Prognostic value of CD73-adenosinergic pathway in solid tumor: A meta-analysis and systematic review

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ABSTRACT

CD73 is a glycosylphosphatidylinositol (GPI) anchored cell surface protein that is encoded by NT5E gene, plays multiple roles in tumor processes. Previous studies have presented a potential value of CD73 served as a detectable biomarker for prognosis of several solid tumors, but the results were more controversially. A comprehensive meta-analysis was conducted to precisely evaluate the prognostic role of CD73 in solid tumors. The included studies were searched in PubMed, Web of Science and EBSCO from Jan 1990 to Jan 2016. Pooled hazard ratios (HR) and corresponding 95% confidence intervals (CI) for overall survival (OS), disease free survival (DFS) were carried out using a fixed or random effects model. Totally, 13 studies about 12,533 patients were included. CD73-high expression was correlating with poor OS (pooled HR = 1.28, 95% CI = 1.19–1.37). In addition, CD73 expression had borderline association with worse DFS (pooled HR = 1.28, 95% CI = 1.01–1.62). Egger’s tests indicated that there was no evidence of significant publication bias. CD73 is an efficient prognostic biomarker in solid tumors, and over-expression of CD73 is associated with inverse OS or DFS. But this predictive value and target therapy for clinical practice yet needs advanced research.

INTRODUCTION

CD73, also known as ecto-5'-nucleotidase (NT5E), is a glycosylphosphatidylinositol linked cell surface enzyme found in normal tissues. Originally, CD73 was defined as a lymphocyte differentiation antigen and an adhesion molecule for lymphocytes binding to endothelium [1]. Recent studies implied that CD73 was over-expressed on various kinds of solid malignant tumors (i.e., breast cancer [2], colorectal cancer [3], prostate cancer [4], ovarian cancer [5], and gallbladder cancer [6]). Over-expression of CD73 was driven by tumor hypoxic microenvironment and some soluble inflammatory factors, such as type I IFNs, TNF-α, IL-1β, TGF-β and Wnt activators [7]. Otherwise, in breast cancer, CD73 expression can also be found negatively regulated by estrogen receptor (ER) expression [8]. CD73 catalyzes the conversion of adenosine (ADO) from AMP released to the extracellular environment, and control a variety of physiologic responses, as well as the development of cancer [9].

CD73 has both nonenzymatic and enzymatic functions in tumor environment [10]. For the nonenzymatic functions, CD73 plays a pivotal role for tumor cells proliferation, angiogenesis and apoptosis, via modulating extrinsic signaling, like EGFR/Akt, VEGF/Akt pathway and impairing antitumor immunity [11, 12]. CD73 endows tumor cells “invasive phenotype”, by reducing cell-cell adhesion, and inducing epithelial-mesenchymal transition (EMT) through the regulation of cadherin-1 and vimentin. So CD73 significantly contributes to tumor metastasis [7]. The enzymatic functions refer to CD73-generated adenosine playing an important role in tumor immune tolerance. Extracellular ADO can exert effect on tumor immune microenvironment through multiple pathways [1, 13]: 1) Limiting cytotoxic activity of effective immune cells: ADO significantly reduces CD8+
T cells homing and inhibits NK cells by interfering with the process of granule exocytosis and reducing the ability of NK cells to adhere to tumor cells [14, 15]. 2) Enhancing immunosuppressive effects: ADO inhibits M1 macrophage activation and induces M2 macrophage polarization via A2A and A2B receptors [16–18]. Myeloid-derived suppressor cells (MDSC) perform a vital role in damaging tumor immune surveillance. ADO promotes the expansion and accumulation of the MDSC in tumor milieu by engaging A2B receptors on myeloid precursor cells [19]. 3) Inducing anomalous differentiation and weakening the function of antigen presenting cells: ADO bonding with A2B receptor can alter dendritic cells phenotype, decrease the level of tumor antigen presentation, and increase vascular endothelial growth factor (VEGF) production [20, 21]. Taken together, CD73-adenosinergic pathway involves in creating a tumorigenic microenvironment by regulating the tumor itself and immune system (Figure 1).

