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**ABOUT THE COVER**

The cover shows the results of an unsupervised two-dimensional hierarchical clustering analysis of the 5,000 most variable genes classifying breast tumors into ER\(^+\), HER2\(^+\), or triple-negative (ER\(^-\), PR\(^-\), HER2\(^-\)) subtypes. The analysis was performed on 800 breast tumors obtained from the Karolinska Institutet, Sweden. The RAS gene expression signature was high in the triple-negative subtype and low in the ER\(^+\) subtype. For details, see the article by Di Cosimo and colleagues on page 49 of this issue.