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Evaluation of the expression of CD200 and CD56 in CD34-positive adult acute myeloid leukemia and its effect on the response to induction of chemotherapy

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Abstract:

BACKGROUND: Acute myeloid leukemia (AML) is characterized by an excess number of myeloid cells in the marrow with maturation arrest and infiltration of bone marrow (BM) and other tissues by myeloblasts, resulting in BM failure.

OBJECTIVES: The main goal of the present study is to investigate CD200 and CD56 aberrant expression in CD34-positive blasts, in newly diagnosed adult AML patients and their relation with the clinical and hematological parameters, as well as to identify their prognostic significance after induction therapy.

MATERIALS AND METHODS: This was a prospective cross-sectional study on thirty patients with newly diagnosed AML, who were tested for the expression of CD200, CD56 using multicolor flow cytometry and re-evaluated after induction therapy regimen.

RESULTS: CD200 and CD56 were aberrantly expressed in 53.3% and 20.0%, respectively, while coexpression of both markers was observed in 13.3%. Interestingly, both markers were expressed more in monocytic subtypes. Significantly, the induction failure in CD200 + patients was 75%, while it was 66.7% in CD56+ patients.

CONCLUSION: The findings of this research provide insights that CD200 and CD56 were closely related to bad prognostic parameters, including high total white blood cell count, low platelet's counts, and low response to induction therapy.

Keywords:

Acute myeloid leukemia, CD200, CD34, CD56, flow cytometry

Malignant proliferation of myeloid progenitor cells that gradually replace normal hematopoiesis in the bone marrow (BM) is known as acute myeloid leukemia (AML).^[1]

AML accounts for approximately 80% of acute leukemia (AL) in an adult. Clinically, patients presented with signs or symptoms of BM failure although signs of leukostasis or neurological dysfunction may be present.^[2] The etiology of majority of AML cases remains unexplained.

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However, many hematological disorders such as myeloproliferative or myelodysplastic disorder and aplastic anemia may represent predisposing factors in minority of cases.^[3]

Genetically, AML resulted from multistep of collaborating mutations in hematopoietic precursors; two hits model is implicated in leukemogenesis: Class 1 includes mutations which activate signal transduction pathways resulting in enhanced proliferation and/or survival of leukemic progenitor cells.

Class 2 include mutations that affect transcription factors or components of the transcriptional coactivation complex.^[4]

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Aberrant immunophenotyping expression

Aberrant immunophenotyping expression is a deviation from the normal pattern of antigen expression.^[5] Abnormal antigenic expression in AL can be grouped into four basic categories:^[6]

- 1. Abnormally increased or decreased levels of expression (intensities) of antigens
- 2. Asynchronous antigen expression, i.e., expression of antigens normally expressed by the cell type or lineage but at an inappropriate time during maturation)
- 3. Abnormally homogeneous expression of one or more antigens by a population that normally exhibits more heterogeneous expression
- 4. Gain of antigens not normally expressed by cell type or lineage such as CD7 expression on myeloid blasts cells.

CD200

it is a cell surface glycoprotein, expressed normally on broad range of cells such as dendritic cells, macrophages, mast cells, neutrophils, and also B- and T-cells.^[7,8] CD200 plays vital role in maintaining self-tolerance and autoimmunity.

Its binding with CD200R leads to induce an immunosuppressive signal which favors the tumor growth. Initially, CD200 was reported in chronic lymphocytic leukemia (CLL), where it has a role in differential diagnosis of mantle cell lymphoma.^[9] CD200 is normally not present on plasma cells; on the other hand, plasma cell myelomas (PCMs) expressed CD200 in more than 70%, while loss of CD200 expression in PCM may be associated with more clinically aggressive disease.^[10]

In AML, CD200 aberrant expression has been proposed as bad prognostic factor, may be due to suppression of natural killer (NK) activity, its overexpression associated with a worse prognosis even with the presence of favorable biological markers, such as Flt3 wild-type, mutated nucleolar protein nucleophosmin and negative expression of CD34 and CD56.^[9]

