

Classification of melanoma for pathologists

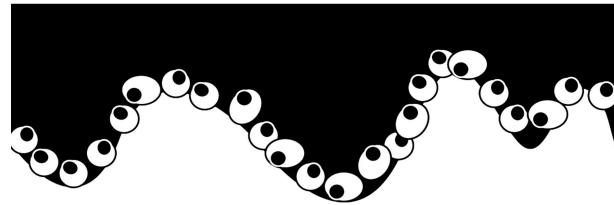
Boris C. Bastian

*Departments of Dermatology and Pathology,
Helen Diller Family Comprehensive Cancer Center
University of California, San Francisco*

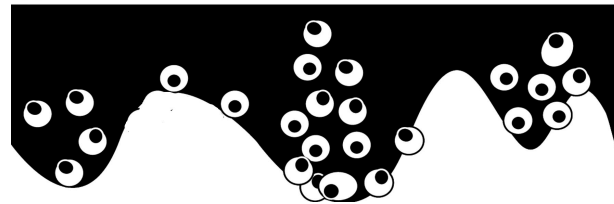
[illegible]

Histopathological phenotypes

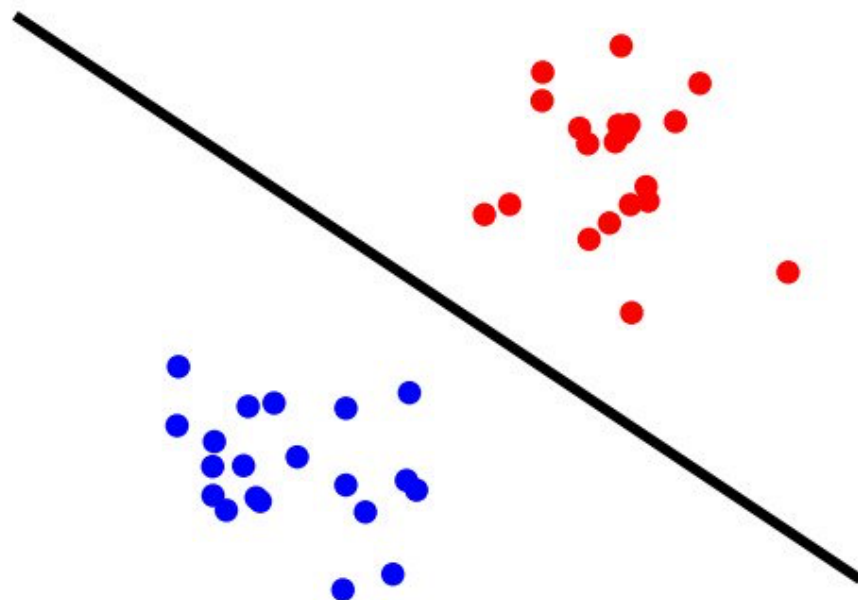
Lentiginous



Pagetoid



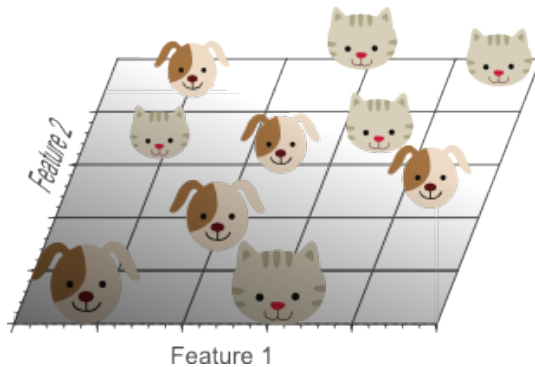
Classification



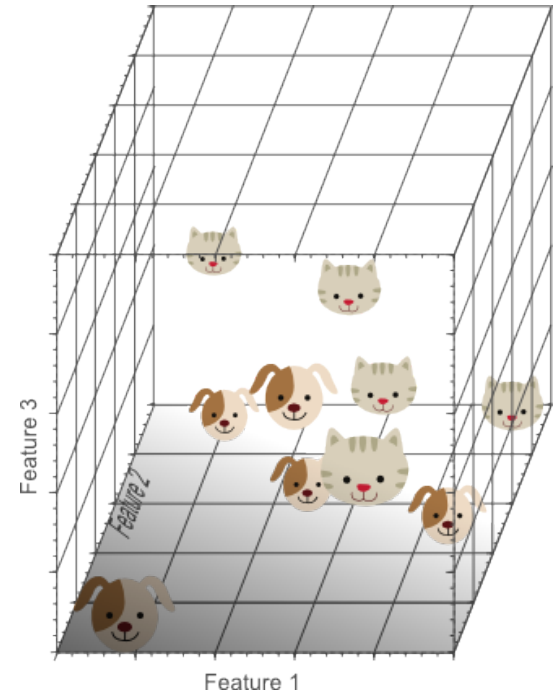
Dimensionality and classification



