



Immunotherapy in thymic epithelial tumors: tissue predictive biomarkers for immune checkpoint inhibitors

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Abstract

Thymic epithelial tumors (TETs) are rare malignant neoplasms arising in the thymus gland. Nevertheless, TETs, including thymomas (TMs), thymic carcinomas (TCs), and thymic neuroendocrine neoplasms (TNENs), are the most common mediastinal malignancies overall. A multidisciplinary approach is required for the appropriate diagnostic and therapeutic management of TETs. To date, the main therapeutic strategies are largely depended on the stage of the tumor and they include surgery with or without neoadjuvant or adjuvant therapy, represented by platinum-based chemotherapy, radiotherapy or chemoradiotherapy. Immune checkpoint inhibitors (ICIs) are ongoing under evaluation in the advanced or metastatic diseases despite the challenges related to the very low tumor mutation burden (TMB) and the high incidence of immune-related adverse events in TETs. In this regard, predictive impact of tissue biomarkers expression such as programmed cell death ligand-1 (PD-L1), and other emerging biomarkers, as well as their optimal and shared interpretation are currently under evaluation in order to predict response rates to ICIs in TETs.

Keywords

Thymic epithelial tumors, immunotherapy, thymoma, thymic carcinoma

Introduction

Thymic epithelial tumors (TETs) are rare thoracic neoplasms arising from epithelial cells of the thymus gland [1]. Although less common than other thoracic neoplasms, such as pulmonary and pleural neoplasms [2–4]. TETs are the most frequently encountered tumors in the prevascular mediastinum [5]. TETs are a



basket of different tumors with different clinical, histopathological, immunophenotypical, molecular and biological features at the base of clinical-pathological differences [6–9]. TETs share the lowest tumor mutation burden (TMB) among adult solid tumors [7, 10] and they, particularly thymomas (TMs), are often associated with peculiar autoimmune diseases, particularly myasthenia gravis, pure red cell aplasia and Good’s syndrome [11]. The choice of the optimal therapeutic treatment of TETs is primarily based on staging and histotype; in detail, surgery is the treatment of choice, being the only curative strategy in localized diseases, with eventually combined radiotherapy and/or chemotherapy on the base of surgical radicality, histotype and stage disease, while platinum-based chemotherapy is the standard of care for locally-advanced or metastatic TETs [12, 13]. However, therapeutic strategies for relapsed or refractory TETs are limited with different targeted agents (epidermal growth factor receptor inhibitors, inhibitors of angiogenesis, c-kit inhibitors, histone deacetylase inhibitors), since no real benefit has been shown in these clinical settings [14–17]. These data and the low TMB of tumors, limiting the identification of new therapeutic targets, explain the need to research new therapeutic strategies. In this regard, immune checkpoint inhibitors (ICIs) are a promising option just likes for other advanced stage malignant neoplasms [18–21].

World Health Organization 2021 classification of TETs: a short summary

TETs arising from thymic epithelial cells, particularly thymic cortical epithelial cells (cTECs) or thymic medullary epithelial cells (mTECs) include TMs, thymic carcinomas (TCs) and thymic neuroendocrine neoplasms (TNENs) (Table 1) [6, 22]. TMs are the most common neoplastic type accounting for more than 50% of TETs, while TCs and TNENs represent approximately 14–22% and 2–5%, respectively [23]. TMs are characterized by thymus-like differentiation as they variably show organotypical features such as lobulated architecture, perivascular spaces, medullary differentiation and intratumoral infiltration of immature T-lymphocytes while Hassall corpuscles are only occasionally identified [6, 24, 25]. TMs are a heterogeneous group of neoplasm with different molecular, histopathological, immunophenotypical and clinical features [26]. TMs are variably encapsulated and well circumscribed masses, except for type B3 TM which shows smooth invasive fronts with invasion in mediastinal fat or adjacent organs; neoplastic proliferation shows lobulated architecture due to the presence of thick fibrous septa and it is organized according to several growth patterns, with bland and spindle/oval cytomorphology and few or no admixed immature T-cells in type A TM, except for s atypical type A TM, and variably atypical polygonal neoplastic cells, as single cellular elements or arranged in clusters (≥ 3 contiguous cells), with dense immature T-lymphocyte infiltrate or few and scattered immature T lymphocytes in type B TMs (B1–B3). Furthermore, some TMs, such as metaplastic TM or micronodular TM with lymphoid stroma, show histopathological features that do not fit well with the most common histotypes and so they are classified separately. Finally, more than one histological subtype defines mixed cases of TM and the diagnosis should list the predominant pattern followed by any minor components in 10% increments, except for type AB TM which is considered a distinct TM subtype (Figure 1) [27–29]. TC is a very rare mediastinal tumor characterized by morphological features of obvious malignant biological behavior and with peculiar immunophenotypic expression of *CD5* and *CD117* (c-kit), unlike conventional carcinomas of other anatomical sites. Several histological types of TC are recognized (Figure 1) [6, 30]. TNENs are very rare thymic neoplasm accounting for 2–5% of all thymic tumors and they are currently classified just like their pulmonary counterpart, with the same nomenclature and the same diagnostic criteria (mitotic index/2 mm², presence/absence of necrosis and cytomorphological features) of lung neutrophil extracellular traps (NETs) and neuroendocrine carcinoma (NECs) [31].

