

# Diagnostic Histological Pitfall of Dedifferentiated Melanoma

Subjects: Allergy

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Dedifferentiated melanoma is a particular form of malignant melanoma with a progressive worsening of the patient's clinical outcome.

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## 1. Introduction

Classically, melanoma has always been considered as “the great mime” for its intrinsic ability to disguise itself in different guises and imitate other types of neoplastic and non-neoplastic lesions <sup>[1]</sup>. This peculiarity constitutes the basic reason why the routine dermatopathologist must always resort to adequate immunohistochemical markers to exclude or confirm the diagnosis of malignant melanoma <sup>[1][2]</sup>. All this is not true in the case of a particular form of malignant melanoma, defined by various authors as “Dedifferentiated Melanoma” (DM) due to the characteristic of losing some or all of the melanocytic immunohistochemical markers <sup>[3]</sup>. The clinical and histopathological difficulty in recognizing and correctly diagnosing this entity has already been previously reported, although only in recent years has the advent of molecular biology and next generation sequences (NGS) contributed, in a fundamental way, to a better understanding of the dedifferentiated phenotype.

## 2. Diagnostic Histological Pitfall of Dedifferentiated Melanoma

The potential diagnoses were basically two: a “collision” lesion consisting of a melanoma and a malignant fibrohistiocytic neoplasm (such as for example the Atypical Fibroxanthoma), or a malignant melanoma that underwent almost total dedifferentiation, so as to almost entirely lose the common markers of melanocyte differentiation. A careful analysis and integration of morphology and immunohistochemistry allowed to describe the presence of focal clusters of SOX-10 and S-100 protein positive melanocytes within the nodular lesion proper (area of transition). For this reason, this lesion was diagnosed as entirely melanoma with a dedifferentiated component, strongly expressing CD10.

Furthermore, for definitive confirmation, we performed Next Generation Sequencing analyses at an external center that revealed BRAFV600K mutation, supporting the diagnosis of dedifferentiated melanoma.

By virtue of the rarity and the diagnostic challenges that DM poses, we have conducted a careful review of the current literature, in order to improve the characterization, understanding, and knowledge of this potential diagnostic pitfall.

A malignant melanoma may be able to simulate various and different neoplasms: the potential of being misunderstood with other malignancies is well known <sup>[1][2]</sup>. In this corollary, dedifferentiated melanoma turns out to be a very aggressive form, with little tendency to medical response <sup>[3]</sup>. Although rare, the dedifferentiation mechanism has been studied in different sets of neoplastic pathologies, and it is recognized quite clearly that in addition to creating difficulties in the correct histopathological diagnosis of the entity, DM poses problems of therapeutic response both to traditional therapy and to immunotherapy (so-called cross-resistance) <sup>[4]</sup>. Although the histological diagnosis of malignant melanoma is known to be difficult, this is even more true in the case of DM, as the morphological characteristics on the one hand and the loss of one or all of the markers of melanocytic differentiation (such as Melan-A, HMB-45, SOX-10 and MITF), poses major diagnostic challenges for, among others, high-grade sarcomas or carcinomas <sup>[5][6]</sup>. Additionally, in our case, the dedifferentiation within the morphologically characterizable malignant melanoma created some diagnostic questions regarding the possibility that it was a collision lesion rather than a melanoma with a real portion of dedifferentiation. This issue has been extensively discussed in the literature <sup>[5][6][7][8]</sup>, and a case has been reported very recently by *Saldana et al.* <sup>[9]</sup> in which the amplification in FISH of the MDM2 gene was described for the first time in a lesion of a 73-year-old

subject, which had led, in the first instance, to hypothesize a liposarcoma. A careful analysis involving also the determination of BRAFV600 allowed to reach the correct diagnosis of DM.

In the last 10 years, various authors have tried to shed light on what may be the potential biological pathways that melanocyte cells follow until they lose the common immunohistochemical markers [4][5][6][7][8][9][10][11][12][13][14][15][16][17][18][19][20][21][22][23][24][25][26][27][28][29][30][31][32]. Therefore, the concept of “phenotypic plasticity” of melanocyte cells has been developed, as it has been demonstrated that the microenvironment where melanocytes operate is able to bi-directionally influence the following phenotype: in particular, the study of melanocytes has been deepened, inducing the transcription factor (MITF) whose expression (also detectable in immunohistochemistry) was correlated with a different biological behavior of melanoma cells. Depending on a greater or lesser expression of MITF, the clones of melanoma have recently been differentiated into: highly proliferative/minimally invasive and low proliferative/highly invasive [4][9]. In addition to MITF, other melanocytic genes (such as TYR, DCT, MART-1) are also upregulated in the proliferative phenotype of melanomas. Conversely, in the invasive genetic signature MITF and other genes (such as INHBA, COL5A1 and SDRPINE1) are involved in modifying the extracellular environment [4][10][11][12][13][14][15][16][17][18].

Furthermore, various authors have shown how the dedifferentiation mechanism is a predictor of poor response to target therapy: for example, although the discovery that the BRAF mutation may be the occasion for molecular targeted therapy, patients with DM appear to have little clinical benefit, both in terms of PFS and OS [19][20][21][22].

From a strictly dermatopathological point of view, a careful evaluation of cell morphology and an inconclusive immunohistochemistry for a specific entity are the starting points from which to start: various authors [23][24][25][26], for example, consider it important, when it is not possible to be sure only with routine diagnostic techniques, carry out molecular investigations for the BRAF mutation, so as to be sure of being faced with a case of DM [27][28][29]. Alkhasawneh et al., in 2019, reported the case of a 52-year-old woman, previously operated on for breast cancer and previous melanoma resulting in pT1b, who presented with a chest lesion that had entirely lost all melanocytic immunohistochemical markers, and expressed (aberrantly) only GATA-3. The diagnosis of DM was made only after careful analysis of the mutation for BRAFV600K [30].

Finally, it is important to underline how the discovery and deepening of the mechanisms of regulation of melanogenesis in mammals [31][32] have clarified that melanocytes are endowed with both a responsive function towards signal molecules (paracrine regulation) and autocrine regulation. This field is very interesting in trying to understand how in malignant melanomas these “physiological” pathways are dysregulated, and we receive lesions with large and abundant extruded melanin pigment.

### 3. Conclusions

Dedifferentiated melanoma is a unique clinical/biological entity, which continues to pose significant diagnostic challenges. It is quite understood that there are difficulties in differential diagnostics with other malignant neoplastic lesions such as undifferentiated carcinomas and sarcomas; dedifferentiation is only a single epiphenomenon of an underlying biological heterogeneity that governs the behavior and clinical aggression of the disease; however, dedifferentiation is a marker of cross-resistance to target therapy and immunotherapy.

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