

RESEARCH NOTE

Open Access



Cytotoxic T-lymphocyte-associated 4 protein expression is associated with a high international prognostic score in advanced-stage classical Hodgkin lymphoma

Flora Dameria Pangaribuan¹, Maria Francisca Ham^{2,3}, Mutiah Mutmainnah⁴ and Agnes Stephanie Harahap^{2,3*}

Abstract

Objective Twenty percent of all classical Hodgkin lymphoma (CHL) cases relapse and recur, especially in advanced stages with a high International Prognostic Score (IPS). Cytotoxic T-lymphocyte-associated protein 4 (CTLA-4) is a regulatory molecule that can inhibit the immune response and is related to tumor aggressiveness. This study aimed to determine the relationship between CTLA-4 expression in advanced-stage CHL and IPS, identifying it as a potential therapy target.

Results In advanced-stage CHL, the group with a high IPS exhibited significantly higher mean CTLA-4 expression compared to the group with a low IPS ($p=0.003$). The group with Hb level < 10.5 g/dl, leukocyte count $> 15,000/\mu\text{L}$, lymphocyte count $< 8\%$, albumin level < 4 g/dl, and stage 4 exhibited higher CTLA-4 expression than the other group, although only leukocyte count and stage showed statistical significance ($p=0.004$ and $p=0.020$). Mean CTLA-4 expression was 239.84 ± 76.36 for nodular sclerosis, 293.95 ± 147.94 for mixed cellularity, 271.4 ± 23.56 for lymphocyte depleted, and 225.2 for lymphocyte-rich subtypes. The results suggest that CTLA-4 expression is associated with adverse prognostic factors in the IPS for advanced-stage CHL, supporting the notion that immune checkpoints play a role in cancer progression.

Keywords Advanced-stage, Cytotoxic T-lymphocyte-associated protein 4, Hodgkin lymphoma, International prognostic score

Introduction

Classical Hodgkin lymphoma (CHL) is a lymphoid malignancy with a challenging diagnosis and treatment. Although most CHL tumours show good prognosis, with an 80% recovery rate, some cases present with relapse and usually are diagnosed at advanced stages [1]. The International Prognostic Score (IPS) uses risk factors in an advanced-stage CHL case to determine treatment plans and predict relapse or recurrence. These risk factors include age, sex, stage, haemoglobin (Hb), leukocyte count, lymphocyte count and serum albumin level [2–5]. Advanced-stage CHL with a higher IPS has a higher risk

*Correspondence:

Agnes Stephanie Harahap
agnes.stephanie01@ui.ac.id

¹ Anatomical Pathology Department, Persahabatan Hospital, Jakarta 13230, Indonesia

² Anatomical Pathology Department, Faculty of Medicine Universitas Indonesia, Dr. Cipto Mangunkusumo National Central General Hospital, Jakarta 10430, Indonesia

³ Faculty of Medicine Universitas Indonesia, Human Cancer Research Center-Indonesian Medical Education and Research Institute, Jakarta 10430, Indonesia

⁴ Faculty of Medicine, Universitas Muhammadiyah Palembang, Palembang 30263, Indonesia



© The Author(s) 2024. **Open Access** This article is licensed under a Creative Commons Attribution 4.0 International License, which permits use, sharing, adaptation, distribution and reproduction in any medium or format, as long as you give appropriate credit to the original author(s) and the source, provide a link to the Creative Commons licence, and indicate if changes were made. The images or other third party material in this article are included in the article's Creative Commons licence, unless indicated otherwise in a credit line to the material. If material is not included in the article's Creative Commons licence and your intended use is not permitted by statutory regulation or exceeds the permitted use, you will need to obtain permission directly from the copyright holder. To view a copy of this licence, visit <http://creativecommons.org/licenses/by/4.0/>. The Creative Commons Public Domain Dedication waiver (<http://creativecommons.org/publicdomain/zero/1.0/>) applies to the data made available in this article, unless otherwise stated in a credit line to the data.

of recurrence or relapse [6]. Advanced-stage CHL with IPS 0–3 shows 83% overall survival and IPS ≥ 4 59% [7].

