



Expression of CD133 and Aldehyde Dehydrogenase 1A1 in Borderline Ovarian Tumor and Their Correlation with International Federation of Gynecology and Obstetrics

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Abstract

Edited by: Sinisa Stojanoski Citation: Dewayani BM, Agustina H, Puspita D, Hernowo BS, Suryanti S. Expression of CD133 and Aldehyde Dehydrogenase 1A1 in Borderline Ovarian Tumor and their Correlation with International Federation of Gynecology and Obstetrics. Open Access Maced J Med Sci. 2020 Mar 25; 8(A):160-163. Sci. 2020 Mar 25; 8(A):160-163. https://doi.org/10.3889/com/ims.2020.347. Keywords: Borderline ovarian tumor; International Federation of Gynecology and Obstetrics stage; CD133. Aldehyde Dehydrogenase 1A1. *Correspondence: Brigitta M. Devayani, Department of American Contention and American Links Factoria Anatomic Pathology, Padjadjaran University, Faculty of Medicine, Bandung, Indonesia. Phone: +6281321778224. E-mail: bm.dewayani@gmail.com Received: 25-Jul-2019 Revised: 21-Jan-2020 Accepted: 27-Feb-2020 Accepted: 27-Feb-2020 Copyright: © 2020 Birgitta M. Dewayani, Hasrayati Agustina, Desi Puspila, Bethy S. Hernowo, Sri Suryanti Funding: Authors would like to declare that this study supported by Padjadjaran University internal grant Competing Interests: The authors have declared that no competing Interests exist Open Access: This is an open-access article distributed under the terms of the Creative Commons Attribution-NunCommercial 4.0 International License (CC BXNC 4.0)

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BACKGROUND: Borderline ovarian tumor (BOT) is an epithelial ovarian tumor with atypical epithelial proliferation without stromal invasion. BOTs may have an aggressive fashion with associated microinvasion feature, peritoneal "implants," regional lymphadenopathy, and higher International Federation of Gynecology and Obstetrics (FIGO) stage. CD133 is cancer stem cells (CSCs) marker that promotes cell proliferation and tumor invasion through induction of nuclear factor kappa B and upregulation of MMP9. Aldehyde dehydrogenase 1A1 (ALDH1A1) is CSCs marker that promotes cell proliferation through the upregulation of c-MYC and cyclin-D1.

AIM: The aim of this study was to investigate the correlation between CD133 and ALDH1A1 expression with FIGO stage. This research was performed as an analytic-observational with a cross-sectional design

METHODS: This research using a paraffin block of patients diagnosed as BOT in Hasan Sadikin Hospital Bandung. Samples were divided in two groups: FIGO stage IA and FIGO stage >IA. All samples were stained by immunohistochemistry CD133 and ALDH1A1. All data were analyzed using the Chi-square test with a significant level 5%

RESULTS: This study showed a statistically significant correlation between CD133 (p = 0.047) and ALDH1A1 (p = 0.042) expression with FIGO stage in BOT. Multivariate analysis showed there was no correlation between CD133 and ALDH1A1 in the affected of FIGO stage in BOT.

CONCLUSION: CD133 and ALDH1A1 in BOT can be considered as a factor to predict the prognosis of BOT through predict FIGO stage.

Introduction

Epithelial ovarian tumor (EOT) consists of benign EOT (cystadenoma), malignant EOT (carcinoma), and borderline ovarian tumor (BOT) [1]. Characteristics of BOTs are atypical epithelial proliferation and without stromal invasion [1], [2], [3]. BOTs comprise up to 15-20% of ovarian epithelial neoplasms. BOTs often occurred in vounger women with an average age is 40 years old [2]. According to the WHO classification, BOT consists of serous BOT, mucinous BOT, endometrioid BOT, and borderline Brenner tumor [1]. BOT often occurred at an earlier stage and has a better prognosis than malignant ovarian tumors. Some BOTs may have an aggressive fashion with associated peritoneal "implants," regional lymphadenopathy, and higher International Federation of Gynecology and Obstetrics (FIGO) stage [1], [4].

Cancer stem cells (CSCs) are a small proportion of tumor cells which are proposed to be able to proliferate and self-renew and invasion extensively [5]. CD133 and aldehyde dehydrogenase 1A1 (ALDH1A1) are CSC marker that can identify in ovarian tumor and correlate with a stage in ovarian cancer [6]. CD133 promotes cell proliferation, tumorigenesis, and tumor invasion and metastasis through induction of nuclear factor kappa B and upregulation of MMP9 [7]. ALDH1A1 promotes cell proliferation through upregulation of c-MYC and cycin-D1 [8]. High tumor cell proliferation, high ability of invasion, implants, and involvement of regional lymphadenopathy in BOT may influence FIGO stage and few cases of recurrences may occur [1]. Stage in BOT currently based on FIGO staging system. FIGO staging system according to involvement of tumor mass, whether unilateral or bilateral, tubal involvement, tumor mass in ascites fluid, peritoneal, lymph node involvement, or distant metastasis [9]. Loizzi et al. found that majority cases BOT had FIGO stage I (85.5%) and the remainder exhibited FIGO stage II (7.3%) and FIGO stage III (7.3%) [3].

