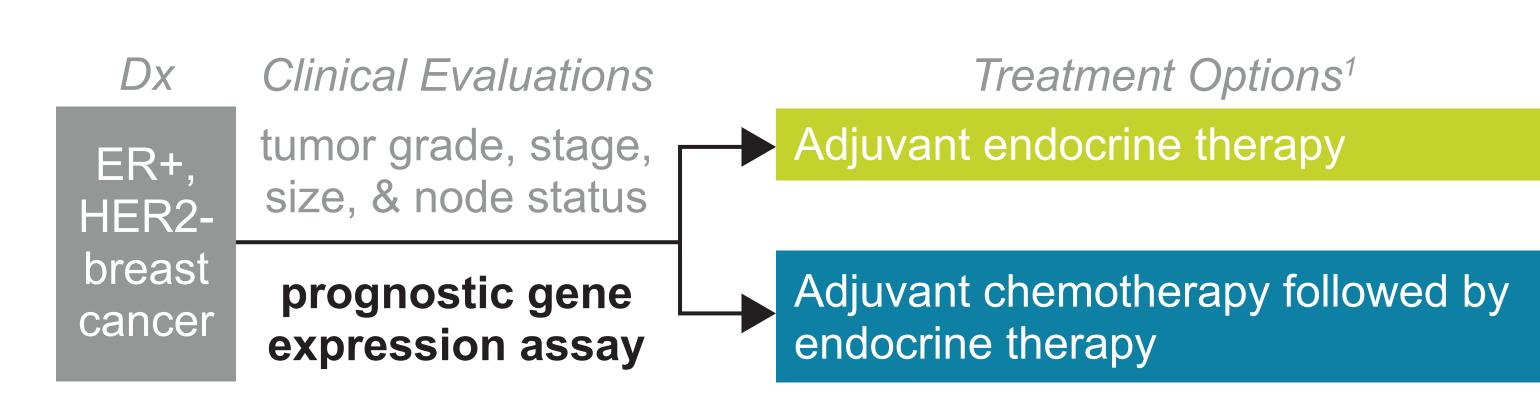
Predicting Expected Absolute Chemotherapy Treatment Benefit in Women with Early-Stage Breast Cancer using a 12-Gene Expression Assay

William Gradishar, MD¹, Mark Robson, MD², Darl D. Flake II, PhD³, Lee Schwartzberg, MD⁴, Priyanka Sharma, MD⁵, Anthony Magliocco, MD⁶, Krystal Brown, PhD³, Alexander Gutin, PhD³, Alexander Gutin, PhD³, Ralf Kronenwett, MD, PhD¬, Johnathan Lancaster, MD, PhD³

1. Feinberg School of Medicine, Northwestern University, Chicago, IL 2. Memorial Sloan Kettering Cancer Center, Memphis, TN 5. University of Kansas City, KS 6. Moffitt Cancer Center, Tampa, FL 7. Myriad International GmBH, Cologne, Germany 1. Feinberg School of Medicine, Northwestern University of Kansas Medical Center, Kansas City, KS 6. Moffitt Cancer Center, Tampa, FL 7. Myriad International GmBH, Cologne, Germany 1. Feinberg School of Medicine, Northwestern University of Kansas Medical Center, Northwestern University of Kansas City, KS 6. Moffitt Cancer Center, Northwestern University of Kansas Medical Center, Northwestern University of Kansas City, KS 6. Moffitt Cancer Center, Northwestern University of Kansas City, KS 6. Moffitt Cancer Center, Northwestern University of Kansas City, KS 6. Moffitt Cancer Center, Northwestern University of Kansas City, KS 6. Moffitt Cancer Center, Northwestern University of Kansas City, KS 6. Moffitt Cancer Center, Northwestern University of Kansas City, KS 6. Moffitt Cancer Center, Northwestern University of Kansas City, KS 6. Moffitt Cancer Center, Northwestern University of Kansas City, KS 6. Moffitt Cancer Center, Northwestern University of Kansas City, KS 6. Moffitt Cancer Center, Northwestern University of Kansas City, KS 6. Moffitt Cancer Center, Northwestern University of Kansas City, KS 6. Moffitt Cancer Center, Northwestern University of Kansas City, KS 6. Moffitt Cancer Center, Northwestern University On Cancer Center, Northwes

BACKGROUND

- Many women with estrogen receptor-positive (ER+), human epidermal growth factor receptor 2-negative (HER2-) breast cancer who have a low risk of distant recurrence can safely forgo adjuvant chemotherapy.1
- Although risk of recurrence has traditionally been evaluated using clinical factors, a prognostic 12-gene expression assay has been previously validated as a superior predictor of the risk of distant recurrence in this population.²⁻⁵



 Here, we estimate the expected absolute chemotherapy benefit for women with early-stage ER+, HER2- breast cancer based on the 12-gene expression assay using a mathematical model.

