

Correlation of Breast Cancer Index (BCI) Risk Classification with Tumor Grade and Ki-67 in a Large Series of Patients with Early-stage, ER+ Breast Cancer

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INTRODUCTION

- Breast Cancer Index (BCI) is a gene expression-based assay that reports both prognostic and predictive results for patients with early-stage, ER+ breast cancer.¹⁻³
- BCI Predictive is based on the HoxB13/IL17BR (H/I) ratio, an endocrine response biomarker, and has been demonstrated to predict likelihood of benefit from extended endocrine therapy.³
- BCI Prognostic is based on the algorithmic combination of H/I and a set of proliferation-based genes (molecular grade index, MGI), and has been shown to significantly stratify patients based on risk of overall (0-10y) and specifically late (post-5 year) recurrence.^{1,2,4}
- MGI is comprised of 5 cell cycle genes (BUB1B, CENPA, NEK2, RACGAP1, and RRM2) and recapitulates tumor grade.⁴
- In this study, correlative analyses of risk stratification by BCI Prognostic and MGI vs tumor grade and Ki67 were performed to characterize their relationship to other markers of proliferation.

MATERIALS & METHODS

- Descriptive analyses were performed using data from the BCI Clinical Correlative Database, which contains demographic information, clinical and pathologic characteristics, and test results for cases submitted for BCI testing in clinical practice.
- Tumor grade and Ki67 information was abstracted from pathology reports; data were available for 1335 and 372 patients with node-negative breast cancer, respectively.
- Ki67 categories (low, intermediate, and high) were based on 10% and 20% IHC expression cutoffs.
- BCI risk categories (low, intermediate, and high) and MGI risk categories (low and high) were based on established clinical cutpoints (5.0825 and 6.5025 for BCI, 0.0 for MGI).
- Pearson correlation was performed to compare BCI Prognostic and MGI to Ki67 as continuous variables; coefficient of determination derived from the analysis of variance (ANOVA) model was used to compare continuous BCI Prognostic and MGI to categorical tumor grade.
- Chi-square test assessed the significance of concordance between BCI Prognostic and MGI risk groups to tumor grade and Ki67 groups.
- Restratification by BCI and MGI were performed within tumor grade and Ki67 expression categories.

RESULTS

Table 1. Patient Characteristics

Total Number of Cases	N=1359	
Age, mean/median	57y / 58y	
ER status (N=905)	Positive	894 (99%)
	Negative	11 (1%)
PR status (N=899)	Positive	801 (89%)
	Negative	94 (10%)
	Borderline	4 (1%)
Her2 status (N=858)	Negative	774 (90%)
	Positive	84 (10%)
Nodal status (N=1359)	Negative	1359 (100%)
	Positive	0 (0%)
Tumor size (N=1347)	≤1 cm	356 (26%)
	>1-≤2 cm	656 (49%)
	>2-≤5 cm	308 (23%)
	>5 cm	27 (2%)
Tumor grade (N=1335)	1	389 (29%)
	2	711 (53%)
	3	235 (18%)
Ki67 (N=372)	Low (≤ 10%)	137 (37%)
	Intermediate (11-20%)	91 (24%)
	High (>20%)	145 (39%)

- Mean age at diagnosis was 57 years.
- The majority of tumors were T1 (75%) and HER2- (90%).
- 99% were ER+, and 89% were PR+.
- Both tumor grade and Ki67 were distributed across low, intermediate, and high classifications.

RESULTS

Table 2. BCI Prognostic Risk Classification vs Grade (N=1335)

	Grade 1 (N=389)	Grade 2 (N=711)	Grade 3 (N=235)
BCI Low Risk (N=712)	305	365	42
BCI Intermediate Risk (N=341)	69	216	56
BCI High Risk (N=282)	15	130	137

Table 3. BCI Prognostic Risk Classification vs Ki67 (N=372)

	Ki67 Low (N=137)	Ki67 Inter (N=91)	Ki67 High (N=145)
BCI Low Risk (N=196)	109	47	40
BCI Intermediate Risk (N=102)	21	36	45
BCI High Risk (N=74)	6	8	60