CD56

it is cell surface glycoprotein, expressed generally in lymphocytes, monocytes, and NK cytotoxic lymphocytes (NK cells), which plays an important role in innate immune system. Plasmocytes normally are CD56-negative, and when it is positive, it can be applied to distinguish multiple myeloma from reactive plasmocytes or monoclonal gammopathies of undetermined significance.^[11,12]

CD56 antigen expression in AML with favorable prognosis such as in t (8;21), may impact on complete remission duration and extramedullary manifestations.^[13]

Incidence of central nervous system disease is highly associated with CD56 antigen especially in patients with AML, myeloid/NK AL, acute lymphocytic leukemia, and lymphoma. This can be attributed to the important function of CD56 in neuronal growth and migration through cell-to-cell adhesion.^[12]

CD34

it is a glycoprotein, expressed commonly on hematopoietic progenitor cells; it acts as a cell–cell adhesion factor.^[14] CD34 antigen is present on both myeloid and lymphoid leukemic cells.^[15]

It has an important role in identification of blasts in hypoplastic marrows to differentiate myelodysplastic syndrome (CD34+ blasts present) and AL from aplastic anemia (no blasts/low marrow CD34+ cells) and also to determine disease processes or vascularization of tumors.^[16]

Materials and Methods

This cross-sectional study was conducted on thirty adults newly diagnosed *de novo* AML patients from January 2017 to June 2017. This study was approved by the Ethical Committee at the College of Medicine/Al-Nahrain University. Patients were taken from the Hematology Department of Baghdad Teaching Hospital of the Medical City and Al-Imamein Kadhimein Medical City.

Written consent was taken from the patients before starting the study. Diagnosis was based on morphology and cytochemistry (Sudan Black and periodic acid–Schiff) of the peripheral blood (PB) and/or BM aspiration samples by an expert hematopathologist; suspected cases were sent for flow cytometric study to confirm diagnosis. 2.5 ml of ethylenediaminetetraacetic acid blood specimens was transferred to a cool box (within 6 h) to a private laboratory to investigate the expression of surface marker antigens (CD200, CD56, and CD34) using four-color flow cytometer (Partec Cyflow[®] Cube 6, Germany).

For gating, we depend on forward scatter/side scatter gate. The device software is based on the Windows TM operating system for multiparametric data acquisition, display, data analysis, and instrument control.^[17] Antigen expression was considered to be positive when the percentage of positive blast cells was $\geq 20\%$. After 3–4 weeks, PB and BM aspiration were collected from the patients to assess their response after induction therapy.

All patients were evaluated for CR achievement. CR was defined by Cheson *et al.*;^[18] patients were classified into CR (BM blasts <5%; absence of blasts with Auer rods; absolute neutrophil count >1.0 × 10^9 /L; and platelet count >100 × 10^9 /L), resistant disease (persistent

leukemia by blood and/or BM examination), and death during induction. In the present study, M3 subtype of the French-American-British (FAB) classification was excluded to decrease bias due to CD34 expression.

Statistical analysis was performed using Microsoft Excel and GraphPad Prism 6. Categorical data formulated as count and percentage. Chi-square test describes the association of these data. Numerical data with normal distribution was described as mean and standard deviation (SD), independent sample t-test used for comparison between two groups. While for abnormal distribution, Mann–Whitney test used for comparison between two groups. The lower level of accepted statistical significant difference is bellow or equal to 0.05.

Results

In the current study, functional connectivity was done to detect the aberrant expression of CD200 and CD56 in adult AML patients. The mean age of the patients was 40.93 ± 15.63 SD years, with a median of 38 years; the range was 17–76 years. Fifty-three percent of the cases were in the age group of 21–40 years. AML was observed more in males than in females with male:female ratio of 3:2. The two most frequent signs were pallor and fever (60% and 46.7%, respectively).

The classification was applied after excluding M3 subtype; M2 subtype was the major portion of the AML subtypes (36.7%), followed by M5 (26.7%), M1 (20%), M4 (13.3%), and M7 (3.3%). Regarding aberrant expression, CD200 was expressed in 16 out of 30 cases (53.3%) and CD56 was expressed in 6 out of 30 cases (20%), whereas simultaneous co expression of both markers was detected in 4 out of 30 cases (13.3%) [Figure 1].