CHL exhibits characteristic morphological features marked by Reed-Sternberg cells or Hodgkin cells with beneficial enrichment of inflammatory cells such as lymphocytes, eosinophils, histiocytes and other immune cells as the background, constituting the tumour micro-environment (TME) [8, 9]. The TME is constructed by the interaction of tumour cells and normal cells, such as stromal cells, fibroblasts, immune cells, and the extracellular matrix. The tumour cells and TME of CHL produce molecules that induce cell proliferation, survival, and immune escape to promote tumour cell growth [10]. A significant component of the TME in CHL is cytotoxic T-lymphocyte-associated protein 4 (CTLA-4), a regulatory molecule that can inhibit the immune response, therefore decreasing the ability of the immune system to destroy tumour cells and is related to tumour aggressiveness. Expression and interaction of CTLA-4 and the TME are thought to correlate with the stage of CHL, as they can reflect the extent of immune evasion by the tumour [11, 12].

Therapeutic management of CHL continues to develop. One of the new therapies to combat relapse/recurrent CHL is through the immune system regulation pathway and immunophysiological interaction in the TME, such as use of immune checkpoint inhibitors (ICIs). This therapy aims to restore the patient's natural immune response against tumour cells. CTLA-4 inhibitors are ICIs that have a promising future in advanced-stage CHL therapy [13, 14]. Studies to date regarding the prognosis of malignancy cases treated with CTLA-4 show differing conclusions. The connection between CTLA-4 and the IPS in CHL is part of a broad landscape in immune system interactions. CTLA-4 can influence the behaviour of CHL, which is reflected in prognostic tools such as the IPS. This study will elucidate the immunoexpression profile of CTLA-4 in advanced-stage CHL and its correlation with the IPS.

Material and methods

Ethics

This study was approved by The Ethics Committee of Universitas Indonesia, Jakarta, Indonesia, in accordance with the Declaration of Helsinki with protocol number KET-127/UN2.F1/ETIK.PPM.00.02/2022. Informed consent was waived by The Ethics Committee of Universitas Indonesia, Jakarta, Indonesia (letter number ND-884/UN2.F1/ETIK/PPM.00.02/2023).

Sample collection

This was a retrospective study that was performed at the Anatomical Pathology Department, Faculty of Medicine

Universitas Indonesia, Dr. Cipto Mangunkusumo Hospital, Jakarta, Indonesia. A total of 40 advanced-stage CHL cases available on the Anatomical Pathology Department archives from the 1st of January 2013 to the 30th of September 2021 were enrolled by using consecutive sampling techniques. All cases were confirmed by comprehensive immunohistochemistry (IHC) examination. *Cotswolds-modified* Ann Arbor was used as the staging system. Stage 2 with B-symptoms and/or bulky disease, stage 3, and stage 4 are categorized as advanced stage. The IPS is determined by factors such as age, sex, Hb value, leukocyte count, lymphocyte count, albumin value, and clinical stage. The samples were divided into two groups: advanced-stage CHL with low IPS (cases with 0–3 IPS factors) and high IPS (cases with > 4 IPS factors). Patients' clinical data, including age, sex, laboratory findings, imaging results, therapy history, and clinical stage, were taken from medical records.

CTLA-4 immunohistochemistry examination

Formalin-fixed paraffin-embedded tissue blocks (FFPE) was cut (3–5 μm), placed on a glass slide, and heated above a slide warmer at 56.5–60.0 $^{\circ}\text{C}$ for 60 min. Incubation was performed in Tris-EDTA with a decloaking chamber followed by submerging in peroxidase blocking solution and protein block solution. The slides were then incubated with an anti-CTLA-4 primary antibody (Biocare Medical, LLC. UMAB249 clone, CA, USA) at a 1:150 dilution for 1 h. Postprimary (Rabbit anti mouse IgG ($< 10 \mu\text{g}/\text{mL}$) in 10% (v/v) animal serum) and secondary (NovolinkTM Polymer. Anti-rabbit Poly-HRP-IgG ($< 25 \mu\text{g}/\text{mL}$) containing 10% (v/v) animal serum in tris-buffered saline/0.09% ProClinTM 950) antibodies by Novolink detection kit[®] RE7140-CE (Leica Biosystems, Newcastle, UK) were then added for 30 min. Negative and positive controls were included with each preparation. Negative controls were performed by omitting the antibody step in each case. The positive control for CTLA-4 was tonsil FFPE and taken from the laboratory archive.