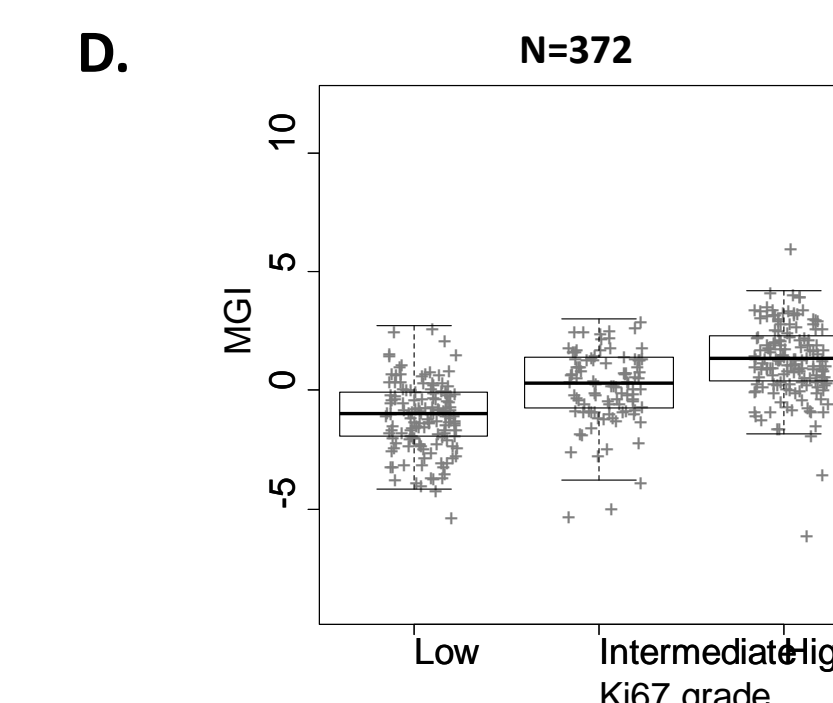
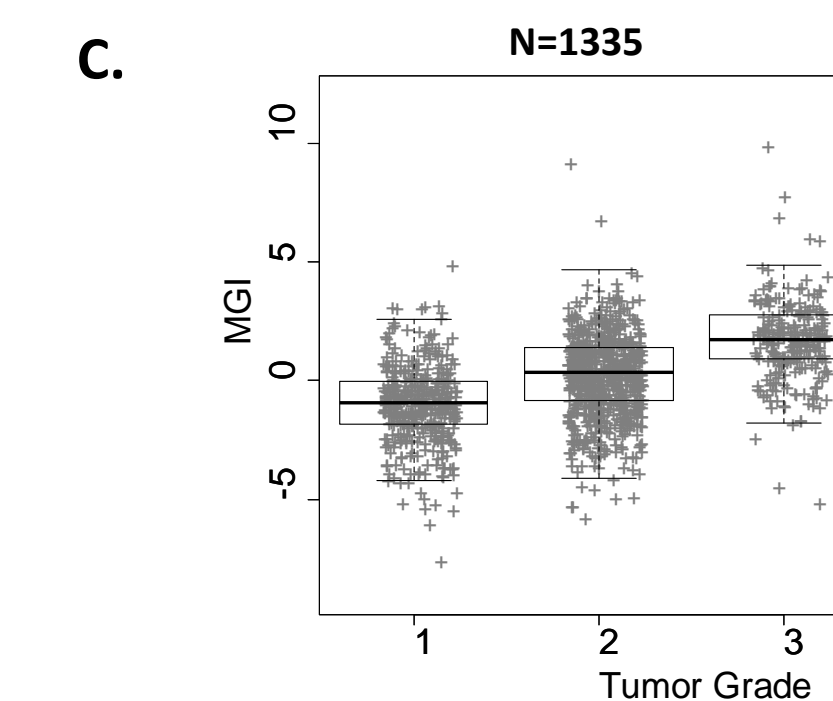
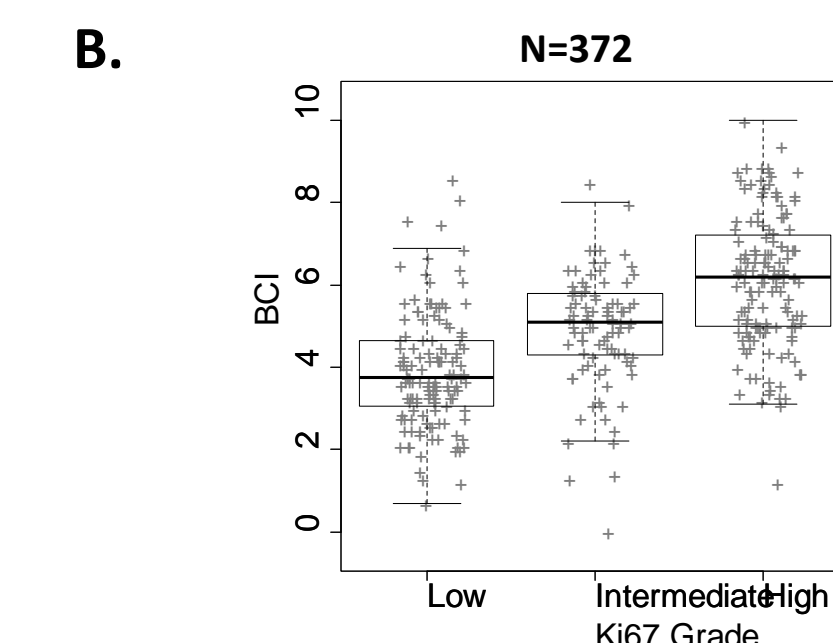
