

Clinical research

Molecular classification of infiltrating ductal carcinomas in Western Algeria

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Abstract

Introduction: Breast carcinoma is a heterogeneous pathology. These subtypes are known to vary in terms of risk factors, natural histories and responses to therapies. The prognostic classification based on gene expression and genomic profiling was implemented to refine therapeutic indications and improve patient survival. Our goal is to classify infiltrating ductal carcinomas according to their molecular profiles and analyze the different clinicopathological variables of these molecular groups.

Material and methods: We conducted a 24-month retrospective study involving 50 patients recruited from the Oran University Hospital and the Oran Regional Military Hospital. Tumors were analyzed histologically and classified after an immunohistochemical study in groups: luminal A, luminal B, Her2+ and basal-like.

Results: Our study showed the predominance of molecular subtype luminal B (36%) of which 55.56% of these tumors were of Scarff-Bloom-Richardson (SBR) grade II and 44% were of grade III. 55.56% were T2 size followed by T1 size with a percentage of about 33.33%. Axillary ganglionic metastases were found in 88.89% of cases, followed by luminal A (32%) with 62.5% grade III, and 37.5% grade SBR II. Thus 37.5% of these tumors were of stage T2 and T4. Axillary ganglionic metastases were present in 87.5%. The HER2 type (20%) with 60% SBR II grade and 40% grade III. All basal-like tumors represented the highest SBR grade III with a percentage of 100%.

Conclusions: Molecular classification is crucial in the direction of treatment.

Key words: breast cancer, molecular subtypes, hormonal receptors (RH), HER2 status, proliferation index (Ki67).

Introduction

Breast cancer is the most common cancer in women. This pathology represents a real major public health problem [1]. According to the latest estimates in 2018, the incidence has increased by more than 24.2% and mortality by 15% worldwide.

In Algeria, the rate of breast cancer is around 180,537 new cases per year, the percentage of which is 16.3%. The current therapeutic strategy is based on clinical and anatomopathological classification studies (TNM and pTNM), which make it possible to assess the locoregional extension at the time of surgery, but also to establish the stage of cancer progression [2].

The TNM classification includes clinicopathological parameters such as: tumor size (T), presence of lymph nodes (N) and presence of metastases (M) [3].

The predominant function of histological classification is to subdivide tumors according to their pathological nature into non-infiltrating tumors (ductal carcinoma in situ and lobular carcinoma in situ) and into infiltrating tumors with metastatic potential [4].

Given the increasing availability of new anti-tumor molecules, it is crucial to improve the prognostic classification of breast cancer to refine the therapeutic indications and improve patient survival, through a more detailed and objective molecular characterization of the pathology [4].

The objective of our present study is to classify the infiltrating mammary carcinomas collected at the University Hospital of Oran “November 1, 1954” and at the Regional Military University Hospital of Oran (HMRUO) according to their molecular profiles, then analyze the different clinicopathological variables of these tumors.

Material and methods

Population of study

This is a retrospective study spread over 24 months, comprising 50 patients aged between 35 and 83 years from West Algeria collected from the university hospital establishment of Oran (EHUO) “November 1, 1954” and the Regional Military University Hospital of Oran (HMRUO) for diagnosis, monitoring and management.

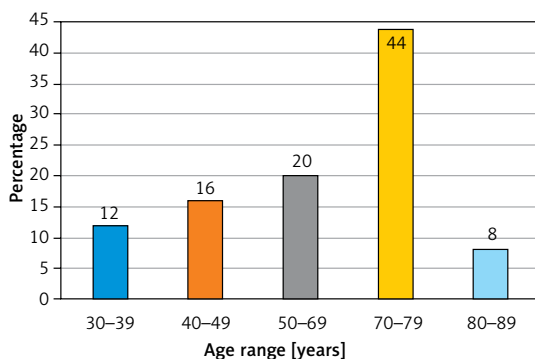


Figure 1. Graphical distribution of patients according to age

Methods

The study was performed on biopsies and surgical specimens fixed in 10% formalin and embedded in paraffin. Our tumor samples were analyzed histologically and classified after an immunohistochemical study into groups: luminal A, luminal B, HER2 and basal-like.