Evaluation of CTLA-4 expression was performed by researchers blinded to the clinical data. Lymphocytes and histiocytes that showed brown staining in the cell membrane or cytoplasm were considered positive. Photographs were taken of five well-stained high-power fields and the average number of positive cells per field was obtained and analysed (Fig. 1).

Data analysis

All data were analysed using SPSS software version 25.0. Continuous variables are presented as the mean due to a normal data distribution using the Kolmogorov-Smirnov test. The statistical analysis was performed using

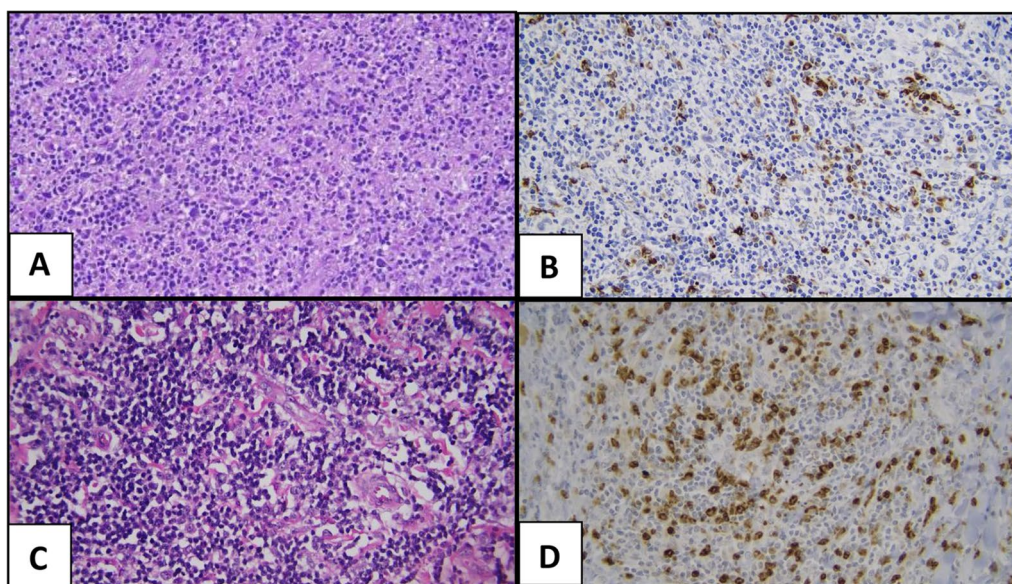


Fig. 1 Expression of CTLA-4 in Low and High IPS Groups of Advanced-stage classical Hodgkin Lymphoma. **A** Morphology of a case with low IPS (HE 400x). **B** Expression of CTLA-4 in the low IPS case (IHC 400x). **C** Case with high IPS (HE 400x). **D** Expression of CTLA-4 in the high IPS case (IHC 400x)

appropriate bivariate tests (unpaired T test). A p value of 0.05 was used as the threshold for statistical significance.

Results

This study included 40 advanced-stage CHL cases involving 21 males (52.5%) and 19 females (47.5%). Age ranged from 12 to 70 years old: 35 (87.5%) of the patients were ≤ 45 years old. Nodular sclerosis (NS) was the predominant CHL subtype (62.5%), followed by mixed cellularity (MC) (27.5%), lymphocyte depleted (LD) (7.5%), and lymphocyte rich (LR) (2.5%). The tumor location were in the lymph nodes and extra nodal sites such as mediastinum, lung, and spleen. The predominant lymph nodes affected were the cervical region. The cases were divided into advanced-stage CHL with low IPS score and advanced-stage CHL with high IPS score. The high IPS score group showed lower Hb levels, lymphocyte counts, and albumin levels and higher leukocyte counts than the low IPS score group. It was notable that more advanced disease stages increased as the IPS increased, as highlighted by none at stage 2 and 80% at stage 4 in the high IPS group, in contrast to 15% at stage 2 and a lower number at stage 4 (45%) cases in the low IPS group. The demographic and clinicopathologic characteristics of each group in this study are presented in Table 1.

