

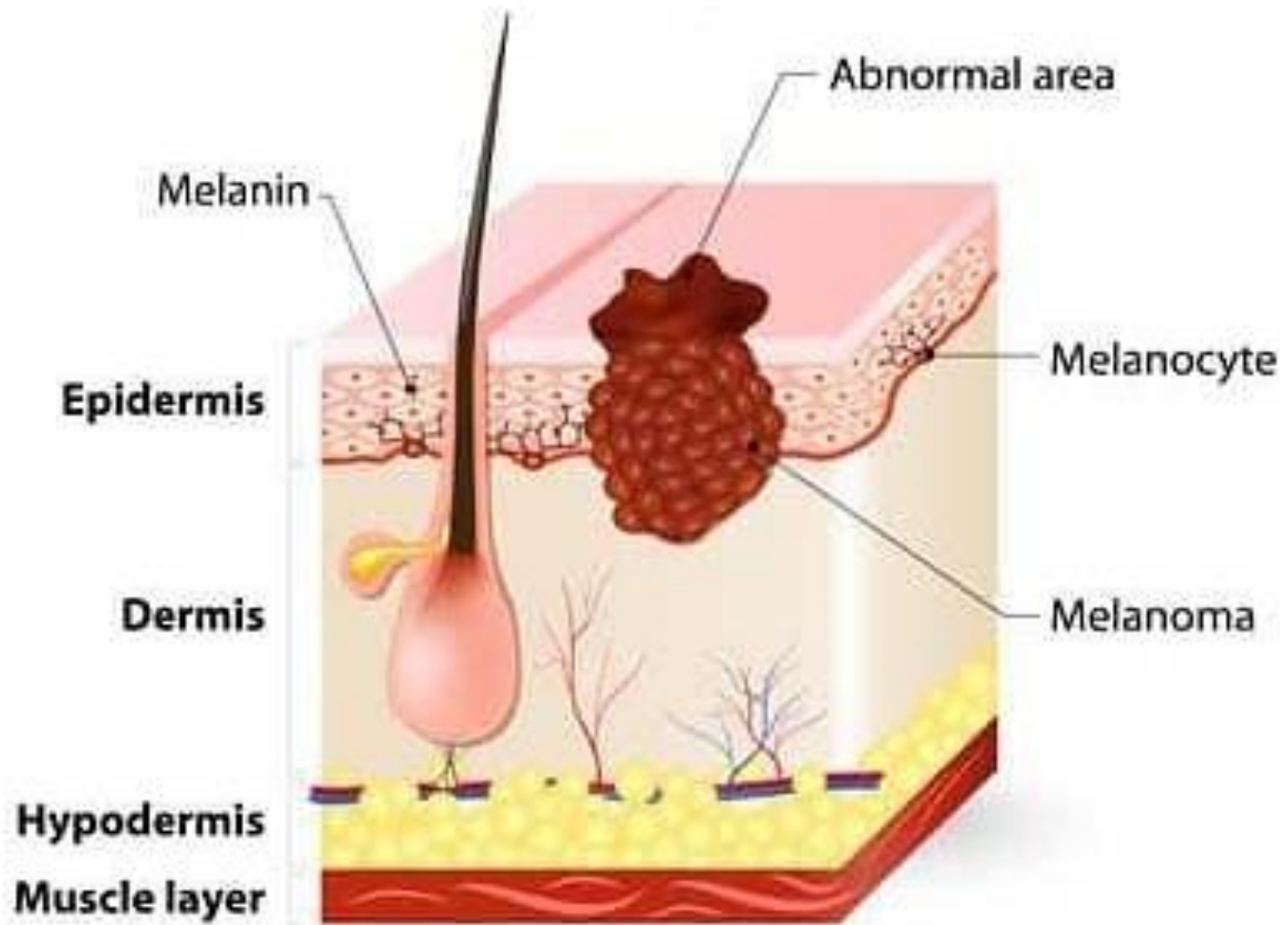


A stochastic clonal selection for modeling intra tumor heterogeneity in human melanoma and clinical implications

Larissa A. C. Carvalho

lari.anastacio@gmail.com

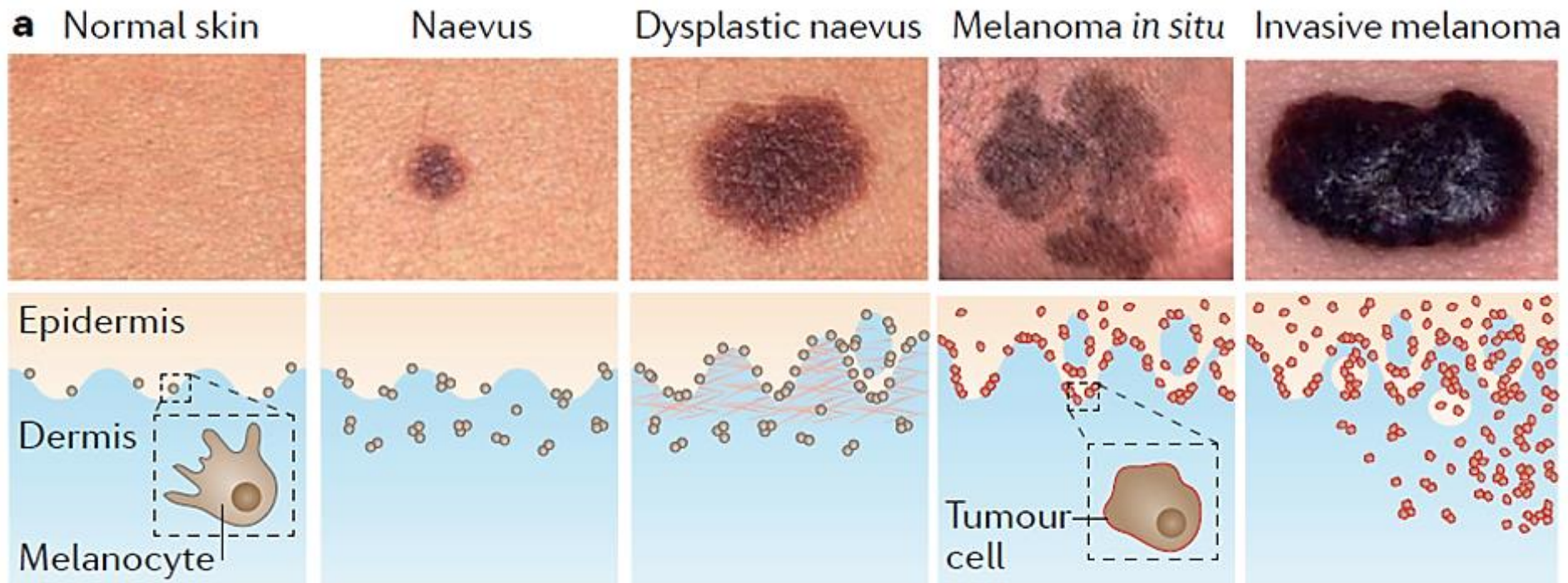
Melanoma of the skin



Melanoma progression: mutational status

Mutations:
BRAF, MYC, NRAS

Senescence suppression
Apoptosis suppression
Telomeres alterations





Melanoma skin cancer rates are on the rise

- Skin cancer: **30% of all tumors**
- Melanoma: **4-6% of all cases** but the most aggressive (**80% of skin cancer deaths**)

