

MicroRNA-223 is Associated with Resistance Towards Platinum-based Chemotherapy and Worse Prognosis in Indonesian Triple-negative Breast Cancer Patients

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Ibnu Purwanto¹
Didik Setyo Heriyanto²
Irianiwati Widodo²
Mohammad Hakimi³
Mardiah Suci Hardianti¹
Teguh Aryandono⁴
Sofia Mubarika Haryana⁵

¹Division of Hematology and Medical Oncology, Department of Internal Medicine, Faculty of Medicine, Public Health, and Nursing, Gadjah Mada University/Dr Sardjito Hospital, Yogyakarta, Indonesia; ²Department of Anatomical Pathology, Faculty of Medicine, Public Health, and Nursing, Gadjah Mada University/Dr Sardjito Hospital, Yogyakarta, Indonesia; ³Department of Clinical Epidemiology and Biostatistics Unit, Faculty of Medicine, Public Health, and Nursing, Gadjah Mada University/Dr Sardjito Hospital, Yogyakarta, Indonesia; ⁴Department of Surgical Oncology, Faculty of Medicine, Public Health, and Nursing, Gadjah Mada University/Dr Sardjito Hospital, Yogyakarta, Indonesia; ⁵Department of Histology and Cell Biology, Faculty of Medicine, Public Health, and Nursing, Gadjah Mada University/Dr Sardjito Hospital, Yogyakarta, Indonesia

Correspondence: Ibnu Purwanto
Division of Hematology and Medical Oncology, Department of Internal Medicine, Faculty of Medicine, Public Health, and Nursing, Gadjah Mada University/Dr Sardjito Hospital, Jalan Farmako, Sekip Utara, Yogyakarta 55281, Indonesia
Tel/Fax (+62 274) 560300
Email ibnupurwanto@ugm.ac.id

Purpose: Determining the optimal strategy to implement systemic treatment modalities has been challenging in triple-negative breast cancer (TNBC). We aim to investigate the role of microRNA-223 (miR-223) as prognostic factor and predictor of response toward chemotherapy in TNBC.

Patients and Methods: We retrospectively analyzed the association of pretreatment miR-223 expression with clinicopathologic characteristics and 36-month overall survival (OS) of 53 all stages TNBC patients. Tumor level of miR-223 was measured using real-time quantitative polymerase chain reaction (expressed in fold change). Cutoff value for miR-223 was determined by using receiver operating curve (ROC). Kaplan–Meier curve was used to perform survival analysis.

Results: The optimum cutoff value for miR-223 was 23.435 (AUC: 0.706, 95%CI: 0.565–0.848; p :0.01; sensitivity: 78.6%; specificity: 56%) and was used to categorize miR-223 expression into over- and underexpressed group. Overexpression of miR-223 was associated with increased expression of EGFR (69.7% vs 35%, p : 0.022) and lower 36-month OS (33.3% vs 70%; median OS±SE (months): 25.66±1.58 vs 30.23±1.99; log rank p <0.05). Worse survival is observed in miR-223 overexpressed group receiving platinum-based chemotherapy compared to miR-223 underexpressed group (mean OS (95%CI) months: 24.7 (20.3–29.1) vs 34.3 (31.2–37.4); p <0.01), while no significant difference observed in non-platinum containing regimen. No significant association was observed between miR-223 expression with other clinicopathologic characteristics.

Conclusion: Overexpression of miR-223 is associated with increased expression of EGFR, worse prognosis, and resistance toward platinum-based chemotherapy in Indonesian TNBC patients.

Keywords: miR-223, chemotherapy, prognosis, EGFR, TNBC

Introduction

Triple-negative breast cancer (TNBC), defined by the absence of estrogen (ER), progesterone (PR), and HER2 receptors, is associated with more aggressive disease characteristic, poor response toward therapy and shorter survival. Despite increased prominence of immunotherapy in TNBC, systemic chemotherapy is still the mainstay of treatment. Unfortunately, only 20% of all TNBC respond to systemic chemotherapy and little is understood behind the mechanism of chemotherapy resistance or susceptibility in these patients.¹ Currently we are still unable to decide which patient might or might not benefit from receiving systemic chemotherapy.