To the best of our knowledge, no data have been reported until now about the role of the immunohistochemically assessed expression of CD133 and ALDH1A1 of BOT in a large, single Institution series in Indonesia.

Subjects and Methods

This study uses an analytic observational method with a cross-sectional study design and retrospective data retrieval/collection. Ethical clearance has been approved/assessed by Health Research Ethic Commission, Padjadjaran University, assessment number 1144/UN6.KEP/EC/2018. The inclusion criteria: The samples were obtained from patients registered at Hasan Sadikin Hospital, were diagnosed with BOT (serous BOT, mucinous BOT, endometrioid BOT, and borderline Brenner tumor) during January 2013 to July 2018. All paraffin blocks from all samples were good and all samples have complete data about FIGO stage in the medical record.

All samples were classified as FIGO stage IA (unilateral involvement) and FIGO stage >IA (bilateral involvement or involvement to another organ). Analysis of expression: Samples from paraffin blocks were prepared for immunohistochemistry (IHC) analysis; IHC analysis was performed based on the protocol provided by the anatomical pathology laboratory. The slides were visualized under the microscope with three reviewers that have no knowledge about clinicopathologic data.

Histoscore calculation

The positive result was shown/visualized as brown staining on the tumor cell. Analysis on CD133 was evaluated by brown staining in the cytoplasmic membrane of tumor cells and ALDH1A1 expression was evaluated by brown staining assessed in the cytoplasm. The stain intensity and distribution measured under the microscope was then converted into histoscore and categorized in high and low expression. This study used antibody CD133 (Polyclonal antibody, Elabscience) with dilution 1:200 and antibodyALDH1A1 (Polyclonal antibody, Santacruz) with dilution 1:800. The intensity was scored as having no expression (0), or weak (1), moderate (2), or strong (3) tumor-cell staining. Distribution was scored as having no expression (0), or <10% (1), 10-50% (2), or >50% of (3) tumor-cell staining.

Histoscores obtained from the samples were represented on a scale of 0-9. Histoscores were classifying to weak (histoscore ≤ 6) and strong (histoscore > 6).

This research was performed as an analyticobservational with a cross-sectional design. Statistical analysis was performed with SPSS using the Chisquare test with a significant level 5%. p <0.05 was considered to be statistically significant and in the test of independence (multivariate analysis), a p >0.05 was considered to show independence between CD133 and ALDH1A1.

Results

During January 2013–July 2018, there were 101 cases of BOT which were registered; however, only 84 samples were matched to inclusion criteria. Age ranging from 18 to 71 years (median 42 years) and 47 patients (56%) was over 40 years of age. The most common histopathology type was mucinous BOT in 59 patients (70.2%) and followed by serous BOT in 22 patients (26.2%) (Table 1).

Variable	n=84 (%)
Age (year)	
Mean±Std.	41.13±13.025
Median	42.00
Range (min–max)	18.00-71.00
<40	37 (44.0%)
≥40	47 (56.0%)
Type of histopathology	
Mucinous BOT	59 (70.2%)
Serous BOT	22 (26.2%)
Endometrioid BOT	2 (2.4%)
Borderline Brenner tumor	1 (1.2%)
Parity	
PO	15 (17.9%)
P≥1	69 (82.1%)

Characteristic of this research subject based on age, type of histopathology, and parity. BOT: Borderline ovarian tumor.

From this study, we found 66 patients with FIGO stage IA. Fourteen patients (21.2%) showed low expression of CD133 and 52 cases (78.8%) showed high expression of CD133 (Figure 1a). Moreover, 18 patients with FIGO stage more than IA showed low expression in 8 cases (44.4%) and high expression in 10 cases (55.6%). From statistical analysis showed the significant correlation between CD133 (p < 0.05) expression with FIGO stage in BOT. Expression of ALDH1A1 in FIGO stage IA showed low expression in 11 patients (16.7%) and high expression in 55 patients (83.3%) (Figure 1b). Patients with FIGO stage >IA showed low expression in 7 patients (38.9%) and high expression in 11 patients (61.1%). From statistical analysis showed the significant correlation between ALDH1A1 (p < 0.05) expression with FIGO stage in BOT (Table 2). Multivariate analysis showed there was no correlation between CD133 and ALDH1A1 in the affected of FIGO stage in BOT (Table 3).