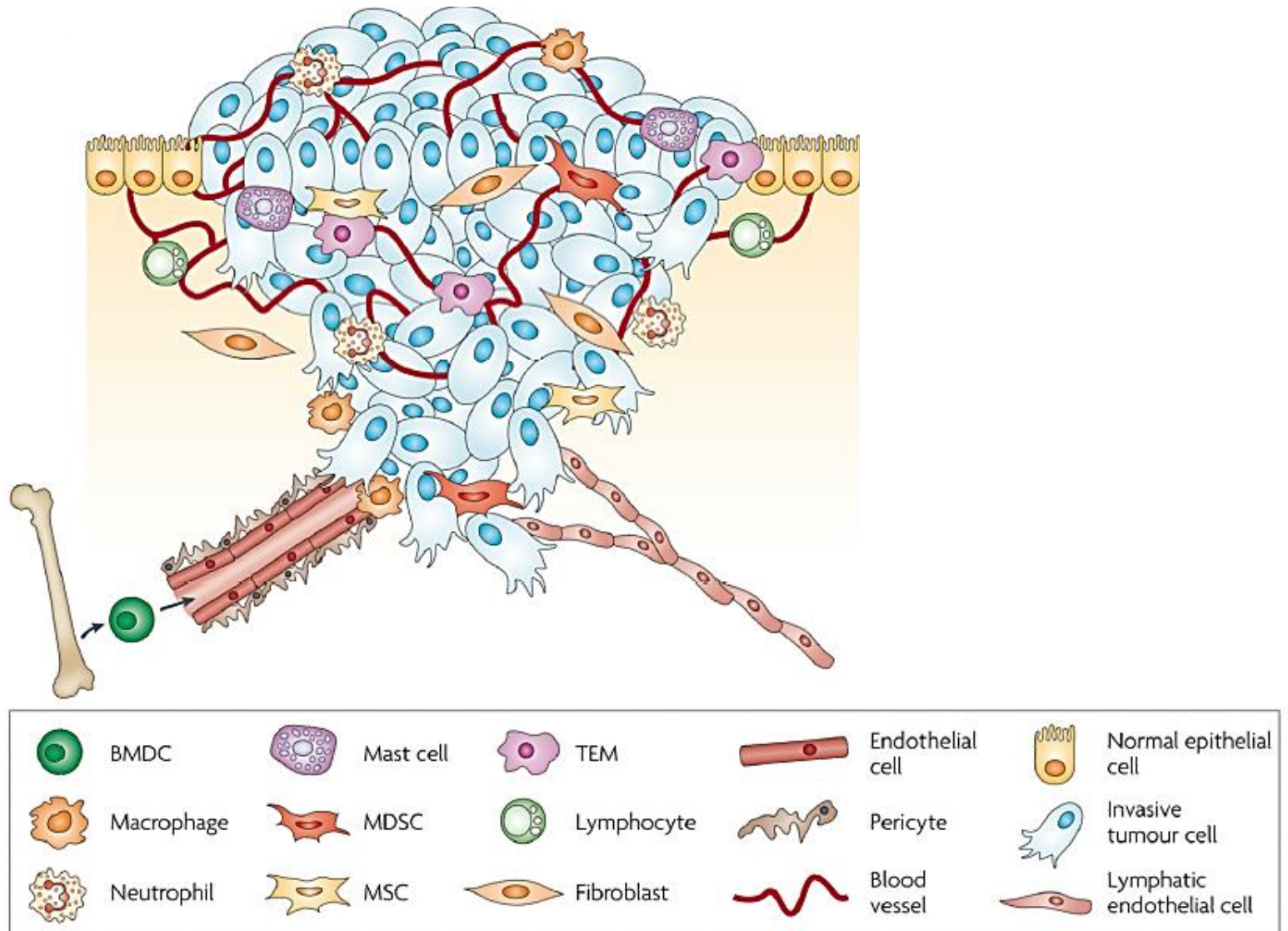
Change in melanoma incidence since 90's

+128%



			Males	Females			
Prostate	164,690	19%			Breast	266,120	30%
Lung & bronchus	121,680	14%			Lung & bronchus	112,350	13%
Colon & rectum	75,610	9%			Colon & rectum	64,640	7%
Urinary bladder	62,380	7%			Uterine corpus	63,230	7%
Melanoma of the skin	55,150	6%			Thyroid	40,900	5%
Kidney & renal pelvis	42,680	5%			Melanoma of the skin	36,120	4%
Non-Hodgkin lymphoma	41,730	5%			Non-Hodgkin lymphoma	32,950	4%
Oral cavity & pharynx	37,160	4%			Pancreas	26,240	3%
Leukemia	35,030	4%			Leukemia	25,270	3%
Liver & intrahepatic bile duct	30,610	4%			Kidney & renal pelvis	22,660	3%
All Sites	856,370	100%	All Sites	878,980	100%		

Tumor microenvironment



Joyce e Pollard, *Nature Reviews Cancer*, 2008.

Romano et al., *International Journal of Molecular Science*, 2017

Modeling melanoma *in vivo*



Genetically engineered mouse

- Fully functional immune system
- Complete microenvironment
- Genetic manipulation

- Mice rarely develop melanoma spontaneously
- Differences in histology
- Mouse cells



Xenograft

- Human cells
- Complete microenvironment
- Study of spontaneous metastasis

- Lacks the immune system
- Differences in physiology
- Mouse microenvironment

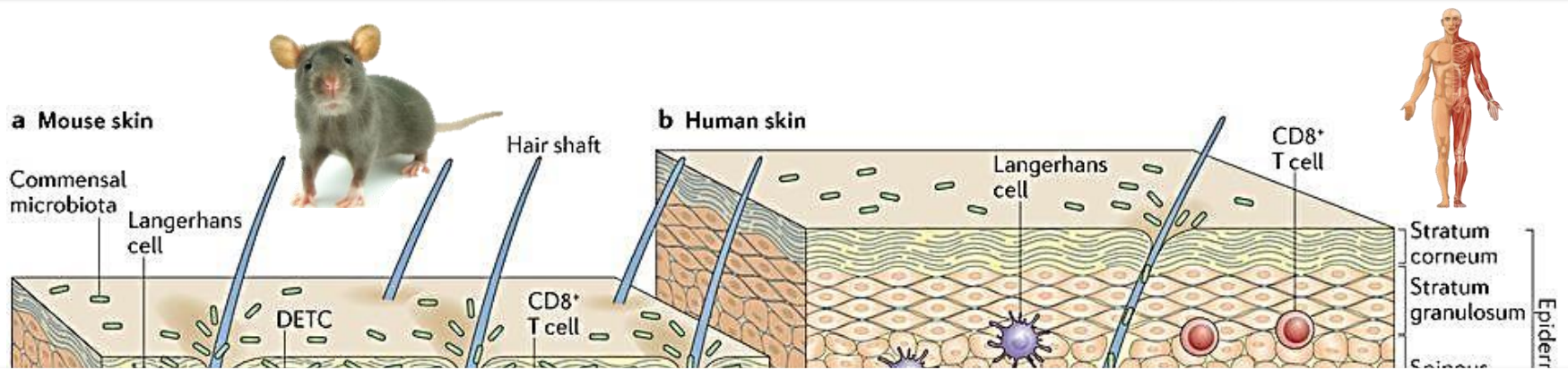


Zebrafish

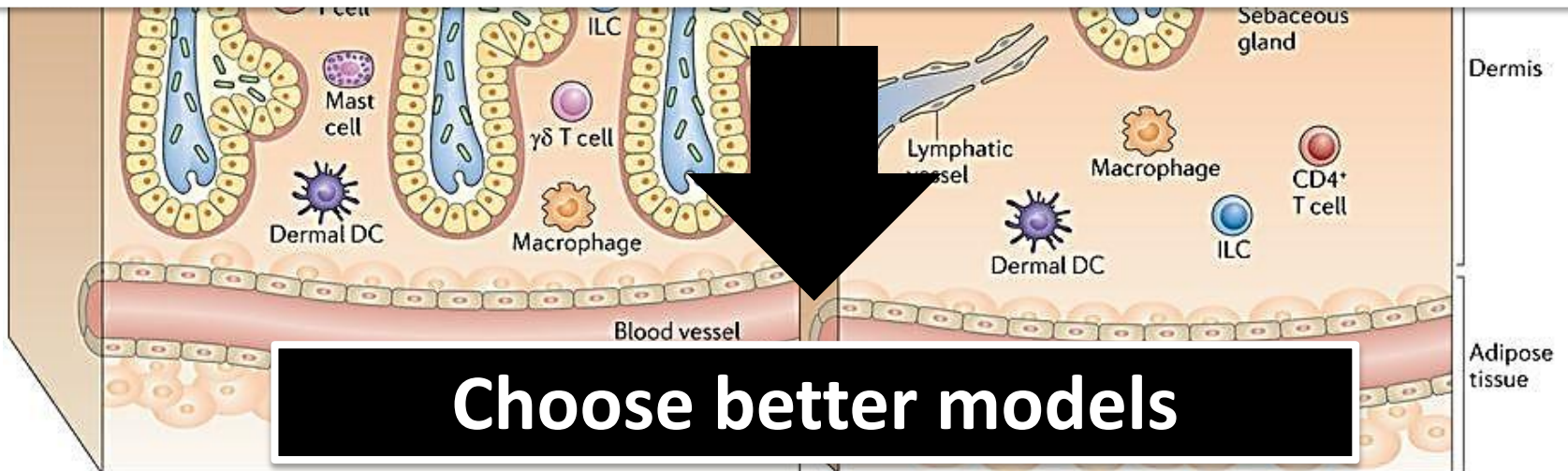
- Shorter generation time, large number of progeny
- Develop spontaneous melanoma
- Simple genetic manipulation