MicroRNA-223 (miR-223) has pleiotropic effect in malignancy, meaning it can act both as an oncogenic-microRNA (oncomiR) or as a tumor suppressor, depending on the context.² MicroRNA-223 is also known to contribute in resistance to chemotherapy in many different malignancies, including resistance to platinum- and anthracycline-based regimens, the two most commonly used regimens in breast cancer therapy.^{3,4} Evidence on the role of miR-223 in breast cancer, especially in TNBC, is still limited and showing conflicting results.^{5–8} Furthermore, most of these studies were preclinical studies limiting their implementation in clinical settings. We aim to investigate the role of miR-223 in Indonesian TNBC patients, including its prognostic role, its association with unfavorable clinical features, response toward chemotherapy, as well as association with other biomarkers, such as cytokeratin 5/6 (CK5/6), EGFR, and p53 mutant.

Patients and Methods

Patient Populations

This was a retrospective study conducted at Dr Sardjito Hospital, Yogyakarta, Indonesia. We consecutively included stage I–IV TNBC patients diagnosed between 2014 and 2017. Out of 272 detected patients, 219 were excluded due to incomplete data, leaving 53 patients included in this study. Clinical data was retrieved from medical records. This study has been approved by the IRB Ethics Committee Faculty of Medicine, Public Health, and Nursing, Gadjah Mada University/Dr Sardjito Hospital, Yogyakarta, Indonesia with approval numbers: KE/FK/0751/EC/2018 (first approval) and KE/0286/03/2020 (latest update).

Pathology Assessment

Tumor samples were obtained from formalin-fixed paraffin-embedded (FFPE) tissue stored at the Department of Anatomical Pathology, Dr Sardjito General Hospital, Yogyakarta, Indonesia; Waskhita Laboratory; Panti Rapih General Hospital; and CITO Laboratory.

Real-time Quantitative Polymerase Chain Reaction

Expression of miR-223 was determined by using real-time quantitative polymerase chain reaction (qRT-PCR). For preparation of RNA from FFPE samples, the (QIAGEN GmbH, Hilden, Germany) RNeasy FFPE kit was used according to manufacturer's recommendations. Three

sections of 10 μ m FFPE thickness were used per preparation. For quantitative PCR (qPCR), RNA/sample was amplified using the (Genes Laboratories Inc., Seongnam, South Korea) NEXpro™ qRT-PCR Master (SYBR) (Cat. no. NexQ-7000) in (Bioneer, Daejeon, South Korea) a Step One Real Time PCR System (Bioneer Exicycler™ 96 Real Time Quantitative Thermal Block). MicroRNA-223 forward primer, 5'-AGC CGT GTCAGTTTG TCA AAT-3', and miR-223 reverse primer, 5'-GTGCAGGGTCCGAGG TC-3'. U6 was used as internal control. U6 forward primer, 5'-CTCGCTTCG GCAGCA CA-3' and U6 reverse primer, 5'-AACGCTTCACGA ATTTGCGT-3'. *GAPDH* has been used for normalization of gene expression data. The cycling conditions were as follows: initial denaturation at 95°C for 10 min, followed by 40 cycles at 95°C for 20 seconds, annealing at 60°C for 40 seconds and extension at 72°C for 60 seconds.

Immunohistochemistry

Block paraffin samples were cut 4 μ m in thickness to analyze the expression of p53 mutant, EGFR and CK5/6 by immunohistochemistry method. Antibodies used in this study were (Abcam, Cambridge, USA) Clone Ab 32,049, rabbit mAb p53 mutant (dilution 1:50), (Novocastra, Newcastle, UK) Clone NCL-L-EGFR, mouse mAb EGFR (dilution 1:20), and (Cell Marque, Trappes, France) Clone D5 and 16B4, RTU, mouse mAb CK5/6, with chromogen DAB. In this study, the membranous expression of EGFR and CK5/6 in more than 10% of the tumor cells, nuclei expression of p53 mutant in more than 1% of the tumor cells were considered positive (Figure 1).⁹ The expression of those markers was examined by two senior pathologists independently. If there was any disagreement between the two pathologists, discussion was carried out and consensus had to be made between them.

Statistical Analysis

Statistical analysis was performed using IBM SPSS version 24 software. Association between tumor subtype and clinicopathologic features was analyzed using chi-squared test or Fisher's exact test if assumptions for chi-squared test were violated. All of the reported *p*-values are two-sided, with significance level set at *p*<0.05. Cutoff value for miR-223 expression was determined by using receiver operating curve (ROC) to categorize miR-223 expression into over- and underexpressed groups. Survival analysis was performed using Kaplan–Meier curve. Cutoff for overall survival (OS) analysis was set at 36 months.

