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The relationship between PD-L1 expression and clinicopathological characteristics and prognosis of Vietnamese gastric cancer patients

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ABSTRACT

Background & objectives: Programmed Death Ligand 1 (PD-L1) is a putative biomarker response to an immune checkpoint blockade that is related to poor outcomes as well as treatment strategy of numerous carcinomas, including gastric cancer. However, there is still a lack of other biomarkers that can predict patient prognosis in clinical settings. For this reason, we investigated PD-L1 expression and Combined Positive Score (CPS), as well its association with histopathological characteristics in gastric cancer patients. **Methods**: We collected clinicopathological data from 54 patients with gastric cancer who underwent surgery at 103 Military Medical Hospital, Vietnam, from December 2018 to November 2020. Immunohistochemistry staining of the tumor specimens for PD-L1 expression and CPS were evaluated and accessed relationship to patient characteristics and overall survival. **Results**: PD-L1 with strong and moderate expression was 1.9% and 20.3%, and CPS \geq 1 was 46.3%. PD-L1 expression and CPS had no statistical relationship with histopathological characteristics, with the exception that tumor location, which was significantly related to PD-L1 expression (p = 0.001). PD-L1 positivity and CPS \geq 1 were related to worse overall survival of gastric cancer patients. **Conclusions**: Our data indicated that PD-L1 and CPS are independent prognostic markers and indicators for use in targeted therapy for gastric cancer patients.

Key words: combined positive score, gastric cancer, histopathological characteristics, PD-L1 expression

INTRODUCTION

Gastric cancer (GC) is the fifth most common cancer globally¹. Most cases have been reported in Eastern Asia, particularly in China, Japan, and South Korea². According to the 2020 GLOBOCAN, gastric cancer was ranked as the fourth most common cancer in Vietnam after liver, lung, and breast cancers, with an average number of new cases of 17,906, with the 5-year prevalence rate at 24.64 per 100,000³. The causes of gastric cancer are still unclear, but some factors have been shown to increase risk. The most common risk factor is the microbe Helicobacter pylori, which contributes to cancer development⁴. Diet and lifestyle risk factors include obesity, a diet high in sodium and low in vegetables, smoked foods, smoking, and alcohol consumption. The cancer can also develop from some stomach diseases, such as gastroesophageal reflux, chronic gastritis, and stomach polyps. Hereditary factors also increase risk of contracting gastric cancer⁵.

Cancer immunotherapy has shown promising results in recent years. In some types of cancer, interaction between programmed death-ligand 1 (PD-L1) in cancer cells and programmed death-1 (PD-1) in T cells inhibits T cell function, thus helping to prevent the cancer cells from evading the immune system⁶. Currently, there are several approaches to locking interaction between PD-L1 and PD-1, such as the use of gene therapy or antagonistic monoclonal antibodies. Previously, a study by Su et al. used gene therapy to reprogram T cells by eliminating the PD-1 gene using the CRISPR-Cas9 technique; the results showed a significant reduction of PD-1 expression but did not affect the viability of T cells, suggesting a new strategy for cancer treatment⁷. Alternatively, the use of monoclonal antibodies to block immune checkpoint molecules has been shown to be a promising therapeutic strategy against several types of cancer^{8,9} and the immune regulatory PD-1/PD-L1 axis has been used for immunotherapy for gastric cancer¹⁰. The National Comprehensive Cancer Network (NCCN) guidelines suggest that immunotherapeutic strategies be applied to patients with advanced-stage GC¹¹. Some clinical trials have indicated that monoclonal

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antibodies targeting PD-L1 or its receptor PD-1 inhibit the PD-1/PD-L1 signaling pathway and enhance T-cell function, which improves outcomes in patients with GC, particularly in the advanced stage¹²⁻¹⁴. In a trial from CheckMate 032 (GC/GEJC cohort), there was a strong association between the combined assessment of PD-L1 expression with the CPS and the effectiveness of anti-PD-1 therapy¹⁵.

Beyond assessment PD-L1 expression and CPS, there are currently no biomarkers for evaluating the clinical efficacy of anti-PD-1/PD-L1 immunotherapy. In Vietnam, the evaluation of PD-L1 expression and CPS in gastric cancer patients is limited, thus constraining the access to anti-PD-1/PD-L1 therapy. Additionally, determining the association between PD-L1 expression and pathological features is essential but still controversial and not well-elucidated. Therefore, we investigate the relationship between PD-L1 expression with clinicopathological characteristics and the survival of Vietnamese patients with gastric cancer.

