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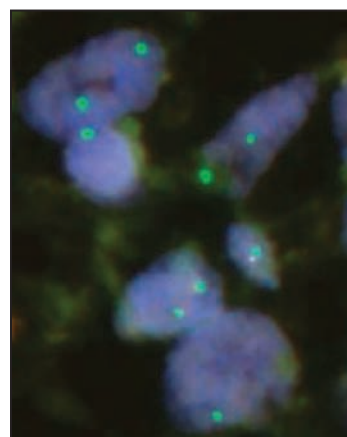
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Genetic Variants of *UGT1A6* Influence Risk of Colorectal Adenoma Recurrence.

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Array-based CGH identifies multiple genomic alterations in MSI-H and MSS colorectal tumors. A small, previously uncharacterized deletion at 16p13.2 was found in 35% of MSI-H and 21% of MSS tumors. Figure shows homozygous deletion of this region by dual-color FISH. Nuclei (blue) show two signals for 16 centromere (green) and the absence of signals for a BAC at 16p13.2 (labeled in red). FISH to an adjacent BAC control showed two red signals. For further details, please see Trautmann *et al.* on page 6379 of this issue.



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