

South Beach
Symposium
medical + aesthetic dermatology

CELEBRATING 20 YEARS
OF PREMIER MEDICAL & AESTHETIC
DERMATOLOGY EDUCATION

REIMAGINING
MEDICAL AND
AESTHETIC
DERMATOLOGY



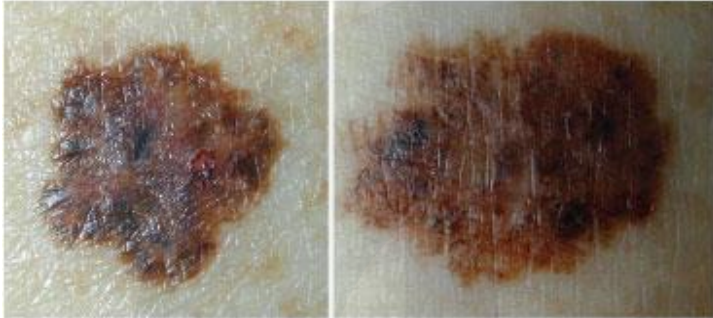
Non-Invasive Detection for Melanoma

REVOLUTIONIZING DERMATOLOGY EDUCATION

Laura K. Ferris, MD, PhD

Professor of Dermatology
University of Pittsburgh

Making biopsy decisions for pigmented lesions



Non-melanoma

Melanoma

How good are we?

- Reader studies:
 - ❖ Sensitivity 65-82%
- NNB: 2.2 – 30.5 (mean 13.2 for US Dermatology practitioners)



Approaches

Genetic analysis

Physical properties

- Visual
- Conduction

Gene expression profiling (GEP)

Benign



Applications in melanoma:
Non-invasive test to help:

1. Distinguish melanoma from nevi

2. Classify histologically equivocal biopsied melanocytic lesions

3. Predict which tumors are at highest risk of metastasis

Malignant

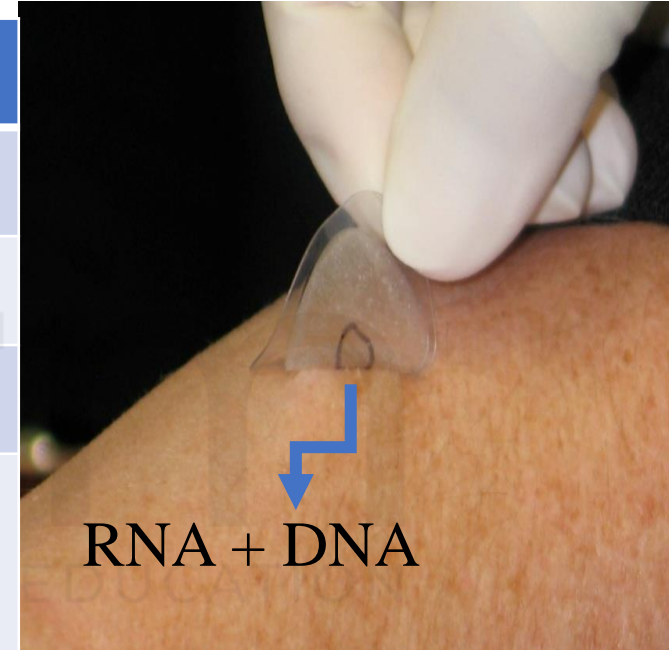


PLA for non-invasive diagnosis of melanoma

J Am Acad Dermatol 2017;76:114-20.

JAMA Dermatol. 2017 Jul 1;153(7):675-680

Genetic finding	% histologic melanoma
LINC (RNA) +	7%
PRAME (RNA) +	50%
LINC <i>and</i> PRAME (RNA) +	93%
TERT promoter mutation (DNA) present	79%



SENSITIVITY:

- ▶ 97% of melanomas express LINC, PRAME, and/or TERT

SPECIFICITY:

- ▶ 48% of non-melanomas are negative for LINC, PRAME *AND* TERT

ARTICLE

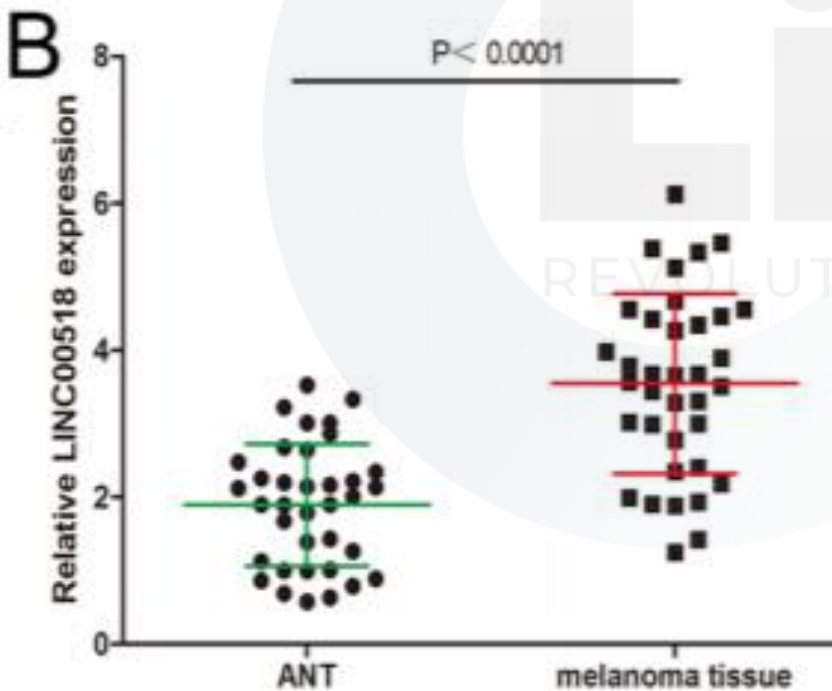
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LINC00518