Numerous in vitro studies demonstrated that CD73 expression was associated with tumor proliferation, invasiveness, angiogenesis, and therapy resistance. But the prognosis role of CD73 in different human solid tumors is controversially according to the following studies. For epithelial ovarian carcinomas, Oh et al. found CD73-high expression predicted better prognosis, lower stage, and higher degree of differentiation [22]. The similar result has also been reported in nonmuscle-invasive urothelial bladder cancer [23]. Nevertheless, other researches in colorectal, gastric, gallbladder, prostate and triple negative breast cancer, proved CD73 was an unfavorable prognostic marker [2, 5, 24, 25]. For purpose of more precisely evaluate the prognostic value of CD73-adenosinergic pathway in solid tumor, a systematic review and meta-analysis was conducted according to previous published studies.

**RESULTS**

**Search results and characteristics of eligible studies**

Initially, 1039 potential studies were yielded utilizing the electronic databases search, of which 13 articles met the inclusion criteria (Figure 2). Those relevant relevant articles were screened for eligibility by duplication and language, and 760 records were excluded. 256 articles were excluded through title and abstract screening. Another ten publications were further excluded after full-text screening. Finally, 13 publications full met the predefined criteria for this meta-analysis [2–6, 22–29]. Relevant characteristics of the eligible studies are summarized in Table 1.
The including articles were observational retrospective studies consisting of approximately 12,533 solid tumors patients, which evaluated the association between expression levels of CD73 and survival parameters. The number of participants ranged from 67 to 6209, and the median follow-up ranged from 18 to 110.6 months. 10/13 of the included studies got a quality score ≥ 6 NOS assessment.

### Evaluation of CD73 expression

The description of detection and definition method of CD73 used in eligible researches was generalized in Supplementary Table 1. In those including studies, three studies used CD73 gene expression, immunohistochemical staining (IHC) was applied in nine studies, and one study (Turcotte et al.) both using gene and protein detection. Among the studies defined CD73 positive, median expression of CD73 staining was 50.77%, and CD73 expression in solid tumors ranged from 26.4% to 74%. Most studies used tumor cell staining for determination of CD73 expression, however, Leclerc et al. and Zhang et al. also used tumor stroma staining.

### CD73 predicts poor OS

Twelve studies provided the association between CD73 and overall survival, including two studies for ovarian cancer, one study for renal cancer, five studies for gastrointestinal cancer, three studies for breast cancer, and one study about prostate cancer. The combined analysis from the published data showed that CD73-high expression was significantly associated with worse OS (pooled HR = 1.28, 95% CI = 1.19–1.37). We also found heterogeneity existed ($I^2 = 67.3\%$, $p = 0.000$) (Figure 3A). Next we conducted subgroup meta-analysis according to CD73 detection methods (IHC and gene expression). IHC evaluated in eight studies demonstrated that over-expression of CD73 was correlated with poor OS (pooled HR = 2.09, 95% CI = 1.57–2.79) (Figure 3B). And CD73 gene expression was used in four studies, and high level of CD73 was also related with worse OS (pooled HR = 1.24, 95% CI = 1.16–1.33). Heterogeneity was also found in this analysis ($I^2 = 58.5\%$, $p = 0.065$) (Figure 3D). Sensitivity analysis was conducted, and we found this heterogeneity was influenced by study sample size. After excluding Cushman et al.’s study, the heterogeneity decreased ($I^2 = 0.0\%$, $p = 0.919$) (Figure 3E).

Subset analyses based on cancer types were also conducted. Five studies for gastrointestinal cancer using IHC detection demonstrated that CD73 over-expression implied an unfavorable OS (pooled HR = 2.61, 95% CI = 1.87–3.63) (Figure 3C). As well as in breast cancer, CD73-high expression was associated with reduced OS (pooled HR = 1.33, 95% CI = 1.19–1.49) (Figure 3F).