- Differences in physiology

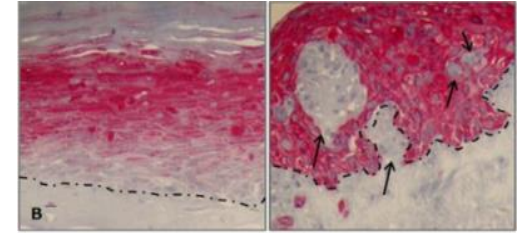
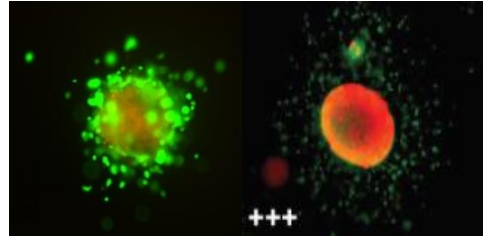
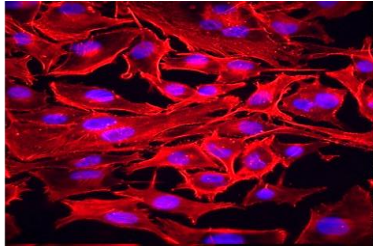
Human and mouse skin are different



The average rate of successful translation from animal models to clinical cancer trials is less than 8%



Modeling melanoma *in vitro*



2D cell culture

- Patient cells or cell line
- Simplicity, convenience
- Preliminary drug screening: drugs that do not work in 2D cultures have no effect in more realistic models

- Lacks the microenvironment

3D spheroid

- Oxygen/nutrient gradient with a hypoxic zone and a central necrosis
- Interaction between melanoma cells and their stroma

- Lacks the whole system

Organotypic skin reconstructs

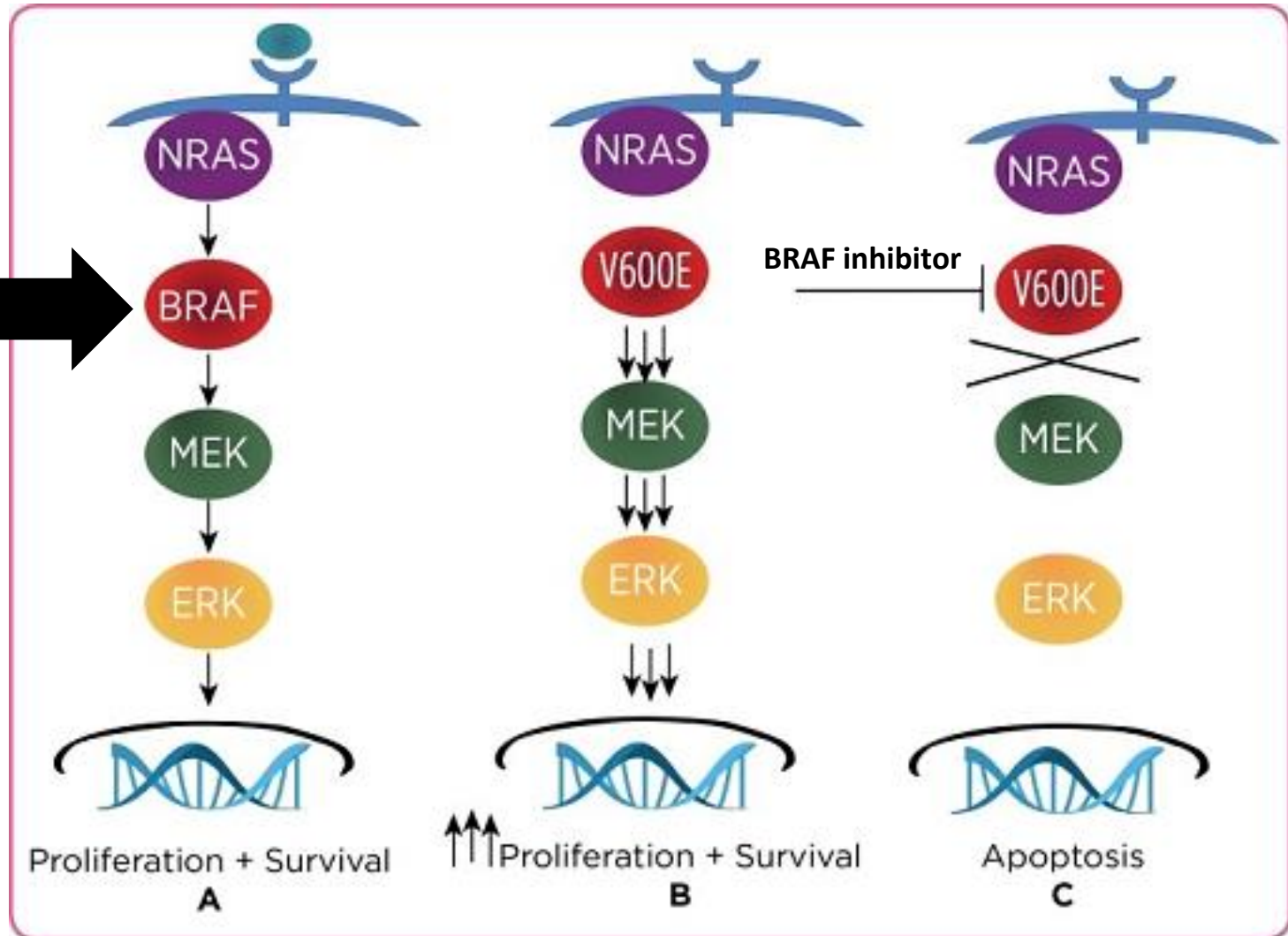
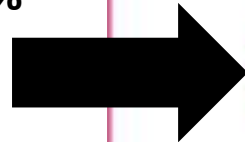
- Closely resembles histologically human skin
- Cells from patients keep their properties
- Drug screening

- Long-term development

Oncogene targeted therapy: BRAF inhibitor

MAPKinase pathway

Mutated in 60%
of melanomas



Melanoma treatment – Resistance

No treatment



15 weeks - Remission

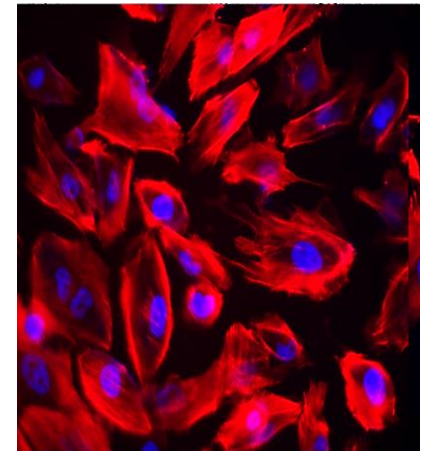
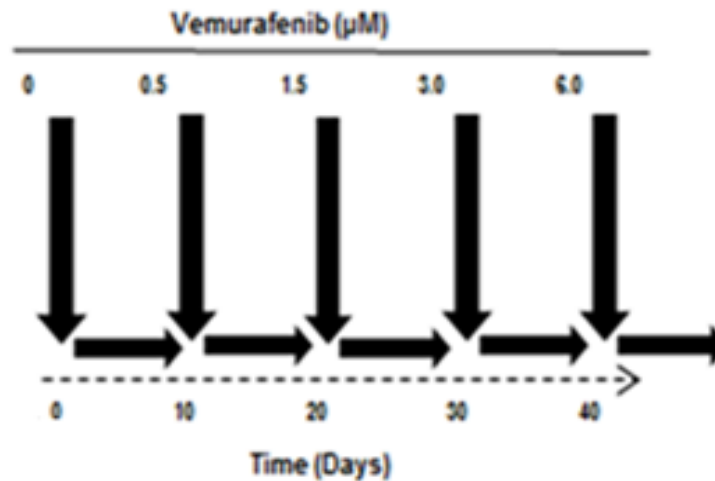
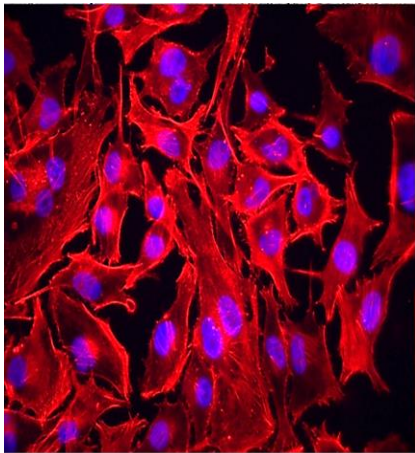


23 weeks - Relapse



How to model resistance *in vitro*?