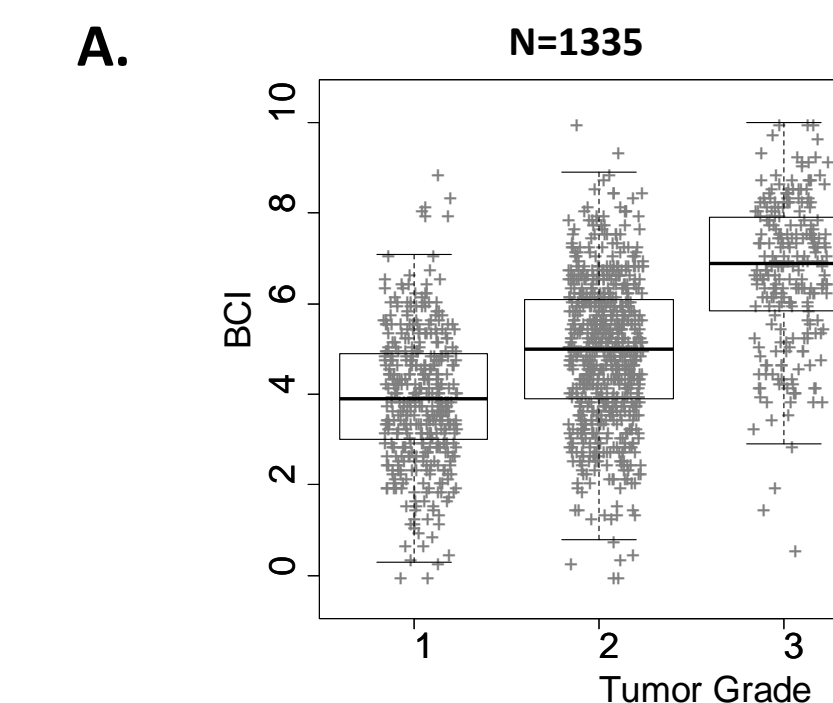
Table 4. MGI Risk Classification vs Grade (N=1335)