Table 1. World Health Organization (WHO) classification of TETs, 5th edition [6]

TM	TC	TNENs
Type A TM	Squamous cell carcinoma of the thymus	Typical carcinoid of the thymus
Type AB TM	Basaloid carcinoma of the thymus	Atypical carcinoid of the thymus
Type B1 TM	Lymphoepithelial carcinoma of the thymus	Small cell carcinoma of the thymus

Table 1. World Health Organization (WHO) classification of TETs, 5th edition [6] (*continued*)

TM	TC	TNENs
Type B2 TM	Clear cell carcinoma of the thymus	Large cell neuroendocrine carcinoma of the thymus
Type B3 TM	Low-grade papillary adenocarcinoma of the thymus	-
Micronodular TM with lymphoid stroma	Mucoepidermoid carcinoma of the thymus	-
Metaplastic TM	TC with adenoid cystic carcinoma-like features	-
Lipofibroadenoma of the thymus	Enteric-type adenocarcinoma of the thymus	-
	Adenocarcinoma not otherwise specified (NOS) of the thymus	
	Adenosquamous carcinoma of the thymus	
	Sarcomatoid carcinoma of the thymus	
	Undifferentiated carcinoma of the thymus	
	TC NOS	

-: blank cells

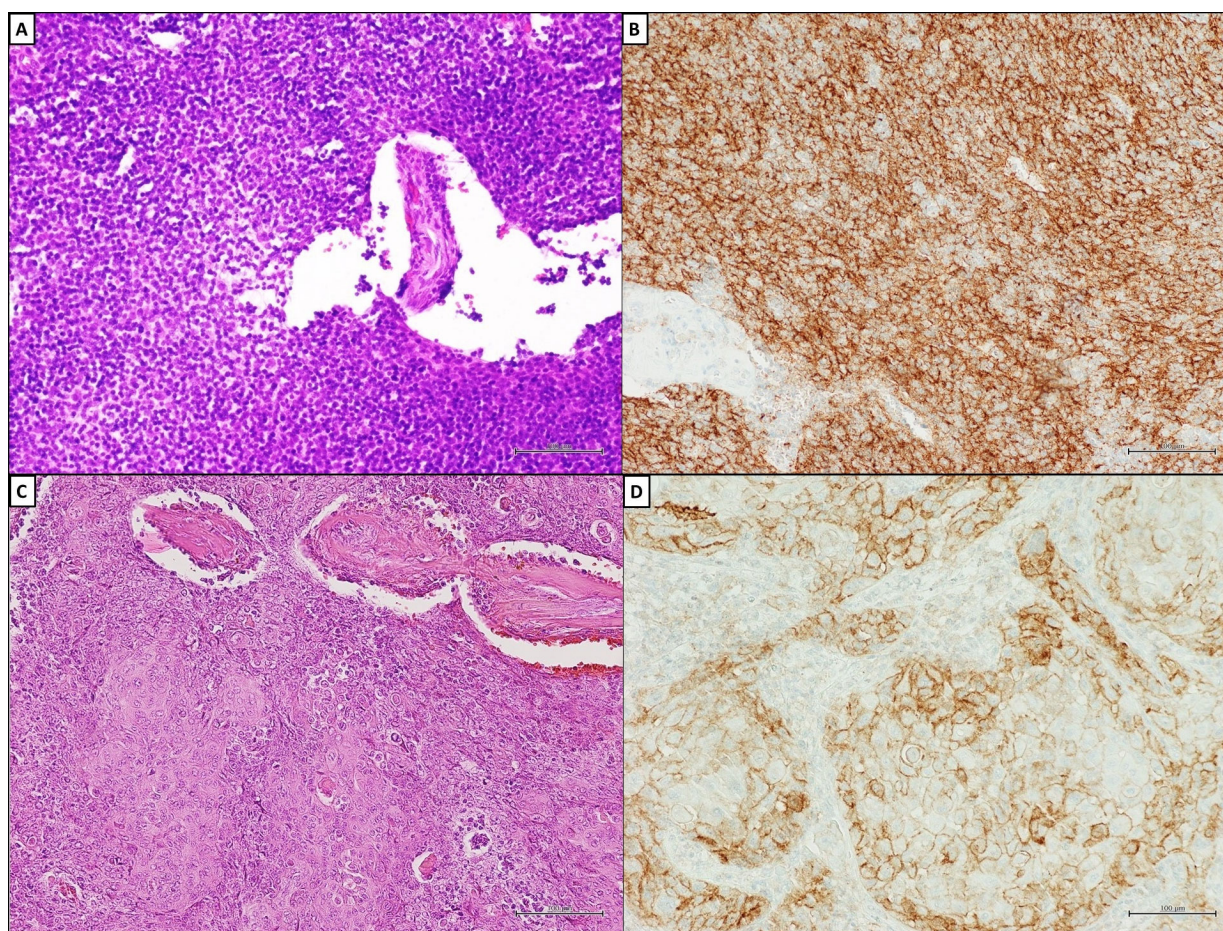


Figure 1. Programmed cell death ligand-1 (PD-L1) expression on tumor cells of TETs. (A) Histopathological appearance of type B2 TM; (B) diffuse and strong immunohistochemical (IHC) membranous staining (MS) for PD-L1 (clone SP263; Ventana Medical Systems) in $\geq 50\%$ of neoplastic cells in type B2 TM and squamous cell carcinoma of the thymus; (C) squamous cell carcinoma of the thymus (hematoxylin and eosin); (D) squamous cell carcinoma of the thymus. A–D, scale bar = 100 μm ; original magnifications $\times 200$. Courtesy of Pathology Unit of Università degli Studi della Campania “L. Vanvitelli”

Immunotherapy in TETs

Potential role of ICI in TETs

Development of neoplasms is partially related to the inability of the immune system to eliminate neoplastic cells in the early stages of the disease [32]. Therefore, multiple immunotherapeutic strategies have been

developed in order to activate and increase antitumor immunity [33]. ICIs are the most common form of immunotherapy strategy used in the clinical practice and they act by targeting programmed death 1 (PD-1)/PD-L1 axis or cytotoxic T-lymphocyte-associated protein 4 (CTLA-4) [34–37]. The efficacy of ICIs can be predicted by the evaluation of appropriate biomarkers of therapy response [38]. The level of PD-L1 expression in tumor cells and TMB are established as predictive biomarkers of therapeutic response in many solid tumors [19, 39, 40]. Recently, investigative efforts have been underway to identify predictive biomarkers of response to ICIs in TETs [41]. In this regard, TETs are characterized by the lowest TMB among all solid adult neoplasms [7, 10] and they are often associated with immune-mediated diseases [11]. These features are of great interest as they could negatively influence the use of ICIs in clinical practice. TMB reflects the number of non-synonymous single nucleotide variants (nsSNVs) in a neoplasm and therefore it is an expression of neo-antigens which can trigger an anti-tumor immune reaction, so determining the therapeutic response to ICIs; although a low TMB is associated with a low response rate to ICIs, the initial experience highlighted a surprisingly higher response rate to ICIs in TETs. The discordance between low TMB and a higher response rate is likely related to the specific biological function of the thymic gland and primarily to its role in T-cell development [42–45]. Moreover, the activation of antitumor immunity can also increase the risk of developing immune-related adverse events [46]. PD-L1 expression in TETs has been widely studied and overall data support its promising role as response predictor [8, 43, 44, 47–67]. Therefore, PD-L1 expression on tumor cells and its expression level appear to be the most promising biomarkers for immunotherapy response in TETs to date [41, 68, 69].