Generation of BRAFi resistant cells



Why does melanoma resistance occur?

Acquired resistance

- Activation of different pathways to overcome the drug

Intrinsic resistance

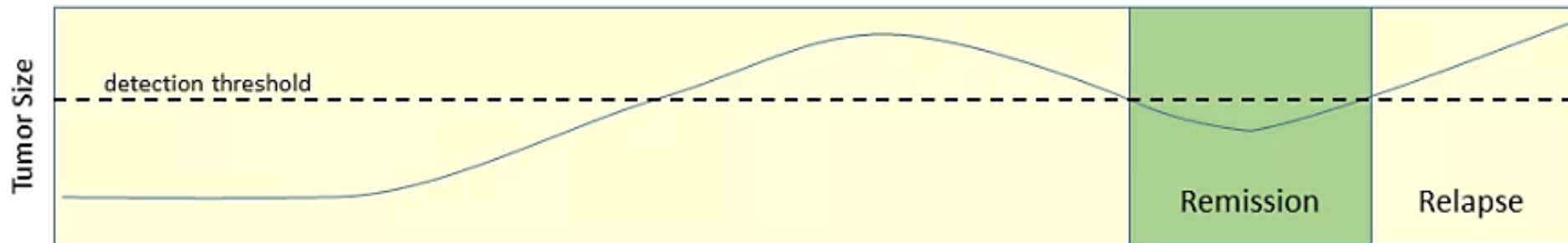
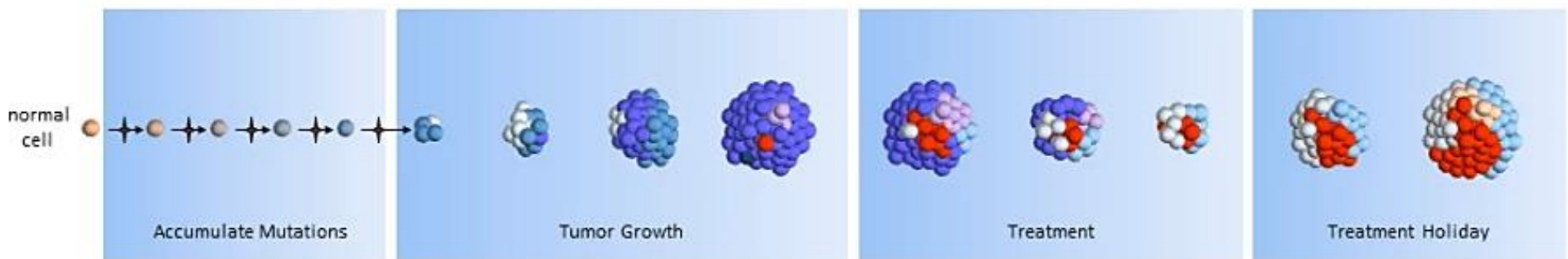
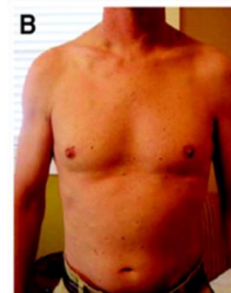
- Cells are resistant to the drug before the use of it