METHODS

Patient cohort

We retrospectively enrolled 54 GC patients who had undergone curative gastrectomy at 103 Military Medical Hospital, Vietnam, between December 2018 and November 2020. All patients were diagnosed using pathological results of H&E staining specimens to identify gastric cancer and also underwent gastrectomy and scraping of regional nodes. These patients were not treated with chemoradiotherapy beforehand and were monitored after surgery. We selected patients with recurrent GC, metastatic cancer to the stomach, or a combination of these and other cancers. The Ethics Committee of 103 Military Medical Hospital approved all study procedures (code: 140/2016/IRB-MH103).

Patient characteristics

We collected data on patient features including tumor location, tumor size, gross appearance (based on the Bormann classification), histologic type (based on the Lauren classification), histopathology classification (based on the WHO classification of Tumors of the Digestive System, 2010¹⁶), tumor differentiation, depth of invasion (T stage according to the 8th edition of the UICC/AJCC TNM classification¹⁷), lymph node status, and vascular invasion.

PD-L1 Immunohistochemistry

Tumor specimens in a paraffin-embedded block were cut into 3 μ m thick sections. We carried out immunohistochemical (IHC) staining for PD-L1 (clone

73-10, rabbit anti-human monoclonal antibody, 10 mg/ml, Leica, UK) on an automated slide stainer (Leica ST5010 Auto Stainer XL, Leica Biosystems), following manufacturer instructions. The samples were deparaffinized in xylene and rehydrated using a series of ethanol. Sections were incubated with the primary antibody for 30 minutes, then the secondary antibody of biotin-labeled anti-rabbit IgG was applied for 8 minutes. After washing in PBS, signals were visualized by incubation with polymer solution for 8 minutes and then 3-3'-diaminobenzidine (DAB) for 30 seconds. Slides were counterstained with hematoxylin for 6 minutes before mounting. After each step, wash through Bond-was solution two times in 2 minutes. Positive and negative controls were used for IHC reactions. Tonsils were used as the positive control, as PD-L1 should show strong staining in the crypt epithelium and weak to moderate staining of the follicular macrophages in the germinal centers.

Evaluation of PD-L1 expression and CPS scoring

We evaluated the expression of PD-L1 and calculated CPS following the protocol of PD-L1 IHC 22C3 pharmDx, which is FDA-approved for *in vitro* diagnostic use¹⁸. We randomly examined four fields on the tumor site slide of each specimen at 400X magnification. The expression of PD-L1 was recorded according to the percentage of tumor-stained cells per total viable tumor cells. We defined less than 1%, 1 — 49%, and \geq 50% as weak, moderate, and strong expressions, respectively. For calculating CPS, we used the formula as follows:

 $CPS = \frac{PD-L1 \ staining \ cells^*}{Total \ viable \ tumor \ cells} X \ 100$ *(tumor cells, lymphocytes, macrophages)

Based on CPS, we divided samples into two groups: CPS < 1 and CPS \geq 1.

Statistical analysis

After collecting sufficient information, the data for categorical variables was shown by frequency and percentage. We used a chi-square test to analyze correlations between PD-L1 expression, CPS, and clinicopathological features. We used the Kaplan-Meier method to construct the survival curves and a logrank test for survival comparison. Statistical analysis was performed using SPSS software ver. 20.0 (IBM, Armonk, NY, USA). A p-value < 0.05 was considered statistically significant.

