




Response to de Nonneville, Finetti, Mamessier, and Bertucci

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We appreciate the insightful correspondence from de Nonneville and colleagues (1) regarding our article “NDRG1 in Aggressive Breast Cancer Progression and Brain Metastasis.” Through their *in silico* analysis of a very large independent clinical cohort of patients with breast cancer (n = 8982), de Nonneville et al. (1) provided data that further validate our clinical findings that NDRG1 is a predictor of poor outcome in patients with aggressive breast cancer. Analysis of the independent cohort also demonstrated that NDRG1 is associated with features of aggressive breast cancer such as enrichment of tumor stemness, tumor invasion and metastasis signatures, further corroborating our preclinical observations. Importantly, de Nonneville et al. (1) expanded their analysis to include breast tumor subtypes considered less aggressive, such as estrogen receptor (ER)+/HER2– tumors; they found that NDRG1 also positively correlates with aggressiveness and poor survival outcome in patients with these tumors. We further stratified the MD Anderson cohort of breast cancer patients we reported in our work (see our article, Figure 6) to assess the prognostic significance of NDRG1 in an independent collection of ER+/HER2– tumors immunostained for NDRG1. We observed a tendency for a worse overall survival in NDRG1-high tumors compared with NDRG1-low tumors in this group (hazard ratio = 2.78, 95% confidence interval = 0.68 to 11.32; 2-sided log-rank test P = .08; Figure 1), consistent with the findings from de Nonneville et al. (1).

We would like to emphasize that our suggestion that “NDRG1 has a context-dependent function in breast cancer” was based on the current literature; we did not perform functional experiments using less aggressive ER+ cell lines. Liu et al. (2) demonstrated that overexpressing NDRG1 reduced cellular invasion, adhesion, and anoikis resistance in MCF-7, an ER+/HER2– cell line. Similarly, Godbole et al. (3) found that silencing NDRG1 increased migration in T47D cells, another ER+/HER2– breast cancer cell line.

In summary, we thank de Nonneville and colleagues for a very informative analysis that provides further confirmation of our findings that NDRG1 is a tumor promoter and predictor of poor outcome in aggressive breast cancer subtypes, and for expanding the analysis to less aggressive ER+/HER2– breast tumors. Clearly, these new findings support the need for designing experiments to establish the role of NDRG1 in

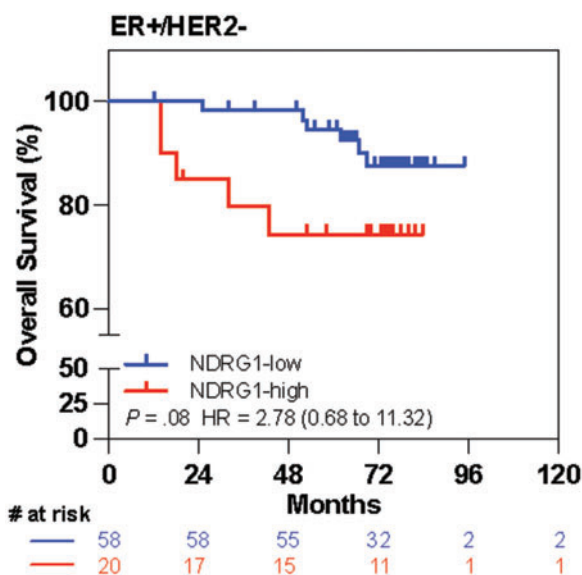


Figure 1. NDRG1 expression correlation with overall survival in ER+/HER2– patients. Kaplan-Meier estimate of overall survival according to NDRG1 expression in ER+/HER2– patients. P value from 2-sided log-rank test, and hazard ratio (HR) with 95% confidence interval (P = .08; HR = 2.78, 95% CI = 0.68 to 11.32) is shown. CI = confidence interval; ER = estrogen receptor.

progression, metastasis, and therapy response in preclinical models of ER+ breast cancer.

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Data Availability

The data that support the findings are available from the corresponding author (bgdebeb@mdanderson.org) upon reasonable request.

References

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