Table 2 reveals that the average CTLA-4 expression in the ≤ 45 years old, female, Hb < 10.5 g/dl, leukocyte $> 15,000/\mu\text{L}$, lymphocyte $< 8\%$, and albumin < 4 g/dl group was higher than that in the opposite group (Table 2). These cut-off points were determined

according to the IPS criteria. Crucially, there was notably higher CTLA-4 expression in stage 4 disease than in disease less than stage 4, suggesting that CTLA-4 expression might correlate with more advanced disease. This result was supported by the fact that the advanced-stage CHL with high IPS group had a significantly higher average CTLA-4 expression than the low IPS group (p value 0.003) (Fig. 2).

A detailed breakdown of CTLA-4 expression in CHL subtypes is presented in Table 3. The findings revealed that CTLA-4 expression was notably highest in the MC subtype, followed by the LD, NS, and LR subtypes. A scatter plot of CHL subtypes displayed in Fig. 3.

Discussion

Although CHL generally has good prognosis, some CHL cases may be refractory to treatment and are usually diagnosed at an advanced stage and prone to recurrence. The IPS has a crucial role in CHL management by including several prognostic indicators to stratify and modify therapy and predict relapse or recurrence. Each adverse prognostic indicator that is present is given a point, and the total score correlates with the expected CHL outcome. Advanced-stage CHL with higher IPS will have poorer prognosis and higher risk of recurrence or relapse. In contrast, lower IPS indicates a more favourable prognosis; hence, each patient can be treated accordingly [6].

CTLA-4 is a regulatory molecule that inhibits T-cell activation. The correlation between IPS and CTLA-4 expression is important as one of the putative ICIs

Table 1 Demographic and clinicopathologic characteristics of advanced-stage CHL patients

Variable	Advanced-stage CHL	
	Low IPS N = 20	High IPS N = 20
Age (years, mean ± SD)	32.55 ± 14.13	30.80 ± 13.63
Sex, n (%)		
Male	9 (45%)	12 (60%)
Female	11 (55%)	8 (40%)
Subtype, n (%)		
Nodular sclerosis	13 (65%)	12 (60%)
Mixed cellularity	5 (25%)	6 (30%)
Lymphocyte depleted	1 (5%)	2 (10%)
Lymphocyte rich	1 (5%)	0
Haemoglobin (g/dl, median (min–max))	11.6 (2.7–14.1)	9.5 (7.4–14.6)
Leukocyte (/μL, mean ± SD)	11.216 ± 5320.96	21.659.8 ± 15.310.17
Lymphocyte (% , median (min–max))	13.5 (6.1–41.4)	4.3 (1.3–12.2)
Albumin (g/dl, mean ± SD)	3.67 ± 0.53	2.75 ± 0.52
Stage, n (%)		
2	3 (15%)	0
3	8 (40%)	4 (20%)
4	9 (45%)	16 (80%)

IPS: International Prognostic Score; CHL: Hodgkin lymphoma; Min: minimum; Max: maximum; SD: standard deviation

because it may shed light on the mechanism of immune evasion disease outcomes. It has been observed that CTLA-4 is expressed differentially in patients with variable IPS levels, which may be related to the disease's aggressiveness. There seems to be a correlation between higher CTLA-4 expression and higher IPS, which suggests that higher IPS CHL may have more severe immune suppression, indicating disease severity and progression [2–5, 7]. In this study, only leukocytes count and tumor stage that were significantly associated with CTLA-4 expression with p-values 0.004 and 0.020, respectively.