| Characteristics | | (%) |
|-----------------------------|----------------------------|-----------|
| Tumor location | Body | 3 (5.5) |
| | Lesser curvature | 17 (31.5) |
| | Pylorus | 34 (63) |
| Tumor size | < 5 cm | 43 (79.6) |
| | \geq 5 cm | 11 (20.4) |
| Bormann classification | Polyp | 3 (5.6) |
| | Fungating | 6 (11.1) |
| | Ulcerous-infiltrative | 37 (68.5) |
| | Diffuse-infiltrative | 8 (14.8) |
| Lauren classification | Diffuse | 33 (61.1) |
| | Intestinal | 8 (14.8) |
| | Mixed | 13 (24.1) |
| WHO classification | Tubular adenocarcinoma | 40 (74.1) |
| | Mucinous adenocarcinoma | 3 (5.6) |
| | Signet ring cell carcinoma | 3 (5.6) |
| | Undifferentiated carcinoma | 8 (14.8) |
| Tumor differentiation | High | 5 (9.3) |
| | Mediated | 22 (40.7) |
| | Poor | 17 (31.5) |
| | Undifferentiated | 10 (18.5) |
| Depth of invasion (T stage) | T1 | 2 (3.7) |
| | T2 | 11 (20.4) |
| | Т3 | 22 (40.7) |
| | T4 | 19 (35.2) |
| Lymph node metastasis | Positive | 27 (50) |
| | Negative | 27 (50) |
| Vascular invasion | Positive | 3 (5.6) |
| | Negative | 51 (94.4) |

Table 1: Clinicopathological features of gastric cancer

 $\mathbf{n}:$ number; **WHO**: World Health Organization

Table 2: PD-L1 expression and CPS calculation

| Parameters | | (%) |
|------------------|----------|-----------|
| PD-L1 expression | Weak | 42 (77.8) |
| | Moderate | 11 (20.3) |
| | Strong | 1 (1.9) |
| CPS | < 1 | 29 (53.7) |
| | ≥ 1 | 25 (46.3) |