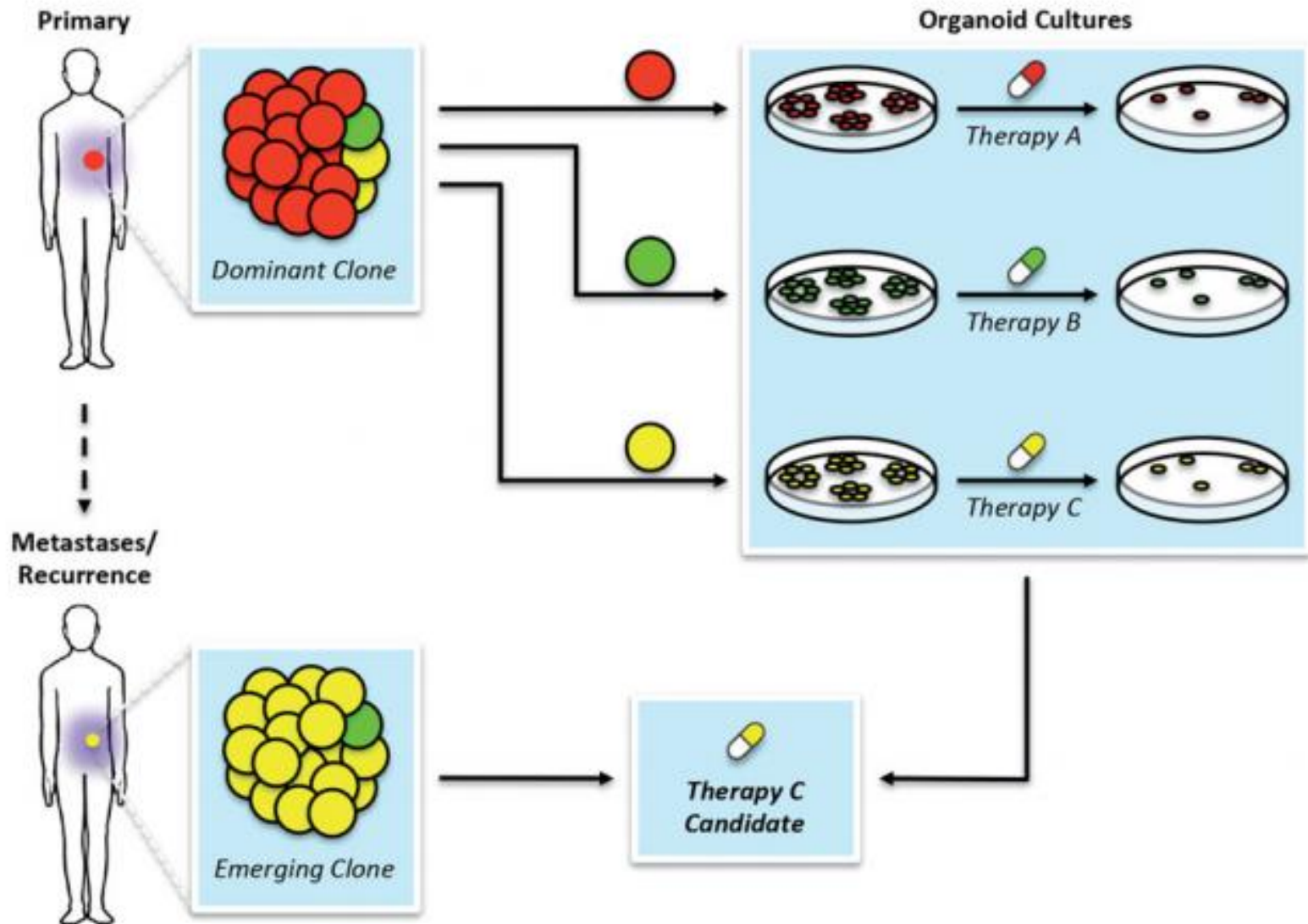
Source: clipground

Intratumor heterogeneity

Tumors comprises heterogeneous subpopulations

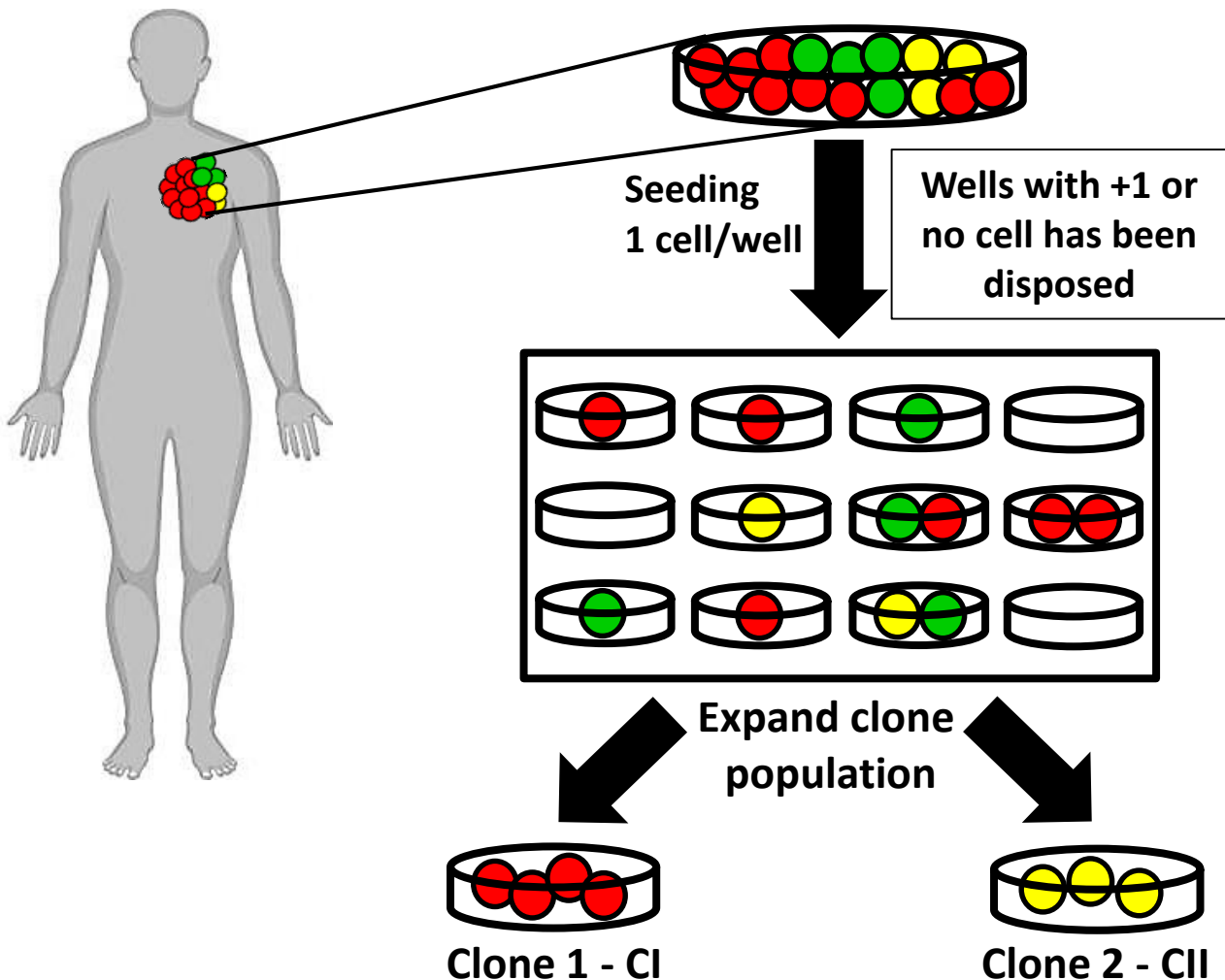


Clinical impact: from *in vitro* to patient



How to model heterogeneity *in vitro*?

Parental heterogeneous population with
no drug treatment –
WM164 cell line

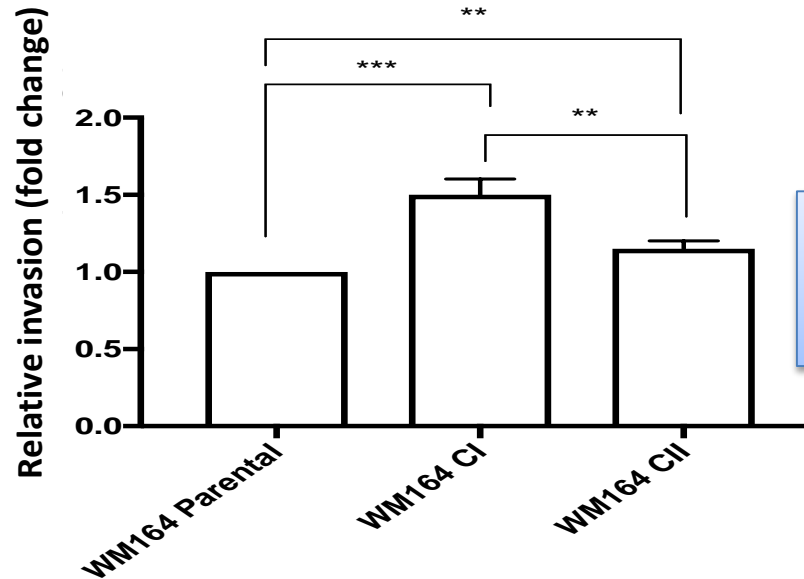
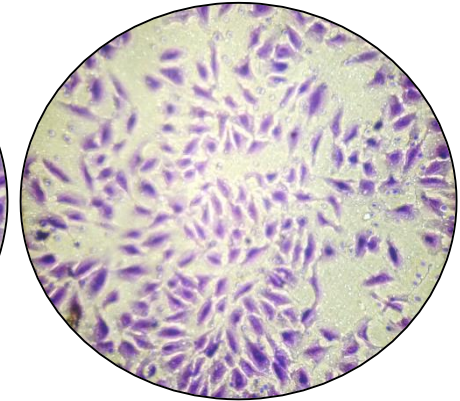
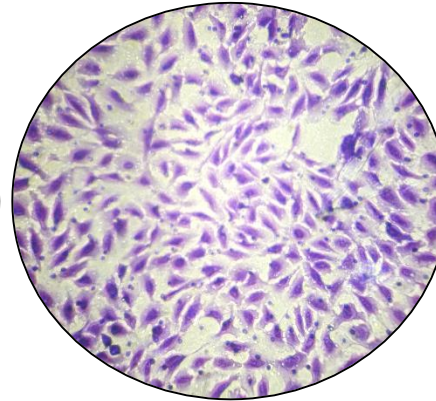
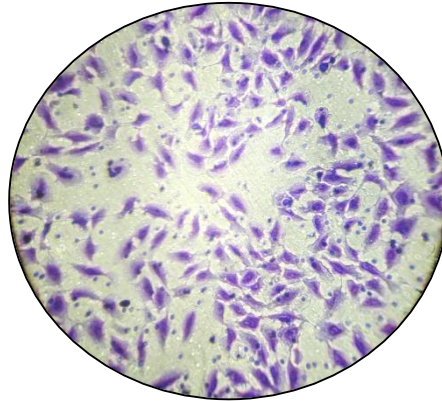
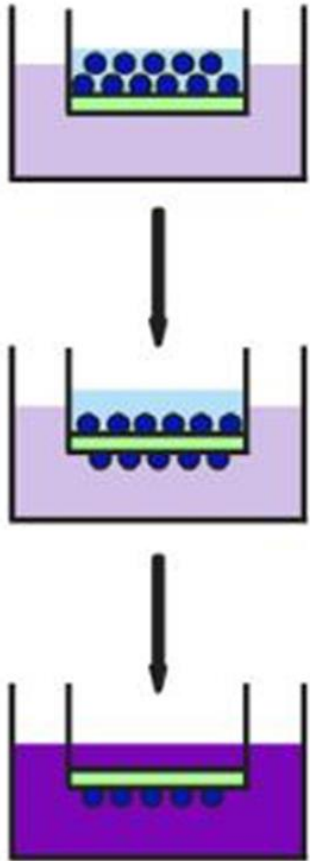