The MC and LD subtypes had the two highest average CTLA-4 expression, 293.9 ± 147.9 cells/FOV and 271.4 ± 23.6 cells/FOV, respectively, while the LR subtype had the lowest average CTLA-4 expression (225.2 cells/FOV). These variations may reflect the distinct histological composition and TME of each subtype. Some studies have reported that the MC and LD subtypes of CHL generally have worse prognosis than the NS and LR subtypes. This is because the MC and LD subtypes are more common in patients of older ages, males, and advanced-stage patients and are associated with HIV and EBV infection. The MC subtype showing the highest CTLA-4 expression presumably correlates with a more immunosuppressive component due to the higher diversity of TME

Table 2 CTLA-4 expression in advanced-stage CHL and its relationship with IPS

Variable	CTLA-4 expression (mean ± SD)	p value
Age		
≤ 45 years old	258.59 ± 98.49	0.757
> 45 years old	243.64 ± 115.25	
Sex		
Female	271.2 ± 107.67	0.387
Male	243.62 ± 91.56	
Haemoglobin		
< 10.5 g/dl	285.85 ± 115.31	0.077
≥ 10.5 g/dl	230.36 ± 75.46	
Leukocyte		
≤ 15,000 μ/L	219.11 ± 74.31	0.004
> 15,000 μ/L	307.6 ± 107.79	
Lymphocyte		
< 8%	279.62 ± 122.07	0.107
≥ 8%	228.74 ± 51.68	
Albumin		
< 4 g/dl	261.06 ± 103.57	0.414
≥ 4 g/dl	217.65 ± 24.07	
Stage		
< 4	210.23 ± 84.94	0.020
4	284.62 ± 98.12	
IPS		
Low IPS	210.11 ± 52.41	0.003
High IPS	303.33 ± 113.52	

Unpaired T test; CTLA-4: cytotoxic T-lymphocyte-associated protein 4; SD: standard deviation

components that may modify the immune response. In contrast, the LR subtype consists of a more homogenous lymphocyte background, suggesting a less suppressed immune response, scarce tumour cells, and better prognosis, as reflected by lower CTLA-4 expression. The NS and LR subtypes are more common in cases involving early stages, young age, and low tumour cell abundance [2, 15, 16].

This study found that extranodal lesions were more common in the high IPS group. The high IPS group also tended to have more stage 4 disease. In line with the increase in CHL stage, CTLA-4 expression was increased and marked more aggressive and extensive disease. This implies that immunological checkpoints such as CTLA-4 are upregulated when CHL progresses, potentially aiding tumour cells in immune evasion and enabling them to multiply and spread with reduced immune system interference [17–21]. In this study, stage 4 cases had significantly higher average CTLA-4 expression than cases