PD-1/PD-L1 expression in TETs

PD-L1 expression on TETs tumor samples staining has been evaluated in many papers (Table 2) [8, 47–66] as well as its predictive value in patients with TETs who had progressed after at least one line of chemotherapy [43, 44, 70]. High PD-L1 and PD-1 IHC expression was widely observed in TETs. In detail, PD-L1 expression ranging from 23–92% of tumor cells in TMs and 36–100% of tumor cells in TCs [69]; variations in protein expression levels are related to various conditions, particularly disease stage (Masaoka stages) and subtype of TETs, with higher levels observed in thymic [70]. Rouquette et al. [67] compared the most used PD-L1 antibodies (i.e., clone E1L3N, clone 22C3, clone SP142, and clone SP263) in a cohort of 103 TETs (53 B3 TMs and 50 TCs) and observed a good concordance, using both 1% and 50% cut-off. However, no cut-off value for PD-L1 positivity has been uniformly and unequivocally recognized, to date. In this regard, it is interesting to report two clinical trials, which studied the effectiveness of ICIs in progressing after at one line of chemotherapy TETs [43, 44]. In their phase II clinical trial, Giaccone et al. [44] aimed to study the effectiveness of pembrolizumab and they used clone 22C3 in order to define PD-L1 expression of archival formalin-fixed paraffin-embedded tissues. They classified PD-L1 IHC expression as high, if at least 50% of the tumour cells stained positive; low if protein expression was observed in 1–49% of tumor cells and negative if no tumour cells in the sample expressed PD-L1. Progression free-survival (PFS) and overall survival (OS) were longer in patients with high PD-L1 expression than those with low or no expression, suggesting that PD-L1 expression in at least 50% of tumor cells correlated with a better response to ICIs (Table 3). Cho et al. [43] likewise investigated PD-L1 expression by immunohistochemistry using the same antibody (clone 22C3) and a similar interpretation protocol (PD-L1) positivity was defined by membranous PD-L1 staining in $\geq 1\%$ of tumor and associated inflammatory cells or positive staining of stroma; PD-L1 expression was classified as high if at least 50% of the tumor cells, inflammatory cells, or stroma cells stained positive while PD-L1 expression in 0% to 49% of cells was classified as low expression). The trial results demonstrated that high PD-L1 immunohistochemistry expression was significantly associated with better response to pembrolizumab in TETs; indeed, overall response rate (ORR) was 35.7% in patients with PD-L1 expression levels higher than 50% (Table 3). Overall data suggest that ICIs (pembrolizumab) yielded encouraging antitumor activity with durable time in refractory, metastatic or relapsed TETs and that the best response rates are obtained with a PD-L1 expression greater than 50%, reporting a significant correlation between high PD-L1 levels and better and more durable response to ICIs.

Table 2. PD-L1 expression on TETs

Reference	TM	TC	Antibody of PD-L1	IHC staining criteria of positivity	TM positivity rate	TC positivity rate
Weissferdt et al. [47]	74/100 (74.00%)	26/100 (26.00%)	Clone EPR4877	MS > 5%	64.00%	54.00%
Arbour et al. [48]	12/23 (52.17%)	11/23 (47.83%)	Clone E1L3N	MS > 25%	92.00%	36.00%
Suster et al. [49]	-	21	Clone SP142	MS > 50%	NA	71.40%
Higuchi et al. [50]	31/39 (79.50%)	8/39 (20.50%)	Clone 28-8	MS ≥ 1%	51.60%	62.50%
Wei et al. [51]	100/169 (59.20%)	69/169 (40.80%)	Clone E1L3N	MS > 50%	36.00%	37.00%
Duan et al. [52]	20/33 (60.60%)	13/33 (39.40%)	Clone Ab58810	Intensity of staining score (1–3); median value of all scores as the cut-off value	65.00%	46.20%
Funaki et al. [53]	-	43	Clone SP142	MS > 50%	NA	60.50%
Katsuya et al. [54]	101/139 (72.60%)	38/139 (27.40%)	Clone E1L3N	H-score: score 3 [staining intensity (0–3) × % of positive cells (0–100%)]	23.00%	70.00%
Padma et al. [55]	65/69 (94.20%)	4/69 (5.80%)	Clone 15	Score 3 (intensity of MS 0–3)	68.00%	75.00%
Marchevsky et al. [56]	38/46 (82.60%)	8/46 (17.40%)	Clone SP142	MS ≥ 6%	92.00%	50.00%
Enkner et al. [8]	37/72 (51.30%)	35/72 (48.70%)	Clone E1L3N	H-score (cut-off value NA)	89.00%	53.00%
Katsuya et al. [57]	12/30 (40.00%)	18/30 (60.00%)	Clone E1L3N	Score ≥ 1 (intensity of MS 0–3)	67.00%	41.00%
Yokoyama et al. [58]	82	-	Clone EPR1161	Youden's index > 38%	53.70%	NA
Tiseo et al. [59]	87/107 (81.30%)	20/107 (18.70%)	Clone E1L3N	H-score: score 3 (intensity of MS 0–3)	18.00%	65.00%
Owen et al. [60]	32/35 (91.40%)	3/35 (8.60%)	Clone 22C3	Score 1 (intensity of MS 0–5)	81.00%	100.00%
Hakiri et al. [61]	81	-	Clone SP142	MS ≥ 1%	27.00%	NA
Guleria et al. [62]	84	-	Clone SP263	MS > 25%	82.00%	NA
Chen et al. [63]	40/70 (57.00%)	30/70 (43.00%)	Clone SP142	MS ≥ 5%	37.50%	76.70%
Bagir et al. [64]	38/44 (86.30%)	6/44 (13.70%)	Clone AM26531AF-N	MS > 5%	81.60%	83.30%
Ishihara et al. [65]	55/66 (83.30%)	11/66 (16.70%)	Clone SP263	MS > 25%	92.70%	72.70%
Berardi et al. [66]	63/68 (92.60%)	5/68 (7.40%)	Clone 28-8	MS > 1%	Overall: 25.00%	

