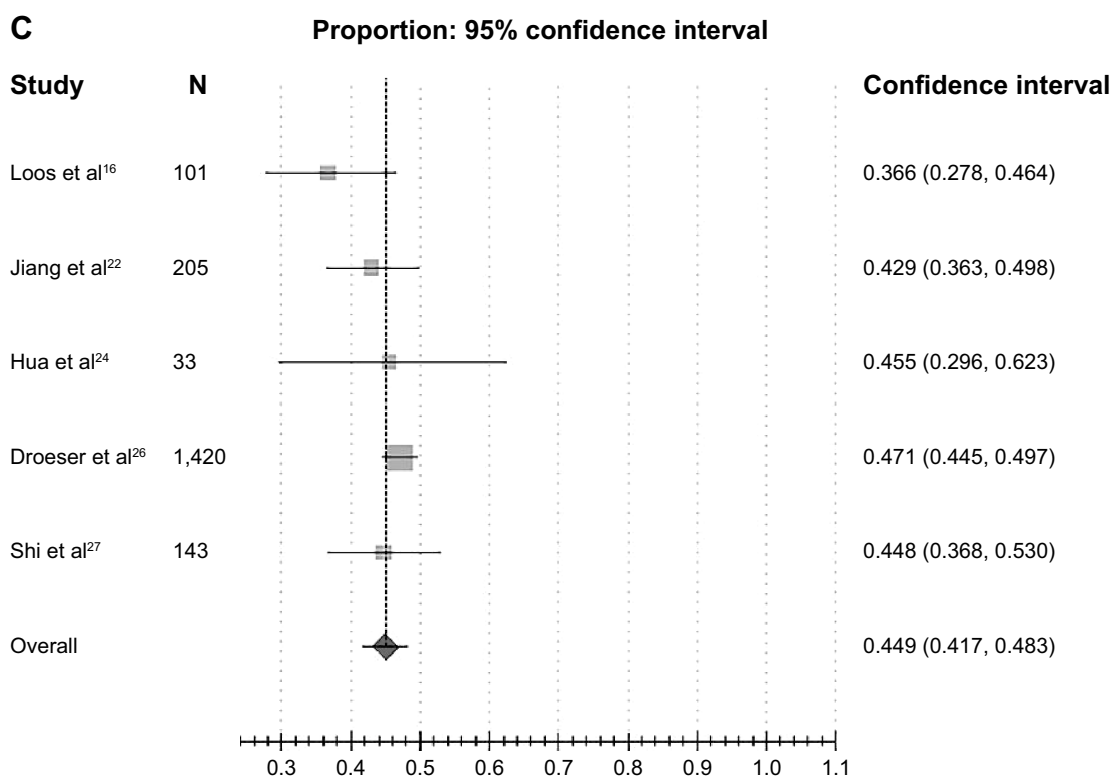
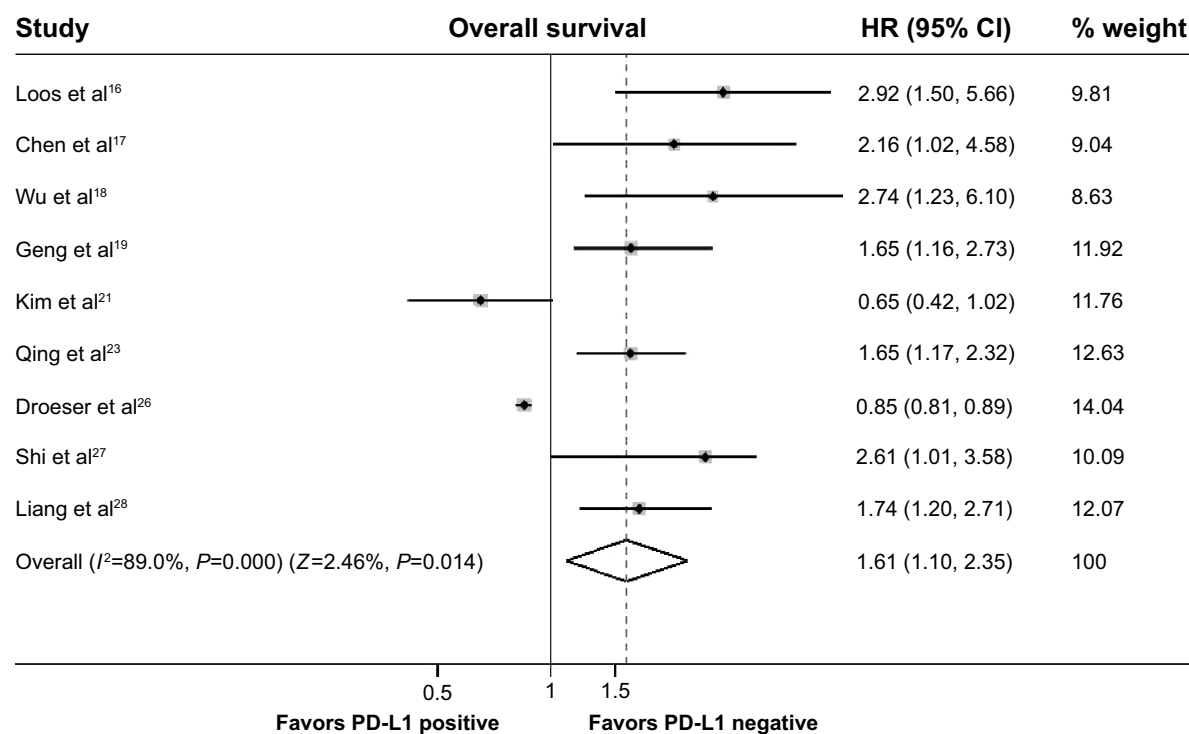