1 Dimension

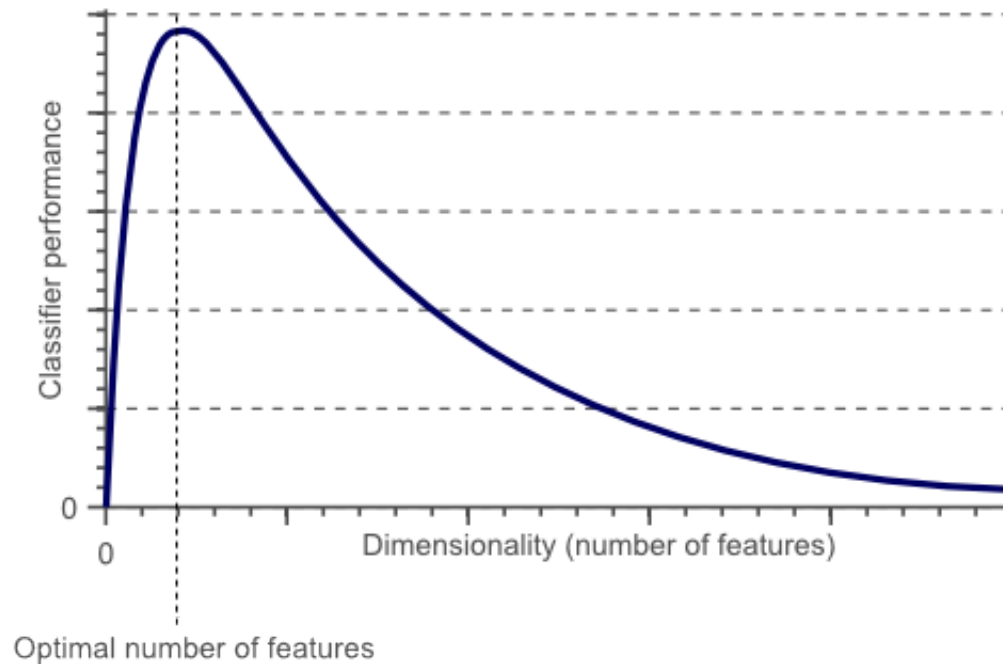


2 Dimensions



3 Dimensions

Dimensionality and classification

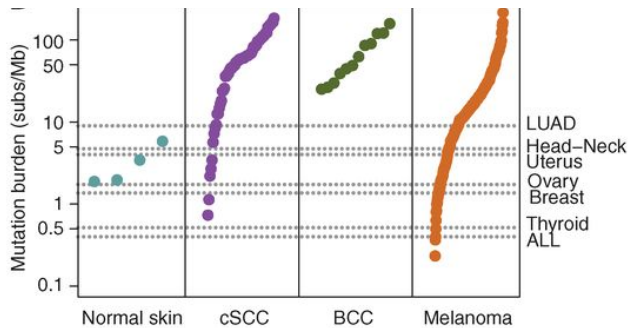


Vincent Spruyt

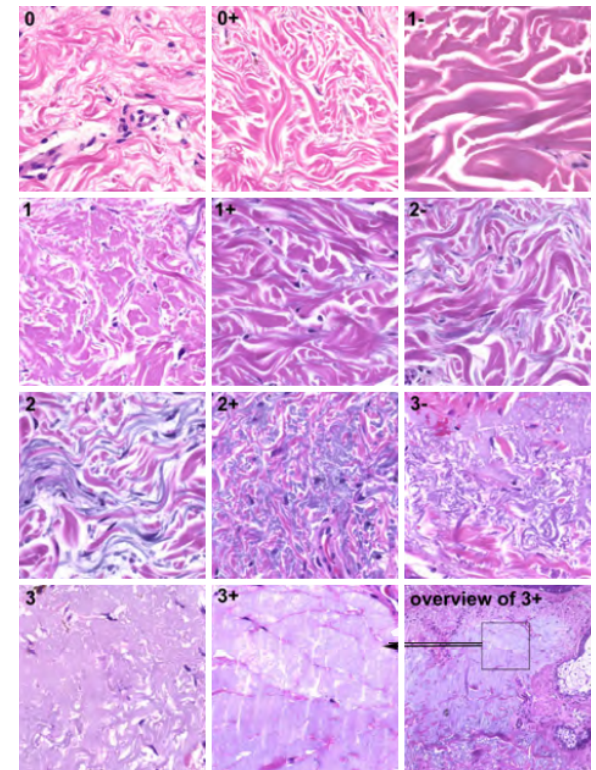
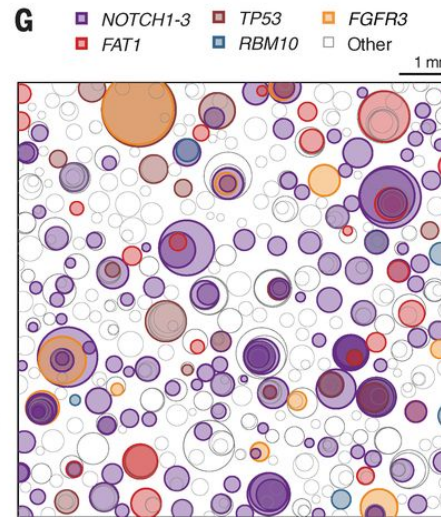
Age of onset



Degree of sun exposure



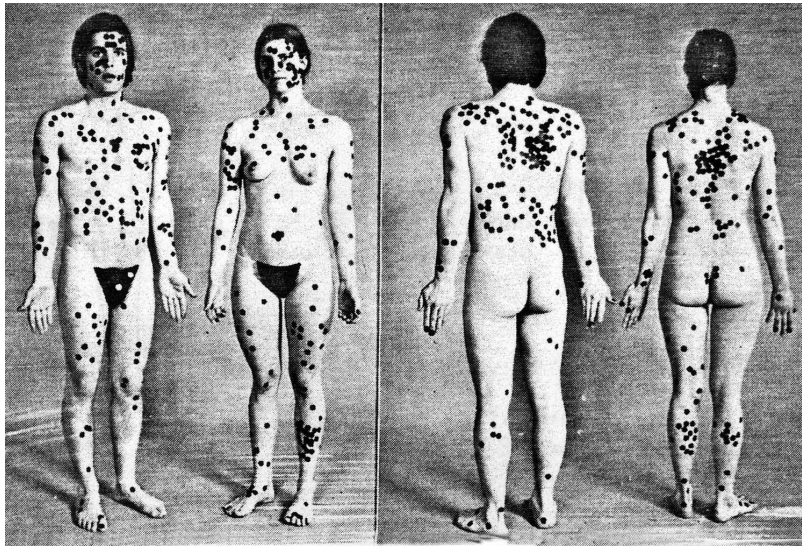
Martincorena et al. Science 2015



Landi, Bauer et al. Science 2015

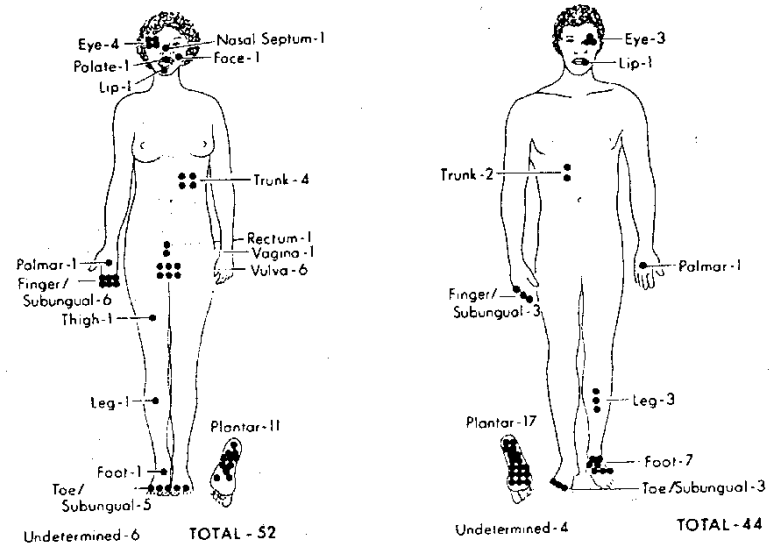
Host characteristics influence melanoma phenotypes