The analysis concerned the following parameters: age, TNM tumor stage [3], Scarff-Bloom-Richardson (SBR) [5] histological grade, lymph node involvement, distant metastases, receptor status, hormones, the HER2 receptor (human epidermal growth factor) and the proliferation index Ki67.

The immunohistochemistry technique was performed with the Leica BOND-MAX device in fully automated mode and the antibodies used were: RE (clone BV3; Ref: Mob 124 Mob195); RP (BV7 clone; Ref: RMABO28); HER2 (c-erbB2) (BV5 clone; Ref: RMABO26); Ki67 (SP6 clone; Ref: RMABO04).

Statistical analysis

Statistical analysis and data processing were performed with the software IBM SPSS version 20. The χ^2 test were used to determine the correlations. A *p*-value < 0.05 was considered statistically significant. The data were reported as clinical stage of tumor, lymph nodes status, metastatic status, SBR grade, ER/PR status, and HER2 expression.

In ethics and professional conduct, all of our patients responded to the medical questionnaire with informed consent.

Results

The careful combination of histological and immunohistochemical parameters results in better classification compared to their individual use, thereby confirming that molecular mechanisms and clinical outcomes are linked to ensure a better understanding of the mechanisms of breast cancer.

Descriptive study of the sample

Our population included patients between 35 and 83 years old from western Algeria. According to our study, patients aged less than 69 years present the lowest frequencies, with respectively 12%, 16% and 20%. These rates increased from the fourth age group 70–79 years with a frequency of 44%, but the age group 80–89 years noted a low frequency of 8% (Figure 1).

Grade III SBR (Scarff-Bloom-Richardson) tumors in our study population rank first, with 56% of cases, followed by grade II tumors at 44%.

Tumor size was an important prognostic factor. The latter was evaluated in the 50 patients that we followed and who presented tumors in stage

T2, the frequency of which was 48%. Concerning size, T1 had a percentage of 28%, followed by size T4 with 16%, followed by size T3, which was the least frequent, with a percentage of 8%. Out of 50 lymph node dissections examined, 80% of the lymph nodes were invaded by the tumor process, of which we found 24 (48%) of them presented with stage N1, 12 (24%) patients presented with stage N2, 4 other patients presented with stage N3 with percentages of 8% and 20% of the patients were carriers of healthy nodes classified N0 (Table I).

The degree of positivity of hormone receptors was proportional to cell differentiation, which measured the receptivity of the tumor to hormone therapy. In our series, 68% of patients expressed estrogen receptors, and also 56% of cases were positive for progesterone receptors.

There was overexpression of the HER2 protein, the percentage of which was 56% versus 44% of patients with a negative HER2 profile (Table I).

For all of our patients, we found 44% of cases with a Ki67 status less than 14 (ki67 low), and 56% of cases with a ki67 status greater than 14 (ki67 high) (Table I).

Distribution of invasive ductal carcinomas according to molecular classification

We were able to identify for each tumor the molecular class according to its immunohistochemical profile (study of hormone receptors (RH), membrane receptor (HER2) and nuclear marker (Ki67)). The results showed that 32% of tumors were luminal group A, 36% were from the luminal B group, 20% were from the HER2 group and 12% of the tumors were from the basal-like group (Figure 2).

In our study, luminal A and luminal B groups were found to be the most common molecular subtypes in patients over 70 years of age (Figure 3).

By correlating the molecular type of tumors with the histological grade, in the majority of cases the histological grade (SBR) of luminal type B and HER2 patients in our research was moderate grade (SBR II) (55.56% and 60% respectively), and

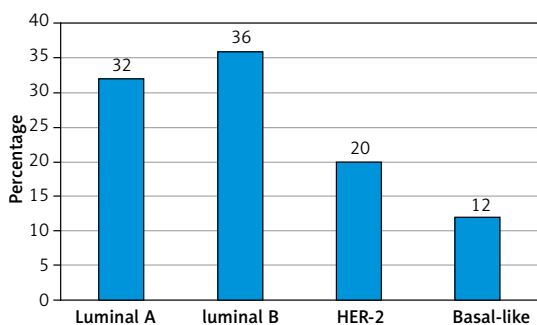


Figure 2. Graphical distribution of patients according to molecular subtypes

Table I. Characteristics of clinicopathological parameters of the study population

Characteristics	Numbers of cases (%)
Tumor site:	
Left breast	18 (36)
Right breast	18 (36)
SBR grade:	
I	0 (0)
II	22 (44)
III	28 (56)
Tumor size:	
T1	14 (28)
T2	24 (48)
T3	4 (8)
T4	8 (16)
Lymph nodes status:	
N0	10 (20)
N1	24 (48)
N2	12 (24)
N3	4 (8)
Metastatic status:	
M0	14 (28)
M1	2 (4)
Mx	34 (68)
ER positive	34 (38)
PR positive	28 (56)
HER2 positive	28 (56)
Ki67 (low)	22 (44)
Ki67 (high)	28 (56)

the luminal A and basal-like subtypes had a high grade (SBR III) (62.5% and 100% respectively with $p = 0.072$) (Table II).

The comparison of the average tumor sizes in our study population showed that there was a difference in tumor size in each molecular subtype such as: the HER2 subtype which had a tumor size of 77.5 mm. Regarding the Basal-like subtype was tumor size of

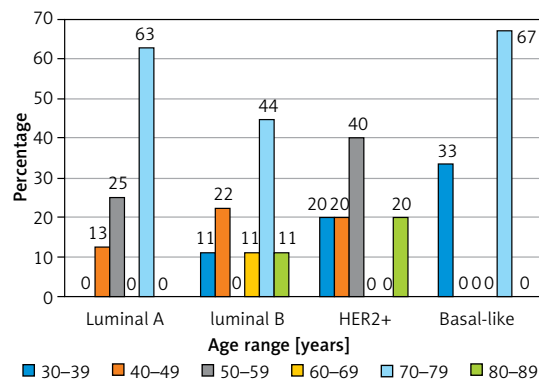


Figure 3. Graphical distribution of tumors by age according to molecular group

50 mm. T2 size was predominant in the three molecular subtypes (HER2, luminal B and luminal A, respectively 60%, 55.5% and 37.5%, $p = 0.013$).

Basal and luminal B tumors had the highest nodal invasion rates of 100% and 89%, respectively, followed by luminal A type of 88%. For HER2 subtype tumors, the percentage corresponded to 60% (Table II).

Discussion

Breast cancer is a major public health concern due to its increasing incidence. It is generally known that the risk of developing breast cancer increases with age.

The analysis of our results shows that the age group most affected is that between 70 and 79 years with a percentage of 44%, unlike the Saudi study which showed that 54.3% of cases were in women under 50 [6]. This can be explained by the absence of a screening program for this age group in Algeria.

The SBR grade is a prognostic factor independent of tumor size and lymph node involvement,

for which the Elston and Ellis grade is currently recommended in 2001. In our study population, SBR grade III tumors are the most frequent, with a percentage of 56% of cases. Our results are consistent with the study of Tahari *et al.* [7], as well as with the Indonesian study in 2019 which reported a percentage of 68.48% of cases with histological grade III and 29.09% with SBR grade II [8].

In the literature, size T2 was the majority in several studies, in particular Ries *et al.* [9], who noted 60% of cases of T2 status, which is consistent with our results, unlike the Moroccan study of Bouzid and his team [10].

The N1 lymph node status in our population is in the majority; these statistical values are consistent with the study by Tahari Z and his team [7], who noted a high percentage of stage N1.

The presence of hormone receptors is a testament to the hormone dependence of breast cancer.

The IHC protein profile study revealed estrogen and progesterone receptor positivity in 68% and

Table II. Correlation of clinicopathological parameters with breast cancer subtypes

Molecular subtypes	Luminal A N = 16 n = %	Luminal B N = 18 n = %	HER2 neu N = 10 n = %	Basal-like N = 6 n = %	P-value
Mean age	64.25	61.55	56.2	61	
Mean tumor size [mm]	47.5	32.5	77.5	50	
Clinical stage of the tumor:					0.013
T1	25% (4)	33.33% (6)	40% (4)	0% (0)	
T2	37.5%	55.56% (10)	60% (6)	33.33% (2)	
T3	0% (0)	11.11% (2)	0% (0)	33.33% (2)	
T4	37.5% (6)	0% (0)	0% (0)	33.33% (2)	
Lymph nodes status:					0.117
Positive (+)	87.5% (14)	88.89% (16)	60% (6)	100% (6)	
Negative (-)	12.5% (2)	11.11% (2)	40% (4)	0% (0)	
Metastatic status:					0.219
Positive (+)	12.5% (2)	0% (0)	0% (0)	0% (0)	
Negative (-)	87.5% (14)	100% (18)	100% (10)	100% (6)	
SBR grad:					0.072
I	0% (0)	0% (0)	0% (0)	0% (0)	
II	37.5% (6)	55.56% (10)	60% (6)	0% (0)	
III	62.5% (10)	44.44% (8)	40% (4)	100% (6)	
Estrogen receptors:					
Positive (+)	100% (16)	100% (16)	0% (0)	0% (0)	
Negative (-)	0% (0)	0% (0)	100% (10)	100% (6)	
Progesterone receptors:					
Positive (+)	72% (12)	78% (14)	0% (0)	0% (0)	
Negative (-)	25% (4)	22% (4)	100% (10)	100% (6)	
HER2:					
Positive (+)	12.5% (2)	88.8% (16)	100% (10)	0% (0)	
Negative (-)	87.5% (14)	11.2% (2)	0% (0)	100% (6)	

56%, respectively. These results are of the same order as among the French (68% RE +) [11].

We also found 56% of cases that expressed progesterone receptors; these results are consistent with Ayadi's [12] study which found 70% of positive progesterone.

Our percentage of HER2 profile positivity correlates with several recently published studies, such as the Indonesian study of Swandari [8], which notes 57.61% positivity.

Our Ki67 biomarker results are lower than those obtained in the literature, in particular the Indonesian study [8] with a percentage of Ki67 > 14 which exceeds 83% of the total.

This study noted a predominance of the luminal B molecular subtype (36%). The result is similar to other studies established in regions of Southeast Asia (luminal B subtype 56.5%) [13].

Our results showed a majority incidence rate of the luminal A molecular subtype in our patients aged ≥ 70 years, the percentage of which was around 63%. Compared to the literature, our results agree with the Indian study established in 2019 [14], showed that the incidence rate of luminal A subtype was high in patients aged ≥ 70 years with a percentage of 72%.

The predominant histological grade of luminal B and HER2 type patients in this research was moderate grade (SBR II), and for luminal A and basal types it was high grade (SBR III), unlike Swandari's study [8], in which the patients with a luminal B and HER2 profile mainly present the SBR III grade, with a percentage of 66.7% and 80.6% respectively.

In our series, the HER2 subtype presents the largest tumor size (77.5 mm), followed by the basal-like type, with an average tumor size of 50 mm. Contrary to what has been found in the literature, notably the Saudi study published in 2019 which noted that the majority of tumor sizes (of the HER2 and basal-like subtypes) varied between 20 and 50 mm [6]. This can be explained by the difference in therapeutic management between countries or by the delay in diagnosis in Algeria.

Data obtained in our research indicate that luminal group B tumors would appear to have a worse prognosis and a very high potential for malignancy.

In conclusion, our knowledge of the heterogeneity of breast cancer at the histological, phenotypical and molecular level has been enriched thanks to the immunohistochemical analyses which have made it possible to approach the molecular classification by means of the genetic signatures described by Peru and his collaborators. The determination of molecular groups is therefore crucial for a better therapeutic strategy.

In our present study, the most frequent molecular phenotype is luminal B (36%) followed by lu-

minal A (32%), HER2 (20%) and finally basal-like (12%).

The results obtained, in which tumors of the HER2 and basal-like phenotype were associated with clinical and histopathological characteristics which are more aggressive than tumors of the luminal phenotype, are encouraging.

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Conflict of interest

The authors declare no conflict of interest.

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