NA: not available; -: blank cells

Table 3. Completed clinical trials with ICIs in TETs

Reference	Treatment	TM/TC	PD-L1 cut-off	PD-L1 positive cases	Primary endpoint	mOS	mPFS	ORR
Giaccone et al. [44]	Pembrolizumab	0/40	PD-L1 ^{high} : ≥ 50%	10/37 (27%)	ORR	Un	4.2	-
			PD-L1 ^{low} : 1–49%	27/37 (73%)				
Cho et al. [43]	Pembrolizumab	7/26	PD-L1 ^{high} : ≥ 50%	14/24 (58.3%)	ORR	-	-	35.7%
			PD-L1 ^{low} : 1–49%	10/24 (41.7%)				

mPFS: median PFS; mOS: median OS; Un: unachieved; NR: no response; -: blank cells

Emerging biomarkers for immunotherapy in TETs

Inhibition of the PD-1/PD-L1 axis is the most promising and most studied immunotherapeutic strategy in TETs. However, there are other immune checkpoints that could be targeted, such as, B7-H3, B7-H4, T cell

immunoglobulin and mucin domain-containing protein 3 (TIM-3), and several co-stimulatory molecules, such as CD137, glucocorticoid-induced tumour necrosis factor receptor family-related protein (GITR), inducible co-stimulator (ICOS), regulating T-cell mediated anti-tumor response [36, 48, 71]. In detail, B7-H4 protein belongs to the B7 family. B7-H4 is a negative co-stimulatory molecule and allows tumor cells to escape immune surveillance; it also plays an essential role in the formation of the tumor microenvironment. B7-H4 protein usually has low expression in normal tissues but higher expression several solid neoplasms [72–74]. High expression of B7-H4 protein by IHC staining (anti-B7-H4 monoclonal antibody, clone EP1165) is positively correlated with high regulatory T cells and forkhead box protein P3 (FOXP3) expression in the microenvironment, thus it can indicate the suppressive immune microenvironment and this relation could predict poor prognosis in patients with TETs [71, 75]. The expression of tissue biomarkers such as TIM-3, CD137, GITR, ICOS and CTLA-4 on tumor infiltrating lymphocytes (TILs) of TETs has been recently evaluated. Arbour et al. [48] observed an expression of TIM-3 and GITR in all evaluated TETs samples while ICOS and CTLA-4 were positive in almost all the samples (91%), with a moderate to high expression of these biomarkers. These data suggest a synergistic action of anti-TIM-3 or CD137 agonist with anti-PD-1/PD-L1 blockade, highlighting the potential need to evaluate the tissue expression of these biomarkers [48]. Moreover, recent results have been shown a Wilms' tumour 1 (WT1) IHC expression on tumor cells, underlining the value of WT1 peptide as an immunotherapy target, particularly WT1 peptide vaccination as a new avenue for treatment of advanced or recurrent TETs [76]. The *WT1* gene is a tumor suppressor and it's overexpressed in several solid and non-solid neoplasms [77–83]. WT1 protein plays several oncogenic roles including involvement in cancer cell growth [84], resistance to apoptosis [85], enhancement of cell migration [86] and tumor vascularization [87]. WT1-targeting immunotherapeutic strategies have been developed in the past years [88–91]. Oji et al. [92] conducted the first phase II clinical trial of WT1 peptide vaccine immunotherapy for advanced TETs. In their report WT1 expression was assessed with IHC staining using a monoclonal anti-WT1 antibody (clone 6f-H2) and samples were scored as positive (WT1 overexpression) when more than 10% of tumor cells were stained in either their cytoplasm or nucleus. Their interesting results showed as WT1 was overexpressed in the majority of TETs (11 of 13 TCs and 4 of 5 TMs) and that vaccination with WT1 peptide induced WT1-specific immune responses and WT1 peptide immunotherapy had clinical potential with a stable disease rate of 75.0% both in patients with TCs and TMs at 3 months. Thus, although the data is still limited, WT1 overexpression in TETs provides an opportunity to develop specific cancer vaccines [93]; however, the impact of this therapeutic strategy on the development of autoimmune-related complications is not yet known. Therefore, future clinical studies are needed to demonstrate the real therapeutic value of WT1 peptide-based immunotherapy and to study the association of WT1 peptide vaccine with the development of autoimmune-related complications in TETs [76, 94].

Conclusions

Immunotherapy is currently not a standard-of-care in TETs but ICIs have demonstrated encouraging clinical activity in relapsed and refractory TETs, although their administration is associated with a high risk of developing or precipitating immune-related adverse events in this clinical setting. Despite the rarity of the tumors, many papers demonstrated significant expression levels of PD-L1 on TETs cells, both as a percentage of immunopositive tumor cells and as intensity of expression. Most of these papers have however evaluated only the percentage of staining cells to define positivity cut-offs, just like the completed clinical trials. In addition, the expression of co-inhibitory immune checkpoints and co-stimulatory molecules regulating antitumor response on TETs tissue samples has been evaluated. Taken together, all these data provide a rationale for using ICIs for treatment of TETs and defining a standardized, univocal and reproducible evaluation protocol for predictive tissue biomarkers, particularly PD-L1, in order to pave the way for the personalized use of ICIs inhibitors in TETs.