Table 1: Characteristics of included studies

<table>
<thead>
<tr>
<th>Study, (Author/year)</th>
<th>Country</th>
<th>Tumor type</th>
<th>Tumor stage</th>
<th>Number of Patients</th>
<th>Male/Female</th>
<th>Age, Median (Range)</th>
<th>Follow-up, months (Range)</th>
<th>Survival Data</th>
<th>NOS Score</th>
</tr>
</thead>
<tbody>
<tr>
<td>Wettstein/2015 [23]</td>
<td>Switzerland</td>
<td>Urothelial Bladder Cancer</td>
<td>Nonmuscle Invasive</td>
<td>174</td>
<td>131/43</td>
<td>69.5 (32–92)</td>
<td>110.6 (32.4–226.8)</td>
<td>DFS</td>
<td>8</td>
</tr>
<tr>
<td>B</td>
<td></td>
<td></td>
<td></td>
<td>1581</td>
<td>0/1581</td>
<td>NR</td>
<td>NR</td>
<td>OS</td>
<td></td>
</tr>
<tr>
<td>Yu/2015 [26]</td>
<td>China</td>
<td>Renal cell cancer</td>
<td>I-IV</td>
<td>189</td>
<td>119/70</td>
<td>58 (35–87)</td>
<td>78 (1–118)</td>
<td>OS, DFS</td>
<td>8</td>
</tr>
<tr>
<td>Zhang/2015 [24]</td>
<td>China</td>
<td>Rectal cancer</td>
<td>I-IV</td>
<td>90</td>
<td>60/30</td>
<td>64.4 ± 12.5</td>
<td>88.5 (83–98)</td>
<td>OS</td>
<td>7</td>
</tr>
<tr>
<td>Oh/2012 [22]</td>
<td>Korea</td>
<td>Epithelial ovarian cancer</td>
<td>I-IV</td>
<td>167</td>
<td>0/167</td>
<td>50.3 ± 13.5</td>
<td>1–120</td>
<td>OS, DFS</td>
<td>8</td>
</tr>
<tr>
<td>Zhi/2012 [28]</td>
<td>China</td>
<td>Breast cancer</td>
<td>I-IV</td>
<td>2898</td>
<td>0/2898</td>
<td>NR</td>
<td>NR</td>
<td>OS, DFS</td>
<td>6</td>
</tr>
<tr>
<td>Supernat/2012 [29]</td>
<td>Poland</td>
<td>Breast cancer</td>
<td>I-III</td>
<td>136</td>
<td>0/136</td>
<td>58.4 (27–86)</td>
<td>21.6 (1.2–42)</td>
<td>OS, DFS</td>
<td>7</td>
</tr>
<tr>
<td>Wu/2012 [3]</td>
<td>China</td>
<td>Colorectal cancer</td>
<td>I-IV</td>
<td>135</td>
<td>75/60</td>
<td>59.5 ± 14.0</td>
<td>74.8 ± 36.2</td>
<td>OS</td>
<td>7</td>
</tr>
<tr>
<td>Validation cohort</td>
<td></td>
<td></td>
<td></td>
<td>223</td>
<td>120/103</td>
<td>57.9 ± 13.8</td>
<td>52.0 ± 21.1</td>
<td>OS</td>
<td>7</td>
</tr>
<tr>
<td>Training cohort</td>
<td>China</td>
<td>Colorectal cancer</td>
<td>I-IV</td>
<td>103</td>
<td>57/46</td>
<td>61.1 (22–83.3)</td>
<td>69.2</td>
<td>OS, DFS</td>
<td>6</td>
</tr>
<tr>
<td>Cushman/2015 [27]</td>
<td>USA</td>
<td>Colorectal cancer</td>
<td>IV</td>
<td>150</td>
<td>55/95</td>
<td>64.4 ± 12.5</td>
<td>120 ± 42</td>
<td>OS, DFS</td>
<td>7</td>
</tr>
</tbody>
</table>
CD73 implies unfavorable DFS