Caucasian



Melanoma Clinical Cooperative Group data

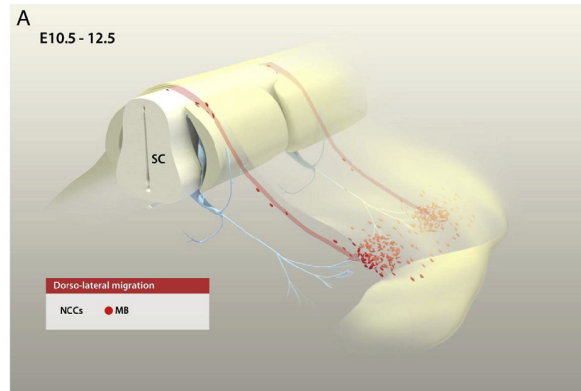
African



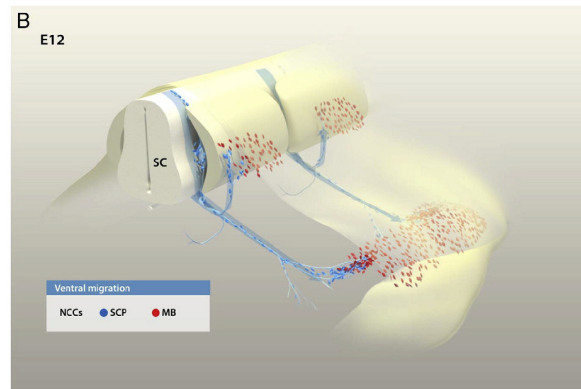
Krementz ET et al. Ann Surg 1976

Cells of origin and their differentiation states

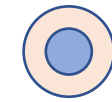
Dorsolateral
developmental
pathway



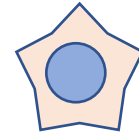
Ventral
developmental
pathway



Stem cell



Progenitor I



Progenitor II



Differentiated
Cell



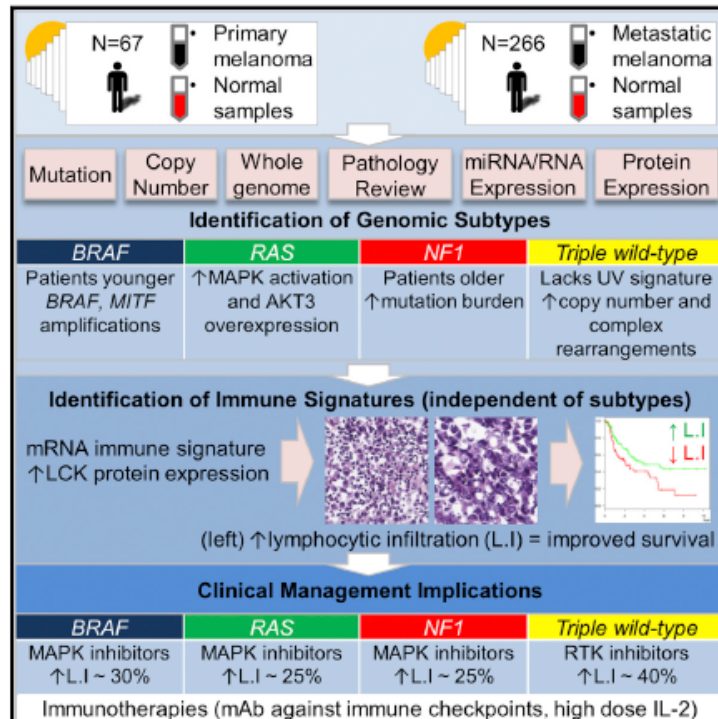
Somatic mutations

Cell

resource

Genomic Classification of Cutaneous Melanoma

Graphical Abstract



Authors

The Cancer Genome Atlas Network

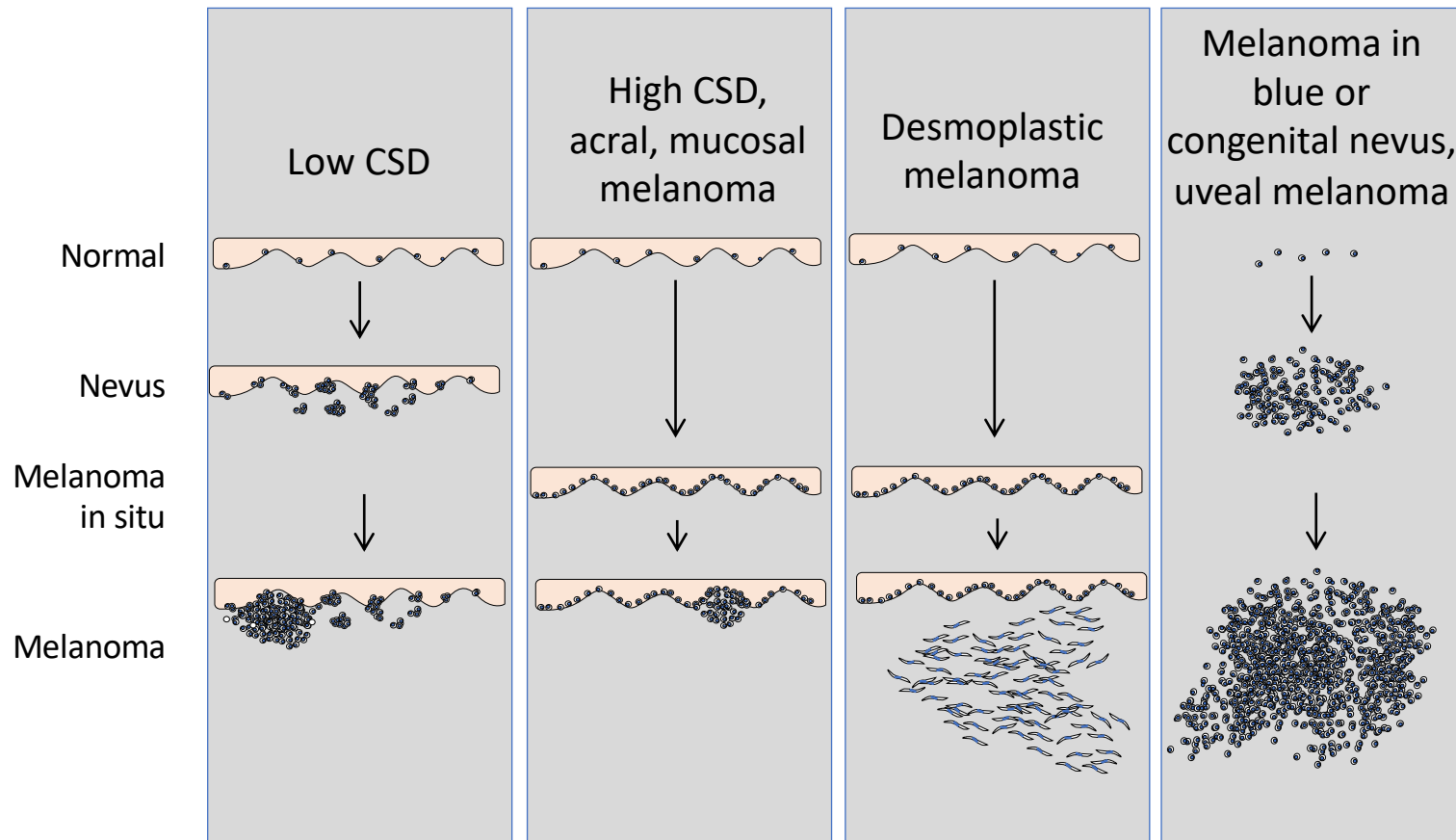
Correspondence

irwatson@mdanderson.org (I.R.W.),
jgershen@mdanderson.org (J.E.G.),
lchin@mdanderson.org (L.C.)

