

Biologic factors of breast cancer and DFS

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Introduction

Novel molecular characterization of breast cancer with cellular markers has allowed a new classification that offers prognostic value, with predictive categories of disease aggressiveness.(1) Biological factors have predictive and prognostic value in breast cancer patients.(2)

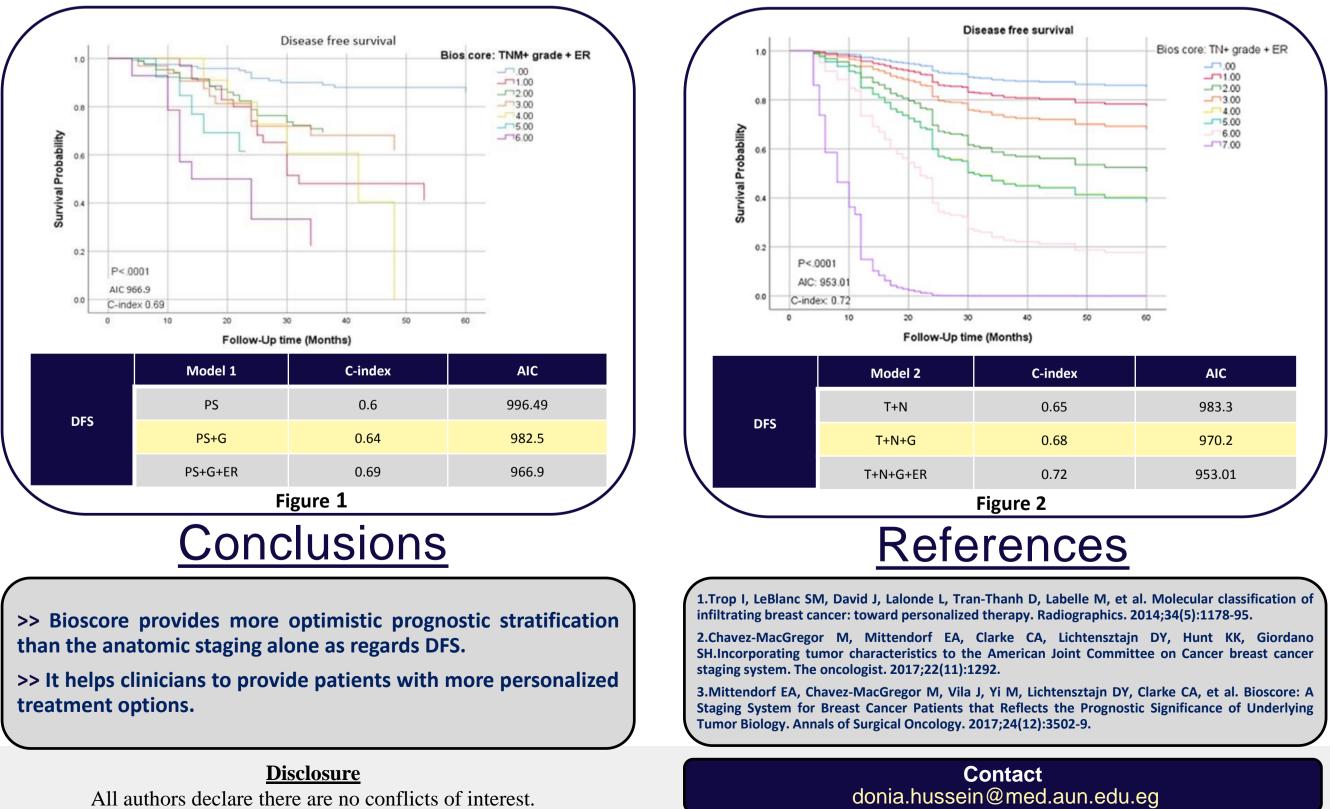
Bioscore staging model was proposed by MD Anderson team. In this model incorporation of biologic factors into AJCC staging system had a prognostic impact on patients survival.(3)

Our work evaluates the prognostic value of the bioscore among non-metastatic female breast cancer patients concerning disease-free survival (DFS).

- Significant factors in the first multivariate model: PS3, G3, and ER -ve.

Methods

- ◆ We reviewed the clinical data of 317 female patients with non-metastatic surgically treated breast cancer from January 2015 to December 2018 presented at Clinical Oncology Department, Assiut University Hospital, Egypt.
- ✤ The biological variables include; pathologic stage (PS), T stage (T), nodal stage (N), grade (G), estrogen receptor (ER), progesterone receptors (PR), and human epidermal growth factor receptor (HER2) status.
- ◆ Univariate & two multivariate analyses were performed to identify variables associated with disease-free survival (DFS).
- ✤ Bioscore calculation by giving a score of 0-4 for each factor according to the hazard ratio magnitude.
- Multiple staging system models were built for significant factors in both univariate and multivariate analyses: PS, PS + G, PS + G + E, T + N, T + N + G, T + N + G + E.
- ◆ The first one used the PS, which takes into account T and N stage as a variable while the second included T and N stages as separate variables.
- Model performance was quantified using Harrell's concordance index (C-index) and the Akaike Information Criterion (AIC) was used to compare model fits.



Results

Significant factors in the univariate analysis: PS3, T2, T3, T4, N3, G2, G3, ER -ve, PR -ve, and HER2 -ve with Hazard Ratio (HR); 4.77, 2.52, 2.80, 5.59, 2.74, 6.92, 16.80, 3.08, 2.11, 0.53 respectively with significant P value (< 0.05).

Significant factors in the second multivariate model: T2, T4, N3, G3, and ER -ve.

Two sets of models were built to determine the utility of combining variables.

Models incorporating G and E status had the highest C-index (0.72) for (T+N + G + ER) in comparison with (0.69)

for (PS+ G + ER) and the lowest AIC (953.01) for (T + N + G + E) and (966.9) for (PS + G + E) as in fig.1,2.