Six studies assessing CD73 expression by IHC talked about the solid tumors DFS. In total, 1159 patients were in the pooled analysis. CD73 positive expression had borderline association with unfavorable DFS (pooled HR = 1.28, 95% CI = 1.01–1.62), and significant heterogeneity ($I^2 = 71.4\%$, $p = 0.004$) (Figure 4A). Sensitivity analysis found the heterogeneity was generated from solid tumor categories. And subgroup analysis found CD73 overexpressed in ovarian cancer was significantly associated with poor DFS (pooled HR = 1.49, 95% CI = 1.14–1.95) (Figure 4B).

Sensitivity analysis and publication bias

Sensitivity analyses were carried out through sequential removal of individual studies to assess the potential heterogeneity of each study on the pooled HRs (data not shown). Egger’s tests and symmetric funnel plots indicated no evidence of significant publication bias in this meta-analysis (Figure 5). However, because of the limited studies included, there might still have possible publication bias in the current meta-analysis difficult to confirm.

DISCUSSION

CD73-adenosinergic pathway has been involved in the pathophysiology of substantial solid tumors. CD73 is thought as a key point in this pathway, but the prognostic role across different solid tumors is still in contradiction. This comprehensive meta-analysis of published data was conducted to assess expression of CD73 and their connection with solid tumors prognosis (for studies that evaluated CD73 by IHC or gene detection). We found that CD73-high expression is a marker of poor OS and DFS in solid tumors.

Among the evaluated tumor types, over-expression of CD73 in gastrointestinal cancer was the one mostly linked with a worse outcome. But in breast cancer, the association of CD73 with long-term survival was controversial, and our analysis results found CD73 was
boundary linked to worse survival. It may influenced by strong heterogeneity of breast cancer. Supernat et al. using IHC test showed the positive CD73 staining predicts longer DFS and OS on breast cancer. However, this study did not distinguish the breast cancer molecular subtypes [29]. Interestingly, Loi’s group through analyzing 44 publicly available microarray datasets about 6,209 breast cancer samples, they implied CD73 gene expression was associated with worse prognosis in triple negative breast cancer patients (HR = 1.5, 95% CI = 1−2.1, p = 0.029), but not in Luminal type (HR = 0.96, 95% CI = 0.77−1.2, p = 0.7) or HER2 positive type (HR = 1.0, 95% CI = 0.71−1.5, p = 0.86) [2]. The above two studies indicated CD73 may play different role in different subtypes of breast cancer. All of these suggested that CD73 as a prognostic biomarker for clinical use may depends on the tumor types and subtypes.

As we all known that CD73 expresses in both tumor cells and stromal cells, but the CD73 expression in tumor stromal cells was less well investigated. Only

Figure 3: Forest plots showing results of studies on the impact of CD73 expression of overall survival (OS). (A) Impact of CD73 expression on OS of patients with solid tumors; (B) Impact of CD73 expression by immunohistochemistry (IHC) on OS of patients with solid tumors; (C) Impact of CD73 expression by immunohistochemistry (IHC) on OS of patients with gastrointestinal cancer; (D) Impact of CD73 expression by CD73 gene detection on OS of patients with solid tumors; (E) Sensitivity analysis was conducted, and excluding Cushman et al., study, the impact of CD73 expression by CD73 gene detection on OS of patients with solid tumors; (F) Impact of CD73 expression by CD73 gene detection on OS of patients with breast cancer.

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two studies involved the stromal CD73, Leclerc’s study showed that positive CD73 staining in prostate cancer stroma was associated with longer biochemical recurrence free survival in univariate analysis (but not significant in multivariate analysis) [4]. A rectal adenocarcinoma research by Zhang et al. had also indicted that higher stromal CD73 expression was linked to early tumor stages and favorable OS. However our meta-analysis may provide strong supporting evidence for CD73 associated with poor OS and DFS [24]. This contradiction suggested that the prognostic role of CD73 may also influenced by the location of CD73 expression.