CPS: combined positive score; **PD-L1**: programmed death ligand 1

Table 3: Correlation of PDL1 expression with clinicopathological charecteristics

| able 3: Correlation of PDLT expression with clinicopathological charecteristics | | | | | | | | |
|---|----------------------|---------------------|----------|-------|----------------------|----------------------|-------|--|
| Characteristics | | PD-L1, n | | р | CPS, n | | р | |
| | X.X 1 | (%) | 0. | | (%) | | | |
| | Weak | Moderate | Strong | | < 1 | <u>≥ 1</u> | | |
| | | | | | | | | |
| Tumour location | | | | | | | | |
| Cardia | 2 (66.7) | 0 (0.0) | 1 (33.3) | | 2 (66.7) | 1 (33.3) | 0.442 | |
| Lesser curvature | 13 (76.5) | 4 (23.5) | 0 (0.0) | 0.001 | 7 (41.2) | 10 (58.8) | | |
| Polyrus | 27 (79.4) | 7 (20.6) | 0 (0.0) | | 20 (58.8) | 14 (41.2) | | |
| Tumour size | | | | | | | | |
| < 5cm | 33 (76.7) | 9 (20.9) | 1 (2.3) | 0.853 | 23 (53.5) | 20 (46.5) | 0.950 | |
| \geq 5cm | 9 (81.8) | 2 (18.2) | 0 (0.0) | | 6 (54.5) | 5 (45.5) | | |
| Bormann classification | | | | | | | | |
| Polyp type | 2 (66.7) | 1 (33.3) | 0 (0.0) | 0.473 | 2 (66.7) | 1 (33.3) | 0.178 | |
| Fungating | 6 (100.0) | 0 (0.0) | 0 (0.0) | | 2 (33.3) | 4 (66.7) | | |
| Ulcerous-infiltrative | 26 (70.3) | 10 (27.0) | 1 (2.7) | | 23 (62.2) | 14 (37.8) | | |
| Diffuse-infultrative | 8 (100.0) | 0 (0.0) | 0 (0.0) | | 2 (25.0) | 6 (75.0) | | |
| Lauren classification | | | | | | | | |
| Diffuse | 26 (78.8) | 6 (18.2) | 1 (3.0) | 0.759 | 18 (54.5) | 15 (45.5) | 0.950 | |
| Intestinal | 7 (87.5) | 1 (12.5) | 0 (0.0) | | 4 (50.0) | 4 (50.0) | | |
| Mixed | 9 (69.2) | 4 (30.8) | 0 (0.0) | | 6 (50.0) | 6 (50.0) | | |
| WHO classification | | | | | | | | |
| Tubular adenocarcinoma | 30 (75.0) | 9 (22.5) | 1 (2.5) | 0.903 | 20 (50.0) | 20 (50.0) | 0.383 | |
| Mucinous denocarci- | 3 (100.0) | 0 (0.0) | 0 (0.0) | | 3 (100.0) | 0 (0.0) | | |
| noma | | | | | | | | |
| Signet ring cell carcinoma | 3 (100.0) | 0 (0.0) | 0 (0.0) | | 2 (66.7) | 1 (33.3) | | |
| Undifferentiated arci- | 6 (75.0) | 2 (25.0) | 0 (0.0) | | 4 (50.0) | 4 (50.0) | | |
| noma | | · · · · | · · / | | · · · · | | | |
| Tumour differentiation | | | | | | | | |
| High | 5 (100.0) | 0 (0.0) | 0 (0.0) | 0.586 | 3 (60.0) | 2 (40.0) | 0.986 | |
| Mediated | 18 (81.8) | 4 (18.2) | 0 (0.0) | | 12 (54.5) | 10 (45.5) | | |
| Poor | 11 (64.7) | 5 (29.4) | 1 (5.9) | | 9 (52.9) | 8 (47.1) | | |
| Undifferentiated | 8 (80.0) | 2(20.0) | 0(0.0) | | 5(50.0) | 5(50.0) | | |
| Depth of invasion (T | 0 (0010) | 2 (2010) | 0 (010) | | 0 (0010) | 0 (0010) | | |
| stage) | | | | | | | | |
| T1 | 2(100.0) | 0 (0 0) | 0(00) | 0.873 | 1(500) | 1(500) | 0.902 | |
| T2 | 9 (81 8) | 2(18.2) | 0(0.0) | 0.075 | 6(545) | 5(45.5) | 0.902 | |
| T2 T3 | 17(773) | $\frac{2}{4}(18.2)$ | 1(45) | | 13(591) | 9 (40.9) | | |
| TJ | 1/(73.7) | $\frac{1}{5}(26.3)$ | 1(4.3) | | 9(474) | 10(52.6) | | |
| Lymph node metectasis | 14(75.7) | 5 (20.5) | 0 (0.0) | |) (47.4) | 10 (32.0) | | |
| Dositive | 22 (81 5) | A(14.8) | 1(37) | 0.394 | 13(491) | 14(501) | 0.412 | |
| Nagativa | 22(01.3) 20(74.1) | 7(250) | 1(3.7) | 0.364 | 15(40.1) 16(50.2) | 14(39.1) 11(40.7) | 0.413 | |
| Vacaular invesion | 20 (74.1) | 7 (23.9) | 0 (0.0) | | 10 (39.3) | 11 (40.7) | | |
| Desitive | 1 (22.2) | 2(66.7) | 0(0,0) | 0.122 | 1 (22.2) | 2(66.7) | 0.467 | |
| Nagativa | 1(33.3) | 2(00.7) | 1(2.0) | 0.122 | 1(33.3) | 2(00.7) | 0.40/ | |
| negative | 41 (80.4) | 9(1/.0) | 1(2.0) | | 28 (34.9) | 23 (45.1) | | |

CPS: combined positive score; n: number; PD-L1: programmed death ligand 1; WHO: World Health Organization.

RESULTS

Patients' clinicopathologic data and relationship with PD-L1 expression

We summarize the pathological characteristics of patients in Table 1. Results demonstrated that tumor location was the most common site in pylorus with 63% prevalence, followed by curvature at 31.5% and 5% in the body. Tumor size < 5 cm accounted for 79.6% of patients, whereas tumor size of \geq 5 cm accounted for 20.4%. Borman classification was ulcerous-infiltrative for 68.5% of patients, diffuseinfiltrative for 14.8%, fungating for 11.1%, and polypoid for 5.6%. Lauren classification was diffuse for 61%, intestinal for 14.8%, and mixed for 24.1%. The WHO classification was tubular adenocarcinoma for 74.1%, mucinous adenocarcinoma, signet-ring cell carcinoma for 5.6%, and undifferentiated carcinoma for 14.8%. The tumor differentiation was high at 9.3%, moderate at 40.7%, low at 31.5%, and undifferentiated at 18.5%. Depth of invasion was T1 at 3.7%, T2 at 20.4%, T3 at 40.7%, and T4 at 35.2%. Lymph node metastasis was 50% for both positive and negative. Vascular invasion was 5.6% for positive and 94.4% for negative.