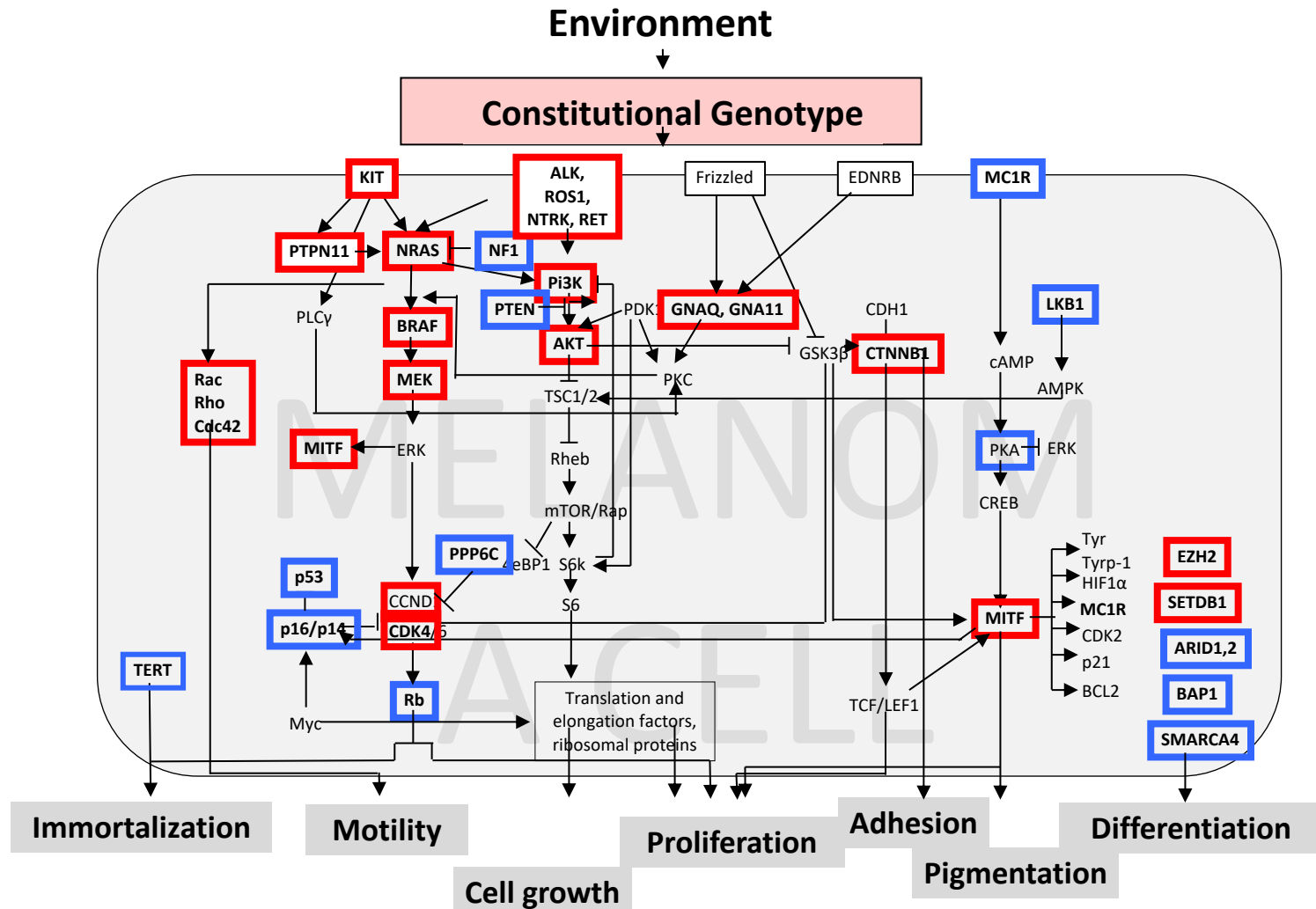
In Brief

An integrative analysis of cutaneous melanomas establishes a framework for genomic classification into four subtypes that can guide clinical decision-making for targeted therapies. A subset of each of the genomic classes expresses considerable immune infiltration markers that are associated with improved survival, with potential implications for immunotherapy.

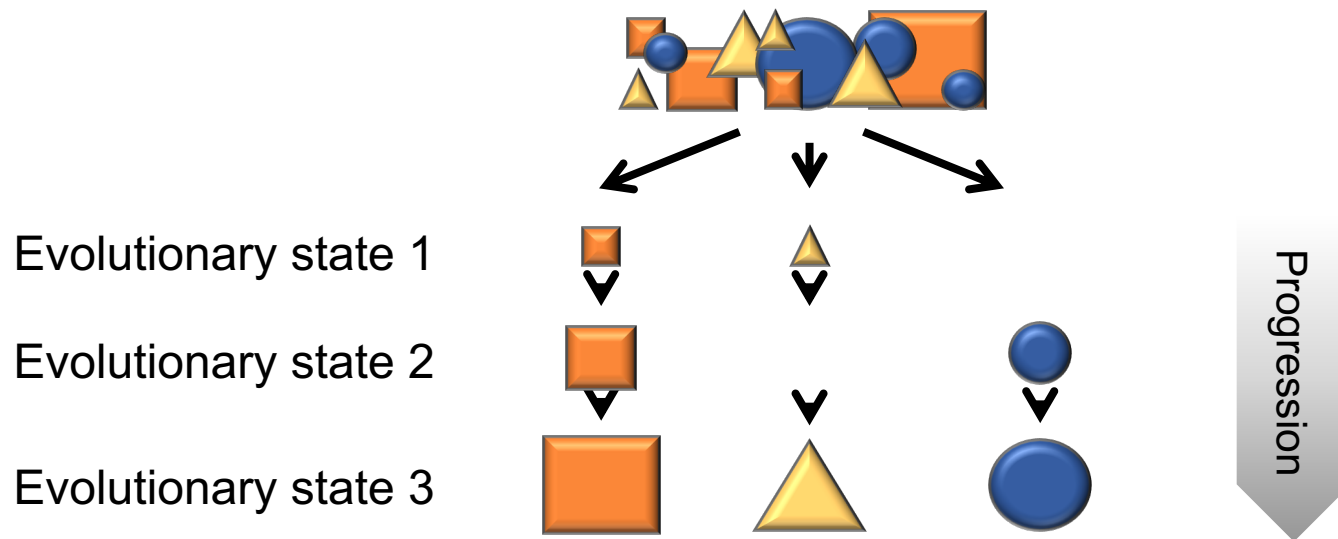
Evolution of primary melanomas



Genotype → phenotype



Integrated taxonomy of melanocytic neoplasms



Evolution of cutaneous melanoma

Melanocyte	Nevus	Intermediate Neoplasm	Melanoma in situ	Invasive Melanoma	Metastasis
1. MAP-kinase pathway mutation					
TERT promoter mutation					
1. G1/S checkpoint mutation					
SWI/SNF mutation					
2. G1/S checkpoint mutation					
2. MAP-kinase pathway mutation					
3. MAP-kinase pathway mutation					
TP53 mutation					
PTEN mutation					
Diagnostic Accuracy					

