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Diffusion-Weighted MRI for Lymphoma Staging—Response
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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	sup+**, HER2	sup+**, or triple-negative (ER	sup−, PR	sup−, HER2	sup−) 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	sup+** subtype. For details, see the article by Di Cosimo and colleagues on page 49 of this issue.