CD73-adenosinergic pathway promote tumor progress not only by regulating the tumor cells proliferation and angiogenesis, but also promoting the tumor to form a suppressive milieu by inhibiting CD8+ T cells, NK cells function, and increasing the generation of MDSCs. This involved tumorigenic microenvironment provide basic research evidences for our meta-analysis results that CD73 over-expression was associated with worse survival outcomes in several solid tumors. And reminds us that blocking this protein in CD73-adenosinergic pathway may be a useful therapeutic target for clinical intervention for different solid tumors in future [30].

Figure 4: Meta-analysis of impact of CD73 expression on disease free survival (DFS) of patients with solid tumors. (A) Impact of CD73 expression on DFS of patients with solid tumors; (B) Impact of CD73 expression on DFS of patients with ovarian cancer.
Recently, many preclinical and phase I trials of anti-CD73 have been performed [11, 31–34] (Table 2). MEDI9447 (a monoclonal antibody specific for the ectoenzyme, CD73) can reduce immunosuppression via increasing CD8+ T cells, inhibiting myeloid-derived suppressor cells and regulatory T cells in the tumor microenvironment, finally inhibit tumor progress. What’s more, Studies also showed that anti-CD73 not only exhibited the activity as a single agent, but also enhanced the activity of PD-1 blockade [31]. So CD73 as therapeutic target is becoming promising and complex.

Several limitations may exist in our meta-analysis. Firstly, the method and cut-off values for assessing CD73 expression are inconsistent. Secondly, due to the limited number of studies, we were incapable to perform detail subgroup analyses to avoid the tumor heterogeneity, so it may exists some heterogeneity. Finally, this is a literature based analysis, small and negative results researches may not be published, which may account for publication bias.

Despite the above limitations, this study was the first meta-analysis demonstrated that CD73-high expression in solid tumor tissues was associated with a worse prognosis, suggesting that CD73 could be used as a predict biomarker and targeting CD73 might be a promising therapeutic approach for solid tumors. Further data are required for the potential impact of CD73-adenosinergic pathway in tumors, especially under different molecule subtypes and different expression location. If anti-CD73 therapy can be proved effective finally, it might challenge the paradigm of adjuvant therapy for solid tumor. Future research, especially randomized controlled trials (RCT), is desirable to confirm this therapy to reduce cancer mortality.

**MATERIALS AND METHODS**

**Literature search and selection criteria**

This study based on Preferred Reporting Items for Systematic Reviews and Meta-Analyses guidelines. And according to the existing literature, so ethical committee or institutional review board approvals were not necessary.

Literature search was conducted independently by two investigators via PubMed, Web of Science and EBSCO published between January 1990 and

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**Figure 5:** Funnel plot for the evaluation of potential publication bias in the impact of CD73 on overall survival (A) and disease free survival (B) of patients with solid tumors.
January 2016. The following MeSH terms were used in search strategy: (“NT5E” OR “Ecto-5’-nucleotidase” OR “CD73”) AND (“CANCER” OR “TUMOR” OR “CARCINOMA”). Studies were selected for further analyze based on careful reading of the online titles and abstracts.

Publications meet all of the following criteria were included: a) studies investigated the association between CD73 and prognostic parameters of solid tumor; b) contained sufficient published data to calculate hazard ratio (HR) and 95% confidence interval (95% CI); and c) original research with English full text. While, duplicated articles or overlapped data were excluded.

Data extraction and assessment of study quality

Two investigators independently evaluated and selected the articles and extracted information in a standardized manner and subsequently resolved disagreements by discussion with another author. The following relevant data were extracted from eligible articles: first author’s name, year, country; number of analyzed patients, assessment methods for CD73 expression, follow-up time, tumor site and stage, and most importantly, the long-term survival (e.g., overall survival (OS) and disease-free survival (DFS)). However, the cut-off value for CD73 varied among including publications.