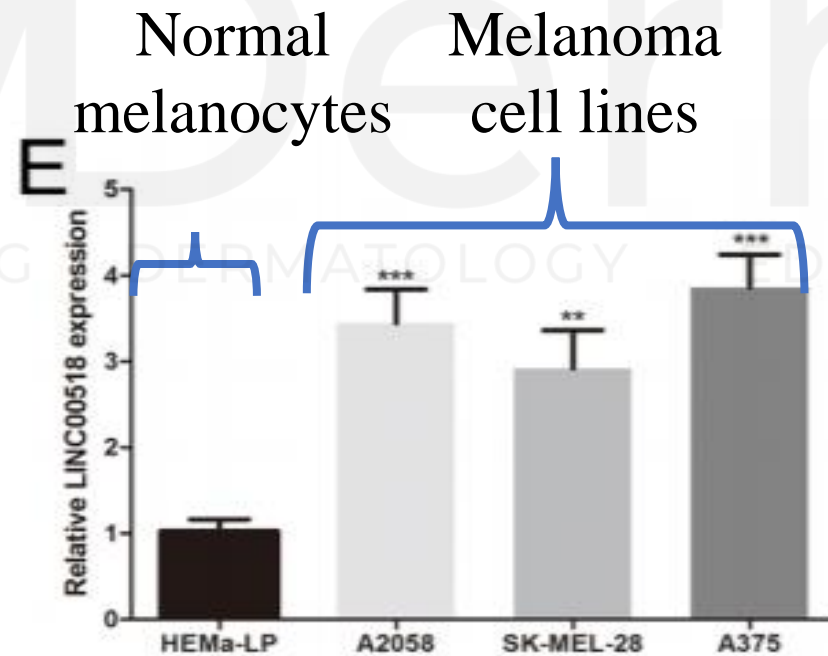
- promotes melanoma invasion and metastasis
- Higher expression associated with lower melanoma survival

Long noncoding RNA LINC00518 acts as a competing endogenous RNA to promote the metastasis of malignant melanoma via miR-204-5p/AP1S2 axis

Wenkang Luan¹, Yuting Ding², Shaojun Ma¹, Hongru Ruan¹, Jinlong Wang¹ and Feng Lu¹



Primary melanoma



Article 2020 Dec 21;12(12):3867

LncRNA *LINC00518* Acts as an Oncogene in Uveal Melanoma by Regulating an RNA-Based Network

Cristina Barbagallo ¹, Rosario Caltabiano ², Giuseppe Broggi ², Andrea Russo ³, Lidia Puzzo ², Teresio Avitabile ³, Antonio Longo ³, Michele Reibaldi ³, Davide Barbagallo ¹, Cinzia Di Pietro ¹, Michele Purrello ¹ and Marco Ragusa ^{1,*}



ORIGINAL RESEARCH
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BBA - Molecular Basis of Disease 1865 (2019) 708–723

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ELSEVIER

BBA - Molecular Basis of Disease

journal homepage: www.elsevier.com/locate/bbadis

Down-regulated expression of LINC00518 prevents epithelial cell growth and metastasis in breast cancer through the inhibition of CDX2 methylation and the Wnt signaling pathway

Hong-Bin Wang^{a,1}, Hong Wei^{b,1}, Jin-Song Wang^a, Lin Li^a, An-Yue Chen^a, Zhi-Gao Li^{a,*}

LINC00518 Promotes Cell Proliferation by Regulating the Cell Cycle of Lung Adenocarcinoma Through miR-185-3p Targeting MECP2

Xu Han[†], Jixiang Wu[†], Yajun Zhang^{*}, Jianxiang Song, Zhan Shi and Huiwen Chang

PRAME (PReferentially expressed Antigen in MElanoma) in melanoma vs. nevi (IHC)

Melanoma Type	In Situ Only	Invasive	Total
Superficial spreading	12/12	37/41	49/53
Lentigo maligna	24/27	15/17	39/44
Acral	7/7	10/11	17/18
Nodular	NA	9/10	9/10
Other*	2/2	6/8	8/10
Subtotal [†]	45/48	77/87	122/135
Desmoplastic [‡]	NA	7/20	7/20
Total	45/48	84/107	129/155

Type of Melanocytic Nevus	Diffuse (4+) IHC PRAME Expression	Focal (1 or 2+) IHC PRAME Expression
Common acquired nevus	0/40	4/40 (1+)
Dysplastic (Clark's) nevus	0/60	10/60 (1+)
		1/60 (2+)
Blue nevus	0/10	0/10
Spitz nevus	1/10	1/10 (1+)
Deep penetrating nevus	0/3	0/3
Traumatized/ recurrent nevus	0/15	1/15 (2+)
		1/15 (1+)
Congenital nevus	0/2	0/2
Nodal nevus	0/5	0/5
Total	1/145	18/145

83% PRAME +
(90% of non-desmoplastic melanoma)

13% PRAME +

Utility of *TERT* promoter mutations for cutaneous primary melanoma diagnosis

Nancy E. Thomas, MD, PhD^{*,†}, Sharon N. Edmiston, BS^{*,†}, Yihsuan S. Tsai, PhD[†], Joel S. Parker, PhD^{†,‡}, Paul B. Googe, MD^{*,§}, Klaus J. Busam, MD[¶], Glynis A. Scott, MD^{||,*}, Daniel C. Zedek, MD^{*,§}, Eloise A. Parrish, MS[†], Honglin Hao^{*}, Nathaniel A. Slater, MD^{*}, Michelle V. Pearlstein, MD^{*}, Jill S. Frank, MS^{†,††}, Pei Fen Kuan, PhD^{‡‡}, David W. Ollila, MD^{†,††}, and Kathleen Conway, PhD^{*,†,§§}

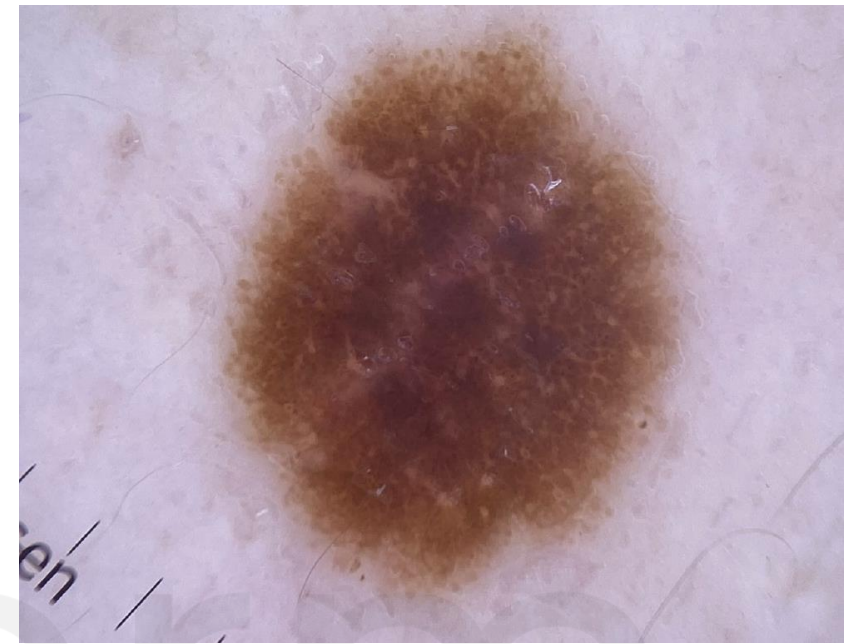
- *TERT* (telomerase reverse transcriptase) – maintains telomeres; mutations lead to uncontrolled replication and proliferation of cancer cells
- *TERT* promotor mutations identified in:
 - 67/87 (77.9%) melanomas
 - 1/72 (1.4%) nevi

Lower rates of *TERT* positivity in:

- Acral melanoma
- Non-white race
- Younger patients
- Non-sun damaged skin
- Lower extremity lesions

Am J Dermatopathol. 2019
April ; 41(4): 264–272.