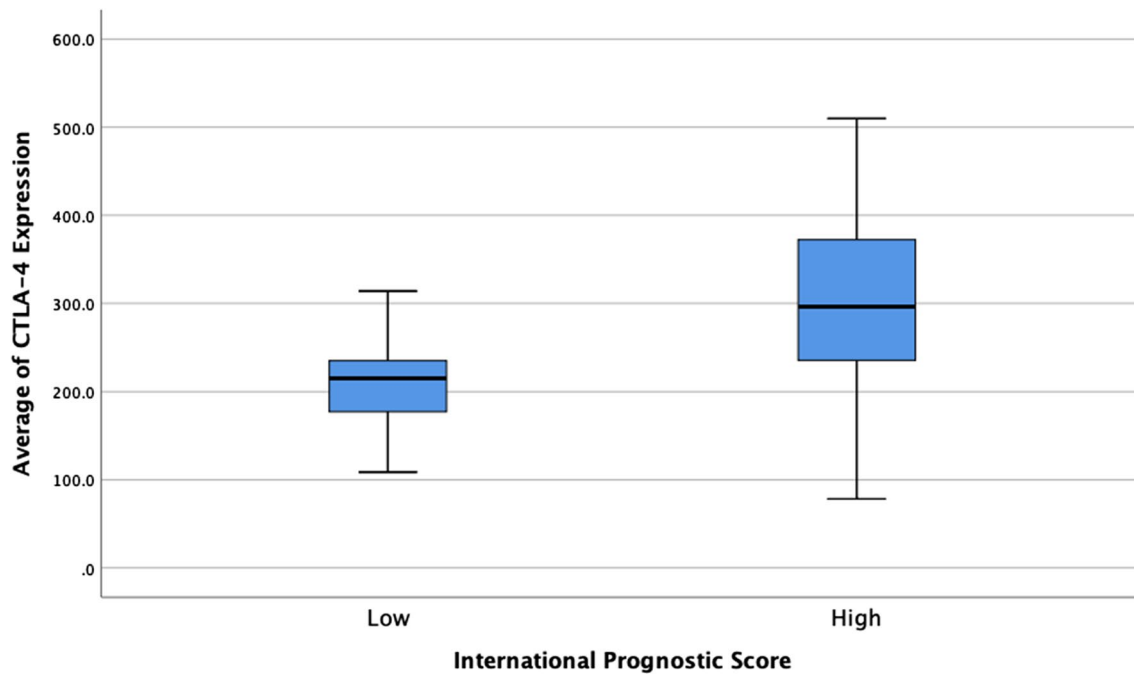


Fig. 2 CTLA-4 expression in the low and high IPS group

Table 3 Average CTLA-4 expression in every CHL subtype

Variable	Average of CTLA-4 expression (mean ± SD)
CHL subtypes	
Nodular sclerosis	239.84 ± 76.36
Mixed cellularity	293.95 ± 147.94
Lymphocyte depleted	271.4 ± 23.56
Lymphocyte rich	225.2

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

with staging below 4, with 284.6 ± 98.1 cells/FOV and 210.2 ± 84.9 cells/FOV, respectively.

Advanced-stage CHL with low IPS was found to have significantly lower CTLA-4 expression than the high IPS group (210.1 ± 52.4 cells/FOV vs. 303.3 ± 113.5 cells/FOV). This finding is similar to the study by Chen Y et al. [22], in which there was a significant difference between the average optical density of CTLA-4 in high- and low-risk groups of DLBCL patients ($p < 0.05$). High IPS and increased expression of CTLA-4 in advanced-stage CHL align with the theory of immunosuppressive status in more aggressive disease states. As an

ICI, CTLA-4 decreases the immune response, and its increased expression can help the tumour to modify the immune response, hence causing a greater disease burden [17–21].

In conclusion, our findings suggest that CTLA-4 expression correlates with the worst prognostic factors in the IPS for advanced-stage CHL, supporting the notion that immune checkpoints play a role in cancer progression. This result supports the potential use of ICI inhibitors in CHL patients with high CTLA-4 expression and adverse IPS factors as personalized treatment based on immunological profiles. Additionally, it may lead to integration of immunological markers into prognostic scoring systems for CHL in the future.

Limitations

The study was limited by the small sample sizes across each subtype. It did not employ additional modalities such as flow cytometry or Western blot, nor did it include double staining of IHC markers for CTLA-4 and other markers of T lymphocytes, macrophages, and granulocytes. Additionally, the study had limited clinical data, such as survival outcomes. We also suggest that future studies use larger sample sizes to analyze subgroups based on histopathology and other factors that encompass the IPS.