Shain et al. NEJM 2015; Shain & Bastian Nature Rev Cancer 2016

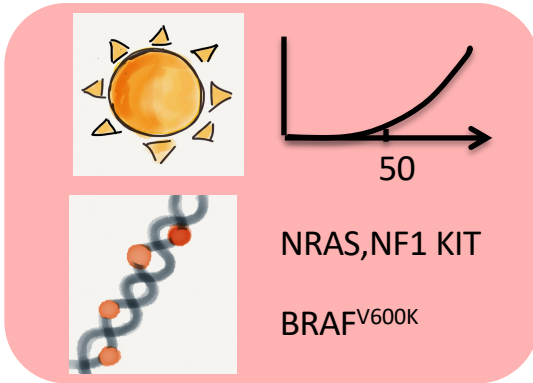
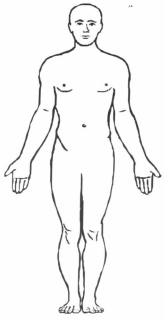
WHO Classification of Melanoma

3rd Edition

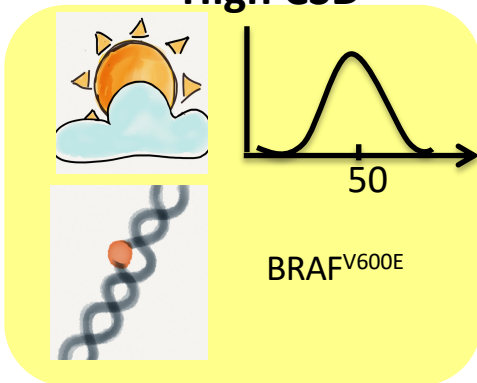
- Superficial spreading melanoma
- Nodular melanoma
- Lentigo maligna melanoma
- Desmoplastic melanoma
- Nevoid melanoma
- Acral-lentiginous melanoma
- Mucosal melanoma
- Uveal melanoma
- Melanoma of childhood
- Melanoma arising from giant congenital nevus
- Melanoma arising from a blue nevus
- Persistent melanoma

4th Edition

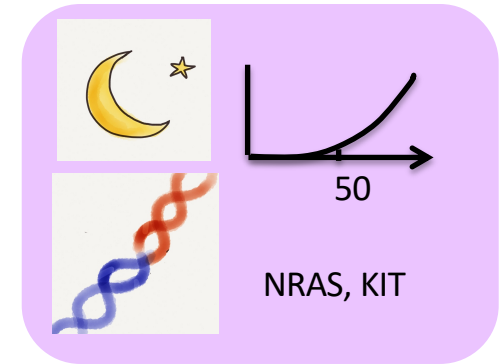
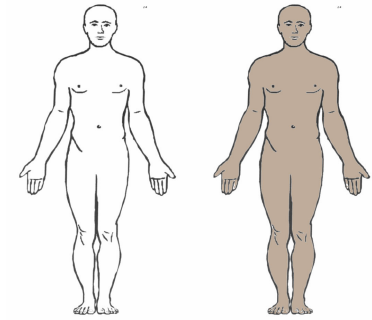
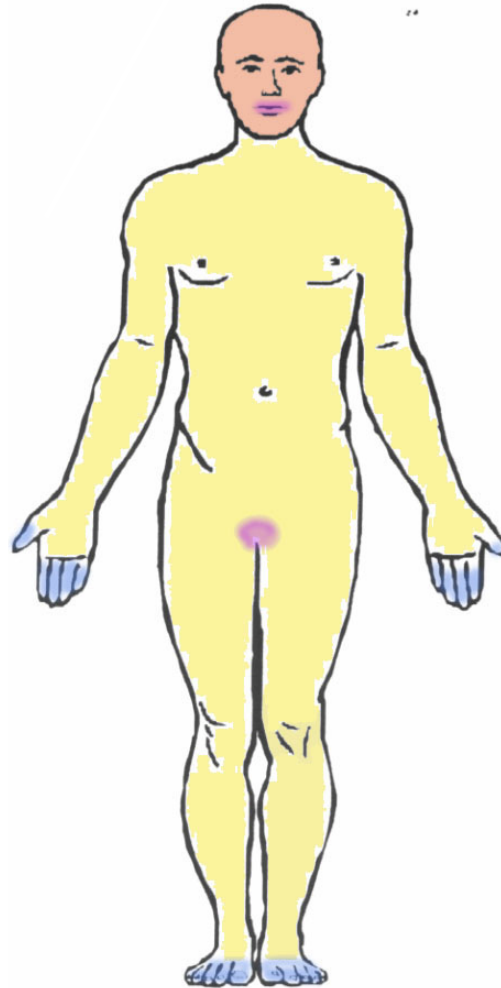
- Low CSD melanoma (SSM)
- High CSD melanoma (LMM)
- Desmoplastic melanoma
- Acral melanoma
- Mucosal melanoma
- Uveal melanoma
- Spitz melanoma
- Melanoma arising from giant congenital nevus
- Melanoma arising from a blue nevus



High CSD



Low CSD

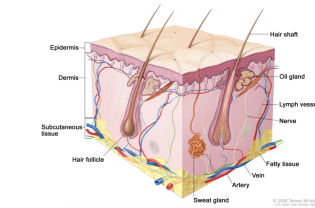
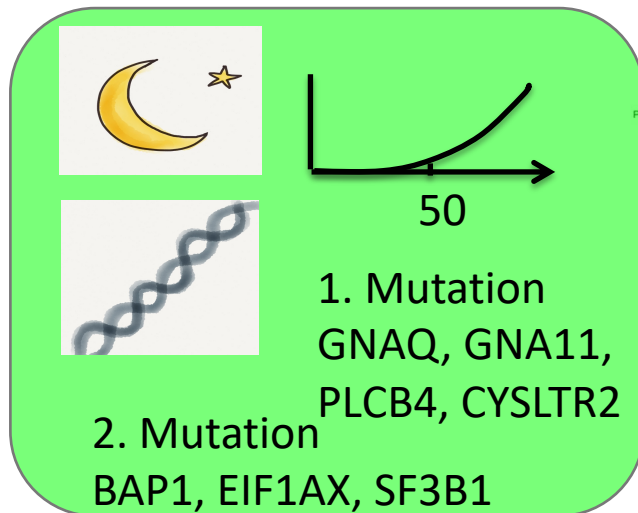


Mucosa

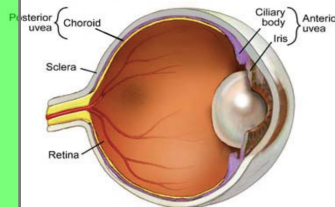


Acral

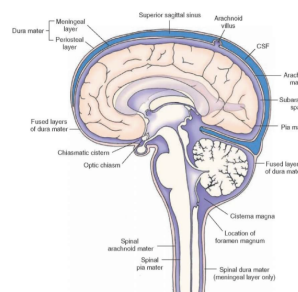
Neoplasms originating from extraepithelial melanocytes



