

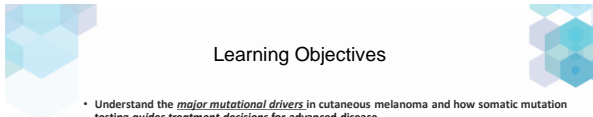


Current and Emerging Biomarkers in Melanoma

Allie H. Grossmann, MD PhD
Associate Professor of Pathology, Division of Anatomic Pathology
Medical Director, Solid Tumor Molecular Oncology, ARUP Laboratories
Investigator, Huntsman Cancer Institute
University of Utah



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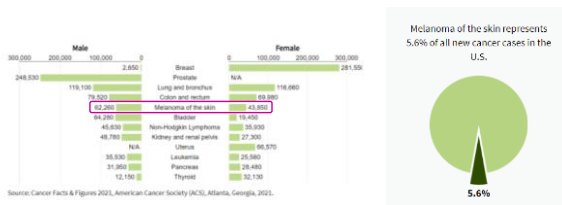
Learning Objectives

- Understand the **major mutational drivers** in cutaneous melanoma and how somatic mutation testing **guides treatment decisions** for advanced disease.
- Understand the **diagnostic utility** of somatic mutation testing for resolving diagnostic uncertainty in metastatic melanoma.
- Realize **unmet clinical needs** where molecular/genomic biomarkers may have utility
 - Improving relapse risk stratification of Stage II-III patients.
 - Predicting survival benefit and immune related adverse events with immune checkpoint blockade.
- Review recent clinical trial and preclinical studies that define a new paradigm for **combining immune checkpoint blockade with targeted therapy**
- Discuss **investigational biomarkers** for melanoma staging and predicting therapeutic response
 - Liquid biopsy, circulating tumor DNA (ctDNA)
 - Inflammatory gene expression profiling of the tumor
 - Tumor mutation burden

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Melanoma is a fairly common cancer

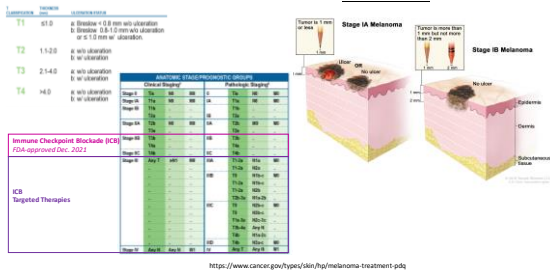
NCI SEER Cancer Database



<https://seer.cancer.gov/statfacts/html/melan.html>

3

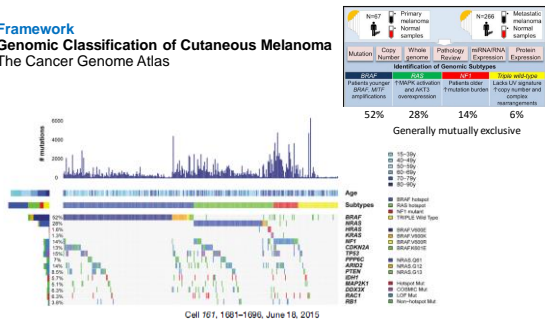
Highly aggressive disease:
risk of metastasis is measured in millimeters



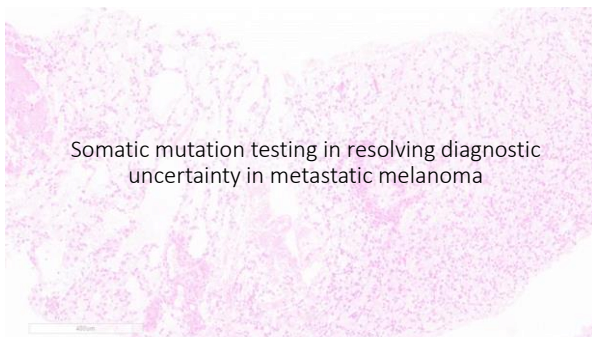
<https://www.cancer.gov/types/skin/hg/melanoma-treatment-pdq>

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Framework Genomic Classification of Cutaneous Melanoma The Cancer Genome Atlas



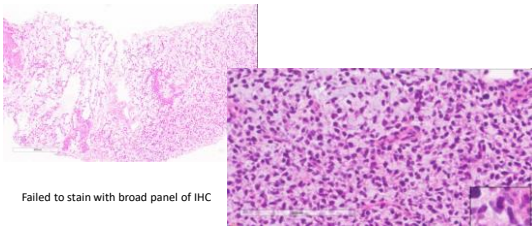
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Somatic mutation testing in resolving diagnostic
uncertainty in metastatic melanoma

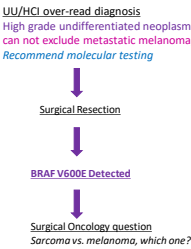
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62yo male referred to HCI, large axillary mass
outside dx = sarcoma (extraskeletal myxoid chondrosarcoma vs. myxoid liposarcoma vs. other)
now growing quickly through radiation (unlike sarcoma), referred to UU/HCI Sarcoma Center



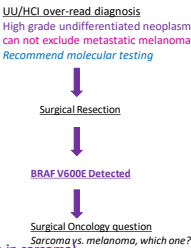
7

Electronic Health Record (outside records review)
history of melanoma, ipsilateral arm, 18 mos prior



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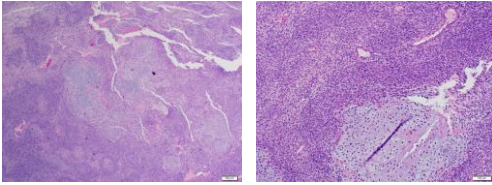
Electronic Health Record (outside records review)
history of melanoma, ipsilateral arm, 18 mos prior