Abbreviations

CTLA-4: cytotoxic T-lymphocyte-associated protein 4

GITR: glucocorticoid-induced tumour necrosis factor receptor family-related protein

ICIs: immune checkpoint inhibitors

ICOS: inducible co-stimulator

IHC: immunohistochemical

MS: membranous staining

ORR: overall response rate

PD-1: programmed death 1

PD-L1: programmed cell death ligand-1

TCs: thymic carcinomas

TETs: thymic epithelial tumors

TIM-3: T cell immunoglobulin and mucin domain-containing protein 3

TMB: tumor mutation burden

TMs: thymomas

TNENs: thymic neuroendocrine neoplasms

WT1: Wilms' tumour 1

Declarations

Author contributions

SL: Conceptualization, Methodology, Investigations, Writing—original draft, Writing—review & editing. MA and SC: Methodology, Investigations. RF: Conceptualization, Methodology, Investigations, Writing—original draft, Writing—review & editing, Supervision.

Conflicts of interest

The authors declare that they have no conflicts of interest.

Ethical approval

According to Italian law, the content of this article does not require ethical approval.

Consent to participate

According to Italian law, the content of this article does not require consent to participate. The images could be used for diulgative purpose for the italian law, since they are totally anonymized.

Consent to publication

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Availability of data and materials

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References

1. Rioja P, Ruiz R, Galvez-Nino M, Lozano S, Valdiviezo N, Olivera M, et al. Epidemiology of thymic epithelial tumors: 22-years experience from a single-institution. *Thorac Cancer*. 2021;12:420–5.
2. Lucà S, Zannini G, Morgillo F, Della Corte CM, Fiorelli A, Zito Marino F, et al. The prognostic value of histopathology in invasive lung adenocarcinoma: a comparative review of the main proposed grading systems. *Expert Rev Anticancer Ther*. 2023;23:265–77.
3. Bade BC, Dela Cruz CS. Lung cancer 2020: epidemiology, etiology, and prevention. *Clin Chest Med*. 2020;41:1–24.
4. Bibby AC, Tsim S, Kanellakis N, Ball H, Talbot DC, Blyth KG, et al. Malignant pleural mesothelioma: an update on investigation, diagnosis and treatment. *Eur Respir Rev*. 2016;25:472–86.
5. Hsu CH, Chan JK, Yin CH, Lee CC, Chern CU, Liao CI. Trends in the incidence of thymoma, thymic carcinoma, and thymic neuroendocrine tumor in the United States. *PLoS One*. 2019;14:e0227197.
6. WHO classification of tumours [Internet]. IARC; c1965-2022 [cited 2021 Apr 21]. Available from: <http://tumourclassification.iarc.who.int>
7. Radovich M, Pickering CR, Felau I, Ha G, Zhang H, Jo H, et al. The integrated genomic landscape of thymic epithelial tumors. *Cancer Cell*. 2018;33:244–58.e10.
8. Enkner F, Pichlhöfer B, Zaharie AT, Krunic M, Holper TM, Janik S, et al. Molecular profiling of thymoma and thymic carcinoma: genetic differences and potential novel therapeutic targets. *Pathol Oncol Res*. 2017;23:551–64.
9. Girard N, Basse C, Schrock A, Ramkissoon S, Killian K, Ross JS. Comprehensive genomic profiling of 274 thymic epithelial tumors unveils oncogenic pathways and predictive biomarkers. *Oncologist*. 2022;27:919–29.
10. Wang Y, Thomas A, Lau C, Rajan A, Zhu Y, Killian JK, et al. Mutations of epigenetic regulatory genes are common in thymic carcinomas. *Sci Rep*. 2014;4:7336.
11. Marx A, Willcox N, Leite MI, Chuang WY, Schalke B, Nix W, et al. Thymoma and paraneoplastic myasthenia gravis. *Autoimmunity*. 2010;43:413–27.
12. Tartarone A, Lerose R, Lettini AR, Tartarone M. Current treatment approaches for thymic epithelial tumors. *Life (Basel)*. 2023;13:1170.
13. Merveilleux du Vignaux C, Dansin E, Mhanna L, Greillier L, Pichon E, Kerjouan M, et al. Systemic therapy in advanced thymic epithelial tumors: insights from the RYTHMIC prospective cohort. *J Thorac Oncol*. 2018;13:1762–70.
14. Rajan A, Giaccone G. Treatment of advanced thymoma and thymic carcinoma. *Curr Treat Options Oncol*. 2008;9:277–87.
15. Thomas A, Rajan A, Berman A, Tomita Y, Brzezniak C, Lee MJ, et al. Sunitinib in patients with chemotherapy-refractory thymoma and thymic carcinoma: an open-label phase 2 trial. *Lancet Oncol*. 2015;16:177–86. Erratum in: *Lancet Oncol*. 2015;16:e105.
16. Zucali PA, De Pas T, Palmieri G, Favaretto A, Chella A, Tiseo M, et al. Phase II study of everolimus in patients with thymoma and thymic carcinoma previously treated with cisplatin-based chemotherapy. *J Clin Oncol*. 2018;36:342–9.
17. Chen Y, Gharwan H, Thomas A. Novel biologic therapies for thymic epithelial tumors. *Front Oncol*. 2014;4:103.
18. Ruiz-Cordero R, Devine WP. Targeted therapy and checkpoint immunotherapy in lung cancer. *Surg Pathol Clin*. 2020;13:17–33.
19. Lucà S, Franco R, Napolitano A, Soria V, Ronchi A, Zito Marino F, et al. PATZ1 in non-small cell lung cancer: a new biomarker that negatively correlates with PD-L1 expression and suppresses the malignant phenotype. *Cancers (Basel)*. 2023;15:2190.
20. Reck M, Remon J, Hellmann MD. First-line immunotherapy for non-small-cell lung cancer. *J Clin Oncol*. 2022;40:586–97. Erratum in: *J Clin Oncol*. 2022;40:1265.