Figure 2 (Continued)



**Figure 2** Single-arm meta-analysis of positive expression rate of programmed cell death I ligand I (PD-LI) on tumor cells in gastrointestinal tract cancer according to immunohistochemistry evaluation method and cut-off value.

**Notes:** (A) Percentage (cut-off value 10%); (B) H-score (cut-off value <50); (C) H-score (cut-off value >50).

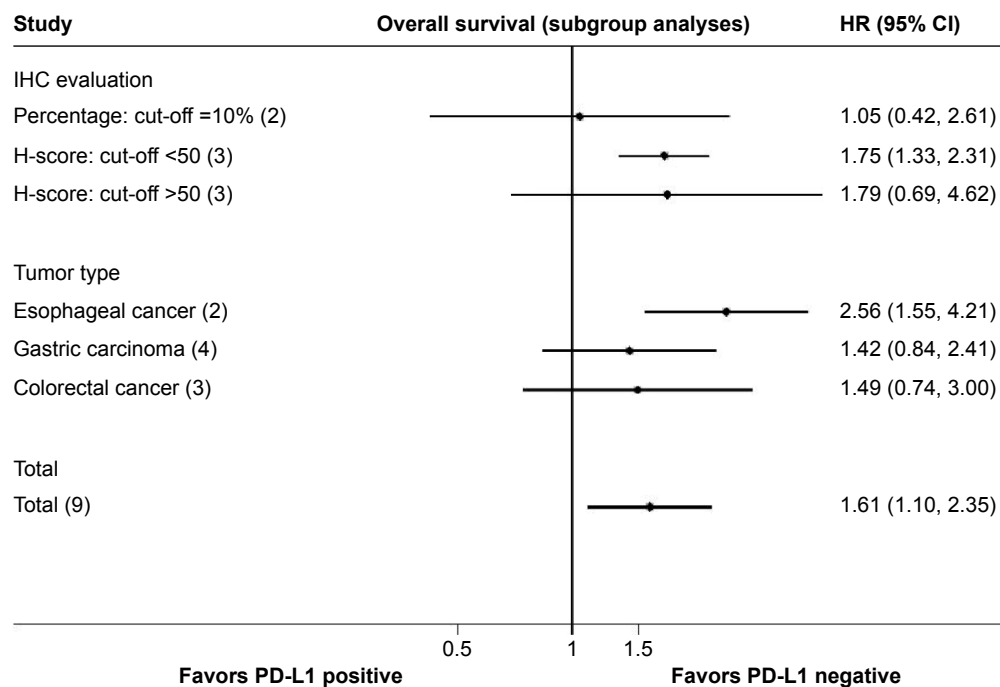


**Figure 3** Total meta-analysis of programmed cell death I ligand I (PD-LI) positive versus PD-LI negative on tumor cells of gastrointestinal tract cancer patients in terms of overall survival.