Characterization of the clones: Invasion

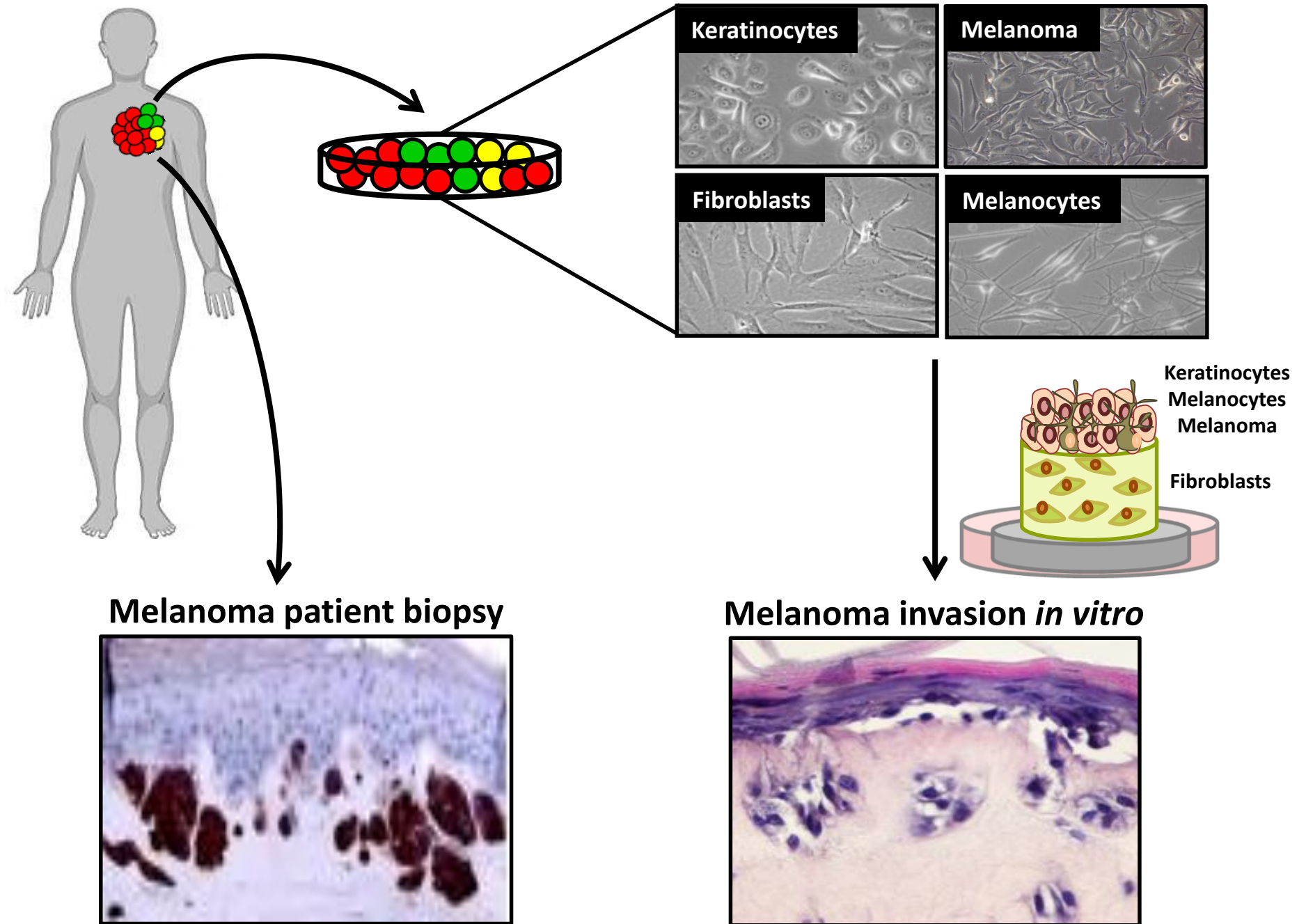
WM164
parental

WM164 CI

WM164 CII

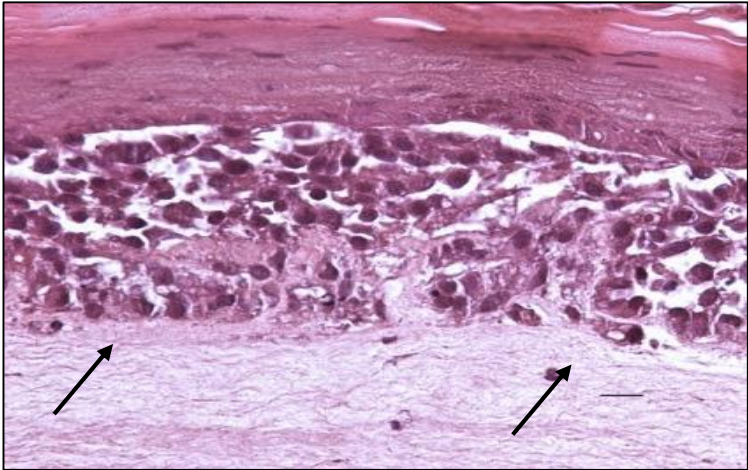


CI is more invasive than CII

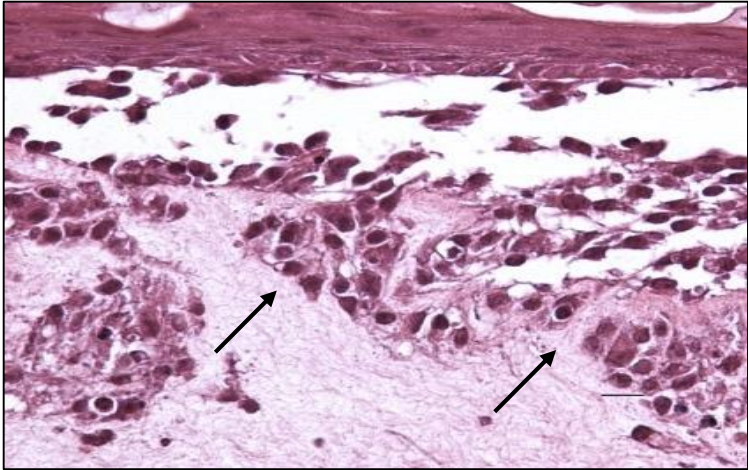


Reconstructed human skin: Invasion

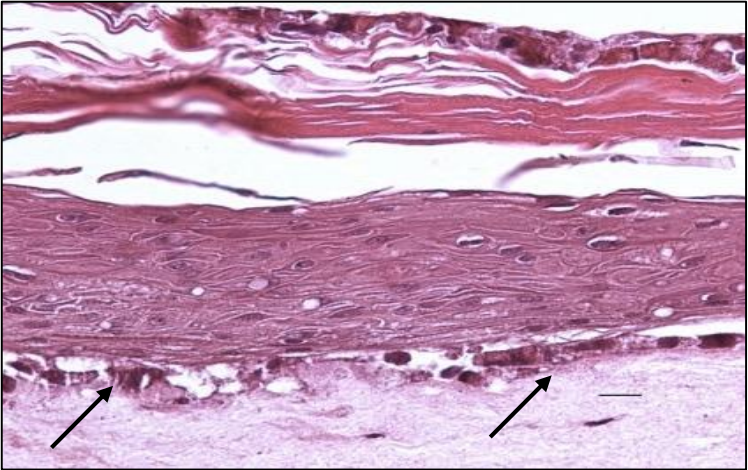
WM164 Parental



WM164 CI

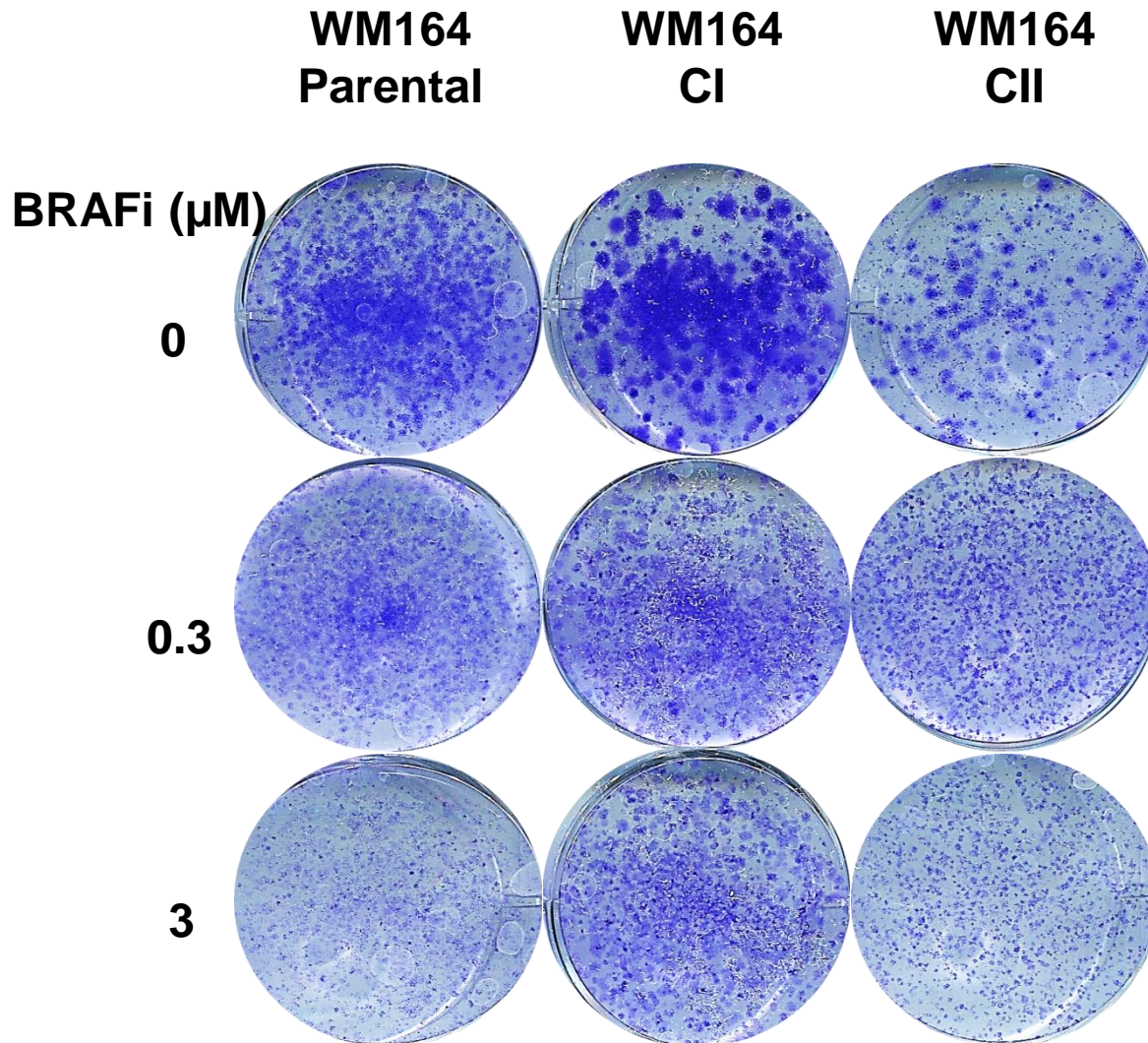


WM164 CII



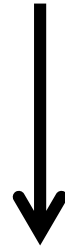
CI is more invasive than CII

Characterization of the clones: intrinsic resistance to BRAFi



**CI is more resistant
than CII**

BRAF mutation

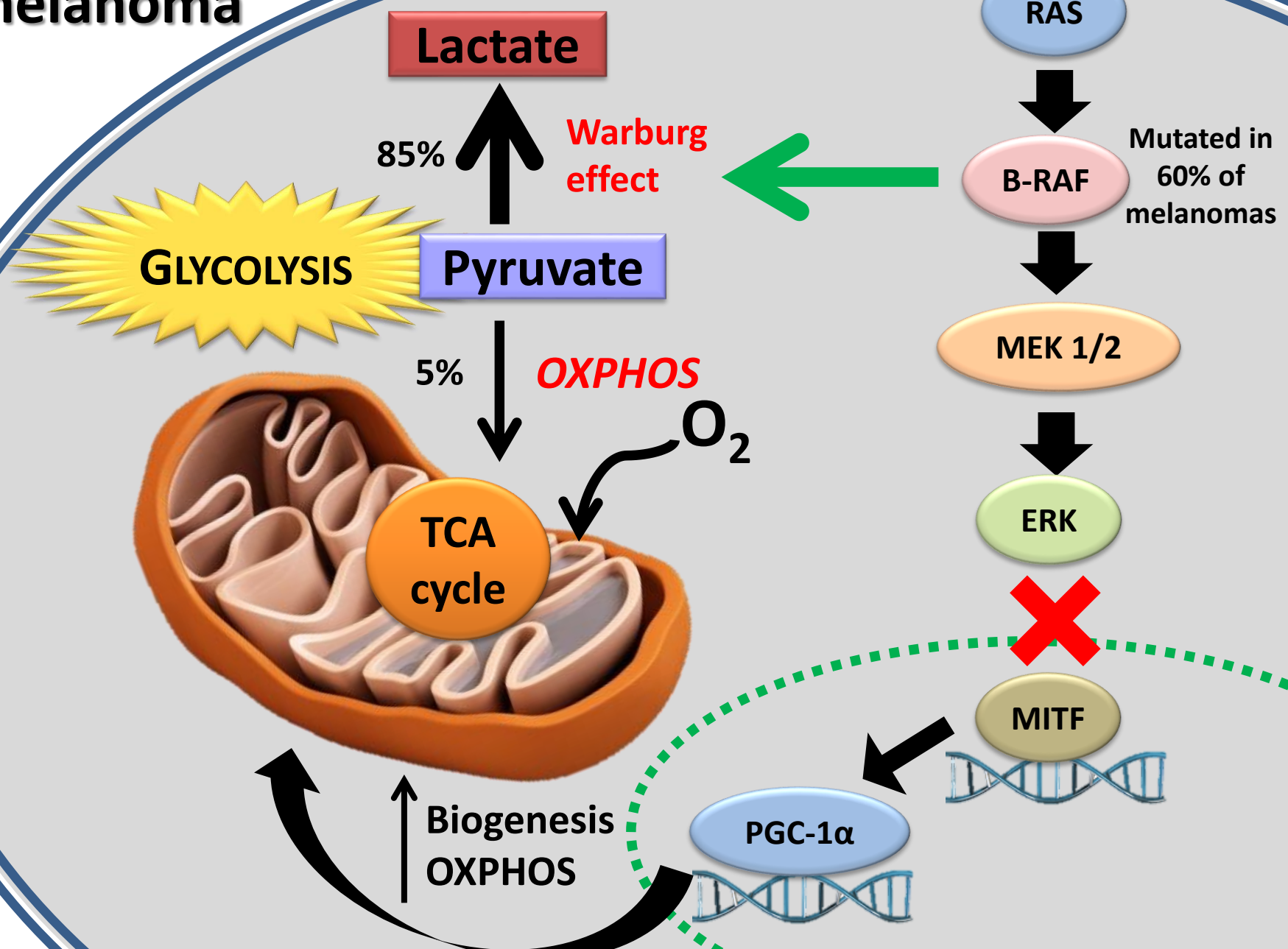


Metabolic reprogramming

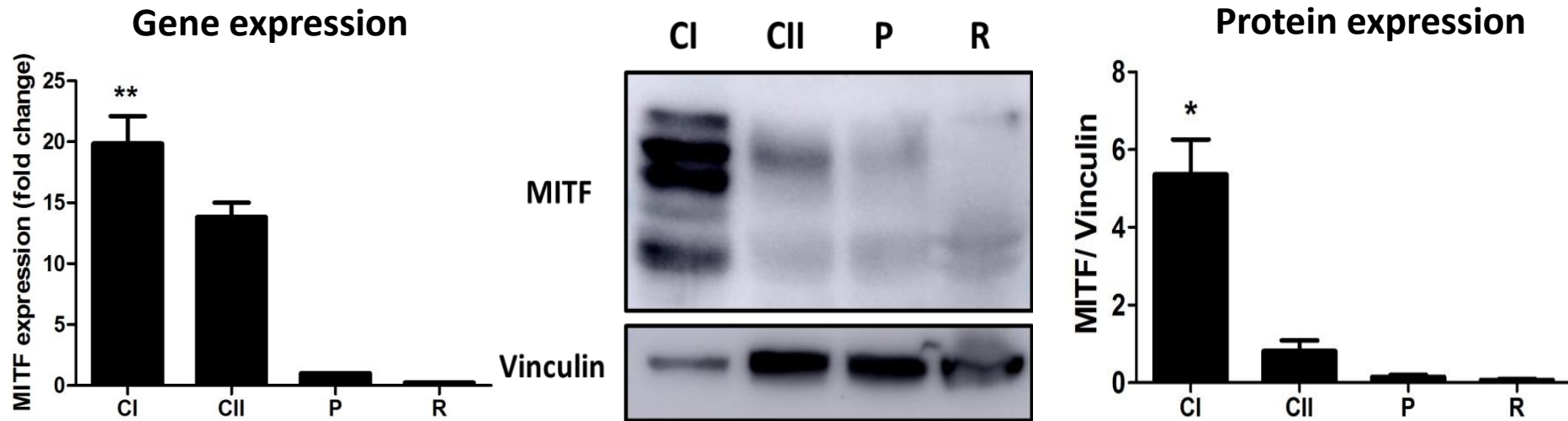