Regarding relation of both aberrant markers with classical prognostic parameters, high white blood cell (WBC) above 30×10^{9} /L and moderate low platelet count ($33.19 \pm 13.46 \times 10^{9}$ /L) were significantly related to aberrant CD200 expression. Similarly, CD56 expression was significantly related to high WBC above 30×10^{9} /L and very low platelet count ($25.83 \pm 7.19 \times 10^{9}$ /L) as well as high PB and BM blast percentage [Tables 1].

CD200 expressed more in cases with monocytic differentiation, it was found in M5 as 6 out of 8 cases (75%) and in all four M4 cases (100%). On the other hand, CD56 was expressed in monocytic subtypes in M5 as 2 out of 8 cases (25%) and in M4 as 2 out of 4 cases (50%).

In all AML patients included in this study, CR was achieved in 14/30 patients (46.6%) at the end of induction therapy. Twelve out of 16 patients (75%) who express CD200 did not respond to induction therapy (P < 0.005).

On the other hand, four out of six cases (66.6%) who expressed CD56 did not respond to induction therapy (P > 0.005) [Table 2].

Discussion

The mean age and range of AML patients included in this study were 40.93 \pm 15.63 (mean \pm SD) years and 17–67 years; those results were in accordance to Iraqi^[19-21] and non-Iraqi studies.^[22]

Furthermore, AML was observed more in male than in female, with a male:female ratio of 3:2, which was in consistent with that reported by the Iraqi Ministry of Health.^[23] The most common presenting features for adult AML were pallor and fever, while lymphadenopathy and gingival hypertrophy were the least presenting features. Those findings were in agreement with Iraqi^[24] and other worldwide studies.^[25]

Moreover, M2 subtype was found to be the most common FAB subtype (36.7%) in accordance with Alwan *et al.*,^[24] who found that M2 (38%) most frequent subtype.

The present study revealed that CD200 was expressed in 53.3%, approximately similar to Damiani *et al.*^[9] and Tiribelli *et al.*,^[26] who reported that CD200 expression was found in 49% and 48%, respectively. However, CD56 was expressed in 20% of cases, which was comparable to the result obtained from El-Sissy *et al.*,^[27] who reported that CD56 was expressed in (20.3%).

CD200 aberrantly expressed more in monocytic subtypes (M4–M5) which was 10 out of 12 cases (83.3%),^[28,29] followed by M1–M2 subtypes as 6 out of 17 cases (35.2%).

Regarding CD56, it was more expressed in monocytic subtype as four out of 12 cases (33.3%); this was in agreement with Di Bona *et al.*^[30] and Graf *et al.*^[31]

Regarding extramedullary manifestations which include (hepatosplenomegaly, lymphadenopathy, and gingival hypertrophy), CD200 was positively expressed in 6 out of 9 cases.



Figure 1: Expression of CD200 and CD56 in AML Patients

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To the best of our knowledge, no previous studies addressed the correlation of the extramedullary infiltrates with CD200 expression.

While CD56 expression was positively expressed in 4 out of 9 cases, this result was consistent with that obtained by Chang *et al.*^[32] Since both markers were expressed more in monocytic subtypes with the presence of extramedullary manifestations, they will confirm the unfavorable role of both markers.

However, this study showed that there was no significant correlation between both markers and patient's age whether above or below 60 years and gender.^[26]

Regarding the hematological parameters, CD200 was observed more in cases with WBC count $\geq 30 \times 10^{9}/L$ than those $<30 \times 10^{9}/L$ in comparable to Damiani *et al.*,^[9] Furthermore, CD200 was significantly expressed in cases with low platelets count; this was in agreement with other study on hematological malignancies such as CLL.^[33]

The present work did not reveal any significant differences in CD200 expressions with respect to the PB and BM blasts percentage. To the best of our knowledge, no previous studies addressed the correlation of the PB blasts and BM blasts percentage with CD200 expression.