IHC results for PD-L1 expression are shown in **Figure 1** and summarized in **Table 2** with rates of weak expression at 77.8%, moderate expression at 20.3%, and strong expression at 1.9%. The corresponding rankings of CPS < 1 accounted for 53.7%, while CPS \geq 1 accounted for 46.3% of patients.

We summarize the relationship between PD-L1 expression and CPS and pathological characteristics of GC patients in **Table 3**. The results show no statistically significant relationship between PD-L1 expression and CPS and tumor size, Bormann classification, Lauren classification, WHO classification, tumor differentiation, depth of invasion, lymph node metastasis, or vascular invasion. Tumor location was the exception, being significantly related to PD-L1 expression (p = 0.001, chi-square test).

Impact of PD-L1 expression and CPS on the survival of patients with gastric cancer

We evaluated PD-L1 expression and CPS with regard to overall survival in gastric cancer patients. This analysis defined samples with PD-L1 strong and moderate expression as positive and samples with weak expression as negative. Based on PD-L1 expression and CPS, overall survival is represented by the Kaplan– Meier curves in Figure 2. PD-L1-positive patients had lower overall survival than PD-L1-negative patients; the difference was statistically significant, with p = 0.001 (**Figure 2A**). Patients with CPS ≥ 1 had lower overall survival than patients with CPS < 1; the difference was statistically significant, with p = 0.047(**Figure 2 B**). The results showed a relationship between the expression of PD-L1 and CPS and the prognosis of gastric cancer patients.

DISCUSSION

In this study, we divided PD-L1 expression into weak, moderate, and strong rankings, with ratios of 77.8%, 20.3%, and 1.9%, respectively. For comparison with other studies, we defined samples with PD-L1 strong and moderate expression as positive, with a combined percentage of 22.2% presence in patients (Table 2). Our results are similar results found in studies by Tatsuro Tamura et al. (29.6%)¹⁴, Kawazoe et al. (24.8%)¹⁹, and Boger *et al.* (23.73%)²⁰. Our figure is, however, higher than that shown in some studies, such as Kang et al. (15.38%)²¹ and Dai et al. (14.32%)²². Conversely, PD-L1 positivity was shown to be much higher in some other studies, such as those by Lin Zhang et al. (50.8%)²³ or KEYNOTE-059 (57.1%)²⁴. Despite many studies on PD-L1 expression, only a few authors accounted for both PD-L1 positivity and CPS in patients with GC. Some recent reports have suggested immunotherapy for patients with advanced gastric cancer and CPS $\geq 1^{25,26}$. CPS measures PD-L1 staining in both tumor and immune cells. In immune cells, PD-L1 expression is essential for inducing immune tolerance²⁷. Therefore, instead of only assessing PD-L1 positivity in tumor cells, CPS calculation also measures the colorization of immune function cells, such as lymphocytes and macrophages. Thus, the CPS ≥ 1 ratio was higher than the PD-L1 positivity ratio in our study. Patients with CPS \geq 1 in our study accounted for 46.3% of participants (Table 2), which is lower than the results found by Shitara et al. (67%)²⁶ or Yamashita et al. (71.7%)²⁸. The percentages of PD-L1 positivity and CPS > 1 in our study differ from other studies and may depend on factors like patient cohort, genotype differences, histopathological characteristics, immunohistochemical staining methods used, or cut-off values for distinguishing between positive and negative PD-L1 expression.

Our results showed no relationship between the expression levels of PD-L1, CPS, and histopathological characteristics in gastric cancer patients, except regarding tumor location, which was significantly related to the expression level of PD-L1 (p = 0.001, chi-square test, **Table 3**). Results from other studies suggest that PD-L1 overexpression is associated with



Figure 1: Immunohistochemical staining for PD-L1 expression in gastric cancer tissue. A: PD-L1 negative. **C-D**: The expression of PD-L1 in gastric cancer according to the intensity weak (**B**), moderate (**C**), and strong (at x400 magnification).