0/529 sarcomas BRAF V600E (absent-exceedingly rare in sarcoma)

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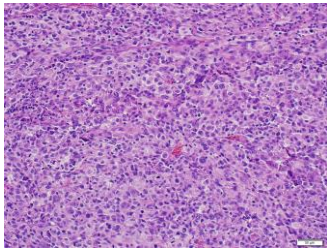
47yo male with axillary mass, and liver masses, transferred care to UU/HCI



Outside diagnosis (3 different reports) = Mesenchymal Chondrosarcoma
NQ molecular confirmation with HEY1-NCOA2 testing

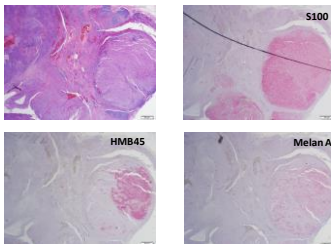
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Multiple nodules with distinct epithelioid morphology + pleomorphism



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Immunostains → Melanoma



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NGS testing confirms diagnosis of metastatic melanoma

NRAS c.34G>C, p.Gly12Arg (p.G12R)

- Interpretation: This *NRAS* (p.Gly12Arg) mutation activates the MAPK pathway (Rajalingam et al., 2007), and it has been reported in melanoma patients (COSMIC database, accessed December 8, 2015). Patients with *NRAS*-mutant melanoma may benefit from systemic immunotherapy (Johnson et al., 2015) as well as treatment with MEK inhibitors (Ascierto et al., 2013; Grimaldi et al., 2014; Thumar et al., 2014).

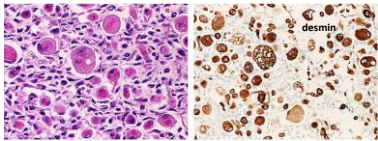
This patient received combo nivolumab/ipilimumab immunotherapy and the liver metastases regressed!

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Resolving diagnostic uncertainty in melanoma

melanomas frequently dedifferentiate when metastatic and/or can display a variety of misleading mesenchymal features

- Spindled, pleomorphic, small round/primitive blue cell, rhabdoid
- Myxoid, osteocartilagenous, lipoblastic metaplasia



Am J Surg Pathol • Volume 45, Number 2, February 2021

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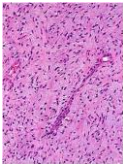
Dedifferentiated and Undifferentiated Melanomas
Report of 35 New Cases With Literature Review
and Proposal of Diagnostic Criteria

Abbas Agaimy, MD,* Robert Stoehl, PhD,* Annkathrin Hornung, MD,† Judith Popp, MD,‡
Michael Erdmann, MD,§ Lucie Heitzlerling, MD,¶ and Arndt Hartmann, MD*

Am J Surg Pathol • Volume 45, Number 2, February 2021

- n=35 unpublished cases, n=50 previously published cases, n=85 total
- negative for S100, SOX10, Melan-A, HMB45, pan-melanoma IHC
- Initial diagnoses (known in 66 cases)
 - undifferentiated/unclassified pleomorphic sarcoma (n=30)
 - undifferentiated epithelioid malignancy (n=7)
 - Pleomorphic rhabdomyosarcoma (n=5)
 - other specific sarcoma types (n=6)
 - poorly differentiated carcinoma (n=2)
 - collision tumor (n=2)
 - atypical fibroxanthoma (n=2)
 - reactive osteochondromatous lesion (n=1)
- 16.6% diagnosis of melanoma was considered

- Axilla, inguinal or other nodal basin, variety of visceral organs and body cavities, soft tissue, bone



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Abbas Agaimy, MD,* Robert Stoeck, PhD,* Annakathrin Hornung, MD,† Judith Popp, MD,†
Michael Erdmann, MD,† Lucie Heinzerling, MD,†‡ and Arndt Hartmann, MD*
Am J Surg Pathol • Volume 45, Number 2, February 2021

Melanoma compatible somatic mutation detected in 73% of cases

| Tumor Category | Male/Female Ratio (%) | Age, Median (Range) | BRF Mutations, n (%) | NRAS Mutations, n (%) | NRAS +V600E/AB BRF Mutations, n (%) | NF1 Mutations/Total Mutations, n (%) |
|---|-----------------------|---------------------|----------------------|-----------------------|-------------------------------------|--------------------------------------|
| Dedifferentiated primary melanoma | 1.8:1 | 66 (47-74) | 214 (14) | 214 (14) | 92 (6) | 327* |
| Undifferentiated metastatic melanoma with known primary | 1.5:1 | 67 (34-88) | 2048 (41) | 1548 (31) | 120 (3); V600K | 0/0* |
| Undifferentiated metastatic melanoma of unknown primary | 6.5:1 | 59 (35-86) | 313 (20) | 915 (61) | 23 (6); both V600K | 1/1 |
| DM | 3:1 | 64 (24-88) | 2577 (32.5) | 2677 (33.8) | 325 (2.1) | 3/3 |

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Authors proposed criteria for the diagnosis undifferentiated metastatic melanoma

Am J Surg Pathol • Volume 45, Number 2, February 2021

[illegible]

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helpful clues to aid in the diagnosis
undifferentiated metastatic melanoma

Am J Surg Pathol • Volume 45, Number 2, February 2021

[illegible]

- History of melanoma
- *Remote* h/o melanoma
 - Higher risk stage?
- Axilla, groin or other LN basin!!!
- Obviously in a LN!!!
- Melanoma-compatible mutation detected

