

Real-world evidence confirms risk stratification of the 31-GEP and i31-GEP in prospectively tested patients with stage I-III cutaneous melanoma

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Background

- Current American Joint Committee on Cancer (AJCC 8th edition) guidelines bin patients with cutaneous melanoma (CM) into risk categories based on the pathological tumor data of Breslow thickness, ulceration status, and sentinel lymph node status.¹ However, this does not consider the heterogenous nature of CM or the tumor's molecular biology.
- The 31-gene expression profile test is prospectively validated to identify patients considered high or low risk by AJCC with low or high-risk tumor biology who may be over- or undertreated by current guidelines and has been shown to have a positive impact on patient outcomes.²⁻⁷
- To further advance personalized patient care, the 31-GEP result was integrated with clinical and pathological factors (i31-GEP for risk of recurrence, ROR) to provide a more personalized, precise risk of tumor recurrence to guide clinical management of CM.⁸

Objective

- Validate the 31-GEP and i31-GEP ROR in a prospectively tested, real world cohort of patients with stage I-III CM to demonstrate added value of 31-GEP testing in clinical care.

Methods

Patients with stage I-III CM enrolled in the CONNECTION study were prospectively tested with the 31-GEP between 2013 and 2017 (n=1,831). Kaplan-Meier analysis with the log-rank test was used to estimate survival differences between low (Class 1A), intermediate (Class 1B/2A), and high (Class 2) risk groups and the i31-GEP risk groups. The i31-GEP ROR combines Breslow thickness, ulceration, SLN status, mitotic rate, tumor location, age, and the 31-GEP to provide a personalized estimate of recurrence-free survival (RFS).⁸ While guidelines have not established an ROR threshold for determining when to escalate or de-escalate care, the NCCN uses stage IIA versus IIB to differentiate management intensity.⁹ This cut-point translates to a 5-year RFS rate of 69.8% and was used here. Cox multivariable regression analysis was used to identify predictors of recurrence.

