Using a **CLINICOPATHOLOGIC** and **GENE EXPRESSION PROFILE** (CP-GEP) model to predict PROGNOSIS in STAGE I-II MELANOMA

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To validate the performance of the CP-GEP model in predicting prognosis in stage I-II melanoma

AIM

BACKGROUND

EARLY-STAGE MELANOMA – A CLINICAL CHALLENGE¹⁻³





- >80% is stage I-II (without metastasis)
- Stage I-II present notable heterogenous survival outcomes - Depending on factors beyond stages?

ADJUVANT IMMUNOTHERAPY - DILEMMA^{4,5}

- Improved RFS in stage II substages (phase III trial)
- Risk of severe adverse effects
- Potential financial strain on healthcare systems

REFINED risk stratification of stage I-II is NEEDED to TAILOR treatment and surveillance **BUT HOW ?**

The CP-GEP model

Developed and validated to **PREDICT** risk of **SENTINEL NODE METASTASIS**⁶⁻¹⁰



CAN CP-GEP PREDICT RISK OF RECURRENCE AND DEATH?

	RFS				05				MSS			
_	5-year		10-year		5-year		10-year		5-year		10-year	
_	%	(95% CI)	%	(95% CI)	%	(95% CI)	%	(95% CI)	0/ /0	(95% CI)	%	(95% CI)
Total (N=438)	87	(84-90)	79	(75-83)	89	(86-92)	80	(76-84)	96	(93-97)	95	(92-97)
CP-GEP High (N=239)	83	(77-87)	75	(69-81)	87	(82-90)	76	(70-81)	93	(89-96)	92	(87-95)
CP-GEP Low (N=199)	92	(87-95)	84	(78-89)	93	(88-95)	85	(79-89)	98	(95-100)	98	(95-100)

Box 3 Kaplan-Meier curves, Hazard ratios and 5-year and 10-year RFS, OS and MSS at a median follow-up of 115 months, stratified by CP-GEP result (High or Low risk).

METHOD



CP-GEP may have potential as a

CONCLUSION

PROGNOSTIC TOOL for

EARLY-STAGE MELANOMA

prognostication TO BE CONTINUED...

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