depth of invasion (T stage)²⁰, lymph node metastasis^{29,30}, and vascular invasion³¹. In contrast, there has not been any demonstrated association of PD-L1 expression with tumor location³², tumor size³³, tumor differentiation³⁴, Lauren classification²², or invasion the lymphatic system²¹. This suggests that the relationship between PD-L1 expression and pathological characteristics varies widely between studies. This study also showed that PD-L1 with weak expression had significantly more prolonged survival than PD-L1 with moderate and strong expression (p = 0.001, Figure 2 A). Other studies have shown similar results; patients with PD-L1 overexpression have been shown to have a shorter survival time and a worse prognosis than patients with weak or negative PD-L1 expression³⁵⁻³⁷. Our results show that patients

with CPS < 1 had significantly greater prolonged survival than patients with CPS \geq 1 (p = 0.047, Figure 2 B). This was similar to the results from Yamashita's study²⁸, which also showed that these patients had a significantly shorter overall survival than those with CPS < 1 (p = 0.017). CPS has better prognostic significance than TPS (Tumor Positive Score), suggesting that, in addition to PD-L1-stained cells, the presence of lymphocytes surrounding the tumors and in the stroma have a significant effect on the overall survival of the patient. Therefore, CPS is an essential indicator for immunotherapy. Schoemig-Markiefka et al. reported that the association between expression of PD-L1 and CPS was related to the assessment of responsiveness to PD-L1 inhibitor target therapy, in which $CPS \ge 1$ increases the response to PD-L1 inhibitor ³⁸.



Figure 2: Kaplan-Meier survival curve for overall survival after gastrectomy. A: The overall survival time (OS) of patients with positive PD-L1 was significantly shorter than patients with negative PD-L1 (p = 0.001). **B:** OS of patients with CPS ≥ 1 was significantly shorter than patients with CPS < 1 (p = 0.047).

The results of our study show that the combination of assessment of PD-L1 expression and CPS can be a factor in helping predict and consider immunotherapy in gastric cancer patients. We believe these findings will help us improve therapeutic strategies in future clinical trials for treatment with PD-1/PD-L1 inhibitors. This study also had some limitations. The sample number was small, resulting in an analytical evaluation was not comprehensive. Additionally, the cut-off values for distinguishing between positive and negative PD-L1 expression still hold room for improvement.

CONCLUSIONS

Overexpression of PD-L1 and CPS \geq 1 contribute to poor prognosis and reduced survival in gastric cancer patients. Our study indicates that CPS and PD-L1 expression are independent prognostic indicators, and that combination of these two indicators can improve strategies for targeting PD-1/PD-L1 inhibitors in therapy for gastric cancer patients.

ABBREVIATIONS

AJCC: American Joint Committee on Cancer; DAB: 3-3'-**CPS**: Combined positive score; diaminobenzidine; FDA: Food and Drug Administration; GC: Gastric cancer; GC/GEJC: Gastric or gastroesophageal junction cancer; GLOBOCAN: Global Cancer Incidence, Mortality And Prevalence; IBM: International Business Machines Corporation; IHC: Immunohistochemical; NCCN: National Comprehensive Cancer Network; NY: New York; PBS: Phosphate buffered saline; PD-1: Programmed death 1; PD-L1: Programmed Death Ligand 1; SPSS: Statistical Package for the Social Sciences; TNM: Tumor-node-metastasis; UICC: Union for International Cancer Control; UK: United Kingdom; USA: United States of America

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AUTHOR'S CONTRIBUTIONS

Conceived and designed the study: TND NMH. Patient characteristics data and sample collection: TND NMH DTT TDT NTL NKT PVT NBD NMH DTC. Analyzed and interpreted the data: TND NMH TDT NKT DTC. Contributed to drafting and writing the article: TND NMH DTT TDT DTC. Involved in all aspects of the work through correspondence: NMH DTC. All authors read and approved the final manuscript.

FUNDING

None.

AVAILABILITY OF DATA AND MATERIALS

Data and materials used and/or analyzed during the current study are available from the corresponding author on reasonable request.

ETHICS APPROVAL AND CONSENT TO PARTICIPATE

All patients provided informed consent for participation in the study. The Ethics Committee of 103 Military Medical Hospital (code: 140/2016/IRB-MH103) approved all study procedures.

CONSENT FOR PUBLICATION

Not applicable.

COMPETING INTERESTS

The authors declare that they have no competing interests.

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