21. Zhang Y, Zhang Z. The history and advances in cancer immunotherapy: understanding the characteristics of tumor-infiltrating immune cells and their therapeutic implications. *Cell Mol Immunol.* 2020;17:807–21.
22. Alves NL, Takahama Y, Ohigashi I, Ribeiro AR, Baik S, Anderson G, et al. Serial progression of cortical and medullary thymic epithelial microenvironments. *Eur J Immunol.* 2014;44:16–22. Erratum in: *Eur J Immunol.* 2014;44:2197.
23. Willner J, Zhou F, Moreira AL. Diagnostic challenges in the cytology of thymic epithelial neoplasms. *Cancers (Basel).* 2022;14:2013.
24. Marx A, Chan JK, Coindre JM, Detterbeck F, Girard N, Harris NL, et al. The 2015 World Health Organization classification of tumors of the thymus: continuity and changes. *J Thorac Oncol.* 2015;10:1383–95.
25. Marx A, Chan JKC, Chalabreysse L, Dacic S, Detterbeck F, French CA, et al. The 2021 WHO classification of tumors of the thymus and mediastinum: what is new in thymic epithelial, germ cell, and mesenchymal tumors? *J Thorac Oncol.* 2022;17:200–13.
26. Kuhn E, Pescia C, Mendogni P, Nosotti M, Ferrero S. Thymic epithelial tumors: an evolving field. *Life (Basel).* 2023;13:314.
27. Moran CA, Weissferdt A, Kalhor N, Solis LM, Behrens C, Wistuba II, et al. Thymomas I: a clinicopathologic correlation of 250 cases with emphasis on the World Health Organization schema. *Am J Clin Pathol.* 2012;137:444–50.
28. Moran CA, Walsh G, Suster S, Kaiser L. Thymomas II: a clinicopathologic correlation of 250 cases with a proposed staging system with emphasis on pathologic assessment. *Am J Clin Pathol.* 2012;137:451–61.
29. Weissferdt A. Common thymomas: classification, histology, staging and prognosis. *Diagn Histopathol.* 2023;29:94–104.
30. Alqaidy D, Moran CA. Thymic carcinoma: a review. *Front Oncol.* 2022;12:808019.
31. Bohnenberger H, Dinter H, König A, Ströbel P. Neuroendocrine tumors of the thymus and mediastinum. *J Thorac Dis.* 2017;9:S1448–57.
32. Forde PM, Reiss KA, Zeidan AM, Brahmer JR. What lies within: novel strategies in immunotherapy for non-small cell lung cancer. *Oncologist.* 2013;18:1203–13.
33. Yang Y. Cancer immunotherapy: harnessing the immune system to battle cancer. *J Clin Invest.* 2015;125:3335–7.
34. Mandal R, Chan TA. Personalized oncology meets immunology: the path toward precision immunotherapy. *Cancer Discov.* 2016;6:703–13.
35. Hargadon KM, Johnson CE, Williams CJ. Immune checkpoint blockade therapy for cancer: an overview of FDA-approved immune checkpoint inhibitors. *Int Immunopharmacol.* 2018;62:29–39.
36. Pardoll DM. The blockade of immune checkpoints in cancer immunotherapy. *Nat Rev Cancer.* 2012;12:252–64.
37. Ribas A, Wolchok JD. Cancer immunotherapy using checkpoint blockade. *Science.* 2018;359:1350–55.
38. Song Y, Li Z, Xue W, Zhang M. Predictive biomarkers for PD-1 and PD-L1 immune checkpoint blockade therapy. *Immunotherapy.* 2019;11:515–29.
39. Herbst RS, Soria JC, Kowanetz M, Fine GD, Hamid O, Gordon MS, et al. Predictive correlates of response to the anti-PD-L1 antibody MPDL3280A in cancer patients. *Nature.* 2014;515:563–7.
40. Lin H, Wei S, Hurt EM, Green MD, Zhao L, Vatan L, et al. Host expression of PD-L1 determines efficacy of PD-L1 pathway blockade-mediated tumor regression. *J Clin Invest.* 2018;128:1708. Erratum in: *J Clin Invest.* 2018;128:805–15.
41. Ao YQ, Gao J, Wang S, Jiang JH, Deng J, Wang HK, et al. Immunotherapy of thymic epithelial tumors: molecular understandings and clinical perspectives. *Mol Cancer.* 2023;22:70.