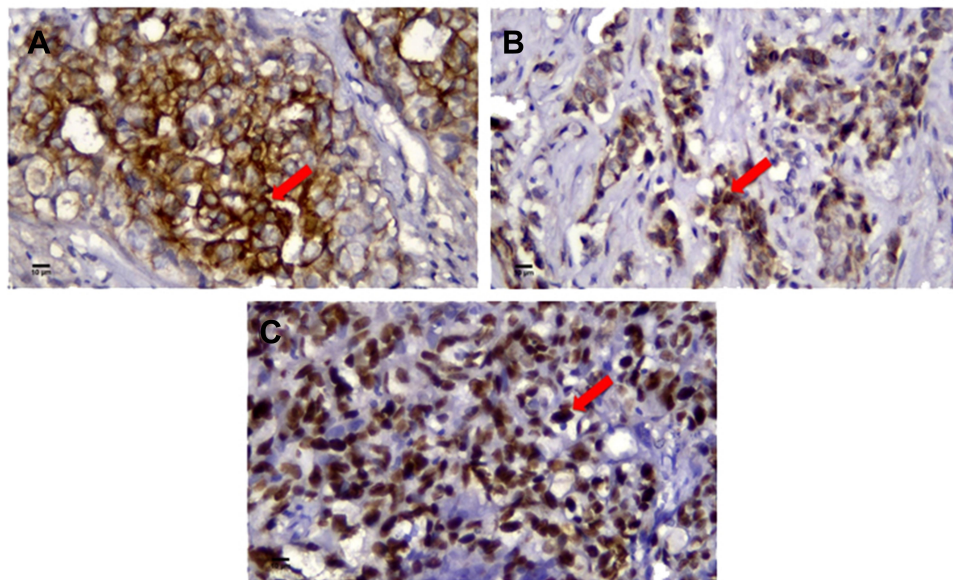


Figure 1 Immunohistochemistry of (A) CK5/6 (red arrow: positively stained in cytoplasmic membrane and cytoplasm of tumor cell). (B) EGFR (red arrow: positively stained in cytoplasmic membrane and cytoplasm of tumor cell). (C) p53 mutant (red arrow: positively stained in nucleus of tumor cell). Bar: 10 µm. Magnification: 400 times.

Results

According to the ROC curve, the optimum cutoff value for miR-223 was 23.435 (AUC: 0.706, 95%CI: 0.565–0.848; *p*: 0.01; sensitivity: 78.6%; specificity: 56%). This cutoff was used to categorize miR-223 expression into over- and underexpressed groups (Figure 2, Table 1). Out of 53 included subjects, 20 were classified into miR-223

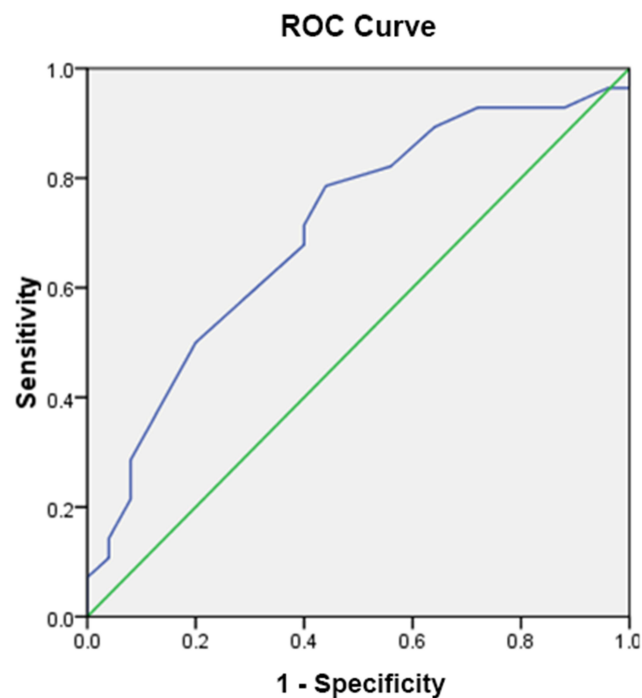


Figure 2 ROC Curve of miR-223 to Predict 36-months OS with AUC of 0.706.