38 yo F, h/o MMIS
10 biopsies in past 13 months



PLA:

LINC: Detected

PRAME: Not Detected

Pathology:

EARLY EVOLVING MALIGNANT MELANOMA
IN-SITU ARISING IN ASSOCIATION WITH A
DYSPLASTIC COMPOUND NEVUS

Lesion on cheek, no change per patient



PLA:

LINC: Detected

PRAME: Not Detected

TERT promoter

mutation: Not Detected



Pathology:

EARLY EVOLVING MALIGNANT MELANOMA
IN-SITU ARISING IN ASSOCIATION WITH A
DYSPLASTIC COMPOUND NEVUS

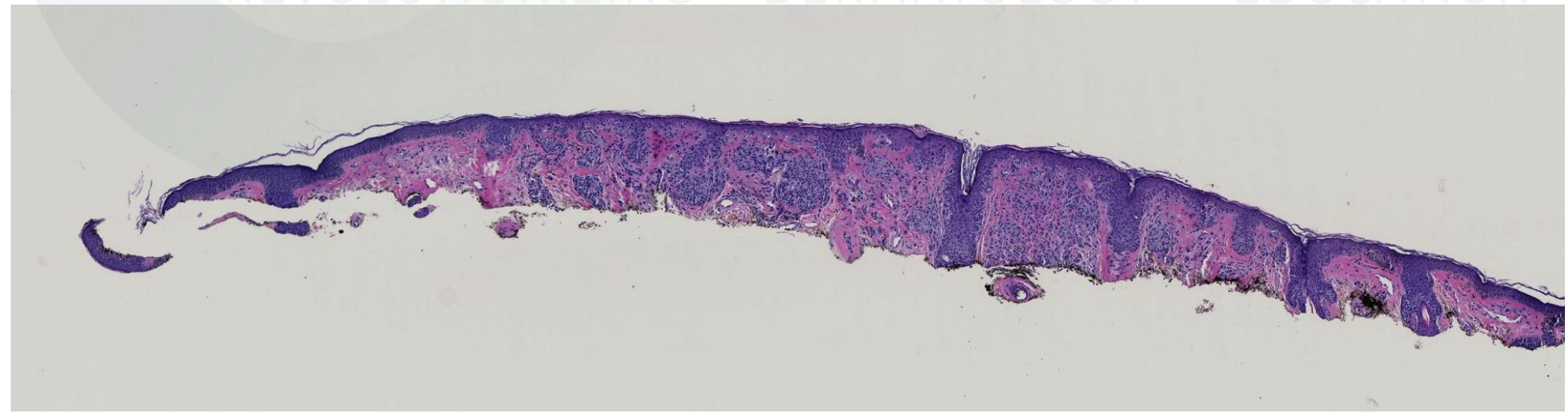


GENE EXPRESSION RISK STATUS: MODERATE (ORANGE)	LINC00518: PRAME:	Not Detected Detected ←
MUTATION RISK STATUS: -	TERT Promoter:	Not Detected

FINAL DIAGNOSIS:

SKIN, LEFT CHEEK, SHAVE:

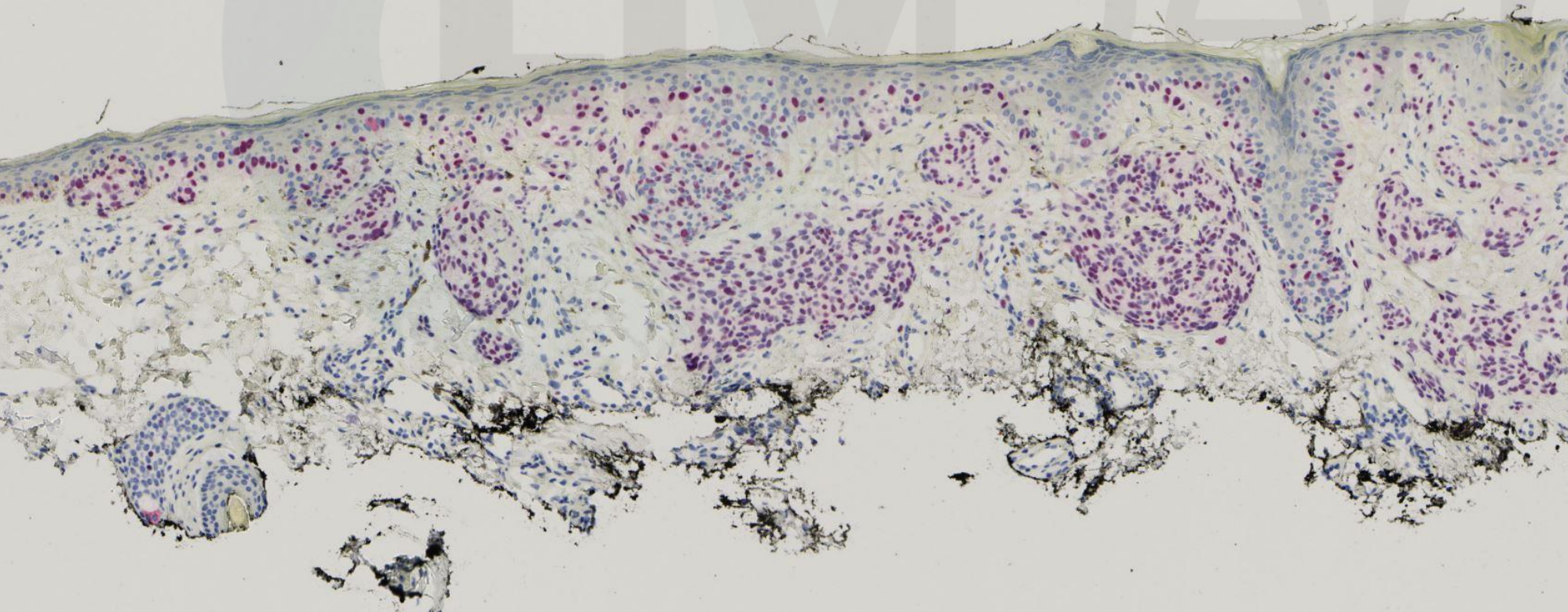
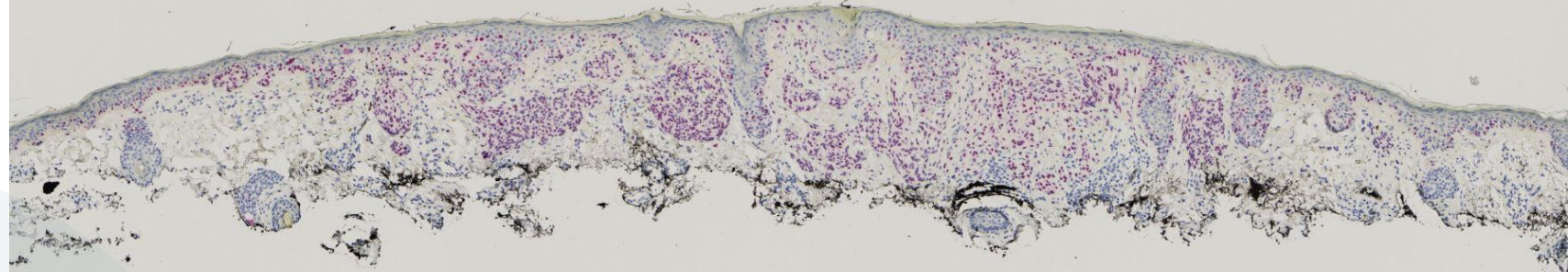
- A. MALIGNANT MELANOMA, LENTIGO MALIGNA TYPE
- B. THE DEPTH OF INVASION (Breslow's thickness) IS 0.5 mm AT LEAST.
- C. SURFACE ULCER IS NOT IDENTIFIED.
- D. MITOTIC COUNT IS LESS THAN 1 PER 1mm².