**Note:** Weights are from random-effects analysis.

**Abbreviations:** CI, confidence interval; HR, hazard ratio.





**Figure 4** Subgroup analysis of programmed cell death 1 ligand 1 (PD-L1) positive versus PD-L1 negative on tumor cells of gastrointestinal tract cancer patients in terms of overall survival.

**Note:** Weights are from random-effects analysis.

**Abbreviations:** CI, confidence interval; HR, hazard ratio; IHC, immunohistochemistry.

some survival data of the HRs from the available data or Kaplan–Meier curves, which may have compromised the precision of the data. Secondly, cut-off values distinguishing positive and negative expression of PD-L1 determined by IHC evaluation and the primary antibodies varied in different types of tumors, which might cause heterogeneity of the overall results. The subgroup results should have addressed some concerns. Thirdly, we were not able to evaluate the expression rate of PD-L1 stratified by tumor type in subgroup analysis due to inconsistent IHC evaluations. Fourthly, we were unable to determine whether a correlation exists between positive PD-L1 expression and the clinical characteristics of gastrointestinal tract cancer. In addition, most of the eligible studies failed to provide data regarding progression-free survival or recurrence-free survival, so we only extracted OS data in our meta-analysis. Besides, scientists might prefer to only report the positive results of a prognostic biomarker, leading to the existence of potential publication bias. Further studies are warranted to address the above issues.

Regardless of the described limitations, this comprehensive analysis statistically confirmed almost half of all gastrointestinal tract cancer patients positively expressed PD-L1 in tumor cells and those PD-L1 positive patients were associated with significantly shorter OS, especially

in esophageal cancer. A recent study has confirmed that positive PD-L1 expression is correlated with the improved efficacy of pembrolizumab, a drug marketed by Merck that targets the PD-1 receptor, in patients with advanced non-small-cell lung cancer.<sup>31</sup> These results hint that, in clinical trials using anti-PD-L1 or anti-PD-1 antibodies as cancer immunotherapy, enrollment might also be preferentially carried out on patients with gastrointestinal tract cancer, especially esophageal cancer. Furthermore, more efforts should be made to investigate which cut-off value is the best for differentiating real PD-L1-positive patients with gastrointestinal tract cancer who would get survival benefit from PD-1/PD-L1 blockers.

## Conclusion

In this meta-analysis, we found the positive expression rate of PD-L1 was nearly 50%, no matter which method for IHC evaluation we chose. Additionally, we found positive PD-L1 expression status in tumor cells is a risk factor for prognosis in gastrointestinal tract cancer, especially in esophageal cancer.

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## Disclosure

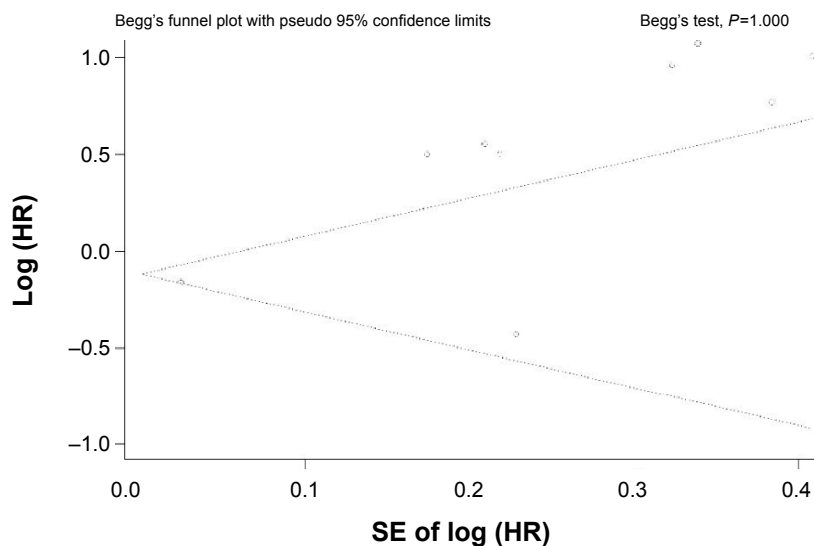
The authors declare no conflicts of interest in this work.

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## Supplementary material



**Figure S1** Graphical funnel plots with Begg's test of the meta-analysis.

**Abbreviations:** HR, hazard ratio; SE, standard error.

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