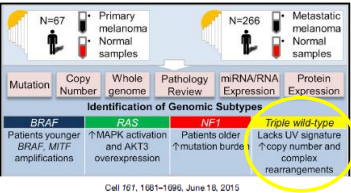
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Resolving diagnostic uncertainty in melanoma

- Detection of **BRAF** and **NRAS** mutations (>70% cutaneous melanoma) can help distinguish undifferentiated melanomas, or melanomas mimicking mesenchymal neoplasms, from soft tissue, bone or visceral sarcomas. **Exceptionally rare in sarcoma.**
 - More challenging to distinguish undifferentiated carcinoma from melanoma by mutation
- **KIT** mutations would not be surprising in metastatic melanoma from older patient with chronic sun damage – and/or could suggest acral, mucosal origin (assuming ruled out GIST)
- **NF1** mutations **do** occur in both melanoma and sarcoma (especially MPNST) – more limited diagnostic utility

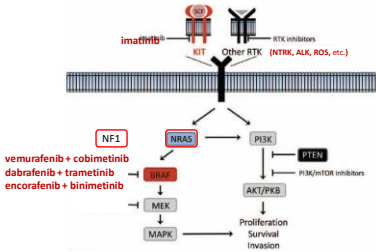
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wild type result does Not exclude melanoma



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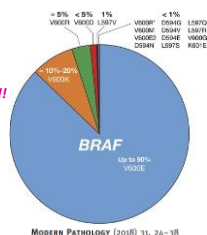
Testing for Actionable Mutations



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Actionable/potentially actionable mutations are common:
somatic mutation testing is standard of care

- **BRAF^{V600E/K} (~50%)**
 - **RAF and MEK inhibitors**
 - Adjuvant Stage III
 - Unresectable Stage III
 - Stage IV
 - **Contraindicated in BRAF wild type melanoma!!!**

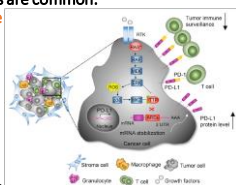


MODERN PATHOLOGY (2018) 31, 24–38

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- **NRAS^{G12R/K/L} (15-30%)**
 - Major unmet clinical need, ongoing trials
 - Correlates with poor survival
 - Minority respond to targeted MEK inhibition
 - **Immune Checkpoint Blockade = First line therapy**

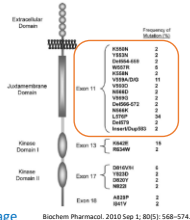


Immunity 47, December 19, 2017

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 - Minority respond to targeted MEK inhibition
 - Immune Checkpoint Blockade = First line therapy
- **KIT exon 11 (10-15% of acral, mucosal melanoma)**
 - Also enriched in melanoma with chronic sun damage
 - Targeted therapy responses are limited and not durable
- **NTRK, ALK, ROS fusions (<1%)**



Biochem Pharmacol. 2010 Sep 1; 80(5):568–574.

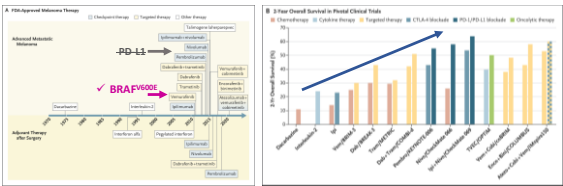
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NCCN Clinical Practice Guidelines (version 2.2022)
Indications for Somatic Mutation Testing

- **Stage III**
 - Eligibility for RAF + MEK inhibitors as adjuvant therapy (*BRAF*^{V600}-mutant)
 - Ongoing trials for neoadjuvant RAF + MEK inhibition (*BRAF*^{V600}-mutant)
 - **Stage IV** – newly diagnosed and relapsed, eligibility for targeted tx
(Retesting after progression on targeted therapy is not recommended)
- Broad panel testing (such as NGS) is recommended if feasible or when initial single gene testing for *BRAF* is negative/not detected.

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Despite major advances in the treatment of advanced-stage melanoma:
NO new standard-of-care biomarkers since 2011



Brendan D. Curti, M.D., and Mark B. Faries, M.D.
N ENGL J MED 384:23 NEJM.ORG JUNE 10, 2021

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Unlike NSCLC and other carcinomas,
PD-L1 testing is NOT required in melanoma

- Tumor PD-L1 staining can identify patients more likely to respond
 - *but* patients with PD-L1 negative tumors may still respond and benefit from anti-PD-1 immunotherapy.
- Stage IIB,IIC, III, IV melanoma are eligible for anti-PD-1 therapy

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Important Clinical Question:
Most effective method for combination treatment?

Immune Checkpoint Blockade (ICB) + targeted therapy

DREAM-seq, NCT02224781

Two-year outcome results reported at the ASCO Annual Meeting, June 2021

SECOMBIT, NCT02631447

ImmunoCobiVem; NCT02902029

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ASCO Plenary Series

DREAMseq (Doublet, Randomized Evaluation in Advanced Melanoma Sequencing) a Phase III Trial: ECOG-ACRIN EA6134

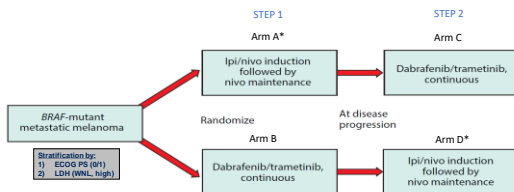
Michael B. Atkins¹, Sandra Lee², Bartosz Chmielowski³, Antoni Ribas³, Ahmad A. Tarhini⁴, Thach-Giao Truong⁵, Diwakar Davar⁶, Mark O'Rourke⁷, Brendan D. Curti⁸, Joanna M. Brell⁹, Kari L. Kendra¹⁰, Alexandra P. Ikeguchi¹¹, Jedd D. Wolchok¹², John M. Kirkwood⁶

¹Georgetown Lombardi Comprehensive Cancer Center, Washington DC; ²Dana-Farber Cancer Institute, Boston MA; ³Umsion Comprehensive Cancer Center, University of California Los Angeles, Los Angeles CA; ⁴Lee Moffitt Cancer Center and Research Institute, Tampa FL; ⁵Seisler Permanente Northern California, Vallejo CA; ⁶Pittsburgh Cancer Institute, Pittsburgh PA; ⁷Greenville Health System Cancer Institute, Greenville SC; ⁸Providence Cancer Institute, Portland OR; ⁹Metrolife Medical Center, Cleveland OH; ¹⁰Ohio State University Comprehensive Cancer Center, Columbus OH; ¹¹University of Oklahoma Medical Center, Oklahoma City OK; ¹²Memorial Sloan Kettering Cancer Center, New York NY

Slide Courtesy Michael B. Atkins, MD

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DREAMseq Trial Treatment Schema

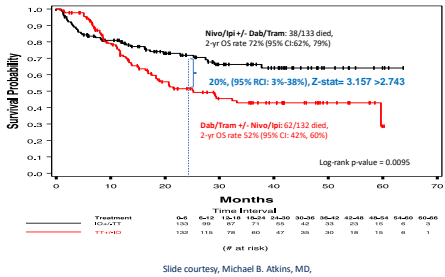


*Nivo/Ipi induction = 12 wks; nivo maintenance = 72 wks

Slide courtesy Michael B. Atkins, MD

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Improved Overall Survival (OS) leading with Nivo/Ipi



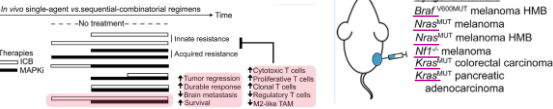
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Recent preclinical studies suggest a promising new
combo treatment paradigm for multiple cancer types

Anti-PD-1/L1 lead-in before MAPK
inhibitor combination maximizes
antitumor immunity and efficacy

Yajie Wang,^{1,2} Siue Li,^{1,2} Zhenzao Yang,^{1,2} Alan P. Algaz,^{1,3} Shirley H. Lonell,¹ Yan Wang,¹ Megan Othman,¹
Aiyang Hong,¹ Xinyan Wang,¹ Chris E. Randsberg,¹ Allison M. Jones,¹ Marcus W. Boisenberg,¹ Stephanie D. Dymov,¹
Alan J. Tackert,¹ Henry Lopez,¹ Clayton Yates,^{1,2} David B. Solit,¹ Antoni Ribas,^{1,2,3,4,5,6,7,8} Marco Piva,^{1,2,3,4,5,6,7,8}
Galen Montesano,^{1,2,3,4,5,6,7,8} and Roger S. Li,^{1,2,3,4,5,6,7,8}

Cancer Cell 39, 1375-1387, October 11, 2021



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Chasing with Biomarkers,
charting unknown waters

Which patients are likely to
receive benefit from ICB?

Which patients are *not* likely to
receive benefit from ICB?



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Management of Immune-Related Adverse Events in Patients Treated With Immune Checkpoint Inhibitor Therapy: ASCO Guideline Update

Rosen J, Schreiber MD, Lawrence BL, et al. *J Clin Oncol* 39:4073-4126. © 2021

- Rash or Inflammatory Dermatitis
- Bullous Dermatoses
- SCAR (Stevens-Johnson syndrome, toxic epidermal necrolysis, acute generalized exanthematous pustulosis) and drug reaction with eosinophilia and systemic symptoms or drug-induced hypersensitivity syndrome
- Hemolytic Anemia, aplastic anemia
- HUS
- Acquired TTP
- Lymphopenia
- ITP
- Acquired hemophilia A

- Colitis
- Hepatitis
- Pneumonitis
- Endocrinopathies (adrenal, thyroid, pituitary, diabetes)
- Autoimmune arthritis
- Myositis, polymyositis-like syndrome
- Nephritis or acute kidney injury
- Myocarditis, Pericarditis, Arrhythmias, Impaired Ventricular Function With Heart Failure, and Vasculitis
- Venous Thromboembolism
- Uveitis or iritis, episcleritis
- Myasthenia Gravis
- Guillain-Barre syndrome
- Peripheral Neuropathy
- Autonomic neuropathy
- Aseptic meningitis
- encephalitis
- Demyelinating Diseases, Including Multiple Sclerosis, Transverse Myelitis, ADEM, ON, and NMO
- Infusion reaction

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Mounting Evidence within Tumors: immunogenicity and inflammation

- immune cell infiltration
 - activated T cells vs. dysfunctional T cells
 - immunosuppressive regulatory T cells and M2-like tumor associated macrophages
- tumor immunogenicity: tumor mutation burden (TMB), neoantigen load, neoantigen heterogeneity
- expression of genes involved in antigen presentation
- specific gene mutations associated with resistance
- adaptive immune resistance, PD-L1 and LAG-3 expression
- inflammatory gene expression (particularly the IFN γ pathway)

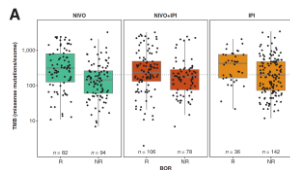
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TMB and Inflammatory Gene Expression Associated with Clinical Outcomes following Immunotherapy in Advanced Melanoma

Retrospective study

CheckMate066 (NCT01721772)
CheckMate067 (NCT01844505)

- whole exome sequencing
 - germline
 - Pre-treatment tumor
- somatic missense mutations
- calculated median for each trial cohort (mutations/exome)
- TMB^{HIGH} > median
- TMB^{LOW} < median



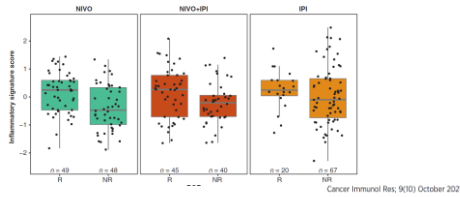
High variance TMB among both Responders vs. Non-responders

Cancer Immunol Res; 9(10) October 2021

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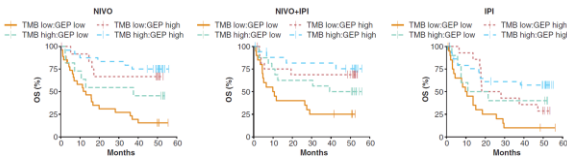
High Variance in Tumor Inflammation Score (TIS) among both responders and nonresponders

- RNA sequencing (RNA-seq), FFPE tumors, pretreatment
- CD274 (PD-L1), CD8A, LAG3, STAT1



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Overall Survival is Stratified by TMB and TIS



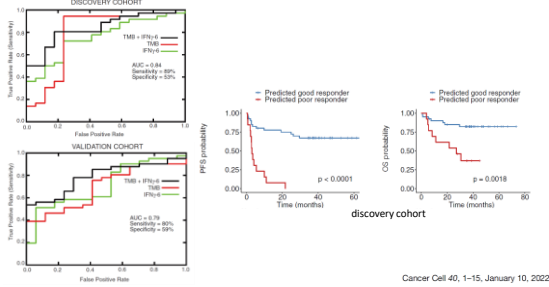
Cancer Immunol Res 2021;9:1020-33

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Multiomic profiling of checkpoint inhibitor-treated melanoma: Identifying predictors of response and resistance, and markers of biological discordance

Felicity Newell,^{1,2,3} Ines Pires da Silva,^{1,2,3,4,5,6,7,8,9,10,11,12,13,14,15,16,17,18,19,20,21,22,23,24,25,26,27,28,29,30,31,32,33,34,35,36,37,38,39,40,41,42,43,44,45,46,47,48,49,50,51,52,53,54,55,56,57,58,59,60,61,62,63,64,65,66,67,68,69,70,71,72,73,74,75,76,77,78,79,80,81,82,83,84,85,86,87,88,89,90,91,92,93,94,95,96,97,98,99,100,101,102,103,104,105,106,107,108,109,110,111,112,113,114,115,116,117,118,119,120,121,122,123,124,125,126,127,128,129,130,131,132,133,134,135,136,137,138,139,140,141,142,143,144,145,146,147,148,149,150,151,152,153,154,155,156,157,158,159,160,161,162,163,164,165,166,167,168,169,170,171,172,173,174,175,176,177,178,179,180,181,182,183,184,185,186,187,188,189,190,191,192,193,194,195,196,197,198,199,200,201,202,203,204,205,206,207,208,209,210,211,212,213,214,215,216,217,218,219,220,221,222,223,224,225,226,227,228,229,230,231,232,233,234,235,236,237,238,239,240,241,242,243,244,245,246,247,248,249,250,251,252,253,254,255,256,257,258,259,260,261,262,263,264,265,266,267,268,269,270,271,272,273,274,275,276,277,278,279,280,281,282,283,284,285,286,287,288,289,290,291,292,293,294,295,296,297,298,299,300,301,302,303,304,305,306,307,308,309,310,311,312,313,314,315,316,317,318,319,320,321,322,323,324,325,326,327,328,329,330,331,332,333,334,335,336,337,338,339,340,341,342,343,344,345,346,347,348,349,350,351,352,353,354,355,356,357,358,359,360,361,362,363,364,365,366,367,368,369,370,371,372,373,374,375,376,377,378,379,380,381,382,383,384,385,386,387,388,389,390,391,392,393,394,395,396,397,398,399,400,401,402,403,404,405,406,407,408,409,410,411,412,413,414,415,416,417,418,419,420,421,422,423,424,425,426,427,428,429,430,431,432,433,434,435,436,437,438,439,440,441,442,443,444,445,446,447,448,449,450,451,452,453,454,455,456,457,458,459,460,461,462,463,464,465,466,467,468,469,470,471,472,473,474,475,476,477,478,479,480,481,482,483,484,485,486,487,488,489,490,491,492,493,494,495,496,497,498,499,500,501,502,503,504,505,506,507,508,509,510,511,512,513,514,515,516,517,518,519,520,521,522,523,524,525,526,527,528,529,530,531,532,533,534,535,536,537,538,539,540,541,542,543,544,545,546,547,548,549,550,551,552,553,554,555,556,557,558,559,560,561,562,563,564,565,566,567,568,569,570,571,572,573,574,575,576,577,578,579,580,581,582,583,584,585,586,587,588,589,590,591,592,593,594,595,596,597,598,599,600,601,602,603,604,605,606,607,608,609,610,611,612,613,614,615,616,617,618,619,620,621,622,623,624,625,626,627,628,629,630,631,632,633,634,635,636,637,638,639,640,641,642,643,644,645,646,647,648,649,650,651,652,653,654,655,656,657,658,659,660,661,662,663,664,665,666,667,668,669,670,671,672,673,674,675,676,677,678,679,680,681,682,683,684,685,686,687,688,689,690,691,692,693,694,695,696,697,698,699,700,701,702,703,704,705,706,707,708,709,710,711,712,713,714,715,716,717,718,719,720,721,722,723,724,725,726,727,728,729,730,731,732,733,734,735,736,737,738,739,740,741,742,743,744,745,746,747,748,749,750,751,752,753,754,755,756,757,758,759,760,761,762,763,764,765,766,767,768,769,770,771,772,773,774,775,776,777,778,779,780,781,782,783,784,785,786,787,788,789,790,791,792,793,794,795,796,797,798,799,800,801,802,803,804,805,806,807,808,809,810,811,812,813,814,815,816,817,818,819,820,821,822,823,824,825,826,827,828,829,830,831,832,833,834,835,836,837,838,839,840,841,842,843,844,845,846,847,848,849,850,851,852,853,854,855,856,857,858,859,860,861,862,863,864,865,866,867,868,869,870,871,872,873,874,875,876,877,878,879,880,881,882,883,884,885,886,887,888,889,890,891,892,893,894,895,896,897,898,899,900,901,902,903,904,905,906,907,908,909,910,911,912,913,914,915,916,917,918,919,920,921,922,923,924,925,926,927,928,929,930,931,932,933,934,935,936,937,938,939,940,941,942,943,944,945,946,947,948,949,950,951,952,953,954,955,956,957,958,959,960,961,962,963,964,965,966,967,968,969,970,971,972,973,974,975,976,977,978,979,980,981,982,983,984,985,986,987,988,989,990,991,992,993,994,995,996,997,998,999,1000,1001,1002,1003,1004,1005,1006,1007,1008,1009,1010,1011,1012,1013,1014,1015,1016,1017,1018,1019,1020,1021,1022,1023,1024,1025,1026,1027,1028,1029,1030,1031,1032,1033,1034,1035,1036,1037,1038,1039,1040,1041,1042,1043,1044,1045,1046,1047,1048,1049,1050,1051,1052,1053,1054,1055,1056,1057,1058,1059,1060,1061,1062,1063,1064,1065,1066,1067,1068,1069,1070,1071,1072,1073,1074,1075,1076,1077,1078,1079,1080,1081,1082,1083,1084,1085,1086,1087,1088,1089,1090,1091,1092,1093,1094,1095,1096,1097,1098,1099,1100,1101,1102,1103,1104,1105,1106,1107,1108,1109,1110,1111,1112,1113,1114,1115,1116,1117,1118,1119,1120,1121,1122,1123,1124,1125,1126,1127,1128,1129,1130,1131,1132,1133,1134,1135,1136,1137,1138,1139,1140,1141,1142,1143,1144,1145,1146,1147,1148,1149,1150,1151,1152,1153,1154,1155,1156,1157,1158,1159,1160,1161,1162,1163,1164,1165,1166,1167,1168,1169,1170,1171,1172,1173,1174,1175,1176,1177,1178,1179,1180,1181,1182,1183,1184,1185,1186,1187,1188,1189,1190,1191,1192,1193,1194,1195,1196,1197,1198,1199,1200,1201,1202,1203,1204,1205,1206,1207,1208,1209,1210,1211,1212,1213,1214,1215,1216,1217,1218,1219,1220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Performance of combined TMB and IFN γ expression signature



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Conclusions

- TMB, neoantigen load, IFN γ expression signature, PD-L1 expression, and presence of CD8+ T cells in the tumor microenvironment are associated with *response* to ICB
- TMB and IFN γ expression signature are *independent* predictive factors
- potential predictive value of combined TMB and inflammatory gene signatures needs to be validated in *prospective* studies using predefined cutoffs

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ImmunoMATCH:

next generation NCI precision medicine trials
prospective molecular profiling and biomarker stratification

S2101 BiCaZO: A Study Combining Two Immunotherapies (Cabozantinib and Nivolumab) to Treat Patients With Advanced Melanoma or HNSCC, an ImmunoMATCH Pilot Study

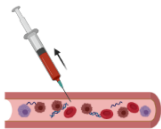
Hypothesis

- TMB and TIS will be feasible for upfront patient stratification
- Combination of Anti-PD1 and VEGFRI are effective and the response rate will be different among tumors with different TMB and TIS

Objectives

- feasibility of 14 day TAT for biomarkers
- Obtain preliminary evidence of clinical activity in pre-defined molecular subgroups (ORR, PFS, OS)

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Liquid Biopsy circulating tumor DNA (ctDNA)

1. Monitoring and predicting treatment efficacy in Stage IV patients
Lancet Oncol 2021; 22: 370-80

2. Predicting relapse and survival in Stage III patients
Annals of Oncology 30: 804-814, 2019

3. Predicting relapse and survival in Stage II/III patients
Annals of Oncology 29: 490-496, 2018

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Circulating tumour DNA in patients with advanced melanoma treated with dabrafenib or dabrafenib plus trametinib: a clinical validation study

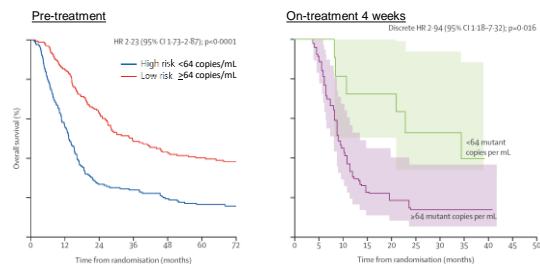
Lancet Oncol 2021; 22: 370-80

Mahrubh M Syeds, Jennifer M Wiggins, Broderick C Corless, Georgina V Long, Keith T Flaherty, Dirk Schadendorf, Paul D Nathan, Caroline Robert, Antoni Ribes, Michael A Davies, Jean Jacques Grob, Eduard Gosal, Matthew Squires, Mahtab Markes, James Garrett, Jan C Brase, David Polsky

- Retrospective study, unresectable or metastatic *BRAF*-mutant melanoma
 - Advanced stage → expect tumor shedding and detectable ctDNA pre-treatment
- phase 3 COMBI-d and phase 2 COMBI-MB trials
 - dabrafenib ± trametinib
- Measured *BRAF*^{V600E/K} ctDNA by droplet digital PCR, n=345 patients
 - Detected in 90% of patients
- Serially collected blood - before treatment and on treatment week 4
- Biomarker study funded by Novartis, testing performed by NYU

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ctDNA testing stratified high vs. low risk for progression and prognosticates overall survival in both baseline and very early on-treatment



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Unmet clinical need: Improving risk stratification for Stage III melanoma

- Stage III patients are eligible for adjuvant ICB therapy
 - Costly
 - Potential for immune related adverse events (irAEs)
- Clinical goal for biomarker development and validations
 - Ideally - avoid unnecessary treatment in patients who are cured by surgery alone
 - 40%–90% of patients with resected stage III disease treated with curative intent will relapse within 5 years
 - identify those at highest risk of relapse, where the benefits of systemic therapy may outweigh the risk of irAEs

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Prediction and monitoring of relapse in stage III melanoma using circulating tumor DNA

Annals of Oncology 30: 804–814, 2019

L. Tan^{1,2*}, S. Sandhu^{1,2*}, R. J. Lee^{3,4*}, J. Li^{1,2}, J. Callahan¹, S. Floun¹, N. Dhomen¹, P. Middlehurst¹, A. Wallace⁵, J. Raleigh⁶, A. Hatzimichailis¹, M. A. Henderson^{1,2}, M. Shackleton⁷, A. Haydon², V. Marai^{8,9}, D. E. Gyorki^{1,2}, D. Oudis^{8,9}, M. A. Dawson^{1,2,9}, R. J. Hicks^{1,2}, P. Lorigan^{8,9}, G. A. McArthur^{1,2}, R. Marais^{8,9}, S. Q. Wong¹⁰ & S.-J. Dawson^{1,2,9*}

- Tumor: mutations identified in 99/133 (74%) patients
 - *BRAF*, *NRAS*, *TERT* promoter
 - Blood: 315 prospectively collected plasma specimens
 - Pre-Op baseline
 - Post-Op
- ctDNA Assay = droplet digital PCR (ddPCR)
- ctDNA was detected in 37 of 99 (37%) individuals
- 53 of 99 (54%) had relapsed with median follow up of 18 months (range: 2–58 months) (none had received adjuvant systemic therapy)

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ctDNA detection increases with increasing T stage (Breslow/primary tumor thickness, ulceration, lymph node stage)

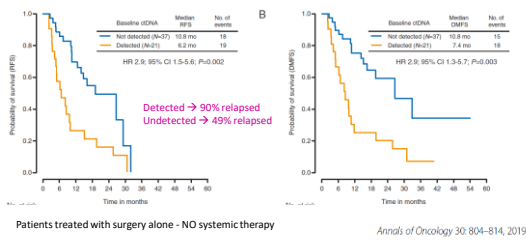
Table 1. Clinicopathological characteristics of patients in the MIV cohort according to baseline and postoperative ctDNA status

| Characteristics | ctDNA baseline | | | P | ctDNA postoperative | | | P | Total N (%) |
|--------------------------------|---------------------|-------------------|----------------|--------------|---------------------|-------------------|----------------|-------------|----------------|
| | Undetected N (%) | Detected N (%) | Total N (%) | | Undetected N (%) | Detected N (%) | Total N (%) | | |
| Age ^a | | | | | | | | | |
| <70 years | 35 (67) | 17 (33) | 52 (76) | 0.55 | 35 (73) | 13 (27) | 48 (74) | 0.74 | 52 (76) |
| ≥70 years | 9 (56) | 7 (44) | 16 (24) | | 9 (57) | 3 (25) | 12 (24) | | 16 (24) |
| Sex ^a | | | | | | | | | |
| Male | 29 (63) | 19 (40) | 48 (71) | 0.28 | 48 (71) | 19 (29) | 67 (70) | 0.77 | 45 (66) |
| Female | 15 (75) | 5 (25) | 20 (29) | | 17 (74) | 6 (26) | 23 (34) | | 23 (34) |
| AJCC substage ^a | | | | | | | | | |
| IA | 7 (100) | 0 (0) | 7 (10) | 0.02 | 7 (10) | 0 (0) | 7 (10) | 0.06 | 8 (11) |
| IB | 16 (78) | 4 (18) | 20 (24) | | 23 (84) | 4 (16) | 27 (33) | | 24 (32) |
| II | 21 (53) | 17 (40) | 38 (46) | | 38 (56) | 23 (32) | 61 (69) | | 37 (48) |
| Breslow thickness ^a | | | | | | | | | |
| ≤2.0 mm | 23 (78) | 6 (21) | 29 (44) | 0.046 | 29 (44) | 7 (21) | 36 (55) | 0.09 | 31 (53) |
| >2.0–4.0 mm | 9 (68) | 4 (31) | 13 (21) | | 7 (88) | 1 (12) | 8 (14) | | 8 (14) |
| >4.0 mm | 11 (52) | 10 (48) | 21 (33) | | 11 (52) | 8 (42) | 19 (33) | | 19 (33) |
| Ulceration ^a | | | | | | | | | |
| Absent | 28 (74) | 10 (26) | 38 (46) | 0.39 | 38 (62) | 4 (11) | 42 (50) | 0.02 | 35 (62) |
| Present | 14 (61) | 12 (55) | 26 (40) | | 23 (88) | 8 (31) | 31 (58) | | 21 (38) |

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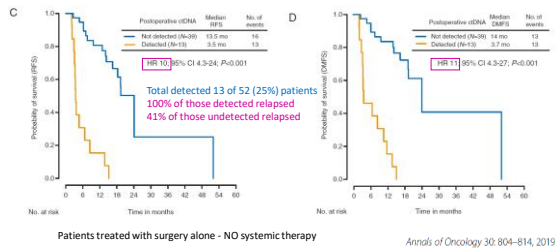
48

**PRE-operative ctDNA detection Stage III melanoma:
reduced relapse free and distant metastasis free survival**



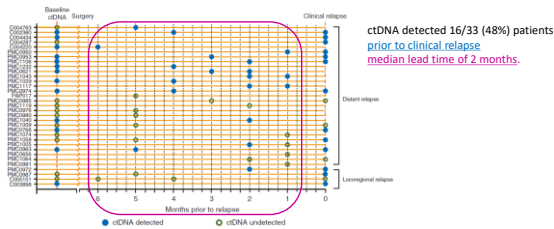
49

**POST-operative ctDNA detection Stage III melanoma
reduced relapse free and distant metastasis free survival**



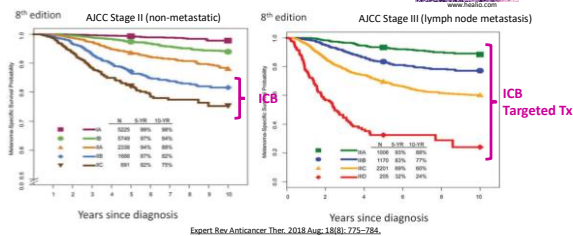
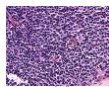
50

**Serial postoperative liquid biopsies
ctDNA was detected prior to relapse in half the patients**



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High Risk Stage II-III Melanoma: improve risk stratification Stage IIB-C, IIIA-B?



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Can ctDNA distinguish relapsers from nonrelapsers within high risk, resected, Stage II/III melanoma patients?

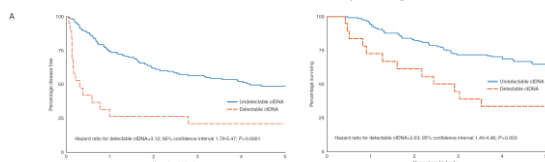
- Retrospective study
- Stage IIB, IIC, III melanoma
- Single plasma collection within 12 weeks after surgery (trial setting)
- ddPCR BRAF^{V600E} and NRAS^{Q61K/L}
- detectable ≥ 1 copy of mutant DNA/2mL plasma

| Characteristic | Total N (%) | Undetectable ctDNA N (%) | Detectable ctDNA N (%) |
|-----------------|----------------|--------------------------------|------------------------------|
| Disease stage | | | |
| I/IIA | 36 (23) | 33 (23) | 3 (16) |
| IIB | 29 (18) | 27 (19) | 2 (11) |
| IIC | 59 (37) | 51 (36) | 8 (42) |
| III | 37 (23) | 31 (22) | 6 (32) |
| Mutation status | | | |
| BRAF V600E | 132 (82) | 117 (82) | 15 (79) |
| NRAS Q61K/L | 29 (18) | 25 (18) | 4 (21) |
| Total | 161 (100) | 142 (88) | 19 (12) |

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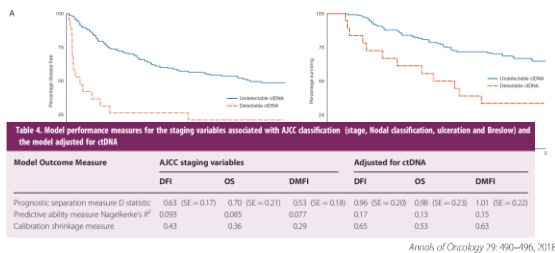
Detection of ctDNA: reduced disease-free and overall survival (5yr) Stage II/III melanoma



Annals of Oncology 29: 490-496, 2018

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Detection of ctDNA improves prognostication of Stage II/III melanoma



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Summary

- Somatic mutation testing, particularly for *BRAF*, remains essential for, and will continue to guide, SOC therapy for cutaneous melanoma
- Panel testing is recommended, if feasible, to cover actionable mutations
 - BRAF > NRAS > KIT > NF1 > NTRK/ROS/ALK*
- Molecular testing may help resolve diagnostic uncertainty with metastatic melanoma
- Recent clinical trial data demonstrates improved efficacy of combo therapy
 - ICB lead followed by targeted therapy
 - Emerging data – may be relevant to other cancers
- Emerging evidence suggest genomic markers of tumor immunogenicity (TMB) and inflammation (CD8 infiltration, IFN γ gene expression signatures) identifies patients who are most likely to benefit from ICB, prospective clinical trials pending
- Liquid biopsy/ctDNA testing may improve disease monitoring and risk stratification, prospective clinical trials needed

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Sapna Patel, MD, Chair SWOG Melanoma Committee, Medical Oncology, MDACC

Kenneth Grossmann, MD PhD, Senior Principle Scientist, Clinical Director, Merck



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