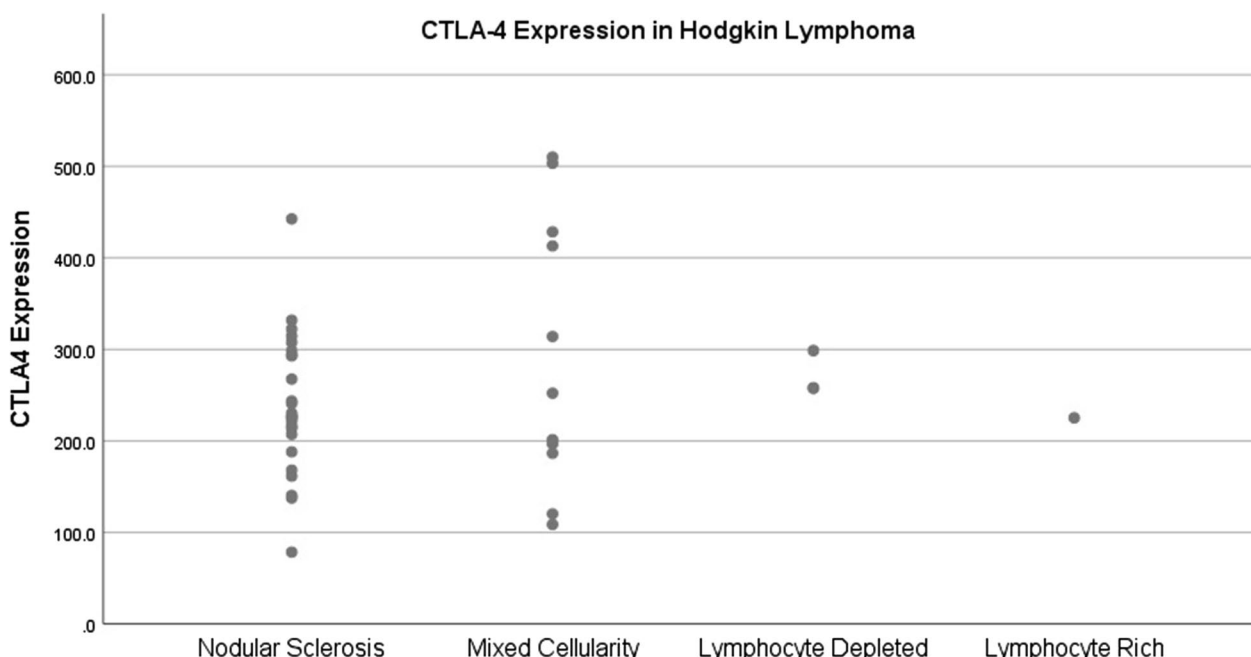


Fig. 3 A scatter plot of CTLA-4 expression in CHL subtypes

Abbreviations

Ccl-20	C–C Motif Chemokine Ligand 20
Ccl-5	C–C Motif Chemokine Ligand 5
CHL	Classical Hodgkin Lymphoma
CTLA-4	Cytotoxic T-Lymphocyte-Associated Protein 4
EBV	Epstein–Barr virus
FFPE	Formalin-fixed paraffin block-embedded tissue
Hb	Haemoglobin (Hb)
HIV	Human immunodeficiency virus
ICIs	Immune checkpoint inhibitors
IHC	Comprehensive Immunohistochemistry
Il-10	Interleukin 10
Il-6	Interleukin 6
IPS	International Prognostic Score
LD	Lymphocytes depleted
LR	Lymphocyte rich
MC	Mixed cellularity
NLPHL	Nodular lymphocytic predominant Hodgkin lymphoma
NS	Nodular sclerosis
Tgf-β	Transforming Growth Factor Beta
TME	Tumour microenvironment
Tnf-α	Tumour Necrosis Factor Alpha

Author contributions

All authors designed the study. Flora Dameria Pangaribuan and Agnes Stephanie Harahap collected the data. Flora Dameria Pangaribuan, Maria Francisca Ham, Mutiah Mutmainnah, and Agnes Stephanie Harahap developed the methodology and performed the data analysis, results presentation, and interpretation. Flora Dameria Pangaribuan, Mutiah Mutmainnah, and Agnes Stephanie Harahap drafted the manuscript. All authors reviewed and approved the final manuscript.

Funding

No special funding was received for the study.

Availability of data and materials

The datasets generated and/or analyzed during the current study are available from the corresponding author upon reasonable request.

Declarations

Ethics approval and consent to participate

This study was approved by The Ethics Committee of Universitas Indonesia, Jakarta, Indonesia, with protocol number KET-127/UN2.F1/ETIK/PPM.00.02/2022. Informed consent was waived by The Ethics Committee of Universitas Indonesia, Jakarta, Indonesia (letter number ND-884/UN2.F1/ETIK/PPM.00.02/2023).

Consent for publication

Not applicable.

Competing interests

The authors declare that they have no competing interests.