underexpressed subgroup and 33 were classified into the miR-223 overexpressed group. The mean age of the subjects was 53 years old, while the median age was 50.28. Subjects were divided into two groups according to its median value, 23 (43.4%) subjects were below 50 years old and 30 (56.6%) subjects more than or equal to 50 years old. The mean and median BMI in this study were 24. Thirty-four (64.2%) subjects had BMI <25 and 19 (35.8%) subjects had BMI ≥25. Seven (13.2%) subjects presented with early stage (I–IIA), 38 (71.7%) subjects with locally advanced stage (IIB–IIIC), while eight (15.1%) subjects with stage IV disease. Thirteen (24.5%) subjects presented with tumor size (T) <5 cm while 40 (75.5%) subjects with T ≥5 cm. Thirty-six (67.9%) subjects presented with no nodal involvement (N), while 17 (32.1%) subjects had at least one nodal metastasis. Distant metastasis was found in eight (15.1%) subjects. There was no significant difference in age, BMI, disease stage, tumor size, nodal status, as well as distant metastasis between miR-223 underexpressed and miR-223 overexpressed subgroup. Subjects in miR-223 overexpressed subgroup had higher frequency

Table 1 AUC Value of miR-223 ROC

Area	Standard Error	p-value	95%CI	
			Lower Bound	Upper Bound
0.706	0.072	0.010	0.565	0.8484

EGFR expression (69.7% vs 35%; $p<0.05$), while no significant difference in expression of p53 mutant (35% vs 45.5%; $p:0.454$) and CK5/6 (50% vs 60.6%; $p:0.57$) between the two subgroups (Table 2).

Out of 53 subjects included, 25 subjects (47.2%) were still alive at 36-months follow-up, while 28 subjects (52.8%) were deceased. Twenty-two out of 33 subjects (41.5%) in miR-223 overexpressed subgroup were deceased, which was worse compared to six out of 20 subjects (30%) in miR-223

underexpressed subgroup. Mean OS of miR-223 overexpressed subgroup was 25.7 months (95%CI: 22.6–28.7 months), which was significantly shorter compared to 30.2 months (95%CI: 26.3–34.1 months) in miR-223 underexpressed subgroup ($p<0.05$) (Figure 3, Table 3). The overall median OS of this study was 31.17 months. When assessing OS according to chemotherapy regimen, there was significant difference between miR-223 overexpressed and miR-223 underexpressed subgroup who received platinum

Table 2 Clinicopathologic Profile of TNBC Patients with miR-223 Underexpression and Overexpression

Parameter	miR-223 Underexpressed	miR-223 Overexpressed	p-value
	(n=20)	(n=33)	
EGFR			
Positive	7	23	0.013 ^a
Negative	13	10	
CK5/6			
Positive	10	20	0.57
Negative	10	13	
p53 Mutant			
Positive	7	15	0.454
Negative	13	18	
Stage			
Early	3	4	0.209
Locally- advanced	16	22	
Metastatic	1	7	
Tumor size			
<5 cm	4	9	0.555
≥5 cm	16	24	
Nodal status			
Positive	12	24	0.336
Negative	8	9	
Distant metastasis			
Present	1	7	0.113
Absent	19	26	
Age			
<50 y.o	6	17	0.126
≥50 y.o	14	16	
BMI			
<25	11	23	0.279
≥25	9	10	
Chemotherapy regimen			
Platinum	9	14	0.559
Non-platinum	11	17	
Missing	0	2	

Note: ^aStatistically significant.

Abbreviations: CK5/6, cytokeratin 5/6; BMI, body mass index.