METHODS

COHORT

 This cohort included treatment-naïve patients with ER+, HER2- breast cancer who were tested with the 12-gene expression assay between October 2014 and December 2017 (Myriad Genetic Laboratories or Myriad GmbH) as part of normal clinical operations (N=2,205).

12-GENE EXPRESSION ASSAY

 The RNA expression of 12 genes was measured by qRT-PCR in formalin-fixed paraffin-embedded breast resections.



Hormone Receptor Normalization Marker of DNA Target Genes

Contamination

- A 12-gene molecular score was calculated as the linear combination of the normalized target gene expression.
- A molecular-clinical score (EPclin score) was calculated by combining the 12-gene molecular score with tumor size and the number of positive lymph nodes.
 - EPclin <3.3: low risk for distant recurrence
 - EPclin ≥3.3: high risk for distant recurrence

STATISTICAL METHODS

1. Risk of Distant Recurrence Without Chemotherapy

• The 10-year risk of distant recurrence in patients treated with endocrine therapy only ('untreated') was obtained from published results of the 12-gene expression assay in the ABCSG-6 and ABCSG-8 trials.²

2. Risk of Distant Recurrence With Chemotherapy

• The 10-year risk of distant recurrence in patients treated with endocrine therapy and adjuvant chemotherapy ('treated') was modeled using the overall chemotherapy treatment benefit to reduce recurrence in ER+, HER2breast cancer from a meta-analysis by the EBCTCG $(HR_{overall} = 0.7).6,7$

$$r_{\text{treated}} = 1 - (1 - r_{\text{untreated}})^{\text{HR}_{\text{treatment}}}$$

$$\log(\text{HR}_{\text{treatment}}) = \alpha - \beta(\text{EPclin} - 1)$$

Risk of 10-year distant metastasis in untreated patients from Part 1

Risk of 10-year distant metastasis in treated patients HR_{treatment} Relative chemotherapy benefit for an EPclin score

Strength of interaction between HR_{treatment} and EPclin Patient EPclin score

- A) Risk of recurrence modeled when there is no interaction between EPclin score and HR_{treatment}
 - $-\beta = 0 (\beta_{\min})$
 - α calculated such that $HR_{overall} = 0.7$

B) Risk of recurrence modeled for the maximum interaction between EPclin score and HR_{treatment}

- $-\alpha=0$
- β calculated such that HR_{overall} = 0.7 (β _{max})
- C) Risk of recurrence modeled when the interaction between EPclin score and HR_{treatment} is varied
 - β varied from min (Part 2A) to max (Part 2B)
 - α calculated such that HR_{overall} = 0.7