Received: 3 January 2024 Accepted: 1 July 2024

Published online: 08 July 2024

References

1. Calabretta E, d'Amore F, Carlo-Stella C. Immune and inflammatory cells of the tumor microenvironment represent novel therapeutic targets in classical hodgkin lymphoma. *Int J Mol Sci.* 2019;20(21):1–19.
2. Shanbhag S, Ambinder RF. Hodgkin lymphoma: a review and update on recent progress. *CA Cancer J Clin.* 2018;68(2):116–32.
3. Hoppe RT, Advani RH, Ai WZ, Ambinder RF, Armand P, Bello CM, et al. Hodgkin lymphoma, version 2. *JNCCN J Natl Compr Cancer Netw.* 2020;18(6):755–81.
4. Hasenclever D, Diehl V, Armitage JO, Assouline D, Björkholm M, Brusa-Molinolo E, et al. A prognostic score for advanced hodgkin's disease. *N Engl J Med.* 1998;339(21):1506–14.
5. Klimm B, Goergen H, Fuchs M, Von Tresckow B, Böll B, Meissner J, et al. Impact of risk factors on outcomes in early-stage hodgkin's lymphoma: an analysis of international staging definitions. *Ann Oncol.* 2013;24(12):3070–6.

6. Potre O, Pescaru M, Sima A, Ionita I, Tudor R, Borsi E, et al. Evaluation of the relapse risk and survival rate in patients with hodgkin lymphoma: a monocentric experience. *Med*. 2021;57(10):1–7.
7. Moccia AA, Donaldson J, Chhanabhai M, Hoskins PJ, Klasa RJ, Savage KJ, et al. International prognostic score in advanced-stage hodgkin's lymphoma: altered utility in the modern era. *J Clin Oncol*. 2012;30(27):3383–8.
8. Eberle FC, Mani H, Jaffe ES. Histopathology of hodgkin's lymphoma. *Cancer J*. 2009;15(2):129–37.
9. Piris MA, Medeiros LJ, Chang KC. Hodgkin lymphoma: a review of pathological features and recent advances in pathogenesis. *Pathology*. 2020;52(1):154–65.
10. Aldinucci D, Celegato M, Casagrande N. Microenvironmental interactions in classical hodgkin lymphoma and their role in promoting tumor growth, immune escape and drug resistance. *Cancer Lett*. 2016;380(1):243–52.
11. Guo Q, Huang F, Goncalves C, Del Rincón SV, Miller WH. Translation of cancer immunotherapy from the bench to the bedside. *Adv Cancer Res*. 2019;143:1–62.
12. Buchbinder E, Hodi FS. Cytotoxic T lymphocyte antigen-4 and immune checkpoint blockade. *J Clin Invest*. 2015;125(9):3377–83.
13. Meti N, Esfahani K, Johnson N. The role of immune checkpoint inhibitors in classical hodgkin lymphoma. *Cancers (Basel)*. 2018;10(6):1–15.
14. Houot R, Merryman RW, Morschhauser F. Total immunotherapy for hodgkin lymphoma. *Lancet Haematol*. 2020;7(9):e629–30.
15. Mani H, Jaffe ES. Hodgkin lymphoma: an update on its biology with new insights into classification. *Clin Lymphoma Myeloma*. 2009;9(3):206–16.
16. Kaseb H, Babiker HM. Hodgkin lymphoma. *StatPearls: Treasure Island*; 2023.
17. Zhao Y, Yang W, Huang Y, Cui R, Li X, Li B. Evolving roles for targeting CTLA-4 in cancer immunotherapy. *Cell Physiol Biochem*. 2018;47(2):721–34.
18. Grosso JF, Jure-Kunkel MN. CTLA-4 blockade in tumor models: an overview of preclinical and translational research. *Cancer Immun*. 2013;13:660–9.
19. Rudd CE, Taylor A, Schneider H. CD28 and CTLA-4 coreceptor expression and signal transduction. *Immunol Rev*. 2009;229(1):12–26.
20. Buchbinder EI, Desai A. CTLA-4 and PD-1 pathways: similarities, differences, and implications of their inhibition. *Am J Clin Oncol*. 2016;39(1):98–106.
21. Rowshanravan B, Halliday N, Sansom DM. CTLA-4: a moving target in immunotherapy. *ASH*. 2018;131(1):58–67.
22. Chen Y, Li M, Cao J, Cai G, Li X, Liu Y, et al. CTLA-4 promotes lymphoma progression through tumor stem cell enrichment and immunosuppression. *Open Life Sci*. 2021;16(1):909–19.

Publisher's Note

Springer Nature remains neutral with regard to jurisdictional claims in published maps and institutional affiliations.