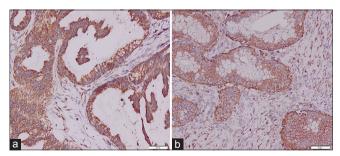


Figure 1: CD133 and aldehyde dehydrogenase 1A1 (ALDH1A1) expression of borderline ovarian tumor in this study. (a) CD133 strong expression in this study, tumor cell stained with CD133 antibody in the cytoplasmic membrane (200×). (b) ALDH1A1 strong expression in this study, tumor cell stained with ALDH1A1 antibody in the cytoplasm (200×)

Table 2: CD133 and ALDH1A1data on FIGO stage of BOT group

FIGO stage of BOT		p value
IA (n=66) %	>IA (n=18) %	
		0.047**
14 (21.2)	8(44.4)	
52 (78.8)	10(55.6)	
		0.042**
11(16.7)	7(38.9)	
55 (83.3)	11(61.1)	
	IA (n=66) % 14 (21.2) 52 (78.8) 11(16.7)	IA (n=66) % >IA (n=18) % 14 (21.2) 8(44.4) 52 (78.8) 10(55.6) 11(16.7) 7(38.9)

in BOT. BOT's Borderline ovarian tumor, FIGO: International Federation of Gynecology and Obstetrics, ALDH1A1: Aldehyde dehydrogenase 1A1.

Discussion

Prognostic factors in BOT include age, FIGO stage, residual disease following surgery, type of peritoneal implants (with or without invasion), presence of microinvasion, micropapillary pattern, and the CA-125 value. Prognosis of BOT overall has a positive prognosis because over 80% of cases are diagnosed at an early stage of the disease and radical surgery and surgical staging are the standard of care for this disease includes hysterectomy, bilateral salpingo-oophorectomy, omentectomy, multiple biopsies, and peritoneal cytology [2].

Table 3: Multivariate analysis correlation CD133 and ALDH1A1expression with FIGO stage of BOT

Variable	Estimation rate	SE	p value
CD133	-0.972	0.577	0.092
ALDH1A1	-1.035	0.602	0.602
Constant	2.128	1.354	0.116

dehydrogenase 1A1.

FIGO stage for ovarian carcinoma includes BOT which is the most superior staging system for predict patient prognosis through discriminating survival outcomes [10]. FIGO stage is according to the involvement of tumor mass, whether unilateral or bilateral, tubal involvement, tumor mass in ascites fluid, peritoneal, lymph node involvement, or distant metastasis [9]. The primary route of ovarian cancer expanding/metastasis includes intraperitoneal implantation of exfoliated tumor cells and spreading through retroperitoneal lymphatic channels [10].

CSCs are a small proportion of tumor cells which are proposed to be able to proliferate and selfrenew and invasion extensively [5]. CD133 is one of CSCs marker in ovarian cancer [6]. Long *et al.* found that the presence of CD133+ ovarian CSC can undergo an epithelial–mesenchymal transition-like process and display enhanced metastatic capacity of tumor cells *in vitro* and *in vivo* through secreting soluble mediators. CCL5 is chemokine which is one of the soluble mediators produced by CD133+ ovarian CSC. Signaling of CCL5 is an important thing for the invasive capability of CSC and tumor cells [11].

This study performed CD133 expression analysis on FIGO stage and microinvasion feature of BOT. The result from IHC staining confirms that there are 55.6% of patients with FIGO stage more than IA showed high expression of CD133. From statistical analysis showed the significant correlation between CD133 (p < 0.047) expression with FIGO stage in BOT. High expression of CD133 results in high stage FIGO stage (FIGO stage more than IA).

ALDH1A1 is a marker of CSC that promotes self-renewal, differentiation, and self-protection of cells. High expressions of ALDH1A1 correlated with poor cancer prognosis but not correlate with highly malignant phenotypes [8]. Hence, in this study, we performed ALDH1A1 expression analysis on FIGO stage of BOT. The result from IHC staining confirms that there are 61.1% of patients with FIGO stage more than IA showed high expression of ALDH1A1. From statistical analysis showed the significant correlation between ALDH1A1 (p<0.042) expression with FIGO stage in BOT. High expression of ALDH1A1 results in high stage FIGO stage (FIGO stage more than IA).

This study also performed multivariate analysis correlation CD133 and ALDH1A1 expression with FIGO staging. There was no correlation between CD133 and ALDH1A1 expression with FIGO stage.

Conclusion

There was a significant correlation between CD133 with FIGO staging and correlation between ALDH1A1 with FIGO stage.

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