	Grade 1 (N=389)	Grade 2 (N=711)	Grade 3 (N=235)
MGI Low Risk (N=624)	294	295	35
MGI High Risk (N=711)	95	416	200

Table 5. MGI Risk Classification vs Ki67 (N=372)

	Ki67 Low (N=137)	Ki67 Inter (N=91)	Ki67 High (N=145)
MGI Low Risk (N=165)	104	40	21
MGI High Risk (N=207)	32	51	124

Figure 1. Correlation of BCI/MGI vs Grade/Ki67



RESULTS

- BCI Prognostic and MGI (as continuous variables) correlated weakly with tumor grade (coefficient of determination = 0.26 and 0.22, respectively) and continuous Ki67 ($r^2 = 0.35$ and 0.33 , respectively).
- Discordance between BCI Prognostic and both tumor grade and Ki67 were statistically significant ($p < 0.00001$ and $p < 0.00001$, respectively).
- BCI Prognostic classified 4% of Grade 1 tumors as high risk and 18% of Grade 3 tumors as low risk. In addition, BCI classified 51% of Grade 2 tumors as low risk and 18% as high risk (Table 2).
- BCI Prognostic classified 4% of low Ki67 patients as high risk and 28% of high Ki67 patients as low risk. In addition, BCI classified 52% of Ki67 Intermediate as low risk and 9% as high risk (Table 3).
- Discordance between MGI and both tumor grade and Ki67 were significant ($p < 0.00001$ and $p < 0.00001$, respectively).
- MGI classified 24% of Grade 1 tumors as high risk and 15% of Grade 3 tumors as low risk. In addition, MGI classified 41% of Grade 2 grade tumors as low risk and 59% as high risk (Table 4).
- MGI classified 24% of low Ki67 patients as high risk and 14% of high Ki67 patients as low risk. In addition, MGI classified 44% of Ki67 intermediate patients as low risk and 56% as high risk (Table 5).

CONCLUSIONS

- Data from this large retrospective analysis indicate that BCI Prognostic and MGI are capturing distinct information related to tumor proliferative status compared with tumor grade and Ki67.
- The algorithmic combination of the mitogenic (MGI) and endocrine response (H/I) gene signatures within BCI Prognostic has been demonstrated to be a superior prognostic factor to clinicopathologic variables in previous multivariate analyses.¹

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Intended Uses and Limitations The Breast Cancer Index (BCI) Risk of Recurrence & Extended Endocrine Benefit Test is intended for use in patients diagnosed with estrogen receptor-positive (ER+), lymph node-negative (LN-) or lymph node positive (LN+; with 1-3 positive nodes) early-stage, invasive breast cancer, who are distant recurrence-free. BCI provides: 1) a quantitative assessment of the likelihood of both late (post-5 years) and overall (0-10 year) distant recurrence following an initial 5 years of endocrine therapy (LN- patients) or 5 years of endocrine therapy plus adjuvant chemotherapy (LN+ patients), and 2) prediction of likelihood of benefit from extended (>5 year) endocrine therapy. BCI results are adjunctive to the ordering physician's workup; treatment decisions require correlation with all other clinical findings. This test was developed and its performance characteristics determined by Biotheranostics, Inc. It has not been cleared or approved by the U.S. Food and Drug Administration. This test is used for clinical purposes. It should not be regarded as investigational or for research. How this information is used to guide patient care is the responsibility of the physician. Biotheranostics is certified under the Clinical Laboratory Improvement Amendments of 1988 to perform high complexity clinical laboratory testing.

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Highlights:

- In this study, correlative analyses of risk stratification by BCI Prognostic and Molecular Grade Index (MGI; 5 genes making up the proliferative component of BCI) vs tumor grade and Ki67 were performed to characterize their relationship to other markers of proliferation.
- Discordance between both BCI Prognostic and MGI to tumor grade and Ki67 were statistically significant.

	Grade 1	Grade 2	Grade 3	Ki67 Low	Ki67 Intermediate	Ki 67 High
Reclassification by BCI	4% BCI High	51% BCI Low 18% BCI High	18% BCI Low	4% BCI High	52% BCI Low 9% BCI High	28% BCI Low
Reclassification by MGI	24% MGI High	41% MGI Low 59% MGI High	15% MGI Low	24% MGI High	44% MGI Low 56% MGI High	14% MGI Low

- Data from this large retrospective analysis indicate that BCI Prognostic and MGI are capturing distinct information related to tumor proliferative status compared with tumor grade and Ki67.
- The algorithmic combination of the mitogenic (MGI) and endocrine response (H/I) gene signatures within BCI Prognostic has been demonstrated to be a superior prognostic factor to clinicopathologic variables in previous multivariate analyses.