Errata


On page 729, in the first column, the last sentence of the second paragraph should read:

Because the authors have reported that only 3% of patients have more than 1 alteration, it will be interesting to discern whether all 3 molecular tests need to be used in the ProMisE model to achieve optimal performance or whether the features most associated with a poor prognosis (i.e., MMR and p53 abn) can be used.

The change has been denoted in bold.
The authors regret this error.
DOI: 10.1002/cncr.30889


The following funding support should be added to the existing information:

Grant P30 CA008748 - Cancer Center Support Grant; Behavioral Research; Bioinformatics; Biostatistics; CTRP Supplement; DEVELOPMENTAL FUNDS; Developmental Funds; Genomics; Microchemistry & Proteomics; PLANNING and EVALUATION; Planning & Evaluation; Web Survey Core.
The authors regret this omission.
DOI: 10.1002/cncr.30890


The following funding support should be added to the existing information:

Grant P30 CA008748 - Cancer Center Support Grant; Behavioral Research; Bioinformatics; Biostatistics; CTRP Supplement; DEVELOPMENTAL FUNDS; Developmental Funds; Genomics; Microchemistry & Proteomics; PLANNING and EVALUATION; Planning & Evaluation; Web Survey Core.
The authors regret this omission.
DOI: 10.1002/cncr.30891