3. Absolute Benefit from Chemotherapy

Absolute Benefit = $r_{untreated} - r_{treated}$

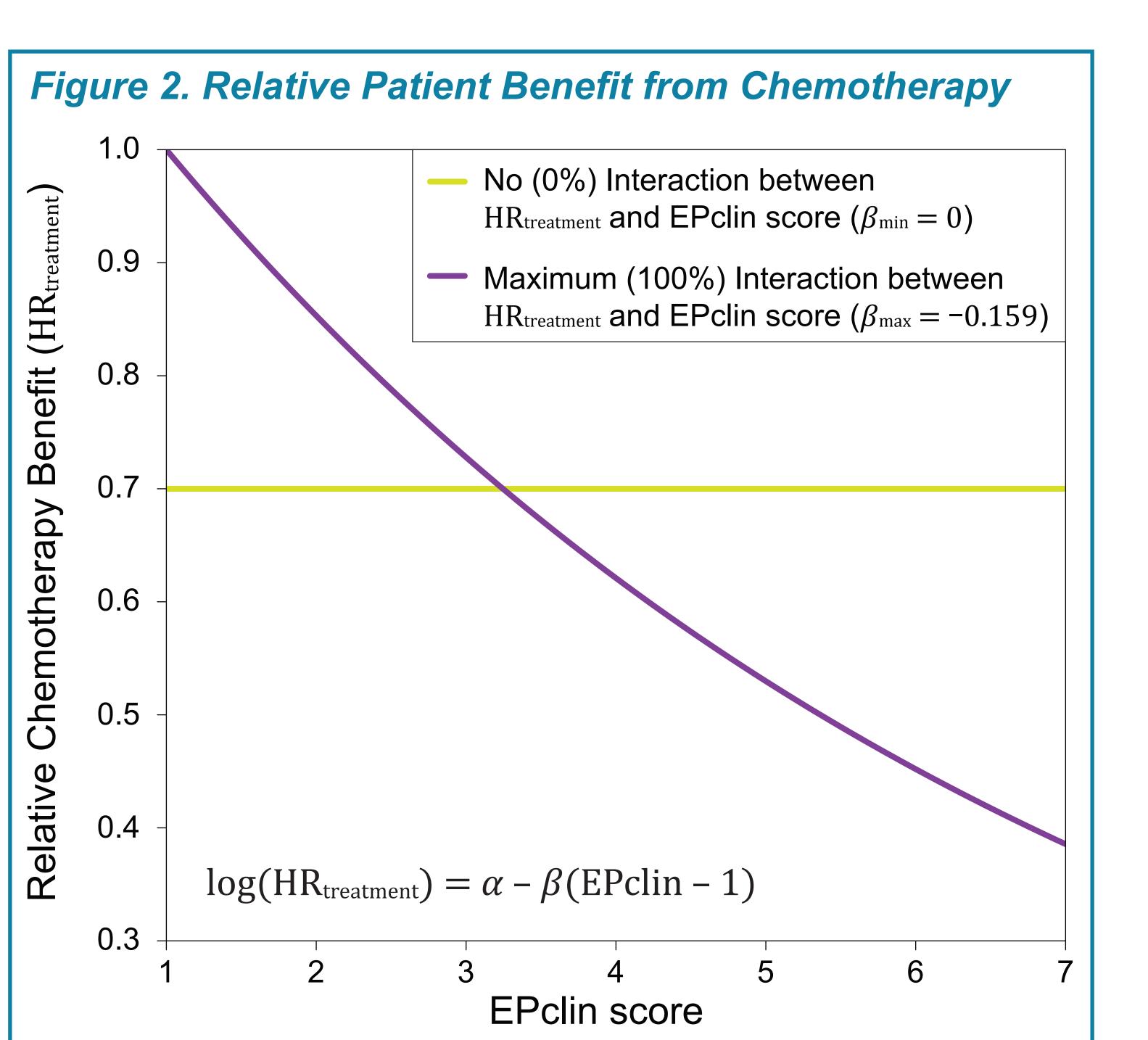
• The EPclin scores for patients in this cohort ranged from 1.3 to 6.2 (Figure 1).

- 58% (1,286/2,205) were in the low risk category.
- 42% (919/2,205) were in the high risk category.

Figure 1. Distribution of EPclin Scores High Risk Median: 3.2 SD: 0.79 Range: 1.3, 6.2 EPclin score

• HR_{treatment} as a function of EPclin score is given in Figure 2

- There is no assumed interaction between treatment benefit and EPclin (0% interaction strength, β_{min}).
- The interaction between treatment benefit and EPclin is maximized (100% interaction strength, β_{max}).



