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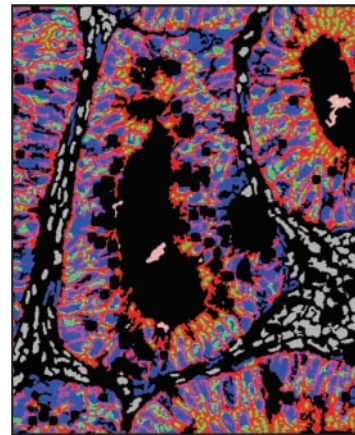
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About the Cover

A colon cancer tissue microarray containing 583 stage I-IV patients was stained for pan-cadherin, DAPI, and Met protein. Using novel automated curvature and intensity-based image analysis, epithelial membrane (*red*), cytoplasm (*green*) and nuclei (*blue*) were segmented. Stromal nuclei are shown in gray. The probability distribution of Met protein within each compartment was determined. A higher distribution of Met in the cytoplasm relative to the membrane was a significant predictor of survival in stage I and II patients. For further details, please see Ginty and colleagues on page 3814 in this issue.



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