Variation in the relation between CD200 expression and hematological parameters can be explained by the direct interaction between CD200-positive cancer cells and myeloid cells, which can be attributed to the high expression of CD200R on tumor-associated myeloid cells which is considered as a source of many soluble factors and enzymes such as vascular endothelial growth factor, tumor necrosis factor-alpha, interleukin (IL)-1 β , IL-6, transforming growth factor beta, IL-10, and nitric-oxide that had variable effect on hematopoietic cells.^[34]

In the current study, CD56 was significantly more expressed in cases with WBC $\geq 30 \times 10^9$, in comparable to the result obtained from Iriyama *et al.*,^[35] who found that WBC count was higher in CD56-positive cases but does not reach the significant level.

Moreover, CD56 was significantly observed in cases with low platelets count and high PB blasts and BM blasts cells. These results were comparable to that published by Di Bona *et al.*^[30] and Hsiao *et al.*;^[36] this variation in hematological parameters may be explained as CD56 expression is associated with an abnormal overexpression of the full-length p48 RUNX1 isoform in AML cells which block hematopoietic differentiation and enhances self-renewal of hematopoietic stem cells.^[37,38]

Regarding the initial response to the induction therapy, CD200 was a significantly correlated with nonresponsiveness to the induction therapy. The mechanism of the role that played by CD200 to produce its negative effect on the outcome is partially defined but may be due to memory T-cells suppression and

Table 1: Correlation between CD200 and CD56 with prognostic parameters

Prognostic parameters	CD200		Р	CD56		Р
	Positive, n (%)	Negative, n (%)		Positive, n (%)	Negative, n (%)	
Age (year)						
≥60	5 (31.2)	2 (14.3)	0.399	5 (31.2)	2 (14.3)	0.399
<60	11 (68.8)	12 (85.7)		11 (68.8)	12 (85.7)	
Gender						
Male	10 (62.5)	8 (57.1)	1.000	10 (62.5)	8 (57.1)	1.000
Female	6 (37.5)	6 (42.9)		6 (37.5)	6 (42.9)	
WBCs (×10 ⁹ /L)						
≥30	11 (68.8)	0	<0.001	11 (68.8)	0	<0.001
<30	5 (31.2)	14 (100)		5 (31.2)	14 (100)	
Extramedullary manifestation						
Yes	6 (66.6)	3 (33.3)	0.440	4 (44.4)	5 (55.5)	0.049
No	10 (47.6)	11 (52.3)		2 (9.5)	19 (90.4)	
Platelets (×10 ⁹ /L)	33.19±13.46	66.36±42.52	0.013	33.19±13.46	66.36±42.52	0.013
Blast BM (%)	63.63±17.35	56.0±16.91	0.235	63.63±17.35	56.0±16.91	0.235
Blast PB (%)	39.19±20.8	34.43±12.02	0.458	39.19±20.8	34.43±12.02	0.458

WBC=White blood cells, BM=Bone marrow, PB=Peripheral blood

Table 2: Correlation between CD200 and CD56 expression and complete remission achievement

Parameters	CD200		Р	CD56		Р
	Positive, n (%)	Negative, n (%)		Positive, n (%)	Negative, n (%)	
Yes	4 (25.0)	10 (71.4)	0.026	2 (33.3)	12 (50.0)	0.657
No	12 (75.0)	4 (28.6)		4 (66.7)	12 (50.0)	

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NK activity reduction in CD200-positive AML patients, especially in the NK cells with high lytic activity.^[9,39]

In concerning to the relation between CD56 and CR, there were 4 out of 6 cases (66.6) expressing CD56 being achieved complete response to induction therapy (P = 0.657), which is comparable to Raspadori *et al.*^[11] In the present study, the all four cases who coexpressed both CD200 and CD56 did not achieve CR; this clarifies the bad prognostic impact of both markers on AML patients when coexpressed together.

Conclusion

CD200 and CD56 expressions were detected in 53.3% and 20% of adult AML cases, respectively.

- CD200 and CD56 were expressed mostly in monocytic AML subtypes.
- Both markers were closely related to bad prognostic parameters including high WBC and low platelets count. Whereas high peripheral blood and bone marrow blast count and extramedullary manifestations where mostly presented with CD56 positive AML cases.
- In view of close association of both markers expression with low response to induction therapy, thus we may propose that both markers expression particularly CD200 in AML patients; could be regarded as an auxiliary poor prognostic marker.

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Nil.

Conflicts of interest

There are no conflicts of interest.

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