42. Samstein RM, Lee CH, Shoushtari AN, Hellmann MD, Shen R, Janjigian YY, et al. Tumor mutational load predicts survival after immunotherapy across multiple cancer types. *Nat Genet.* 2019;51:202–6.
43. Cho J, Kim HS, Ku BM, Choi YL, Cristescu R, Han J, et al. Pembrolizumab for patients with refractory or relapsed thymic epithelial tumor: an open-label phase II trial. *J Clin Oncol.* 2019;37:2162–70.
44. Giaccone G, Kim C, Thompson J, McGuire C, Kallakury B, Chahine JJ, et al. Pembrolizumab in patients with thymic carcinoma: a single-arm, single-centre, phase 2 study. *Lancet Oncol.* 2018;19:347–55.
45. Heery CR, O’Sullivan-Coyne G, Madan RA, Cordes L, Rajan A, Rauckhorst M, et al. Avelumab for metastatic or locally advanced previously treated solid tumours (JAVELIN Solid Tumor): a phase 1a, multicohort, dose-escalation trial. *Lancet Oncol.* 2017;18:587–98.
46. June CH, Warshauer JT, Bluestone JA. Is autoimmunity the Achilles’ heel of cancer immunotherapy? *Nat Med.* 2017;23:540–7. Erratum in: *Nat Med.* 2017;23:1004.
47. Weissferdt A, Fujimoto J, Kalhor N, Rodriguez J, Bassett R, Wistuba II, et al. Expression of PD-1 and PD-L1 in thymic epithelial neoplasms. *Mod Pathol.* 2017;30:826–33.
48. Arbour KC, Naidoo J, Steele KE, Ni A, Moreira AL, Rekhtman N, et al. Expression of PD-L1 and other immunotherapeutic targets in thymic epithelial tumors. *PLoS One.* 2017;12:e0182665.
49. Suster D, Pihan G, Mackinnon AC, Suster S. Expression of PD-L1/PD-1 in lymphoepithelioma-like carcinoma of the thymus. *Mod Pathol.* 2018;31:1801–6.
50. Higuchi R, Goto T, Hirotsu Y, Nakagomi T, Yokoyama Y, Otake S, et al. PD-L1 expression and tumor-infiltrating lymphocytes in thymic epithelial neoplasms. *J Clin Med.* 2019;8:1833.
51. Wei YF, Chu CY, Chang CC, Lin SH, Su WC, Tseng YL, et al. Different pattern of PD-L1, IDO, and FOXP3 Tregs expression with survival in thymoma and thymic carcinoma. *Lung Cancer.* 2018;125:35–42.
52. Duan J, Liu X, Chen H, Sun Y, Liu Y, Bai H, et al. Impact of PD-L1, transforming growth factor- β expression and tumor-infiltrating CD8+ T cells on clinical outcome of patients with advanced thymic epithelial tumors. *Thorac Cancer.* 2018;9:1341–53.
53. Funaki S, Shintani Y, Fukui E, Yamamoto Y, Kanzaki R, Ose N, et al. The prognostic impact of programmed cell death 1 and its ligand and the correlation with epithelial-mesenchymal transition in thymic carcinoma. *Cancer Med.* 2019;8:216–26.
54. Katsuya Y, Fujita Y, Horinouchi H, Ohe Y, Watanabe S, Tsuta K. Immunohistochemical status of PD-L1 in thymoma and thymic carcinoma. *Lung Cancer.* 2015;88:154–9.
55. Padda SK, Riess JW, Schwartz EJ, Tian L, Kohrt HE, Neal JW, et al. Diffuse high intensity PD-L1 staining in thymic epithelial tumors. *J Thorac Oncol.* 2015;10:500–8.
56. Marchevsky AM, Walts AE. PD-L1, PD-1, CD4, and CD8 expression in neoplastic and nonneoplastic thymus. *Hum Pathol.* 2017;60:16–23.
57. Katsuya Y, Horinouchi H, Asao T, Kitahara S, Goto Y, Kanda S, et al. Expression of programmed death 1 (PD-1) and its ligand (PD-L1) in thymic epithelial tumors: Impact on treatment efficacy and alteration in expression after chemotherapy. *Lung Cancer.* 2016;99:4–10.
58. Yokoyama S, Miyoshi H, Nishi T, Hashiguchi T, Mitsuoka M, Takamori S, et al. Clinicopathologic and prognostic implications of programmed death ligand 1 expression in thymoma. *Ann Thorac Surg.* 2016;101:1361–9.
59. Tiseo M, Damato A, Longo L, Barbieri F, Bertolini F, Stefani A, et al. Analysis of a panel of druggable gene mutations and of ALK and PD-L1 expression in a series of thymic epithelial tumors (TETs). *Lung Cancer.* 2017;104:24–30.
60. Owen D, Chu B, Lehman AM, Annamalai L, Yearley JH, Shilo K, et al. Expression patterns, prognostic value, and intratumoral heterogeneity of PD-L1 and PD-1 in thymoma and thymic carcinoma. *J Thorac Oncol.* 2018;13:1204–12.
61. Hakiri S, Fukui T, Mori S, Kawaguchi K, Nakamura S, Ozeki N, et al. Clinicopathologic features of thymoma with the expression of programmed death ligand 1. *Ann Thorac Surg.* 2019;107:418–24.