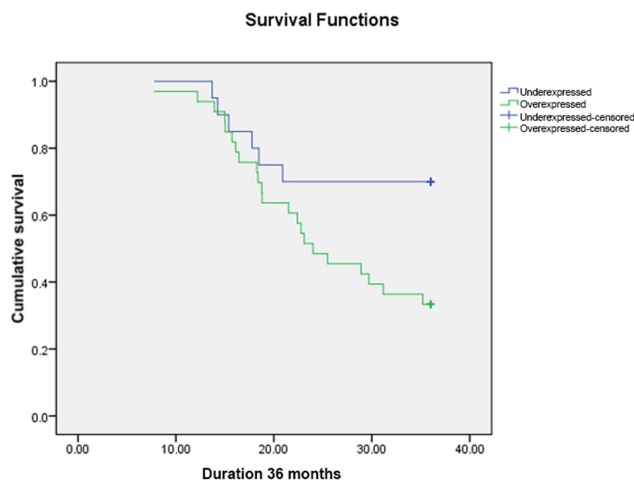


Figure 3 Kaplan–Meier Overall Survival Curve of All TNBC Patients According to miR-223 Expression.

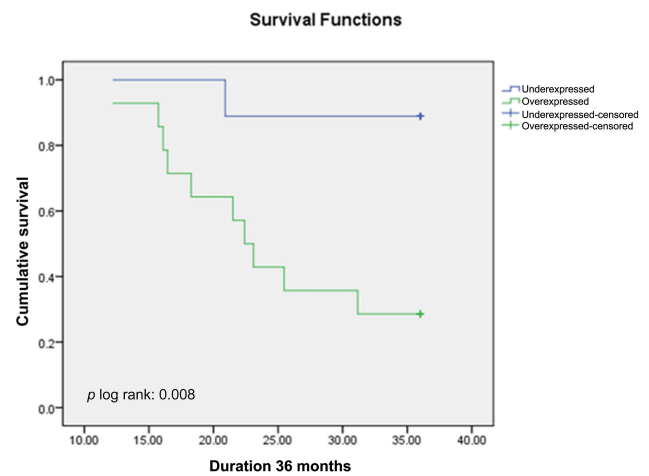


Figure 4 Kaplan–Meier Overall Survival Curve of TNBC Patients Treated with Platinum-containing Regimen According to miR-223 Expression.

containing regimen (mean OS (95%CI) months: 24.7 (20.3–29.1) vs 34.3 (31.2–37.4); $p < 0.01$) (Figure 4), while there was no significant difference between the two sub-groups who received non-platinum containing regimen (mean OS (95%CI) months: 26.2 (21.7–30.8) vs 26.8 (20.9–32.8); $p: 0.69$) (Figure 5). Research data is available in [supplementary file 1](#).

Discussion

In our cohort, overexpression of miR-223 was associated with worse prognosis (mean OS: 25.7 vs 30.2 months). Furthermore, EGFR was more frequently expressed in the miR-223 overexpressed group compared to the miR-223 underexpressed group (69.7% vs 35%; $p < 0.05$). Expression of EGFR is generally associated with basal-likeness which represents more aggressive subtype of TNBC.¹⁰ Our finding differs from previous preclinical studies which observed miR-223 as tumor suppressor in TNBC cell lines.^{5,7} On the contrary, a retrospective clinical study observing serum level of circulating miRNA in 60 TNBC patients showed that miR-223 level is significantly elevated in relapsing

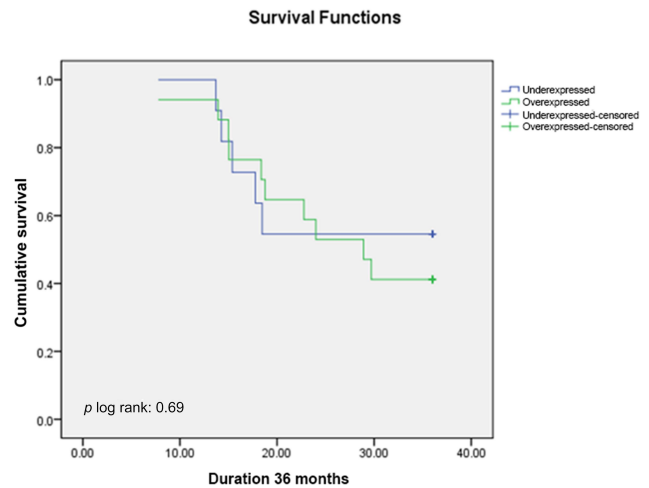


Figure 5 Kaplan–Meier Overall Survival Curve of TNBC Patients Treated with Non-platinum-containing Regimen According to miR-223 Expression.

group compared to nonrelapsing group, suggesting its role as oncomiR, which is consistent with our study.⁶

The prognostic role of miR-223 in our cohort was especially apparent in patients receiving platinum-based chemotherapy, while no survival difference was observed in patients receiving non-platinum chemotherapy regimen.