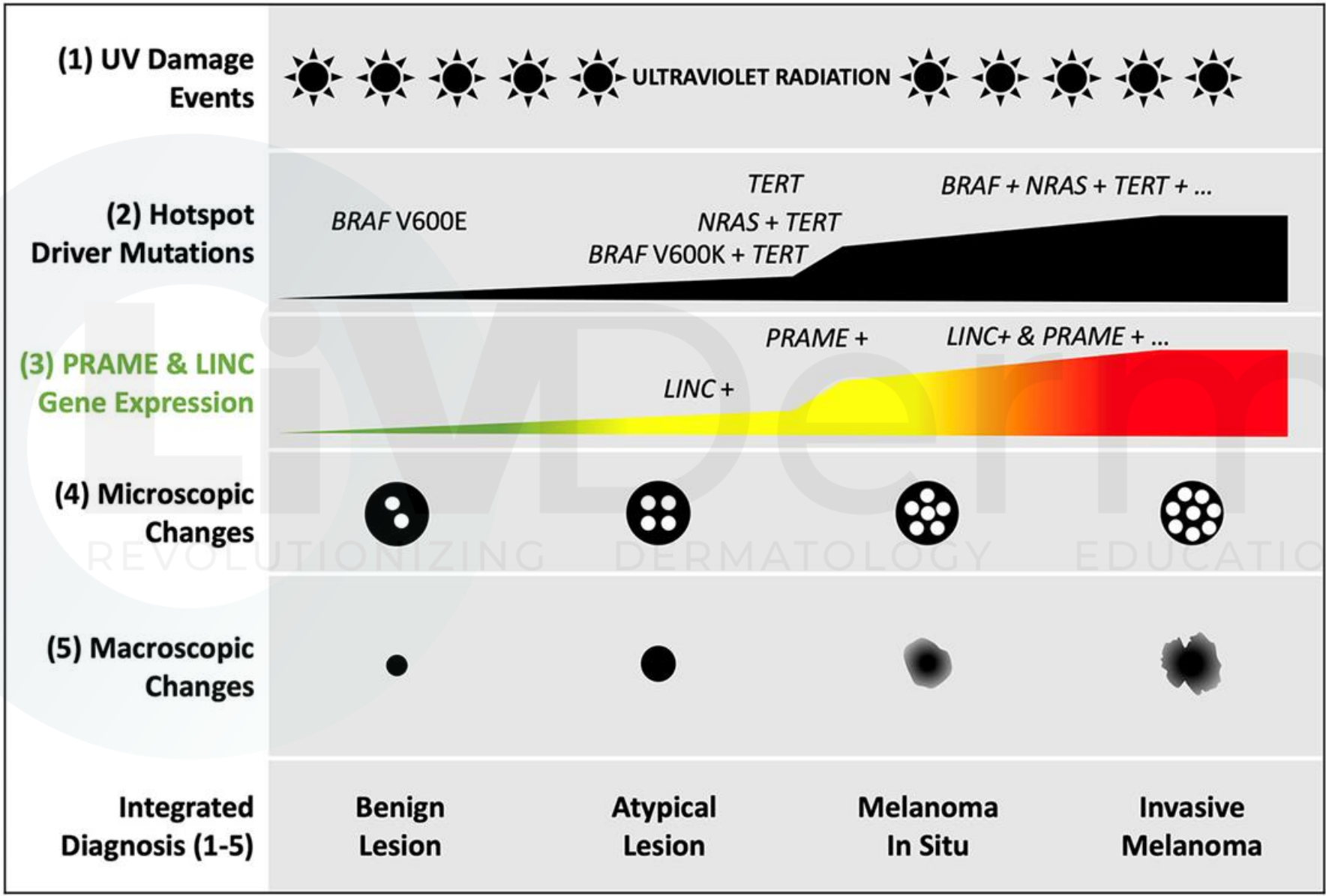
REVOLUTIONIZING DERMATOLOGY EDUCATION

PRAME IHC

Courtesy of Arivarasan
Karunamurthy, MD



CATION



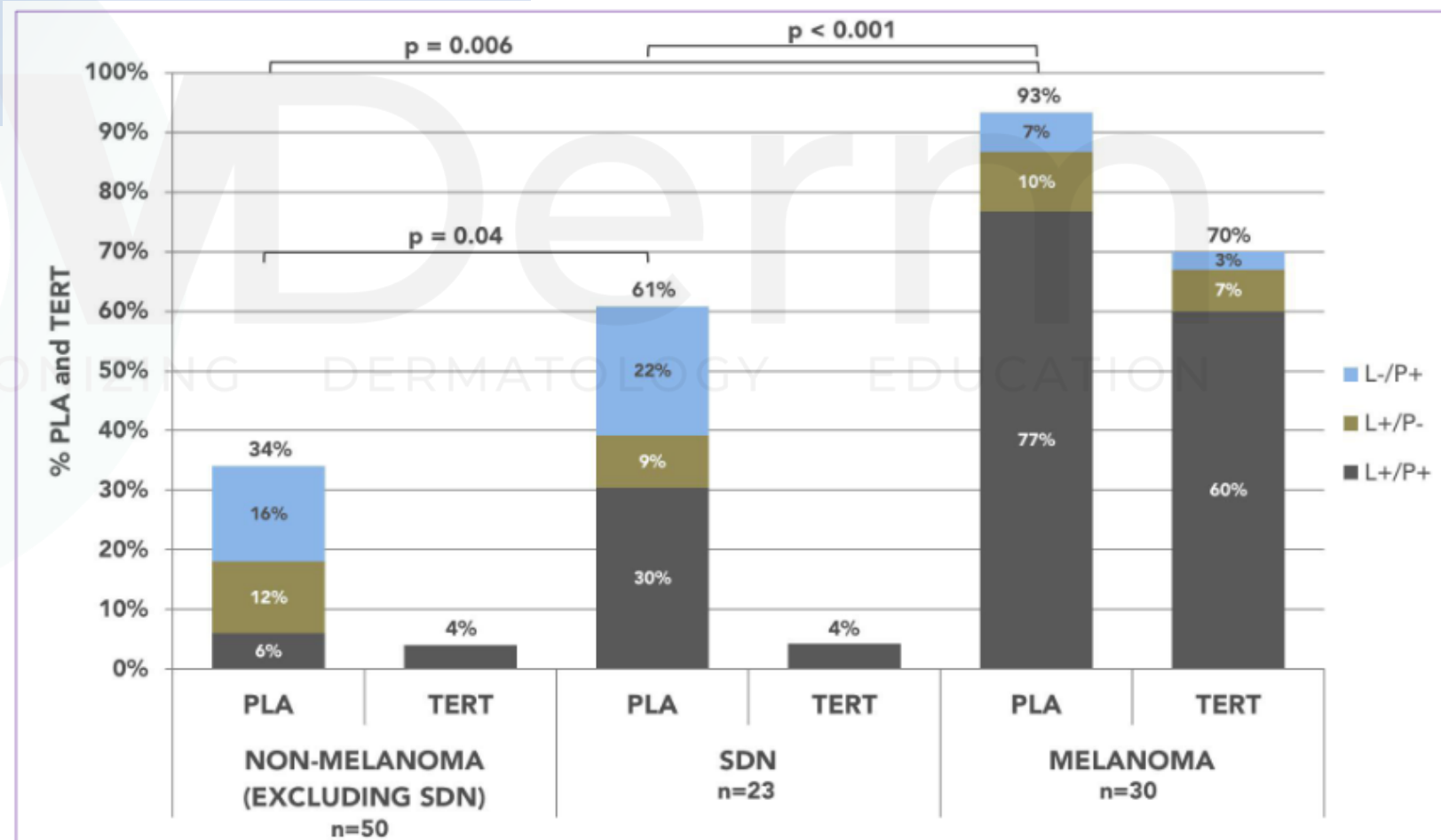
Risk Stratification of Severely Dysplastic Nevi by Non-Invasively Obtained Gene Expression and Mutation Analyses (SKIN, March 2020)

Severely dysplastic nevi:

- Commonly express LINC and/or PRAME
- *Rarely* carry TERT promoter mutation