RESULTS

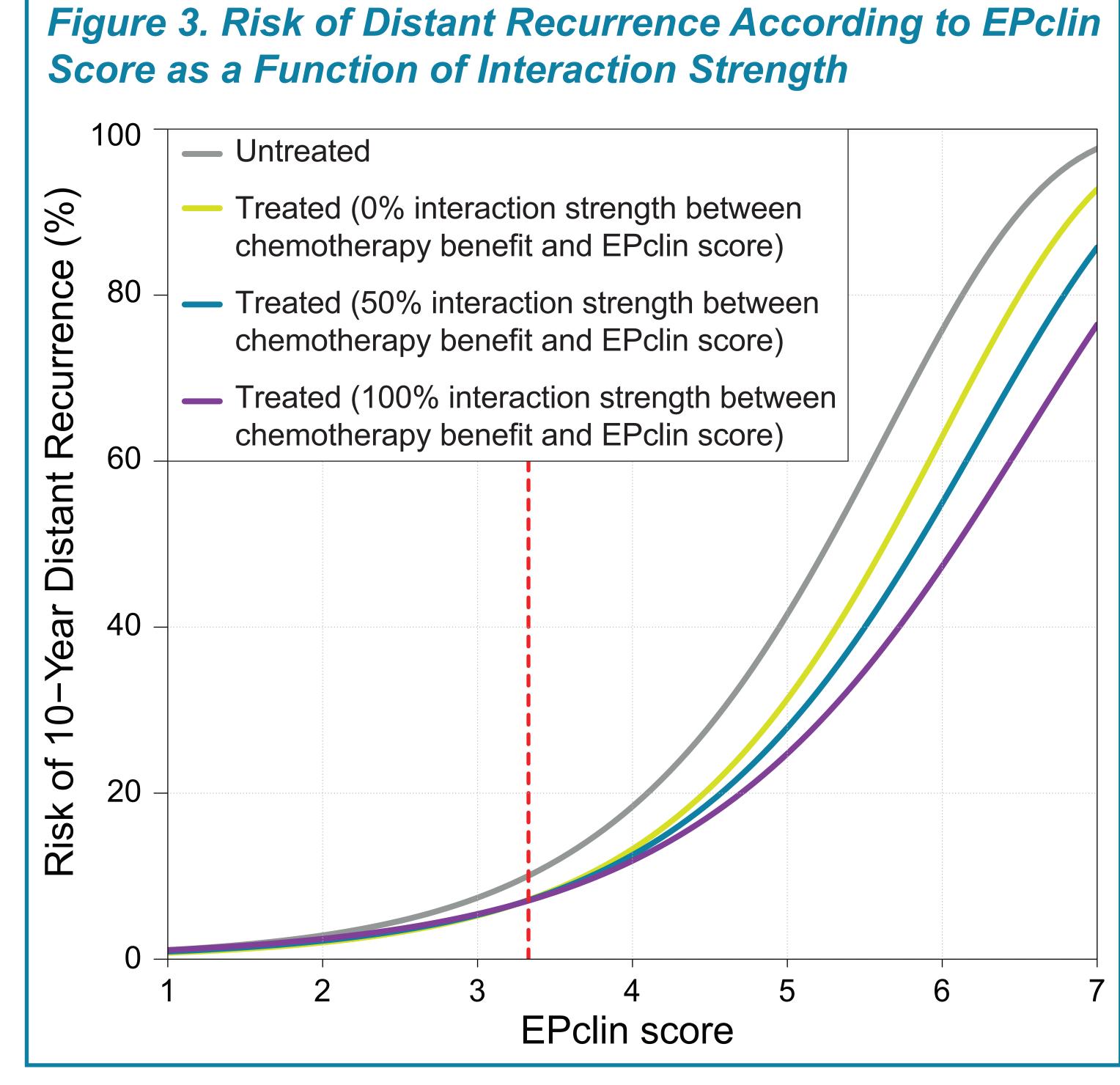


Figure 4. Absolute Chemotherapy Benefit

— Patients with high (≥3.3) EPclin score

Patients with low (<3.3) EPclin score

The interaction strength between HR_{treatment}

interaction (0%, $\beta_{\min} = 0$) to the maximum

possible interaction (100%, $\beta_{\text{max}} = -0.159$).

and EPclin score was varied from no

Interaction Strength (%)

- The risk of distant metastasis was calculated for a variety of interaction strengths between HR_{treatment} and EPclin score (Figure 3).
 - There was little separation between the treated and untreated risk curves for low EPclin scores.
 - There was much more separation between the treated and untreated risk curves for high EPclin scores.
- The effect of interaction strength on absolute chemotherapy benefit was rather moderate (Figure 4).
- The absolute chemotherapy benefit in high risk patients ranged from 7.3% (β_{max}) to 5.3% (β_{min}) (Figure 4).
 - This corresponded with mean 10-year risks of distant recurrence ranging from 14.7% (β_{max}) to 12.8% (β_{min}).
- The absolute chemotherapy benefit in low risk patients ranged from 1.5% (β_{max}) to 1.8% (β_{min}) (Figure 4).
 - This corresponded with mean 10-year risks of distant recurrence ranging from 4.6% (β_{max}) to 4.3% (β_{min}).

SUMMARY

- A mathematical model was used to estimate the expected absolute chemotherapy benefit among patients with earlystage ER+, HER2- breast cancer based on the test results of a 12-gene expression assay.
- This was done using the overall adjuvant chemotherapy benefit as determined in a meta-analysis by the EBCTCG in order to overcome the potential bias inherent to individual trials that result in variable observed treatment benefits.
- In this analysis, the 12-gene expression assay was estimated to predict absolute benefit from adjuvant chemotherapy in women with ER+, HER2- early stage breast cancer, regardless of which EPclin score cohorts accrued maximal relative treatment benefit.

REFERENCES

- . Gradishar W, et al. NCCN Clinical Practice Guidelines in Oncology: Breast Cancer. 2017; Version 1.2018.
- 2. Filipits M, et al. Clin Cancer Res. 2011;17:6012-6020.
- 3. Dubsky P, et al. *Br J Cancer*. 2013;109:2959-2964.
- 4. Buus R, et al. *J Natl Cancer Inst*. 2016;108:djw149.
- 5. Martin M, et al. *Breast Cancer Res.* 2014;16:R38. 6. Early Breast Cancer Trialists Collaborative Group. N Engl J Med. 1988; 319:1681-1692.
- 7. Hayes DF. *J Clin Oncol*. 2012;30:1264-1267.

Presented at ASCO on June 2, 2018