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ABOUT THE COVER

Expression of the miR-34 family was found to be frequently reduced in human epithelial ovarian cancer, particularly so in tumors with p53 mutations. The figure shows miR-34a expression (dark blue) in ovarian serous adenocarcinoma as determined by in situ hybridization with locked nucleic acid–modified probes. Immunohistochemistry in serial sections revealed significant inverse correlation between miR-34a and its target MET, an oncogene commonly overexpressed in advanced stages of cancer. For details, see the article by Corney and colleagues on page 1119 of this issue.