References

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Results

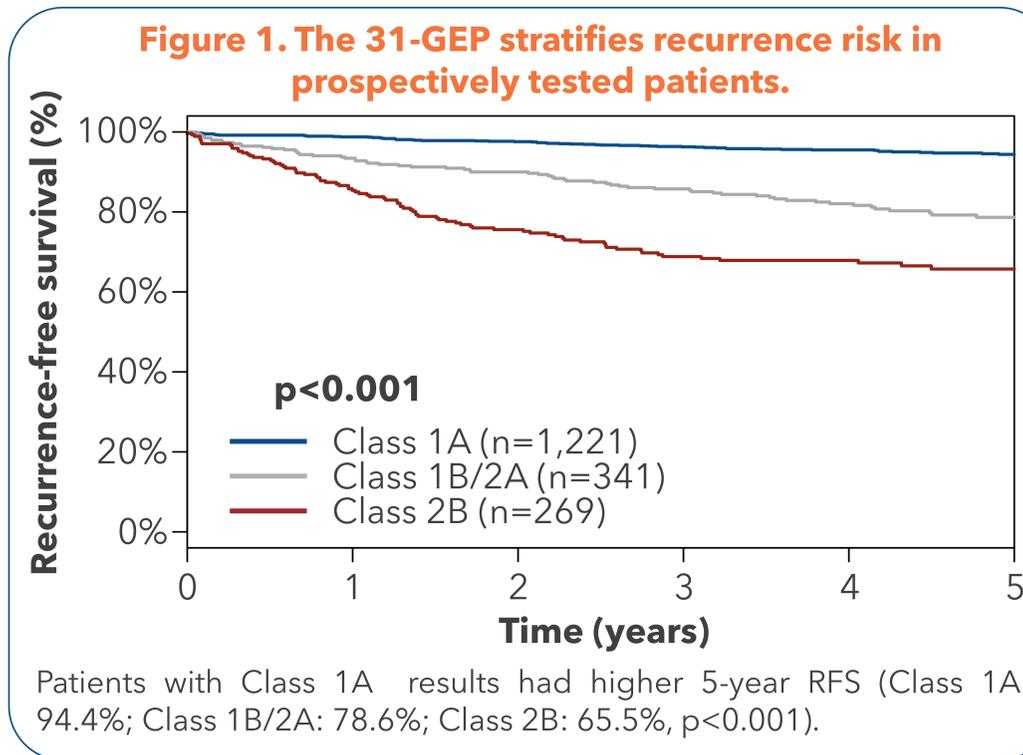
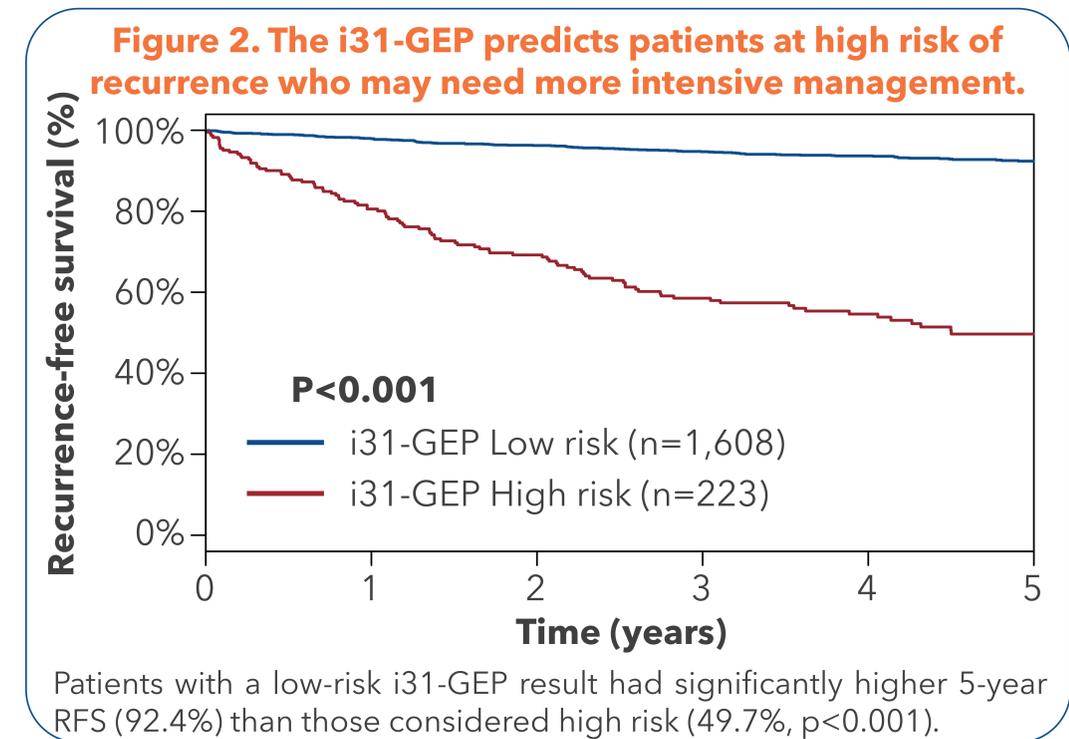


Table 1. Multivariable analysis demonstrates independent and significant prognostic information

| Recurrence-free survival | Multivariable HR | p-value |
|---------------------------------|------------------|---------|
| 31-GEP Class 1A | Reference | -- |
| 31-GEP Class 1B/2A | 2.07 | <0.001 |
| 31-GEP Class 2B | 2.40 | <0.001 |
| Age (continuous) | 1.02 | <0.001 |
| SLN negative | Reference | -- |
| SLN unknown | 0.71 | 0.085 |
| SLN positive | 4.54 | <0.001 |
| Breslow thickness (continuous) | 1.09 | 0.027 |
| Ulceration absent | Reference | -- |
| Ulceration present | 1.57 | 0.005 |
| Mitotic rate <2/mm ² | Reference | -- |
| Mitotic rate ≥2/mm ² | 1.47 | 0.016 |



Clinical Impact

- Identifying high-risk patients allows for earlier management decisions while patients have a lower tumor burden.
- Baseline high-risk patient detection may promote earlier administration of immunotherapy and potentially better therapeutic outcomes

Conclusions

- This real-world evidence study of prospectively tested patients confirms the independent performance of both the 31-GEP and the i31-GEP in improving treatment pathway decision accuracy.

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