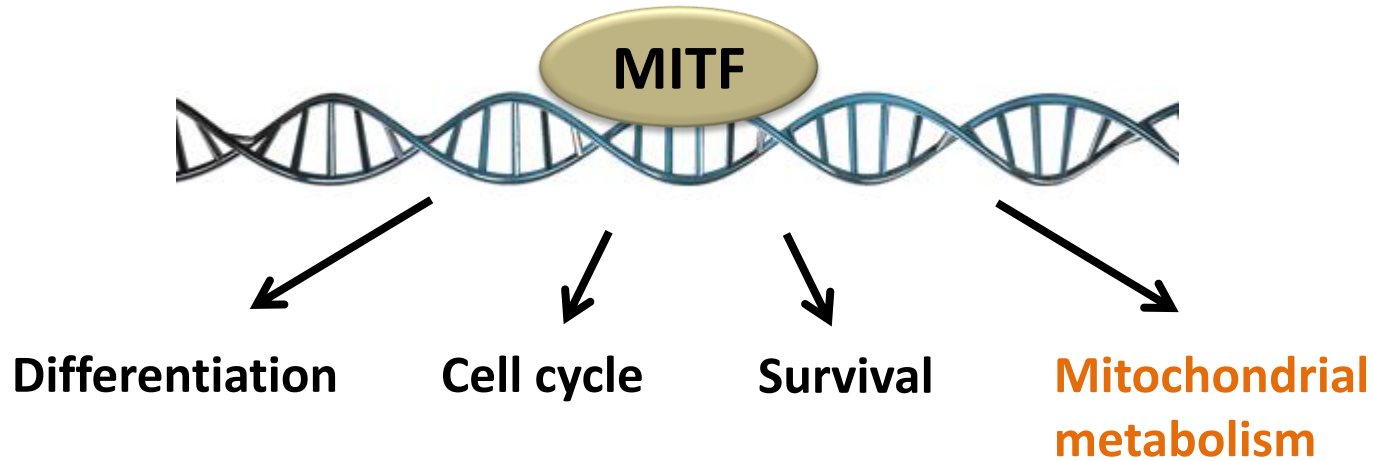
Can mitochondria be the main responsible for the tumor heterogeneity and resistance?

Oxidative metabolism in melanoma

MAPKinase pathway



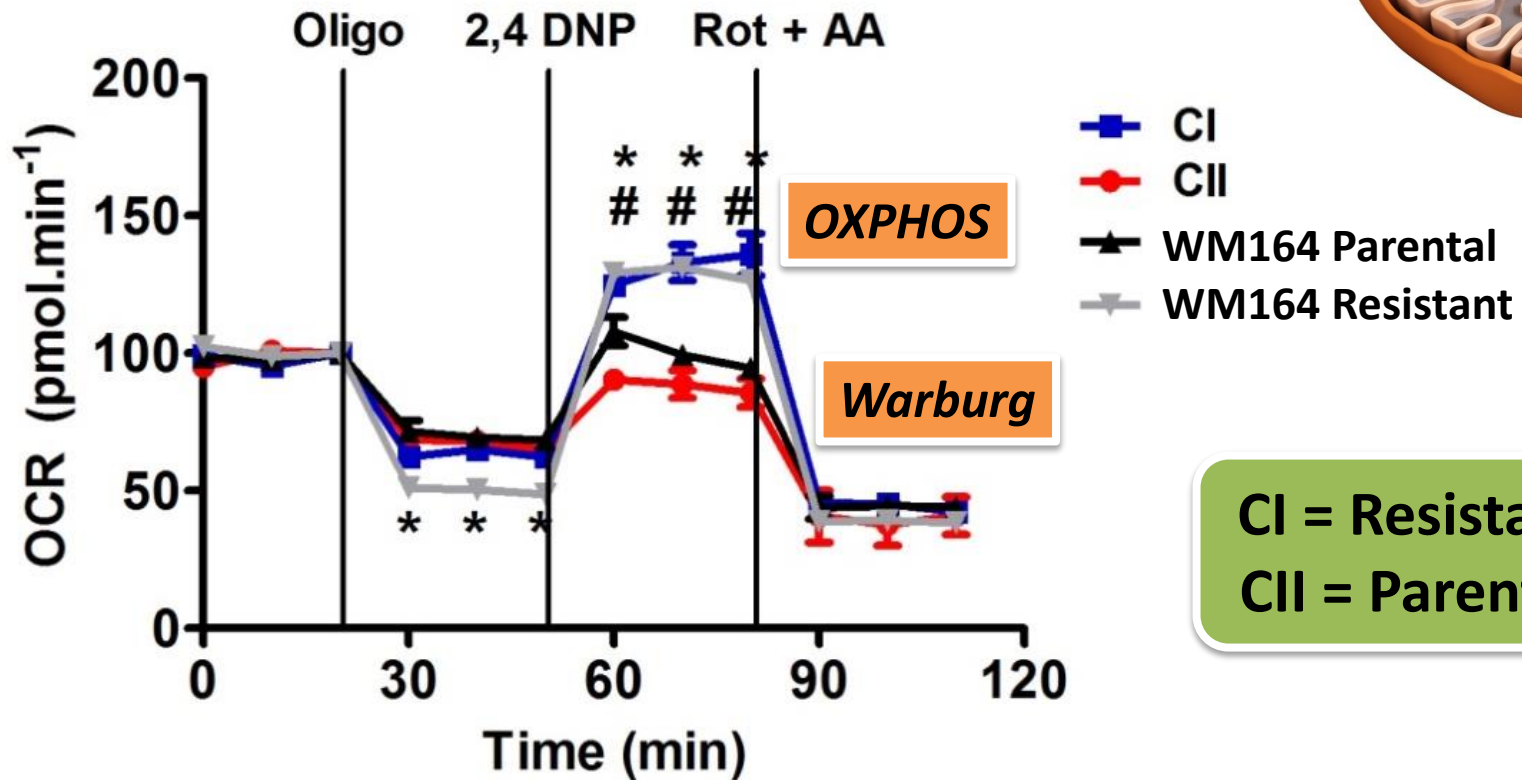
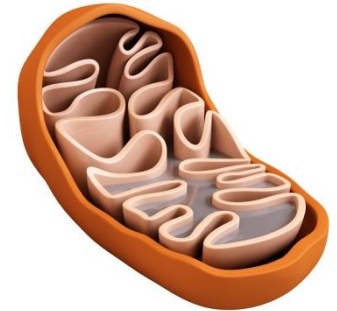
The major transcriptional melanocyte regulator



CI showed higher expression of MITF

Mitochondrial metabolism

Oxygen consumption rate



CI = Resistant
CII = Parental

CI and Resistant showed identical profiles

Can we develop better tools to detect melanoma earlier?

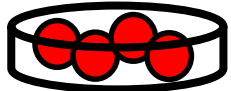
Will identification of melanoma clones dictate therapy?

