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Female melanoma: the role of ovarian stimulation and antioestrogens cancer therapy

A retrospective study with immunohistochemical oestrogen receptors alpha and beta analysis.

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Thesis outline

This thesis is organised as follows: Chapter 1 outlines the research background, objectives and research scope. A comprehensive review of the literature regarding the melanoma sex disparities and the role of oestrogens and oestrogen receptors on female melanoma is discussed in Chapter 2. Details of the methods are discussed in Chapter 3, such as the statistical analysis of the population study and the characterisation techniques for oestrogen receptors immunohistochemical staining. Chapter 4 highlights the results of our research. In Chapter 5, a discussion on the findings of the research is carried out. Finally, Chapter 6 concludes the thesis and provide some recommendations for future prospectives.

Chapter 1

1.0 Research background:

Cutaneous melanoma (CM) is the most lethal type of cutaneous cancer¹. In the last two decades, the incidence of CM has significantly risen in fair-skinned populations². In 2019, it was estimated that 5.699 new CM cases occurred among females in Italy. CM represents the third most common cancer in Italian women under 49 years old³. The hypothesis that sex hormones can have a role in melanoma genesis was postulated based on two main epidemiologic observations: the difference in onset age of melanoma among genders and the better survival of women. In the last decades, many molecular studies performed on animal models and cell lines confirmed that sex hormones have a part in melanogenesis and melanoma genesis⁴. However, these findings seem not to correlate with epidemiological data directly, and many controversies are present regarding the role of exogenous oestrogens intake and endogenous hormonal status in female melanoma^{5,6}.

In recent years, oncology research focused its attention on the identification of oestrogen receptors in various tissues, some of them believed not to be responsive to sex hormones, but the site of indirect estrogenic action⁷. In some of these, such as the prostate and brain, the predominant oestrogen receptor is ER β (Estrogen Receptor beta). On the contrary, its reduced expression has been found in the tumorigenesis of other organs, including breast, colon and ovary⁸⁻¹⁰. ER β has an antiproliferative, proapoptotic action and is also involved in the regulation of the expression of numerous molecules of adhesion.^{11,12} As well-known, the skin is also an oestrogen-responsive tissue. Some descriptive studies documented the presence of ER β in the skin and as a predominant receptor in melanocytes both in dysplastic naevi and melanomas.¹³⁻¹⁵ Moreover, in melanoma ER β , seems to be inversely correlated with Breslow thickness in melanoma: in particular, thick melanomas (> 1 mm) display a lower expression of the receptor than thin ones (< 1 mm)^{16,17}. This inverse correlation suggests that the loss of ER β may influence the aggressiveness of melanoma and its progression, thus identifying a potential prognostic marker. However, the real significance of this correlation is still under investigation.

On the other hand, ER α was not detected with immunohistochemistry in benign or malignant melanocytic lesions even if its presence was revealed with mRNA reverse-transcriptase polymerase chain reaction¹⁷. For this reason, ER α presence in primary CM is still an unresolved issue.

In a preliminary analysis, we investigate, through immunohistochemistry, the presence of ER α and progesterone receptor (PR) in 28 female patients. Of these, 14 patients had received a diagnosis of melanoma (7 patients) or melanocytic nevus with severe atypia (7 patients) in pregnancy or the six months after childbirth or following repeated IVF courses in the previous year. The remaining 14 patients, of the same age and ethnicity, had removed a melanoma (7 patients) or a melanocytic nevus with severe histological atypia (7 patients). This latter did not have any history of hormonal stimulation or oral oestrogen-progestogen intake. Upon immunohistochemical analysis, we found that only two women showed a significant cytoplasmatic positivity for ER α . Upon immunohistochemical investigation, the CMs that were strongly positive were of the nodular type and with Breslow thickness> 1 mm: 1.2 mm and 1.4 mm, respectively. The remaining cases did not show positivity for ER α and PR¹⁸. These preliminary results may suggest the existence of a correlation in women between cytoplasmatic ER α and the onset of melanoma with a histopathological feature of aggressiveness, the nodular histotype.

1.1 Objectives

The primary objective of our study is to investigate the clinical and histopathological features of melanoma in this female population, focusing on two main subgroups: women in fertile age and women in menopausal age. We evaluated all female patients with histological diagnosis of cutaneous melanoma diagnosed from 01/2008 to 12/2018 at the Sant'Orsola-Malpighi Hospital and evaluated at least once at the Melanoma Unit of the Dermatology Department, University of Bologna.

The secondary objective is to evaluate the expression of ER α and ER β by immunohistochemical analysis (anti-ER α and anti-ER β antibodies) in this cohort. The investigation involves women who underwent to hormonal stimulations for medically assisted procreation (MAP), women in antioestrogens cancer therapy for breast cancer and two control groups matched for age and stage.

The tertiary objective is to evaluate the expression of ER β and ER α for each group in correlation with the Breslow thickness and other relevant histopathological (histotype, ulceration, regression, mitotic index), clinical (age, skin photodamaging, number of nevi, melanoma location) and dermoscopic characteristics (atypical network, blue-white veil, atypical vascular pattern, irregular streaks, irregular dots/globules, irregular blotches and regression structures).

1.2 Research scope

In this study, we planned to obtain data regarding the ERs presence in melanoma cells of women underwent to hormonal stimulations for medically assisted procreation (MAP) and women in antioestrogens cancer therapy. We look for any possible correlation between clinical and histopathological characteristics and these two different hormonal scenarios among our study population. These data will validate our preliminary results regarding the cytoplasmatic presence of ER α in the nodular melanoma of the patients with a history of ovarian stimulations. We also plan to verify if the expression of ER β by immunohistochemical analysis (anti-ER β antibodies) correlate inversely with the Breslow thickness or with other histopathological prognostic characteristics (histotype, ulceration, regression, mitotic index). Our main aim is to try to shed some lights on the oestrogen receptors presence and their role in female melanoma hormonal status.

2.0 Women melanoma and melanoma sex disparities

Melanoma has a different behaviour among gender regarding the age of incidence, body site affected and survival^{19,20}. In particular, women are more like to develop melanoma in the premenopausal age, while the incidence in the postmenopausal period is significantly lower. It is well known that females more commonly have melanoma on the lower extremities than men who often display it in the trunk, and this is true regardless of CM invasiveness²¹. Moreover, women have a better prognosis with a survival advantage over men²²; this makes the female gender an almost protective factor against death from melanoma²³. Indeed, this benefit in survival seems to be not linked the known prognostic factors such as Breslow thickness, mitoses, age at diagnosis, morphology, body part affected, ulceration, lymph node analysis as demonstrated in observational studies and clinical trials²⁴⁻²⁶. For these reasons, the female gender is considered worldwide an independent favourable prognostic factor for melanoma survival²⁷.

2.1 Behavioural factors

Many behavioural differences among gender were investigated to assess their role in these melanoma sex disparities. Differences towards sun exposure and health care were the main investigated factors. Women are generally more prone to sunbathing²⁸. On the other hand, men are more likely to adopt riskier behaviour concerning preventive care and medical compliance²⁹.

Even nutritional habits may have a role as shown in a recent work by Lukic M. et al., where the authors found that a moderate intake of filtered coffee reduces the risk of CM among a large cohort of Norwegian women³⁰.

Regarding the impact of ultraviolet (UV) exposure, a vast population study finds that UV is a significant causative factor for melanoma but only for women older than 44 years. In this study, the UV factor was not able to fully explain the incidence of melanoma in younger women³¹. Moreover, the latest findings suggest a relationship between UV index and male gender, not with female one³².

2.2 Genetic factors

Numerous recent studies indicated that these gender melanoma differences are more likely due to biological rather than the overcited behavioural factors, also considering that these disparities are observed worldwide, regardless of sun exposure levels.

Biological mechanisms proposed in melanoma sex differences are mainly linked to immune system actions, oxidative stress reaction ability, vitamin D metabolism, genetic sex chromosome expression and hormonal regulation³³.

Regarding genetics, it is meaningful to highlight the role of the selective inactivation of the X chromosome in female body cells that results in somatic mosaicism with a higher genetic diversification. X chromosome alone has many oncogenes and tumour suppressor genes that are involved in cancerogenesis^{33,34}.

Indeed, from childhood, there are sex differences in mole distribution, under genetic control in women, that reflect the distinctive melanoma distribution among sexes in adulthood^{19,35}. Epigenetic effects linked to X inactivation are likely to influence the nevi distribution³⁶. Moreover, in patients affected by Turner syndrome (XO) is expected the presence of a large number of moles on limbs and a higher risk for melanoma development ^{19,37}.