Table 3 Mean OS of TNBC Patients According to miR-223 Expression

miR-223 Expression	Mean OS (Month)	Standard Error (Month)	95%CI	
			Lower Bound (Month)	Upper Bound (Month)
Underexpressed	30.225	1.997	26.311	34.139
Overexpressed	25.655	1.578	22.561	28.748
Overall	27.379	1.275	24.880	29.879

Abbreviation: OS, overall survival.

This might suggest resistance to platinum as one of the underlying mechanisms. Research has shown that platinum-based chemotherapy results in better outcome in TNBC compared to non-TNBC.^{11,12} Mechanism behind the higher efficacy of this regimen is thought to be caused by homologous recombination deficiency (HRD) which is a commonly observed characteristic in both *BRCA* mutated and sporadic-TNBC with “*BRCAness*” profile.^{13–16} However, molecular heterogeneity of TNBC causes varied response to chemotherapy amongst these patients. Roughly only 20% of all TNBC patients show excellent response, while the rest exhibit little to no benefit.¹ Resistance to chemotherapy in TNBC, including to platinum-based regimen, has been observed in previous studies.^{17,18} The main mechanisms of chemotherapy resistance in TNBC are (1) ATP-binding cassette (ABC) transporter-mediated drug efflux; (2) cancer stem cells (CSC); (3) hypoxia; (4) avoidance of apoptosis; (5) various signaling pathways, including NF- κ B (nuclear factor kappa-light-chain-enhancer of activated B cells), PTEN and PI3K-AKT-mTOR pathway, JAK/STAT pathway, and receptor tyrosine kinase; (6) mRNA (although evidence on miR-223 in this subject is lacking); and (7) TNBC heterogeneity.¹⁸ Although there is currently no evidence of platinum resistance associated with miR-223 in TNBC, such observations have been made in other types of malignancy. MicroRNA-223 has been reported to cause platinum resistance in non-small-cell lung cancer (NSCLC) through regulation of autophagy, and through deregulation of cell cycle in gastric cancer.^{3,19} A preclinical study to evaluate the mechanism of miR-223 causing platinum resistance through TNBC cell-line or a xenograft model can improve our understanding on this subject.

Determining the optimal strategy to implement systemic treatment modalities has been challenging in TNBC. Due to the heterogeneity of TNBC, there might not be a “one for all” solution. Instead, the more realistic approach is to identify and decipher the role of various biomarkers with strategic roles in the disease development and resistance/susceptibility towards various therapeutic modalities. Although we are yet unable to define the precise mechanism behind miR-223 as a contributor toward platinum resistance, we were able to define which group of patients would receive benefit from platinum-based chemotherapy and which group would not according to miR-223 expression. However, this is a small sample-sized retrospective study which ideally requires a verification cohort in a prospective study with sizable sample to validate its result.

Conclusion

In summary, miR-223 overexpression is associated with worse prognosis, increased expression of EGFR, and resistance to platinum-based chemotherapy in Indonesian TNBC patients.

Ethics Approval and Informed Consent

Written informed consent was obtained from the patients involved, including permission to use clinical data and re-examination of tissue specimen. This study was conducted in accordance with the Declaration of Helsinki. Ethical clearance was approved by the IRB Ethics Committee Faculty of Medicine, Public Health, and Nursing, Gadjah Mada University/Dr Sardjito Hospital, Yogyakarta, Indonesia with approval numbers: KE/FK/0751/EC/2018 (first approval) and KE/0286/03/2020 (latest update).

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Author Contributions

All authors made a significant contribution to the work reported, whether that is in the conception, study design, execution, acquisition of data, analysis and interpretation, or in all these areas; took part in drafting, revising or critically reviewing the article; gave final approval of the version to be published; have agreed on the journal to which the article has been submitted; and agree to be accountable for all aspects of the work.

Disclosure

The authors report no conflicts of interest in this work.

The initial result of this study was presented at the 2020 ESMO Asia Virtual Congress as a poster presentation. The poster’s abstract was published in the *Annals of Oncology*: [https://www.annalsofoncology.org/article/S0923-7534\(20\)42552-9/fulltext](https://www.annalsofoncology.org/article/S0923-7534(20)42552-9/fulltext).

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