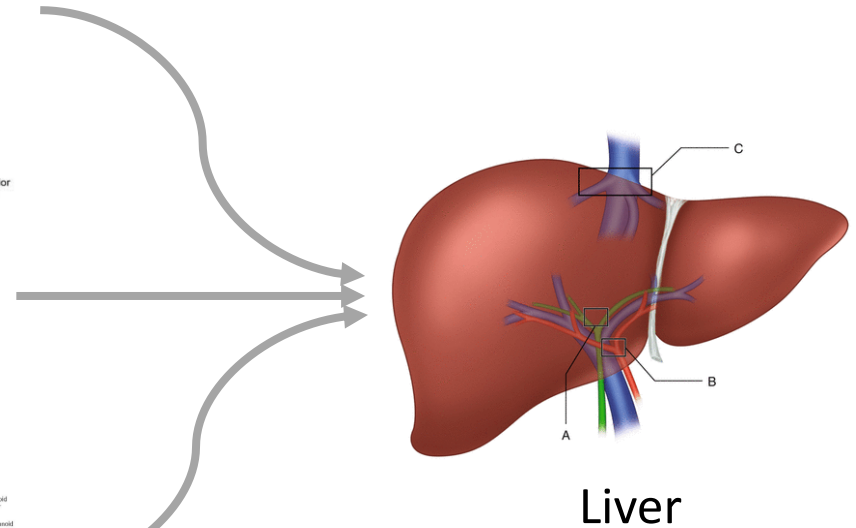
Dermis



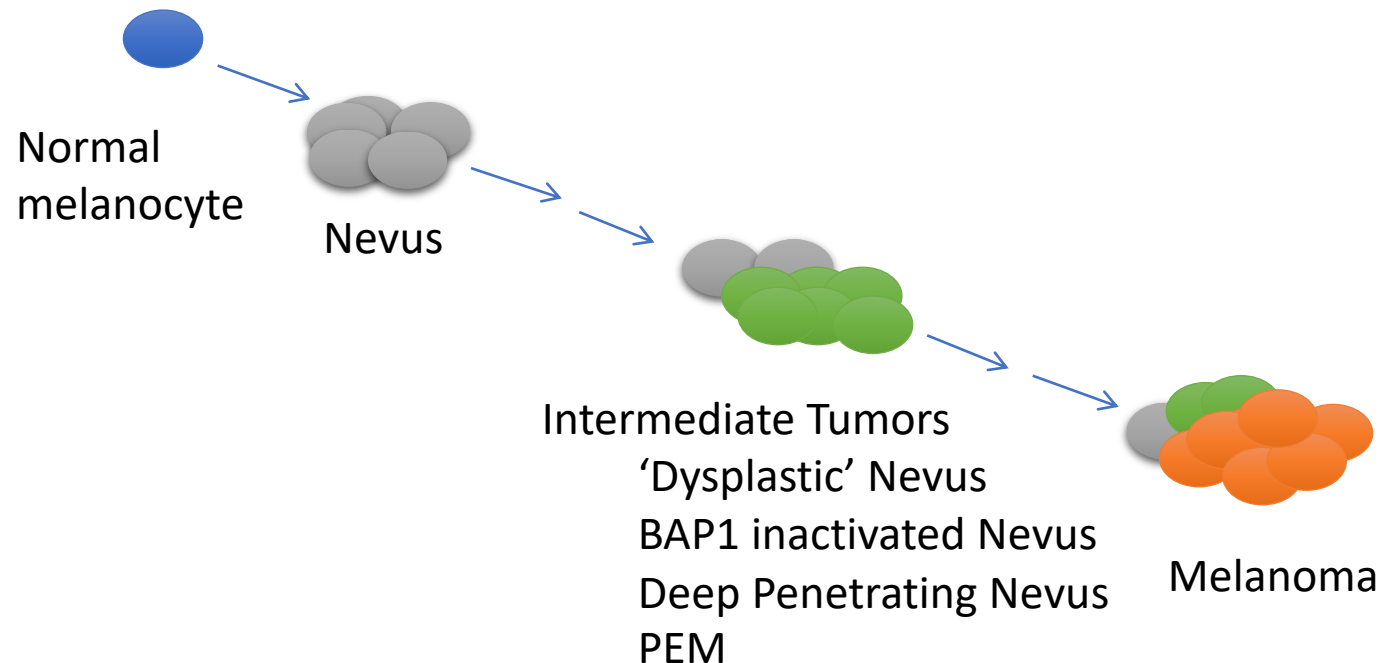
Choroid



Leptomeninges

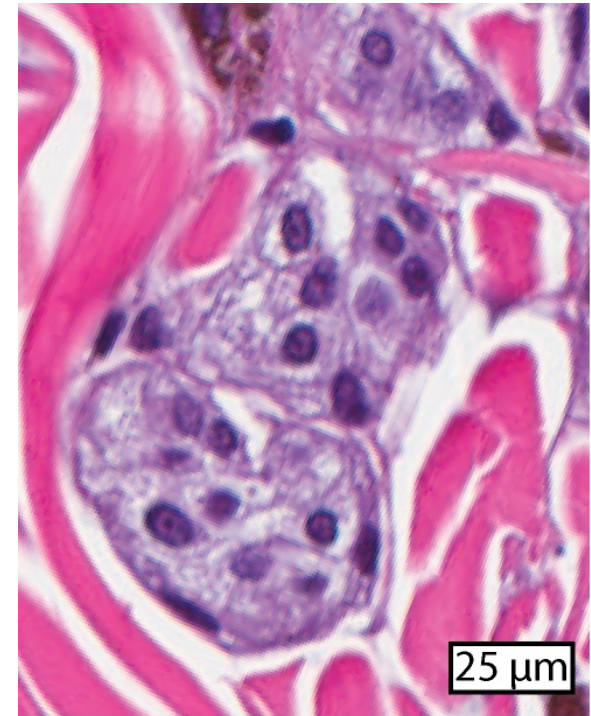
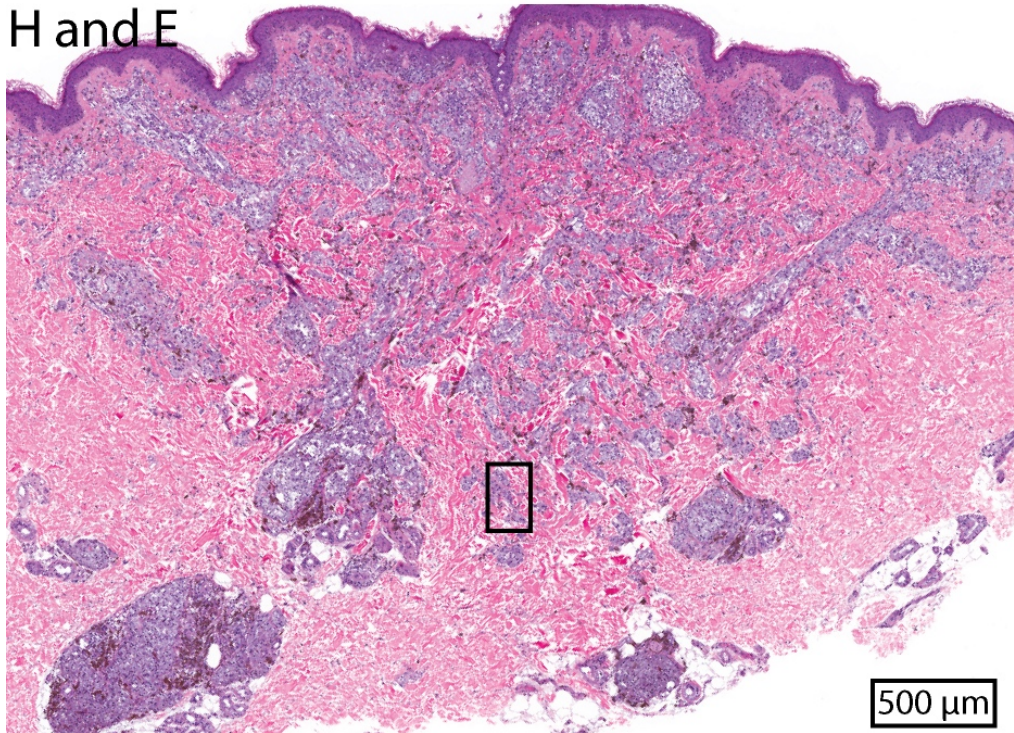