62. Guleria P, Husain N, Shukla S, Kumar S, Parshad R, Jain D. PD-L1 immuno-expression assay in thymomas: study of 84 cases and review of literature. *Ann Diagn Pathol*. 2018;34:135–41.
63. Chen Y, Zhang Y, Chai X, Gao J, Chen G, Zhang W, et al. Correlation between the expression of PD-L1 and clinicopathological features in patients with thymic epithelial tumors. *Biomed Res Int*. 2018;2018:5830547.
64. Bagir EK, Acikalin A, Avci A, Gumurdulu D, Paydas S. PD-1 and PD-L1 expression in thymic epithelial tumours and non-neoplastic thymus. *J Clin Pathol*. 2018;71:637–41.
65. Ishihara S, Okada S, Ogi H, Kodama Y, Shimomura M, Tsunozuka H, et al. Programmed death-ligand 1 expression profiling in thymic epithelial cell tumors: clinicopathological features and quantitative digital image analyses. *Lung Cancer*. 2020;145:40–7.
66. Berardi R, Goteri G, Brunelli A, Pagliaretta S, Paolucci V, Caramanti M, et al. Prognostic relevance of programmed cell death protein 1/programmed death-ligand 1 pathway in thymic malignancies with combined immunohistochemical and biomolecular approach. *Expert Opin Ther Targets*. 2020;24:937–43.
67. Rouquette I, Taranchon-Clermont E, Gilhodes J, Bluthgen MV, Perallon R, Chalabreysse L, et al. Immune biomarkers in thymic epithelial tumors: expression patterns, prognostic value and comparison of diagnostic tests for PD-L1. *Biomark Res*. 2019;7:28.
68. Jakopovic M, Bitar L, Seiwerth F, Marusic A, Krpina K, Samarzija M. Immunotherapy for thymoma. *J Thorac Dis*. 2020;12:7635–41.
69. Zhao C, Rajan A. Immune checkpoint inhibitors for treatment of thymic epithelial tumors: how to maximize benefit and optimize risk? *Mediastinum*. 2019;3:35.
70. Chen HF, Wu LX, Li XF, Zhu YC, Pan WW, Wang WX, et al. PD-L1 expression level in different thymoma stages and thymic carcinoma: a meta-analysis. *Tumori*. 2020;106:306–11.
71. Yan X, Feng J, Hong B, Qian Y. The expression of PD-L1 and B7-H4 in thymic epithelial tumor and its relationship with tumor immune-infiltrating cells. *Front Oncol*. 2021;11:662010.
72. Shen L, Qian Y, Wu W, Weng T, Wang FXC, Hong B, et al. B7-H4 is a prognostic biomarker for poor survival in patients with pancreatic cancer. *Hum Pathol*. 2017;66:79–85.
73. Zheng C, Yang R. RCD24, B7-H4 and PCNA expression and clinical significance in ovarian cancer. *J BUON*. 2019;24:715–9.
74. Kim NI, Park MH, Kweon SS, Lee JS. B7-H3 and B7-H4 expression in breast cancer and their association with clinicopathological variables and T cell infiltration. *Pathobiology*. 2020;87:179–92.
75. Qi Y, Huang X, Ji C, Wang C, Yao Y. The co-inhibitory immune checkpoint proteins B7-H1(PD-L1) and B7-H4 in high grade glioma: from bench to bedside. *Transl Oncol*. 2024;39:101793.
76. Takahashi N, Zhao C, Rajan A. WT1 as an immunotherapy target for thymic epithelial tumors: a novel method to activate anti-tumor immunity. *Mediastinum*. 2019;3:11.
77. Call KM, Glaser T, Ito CY, Buckler AJ, Pelletier J, Haber DA, et al. Isolation and characterization of a zinc finger polypeptide gene at the human chromosome 11 Wilms' tumor locus. *Cell*. 1990;60:509–20.
78. Inoue K, Sugiyama H, Ogawa H, Nakagawa M, Yamagami T, Miwa H, et al. WT1 as a new prognostic factor and a new marker for the detection of minimal residual disease in acute leukemia. *Blood*. 1994;84:3071–9.
79. Oji Y, Miyoshi S, Maeda H, Hayashi S, Tamaki H, Nakatsuka S, et al. Overexpression of the Wilms' tumor gene WT1 in de novo lung cancers. *Int J Cancer*. 2002;100:297–303.
80. Loeb DM, Evron E, Patel CB, Sharma PM, Niranjana B, Buluwela L, et al. Wilms' tumor suppressor gene (*WT1*) is expressed in primary breast tumors despite tumor-specific promoter methylation. *Cancer Res*. 2001;61:921–5.
81. Oji Y, Yamamoto H, Nomura M, Nakano Y, Ikeba A, Nakatsuka S, et al. Overexpression of the Wilms' tumor gene *WT1* in colorectal adenocarcinoma. *Cancer Sci*. 2003;94:712–7.

82. Oji Y, Nakamori S, Fujikawa M, Nakatsuka S, Yokota A, Tatsumi N, et al. Overexpression of the Wilms' tumor gene *WT1* in pancreatic ductal adenocarcinoma. *Cancer Sci.* 2004;95:583–7.
83. Oji Y, Suzuki T, Nakano Y, Maruno M, Nakatsuka S, Jomgeow T, et al. Overexpression of the Wilms' tumor gene *WT1* in primary astrocytic tumors. *Cancer Sci.* 2004;95:822–7.
84. Algar EM, Khromykh T, Smith SI, Blackburn DM, Bryson GJ, Smith PJ. A *WT1* antisense oligonucleotide inhibits proliferation and induces apoptosis in myeloid leukaemia cell lines. *Oncogene.* 1996;12:1005–14.
85. Ito K, Oji Y, Tatsumi N, Shimizu S, Kanai Y, Nakazawa T, et al. Antiapoptotic function of 17AA(+)WT1 (Wilms' tumor gene) isoforms on the intrinsic apoptosis pathway. *Oncogene.* 2006;25:4217–29.
86. Jomgeow T, Oji Y, Tsuji N, Ikeda Y, Ito K, Tsuda A, et al. Wilms' tumor gene *WT1* 17AA(-)/KTS(-) isoform induces morphological changes and promotes cell migration and invasion *in vitro*. *Cancer Sci.* 2006;97:259–70.
87. Wagner N, Michiels JF, Schedl A, Wagner KD. The Wilms' tumour suppressor *WT1* is involved in endothelial cell proliferation and migration: expression in tumour vessels *in vivo*. *Oncogene.* 2008;27:3662–72.
88. Oka Y, Tsuboi A, Taguchi T, Osaki T, Kyo T, Nakajima H, et al. Induction of *WT1* (Wilms' tumor gene)-specific cytotoxic T lymphocytes by *WT1* peptide vaccine and the resultant cancer regression. *Proc Natl Acad Sci U S A.* 2004;101:13885–90.
89. Keilholz U, Letsch A, Busse A, Asemissen AM, Bauer S, Blau IW, et al. A clinical and immunologic phase 2 trial of Wilms tumor gene product 1 (*WT1*) peptide vaccination in patients with AML and MDS. *Blood.* 2009;113:6541–8.
90. Van Tendeloo VF, Van de Velde A, Van Driessche A, Cools N, Anguille S, Ladell K, et al. Induction of complete and molecular remissions in acute myeloid leukemia by Wilms' tumor 1 antigen-targeted dendritic cell vaccination. *Proc Natl Acad Sci U S A.* 2010;107:13824–9.
91. Maslak PG, Dao T, Krug LM, Chanel S, Korontsvit T, Zakhaleva V, et al. Vaccination with synthetic analog peptides derived from *WT1* oncoprotein induces T-cell responses in patients with complete remission from acute myeloid leukemia. *Blood.* 2010;116:171–9.
92. Oji Y, Inoue M, Takeda Y, Hosen N, Shintani Y, Kawakami M, et al. *WT1* peptide-based immunotherapy for advanced thymic epithelial malignancies. *Int J Cancer.* 2018;142:2375–82.
93. Ballman M, Zhao C, McAdams MJ, Rajan A. Immunotherapy for management of thymic epithelial tumors: a double-edged sword. *Cancers (Basel).* 2022;14:2060.
94. Suzuki M, Hishida T, Asakura K, Asamura H. *WT1* peptide-based immunotherapy for refractory thymic epithelial malignancies. *Mediastinum.* 2019;3:12.