In vitro ----- non-animal model -----> *Patient*

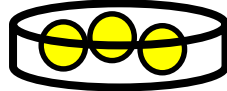
Predict metastasis and resistance

Choose the best therapy

CI



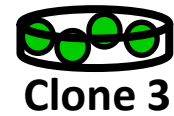
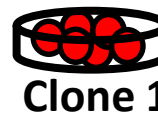
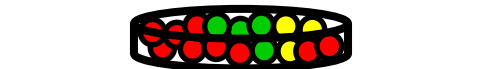
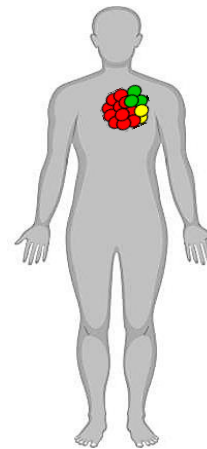
CII



Invasion
Resistance
MITF
OXPHOS

Invasion
Resistance
MITF
OXPHOS

Isolation of clones



Clone 1

Clone 3

Clone 2

Best therapy



Acknowledgments

Skin Biology Group – School of Pharmaceutical Sciences



Dr. Manoela Tiago



Prof. Dr. Keiran Smalley
Dr. Inna Smalley



Prof. Dr. Nadja C. Souza-Pinto
Dr. Mateus P. Mori

University of São Paulo



Financial support:

