



## Consensus Statement Evidentiary Tables :

### Assessing the Evidence for and Utility of Gene Expression Profiling of Primary Cutaneous Melanoma\*

Levels of evidence adopted from *American Association of Clinical Endocrinologists and American College of Endocrinology Protocol for Standardized Production of Clinical Practice Guidelines, Algorithms, and Checklists-2017 Update*.

*Mechanick JL, Pessah-Pillack R, Camacho P, Correa R, Figaro MK, Garber JR, et al. American Association of Clinical Endocrinologists and American College of Endocrinology Protocol for Standardized Production of Clinical Practice Guidelines, Algorithms, and Checklists-2017 Update. Endocr Pract. 2017; 8:1006-1021.*

## Bookmarks to Tables for Each Question

[Question 1](#)

[Question 2](#)

[Question 3 + Future Directions for Research](#)

## Question 1

In adult patients with AJCC pT1a-pT4b primary cutaneous melanoma, does GEP testing improve patient selection and decision-making for sentinel lymph node biopsy as compared to the use of conventional clinical and pathologic factors alone?

Summary of Evidence Table Question 1, 2						
Level of Evidence*^  Choose one:	Study Citation Author(s), Year, Title, Journal, Volume, Page #s	Funding Source	Study Methodology	Objective & Methods	Results	Conclusions
<div> <div> <input type="checkbox"/> EL 1; RCT  <input type="checkbox"/> EL 1; MRCT </div> <div> <input type="checkbox"/> EL 2; MNRCT  <input type="checkbox"/> EL 2; NMA  <input type="checkbox"/> EL 2; NRCT  <input type="checkbox"/> EL 2; PCS  <input type="checkbox"/> EL 2; RCCS  <input type="checkbox"/> EL 2; NCCS  <input type="checkbox"/> EL 2; CSS  <input type="checkbox"/> EL 2; ES  <input type="checkbox"/> EL 2; OLES  <input type="checkbox"/> EL 2; PHAS </div> <div> <input type="checkbox"/> EL 3; DS  <input type="checkbox"/> EL 3; ECON  <input checked="" type="checkbox"/> EL 3; CCS  <input type="checkbox"/> EL 3; SCR  <input type="checkbox"/> EL 3; PRECLIN  <input type="checkbox"/> EL 3; BR </div> <div> <input type="checkbox"/> EL 4; NE  <input type="checkbox"/> EL 4; O </div> </div>	<p>Amaral T, Sinnberg T, Chatziioannou E, et al. Identification of Stage I/II Melanoma Patients at High Risk for Recurrence Using a Model Combining Clinicopathologic Factors with Gene Expression Profiling (CP_GEP). <i>Eur J Cancer</i>. 2023;182:155-162.</p> <p><i>Country:</i> Germany, Netherlands</p>	<p><i>Source:</i> The study was partially funded by SkylineDx.</p> <p>4 authors: SkylineDx employees</p> <p>2 authors: institutional funding from SkylineDx in relation to the submitted work.</p> <p>2 authors: Institutional financial support from Neracare</p> <p>5 authors: stock/ownership interest in SkylineDx</p> <p><u>1 author: leadership SkylineDx</u></p> <p><u>1 author: consulting or advisory role</u></p>	<p><i>Methodology:</i> Blinded retrospective single-center study of stage I/II patients Single center retrospective review of a prospective database.</p>	<p><i>Stated Objective:</i> To validate CP-GEP for the identification of high-risk for disease recurrence stage I/II patients.</p> <p><input type="checkbox"/> Prospective <input checked="" type="checkbox"/> Retrospective</p> <p><i>Study Population and Setting:</i> Stage I/II, age &gt;18, 2000-2017, with negative SLNB</p> <p><i>N:</i> 543 patients with SLNB+ 83 control without SLNB, of which 80 were analyzed</p> <p><i>Intervention:</i> CP-GEP with clinicopathologic features age and Breslow thickness with the expression of (ITGB3, PLAT, SERPINE2, GDF15, TGFB1, LOXL4, CXCL8 and MLANA) corrected by (RLP0 and ACTB)</p> <p>CP-GEP combined protocol to predict recurrence</p>	<p><i>Results:</i> W/ SLNB CP-GEP stratified 424 patients (78% cohort) into RFS rates of 77.8% for high risk (195 patients) and 93% for low risk (229 patients). HR 3.53 ( p, 0.001).</p> <p>RFS stage I 90.7%, stage II 66.1% DMFS stage I 96.0%, stage II 82.2% OS stage I 95.6%, stage II 79.0%</p> <p>For stage I/II: CP-GEP classified 311 as high risk with HR 4.73 with p-value &lt;0.001</p> <p>Subgroup I/IIA: CP-GEP stratified low v high-risk with a HR of 3.53 (p &lt;0.001).</p> <p>RFS I/IIA for high-risk (195 patients) was 77.8% and 93.0% for low-risk.</p> <p>W/O SLNB</p>	<p><i>Describe conclusions relative to question:</i></p> <p>CP-GEP predicts high risk patients in stage I/II and could be used to replace SLNB and to guide adjuvant treatments.</p> <p>CP-GEP could replace SLNB,</p> <p>CP-GEP enhances traditional histologic classification of patients.</p> <p><i>Critiques of Methodology:</i> Group stages I/II and don't separate to be able to discern clinicopathologic characteristics vs CP-GEP. The stage I/II vs I/IIA</p>

				<p>Outcome Measures: Kaplan-Meier curves to determine prognostic value. Primary endpoint was RFS also DMFS and OS.</p> <p>Recurrence after negative SLNB</p> <p><i>Follow-Up:</i> Median follow up of 83.63 months for patients with negative SLNB; 40.77 mo for patients without SLNB</p> <hr/> <p><i>Notes</i></p>	<p>Classified 11/80 high-risk which captured 6/7 recurrences.</p> <p>The five-year RFS for stage I/II patients was 79.9% (95% CI: 76.0%-83.2%). CP-GEP identified 311 patients as high risk for disease recurrence with a HR of 4.73 with a p-value &lt;0.001, capturing 83 out of 98 reported relapses. For subgroup stage I/IIA, CP-GEP was able to significantly stratify CP-GEP low-risk and high-risk patients with a HR of 3.53 (p value &lt;0.001) for five-year RFS. For stage I/IIA, the five-year RFS rates for CP-GEP high-risk patients were 77.8% (95% CI: 70.9%-83.3%) and 93.0% (95% CI: 88.5%-95.8%) for CP-GEP low-risk patients. "Compared to AJCC low-risk (stage I/IIA) patients with an RFS rate of 86.0% (95% CI: 82.0%-89.1%), CP-GEP was able to split 195 high-risk patients who had a worse five year RFS survival of 77.8%"</p> <hr/> <p>Notes: Wo SLNB high-risk curves are dramatically different, does this argue for doing a SLNB.</p>	<p>Initial curves do not separate by histologic stage. Group "Stage I/II" together.</p>
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					<p>Without SLNB5/11 classified as high-risk without recurrence.</p> <p>They group the analyses into “who would qualify for adjuvant therapy vs who wouldn’t” There is no subset analysis by the actual histologic stages.</p>	
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		<p>metastasis of melanoma</p> <p>1 author: research funding from SkylineDx</p>	<p>lower than the cutoff were classified as negative.</p> <p>Model validated by repeated cross-validation or bootstrapping.</p> <p>GEP defined from 11 genes that differentiated the patients with and without nodal metastasis detected by SLNB within 90 days of primary diagnosis: ADAM metalloproteinase domain 12 (ADAM12), interleukin 8 (CXCL8), growth differentiation factor 15 (GDF15), integrin-<math>\beta</math>3 (ITGB3), galectin 1 (LGALS1), lysyl oxidase like 4 (LOXL4), melanoma antigen recognized by T cells 1 (MLANA), tissue-type plasminogen activator (PLAT), protein kinase C-<math>\beta</math> (PRKCB), glia-derived nexin (SERPINE2), and transforming growth factor-<math>\beta</math> (TGF-<math>\beta</math>) receptor 1 (TGFR1).</p>	<p>CP-GEP model GEP profile plus standard histologic parameters, creative modeling.</p> <p>(Clinicopathologic variables with eight-gene GEP) <i>Outcome Measures:</i> SLN positivity within 90 days of diagnosis.</p> <p>SLN metastasis within 90 days of melanoma diagnosis</p> <p><i>Follow-Up:</i> Appears to be 90 day follow up only (no subsequent follow up for false negative, in-basin recurrence)</p> <hr/> <p><i>Notes:</i> Excluded high risk (T4 patients and performed SLNB in T1a patients that have "high risk features," lower than many institutional thresholds</p>	<p>T1b SLNB reduction rate 80% T2a SLNB reduction rate 48% with NNP 95%</p> <p>Compare MSKCC nomogram which performed similarly to CP model but out-performed by the CP-GEP AUC</p> <p>This is a complicated statistical analysis, but the modeling seems effective. Not sure that it proves to be statistically more powerful than clinicopathological criteria alone, in real-world use. A statistician would be helpful for this study.</p> <hr/> <p>Notes: evaluated multiple CP factors to predict SLN+ and showed age and Breslow depth most predictive</p> <p>Stratified by T stage (table 4).</p> <hr/> <p><i>Notes:</i></p>	<p>on in-nodal basin recurrences therefore the accuracy of this test longitudinally is unclear.</p> <p><i>Critiques of Methodology:</i> Sophisticated statistical analysis.</p> <p>AUC greatest for combined model CP-GEP but did not analyze if statistically significant improvement.</p> <p>It is a statistical model. The discussion implies that the only benefit of SLNB is to determine adjuvant therapy, but there are other benefits.</p>
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			<p>Logistic regression modeling was used to develop a novel model combining CP factors (i.e.: Breslow depth and patient age) and a GEP.</p> <p>CP-GEP model was based on the expression of MLANA, a melanosome marker, and seven genes</p> <p>Functionally linked to EMT and with specific roles in angiogenesis/hypoxia and coagulation: GDF15,</p> <p>CXCL8,</p> <p>LOXL4, TGFB1, ITGB3, PLAT, and SERPINE2</p> <p>Retrospective analysis of prospectively collected specimens with follow up</p>			
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**Summary of Evidence Table**  
**Question 1, 2, 3**

Level of Evidence*^  Choose one:	Study Citation Author(s), Year, Title, Journal, Volume, Page #s	Funding Source	Study Methodology	Objective & Methods	Results	Conclusions
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		advisory board				
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						the majority (68%) also had sentinel lymph node biopsy done.
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						<ul style="list-style-type: none"><li>-Who gave the lecture on clinical validity of the test before the study??- that could significantly bias results</li><li>-Authors do not recognize these limitations in the manuscript</li><li>-They gave the residents an answer of “onc” and imaging referral which would not be currently recommended for the thin Breslow depths noted; confusing</li></ul>
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## Summary of Evidence Table

### Question 1, 2, 3

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<div> <input type="checkbox"/> EL 1; RCT  <input type="checkbox"/> EL 1; MRCT    <input type="checkbox"/> EL 2; MNRCT  <input type="checkbox"/> EL 2; NMA  <input type="checkbox"/> EL 2; NRCT  <input type="checkbox"/> EL 2; PCS  <input type="checkbox"/> EL 2; RCCS  <input type="checkbox"/> EL 2; NCCS  <input type="checkbox"/> EL 2; CSS  <input type="checkbox"/> EL 2; ES  <input type="checkbox"/> EL 2; OLES  <input type="checkbox"/> EL 2; PHAS    <input type="checkbox"/> EL 3; DS  <input type="checkbox"/> EL 3; ECON  <input type="checkbox"/> EL 3; CCS  <input type="checkbox"/> EL 3; SCR  <input type="checkbox"/> EL 3; PRECLIN  <input type="checkbox"/> EL 3; BR    <input checked="" type="checkbox"/> EL 4; NE  <input type="checkbox"/> EL 4; O </div>	<div> <div>Strong</div> <div>Intermediate</div> <div>Weak</div> <div>No Evidence</div> </div> <p>Farberg AS, Marson JW, Glazer A, et al. Expert Consensus on the Use of Prognostic Gene Expression Profiling Tests for the Management of Cutaneous Melanoma: Consensus from the Skin Cancer Prevention Working Group. <i>Dermatol Ther.</i> 2022;12(4):807-823. (81)</p> <p>Country: USA</p>	<p>Source: No funding or sponsorship was received for this study or publication of this article.</p> <p>3 authors=consultants for Castle Biosciences</p> <p>1 author=advisory board member and speaker for Castle Biosciences</p>	<p>Methodology: Modified Delphi technique for consensus statements.</p> <p>Review – no new data.</p> <p>Skin Cancer Prevention Working Group are authors.</p>	<p>Stated Objective: Assess applications for GEP in melanoma management.</p> <p><input type="checkbox"/> Prospective <input type="checkbox"/> Retrospective</p> <p>Study Population and Setting:</p> <p>N:</p> <p>Intervention:</p> <p>Outcome Measures:</p> <p>Follow-Up:</p> <p>Notes:</p>	<p>Results: Consensus statements support role for GEP above and beyond AJCC and NCCN.</p> <hr/> <p>Notes: Consensus statements support role for GEP above and beyond AJCC and NCCN. The statements are non-specific.</p>	<p>Describe conclusions relative to question:</p> <p>Consensus statements support role for GEP.</p> <p>“GEP tests provide additional, reproducible information for dermatologists to consider within the larger framework of the eighth edition of the AJCC and NCCN cutaneous melanoma guidelines when counseling regarding prognosis and when considering a sentinel lymph node biopsy.”</p> <p>Critiques of Methodology: No formal accompanying systematic review/meta-analysis to support the consensus</p>



						<p>process (only 8 panelists).</p> <p>Review of existing data.</p>
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**Summary of Evidence Table**  
**Question 1, 2**

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						<p><i>2 tumors metastasized, whereas 1 (0.5%) of 201 Stage I Class 1 tumors metastasized.</i></p> <p>This implies that that the GEP is not discerning in Stage I patients since Class differentiation did not predict any pattern of disease recurrence.</p> <p>Discussion trying to explain why their findings might not reflect the ultimate utility of the GEP.</p> <p>Retrospective Enriched for subsequent testing after recurrence in some of the patients, number of patients in this subcohort nor method for identification not stated</p> <p>With median f/u 23 months 5 yr KM survival analysis seems invalid.</p> <p>Class 1 median Breslow 0.6 mm vs Class 2 = 2.2 mm</p> <p>No analysis comparing value of 31-GEP beyond AJCC stage, Breslow Depth</p>
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		<p>with resources and use of facilities at the Veterans Affairs Palo Alto Health Care System in Palo Alto CA. Contents do not represent the views of the US Dept of Veterans Affairs or the US Gov't. The funders had no role in the design and conduct of the study; collection, management, analysis and interpretation of data, preparation, review or approval of the manuscript; and decision to submit for publication.</p> <p>1 author: nonfinancial support from Castle Biosciences outside submitted work.</p> <p>1 author: grants from Castle Biosciences outside</p>		<p>Follow-Up:</p> <hr/> <p>Notes:</p>	<p>with SLNB and adjuvant therapy. The MPWG members favor conducting retrospective studies that evaluate multiple GEP testing platforms on fully annotated archived samples before embarking on costly prospective studies and recommend avoiding routine use of GEP testing to direct patient management until prospective studies support their clinical utility.</p> <hr/> <p>Notes:</p>	
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		<p>submitted work</p> <p>1 author: personal fees outside the submitted work Castle Biosciences</p> <p>1 author: personal fees from Neracare outside submitted work</p> <p>1 author: manuscripts and abstracts published using the test with company support of the assay—all publications were peer review and no personal of institutional payment or compensation was rec'd. Castle Biosciences</p> <p>1 author: served as an investigator for Castle Biosciences (no personal financial compensation)</p>				
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				<div>Follow-Up: n/a unclear</div> <div></div>		
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		<p>conduct of the study; collection, management, analysis, and interpretation of the data; preparation, review, or approval of the manuscript and decision to submit the manuscript for publication.</p> <p>2 authors: research support from Castle Biosciences outside this project</p> <p>1 author: research support from Castle both during and outside this work</p> <p>1 author: grant support from Skyline Dx outside this work</p> <p>1 author: nonfinancial support from SkylineDx outside the work</p>				<p>i-31-GEP) purported to predict SLN positivity.”</p> <p><i>Critiques of Methodology:</i></p> <p>Other than funding support, no concerns. Appropriately designed and performed consensus conference.</p>
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<div><div><div><div><div><input type="checkbox"/>EL 1; RCT</div><div><input type="checkbox"/>EL 1; MRCT</div></div><div><div><div><input type="checkbox"/>EL 2; MNRCT</div><div><input type="checkbox"/>EL 2; NMA</div><div><input type="checkbox"/>EL 2; NRCT</div><div><input type="checkbox"/>EL 2; PCS</div><div><input type="checkbox"/>EL 2; RCCS</div><div><input type="checkbox"/>EL 2; NCCS</div><div><input type="checkbox"/>EL 2; CSS</div><div><input type="checkbox"/>EL 2; ES</div><div><input type="checkbox"/>EL 2; OLES</div><div><input type="checkbox"/>EL 2; PHAS</div></div><div><div><div><input checked="" type="checkbox"/>EL 3; DS</div><div><input type="checkbox"/>EL 3; ECON</div><div><input type="checkbox"/>EL 3; CCS</div><div><input type="checkbox"/>EL 3; SCR</div><div><input type="checkbox"/>EL 3; PRECLIN</div><div><input type="checkbox"/>EL 3; BR</div></div><div><div><div><input type="checkbox"/>EL 4; NE</div><div><input type="checkbox"/>EL 4; O</div></div></div></div></div><div><div>Strong</div><div>Intermediate</div><div>Weak</div><div>No Evidence</div></div></div></div></div>	<p>Marchetti MA, Dusza SW, Bartlett EK. Utility of a Model for Predicting the Risk of Sentinel Lymph Node Metastasis in Patients With Cutaneous Melanoma. <i>JAMA Dermatol.</i> 2022;158(6):680-683 (89)</p> <p>Country: USA</p>	<p><i>Source:</i> This research was funded in part through the Memorial Sloan Kettering Cancer Center institutional National Institutes of Health/National Cancer Institute Cancer Center Support Grant P30 CA008748. The NIH had no role in the design and conduct of the study; collection, k management, analysis, and interpretation of the data, preparation, review, or approval of the manuscript; and decision to submit the manuscript for publication.</p> <p>1 author: institutional research support from SkylineDx outside the submitted work</p>	<p><i>Methodology:</i> Decision analytic study based on i31-GEP model (presented as citation #78, Whitman et al.).</p> <p>Use of mathematical model on existing dataset.</p> <p>“Raw data and classification measures of the i31-GEP-SLNB model by American Joint Committee on Cancer (AJCC) T category were extracted from Whitman et al2 for patients with T1a-HR, T1b, and T2 disease;”</p>	<p><i>Stated Objective:</i> “To determine if use of the i31-GEP-SLNB model is associated with clinical benefit when used to select patients for SLN biopsy.”</p> <p><input type="checkbox"/>Prospective <input checked="" type="checkbox"/>Retrospective</p> <p><i>Study Population and Setting:</i> See methodology.</p> <p>N: 1097</p> <p><i>Intervention:</i> Use of mathematical modeling on existing dataset.</p> <p><i>Outcome Measures:</i> “To determine if use of the i31-GEP-SLNB model is associated with clinical benefit when used to select patients for SLN biopsy.”</p> <p><i>Follow-Up:</i> Not stated</p> <hr/> <p><i>Notes:</i></p>	<p><i>Results:</i> “Compared with other SLN biopsy selection strategies, use of the i31-GEP-SLNB model had greater net benefit for patients with T1b (+0.012), T2a (+0.002), and T2b melanoma (+0.002) but not for those with high-risk T1a (−0.003) disease. The improvement in relative utility was +22%in patients with T1b, +1% in T2a, and +2%in T2b melanoma.”</p> <hr/> <p><i>Notes:</i></p>	<p><i>Describe conclusions relative to question:</i></p> <p>Appropriate</p> <p><i>Critiques of Methodology:</i> Should not be included as source GEP article</p> <p>Computational modeling</p>

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			reduction rate and corresponding 95% confidence intervals.	<p>NPV, PPV SLNB reduction rate</p> <p>Time order to report= order to revision + revision to shipment + shipment to report.</p> <p><i>Follow-Up:</i> Na</p> <hr/> <p><i>Notes:</i> N should include 5 patients that CP-GEP was invalid</p>	<p>45.8% (95% CI: 38.6–53.2), a NPV of 96.7% (95% CI: 90.6–99.3) and a PPV of 23.7% (95% CI: 16.8–31.8). The SLNB reduction rate was 39.2% (95% CI: 32.7–45.9).</p> <p>of CP-GEP low risk cases where found, the model yielded a sensitivity of 84.2% (95% CI: 60.4–96.6), a specificity of 57.3% (95% CI: 48.8–65.6), a NPV of 96.5% (95% CI: 90.0–99.2) and PPV of 20.7% (95% CI: 12.4–31.5). In this group, a SLNB reduction rate of 52.5% (95% CI: 44.5–60.5) was found.</p> <hr/> <p>Notes:</p>	<p>high low risk and whether +/- SLN.</p> <p>Only 112 T2 pt</p> <p>All T4 classified as high risk, and majority T3 so unclear benefit of GEP offer over histopathology.</p> <p>Utility Need follow up for in-nodal basin recurrence.</p>
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<input type="checkbox"/> EL 1; RCT <input type="checkbox"/> EL 1; MRCT  <input type="checkbox"/> EL 2; MNRCT <input type="checkbox"/> EL 2; NMA <input type="checkbox"/> EL 2; NRCT <input checked="" type="checkbox"/> EL 2; PCS <input type="checkbox"/> EL 2; RCCS <input type="checkbox"/> EL 2; NCCS <input type="checkbox"/> EL 2; CSS <input type="checkbox"/> EL 2; ES <input type="checkbox"/> EL 2; OLES <input type="checkbox"/> EL 2; PHAS  <input type="checkbox"/> EL 3; DS <input type="checkbox"/> EL 3; ECON <input type="checkbox"/> EL 3; CCS <input type="checkbox"/> EL 3; SCR <input type="checkbox"/> EL 3; PRECLIN <input type="checkbox"/> EL 3; BR  <input type="checkbox"/> EL 4; NE <input type="checkbox"/> EL 4; O	<p><b>Strong</b></p> <p>Thorpe RB, Covington KR, Caruso HG, et al. Development and Validation of a Nomogram Incorporating Gene Expression Profiling and Clinical Factors for Accurate Prediction of Metastasi in Patients with Cutaneous Melanoma following Mohs Micrographic Surgery. J. Am. Acad. Dermatol. 2022;86(4):846-853. (96)</p> <p><b>Intermediate</b></p> <p><b>Weak</b></p> <p><b>No Evidence</b></p> <p>Country: USA 9 Mohs private practices</p>	<p><i>Source:</i></p> <ol style="list-style-type: none"> <li>1. Statistician provided by Castle Biosciences</li> <li>2. IRB funded by Skin Cancer Trust for independent Research and Education, a 501c3 managed by Drs. Zitelli and Brodland</li> </ol> <p>4 authors: employees of Castle Biosciences</p> <p>4 authors: stockholders at Castle Biosciences</p>	<p><i>Methodology:</i></p> <p>Prospective testing with 31-GEP of patients with melanoma undergoing melanoma surgery at Mohs surgery center.</p> <p>Melanoma patients with complete clinical and pathologic information including 31-GEP enrolled from the nine centers</p> <p>Years of enrollment not stated.</p> <p>All pts had Mohs with six mm initial margins with further margins until clear by MART1 IHC.</p>	<p><i>Stated Objective:</i></p> <p>To develop a nomogram for predicting metastasis using 31-GEP and T stage.</p> <p>To define a nomogram for practical clinical use using 31-GEP.</p> <p><input checked="" type="checkbox"/> Prospective <input type="checkbox"/> Retrospective</p> <p><i>Study Population and Setting:</i></p> <p>Patients undergoing melanoma surgery at nine Mohs surgery centers.</p> <p>Nine Mohs Private Derm Practices</p> <p>N: 1124 enrolled – data presented is from 684 patients followed for min 1 yr or a “metastatic event” as those used for nomogram development (p847).</p> <p><i>Intervention:</i> Mohs + 31-GEP</p> <p><i>Outcome Measures:</i></p> <p>Nomogram development for predicting metastasis using</p> <p>Risk of metastasis using a nomogram</p> <p><i>Follow-Up:</i></p>	<p><i>Results:</i></p> <p>31-GEP test and T stage offers simplest nomogram with lowest Bayesian information score— validated in a separate cohort of 905 patients.</p> <p>Class 2B patients do worse (53/684 = 7.7% of the cohort Don't see where the nomogram considering Breslow depth and 31-GEP adds value over either alone (see Table II) except maybe T2-T3a and shifts %s in clinically unmeaningful way – wouldn't change f/u or rx recommendations.</p> <p><i>Notes:</i></p> <p>Mohs for invasive melanoma is unsupported by RCT evidence. Graphs present five year survival without numbers at risk data &amp; P value doesn't state what the comparison is (class IA,1B,2A all appear to overlap on the RFS and DMFS curves) Demographics in supplemental table mean BT 0.5 mm (0.1-13.0 mm)</p>	<p><i>Describe conclusions relative to question:</i></p> <p>31-GEP test and T stage can provide prognostic information for metastasis.</p> <p>Question biopsy as non-invasive (need to do it to get T category which they refer to as T stage).</p> <p><i>Critiques of Methodology:</i></p> <ol style="list-style-type: none"> <li>1. SLN biopsy not done for many patients and not included in the nomogram</li> <li>2. Limited to patients undergoing surgery at Mohs surgery centers which may reflect a selection bias (very few T2B</li> </ol>

				<p>At least one year f/u or metastatic event; median f/u 3.2 years</p> <p>Median 3.2 years</p> <hr/> <p><i>Notes:</i></p>	<p>and correlate with 31-GEP class for entire 1124 pt cohort not the 684 used to develop nomogram.</p>	<p>and higher lesions)</p> <p>3. Relatively short f/u particularly for earlier T stage lesions</p> <p>Study design has flaws:  Patient selection  No SLNB in 654 of the 684 pts  No prospective validation cohort for nomogram – used an “archival cohort of 901 stage I-III patients w/ median 6.1 yrs f/u from 22 centers (no ref) to “validate” which is different from this study population.</p>
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		<p>unrelated to this study from Castle Bioscience</p> <p>13 authors: rec'd funding for sample and/or clinical data acquisition and processing for this study</p> <p>2 authors: rec'd travel funding to present data related to this study at a meeting and/or other purposes</p>	<p>than 5 years of follow-up or a documented recurrence of melanoma [24,25,27,32].</p> <p>These patients represent a subset of the 946 patient retrospective cohort described above."</p>	<p>Outcome Measures: Sentinel Node status</p> <p>Follow-Up: 5 years</p> <p>____This is a very curated dataset_</p> <p>See Table 1.</p> <p>_____ <i>Notes:</i></p>		<p>impactful in predicting SLN status.</p> <p><b>Not</b> sure that the indications to perform a SLNB for T1a melanomas are widely used in other institutions.</p> <p>Methodology: Cohorts are not clearly defined.</p>
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		<div>1 author: leadership Castle Bioscience</div> <div>1 author: honoraria Castle Bioscience</div> <div>1 author: expert testimony Castle Bioscience</div>		<i>Notes:</i>	Results from development cohort were previously reported (Vetto, Future Oncol 2019).	
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			<p>model incorporating the 31-GEP result, Breslow thickness, ulceration, mitotic rate, LVI, transected base, presence of regression, presence of TILs, microsatellites, patient preference, patient age, and patient comorbidities</p> <p>SLNB reduction rates were analyzed based on a previous algorithm indicating that patients with a 31-GEP Class 1A result who are either 65 years old (primary analysis) or 55 years old (secondary analysis) with T1- T2 tumors have a &lt; 5% risk of SLN positivity</p> <p>172 patients needed</p>	<p>SLNB reduction rate compared to historical control</p> <p>Surgeon report of utility of 31 GEP in decision to perform or not perform SNLB</p> <p>What proportion of SLNB eligible patients decision for SLNB was influenced by DecisionDx test</p> <p>Follow-Up: n/a</p> <hr/> <p>Notes: The title of the journal is informative: Medical Research and Opinion.</p>	<p>forgo SLNB with 70/100 not performed</p> <p>Clinicians 89.1% surg onc, 7.8% dermatologists, 3.1% med onc.</p> <p>32.6% (63/191) influenced to perform with 92/1% performed.</p> <p>SLNB performance rate 59.1% lower than baseline 78%</p> <p>SLNB positivity class 1A 3.0%, comparison to Class1B/2A performance rate 80.6%, positivity rate 13.8%; and class 2B SLNB performance 94.7% and positivity 22.2%</p> <p>Pt &gt; 55 yo performance class IA SLNB 44.6 %, + 2.2% 1B/2A 77.4%, 16.7% 2B 100%, 26.7%</p> <p>Pt &gt; 65 yo performance class; SLNB rate, positive IA SLNB 48.0 %, + 0% 1B/2A 71.4%, +26.7% 2B 100%, +33.3%</p> <p>52% of clinical decisions were influenced by the results of the 31-GEP test</p> <hr/> <p>Notes: Lowest risk group Class IA doesn't make sense influenced 88.9% (16/18) to perform SLNB. What was time to treatment</p>	<p>Tumor location not variable</p> <p>Do not explain survey of clinicians in 31-GEP and other factors influencing decision.</p> <p>SLNB reduction rates analyzed using algorithm based on prior 31-GEP study.</p> <p>Grouped class 1B/2A.</p> <p>No patient outcomes are tied to decision making. The clinician is what is being studied here and unclear how many clinicians were included in study</p> <p>Bias in case selection – they were all patients who had the test done No comparison to other methods of predicting SLN positivity</p> <p>There is no outcome data—specifically no data on which patients who did not</p>
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					<p>considering that Castel ordered at clinic visit 1.</p> <p>Factors such as LVI, microsatellites, ulceration had no impact on SLNB decision.</p> <p>Sensitivity analysis oddly reported with rate of SLNB.</p> <p>Not clear what an appropriate historical control is for this cohort.</p> <p>The question here appears to be when the test was ordered and resulted prior to therapeutic operation for melanoma did the test results influence the operative plan (SLNB planned or omitted)</p> <hr/> <p>Notes:</p>	<p>undergo SLNB developed regional nodal disease—does not therefore really provide meaningful data on the accuracy of this assay</p>
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					<div><div>SORT is Strength of Recommendation Taxonomy (A, B, or C)</div></div>	<div><div>particularly for T1 lesions</div><div>This is a consensus panel (recommendations made as the National Society for Cutaneous Medicine)</div></div>
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## Question 2

Does GEP testing improve current risk-stratification of adult patients with AJCC pT1a-pT4b primary cutaneous melanoma sufficiently to recommend its utilization to guide decision-making for surveillance imaging and follow-up?

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Level of Evidence* <sup>^</sup>  Choose one:	Study Citation Author(s), Year, Title, Journal, Volume, Page #s	Funding Source	Study Methodology	Objective & Methods	Results	Conclusions
<div> <div> <input type="checkbox"/> EL 1; RCT  <input type="checkbox"/> EL 1; MRCT    <input type="checkbox"/> EL 2; MNRCT  <input type="checkbox"/> EL 2; NMA  <input type="checkbox"/> EL 2; NRCT  <input type="checkbox"/> EL 2; PCS  <input checked="" type="checkbox"/> EL 2; RCCS  <input type="checkbox"/> EL 2; NCCS  <input type="checkbox"/> EL 2; CSS  <input type="checkbox"/> EL 2; ES  <input type="checkbox"/> EL 2; OLES  <input type="checkbox"/> EL 2; PHAS    <input type="checkbox"/> EL 3; DS  <input type="checkbox"/> EL 3; ECON  <input type="checkbox"/> EL 3; CCS  <input type="checkbox"/> EL 3; SCR  <input type="checkbox"/> EL 3; PRECLIN  <input type="checkbox"/> EL 3; BR    <input type="checkbox"/> EL 4; NE  <input type="checkbox"/> EL 4; O </div> <div> <div>Strong</div> <div>Intermediate</div> <div>Weak</div> <div>No Evidence</div> </div> </div>	<p>Amaral TM, Hoffman MC, Sinnberg T, et al. Clinical Validation of a Prognostic 11-Gene Expression Profiling Score in Prospectively Collected FFPE Tissue of Patients with AJCC v8 Stage II Cutaneous Melanoma. <i>Eur J Cancer</i>. 2020 Jan;125:38-45. Epub 2019 Dec 12. (111)</p> <p>Country: Germany</p>	<p>Source: None declared;</p> <p>3 authors rec'd grants and/or personal fees from Neracare while conducting the study.</p> <p>1 author rec'd grants from Neracare outside the study.</p> <p>1 author is a Neracare employee since 2017.</p>	<p>Methodology: Retrospective cohort study.</p> <p>Stage II melanoma pts 2000-2016 with tissue for testing from German registry.</p>	<p>Stated Objective: Assess the prognostic utility of 11-gene signature in a dichotomized manner.</p> <p><input type="checkbox"/> Prospective <input checked="" type="checkbox"/> Retrospective</p> <p>Study Population and Setting: Stage II patients with melanoma between 2000 and 2016; German registry</p> <p>N: 246 24 of 302 with tissue from 1755 stage II melanoma pts in registry.</p> <p>Intervention: 11-GEP</p> <p>Outcome Measures: Melanoma specific survival Distant metastasis free survival Recurrence free survival</p> <p>Follow-Up: 41 months</p> <p>Notes: Analyzed MelaGenix score as dichotomized and continuous variable but modeling with dichotomized variable as low vs high risk</p>	<p>Results: 11-GEP signature was significantly associated with MSS, DMFS, and RFS and independently associated with MSS when accounting for tumor thickness and age.</p> <p>11-GEP predictive of 5 and 10 year MSS, p=0.18</p> <p>Notes:</p>	<p>Describe conclusions relative to question: 11-GEP signature can independently predict MSS when accounting for tumor thickness and age in stage II patients and may help in selection of adjuvant therapies.</p> <p>Authors conclude 11-GEP independent predictor of MSS but in MVA Breslow depth and age also highly significant with tighter Cis for tumor thickness, therefore more significant P value AND that patients with high score</p>

						<p>should have adjuvant systemic therapy which cannot be concluded from these data.</p> <p><i>Critiques of Methodology:</i> Not adjusting for other factors such as ulceration 20% of patients did not undergo SLN biopsy.</p> <p>20% of patients didn't have SLN surgery.</p> <p>No information on characteristics of study cohort vs untested stage II patients.</p> <p>Treatment with immune checkpoint blockade not controlled for.</p> <p>Is 10 year endpoint valid for data set with 3.5 year median f/u?</p>
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Summary of Evidence Table Question 1, 2						
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<div> <div> <input type="checkbox"/> EL 1; RCT  <input type="checkbox"/> EL 1; MRCT    <input type="checkbox"/> EL 2; MNRCT  <input type="checkbox"/> EL 2; NMA  <input type="checkbox"/> EL 2; NRCT  <input type="checkbox"/> EL 2; PCS  <input type="checkbox"/> EL 2; RCCS  <input type="checkbox"/> EL 2; NCCS  <input type="checkbox"/> EL 2; CSS  <input type="checkbox"/> EL 2; ES  <input type="checkbox"/> EL 2; OLES  <input type="checkbox"/> EL 2; PHAS    <input type="checkbox"/> EL 3; DS  <input type="checkbox"/> EL 3; ECON  <input checked="" type="checkbox"/> EL 3; CCS  <input type="checkbox"/> EL 3; SCR  <input type="checkbox"/> EL 3; PRECLIN  <input type="checkbox"/> EL 3; BR    <input type="checkbox"/> EL 4; NE  <input type="checkbox"/> EL 4; O </div> <div> <div>Strong</div> <div>Intermediate</div> <div>Weak</div> <div>No Evidence</div> </div> </div>	<p>Amaral T, Sinnberg T, Chatziioannou E, et al. Identification of Stage I/II Melanoma Patients at High Risk for Recurrence Using a Model Combining Clinicopathologic Factors with Gene Expression Profiling (CP_GEP). <i>Eur J Cancer</i>. 2023;182:155-162.</p> <p><i>Country:</i> Germany Germany/Netherlands</p>	<p><i>Source:</i> <u>The study was partially funded by SkylineDx.</u></p> <p><u>4 authors: SkylineDx employees</u></p> <p><u>2 author: institutional funding from SkylineDx in relation to the submitted work.</u></p> <p><u>2 author: Institutional financial support from Neracare</u></p> <p><u>5 author: stock/ownership interest in SkylineDx</u></p> <p><u>1 author: leadership SkylineDx</u></p> <p><u>1 author: consulting or advisory role</u></p>	<p><i>Methodology:</i> Blinded retrospective single-center of patients with stage I/II patients.</p> <p>Single center RP review of a prospective database.</p>	<p><i>Stated Objective:</i> To validate CP-GEP for the identification of high-risk for disease recurrence stage I/II patients.</p> <p><input type="checkbox"/> Prospective <input checked="" type="checkbox"/> Retrospective</p> <p><i>Study Population and Setting:</i> Stage I/II, age &gt;18, bw 2000-2017, with negative SLNB</p> <p>Patients with Stage I/II melanoma, SLN negative</p> <p><i>N:</i> 543 patients with SLNB+ 83 control without SLNB, of which 80 analyzed</p> <p><i>Intervention:</i> CP-GEP with clinicopath features age and Breslow thickness with the expression of (ITGB3, PLAT, SERPINE2, GDF15, TGFB1, LOXL4, CXCL8 and MLANA) corrected by (RLP0 and ACTB)</p> <p>CP-GEP combined protocol to predict recurrence</p>	<p><i>Results:</i> W SLNB CP-GEP stratified 424 patients (78% cohort) into RFS rates of 77.8% for high risk (195 patients) and RFS rate of 93% for low risk (229 patients). HR 3.53 ( p, 0.001).</p> <p>RFS stage I 90.7%, stage II 66.1% DMFS stage I 96.0%, stage II 82.2%. OS stage I 95.6%, stage II 79.0%</p> <p>For stage I/II: CP-GEP classified 311 as high risk with HR 4.73 with p-value &lt;0.001</p> <p>Subgroup I/IIA: CP-GEP stratified low v high-risk with a HR of 3.53 ( p-value &lt;0.001).</p> <p>RFS I/IIA for high-risk (195 patients) was 77.8% and 93.0% for low-risk.</p> <p>W/o SLNB Classified 11/80 high-risk which captured 6/7 recurrences.</p> <p>The five-year RFS for stage I/II patients was 79.9%</p>	<p><i>Describe conclusions relative to question:</i></p> <p>CP-GEP predicts high risk pt in stage I/II and could be used to replace SLNB and be used to guide adjuvant treatments.</p> <p>CP-GEP could replace SLNB</p> <p>CP-GEP enhances traditional histologic classification of patients.</p> <p><i>Critiques of Methodology:</i> Group stages I/II and do not separate to be able to discern clinicopathologic characteristics vs CP-GEP. The stage I/II vs I/IIA</p>

				<p>Outcome Measures: Kaplan-Meier curves to determine prognostic value. Primary endpoint was RFS also DMFS and OS.</p> <p>Recurrence after negative SLNbx</p> <p><i>Follow-Up:</i> Median follow up of 83.63 months for patients with negative SLNB outcome</p> <p>40.77 mo for patients wo SLNB</p> <hr/> <p><i>Notes</i></p>	<p>(95% CI: 76.0%e83.2%). CP-GEP identified 311 patients as high risk for disease recurrence with a HR of 4.73 with a p-value &lt;0.001, capturing 83 out of 98 reported relapses. For subgroup stage I/IIA, CP-GEP was able to significantly stratify CPGE low-risk and high-risk patients with a HR of 3.53 (p value &lt;0.001) for five-year RFS. For stage I/IIA, the five-year RFS rates for CP-GEP high-risk patients were 77.8% (95% CI: 70.9%e83.3%) and 93.0% (95% CI: 88.5%e95.8%) for CP-GEP low-risk patients. “Compared to AJCC low-risk (stage I/IIA) patients with an RFS rate of 86.0% (95% CI: 82.0%e89.1%), CP-GEP was able to split 195 high-risk patients who had a worse five year RFS survival of 77.8%”</p> <hr/> <p>Notes: : Wo slnb high-risk curves are dramatically different, does this argue for doing a slnb.</p> <p>Wo slnb 5/11 classified as high-risk without recurrence.</p> <p>They group the analyses into “who would qualify for adjuvant therapy vs who wouldn’t” There is no</p>	<p>Initial curves again do not separate by histologic stage. Group “Stage I/II” together.</p>
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					subset analysis by the actual histologic stages which aids their analysis but doesn't accurately reflect our ability to discern.	
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## Summary of Evidence Table

### Question 2

Level of Evidence*^  Choose one:	Study Citation Author(s), Year, Title, Journal, Volume, Page #s	Funding Source	Study Methodology	Objective & Methods	Results	Conclusions
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		<p>The Connecticut Tumor Registry is supported by Federal funds from the National Cancer Institute, National Institutes of Health, Department of Health and Human Services, under Contract no. HHSN261201800002I.</p> <p>University of Hawaii Cancer Center, HI through National Cancer Institute, Surveillance, Epidemiology and End Results Program Contract Award HHSN261201300009I. : Department of Population and Public Health Sciences, Keck School of Medicine, University of Southern California, Los Angeles, CA through National Cancer Institute, Surveillance, Epidemiology and End Results Program Contract Award HHSN261201800032I</p> <p>: Cancer Data Registry of Idaho, Idaho Hospital Association, Boise, Idaho through National Cancer Institute, Surveillance, Epidemiology and End Results Program Contract Award HHSN261201800006I and Centers for Disease Control and Prevention 1NU58DP006270.</p> <p>Bureau of Cancer Epidemiology, New York State Department of Health, Albany, NY through National Cancer Institute, Surveillance, Epidemiology and End Results Program Contract Award HHSN261201800005I and Centers for Disease Control and Prevention NU58DP006309</p> <p>: Division of Public Health Sciences Fred Hutchinson Cancer Center, through National Cancer Institute, Surveillance, Epidemiology and</p>		<p><i>Intervention:</i> 31-GEP 31 GEP vs no testing 2016-2018</p> <p><i>Outcome Measures:</i> MSS, OS</p> <p><i>Follow-Up:</i> Not specifically reported, limited to 3 years by inclusion criteria</p> <hr/> <p><i>Notes:</i></p>	<p>methodology, the conclusions of the research on 31-GEP's effectiveness are potentially weakened. More rigorous studies are essential to confirm the findings.</p>	<p>The study relied on a three-year f/u for survival outcomes, potentially skewing data due to the death of higher-stage patients within this short time frame. Absence of data on specific outcomes such as recurrence-free and distant metastasis-free survival. The study also lacked details on results specific to clinical substages and did not include clinical stage as a factor in multivariable analyses. Selection criteria for the 31-GEP patient cohort (comprising 4,687 individuals) is vague, raising concerns about selection bias and the cohort's representation of the wider melanoma patient demographic in the US.</p>
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		<p>End Results Program Contract Award HHSN2612018000041</p> <p>Emory University, Atlanta GA. through National Cancer Institute, Surveillance, Epidemiology and End Results Program Contract Award HHSN261201800003I and Centers for Disease Control and Prevention 6NU58DP006352-05-01</p> <p>Emory University, Atlanta, GA, through National Cancer Institute, Surveillance, Epidemiology and End Results Program Contract Award HHSN261201800014I.</p> <p>School of Public Health Louisiana State University Health New Orleans, through National Cancer Institute, Surveillance, Epidemiology and End Results Program Contract Award HHSN261201800007I/ HHSN26100002.</p> <p>7 authors: employees Castle Biosciences 7 author: stock &amp; other ownership interests Castle Biosciences</p> <p>1 author: research funding Castle Biosciences</p> <p>1 author: consulting or advisory role Castle Biosciences</p> <p>3 author: patents, royalties, other intellectual property Castle Biosciences</p> <p>3 author: travel, accommodations, expenses Castle Biosciences</p> <p>1 author: leadership Castle Biosciences</p>				
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Summary of Evidence Table Question 2						
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		1 peer reviewer= is a consultant to Castle Biosciences				management changes were beneficial to patients based on outcome.
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		3 author: Castle Bioscience advisory board				
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Summary of Evidence Table Question 1, 2, 3						
Level of Evidence*^  Choose one:	Study Citation Author(s), Year, Title, Journal, Volume, Page #s	Funding Source	Study Methodology	Objective & Methods	Results	Conclusions
<div> <div> <input type="checkbox"/> EL 1; RCT  <input type="checkbox"/> EL 1; MRCT  <input type="checkbox"/> EL 2; MNRCT  <input type="checkbox"/> EL 2; NMA  <input type="checkbox"/> EL 2; NRCT  <input type="checkbox"/> EL 2; PCS  <input type="checkbox"/> EL 2; RCCS  <input type="checkbox"/> EL 2; NCCS  <input type="checkbox"/> EL 2; CSS  <input type="checkbox"/> EL 2; ES  <input type="checkbox"/> EL 2; OLES  <input type="checkbox"/> EL 2; PHAS  <input checked="" type="checkbox"/> EL 3; DS  <input type="checkbox"/> EL 3; ECON  <input type="checkbox"/> EL 3; CCS  <input type="checkbox"/> EL 3; SCR  <input type="checkbox"/> EL 3; PRECLIN  <input type="checkbox"/> EL 3; BR  <input type="checkbox"/> EL 4; NE  <input type="checkbox"/> EL 4; O </div> <div> <div>Strong</div> <div>Intermediate</div> <div>Weak</div> <div>No Evidence</div> </div> </div>	<p>Cook RW, Middlebrook B, Wilkinson J, et al. Analytic validity of DecisionDx-Melanoma, a gene expression profile test for determining metastatic risk in melanoma patients. <i>Diagn Pathol.</i> Feb 13, 2018;13(1):13.</p> <p><i>Country:</i> USA</p>	<p><i>Source:</i> This study was sponsored by Castle Biosciences</p> <p>7 authors: employees of Castle Biosciences</p> <p>7 authors: hold stock in the company</p>	<p><i>Methodology:</i></p> <p>Lab validation study</p> <p>Inter-assay, inter-instrument, and inter-operator studies to evaluate reliability of test results, sample stability and reagent stability.</p>	<p><i>Stated Objective:</i> Assess reproducibility of Decision DX assay</p> <p><input type="checkbox"/> Prospective <input checked="" type="checkbox"/> Retrospective</p> <p><i>Study Population and Setting:</i></p> <p><i>N:</i> 8244 samples run (March 2013-June 2016) 168 samples run twice to assess reproducibility</p> <p>168 specimens (de-identified of patient info)</p> <p><i>Intervention:</i> none</p> <p><i>Outcome Measures:</i> Reproducibility of Decision Dx assay</p> <p><i>Follow-Up:</i></p> <hr/> <p><i>Notes:</i> Based on existing fixed tissue samples</p>	<p><i>Results:</i> Among all samples, the technical success of the assay was 98% in the 85% of samples that met pre-specified tumor content parameters.</p> <p>Concordance among assays (in 168 subset of samples for which repeat assay was performed) on two consecutive days was 99%. Inter-instrumental concordance was 95%.</p> <p>Analytic validity based on inter-assay, inter-operator, and inter-instrument reliability measurements show technical success of the test (99% rate).</p> <hr/> <p><i>Notes:</i></p>	<p><i>Describe conclusions relative to question:</i></p> <p>The assay is reliable and reproduceable.</p> <p><i>Critiques of Methodology:</i></p>

**Summary of Evidence Table**  
**Question 2**

Level of Evidence*^	Study Citation Author(s), Year, Title, Journal, Volume, Page #s	Funding Source	Study Methodology	Objective & Methods	Results	Conclusions
Choose one:						
<div><div><div><input type="checkbox"/> EL 1; RCT <input type="checkbox"/> EL 1; MRCT</div><div><input type="checkbox"/> EL 2; MNRCT <input type="checkbox"/> EL 2; NMA <input type="checkbox"/> EL 2; NRCT <input type="checkbox"/> EL 2; PCS <input checked="" type="checkbox"/> EL 2; RCCS <input type="checkbox"/> EL 2; NCCS <input type="checkbox"/> EL 2; CSS <input type="checkbox"/> EL 2; ES <input type="checkbox"/> EL 2; OLES <input type="checkbox"/> EL 2; PHAS</div><div><input type="checkbox"/> EL 3; DS <input type="checkbox"/> EL 3; ECON <input type="checkbox"/> EL 3; CCS <input type="checkbox"/> EL 3; SCR <input type="checkbox"/> EL 3; PRECLIN <input type="checkbox"/> EL 3; BR</div><div><input type="checkbox"/> EL 4; NE <input type="checkbox"/> EL 4; O</div></div><div>Strong   &lt;</div></div>						

			and did not have routine imaging surveillance for recurrence.			<p>for follow up without imaging. Unclear if imaging alone without GEP would have had same results.</p> <p>Retrospective Differing f/u Uncontrolled survival analysis Excluded 25 patients from experimental group who recurred but didn't follow imaging protocol</p>
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Summary of Evidence Table  
Question 1, 2

Level of Evidence*^  Choose one:	Study Citation Author(s), Year, Title, Journal, Volume, Page #s	Funding Source	Study Methodology	Objective & Methods	Results	Conclusions
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<div> <input type="checkbox"/>EL 1; RCT  <input type="checkbox"/>EL 1; MRCT   <input type="checkbox"/>EL 2; MNRCT  <input type="checkbox"/>EL 2; NMA  <input type="checkbox"/>EL 2; NRCT  <input type="checkbox"/>EL 2; PCS  <input type="checkbox"/>EL 2; RCCS  <input type="checkbox"/>EL 2; NCCS  <input type="checkbox"/>EL 2; CSS  <input type="checkbox"/>EL 2; ES  <input type="checkbox"/>EL 2; OLES  <input type="checkbox"/>EL 2; PHAS   <input type="checkbox"/>EL 3; DS  <input type="checkbox"/>EL 3; ECON  <input type="checkbox"/>EL 3; CCS  <input type="checkbox"/>EL 3; SCR  <input type="checkbox"/>EL 3; PRECLIN  <input type="checkbox"/>EL 3; BR   <input checked="" type="checkbox"/>EL 4; NE  <input type="checkbox"/>EL 4; O </div>	<div> <div>Strong</div> <div>Intermediate</div> <div>Weak</div> <div>No Evidence</div> </div>	<p>Farberg AS, Glazer AM, White R, Rigel DS. Impact of a 31-gene Expression Profiling Test for Cutaneous Melanoma on Dermatologists' Clinical Management Decisions. <i>J Drugs Dermatol</i>. 2017;16(5):428-431. (16)</p> <p>Country: USA</p>	<p><i>Source:</i> None noted</p> <p>2 authors=consultants to Castle Biosciences, Inc.</p> <p>1=author participated in research fellowship partially funded by Castle Biosciences.</p>	<p><i>Methodology:</i></p> <p>Gave Derm residents patient vignettes to assess clinical decision making, incorporating 31-GEP. They were given a presentation before on “clinical validity” for GEP.</p> <p>Survey study of dermatology residents provided case-based scenarios involving management of 31-GEP results.</p>	<p><i>Stated Objective:</i> To determine impact of 31-GEP test on clinical decision making.</p> <p><input checked="" type="checkbox"/>Prospective <input type="checkbox"/>Retrospective</p> <p><i>Study Population and Setting:</i> Survey study of case-based scenarios provided to dermatology residents.</p> <p>Derm residents at a national conference given clinical scenarios involving 31-GEP test (Stage IA, IB, IIA pts).</p> <p><i>N:</i> 169 dermatology resident physicians of 172 attendees responded to survey questions</p> <p><i>Intervention:</i> 31-GEP results given or not with clinical info.</p> <p>Six cutaneous melanoma cases with GEP information varied.</p> <p><i>Outcome Measures:</i> Management decisions (SLN bx, onc referral)</p> <p>Clinical recommendation</p> <p><i>Follow-Up:</i> N/A</p> <p><i>Notes:</i></p>	<p><i>Results:</i> Approximately 50% of respondents changed clinical decision making when faced with a Class II GEP result.</p> <p>Most respondents gave 1.0 mm Breslow as guiding modality for SLNbx.</p> <p>Basically states that when given 31-GEP results, specifically class 2 result, Breslow depth to guide SLNbx changed (more likely to refer thinner Breslow depth if high risk GEP class). Also more likely to recommend imaging.</p> <p><i>Notes:</i></p> <p><i>Describe conclusions relative to question:</i> “GEP result had a significant and appropriate impact on clinical decision making” (there was no evidence provided that this was appropriate).</p> <p>The 31-GEP test had a “significant and appropriate” impact on management while remaining within established guidelines.</p> <p><i>Critiques of Methodology:</i> Provides no information regarding validity of GEP</p> <p>Were they trying to get the “right” answer based on who was presenting info to them? -Who gave the lecture on clinical validity of the test before the study??- that could significantly bias results</p>
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						<ul style="list-style-type: none"><li>-Authors do not recognize these limitations in the manuscript</li><li>-They gave the residents an answer of “onc” and imaging referral which would not be currently recommended for the thin Breslow depths noted; confusing</li></ul>
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						<div>process (only 8 panelists).</div> <div>Review of existing data.</div>
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**Summary of Evidence Table**  
**Question 2,3**

Level of Evidence*^  Choose one:	Study Citation Author(s), Year, Title, Journal, Volume, Page #s	Funding Source	Study Methodology	Objective & Methods	Results	Conclusions
<div> <div> <input type="checkbox"/> EL 1; RCT  <input type="checkbox"/> EL 1; MRCT    <input type="checkbox"/> EL 2; MNRCT  <input type="checkbox"/> EL 2; NMA  <input type="checkbox"/> EL 2; NRCT  <input type="checkbox"/> EL 2; PCS  <input type="checkbox"/> EL 2; RCCS  <input type="checkbox"/> EL 2; NCCS  <input type="checkbox"/> EL 2; CSS  <input type="checkbox"/> EL 2; ES  <input type="checkbox"/> EL 2; OLES  <input type="checkbox"/> EL 2; PHAS    <input type="checkbox"/> EL 3; DS  <input type="checkbox"/> EL 3; ECON  <input checked="" type="checkbox"/> EL 3; CCS  <input type="checkbox"/> EL 3; SCR  <input type="checkbox"/> EL 3; PRECLIN  <input type="checkbox"/> EL 3; BR    <input type="checkbox"/> EL 4; NE  <input type="checkbox"/> EL 4; O </div> <div> <div>Strong</div> <div>Intermediate</div> <div>Weak</div> <div>No Evidence</div> </div> </div>	<p>Ferris LK, Farberg AS, Middlebrook B, et al. Identification of high-risk cutaneous melanoma tumors is improved when combining the online American Joint Committee on Cancer Individualized Melanoma Patient Outcome Prediction Tool with a 31-gene expression profile-based classification. <i>J Am Acad Dermatol.</i> May 2017;76(5):818-825.e3.</p> <p>Country: USA</p>	<p>Source: Funded by Castle Biosciences which provided financial compensation to those centers contributing cutaneous melanoma tissue to the study.</p> <p>6 authors: employees of Castle Biosciences</p> <p>4 authors: served as consultants to Castle Bioscience</p>	<p>Methodology: RNA isolated to run 31 gene test.</p> <p>Five-year survival for each patient using AJCC Individualized Melanoma Patient Outcome Prediction Tool</p> <p>Survival cutoff for low and high-risk determined by stage IIA (79%) and stage IIB (68%) cohorts.</p> <p>K-M analysis and Cox proportional hazards survival analysis performed.</p> <p>205 cases of stage I/II CM with available tissue were collected from six US centers.</p>	<p>Stated Objective: To compare accuracy of GEP with risk determined using the web-based AJCC Individualized Melanoma Patient Outcome Prediction Tool.</p> <p>Compare recurrence risk prediction by AJCC Stage vs. 31-GEP class.</p> <p><input type="checkbox"/> Prospective <input checked="" type="checkbox"/> Retrospective</p> <p>Study Population and Setting: Stage I/II CM. 205 retrospectively collected.</p> <p>N: 205</p> <p>Intervention: GEP and AJCC prediction tool GEP vs conventional staging</p> <p>Outcome Measures: RFS, DMFS, OS</p> <p>Follow-Up: 6.9 y f/u 6.9 years (0.1-15.4)</p> <p>Notes:</p>	<p>Results: 43 (21%) cases had discordant GEP and AJCC classification (with 79% cutoff). 11/43 discordant cases classified as high risk GEP but low by AJCC.</p> <p>Sensitivity GEP 82%, 81% and 78% for RFS, DMFS and MSS. AJCC 70%, 69% and 60%.</p> <p>Specificity GEP 77%, 69%, 69% and AJCC 83%, 76%, 74%</p> <p>Increased sensitivity in combining GEP + AJCC with decreased specificity.</p> <p>Multivariate cox regression GEP w AJCC tool indicated GEP more significantly associated with DM and death than binary AJCC.</p> <p>HR GEP vs AJCC 79% Recurrence- 5.9 v 3.6 DMFS- 5.3 v 3.0, Death -5.3 vs 2.2</p> <p>Use of GEP test with AJCC stage improved sensitivity to detect recurrence and to a lesser degree death at the cost of specificity compared with AJCC alone.</p>	<p>Describe conclusions relative to question:</p> <p>To combine GEP with AJCC may help identify pt benefit from increased surveillance and administer therapies.</p> <p>GEP can improve detection of recurrence/mets when used in combination with AJCC.</p> <p>Somewhat overstated based on limitations of dataset.</p> <p>Critiques of Methodology: Using AJCC tool as binary high-risk vs low-risk to fit comparison.</p> <p>Unconventional and narrow to define sensitivity and specificity of</p>

					<div>Notes:</div>	<p>test using definition of low-risk group from stage IIA AJCC 79% survival and high-risk 68% survival based off of stage IIB.</p> <p>These patients were previously published in the development of the model. Unclear how this subset was selected. Stage was dichotomized at IIA- vs IIB+ somewhat arbitrarily.</p>
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				<p>Recurrence-free survival – primary DMFS and DSS - secondary</p> <p><i>Follow-Up:</i> At least 5-years or documented recurrence</p> <p>Minimum 5 years or event</p> <hr/> <p><i>Notes:</i></p>	<p>melanoma-specific mortality events.</p> <p>-31-GEP Class 2A not significant predictor RFS, and DMFS.</p> <p>Stage I - IIA Class 1A had significantly better 5-year RFS, DMFS and MSS than 2B (p&lt;.0001).</p> <p>For stage I-IIA, 31-GEP class 2B most significant predictor of RFS and DMFS in multivariate analysis including thickness, ulceration, and mitotic rate. Thickness was only significant for RFS.</p> <p>Multivariate analysis for MSS and only Class 2 significant predictor of MSS.</p> <p>Majority thin tumors T1 were low-risk Class 1 89.3% (251/281) however 5.3% Class 2 B. (2.0% of T1a and 13.9% of T1b).</p> <p>Thin tumor Class 1A v 2B RFS 96.8% and 64.6% respectively (p&lt;0.001). DMFS 97.2% v 84.4% (p=0.007). Only one death in this group so MSS not done.</p> <p>Thin tumor Cox multivariate analysis of thickness, mitoses, ulceration and SLNB positivity showed 31-GEP Class 2B only independent and significant predictor of</p>	<p><i>Critiques of Methodology:</i> Retrospective -size of cohort especially as analyze smaller groups such as in stage or ulceration and fewer events.</p> <p>31-GEP most significant single variable for predictor but what about combination of thickness, ulceration and mitosis as is clinically used. Limitations discussed included incomplete pathologic staging data owing to variation in reporting standards.</p> <p>Only 429 (63%) staged with SNB.</p> <p>Authors have conflict of interest – on advisory board for Castle Bioscience.</p>
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					<div>RFS. n=57; HR 9.34, p=.004).</div> <div>Notes:</div>	
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				<p>Primary: RFS, DMFS, OS, MSS Secondary endpoint: 31-GEP predicted outcome in combination with node status to determine prognostic value added.</p> <p>DSS/DMFS/DSS &amp; OS</p> <p><i>Follow-Up:</i> ≥ 5 years required, median length 7.1 years</p> <p>5 years or event</p> <hr/> <p><i>Notes:</i></p>	<p>.01). Class 2 predictor of DM (p=.04).</p> <p>Binary GEP combined with node status for Kaplan-Meier curves. Class 1/ node-negative v Class 2 / node negative RFS 83% v 37%, DMFS 85% v 45%, OS 91% v 62% and MSS ws 96% v 78%.</p> <p>Class 1/node-positive v Class 2/node positive RFS 29% v 19%, DMFS 43% v 23%, OS 100% v 36% and MSS ws 100% v 50%.</p> <p>Molecular class sensitivity for prediction recurrence, DM, death any cause, and melanoma-specific death 74%, 74%, 80%, and 88%. compared to node status 41%, 40%, 43%, and 52%.</p> <p>NPV molecular class 76%, 78%, 87%, and 96% compared to node status 64%, 67%, 76%, and 90%.</p> <p>Combining molecular and node increased the accuracy of identifying high-risk sensitivity of 81%, 80%, 82% and 88%. NPV 81%, 82%, 88% and 96%.</p> <hr/> <p>Notes: Figure 2 Class1/node+ 1 event and 0 events</p>	<p>Size of cohort too small for subgroup analysis</p> <p>Limitation discussed in paper is low sample size impacting multivariate analysis w low MSS events corrected by using only GEP class and AJCC stage but then doesn't control for all clinicopathologic factors.</p> <p>Statement that this cohort has some features more aggressive than clinical population should be expounded upon</p> <p>What does that say of selection bias</p> <p>Same cohort has been used in Gastman BR, Gerami P, Kurley SJ, Cook RW, Leachman S, Vetto JT. Identification of patients at risk of metastasis using a prognostic 31-gene expression</p>
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						<p>profile in subpopulations of melanoma patients with favorable outcomes by standard criteria. JI Am Acad Dermatol. 2019;80(1):149-157.e144</p> <p>Small cohort of patients</p> <p>Specificity and sensitivity data quoted for outcome? appropriate?</p>
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**Question 2**

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<div> <input type="checkbox"/> EL 1; RCT  <input type="checkbox"/> EL 1; MRCT  <input type="checkbox"/> EL 2; MNRCT  <input type="checkbox"/> EL 2; NMA  <input type="checkbox"/> EL 2; NRCT  <input type="checkbox"/> EL 2; PCS  <input checked="" type="checkbox"/> EL 2; RCCS  <input type="checkbox"/> EL 2; NCCS  <input type="checkbox"/> EL 2; CSS  <input type="checkbox"/> EL 2; ES  <input type="checkbox"/> EL 2; OLES  <input type="checkbox"/> EL 2; PHAS  <input type="checkbox"/> EL 3; DS  <input type="checkbox"/> EL 3; ECON  <input type="checkbox"/> EL 3; CCS  <input type="checkbox"/> EL 3; SCR  <input type="checkbox"/> EL 3; PRECLIN  <input type="checkbox"/> EL 3; BR  <input type="checkbox"/> EL 4; NE  <input type="checkbox"/> EL 4; O </div>	<div> <p><b>Strong</b></p> <p>Gerami P, Cook RW, Russell MC, et al. Gene expression profiling for molecular staging of cutaneous melanoma in patients undergoing sentinel lymph node biopsy. <i>J Am Acad Dermatol.</i> 2015;72(5):780-785.e783. (7)</p> <p><i>Country:</i> USA</p> <p><b>Intermediate</b></p> <p><b>Weak</b></p> <p><b>No Evidence</b></p> </div>	<div> <p><i>Source:</i></p> <p>Partially supported by Castle BioSciences Inc.</p> <p>4 authors: employees of Castle Biosciences</p> <p>3 authors: consultants to Castle Biosciences</p> <p>3 authors: speakers for Castle Biosciences</p> </div>	<div> <p><i>Methodology:</i></p> <p>Case series of patients with GEP 31-GEP result.</p> <p>RT-PCR; 31 genes.</p> <p>SLN outcome from clinical data.</p> <p>DFS DMFS OS</p> </div>	<div> <p><i>Stated Objective:</i> Evaluate prognostic value of GEP in combination with SLNB.</p> <p><input type="checkbox"/> Prospective <input checked="" type="checkbox"/> Retrospective</p> <p><i>Study Population and Setting:</i> Multicenter retrospective cohort of patients with GEP available. Patients who underwent SLNB and had at least 5 years follow up were included.</p> <p><i>N:</i> 217 cutaneous melanoma undergoing SLNBx.</p> <p>1998-2009</p> <p><i>Intervention:</i> Castle 31 gene GEP</p> <p><i>Outcome Measures:</i> DFS DMFS OS</p> <p><i>Follow-Up:</i> Not reported in abstract.</p> <p><i>Notes:</i> This cohort is a subset of pts from a previously published cohort. Not clearly stated.</p> </div>	<div> <p><i>Results:</i></p> <p>In patients with both GEP class and SLN status available, GEP class II was associated with adverse outcomes in both SLN+ and SLN- patients.</p> <p>SLNB+=58/217 SLNB- 159/217</p> <p>37 of 58 SLNB+ developed a metastatic event.</p> <p>70 of 159 SLNB- developed a metastatic event.</p> <p>Class 2 high risk: 141 Class I low risk: 76</p> <p>91 of 141 Class 2 to metastatic disease.</p> <p>16 of 76 Class 1 to metastatic disease.</p> <p>PPV of SLNB for predicting distant mets: 55% and NPV 67%.</p> <p>PPV GEP for distant mets: 50% and NPV 82%.</p> <p>OS at 5 years for SLNB+ 62%; 70% for SLN-.</p> <p>SLNB+ and GEP class 2 were predictors of DFS, DMFS, OS.</p> <p>GEP with SLNB:</p> </div>	<div> <p><i>Describe conclusions relative to question:</i></p> <p>In combination with SLNB, GEP will help identify high risk SLN- patients.</p> <p>GEP an objective tool to predict metastatic risk in SLNB-eligible pts.</p> <p>GEP outcome a more significant and better predictor of each endpoint compared to SLNB.</p> <p>With SLNB, GEP improved prognostication.</p> <p><i>Critiques of Methodology:</i> Bi-variate analysis (SLN and GEP only). Limited sample size.</p> <p>Their SLN- group had a 30% risk of metastatic</p> </div>

					<p>Class 1/SLNB- (67 pts): DFS and DMFS 83% and 86%.</p> <p>Class 1/SLNB+ (9 pts): DFS 53% DMFS 53% OS 77%.</p> <p>Class 2/SLNB+ (49 pts): DFS 33% DMFS 42%, OS 57%.</p> <p>Class 2/SLNB (58): DFS 35%, DMFS 49%, OS 55%.</p> <hr/> <p><i>Notes:</i></p>	<p>events which is high – lower OS also than expected.</p> <p>Time span is older cohort (early 2000's).</p> <p>The Breslow depth for SLNB- was 2.3 (fairly high risk group to begin with); the SLNB was 4mm- thick melanomas. Would be better to have more detailed TNM staging described instead of how they presented it and more clinicopathologic features (site of melanoma etc)..</p>
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				<p>Follow-Up: At least 5 years follow-up or metastatic event—N=34 &lt;5 years follow-up accepted in initial preliminary analysis 5 year min or until recurrence; median f/u except 34 patients who had &lt;3 yrs f/u data.</p> <hr/> <p><i>Notes:</i> Tumor density on slide&gt;60% required. Signature genes enumerated.</p>		
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						<p>Problematic methodology in a systematic review and meta-analysis of DecisionDx-Melanoma Journal of the American Academy of Dermatology, Volume 83, Issue 5, November 2020, Pages e357-e358 Michael A. Marchetti, Stephen W. Dusza, Edmund K. Bartlett</p> <p>Search criteria included “31-gene” which is specific for their assay (not unbiased look at GEP in general). Their cohorts were heterogeneous across stages I – III.</p>
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			<p>cases identified as having metastatic disease whose primary tumors were then retrieved for 31-GEP if not done previously</p>	<p><i>Follow-Up:</i> 23 months Median &lt;2 years Mean 23 months</p> <hr/> <p><i>Notes:</i> Also analyze patient and tumor features associated with 31-GEP class and performance of 1B and 2A categories in terms of clinical utility.</p> <p>SLN+ not considered as an event for this study.</p> <p>No data on untested pts.</p>	<p>13 patients developed metastases, GEP identified 77% of these.</p> <p>99% NPV of the GEP “only 3 of the 214 Class I patients metastasized.”</p> <p>24 of 37 stage II patients were Class II and 10 metastasized, (not surprising), on 42%.</p> <p>2 of 13 stage II that were Class I metastasized.</p> <p>Of 256: 214 (84%) Class 1, (193 Class 1A, 21 Class 1B) 42 (16%) Class 2, (16 Class 2A, 26 Class 2B)</p> <p>Of 256: 13 developed metastasis (10 of 13 Class 2 on testing).</p> <p>3-year MFS rate 98% for Class 1 patients and 74% for Class 2 patients, with 5-year MFS rates of 93% and 69%, respectively (p &lt; .00001)</p> <hr/> <p>Notes: No comparison based on histologic staging beyond stage I and Stage II (no substaging comparisons).</p> <hr/> <p>Notes:</p>	<p>Relatively short follow-up. Only 13 metastatic events in the entire cohort making any robust multivariate analysis not possible.</p> <p>Unclear what the treatment was of the primary (WLE? SLNB?) and type of recurrence is not stated.</p> <p>Short f/u time.</p> <p>Mention in Data Collection that separate IRB to retrospectively test tumors from a subcohort of patients with known metastatic disease after initial excision.</p> <p>Flaws in study with key treatment and other data elements (AJCC stage, e.g.) missing and very short f/u and &gt;80% of patients studied at very low risk based on conventional pathology risk factors – in the population of interest N is small.</p> <p><i>Critiques of Methodology:</i> Quoted from the paper:</p> <p><i>At the time of initial CM diagnosis, 219 (86%) of the</i></p>
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						<p><i>tested tumors were Stage I. None of the 18 Stage I Class 2 tumors metastasized, whereas 1 (0.5%) of 201 Stage I Class 1 tumors metastasized.</i></p> <p>This implies that that the GEP is not discerning in Stage I patients since Class differentiation did not predict any pattern of disease recurrence.</p> <p>Discussion tries to explain why their findings might not reflect the ultimate utility of the GEP.</p> <p>Retrospective</p> <p>Enriched for subsequent testing after recurrence in some of the patients, number of patients in this subcohort nor method for identification not stated</p> <p>With median f/u 23 months 5 yr KM survival analysis seems invalid.</p> <p>Class 1 median Breslow 0.6 mm vs Class 2 = 2.2 mm</p> <p>No analysis comparing value of 31-GEP beyond AJCC stage, Breslow Depth</p>
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		<p>work supported with resources and use of facilities at the Veterans Affairs Palo Alto Health Care System in Palo Alto CA. Contents do not represent the views of the US Dept of Veterans Affairs or the US Gov't. The funders had no role in the design and conduct of the study; collection, management, analysis and interpretation of data, preparation, review or approval of the manuscript; and decision to submit for publication.</p> <p>1 author: nonfinancial support from Castle Biosciences outside submitted work.</p> <p>1 author: grants from Castle</p>		<p>Outcome Measures: N/A</p> <p>Follow-Up:</p> <hr/> <p>Notes:</p>	<p>performing clinical trials incorporating GEP testing with SLNB and adjuvant therapy. The MPWG members favor conducting retrospective studies that evaluate multiple GEP testing platforms on fully annotated archived samples before embarking on costly prospective studies and recommend avoiding routine use of GEP testing to direct patient management until prospective studies support their clinical utility.</p> <hr/> <p>Notes:</p>	
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		<p>Biosciences outside submitted work</p> <p>1 author: personal fees outside the submitted work Castle Biosciences</p> <p>1 author: personal fees from Neracare outside submitted work</p> <p>1 author: manuscripts and abstracts published using the test with company support of the assay—all publications were peer review and no personal of institutional payment or compensation was rec'd. Castle Biosciences</p> <p>1 author: served as an investigator for Castle Biosciences (no personal financial compensation)</p>				
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					<p>recurrence-free survival, along with Breslow, mitotic rate and sentinel node status.</p> <p>Ulceration status not significant independent predictor of RFS, suggesting class 2 is surrogate marker of ulceration (which in turn would be called “poorly differentiated” if it were any other tumor)</p> <hr/> <p><i>Notes:</i></p>	<p>recommendations could be made.</p> <p><i>Critiques of Methodology:</i></p> <ul style="list-style-type: none"><li>·Short f/u &lt; than 2 years overall will not capture all recurrence/survival events.</li></ul> <p>No new evidence from this alone.</p> <p>Short F/u &lt; 2 years.</p>
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## Summary of Evidence Table Question 2

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<div> <input type="checkbox"/> EL 1; RCT  <input type="checkbox"/> EL 1; MRCT    <input type="checkbox"/> EL 2; MNRCT  <input type="checkbox"/> EL 2; NMA  <input type="checkbox"/> EL 2; NRCT  <input checked="" type="checkbox"/> EL 2; PCS  <input type="checkbox"/> EL 2; RCCS  <input type="checkbox"/> EL 2; NCCS  <input type="checkbox"/> EL 2; CSS  <input type="checkbox"/> EL 2; ES  <input type="checkbox"/> EL 2; OLES  <input type="checkbox"/> EL 2; PHAS    <input type="checkbox"/> EL 3; DS  <input type="checkbox"/> EL 3; ECON  <input type="checkbox"/> EL 3; CCS  <input type="checkbox"/> EL 3; SCR  <input type="checkbox"/> EL 3; PRECLIN  <input type="checkbox"/> EL 3; BR    <input type="checkbox"/> EL 4; NE  <input type="checkbox"/> EL 4; O </div>	<div> <div>Strong</div> <div>Intermediate</div> <div>Weak</div> <div>No Evidence</div> </div> <p>Hsueh EC, DeBloom JR, Lee JH, et al. Long-Term Outcomes in a Multicenter, Prospective Cohort Evaluating the Prognostic 31-Gene Expression Profile for Cutaneous Melanoma. <i>JCO Precis Oncol.</i> Apr 6, 2021;5:</p> <p>Country: USA</p>	<p><i>Source:</i> This study was partially sponsored by Castle Biosciences, which provided financial compensation to those who provided melanoma tissue for this study.</p> <p>4 authors: employees of Castle Biosciences</p> <p>4 author: stock and other ownership interests</p> <p>1 author: Castle Biosciences speakers bureau</p> <p>1 author: Consulting or advisory role with Castle Biosciences</p> <p>2 author: patent, royalty, other intellectual property in GEP tests or Castle</p>	<p><i>Methodology:</i> Used INTEGRATE/EXPAND Patients with Stage I-III CM.</p> <p>Multicenter registry INTEGRATE and EXPAND RFS, DMFS, OS assessed using Kaplan-Meier and Cox regression analysis.</p>	<p><i>Stated Objective:</i> To test the hypotheses that the 31-GEP provides prognostic value for patients with stage I-III CM, and that patients with stage I-IIA melanoma and class 2 31-GEP results have metastatic risk similar to patients for whom surveillance is recommended.</p> <p>To assess if the 31-GEP add prognostic value to clinicopathologic features.</p> <p><input checked="" type="checkbox"/> Prospective <input type="checkbox"/> Retrospective</p> <p><i>Study Population and Setting:</i> Stage I- III CM</p> <p>11 centers CM stage I-III pts 16 years old with GEP results</p> <p><i>N:</i> 372 assessed for eligibility. 334 met enrollment criteria and had test results were enrolled. Eleven were excluded from analysis, leaving 323 who met enrollment and analysis inclusion criteria.</p> <p><i>Intervention:</i> All had castle 31-GEP test</p>	<p><i>Results:</i> The 31-GEP was significant for RFS, DMFS, and OS in a univariate analysis and was a significant, independent predictor of RFS, DMFS, and OS in a multivariable analysis.</p> <p>GEP class 2 results were significantly associated with lower 3-year RFS, DMFS, and OS in all patients and those with stage I-IIA disease.</p> <p>Patients with stage I-IIA CM and a class 2 result had recurrence, distant metastasis, and death rates similar to patients with stage IIB-III CM.</p> <p>Class 2 were associated with high-risk CP features as compared to class 1 including age, male, Breslow, ulceration and + SLN.</p> <p>Class 2 v Class 1: 3 -year RFS 66% v 95% DMFS 79% v 97% OS 81% v 95%</p> <p>univariate analysis including class 2 (v class 1) 31-</p>	<p><i>Describe conclusions relative to question:</i> Pts with Stage I-IIA CM and class 2 GEP may be candidates for more intense f/u.</p> <p>31-GEP is a significant, independent prognostic factor for 3-year recurrence-free survival, distant metastasis-free survival, and overall survival in a prospectively enrolled cohort of patients diagnosed with stage I-III melanoma. Patients diagnosed with stage I-IIA melanoma who receive a high-risk 31-GEP result (class 2) have</p>



		<p>Bioscience related patents</p> <p>4 author: travel, accommodations, expenses, Castle Biosciences</p>		<p><i>Outcome Measures:</i> RFS, DMFS, OS</p> <p><i>Follow-Up:</i> 3.2 years</p> <hr/> <p><i>Notes:</i></p>	<p>GEP results, continuous age, male (v female) sex, head and neck (v non-head and neck) tumor location, continuous mitotic rate, and high-risk stage IIB-III (v stage I-IIA) AJCC 8th ed staging, only GEP class 2 and high-risk AJCC stages reached a threshold of P , .01 for</p> <p>all three end points; whereas, mitotic rate was significant for DMFS, and age was significant for OS.</p> <p>Similarly, subclass analysis of the 31-GEP demonstrated that classes 1B, 2A, and 2B were significant</p> <p>predictors of recurrence risk; whereas, classes 2A and 2B were significant predictors of distant metastasis, and class 2B was a significant predictor of mortality</p> <p>In a multivariable analysis, 31-GEP class 2 (hazard ratio [HR], 4.34 [95% CI, 2.10 to 8.96], P , .001) and AJCC stage IIB-III (HR, 2.98 [95% CI, 1.48 to 6.02], P = .002) were inde-</p>	<p>survival outcomes like those with stage IIB-III melanoma, for whom national guidelines recommend more intense follow-up.</p> <p><i>Critiques of Methodology:</i> Relatively short f/u</p> <p>Doesn't break down by stage</p>
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					<p>pendent, significant predictors of RFS. For 3-year DMFS, 31-</p> <p>GEP class 2 (HR, 5.45 [95% CI, 2.09 to 14.25], P , .001), AJCC stage IIB-III (HR, 2.81 [95% CI, 1.16 to 6.58], P = .023), and mitotic rate (HR, 1.04 [95% CI, 1.01 to 1.07], P = .007) were significant. 31-GEP class 2 (HR, 3.13 [95% CI, 1.23 to 7.96], P = .016), AJCC stage IIB-III (HR, 3.89 [95% CI, 1.60 to 9.50], P = .003), and age (HR, 1.08 [95% CI, 1.04 to 1.13], P , .001) were significant for OS</p> <p>identifying high-risk patients by either 31-GEP class 2 result or AJCC high-risk category, sensitivity was enhanced for 3-year RFS (76%), DMFS (88%), and OS (76%) compared with AJCC alone with sensitivities of 57% (RFS), 62% (DMFS), and 60% (OS) or 31-GEP status alone with sensitivities of 64% (RFS), 69% (DMFS), and 68% (OS). Class 2 31-GEP results identified AJCC stage I-IIA patients with increased risk for recurrence, distant</p>	
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					<p>metastasis, and death with 44%, 70%, and 40% sensitivity, respectively; whereas a class 1 result confirmed a</p> <p>low risk of recurrence, distant metastasis, and death in this population with an NPV of 92%, 98%, and 95%, respectively</p> <hr/> <p><i>Notes:</i> multivariate HR for OS almost identical</p>	
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				<p><i>Follow-Up:</i> n/a unclear</p> <hr/> <p><i>Notes:</i> Could be considered retrospective cohort study vs. data science.</p>		
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					<div>Notes: 25% of patients did not have SLNB.</div>	
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		<p>study; collection, management, analysis, and interpretation fo the data; preparation, review, or approval of the manuscript and decision to submit the manuscript for publication.</p> <p>2 authors: research support from Castle Biosciences outside this project</p> <p>1 author: research support from Castle both during and outside this work</p> <p>1 author: grant support from Skyline Dx outside this work</p> <p>1 author: nonfinancial support from SkylineDx outside the work</p>				<p>i-31-GEP) purported to predict SLN positivity.”</p> <p><i>Critiques of Methodology:</i></p> <p>Other than funding support, no concerns. Appropriately designed and performed consensus conference.</p>
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					<p>Univariate analysis Breslow thickness, ulceration, SLNB and GEP significantly associated with RFS and DMFS (p&lt;0.001).</p> <p>Multivariate analysis only SLNB and GEP class were statistically associated with both RFS (p=0.008 and 0.0001) and DMFS (p=0.0019 and 0.001).</p> <p>HR Class 2 9.2 (p&lt;0.001 CI 3.0-28.5) for RFS and 19 (p&lt;0.001 CI 2.12-170.5).</p> <p>SLNB result was also associated with RFS and DMFS (P&lt;0.02, HR=3.5, 3.7, 95% CI=1.4-9.1, 1.2-11.3).</p> <p>Breslow significant for RFS HR 1.15 ( p=0.0015, CI 1.01-1.31) not significant for DMFS in multivariate.</p> <p>Ulceration not significant.</p> <p>GEP subclass additional stratification w 3-yr RFS and DMFS rates for subclass 2B of 39.5% and 59.6%.</p> <p>Notes:</p>	Investigators not blinded to GEP results.
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						data from 4 fairly dated studies).
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## Summary of Evidence Table Question 2

Level of Evidence* <sup>^</sup>  Choose one:	Study Citation Author(s), Year, Title, Journal, Volume, Page #s	Funding Source	Study Methodology	Objective & Methods	Results	Conclusions
<div> <input type="checkbox"/> EL 1; RCT  <input type="checkbox"/> EL 1; MRCT  <input checked="" type="checkbox"/> EL 2; MNRCT  <input type="checkbox"/> EL 2; NMA  <input type="checkbox"/> EL 2; NRCT  <input type="checkbox"/> EL 2; PCS  <input type="checkbox"/> EL 2; RCCS  <input type="checkbox"/> EL 2; NCCS  <input type="checkbox"/> EL 2; CSS  <input type="checkbox"/> EL 2; ES  <input type="checkbox"/> EL 2; OLES  <input type="checkbox"/> EL 2; PHAS    <input type="checkbox"/> EL 3; DS  <input type="checkbox"/> EL 3; ECON  <input type="checkbox"/> EL 3; CCS  <input type="checkbox"/> EL 3; SCR  <input type="checkbox"/> EL 3; PRECLIN  <input type="checkbox"/> EL 3; BR    <input type="checkbox"/> EL 4; NE  <input type="checkbox"/> EL 4; O </div> <div> <div>Strong</div> <div>Intermediate</div> <div>Weak</div> <div>No Evidence</div> </div>	<p>Marchetti M, Coit D, Dusza S, et al. Performance of Gene Expression Profile Tests for Prognosis in Patients With Localized Cutaneous Melanoma: A Systematic Review and Meta-analysis. JAMA Dermatol. 2020 Sep 1;156(9):953-962. (98)</p> <p>Country: USA + international</p>	<p>Source: Funded in part by Memorial Sloan Kettering Cancer Center institutional National Institutes of Health/ National Cancer Center Support Grant P30 CA008748</p> <p>The funding source had no role in the design and conduct of the study; collection, management, analysis and interpretation of the data; preparation, review, or approval of the manuscript; and decision to submit the manuscript for publication.</p>	<p>Methodology: Meta-analysis/systematic review.</p> <p>Systematic review/meta-analysis</p> <p>Adaptation of the Checklist for Critical Appraisal and Data Extraction for Systematic Reviews of Prediction Modeling Studies (CHARMS-PF)</p> <p>Literature Review of 2015-2019 articles indexed in PubMed plus relevant ASCO 2017-2019 abstracts reviewed by 2 rounds on line survey and in-person 2 hours meeting with review of survey results, relevant articles.</p>	<p>Stated Objective: Performance of commercially available GEP tests in predicting outcomes in patients with stage I/II melanoma.</p> <p><input checked="" type="checkbox"/> Prospective <input checked="" type="checkbox"/> Retrospective</p> <p>Systematic review and Meta-analysis included retrospective and retrospective studies</p> <p>Study Population and Setting: 7 studies (5 DecisionDx; 2 MelaGenix) 1450 study participants</p> <p>Meta-analysis and expert consensus by Melanoma Prevention Working Group + other international melanoma specialists (interdisciplinary including patient advocates).</p> <p>N: 1450 patients</p> <p>Intervention: GEP testing</p> <p>Outcome Measures: Melanoma recurrence Proportion of patients with melanoma recurrence accurately predicted by GEP Test.</p>	<p>Results:</p> <p>Among patients with recurrence Decision Dx correctly classified 29% and 82% with stage I and II disease respectively.</p> <p>Among patients with recurrence MelaGenix correctly classified 32% and 76% with stage I and II disease respectively.</p> <p>Without recurrence.</p> <p>Decision Dx 90% and 44% with stage I and II disease Respectively.</p> <p>MelaGenix 77% and 43% with stage I and II disease respectively.</p> <p>Before GEP testing is routinely used, the clinical benefit must be established through further clinical investigations.</p> <p>For Decisions Dx: In patients with recurrence, accurately predicted recurrence in 29% stage I and 82% stage II;</p>	<p>Describe conclusions relative to question:</p> <p>GEP testing poor in predicting recurrence in patients with Stage I disease.</p> <p>GEP assays appear to have poor predictive capacity for patients with stage I melanoma.</p> <p>Prognostic ability was felt to be poor for stage I disease for both GEPs evaluated.</p> <p>Prognostication was better in stage II disease</p> <p>Critiques of Methodology:  Heterogeneity in study</p>

				<p><i>Follow-Up:</i></p> <hr/> <p><i>Notes:</i> 7 studies (including 2 GEPs)</p>	<p>In patients without recurrence, accurately predicted this outcome in 90% and 44% for stage I and II respectively.</p> <p>For Melagenix: In patients with recurrence, accurately predicted recurrence in 32% and 76% for stage I and II respectively; In patients without recurrence, accurately predicted this outcome in 77% and 43% for stage I and II respectively.</p> <p>Primary endpoint was whether GEP correctly classified risk of recurrence.</p> <p>Overall, prognostic ability was felt to be poor for stage I disease for both GEPs evaluated.</p> <p>Among patients with recurrence, DecisionDx-Melanoma correctly classified 29% with stage I disease and 82% with stage II disease. Among patients without recurrence, the test correctly classified 90% with stage I disease and 44% with stage II disease.</p> <p>Among patients with recurrence, MelaGenix correctly classified 32% with stage I disease and 76% with stage II disease. Among patients</p>	<p>designs included and follow-up and reporting.</p> <p>Relatively short follow-up of several studies;</p> <p>Quality of evidence assessment of studies included could be subject to biases despite use of structured tools.</p> <p>Heterogeneous studies of varying quality and of varying length of patient f/u.</p> <p>Meta-analysis of HRs was not able to be done. Potential overlap of cohorts in studies.</p>
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				<div>RFS, DMFS DFS</div> <div>Follow-Up: 3.3 years</div> <div><hr/><i>Notes:</i></div>		<div>overstated. No SLNB results presented. No comparison with mitotic rate. Class 2B tumors – 50% ulcerated – unsurprising worse outcome. Survival data but no Cox regression analysis</div>
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		<p>2006-018702 (GenoMEL) and by the European Commission under the 7th Framework Programme, Diagnostix; The National Cancer Institute (NCI) of the US National Institute of Health (NIH) (CA83115), a grant from "Fundacio La Marató de TV3, 201331-30", Catalonia, Spain; CERCA Programme / Generalitat de Catalunya, and a grant from "Fundacion Científica de la Asociación Española Contra el Cáncer", Spain. Part of the work was carried out at the Esther Koplowitz Center, Barcelona</p> <p>The sponsors had no role in the design and conduct of the study, nor in the collection, analysis and interpretation</p>		<p>Recurrence Disease-free Recurrence</p> <p><i>Follow-Up:</i> 26 months : median 26 months (IQR 22-30 months) 26 months (median) based on "routine" surveillance protocols.</p> <hr/> <p><i>Notes:</i></p>	<hr/> <p><i>Notes:</i> 86/88 test successful.</p> <p>Type of relapse not stated.</p>	<p>Relatively short follow-up; Cohort limited to Spanish Caucasians;</p> <p>Excluded SLN+ patients;</p> <p>Table 2 – MVA Age is not significant in UV but included in MVA -is MVA valid?</p> <p>No power calculation</p> <p>Small patient numbers (7 patients developed metastases)—small numbers for robust multivariate analysis. Only 7/33 patients class II relapse (21.2%).</p> <p>The title suggests a prospective study, but this is a retrospective study (using prospectively captured data of patients who underwent GEP testing 2015-2016 and were followed for a median of 26 months).</p>
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		<p>of data, nor in the preparation, review and approval of the manuscript nor in the decision to submit the manuscript for publication.</p> <p>The authors declare that they have no COI.</p>				<p>Small cohort (does not appear to include consecutive patients—there may be some patient selection bias)</p>
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			clinical use trial, partnering with cooperative oncology groups to design adjuvant trials for node negative pt that include stratification by GEP class.		<p>Stratifying for T-stage and GEP found T-stage accounted for 43% of the decision on follow-up, GEP for 42%, and ulceration for 15%.</p> <hr/> <p><i>Notes:</i> Text of stage II, Class 2 unclear exactly what 64 v 36% is. Which is surgery + adj trial and what is the other category ie derm alone, surgery alone. Does not match exactly w Fig. 3 with appears to be 32% Surgical oncology.</p>	<p>Authors are employees of Castle Bioscience – not independent</p> <p>No follow-up therefore false-negative rate not available.</p>
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Summary of Evidence Table  
Question 1, 2, 3

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				<p>At least one year f/u or metastatic event; median f/u 3.2 years</p> <p>Median 3.2 years</p> <hr/> <p><i>Notes:</i></p>	<p>and correlate with 31-GEP class for entire 1124 pt cohort not the 684 used to develop nomogram.</p>	<p>No SLNB in 654 of the 684 pts No prospective validation cohort for nomogram – used an “archival cohort of 901 stage I-III patients w/ median 6.1 yrs f/u from 22 centers (no ref) to “validate” which is different from this study population.</p>
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				<p>Annual PET-CT + Brain MRI</p> <p><i>Outcome Measures:</i> Metastases detected by PETCT % with distant disease identified on screening PET-CT + brain MRI</p> <p><i>Follow-Up:</i> Within 3 years median 16 months</p> <hr/> <p><i>Notes:</i></p>		<p>potential selection biases.</p> <p>Incidence of metastases is fairly high in stage II in patients not undergoing Decision Dx; the majority of recurrences were in stage IIC patients (5/8) which would be expected without GEP Testing.</p> <p>Relatively short follow-up.</p> <p>Retrospective based on test receipt and imaging. Short f/u 16 months median. No comparator group. Rec overstated based on these data.</p>
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		<p>2 author: speakers bureau Castle Biosciences</p> <p>2 authors are a consultant for Castle Biosciences</p> <p>1 author: advisory board member Castle Bioscience</p>			<p>Notes:</p>	<p>Data only selectively presented</p> <p>The source of the patients is unclear. Presumably many have been previously published.</p> <p>Selective reporting of data. No crosstab of tumor characteristics and GEP-class.</p> <p>Details of this cohort are very poorly described.</p>
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Summary of Evidence Table Question 2						
Level of Evidence*^  Choose one:	Study Citation Author(s), Year, Title, Journal, Volume, Page #s	Funding Source	Study Methodology	Objective & Methods	Results	Conclusions
<input type="checkbox"/> EL 1; RCT <input type="checkbox"/> EL 1; MRCT  <input type="checkbox"/> EL 2; MNRCT <input type="checkbox"/> EL 2; NMA <input type="checkbox"/> EL 2; NRCT <input type="checkbox"/> EL 2; PCS <input checked="" type="checkbox"/> EL 2; RCCS <input type="checkbox"/> EL 2; NCCS <input type="checkbox"/> EL 2; CSS <input type="checkbox"/> EL 2; ES <input type="checkbox"/> EL 2; OLES <input type="checkbox"/> EL 2; PHAS  <input type="checkbox"/> EL 3; DS <input type="checkbox"/> EL 3; ECON <input type="checkbox"/> EL 3; CCS <input type="checkbox"/> EL 3; SCR <input type="checkbox"/> EL 3; PRECLIN <input type="checkbox"/> EL 3; BR  <input type="checkbox"/> EL 4; NE <input type="checkbox"/> EL 4: O	<p><b>Strong</b></p> <p>Zager JS, Gastman BR, Leachman S, et al. Performance of a Prognostic 31-Gene Expression Profile in an Independent Cohort of 523 Cutaneous Melanoma Patients. <i>BMC Cancer</i>. 2018;18(1):130. (36)</p> <p><i>Country:</i> USA</p>	<p><i>Source:</i></p> <p>This study was sponsored by Castle BioSciences, Inc., which provided funding for tissue and clinical data retrieval to contributing centers.</p> <p>10 authors: previously served as paid consultants to Castle Biosciences</p> <p>1 author: services as a paid clinical consultant to Castle Bioscience</p> <p>4 authors: Employees to Castle Bioscience</p> <p>4 authors: hold stock options for</p>	<p><i>Methodology:</i></p> <p>GEP classified primary melanoma tumors as Class 1 or Class 2 which was correlated to clinical outcome and assessed along w AJCC v7 staging. Multi-center.</p> <p>Multicentre study. Archival tissue sent for analysis.</p>	<p><i>Stated Objective:</i></p> <p>Predictive survival value of GEP for melanoma patients.</p> <p>To evaluate 31-GEP's prognostic accuracy in independent cohort of CM patients.</p> <p><input type="checkbox"/> Prospective <input checked="" type="checkbox"/> Retrospective</p> <p><i>Study Population and Setting:</i></p> <p>Primary cutaneous melanoma patients staged with SNB</p> <p>Multi-center (16) primary melanoma tumors</p> <p>Inclusion criteria: Biopsy confirmed stage I–III cutaneous melanoma diagnosed between 2000 and 2014, with at least five years of follow-up, unless there was an earlier documented recurrence or metastatic event.</p> <p><i>N:</i> 523</p> <p><i>Intervention:</i></p> <p>31-GEP GEP staging of primary SNB</p> <p><i>Outcome Measures:</i></p>	<p><i>Results:</i></p> <p>5-year RFS, DMFS and MSS for Class 1 was 88%, 93% and 98% respectively and Class 2 was 52%, 60% and 78% (p&lt;0.001).</p> <p>Stage I pt 5-year RFS class 1- 96%, Class 2- 85%. RFS class 1A- 98%, Class 2B- 73%. (p&lt;0.001).</p> <p>Stage I pt 5-year DMFS class 1- 97%, Class 2- 90%. RFS class 1A- 98%, Class 2B-87%. (p=0.05).</p> <p>Stage I pt 5-year MSS class 1A- 100%, Class 2B-93%. (p&lt;0.01).</p> <p>Stage II pt 5-year RFS class 1- 74%, Class 2- 55%. p=0.043. RFS class 1A- 77%, Class 2B-50%. (p=0.13).</p> <p>Stage I pt 5-year DMFS class 1- 90%, Class 2- 63%. RFS class 1A- 95%, Class 2B- 57%. (p&lt;0.001).</p> <p>Stage I pt 5-year MSS class 1A- 100%, Class 2B-82%. (p=.13).</p> <p>30/43 (70%) patients stage I and II with distant mets were Class 2.</p>	<p><i>Describe conclusions relative to question:</i></p> <p>31-GEP is an accurate predictor of metastatic risk.</p> <p>Class II GEP result is significant independent predictor of survival in multivariate analysis. However, SLNB status and Breslow thickness still significant. High Risk GEP seems to be replacing ulceration status and mitotic rate in primary risk factors. Predictor of low risk disease In SLN negative patients.</p> <p><i>Critiques of Methodology:</i></p> <p>Retrospective</p> <p>Size of cohort especially as</p>

		Castle Bioscience		<p>Recurrence-free (RFS) and distant metastasis-free (DMFS) survival. MSS-secondary endpoint. 5-year RFS and DMFS rates</p> <p><i>Follow-Up:</i> 5 years +, unless earlier documented recurrence.</p> <p>Minimum 5 years</p> <hr/> <p><i>Notes:</i></p>	<p>9/11 (82%) patients stage I and II who died were Class 2.</p> <p>Stage IIIA 5-year RFS class 1- 72%, Class 2- 51%, p=0.015. DMFS class 1- 80%, Class 2- 54%. (p0.019). MSS class 1- 100%, Class 2- 67%. (p0.009).</p> <p>In univariate analysis, DMFS Hazard ratio Class 1 was 5.4 and Class 2 was 6.6 p &lt;0.001. Breslow thickness, ulceration, mitotic rate and SLN status also significant.</p> <p>Multivariate model HR for RFS Breslow 1.2, p &lt;0.001; mitotic rate 0.9, NS; ulceration 1.4, NS; SLN positive 2.5, p&lt;0.001; GEP Class 2, 2.1, p=0.003.</p> <p>HR for DMFS Breslow 1.3, p &lt;0.001; mitotic rate 0.9, NS; ulceration 1.2, NS; SLN positive 3.0, p&lt;0.001; GEP Class 2, 2.7, p=0.002.</p> <p>SLNB status N=337 both SLN status and GEP. 5-yr RFS SLN-neg/ Class 1 87% SLN-neg/ Class 2 67%</p> <p>5-yr DMFS SLN-neg/ Class 1 93% SLN-neg/ Class 2 75%</p> <p>5-yr MSS</p>	<p>analyze smaller groups such as in stage or ulceration and fewer events</p> <p>Classification Breslow thickness <math>\leq</math> 1mm v &gt; 1 mm. Should look at continuous range or more categories ie 1-2, 2-3, &gt;4.</p> <p>Statistical methodology most appropriate that I have seen so far.</p>
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					<p>SLN-neg/ Class 1 98% SLN-neg/ Class 2 92%</p> <p>RFS/DMFS/MSS SLN-pos/ Class 1: 61%, 74%, 93%.</p> <p>SLN-pos/ Class 2: 37%, 44% and 63%.</p> <p>SLN-neg/Class 1A v SLN- neg/Class 2b 90% v 60%, 96% vs 69% and 100% vs 88%</p> <p>GEP v SLNB accuracy <i>sensitivity</i> R/DM/MSS Class 2 70%, 75%, and 85%, compared to SLNB- positivity 66%, 67% and 79%</p> <p>GEP v SLNB accuracy <i>specificity</i> R/DM/MS mortality Class 1 71%, 69%, and 64%, compared to SLNB- negativity 65%, 62% and 58%.</p> <p>Positive predictive value (PPV) of Class 2 and SLN- positivity were 48%, and 52% recurrence, 40% and 42% and 19% and 21% for MSS. The PPV of class 2B was 55% for recurrence, 45% for DM and 24% for MS mortality.</p> <p>NPV for Class 1 and SLN- negative were 87% and 76% for recurrence, 91% and 82% for DM and 98% and 95% for MS mortality. NPV Class 1A was 89% for recurrence, 94% for DM and 99% for MS mortality.</p>	
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					<p>Improved sensitivity with SLNB +GEP 88% for all recurrences, 91% distant mets.</p> <p>The 5-year RFS rates for Class 1 and Class 2 were 88% and 52%, respectively, and DMFS rates were 93% versus 60%, respectively (P &lt; 0.001). The GEP was a significant predictor of RFS and DMFS in univariate analysis (hazard ratio [HR] = 5.4 and 6.6, respectively, P &lt; 0.001 for each), along with Breslow thickness, ulceration, mitotic rate, and sentinel lymph node (SLN) status (P &lt; 0.001 for each). GEP, tumor thickness and SLN status were significant predictors of RFS and DMFS in a multivariate model that also included ulceration and mitotic rate (RFS HR = 2.1, 1.2, and 2.5, respectively, P &lt; 0.001 for each; and DMFS HR=2.7, 1.3 and 3.0, respectively, P &lt; 0.01 for each).</p> <p>Notes: Class IB and 2A not discussed much in paper; however, curves appear to overlap.</p>	
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					<div>SORT is Strength of Recommendation Taxonomy (A, B, or C)</div>	<div>(recommendations made as the National Society for Cutaneous Medicine)</div>
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### Question 3 + Future Research Directions

- 1) In adult patients with primary cutaneous melanoma, does GEP testing provide additional information and improve risk-stratification, beyond current diagnostic standards, to influence decisions for the utilization and the utility of adjuvant therapy?

#### Future Research Directions

What further research is needed to inform indications for GEP testing in the clinical care of patients with AJCC pT1a-pT4b (cNOM0) primary cutaneous melanoma?

Summary of Evidence Table Question 1, 2, 3						
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		<div>Bioscience investigator</div> <div>2 author: Castle Bioscience honoraria</div> <div>3 author: Castle Bioscience advisory board</div>		<div>Notes:</div>	<div>practice, expert opinion, or case series</div>	
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Summary of Evidence Table Question 1, 2, 3						
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## Summary of Evidence Table

### Question 1, 2, 3

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Choose one:						
<div><div><div><input type="checkbox"/>EL 1; RCT</div><div><input type="checkbox"/>EL 1; MRCT</div><div><input type="checkbox"/>EL 2; MNRCT</div><div><input type="checkbox"/>EL 2; NMA</div><div><input type="checkbox"/>EL 2; NRCT</div><div><input type="checkbox"/>EL 2; PCS</div><div><input type="checkbox"/>EL 2; RCCS</div><div><input type="checkbox"/>EL 2; NCCS</div><div><input type="checkbox"/>EL 2; CSS</div><div><input type="checkbox"/>EL 2; ES</div><div><input type="checkbox"/>EL 2; OLES</div><div><input type="checkbox"/>EL 2; PHAS</div><div><input type="checkbox"/>EL 3; DS</div><div><input type="checkbox"/>EL 3; ECON</div><div><input type="checkbox"/>EL 3; CCS</div><div><input type="checkbox"/>EL 3; SCR</div><div><input type="checkbox"/>EL 3; PRECLIN</div><div><input type="checkbox"/>EL 3; BR</div><div><input checked="" type="checkbox"/>EL 4; NE</div><div><input type="checkbox"/>EL 4; O</div></div><div><div>Strong</div><div>Intermediate</div><div>Weak</div><div>No Evidence</div></div></div>	<div>Farberg AS, Marson JW, Glazer A, et al. Expert Consensus on the Use of Prognostic Gene Expression Profiling Tests for the Management of Cutaneous Melanoma: Consensus from the Skin Cancer Prevention Working Group. <i>Dermatol Ther.</i> 2022;12(4):807-823. (81)</div> <div>Country: USA</div>	<div>Source: No funding or sponsorship was received for this study or publication of this article.</div> <div>3 authors=consultants for Castle Biosciences</div> <div>1 author=advisory board member and speaker for Castle Biosciences</div>	<div>Methodology: Modified Delphi technique for consensus statements.</div> <div>Review – no new data.</div> <div>Skin Cancer Prevention Working Group are authors.</div>	<div>Stated Objective: Assess applications for GEP in melanoma management.</div> <div><input type="checkbox"/>Prospective <input type="checkbox"/>Retrospective</div> <div>Study Population and Setting:</div> <div>N:</div> <div>Intervention:</div> <div>Outcome Measures:</div> <div>Follow-Up:</div> <div>Notes:</div>	<div>Results: Consensus statements support role for GEP above and beyond AJCC and NCCN.</div> <div>Notes: Consensus statements support role for GEP above and beyond AJCC and NCCN. The statements are non-specific.</div>	<div>Describe conclusions relative to question:</div> <div>Consensus statements support role for GEP.</div> <div>“GEP tests provide additional, reproducible information for dermatologists to consider within the larger framework of the eighth edition of the AJCC and NCCN cutaneous melanoma guidelines when counseling regarding prognosis and when considering a sentinel lymph node biopsy.”</div> <div>Critiques of Methodology: No formal accompanying systematic review/meta-analysis to support the consensus</div>

						<p>process (only 8 panelists).</p> <p>Review of existing data.</p>
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**Summary of Evidence Table**  
**Question 2,3**

Level of Evidence*^  Choose one:	Study Citation Author(s), Year, Title, Journal, Volume, Page #s	Funding Source	Study Methodology	Objective & Methods	Results	Conclusions
<div> <input type="checkbox"/> EL 1; RCT  <input type="checkbox"/> EL 1; MRCT    <input type="checkbox"/> EL 2; MNRCT  <input type="checkbox"/> EL 2; NMA  <input type="checkbox"/> EL 2; NRCT  <input type="checkbox"/> EL 2; PCS  <input type="checkbox"/> EL 2; RCCS  <input type="checkbox"/> EL 2; NCCS  <input type="checkbox"/> EL 2; CSS  <input type="checkbox"/> EL 2; ES  <input type="checkbox"/> EL 2; OLES  <input type="checkbox"/> EL 2; PHAS    <input type="checkbox"/> EL 3; DS  <input type="checkbox"/> EL 3; ECON  <input checked="" type="checkbox"/> EL 3; CCS  <input type="checkbox"/> EL 3; SCR  <input type="checkbox"/> EL 3; PRECLIN  <input type="checkbox"/> EL 3; BR    <input type="checkbox"/> EL 4; NE  <input type="checkbox"/> EL 4; O </div> <div> <div>Strong</div> <div>Intermediate</div> <div>Weak</div> <div>No Evidence</div> </div>	<p>Ferris LK, Farberg AS, Middlebrook B, et al. Identification of high-risk cutaneous melanoma tumors is improved when combining the online American Joint Committee on Cancer Individualized Melanoma Patient Outcome Prediction Tool with a 31-gene expression profile-based classification. <i>J Am Acad Dermatol.</i> May 2017;76(5):818-825.e3.</p> <p>Country: USA</p>	<p>Source: Funded by Castle Biosciences which provided financial compensation to those centers contributing cutaneous melanoma tissue to the study.</p> <p>6 authors: employees of Castle Biosciences</p> <p>4 authors: served as consultants to Castle Bioscience</p>	<p>Methodology: RNA isolated to run 31 gene test.</p> <p>Five-year survival for each patient using AJCC Individualized Melanoma Patient Outcome Prediction Tool</p> <p>Survival cutoff for low and high-risk determined by stage IIA (79%) and stage IIB (68%) cohorts.</p> <p>K-M analysis and Cox proportional hazards survival analysis performed.</p> <p>205 cases of stage I/II CM with available tissue were collected from six US centers.</p>	<p>Stated Objective: To compare accuracy of GEP with risk determined using the web-based AJCC Individualized Melanoma Patient Outcome Prediction Tool.</p> <p>Compare recurrence risk prediction by AJCC Stage vs. 31-GEP class.</p> <p><input type="checkbox"/> Prospective <input checked="" type="checkbox"/> Retrospective</p> <p>Study Population and Setting: Stage I/II CM. 205 retrospectively collected.</p> <p>N: 205</p> <p>Intervention: GEP and AJCC prediction tool GEP vs conventional staging</p> <p>Outcome Measures: RFS, DMFS, OS</p> <p>Follow-Up: 6.9 y f/u 6.9 years (0.1-15.4)</p> <p>Notes:</p>	<p>Results: 43 (21%) cases had discordant GEP and AJCC classification (with 79% cutoff). 11/43 discordant cases classified as high risk GEP but low by AJCC.</p> <p>Sensitivity GEP 82%, 81% and 78% for RFS, DMFS and MSS. AJCC 70%, 69% and 60%.</p> <p>Specificity GEP 77%, 69%, 69% and AJCC 83%, 76%, 74%</p> <p>Increased sensitivity in combining GEP + AJCC with decreased specificity.</p> <p>Multivariate cox regression GEP w AJCC tool indicated GEP more significantly associated with DM and death than binary AJCC.</p> <p>HR GEP vs AJCC 79% Recurrence- 5.9 v 3.6 DMFS- 5.3 v 3.0, Death -5.3 vs 2.2</p> <p>Use of GEP test with AJCC stage improved sensitivity to detect recurrence and to a lesser degree death at the cost of specificity compared with AJCC alone.</p>	<p>Describe conclusions relative to question:</p> <p>To combine GEP with AJCC may help identify pt benefit from increased surveillance and administer therapies.</p> <p>GEP can improve detection of recurrence/mets when used in combination with AJCC.</p> <p>Somewhat overstated based on limitations of dataset.</p> <p>Critiques of Methodology: Using AJCC tool as binary high-risk vs low-risk to fit comparison.</p> <p>Unconventional and narrow to define sensitivity and specificity of test using definition of</p>



					<div>Notes:</div>	<div>low-risk group from stage IIA AJCC 79% survival and high-risk 68% survival based off of stage IIB.</div> <div>These patients were previously published in the development of the model. Unclear how this subset was selected. Stage was dichotomized at IIA- vs IIB+ somewhat arbitrarily.</div>
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		<p>with resources and use of facilities at the Veterans Affairs Palo Alto Health Care System in Palo Alto CA. Contents do not represent the views of the US Dept of Veterans Affairs or the US Gov't. The funders had no role in the design and conduct of the study; collection, management, analysis and interpretation of data, preparation, review or approval of the manuscript; and decision to submit for publication.</p> <p>1 author: nonfinancial support from Castle Biosciences outside submitted work.</p> <p>1 author: grants from Castle Biosciences outside</p>		<p>Follow-Up:</p> <hr/> <p>Notes:</p>	<p>with SLNB and adjuvant therapy. The MPWG members favor conducting retrospective studies that evaluate multiple GEP testing platforms on fully annotated archived samples before embarking on costly prospective studies and recommend avoiding routine use of GEP testing to direct patient management until prospective studies support their clinical utility.</p> <hr/> <p>Notes:</p>	
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		<p>submitted work</p> <p>1 author: personal fees outside the submitted work Castle Biosciences</p> <p>1 author: personal fees from Neracare outside submitted work</p> <p>1 author: manuscripts and abstracts published using the test with company support of the assay—all publications were peer review and no personal of institutional payment or compensation was rec'd. Castle Biosciences</p> <p>1 author: served as an investigator for Castle Biosciences (no personal financial compensation)</p>				
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<input type="checkbox"/> EL 1; RCT <input type="checkbox"/> EL 1; MRCT  <input type="checkbox"/> EL 2; MNRCT <input type="checkbox"/> EL 2; NMA <input type="checkbox"/> EL 2; NRCT <input type="checkbox"/> EL 2; PCS <input type="checkbox"/> EL 2; RCCS <input type="checkbox"/> EL 2; NCCS <input type="checkbox"/> EL 2; CSS <input type="checkbox"/> EL 2; ES <input type="checkbox"/> EL 2; OLES <input type="checkbox"/> EL 2; PHAS  <input type="checkbox"/> EL 3; DS <input type="checkbox"/> EL 3; ECON <input type="checkbox"/> EL 3; CCS <input type="checkbox"/> EL 3; SCR <input type="checkbox"/> EL 3; PRECLIN <input type="checkbox"/> EL 3; BR  <input checked="" type="checkbox"/> EL 4; NE <input type="checkbox"/> EL 4; O	<p><b>Strong</b></p> <p>Svoboda RM, Glazer AM, Farberg AS, Rigel DS. Factors Affecting Dermatologists' Use of a 31-Gene Expression Profiling Test as an Adjunct for Predicting Metastatic Risk in Cutaneous Melanoma. J. Drugs Dermatol. 2018;17(5):544-547. (35)</p> <p><b>Intermediate</b></p> <p><i>Country:</i> USA</p> <p><b>Weak</b></p> <p><b>No Evidence</b></p>	<p><i>Source:</i></p> <p>This study was funded in part by a grant from Castle Biosciences Inc.</p> <p>1 author: consultant to Castle Bioscience</p> <p>1 author: served on advisory board for Castle Bioscience</p> <p>2 authors: participated in research fellowship that was partially funded by Castle Bioscience</p>	<p><i>Methodology:</i></p> <p>Dermatologists answered questionnaire with four clinical vignettes to determine the impact of Breslow thickness, ulceration and SLNB status decision to order GEP test.</p> <p>Survey-based study performed at a conference.</p>	<p><i>Stated Objective:</i> Clinical factors that impact dermatologists' decisions to utilize 31-GEP.</p> <p><input checked="" type="checkbox"/> Prospective <input type="checkbox"/> Retrospective</p> <p><i>Study Population and Setting:</i> Dermatology questionnaire at a national conference.</p> <p>Panel of clinicians. Survey</p> <p><i>N:</i> 181/187</p> <p><i>Intervention:</i> Questionnaire with four clinical vignettes.</p> <p>Opinion regarding management</p> <p><i>Outcome Measures:</i> Percentage of respondents who would order 31-GEP in clinical scenarios.</p> <p>Recommended treatment pathway.</p> <p><i>Follow-Up:</i> N/A</p> <p><i>Notes:</i></p>	<p><i>Results:</i> A majority of patients would recommend GEP test."</p> <p>Breslow thickness <math>\geq 0.5</math> mm, majority dermatologists would order GEP. Ulceration was associated with a statistically significant increase to recommend for all but the thickest <math>\geq 2.1</math> mm. For thin tumor (0.26 mm) ulceration significantly changed from 22% to 67%, <math>p &lt; 0.001</math>). A negative SLNBx only associated with statistically significant increase in the percentage for the thinnest tumors (22% to 34%, <math>p = 0.033</math>).</p> <p>Impact of GEP test result on T1b 0.76-1.0 mm melanoma: Class 1 result -91% respondents reported less likely to recommend SLNB. Class 2 - 81% would make more likely to recommend SLNB.</p> <p><i>Notes:</i></p>	<p><i>Describe conclusions relative to question:</i> Not relevant to current clinical practice.</p> <p>Ulceration most important factor deciding to order 31-GEP.</p> <p><i>Critiques of Methodology:</i> Authors do not state whether the survey was taken at a session paid for by Castle Bioscience.</p> <p>Authors have conflict of interest – was the survey voluntary</p> <p>-Survey of dermatologists' ordering pattern does not evaluate the validity of the test. - Selection bias in those willing to take the survey</p>

							<div>-was survey voluntary or was there an incentive.</div>
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