Melanoma

- Most express LINC and/or PRAME *AND* TERT promoter mutation



What about negative PLA lesions?

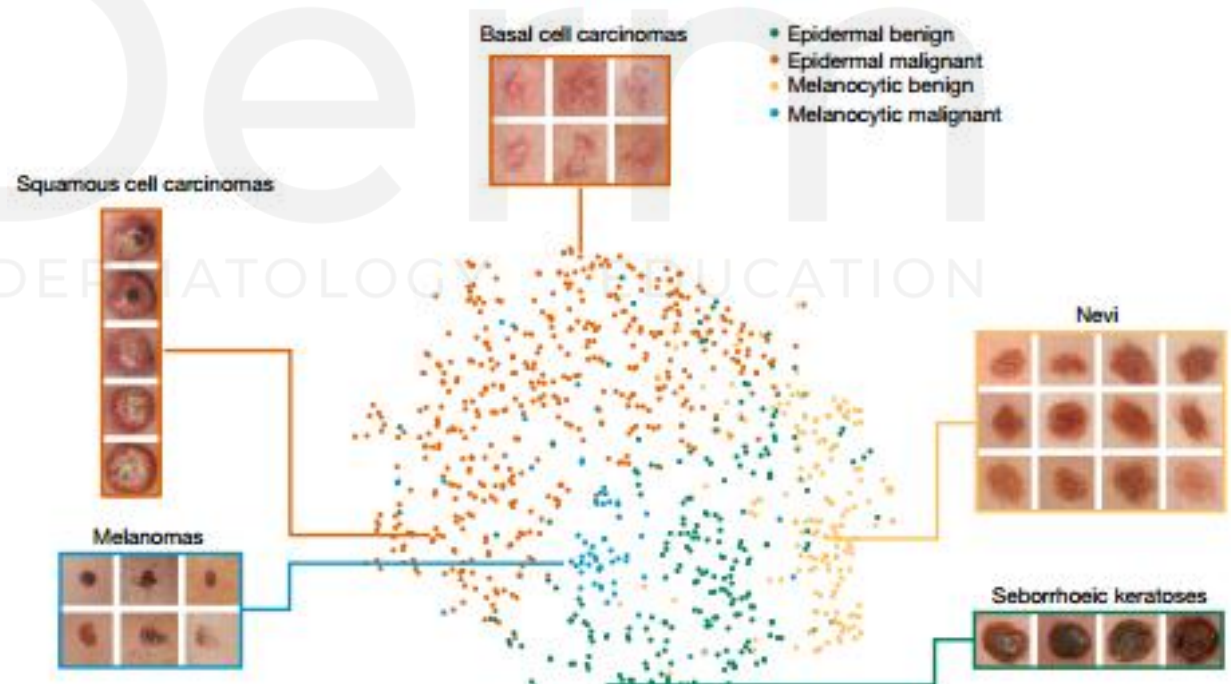
- 1781 PLA negative patients
 - Clinical follow up on 69%
 - 10 (0.8%) diagnosed in next ~ 1 yr with melanoma (in situ, stage I)
- 304 PLA negative patients, retested 6-12 mo later
 - 34 (11%) were PLA positive, all biopsied
 - 3 (1%) melanomas (all in situ)

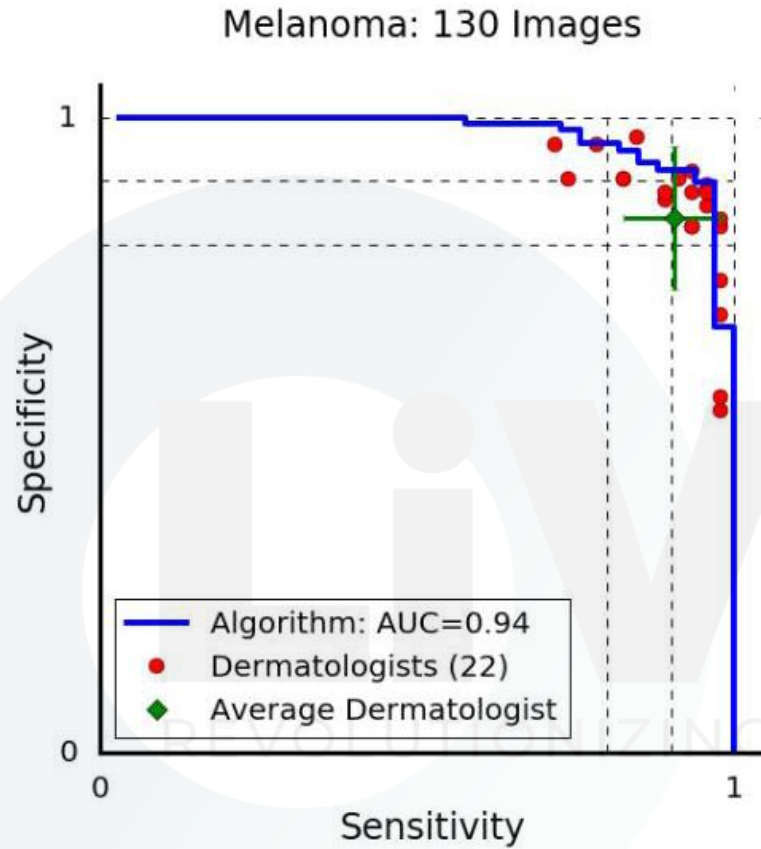
Negative predictive value >99%

Dermatologist-level classification of skin cancer with deep neural networks

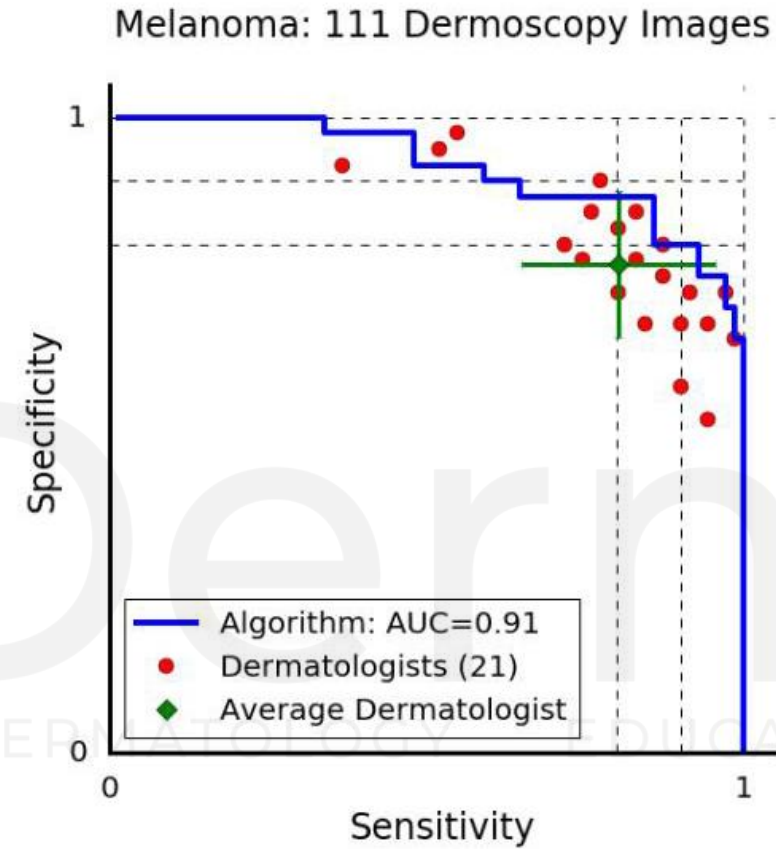
- Trained on >100,000 skin lesion images of >2,000 diseases
- Compared accuracy to dermatologists
 - Melanoma vs nevi
 - BCC / SCC vs SK
 - NOT SK vs melanoma!

Clinical or dermatoscopic images





33 melanomas, 97 benign nevi



71 melanomas, 40 benign nevi

Red dot above the curve = Dermatologist outperformed computer
 Red dot below the curve = Computer outperformed the dermatologist

No data on lesion thickness

Sensitivity	Specificity	Negative predictive value	Positive predictive value
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Discrimination of malignant from benign pigmented skin lesion

All human readers (n=511)

Melanoma	73.1%	92.8%	96.4%	56.6%
	(65.8–79.6)	(91.3–94.2)	(95.3–97.4)	(49.7–63.2)

Expert readers (n=27)

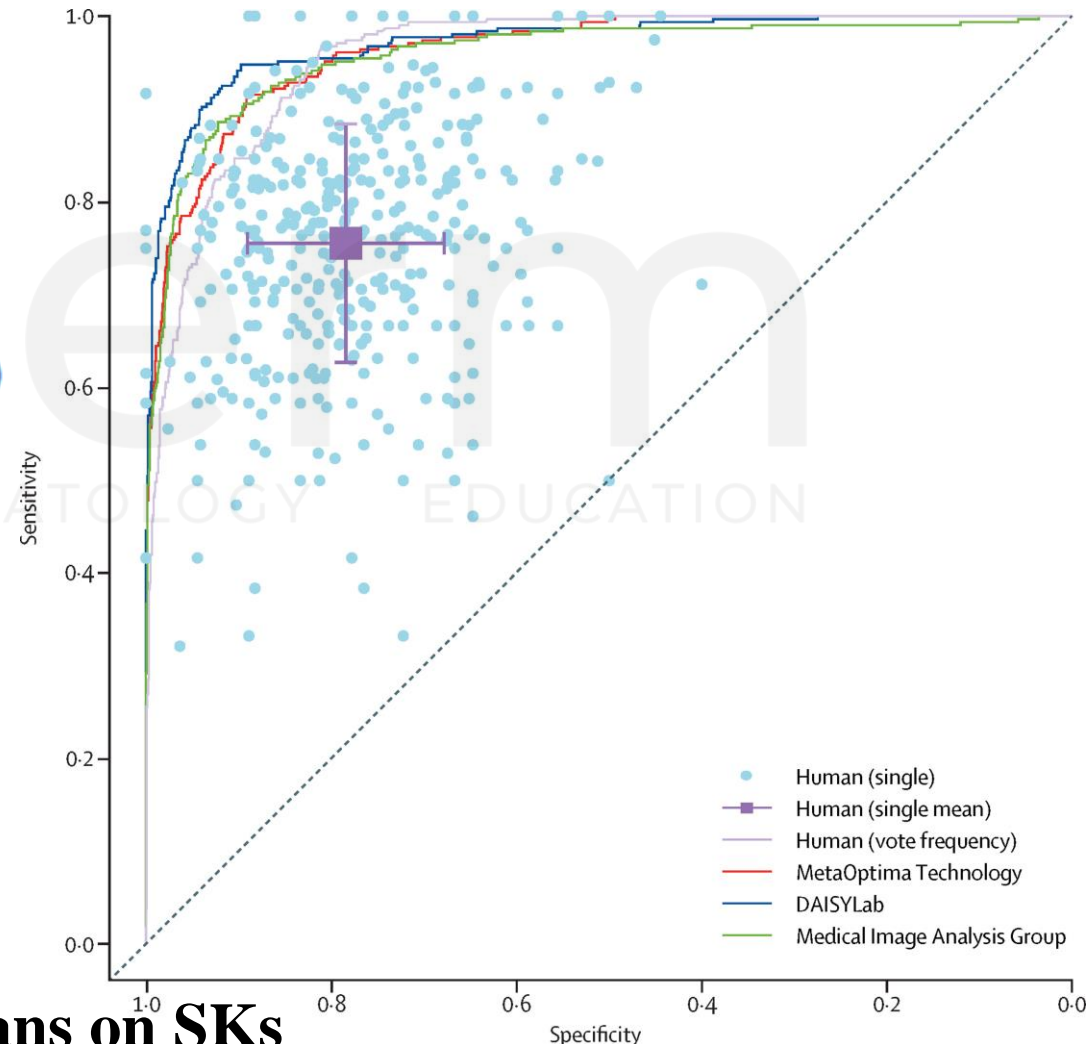
Melanoma	67.8%	94.0%	95.8%	58.9%
	(60.3–74.8)	(92.5–95.2)	(94.6–96.8)	(51.7–65.8)

All algorithms (n=139)

Melanoma	67.3%	97.0%	95.9%	74.2%
	(59.7–74.2)	(96.0–97.9)	(94.7–96.9)	(66.6–80.9)

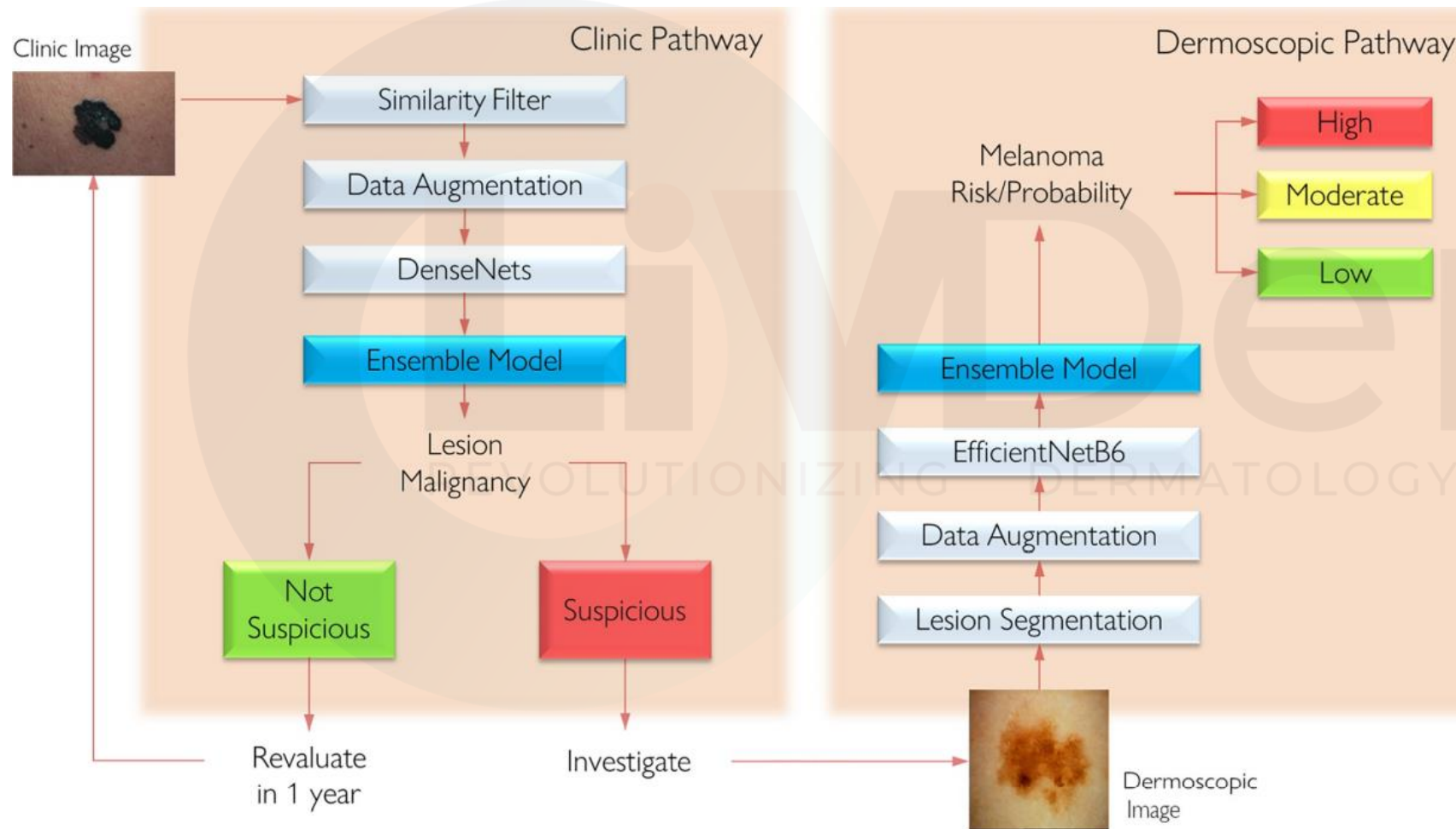
Top three algorithms (n=3)

Melanoma	81.9%	96.2%	97.6%	73.6%
	(75.4–87.3)	(95.1–97.2)	(96.7–98.4)	(66.9–79.6)



Top 3 algorithms also performed equivalent to humans on SKs

AI for melanoma diagnosis in primary care- the future?



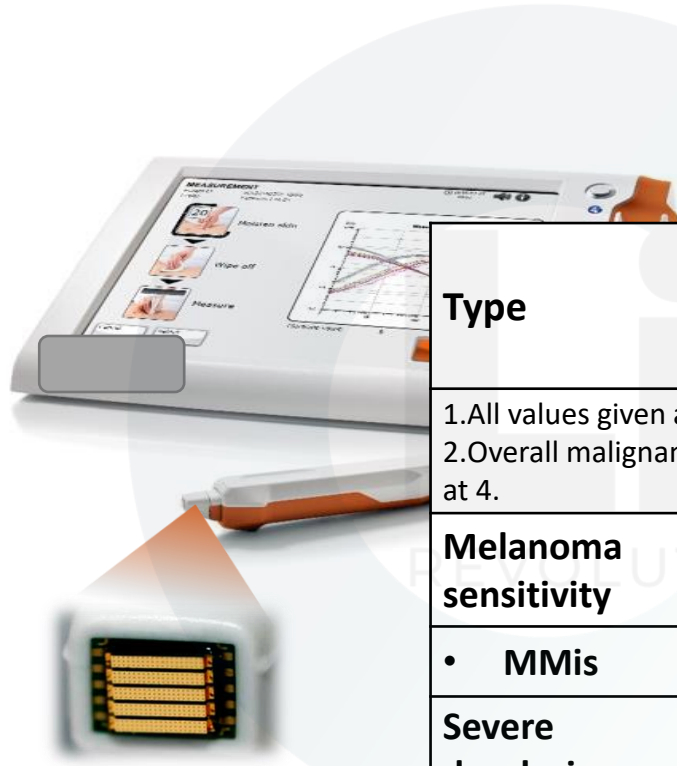
Trained using database of available and generated clinical and dermoscopic images

- Sensitivity : 90%
- Specificity: 85-89%
- PPV: 59-65%
- NPV: 97%

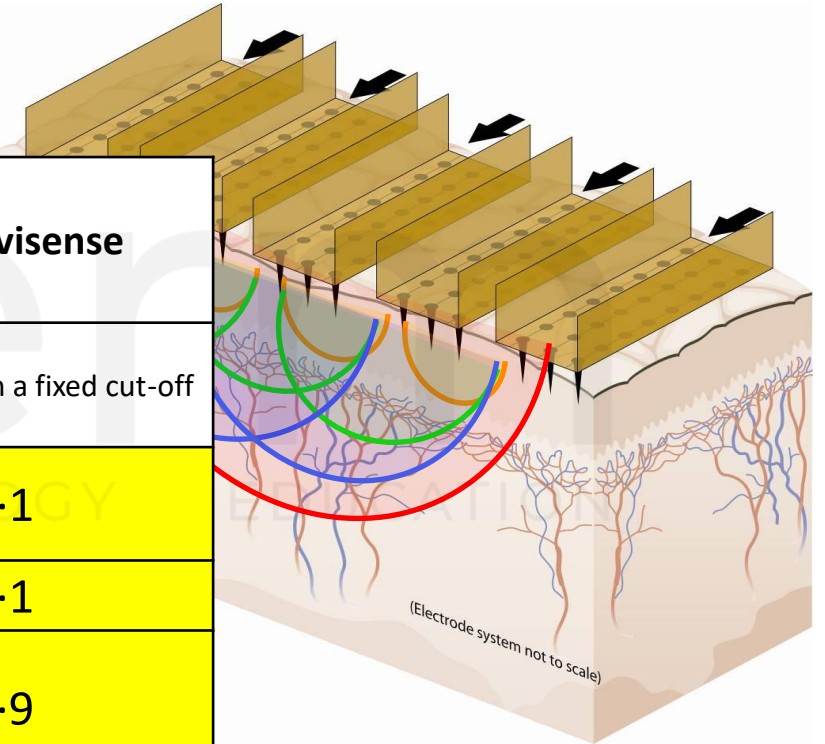
CAD system pathways developed to provide outputs for clinical and dermoscopy images to primary care physicians.

Electrical Impedance Spectroscopy (EIS)

FDA approved 2017 for use by dermatologists



Type	Seven-point	Malignancy grading	Local pathology	Nevisense
1.All values given as a percentage. NA, not applicable. 2.Overall malignancy grading as determined by the visual classification board with a fixed cut-off at 4.				
Melanoma sensitivity	49.2	70.6	84.5	97.1
• MMis	28.7	55.4	73.3	94.1
Severe dysplasia sensitivity	12.1	38.3	NA	83.9
Overall specificity	94.2	81.4	98.0	35.8



Br J Dermatol. 2014.
Nov;171(5):1099-107.
FDA report.

NOTE: sensitivity 57.1% in patients <30 years of age (small n)

MELAFIND- What can we learn?

- Sensitivity comes at the cost of specificity
- Limited utility: Recommends biopsy of about 90% of lesions
- Extremely expensive optics/ machine
- Fixed classifier- cannot “learn” in real time → need FDA reapproval
- Product no longer available / supported



	Sensitivity	Specificity
MelaFind	97%	9%
Readers	72%	51%

Threshold	Sensitivity (95% CI)	Specificity (95% CI)
≥ -3	100.0% (97.9-100%)	0.8% (0.4-1.4%)
≥ -2	99.4% (96.9-100%)	1.3% (0.8-2.0%)
≥ -1	98.9% (95.9-99.9%)	3.6% (2.7-4.7%)
≥ 0	98.3% (95.1-99.6%)	10.8% (9.2-12.5%)
≥ 1	93.1% (88.3-96.4%)	29.8% (27.4-32.2%)
≥ 2	75.4% (78.4-81.6%)	60.0% (57.4-62.5%)
≥ 3	54.3% (46.6-61.8%)	81.8% (79.7-83.8%)
≥ 4	30.9% (24.1-38.3%)	91.4% (89.9-92.8%)
≥ 5	15.4% (10.4-21.7%)	96.1% (95.0-97.0%)
≥ 6	7.4% (4.0-12.4%)	98.8% (98.2-99.4%)

Human vs Machine – key differences in determining if a lesion is benign or malignant

Parameter	Human	Machine
Context	Consider all lesions on the skin, patient history, risk factors	Lesion in isolation; only the lesion the user chooses to evaluate
Objectivity	<ul style="list-style-type: none">• Shaded by experience, fear of missing melanoma, incentive to biopsy or not• Favor biopsy of benign over missing malignant	<ul style="list-style-type: none">• Objective• Can choose to maximize sensitivity vs. specificity
Learning	Years: one patient / paper/ textbook at a time	Can train classifier in hours / days
Features evaluated	<ul style="list-style-type: none">• Uses set criteria to evaluate a lesion• Can usually explain “why”	<ul style="list-style-type: none">• Can identify and use new features and process large amounts and layers of data• Cannot always explain “why”

Sensitivity and specificity are just one factor: must consider how tools designed to improve melanoma detection should best be integrated into clinical practice!