Table 2: Pre-clinical and clinical trials evaluating anti–CD73 therapeutic strategies

<table>
<thead>
<tr>
<th>Study</th>
<th>Phase/Condition</th>
<th>Experimental arm(s)</th>
<th>Efficiency</th>
<th>Reference</th>
</tr>
</thead>
</table>
| MedImmune/2016 | **Phase I (NCT02503774)** Advanced Solid Tumors | **Arm A**: MEDI9447  
**Arm B**: MEDI9447+MEDI4736 | Recruiting  | www.clinicaltrials.gov |
| Hay/2015     | Mice model CT26 colon cancer | Arm A: Untreated  
Arm B: Isotype Mix  
Arm C: MEDI9447  
Arm D: Anti-PD1  
Arm E: MEDI9447+Anti-PD1 | Enhances anti-tumor activity of anti-PD1 | [31] |
| Stagg/2010   | Mice model 4T1.2 breast cancer | Arm A: Ig  
Arm B: anti-CD73 mAb (TY/23) | Inhibits breast tumor growth and metastasis | [30] |
| Terp/2013    | Mice model LM3 hepatocellular cancer | Arm A: anti-CD73 AD2 mAb  
Arm B: control mAb | Inhibits the ability of circulating tumor cells to extravasate and colonize, leading to inhibition of metastasis | [32] |
| Allard/2013  | Mice model MC38-OVA colon cancer, RM-1 prostate cancer, 4T1.2 breast cancer | Arm A: Ig  
Arm B: anti-CD73 mAb (TY/23)  
Arm C: anti-PD-1 mAb (RMP1-14)  
Arm D: anti-CTLA-4 mAb (UC10-4F10)  
Arm E: TY/23+RMP1-14  
Arm F: TY/23+ UC10-4F10 | Enhances the therapeutic activity of anti-PD-1 and anti-CTLA-4 mAbs | [33] |
| Wang/2011    | Mice model ID8 ovarian cancer | Arm A: Untreated  
Arm B: APCP  
Arm C: T cells  
Arm D: T cells+APCP*  
Arm E: anti-CD73 mAb (TY/23)  
Arm F: T cells+TY/23 | Inhibits tumor growth and augments the efficacy of adoptive T cell therapy | [11] |

*APCP: selective inhibitor α,β-methylene adenosine 5’-diphosphate.
so CD73-high expression was defined refer to the original articles. In order to avoid bias, the HR was extracted preferentially from multivariable analyses. If not available, HR from univariate analyses also extracted. Kaplan–Meier curves were evaluated using GetData Graph Digitizer 2.26 (http://getdatagraph-digitizer.com) for the articles did not provide survival data directly.

Two independent authors conducted the quality assessment for each eligible study using Newcastle–Ottawa Scale (NOS). NOS score higher than six were indicated as high-quality studies.

**Statistical analysis**

Survival outcomes were the primary end-points in this study, so the data synthesized using HR and 95% CI to evaluate the impact of CD73 expression on long-term survival of solid tumor. The heterogeneity of pooled data was measured using the Cochran $Q$-test and I-squared test. When $I^2 < 50\%$ and $p > 0.10$ were considered as no heterogeneity and a fixed-effects model (Mantel-Haenszel) was performed. Otherwise, random-effects model (DerSimonian and Laird) was adopted. The sources of inter-study heterogeneity were explored using subgroup analysis and sensitivity analysis. A funnel plot with Begg’s and Egger’s test was applied to assess the potential publication bias.

Statistical analyses were performed using STATA 12.0 software (STATA Corporation, College Station, TX, http://www. stata.com). And statistical significance was defined as $p$-value $< 0.05$.

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**CONFLICTS OF INTEREST**

The authors declare no conflicts of interest.

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