Genetic evolution of melanocytic neoplasms



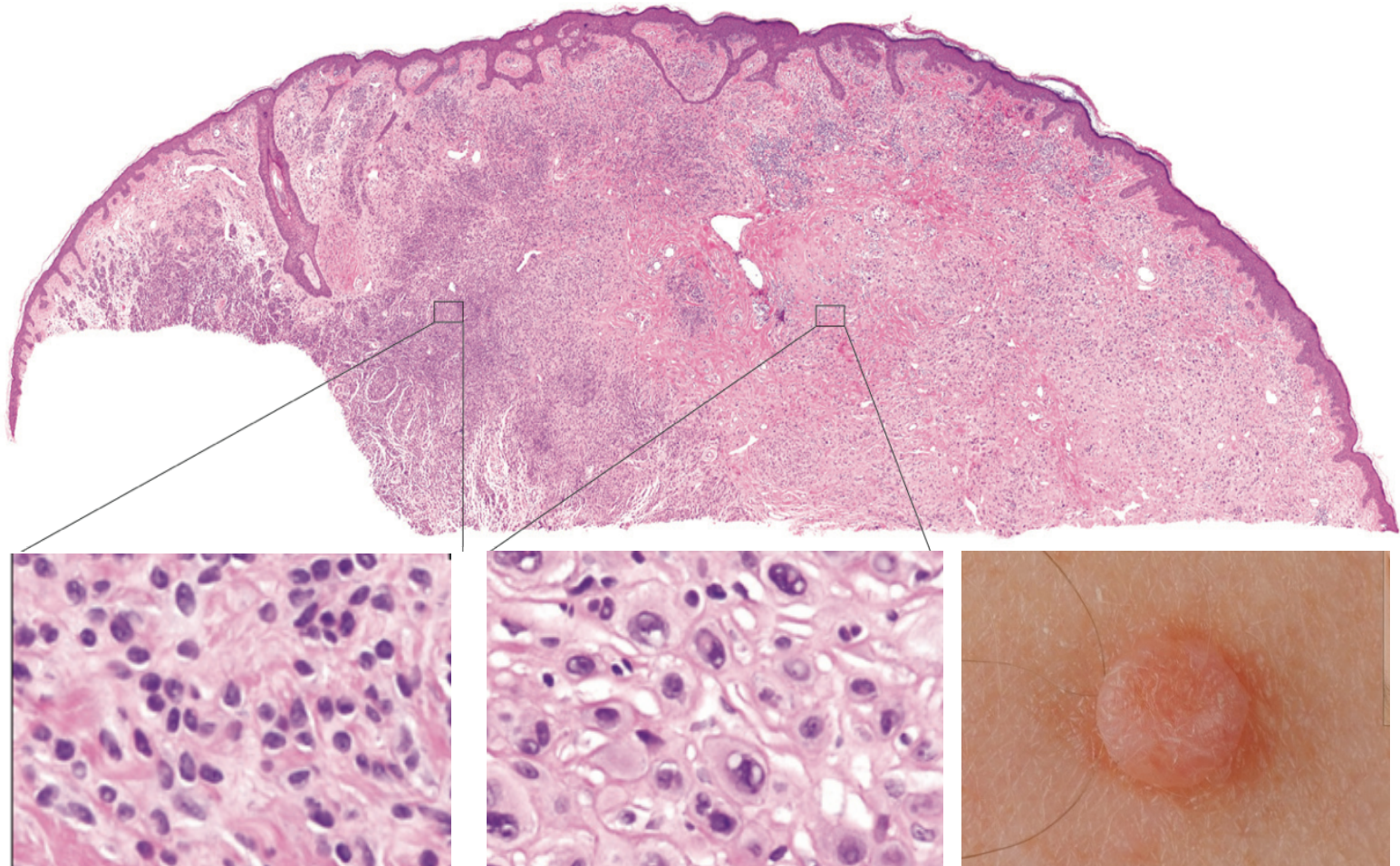
Deep penetrating nevus

H and E



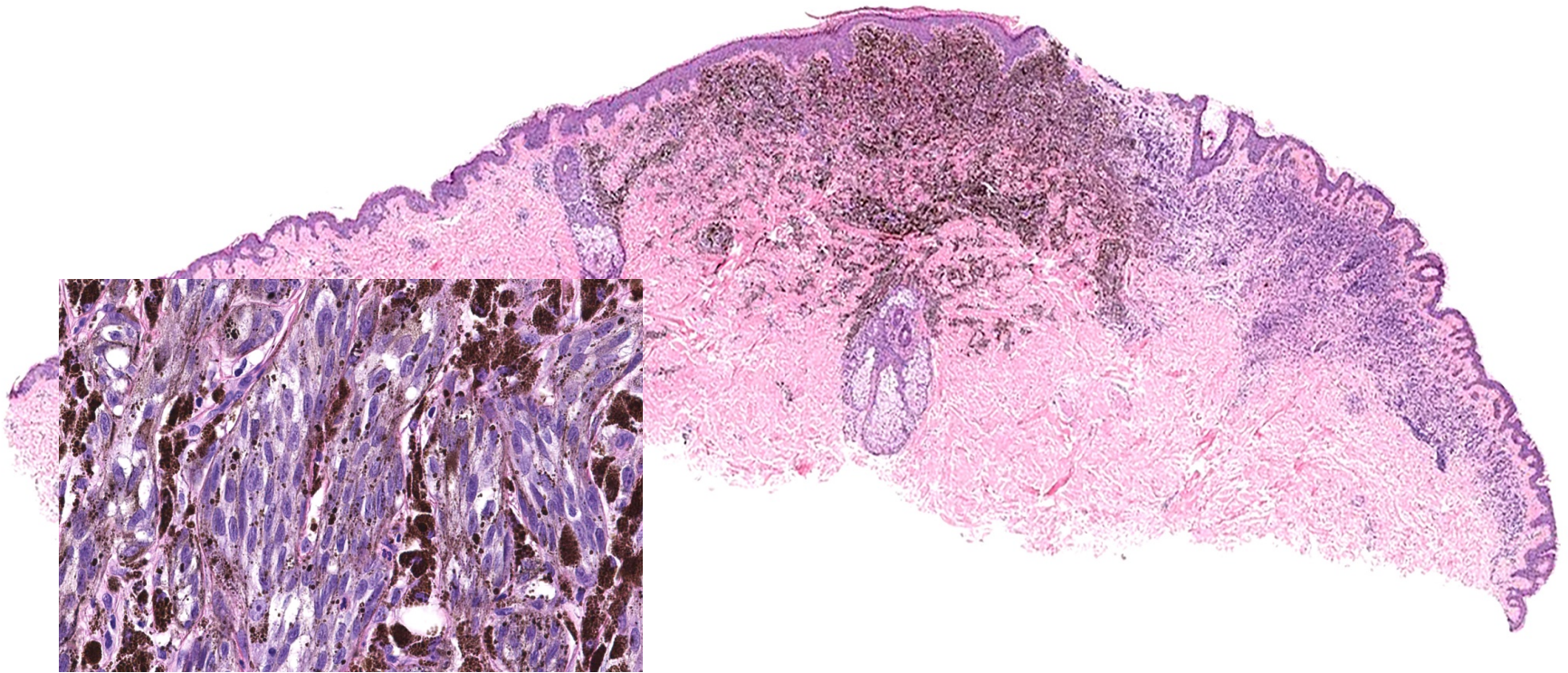
DPN = MAP-kinase pathway + WNT pathway

BAP1-inactivated Spitzoid tumor



BAP1oma = MAP-kinase pathway + BAP1^{-/-}

Pigmented epithelioid melanocytoma



PEM = MAP-kinase pathway + PKA pathway

WHO “Pathways” to melanomas

Pathway I	Low-CSD melanoma/superficial spreading melanoma
Pathway II	High-CSD melanoma/lentigo maligna melanoma
Pathway III	Desmoplastic melanoma
Pathway IV	Malignant Spitz tumour (Spitz melanoma)
Pathway V	Acral melanoma
Pathway VI	Mucosal melanoma
Pathway VII	Melanoma arising in congenital naevus
Pathway VIII	Melanoma arising in blue naevus
Pathway IX	Uveal melanoma

Revised WHO classification, 4th Edition

	Low UV radiation exposure / CSD				High UV radiation exposure / CS	
Pathway	I				II	III
Endpoint of pathway	Low-CSD melanoma / SSM				High-CSD melanoma / LMM	Desmoplastic melanoma
Benign neoplasms (naevi)	Naevus				? IMP	? IMP
Intermediate/low-grade dysplasias and melanocytomas	Low-grade dysplasia	BIN	DPN		? IAMP/dysplasia	? IAMP/dysplasia
Intermediate/high-grade dysplasias and melanocytomas	High-grade dysplasia / MIS	<i>BAP1</i> -inactivated melanocytoma / MELTUMP	Deep penetrating melanocytoma / MELTUMP	PEM / MELTUMP	Lentigo maligna (MIS)	MIS
Malignant neoplasms	Low-CSD melanoma / SSM (VGP)	Melanoma in BIN (rare)	Melanoma in DPN (rare)	Melanoma in PEM (rare)	LMM (VGP)	Desmoplastic melanoma
Common mutations ^{a,b}	BRAF p.V600E; NRAS <i>TERT;</i> <i>CDKN2A;</i> <i>TP53;</i> <i>PTEN</i>	BRAF or NRAS + <i>BAP1</i>	BRAF , MAP2K1 , or NRAS + CTNNB1 or APC	BRAF + PRKAR1A or PRKCA	NRAS ; BRAF (<i>non-p.V600E</i>); KIT ; <i>NF1</i> <i>TERT;</i> <i>CDKN2A;</i> <i>TP53;</i> <i>PTEN;</i> RAC1	<i>NF1</i> ; <i>ERBB2</i> ; MAP2K1 ; MAP3K1 ; BRAF ; <i>EGFR</i> ; MET <i>TERT</i> ; <i>NFKBIE</i> ; NRAS ; PIK3CA ; PTPN11

BIN, *BAP1*-inactivated naevus; **BN**, blue naevus; **CBN**, cellular blue naevus; **CN**, congenital naevus; **CSD**, cumulative sun damage; **DPN**, deep penetrating naevus; **IAMP**, intraepidermal atypical melanocytic proliferation; **IAMPUS**, intraepidermal atypical melanocytic proliferation of uncertain significance; **IMP**, intraepidermal melanocytic proliferation without atypia; **LMM**, lentigo maligna melanoma; low/high-CSD melanoma, melanoma in skin with a low/high degree of cumulative sun damage; **MELTUMP**, melanocytic tumour of uncertain malignant potential; **MIS**, melanoma in situ; **PEM**, pigmented epithelioid melanocytoma; **SSM**, superficial spreading melanoma; **STUMP**, spitzoid tumour of uncertain malignant potential; **UV**, ultraviolet; **VGP**, vertical growth phase (tumorigenic and/or mitogenic melanoma).

Revised WHO classification, 4th Edition

Low to no (or variable/incidental) UV radiation exposure / CSD

IV	V	VI	VII	VIII	IX
Malignant Spitz tumour / Spitz melanoma	Acral melanoma	Mucosal melanoma	Melanoma in CN	Melanoma in BN	Uveal melanoma
Spitz naevus	? Acral naevus	? Melanosis	CN	Blue naevus	? Naevus?
Atypical Spitz tumour (melanocytoma)	IAMP / dysplasia	Atypical melanosis / dysplasia / IAMPUS	Nodule in CN (melanocytoma)	(Atypical) CBN (melanocytoma)	?
STUMP / MELTUMP	Acral MIS	Mucosal MIS	MIS in CN	Atypical CBN	?
Malignant Spitz tumour / Spitz melanoma (tumorigenic)	Acral melanoma (VGP)	Mucosal lentiginous melanoma (VGP)	Melanoma in CN (tumorigenic)	Melanoma in blue naevus (tumorigenic)	Uveal melanoma
HRAS ; ALK ; ROS1 ; RET ; NTRK1 ; NTRK3 ; BRAF ; MET CDKN2A	KIT ; NRAS ; BRAF ; HRAS ; KRAS ; NTRK3 ; ALK ; NF1 CDKN2A ; TERT ; CCND1 ; GAB2	KIT ; NRAS ; KRAS or BRAF NF1 ; CDKN2A ; SF3B1 ; CCND1 ; CDK4 ; MDM2	NRAS ; BRAF p.V600E (small lesions) ; BRAF	GNAQ ; GNA11 ; CYSLTR2 BAP1 ; EIF1AX ; SF3B1	GNAQ ; GNA11 ; CYSLTR2 ; or PLCB4 SF3B1 ; EIF1AX ; BAP1

Definitions: *Melanocytoma* is a tumorigenic neoplasm of melanocytes that generally has increased cellularity and/or atypia (compared with a common naevus) and an increased (although generally still low) probability of neoplastic progression; *tumorigenic* means forming a mass of neoplastic cells.

^a Common mutations in each pathway are listed. Mutations already identified in benign or borderline low lesions are shown in bold.

^b Blue, loss-of-function mutation; red, gain-of-function mutation; green, change-of-function mutation; orange, amplification; purple, rearrangement; grey, promoter mutation.