Furthermore, in the Spanish population, a sex-specific genetic effect was observed to be associated with sunlight sensitivity, pigmentation and melanoma risk³⁸.

Finally, the mutation burden of CM is significantly higher in male compare to female, mirroring a female immune system better capacity of effective antitumor surveillance³⁹.

2.3 Immune factors

Melanoma is a high immunogenic cancer, as also demonstrated by the higher risk for CM in patients with systemic immunosuppression included transplanted ones⁴⁰. For this reason, an important topic of sex disparities toward melanoma is related to the aged and sex-hormone changes in immune system⁴¹.

Especially after puberty in both adaptative and innate immunity, there are changes between males and females. These impact the incidence and pathogenesis of cancer as well as the efficacy of cancer immunotherapy. Ageing in the innate immune system, with its consequences, seems to be faster in men than in women ^{42,43}. Generally, Ig G and Ig M levels, CD4+ T cells, CD4/CD8 ratio and slightly CD 3+ cells T-lymphocytes are found higher in females; on the contrary, a lower Th1 response and IL-2 production characterised the males immune response⁴⁴⁻⁴⁵.

Sex hormones play a crucial role in the modulation and homeostasis of the immune system, considering that the ERs are expressed by T cytotoxic cells, B cells, dendritic cells and others. The exact mechanisms through which this interaction occurs are complex, as it also involves receptors other than ERs and it is still under investigation⁴⁶.

In mice model oestrogens enhances the production of molecules, through specific gene transcriptions, that are implicated in the prolonged survival and activation of B cells⁴⁷.

Oestradiol is the primary steroid molecule involved in the relationship between oestrogens and the human immune system. It is proved that oestradiol increases Ig G and Ig M levels in peripherical monocytes ^{33,48} and its concentration represent a regulator in the adaptative immune function as a low concentration elicits a Th1 response. Meanwhile, a high level evokes the humoral immunity and a Th2 response⁴⁹.

During the last decades with the beginning of the immunotherapy era in melanoma treatment, it has been found a relationship between sex hormones and the PD-1 signalling pathway. In particular, oestrogens can modulate the programmed cell death ligand 1 (PD-L1) and Treg-linked B7-H1 suppression, providing a better response through anticancer therapy with anti-PD-1 agents^{49,50}.

2.4 Hormonal factors

A large body of the literature maintains the positive function of oestrogen against melanoma progression. Sex hormones, particularly oestrogens, have been investigated as the main mechanism involved in the gender disparities, considering the difference in the cellular hormonal backgrounds. The exposure to oestrogens is physiologically part of the woman life, and it depends on reproductive factors: mainly menarche, parity and menopause. A relationship between endogenous hormones and melanoma is still debated.

Besides, the data from the published studies are often affected by the study design. The majority of them are retrospective cohort study that can have recall and selection bias.

The first works seem to prove a possible influence of female reproductive factors on melanoma⁵¹, although in the last few years, larger scientific studies do not show any relationship ^{5,52,53}. Indeed, recently Olsen CM et al.⁵⁴ have found no association between melanoma incidence and age at menarche, parity, menopausal status at baseline and age at menopause in their prospective cohort study on Australian women. A similar result was displayed by Støer NC et al.⁵⁵ within their large (172.478 women) nationwide population-based cohort in Norway.

Donley GM et al.⁵⁶ instead, demonstrated an increased melanoma risk associated with early age at menarche and late age at menopause, hypothesising that endogenous oestrogen exposure during childhood can increase during life photocarcinogenicity.

About parity, the literature is even more controversial: parity at a younger age is generally considered a protective factor as well as higher parity even if a bias linked to socio-economic confounders and individual lifestyle can be present^{5,52,57-60}.

Regarding benign gynaecological diseases and melanoma risk, there is no sufficient evidence considering the no-large number of epidemiological studies that investigated this topic ⁶¹. A prospective epidemiologic study found a statistically significant increase of melanoma risk among women with endometriosis (relative risk 1.62) and fibroma (relative risk 1.33)⁶². Recently these data were confirmed by the same study group finding also a higher risk for the non-melanoma skin cancers⁶³

The effect of pregnancy in melanoma is still an unresolved issue, especially concerning the disease prognosis during motherhood⁶⁴.

It is known that pregnancy is accompanied by numerous physiological changes, including skin hyperpigmentation in almost 85% to 90% women^{65,66}. In some early reports, changes in the moles have been observed during gestation. In particular, the most frequently observed modification concerned the colour of the nevi, which turned towards darker shades with pregnancy, and their size. These variations were in many cases, self-perceived and were reported in about 30% of women⁶⁷. Is recognised that there is no evidence of physiologic changes in nevi during pregnancy. The most visible changes are found in skin lesions located in the breast and abdomen, and they are due to their growth with normal skin expansion⁶⁸. Only one histopathological study found an increase in dermal mitoses and Ki-67 proliferation index among dermal nevi excised during pregnacy⁶⁹. There are prove of dermatoscopic changes, but they are transient, and none are suggestive of melanoma⁷⁰. In pregnant patients with a history of melanoma and multiple dysplastic nevi, dermatology examinations should be close also in the early post-partum⁷¹.

The scientific literature encompasses many epidemiological studies that looked at the relationship between pregnancy and CM. There are many limitations in data interpretation linked to the type of statistical analysis and study design, the number of participants, the definition of the so-called "pregnancy-associated melanoma" (PAM). Moreover, in case-control studies, age-matched control groups do not take into account confounding factors such as ulceration, melanoma staging, anatomic site of lesions, phototype and sun exposure⁷².

Concerning the CM diagnosis during pregnancy, a large body of literature do not reveal a poorer prognosis in pregnant patients⁷³⁻⁷⁶.

A large Swedish registry-based study compared mortality between PAM (defined as a period ranging from pregnancy to two years after child delivery) and women not diagnosed near childbirth: the cause-specific mortality did not differ between the two groups adjusting for age, period, education, parity, and tumour location (adjusted hazard ratio 1.09, 95% confidence interval 0.83-1.42)⁷⁷.

In a previous review by Lens M and Bataille V^5 of 10 case-control studies emerged that pregnancy does not impact melanoma survival and also not increase the risk of a second primary CM. Moreover, the authors

concluded that there is no reason for physicians to suggest a delay of a subsequent pregnancy in stage I melanoma patients. This advice is accepted in almost all European countries⁷⁸.

On the contrary, a recent single-institution study was PAM is defined as melanoma diagnosis during gestation or within one year after childbearing, maintains pregnancy is a negative factor with a five-, seven-, and ninefold increase in mortality, metastasis, and recurrence, respectively, when compared with controls ⁷⁹.

Conversely, in a large body of the literature, there are no findings that CM prognosis is changed when women melanoma is diagnosed up to five years following childbirth^{77,80,81}, except for one study that shows an increased risk of death in the first year postpartum⁸². A delay in diagnosis may cause this finding.

In women life, exogenous oestrogens sources are mainly represented by oral contraceptives (OCPs), hormone replacement therapy (HRT) and less frequently by ovarian stimulation for MAP.

Starting from the observation that acquired skin hyperpigmentation can also be present in women under OCPs⁸³ and that keratinocytes hold ERs⁸⁴, multiple studies have assessed the risk of melanoma associated with OCPs. Some works display an elevated risk, especially with prolonged use⁸⁵. These data can be partially explained by the behaviour through sun exposure and by the photosensitizing property of these drugs⁸⁶. Furthermore, a current work finds a positive association between the use of OC and HRT and one of the most important phenotypic risk factors of CM: a high melanocytic nevi count⁸⁷.

The two main meta-analyses of epidemiologic observational studies suggested no evidence for a higher melanoma risk with the use of OCPs^{52,88}. A pooled analysis of ten case-control studies by Karagas, M.R. et al.⁸⁹ displayed the same result: no relation between melanoma incidence and duration of oral contraceptive use was found even considering other confounder factors such as hair colour, sun sensitivity, family history of melanoma and sun exposure. The post-hoc analysis performed by the Women's Health Initiative trials on HRT in postmenopausal women shows no increased risk for melanoma and non-melanoma skin cancer⁹⁰. Meanwhile, the large EPIC (European Prospective Investigation into Cancer and Nutrition) prospective cohort study found a weak, statistically nonsignificant association between OCPs or HRT use and melanoma risk⁹¹. Besides, a recent case-control study by De Giorgi V et al⁹² shows that OCPs or HRT have even a protective effect against CM. A clear relationship between the duration of oral contraceptives intake and the age of onset and the incidence of melanoma is still debating.

The association between ovarian stimulation and CM risk is another controversial issue. Actually, its importance is increasing along with the increment of assisted reproductive techniques (ART) use. Worldwide almost 1% of birth are due to ART utilize⁹³. The cancer risk with other hormonal treatments is not comparable because of the high or very high (over ten-fold comparing to normal ovarian cycle⁹⁴) oestradiol levels, progesterone supplementation and antioestrogen (clomiphene) or gonadotrophins intake⁹⁵. A recent work ⁹⁶ analysing a large longitudinal cohort of US women (113,226 women, including 53,859 women without prior ART treatment), showed non-significant higher risks for CM after nearly five years of follow-up compared to the general population, considering also confounding effect of infertility. Most of the published studies had a

small number of cases, and consequently, they are not able to reach statistical significance. Two larger studies with respectively 42 and 112 participants displayed an almost 5% increased risk of CM among women using fertility drugs^{97,98}. A current review⁹⁹, examining potential associations between in vitro fertilization (IVF) and CM, do not reveal a consistent relationship between IVF and CM.

Nevertheless, the authors highlighted as the data indicates a potential increased risk for MM in ever-parous women undergone IVF. Some studies demonstrated an increased CM in women undergoing ovarian stimulation related to clomiphene citrate, a drug commonly used in controlled ovarian hyperstimulation^{100,101}. Meanwhile, the majority of works on fertility drugs and cancer showed no association with CM¹⁰²⁻¹⁰⁴.

Lastly, a possible scenario is characterized by antioestrogen cancer therapy intake. Indeed, epidemiological studies maintain that women with previous breast cancer (BC) have a higher risk of CM, and this risk is greater for patients who do not receive anti-estrogen therapy^{105,106}. Aromatase inhibitors (AIs) are both therapeutic and adjuvant agents in BC. Early studies found that the aromatase enzyme is expressed in melanoma, but no relationships were demonstrated between the presence of the enzyme and prognosis^{107,108}.

2.5 Estrogen receptors and melanoma

Estrogens are known to contribute to skin processes such as wound healing, thickening epidermis, mediating inflammatory disorders and preventing from photoaging ¹⁰⁹. Skin cells are targets of these hormones, and they are known to express ERs that are responsible for cutaneous estrogenic effects through their binding and the consequent activation of the two specific receptors, called estrogen receptors alpha (ER α) and estrogen receptor beta (ER β), which have similar structural and functional characteristics. ERs belong to the nuclear steroid hormone receptor superfamily and exert their role as ligand-activated transcription factor upon homoheterodimers. Structurally, ER α and ER β are respectively 595 and, 530 amino acids and their weight are respectively 67kDa and 59kDa. They are both characterized by DNA binding and a ligand-binding domains¹¹⁰(Figure 1).

In particular, the molecular structure is composed of:

- an N-terminal domain (A / B domain), highly variable both in sequence and in length, containing the AF1 region (Activation Function-1) a regulatory sequence¹¹¹; it includes a zinc-finger structure that is responsible for binding target sequences.

- a highly conserved central domain (DNA binding domain, DBD or C); it is crucial for chromatin binding.

- a hinge domain (D domain); important for translocation in the RE nucleus, it is unmasked upon oestrogens binding and, it connects the C and E domains. Moreover, it can bind the chaperone proteins.

- an E domain, which binds the hormone (ligand-binding domain, LBD).

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- an F domain at the C-terminal region. It contains the AF2 region (Activation Function-2) another regulatory sequence. Together with the E domain, it also has binding sites for coactivators and co-inhibitors.

AF1 can be active without the presence of the steroid hormones. The main architectural difference between the two receptors is that ER β has a shorter amino-terminal domain than ER α . The ERs are encoded by two different genes that are located on chromosome 6 for ER α and chromosome 14 for ER β .

Regarding ER α , besides the full-length isoforms, there are several shorten isoforms that are the results of alternative splicing or different start codons. There are unable by themselves to control genes transcription, but they can arrange heterodimers with the full-length form and inhibit its action. Meanwhile, the shorter isoform (ER- α 36) is responsible for membrane-initiated signaling¹¹².

On the other hand, ER β has five isoforms that differ from the full-length protein for the composition of the C-terminal region. ER β isoforms can abolish ER α function by forming a heterodimer with it¹¹³. All the ERs are mainly located in the nucleus, though some can be associated with the cell surface membrane. These latter are rapidly activated by estrogenic exposure. ER α and ER β are activated with different affinity, by various both natural and synthetic ligands.

The binding between intracellular ERs and their ligands determines the genomic pathway with nuclear translocation and the direct interaction with chromatin at specific target sequences, known as oestrogen response elements (EREs)¹¹⁴. Recent reports assess that in 35% of genes, involved in estrogenic biological effect, there is a lack in ERE-like sequences^{113,115}. In these the mechanism leading to genetic expression is called "transcriptional cross-talk", and it consists on an interaction between oestrogen receptor complexes and transcription factors such as the stimulating protein-1 (Sp-1) and the activator protein-1 (AP-1)^{116,117}. In the non-genomic pathway, instead, the binding is on the cellular membrane, and it triggers a cascade of intracellular signalling events leading to gene expression with or without a direct link between estrogenreceptor complexes and DNA. The changes in the cytosolic signalling, leading to the activation of four different protein-kinase cascades, including the RAS/BRAF/MEK axis^{118,119}. Strikingly, the MAPK pathway contains the molecules that are targeted by the BRAF inhibitors, such as vemurafenib, and the MEK inhibitors, such as dabrafenib, that are the first-line treatment for V600E mutant BRAF-harboring melanomas. To note, BRAF is known as the main protein driver in melanoma growth, occurring in about 50% of CM, especially in young patients with tumour located mainly on sun-exposed areas¹⁰⁸. The biological importance of the interaction between MAPK activation and ERa molecular mechanism is also demonstrated in vivo in animal models. Moreover, a large body of literature maintains strong communication between steroid hormones and growth factor receptors, having a great impact on cellular response to physiological or pathological situations^{108,120}.

The most involved estrogen receptor in *the non-genomic pathway* is the G protein-coupled estrogen receptor (GPER1), identified by molecular cloning methods and isolated almost two decades ago. GPER1 is a G protein-coupled membrane receptor that acts independently from ERs. Their agonists are G1 and 17β -estradiol¹²¹. It is crucial for many cellular processes, and it is also associated with cancers genesis and growing. Most important, in cutaneous tissue it is involved in melanogenesis regulation, and it is expressed in melanoma

cells^{108, 122,123}. In vitro GPER1 agonists are able to inhibit melanoma cell proliferation¹²⁴. Furthermore, when GPER1 is co-expressed with ER β in melanoma, the histopathological parameters are more favourable (lower Breslow thickness, lower mitotic rate and higher presence of peritumoral lymphocyte infiltration)¹²⁵.

There is growing evidence that alters function in estrogen receptors signalling may have a role in cancer genesis, development and evolution. In the same way, it is well known that ER α plays a part in tumorigenesis by increase cellular growth while, on the contrary, ER β has an antiproliferative action. ER α is the most representative ER in the human epidermis, and typically its presence is reduced in both benign and dysplastic nevi, but also in metastatic or primary melanoma, as well as in pregnancy-associated melanoma. It is usually rarely detectable with traditional immunohistochemistry analysis. Its amount in CM cells seems to not match with the pathophysiology of melanoma precursor lesions or melanomas themselves¹²⁶. In contrast, our preliminary immunohistochemical study showed a strong cytoplasmic localization of ER α in melanoma of women who experienced ARTs¹⁸.

Furthermore, the melanoma susceptibility, as well as the response to therapy, can be partially influenced by the presence of ER α single-nucleotide polymorphisms¹²⁷. At the same time, ER β is the predominant ER subtype in all melanocytic lesions: moles, dysplastic naevi and melanoma¹²⁸. In addition, ER β tends to diminish in vitro after UV exposure and is more present within thin melanocytic lesions such as naevi with severe dysplasia and in situ melanoma. ER β has been suggested as a CM marker¹²⁹ due to its exact inverse correlation with the most powerful CM prognostic factor: the Breslow thickness¹³⁰. Data regarding the potential protective function of ER β has also been found in breast, ovarian and prostate tumours as well as in colon cancer¹³¹⁻¹⁴⁰. An immunohistochemical study indicates that ER β is generally more present in women CM, especially pregnant ones, in comparison to men. The work of Schimdt et al. maintain these findings evaluating ER β expression in three categories: pregnant women, non-pregnant women, and men of patients stage and agematched¹³.

3.0 Materials and methods

3.1 Study design

A cross-sectional study is performed at Dermatology section, Department of Experimental, Diagnostic and Specialty Medicine, University of Bologna, Italy. Written informed consent was obtained for all study participants. The Ethics Committee approved the study of the Sant'Orsola-Malpighi Hospital and University of Bologna (protocol number 309/2019Sper.AOUBo).

3.2. Study population

We performed a retrospective analysis of patients with a histologically confirmed diagnosis of invasive or in situ CM received from 01/2008 to 12/2018 at the Sant'Orsola-Malpighi Hospital and evaluated at least once

at the Melanoma Unit of the Dermatology Department, University of Bologna. No restraints were applied to Breslow thickness.

The inclusion criteria were:

- Female gender
- Age >18 years

The exclusion criteria were:

- Patients with unknown primary CM
- Breslow thickness not available

For patients with multiple primary melanomas, the most invasive tumour was included in the analysis. Patient information was recorded through medical data. In particular, the following epidemiological data were evaluated in the study: age at first CM diagnosis, family history of melanoma, Fitzpatrick skin type, date of CM diagnosis, CM body site, previous excision of non-melanoma skin cancer, breast cancer personal history, age at menarche, hormone therapy (both OCPs and HRT) duration, number of pregnancies, age at first pregnancy, number of abortions, use of ARTs, age at menopause. Medical data taken from the patient records were as follows: Breslow thickness, number of mitoses, ulceration, histotype, type of growth, presence of satellite nodules, tumour-infiltrating Lymphocytes (TIL), regression, lymphovascular invasion, presence of metastasis. To investigate the clinical and histopathological characteristics of female melanoma in different women hormonal status, we divided our population study into two main subgroups: women in fertile age versus women within menopausal age.

The secondary objective was to evaluate the expression of ER α and ER β by immunohistochemical analysis (anti-ER α and anti-ER β antibodies) among selected patients of our study population.

3.3 Statistical analysis

All the statistical analyses were performed using R version 4.0.2 (https://www.r-project.org/). Individual and tumour characteristics were summarized through descriptive statistics, i.e. mean and standard deviation (SD) for quantitative variables, and absolute and percentage frequencies for categorical variables.

Tumour characteristics (all categorical) were compared between two different age groups (Age \leq 52 years and Age > 52 years) and between individuals with or without a medical history of breast cancer, by means of a Fisher exact test. A statistically significant association was defined as a p-value of less than 0.05.

Statistically significant associations were also visually displayed in the respective tables.

3.4 Patients selection for immunohistochemical analysis

The selection of patients applicable for immunohistochemical analysis of paraffin-embedded archival tissues was based on:

- Positive anamnesis for hormonal stimulations for medically assisted procreation (MAP) or antioestrogens breast cancer therapy intake
- Paraffin-embedded tissue-blocks available at the Dermatology Unit of the Policlinic Sant'Orsola-Malpighi
- Negative family history of melanoma
- No family cancer syndromes
- No BCRA1 or BCRA2 mutation detected upon breast cancer gene mutation screening

The investigation also involves other two control groups of women respectively on pre and postmenopausal status of the same age and melanoma staging.

3.5 Tissue samples and histological evaluation

The melanoma material included in the study were obtained by surgical excision carried out at the Skin Tumour Center and Melanoma Unit of the Dermatology Unit of the S. Orsola - Malpighi Polyclinic of Bologna.

The tissue, consisting of FFPE blocks, used for the study was that not dedicated to the routine diagnostic process and not reserved for molecular analysis in case of disease progression.

For histological evaluation, tissue samples had been fixed in 10% formalin, embedded in paraffin at pH 7.4 and stained with Haematoxylin and Eosin (H&E).

All the selected cases were analysed and re-staged by an expert pathologist according to 2018 WHO Classification of Skin Cancer and the 8th edition of the AJCC Cancer Staging Manual and the 2020 CAP melanoma reporting protocol ¹⁴¹⁻¹⁴³. The following clinical-pathological features of primary melanoma have been evaluated:

- Melanoma subtype
- Ulceration
- Growth phase
- Breslow thickness (mm)
- Mitotic count
- Tumour infiltrating lymphocytes-TILs
- Regression (<or >75%)

- Angioinvasion
- Perineural infiltration
- Folliculotropism
- Microsatellites
- Margin status
- pT staging

Two slides three µm thicker were obtained from each block and subsequently used for immunocytochemical receptor research.

Once the two slides were obtained for each patient, one for the detection of $ER\alpha$ and one for $ER\beta$ estrogen receptors, the material was transferred to the Unit of Anatomy and Pathological Histology of the S. Orsola - Malpighi Polyclinic where the immunohistochemical analysis was carried out.

3.6 Immunohistochemical staining and molecular analysis

The immunohistochemical analysis was conducted with the support of the VENTANA BenchMark ULTRA automated slide stainer (Ventana Benchmark Ultra System, Diagnostics Roche, Hoffmann-La Roche; Basel, Swiss) with the Opti View DAB IHC Detection Kit (Ventana).

Detection Kit is an indirect, biotin-free system for detecting specific primary mouse and rabbit antibodies bound to an antigen in paraffin-embedded tissue sections that are stained on the VENTANA automated slide stainers and visualized by light microscopy. It also uses a cocktail of enzyme-labelled secondary antibodies, which localize the bound of the primary antibody. The cocktail is recognized by an enzyme-bound tertiary antibody that is visualized with hydrogen peroxide substrate and 3, 3' – diaminobenzidine tetrahydrochloride (DAB) chromogen, which produces a brown precipitate detectable by light microscopy.

The staining protocol consists of several steps during which the reagents are incubated for predetermined periods of time and at specific temperatures. At the end of each incubation phase, the Ventana BenchMark Series instrument washes the sections in order to remove the unbound material and applies a liquid coverslip solution that minimizes the evaporation of the reagents acquired from the slide. The results are interpreted with an optical microscope and are useful for the differential diagnosis of pathophysiological processes, which may or may not be associated with a particular antigen.

In particular, the slides were deparaffinized in Bio-Clear (Bio-Optica, Milan, Italy) and hydrated with decreasing ethanol concentrations and distilled water. The immunostaining protocol included the antigen retrieval by multi-steps pre-treatment with CC1 solution (Ventana), the block of the endogenous peroxidases with 2.0% H202 in distilled water (5 minutes) and the incubation with the antibody for ERa (clone SP1,

Rabbit Monoclonal, Ventana, dilution ready to use-RTU, Ventana Medical Systems, Tucson, Arizona) and ER β (Clone ERb455, Mouse monoclonal, dilution ready to use-RTU, Ventana Medical Systems, Tucson, Arizona) respectively for 32' and 40' at 37° followed by DAB staining (Opti View DAB IHC Detection Kit, Ventana).

The precipitated chromogen indicates the presence of the antigen and the intensity of the reaction is proportional to the amount of antigen present.

If the antigens are present in the cytoplasm of the cells, the staining can involve the entire cytoplasm or only a part of it, depending on the antigenic content.

In our investigation, Nuclear and cytoplasmatic staining was separately evaluated and scored by the percentage of stained tumour cells; the staining was further classified as 0 ($\leq 20\%$), 1 (21%-50%), or 2 ($\geq 50\%$), as previously reported^{13,15,18}.

The most challenging aspect of immunohistochemistry is slide evaluation. In the interpretation of the results, both the specific antigen staining and the non-specific background staining must be evaluated. If the positivity depends on the localization of a particular antigen, various other factors, such as the condition of the tissue, the homogeneity of the fixation and the possible artefacts, can play an important role in an accurate interpretation.

In the ER α analysis, to avoid misleading, in the ten and four of the examined specimens there were positive (nuclear and cytoplasmatic) sebaceous lobules and/or hair follicles (basal cells of infundibulum and isthmus), respectively, and they served as the internal positive control. Conversely, breast carcinoma samples served as the external positive control.

Two cases have been molecularly characterized at our laboratory of oncologic and transplantation molecular pathology. DNA extraction and mutation analyses in oncogenes BRAF, KIT and NRAS with Sanger direct sequencing has been performed as previously described¹⁴⁴.

3.7 Dermatoscopy evaluation

The dermatoscopic evaluation of the select patients was performed by two expert dermatologists trained in dermoscopy. All the examined imagines were collected at the Melanoma Unit of the University of Bologna. The device used for obtaining both clinical and dermatoscopic images was FotoFinder Medicam® 800HD Dermoscope (Fotofinder Systems GmbH, Birbach, Germany), with a 1920 x 1080 resolution. Both polarized and non-polarized light was used for the pictures acquisition. Written consent was obtained by all patients for the use of images for research purpose. The dermatoscopic patterns analysed were atypical network, bluewhite veil, atypical vascular pattern, irregular streaks, irregular dots/globules, irregular blotches, regression structures, hyperpigmentation, distribution of pigment.

4.0 Results

4.1 Epidemiological analysis

We reviewed our database encompasses all melanoma patients evaluated at the Melanoma Unit at the University of Bologna, Italy from 2008 to 2018, finding 1054 females. From these, a total of 810 women with a previous diagnosis of CM met the inclusion criteria and were initially included in the cohort study. The main characteristics of the individuals and the CM were summarized in Table 1 and 2.

The median age at menarche was 12,16 in our population study; meanwhile, the average value of menopausal age was 52,47. The hormone therapy duration, including with this term, both OCPs and HRT, was 7,37 years. The mean value of pregnancies and abortions were respectively 1,25 and 0,36 with age at first pregnancy of 30,78 years. Of the available data, almost 30% reported a positive family history of CM and a total of 67 patients (6,91%) had a multiple primary CM. The females who had melanoma during pregnancy were 9 (1,1%), besides the cases of ovarian stimulation, as part of ARTs, were 6 in total (0,7%). Finally, women with a medical history of breast cancer were 24 (2,9%).

Regarding CM features, the most common location was the trunk and the legs with an identical percentage (almost 27%), following by arms (13,46%), head and neck (7,53%), acral location (4,20%) and finally the external mucosal site with a total of 3 cases (0,37%) in all our cohort. The majority of CM were thin melanomas with Breslow thickness (BT) less or equal to 1 mm (80,62%). The thicker CMs (\geq 2 mm, pT3 according to the 8th edition of the AJCC Cancer Staging Manual) represented just the 9,38% of all the cases. High mitotic index and ulceration were found in less than 8% of patients. The most common histotype was superficial spreading melanoma (ssm) (69,14%), followed by nodular melanoma (nm) (7,04%) and acral lentiginous melanoma (alm) (3,70%). A vertical cancer growth was identified in more than a quarter (25,56%) of the subjects. The TIL was absent in slightly more than half of the subjects (51,36%). Among the others, the brisk TIL was predominant (33,21%). Lymphovascular invasion and extensive regression (>75%) was respectively detected in about 3% and 4% of individuals.

Considering the mean age at menopause in our study population, in order to investigate differences between women in different physiological hormonal status, we divided the population study into two main subgroups: women aged over 52 and women having less than 52 years old. The two groups were both of 405 individuals, precisely 50% of our cohort. Table 3 displays a summary of the tumour characteristics distributed by age group. The location was significant divergent among the two subunits: in particular women in fertile age are more prone to have CM on the trunk. Conversely, females on the postmenopausal period are more likely to develop CM on arms, legs and acral location (Figure 2). The Breslow thickness was statistically (P-value <0,001) higher in women aged over 52 years, compared with the younger ones (Figure 3). Even if no statistical significance was observed regarding the mitotic index, a greater percentage (11,46% versus 6,16%) of postmenopausal females had a high (\geq 3) mitotic index (Figure 4). Furthermore, a statistical relevance (Figure 5) is noticed about CM ulceration, most common in postmenopausal age. Regarding the histotype not statistically meaningful was found, although older females are more likely to develop nodular melanoma (10,32% versus 6,67 %) (Figure 6).

Due to the insufficient number of cases, a statistical comparison between women who underwent to ovarian stimulation (a total of 6 persons) and the other women of the cohort, was not performed. The individual and cancer characteristics of females experienced ARTs are summarized together with immunohistochemical analysis in Table 6.

Concerning the group for women with a history of BC, when compared with the women without BC, we observed a significantly higher percentage of CM with "non-brisk" TILs (Figure 7).

4.2 Histological and Immunohistochemical analysis of selected cases

From a total of 30 selected cases with FFPE blocks accessible at the Dermatopathology unit of the University of Bologna, two patients, belonging to the control groups, were excluded from the analysis due to scarce biological material available. A total of 28 cases were analysed. According to the 2018 WHO Classification of Skin Tumour, the histological subtypes were 24 (85.7%) superficial spreading melanoma (SSM), 3 (10.7%) naevoid melanoma (NEM) and 1 (3.6%) acral lentiginous melanoma (ALM). The Breslow thickness ranged from 0 to 1.9 mm (mean: 0.69 mm). According to the 8th edition of the AJCC Cancer Staging Manual, the pT stages were 2 (7.1%) pTis, 17 (60.7%) pT1a, 2 (7.1%) pT1b, 6 (21.3%) pT2a, 1 (3.8%) pT2b. Sentinel lymph node biopsy was performed by nine patients (32,1%), in one case subclinical micrometastasis was discovered, and the patient underwent to radical lymphadenectomy, with no disease relapses at three years of follow-up. None of the investigated patients had distant metastasis from CM. The mean value of follow-up was four years (range: 2-10). No death from melanoma or other causes occurred.

All melanoma cases showed absent or focal nuclear positivity for ER α (21/28 cases with score 0; 7/28 cases with focal nuclear staining, range: 1-3%). The subjects with slight nuclear presence were three controls, three patients with antioestrogen therapy for BC and one woman with previous ARTs. At the opposite, 12/28 cases showed score 0 for cytoplasmatic ER α , whereas score 1 and 2 was both observed in 16/28 patients (Figure 8). The median value for cytoplasmatic ER α was 30.5%, with a median value of 25% and a range of 0-80%. The staining had a granular-like aspect. All cases with BC had a cytoplasmatic ER α positivity, except one. She was not under antioestrogen therapy at the time of CM onset, and recently, she has developed a third tumour: a clear cell renal cell carcinoma. A genetic mutation screening is actually in progress.

About ER β , positive nuclear expression (score 1 or 2) was detected just in 11/28 subjects, six belonging to the BC group, four controls and one with ovarian stimulation for infertility. Negative presence of ER β receptor characterized the majority of our patients, 17/28 cases. The nuclear staining had a wide range of

values (from 0 to 90%) (Figure 9). On the other hand, almost all of the examined melanomas 22/28 displayed a variable cytoplasmatic expression of ER β (mean value: 59.8%).

All the patient's characteristics and immunohistochemical results divided by groups are listed in Table 5 and 6.

Upon the Pearson correlation analysis, no association was shown between cytoplasmatic ER α and age or Breslow thickness. No statistical correlation was either found between nuclear ER β and age. Regarding nuclear ER β and Breslow thickness, the result was near to achieve the significance, although it did not reach any. (Figure 10)

No association emerged from ERs immunohistochemical expression and other histopathological such as regression, mitotic index or other histological parameters of CM.

The two cases with molecular characterization belong to the women who had ovarian stimulation, and both showed BRAF V600E mutation.

4.3 Dermatoscopic aspects

The dermatoscopic analysis was performed for all 28 patients. All patients belonged to phototype II according to Fitzpatrick classification. In the total of women with previous BC and ARTs, CM was characterized by a marked, dark brown and black atypical network and displayed irregular blotches (100%). Moreover, the presence of irregular streaks was common (71,4%) (Figure 11). Rarely the CMs in this group had an irregular pigmented distribution (28,6%) or an atypical vascular pattern (35,7%). Furthermore, generally, the atypical network was present in 89,3% of all subjects, although, in the control group, the network was less pigmented and irregularly distributed (78,57%). Finally, irregular dots/globules characterized the majority of the controls (92,8%) (Figure 12).

5.0 Discussion

The evidence clearly shows that women have a better outcome for many cancer types, including melanoma. Gender disparities in CM are present worldwide and over time despite dissimilar behaviours and latitudes. Furthermore, the female melanoma prognosis is still better than men when adjusting for the known melanoma prognostic factors¹⁴⁵. These data, reported in the literature, strongly suggest that in addition to cancer-specific characteristics, an important role is played by the host's gender. As well known, the incidence of CM has higher during fertile age, conversely is lower and associated with a poorer prognosis in menopausal era¹⁴⁶. Our cohort retrospective aimed to study female CM, considering both endogenous and exogenous oestrogen sources. Endogenous hormone basis was investigated through epidemiological studies examining the differences between women in fertile age and postmenopausal period, using as a cut-off the mean value of age at menopause. From this first analysis, we found a divergence in CM location: among younger women, CM are more likely to develop in the trunk, on contrary postmenopausal females are more

prone to have CM on extremities. The same result was already reported in a large population-based cohort study of 1347 Norwegian women. In their work, Støer NC et al⁵⁶ assess that this heterogeneity is probably caused by confounding factors as age, as CMs on the trunk are more common in younger people, mostly young women. Melanomas developing at different body sites are associated with distinct patterns of sun exposure. Whiteman DC et al. in 2006¹⁴⁷ stated that trunk melanomas are more likely associated with intermittent patterns of sun exposure, supporting his hypothesis of divergent causal pathways in CM¹⁴⁸. The two CM paths should be related respectively to chronic sun exposure and the presence of few naevi and to intense-intermitted solar exposure and higher mole density with a freckling tendency and a melanocyte instability¹⁴⁹. In addition, regarding UV radiation, it should also be considered the role that estrogens play in modulating melanogenesis and decreasing oxidative stress as demonstrated in many in vitro studies^{150,151}. Thick melanomas, ulceration and high (≥ 3) mitotic index was seen in older women. This more aggressive histopathological profile of CM is principally due to a selection bias and other confounding factors, such as histotype. Although, an influence of lower oestrogens level in down-regulation of immune processes cannot be totally excluded¹⁵². Moreover, it was observed that nodular melanoma is more often found in postmenopausal women, and this fact is confirmed in literature by the reflection of its higher incidence directly related to elder status among patients¹⁵³.

According to the investigation, we intend to realize, in our cohort study, we observed two different opposite situations concerning exogenous oestrogen sources.

One was represented by women who underwent ovarian stimulation for ARTs, and the other one was observed among women following antioestrogen therapy for breast cancer.

The first group observed was a tiny percentage (0,7%) compared to the total number of investigated cases. For this reason, a statistical analysis was not performed. Three of those subjects was treat by follitropin Alfa injection to stimulate ovarian follicular development. This drug was reported to have induced in one case of recurrent malignant melanoma¹⁵⁴. Nevertheless, there is no proven evidence that there is a higher melanoma risk after assisted reproduction⁹⁵

In the second group, there were a higher number of patients investigated, though the percentage was still not high (2,9%). On those patients, a statistical analysis was performed to have a comparison between the rest of the investigated cases. The emerging results lead to a statistical significance regarding the presence of "non-brisk" TILs. TIL has been considered in several studies that have contradictory results on its meaning in the CM prognosis. A recent study by Sinnamon AJ et al¹⁵⁵ showed that TIL is a predictive factor of sentinel lymph node positivity among men, but no association was found in women. In our study, an association between the non-brisk pattern and the group of women with previous BC was observed. According to our finding, further investigations must be needed to evaluate whether different cancers on the same individual can have a similar TILs composition or not. It could be necessary to examine also TILs meaning related to disease prognosis and patients' treatment response¹⁵⁶.

In addition to the epidemiological investigation we have discussed, it was also performed immunohistochemical analysis to evaluate the presence of ER α and ER β among melanoma cells of the women undergone ovarian stimulation for ARTs and of the women following antioestrogen therapy for breast cancer. Results emerging from this analysis show that all women assuming aromatase inhibitors therapy have a widespread cytoplasmatic expression of ER α in melanoma cells. In a previous study of our group, we found a similar result in women with nodular melanoma treated with multiple cycles of hormonal stimulation for in vitro fertilization⁵. The meaning of cytoplasmatic ERa in melanoma is still debated. It could suggest a potentially significant role of oestrogen non-genomic pathway in these patients¹⁵⁷, or it can be a mechanism of ERs modulation in response to aromatase inhibitor therapy¹⁵⁸. Finally, the reason of its presence in a granular shape can be just the evidence of a false positive due to the residual endogen peroxidases, even if endogen peroxidases were accurately inhibited following our protocol for immunohistochemical staining. Concerning ER β , our study shows no correlation between nuclear positivity and Breslow thickness values, contrarily to what was displayed in literature. Undoubtedly the use of different anti-ERß antibodies can partially affect the immunohistochemical analysis. In our study, we used an antibody directed towards isoform 1, that is favoured in many studies¹⁵⁹. Eventually, a dermatoscopic analysis was performed on all patients selected for immunohistochemical analysis to investigate the clinical aspect of their CMs. As it was shown in literature by Auriemma et al. ¹⁶⁰, our work also evidences, among the two particular patients subgroups a darker brown atypical network and irregular blotches that clinically characterized the CM in comparison to control groups.

Our investigation is limited by a number of factors, first including the small amount of available cases. For these reasons, it can be subjected to many biases. However, our work tried to enlighten some of the existing shadows on the role of ERs and hormonal factors in CM in a real-world setting. Based upon our direct experience, confirmed by literature, it can be affirmed that ERs and hormonal factors play a marginal role in the aetiology of female melanoma. Instead, sex-specific genetics and immunity have a leading effect on melanoma genesis, development and progression¹⁶¹.

6.0 Present and future perspective

Drugs target ERs are still under investigation as a therapy for CM, despite tamoxifen (TAM), a non-selective ER antagonist, was first observed, in clinical trials, to have no relevant effect on CM survival rate¹⁶². This finding is probably due to TAM proprieties: at the same time antitumoral and pro-survival agent, depending on ER α /ER β ratio in the target tissue¹⁶³. Moreover, TAM has also demonstrated an agonist action towards GPER in vitro¹²⁴ It is recently showed that endoxifen, a 100 times more powerful active metabolite of TAM, can be a promising new therapeutic agent against CM by blocking ER transcription¹⁶⁴. Besides, selective ER β agonist, LY500307, has been observed to have efficacy against lung metastasis from CM, both in vitro and in vivo. It acts in increasing the innate immunity's ability to suppress cancer through the production of IL-

1β. This latter is a chemotactic molecule produced by cancer cells after the stimulation of their ERβs. This molecular mechanism is another example of the primary role that estrogen signalling can play in CM and how much it can be significant for future anticancer pharmacological research¹⁶⁵. GPER also has a relevant part in immunomodulation operated by estrogen signalling. In 2018 Natale A. et al.¹⁶⁶ proved that GPER action prevents tumour spreading and selects a cancer immune profile that makes CM more vulnerable to immune checkpoint blockade agents.

Finally, an emerging research field is represented by the connections between microRNAs (miRNAs) and estrogen receptor pathways.

Several studies found that miRNAs dysregulation/ specific disease profiles are common in cancer cells, and they are implicated in tumour development, progression but also cancer suppression^{167,168}. In particular, miRNAs expression has demonstrated a regulatory effect on ERs expression. Specifically, miR-206, miR-22 and let-7 have demonstrated to downregulate ER α levels in breast cancer, with a consequent proapoptotic and antiproliferative effect ¹⁶⁹⁻¹⁷¹. Besides estrogen-induced miR-196a expression has a prognostic value in hormone therapy, responsive breast cancer¹⁷². On the other hand, ER α -mediated estrogen signalling downregulates the level of miR-21, miR-26a, miR-140, miR-181b and overexpress miR-206, and miR-190a, miR-191, miR-203 and miR-425¹⁷³. Regarding ER β , less is known. Several studies show that it is closely related to suppression in miR-17, miR-30a, miR-200a and miR-200b expression, and it upregulates of miR-23b, miR-24-1 and miR-27b¹⁷⁴. Recently also some transcriptional factors have been discovered to recognize the DNA ERE-elements and activate miRNAs transcription depending on ERs action¹⁷⁵. The interaction between tissue specificity of miRNA expression and ERs and GPER is even more complex and still under investigation also in endothelial cells¹⁷⁶.

In melanoma, a key role in disease spreading and progression has been proposed for miR-221 and -222 expression, negatively regulated by ER α . Furthermore, miR-221/222 modulate directly KIT oncogene that is typically mutated in acral and mucosal melanoma^{177,178}.

More is to investigate in the composite scenario of ERs, GPER, miRNA expression and their effect on the melanoma microenvironment.

7.0 Tables

Table 1 and Table 2 summarizes, respectively, individual and tumour characteristics.

| Table 1. mulvidual characteristics. | |
|--|--------------|
| Age at diagnosis, years, n (%) | |
| ≤52 | 405 (50.00) |
| >52 | 405 (50.00) |
| Phototype, n(%) | |
| 1 | 50 (6.17) |
| 2 | 548 (67.65) |
| 3 | 212 (26.17) |
| Photodamage, n(%) | |
| No | 191 (23.59) |
| Yes | 619 (76.41) |
| Familiarity, n(%) | |
| No | 578 (71.35) |
| Yes | 232 (28.65) |
| Other cutaneous tumours, n(%) | |
| No | 386 (63.69) |
| Yes | 220 (36.30) |
| NA | 204 |
| Previous breast cancer, n(%) | |
| No | 786 (97.03) |
| Yes | 24 (2.96) |
| Age at menarche, years, mean (sd) | 12,16 |
| Assisted procreation, n(%) | |
| No | 804 (99.25) |
| Yes | 6 (0.74) |
| Hormone therapy, n(%) | |
| No | 183 (36.38) |
| Yes | 320 (63.62) |
| NA | 307 |
| Hormone therapy duration, years, mean (sd) | 7.37 (5.86) |
| Number of pregnancies, mean (SD) | 1.25 (0.97) |
| Age at first pregnancy, years, mean (SD) | 30.78 (6.41) |
| Number of abortions, mean (SD) | 0.36 (0.76) |
| Age at menopause, years, mean (SD) | 52.47 (3.25) |

Table 1. Individual characteristics

Table 2. Tumour characteristics.

| Location, n(%) | |
|----------------|-------------|
| Head / neck | 60 (7.53) |
| Trunk | 221 (27.28) |
| Arms | 109 (13.46) |
| Legs | 219 (27.04) |

| Acrals | 34 (4.20) |
|---------------------------|------------------------|
| Mucous membranes | 3 (0.37) |
| NA | 164 (20.25) |
| Tumour thickness, mm, | . , |
| mean (sd) | |
| ≤1 | 653 (80.62) |
| (1.2] | 76 (9.38) |
| >2 | 76 (9.38) |
| NA | 5 (0.62) |
| Mitotic index, mean (sd) | - () |
| 0 | 497 (61 36) |
| 1-2 | 132 (16 29) |
| >3 | 61(753) |
| NA | 120(14.81) |
| 1NA | 120 (14.01) |
| | (72)(92)(0) |
| INO Vez | 6/3(83.09) |
| Yes | 04 (7.90) 72 (0.01) |
| | /3 (9.01) |
| Histotype, n(%) | |
| ssm | 560 (69.14) |
| nm | 57 (7.04) |
| lm | 9 (1.11) |
| lmm | 8 (0.99) |
| alm | 30 (3.70) |
| desmoplastic | 1 (0.12) |
| rare type | 4 (0.49) |
| NA | 141 (17.41) |
| Type of growth, n(%) | |
| Radial | 476 (58.77) |
| Vertical | 207 (25.56) |
| NA | 127 (15.67) |
| Satellite nodules, n(%) | |
| No | 620 (76.54) |
| Yes | 8 (0.99) |
| NA | 182 (22.47) |
| TIL, n(%) | () |
| No | 416 (51.36) |
| Brisk | 269 (33.21) |
| Non-brisk | 112(13.83) |
| NA | 13(1.61) |
| I ymphoyascular | 15 (1.01) |
| invasion n(%) | |
| No | 636 (78 52) |
| Ves | 26(3.21) |
| | 20(3.21) 148(1827) |
| $\mathbf{D}_{\mathbf{A}}$ | 140 (10.27) |
| N ₁ | 100 (50 27) |
| INO Vez | 408(30.37) |
| Y es | 230 (29.14) |
| >/5% | 33 (4.07) |
| NA | 133 (16.42) |
| Metastases, n(%) | |
| No | 585 (72.22) |
| Yes | 100 (12.35) |
| NA | 125 (15.43) |

| | Age ≤ 52 | Age > 52 | P-value |
|--------------------------|---------------------------|--------------------------|---------|
| Location, n(%) | | | <0.001 |
| Head / neck | 10 (3.15) | 50 (15.20) | |
| Trunk | 137 (43.22) | 84 (25.53) | |
| Arms | 49 (15.46) | 60 (18.24) | |
| Legs | 107 (33.75) | 112 (34.04) | |
| Acrals | 13 (4.10) | 21 (6.38) | |
| Mucous membranes | 1 (0.32) | 2 (0.61) | |
| Tumour thickness, mm, | | | |
| mean (sd) | | | <0.001 |
| ≤1 | 338 (83.87) | 315 (78.36) | |
| (1,2] | 43 (10.67) | 33 (8.21) | |
| >2 | 22 (5.46) | 54 (13.43) | |
| Mitotic index, mean (sd) | | | 0.040 |
| 0 | 250 (73.31) | 247 (70.77) | |
| 1-2 | 70 (20.53) | 62 (17.77) | |
| ≥3 | 21 (6.16) | 40 (11.46) | |
| Ulceration, n(%) | | | 0.001 |
| No | 347 (94.81) | 326 (87.87) | |
| Yes | 19 (5.19) | 45 (12.13) | |
| Histotype, n(%) | | | 0.037 |
| ssm | 290 (87.88) | 270 (79.65) | |
| nm | 22 (6.67) | 35 (10.32) | |
| lm | 2 (0.61) | 7 (2.06) | |
| lmm | 1 (0.30) | 7 (2.06) | |
| alm | 13 (3.94) | 17 (5.01) | |
| desmoplastic | 0 (0.00) | 1 (0.29) | |
| rare type | 2 (0.61) | 2 (0.59) | |
| Type of growth, n(%) | | | 0.740 |
| Radial | 232 (69.05) | 244 (70.32) | |
| Vertical | 104 (30.95) | 103 (29.68) | |
| Satellite nodules, n(%) | | | 0.287 |
| No | 304 (99.35) | 316 (98.14) | |
| Yes | 2 (0.65) | 6 (1.86) | |
| TIL, n(%) | | 010 (50 (0) | 0.920 |
| No | 206 (51.76) | 210 (52.63) | |
| Brisk | 137 (34.42) | 132 (33.08) | |
| Non-brisk | 55 (13.82) | 57 (14.29) | 0.070 |
| Invasion, n(%) | 221 (07 57) | 215(0450) | 0.070 |
| No | 321 (97.57) | 315 (94.59) | |
| Y es | 8 (2.43) | 18 (5.41) | 0.202 |
| Regression, n(%) | 107 (59.91) | 211((1.70)) | 0.382 |
| INO Vac | 197 (38.81) | 211(01./0) 118(24.50) | |
| 1 es | 118(33.22) | 118(34.30) 12(2.80) | |
| >/3% | 20 (5.97) | 15 (3.80) | 0 706 |
| No | 112 (75 72) | 410 (70.00) | 0./80 |
| | $\frac{443}{142} (13.13)$ | 410(70.09) 175(20.01) | |
| 1 68 | 142 (24.27) | 173 (29.91) | |
| | | | |

Table 3. Tumour characteristics by age group. Significant p-values are in bold.

| | No breast cancer | Breast Cancer | P-value |
|--------------------------|------------------|-------------------------|---------|
| Location, n(%) | | | 0.142 |
| Head / neck | 59 (9.41) | 2 (8.33) | |
| Trunk | 210 (33.49) | 12 (50.00) | |
| Arms | 105 (16.75) | 5 (20.83) | |
| Legs | 217 (34.61) | 3 (12.50) | |
| Acrals | 33 (5.27) | 2 (8.33) | |
| Mucous membranes | 3 (0.48) | 0 (0.00) | |
| Tumour thickness, mm, | | | |
| mean (SD) | | | 0.507 |
| ≤1 | 566 (74.73) | 19 (79.17) | |
| (1,2] | 95 (12.58) | 4 (16.67) | |
| >2 | 96 (12.69) | 1 (4.17) | |
| Mitotic index, mean (sd) | | | 0.272 |
| 0 | 480 (71.64) | 19 (79.17) | |
| 1-2 | 131 (19.55) | 2 (8.33) | |
| ≥3 | 59 (8.21) | 3 (12.50) | |
| Ulceration, n(%) | 0) (0.21) | 5 (12.00) | 1.000 |
| No | 654 (91.21) | 22 (91.67) | 11000 |
| Yes | 63 (8 79) | 2 (8.33) | |
| Histotype n(%) | 00 (0179) | 2 (0.00) | 0.940 |
| ssm | 538 (83.41) | 22 (91.67) | 0.0 10 |
| nm | 55 (8 53) | 2 (8.33) | |
| lm | 9(1.40) | 0(0.00) | |
| lmm | 8 (1 24) | 0(0.00) | |
| alm | 30 (4.65) | 0(0.00) | |
| desmonlastic | 1(0.15) | 0(0.00) | |
| rare type | 4(0.62) | 0(0.00) | |
| Type of growth $n(\%)$ | 1 (0.02) | 0 (0.00) | 0.213 |
| Radial | 456 (69 20) | 20 (83 33) | 0.215 |
| Vertical | 203 (30.80) | 4(1767) | |
| Satellite nodules n(%) | 203 (30.00) | 1(17.07) | 1 000 |
| No | 606 (98 70) | 24(100.00) | 1.000 |
| Ves | 8 (1 30) | 0(0.00) | |
| TIL $n(\%)$ | 0 (1.50) | 0 (0.00) | 0.001 |
| No | 406 (52 52) | 10 (41 67) | 0.001 |
| Brisk | 265 (34 28) | 4(16.67) | |
| Non-brisk | 102 (13 20) | 10(41.67) | |
| Invasion $n(\%)$ | 102 (15.20) | 10 (41.07) | 1 000 |
| No | 612 (94 44) | 24(100.00) | 1.000 |
| Ves | 36 (5 56) | 0(0.00) | |
| Regression n(%) | 50 (5.50) | 0 (0.00) | 0 486 |
| No | 395 (60 49) | 13 (54 17) | 0.700 |
| Ves | 225 (34 46) | 11(45.83) | |
| >75% | 223 (34.40) | 0 (0 00) | |
| Vetastases n(%) | 55 (5.05) | 0 (0.00) | 0 3 3 0 |
| No | 565 (85 18) | 20 (82 22) | 0.330 |
| | 06 (11 52) | 20 (03.33) 1 (16 67) | |
| 1 05 | 90 (14.32) | 4(10.07) | |

Table 4. Tumour characteristics by the presence of breast cancer. Significant p-values are in bold

Table 5. Clinical, histological and immunohistochemical data of patients with breast cancer included in the study.

| Patient number | Age at melanoma diagnosis | Age at breast cancer diagnosis | Drug therapy for breast cancer | Duration of breast cancer therapy (years) | Photodamaging | Number of nevi | Melanoma location (0=extremities, 1=trunk and dorsum, 2=head neek, 3=other) | Melanoma histotype (0=superficial spreading melanoma, | Breslow thickness | ERα N in % | ERα N (0<20%, 1=21%- 50%, 2>50%) | ERα C in % | ERα C (0<20%, 1=21%- 50%, 2>50%) | ERβ N in % | ERβ N (0<20%, 1=21%- 50%, 2>50%) | ERβ C in % |
|-------------------|---------------------------------|---|--|---|---------------|-------------------|--|---|----------------------|------------------|--|------------------|--|------------------|--|------------------|
| | | | | | | | licex, 5 other) | melanoma) | | | | | | | | |
| 1 | 69 | 67 | Aromatase inhibitor | 1 | yes | 50-100 | 0 | 0 | 0.4 | 2 | 0 | 55 | 2 | 60 | 2 | 90 |
| 2 | 65 | 64 | Aromatase inhibitor | 1 | yes | <50 | | | 0.6 | 2 | 0 | 60 | 2 | 80 | 2 | 80 |
| 3 | 63 | 62 | (Anastrozole) Aromatase inhibitor (Anastrozole) | 1 | yes | <50 | 1 | 0 | 0.3 | 0 | 0 | 55 | 2 | 80 | 2 | 20 |
| 4 | 48 | 50 | Aromatase inhibitor | 2 | No | 20-30 | 1 | 0 | 0.2 | 0 | 0 | 25 | 1 | 5 | 0 | 20 |
| 5 | 44 | 52 | Trastuzumab and aromatase inhibitor | 1 | No | <50 | | 0 | 0 | 0 | 0 | 60 | 2 | 80 | 2 | 80 |
| 6 | 58 | 49 | (Anatrozole) Interferon at CM diagnosis, then letrozole | 1 | Yes | 5 | 1 | 0 | 0,2 | 0 | 0 | 5 | 0 | 10 | 0 | 70 |
| 7 | 51 | 53 | Aromatase inhibitor (Lestrozole) | 2 | No | 50-100 | 1 | 0 | 1,3 | 2 | 0 | 70 | 2 | 10 | 0 | 90 |
| 8 | 81 | 81 | Aromatase inhibitor (Anastrozole) | <1 | yes | <20 | 2 | 0 | 0.6 | 0 | 0 | 80 | 2 | 60 | 2 | 90 |
| 9 | 63 | 62 | Aromatase inhibitor (Anastrozole) | 1 | yes | 30 | 1 | 0 | 0.3 | 0 | 0 | 40 | 1 | 60 | 2 | 60 |

| Patient number | Age at melanoma diagnosis | Age at ovarian stimulation | Drug therapy for ovarian stimulation | Cycles of ovarian stimulation | Successful pregnancy (yes/no) | Photodamaging | Number of nevi | Melanoma location (0=extremities, 1=trunk and dorsum, | Melanoma histotype (0=superficial spreading melanoma, | Breslow thickness | ERα N in % | ERα N (0<20%, 1=21%- 50%, 2>50%) | ERα C in % | ERα C (0<20%, 1=21%- 50%, 2>50%) | ERβ N in % | ERβ N (0<20%, 1=21%- 50%, 2>50%) | ERβ C in % |
|-------------------|---------------------------------|----------------------------------|---|-------------------------------------|-------------------------------------|---------------|-------------------|---|---|----------------------|------------------|--|------------------|--|------------------|--|------------|
| | | | | | | | | 2=head neck, 3=other) | 1=naevoid melanoma 2= acral melanoma) | | | | | | | | |
| 10 | 39 | 38 | Progesterone, others | 1 | yes | no | 30-50 | | | | | | | | | | |
| | | | unknown | | | | | 1 | 1 | 1.2 | 0 | 0 | 10 | 0 | 0 | 0 | 20 |
| 11 | 48 | 40 | Unknown | 1 | yes | yes | <10 | 1 | 0 | 0.2 | 0 | 0 | 25 | 1 | 5 | 0 | 20 |
| 12 | 43 | 34 | Uncomplete ovarian | 1 | yes | yes | <20 | | | | | | - | | - | | |
| - 10 | | | stimulation | | | | | 0 | 2 | 1 | 0 | 0 | 5 | 0 | 5 | 0 | 55 |
| 13 | 34 | | Follitropin alfa, triptorelin acetate injection, menotropins, canicalix | 3 | no | yes | >50 | 0 | 0 | 1.2 | 2 | 0 | 30 | 1 | 5 | 0 | 100 |
| 14 | 43 | | Follitropin alfa, ganirelix, chorionic gonadotropin, progesterope | 1 | yes | yes | >50 | 0 | 1 | 1.2 | 0 | 0 | 15 | 0 | 90 | 2 | 30 |
| 15 | 30 | 28 | Follitronin | 2 | 20 | Vec | 50-100 | <u> </u> | - | 1.5 | 0 | | 15 | 5 | ,,, | | 50 |
| 15 | 50 | 20 | alfa, | 2 | 10 | yes | 50-100 | | | | | | | | | | |
| | | | ganirelix, | | | | | | | | | | | | | | |
| | | | cnorionic | | | | | | | | | | | | | | |
| | | | progesterone | | | | | 1 | 0 | 0.6 | 0 | 0 | 0 | 0 | 0 | 0 | 70 |

Table 6. Clinical, histological and immunohistochemical data of patients with ovarian stimulation included in the study.

| Dermoscopic features | Cases | Controls | |
|--|---------------|----------------|--|
| | | | |
| Atypical pigmented | 14/14 (100%) | 12/14 (85,7%) | |
| Blue-white veil | 7/14 (50%) | 2/14 (14,3%) | |
| Atypical vascular pattern | 5/14 (35,7%) | 7/14 (50%) | |
| Irregular streaks | 10/14 (71,4%) | 9/14 (64,3%) | |
| Irregular dots/globules | 8/14 (57,1%) | 13/14 (92,8%) | |
| Irregular blotches | 14/14 (100%) | 7/14 (50%) | |
| Regression structures | 7/14 (50%) | 8/14 (57,1%) | |
| Irregular pigment | 4/14 (28,6%) | 11/14 (78,57%) | |
| Pigmentation in more than 50% of the CM | 9/14 (64,3%) | 6/14 (42,8%) | |

Table 7. Dermatoscopic features of CM in patients with ovarian stimulation and antioestrogens cancer therapy versus control groups

8.0 Figures



Figure 1 Molecular structures of estrogen receptors

Figure 2. Location distribution by age group.



Location



Figure 3. Breslow thickness distribution by age group.

Figure 4. Mitotic index distribution by age group.







Figure 6. Histotype distribution by age group.





Figure 7. TIL distribution by the presence of breast cancer.

Figure 8. Diffuse cytoplasmatic positivity for ERα in the melanoma cells (original magnification ×100)



Figure 9. Nuclear positivity for ERβ in the melanoma cells of the same case (original magnification ×100)





Figure 10. The scatter plot of the data, indicating the reasonableness of assuming a linear association between the variables.

Figure 11. Cutaneous melanoma of the trunk in a patient underwent to assisted reproductive treatment



Figure 12. Cutaneous melanoma of the trunk in a patient belonging to the control group



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