Université BORDEAUX Metabolic and immune features as predictive biomarkers of risk of stratification of skin carcinoma



a recherche médical



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No conflics of interest

Introduction

Exposure to ultraviolet (UV) radiation from the sun is the most significant risk factor resulting in non-melanoma skin cancers (NMSCs), including cutaneous squamous cell carcinomas (cSCCs) which their incidence rates are still on the rise.

cSCCs typically manifests as a spectrum of progressively advanced malignancies, ranging from a precursor actinic keratosis (AK) to squamous cell carcinoma (SCC), in situ, invasive cSCC and finally metastatic¹.



Actinic Keratosis (AK)

In situ cSCCs

Invasive cSCCs

Metastasis

Skin carcinogenesis in multiple stages

A better understanding of molecular changes involved in the transformation of this UVB-induced precancerous lesions (actinic keratosis (AK)) to localized tumors and then metastasis will aid in early detection, development of biomarkers and future targeted strategies^{2,3}.

Time

Research Highlights

- Characterize the molecular and metabolic features of cSCCs at different stages of carcinogenesis
- Identify the immunologic landscape of cSCCs at different stages of carcinogenesis
- Uncover relevant biomarkers predicting the cSCCs evolution risk.

Results

1-Very similar metabolic pathways are involved in hyperplasic, AK and peritumoral tissue 2-On the contrary, metabolic profile of tumors are differing from precancerous lesions





Downregulation in lipid biosynthesis and upregulation in glycolysis

- Persistent Purine and Pyrimidine metabolic pathway
- Specific metabolic modifications, of which some persists throughout tumor development, occur at a very early stage of skin carcinogenesis

3- Several immune cell subset are depleted in cSCC compared to AK or enriched



IGF-1 Signaling Glycogen Degradation III **Calcium Signaling** Superpathway of Geranylgeranyldiphosphate Biosynthesis I (v... Tryptophan Degradation III (Eukaryotic) Glutaryl-CoA Degradation Mevalonate Pathway I Glycogen Degradation II Melanocyte Development and Pigmentation Signaling Valine Degradation I Tryptophan Degradation X (Mammalian, via Tryptamine) Tumor Microenvironment Pathway Oxidative Ethanol Degradation III Ethanol Degradation II Autophagy Xenobiotic Metabolism AHR Signaling Pathway Serotonin Degradation



Metabolic profiles differ in subgroups of patients including OXPHOS and glycolysis

